

Inference Methods for Noisy Correlated Responses with Measurement Error

by

Qihuang Zhang

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Examining Committee Membership

The following served on the Examining Committee for this thesis. The decision of the Examining Committee is by majority vote.

External Examiner: Dr. Xin Gao
Professor, York University

Supervisor: Dr. Grace Yun Yi
Professor

Internal Member: Dr. Michael Wallace
Assistant Professor

Internal Member: Dr. Yeying Zhu
Associate Professor

Internal-External Member: Dr. Sharon Kirkpatrick
Associate Professor, School of Public Health and Health Systems

Author's Declaration

I hereby declare that I am the sole author of this thesis. This is a true copy of the thesis, including any required final revisions, as accepted by my examiners.

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Abstract

Studying complex relationships between correlated responses and the associated covariates has attracted much research interest. Numerous approaches have been developed to model correlated responses. However, most available methods rely on a crucial condition that response variables need to be precisely measured. In practice, this condition is often violated due to various reasons related to the data collection, study designs, or the nature of the variables. Without taking care of the feature of mismeasurement in variables, inference results are often biased.

Although measurement error and misclassification have been extensively studied in the literature, many problems of mismeasurement in correlated responses remain unexplored. The first problem of interest concerns measurement error and misclassification in the joint modeling of continuous and binary responses. In Chapter 2, we consider the setting with a bivariate outcome vector which contains a continuous component and a binary component both subject to mismeasurement. We propose an induced likelihood approach and describe an EM algorithm to handle measurement error in continuous response and misclassification in binary response simultaneously. The algorithm is fast and can be easily implemented. Simulation studies confirm that the proposed methods successfully remove the bias induced from the response mismeasurement. We implement the proposed methods to mice data arising from a genome-wide association study.

As a complement to the likelihood-based methods discussed in Chapter 2, in Chapter 3, we explore the bivariate generalized estimation equation method with mixed responses subject to measurement error and misclassification. The generalized estimating equation method enjoys robustness to certain model misspecification as well as consistency in the estimation of the mean structure parameters. However, the consistency property relies on the unbiasedness of estimating functions which can break down in the presence of the measurement error and misclassification in responses. We propose an insertion strategy to simultaneously account for measurement error effects in a continuous response and misclassification effects in a binary response. We consider scenarios where either an internal or an external validation subsample is available.

In Chapter 4, we consider a more complex situation where covariates are of a high dimension and may possess a network structure. We start with the case where data are precisely measured and propose a generalized network structure model together with the development of a two-step inferential procedure. In the first step, we employ a Gaussian graphical model to facilitate the network structure, and in the second step, we incorporate the estimated graphical structure of covariates and develop an estimating equation method.

Furthermore, we extend the development to accommodating mismeasured responses. We consider two cases where the information on mismeasurement is known or a validation sample is available. Theoretical results are established and numerical studies are conducted to justify the performance of the proposed methods.

In contrast to error-prone continuous and binary responses considered in the first three chapters, we investigate error-corrupted count data which particularly involve zero-inflated counts, a problem that has received little attention. Zero-inflated count data arise frequently from cancer genomics studies, and it is often of interest to incorporate the feature of excessive zeros in the analysis. However, measurement error in count responses is barely studied, let alone the zero-inflated Poisson model with measurement error. In Chapter 5, we propose a novel measurement error model which is unique for addressing error-contaminated count data. We show that ignoring the measurement error effects in analyzing the count response may generally lead to invalid inference results, and meanwhile, we identify situations where ignoring measurement error can still yield consistent estimators. Furthermore, we propose a Bayesian method to address the effects of measurement error under the zero-inflated Poisson model. We develop a data-augmentation algorithm that is easy to implement. Simulation studies are conducted to evaluate the performance of the proposed method. We apply our method to analyze a set of prostate adenocarcinoma genomics data.

Finally, in Chapter 6, we examine another type of correlated responses: time series data. We consider the autoregressive model and establish analytical results for quantifying the biases of the parameter estimation if the measurement error effects are neglected. We propose two measurement error models to describe different processes of data contamination. An estimating equation approach is proposed for the estimation of the model parameters with measurement error effects accounted for. We further discuss forecasting using error-prone times series data. This work is motivated by the need of understanding the ongoing evolving situation of the COVID-19 pandemic. It is important to assess how the mortality rate may change over time, but error-contaminate COVID-19 data present a considerable challenge in uncovering the true development path of the disease.

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Dedication

To my parents, Qi Zhang and Yumei Tang

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Chapter 1

Introduction

Modeling correlated responses are commonly conducted in genomics studies and epidemiology, which has attracted extensive research interests. In this thesis, we focus on handling three classes of correlated responses that have wide application scopes in practice. The first class includes the bivariate mixed responses, with a continuous component and a binary component. The second class contains the zero-inflated count data, which consists of two correlated components, with one component controlling the probability of taking value zero and another component modeling the distribution of a count variable. The third class concerns time series data which are autocorrelated.

Most work of modeling correlated responses relies on the critical condition that the variables are precisely measured, although this condition is usually made implicitly. However, measurement error is almost inevitable in practice. For example, in genetics, a genotype can be misclassified due to sequencing errors. In studies of infectious disease, the number of people infected with a certain disease, e.g., COVID-19, can be underestimated due to the asymptomatic infections.

Despite available discussions of measurement error in responses, there has been relatively little work on exploring the measurement error effects on the analysis of correlated responses (Neuhaus, 2002; Chen et al., 2011). In this thesis, we develop inference methods to address the effects due to measurement error and misclassification in different types of correlated responses, including likelihood-based methods, estimating equations methods, and Bayesian methods.

This thesis research is also motivated to tackle the challenges induced by noisy data arising in applications. In genetic association studies, sometimes the research interest lies in

studying the association of a genetic biomarker with mixed responses, which may be error-prone, and this motivates the topics discussed in Chapters 2–3. Meanwhile, understanding the pathway of genetic networks attracts a lot of interest, where numerous candidate genetic variants are associated with multiple traits in a complex manner. This presents a nice scenario of the application of joint models discussed in Chapter 4. Chapter 5 examines the zero-inflated Poisson model which is widely applied to handle cancer genomics and microbiome data to account for excessive zeros in count data. The ongoing pandemic of COVID-19 presents a perfect example of measurement error in time series data discussed in Chapter 6. Although our methods developed in this thesis are motivated by the unique features of individual data, the application scope of our methods is very broad.

To better understand our development in the following chapters, in this chapter, we review relevant topics. The remainder is organized as follows. In Section 1.1, we introduce three classes of correlated responses and the approaches that are often used to handle them. In Section 1.2, we explain the measurement error and misclassification mechanisms. In Section 1.3, we discuss the undirected graphical model under the exponential family setting. In Section 1.4, we explain several basic concepts of genome-wide association studies. Finally, we outline the thesis topics in Section 1.5.

1.1 Modeling Correlated Responses

In this thesis, we are interested in three classes of correlated responses: mixed continuous and discrete responses, count variables with the zero-inflating feature, and time series data. In the sequel, we separately review each class of correlated responses as well as some associated methods.

1.1.1 Mixed Responses and Joint Models

While modeling multiple responses of the same type has been extensively studied in longitudinal studies, modeling mixed type of outcomes, such as continuous and binary responses, has attracted increasing attention. Several models and inference methods were developed, such as generalized estimating equation methods (Liang and Zeger, 1986; Zeger and Liang, 1986), latent variable models (Sammel and Ryan, 1996), and multivariate linear mixed models (MLMM, Sammel et al., 1999). Jointly modeling multiple responses simultaneously has the advantage of boosting the estimation efficiency and the statistical power in testing genetic effects (McCulloch, 2008). There have been various joint models, including those for multiple discrete responses (Chen et al., 2016), latent variable models for

mixed continuous and discrete outcomes (Sammel et al., 1997; Teixeira-Pinto and Normand, 2009; Lin et al., 2014), correlated Probit models (Gueorguieva and Agresti, 2001), estimating function methods (Prentice and Zhao, 1991; Fitzmaurice and Laird, 1995), and the Bayesian framework (Wagner and Tüchler, 2010). Further generalizations were explored for handling clustered data (Catalano, 1997; Lin et al., 2010) and high dimensional data (Faes et al., 2008).

Although numerous methods were proposed to incorporate the correlation among responses, these methods can be roughly classified into two categories: likelihood approaches and estimating equation methods. These methods have their advantages and disadvantages. For example, the estimating function method is robust to the model specification by taking the price of the loss of efficiency. On the other hand, the estimators based on the likelihood methods are the most efficient but rely on the correct specification of the full distribution. In this section, we review the generalized linear mixed model with likelihood theory and the generalized estimation equations with estimating function theory.

Generalized Linear Mixed Model

For $i = 1, \dots, n$ and $j = 1, \dots, m$, let Y_{ij} be the j th response for the i th individual and let $X_i = (X_{i1}^T, \dots, X_{ip}^T)^T$ be the vector of covariates for the i th subject. Write $Y_i = (Y_{i1}, \dots, Y_{im})^T$. The *generalized linear mixed model* (GLMM) can be described in two steps. Assume that conditional on random effects u_i as well as covariates X_i , the Y_{ij} are independent and marginally follow a distribution from the exponential family given by

$$f(y_{ij}|x_i, u_i) = \exp \left\{ \frac{y_{ij}\varphi_{ij} - b(\varphi_{ij})}{a(\psi)} + c(y_{ij}; \psi) \right\}, \quad (1.1)$$

where $a(\cdot)$, $b(\cdot)$ and $c(\cdot)$ are known functions, φ_{ij} is the canonical parameter, and ψ is a dispersion parameter.

Based on the specification in (1.1), given covariates X_i and random effects u_i , the conditional mean $\mu_{uij} = E(Y_{ij}|u_i, X_i)$ equals $b'(\varphi_{ij})$ which is postulated by

$$g(\mu_{uij}) = \beta_0 + \beta_x^T X_{ij} + u_i^T F_{ij},$$

where $b'(\cdot)$ is the first derivative of $b(\cdot)$, $g(\cdot)$ is a link function, $\beta = (\beta_0, \beta_x^T)^T$ is the vector of regression parameters, and F_{ij} is a quantity determined by the study design and correlation among the responses. The random effects u_i are assumed to be independent of the covariates X_i .

Contrary to the name of *random effects*, the components of β are often called *fixed effects*. The β parameter in the generalized linear mixed model has a different interpretation from the generalized linear model with fixed effects only. In GLMM, the parameter β represents the changes of transformed responses associated with one unit change of covariates for an individual, whereas the β in the generalized linear model is interpreted as the changes at the population level.

Generalized Estimating Equation

For $i = 1, \dots, n$ and $j = 1, \dots, m_i$, let $\mu_{ij} = E(Y_{ij}|X_{ij})$ and $v_{ij} = \text{Var}(Y_{ij}|X_{ij})$ be the conditional mean and variance, respectively, given covariates X_{ij} .

The conditional mean μ_{ij} is modeled by

$$g(\mu_{ij}) = \beta_0 + \beta_x^T X_{ij},$$

where $g(\cdot)$ is a prespecified link function and $\beta = (\beta_0, \beta_x^T)^T$ is the vector of regression parameters.

The conditional variance v_{ij} is often modeled by a function of the mean and the dispersion parameters ψ . Namely,

$$v_{ij} = h(\mu_{ij}; \psi),$$

where ψ is the dispersion parameter and $h(\cdot)$ is a specified function characterizing the relationship between the conditional variance v_{ij} and the conditional mean μ_{ij} of Y_{ij} given X_{ij} . For instance, the variance function of the binary response is often specified as $h(\mu_{ij}; \psi) = \mu_{ij}(1 - \mu_{ij})$ where $\psi = 1$.

With the only assumptions on the first two moments, the *generalized estimating equation* (GEE) method is a natural way to estimate β . Let V_i be the conditional variance of Y_i given X_i . Define the estimating function

$$U_i(\beta) = D_i V_i^{-1} (Y_i - \mu_i),$$

where $\mu_i = (\mu_{i1}, \dots, \mu_{im})^T$, and $D_i = \frac{\partial \mu_i^T}{\partial \beta}$.

Then solving

$$\sum_{i=1}^n U_i(\beta) = 0,$$

for β gives a consistent estimator of β , say $\hat{\beta}$, provided regularity conditions (Liang and Zeger, 1986; Prentice and Zhao, 1991). In addition, $\sqrt{n}(\hat{\beta} - \beta)$ is asymptotically normally distributed with mean 0 and covariance matrix

$$\left\{ E \left(\frac{\partial U_i(\beta)}{\partial \beta^T} \right) \right\}^{-1} E\{U_i(\beta)U_i(\beta)^T\} \left\{ E \left(\frac{\partial U_i(\beta)}{\partial \beta^T} \right) \right\}^{-1T}.$$

We comment that the validity of the GEE method hinges on the assumption that

$$E(Y_{ij}|X_i) = E(Y_{ij}|X_{ij})$$

if the working matrix for V_i is not diagonal. A detailed discussion on this assumption can be found in Yi (2017, Section 5.1).

The consistency of the first order generalized estimating equation also requires the mean structure to be correctly specified regardless of whether the covariance structure is correctly specified or not. Sometimes, the association structure may be of scientific interest, and the second-order GEEs are constructed by modeling the second moment (Prentice and Zhao, 1991). Hall and Severini (1998) extended the original GEE model based on quasi-likelihood to improve the efficiency without requiring any covariance specification. Hall (2001) reviewed the relationships between different GEE approaches.

1.1.2 Zero-inflated Count Data and Zero-inflated Poisson Model

Count data arise from many studies of genomics (e.g. Fu et al., 2017) and microbiome (e.g. Xu et al., 2020), and they are commonly modeled by a Poisson distribution. On the other hand, count data may contain excessive zeros, which come from two sources, classified as “*structural zeros*” and “*sampling zeros*”. The “*structural zeros*” refers to that an individual is not “at risk” for the event and hence has no possibility to have a positive count. The “*sampling zeros*”, on the contrary, refers to that the individual is “at-risk” with a positive count, but results in a zero count by chance. For example, the count of the copy number variations (CNVs) is a useful indication of mutations in genes that might be associated with an increased risk of cancer. However, whether or not the CNVs are observed is also determined by whether the relevant pathways are activated. Many subjects have no CNVs simply due to the inactivated pathways, leading to extra “structural” zeros than expected when considering the Poisson distribution.

Viewing data as being generated from a mixture of a point mass at zero and a Poisson distribution, a zero-inflated Poisson model (Lambert, 1992) is commonly used to address

the excessive zero issue in the analysis of count data. It basically consists of two correlated components, where each component models a different aspect of zero-inflated count data. Specifically, one component concerns the probability of an individual sampled from an “at-risk” group and another component models the count variable conditional on the “at-risk” group.

To be specific, for $i = 1, \dots, n$, let Y_i denote the count outcome for subject i taking a non-negative integer value and let X_i denote the associated covariate vector of dimension p_x . For $i = 1, \dots, n$, let $\phi_i = P(A_i = 1|X_i)$ represent the conditional probability of sampling from ‘at-risk’ group, given X_i , and let $\mu_i = E(Y_i|A_i = 1, X_i)$ denote the condition mean of Y_i , given being sampled from the ‘at-risk’ group and the covariate X_i , which are assumed to satisfy $0 < \phi_i < 1$, and $\mu_i > 0$. That is, Y_i is sampled from the “non-at-risk” group with probability $1 - \phi_i$, and sampled from the “at-risk” group with probability ϕ_i , following a Poisson distribution with mean μ_i :

$$\begin{aligned} Y_i &= 0, \text{ with probability } 1 - \phi_i, \\ Y_i &\sim \text{Poisson}(\mu_i), \text{ with probability } \phi_i. \end{aligned}$$

1.1.3 Time Series Data and Autoregressive Model

Time series data arise commonly in epidemiology and infectious disease studies. Such data are taken as the third type of correlated responses in this thesis, where the correlation among the responses is directly reflected by the autocorrelation (or serial correlation). To model time series data, various models have been proposed, such as the classical decomposition model, autoregressive integrated moving average (ARIMA) model, autoregressive conditional heteroskedasticity (ARCH) model, state-space models, etc.

We denote a time series as $\{X_t : t = 1, \dots, T\}$, where X_t is a random variable and T is a positive integer or infinite. Stationarity is an important assumption for many models. The *strictly stationarity* for time series X_t is defined as

$$(X_1, \dots, X_n)^T \stackrel{d}{=} (X_{1+r}, \dots, X_{n+r})^T \quad (1.2)$$

for any positive integer n and r , where $\stackrel{d}{=}$ means those variables have the same joint distribution.

Sometimes, the assumption described in (1.2) is too strict and unrealistic in reality. We may consider a weaker condition for the stationarity assumption, where time series $\{X_t : t = 1, \dots, T\}$ is *weakly stationary* if these two conditions are satisfied:

- (i) $E(X_t)$ is independent of t ,
- (ii) $\text{Cov}(X_t, X_{t+r})$ is independent of t for each r .

In analysis of times series data, $E(X_t)$ and $\text{Cov}(X_t, X_{t+r})$ are important to be quantified. We define the autocovariance function

$$\gamma(h) = \text{Cov}(X_{t+h}, X_t) \text{ for } h = 0, \pm 1, \dots,$$

and the autocorrelation function (ACF) is then defined as

$$\rho(h) = \frac{\gamma(h)}{\gamma(0)}.$$

The autocovariance function and the autocorrelation function provide useful measures for the degree of dependence on the serial variables at different time lags, and thus, play important roles in the forecasting of future values.

Due to the importance of the autocovariance function, its properties under the weakly stationarity assumption are well studied:

- (1) $\gamma(0) \geq 0$;
- (2) $|\gamma(h)| \leq |\gamma(0)|$;
- (3) $\gamma(h) = \gamma(-h)$.

Autoregressive models are useful in analyzing time series data, which study the dependence of X_t on $\{X_{t-p}, \dots, X_{t-1}\}$, and is given by

$$X_t = \phi_0 + \sum_{j=1}^p \phi_j X_{t-j} + \epsilon_t,$$

where p is an integer smaller than T , $(\epsilon_1, \dots, \epsilon_t)^\top$ is independent of $(X_1, \dots, X_t)^\top$ with each ϵ_t having zero mean and a variance, say, σ_ϵ^2 , ϕ_0 is a constant drift, and $\phi = (\phi_1, \dots, \phi_p)^\top$ is the regression coefficient.

1.2 Measurement Error and Misclassification

Measurement error is prevalent in various cases. Sometimes it is because of technique errors. For example, when measuring a length, the last digit of the measurement is usually an estimate. Sometimes it is because of recall bias. In observational epidemiology, people answer questionnaires according to their experience in the past which is error-prone. A detailed discussion on reasons and sources of measurement error is provided by Yi (2017, Section 2.1).

In the literature of measurement error, we often distinguish different types of error-prone variables; the case of error-prone *continuous* variables is called *measurement error* and the case of error-prone *discrete* variables *misclassification*, although sometimes both cases are simply referred to as *measurement error* or *mismeasurement*.

In this section, we review some measurement error models and misclassification models.

1.2.1 Measurement Error

For $i = 1, \dots, n$, let Y_i denote the precisely measured continuous response. Due to measurement error, we do not observe Y_i , but instead, we observe a surrogate Y_i^* . The relationship between the true response Y_i and the observed surrogates Y_i^* can be described by different measurement error models in the same manner as Yi (2017) by introducing a random variable e_i :

1. Classical Additive Error Model:

$$Y_i^* = Y_i + e_i,$$

where the error term e_i is often assumed to be independent of the true response Y_i .

2. Multiplicative Model:

$$Y_i^* = Y_i e_i,$$

where the mean of e_i is assumed to be 1.

3. Linear Regression Model:

$$Y_i^* = \gamma_0 + \gamma_1 Y_i + \gamma_2^T X_i + e_i,$$

where e_i is independent of $\{Y_i, X_i\}$ and is often assumed to follow a normal distribution with mean zero and variance σ_e^2 , and γ_0 , γ_1 , and γ_2 are parameters.

4. General Regression Model:

$$Y_i^* = m(Y_i, X_i; \gamma) + e_i,$$

where $m(\cdot)$ is a prespecified function which can be nonlinear, γ is the vector of regression parameters associated with the measurement error model, and e_i is independent of $\{Y_i, X_i\}$.

Although these measurement error models provide a flexible specification of the relationship between the error-prone variable Y_i and its surrogate version Y_i^* , model identifiability issues may be a problem. To make the inferences meaningful, a specified model $f(y; \theta)$ must be *identifiable*. That is, if two parameters θ_1 and θ_2 make $f(y; \theta_1) = f(y; \theta_2)$ hold for any all possibly observed y (in a set of probability 1), then

$$\theta_1 = \theta_2.$$

Measurement error in covariates has received extensive research interest. On the other hand, less research work has been directed to measurement error in response, partly because the measurement error in response can be ignored in some scenarios, such as the response model described by a linear regression model together with a certain additive measurement error model. However, measurement error in response is not always ignorable if the measurement error process is nonlinear (Yi, 2017, Page 353).

1.2.2 Misclassification

When error-prone variables are discrete, we usually describe it as a misclassification problem. Let Y_i be a binary variable following a Bernoulli distribution. The true response Y_i is not observable but instead we observe the surrogate Y_i^* . Let $\pi_{i0} = P(Y_i^* = 1|Y_i = 0, X_i)$ and $\pi_{i1} = P(Y_i^* = 0|Y_i = 1, X_i)$ be the misclassification probabilities that may depend on the covariates X_i . The relationship between the true response Y_i and the observed surrogate Y_i^* is often modeled by logistic regressions models:

$$\text{logit } \pi_{i1} = \alpha_{01} + \alpha_{x1}^T X_i,$$

and

$$\text{logit } \pi_{i0} = \alpha_{00} + \alpha_{x0}^T X_i,$$

where $\alpha = (\alpha_{01}, \alpha_{x1}^T, \alpha_{00}, \alpha_{x0}^T)^T$ is the vector of the regression parameters.

Misclassification in responses and covariates has been studied in the literature (e.g., Neuhaus, 2002; Ramalho, 2002; Prescott and Garthwaite, 2002; Paulino et al., 2003; Chen et al., 2011; Yi et al., 2015; Shu and Yi, 2017). The misclassification in response will generally lead to biased estimation of parameters if no action is properly taken.

1.3 Undirected Graphical Model

An *undirected graphical model* (UGM), also called a *Markov random field* (MRF) or a *Markov network*, does not require specification for the edge orientations and is natural to be applied for image analysis and spatial statistics.

For $i = 1, \dots, n$, suppose $X_i = (X_{i1}, \dots, X_{ip})^T$ is a random vector for subject i . Let $V_i = \{1, 2, \dots, p\}$ be the index set of the vertices, corresponding to the variables $\{X_{i1}, \dots, X_{ip}\}$, and let $E_i = V_i \times V_i$ denote the set of edges derived from V_i . We use an undirected graph $G_i = (V_i, E_i)$ to describe the relationship among the covariates for subject i , where an edge of vertices s and t represents that X_s and X_t are correlated. Since the distribution of random vector X_i is assumed to be the same for each subject, we consider the graph for each individual to be identical. Namely, $G_1 = \dots = G_n \equiv G$ with $G = (V, E)$.

Markov independence is an important assumption for graphical models. To illustrate this assumption, we first define a cut set $\mathcal{C} \subseteq V$ to be a set of nodes that separate the graph G into two disjoint components \mathcal{A} and \mathcal{B} (Figure 1.1).

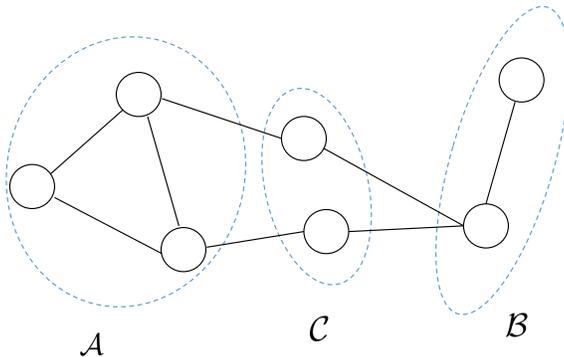


Figure 1.1: *Example of a graph separated by a cut set \mathcal{C}*

Assumption 1 (Markov independence assumption) *For all cut sets $\mathcal{C} \subset V$*

$$X_{\mathcal{A}} \perp\!\!\!\perp X_{\mathcal{B}} | X_{\mathcal{C}},$$

where $X_{\mathcal{A}}$, $X_{\mathcal{B}}$ and $X_{\mathcal{C}}$ are the covariates corresponding to the sets \mathcal{A} , \mathcal{B} and \mathcal{C} , respectively, and $\perp\!\!\!\perp$ represents “is conditionally independent of”.

Based on Assumption 1, a graphical model following the exponential family distribution can be constructed by

$$f(x_i; \theta, \Theta) = \exp \left\{ \sum_{k \in V} \theta_k B(x_{ik}) + \sum_{(s,t) \in E} \theta_{st} B(x_{is}) B(x_{it}) + \sum_{k \in V} C(x_{ik}) - A(\theta, \Theta) \right\}, \quad (1.3)$$

where $\theta = (\theta_1, \dots, \theta_p)^\top$ is the vector of parameters, $\Theta = [\theta_{st}]$ is a $p \times p$ symmetric matrix with diagonal elements to be one, $B(\cdot)$ and $C(\cdot)$ are prespecified functions, and the function $A(\theta, \Theta)$ is the normalizing constant to guarantee (1.3) to be a probability density function.

The density function (1.3) provides a general form, which includes many useful cases. For example, the Gaussian graphical model can be derived with the specification of $B(x_{it}) = \frac{x_{it}}{\sigma_t}$ and $C(x_{it}) = -\frac{x_{it}^2}{2\sigma_t^2}$, where σ_t is a dispersion parameter to scale the covariate, and its formulation is given by

$$f(x_i; \theta, \Theta) = \exp \left\{ \sum_{k \in V} \frac{1}{\sigma_k} \theta_k x_{ik} + \sum_{(s,t) \in E} \frac{1}{\sigma_s \sigma_t} \theta_{st} x_{is} x_{it} - \sum_{k \in V} \frac{1}{\sigma_k^2} x_{ik}^2 - A(\theta, \Theta) \right\}, \quad (1.4)$$

where σ_k is a scale parameter for X_{ik} , and $A(\theta, \Theta)$ is the normalizing constant. When the covariates follow a Bernoulli distribution, the Ising model can be derived from (1.3), given by

$$f(x_i; \theta, \Theta) = \exp \left\{ \sum_{(s,t) \in E} \theta_{st} x_{is} x_{it} - A(\theta, \Theta) \right\}, \quad (1.5)$$

where $B(x_{it}) = x_{it}$, $C(x_{it}) = 0$, and $A(\theta, \Theta)$ is a normalizing constant.

There are two methods for the parameter estimation of θ and Θ . The first method is to estimate the parameter based on the global likelihood. For example, for the Gaussian graphical model, the estimator can be estimated by maximizing the rescaled global log-likelihood $L(\Theta; X)$, which takes the form

$$L(\Theta; X) = \log \text{Det}(\Theta) - \text{Tr}(S\Theta) - \lambda \|\Theta\|_1,$$

where $S = \frac{1}{n} \sum_{i=1}^n x_i x_i^\top$ is the empirical covariance matrix, $\|\Theta\|_1 = \sum_{s \neq t} |\theta_{st}|$ is the ℓ_1 -norm of the off-diagonal entries of Θ , λ is the shrinkage parameter controlling the strength of the penalty, and $\text{Det}(\cdot)$ and $\text{Tr}(\cdot)$ are, respectively, the determinant and trace of a matrix.

In practice, the parameter can be estimated through the *graphical least absolute shrinkage and selection operator* (LASSO) algorithm (Friedman et al., 2008). Due to the complexity of the computation, the algorithm generally takes a long time. This method seems

to be mainly applied for Gaussian graphical models. An alternative method of estimating Θ is based on the idea of the neighborhood-likelihood.

For a given vertex $s \in V$, we use

$$X_{(-s)} = \{X_t : t \in V \setminus \{s\}\}$$

to denote the collection of all other random variables in the graph except X_s . Based on the Markov independence assumption, we define the neighborhood set for s

$$\mathcal{N}(s) = \{t \in V : (s, t) \in E\}$$

to be the set of relevant variables for variable X_s .

As shown in Figure 1.2, the set $\mathcal{N}(s)$ is a cut set that separates $\{s\}$ from the remaining variables.

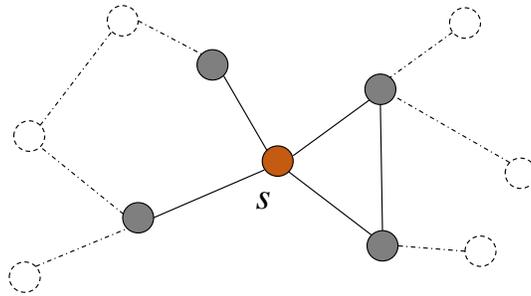


Figure 1.2: The vertices in gray compose the neighborhood set of vertex s , $\mathcal{N}(s)$

Write $\theta_{(-s)} = (\theta_{s1}, \dots, \theta_{s,(s-1)}, \theta_{s,(s+1)}, \dots, \theta_{sp})^T$ and let $\ell(\theta_{(-s)})$ be the log-likelihood for $\theta_{(-s)}$ scaled by $-1/n$,

$$\begin{aligned} \ell(\theta_{(-s)}) &= -\frac{1}{n} \log \left\{ \prod_{i=1}^n P(X_s | X_{(-s)}) \right\} \\ &= -\frac{1}{n} \sum_{i=1}^n \left\{ \theta_s B(X_{is}) + \sum_{t \in \mathcal{N}(s)} \theta_{st} B(X_{is}) B(X_{it}) + C(X_{is}) + D(\theta) \right\}, \end{aligned}$$

where $B(\cdot)$ and $C(\cdot)$ are functions specified the same as in (1.3) and $D(\theta)$ is the log-normalization constant.

Then for $s \in V$, we obtain an estimator for $\theta_{(-s)}$, denoted as $\hat{\theta}_{(-s)}$, by minimizing

$$\ell(\theta_{(-s)}) + \lambda \|\theta_{(-s)}\|_1,$$

where λ is the tuning parameter and $\|\cdot\|_1$ is the ℓ_1 -norm. Although this method is fast in implementation, it has a drawback that the resulting precision matrix Θ might be asymmetric. Namely, $\hat{\theta}_{st} \neq \hat{\theta}_{ts}$ for some $s, t \in V$, where $\hat{\theta}_{st}$ and $\hat{\theta}_{ts}$ represent the estimates for the (s, t) th and the (t, s) th entry of Θ . To overcome this problem, extra rules are applied to the estimates. For example, AND rule decides $\hat{\theta}_{st}$ and $\hat{\theta}_{ts}$ to be nonzero only if both of $\hat{\theta}_{st}$ and $\hat{\theta}_{ts}$ are nonzero; OR rule decides $\hat{\theta}_{st}$ and $\hat{\theta}_{ts}$ to be nonzero if either of $\hat{\theta}_{st}$ and $\hat{\theta}_{ts}$ is nonzero (Meinshausen and Bühlmann, 2006).

1.4 Genome-Wide Association Study

Genome-wide association studies (GWAS), also known as *whole-genome association studies* (WGAS), are observational studies searching for causal genetic variants that are associated with the responses of primary interest by scanning over a genome-wide set of genetic variants in different individuals. GWAS often focuses on the associations between *single-nucleotide polymorphisms* (SNPs) and clinical outcomes, such as diseases.

A genome-wide association study is often conducted in two stages (Wason and Dudbridge, 2012). In the first stage, candidate SNPs are selected through a simple model, such as linear regression, to gain computation speed. A set of candidate SNPs are selected according to the strength of the associations. In this stage, due to a large number of tests, several techniques can be applied to address the multiple testing issue, such as Bonferroni correction and the Benjamini-Hochberg Procedure (Benjamini and Hochberg, 1995), etc.

In the second stage, a more advanced approach is applied to study the association between the responses and the candidate SNPs screened from the first stage. There are two purposes for this stage. Firstly, the results getting from the first stage are rudimentary and need further validation. Secondly, advanced approaches can be used to study more complex research problems, such as identifying the possible genetic pathways and the pleiotropy effects to be introduced in Section 1.4.1. In this thesis, our research interest lies in the second stage of the genome-wide association study.

The genome-wide association studies involve several new data features. One important example is the *population stratification*. In the following subsections, we first introduce the basic concepts of statistical genetics and then review several methods concerning the population stratification.

1.4.1 Basic Concepts in Statistical Genetics

Pleiotropy is a common phenomenon in genetics that one gene is simultaneously associated with multiple traits. It has been widely studied in various types of research including Phenylketonuria (e.g., [Penrose, 1951](#)), Schizophrenia (e.g., [Navarrete et al., 2013](#)), and many others (e.g., [Kraja et al., 2014](#)).

As opposed to the pleiotropy effect, the term *polygenic* refers to the phenomenon that a group of genetic variants is associated with the same phenotype outcome. Examples of human polygenic traits are height, skin color, and eye color.

In genetics, the genes can not only have a complex association with multiple traits but also interact with each other in a complex manner through a collection of molecular regulators. To understand the mechanism of how the genes interact, these interactions are often modeled as a network, which is the so-called *gene regulatory network* (GNR). In the network, each gene will be expressed as a node and an interaction between two genes will be represented by an edge ([Yu et al., 2015](#)).

Although there are different types of graphs, the most commonly used graph in the gene regulatory network is the hub graph. In a hub graph, some of the nodes have a number of links that greatly exceed the average number and these nodes are called hub nodes. In GRN, the genes that are highly connected with other genes are called hub genes. Recently, because the hub genes play a key role in biological processes and are informative to uncover the mechanism of diseases, identifying hub genes has attracted a lot of the research interest ([Akavia et al., 2010](#)).

1.4.2 Population Stratification

The population features, such as the ethnicity for the human being data, can serve as confounders in genetic association studies, due to the non-random mating within populations, which is usually caused by the geological separation. To avoid the spurious association of the genetic variant and the response, several strategies have been developed.

The first method is to adjust the population stratification by the multi-trait mixed model (MTMM). The MTMM is an extension of the linear mixed model by incorporating the relatedness among the subjects into the covariance matrix of random effects. To be specific, the MTMM is defined as

$$Y_i = \beta^T X_i + u_i + \epsilon_i, \quad \text{for } i = 1, \dots, n,$$

where Y_i is a continuous trait, X_i includes the environmental covariates and genotype covariates, ϵ_i is the random error independent of $\{X_i, u_i\}$, β is the vector of regression coefficients, u_i is the random effect representing the confounding effect resulting from the subject dependent populations stratification. Write $u = (u_1, \dots, u_n)^T$. Often, u is assumed to follow $N(0, \sigma_g^2 R)$, where σ_g^2 is a scale parameter and R is an $n \times n$ positive definite matrix representing the pairwise relatedness among subjects. The relatedness matrix R can be determined by various methods according to the different nature of data. For example, when pedigree data are available, R is determined by the kinship matrix (Lange, 2003, Page 82). An alternative approach is to estimate relatedness matrices from genome-wide SNPs. The detail of specifying the relatedness matrix is to be discussed in Section 2.1.1.

The second method is to control the confounding by including the principal components of the relatedness matrix as fixed effects. To conduct the principal component analysis, the relatedness matrix is decomposed using the eigenvalue decomposition (EVD),

$$R = LDL^T,$$

where the columns of L are eigenvectors of R , and D is a diagonal matrix of positive eigenvalues of R . Let $F = RLD^{-\frac{1}{2}}$ be the matrix of principal components of the genetic information for the subjects, where each row of F , denoted as F_i , is the principal components for subject i . Here, $D^{-\frac{1}{2}}$ represents the diagonal matrix whose diagonal elements are the reciprocal and square root of those diagonal elements of D .

The principal components based model can be cast as

$$Y_i = \beta_1^T X_i + \beta_2^T F_i + \epsilon_i,$$

where $\beta = (\beta_1^T, \beta_2^T)^T$ is the vector of regression coefficients, F_i is the first few largest principal components of the relatedness matrix R .

Compared to the principal component based linear model, the MTMM model can be used to account for higher-rank confounding. On the other hand, controlling confounders by including fixed effects can circumvent the intensive computational burden.

1.5 Thesis Topics and the Outline

This thesis tackles several important problems and offers new additions to the literature. The thesis contains seven chapters with the last chapter concluding the thesis and the appendix including additional materials for Chapters 2–6. The topics and the development of Chapters 2–6 are outlined as follows.

1.5.1 Latent Variable Model with Bivariate Mixed Responses Subject to Measurement Error and Misclassification

In genetic association studies, due to the concern of population stratification, a multi-trait mixed model (MTMM) is often considered. In the MTMM, random effects are used not only to model the correlation for the multiple outcomes of the same subject but also to adjust for the relatedness among subjects. The MTMM has been widely applied in genome-wide association studies in various setting (Zhang et al., 2010; Korte et al., 2012; Zhou and Stephens, 2014; Furlotte and Eskin, 2015).

While mixed effects models have been widely used, they do not automatically ensure the inference results to be valid without conditions. A critical condition of using such models hinges on the precise measurements of the variables. Measurement error and misclassification, however, are typical features in genetic studies, but they are often ignored in most applications. Even if the practitioners aware the importance of the measurement, they might still ignore the effect of measurement error in genome-wide association studies due to intensive computational burdens, and accounting for the measurement error and misclassification does not only require great efforts of modeling but also introduce a large number of algorithm implementations. This being said, it is important to accommodate the mismeasurement to obtain valid results for genetic studies. Because of the advances in computer technology, implementing time-consuming algorithms does not seem to be an obstacle as before. A few studies, such as Hossain et al. (2009), Smith et al. (2013), and Rekaya et al. (2016), investigated some methods of analyzing genetic data with misclassification in a variable.

In the literature of response mismeasurements, there has been research exploring either *measurement error* in a continuous response (e.g., Buonaccorsi, 1991; Pepe et al., 1994; Buonaccorsi, 1996), or *misclassification* in a discrete response (e.g., Neuhaus, 2002; Ramalho, 2002; Prescott and Garthwaite, 2002; Paulino et al., 2003; Chen et al., 2011). However, no available work has been directed to deal with mixed responses with mismeasurement in continuous and discrete components, although there were a few studies simultaneously addressing mixed types of mismeasurement in covariates (Spiegelman et al., 2000; Yi et al., 2015; Zhang and Yi, 2019).

In Chapter 2, we consider the problem of joint modeling mixed responses with a continuous and a binary variable respectively subject to measurement error and misclassification. We employ the bivariate regression model with a latent variable which features the dependence of the response components as well as the population stratification. We propose two methods, the induced likelihood method and the EM algorithm approach, to account for both measurement error and misclassification of the responses in inferential procedures. A

general framework is considered for the specification of the mismeasurement processes. We show that both methods yield valid estimation results.

1.5.2 Estimating Equation Approach with Bivariate Mixed Responses Subject to Measurement Error and Misclassification

Correlated mixed types of data, containing both continuous and discrete variables arise commonly from clinical trials and genetic association studies. Many models have been proposed for analyzing such data. In addition to the well-studied likelihood approaches, marginal models have also been widely used to handle correlated mixed types of data due to the advantage of robustness to certain model misspecification since no full distributional assumptions are required. For example, generalized estimating equations, proposed by [Liang and Zeger \(1986\)](#), analyze the longitudinal data marginally and require only modeling of the first and second moments. This method has been further extended by many authors including [Prentice and Zhao \(1991\)](#); [Hall and Severini \(1998\)](#); [Pan \(2001\)](#); [Wang and Long \(2011\)](#); [Wang et al. \(2012\)](#).

Marginal methods are useful in joint modeling of mixed responses, such as a continuous response and a discrete response (e.g., [Liu et al., 2009](#)). However, such methods rely on a crucial condition that the variables must be precisely measured. It is well known that the mismeasurement in responses induces both biased parameter estimation and efficiency loss (e.g., [Neuhaus, 1999, 2002](#); [Chen et al., 2011](#)).

In Chapter 3, we use the bivariate generalized estimating equation to analyze mixed continuous and discrete responses subject to mismeasurement. We develop an insertion strategy to form unbiased estimating functions to accommodate the effects of measurement error and misclassification in responses. We consider different study designs including the main study/internal validation study and the main study/external study ([Spiegelman et al., 2000](#); [Yi et al., 2018](#)). We evaluate the proposed methods both theoretically and numerically.

1.5.3 Generalized Network Mixed Model in Discovering Gene Regulatory Network with Mixed Responses Subject to Measurement Error and Misclassification

In genetic analysis, genes can not only have a complex association with multiple traits but also interact with each other in a complex manner through molecular regulators. To understand how these genes may interact, *gene regulatory networks* (GRN) are often employed to describe the associations among genes, where genes are taken as nodes and an interaction between two genes is featured by an edge (Friedman, 2004). While many methods have been proposed for studying gene regulatory networks, a noticeable limitation is that the computational procedures are problem-specific, which hinders their application scope. To overcome this issue, several studies of applying graphical models to construct gene regulatory networks were motivated (e.g., Li et al., 2012; Yu et al., 2015).

Although graphical models have been developed to construct gene regulatory networks, most available work only focused on the modeling of covariates and did not consider how to model the relationship between network structured covariates and a response variable, let alone for the case with mixed bivariate responses with both continuous and discrete components. With the analysis of mixed responses, generalized estimating equation methods are useful because of its robustness of not requiring the specification of the joint distribution of the response variables as well as its flexibility of accommodating different covariance structures of the responses. The validity of such methods, however, is vulnerable to the mismeasurement of the response variables.

While it is well studied that mismeasurement in a discrete response typically breaks down the usual inference methods and ignoring this feature commonly yields erroneous inference results (e.g., Neuhaus, 1999; Chen et al., 2011), to our knowledge, there has not been research on dealing with error-contaminated mixed responses of both discrete and continuous components, let alone for their relationship with covariates of network structures.

In Chapter 4, we tackle this problem and make the following contributions: (1) we propose a new class of generalized network structured models to delineate the relationship between bivariate responses and covariates of a network structure; (2) we develop a two-stage inferential procedure to identify the network structure for covariates and to address the mismeasurement effects in responses of both continuous and discrete components; (3) we rigorously establish the asymptotic results for the proposed estimators and study the efficiency issues for different methods; (4) our methods offer tools for a broad variety of applications to handle error-prone data with complex association structures. For example,

they can be applied in genetic studies to simultaneously identify the gene regulatory network and study the association between the gene network and mixed type traits with the effects of mismeasurement accounted for.

To be specific, we develop a generalized network structured model which incorporates the graphical structure in the generalized linear models through a two-step procedure. In the first step, we identify the network structure in the covariates via the Gaussian graphical model. In the second step, we build generalized estimating equations to study the association between the bivariate responses and the network structured covariates selected from Step 1, where the effects due to the contamination in the responses are accommodated for valid inferential procedures. We start with a simple situation where the model parameters for the mismeasurement processes are known; this development highlights the idea of how effects of mismeasurement in the mixed responses can be accounted for in combination with the examination with the network structure for covariates. Furthermore, we extend the development to accommodating the cases where the parameters for the mismeasurement models are unknown and must be estimated from an additional validation subsample.

1.5.4 Zero-Inflated Poisson Models with Measurement Error in Response

Research on zero-inflated Poisson models has become active and has attracted various studies from several perspectives. [Rodrigues \(2003\)](#) and [Klein et al. \(2015\)](#) pursued a Bayesian inference analysis with zero-inflated models. [Todem et al. \(2016\)](#) developed a marginal model for the zero-inflated Poisson data. [Xiang et al. \(2007\)](#) and [Yang et al. \(2010\)](#) proposed a score test under the zero-inflated Poisson model.

Measurement error in count data has been scarcely explored, which basically has two challenges. Firstly, the count variable is an integer, and thus the traditional measurement error models, such as the classical additive model is not applicable in this situation. Secondly, the variables are bounded below but unbounded above, because the observed values are always positive.

In Chapter 5, we propose a measurement error model that is unique for error-corrupted count data by incorporating two possible sources of measurement error. We explore the validity of statistical inference when measurement error in count data is ignored. We develop a Bayesian framework to account for the measurement error effects, which avoids the unidentifiability issue through the inclusion of weakly informative priors.

1.5.5 Autoregressive Models with Data Subject to Measurement Error

Time series data are common in the fields of epidemiology, economics, and engineering, and various models and methods have been developed for analyzing such data. The validity of these methods, however, hinges on the condition that time series data are precisely collected. This condition is restrictive in applications. Measurement error is often inevitable. In the study of air pollution, for example, it is difficult or even impossible to precisely obtain the true measurement of the air population.

Some work on time series subject to measurement error is available in the literature. [Tanaka \(2002\)](#) proposed a Lagrange multiplier test to assess the presence of measurement error in time series data. [Staudenmayer and Buonaccorsi \(2005\)](#) explored the classical measurement error model for the autoregressive model. [Tripodis and Buonaccorsi \(2009\)](#) studied measurement error in forecasting using the Kalman filter. [Dedecker et al. \(2014\)](#) considered non-linear AR(1) model with measurement error. Despite available discussions of measurement error in time series, several limitations restrict the application scope of the existing work. Most available methods consider only the autoregressive models without the drift and assume the simplest additive measurement error model. Furthermore, most work involves a complex formulation to adjusted for the measurement error effects, which is not straightforward to implement for practitioners. In addition, to our knowledge, there is no available work addresses measurement error effects on prediction under the autoregressive model.

In Chapter 6, we systematically explore the analysis of error-prone time series data under the autoregressive model. We propose two types of models to delineate measurement error processes: the additive regression models and multiplicative models. These modeling schemes offer us great flexibility in facilitating different applications. We investigate the impact of the naive analysis which ignores the feature of measurement error in the inferential procedures, and we obtain analytical results for characterizing the biases due to the naive analysis. We develop an estimating equation approach to adjust for the measurement error effects on time series analysis. We establish asymptotic results for the proposed estimators and develop the theoretical results for the forecasting of times series in the presence of measurement error. Finally, we describe a block bootstrap algorithm for computing standard errors of the proposed estimators.

Our work is partially motivated by the data of COVID-19, a wide-spread disease that has become a global health challenge and has caused over ten million infections and half million deaths as of August, 2020. Because of the special features of the disease, the data of COVID-19 introduce many new challenges: 1) due to the asymptomatic infected

cases and the patients with light symptoms who do not go to hospitals, the reported cases with COVID-19 are typically smaller than the true number of infected cases; 2) due to the limited test resources, many infected cases are not able to be identified instantly; and 3) the varying incubation periods lead to the delay of the identification of the infections. Consequently, the discrepancy between the reported case number and the true case number can be substantial, and ignoring these features and applying the traditional time series analysis method would no longer produce valid results.

In Chapter 6, we apply the developed methods to analyze the COVID-19 data. We are interested in studying how the mortality rate in a region may change over time and describing the trajectory of the death rate. While the mortality rate of a disease is defined as the death number divided by the case number, the determination of the mortality rate of COVID-19 is challenging. In contrast to the standard definition, [Baud et al. \(2020\)](#) estimated mortality rates by dividing the number of deaths on a given day by the number of patients with confirmed COVID-19 infection 14 days before, with the consideration of the maximum incubation time to be 14 days. Due to the unique features of COVID-19, there does not seem to be a precise way to define the mortality rate of COVID-19. In this chapter, we conduct a sensitivity analysis to assess the severity of the pandemic by using different definitions of the mortality rate and considering different ways of modeling measurement error in the data.

Using the data collected from the dashboard developed by Johns Hopkins University (JHU-CSSE, [Dong et al., 2020](#)), we analyze the mortality rates of COVID-19 and conduct forecasting of the COVID-19 related mortality rate for the four most populated provinces in Canada, British Columbia, Ontario, Quebec, and Alberta.

Chapter 2

Latent Variable Models with Bivariate Mixed Responses Subject to Measurement Error and Misclassification

In this chapter, we focus on the effects of measurement error and misclassification on analysis of the mixed responses postulated with latent variable models. The notation and the setup for the response model, the measurement error model as well as the misclassification models are introduced in Section 2.1. We describe the induced likelihood method in Section 2.2 and the EM algorithm method in Section 2.3. We extend the method to facilitating pedigree data with correlated subjects in Section 2.4. Simulations studies are conducted to evaluate the performance of the two methods in Section 2.5. To illustrate the usage of the methods, in Section 2.6 we conduct numerical analysis using the mice data arising from a genome-wide association study.

2.1 Model Setup

2.1.1 Response Model

Suppose n subjects are recruited independently in the study. For subject $i = 1, \dots, n$, two possibly correlated responses Y_{ij} are measured for $j = 1, 2$, where Y_{i1} is a continuous

variable, and Y_{i2} is a binary variable. Write $Y_i = (Y_{i1}, Y_{i2})^T$. Let $X_i = (X_{i1}, \dots, X_{ip_x})^T$ denote the covariate vector for subject i , where p_x is the dimension of the covariates. To facilitate the association structure between the mix-type responses Y_{i1} and Y_{i2} , we introduce a latent variable u_i . Conditional on random effects u_i and covariates X_i , we assume that Y_{i1} and Y_{i2} are independent, each having a probability density or mass function from the exponential family

$$f(y_{ij}|u_i, x_i) = \exp [\{y_{ij}\eta_j - b_j(\eta_j)\}/d_j(\phi_j) + c_j(y_{ij}, \phi_j)],$$

for $i = 1, \dots, n$ and $j = 1, 2$ where $b_j(\cdot)$, $c_j(\cdot)$ and $d_j(\cdot)$ are known functions, η_j is a canonical parameter, and ϕ is a dispersion parameter.

Let $\mu_{ij} = E(Y_{ij}|u_i, X_i)$ be the conditional mean of response Y_{ij} for $j = 1, 2$, and then $\mu_{ij} = b'_j(\eta_j)$. To explicitly describe the dependence of μ_{ij} on random effects and the covariates, we consider a bivariate generalized linear mixed model

$$\begin{bmatrix} g_1(\mu_{i1}) \\ g_2(\mu_{i2}) \end{bmatrix} = \begin{bmatrix} \beta_1^T X_i \\ \beta_2^T X_i \end{bmatrix} + \begin{bmatrix} u_i \\ u_i \end{bmatrix}, \quad (2.1)$$

where $g_1(\cdot)$ and $g_2(\cdot)$ are the link functions, determined by $g_1^{-1}(\cdot) = b'_1(\cdot)$ and $g_2^{-1}(\cdot) = b'_2(\cdot)$; $\beta = (\beta_1^T, \beta_2^T)^T$ is the vector of regression coefficients that is of primary interest; and u_i is a random effect. For the continuous response Y_{i1} with a normal distribution, $b'_1(t)$ is taken as t ; and for the binary response Y_{i2} , $b'_2(t) = \exp(t)/(1 + \exp(t))$, yielding that $g_1(t) = t$ and $g_2(t) = \log \frac{t}{1-t}$, respectively, where t represents the argument of functions.

Write $u = (u_1, \dots, u_n)^T$. Often, u is assumed to follow a normal distribution $N(0, \sigma_g^2 R)$, where σ_g^2 is an unknown scale parameter, and $R = [R_{jk}]_{n \times n}$ is a specified positive definite matrix with the (j, k) element R_{jk} determined by the study design, where $j, k = 1, \dots, n$. In applications, a particular specification of R may be imposed to feature a problem-specific association structure. For instance, to reflect the independence among different subjects, [Sammel et al. \(1997\)](#) set R as a diagonal matrix with the diagonal elements being given (such as 1 or other values). In the development in Sections 2-4, we consider the case where R is a given diagonal matrix, and in Section 5, we extend the diagonal R to a blockwise diagonal matrix to reflect the correlation among the subjects.

Model (2.1) is useful for characterizing the dependence of mix-type responses on covariates ([Sammel et al., 1997](#)). This model can be conveniently used to analyze genetic data with mix-type responses, where the genotype information may be summarized as the covariates $X_i = (X_{i1}, \dots, X_{ip_x})^T$, where for $k = 1, \dots, p_x$, covariate X_{ik} can be continuous (e.g., representing a continuous measurement of an environmental effect), or binary

(e.g., representing a clinical treatment); X_{ik} can be ordinal referring to, for example, the genotype.

For instance, we observe the genotypes through genetic markers (called single nucleotide polymorphisms, SNPs) in each locus (the location of a gene on the genome). For $k = 1, \dots, p_x$, let $X_{ik}^{(1)}$ and $X_{ik}^{(2)}$ stand for the nucleotides of SNP k for subject i inherited from the father and mother respectively. Each SNP consists of two nucleotides, each being one of the two types of alleles, “ A_1 ” and “ A_2 ”. Hence, all possible forms of a SNP are “ A_1A_1 ”, “ A_1A_2 ” and “ A_2A_2 ”, derived from different combinations of the alleles. Then the covariate X_{ik} , representing k th SNP for subject i , is coded according to the nucleotide level of A_2 , given by

$$X_{ik} = I(X_{ik}^{(1)} = A_2) + I(X_{ik}^{(2)} = A_2), \quad (2.2)$$

yielding an ordinal variable X_{ik} taking values of 0, 1 and 2, where $I(\cdot)$ is an indicator function.

2.1.2 Measurement Error and Misclassification Models

For $i = 1, \dots, n$, suppose that the response variables Y_{i1} and Y_{i2} are subject to mismeasurement and that their precise measurements may not be observed for every subject. Let Y_{i1}^* and Y_{i2}^* denote the observed measurements of Y_{i1} and Y_{i2} , respectively; they are also called surrogate measurements of Y_{i1} and Y_{i2} . Let $Z_i = (Z_{i1}, \dots, Z_{ip_z})^T$ denote the covariate vector involved in the measurement error and misclassification process for subject i where p_z is the dimension of Z_i . For ease of exposition, we assume that Z_i is a subset of X_i ; if this is not the case, we can modify our initial definition of X_i to include Z_i as its part.

To describe the mismeasurement processes, we consider the factorization

$$f(y_{i1}^*, y_{i2}^* | y_{i1}, y_{i2}, u_i, x_i) = f(y_{i1}^* | y_{i2}^*, y_{i1}, y_{i2}, u_i, x_i) f(y_{i2}^* | y_{i1}, y_{i2}, u_i, x_i). \quad (2.3)$$

We assume that

$$f(y_{i1}^* | y_{i2}^*, y_{i1}, y_{i2}, x_i, u_i) = f(y_{i1}^* | y_{i1}, y_{i2}, x_i) = f(y_{i1}^* | y_{i1}, y_{i2}, z_i), \quad (2.4)$$

and
$$f(y_{i2}^* | y_{i1}, y_{i2}, x_i, u_i) = f(y_{i2}^* | y_{i1}, y_{i2}, x_i) = f(y_{i2}^* | y_{i2}, x_i) = f(y_{i2}^* | y_{i2}, z_i). \quad (2.5)$$

Assumptions (2.4) and (2.5) basically say that conditional on the true responses and the covariates, surrogate measurements and random effects are independent. The assumptions also suggest that Z_i completely reflects the dependence on the covariates when featuring the measurement error and misclassification processes. While the last two equalities in

(2.5) are not needed to assume, having them offers us a convenient way to model the misclassification probabilities; see Yi et al. (2015).

Let $\pi_{i0} = P(Y_{i2}^* = 1|Y_{i2} = 0, Z_i)$ and $\pi_{i1} = P(Y_{i2}^* = 0|Y_{i2} = 1, Z_i)$ be the misclassification probabilities that may depend on the covariates Z_i . We consider logistic models for the misclassification process,

$$\begin{aligned} \text{logit } \pi_{i1} &= \alpha_{01} + \alpha_{z1}^T Z_i, \\ \text{and} \quad \text{logit } \pi_{i0} &= \alpha_{00} + \alpha_{z0}^T Z_i, \end{aligned} \tag{2.6}$$

where $\alpha = (\alpha_{01}, \alpha_{z1}^T, \alpha_{00}, \alpha_{z0}^T)^T$ is the vector of the regression parameters.

For the continuous response Y_{i1} , we consider a regression model which facilitates possible dependence of Y_{i1}^* on $\{Y_{i1}, Y_{i2}, Z_i\}$,

$$Y_{i1}^* = m(Y_{i1}, Y_{i2}, Z_i; \gamma) + e_i, \tag{2.7}$$

where e_i is the random error independent of $\{Y_{i1}, Y_{i2}, X_i, u_i\}$ and has zero mean and constant variance σ_e^2 , γ is the vector of regression coefficients, and $m(\cdot)$ is the mean function that can be linear or nonlinear.

Often, an additive model is considered for (2.7), given by

$$Y_{i1}^* = \gamma_0 + \gamma_1 Y_{i1} + \gamma_2 f(Y_{i2}) + \gamma_3 Z_{i3} + e_i, \tag{2.8}$$

where $f(\cdot)$ is a function of the binary response Y_{i2} , $\gamma = (\gamma_0, \gamma_1, \gamma_2, \gamma_3)^T$ is the vector of parameters, and a normal distribution is assumed for e_i . We comment that model (2.8) offers the flexibility and convenience of featuring the dependence of surrogate variable Y_{i1}^* on the true responses and covariates, but model identifiability may be a concern if no care is taken. In Appendix A.1, we outline the discussion on this aspect.

2.2 Estimation Procedures

2.2.1 Induced Likelihood for the Observed Data

To see how the distribution of the observed Y_i^* is different from that of Y_i , we derive the conditional distribution of Y_i^* given $\{u_i, X_i\}$. Indeed,

$$\begin{aligned}
f(y_i^*|u_i, x_i) &= \int_{y_{i1}} \sum_{y_{i2}} f(y_{i1}^*, y_{i2}^*, y_{i1}, y_{i2}|u_i, x_i) dy_{i1} \\
&= \int_{y_{i1}} f(y_{i1}^*|y_{i1}, y_{i2} = 1, z_i) f(y_{i2}^*|y_{i2} = 1, z_i) f(y_{i1}|u_i, x_i) f(y_{i2} = 1|u_i, x_i) dy_{i1} \\
&\quad + \int_{y_{i1}} f(y_{i1}^*|y_{i1}, y_{i2} = 0, z_i) f(y_{i2}^*|y_{i2} = 0, z_i) f(y_{i1}|u_i, x_i) f(y_{i2} = 0|u_i, x_i) dy_{i1},
\end{aligned} \tag{2.9}$$

where in the second equality, we use (2.4), (2.5), and the conditional independence of Y_{i1} and Y_{i2} given $\{u_i, X_i, Z_i\}$. Using the model formulations in Section 2.1, we have the following expressions for the terms of (2.9):

$$\begin{aligned}
f(y_{i1}^*|y_{i1}, y_{i2}, z_i) &= \frac{1}{\sqrt{2\pi\sigma_e^2}} \exp \left[-\frac{\{y_{i1}^* - m(y_{i1}, y_{i2}, z_i; \gamma)\}^2}{2\sigma_e^2} \right]; \\
f(y_{i2}^*|y_{i2} = q, z_i) &= \left\{ \frac{\exp(\alpha_{0q} + \alpha_{zq}^T z_i)}{1 + \exp(\alpha_{0q} + \alpha_{zq}^T z_i)} \right\}^{q(1-y_{i2}^*) + (1-q)y_{i2}^*} \\
&\quad \times \left\{ \frac{1}{1 + \exp(\alpha_{0q} + \alpha_{zq}^T z_i)} \right\}^{qy_{i2}^* + (1-q)(1-y_{i2}^*)} \quad \text{with } q = 0, 1; \\
f(y_{i1}|x_i, u_i) &= \frac{1}{\sqrt{2\pi\sigma^2}} \exp \left\{ -\frac{(y_{i1} - \beta_1^T x_i - u_i)^2}{2\sigma^2} \right\}; \\
f(y_{i2}|x_i, u_i) &= \left\{ \frac{\exp(\beta_2^T x_i + u_i)}{1 + \exp(\beta_2^T x_i + u_i)} \right\}^{y_{i2}} \left\{ \frac{1}{1 + \exp(\beta_2^T x_i + u_i)} \right\}^{1-y_{i2}}.
\end{aligned}$$

Consequently, the conditional distribution of Y_i^* , given X_i , is given by

$$\begin{aligned}
f(y_i^*|x_i) &= \int f(y_i^*, u_i|x_i) du_i \\
&= \int f(y_i^*|u_i, x_i) f(u_i|x_i) du_i,
\end{aligned} \tag{2.10}$$

where $f(u_i|x_i) = \frac{1}{(2\pi\sigma_g^2 R_{ii})^{\frac{1}{2}}} \exp\left(-\frac{u_i^2}{2\sigma_g^2 R_{ii}}\right)$, with R_{ii} being the i th diagonal elements of matrix R , and $f(y_i^*|u_i, x_i)$ is determined by (2.9).

Let $\theta = (\beta^\top, \gamma^\top, \alpha^\top, \sigma^2, \sigma_e^2, \sigma_g^2)^\top$. Inference about θ can be carried out using the likelihood for the observed data, given by

$$L(\theta) = \prod_{i=1}^n f(y_i^* | x_i), \quad (2.11)$$

where $f(y_i^* | x_i)$ is determined by (2.10) with the dependence on parameter θ suppressed in the notation.

Maximizing $L(\theta)$ with respect to θ gives the maximum likelihood estimator, say $\hat{\theta}$, of θ . Under regularity conditions, this is equivalent to solving

$$\sum_{i=1}^n S_i(\theta) = 0, \quad (2.12)$$

where $S_i(\theta) = \frac{\partial}{\partial \theta} \log f(y_i^* | x_i)$. Typically, (2.12) does not have an analytic solution; solving (2.12) usually requires a numerical method, such as the Newton-Raphson method.

Under regularity conditions, $\hat{\theta}$ is a consistent estimator of θ , and

$$\sqrt{n}(\hat{\theta} - \theta) \xrightarrow{d} N(0, I^{-1}(\theta)) \quad \text{as } n \rightarrow \infty,$$

where

$$I(\theta) = E \{ S_i(\theta) S_i^\top(\theta) \}. \quad (2.13)$$

2.2.2 Implementation

The estimation of the parameter θ is conducted by maximizing (2.11). We realize this by using the Newton-Raphson algorithm in combination with the Monte Carlo method. Let N_u and N_y be prespecified large integers.

Step 1. Take an initial parameter value $\theta^{(0)} = (\beta^{(0)\top}, \gamma^{(0)\top}, \alpha^{(0)\top}, \sigma^{2(0)}, \sigma_e^{2(0)}, \sigma_g^{2(0)})^\top$ and set the iteration index $t = 0$.

Step 2. At iteration $(t + 1)$, for $i = 1, \dots, n$, independently generate a sequence of values, $\{u_i^{[1]}, \dots, u_i^{[N_u]}\}$, from $N(0, \sigma_g^{2(t)} R_{ii})$, where $\sigma_g^{2(t)}$ is the parameter σ_g^2 evaluated at the t th iteration and R_{ii} is the (i, i) entry of matrix R . For $i = 1, \dots, n$ and $a = 1, \dots, N_u$, generate $\{y_{i1}^{[a,1]}, \dots, y_{i1}^{[a, N_y]}\}$ from $N(\beta^{(t)\top} X_i + u_i^{[a]}, \sigma^{2(t)})$, where $\sigma^{2(t)}$ and $\beta^{(t)}$ are, respectively, the parameter σ^2 and β evaluated at the t th iteration.

Step 3. The likelihood function in (2.11) is approximated by

$$\tilde{L}(\theta) = \prod_{i=1}^n \left\{ \frac{1}{N_u} \sum_{a=1}^{N_u} f(y_i^* | u_i^{[a]}, x_i) \right\}, \quad (2.14)$$

where

$$\begin{aligned} f(y_i^* | u_i^{[a]}, x_i) &= \frac{1}{N_y} \sum_{b=1}^{N_y} f(y_{i1}^* | y_{i1}^{[a,b]}, y_{i2} = 1, z_i) f(y_{i2}^* | y_{i2} = 1, z_i) f(y_{i2} = 1 | u_i^{[a]}, x_i) \\ &+ \frac{1}{N_y} \sum_{b=1}^{N_y} f(y_{i1}^* | y_{i1}^{[a,b]}, y_{i2} = 0, z_i) f(y_{i2}^* | y_{i2} = 0, z_i) f(y_{i2} = 0 | u_i^{[a]}, x_i). \end{aligned}$$

Step 4. Compute $\tilde{S}(\theta^{(t)}) = \frac{\partial}{\partial \theta} \log \tilde{L}(\theta)$ and $\tilde{I}(\theta^{(t)}) = -\frac{\partial^2}{\partial \theta \partial \theta^T} \log \tilde{L}(\theta)$, and update $\theta^{(t+1)}$ by

$$\theta^{(t+1)} = \tilde{I}(\theta^{(t)})^{-1} \tilde{S}(\theta^{(t)}) + \theta^{(t)}.$$

Step 5. Check if the $\theta^{(t+1)}$ converges by evaluating $|(\theta^{(t+1)} - \theta^{(t)}) / (\theta^{(t)} + c_1)| < c_2$, where c_1 and c_2 are prespecified small tolerance values. Otherwise, $t := t + 1$, and go back to Step 2.

The Monte Carlo approximation from Step 2 to Step 4 has the computation complexity of order $O(nN_uN_y)$. For an accurate approximation, the computation typically requires a large number of replicates such as 10000 for N_u and N_y . Alternatively, one may employ the Gaussian quadrature method (James, 1980) to approximate the integral in (2.11), as discussed in Appendix A.2.

2.3 EM Algorithm

In this section, we consider an alternative to estimating model parameters using the EM algorithm. The log-likelihood for the complete data $\{(Y_{i1}, Y_{i2}, Y_{i1}^*, Y_{i2}^*, u_i) : i = 1, \dots, n\}$, given X_i , is

$$\sum_{i=1}^n \log f(y_{i1}^*, y_{i2}^*, y_{i1}, y_{i2}, u_i | x_i) = \sum_{i=1}^n \log f(y_{i1}, y_{i2}, u_i | y_{i1}^*, y_{i2}^*, x_i) + \sum_{i=1}^n \log f(y_{i1}^*, y_{i2}^* | x_i).$$

Thus, the E-step of the EM algorithm evaluates

$$Q(\theta, \theta^{(t)}) = \sum_{i=1}^n Q_i(\theta, \theta^{(t)}),$$

where

$$Q_i(\theta, \theta^{(t)}) = E_{Y_{i1}, Y_{i2}, u_i | Y_{i1}^*, Y_{i2}^*, X_i; \theta^{(t)}} \{ \log f(Y_{i1}^*, Y_{i2}^*, Y_{i1}, Y_{i2}, u_i | X_i; \theta) \},$$

and the expectation is taken with respect to the conditional distribution of (Y_{i1}, Y_{i2}, u_i) given $\{Y_{i1}^*, Y_{i2}^*, X_i\}$ with θ set as $\theta^{(t)}$, the estimate of θ at iteration t .

The M-step is to maximize $Q(\theta, \theta^{(t)})$ with respect to θ , which is equivalent to solving

$$\sum_{i=1}^n \frac{\partial Q_i(\theta, \theta^{(t)})}{\partial \theta} = 0 \quad (2.15)$$

for θ , provided regularity conditions. To obtain the solution of (2.15), we may implement the Newton-Raphson algorithm. An updated estimate $\theta^{(t+1)}$ for θ at iteration t is given by

$$\theta^{(t+1)} = \theta^{(t)} + \left\{ \sum_{i=1}^n \frac{\partial^2}{\partial \theta \partial \theta^T} Q_i(\theta, \theta^{(t)}) \right\}^{-1} \left\{ \sum_{i=1}^n \frac{\partial}{\partial \theta} Q_i(\theta, \theta^{(t)}) \right\}, \quad (2.16)$$

where $t = 0, 1, 2, \dots$, and $\theta^{(0)}$ is an initial value of θ .

The integral involved in (2.16) can be approximated by numeric methods, such as the Monte Carlo or Gaussian quadrature algorithm. Repeat through the E and M steps until convergence of $\{\theta^{(t+1)} : t = 0, 1, \dots\}$, and let $\hat{\theta}$ denote the resulting limit. The variance estimate of $\hat{\theta}$ can be obtained following [Louis \(1982\)](#) or using the bootstrap procedure.

2.4 Extension to Handling Clustered Data

In genome-wide association studies (GWAS), facilitating the genetic relatedness for subjects in the same clusters (e.g., families) is important to reflect the cluster structure of data. To this end, we extend the preceding development by allowing the matrix R in Section 2.1.1 to feature the inherent relatedness within the same cluster or family.

Let

$$R = \begin{pmatrix} R_1 & & & \\ & R_2 & & \\ & & \ddots & \\ & & & R_{n_f} \end{pmatrix}$$

be a blockwise-diagonal matrix which delineates the population stratification based on the pedigree information, where n_f is the total number of families (or clusters) and R_i is the $n_i \times n_i$ relatedness matrix of family i . For $i = 1, \dots, n_f$, the (i_1, i_2) element $R_{i_1 i_2}$ of R_i may be, for example, defined as the kinship coefficient (Lange, 2003, Page 82) for subjects i_1 and i_2 in the family i , which is the weighted summation of the probabilities of each allele pair for subjects i_1 and i_2 to be *identical by descent* (IBD) at the same locus k . Here an allele pair is taken as *identical by descent* if the pair has the same type of nucleotide and is inherited from the same ancestor.

To be specific, for $k = 1, \dots, p_x$,

$$R_{i_1 i_2} = \frac{1}{4} \sum_{l=1}^2 \sum_{s=1}^2 P \left(X_{i_1 k}^{(l)} = X_{i_2 k}^{(s)} \right),$$

where $P(X_{i_1 k}^{(l)} = X_{i_2 k}^{(s)})$ is assumed to be identical for all the k and represents the common probability that the two alleles, $X_{i_1 k}^{(l)}$ and $X_{i_2 k}^{(s)}$, are inherited from the same ancestor for $l = 1, 2$ and $s = 1, 2$. Here $X_{i_1 k}^{(1)}$ and $X_{i_1 k}^{(2)}$ represent the nucleotides inherited from the father and mother of subject i , respectively; in applications, the probabilities $P(X_{i_1 k}^{(l)} = X_{i_2 k}^{(s)})$ are often determined by pedigree data. For instance, $R_{i_1 i_2} = 0.5$ if i_1 and i_2 are monozygotic twins and $R_{i_1 i_2} = 0.25$ if i_1 and i_2 has a parent-offspring relationship.

For $i = 1, \dots, n_f$, let $Y_{i11}, \dots, Y_{in_i1}$ be the continuous responses and $Y_{i12}, \dots, Y_{in_i2}$ be the binary responses of n_i subjects in the i th family, and let $Y_{i11}^*, \dots, Y_{in_i1}^*$ and $Y_{i12}^*, \dots, Y_{in_i2}^*$ be their corresponding surrogate measurements. For $i = 1, \dots, n_f$ and $r = 1, \dots, n_i$, we write $Y_{ir} = (Y_{ir1}, Y_{ir2})^T$, $Y_{ir}^* = (Y_{ir1}^*, Y_{ir2}^*)^T$, and $Y_i^* = (Y_{i1}^{*T}, \dots, Y_{in_i}^{*T})^T$. Then for $i = 1, \dots, n_f$ and $r = 1, \dots, n_i$, the response model (2.1) and the mismeasurement models in Section 2.1.2 are used to describe Y_{ir} and Y_{ir}^* where the random effect in (2.1) is now denoted as u_{ir} .

With this setup, the conditional distribution (2.10) is now modified to be

$$f(y_i^* | x_i) = \int \prod_{r=1}^{n_i} f(y_{ir}^* | \tilde{u}_i, x_i) f(\tilde{u}_i) d\tilde{u}_i,$$

where $\tilde{u}_i = (u_{i1}, \dots, u_{in_i})^T$ follows a multivariate normal distribution with mean zero and covariance matrix $\sigma_g^2 R_i$. Then, the inference about the model parameter θ can be carried out using the same procedure in Section 2.2 or 2.3, and the asymptotic distribution for the resulting estimator can be established in a similar manner.

2.5 Simulation Studies

We conduct simulation studies to evaluate the performance of the proposed method in terms of parameter estimates and associated variance estimates. In contrast, we also consider three naive methods. In the first naive method (Naive Method 1), we ignore both misclassification and measurement error in response variables and estimate the parameters of the response model by fitting a generalized linear model using R function `glm()` directly to the observed response measurements; in the second naive method (Naive Method 2), we ignore misclassification in the binary response but account for continuous response measurement error; and in the third analysis (Naive Method 3), we ignore continuous response measurement error but just address misclassification in the binary response.

The sample size is set as $n = 1000$ and we consider model (2.1) with $p_x = 2$ and model (2.6) with $p_z = 1$ and covariates Z_i independently generated from the uniform distribution $U(0, 2)$. To generate covariates X_i for model (2.1), we consider two scenarios with different nature in X_i . In Scenario 1, covariates are continuous where X_{i1} and X_{i2} are independently generated from $U(-3, 4)$ and $N(0, 1)$, respectively. In Scenario 2, covariates are ordinal representing a genotype shown in model (2.2); specifically, $X_{ij} = X_{ij}^{(1)} + X_{ij}^{(2)}$ for $j = 1, 2$ where $X_{i1}^{(1)}$ and $X_{i1}^{(2)}$ are independently generated from Bernoulli(0.2), and $X_{i2}^{(1)}$ and $X_{i2}^{(2)}$ are independently generated from Bernoulli(0.5); here different success probabilities of the Bernoulli distribution are chosen to reflect different minor allele frequencies (MAF) of genotypes.

2.5.1 Performance of the Methods in Sections 2.2 and 2.3: Simulation Design

For $i = 1, \dots, n$, the random effects u_i , featuring the correlation between the continuous and discrete responses Y_{i1} and Y_{i2} , are independently generated from $N(0, \sigma_g^2 R_{ii})$, where $R_{ii} = 1$ for $i = 1, \dots, 500$, and $R_{ii} = 2$ for $i = 501, \dots, 1000$, and σ_g is set as 0.8. The response vector $Y_i = (Y_{i1}, Y_{i2})^T$ is then generated from the joint distribution

$$f(y_{i1}, y_{i2} | u_i) = \frac{1}{\sqrt{2\pi}} \exp \left[-\frac{1}{2} \{y_{i1} - g_1(\mu_{i1})\}^2 \right] g_2(\mu_{i2})^{y_{i2}} \{1 - g_2(\mu_{i2})\}^{1-y_{i2}},$$

where $g_1(\mu_{i1})$ and $g_2(\mu_{i2})$ are specified as in model (2.1) with $g_1(t) = t$ and $g_2(t) = \log\left(\frac{t}{1-t}\right)$, and the coefficient $\beta = (\beta_{10}, \beta_{20}, \beta_{11}, \beta_{21}, \beta_{12}, \beta_{22})^T$ is set as $(0.7, 1.5, 0.7, -1.2, 1, 1)^T$.

The surrogate measurement Y_{i1}^* is generated from the measurement error model (2.8) with $\gamma_0 = 0$, $\gamma_1 = 1$, $\gamma_3 = 0$ and $f(x) = 2x - 1$ to transform the values of Y_{i2} from $\{0, 1\}$

into $\{-1, 1\}$. For the misclassification of Y_{i2} , we generate the surrogate measurement Y_{i2}^* from the model

$$\text{logit } \pi_{i1} = \alpha_{01} + \alpha_{z1}Z_i,$$

and

$$\text{logit } \pi_{i0} = \alpha_{00} + \alpha_{z0}Z_i,$$

where $\alpha = (\alpha_{01}, \alpha_{z1}, \alpha_{00}, \alpha_{z0})^T$ is the vector of parameters to be specified.

We consider four settings with different degrees of measurement error and misclassification rates. Settings 1 and 2 differ in the value of γ_2 , with $\gamma_2 = 0.001$ for Setting 1, and $\gamma_2 = 1.0$ for Setting 2; in these two settings, σ_e is set 0.25 or 0.50 to reflect increasing degrees of measurement error in Y_{i1} and α is set as $(-1.386, 0, -1.386, 0)^T$. In Settings 3 and 4, we take $\sigma_e = 1.0$ and $\gamma_2 = 1.0$ but consider different values for α ; in Setting 3, we let $\alpha_{z1} = \alpha_{z0} = 0$ and set $\alpha_{01} = \alpha_{00}$ to be -4.595 or -2.197 , respectively, yielding 1% and 10% misclassification proportions; and in Setting 4, we set $\alpha_{01} = \alpha_{00} = -1$ and let $\alpha_{z1} = \alpha_{z0}$ take a value of -3.5 or -1.2 , leading to about 1% and 10% misclassification proportions, respectively.

2.5.2 Performance of the Method in Section 2.4: Simulation Design

In this simulation study, we consider the case where subjects are correlated by pairs. For $i = 1, \dots, 500$, the random effects $(u_{i1}, u_{i2})^T$ are generated from a bivariate normal distribution with mean zero and covariance matrix $\sigma_g^2 R_i$, where $\sigma_g = 0.8$ and $R_i = \begin{pmatrix} 1 & \rho \\ \rho & 1 \end{pmatrix}$. Here, ρ is taken as 0.25 or 0.5, respectively, to possibly represent the parent-offspring relationship or monozygotic twin relationship of pairs (Lange, 2003). The response vector for cluster i , $Y_i = (Y_{i11}, Y_{i12}, Y_{i21}, Y_{i22})^T$, is then generated from the joint distribution

$$f(y_{i11}, y_{i12}, y_{i21}, y_{i22} | u_i) = \frac{1}{2\pi} \exp \left[-\frac{1}{2} \{y_{i11} - g_1(\mu_{i11})\}^2 - \frac{1}{2} \{y_{i21} - g_1(\mu_{i21})\}^2 \right] \\ \times g_2(\mu_{i2})^{y_{i12}} \{1 - g_2(\mu_{i12})\}^{1-y_{i12}} g_2(\mu_{i22})^{y_{i22}} \{1 - g_2(\mu_{i22})\}^{1-y_{i22}},$$

where $g_1(\mu_{i11})$, $g_1(\mu_{i21})$, $g_2(\mu_{i12})$ and $g_2(\mu_{i22})$ are specified as in model (2.1) with $g_1(t) = t$ and $g_2(t) = \log\left(\frac{t}{1-t}\right)$, and the coefficient $\beta = (\beta_{10}, \beta_{20}, \beta_{11}, \beta_{21}, \beta_{12}, \beta_{22})^T$ is set as $(0.7, 1.5, 0.7, -1.2, 1, 1)^T$. We comment that the correlation among the components of Y_i is facilitated by the inclusion of random effects $(u_{i1}, u_{i2})^T$ in $(\mu_{i11}, \mu_{i21}, \mu_{i12}, \mu_{i22})^T$.

The surrogate Y_{i1}^* and Y_{i2}^* are generated in the same way as in Section 2.5.1, with $\sigma_e = 0.25$, $\gamma_2 = 1$, $\alpha_{z1} = \alpha_{z0} = 0$ and $\alpha_{z1} = \alpha_{z0} = -1.386$, yielding a misclassification rate about 20%.

2.5.3 Simulation Results

We report the results obtained from the proposed methods and Naive Methods 1–3 in Tables 2.1–2.5 here.

In Tables 2.1–2.4, we report the simulation results for Section 2.5.1 and in Table 2.5 we display the results for Section 2.5.2, where “Bias” is calculated as $\frac{1}{N} \sum_{i=1}^N \hat{\theta}_k - \theta_k$, “SEE” is the empirical standard error, “SEM” is calculated by (2.13), and “CR” stands for the coverage rate of 95% confidence intervals for a parameter. That is for parameter θ_k , its *CR* is given by

$$\frac{1}{N} \sum_{i=1}^N I(\hat{\theta}_k^{(L)} < \theta_k < \hat{\theta}_k^{(U)}),$$

where N is the number of simulation studies repeated, $\hat{\theta}_k^{(L)}$ and $\hat{\theta}_k^{(U)}$ are calculated as $\hat{\theta}_k - \text{sd}(\hat{\theta}_k) \times Z_{0.975}$ and $\hat{\theta}_k + \text{sd}(\hat{\theta}_k) \times Z_{0.975}$, respectively. Here, $\hat{\theta}_k$ is an estimate of θ_k , $\text{sd}(\hat{\theta}_k)$ is the associated standard error, and $Z_{0.975}$ is the 0.975 quantile of $N(0, 1)$.

Simulation results demonstrate that in the presence of mismeasurement in the response components, the naive methods incur various kinds of biases although they may differ in the magnitude for different settings. The naive methods generally produce large finite sample biases and unreliable coverage rates for 95% confidence intervals that way off the nominal level 95%. On the other hand, the two methods that correct for the mismeasurement effects work very well for various settings regardless of whether the subjects are independent or clustered. These methods yield small finite sample biases for the point estimates and fairly good coverage rates for 95% confidence intervals.

2.6 Analysis of Mice SNPs Data

In this section, we illustrate our methods by analyzing the outbred Carworth Farms White (CFW) mice data arising from a genome-wide association study. (Parker et al., 2016a,b) This study provided measurements for 1200 mice on complex traits, including behavioral, physiological and gene expression traits. The original data contain measurements of 99787 SNPs for 1200 mice. Mice with missing responses and the SNPs with the minor allele frequency (MAF) lower than 0.05 are removed because such SNPs have low heterozygosity and often lead to false-positive results in association tests (Anderson et al., 2010). We examine the subset with 1157 mice and 77838 SNPs.

For $i = 1, \dots, 1157$, let Y_{i1} denote the true length of the tibia bone (in *mm*) and let Y_{i2} be a binary outcome where “0” represents a healthy bone and “1” stands for an abnormal

bone. The surrogate Y_{i1}^* is obtained in the laboratory and may differ from the true length Y_{i1} , and Y_{i2}^* is measured by a subjective classification rule based on the 90 percentile of bone-mineral density (BMD).

To analyze how the true responses are associated with the SNPs using the proposed method with mismeasurement effects accounted for, we carry out three steps of analysis. The first two steps are performed to screen unimportant SNPs to reduce the dimension of SNPs that is substantially larger than the sample size. The third step is to carry out a refined, post-screening analysis by applying the proposed method to the bivariate generalized linear mixed model (2.1) with measurement error effects taken into account.

In Step 1, we conduct the principal component analysis (PCA) (Price et al., 2006). Let G denote the $n \times n$ genomic relationship matrix using the genetic data following the discussion of Section 3.2 in VanRaden (2008). Then we express G using the eigenvalue decomposition (EVD),

$$G = LDL^T,$$

where the columns of L are the eigenvectors of G , and D is the diagonal matrix of the positive eigenvalues of G . Let $F = GLD^{-\frac{1}{2}}$ be the matrix of principal components of the genetic information, with the i th row, denoted as F_i , representing the principal components for subject i . According to the scree plot in Figure 2.1 and using the ‘‘elbow’’ criterion, we include the first five principal components, denoted as $F_{i1}, F_{i2}, F_{i3}, F_{i4}$ and F_{i5} , for subject i as the fixed effects when building the response model.

In Step 2, we conduct a genomewide screening procedure by examining each SNP one at a time using a model similar to (2.1). To adjust for the population stratification, we also include five largest principal components $PC_i = (F_{i1}, F_{i2}, F_{i3}, F_{i4}, F_{i5})^T$ for subject i . Let W_{ij} be the j th SNP for subject i and $j = 1, \dots, p_{\text{SNP}}$, where p_{SNP} is the dimension of SNPs. We repeat the screening for $j = 1, \dots, p_{\text{SNP}}$ by respectively considering the model with error effects adjusted:

$$g_1(\mu_{i1}) = \beta_{10}^* + \beta_{11}^* W_{ij} + PC_i^T \cdot \beta_{PC1}^* + u_i^*; \quad (2.17)$$

$$g_2(\mu_{i2}) = \beta_{20}^* + \beta_{21}^* W_{ij} + PC_i^T \cdot \beta_{PC2}^* + u_i^*; \quad (2.18)$$

where $g_1(\cdot)$ is set as the identity function, $g_2(t) = \log\left(\frac{t}{1-t}\right)$, u_i^* is the random effect, and $\beta_{10}^*, \beta_{11}^*, \beta_{20}^*, \beta_{21}^*, \beta_{PC1}^*$ and β_{PC2}^* are parameters.

To feature the misclassification of Y_{i2}^* , we use model (2.6) with Z_i taken as the bone-mineral density (BMD). Regarding the measurement error in Y_{i1} , following the discussion in Appendix A.1, we consider model (2.8) with $\gamma_0 = 0$, $\gamma_1 = 1$, $\gamma_3 = 0$ and $f(x) = 2x - 1$ for transforming the values of Y_{i2} from $\{0, 1\}$ into $\{-1, 1\}$. Then we perform the Wald

test to (2.17) with the null hypothesis $H_0 : \beta_{11}^* = 0$ and to (2.18) with the null hypothesis $H_0 : \beta_{21}^* = 0$, respectively, where we employ the induced likelihood method and the EM algorithm as opposed to the naive method without addressing error-in-variables.

In Figure 2.2, we report the Manhattan plot for each method which displays the resulting distribution of the SNP significant level, where SNPs are placed on the x-axis according to their chromosomal position, and the $-\log_{10}$ of the SNP associated p-values obtained from the Wald test are recorded on the y-axis. Using the significance level 10^{-6} as a threshold, we retain three SNPs, *rs31681083* (chromosome 8), *rs33030459* (chromosome 9) and *rs265727287* (chromosome 12) for our post-selection analysis in Step 3.

Finally, in Step 3, we build a final model with form (2.1) where X_i include the three selected SNPs, *rs31681083* (X_{i1}), *rs33030459* (X_{i2}) and *rs265727287* (X_{i3}), as well as the body weight of a mouse (X_{i4}) and the five largest principal components $\text{PC} = (F_{i1}, F_{i2}, F_{i3}, F_{i4}, F_{i5})^T$. That is,

$$\begin{bmatrix} g_1(\mu_{i1}) \\ g_2(\mu_{i2}) \end{bmatrix} = \begin{bmatrix} \beta_{10} + \beta_{11}X_{i1} + \beta_{12}X_{i2} + \beta_{13}X_{i3} + \beta_{14}X_{i4} + \text{PC}^T \cdot \beta_{\text{PC1}} \\ \beta_{20} + \beta_{21}X_{i1} + \beta_{22}X_{i2} + \beta_{23}X_{i3} + \beta_{24}X_{i4} + \text{PC}^T \cdot \beta_{\text{PC2}} \end{bmatrix} + \begin{bmatrix} u_i \\ u_i \end{bmatrix} \quad (2.19)$$

with $g_1(t) = t$ and $g_2(t) = \log\left(\frac{t}{1-t}\right)$ as well as random effects u_i .

We apply the induced likelihood method and the EM algorithm in contrast the naive method ignoring the error effects to fit model (2.19), and present the results in Table 2.6. Both the induced likelihood method and the EM algorithm produce fairly close results, and they suggest the same evidence for significance or insignificance of each covariates in model (2.19). At the significance level 0.05, the SNPs *rs31681083* (X_{i1}), *rs33030459* (X_{i2}) and *rs265727287* (X_{i3}) are significantly associated with tibia length, and the SNPs *rs31681083* (X_{i1}) and *rs33030459* (X_{i2}) are significantly associated with the bone condition. It is also observed that the bodyweight (X_{i4}) is significantly associated with both tibia length and bone condition. However, in the naive analysis which disregards mismeasurement effects, we obtain different evidence that *rs31681083* (X_{i1}), *rs33030459* (X_{i2}) and bodyweight (X_{i4}) are not significantly associated with the bone condition. It also shows the opposite evidence for the effect of SNP *rs33030459* (X_{i2}) on the bone condition from that revealed by the methods of accommodating mismeasurement effects.

The analyses also reveal evidence of misclassification in the binary response Y_{i2} , reflected by the estimation results of α_{z0} and α_{z1} in Table 2.6. For healthy bones, a lower BMD is associated with a higher probability of misclassification as the estimate of α_{z0} is negative, and for abnormal bones, a higher BMD is associated with a higher probability of misclassification as the estimate of α_{z1} is positive. In addition, the estimate of γ_2 is significantly negative, suggesting the measurement error in tibia length (Y_{i1}) is negatively dependent on the true bone condition (Y_{i2}).

Table 2.1: Simulation results: Settings 1 and 2 of the Scenario 1 in the first simulation study

σ_e	Naive Method										Proposed Methods														
	Naive Method 1					Naive Method 2					Naive Method 3					Induced Likelihood Method					EM Algorithm				
	Bias	SEE	SEM	CR%		Bias	SEE	SEM	CR%		Bias	SEE	SEM	CR%		Bias	SEE	SEM	CR%		Bias	SEE	SEM	CR%	
	Setting 1: $\gamma = 0.01$																								
	β_{10}	0.001	0.048	0.049	95.7	0.001	0.048	0.047	95.0	0.002	0.047	0.049	95.9	0.001	0.055	0.056	95.8	0.001	0.055	0.055	95.8	0.001	0.055	0.055	95.8
	β_{11}	0.001	0.023	0.024	95.6	0.020	0.028	0.034	97.1	0.000	0.022	0.024	96.7	0.001	0.058	0.055	92.5	0.002	0.059	0.055	92.4	0.000	0.059	0.055	92.4
	β_{12}	0.001	0.047	0.048	95.7	-0.002	0.047	0.046	94.1	0.001	0.047	0.047	95.4	0.001	0.055	0.054	95.2	0.000	0.055	0.054	95.7	0.000	0.055	0.054	95.7
	β_{20}	-0.470	0.074	0.073	0.0	-0.469	0.076	0.084	0.0	0.050	0.400	0.356	93.4	0.044	0.415	0.360	93.4	0.050	0.414	0.360	93.7	0.050	0.414	0.360	93.7
0.25	β_{21}	1.016	0.039	0.038	0.0	0.931	0.042	0.048	0.0	-0.084	0.370	0.299	89.9	-0.075	0.393	0.327	91.9	-0.085	0.388	0.329	92.5	-0.085	0.388	0.329	92.5
	β_{22}	-0.726	0.070	0.072	0.0	-0.716	0.073	0.082	0.0	0.054	0.351	0.299	92.2	0.047	0.361	0.310	91.0	0.055	0.359	0.312	91.8	0.055	0.359	0.312	91.8
	γ_2	-	-	-	-	-1.170	0.027	0.119	0.0	-	-	-	-	-0.004	0.161	0.143	92.7	-0.001	0.157	0.143	92.8	-0.001	0.157	0.143	92.8
	α_{00}	-	-	-	-	-	-	-	-	-0.040	0.248	0.211	94.9	-0.079	0.409	0.342	95.1	-0.076	0.594	1.164	95.2	-0.076	0.594	1.164	95.2
	α_{01}	-	-	-	-	-	-	-	-	-0.063	0.827	0.205	95.2	-0.078	0.396	0.327	95.4	-0.079	0.984	0.283	95.1	-0.079	0.984	0.283	95.1
	σ_g	-	-	-	-	-0.191	0.027	0.138	100	0.001	0.098	0.039	54.4	0.006	0.125	0.119	95.1	0.004	0.127	0.118	93.8	0.004	0.127	0.118	93.8
	β_{10}	0.002	0.050	0.051	95.8	-0.001	0.050	0.048	95.0	0.002	0.049	0.051	96.0	0.001	0.058	0.058	95.8	0.001	0.058	0.058	95.9	0.001	0.058	0.058	95.9
	β_{11}	0.001	0.024	0.025	95.5	0.003	0.030	0.034	98.0	0.001	0.023	0.025	96.7	0.002	0.061	0.058	92.6	0.003	0.062	0.058	92.4	0.003	0.062	0.058	92.4
	β_{12}	0.001	0.049	0.050	95.4	-0.001	0.049	0.047	94.0	0.001	0.048	0.049	95.6	0.000	0.057	0.057	95.5	-0.000	0.058	0.057	95.5	-0.000	0.058	0.057	95.5
	β_{20}	-0.470	0.074	0.073	0.0	-0.468	0.077	0.084	0.0	0.051	0.407	0.360	94.0	0.048	0.419	0.365	94.0	0.051	0.418	0.364	94.2	0.051	0.418	0.364	94.2
0.5	β_{21}	1.016	0.039	0.038	0.0	0.933	0.042	0.049	0.0	-0.086	0.383	0.306	90.9	-0.081	0.402	0.336	91.5	-0.087	0.399	0.337	92.3	-0.087	0.399	0.337	92.3
	β_{22}	-0.726	0.070	0.072	0.0	-0.715	0.073	0.082	0.0	0.056	0.360	0.303	91.7	0.052	0.372	0.316	90.8	0.057	0.369	0.317	91.1	0.057	0.369	0.317	91.1
	γ_2	-	-	-	-	-1.167	0.029	0.123	0.0	-	-	-	-	-0.002	0.165	0.151	92.2	0.001	0.166	0.151	92.1	0.001	0.166	0.151	92.1
	α_{00}	-	-	-	-	-	-	-	-	-0.049	0.344	0.213	95.0	-0.074	0.406	0.355	95.1	-0.089	0.747	0.715	95.1	-0.089	0.747	0.715	95.1
	α_{01}	-	-	-	-	-	-	-	-	-0.056	0.532	0.210	94.7	-0.074	0.395	0.331	95.4	-0.102	0.980	1.126	95.1	-0.102	0.980	1.126	95.1
	σ_g	-	-	-	-	-0.099	0.023	0.140	100	0.002	0.104	0.042	54.4	0.005	0.129	0.129	95.7	0.002	0.134	0.128	95.2	0.002	0.134	0.128	95.2
	Setting 2: $\gamma = 1$																								
	β_{10}	0.167	0.058	0.058	19.4	0.164	0.057	0.058	20	0.167	0.057	0.057	17.5	0.001	0.057	0.058	95.9	0.000	0.057	0.058	95.6	0.000	0.057	0.058	95.6
	β_{11}	-0.348	0.025	0.028	0.0	-0.336	0.027	0.045	0.0	-0.350	0.025	0.032	0.0	-0.001	0.047	0.049	95.4	0.002	0.049	0.049	94.5	0.002	0.049	0.049	94.5
	β_{12}	0.186	0.056	0.056	9.2	0.182	0.055	0.057	10	0.186	0.057	0.055	8.7	0.001	0.055	0.056	95.2	0.001	0.055	0.056	95.4	0.001	0.055	0.056	95.4
	β_{20}	-0.470	0.074	0.073	0.0	-0.466	0.078	0.090	0.0	-0.074	0.255	0.303	96.5	0.016	0.194	0.193	95.7	0.017	0.192	0.192	96.0	0.017	0.192	0.192	96.0
0.25	β_{21}	1.016	0.039	0.038	0.0	0.925	0.044	0.056	0.0	0.144	0.198	0.178	79.2	-0.028	0.163	0.165	95.0	-0.025	0.163	0.164	94.9	-0.025	0.163	0.164	94.9
	β_{22}	-0.726	0.070	0.072	0.0	-0.714	0.074	0.088	0.0	-0.141	0.227	0.231	87.9	0.015	0.196	0.197	95.7	0.014	0.198	0.196	95.2	0.014	0.198	0.196	95.2
	γ_2	-	-	-	-	-1.021	0.035	0.163	0.0	-	-	-	-	-0.004	0.115	0.121	95.8	0.005	0.121	0.120	94.6	0.005	0.121	0.120	94.6
	α_{00}	-	-	-	-	-	-	-	-	-0.266	0.219	0.209	84.9	-0.002	0.132	0.132	94.6	-0.008	0.134	0.133	94.7	-0.008	0.134	0.133	94.7
	α_{01}	-	-	-	-	-	-	-	-	-0.253	0.215	0.204	84.8	-0.004	0.131	0.131	94.6	-0.008	0.133	0.131	94.7	-0.008	0.133	0.131	94.7
	σ_g	-	-	-	-	0.017	0.031	0.173	100	0.014	0.084	0.039	0.0	0.007	0.118	0.122	95.6	-0.001	0.127	0.122	94.2	-0.001	0.127	0.122	94.2
	β_{10}	0.167	0.060	0.060	21.9	0.162	0.058	0.062	26	0.168	0.059	0.059	19.3	0.003	0.059	0.060	95.7	0.000	0.060	0.061	95.5	0.000	0.060	0.061	95.5
	β_{11}	-0.348	0.026	0.029	0.0	-0.343	0.030	0.049	0.0	-0.350	0.026	0.033	0.0	-0.003	0.049	0.052	95.9	0.003	0.052	0.052	94.8	0.003	0.052	0.052	94.8
	β_{12}	0.186	0.057	0.058	9.8	0.182	0.057	0.061	14	0.186	0.057	0.057	10.2	0.003	0.057	0.058	95.3	0.001	0.057	0.058	95.5	0.001	0.057	0.058	95.5
	β_{20}	-0.470	0.074	0.073	0.0	-0.462	0.080	0.091	0.0	-0.061	0.264	0.307	96.8	0.014	0.198	0.201	95.9	0.016	0.198	0.199	96.1	0.016	0.198	0.199	96.1
0.5	β_{21}	1.016	0.039	0.038	0.0	0.910	0.049	0.059	0.0	0.124	0.209	0.182	80.9	-0.034	0.169	0.173	95.4	-0.026	0.171	0.172	94.9	-0.026	0.171	0.172	94.9
	β_{22}	-0.726	0.070	0.072	0.0	-0.709	0.076	0.089	0.0	-0.127	0.236	0.234	87.6	0.017	0.203	0.203	95.4	0.014	0.205	0.202	95.0	0.014	0.205	0.202	95.0
	γ_2	-	-	-	-	-1.024	0.042	0.179	0.0	-	-	-	-	-0.010	0.120	0.128	96.2	0.006	0.129	0.128	94.8	0.006	0.129	0.128	94.8
	α_{00}	-	-	-	-	-	-	-	-	-0.259	0.232	0.212	85.9	-0.001	0.136	0.135	94.7	-0.009	0.137	0.135	94.7	-0.009	0.137	0.135	94.7
	α_{01}	-	-	-	-	-	-	-	-	-0.248	0.220	0.206	85.8	-0.004	0.134	0.133	94.8	-0.008	0.136	0.133	94.6	-0.008	0.136	0.133	94.6
	σ_g	-	-	-	-	-0.034	0.037	0.189	100	0.427	0.093	0.041	0.0	0.016	0.120	0.130	96.2	-0.002	0.134	0.132	95.2	-0.002	0.134	0.132	95.2

Table 2.2: Simulation results: Settings 3 and 4 of the Scenario 1 in the first simulation study

α_0	α_z	Naive Method 1						Naive Method 2						Naive Method 3						Induced Likelihood Method						Proposed Methods					
		Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%						
Setting 3																															
		β_{10}	0.168	0.066	0.066	27.9	0.138	0.076	0.076	54.2	0.169	0.067	0.064	25.1	0.001	0.055	0.056	95.8	0.004	0.062	0.064	95.8	0.004	0.062	0.064	95.8					
		β_{11}	-0.348	0.029	0.032	0.0	-0.200	0.075	0.075	19.2	-0.349	0.029	0.036	0.0	0.001	0.058	0.055	95.2	0.002	0.061	0.061	95.2	0.002	0.061	0.061	95.6					
		β_{12}	0.186	0.063	0.064	17.5	0.145	0.074	0.074	50.8	0.186	0.064	0.062	16.3	0.001	0.055	0.054	95.2	0.000	0.063	0.064	95.1	0.000	0.063	0.064	95.1					
		β_{20}	-0.141	0.100	0.100	69.7	-0.122	0.107	0.134	89.9	0.117	0.145	0.172	93.3	0.044	0.415	0.360	93.4	0.009	0.136	0.140	95.1	0.009	0.136	0.140	95.1					
	-4.597	0.0	β_{21}	0.306	0.072	0.071	2.9	0.113	0.107	0.117	86.6	-0.257	0.104	0.119	40.3	-0.075	0.393	0.327	91.9	-0.020	0.122	0.126	95.9	-0.020	0.122	0.126	95.9				
		β_{22}	-0.210	0.104	0.103	46.1	-0.179	0.116	0.140	77.0	0.127	0.152	0.161	90.8	0.047	0.361	0.310	91.0	0.013	0.143	0.143	95.5	0.013	0.143	0.143	95.5					
		γ_2	-	-	-	-	-0.604	0.185	0.196	9.3	-	-	-	-	-0.004	0.161	0.143	92.7	0.001	0.153	0.153	95.5	0.001	0.153	0.153	95.5					
		α_{00}	-	-	-	-	-	-	-	-	-2.512	5.326	1.705	99.7	-	-	-	-	-	-	-	-	-	-	-	-					
		α_{01}	-	-	-	-	-	-	-	-	-2.292	4.636	1.515	98.4	-	-	-	-	-	-	-	-	-	-	-	-					
		σ_g	-	-	-	-	0.255	0.104	0.184	97.0	0.677	0.052	0.042	0.0	0.006	0.125	0.119	95.4	-0.732	4.683	2.084	94.3	-0.732	4.683	2.084	94.3					
		β_{10}	0.168	0.066	0.066	27.9	0.166	0.065	0.073	35.1	0.169	0.065	0.065	25.7	0.009	0.065	0.067	95.6	0.002	0.067	0.067	95.6	0.002	0.067	0.067	95.2					
		β_{11}	-0.348	0.029	0.032	0.0	-0.308	0.030	0.068	0.5	-0.349	0.029	0.036	0.0	-0.014	0.059	0.061	95.6	0.003	0.063	0.062	94.7	0.003	0.063	0.062	94.7					
		β_{12}	0.186	0.063	0.064	17.5	0.182	0.063	0.073	28.9	0.186	0.063	0.063	17.0	0.011	0.063	0.065	95.4	0.001	0.064	0.065	95.6	0.001	0.064	0.065	95.6					
		β_{20}	-0.350	0.085	0.082	1.9	-0.349	0.087	0.108	7.2	0.043	0.208	0.225	96.4	0.018	0.184	0.181	95.2	0.012	0.180	0.179	95.4	0.012	0.180	0.179	95.4					
	-2.197	0.0	β_{21}	0.752	0.050	0.048	0.0	0.644	0.052	0.082	0.1	-0.098	0.162	0.149	91.2	-0.036	0.155	0.160	96.5	-0.022	0.159	0.160	95.2	-0.022	0.159	0.160	95.2				
		β_{22}	-0.545	0.080	0.082	0.1	-0.537	0.082	0.108	0.0	0.023	0.190	0.194	95.5	0.022	0.171	0.179	96.2	0.012	0.177	0.178	95.1	0.012	0.177	0.178	95.1					
		γ_2	-	-	-	-	-0.915	0.033	0.217	0.0	-	-	-	-	-0.047	0.145	0.155	94.8	0.004	0.158	0.155	94.1	0.004	0.158	0.155	94.1					
		α_{z0}	-	-	-	-	-	-	-	-	-0.287	0.311	0.285	92.9	-	-	-	-	-	-	-	-	-	-	-	-					
		α_{z1}	-	-	-	-	-	-	-	-	-0.276	0.285	0.274	92.9	-0.070	0.212	0.213	95.4	-0.022	0.217	0.211	95.4	-0.022	0.217	0.211	95.4					
		σ_g	-	-	-	-	0.212	0.039	0.213	100.0	0.585	0.085	0.045	0.0	0.046	0.138	0.156	95.2	-0.002	0.161	0.160	96.1	-0.002	0.161	0.160	96.1					
Setting 4																															
		β_{10}	0.166	0.063	0.066	28.1	0.158	0.065	0.075	44.8	0.166	0.065	0.064	27.1	0.006	0.063	0.065	95.1	-0.000	0.064	0.065	95.1	-0.000	0.064	0.065	95.1					
		β_{11}	-0.348	0.029	0.032	0.0	-0.286	0.044	0.075	1.1	-0.349	0.029	0.036	0.0	-0.009	0.059	0.061	94.8	0.002	0.061	0.061	94.3	0.002	0.061	0.061	94.3					
		β_{12}	0.184	0.063	0.064	15.7	0.172	0.065	0.076	36.4	0.185	0.064	0.062	14.5	0.006	0.062	0.064	93.9	-0.001	0.065	0.064	93.4	-0.001	0.065	0.064	93.4					
		β_{20}	-0.243	0.095	0.091	23.9	-0.237	0.099	0.122	51.7	0.125	0.165	0.176	92.0	0.023	0.153	0.145	93.9	0.011	0.148	0.143	94.3	0.011	0.148	0.143	94.3					
		β_{21}	0.526	0.062	0.058	0.0	0.393	0.084	0.097	3.7	-0.263	0.114	0.123	41.9	-0.029	0.126	0.128	95.0	-0.016	0.128	0.127	94.8	-0.016	0.128	0.127	94.8					
	-1.000	-3.5	β_{22}	-0.382	0.091	0.092	1.7	-0.369	0.095	0.122	12.3	0.131	0.156	0.172	91.9	0.022	0.142	0.149	96.0	0.007	0.145	0.148	95.4	0.007	0.145	0.148	95.4				
		γ_2	-	-	-	-	-0.797	0.089	0.212	0.0	-	-	-	-	-0.033	0.149	0.154	95.0	0.003	0.154	0.155	94.7	0.003	0.154	0.155	94.7					
		α_{z0}	-	-	-	-	-	-	-	-	-0.069	0.424	0.495	98.4	-0.090	0.393	0.458	98.3	0.028	0.437	0.454	96.1	0.028	0.437	0.454	96.1					
		α_{z1}	-	-	-	-	-	-	-	-	-0.364	0.915	1.680	98.5	-0.042	0.398	1.406	94.6	-0.383	1.661	1.485	93.8	-0.383	1.661	1.485	93.8					
		σ_g	-	-	-	-	-	-	-	-	-0.406	0.914	1.676	99.2	-0.106	0.988	1.415	95.5	-0.493	1.803	1.516	95.4	-0.493	1.803	1.516	95.4					
		β_{10}	0.166	0.063	0.066	28.1	0.165	0.063	0.073	38.1	0.166	0.064	0.065	27.1	0.009	0.064	0.067	95.7	0.000	0.066	0.067	95.8	0.000	0.066	0.067	95.8					
		β_{11}	-0.348	0.029	0.032	0.0	-0.311	0.030	0.067	0.0	-0.349	0.029	0.036	0.0	-0.015	0.057	0.062	96.1	0.001	0.063	0.062	94.2	0.001	0.063	0.062	94.2					
		β_{12}	0.184	0.063	0.064	15.7	0.180	0.063	0.073	27.9	0.185	0.063	0.063	15.5	0.006	0.065	0.065	93.5	0.000	0.067	0.066	93.7	0.000	0.067	0.066	93.7					
		β_{20}	-0.372	0.085	0.080	1.2	-0.371	0.088	0.106	6.2	0.058	0.212	0.226	96.6	0.022	0.180	0.183	95.8	0.013	0.187	0.179	94.7	0.013	0.187	0.179	94.7					
		β_{21}	0.806	0.051	0.045	0.0	0.699	0.052	0.074	0.0	-0.113	0.162	0.151	90.8	-0.041	0.159	0.163	95.0	-0.017	0.171	0.160	94.0	-0.017	0.171	0.160	94.0					
	-1.000	-1.2	β_{22}	-0.585	0.081	0.079	0.0	-0.576	0.083	0.104	0.0	0.034	0.198	0.199	95.9	0.025	0.181	0.183	96.8	0.009	0.181	0.180	94.7	0.009	0.181	0.180	94.7				
		γ_2	-	-	-	-	-0.939	0.030	0.221	0.0	-	-	-	-	-0.045	0.146	0.156	94.5	0.000	0.158	0.157	94.5	0.000	0.158	0.157	94.5					
		α_{z0}	-	-	-	-	-	-	-	-	-0.035	0.324	0.328	97.0	-0.015	0.288	0.289	96.7	-0.009	0.347	0.296	96.4	-0.009	0.347	0.296	96.4					
		α_{z1}	-	-	-	-	-	-	-	-	-0.299	0.501	0.491	97.2	-0.023	0.358	0.366	95.2	-0.060	0.516	0.397	95.5	-0.060	0.516	0.397	95.5					
		σ_g	-	-	-	-	-	-	-	-	-0.310	0.515	0.484	95.8	-0.014	0.350	0.357	95.2	-0.092	0.767	0.408	95.6	-0.092	0.767	0.408	95.6					

Table 2.4: Simulation Results: settings 3 and 4 of the scenario 2 in the first simulation study

α_0	α_z	Naive Method 1						Naive Method 2						Naive Method 3						Induced Likelihood Method						Proposed Methods								
		Bias	SEM	CR%	CR%	SEM	CR%	Bias	SEM	CR%	CR%	SEM	CR%	Bias	SEM	CR%	CR%	SEM	CR%	Bias	SEM	CR%	CR%	SEM	CR%	Bias	SEM	CR%	CR%	SEM	CR%			
Setting 3																																		
		β_{10}	0.521	0.128	0.123	1.6	0.414	0.151	0.171	27.5	0.505	0.131	0.115	0.8	-0.009	0.143	0.143	95.1	-0.003	0.143	0.143	95.2	-0.003	0.143	0.143	95.1	-0.003	0.143	0.143	95.2	-0.003	0.143	0.143	95.2
		β_{11}	-0.316	0.122	0.116	23.9	-0.318	0.131	0.141	34.3	-0.272	0.126	0.107	20.5	0.011	0.116	0.118	95.3	0.016	0.116	0.118	95.6	-0.004	0.115	0.118	95.6	-0.004	0.115	0.118	95.6	-0.004	0.115	0.118	95.6
		β_{12}	0.234	0.094	0.093	28.5	0.134	0.105	0.117	82.8	0.228	0.098	0.090	28.6	0.001	0.092	0.092	94.6	-0.004	0.092	0.092	94.7	-0.004	0.091	0.092	94.7	-0.004	0.091	0.092	94.7	-0.004	0.091	0.092	94.7
		β_{20}	-0.302	0.145	0.146	46.2	-0.157	0.159	0.227	95.5	0.378	0.221	0.301	80.0	-0.015	0.208	0.228	96.7	-0.036	0.208	0.228	96.7	-0.036	0.208	0.228	96.7	-0.036	0.208	0.228	96.7	-0.036	0.208	0.228	96.7
-4.595	0.0	β_{21}	0.233	0.146	0.139	60.5	0.256	0.160	0.202	79.0	-0.235	0.228	0.222	83.3	-0.024	0.199	0.193	94.6	-0.042	0.199	0.193	94.7	-0.042	0.199	0.193	94.7	-0.042	0.199	0.193	94.7	-0.042	0.199	0.193	94.7
		β_{22}	-0.190	0.126	0.126	66.2	-0.045	0.138	0.183	98.1	0.183	0.187	0.196	90.4	0.024	0.169	0.172	95.8	0.027	0.169	0.172	95.6	0.027	0.169	0.172	95.6	0.027	0.169	0.172	95.6	0.027	0.169	0.172	95.6
		γ_2	-	-	-	-	-0.638	0.142	0.237	6.1	-	-	-	-	0.014	0.174	0.178	94.8	0.015	0.174	0.178	95.1	0.015	0.174	0.178	95.1	0.015	0.174	0.178	95.1	0.015	0.174	0.178	95.1
		α_{00}	-	-	-	-	-	-	-	-	0.124	0.281	2.012	99.4	-	-	-	92.0	-	-	-	92.1	-	-	-	92.1	-	-	-	92.1	-	-	-	92.1
		α_{01}	-	-	-	-	0.309	0.087	0.224	98.4	0.429	0.607	4.130	99.0	-0.733	2.480	2.833	86.7	-0.733	2.480	2.833	86.7	-0.733	2.480	2.833	86.7	-0.733	2.480	2.833	86.7	-0.733	2.480	2.833	86.7
		σ_g	0.521	0.128	0.123	1.6	0.494	0.127	0.148	5.4	0.521	0.129	0.117	1.0	0.000	0.154	0.153	94.9	-0.005	0.154	0.153	94.8	-0.005	0.156	0.153	94.8	-0.005	0.156	0.153	94.8	-0.005	0.156	0.153	94.8
		β_{11}	-0.316	0.122	0.116	23.9	-0.323	0.121	0.128	27.3	-0.319	0.123	0.109	20.7	0.004	0.123	0.123	94.1	0.009	0.122	0.123	94.3	0.009	0.122	0.123	94.3	0.009	0.122	0.123	94.3	0.009	0.122	0.123	94.3
		β_{12}	0.234	0.094	0.093	28.5	0.206	0.094	0.107	52.5	0.238	0.095	0.092	28.9	0.003	0.096	0.095	94.3	-0.001	0.096	0.095	94.4	-0.001	0.096	0.095	94.4	-0.001	0.096	0.095	94.4	-0.001	0.096	0.095	94.4
		β_{20}	-0.579	0.135	0.134	1.6	-0.514	0.138	0.197	19.1	0.307	0.297	0.354	89.0	-0.006	0.280	0.271	94.5	-0.019	0.280	0.271	94.4	-0.019	0.280	0.271	94.4	-0.019	0.280	0.271	94.4	-0.019	0.280	0.271	94.4
		β_{21}	0.535	0.131	0.126	1.5	0.553	0.134	0.177	8.5	-0.111	0.271	0.248	91.9	-0.020	0.246	0.230	93.8	-0.019	0.245	0.230	93.8	-0.019	0.245	0.230	93.8	-0.019	0.245	0.230	93.8	-0.019	0.245	0.230	93.8
-2.197	0.0	β_{22}	-0.479	0.111	0.109	1.0	-0.408	0.111	0.151	16.7	0.061	0.230	0.224	94.7	0.025	0.206	0.210	96.5	0.025	0.207	0.210	96.2	0.025	0.207	0.210	96.2	0.025	0.207	0.210	96.2	0.025	0.207	0.210	96.2
		γ_2	-	-	-	-	-0.887	0.030	0.232	0.0	-	-	-	-	0.000	0.172	0.182	96.0	0.012	0.175	0.182	95.2	0.012	0.175	0.182	95.2	0.012	0.175	0.182	95.2	0.012	0.175	0.182	95.2
		α_{00}	-	-	-	-	-	-	-	-	-0.297	0.279	0.261	87.4	-0.011	0.176	0.178	96.5	-0.002	0.179	0.177	95.8	-0.002	0.179	0.177	95.8	-0.002	0.179	0.177	95.8	-0.002	0.179	0.177	95.8
		α_{01}	-	-	-	-	0.261	0.038	0.230	100.0	-0.914	0.808	2.012	100.0	-0.039	1.604	1.042	93.7	-0.014	1.604	1.042	93.7	-0.014	1.604	1.042	93.7	-0.014	1.604	1.042	93.7	-0.014	1.604	1.042	93.7
		σ_g	0.494	0.081	0.084	0.0	0.464	0.086	0.143	2.0	0.500	0.086	0.084	0.0	0.002	0.110	0.110	95.3	0.009	0.139	0.143	94.9	0.009	0.139	0.143	94.9	0.009	0.139	0.143	94.9	0.009	0.139	0.143	94.9
		β_{11}	-0.390	0.123	0.121	10.9	-0.390	0.122	0.158	25.6	-0.397	0.132	0.113	9.5	0.002	0.122	0.123	95.2	0.002	0.122	0.123	95.2	0.002	0.122	0.123	95.2	0.002	0.122	0.123	95.2	0.002	0.122	0.123	95.2
		β_{12}	0.305	0.068	0.068	0.6	0.280	0.074	0.105	14.3	0.307	0.070	0.073	0.7	-0.003	0.078	0.078	94.1	-0.006	0.091	0.091	95.3	-0.006	0.091	0.091	95.3	-0.006	0.091	0.091	95.3	-0.006	0.091	0.091	95.3
		β_{20}	-0.426	0.095	0.093	0.8	-0.373	0.107	0.173	36.6	0.394	0.166	0.180	40.3	0.005	0.165	0.163	94.1	-0.017	0.218	0.216	94.3	-0.017	0.218	0.216	94.3	-0.017	0.218	0.216	94.3	-0.017	0.218	0.216	94.3
		β_{21}	0.343	0.130	0.126	22.3	0.343	0.133	0.198	61.7	-0.304	0.224	0.220	72.0	-0.007	0.189	0.186	95.1	-0.023	0.196	0.193	95.4	-0.023	0.196	0.193	95.4	-0.023	0.196	0.193	95.4	-0.023	0.196	0.193	95.4
-1.000	-3.5	β_{22}	-0.287	0.078	0.079	6.2	-0.246	0.088	0.133	54.1	0.238	0.134	0.143	63.4	0.012	0.127	0.125	95.0	0.021	0.179	0.173	95.2	0.021	0.179	0.173	95.2	0.021	0.179	0.173	95.2	0.021	0.179	0.173	95.2
		γ_2	-	-	-	-	-0.709	0.067	0.277	2.1	-	-	-	-	0.001	0.170	0.169	94.7	0.002	0.170	0.175	94.6	0.002	0.170	0.175	94.6	0.002	0.170	0.175	94.6	0.002	0.170	0.175	94.6
		α_{00}	-	-	-	-	-	-	-	-	-0.041	0.531	0.504	95.2	-	-	-	95.1	-	-	-	94.1	-	-	-	94.1	-	-	-	94.1	-	-	-	94.1
		α_{z0}	-	-	-	-	-	-	-	-	-0.866	2.236	2.096	98.1	-0.089	1.777	1.444	92.8	-1.329	2.009	1.832	94.9	-1.329	2.009	1.832	94.9	-1.329	2.009	1.832	94.9	-1.329	2.009	1.832	94.9
		α_{01}	-	-	-	-	-	-	-	-	-0.475	1.250	1.321	98.6	-0.416	5.080	2.927	90.1	-0.561	5.178	6.885	90.4	-0.561	5.178	6.885	90.4	-0.561	5.178	6.885	90.4	-0.561	5.178	6.885	90.4
		α_{z1}	-	-	-	-	-	-	-	-	-0.146	5.021	4.139	99.7	-0.013	0.179	0.179	95.9	-0.011	0.165	0.173	96.7	-0.011	0.165	0.173	96.7	-0.011	0.165	0.173	96.7	-0.011	0.165	0.173	96.7
		σ_g	0.494	0.081	0.084	0.0	0.400	0.058	0.280	99.7	0.800	0.045	0.050	0.0	0.002	0.116	0.116	94.9	0.010	0.142	0.147	95.1	0.010	0.142	0.147	95.1	0.010	0.142	0.147	95.1	0.010	0.142	0.147	95.1
		β_{10}	0.494	0.081	0.084	0.0	0.481	0.080	0.127	0.3	0.499	0.083	0.087	0.0	0.002	0.116	0.116	94.9	0.010	0.142	0.147	95.1	0.010	0.142	0.147	95.1	0.010	0.142	0.147	95.1	0.010	0.142	0.147	95.1
		β_{11}	-0.390	0.123	0.121	10.9	-0.391	0.122	0.146	19.3	-0.394	0.125	0.116	10.2	0.001	0.130	0.128	94.6	0.001	0.130	0.128	94.6	0.001	0.130	0.128	94.6	0.001	0.130	0.128	94.6	0.001	0.130	0.128	94.6
		β_{12}	0.305	0.068	0.068	0.6	0.294	0.067	0.094	5.5	0.307	0.069	0.075	1.0	0.001	0.081	0.081	95.0	-0.004	0.093	0.093	95.2	-0.004	0.093	0.093	95.2	-0.004	0.093	0.093	95.2	-0.004	0.093	0.093	95.2
		β_{20}	-0.646	0.087	0.087	0.0	-0.612	0.090																										

Table 2.5: Simulation results: the second simulation study

	Naive Method 1						Naive Method 2						Naive Method 3						Induced Likelihood Method						Proposed Methods														
	Bias		SEM		CR%		Bias		SEM		CR%		Bias		SEM		CR%		Bias		SEM		CR%		Bias		SEM		CR%										
	SEE	CR%	SEE	CR%	SEE	CR%	SEE	CR%	SEE	CR%	SEE	CR%	SEE	CR%	SEE	CR%	SEE	CR%	SEE	CR%	SEE	CR%	SEE	CR%	SEE	CR%	SEE	CR%											
Parent-offspring relationship																																							
β_{10}	0.552	0.104	0.098	0.0	0.295	0.111	0.118	27.6	0.556	0.109	0.098	0.0	0.008	0.124	0.125	95.8	0.025	0.123	0.132	96.6																			
β_{11}	-0.323	0.091	0.093	5.8	-0.172	0.089	0.098	58.2	-0.325	0.094	0.086	4.8	0.000	0.096	0.097	96.2	-0.013	0.096	0.102	95.9																			
β_{12}	0.224	0.074	0.074	15.0	0.116	0.071	0.078	69.4	0.223	0.079	0.072	14.4	-0.007	0.075	0.074	93.9	0.007	0.074	0.077	95.5																			
β_{20}	-0.532	0.145	0.136	4.6	-0.429	0.167	0.169	27.2	0.251	0.269	0.238	94.1	-0.005	0.265	0.239	92.1	0.025	0.300	0.291	93.3																			
β_{21}	0.516	0.128	0.127	1.8	0.458	0.143	0.151	13.8	0.027	0.226	0.241	97.5	-0.007	0.217	0.212	95.4	-0.036	0.252	0.257	95.9																			
β_{22}	-0.467	0.119	0.110	1.6	-0.428	0.128	0.127	9.6	-0.070	0.202	0.215	94.3	0.011	0.195	0.195	95.8	0.046	0.233	0.238	96.5																			
γ_2	-	-	-	-	-0.415	0.113	0.177	25.2	-	-	-	-	0.001	0.142	0.149	94.5	-0.041	0.113	0.151	98.2																			
α_{00}	-	-	-	-	-	-	-	-	-0.484	0.271	0.289	71.1	-0.023	0.159	0.156	94.9	-0.006	0.107	0.106	95.3																			
α_{01}	-	-	-	-	-	-	-	-	-2.163	0.875	3.955	100.0	-0.148	0.773	1.036	94.1	-0.103	1.108	0.604	95.3																			
σ_g	-	-	-	-	-0.209	0.187	0.619	99.8	0.701	0.045	0.064	0.0	-0.020	0.209	0.338	97.0	0.157	0.120	0.200	95.5																			
Monozygotic twin relationship																																							
β_{10}	0.552	0.109	0.098	0.0	0.335	0.109	0.099	9.4	0.552	0.107	0.104	0.0	0.007	0.110	0.109	94.8	-0.001	0.118	0.119	96.1																			
β_{11}	-0.323	0.093	0.093	5.8	-0.196	0.087	0.088	40.8	-0.324	0.092	0.084	5.0	0.001	0.088	0.089	95.0	0.001	0.095	0.096	94.8																			
β_{12}	0.223	0.076	0.074	17.0	0.133	0.070	0.072	56.4	0.223	0.075	0.070	12.5	-0.006	0.071	0.069	92.6	0.000	0.073	0.073	94.1																			
β_{20}	-0.531	0.144	0.136	3.6	-0.382	0.167	0.163	35.2	0.083	0.250	0.368	98.8	-0.001	0.255	0.229	91.8	0.018	0.291	0.277	93.2																			
β_{21}	0.516	0.130	0.127	1.4	0.432	0.149	0.149	19.0	0.142	0.215	0.231	89.9	-0.005	0.210	0.203	95.2	-0.025	0.258	0.243	94.2																			
β_{22}	-0.470	0.118	0.110	1.8	-0.414	0.131	0.126	10.4	-0.169	0.187	0.201	85.1	-0.001	0.189	0.188	95.6	0.018	0.225	0.226	95.7																			
γ_2	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-																			
α_{00}	-	-	-	-	-	-	-	-	-0.871	0.523	0.561	74.2	-0.024	0.158	0.156	94.8	-0.002	0.099	0.104	96.0																			
α_{01}	-	-	-	-	-	-	-	-	-2.732	0.315	5.631	100.0	-0.164	0.782	1.087	93.8	-0.091	0.607	0.419	95.9																			
σ_g	-	-	-	-	-0.034	0.110	0.133	98.8	0.597	0.089	0.095	0.0	-0.005	0.105	0.106	95.8	0.096	0.099	0.108	85.6																			

Table 2.6: Analysis results for the mice SNPs data

Parameter	Naive Analysis			Induced Likelihood Method			EM Algorithm		
	Estimate	S.E	p-value	Estimate	S.E	p-value	Estimate	S.E	p-value
Estimates for Response Models - Tibia Length									
β_{10}	17.141	0.121	<0.001	17.451	0.161	<0.001	17.469	0.166	<0.001
β_{11}	-0.058	0.022	0.009	-0.152	0.035	<0.001	-0.154	0.036	<0.001
β_{12}	-0.109	0.032	0.001	-0.204	0.043	<0.001	-0.210	0.045	<0.001
β_{13}	0.113	0.020	<0.001	0.131	0.026	<0.001	0.131	0.027	<0.001
β_{14}	0.047	0.004	<0.001	0.037	0.006	<0.001	0.036	0.006	<0.001
β_{15}	-0.002	0.001	0.005	-0.002	0.001	0.027	-0.003	0.001	0.013
β_{16}	0.002	0.001	0.156	-0.001	0.002	0.505	-0.002	0.002	0.355
β_{17}	0.002	0.001	0.207	-0.004	0.002	0.027	-0.005	0.002	0.024
β_{18}	0.000	0.001	0.695	-0.008	0.002	<0.001	-0.008	0.002	<0.001
β_{19}	-0.003	0.001	0.047	-0.008	0.002	<0.001	-0.008	0.002	<0.001
Estimates for Response Models - Tibia Length									
β_{20}	-1.364	1.052	0.195	6.360	2.696	0.018	6.137	2.575	0.017
β_{21}	-0.305	0.188	0.105	-1.590	0.532	0.003	-1.513	0.501	0.003
β_{22}	0.097	0.272	0.720	-2.245	0.851	0.008	-2.187	0.799	0.006
β_{23}	-0.170	0.171	0.320	0.363	0.393	0.356	0.336	0.376	0.371
β_{24}	-0.020	0.038	0.605	-0.219	0.097	0.024	-0.207	0.093	0.025
β_{25}	-0.004	0.008	0.617	0.001	0.015	0.946	-0.003	0.014	0.832
β_{26}	-0.002	0.010	0.830	-0.045	0.021	0.035	-0.048	0.021	0.019
β_{27}	-0.040	0.011	<0.001	-0.102	0.031	0.001	-0.096	0.030	0.001
β_{28}	-0.022	0.010	0.029	-0.138	0.038	<0.001	-0.132	0.036	<0.001
β_{29}	0.028	0.011	0.011	-0.102	0.031	0.001	-0.095	0.029	0.001
Estimates for Mismeasurement Models									
γ_2	-	-	-	-0.232	0.045	<0.001	-0.248	0.052	<0.001
α_{00}	-	-	-	17.794	3.954	<0.001	17.490	3.529	<0.001
α_{01}	-	-	-	-12.210	1.467	<0.001	-12.329	1.534	<0.001
α_{z0}	-	-	-	-0.171	0.039	<0.001	-0.168	0.035	<0.001
α_{z1}	-	-	-	0.099	0.013	<0.001	0.100	0.014	<0.001
σ_g	-	-	-	0.021	2.123	0.992	0.181	0.282	0.522

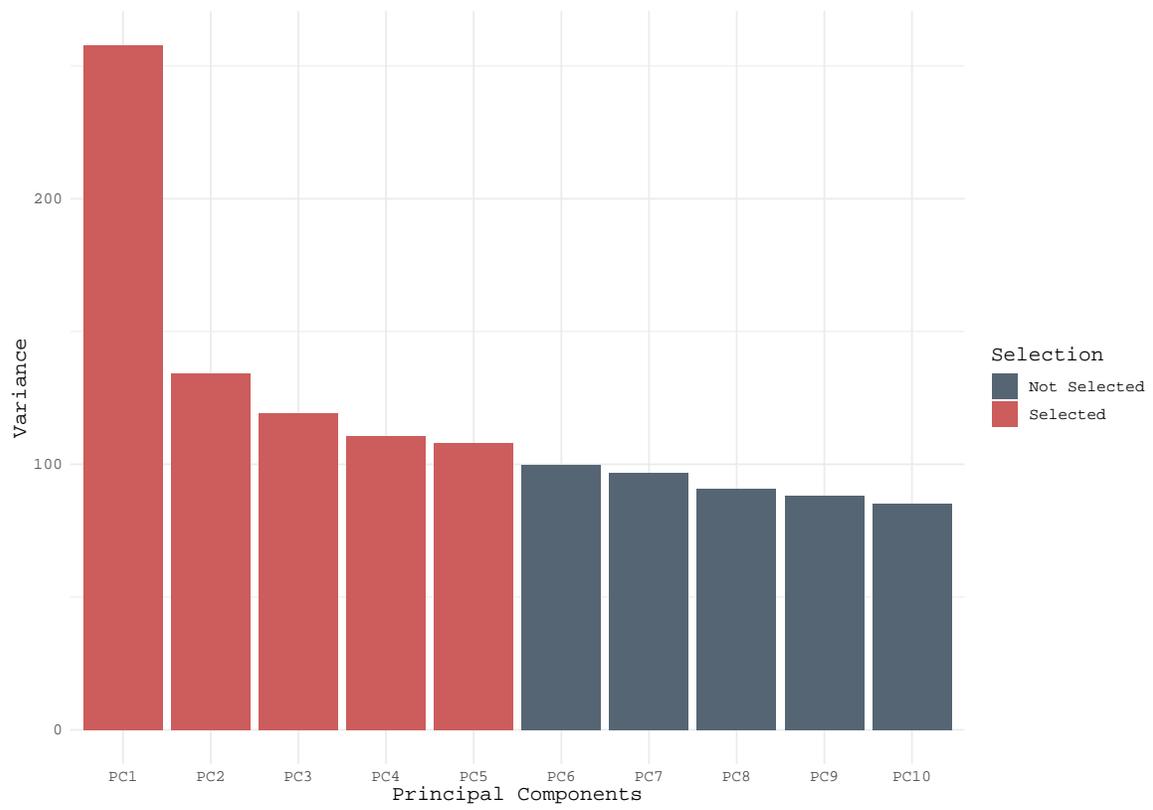


Figure 2.1: *The scree plot of the principal component analysis*

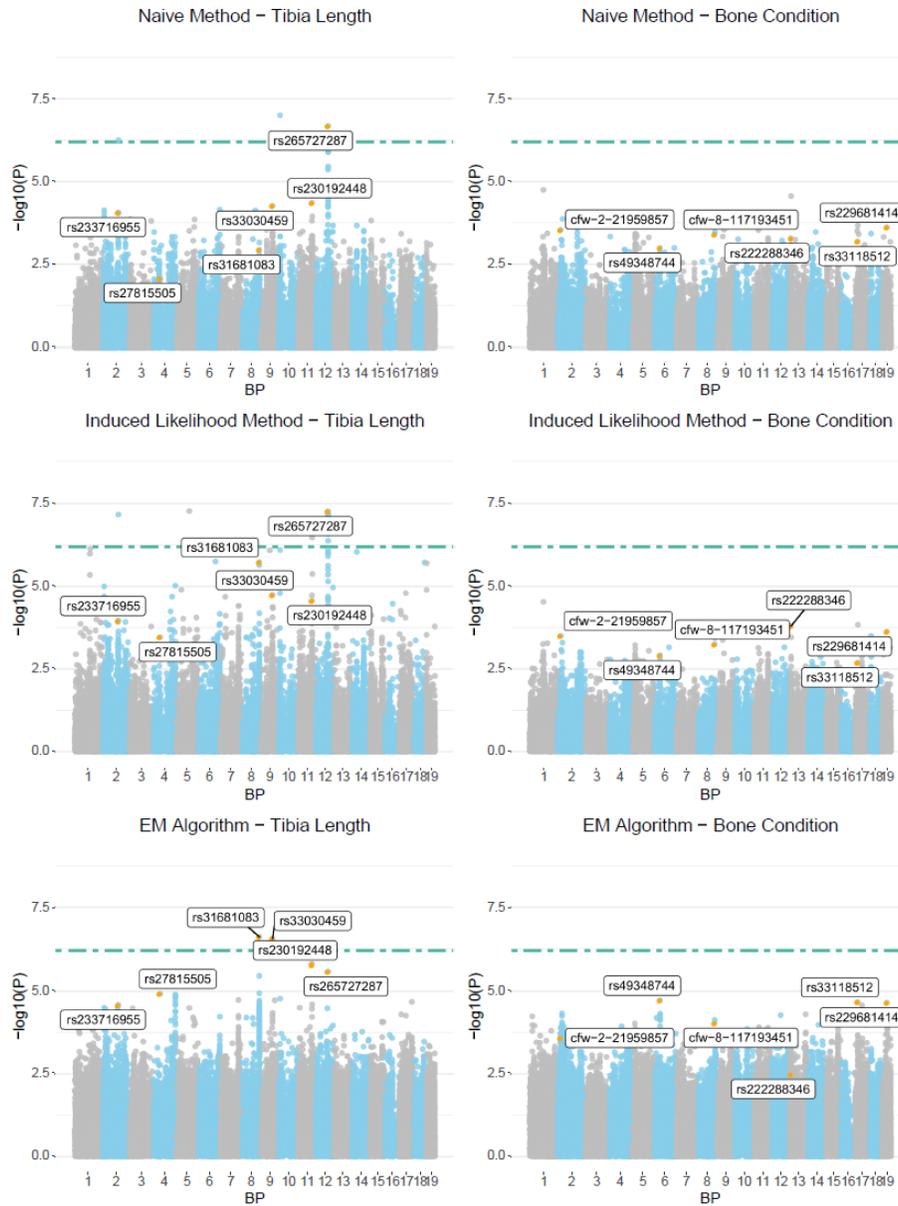


Figure 2.2: The Manhattan plots of the genome-wide association studies for the three methods and two responses. The x-axis shows the base-pair position (BP, the location of a SNP) on genome which is divided as 19 chromosomes labeled from 1 to 19. The y-axis is the $-\log_{10}$ scale of the p-value. Horizontal green dash lines mark the significant level 10^{-6} . The orange dots show the SNPs discussed in the text and their labels are marked in the small boxes.

Chapter 3

Estimating Equation Approach with Bivariate Mixed Responses Subject to Measurement Error and Misclassification

In contrast to likelihood-based approaches developed in Chapter 2, in this chapter, we confine our attention to the marginal modeling, where we explore estimation equation approaches to handle measurement error and misclassification in responses. In Section 3.1, we present the basic notation and the model setup. In Section 3.2, we first introduce the measurement error model and the misclassification model, and then we develop an insertion strategy for estimation of the model parameters to account for the effects of measurement error and misclassification in responses. In Section 3.3, we extend the method to the scenario where either external or internal validation data are available. Simulations studies are conducted in Section 3.4 to evaluate the performance of the proposed methods. In Section 3.5 we apply the proposed method to analyze the mice data arising from a genome-wide association study.

3.1 Model Setup and Framework

3.1.1 Response Model

We consider the case with bivariate responses for which one component is continuous and one component is binary. For $i = 1, \dots, n$, let $Y_i = (Y_{i1}, Y_{i2})^T$, where Y_{i1} denotes the continuous response, and Y_{i2} represents the binary response, and n is the number of subjects. Let $X_i = (X_{i1}, \dots, X_{ip})^T$ denote the covariate vector for subject i , where p is a positive integer. For $i = 1, \dots, n$ and $j = 1, 2$, let $\mu_{ij} = E(Y_{ij}|X_i)$ be the conditional mean of the Y_{ij} , given X_i , and let $v_{ij} = \text{Var}(Y_{ij}|X_i)$ be the conditional variance of Y_{ij} given covariates X_i .

We assume Y_i and $Y_{i'}$ are independent for any $i \neq i'$, but Y_{i1} and Y_{i2} could be correlated. A bivariate generalized linear model is employed to characterize the dependence of μ_{ij} on X_i for $j = 1, 2$:

$$\begin{aligned} g_1(\mu_{i1}) &= \beta_1^T X_i; \\ g_2(\mu_{i2}) &= \beta_2^T X_i, \end{aligned} \quad (3.1)$$

where $\beta = (\beta_1^T, \beta_2^T)^T$ is the vector of regression parameters, and $g_1(\cdot)$ and $g_2(\cdot)$ are link functions. For example, one may specify $g_1(t) = t$ and $g_2(t) = \log\{t/(1-t)\}$.

We assume that for $j = 1, 2$,

$$v_{ij} = h(\mu_{ij}; \psi_j), \quad (3.2)$$

where ψ_j is the dispersion parameter and $h(\cdot)$ is a specified function characterizing the relationship between the conditional variance v_{ij} and the conditional mean μ_{ij} of Y_{ij} given X_i . For instance, the variance functions of the continuous and binary response are often specified, respectively, as

$$v_{i1}(\mu_{i1}) = \psi_1,$$

and

$$v_{i2}(\mu_{i2}) = \mu_{i2}(1 - \mu_{i2}),$$

where ψ_1 is often further reparameterized as σ^2 because of its non-negative property.

3.1.2 Estimating Equation Method

Let $V_{i1} = \text{Var}(Y_i|X_i)$ be the conditional covariance matrix of the response vector Y_i , given X_i . The covariance matrix V_{i1} is decomposed as

$$V_{i1} = B_i^{\frac{1}{2}} C_i B_i^{\frac{1}{2}}, \quad (3.3)$$

where $B_i = \text{diag}\{v_{ij} : j = 1, 2\}$ and C_i is the correlation matrix $\begin{pmatrix} 1 & \lambda \\ \lambda & 1 \end{pmatrix}$ of the response vector Y_i , given X_i , with the parameter λ bounded in $[-1, 1]$. Let $\phi = (\psi_1, \lambda)^\top$, and $\theta = (\beta^\top, \phi^\top)^\top$.

For $i = 1, \dots, n$, let

$$U_{i1}(\theta) = D_{1i}^\top V_{i1}^{-1}(Y_i - \mu_i), \quad (3.4)$$

where $D_{i1} = \frac{\partial \mu_i}{\partial \beta^\top}$ is a $2 \times 2p$ matrix. Then $U_{i1}(\theta)$ is an unbiased estimating function which can be used to estimate β if the parameter ϕ were known.

To estimate ϕ , we construct a second set of estimating functions. For $i = 1, \dots, n$ and $j, k = 1, 2$, let v_{ijk} denote the (j, k) th element of V_{i1} . Define $\xi_i = (v_{ijk} : 1 \leq j \leq k \leq 2)^\top$, and $S_i = \{(Y_{ij} - \mu_{ij})(Y_{ik} - \mu_{ik}) : 1 \leq j \leq k \leq 2\}^\top$. Let

$$U_{i2}(\theta) = D_{2i}^\top V_{i2}^{-1}(S_i - \xi_i), \quad (3.5)$$

where $D_{i2} = \frac{\partial \xi_i}{\partial \phi^\top}$, and V_{i2} is a 3×3 weight matrix. Then $U_{i2}(\theta)$ is an unbiased estimating function of ϕ for any given β . This estimating function is the most efficient in the class of all estimating functions of form (3.5) if the weight matrix V_{i2} is set as the covariance matrix of S_i . However, such a specification requires the modeling of the third and fourth moments of Y_{ij} , which is often difficult or of no interest. In practice, V_{i2} is often specified as a diagonal matrix (e.g., [Hall, 2001](#); [Yi and Cook, 2002](#)). Although such a specification may incur some efficiency loss, it allows us to keep the model assumptions minimal, thus protecting us against model misspecification.

Let $U_i(\theta) = (U_{i1}^\top(\theta), U_{i2}^\top(\theta))^\top$. By the estimating function theory (e.g., [Liang and Zeger, 1986](#); [Godambe, 1991](#); [Newey and McFadden, 1994](#); [Yi, 2017](#), Section 1.3.2), under regularity conditions, solving

$$\sum_{i=1}^n U_i(\theta) = 0$$

for θ gives a consistent estimator, say, $\tilde{\theta}$, of θ , and $\sqrt{n}(\tilde{\theta} - \theta)$ has an asymptotic normal distribution with mean zero and covariance matrix

$$\left\{ E \left(\frac{\partial U_i(\theta)}{\partial \theta^\top} \right) \right\}^{-1} E\{U_i(\theta)U_i^\top(\theta)\} \left\{ E \left(\frac{\partial U_i(\theta)}{\partial \theta^\top} \right) \right\}^{-1\top}.$$

3.2 Methodology

3.2.1 Measurement Error and Misclassification Models

Suppose that for $i = 1, \dots, n$, the response variables Y_{i1} and Y_{i2} are subject to mismeasurement and their precise measurements are not observed for every subject, but instead,

surrogate measurements Y_{i1}^* and Y_{i2}^* are observed, respectively, for Y_{i1} and Y_{i2} .

To describe the mismeasurement processes, we consider the factorization

$$f(y_{i1}^*, y_{i2}^* | y_{i1}, y_{i2}, x_i) = f(y_{i1}^* | y_{i2}^*, y_{i1}, y_{i2}, x_i) f(y_{i2}^* | y_{i1}, y_{i2}, x_i), \quad (3.6)$$

for which we assume that

$$f(y_{i1}^* | y_{i2}^*, y_{i1}, y_{i2}, x_i) = f(y_{i1}^* | y_{i1}, y_{i2}, x_i)$$

and

$$f(y_{i2}^* | y_{i1}, y_{i2}, x_i) = f(y_{i2}^* | y_{i2}, x_i). \quad (3.7)$$

Let $Z_i = (Z_{i1}, \dots, Z_{ip_z})^T$ be the covariates involved in the misclassification. For ease of exposition, we assume that Z_i is a subset of X_i ; if this is not the case, we can modify our initial definition of X_i to include Z_i as its part. Let $\pi_{i0} = P(Y_{i2}^* = 1 | Y_{i2} = 0, Z_i)$ and $\pi_{i1} = P(Y_{i2}^* = 0 | Y_{i2} = 1, Z_i)$ be the misclassification probabilities that may depend on the covariates. We consider logistic models,

$$\text{logit } \pi_{i1} = \alpha_{01} + \alpha_{z1}^T Z_i$$

and

$$\text{logit } \pi_{i0} = \alpha_{00} + \alpha_{z0}^T Z_i, \quad (3.8)$$

where $\alpha = (\alpha_{01}, \alpha_{z1}^T, \alpha_{00}, \alpha_{z0}^T)^T$ is the vector of the regression parameters.

For the continuous response Y_{i1} , we consider a regression model which facilitates possible dependence of Y_{i1}^* on $\{Y_{i1}, Y_{i2}, Z_i\}$

$$Y_{i1}^* = \gamma_0 + \gamma_1 Y_{i1} + \gamma_2 Y_{i2} + \gamma_3^T Z_i + e_i, \quad (3.9)$$

where e_i is the random error which is independent of $\{Y_{i1}, Y_{i2}, Z_i\}$ and has zero mean and constant variance σ_e^2 , $\gamma = (\gamma_0, \gamma_1, \gamma_2, \gamma_3^T)^T$ is the vector of regression coefficients. Often, a normal distribution is assumed for the e_i .

Let $\eta = (\gamma^T, \alpha^T)^T$ denote the vector of parameters associated with the models (3.8) and (3.9).

3.2.2 Estimating Equation Method in the Presence of Mismeasurement

Without addressing measurement error and misclassification in the response, simply replacing Y_{ij} with Y_{ij}^* in the estimating functions (3.4) and (3.5) results in estimating functions that are no longer unbiased, and the resultant estimators may be inconsistent. To account

for the mismeasurement effects, we develop a two-step procedure to construct new estimating functions, say $U_{i1}^{**}(\theta)$ and $U_{i2}^{**}(\theta)$, which are expressed in terms of the observed measurements Y_{i1}^* and Y_{i2}^* together with the covariates and the model parameters and satisfy

$$E\{U_{i1}^{**}(\theta)\} = 0, \text{ and } E\{U_{i2}^{**}(\theta)\} = 0.$$

To this end, we develop a two-step procedure to correct for the effects of misclassification in Y_{i2} and those of measurement error in Y_{i1} sequentially. In Step 1, we define

$$Y_{i2}^{**} = \frac{Y_{i2}^* - \pi_{i0}}{1 - \pi_{i0} - \pi_{i1}},$$

where π_{i0} and π_{i1} are the misclassification rates postulated by (3.8). It is readily seen that $E(Y_{i2}^{**}|Y_{i2}, X_{i1}) = Y_{i2}$.

Then we modify (3.4) and (3.5) by replacing Y_{i2} and Y_{i2}^{**} and define

$$U_{i1}^*(\theta) = D_{1i}^T V_{i1}^{-1} \begin{pmatrix} Y_{i1} - \mu_{i1} \\ Y_{i2}^{**} - \mu_{i2} \end{pmatrix}, \text{ and } U_{i2}^*(\theta) = D_{2i}^T V_{i2}^{-1} \begin{pmatrix} Y_{i1}^2 - 2\mu_{i1}Y_{i1} - \mu_{i1}^2 - \xi_{i1} \\ Y_{i1}Y_{i2}^{**} - \mu_{i1}Y_{i2}^{**} - \mu_{i2}Y_{i1} - \xi_{i2} \\ Y_{i2}^{**} - 2\mu_{i2}Y_{i2}^{**} + \mu_{i2}^2 - \xi_{i3} \end{pmatrix}, \quad (3.10)$$

for which we use $Y_{i2}^2 = Y_{i2}$ for a binary variable Y_{i2} taking the value of either 0 or 1.

In Step 2, we further modify (3.10) by replacing Y_{i1} with the observed variables in order to obtain $U_{i1}^{**}(\theta)$ and $U_{i2}^{**}(\theta)$. To this end, define

$$Y_{i1}^{**} = \frac{Y_{i1}^* - \gamma_0 - \gamma_2 Y_{i2}^{**} - \gamma_3^T Z_i}{\gamma_1},$$

$$Y_{i11}^{**} = Y_{i1}^{**2} - \frac{\sigma_e^2}{\gamma_1^2} - \frac{\gamma_2^2}{\gamma_1^2} \Delta_i,$$

and

$$Y_{i12}^{**} = Y_{i1}^{**} Y_{i2}^{**} + \frac{\gamma_2}{\gamma_1} \Delta_i,$$

where

$$\Delta_i = \frac{\Delta_{i0}^{1-Y_{i2}^*} \Delta_{i1}^{Y_{i2}^*} - \Delta_{i0} \pi_{i1} - \Delta_{i1} \pi_{i0}}{1 - \pi_{i1} - \pi_{i0}},$$

$$\Delta_{i0} = \frac{\pi_{i0} - \pi_{i0}^2}{(1 - \pi_{i1} - \pi_{i0})^2},$$

and

$$\Delta_{i1} = \frac{\pi_{i1} - \pi_{i1}^2}{(1 - \pi_{i1} - \pi_{i0})^2}.$$

Let $U_i^{**}(\theta)$ be $U_i^*(\theta) = (U_{i1}^{*\text{T}}(\theta), U_{i2}^{*\text{T}}(\theta))^{\text{T}}$ with $Y_{i1}, Y_{i1}^2, Y_{i1}Y_{i2}^{**}$ replaced by $Y_{i1}^{**}, Y_{i11}^{**}, Y_{i12}^{**}$, respectively. In Appendix B.1, we show that $E(Y_{i1}^{**}|Y_{i1}, Y_{i2}) = Y_{i1}$, $E(Y_{i11}^{**}|Y_{i1}, Y_{i2}, X_i) = Y_{i1}^2$, and $E(Y_{i12}^{**}|Y_{i1}, Y_{i2}, X_i) = Y_{i1}Y_{i2}$, thus yielding

$$E[U_i^{**}(\theta)|Y_{i1}, Y_{i2}, X_i] = U_i(\theta).$$

The unbiasedness of $U_i^{**}(\theta)$ is immediate from that of $U_i(\theta)$, thus $U_i^{**}(\theta)$ may be used to obtain a consistent estimator of θ because it is expressed in terms of the observed data. To do so, we note that however, parameter η for the misclassification and measurement error models are involved in $U_i^{**}(\theta)$. To explicitly spell out the dependence on η , we write $U_i^{**}(\theta)$ as $U_i^{**}(\theta, \eta)$. If η is known, say taking a value η_0 , then by the estimating function theory, under regularity conditions (e.g., Godambe, 1991; Newey and McFadden, 1994; Yi, 2017, Section 1.3.2), solving

$$\sum_{i=1}^n U_i^{**}(\theta, \eta_0) = 0 \quad (3.11)$$

gives a consistent estimator, say $\hat{\theta}$, of $\theta = (\beta^{\text{T}}, \phi^{\text{T}})^{\text{T}}$, and that $\sqrt{n}(\hat{\theta} - \theta)$ has an asymptotic normal distribution with mean zero and covariance matrix

$$\left\{ E \left(\frac{\partial U_i^{**}(\theta, \eta_0)}{\partial \theta^{\text{T}}} \right) \right\}^{-1} E \{ U_i^{**}(\theta, \eta_0) U_i^{**\text{T}}(\theta, \eta_0) \} \left\{ E \left(\frac{\partial U_i^{**}(\theta, \eta_0)}{\partial \theta^{\text{T}}} \right) \right\}^{-1\text{T}}.$$

3.3 Estimation Methods with Validation Data

In many applications, the parameter η for the measurement error and misclassification models is usually unknown, and is estimated from additional validation data. We now consider two types of validation studies, *internal validation* and *external validation*. Let \mathcal{M} denote the index set of the subjects in the main study, where $\{(y_{i1}^*, y_{i2}^*, x_i) : i \in \mathcal{M}\}$ is available. Let \mathcal{V} represent the index set of the subjects in the validation data. For internal validation, the validation data contain $\{(y_{i1}^*, y_{i2}^*, y_{i1}, y_{i2}, x_i) : i \in \mathcal{V}\}$ with $\mathcal{V} \subset \mathcal{M}$; for external validation, the validation data contain $\{(y_{i1}^*, y_{i2}^*, y_{i1}, y_{i2}, z_i) : i \in \mathcal{V}\}$ with $\mathcal{M} \cap \mathcal{V} = \emptyset$. Let m denote the size of the validation subsample \mathcal{V} .

3.3.1 External Validation

Estimation of η can be carried out by maximizing the conditional likelihood function

$$L(\eta) = \prod_{i=1}^n L_i(y_{i1}^*, y_{i2}^* | y_{i1}, y_{i2}, x_i; \eta),$$

with respect to η , where $L_i(y_{i1}^*, y_{i2}^* | y_{i1}, y_{i2}, x_i; \eta)$ is the likelihood function contributed from the i th individual and is determined by (3.8) and (3.9).

Let

$$S_i(\eta) = \partial \overline{\log} L_i(y_{i1}^*, y_{i2}^* | y_{i1}, y_{i2}, x_i; \eta) / \partial \eta \quad (3.12)$$

denote the score function of parameter η .

With external validation data, we consider estimation function

$$U^{(E)}(\theta, \eta) = \sum_{i \in \mathcal{M}} \begin{pmatrix} U_{i1}^{**}(\theta, \eta) \\ U_{i2}^{**}(\theta, \eta) \\ 0 \end{pmatrix} + \sum_{i \in \mathcal{V}} \begin{pmatrix} 0 \\ 0 \\ S_i(\eta) \end{pmatrix}, \quad (3.13)$$

where $S_i(\eta)$ is the score function determined by (3.12). Then solving

$$U^{(E)}(\theta, \eta) = 0$$

gives an estimator of $(\theta^T, \eta^T)^T$, denoted as $(\hat{\theta}_E^T, \hat{\eta}_E^T)^T$.

Assume that regularity conditions hold and that the ratio m/n approaches a positive constant ρ as $n \rightarrow \infty$. In Appendix B.2, we show that $(\hat{\theta}_E^T, \hat{\eta}_E^T)^T$ is a consistent estimator of $(\theta^T, \eta^T)^T$, and $\sqrt{n} \left\{ (\hat{\theta}_E^T, \hat{\eta}_E^T)^T - (\theta^T, \eta^T)^T \right\}$ has an asymptotic normal distribution with mean zero and covariance matrix $\frac{1}{1+\rho} \Gamma_E^{-1} \Sigma_E (\Gamma_E^{-1})^T$, where

$$\begin{aligned} \Gamma_E &= -\frac{1}{1+\rho} \begin{bmatrix} E \left(\frac{\partial U_{i1}^{**}}{\partial \theta^T} \right) & E \left(\frac{\partial U_{i1}^{**}}{\partial \eta^T} \right) \\ E \left(\frac{\partial U_{i2}^{**}}{\partial \theta^T} \right) & E \left(\frac{\partial U_{i2}^{**}}{\partial \eta^T} \right) \\ 0 & 0 \end{bmatrix} - \frac{\rho}{1+\rho} \begin{bmatrix} 0 & 0 \\ 0 & 0 \\ 0 & E \left(\frac{\partial S_i}{\partial \eta^T} \right) \end{bmatrix}; \\ \Sigma_E &= \frac{1}{1+\rho} \begin{bmatrix} E \left(U_{i1}^{**} U_{i1}^{**T} \right) & E \left(U_{i1}^{**} U_{i2}^{**T} \right) & 0 \\ E \left(U_{i2}^{**} U_{i1}^{**T} \right) & E \left(U_{i2}^{**} U_{i2}^{**T} \right) & 0 \\ 0 & 0 & 0 \end{bmatrix} + \frac{\rho}{1+\rho} \begin{bmatrix} 0 & 0 & 0 \\ 0 & 0 & 0 \\ 0 & 0 & E \left(S_i S_i^T \right) \end{bmatrix}. \end{aligned} \quad (3.14)$$

3.3.2 Internal Validation

To account for the effects of measurement error and misclassification in responses, we construct the estimating functions

$$U^{(I)}(\theta, \eta) = \sum_{i \in \mathcal{M} \setminus \mathcal{V}} \begin{pmatrix} U_{i1}^{**}(\theta, \eta) \\ U_{i2}^{**}(\theta, \eta) \\ 0 \end{pmatrix} + \sum_{i \in \mathcal{V}} \begin{pmatrix} U_{i1}(\theta, \eta) \\ U_{i2}(\theta, \eta) \\ S_i(\eta) \end{pmatrix}, \quad (3.15)$$

where $U_{i1}(\theta, \eta)$ and $U_{i2}(\theta, \eta)$ are the estimating functions under the true model as (3.4) and (3.5), and $S_i(\eta)$ is the score function determined by (3.12). Here and elsewhere, 0 may represent the real number zero, a zero vector, or a zero matrix whose meaning is clear in each context. One can obtain an estimator, $(\hat{\theta}_I^T, \hat{\eta}_I^T)^T$ for $(\theta^T, \eta^T)^T$, by solving equation

$$U^{(I)}(\theta, \eta) = 0 \quad (3.16)$$

with respect to θ and η .

Since $S_i(\eta)$ does not depend on θ , solving (3.16) is equivalent to a two-step procedure. First obtain $\hat{\eta}_I$ by solving $\sum_{i \in \mathcal{V}} S_i(\eta) = 0$. Then solve the equation

$$U^{(I)}(\theta, \eta) = \sum_{i \in \mathcal{M} \setminus \mathcal{V}} \begin{pmatrix} U_{i1}^{**}(\theta, \hat{\eta}_I) \\ U_{i2}^{**}(\theta, \hat{\eta}_I) \end{pmatrix} + \sum_{i \in \mathcal{V}} \begin{pmatrix} U_{i1}(\theta, \hat{\eta}_I) \\ U_{i2}(\theta, \hat{\eta}_I) \end{pmatrix} = 0,$$

to obtain an estimator of θ , denoted as $\hat{\theta}_I = (\hat{\beta}_I^T, \hat{\phi}_I^T)^T$.

Assume that regularity conditions hold and that the ratio m/n approaches a positive constant ρ as $n \rightarrow \infty$. In Appendix B.3, we show that $(\hat{\theta}_I^T, \hat{\eta}_I^T)^T$ is a consistent estimator of $(\theta^T, \eta^T)^T$, and $\sqrt{n} \left\{ (\hat{\theta}_I^T, \hat{\eta}_I^T)^T - (\theta^T, \eta^T)^T \right\}$ has an asymptotic normal distribution with mean zero and covariance matrix $\Gamma_I^{-1} \Sigma_I (\Gamma_I^{-1})^T$, where

$$\begin{aligned} \Gamma_I &= - (1 - \rho) \begin{bmatrix} E \left(\frac{\partial U_{i1}^{**}}{\partial \theta^T} \right) & E \left(\frac{\partial U_{i1}^{**}}{\partial \eta^T} \right) \\ E \left(\frac{\partial U_{i2}^{**}}{\partial \theta^T} \right) & E \left(\frac{\partial U_{i2}^{**}}{\partial \eta^T} \right) \\ 0 & 0 \end{bmatrix} - \rho \begin{bmatrix} E \left(\frac{\partial U_{i1}}{\partial \theta^T} \right) & 0 \\ E \left(\frac{\partial U_{i2}}{\partial \theta^T} \right) & 0 \\ 0 & E \left(\frac{\partial S_i}{\partial \eta^T} \right) \end{bmatrix}; \\ \Sigma_I &= (1 - \rho) \begin{bmatrix} E (U_{i1}^{**} U_{i1}^{**T}) & E (U_{i1}^{**} U_{i2}^{**T}) & 0 \\ E (U_{i2}^{**} U_{i1}^{**T}) & E (U_{i2}^{**} U_{i2}^{**T}) & 0 \\ 0 & 0 & 0 \end{bmatrix} \\ &+ \rho \begin{bmatrix} E (U_{i1} U_{i1}^T) & E (U_{i1} U_{i2}^T) & E (U_{i1} S_i^T) \\ E (U_{i2} U_{i1}^T) & E (U_{i2} U_{i2}^T) & E (U_{i2} S_i^T) \\ E (S_i U_{i1}^T) & E (S_i U_{i2}^T) & E (S_i S_i^T) \end{bmatrix}. \end{aligned} \quad (3.17)$$

3.3.3 Weighted Estimator with Internal Validation Data

Estimation of $(\theta^T, \eta^T)^T$ based on (3.15) basically treats the validation data and non-validation data equally. To improve the efficiency of parameter estimation, we may attach suitable weights to adjust contributions from the validation sample and the non-validation sample.

Let $W = \text{diag}(w_1, \dots, w_q)$ be a diagonal matrix, where $0 \leq w_j \leq 1$ for $j = 1, \dots, p_\theta$ and $w_j = 0$ for $j = (p_\theta + 1), \dots, q$. Here $q = p_\theta + p_\eta$, and p_θ and p_η represent, respectively, the dimension of θ and η . We modify the estimating function (3.15) as

$$U^{(W)}(\theta, \eta) = \sum_{i \in \mathcal{M} \setminus \mathcal{V}} W \begin{pmatrix} U_{i1}^{**}(\theta, \eta) \\ U_{i2}^{**}(\theta, \eta) \\ 0 \end{pmatrix} + \sum_{i \in \mathcal{V}} (I_q - W) \begin{pmatrix} U_{i1}(\theta, \eta) \\ U_{i2}(\theta, \eta) \\ S_i(\eta) \end{pmatrix},$$

where $U_{i1}(\theta, \eta)$, $U_{i2}(\theta, \eta)$, and $S_i(\eta)$ are defined in the same way as in (3.15), and I_q is the $q \times q$ identity matrix. An estimator of $(\theta^T, \eta^T)^T$, denoted $(\hat{\theta}_w^T, \hat{\eta}_w^T)^T$, is obtained by solving the equation

$$U^{(W)}(\theta, \eta) = 0$$

for θ and η .

Assume regularity conditions hold and that the ratio m/n approaches a positive constant ρ as $n \rightarrow \infty$. Similar to the estimator obtained from (3.15), we can show that $(\hat{\theta}_w^T, \hat{\eta}_w^T)^T$ is a consistent estimator of $(\theta^T, \eta^T)^T$, and $\sqrt{n} \left\{ (\hat{\theta}_w^T, \hat{\eta}_w^T)^T - (\theta^T, \eta^T)^T \right\}$ has an asymptotic normal distribution with mean zero and covariance matrix $\Gamma_W^{-1} \Sigma_W \Gamma_W^{-1T}$, where

$$\begin{aligned} \Gamma_W &= - (1 - \rho) W \begin{bmatrix} E \left(\frac{\partial U_{i1}^{**}}{\partial \theta^T} \right) & E \left(\frac{\partial U_{i1}^{**}}{\partial \eta^T} \right) \\ E \left(\frac{\partial U_{i2}^{**}}{\partial \theta^T} \right) & E \left(\frac{\partial U_{i2}^{**}}{\partial \eta^T} \right) \\ 0 & 0 \end{bmatrix} - \rho (I - W) \begin{bmatrix} E \left(\frac{\partial U_{i1}}{\partial \theta^T} \right) & 0 \\ E \left(\frac{\partial U_{i2}}{\partial \theta^T} \right) & 0 \\ 0 & E \left(\frac{\partial S_i}{\partial \eta^T} \right) \end{bmatrix}; \\ \Sigma_W &= (1 - \rho) W \begin{bmatrix} E \left(U_{i1}^{**} U_{i1}^{**T} \right) & E \left(U_{i1}^{**} U_{i2}^{**T} \right) & 0 \\ E \left(U_{i2}^{**} U_{i1}^{**T} \right) & E \left(U_{i2}^{**} U_{i2}^{**T} \right) & 0 \\ 0 & 0 & 0 \end{bmatrix} W \\ &+ \rho (I - W) \begin{bmatrix} E \left(U_{i1} U_{i1}^T \right) & E \left(U_{i1} U_{i2}^T \right) & E \left(U_{i1} S_i^T \right) \\ E \left(U_{i2} U_{i1}^T \right) & E \left(U_{i2} U_{i2}^T \right) & E \left(U_{i2} S_i^T \right) \\ E \left(S_i U_{i1}^T \right) & E \left(S_i U_{i2}^T \right) & E \left(S_i S_i^T \right) \end{bmatrix} (I - W). \end{aligned}$$

The optimal weights can be obtained by minimizing $\text{Tr} \left(\Gamma_W^{-1} \Sigma_W \Gamma_W^{-1T} \right)$ with respect to $\{w_1, \dots, w_\theta\}$ with the constraints $w_j = 0$ for $j = (p_\theta + 1), \dots, q$, where $\text{Tr}(A)$ is the trace of matrix A . Although the idea is straightforward, this optimization is computationally difficult. Alternatively, we develop an optimal weighted estimator based on linear combinations of two simple estimators discussed as follows.

The first estimator of $(\theta^T, \eta^T)^T$ is obtained using the validation data only. Let $(\hat{\theta}_I^{(0)T}, \hat{\eta}_I^{(0)T})^T$

be the resulting estimator by solving the equation

$$\sum_{i \in \mathcal{V}} \begin{pmatrix} U_{i1}(\theta, \eta) \\ U_{i2}(\theta, \eta) \\ S_i(\eta) \end{pmatrix} = 0 \quad (3.18)$$

for θ and η .

The second estimator of θ , denoted as $\widehat{\theta}_I^{(1)}$, solves the estimating equation constructed from the non-validation data,

$$\sum_{i \in \mathcal{M} \setminus \mathcal{V}} \begin{pmatrix} U_{i1}^{**}(\theta, \widehat{\eta}_I^{(0)}) \\ U_{i2}^{**}(\theta, \widehat{\eta}_I^{(0)}) \end{pmatrix} = 0 \quad (3.19)$$

for θ , where $U_{ij}^{**}(\theta, \widehat{\eta}_I^{(0)})$ is determined by $U_{ij}^{**}(\theta, \eta)$ in (3.11) with η replaced by $\widehat{\eta}_I^{(0)}$.

Under regularity conditions, both $\widehat{\theta}_I^{(0)}$ and $\widehat{\theta}_I^{(1)}$ are consistent estimators for θ . We consider a weighted estimator to be a linear combination of $\widehat{\theta}_I^{(0)}$ and $\widehat{\theta}_I^{(1)}$:

$$\widetilde{\theta}_I(\Omega) = \Omega \widehat{\theta}_I^{(1)} + (I_{p_\theta} - \Omega) \widehat{\theta}_I^{(0)}, \quad (3.20)$$

where $\Omega = \text{diag}(\omega_1, \dots, \omega_{p_\theta})$ is a diagonal matrix with constants $0 \leq \omega_j \leq 1$ for $j = 1, \dots, p_\theta$.

To find the optimal weights, we target to minimize the asymptotic variance for each element of $\widetilde{\theta}_I(\Omega)$. For $r = 1, \dots, p_\theta$, let $\widetilde{\theta}_{Ir}(\Omega)$, $\widehat{\theta}_{Ir}^{(0)}$ and $\widehat{\theta}_{Ir}^{(1)}$ be the r th component of $\widetilde{\theta}_I(\Omega)$, $\widehat{\theta}_I^{(0)}$ and $\widehat{\theta}_I^{(1)}$, respectively. The variance of $\widetilde{\theta}_{Ir}(\Omega)$ is given by

$$\begin{aligned} \text{Var}(\widetilde{\theta}_{Ir}(\Omega)) &= \omega_r^2 \left(\text{Var}(\widehat{\theta}_{Ir}^{(0)}) + \text{Var}(\widehat{\theta}_{Ir}^{(1)}) - 2\text{Cov}(\widehat{\theta}_{Ir}^{(0)}, \widehat{\theta}_{Ir}^{(1)}) \right) \\ &\quad - 2\omega_r \left(\text{Var}(\widehat{\theta}_{Ir}^{(0)}) - \text{Cov}(\widehat{\theta}_{Ir}^{(0)}, \widehat{\theta}_{Ir}^{(1)}) \right) + \text{Var}(\widehat{\theta}_{Ir}^{(1)}), \end{aligned}$$

which is minimized at

$$\omega_r^* = \frac{\text{Var}(\widehat{\theta}_{Ir}^{(0)}) - \text{Cov}(\widehat{\theta}_{Ir}^{(0)}, \widehat{\theta}_{Ir}^{(1)})}{\text{Var}(\widehat{\theta}_{Ir}^{(0)}) + \text{Var}(\widehat{\theta}_{Ir}^{(1)}) - 2\text{Cov}(\widehat{\theta}_{Ir}^{(0)}, \widehat{\theta}_{Ir}^{(1)})}.$$

Let $\Omega^* = \text{diag}(\omega_1^*, \dots, \omega_{p_\theta}^*)$. Then the estimator $\widetilde{\theta}_I^* = \Omega^* \widehat{\theta}_I^{(1)} + (I - \Omega^*) \widehat{\theta}_I^{(0)}$ is the optimal estimator among the linear combinations of form (3.20).

In practice, ω_r^* is estimated by

$$\widehat{\omega}_r^* = \frac{\widehat{\text{Var}}(\widehat{\theta}_{I_r}^{(1)}) - \widehat{\text{Cov}}(\widehat{\theta}_{I_r}^{(0)}, \widehat{\theta}_{I_r}^{(1)})}{\widehat{\text{Var}}(\widehat{\theta}_{I_r}^{(0)}) + \widehat{\text{Var}}(\widehat{\theta}_{I_r}^{(1)}) - 2\widehat{\text{Cov}}(\widehat{\theta}_{I_r}^{(0)}, \widehat{\theta}_{I_r}^{(1)})},$$

where $\widehat{\text{Var}}(\widehat{\theta}_{I_r}^{(1)})$, $\widehat{\text{Var}}(\widehat{\theta}_{I_r}^{(0)})$ and $\widehat{\text{Cov}}(\widehat{\theta}_{I_r}^{(0)}, \widehat{\theta}_{I_r}^{(1)})$ are estimates for $\text{Var}(\widehat{\theta}_{I_r}^{(1)})$, $\text{Var}(\widehat{\theta}_{I_r}^{(0)})$ and $\text{Cov}(\widehat{\theta}_{I_r}^{(0)}, \widehat{\theta}_{I_r}^{(1)})$ by stacking the estimating functions in (3.18) and (3.19). The details are presented in Appendix B.4.

We comment that in practice, the resulting weights $\widehat{\omega}_r^*$ may not satisfy the constraint that $0 \leq \widehat{\omega}_r^* \leq 1$. If $\widehat{\omega}_r^* < 0$, we set $\widehat{\omega}_r^*$ to be 0 and if $\widehat{\omega}_r^* > 1$, we specify $\widehat{\omega}_r^*$ to be 1.

3.4 Simulation Studies

We conduct simulation studies to evaluate the performance of the proposed methods in terms of parameter estimation and associated variance estimation. Similar to the simulation studies in Chapter 2, for the sake of comparison, we consider three naive methods, where either measurement error or misclassification, or both are ignored.

We consider the sample size $n = 1000$. The X_{i1} is independently generated from $U(-3, 4)$, and the X_{i2} is independently generated from $N(0, 1)$. The response vector $Y_i = (Y_{i1}, Y_{i2})^T$ is generated from the model

$$\begin{aligned} g_1(\mu_{i1}) &= \beta_{10} + \beta_{11}X_{i1} + \beta_{12}X_{i2}, \\ g_2(\mu_{i2}) &= \beta_{20} + \beta_{21}X_{i1} + \beta_{22}X_{i2}, \end{aligned}$$

where the coefficient vector $\theta = (\beta_{10}, \beta_{20}, \beta_{11}, \beta_{21}, \beta_{12}, \beta_{22})^T$ is set as $(0.7, 0.7, 1.5, -1.5, -1, 1)^T$, $g_1(t) = t$, and $g_2(t) = \log\left(\frac{t}{1-t}\right)$. That is, Y_{i1} is generated by $N(\mu_{i1}, \sigma^2)$ where σ^2 is set as 1, and Y_{i2} is independently generated from Bernoulli(μ_{i2}).

The surrogate measurement Y_{i1}^* is generated from the measurement error model, $Y_{i1}^* = Y_{i1} + \gamma Y_{i2} + e_i$, where e_i is a centered normal random error with variance σ_e^2 and is independent of $\{Y_{i1}, Y_{i2}\}$. For the misclassification of Y_{i2} , we generate the surrogate measurement Y_{i2}^* by misclassification models (3.8). The values of α , γ , σ_e^2 are specified in Section 3.4.1.

For each estimator, we report the finite sample biases (denoted as ‘‘bias’’), the standard error (denoted as ‘‘SEE’’), the model-based standard error (denoted as ‘‘SEM’’), or the coverage rate (denoted as ‘‘CR’’).

3.4.1 Simulation 1: Evaluation for the Case with Known Mismeasurement Parameters

In this subsection, we consider the case where the parameters of measurement error and misclassification models are known to the method in Section 3.2.2, taking the values as in the specifications of generating the random variables.

To study the performance of the methods, we consider three settings. In Setting 1, fix $\alpha = (-1.386, 0, -1.386, 0)^T$ and $\gamma = 0.8$, and compare the performance of the naive models and the proposed model under different degrees of measurement error, where σ_e are set as 0.1, 0.5, 0.7. In Setting 2, fix $\sigma_e = 0.1$ and $\gamma = 0.8$, and evaluate the performance using the data simulated with different misclassification rates, where $\alpha = (\alpha_{01}, \alpha_{x1}, \alpha_{00}, \alpha_{x0})^T$ is set as $(-4.595, 0, -4.595, 0)^T$, $(-2.197, 0, -2.197, 0)^T$, or $(-1.386, 0, -1.386, 0)^T$, yielding the misclassification rates π_0 and π_1 as 1%, 10% or 20%, respectively. In Setting 3, fix $\sigma_e = 0.1$ and $\alpha = (-1.386, 0, -1.386, 0)^T$, and evaluate the methods for different measurement error mechanisms which are independent of the binary outcome Y_{i2} ($\gamma = 0$), negatively associated with the binary outcome ($\gamma = -0.8$), or positively associated with the binary outcome ($\gamma = 0.8$).

The results are presented in Tables 3.1–3.3. Different naive methods may perform differently in both the point estimation and the variance estimation, but they all produce large biases in the point estimation and poor coverage rates. Conversely, the proposed method successfully corrects the biases due to the response mismeasurement, yielding reasonably small finite sample biases and coverage rates in good agreement with the nominal value 95%.

3.4.2 Simulation 2: Evaluation of the Case with Validation Data

In this simulation study, we compare the performance of the methods for three scenarios. In the first scenario, we consider the same case as in Section 3.4.1 where the mismeasurement parameter η is known. In the second and the third scenarios, we evaluate the performance for the methods described in Sections 3.3.1 and 3.3.2 where η is unavailable but estimated from either external validation data or internal validation data. We also display the results of the method using the true measurements Y_{i1} and Y_{i2} for comparisons.

We consider the same three settings as in Simulation 1. The results are reported in Tables 3.4–3.6. As expected, the method using true response measurements produces the best results with the smallest finite sample biases and model-based standard errors as well as the best coverage rates of 95% confidence intervals. On the other hand, the proposed

methods perform quite well for different scenarios. Finite sample biases are close to those produced from the method with the true response measurements; model-based standard errors agree fairly well with empirical standard errors and coverage rates of 95% confidence intervals are in good agreement with the nominal level 95%.

3.4.3 Simulation 3: Evaluation of the Proposed Method with Internal Validation Data with Different Sample Sizes and Different Weights

In this subsection we compare the estimator described in Section 3.3.2 and the weighted estimators described in Section 3.3.3. We also consider four different weights to compare the estimates of using validation data only ($\hat{\theta}_1^{(0)}$), using non-validation data only ($\hat{\theta}_1^{(1)}$), using equal weights for validation data and non-validation data ($\hat{\theta}_1$), and optimal weighted estimator ($\hat{\theta}_1^*$). Our assessment focuses on examining the impact on the performance of the proposed estimator of the sample size, the sample size ratio between the validation data and non-validation data, and the weight choice. We consider two scenarios. In Scenario 1, we fix the total sample size to be 1500 and let the sample size ratio vary as 2:1, 1:1, 1:2. In Scenario 2, we fix the ratio to be 1:2 and let the total sample size be 1500 and 3000.

The results for Scenario 1 are presented in Figures 3.1–3.2 and the results for Scenario 2 are presented in Table 3.7. It is clear that the estimator with optimal weights described in Section 3.3.3 and the estimator described in Section 3.3.2 perform the best among all the estimators in terms of both finite sample biases and standard errors. Moreover, the former estimator greatly outperforms the latter one. The efficiency gain of using the estimator with optimal weights over the estimator in Section 3.3.2 can be as large as 58%, shown by the estimate of β_{21} with $n = 1500$.

3.5 Application to Mice SNPs Data

To illustrate the usage of the proposed method, we analyze data arising from a genome-wide association study of outbred Carworth Farms White mice data (Parker et al., 2016b). This study provided measurements with complex traits, including behavioral, physiological, and gene expression traits.

For $i = 1, \dots, 1128$, let Y_{i1} be the weight of the tibialis anterior muscle (in *mg*), and let Y_{i2} be the binary outcome where “0” represents a healthy tibia bone and “1” stands

for abnormal tibia bone, which is defined as the 90% quantile of the bone-mineral density. Due to the concern of data quality, the true measurements of the responses Y_{i1} and Y_{i2} for 464 subjects are not available but their surrogate measurements Y_{i1}^* and Y_{i2}^* are available, where Y_{i1}^* is the predicted tibialis anterior muscle weights based on muscle from other body parts of the mice and Y_{i2}^* is the bone condition judged by subjective observations from technicians. Precise measurements of the responses Y_{i1} and Y_{i2} together with their surrogates Y_{i1}^* and Y_{i2}^* are available for the remaining 664 subjects, which are taken as the validation data. Covariates include a continuous variable measuring the SNPs *rs27338905* (X_{i1}), and the first two principal components of genetics data (X_{i2}) and (X_{i3}) for subject i which are described below in detail. Our main interest lies in studying the association of SNPs *rs27338905* with two physiological traits. We employ the model (3.1) with $g_1(t) = t$ and $g_2(t) = \log\{t/(1-t)\}$ to facilitate the dependence of the responses on the covariates.

To account for the effect of population stratification (Price et al., 2006), similar to Section 2.6 in Chapter 2, we conduct principal component analysis. According to the scree plot in figure 3.3 and based on the “elbow” criterion, we include the first two principal components (X_{i2}) and (X_{i3}) as fixed effects in the response model.

We consider two settings for the misclassification and measurement error models. In Setting 1, we consider the body weight (X_{i4}) to be the covariates in model (3.8) to feature the misclassification of Y_{i2}^* . For the measurement error model, we consider (3.9) with the covariates chosen to be the body weight (X_{i4}). In Setting 2, we consider model (3.8) to be postulated by constants α_{00} and α_{01} , and an additional constraint that $\gamma_2 = 0$ is imposed for the measurement error model (3.9).

We analyze the data using the proposed optimal weighted estimator with internal validation data, described in Section 3.3.3. We compare the results with the naive model where the mismeasurement is ignored. The results are presented in Table 3.8. The proposed method with two settings produces similar estimation results. Under the significance level of 0.05, the estimates of β_{11} and β_{21} suggest that SNP *rs27338905* is significantly associated with the weight of tibialis anterior muscle but is not associated with the bone condition. The estimates of β_{12} , β_{13} , β_{22} and β_{23} show that the effects of population stratification are not significant. However, the naive method produces somewhat different findings; there is no evidence showing the effects of SNP *rs27338905* on the weight of the tibialis anterior muscle.

Regarding the results for the parameters of the mismeasurement models, the measurement error process is not influenced by the bone condition (Y_{i2}), as indicated by the estimates of γ_2 . The bodyweight of mice is only involved in the measurement error process but not the misclassification process because the estimate of γ_3 is significant but the

estimates of α_{x1} and α_{x0} are not. This suggests that the simpler specification of measurement models in Setting 2 is perhaps adequate and there is no need to consider the more complicated models in Setting 1.

Table 3.1: Results for Simulation 1 where parameters for mismeasurement models are assumed known: Setting 1 with different degrees of measurement error ($\sigma_e = 0.1, 0.5$ and 0.7)

σ_e	Naive Method																
	Naive Scenario 1				Naive Scenario 2				Naive Scenario 3								
	Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%					
0.1	β_{10}	0.469	0.034	0.034	0.0	0.012	0.031	0.034	94.3	0.466	0.040	0.040	0.4	-0.004	0.034	0.034	94.7
	β_{11}	-0.146	0.016	0.016	0.0	-0.027	0.016	0.016	61.0	-0.137	0.023	0.020	3.8	0.001	0.016	0.017	93.8
	β_{12}	0.075	0.033	0.033	37.1	0.012	0.033	0.033	93.9	0.063	0.049	0.042	62.8	-0.000	0.036	0.034	94.2
	β_{20}	-0.314	0.085	0.084	4.9	-0.313	0.082	0.084	4.6	0.015	0.190	0.152	90.3	0.047	0.201	0.204	96.9
	β_{21}	0.686	0.054	0.050	0.0	0.684	0.052	0.054	0.0	-0.018	0.195	0.180	92.4	-0.086	0.220	0.249	96.1
	β_{22}	-0.511	0.089	0.084	0.1	-0.509	0.087	0.086	0.1	0.002	0.212	0.180	90.5	0.068	0.243	0.253	97.1
σ	-	-	-	-	0.029	0.022	0.023	77.1	0.044	0.023	0.023	59.4	0.001	0.024	0.025	94.9	
ξ	-	-	-	-	-0.137	0.033	0.034	2.4	0.303	0.057	0.049	0.1	-0.002	0.060	0.063	95.3	
0.5	β_{10}	0.469	0.037	0.037	0.0	0.012	0.035	0.037	94.1	0.467	0.043	0.043	0.3	-0.004	0.036	0.038	95.3
	β_{11}	-0.146	0.018	0.018	0.0	-0.027	0.018	0.018	66.6	-0.137	0.025	0.022	4.8	0.001	0.019	0.018	93.6
	β_{12}	0.075	0.037	0.036	45.4	0.012	0.036	0.036	93.8	0.063	0.051	0.046	67.7	-0.001	0.038	0.037	93.3
	β_{20}	-0.455	0.076	0.074	0.0	-0.313	0.082	0.084	4.8	0.020	0.191	0.154	90.3	0.047	0.200	0.204	96.8
	β_{21}	0.986	0.041	0.039	0.0	0.684	0.052	0.054	0.0	-0.029	0.199	0.185	92.8	-0.085	0.220	0.249	96.1
	β_{22}	-0.712	0.077	0.073	0.0	-0.509	0.087	0.086	0.1	0.013	0.217	0.184	91.0	0.068	0.244	0.252	97.1
σ	-	-	-	-	0.029	0.028	0.028	83.5	0.155	0.026	0.026	0.0	0.002	0.029	0.030	94.6	
ξ	-	-	-	-	-0.137	0.035	0.037	4.5	0.275	0.056	0.048	0.1	-0.002	0.065	0.069	95.4	
0.7	β_{10}	0.470	0.040	0.041	0.0	0.013	0.038	0.040	94.0	0.466	0.046	0.046	0.4	-0.004	0.040	0.041	95.1
	β_{11}	-0.146	0.020	0.020	0.0	-0.027	0.019	0.019	70.9	-0.136	0.026	0.023	5.2	0.001	0.020	0.020	94.0
	β_{12}	0.075	0.040	0.040	51.6	0.011	0.038	0.039	93.8	0.059	0.053	0.049	70.8	-0.001	0.041	0.040	93.3
	β_{20}	-0.456	0.076	0.074	0.0	-0.313	0.082	0.084	4.9	0.024	0.191	0.157	91.0	0.047	0.200	0.203	96.8
	β_{21}	0.986	0.041	0.039	0.0	0.684	0.052	0.054	0.0	-0.036	0.201	0.188	92.9	-0.085	0.221	0.249	96.1
	β_{22}	-0.712	0.078	0.073	0.0	-0.509	0.087	0.086	0.1	0.019	0.221	0.187	91.6	0.067	0.243	0.252	97.1
σ	-	-	-	-	0.028	0.033	0.034	87.0	0.255	0.028	0.029	0.0	0.002	0.034	0.035	94.6	
ξ	-	-	-	-	-0.137	0.038	0.041	7.4	0.253	0.056	0.048	0.4	-0.002	0.070	0.075	95.7	

Table 3.2: Results for Simulation 1 where parameters for mismeasurement models are assumed known: Setting 2 with different misclassification levels ($\alpha = -4.595, -2.197$ and -1.386)

α_0	Naive Method																
	Naive Scenario 1				Naive Scenario 2				Naive Scenario 3								
	Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%					
-4.595	β_{10}	0.469	0.037	0.037	0.0	0.000	0.031	0.033	94.5	0.467	0.034	0.034	0.0	-0.001	0.031	0.033	94.3
	β_{11}	-0.146	0.018	0.018	0.0	-0.003	0.016	0.016	93.6	-0.144	0.016	0.016	0.0	-0.000	0.016	0.016	94.6
	β_{12}	0.075	0.037	0.036	45.5	0.001	0.032	0.032	95.6	0.073	0.034	0.034	42.3	0.000	0.032	0.032	95.5
	β_{20}	-0.045	0.104	0.108	93.1	-0.045	0.100	0.107	93.1	-0.006	0.109	0.112	94.7	0.015	0.114	0.121	96.2
	β_{21}	0.109	0.085	0.084	71.8	0.109	0.082	0.088	74.0	0.023	0.095	0.099	94.2	-0.020	0.104	0.111	95.5
-2.197	β_{22}	-0.082	0.116	0.113	88.1	-0.081	0.113	0.114	88.4	-0.027	0.121	0.120	91.9	0.017	0.128	0.133	94.3
	σ	-	-	-	-	0.002	0.022	0.023	95.3	0.037	0.022	0.023	65.2	-0.001	0.022	0.023	95.3
	ξ	-	-	-	-	-0.017	0.040	0.041	92.4	0.302	0.040	0.040	0.0	-0.001	0.043	0.045	95.5
	β_{10}	0.469	0.037	0.037	0.0	0.012	0.031	0.034	94.3	0.466	0.040	0.040	0.4	-0.004	0.034	0.034	94.7
	β_{11}	-0.146	0.018	0.018	0.0	-0.027	0.016	0.016	61.0	-0.137	0.023	0.020	3.8	0.001	0.016	0.017	93.8
-1.386	β_{12}	0.075	0.037	0.036	45.5	0.012	0.033	0.033	93.9	0.063	0.049	0.042	62.8	-0.000	0.036	0.034	94.2
	β_{20}	-0.180	0.165	0.096	49.0	-0.313	0.082	0.084	4.6	0.015	0.190	0.152	90.3	0.047	0.201	0.204	96.9
	β_{21}	0.398	0.297	0.067	35.9	0.684	0.052	0.054	0.0	-0.018	0.195	0.180	92.4	-0.086	0.220	0.249	96.1
	β_{22}	-0.296	0.238	0.099	44.1	-0.509	0.087	0.086	0.1	0.002	0.212	0.180	90.5	0.068	0.243	0.253	97.1
	σ	-	-	-	-	0.029	0.022	0.023	77.1	0.044	0.023	0.023	59.4	0.001	0.024	0.025	94.9
ξ	-	-	-	-	-0.137	0.033	0.034	2.4	0.303	0.057	0.049	0.1	-0.002	0.060	0.063	95.3	
-1.386	β_{10}	0.469	0.037	0.037	0.0	0.026	0.033	0.034	88.7	0.460	0.059	0.048	7.9	-0.007	0.040	0.039	94.4
	β_{11}	-0.146	0.018	0.018	0.0	-0.057	0.016	0.017	6.3	-0.119	0.033	0.025	17.3	0.005	0.020	0.019	93.8
	β_{12}	0.075	0.037	0.036	45.5	0.028	0.034	0.033	86.4	0.048	0.071	0.055	65.1	-0.021	0.044	0.040	93.8
	β_{20}	-0.272	0.192	0.089	32.7	-0.455	0.075	0.074	0.0	0.061	0.292	0.192	84.2	0.122	0.330	0.338	96.5
	β_{21}	0.594	0.369	0.058	23.9	0.984	0.039	0.041	0.0	-0.111	0.320	0.249	84.6	-0.244	0.361	0.448	94.0
β_{22}	-0.435	0.280	0.090	29.4	-0.709	0.075	0.074	0.0	0.074	0.339	0.243	83.2	0.195	0.393	0.433	95.4	
σ	-	-	-	-	0.054	0.023	0.024	35.4	0.119	0.027	0.023	47.7	0.036	0.029	0.029	89.2	
ξ	-	-	-	-	-0.228	0.029	0.031	0.0	0.289	0.081	0.057	2.9	-0.008	0.080	0.093	95.8	

Table 3.3: Results for Simulation 1 where parameters for mismeasurement models are assumed known: The setting 3 with different values of γ ($\gamma = 0, -0.8$ and 0.8)

γ	Naive Method																	
	Naive Scenario 1				Naive Scenario 2				Naive Scenario 3				Proposed					
	Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%		
0.0	β_{10}	-0.001	0.033	0.033	94.2	-0.001	0.031	0.033	94.2	-0.003	0.032	0.033	94.1	-0.003	0.032	0.033	94.4	
	β_{11}	-0.000	0.016	0.016	94.4	-0.000	0.016	0.016	94.3	0.000	0.017	0.016	93.0	0.000	0.017	0.016	93.4	
	β_{12}	0.000	0.032	0.032	94.9	0.000	0.032	0.032	95.1	-0.000	0.034	0.032	93.8	-0.000	0.034	0.032	94.4	
	β_{20}	-0.314	0.085	0.084	4.9	-0.314	0.082	0.084	4.6	0.047	0.200	0.168	92.6	0.047	0.201	0.204	96.9	
	β_{21}	0.686	0.054	0.050	0.0	0.686	0.052	0.054	0.0	-0.086	0.221	0.207	94.2	-0.086	0.221	0.249	96.1	
	β_{22}	-0.511	0.089	0.084	0.1	-0.511	0.087	0.086	0.1	0.068	0.241	0.209	93.8	0.068	0.241	0.253	97.1	
	σ	-	-	-	-	-0.001	0.022	0.023	94.7	0.006	0.022	0.022	94.6	0.001	0.022	0.023	93.9	
	ξ	-	-	-	-	-0.000	0.032	0.034	96.2	-0.000	0.056	0.047	90.8	-0.000	0.056	0.059	96.5	
	-0.8	β_{10}	-0.470	0.033	0.034	0.0	-0.014	0.032	0.034	91.5	-0.467	0.043	0.040	0.3	-0.002	0.034	0.034	94.7
		β_{11}	0.146	0.016	0.016	0.0	0.027	0.016	0.016	59.8	0.137	0.024	0.020	3.8	-0.000	0.017	0.017	93.2
β_{12}		-0.075	0.034	0.033	39.8	-0.012	0.033	0.033	92.2	-0.064	0.050	0.043	63.5	0.000	0.034	0.034	93.8	
β_{20}		-0.314	0.085	0.084	4.9	-0.314	0.082	0.084	5.1	0.018	0.185	0.152	90.3	0.046	0.200	0.204	96.7	
β_{21}		0.686	0.054	0.050	0.0	0.684	0.052	0.054	0.0	-0.018	0.197	0.180	92.2	-0.085	0.221	0.249	96.1	
β_{22}		-0.511	0.089	0.084	0.1	-0.509	0.087	0.086	0.1	0.003	0.212	0.181	90.1	0.068	0.240	0.253	97.2	
σ		-	-	-	-	0.028	0.023	0.023	76.9	0.042	0.023	0.023	59.1	-0.000	0.025	0.025	94.7	
ξ		-	-	-	-	0.136	0.032	0.034	1.9	-0.304	0.055	0.048	0.0	0.001	0.060	0.063	95.8	
0.8		β_{10}	0.469	0.034	0.034	0.0	0.012	0.031	0.034	94.3	0.466	0.040	0.040	0.4	-0.004	0.034	0.034	94.7
		β_{11}	-0.146	0.016	0.016	0.0	-0.027	0.016	0.016	61.0	-0.137	0.023	0.020	3.8	0.001	0.016	0.017	93.8
	β_{12}	0.075	0.033	0.033	37.1	0.012	0.033	0.033	93.9	0.063	0.049	0.042	62.8	-0.000	0.036	0.034	94.2	
	β_{20}	-0.314	0.085	0.084	4.9	-0.313	0.082	0.084	4.6	0.015	0.190	0.152	90.3	0.047	0.201	0.204	96.9	
	β_{21}	0.686	0.054	0.050	0.0	0.684	0.052	0.054	0.0	-0.018	0.195	0.180	92.4	-0.086	0.220	0.249	96.1	
	β_{22}	-0.511	0.089	0.084	0.1	-0.509	0.087	0.086	0.1	0.002	0.212	0.180	90.5	0.068	0.243	0.253	97.1	
	σ	-	-	-	-	0.029	0.022	0.023	77.1	0.044	0.023	0.023	59.4	0.001	0.024	0.025	94.9	
	ξ	-	-	-	-	-0.137	0.033	0.034	2.4	0.303	0.057	0.049	0.1	-0.002	0.060	0.063	95.3	

Table 3.4: Results for Simulation 2: Setting 2 with different degree of measurement error when interval validation data is available ($\sigma_e = 0.1, 0.5$ and 0.7)

σ_e	Method with True Measurements										Proposed Method									
	Known Parameter					Internal Validation					External Validation									
	Bias	SEE	SEM	CR%	CR%	Bias	SEE	SEM	CR%	CR%	Bias	SEE	SEM	CR%	CR%					
0.1	β_{10}	-0.000	0.027	0.027	95.1	-0.004	0.034	0.034	94.7	-0.002	0.033	0.034	94.3	0.001	0.028	0.029	95.5			
	β_{11}	-0.000	0.013	0.013	95.2	0.001	0.016	0.017	93.8	-0.002	0.017	0.016	93.7	0.000	0.014	0.014	94.5			
	β_{12}	0.001	0.026	0.026	95.5	-0.000	0.036	0.034	94.2	-0.004	0.034	0.033	94.9	-0.000	0.028	0.027	94.3			
	β_{20}	0.004	0.089	0.092	95.8	0.047	0.201	0.204	96.9	0.037	0.182	0.198	97.0	0.031	0.198	0.208	95.5			
	β_{21}	-0.007	0.075	0.076	95.9	-0.086	0.220	0.249	96.1	-0.071	0.194	0.224	94.9	-0.070	0.231	0.250	95.1			
	β_{22}	0.009	0.099	0.098	94.9	0.068	0.243	0.253	97.1	0.056	0.211	0.219	96.4	0.055	0.230	0.236	94.6			
	σ	-	-	-	-	0.001	0.024	0.025	94.9	0.003	0.022	0.024	95.7	-0.000	0.020	0.020	93.9			
	ξ	-	-	-	-	-0.002	0.060	0.063	95.3	-0.001	0.054	0.055	96.2	-0.003	0.055	0.054	94.3			
	γ_2	-	-	-	-	-	-	-	-	0.000	0.006	0.006	94.9	-0.000	0.004	0.004	95.6			
	σ_e	-	-	-	-	-	-	-	-	-0.000	0.003	0.003	94.5	-0.000	0.002	0.002	94.8			
α_1	-	-	-	-	-	-	-	-	-0.025	0.208	0.215	95.7	-0.006	0.123	0.123	94.4				
α_0	-	-	-	-	-	-	-	-	-0.021	0.204	0.212	95.0	-0.005	0.122	0.121	94.4				
0.5	β_{10}	-0.000	0.027	0.027	95.1	-0.004	0.036	0.038	95.3	-0.003	0.035	0.037	94.6	0.001	0.032	0.034	94.8			
	β_{11}	-0.000	0.013	0.013	95.2	0.001	0.019	0.018	93.6	-0.003	0.018	0.017	93.6	0.000	0.016	0.016	94.2			
	β_{12}	0.001	0.026	0.026	95.5	-0.001	0.038	0.037	93.3	-0.005	0.035	0.035	93.8	-0.001	0.031	0.030	94.3			
	β_{20}	0.004	0.089	0.092	95.8	0.047	0.200	0.204	96.8	0.037	0.182	0.198	97.0	0.030	0.198	0.208	95.4			
	β_{21}	-0.007	0.075	0.076	95.9	-0.085	0.220	0.249	96.1	-0.071	0.194	0.224	94.8	-0.069	0.232	0.250	95.0			
	β_{22}	0.009	0.099	0.098	94.9	0.068	0.244	0.252	97.1	0.056	0.211	0.220	96.4	0.055	0.230	0.236	94.6			
	σ	-	-	-	-	0.002	0.029	0.030	94.6	0.004	0.026	0.027	95.5	-0.000	0.025	0.025	95.4			
	ξ	-	-	-	-	-0.002	0.065	0.069	95.4	-0.002	0.058	0.059	95.9	-0.002	0.060	0.059	94.8			
	γ_2	-	-	-	-	-	-	-	-	0.002	0.032	0.032	94.9	-0.000	0.018	0.018	95.6			
	σ_e	-	-	-	-	-	-	-	-	-0.001	0.016	0.016	94.5	-0.000	0.009	0.009	94.8			
α_1	-	-	-	-	-	-	-	-	-0.025	0.208	0.215	95.7	-0.006	0.123	0.123	94.4				
α_0	-	-	-	-	-	-	-	-	-0.021	0.204	0.212	95.0	-0.005	0.122	0.121	94.4				
0.7	β_{10}	-0.000	0.027	0.027	95.1	-0.004	0.040	0.041	95.1	-0.003	0.038	0.040	95.4	0.001	0.037	0.038	94.7			
	β_{11}	-0.000	0.013	0.013	95.2	0.001	0.020	0.020	94.0	-0.003	0.019	0.019	95.0	0.000	0.017	0.017	95.1			
	β_{12}	0.001	0.026	0.026	95.5	-0.001	0.041	0.040	93.3	-0.006	0.037	0.036	92.8	-0.001	0.034	0.033	94.8			
	β_{20}	0.004	0.089	0.092	95.8	0.047	0.200	0.203	96.8	0.037	0.181	0.198	97.2	0.030	0.198	0.208	95.4			
	β_{21}	-0.007	0.075	0.076	95.9	-0.085	0.221	0.249	96.1	-0.070	0.194	0.224	94.7	-0.069	0.233	0.250	94.9			
	β_{22}	0.009	0.099	0.098	94.9	0.067	0.243	0.252	97.1	0.055	0.210	0.220	96.4	0.054	0.230	0.236	94.6			
	σ	-	-	-	-	0.002	0.034	0.035	94.6	0.004	0.030	0.031	95.2	-0.000	0.030	0.030	94.4			
	ξ	-	-	-	-	-0.002	0.070	0.075	95.7	-0.002	0.061	0.063	96.0	-0.002	0.064	0.064	95.0			
	γ_2	-	-	-	-	-	-	-	-	0.003	0.045	0.045	94.9	-0.001	0.025	0.026	95.6			
	σ_e	-	-	-	-	-	-	-	-	-0.001	0.022	0.022	94.5	-0.000	0.013	0.013	94.8			
α_1	-	-	-	-	-	-	-	-	-0.025	0.208	0.215	95.7	-0.006	0.123	0.123	94.4				
α_0	-	-	-	-	-	-	-	-	-0.021	0.204	0.212	95.0	-0.005	0.122	0.121	94.4				

Table 3.5: Results for Simulation 2: Setting 2 with different misclassification levels with interval validation data ($\alpha = -4.595, -2.197$ and -1.386)

α	Method with True Measurements										Proposed Method									
	Known Parameter					Internal Validation					External Validation									
	Bias	SEE	SEM	CR%	CR%	Bias	SEE	SEM	CR%	CR%	Bias	SEE	SEM	CR%	CR%	Bias	SEE	SEM	CR%	CR%
-4.595	β_{10}	-0.000	0.027	0.027	95.1	95.1	0.031	0.033	94.3	94.3	-0.001	0.031	0.033	94.2	94.2	0.000	0.025	0.027	95.9	95.9
	β_{11}	-0.000	0.013	0.013	95.2	95.2	-0.000	0.016	94.6	94.6	0.000	0.016	0.016	94.2	94.2	0.000	0.013	0.013	95.0	95.0
	β_{12}	0.001	0.026	0.026	95.5	95.5	0.000	0.032	95.5	95.5	0.000	0.032	0.032	95.7	95.7	-0.000	0.025	0.026	95.7	95.7
	β_{20}	0.004	0.089	0.092	95.8	95.8	0.015	0.114	96.2	96.2	0.013	0.114	0.122	96.5	96.5	0.009	0.105	0.104	94.7	94.7
	β_{21}	-0.007	0.075	0.076	95.9	95.9	-0.020	0.104	95.5	95.5	-0.019	0.104	0.109	97.0	97.0	-0.018	0.095	0.097	94.3	94.3
	β_{22}	0.009	0.099	0.098	94.9	94.9	0.017	0.128	94.3	94.3	0.016	0.122	0.130	94.3	94.3	0.015	0.112	0.112	95.2	95.2
	σ	-	-	-	-	-	-0.001	0.022	95.3	95.3	-0.001	0.021	0.023	95.3	95.3	-0.001	0.018	0.019	94.5	94.5
	ξ	-	-	-	-	-	-0.001	0.043	95.5	95.5	-0.001	0.042	0.044	95.1	95.1	-0.000	0.037	0.037	93.8	93.8
	σ_e	-	-	-	-	-	-	-	-	-	0.000	0.006	0.006	94.4	94.4	-0.000	0.004	0.004	95.6	95.6
	σ_1	-	-	-	-	-	-	-	-	-	-0.000	0.003	0.003	94.3	94.3	-0.000	0.002	0.002	94.8	94.8
σ_0	-	-	-	-	-	-	-	-	-	-1.767	0.566	0.693	96.4	96.4	-0.085	0.368	0.385	95.8	95.8	
-2.197	β_{10}	-0.000	0.027	0.027	95.1	95.1	-0.004	0.034	94.7	94.7	-0.002	0.033	0.034	94.3	94.3	0.001	0.028	0.029	95.5	95.5
	β_{11}	-0.000	0.013	0.013	95.2	95.2	0.001	0.016	93.8	93.8	0.002	0.017	0.016	93.7	93.7	0.000	0.014	0.014	94.5	94.5
	β_{12}	0.001	0.026	0.026	95.5	95.5	-0.000	0.036	94.2	94.2	-0.004	0.034	0.033	94.9	94.9	-0.000	0.028	0.027	94.3	94.3
	β_{20}	0.004	0.089	0.092	95.8	95.8	0.047	0.201	96.9	96.9	0.037	0.182	0.198	97.0	97.0	0.031	0.198	0.208	95.5	95.5
	β_{21}	-0.007	0.075	0.076	95.9	95.9	-0.086	0.220	96.1	96.1	-0.071	0.194	0.224	94.9	94.9	-0.070	0.231	0.250	95.1	95.1
	β_{22}	0.009	0.099	0.098	94.9	94.9	0.068	0.243	97.1	97.1	0.056	0.211	0.219	96.4	96.4	0.055	0.230	0.236	94.6	94.6
	σ	-	-	-	-	-	0.001	0.024	94.9	94.9	0.003	0.022	0.024	95.7	95.7	-0.000	0.020	0.020	93.9	93.9
	ξ	-	-	-	-	-	-0.002	0.060	95.3	95.3	-0.001	0.054	0.055	96.2	96.2	-0.003	0.055	0.054	94.3	94.3
	σ_e	-	-	-	-	-	-	-	-	-	0.000	0.006	0.006	94.9	94.9	-0.000	0.004	0.004	95.6	95.6
	σ_1	-	-	-	-	-	-	-	-	-	-0.000	0.003	0.003	94.5	94.5	-0.000	0.002	0.002	94.8	94.8
σ_0	-	-	-	-	-	-	-	-	-	-0.025	0.208	0.215	95.7	95.7	-0.006	0.123	0.123	94.4	94.4	
-1.386	β_{10}	-0.000	0.027	0.027	95.1	95.1	-0.007	0.040	94.5	94.5	-0.003	0.039	0.037	94.7	94.7	0.003	0.036	0.034	95.1	95.1
	β_{11}	-0.000	0.013	0.013	95.2	95.2	0.005	0.020	94.0	94.0	0.008	0.019	0.018	94.6	94.6	0.010	0.017	0.016	94.2	94.2
	β_{12}	0.001	0.026	0.026	95.5	95.5	-0.021	0.044	94.2	94.2	-0.021	0.039	0.035	94.2	94.2	-0.009	0.033	0.031	94.8	94.8
	β_{20}	0.004	0.089	0.092	95.8	95.8	0.122	0.330	96.4	96.4	0.103	0.276	0.313	96.5	96.5	0.102	0.322	0.342	94.6	94.6
	β_{21}	-0.007	0.075	0.076	95.9	95.9	-0.244	0.361	93.9	93.9	-0.224	0.325	0.390	94.2	94.2	-0.245	0.376	0.420	90.9	90.9
	β_{22}	0.009	0.099	0.098	94.9	94.9	0.195	0.393	95.3	95.3	0.167	0.335	0.352	94.9	94.9	0.188	0.362	0.393	92.4	92.4
	σ	-	-	-	-	-	0.036	0.029	89.7	89.7	0.031	0.025	0.026	93.2	93.2	0.037	0.025	0.024	90.5	90.5
	ξ	-	-	-	-	-	-0.008	0.080	95.7	95.7	-0.003	0.069	0.077	96.9	96.9	-0.009	0.072	0.080	94.6	94.6
	σ_e	-	-	-	-	-	-	-	-	-	0.000	0.006	0.006	94.9	94.9	-0.000	0.004	0.004	95.7	95.7
	σ_1	-	-	-	-	-	-	-	-	-	-0.000	0.003	0.003	94.6	94.6	-0.000	0.002	0.002	94.6	94.6
σ_0	-	-	-	-	-	-	-	-	-	-0.010	0.151	0.160	95.9	95.9	-0.007	0.089	0.092	94.9	94.9	
										-0.005	0.153	0.158	95.1	95.1	-0.002	0.085	0.091	95.2	95.2	

Table 3.6: Results for Simulation 2: Setting 3 with different values of γ with validation data ($\gamma = 0.0, -0.8$ and 0.8)

γ	Method with True Measurements										Proposed Method																																																																																																																																																																																																							
	Known Parameter					Internal Validation					External Validation																																																																																																																																																																																																							
	Bias	SEE	SEM	CR%	CR%	Bias	SEE	SEM	CR%	CR%	Bias	SEE	SEM	CR%	CR%	Bias	SEE	SEM	CR%	CR%																																																																																																																																																																																														
0.0	β_{10}	-0.000	0.027	0.027	95.1	-0.003	0.032	0.033	94.4	-0.002	0.031	0.033	94.0	-0.001	0.025	0.027	96.2	β_{11}	-0.000	0.013	0.013	95.2	-0.000	0.017	0.016	93.4	-0.001	0.016	0.016	94.0	-0.000	0.013	0.013	95.3	β_{12}	0.001	0.026	0.026	95.5	-0.000	0.034	0.032	94.4	-0.003	0.033	0.032	94.3	0.001	0.027	0.026	95.2	β_{20}	0.004	0.089	0.092	95.8	0.047	0.201	0.204	96.9	0.037	0.182	0.198	97.0	0.031	0.199	0.208	95.5	β_{21}	-0.007	0.075	0.076	95.9	-0.086	0.221	0.249	96.1	-0.070	0.195	0.225	94.8	-0.070	0.231	0.250	95.1	β_{22}	0.009	0.099	0.098	94.9	0.068	0.241	0.253	97.1	0.056	0.212	0.220	96.4	0.055	0.230	0.237	94.5	σ	-	-	-	-	0.001	0.022	0.023	93.9	0.002	0.022	0.022	94.7	-0.001	0.018	0.018	94.2	ξ	-	-	-	-	-0.000	0.056	0.059	96.5	-0.001	0.050	0.051	95.7	-0.002	0.049	0.048	95.0	γ_2	-	-	-	-	-	-	-	-	-	0.000	0.006	0.006	94.9	-0.000	0.004	0.004	95.6	σ_e	-	-	-	-	-	-	-	-	-	-0.000	0.003	0.003	94.5	-0.000	0.002	0.002	94.8	α_1	-	-	-	-	-	-	-	-	-	-0.025	0.208	0.215	95.7	-0.006	0.123	0.123	94.4	α_0	-	-	-	-	-	-	-	-	-	-0.021	0.204	0.212	95.0	-0.005	0.122	0.121	94.4		
	-0.8	β_{10}	-0.000	0.027	0.027	95.1	-0.002	0.034	0.034	94.7	-0.001	0.033	0.034	94.5	-0.001	0.028	0.029	95.9	β_{11}	-0.000	0.013	0.013	95.2	-0.000	0.017	0.017	93.2	0.000	0.017	0.016	94.1	-0.001	0.013	0.014	95.9	β_{12}	0.001	0.026	0.026	95.5	0.000	0.034	0.034	93.8	-0.001	0.034	0.033	94.5	0.002	0.028	0.028	95.7	β_{20}	0.004	0.089	0.092	95.8	0.046	0.200	0.204	96.7	0.037	0.182	0.198	97.1	0.030	0.199	0.208	95.4	β_{21}	-0.007	0.075	0.076	95.9	-0.085	0.221	0.249	96.1	-0.070	0.195	0.225	94.9	-0.069	0.231	0.250	95.1	β_{22}	0.009	0.099	0.098	94.9	0.068	0.240	0.253	97.2	0.056	0.210	0.219	96.4	0.054	0.230	0.237	94.5	σ	-	-	-	-	-0.000	0.025	0.025	94.7	0.001	0.023	0.024	94.7	-0.001	0.020	0.020	93.6	ξ	-	-	-	-	0.001	0.060	0.063	95.8	-0.000	0.055	0.056	95.8	-0.000	0.054	0.054	94.4	γ_2	-	-	-	-	-	-	-	-	-	0.000	0.006	0.006	94.9	-0.000	0.004	0.004	95.6	σ_e	-	-	-	-	-	-	-	-	-	-0.000	0.003	0.003	94.5	-0.000	0.002	0.002	94.8	α_1	-	-	-	-	-	-	-	-	-	-0.025	0.208	0.215	95.7	-0.006	0.123	0.123	94.4	α_0	-	-	-	-	-	-	-	-	-	-0.021	0.204	0.212	95.0	-0.005	0.122	0.121	94.4	
		0.8	β_{10}	-0.000	0.027	0.027	95.1	-0.004	0.034	0.034	94.7	-0.002	0.033	0.034	94.3	-0.001	0.028	0.029	95.5	β_{11}	-0.000	0.013	0.013	95.2	0.001	0.016	0.017	93.8	0.002	0.017	0.016	93.7	0.000	0.014	0.014	94.5	β_{12}	0.001	0.026	0.026	95.5	-0.000	0.036	0.034	94.2	-0.004	0.034	0.033	94.9	-0.000	0.028	0.027	94.3	β_{20}	0.004	0.089	0.092	95.8	0.047	0.201	0.204	96.9	0.037	0.182	0.198	97.0	0.031	0.198	0.208	95.5	β_{21}	-0.007	0.075	0.076	95.9	-0.086	0.220	0.249	96.1	-0.071	0.194	0.224	94.9	-0.070	0.231	0.250	95.1	β_{22}	0.009	0.099	0.098	94.9	0.068	0.243	0.253	97.1	0.056	0.211	0.219	96.4	0.055	0.230	0.236	94.6	σ	-	-	-	-	0.001	0.024	0.025	94.9	0.003	0.022	0.024	95.7	-0.000	0.020	0.020	93.9	ξ	-	-	-	-	-0.002	0.060	0.063	95.3	-0.001	0.054	0.055	96.2	-0.003	0.055	0.054	94.3	γ_2	-	-	-	-	-	-	-	-	-	0.000	0.006	0.006	94.9	-0.000	0.004	0.004	95.6	σ_e	-	-	-	-	-	-	-	-	-	-0.000	0.003	0.003	94.5	-0.000	0.002	0.002	94.8	α_1	-	-	-	-	-	-	-	-	-	-0.025	0.208	0.215	95.7	-0.006	0.123	0.123	94.4	α_0	-	-	-	-	-	-	-	-	-	-0.021	0.204	0.212	95.0	-0.005	0.122	0.121	94.4

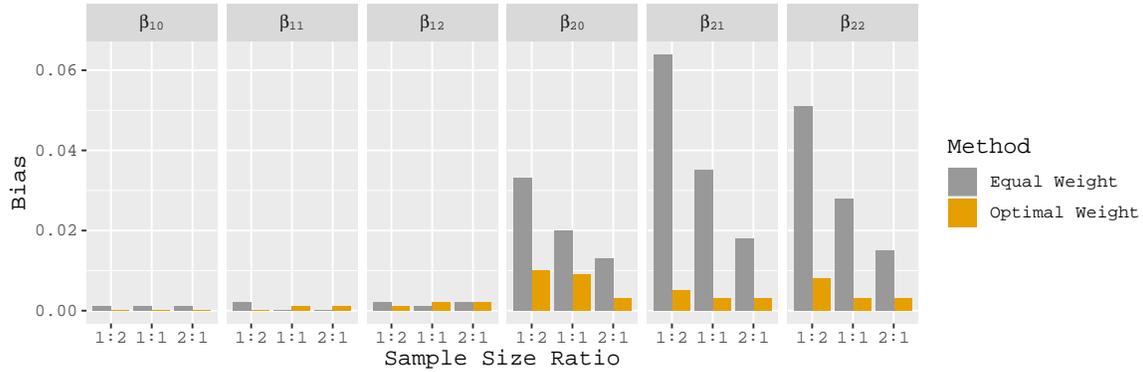


Figure 3.1: Biases of the estimates in Simulation 3: Scenario 1 with different sample size ratios between validation data and non-validation data. Equal Weight: the proposed method with internal validation data described in Section 3.3.2. Optimal Weight: the proposed weighted estimator with optimal weights as described in Section 3.3.3.

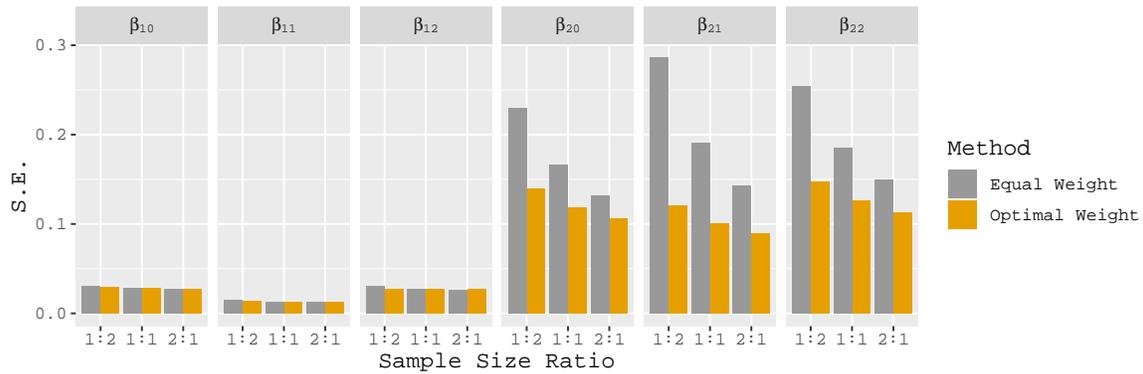


Figure 3.2: Standard error of the estimates in Simulation 3: Scenario 1 with different sample size ratios between validation data and non-validation data. Equal Weight: the proposed method with internal validation data described in Section 3.3.2. Optimal Weight: the proposed weighted estimator with optimal weights as described in Section 3.3.3.

Table 3.7: Results for Simulation 3: Scenario 2 with different weights and sample sizes

Parameter	Estimator	n=1500				n=3000					
		Bias	SEE	SEM	CR%	ARE ^a	Bias	SEE	SEM	CR%	ARE ^a
β_{10}	Naive	0.469	0.028	0.028	0.0	-	0.469	0.019	0.020	0.0	-
	$\hat{\theta}_I^{(0)}$	0.001	0.046	0.046	94.5	63%	-0.002	0.025	0.027	95.2	74%
	$\hat{\theta}_L^{(1)}$	-0.005	0.038	0.037	95.5	78%	0.001	0.028	0.029	95.5	69%
	$\hat{\theta}_I$	-0.001	0.029	0.031	95.4	94%	-0.001	0.022	0.021	94.9	95%
	$\hat{\theta}_I^*$	0.000	0.029	0.029	94.5	-	-0.001	0.019	0.020	96.0	-
β_{11}	Naive	-0.146	0.013	0.013	0.0	-	-0.146	0.009	0.009	0.0	-
	$\hat{\theta}_I^{(0)}$	-0.000	0.022	0.022	94.8	64%	-0.000	0.012	0.013	94.8	69%
	$\hat{\theta}_L^{(1)}$	0.007	0.017	0.017	95.1	82%	0.000	0.014	0.014	94.5	64%
	$\hat{\theta}_I$	0.002	0.013	0.015	94.2	93%	0.000	0.011	0.010	94.5	90%
	$\hat{\theta}_I^*$	-0.000	0.014	0.014	94.8	-	-0.000	0.009	0.009	95.3	-
β_{12}	Naive	0.077	0.026	0.027	19.2	-	0.074	0.019	0.019	2.8	-
	$\hat{\theta}_I^{(0)}$	0.002	0.045	0.045	94.7	6%	-0.001	0.025	0.026	93.7	73%
	$\hat{\theta}_L^{(1)}$	-0.011	0.035	0.034	93.8	79%	-0.000	0.028	0.027	94.3	70%
	$\hat{\theta}_I$	-0.002	0.027	0.031	94.7	87%	-0.002	0.022	0.019	94.9	100%
	$\hat{\theta}_I^*$	0.001	0.028	0.027	95.0	-	-0.001	0.020	0.019	95.0	-
β_{20}	Naive	-0.319	0.067	0.069	0.2	-	-0.321	0.050	0.048	0.0	-
	$\hat{\theta}_I^{(0)}$	0.014	0.155	0.160	95.3	88%	0.003	0.088	0.092	94.8	90%
	$\hat{\theta}_L^{(1)}$	0.081	0.284	0.296	96.1	47%	0.031	0.198	0.208	95.5	40%
	$\hat{\theta}_I$	0.033	0.186	0.230	96.50	61%	0.013	0.148	0.148	96.1	56%
	$\hat{\theta}_I^*$	-0.010	0.130	0.140	95.9	-	-0.005	0.081	0.083	94.2	-
β_{21}	Naive	0.692	0.046	0.041	0.0	-	0.693	0.032	0.029	0.0	-
	$\hat{\theta}_I^{(0)}$	-0.027	0.126	0.133	95.2	91%	-0.007	0.074	0.076	94.0	93%
	$\hat{\theta}_L^{(1)}$	-0.161	0.302	0.358	93.3	34%	-0.070	0.231	0.250	95.1	28%
	$\hat{\theta}_I$	-0.064	0.204	0.287	94.60	42%	-0.033	0.184	0.175	94.5	41%
	$\hat{\theta}_I^*$	0.005	0.118	0.121	94.0	-	0.005	0.072	0.071	93.1	-
β_{22}	Naive	-0.516	0.071	0.068	0.0	-	-0.517	0.050	0.048	0.0	-
	$\hat{\theta}_I^{(0)}$	0.028	0.168	0.171	95.2	86%	0.008	0.095	0.097	95.0	92%
	$\hat{\theta}_L^{(1)}$	0.119	0.287	0.319	94.9	46%	0.055	0.230	0.236	94.6	38%
	$\hat{\theta}_I$	0.051	0.201	0.254	95.9	58%	0.027	0.164	0.160	94.9	56%
	$\hat{\theta}_I^*$	-0.008	0.144	0.147	93.5	-	-0.005	0.087	0.089	94.2	-

^a ARE: Average relative efficiency comparing to the optimal weights.

Table 3.8: The implementation of the GEE in the genetic study under different specification of measurement error variance and misclassification rate

	Naive Method			Proposed Method ^a					
				Setting 1			Setting 2		
	Estimate	S.E.	p-value	Estimate	S.E.	p-value	Estimate	S.E.	p-value
Estimates for Response Model									
β_{10}	57.958	0.540	< 0.001	59.001	0.555	< 0.001	57.508	0.482	< 0.001
β_{11}	0.471	0.375	0.209	1.105	0.388	0.004	0.924	0.339	0.006
β_{12}	0.001	0.013	0.964	-0.000	0.019	0.986	-0.000	0.015	0.997
β_{13}	-0.012	0.022	0.599	-0.008	0.023	0.720	-0.008	0.021	0.692
β_{20}	-2.156	0.540	< 0.001	-2.615	0.313	< 0.001	-2.535	0.286	< 0.001
β_{21}	0.203	0.375	0.588	0.273	0.202	0.176	0.259	0.183	0.157
β_{22}	-0.001	0.013	0.958	0.003	0.009	0.718	0.002	0.008	0.816
β_{23}	0.003	0.022	0.896	-0.009	0.011	0.400	-0.006	0.010	0.520
Estimates for Mismeasurement Models									
γ_0	-	-	-	17.278	1.096	< 0.001	17.285	1.097	< 0.001
γ_1	-	-	-	0.560	0.016	< 0.001	0.560	0.016	< 0.001
γ_2	-	-	-	-	-	-	-0.371	0.390	0.341
γ_3	-	-	-	0.334	0.042	< 0.001	0.334	0.042	< 0.001
σ_e	-	-	-	2.966	0.069	< 0.001	2.967	0.069	< 0.001
α_{01}	-	-	-	-0.201	0.260	0.441	-2.060	2.442	0.399
α_{x1}	-	-	-	-	-	-	0.071	0.093	0.444
α_{00}	-	-	-	-4.446	0.380	< 0.001	0.685	7.498	0.927
α_{x0}	-	-	-	-	-	-	-0.206	0.311	0.509

^a Optimal weights are chosen for the internal validation data.

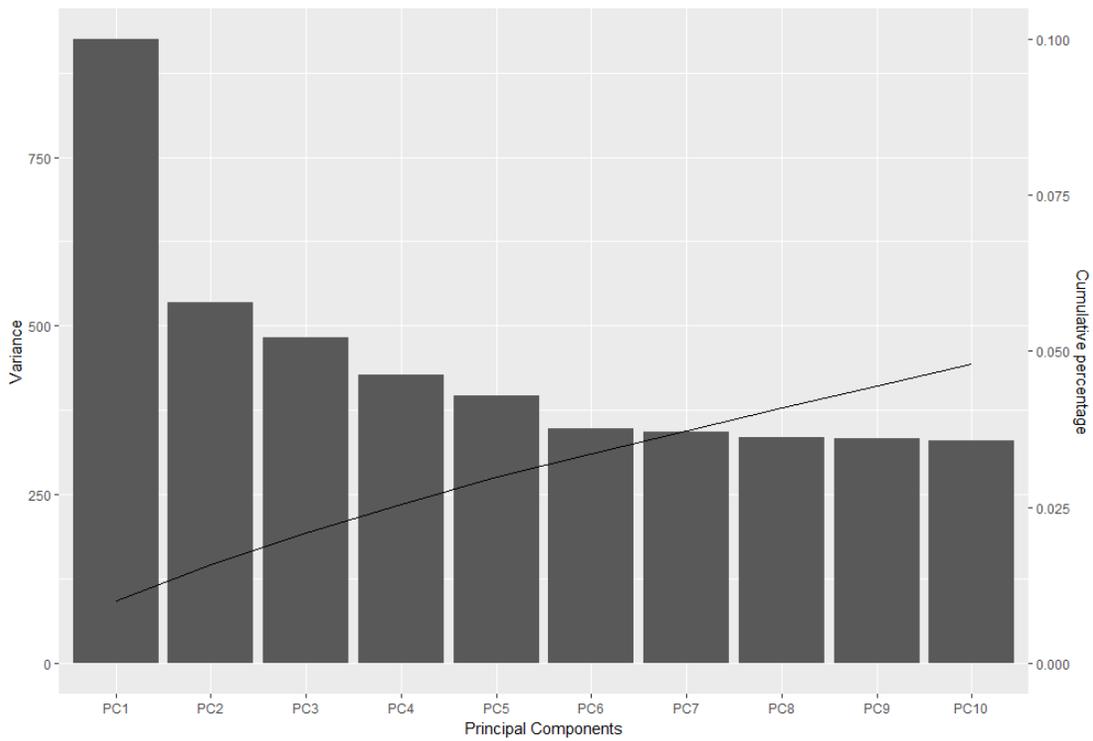


Figure 3.3: *The screeplot of the principal component analysis of the genotype data. The top 10 principal components are presented. The bar refers to the variance of each principal components. The solid line refers to the cumulative percentage of the variance.*

Chapter 4

Generalized Network Structured Models with Mixed Responses subject to Measurement Error and Misclassification

As a continuation of the previous two chapters, in this chapter, we further consider settings where covariates are of a high dimension and are associated with a network structure. In Section 4.1, we start with the saturated response model and discuss the estimation of the model parameters under the framework of generalized estimating equations. In Section 4.2, we develop the generalized network structured model (GNSM), describe a two-step implementation procedure of GNSM, and present the theoretical results for the proposed estimators. In Sections 4.3 and 4.4, we further extend the GNSM to the augmented GNSM by accounting for the effects due to measurement error and misclassification in the response variables, and we also discuss efficiency issues for the proposed estimators. Simulations studies are conducted in Section 4.5 to evaluate the performance of GNSM in regards to both variable selection and parameter estimation. In Section 4.6, we apply the augmented GNSM to a mice data set arising from a genome-wide association study.

4.1 Notation and Framework

Suppose n independent subjects are recruited for the study. For subject $i = 1, \dots, n$, correlated responses Y_{i1} and Y_{i2} are measured, where Y_{i1} denotes the continuous response,

and Y_{i2} denotes the binary response. Define $Y_i = (Y_{i1}, Y_{i2})^T$. Let $X_i = (X_{i1}, \dots, X_{ip})^T$ denote the covariate vector for subject i , where p is the number of covariates. For $i = 1, \dots, n$ and $j = 1, 2$, let $\mu_{ij} = E(Y_{ij}|X_i)$ be the conditional mean of the Y_{ij} , given X_i , and let $v_{ij} = \text{Var}(Y_{ij}|X_i)$ be the conditional variance of Y_{ij} given covariates X_i .

4.1.1 Saturated Response Model

To characterize the relationship among the covariates $\{X_{i1}, \dots, X_{ip}\}$, we use a *graph*, denoted as $G_i = (V_i, \tilde{E}_i)$, where $V_i = \{1, \dots, p\}$ includes all the indices of covariates and $\tilde{E}_i = V_i \times V_i$ is an index set of all pairs of covariates. A covariate X_{ij} is represented by a *vertex* of the graph G_i if $j \in V_i$. A pair of predictors $\{X_{is}, X_{it}\}$ is linked by an *edge* of the graph G_i if $(s, t) \in \tilde{E}_i$, and X_{is} and X_{it} are conditional dependent, given the remaining variables; let E_i denote the set of all pairs (s, t) if X_{is} and X_{it} are linked by an edge. We assume that all subjects have the same covariate dependence structures. Namely, $G_1 = G_2 = \dots = G_n \equiv G$. We now let V , \tilde{E} and E denote the vertices, $V \times V$, and edges of the graph, respectively.

We first consider a saturated model which includes all main effects and interactions,

$$\begin{aligned} g_1(\mu_{i1}) &= \beta_{1,0} + \sum_{k \in V} \beta_{1,k} X_{ik} + \sum_{(s,t) \in \tilde{E}} \beta_{1,st} X_{is} X_{it}; \\ g_2(\mu_{i2}) &= \beta_{2,0} + \sum_{k \in V} \beta_{2,k} X_{ik} + \sum_{(s,t) \in \tilde{E}} \beta_{2,st} X_{is} X_{it}, \end{aligned} \quad (4.1)$$

where $\beta = (\beta_M^T, \beta_P^T)^T$ with $\beta_M = (\beta_{1,0}, \beta_{2,0}, \beta_{1,k}, \beta_{2,k} : k \in V)^T$ and $\beta_P = (\beta_{1,st}, \beta_{2,st} : (s, t) \in \tilde{E})^T$, and $g_1(\cdot)$ and $g_2(\cdot)$ are link functions. For example, one may specify $g_1(t) = t$ and $g_2(t) = \log\{t/(1-t)\}$. Let p_s be the dimension of β .

4.1.2 Estimating Equation

In this chapter, we start with the same estimation procedure as in described in Section 3.1.2. We use the notation $\tilde{U}_{i1}(\beta, \phi)$ and $\tilde{U}_{i2}(\beta, \phi)$ to denote the estimation equations constructed based on the saturated model (4.1).

Let $\mu_i = (\mu_{i1}, \mu_{i2})^T$. For $i = 1, \dots, n$, define the estimating functions

$$\tilde{U}_{i1}(\beta, \phi) = D_{i1}^T V_{i1}^{-1} (Y_i - \mu_i); \quad (4.2)$$

$$\tilde{U}_{i2}(\beta, \phi) = D_{i2}^T V_{i2}^{-1} (S_i - \xi_i), \quad (4.3)$$

where $D_{i1} = \frac{\partial \mu_i}{\partial \beta}$, V_{i1} is given by (3.3), $\xi_i = (v_{ijk} : 1 \leq j \leq k \leq 2)^T$, $D_{i2} = \frac{\partial \xi_i}{\partial \phi^T}$, $S_i = \{(Y_{ij} - \mu_{ij})(Y_{ik} - \mu_{ik}) : 1 \leq j \leq k \leq 2\}^T$, and V_{i2} is a 3×3 weight matrix as defined in Section 3.1.2.

Let $\tilde{U}_i(\beta, \phi) = \left(\tilde{U}_{i1}^T(\beta, \phi), \tilde{U}_{i2}^T(\beta, \phi) \right)^T$. By the estimating function theory (e.g., Liang and Zeger, 1986; Godambe, 1991; Newey and McFadden, 1994; Yi, 2017, Section 1.3.2), under regularity conditions, solving

$$\sum_{i=1}^n \tilde{U}_i(\beta, \phi) = 0$$

for $(\beta^T, \phi^T)^T$ gives an estimator, say, $(\tilde{\beta}^T, \tilde{\phi}^T)^T$, of $(\beta^T, \phi^T)^T$.

4.2 Generalized Network Structured Model

4.2.1 Model Form

To focus on modeling the pairwise associations among the components of X_{ij} , we consider the graphical model (Hastie et al., 2015, Section 11)

$$f(x_i; \Theta) = \exp \left\{ -\frac{1}{2} \sum_{(s,t) \in \tilde{E}} \theta_{st} x_{is} x_{it} - \frac{1}{2} \sum_{k \in V} x_{ik}^2 - A(\Theta) \right\}, \quad (4.4)$$

where $\Theta = [\theta_{st}]$ is a $p \times p$ symmetric matrix with diagonal elements to be one and (s, t) element to be θ_{st} , and $A(\Theta) = -\frac{1}{2} \log \det \left| \frac{\Theta}{2\pi} \right|$ is the normalizing constant. This model basically implies that X_i follows a multivariate Gaussian distribution with zero mean and covariance matrix Σ , where $\Theta = \Sigma^{-1}$; and Θ is also known as the *precision matrix*.

In model (4.4), a nonzero parameter θ_{st} implies that X_{is} and X_{it} are conditionally dependent, given other covariates. In applications, not every paired covariates components in X_i are necessarily correlated. That is, the edge set E is not necessarily identical to \tilde{E} but $E = \{(s, t) \in \tilde{E} : \theta_{st} \neq 0\}$. To feature the dependence of the responses on the covariates with a network structure, we propose a generalized network structured model

$$\begin{aligned} g_1(\mu_{i1}) &= \beta_{1,0} + \sum_{k \in V} \beta_{1,k} X_{ik} + \sum_{(s,t) \in E} \beta_{1,st} X_{is} X_{it}; \\ g_2(\mu_{i2}) &= \beta_{2,0} + \sum_{k \in V} \beta_{2,k} X_{ik} + \sum_{(s,t) \in E} \beta_{2,st} X_{is} X_{it}, \end{aligned} \quad (4.5)$$

where $\beta_{\text{M}} = (\beta_{1,0}, \beta_{2,0}, \beta_{1,k}, \beta_{2,k} : k \in V)^{\text{T}}$ and $\beta_{\text{I}} = (\beta_{1,st}, \beta_{2,st} : (s, t) \in E)^{\text{T}}$ are the regression coefficients. To differentiate the parameters in the saturated model (4.1), we let $\beta_{\text{II}} = (\beta_{1,st}, \beta_{2,st} : (s, t) \in \tilde{E} \setminus E)^{\text{T}}$.

4.2.2 Estimation Procedure

To determine the model form (4.5) as well as to estimate the associated parameters, we need first to determine the set E . This essentially is equivalent to selecting active interaction terms in the saturated model (4.1). In this section, we describe a two-stage procedure. In Stage 1, we determine the dependence structure of the covariates via the graphical model (4.4). In Stage 2, we use the estimating equation method to estimate the associated model parameters. These two stages are respectively described in the following two subsections in detail.

Stage 1: Identification of the Covariates Network Structure E

To identify E , we maximize the penalized log-likelihood function

$$\ell(\Theta) = \sum_{i=1}^n \log f(x_i; \Theta) - \lambda \|\Theta\|, \quad (4.6)$$

where λ is a positive tuning parameter controlling the sparsity of the resulting parameter matrix and $\|\cdot\|$ is a penalizing norm function. A widely used norm is the ℓ_1 -norm, yielding the penalty of the Least Absolute Shrinkage and Selection Operator (Tibshirani, 1996).

Directly maximizing (4.6), such as the Graphical LASSO Algorithm, requires a computationally intensive algorithm (Friedman et al., 2008). In practice, a simpler estimation method is often carried out by a neighborhood-based likelihood derived from (4.4). Specifically, for every $s \in V$, let $X_{i, V \setminus \{s\}}$ denote the $(p-1)$ -dimensional subvector of X_i with its s th component removed, i.e., $X_{i, V \setminus \{s\}} = (X_{i1}, \dots, X_{i, s-1}, X_{i, s+1}, \dots, X_{ip})^{\text{T}}$. Then the conditional probability density function for X_{is} , given $X_{i, V \setminus \{s\}}$, is given by

$$f(x_{is} | x_{i, V \setminus \{s\}}; \theta_{(-s)}) = \exp \left\{ -\frac{1}{2} x_{is} \left(\sum_{t \in V \setminus \{s\}} \theta_{st} x_{it} \right) - \frac{1}{2} x_{is}^2 - D \left(\sum_{t \in V \setminus \{s\}} \theta_{st} x_{it} \right) \right\}, \quad (4.7)$$

where $D(\cdot) = \frac{1}{2} \log 2\pi + \frac{1}{8} (\sum_{t \in V \setminus \{s\}} \theta_{st} x_{it})^2$ is the normalizing constant ensuring the integration of (4.7) equal one, and $\theta_{(-s)} = (\theta_{s1}, \dots, \theta_{s, s-1}, \theta_{s, s+1}, \dots, \theta_{sp})^{\text{T}}$ is a $(p-1)$ -dimensional

vector of parameters indicating the relationship of X_{is} with all other predictors X_{it} for $t \in \{1, \dots, s-1, s+1, \dots, p\}$ associated with (4.7).

Let $\ell(\theta_{(-s)})$ be the log-likelihood for $\theta_{(-s)}$ multiplied by $-\frac{1}{n}$,

$$\begin{aligned} \ell(\theta_{(-s)}) &= -\frac{1}{n} \log \left\{ \prod_{i=1}^n f(x_{is} | x_{i, V \setminus \{s\}}; \theta_{(-s)}) \right\} \\ &= \frac{1}{n} \sum_{i=1}^n \left\{ \frac{1}{2} x_{is} \left(\sum_{t \in V \setminus \{s\}} \theta_{st} x_{it} \right) + D \left(\sum_{t \in V \setminus \{s\}} \theta_{st} x_{it} \right) \right\}. \end{aligned}$$

Then an estimator of $\theta_{(-s)}$ can be obtained as

$$\hat{\theta}_{(-s)} = \arg \min_{\theta_{(-s)}} \left\{ \ell(\theta_{(-s)}) + \lambda \|\theta_{(-s)}\|_1 \right\},$$

where λ is a tuning parameter and $\|\cdot\|_1$ is the ℓ_1 -norm.

The preceding procedure is repeated for all $s \in V$ to yield an estimator $\hat{\theta}_s$ for all $s \in V$. Let $\mathcal{N}(s) = \{t : (s, t) \in E\}$ denote the neighbor set for $s \in V$. To determine an estimated set of edges, we define

$$\hat{\mathcal{N}}(s) = \left\{ t \in V : \hat{\theta}_{st} \neq 0 \right\}$$

as the estimated neighbor set for $s \in V$. It is worth noting that, for $(s, t) \in E$, the estimates $\hat{\theta}_{st}$ and $\hat{\theta}_{ts}$ are not necessarily identical or both equal to zero at the same time although θ_{st} and θ_{ts} are constrained to be equal. Therefore, $s \in \hat{\mathcal{N}}(t)$ does not imply $t \in \hat{\mathcal{N}}(s)$, and vice versa. To overcome this discrepancy, we apply the OR rule (Meinshausen and Bühlmann, 2006; Hastie et al., 2015, Page 255) when determining the inclusion of edge (s, t) in the estimated edge set \hat{E} by either $s \in \hat{\mathcal{N}}(t)$ or $t \in \hat{\mathcal{N}}(s)$. Namely, we take

$$\hat{E} = \left\{ (s, t) : s \in \hat{\mathcal{N}}(t) \text{ or } t \in \hat{\mathcal{N}}(s) \right\}$$

as the estimated set of the edges.

The preceding procedure requires a suitable choice of the tuning parameter λ . Several methods, including the cross-validation (e.g., the BIC), the stability approach to regularization selection (StARS) (Liu et al., 2010), and the rotation information criterion (Zhao et al., 2012), may be employed to determine the optimal value of λ . In this paper, we use the rotation information criterion. To be specific, we take a set of candidate values for λ ,

such as an equally spaced sequence from 0 to a certain positive value. We first arrange the sample data in an array and then shuffle the data by randomly rotating the order of subjects (rows) for each variable (columns). This procedure creates a reshuffled dataset so that the association between paired variables is minimal. Then we implement our method to this reshuffled dataset and find the smallest value of λ such that all edges are regularized to 0. We repeat this procedure several times (such as 10 times) using the R package **huge** (Zhao et al., 2012) and select the resulting smallest value of λ .

Under regular conditions in Meinshausen and Bühlmann (2006), we have that as $n \rightarrow \infty$,

$$P\left(\widehat{E} = E\right) \rightarrow 1.$$

That is, the estimated set of edges \widehat{E} approximates the true network structure E in probability. This results is available in Ravikumar et al. (2010, Section 2.2) and Theorem 5(b) of Yang et al. (2015).

Stage 2: Estimation of Model Parameters

Once the network structure for X is identified, estimation of the model parameters in model (4.5) can proceed in the same manner as in Section 4.1.2, with modifications of estimating functions (4.2) and (4.3) to reflect the difference in the parameters for models (4.1) and (4.5). Let $(\widehat{\beta}_M^T, \widehat{\beta}_I^T)^T$ and $\widehat{\phi}$, respectively, denote the resultant estimators of $(\beta_M^T, \beta_I^T)^T$ and ϕ . Let $U_{i1}(\beta_M, \beta_I, \phi)$ and $U_{i2}(\beta_M, \beta_I, \phi)$ denote, respectively, the estimating functions by modifying $\widetilde{U}_{i1}(\beta, \phi)$ and $\widetilde{U}_{i2}(\beta, \phi)$ in (4.2) and (4.3). Let $U_i(\beta_M, \beta_I, \phi) = (U_{i1}^T(\beta_M, \beta_I, \phi), U_{i2}^T(\beta_M, \beta_I, \phi))^T$.

Then solving

$$\sum_{i=1}^n U_i(\beta_M, \beta_I, \phi) = 0 \tag{4.8}$$

for $(\beta_M^T, \beta_I^T, \phi^T)^T$ gives an estimator, say, $(\widehat{\beta}_M^T, \widehat{\beta}_I^T, \widehat{\phi}^T)^T$.

We comment that the true edge set E in model (4.5) is unknown and is estimated via the procedure in Section 4.2.2, thus inducing extra uncertainty in implementing (4.8) for the estimation of parameters. Consistent with the comments after (4.5) on the expression of the parameters in models (4.1) and (4.5), we set $\widehat{\beta}_{II} = 0$ as the estimator for the subvector of β_{II} which includes the coefficients corresponding to the covariates in the unselected edge set $\widetilde{E} \setminus \widehat{E}$. It is noted that $\widehat{\beta}_{II}$ may not be identical to β_{II} : their dimension can even be

different due to the variability induced in estimating E in Section 4.2.2. We now establish theoretical results for the estimator obtained from preceding Stages 1 and 2.

Theorem 4.1 *Let $\beta_{\text{II}(0)}$ be the true value of β_{II} . Under regularity conditions, there exists some constant $c > 0$ such that*

$$P(\widehat{\beta}_{\text{II}} = \beta_{\text{II}(0)}) \geq 1 - O(\exp(-cn)).$$

This theorem suggests that as $n \rightarrow \infty$, with the probability approaching 1, the estimator $\widehat{\beta}_{\text{II}}$ has the oracle property.

Next, we establish asymptotic properties for the estimator $(\widehat{\beta}_{\text{M}}^{\text{T}}, \widehat{\beta}_{\text{I}}^{\text{T}}, \widehat{\phi}^{\text{T}})^{\text{T}}$ in the following theorem; the proof of the results is presented in Appendix C.3.

Theorem 4.2 *Let $(\beta_{\text{M}(0)}^{\text{T}}, \beta_{\text{I}(0)}^{\text{T}}, \phi_0^{\text{T}})^{\text{T}}$ denote the true value of the parameters $(\beta_{\text{M}}^{\text{T}}, \beta_{\text{I}}^{\text{T}}, \phi^{\text{T}})^{\text{T}}$. Under regularity conditions, we have the following results:*

- (i) $(\widehat{\beta}_{\text{M}}^{\text{T}}, \widehat{\beta}_{\text{I}}^{\text{T}}, \widehat{\phi}^{\text{T}})^{\text{T}}$ is a consistent estimator of $(\beta_{\text{M}(0)}^{\text{T}}, \beta_{\text{I}(0)}^{\text{T}}, \phi_0^{\text{T}})^{\text{T}}$.
- (ii) $\sqrt{n}\{(\widehat{\beta}_{\text{M}}^{\text{T}}, \widehat{\beta}_{\text{I}}^{\text{T}}, \widehat{\phi}^{\text{T}})^{\text{T}} - (\beta_{\text{M}(0)}^{\text{T}}, \beta_{\text{I}(0)}^{\text{T}}, \phi_0^{\text{T}})^{\text{T}}\}$ has the asymptotic normal distribution with mean zero and covariance matrix

$$\Gamma_0^{-1} \Sigma_0 \Gamma_0^{-1\text{T}}, \tag{4.9}$$

where $\Gamma_0 = \left\{ E \left(\frac{\partial U_i(\beta_{\text{M}}, \beta_{\text{I}}, \phi)}{\partial \beta_{\text{M}}^{\text{T}}} \right) \quad E \left(\frac{\partial U_i(\beta_{\text{M}}, \beta_{\text{I}}, \phi)}{\partial \beta_{\text{I}}^{\text{T}}} \right) \quad E \left(\frac{\partial U_i(\beta_{\text{M}}, \beta_{\text{I}}, \phi)}{\partial \phi^{\text{T}}} \right) \right\} \Big|_{\substack{\beta_{\text{M}} = \beta_{\text{M}(0)} \\ \beta_{\text{I}} = \beta_{\text{I}(0)} \\ \phi = \phi_0}}$ and

$$\Sigma_0 = E\{U_i(\beta_{\text{M}(0)}, \beta_{\text{I}(0)}, \phi_0) U_i^{\text{T}}(\beta_{\text{M}(0)}, \beta_{\text{I}(0)}, \phi_0)\}.$$

4.3 Generalized Network Structured Model with Measurement Error and Misclassification

Suppose that the response variables Y_{i1} and Y_{i2} are subject to mismeasurement and their precise measurements are not observed for every subject $i = 1, \dots, n$, but instead, surrogate measurements Y_{i1}^* and Y_{i2}^* are observed, respectively, for Y_{i1} and Y_{i2} . To describe the mismeasurement processes, we consider the same factorization described in (3.6) and the assumptions described in (3.7).

4.3.1 Measurement Error and Misclassification Models

Let $\pi_{i0} = P(Y_{i2}^* = 1|Y_{i2} = 0, Z_i)$ and $\pi_{i1} = P(Y_{i2}^* = 0|Y_{i2} = 1, Z_i)$ be the misclassification probabilities that may depend on the covariates. We consider the same misclassification models (3.8) as described in Section 3.2.1. For the continuous response Y_{i1} , we consider a regression model which facilitates possible dependence of Y_{i1}^* on $\{Y_{i1}, Y_{i2}, Z_i\}$, as given by (3.9). Let $\eta = (\gamma^T, \alpha^T)^T$ denote the vector of parameters associated with (3.8) and (3.9).

4.3.2 Estimation Procedures with a Given Nuisance Parameter

η

The presence of mismeasurement in Y_i does not affect the first step of identifying the network structure in X_i described in Section 4.2.2. However, if no action is taken to address measurement error and misclassification in the responses, simply replacing Y_{ij} with Y_{ij}^* in the estimating functions (4.2) and (4.3) would yield biased estimating functions, and hence, possibly resulting in inconsistent estimators.

To account for the mismeasurement effects, we construct valid estimating functions, say $U_i^*(\beta_M, \beta_I, \phi)$ expressed in terms of the observed measurements Y_{i1}^* and Y_{i2}^* together with the covariates and the model parameters, such that

$$E\{U_i^*(\beta_M, \beta_I, \phi)\} = 0.$$

To this end, we first define that $\Delta_{i0} = \frac{\pi_{i0} - \pi_{i0}^2}{(1 - \pi_{i1} - \pi_{i0})^2}$, $\Delta_{i1} = \frac{\pi_{i1} - \pi_{i1}^2}{(1 - \pi_{i1} - \pi_{i0})^2}$, and $\Delta_i = \frac{\Delta_{i0}^{1-Y_{i2}^*} \Delta_{i1}^{Y_{i2}^*} - \Delta_{i0} \pi_{i1} - \Delta_{i1} \pi_{i0}}{1 - \pi_{i1} - \pi_{i0}}$, where π_{i0} and π_{i1} are the misclassification rates postulated by (3.8). Let $Y_{i2}^{**} = \frac{Y_{i2}^* - \pi_{i0}}{1 - \pi_{i0} - \pi_{i1}}$, $Y_{i1}^{**} = \frac{Y_{i1}^* - \gamma_0 - \gamma_2 Y_{i2}^{**} - \gamma_3^T Z_i}{\gamma_1}$, $Y_{i11}^{**} = Y_{i1}^{**2} - \frac{\sigma_\epsilon^2}{\gamma_1^2} - \frac{\gamma_2^2}{\gamma_1^2} \Delta_i$, and $Y_{i12}^{**} = Y_{i1}^{**} Y_{i2}^{**} + \frac{\gamma_2}{\gamma_1} \Delta_i$. In Section 3.2.2, it has been shown that

$$E(Y_{ik}^{**} | Y_{i1}, Y_{i2}) = Y_{ik} \quad \text{and} \quad E(Y_{i1k}^{**} | Y_{i1}, Y_{i2}, X_i) = Y_{i1} Y_{ik} \quad \text{for } k = 1, 2. \quad (4.10)$$

Let $U_i^*(\beta_M, \beta_I, \phi)$ be $U_i(\beta_M, \beta_I, \phi)$ in (4.8) with $Y_{i1}, Y_{i2}, Y_{i1}^2, Y_{i1} Y_{i2}$ replaced by $Y_{i1}^{**}, Y_{i2}^{**}, Y_{i11}^{**}$ and Y_{i12}^{**} , respectively. Then by (4.10),

$$E[U_i^*(\beta_M, \beta_I, \phi) | Y_{i1}, Y_{i2}, X_i] = U_i(\beta_M, \beta_I, \phi),$$

and thus, by the unbiasedness of $U_i(\beta_M, \beta_I, \phi)$, $U_i^*(\beta_M, \beta_I, \phi)$ is an unbiased estimating function. If the parameter η for the misclassification and measurement error models is known

or estimated from an additional study, then solving

$$\sum_{i=1}^n U_i^*(\beta_M, \beta_I, \phi) = 0 \quad (4.11)$$

for β_M , β_I and ϕ gives an estimator, say $(\widehat{\beta}_M^T, \widehat{\beta}_I^T, \widehat{\phi}^T)^T$, of $(\beta_M^T, \beta_I^T, \phi^T)^T$.

Theorem 4.3 *Under regularity conditions including those of [Newey and McFadden \(1994\)](#), [Yi \(2017, Section 1.3.2\)](#) and [Meinshausen and Bühlmann \(2006\)](#), the estimator $(\widehat{\beta}_M^T, \widehat{\beta}_I^T, \widehat{\phi}^T)^T$ is consistent, and $\sqrt{n}\{(\widehat{\beta}_M^T, \widehat{\beta}_I^T, \widehat{\phi}^T)^T - (\beta_{M(0)}^T, \beta_{I(0)}^T, \phi_0^T)^T\}$ has an asymptotic normal distribution with mean zero and covariance matrix*

$$\Gamma^{-1}\Sigma\Gamma^{-1T}, \quad (4.12)$$

where $\Gamma = \left\{ E \left(\frac{\partial U_i^*(\beta_M, \beta_I, \phi, \eta_0)}{\partial \beta_M^T} \right) \quad E \left(\frac{\partial U_i^*(\beta_M, \beta_I, \phi, \eta_0)}{\partial \beta_I^T} \right) \quad E \left(\frac{\partial U_i^*(\beta_M, \beta_I, \phi, \eta_0)}{\partial \phi^T} \right) \right\} \Big|_{\substack{\beta_M = \beta_{M(0)} \\ \beta_I = \beta_{I(0)} \\ \phi = \phi_0}}$ and

$$\Sigma = E\{U_i^*(\beta_{M(0)}, \beta_{I(0)}, \phi_0, \eta_0)U_i^{*T}(\beta_{M(0)}, \beta_{I(0)}, \phi_0, \eta_0)\}.$$

4.4 Estimation Procedures with Validation Data

4.4.1 External Validation

To incorporate estimation of η in the estimation of $(\beta_M^T, \beta_I^T, \phi^T)^T$, consider the likelihood function contributed from subject i in the validation sample:

$$L_i(y_{i1}^*, y_{i2}^* | y_{i1}, y_{i2}, x_i; \eta) = f(y_{i1}^* | y_{i1}, y_{i2}, z_i) f(y_{i2}^* | y_{i1}, y_{i2}, z_i),$$

where the index $i \in \mathcal{V}$, $f(y_{i1}^* | y_{i1}, y_{i2}, z_i)$ is determined by (3.9) with the form $\frac{1}{\sqrt{2\pi}\sigma_e} \exp\left\{-\frac{(y_{i1}^* - \gamma_0 - \gamma_1 y_{i1} - \gamma_2 y_{i2} - \gamma_3 z_i)^2}{2\sigma_e^2}\right\}$; and determined by (3.8), $f(y_{i2}^* | y_{i1}, y_{i2}, z_i)$ equals

$$\begin{aligned} & \left\{ \frac{\exp(\alpha_{00} + \alpha_{z_0}^T z_i)}{1 + \exp(\alpha_{00} + \alpha_{z_0}^T z_i)} \right\}^{(1-y_{i2})y_{i2}^*} \left\{ \frac{1}{1 + \exp(\alpha_{00} + \alpha_{z_0}^T z_i)} \right\}^{(1-y_{i2})(1-y_{i2}^*)} \\ & \times \left\{ \frac{\exp(\alpha_{01} + \alpha_{z_1}^T z_i)}{1 + \exp(\alpha_{01} + \alpha_{z_1}^T z_i)} \right\}^{y_{i2}(1-y_{i2}^*)} \left\{ \frac{1}{1 + \exp(\alpha_{01} + \alpha_{z_1}^T z_i)} \right\}^{y_{i2}y_{i2}^*}. \end{aligned}$$

Let

$$S_i(\eta) = \partial \log L_i(y_{i1}^*, y_{i2}^* | y_{i1}, y_{i2}, x_i; \eta) / \partial \eta \quad \text{for } i \in \mathcal{V}, \quad (4.13)$$

and construct the estimating function

$$U^{(\text{EV})}(\beta_M, \beta_I, \phi, \eta) = \sum_{i \in \mathcal{M}} \begin{pmatrix} U_i^*(\beta_M, \beta_I, \phi, \eta) \\ 0 \end{pmatrix} + \sum_{i \in \mathcal{V}} \begin{pmatrix} 0 \\ S_i(\eta) \end{pmatrix}, \quad (4.14)$$

where $S_i(\eta)$ is the score function determined by (4.13), and $U_i^*(\beta_M, \beta_I, \phi, \eta)$ is the estimating equation in (4.11) with the dependence on η explicitly spelled out. Then solving

$$U^{(\text{EV})}(\beta_M, \beta_I, \phi, \eta) = 0$$

for β_M, β_I, ϕ and η gives an estimator of β_M, β_I, ϕ and η , denoted as $\widehat{\beta}_M^{(\text{EV})}, \widehat{\beta}_I^{(\text{EV})}, \widehat{\phi}^{(\text{EV})}$, and $\widehat{\eta}^{(\text{EV})}$, respectively.

Since $S_i(\eta)$ does not depend on $(\beta_M^T, \beta_I^T, \phi^T)^T$, solving (4.14) is equivalent to a two-step procedure. First obtain $\widehat{\eta}^{(\text{EV})}$ by solving $\sum_{i \in \mathcal{V}} S_i(\eta) = 0$. Then solve the equation

$$\sum_{i \in \mathcal{M}} U_i^*(\beta_M, \beta_I, \phi, \widehat{\eta}^{(\text{EV})}) = 0$$

for β_M, β_I and ϕ to obtain estimators of β_M, β_I and ϕ , denoted as $\widehat{\beta}_M^{(\text{EV})}, \widehat{\beta}_I^{(\text{EV})}$ and $\widehat{\phi}^{(\text{EV})}$, respectively.

Theorem 4.4 *Assume that regularity conditions in Theorem 4.3 hold and that the ratio m/n approaches a positive constant ρ as $n \rightarrow \infty$, we have the following results:*

- (i) $\sqrt{n} \left\{ (\widehat{\beta}_M^{(\text{EV})T}, \widehat{\beta}_I^{(\text{EV})T}, \widehat{\phi}^{(\text{EV})T}, \widehat{\eta}^{(\text{EV})T})^T - (\beta_{M(0)}^T, \beta_{I(0)}^T, \phi_0^T, \eta_0^T)^T \right\}$ has an asymptotic normal distribution with mean zero and covariance matrix $\frac{1}{1+\rho} \Gamma_{(\text{EV})}^{-1} \Sigma_{(\text{EV})} (\Gamma_{(\text{EV})}^{-1})^T$, where

$$\begin{aligned} \Gamma_{(\text{EV})} &= \frac{1}{1+\rho} \begin{bmatrix} E \left(\frac{\partial U_i^*}{\partial \beta_M^T} \right) & E \left(\frac{\partial U_i^*}{\partial \beta_I^T} \right) & E \left(\frac{\partial U_i^*}{\partial \phi^T} \right) & E \left(\frac{\partial U_i^*}{\partial \eta^T} \right) \\ 0 & 0 & 0 & 0 \end{bmatrix} + \frac{\rho}{1+\rho} \begin{bmatrix} 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & E \left(\frac{\partial S_i}{\partial \eta^T} \right) \end{bmatrix}; \\ \Sigma_{(\text{EV})} &= \frac{1}{1+\rho} \begin{bmatrix} E(U_i^* U_i^{*T}) & 0 \\ 0 & 0 \end{bmatrix} + \frac{\rho}{1+\rho} \begin{bmatrix} 0 & 0 \\ 0 & E(S_i S_i^T) \end{bmatrix}. \end{aligned} \quad (4.15)$$

- (ii) $\sqrt{n} \left\{ (\widehat{\beta}_M^{(\text{EV})T}, \widehat{\beta}_I^{(\text{EV})T}, \widehat{\phi}^{(\text{EV})T})^T - (\beta_{M(0)}^T, \beta_{I(0)}^T, \phi_0^T)^T \right\}$ has an asymptotic normal distribution with mean zero and covariance matrix $(1+\rho) \Gamma_{(\text{EV})\beta}^{-1} \Sigma_{(\text{EV})\beta} \Gamma_{(\text{EV})\beta}^{-1T}$, where

$$\begin{aligned} \Gamma_{(\text{EV})\beta} &= \left\{ E \left(\frac{\partial U_i^*(\beta_M, \beta_I, \phi, \widehat{\eta}^{(\text{EV})})}{\partial \beta_M^T} \right) \quad E \left(\frac{\partial U_i^*(\beta_M, \beta_I, \phi, \widehat{\eta}^{(\text{EV})})}{\partial \beta_I^T} \right) \quad E \left(\frac{\partial U_i^*(\beta_M, \beta_I, \phi, \widehat{\eta}^{(\text{EV})})}{\partial \phi^T} \right) \right\} \Bigg|_{\substack{\beta_M = \beta_{M(0)} \\ \beta_I = \beta_{I(0)} \\ \phi = \phi_0}}; \\ \Sigma_{(\text{EV})\beta} &= E \{ U_i^*(\beta_{M(0)}, \beta_{I(0)}, \phi_0, \widehat{\eta}^{(\text{EV})}) U_i^{*T}(\beta_{M(0)}, \beta_{I(0)}, \phi_0, \widehat{\eta}^{(\text{EV})}) \}. \end{aligned} \quad (4.16)$$

The proof of Theorem 4.4(i) is presented in Appendix C.4, and Theorem 4.4(ii) can be readily derived from Theorem 4.4(i) by matrix calculation.

4.4.2 Internal Validation

With internal validation data, we consider the estimating function

$$U^{(iv)}(\beta_M, \beta_I, \phi, \eta) = \sum_{i \in \mathcal{M} \setminus \mathcal{V}} \begin{pmatrix} U_i^*(\beta_M, \beta_I, \phi, \eta) \\ 0 \end{pmatrix} + \sum_{i \in \mathcal{V}} \begin{pmatrix} U_i(\beta_M, \beta_I, \phi) \\ S_i(\eta) \end{pmatrix}, \quad (4.17)$$

where for $i \in \mathcal{M} \setminus \mathcal{V}$, $U_i^*(\beta_M, \beta_I, \phi, \eta)$ is the estimating equation in (4.11) with the dependence on η explicitly spelled out and for $i \in \mathcal{V}$, $U_i(\beta_M, \beta_I, \phi)$ is the estimating function in (4.8), and $S_i(\eta)$ is determined by (4.13). Then solving equation

$$U^{(iv)}(\beta_M, \beta_I, \phi, \eta) = 0 \quad (4.18)$$

for β_M , β_I , ϕ and η yields estimators for them, respectively denoted as $\widehat{\beta}_M^{(iv)}$, $\widehat{\beta}_I^{(iv)}$, $\widehat{\phi}^{(iv)}$, and $\widehat{\eta}^{(iv)}$.

Similar to that in Section 4.4.1, solving (4.18) is equivalent to a two-step procedure. First obtain $\widehat{\eta}^{(iv)}$ by solving $\sum_{i \in \mathcal{V}} S_i(\eta) = 0$. Then solve the equation

$$\sum_{i \in \mathcal{M} \setminus \mathcal{V}} U_i^*(\beta_M, \beta_I, \phi, \widehat{\eta}^{(iv)}) + \sum_{i \in \mathcal{V}} U_i(\beta_M, \beta_I, \phi) = 0$$

for β_M , β_I and ϕ to obtain estimators of β_M , β_I and ϕ , denoted as $\widehat{\beta}_M^{(iv)}$, $\widehat{\beta}_I^{(iv)}$, $\widehat{\phi}^{(iv)}$, respectively.

Theorem 4.5 *Assume that regularity conditions in Theorem 4.3 hold and that the ratio m/n approaches a positive constant ρ as $n \rightarrow \infty$, we have the following results:*

- (i) $\sqrt{n} \left\{ (\widehat{\beta}_M^{(iv)\top}, \widehat{\beta}_I^{(iv)\top}, \widehat{\phi}^{(iv)\top}, \widehat{\eta}^{(iv)\top})^\top - (\beta_{M(0)}^\top, \beta_{I(0)}^\top, \phi_0^\top, \eta_0^\top)^\top \right\}$ has an asymptotic normal distribution with mean zero and covariance matrix $\Gamma_{(iv)}^{-1} \Sigma_{(iv)} (\Gamma_{(iv)}^{-1})^\top$, where

$$\begin{aligned} \Gamma_{(iv)} &= (1 - \rho) \begin{bmatrix} E \left(\frac{\partial U_i^*}{\partial \beta_M^\top} \right) & E \left(\frac{\partial U_i^*}{\partial \beta_I^\top} \right) & E \left(\frac{\partial U_i^*}{\partial \phi^\top} \right) & E \left(\frac{\partial U_i^*}{\partial \eta^\top} \right) \\ 0 & 0 & 0 & 0 \end{bmatrix} + \rho \begin{bmatrix} E \left(\frac{\partial U_i}{\partial \beta_M^\top} \right) & E \left(\frac{\partial U_i}{\partial \beta_I^\top} \right) & E \left(\frac{\partial U_i}{\partial \phi^\top} \right) & 0 \\ 0 & 0 & 0 & E \left(\frac{\partial S_i}{\partial \eta^\top} \right) \end{bmatrix}; \\ \Sigma_{(iv)} &= (1 - \rho) \begin{bmatrix} E(U_i^* U_i^{*\top}) & 0 \\ 0 & 0 \end{bmatrix} + \rho \begin{bmatrix} E(U_i U_i^\top) & E(U_i S_i^\top) \\ E(S_i U_i^\top) & E(S_i S_i^\top) \end{bmatrix}. \end{aligned} \quad (4.19)$$

(ii) $\sqrt{n} \left\{ (\widehat{\beta}_M^{(iv)\text{T}}, \widehat{\beta}_I^{(iv)\text{T}}, \widehat{\phi}^{(iv)\text{T}})^{\text{T}} - (\beta_{M(0)}^{\text{T}}, \beta_{I(0)}^{\text{T}}, \phi_0^{\text{T}})^{\text{T}} \right\}$ has an asymptotic normal distribution with mean zero and covariance matrix $\Gamma_{(iv)\beta}^{-1} \Sigma_{(iv)\beta} (\Gamma_{(iv)\beta}^{-1})^{\text{T}}$, where

$$\Gamma_{(iv)\beta} = (1 - \rho) \begin{bmatrix} E \left(\frac{\partial U_i^*}{\partial \beta_M^{\text{T}}} \right) & E \left(\frac{\partial U_i^*}{\partial \beta_I^{\text{T}}} \right) & E \left(\frac{\partial U_i^*}{\partial \phi^{\text{T}}} \right) \end{bmatrix} + \rho \begin{bmatrix} E \left(\frac{\partial U_i}{\partial \beta_M^{\text{T}}} \right) & E \left(\frac{\partial U_i}{\partial \beta_I^{\text{T}}} \right) & E \left(\frac{\partial U_i}{\partial \phi^{\text{T}}} \right) \end{bmatrix}, \quad (4.20)$$

$$\Sigma_{(iv)\beta} = (1 - \rho) E (U_i^* U_i^{*\text{T}}) + \rho E (U_i U_i^{\text{T}}) - \rho E (U_i S_i^{\text{T}}) \{ E (S_i S_i^{\text{T}}) \}^{-1} E (S_i U_i^{\text{T}}).$$

The proof of Theorem 4.5(i) is presented in Appendix C.5, and Theorem 4.5(ii) can be readily derived from Theorem 4.5(i) by matrix calculation. We comment that Theorem 4.4(ii) and Theorem 4.5(ii) have appealing implications in that they extend the estimator in Theorem 4.3 to more realistic scenarios with unknown parameter η associated with the measurement error and misclassification models to be estimated from additional data sources. The estimators in Theorem 4.4(ii), Theorem 4.5(ii) and Theorem 4.3 are all consistent estimators of $(\beta_M^{\text{T}}, \beta_I^{\text{T}}, \phi^{\text{T}})^{\text{T}}$, but they differ in the efficiency because of the nuisance parameter η . The estimator $(\widehat{\beta}_M^{(ev)\text{T}}, \widehat{\beta}_I^{(ev)\text{T}}, \widehat{\phi}^{(ev)\text{T}})^{\text{T}}$ in Theorem 4.4(ii) is less efficient than the estimator $(\widehat{\beta}_M^{\text{T}}, \widehat{\beta}_I^{\text{T}}, \widehat{\phi}^{\text{T}})^{\text{T}}$ in Theorem 4.3 if η_0 is set as $\widehat{\eta}^{(E)}$ associated with the asymptotic covariance matrix in Theorem 4.4(ii), because the asymptotic covariance matrix for the former estimator is $(1 + \rho)$ times of that of the latter estimator. On the contrary, the estimator in Theorem 4.5(ii) is more efficient than that in Theorem 4.3, provided certain conditions, as shown in

Theorem 4.6 Let $\Delta = E(U_i S_i^{\text{T}}) \{ E(S_i S_i^{\text{T}}) \}^{-1} E(S_i U_i^{\text{T}})$. Consider Σ_0 , Γ_0 , Σ and Γ that are defined in Theorems 4.2 and 4.3, respectively. Assume the regularity conditions of Theorem 4.5. If

$$\Gamma_0^{-1} \Sigma_0 \Gamma_0^{-1\text{T}} \leq \Gamma^{-1} \Sigma \Gamma^{-1\text{T}}, \quad (4.21)$$

and

$$\Gamma_0^{-1} \Sigma_0 \Gamma_0^{-1\text{T}} + \Gamma_0^{-1} \Sigma \Gamma_0^{-1\text{T}} \leq \Gamma^{-1} \Sigma \Gamma^{-1\text{T}} + \Gamma_0^{-1} \Sigma \Gamma^{-1\text{T}} + \Gamma_0^{-1} \Delta \Gamma_0^{-1\text{T}}, \quad (4.22)$$

then we have

$$Avar\{(\widehat{\beta}_M^{(iv)\text{T}}, \widehat{\beta}_I^{(iv)\text{T}}, \widehat{\phi}^{(iv)\text{T}})^{\text{T}}\} \leq Avar\{(\widehat{\beta}_M^{\text{T}}, \widehat{\beta}_I^{\text{T}}, \widehat{\phi}^{\text{T}})^{\text{T}}\}, \quad (4.23)$$

for every $\rho \in (0, 1]$, where $Avar(\cdot)$ represents the asymptotic covariance matrix of an estimator, and the inequality \leq is the Loewner order.

The proof of Theorem 4.6 is outlined in Appendix C.6. This theorem says that under some conditions, the estimators in Section 4.4.2, with nuisance parameter η estimated from

an internal validation subsample, are more efficient than the estimators in Section 4.3.2, with η being given. Such a result appears somewhat counterintuitive as one may expect estimation of η would induce additional variability for estimators of β_M , β_I and ϕ . However, this phenomenon arises commonly in the context of using estimating functions (instead of the likelihood method) for estimation, as discussed by Newey and McFadden (1994, Chapter 6). The condition (4.21) compares the asymptotic covariance matrix for two estimators derived from different scenarios. This condition requires the estimator in Theorem 4.2, derived from the true response measurements, to be more efficient than the estimator in Theorem 4.3, obtained from surrogates measurements, which is often true when Y_i is less variable than Y_i^* . To understand condition (4.22), we look at the two terms at the left-hand side first where the first term represents the asymptotic covariance matrix in Theorem 4.2(ii). For the first term of the left-hand side of (4.22), we replace the left Γ_0 with Γ and Σ_0 with Σ ; for the second term of the left-hand side of (4.22), we replace the right Γ_0 with Γ , then (4.22) requires that the difference of such changes cannot exceed $\Gamma_0 \Delta \Gamma_0^T$, a non-negative definite matrix which involves the variability of S_i (i.e., $E(S_i S_i^T)$), the covariance between S_i and U_i (i.e., $E(S_i U_i^T)$) and the sensitivity of U_i (i.e., Γ_0). The efficiency gain stated in Theorem 4.6 holds for any value $\rho \in (0, 1]$, meaning that using any reasonably large internal validation subsample always increase efficiency relative to the case with η being given, provided certain conditions discussed earlier.

4.5 Simulation Studies

4.5.1 Simulation 1: Comparison of the GNSM with Ordinary LASSO

In this subsection, we conduct simulation studies to evaluate the performance of the proposed method for the variable selection (Section 4.2.2) and the parameter estimation (Section 4.2.2), where no mismeasurement is present.

To evaluate the performance of the methods, we consider different graphs, displayed in Figure 4.1, with different dependent structures of the covariates, which are characterized by varying degrees of nodes. Here the *degree* of a node is defined as the number of edges connected to this node. In the *hub* graph, two nodes have a higher degree than the other four nodes. The *scale-free* graph is generated by the Barabási-Albert algorithm (Barabási and Albert, 1999), where we start with an initial graph with only two connected nodes and then randomly connect a new node to only one existing node successively. In the *block*

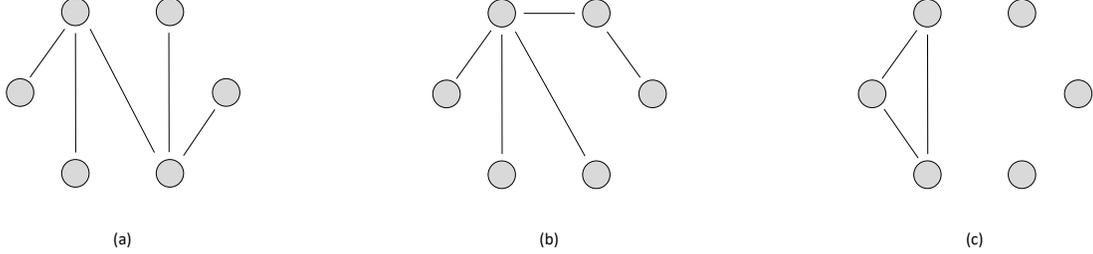


Figure 4.1: *Illustration of different graphs. (a): hub graph. (b): scale-free graph. (c): block graph*

graph, nodes are classified into several blocks in which the degrees of nodes within each block are the same.

The covariates X_i are generated from a multivariate normal distribution with mean zero and covariance matrix $\Sigma = \Theta^{-1}$, where the precision matrix Θ is, respectively, given by

$$\Theta_1 = \begin{bmatrix} 1 & \theta_{12} & \theta_{13} & \theta_{14} & 0 & 0 \\ \theta_{21} & 1 & 0 & 0 & 0 & 0 \\ \theta_{31} & 0 & 1 & 0 & 0 & 0 \\ \theta_{41} & 0 & 0 & 1 & \theta_{45} & \theta_{46} \\ 0 & 0 & 0 & \theta_{54} & 1 & 0 \\ 0 & 0 & 0 & \theta_{64} & 0 & 1 \end{bmatrix}, \Theta_2 = \begin{bmatrix} 1 & \theta_{12} & \theta_{13} & \theta_{14} & 0 & \theta_{16} \\ \theta_{21} & 1 & 0 & 0 & 0 & 0 \\ \theta_{31} & 0 & 1 & 0 & 0 & 0 \\ \theta_{41} & 0 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & \theta_{56} \\ \theta_{61} & 0 & 0 & 0 & \theta_{65} & 1 \end{bmatrix}, \text{ and } \Theta_3 = \begin{bmatrix} 1 & \theta_{12} & \theta_{13} & 0 & 0 & 0 \\ \theta_{21} & 1 & \theta_{23} & 0 & 0 & 0 \\ \theta_{31} & \theta_{32} & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 0 & 0 & 1 \end{bmatrix}$$

for the hub, scale-free, and block graphs. Here the θ_{ij} take a value either 0.2 or -0.2 for the connected edges, and 0 otherwise.

The responses Y_i are generated from the joint distribution

$$f(y_{i1}, y_{i2}) = \left[\Phi \left\{ \frac{g_2(\mu_{i2}) + \rho_c \left(\frac{y_{i1} - g_1(\mu_{i1})}{\sigma} \right)}{\sqrt{1 - \rho_c^2}} \right\} \right]^{y_{i2}} \left[1 - \Phi \left\{ \frac{g_2(\mu_{i2}) + \rho_c \left(\frac{y_{i1} - g_1(\mu_{i1})}{\sigma} \right)}{\sqrt{1 - \rho_c^2}} \right\} \right]^{1 - y_{i2}} \\ \times \frac{1}{\sqrt{2\pi}\sigma} \exp \left\{ -\frac{(y_{i1} - g_1(\mu_{i1}))^2}{2\sigma^2} \right\},$$

where $\Phi(t)$ is the cumulative distribution function for the standard normal distribution, $g_1(\mu_{i1})$ and $g_2(\mu_{i2})$ are specified as in model (4.5) with E indicated by each graph in Figure 4.1, ρ_c determines the correlation among Y_{i1} and Y_{i2} , and we set $g_1(t) = t$ and

$g_2(t) = \Phi^{-1}(t)$. The joint distribution was discussed by [de Leon and Wu \(2011\)](#). We set β_{II} to be a zero vector of dimension 22 or 18. Let β_{M} be of dimension 12, and let β_{I} be of dimension 6 or 10; the values of β_{M} and β_{I} are recorded in [Table 4.1](#) which fall in the intervals $[-0.7, -0.1] \cup [0.1, 0.7]$. The sample size n is set as 50, 200 or 1000. The simulations are run for 1000 times for each parameter configuration.

To show the performance of the methods, we evaluate the results for the network identification described in Stage 1 of [Section 4.2.2](#) and the results of parameter estimation described in Stage 2 of [Section 4.2.2](#) using different measures. For the procedure in Stage 1, we define two measures of variable selection, the true positive rate

$$\text{TPR}(\lambda) = \frac{|\{(s, t) : (s, t) \in E \text{ and } (s, t) \in \widehat{E}\}|}{|\{(s, t) : (s, t) \in E\}|}$$

and false negative rate

$$\text{FNR}(\lambda) = \frac{|\{(s, t) : (s, t) \notin E \text{ and } (s, t) \in \widehat{E}\}|}{|\{(s, t) : (s, t) \notin E\}|},$$

where $|\mathcal{A}|$ is the size of the set \mathcal{A} , and \widehat{E} is the estimated edge set for a given tuning parameter λ . For the estimation procedure in Stage 2, we consider two measures,

$$\|\widehat{\beta} - \beta\|_1 = \sum_{j \in \mathcal{I}} \sum_{t=1}^T |\widehat{\beta}_j^{(t)} - \beta_j^{(t)}| \quad \text{and} \quad \|\widehat{\beta} - \beta\|_2 = \sum_{j \in \mathcal{I}} \sum_{t=1}^T \sqrt{|\widehat{\beta}_j^{(t)} - \beta_j^{(t)}|^2},$$

where $\widehat{\beta}_j^{(t)}$ and $\beta_j^{(t)}$ represent, respectively, the j th component of $\widehat{\beta}$ and β in the t th simulation, T is the total number of simulations, and \mathcal{I} is the index set of β_{M} and β_{I} .

To see how the choice of the tuning parameter λ may affect the results obtained from [\(4.6\)](#), for a given λ in the interval $[0, 0.8]$, we plot $\text{TPR}(\lambda)$ against $\text{FNR}(\lambda)$ in [Figure 4.2](#), where we report the results for the sample size $n=50, 200$ and 1000. [Figure 4.2](#) shows that for a given λ , the performance of the method in [Section 4.2.2](#) improves as the sample size increases.

In [Table 4.2](#), we report the results obtained from the estimation procedure in [Section 4.2.2](#), which clearly demonstrate the improved performance of the proposed GNSM method as the sample size increases regardless of the graph types.

4.5.2 Simulation 2: Augmented GNSM with Measurement Error and Misclassification in Responses

In this subsection, we evaluate the performance of the proposed estimators when the mixed responses are subject to both measurement error and misclassification.

The covariates and the true responses are generated in the same way as in Section 4.5.1. The surrogate measurement Y_{i1}^* is generated from the measurement error model, $Y_{i1}^* = Y_{i1} + \gamma Y_{i2} + e_i$, where γ is set as 0.5, e_i follows a normal distribution with mean zero and variance σ_e^2 and is independent of $\{Y_{i1}, Y_{i2}\}$. We set σ_e^2 to be 0.2 or 0.7, reflecting different degrees of measurement error. The surrogate measurement Y_{i2}^* is generated from the misclassification models (3.8), where Z_i is generated from $\text{Uniform}(-2, 3)$, and the parameter α is set as $(-4, -1)^T$, $(-3, 0)^T$ and $(-3, 1)^T$, respectively yielding the misclassification rate of 1%, 5% and 10%. The sample size n is taken as 1000, and we take the generated data $\{(y_{i1}^*, y_{i2}^*, x_i) : i = 1, \dots, n\}$ as the main study data.

To simulate validation data, we generate a validation sample of size 500 using the same method as for generating the main study data. For the internal validation data, we keep all the measurements $\{(y_{j1}^*, y_{j2}^*, y_{j1}, y_{j2}, x_j, z_j) : j = 1, \dots, 500\}$ as validation data; and for external validation sample, we take $\{(y_{j1}^*, y_{j2}^*, y_{j1}, y_{j2}, z_j) : j = 1, \dots, 500\}$ as validation data.

Simulation studies are run 1000 times for each parameter configuration. We compare the performance of the augmented GNSM (Section 4.3.2) with the naive GNSM (Section 4.2.2) where the effects of measurement error and misclassification are ignored. To display the results, we separately report the results for $g_1(\mu_{i1})$ and $g_2(\mu_{i2})$ which respectively describe the continuous and binary responses. Let \mathcal{I} be the index set of β_M and β_I as in Section 4.5.1. Let $\hat{\beta}^{(t)j}$ denote the estimate for the j th component of $(\beta_M^T, \beta_I^T)^T$ at the t th simulation. We report the average bias (denoted ‘‘avgBias’’) by calculating $\frac{1}{|\mathcal{I}|} \sum_{j \in \mathcal{I}} |\bar{\beta}^j - \beta^j|$, the average empirical standard error (denoted ‘‘avgSEE’’) by calculating $\frac{1}{|\mathcal{I}|} \sum_{j \in \mathcal{I}} \text{esd}^j$, average model standard error (denoted ‘‘avgSEM’’) by calculating $\frac{1}{|\mathcal{I}|} \sum_{j \in \mathcal{I}} \text{msd}^j$, and the average coverage rate (denoted ‘‘avgCR’’) by calculating $\frac{1}{|\mathcal{I}|} \sum_{j \in \mathcal{I}} \text{CR}^j$, where $\bar{\beta}^j = \frac{1}{T} \sum_{t=1}^T \hat{\beta}^{(t)j}$, esd^j is the empirical standard error of the j th estimator, msd^j stands for the standard error of the j th estimator estimated by proposed model, CR^j is computed as

$$\text{CR}_l^j = \frac{1}{T} \sum_{t=1}^T I(\hat{\beta}^{(t)j(L)} < \beta_l^j < \hat{\beta}^{(t)j(U)}),$$

with $\hat{\beta}^{(t)j(L)}$ and $\hat{\beta}^{(t)j(U)}$ respectively representing the lower and upper bounds of the 95% confidence interval at simulation t , and T is the number of simulations taken as 1000 for each setting.

The results are presented in Tables 4.3 and 4.4. Simulation results clearly show that in the presence of mismeasurement in responses, the naive GNSM generally produces large finite sample biases and unreliable coverage rates for 95% confidence intervals. On the contrary, the augmented GNSM method adjusts for the mismeasurement effects and produces good results with small finite sample biases for the point estimates and fairly good coverage rates for 95% confidence intervals.

The estimators produced with the availability of an external validation sample have higher standard errors than those obtained under the scenario where the true parameters are known. On the other hand, the estimators resulted from the interval validation method are the most efficient among the three methods, confirming the results in Theorem 4.6.

4.6 Sensitivity Analysis of Mice SNPs Data

In this section, we apply the proposed method to analyze the outbred Carworth Farms White (CFW) mice data arising from a genome-wide association study (Parker et al., 2016b). The data set includes measurements of 1200 mice on behavioral, physiological, and gene expression traits. It is interesting to study the association between a set of candidate SNPs as well as their possible interactions with two bone morphology traits, defined as the length of the tibia and the bone condition. To be specific, the covariates include 20 candidate SNPs which were shown to be potentially associated with physiological traits, reported in Supplementary Table 2 of Parker et al. (2016b) and were scaled to have zero mean and unit standard error. Let Y_{i1} denote the length of the tibia bone (in *mm*) and let Y_{i2} be a binary outcome where “0” represents a healthy bone and “1” stands for an abnormal bone. The measurements of Y_{i1} and Y_{i2} are error-prone, where measurement error may be involved with the continuous responses Y_{i1} due to laboratory error and variation, and misclassification may occur in classifying the value of Y_{i2} which is based on the 90 percentile of bone-mineral density (BMD) of the sample. Consequently, the available measurements are taken as surrogate measurements, denoted as Y_{i1}^* and Y_{i2}^* , of the true responses Y_{i1} and Y_{i2} .

To analyze the data by accommodating possibly existing association structures in the covariates as well as addressing the mismeasurement effects in responses, we employ the two-step procedure for the proposed augmented generalized structured network model to

conduct inferences. In the first step, we fit a Gaussian graphical model to the covariates using the method of Section 4.2.2 with the optimal λ determined by the rotation information criterion. The identified association structure among the covariates is displayed on the left-hand side of Figure 4.3, which shows only four identified edges. On the right-hand side of Figure 4.3, we plot the sparsity level against the tuning parameter λ , where the sparsity level is defined as the number of selected edges divided by the total number of edges in the saturated graph. It is seen that the sparsity is fairly insensitive to the choice of tuning parameter around the neighbor of our optimal λ .

In the second step, we implement the estimation method described in Section 4.2.2 by incorporating the covariate association structure identified in the first step, where the response model is given by (4.5) with $g_1(t) = t$ and $g_2(t) = \log \frac{t}{1-t}$, and the measurement error model and the misclassification model are specified as (3.9) and (3.8), respectively. To show how inference results may be affected by mismeasurement effects, we conduct sensitivity analysis by considering different degrees of mismeasurement in Y_{i1} and Y_{i2} . For model (3.9), we take $\sigma_e = 0.77$ according to Lynch et al. (2019); in addition, we set σ_e to be 0.72 or 0.82. For model (3.8), we consider $\alpha_0 = \alpha_1 = -2.5, -1.5, \text{ or } -0.5$, respectively yielding tiny (5%), moderate (10%), and substantial (20%) misclassification rates.

The analysis results are presented in Tables 4.5–4.7. The estimation results and the inference conclusions are not sensitive to the different degrees of measurement error and misclassification rates we consider. The SNPs rs25203010 and rs265727287 are significantly associated with tibia length, which is consistent with the finding in Parker et al. (2016b). For the bone condition responses, rs33583459, rs29477109, and rs265727287 are identified to be the significant factors as their p-values are smaller than 0.01. The four interaction terms are strongly associated with the responses, indicating that the network structure plays an important role in studying the relationship between the candidate SNPs and the responses.

Table 4.1: The true parameter values for the data generation model in the simulation studies

Graph	The continuous component																					
	β_{11}	β_{12}	β_{13}	β_{14}	β_{15}	β_{16}	$\beta_{1,(1,2)}$	$\beta_{1,(1,3)}$	$\beta_{1,(1,4)}$	$\beta_{1,(1,5)}$	$\beta_{1,(1,6)}$	$\beta_{1,(2,3)}$	$\beta_{1,(2,4)}$	$\beta_{1,(2,5)}$	$\beta_{1,(2,6)}$	$\beta_{1,(3,4)}$	$\beta_{1,(3,5)}$	$\beta_{1,(3,6)}$	$\beta_{1,(4,5)}$	$\beta_{1,(4,6)}$	$\beta_{1,(5,6)}$	
Hub	0.644	0.675	0.331	0.430	0.139	-0.442	-0.458	0.409	-0.681	0	0											
Scale-free	0.644	0.675	0.331	0.430	0.139	-0.442	-0.458	0.409	-0.681	0	0.518											
Block	0.644	0.675	0.331	0.430	0.139	-0.442	-0.458	0.409	0	0	0											
Hub	0	0	0	0	0	0	0	0	0.662	0.633	0											
Scale-free	0	0	0	0	0	0	0	0	0	0	0.484											
Block	-0.390	0	0	0	0	0	0	0	0	0	0											
Graph	The binary component																					
	β_{21}	β_{22}	β_{23}	β_{24}	β_{25}	β_{26}	$\beta_{2,(1,2)}$	$\beta_{2,(1,3)}$	$\beta_{2,(1,4)}$	$\beta_{2,(1,5)}$	$\beta_{2,(1,6)}$	$\beta_{2,(2,3)}$	$\beta_{2,(2,4)}$	$\beta_{2,(2,5)}$	$\beta_{2,(2,6)}$	$\beta_{2,(3,4)}$	$\beta_{2,(3,5)}$	$\beta_{2,(3,6)}$	$\beta_{2,(4,5)}$	$\beta_{2,(4,6)}$	$\beta_{2,(5,6)}$	
Hub	-0.168	-0.270	-0.579	-0.266	0.143	-0.256	-0.396	-0.599	-0.160	0	0											
Scale-free	-0.168	-0.270	-0.579	-0.266	0.143	-0.256	-0.396	-0.599	-0.160	0	-0.516											
Block	-0.168	-0.270	-0.579	-0.266	0.143	-0.256	-0.396	-0.599	0	0	0											
Hub	0	0	0	0	0	0	0	-0.223	-0.445	0												
Scale-free	0	0	0	0	0	0	0	0	0	-0.665												
Block	-0.658	0	0	0	0	0	0	0	0	0												



Figure 4.2: Results for Simulation 1: The plot of true positive rate against the false negative rate obtained from the proposed GNSM for different values of tuning parameter λ

Table 4.2: Results for Simulation 1: The bias of the estimators of β with different responses types, sample sizes, graph types

n	Graph	Continuous Component		Discrete Component	
		$\ \cdot\ _1$	$\ \cdot\ _2$	$\ \cdot\ _1$	$\ \cdot\ _2$
50	block	6.369	3.562	7.636	3.195
	hub	7.291	3.727	9.274	3.651
	scale-free	6.914	3.420	9.128	3.579
200	block	1.366	0.752	1.345	0.550
	hub	0.772	0.312	1.836	0.684
	scale-free	0.691	0.279	1.806	0.676
1000	block	0.228	0.093	0.561	0.228
	hub	0.273	0.101	0.742	0.274
	scale-free	0.269	0.099	0.755	0.278

Table 4.3: Results for Simulation 2 with $\sigma_e = 0.2$

Graph	π^a	Naive Method			Known Parameter			Proposed Method								
		Internal Validation			Internal Validation			External Validation								
		avgBias	avgSEE	avgCI%	avgBias	avgSEE	avgCI%	avgBias	avgSEE	avgCI%						
		Continuous Response														
Block	1%	0.040	0.042	0.034	67.7	0.001	0.032	0.035	95.7	0.003	0.026	0.026	0.032	0.032	0.033	94.4
	5%	0.040	0.042	0.034	67.4	0.001	0.032	0.035	95.7	0.001	0.026	0.026	0.032	0.032	0.033	94.4
	10%	0.040	0.042	0.034	67.3	0.002	0.033	0.035	95.3	0.001	0.027	0.027	0.033	0.033	0.034	94.6
Hub	1%	0.036	0.042	0.034	70.6	0.001	0.032	0.034	95.9	0.001	0.026	0.026	0.032	0.032	0.033	94.3
	5%	0.037	0.042	0.034	70.5	0.001	0.032	0.034	95.7	0.001	0.026	0.026	0.032	0.032	0.033	94.3
	10%	0.037	0.042	0.034	70.4	0.001	0.032	0.034	95.3	0.001	0.026	0.027	0.033	0.033	0.034	94.3
Scale-free	1%	0.036	0.042	0.034	72.2	0.001	0.032	0.034	95.7	0.000	0.026	0.026	0.032	0.032	0.033	94.3
	5%	0.037	0.042	0.034	72.0	0.001	0.032	0.034	95.6	0.001	0.026	0.026	0.032	0.032	0.033	94.3
	10%	0.037	0.042	0.034	71.9	0.001	0.033	0.035	95.1	0.001	0.026	0.027	0.034	0.034	0.036	94.4
		Discrete Response														
Block	1%	0.021	0.079	0.081	93.7	0.007	0.085	0.089	95.3	0.005	0.069	0.071	0.086	0.086	0.091	95.6
	5%	0.069	0.076	0.079	83.3	0.011	0.095	0.099	95.4	0.004	0.075	0.078	0.098	0.098	0.102	95.6
	10%	0.139	0.073	0.076	53.1	0.018	0.119	0.114	93.6	0.012	0.097	0.102	0.135	0.135	0.148	96.7
Hub	1%	0.019	0.080	0.082	94.0	0.009	0.086	0.089	95.3	0.006	0.069	0.072	0.087	0.087	0.095	94.9
	5%	0.064	0.078	0.079	84.7	0.011	0.097	0.100	95.1	0.007	0.076	0.078	0.100	0.100	0.103	95.4
	10%	0.126	0.075	0.076	59.1	0.020	0.123	0.116	92.9	0.015	0.098	0.104	0.138	0.138	0.152	96.8
Scale-free	1%	0.022	0.080	0.083	93.6	0.006	0.088	0.091	95.2	0.007	0.071	0.073	0.091	0.091	0.098	95.1
	5%	0.071	0.078	0.081	83.4	0.009	0.099	0.102	94.8	0.007	0.078	0.081	0.102	0.102	0.106	95.4
	10%	0.140	0.076	0.077	53.6	0.019	0.126	0.120	93.0	0.016	0.101	0.109	0.143	0.143	0.161	96.9

^a The misclassification rate π is set as $\pi_1 = \pi_0$.

Table 4.4: Results for Simulation 2 with $\sigma_e = 0.7$

Graph	π^a	Naive Method			Known Parameter			Proposed Method									
		Internal Validation			Internal Validation			External Validation									
		avgBias	avgSEM	avgCI%	avgBias	avgSEM	avgCI%	avgBias	avgSEM	avgCI%	avgBias	avgSEM	avgCI%				
Continuous Response																	
Block	1	0.040	0.047	0.041	0.006	0.038	0.041	95.3	0.001	0.030	0.030	0.030	94.7	0.005	0.040	0.040	94.3
	5	0.040	0.047	0.041	0.003	0.038	0.041	95.2	0.001	0.030	0.030	0.030	94.4	0.004	0.040	0.040	94.4
	10	0.040	0.047	0.040	0.007	0.039	0.041	94.9	0.003	0.030	0.030	0.031	94.6	0.011	0.040	0.041	94.4
Hub	1	0.037	0.047	0.040	0.002	0.038	0.040	95.4	0.003	0.030	0.030	0.030	94.5	0.016	0.039	0.039	94.6
	5	0.037	0.047	0.040	0.003	0.038	0.040	95.3	0.006	0.030	0.030	0.030	94.4	0.019	0.039	0.039	94.6
	10	0.037	0.047	0.040	0.006	0.039	0.040	94.9	0.001	0.030	0.030	0.030	94.6	0.016	0.040	0.041	94.5
Scale-free	1	0.036	0.047	0.040	0.001	0.039	0.040	95.2	0.001	0.030	0.030	0.030	94.3	0.001	0.039	0.039	94.1
	5	0.037	0.047	0.040	0.002	0.039	0.040	95.1	0.001	0.030	0.030	0.030	94.4	0.004	0.039	0.039	94.1
	10	0.036	0.047	0.040	0.004	0.040	0.041	94.7	0.001	0.030	0.030	0.031	94.6	0.010	0.040	0.043	94.3
Discrete Response																	
Block	1	0.021	0.079	0.081	0.007	0.086	0.089	95.4	0.005	0.069	0.071	0.071	95.3	0.009	0.086	0.091	95.5
	5	0.069	0.076	0.079	0.011	0.095	0.099	95.4	0.004	0.075	0.078	0.078	95.3	0.009	0.098	0.102	95.6
	10	0.139	0.073	0.076	0.018	0.119	0.114	93.6	0.012	0.097	0.102	0.102	96.3	0.024	0.134	0.148	96.7
Hub	1	0.018	0.080	0.082	0.009	0.086	0.089	95.3	0.006	0.069	0.072	0.072	94.9	0.008	0.088	0.095	94.8
	5	0.063	0.078	0.079	0.011	0.097	0.100	95.1	0.007	0.076	0.078	0.078	94.9	0.012	0.100	0.103	95.4
	10	0.126	0.075	0.076	0.020	0.123	0.116	92.9	0.015	0.098	0.104	0.104	96.0	0.029	0.138	0.151	96.7
Scale-free	1	0.022	0.081	0.083	0.006	0.088	0.091	95.1	0.007	0.071	0.073	0.073	94.7	0.011	0.090	0.098	95.1
	5	0.071	0.078	0.081	0.009	0.099	0.102	94.8	0.007	0.078	0.081	0.081	95.3	0.013	0.102	0.106	95.4
	10	0.139	0.076	0.077	0.019	0.126	0.120	93.0	0.016	0.101	0.109	0.109	96.2	0.031	0.143	0.160	96.9

^a The misclassification rate π is set as $\pi_1 = \pi_0$.

Table 4.5: Results of the sensitivity analysis for the mice SNPs data when $\sigma_e = 0.72$

Response	Tiebia Length						Bone Condition					
	5%		10%		20%		5%		10%		20%	
	β (S.E.)	p-value										
rs45690064	0.106 (0.349)	0.761	0.106 (0.349)	0.761	0.106 (0.349)	0.761	0.028 (0.096)	0.772	0.073 (0.161)	0.650	0.059 (0.203)	0.770
rs27338905	-0.386 (0.347)	0.266	-0.386 (0.347)	0.266	-0.386 (0.347)	0.266	0.095 (0.101)	0.345	0.212 (0.169)	0.209	0.181 (0.213)	0.395
rs32962338	-0.547 (0.362)	0.131	-0.547 (0.362)	0.131	-0.547 (0.362)	0.131	0.182 (0.095)	0.054	0.245 (0.160)	0.125	0.287 (0.203)	0.158
rs33583459	-0.123 (0.345)	0.722	-0.123 (0.345)	0.722	-0.123 (0.345)	0.722	0.655 (0.109)	<0.001	1.281 (0.256)	<0.001	1.146 (0.389)	0.003
rs224051056	0.086 (0.344)	0.802	0.086 (0.344)	0.802	0.086 (0.344)	0.802	0.033 (0.097)	0.734	0.036 (0.156)	0.819	0.060 (0.199)	0.762
rs33217671	-0.230 (0.358)	0.521	-0.230 (0.358)	0.521	-0.230 (0.358)	0.521	0.241 (0.101)	0.017	0.465 (0.190)	0.014	0.439 (0.236)	0.063
rs38916331	-0.045 (0.340)	0.896	-0.045 (0.340)	0.896	-0.045 (0.340)	0.896	0.013 (0.107)	0.903	-0.001 (0.180)	0.997	0.035 (0.222)	0.873
rs47869247	-0.091 (0.354)	0.796	-0.091 (0.354)	0.796	-0.091 (0.354)	0.796	0.035 (0.095)	0.716	0.059 (0.155)	0.704	0.048 (0.196)	0.808
rs217439518	-0.202 (0.345)	0.559	-0.202 (0.345)	0.559	-0.202 (0.345)	0.559	0.062 (0.101)	0.540	0.111 (0.166)	0.505	0.129 (0.208)	0.534
rs29477109	-0.017 (0.355)	0.962	-0.017 (0.355)	0.962	-0.017 (0.355)	0.962	-0.458 (0.107)	<0.001	-0.820 (0.233)	<0.001	-0.783 (0.285)	0.006
rs252503010	-5.065 (1.598)	0.002	-5.065 (1.598)	0.002	-5.065 (1.598)	0.002	1.479 (0.642)	0.021	0.574 (0.466)	0.219	1.128 (0.882)	0.201
rs265727287	8.857 (1.627)	<0.001	8.857 (1.627)	<0.001	8.857 (1.627)	<0.001	-1.879 (0.445)	<0.001	-2.607 (0.586)	<0.001	-2.645 (0.977)	0.007
rs246035173	0.698 (0.481)	0.147	0.698 (0.481)	0.147	0.698 (0.481)	0.147	-0.094 (0.123)	0.445	-0.171 (0.215)	0.427	-0.184 (0.251)	0.463
rs231489766	-0.449 (0.497)	0.366	-0.449 (0.497)	0.366	-0.449 (0.497)	0.366	0.108 (0.137)	0.428	0.095 (0.225)	0.673	0.164 (0.287)	0.569
rs46826545	0.395 (0.411)	0.336	0.395 (0.411)	0.336	0.395 (0.411)	0.336	-0.089 (0.110)	0.419	-0.174 (0.185)	0.348	-0.173 (0.231)	0.452
rs51809856	0.751 (0.455)	0.099	0.751 (0.455)	0.099	0.751 (0.455)	0.099	-0.099 (0.116)	0.390	-0.257 (0.218)	0.338	-0.197 (0.254)	0.440
rs6279141	-0.141 (0.429)	0.743	-0.141 (0.429)	0.743	-0.141 (0.429)	0.743	0.053 (0.106)	0.619	0.058 (0.186)	0.754	0.091 (0.227)	0.688
rs30535702	0.087 (0.357)	0.808	0.087 (0.357)	0.808	0.087 (0.357)	0.808	-0.223 (0.101)	0.028	-0.498 (0.193)	0.010	-0.383 (0.243)	0.115
rs30201629	0.180 (0.381)	0.636	0.180 (0.381)	0.636	0.180 (0.381)	0.636	-0.046 (0.098)	0.641	-0.048 (0.141)	0.731	-0.081 (0.183)	0.655
rs30549753	-0.495 (0.334)	0.138	-0.495 (0.334)	0.138	-0.495 (0.334)	0.138	0.129 (0.090)	0.156	0.094 (0.125)	0.452	0.206 (0.173)	0.236
Interaction 1 ^a	7.178 (0.203)	<0.001	7.178 (0.203)	<0.001	7.178 (0.203)	<0.001	1.699 (0.598)	0.004	-2.082 (0.417)	<0.001	-1.526 (0.338)	<0.001
Interaction 2 ^b	3.249 (0.334)	<0.001	3.249 (0.334)	<0.001	3.249 (0.334)	<0.001	-0.650 (0.170)	<0.001	-0.969 (0.279)	0.001	-1.090 (0.384)	0.004
Interaction 3 ^c	2.612 (0.316)	<0.001	2.612 (0.316)	<0.001	2.612 (0.316)	<0.001	-0.507 (0.150)	0.001	-0.529 (0.190)	0.005	-0.777 (0.318)	0.014
Interaction 4 ^d	2.946 (0.345)	<0.001	2.946 (0.345)	<0.001	2.946 (0.345)	<0.001	-0.459 (0.128)	<0.001	-0.668 (0.229)	0.004	-0.808 (0.315)	0.010

^a rs252503010 × rs265727287

^b rs246035173 × rs231489766

^c rs231489766 × rs46826545

^d rs51809856 × rs6279141

Table 4.6: Results of the sensitivity analysis for the mice SNPs data when $\sigma_e = 0.77$

Response	Tiebia Length						Bone Condition					
	5%		10%		20%		5%		10%		20%	
	β (S.E.)	p-value										
rs45690064	0.106 (0.349)	0.761	0.106 (0.349)	0.761	0.106 (0.349)	0.761	0.028 (0.096)	0.772	0.057 (0.166)	0.732	0.070 (0.205)	0.732
rs27338905	-0.386 (0.347)	0.266	-0.386 (0.347)	0.266	-0.386 (0.347)	0.266	0.095 (0.101)	0.345	0.213 (0.171)	0.214	0.198 (0.217)	0.362
rs32962338	-0.547 (0.362)	0.131	-0.547 (0.362)	0.131	-0.547 (0.362)	0.131	0.182 (0.095)	0.054	0.237 (0.164)	0.149	0.287 (0.206)	0.164
rs33583459	-0.123 (0.345)	0.722	-0.123 (0.345)	0.722	-0.123 (0.345)	0.722	0.655 (0.109)	<0.001	1.555 (0.312)	<0.001	1.174 (0.397)	0.003
rs224051056	0.086 (0.344)	0.802	0.086 (0.344)	0.802	0.086 (0.344)	0.802	0.033 (0.097)	0.734	0.428 (0.158)	0.859	0.056 (0.201)	0.781
rs33217671	-0.230 (0.358)	0.521	-0.230 (0.358)	0.521	-0.230 (0.358)	0.521	0.241 (0.101)	0.017	0.444 (0.190)	0.020	0.463 (0.244)	0.058
rs38916331	-0.045 (0.340)	0.896	-0.045 (0.340)	0.896	-0.045 (0.340)	0.896	0.013 (0.107)	0.903	-0.014 (0.182)	0.938	0.042 (0.225)	0.851
rs47869247	-0.091 (0.354)	0.796	-0.091 (0.354)	0.796	-0.091 (0.354)	0.796	0.035 (0.095)	0.716	0.063 (0.157)	0.690	0.056 (0.199)	0.778
rs217439518	-0.202 (0.345)	0.559	-0.202 (0.345)	0.559	-0.202 (0.345)	0.559	0.062 (0.101)	0.540	0.110 (0.168)	0.513	0.133 (0.210)	0.526
rs29477109	-0.017 (0.355)	0.962	-0.017 (0.355)	0.962	-0.017 (0.355)	0.962	-0.458 (0.107)	<0.001	-0.806 (0.237)	0.001	-0.819 (0.299)	0.006
rs252503010	-5.065 (1.598)	0.002	-5.065 (1.598)	0.002	-5.065 (1.598)	0.002	1.479 (0.642)	0.021	0.622 (0.472)	0.188	0.955 (0.819)	0.244
rs265727287	8.857 (1.627)	<0.001	8.857 (1.627)	<0.001	8.857 (1.627)	<0.001	-1.879 (0.445)	<0.001	-2.686 (0.601)	<0.001	-2.621 (0.938)	0.005
rs246035173	0.698 (0.481)	0.147	0.698 (0.481)	0.147	0.698 (0.481)	0.147	-0.094 (0.123)	0.445	-0.170 (0.221)	0.443	-0.205 (0.257)	0.426
rs231489766	-0.449 (0.497)	0.366	-0.449 (0.497)	0.366	-0.449 (0.497)	0.366	0.108 (0.137)	0.428	0.090 (0.226)	0.692	0.140 (0.285)	0.624
rs46826545	0.395 (0.411)	0.336	0.395 (0.411)	0.336	0.395 (0.411)	0.336	-0.089 (0.110)	0.419	-0.169 (0.188)	0.369	-0.182 (0.234)	0.435
rs51809856	0.751 (0.455)	0.099	0.751 (0.455)	0.099	0.751 (0.455)	0.099	-0.099 (0.116)	0.390	-0.288 (0.228)	0.207	-0.209 (0.261)	0.423
rs6279141	-0.141 (0.429)	0.743	-0.141 (0.429)	0.743	-0.141 (0.429)	0.743	0.053 (0.106)	0.619	0.064 (0.192)	0.740	0.078 (0.230)	0.736
rs30535702	0.087 (0.357)	0.808	0.087 (0.357)	0.808	0.087 (0.357)	0.808	-0.223 (0.101)	0.028	-0.482 (0.194)	0.013	-0.414 (0.252)	0.100
rs30201629	0.180 (0.381)	0.636	0.180 (0.381)	0.636	0.180 (0.381)	0.636	-0.046 (0.098)	0.641	-0.044 (0.146)	0.765	-0.078 (0.183)	0.670
rs30549753	-0.495 (0.334)	0.138	-0.495 (0.334)	0.138	-0.495 (0.334)	0.138	0.129 (0.090)	0.156	0.075 (0.127)	0.552	0.193 (0.172)	0.262
Interaction 1 ^a	7.178 (0.203)	<0.001	7.178 (0.203)	<0.001	7.178 (0.203)	<0.001	-1.699 (0.598)	0.004	-2.069 (0.382)	<0.001	-1.577 (0.353)	<0.001
Interaction 2 ^b	3.249 (0.334)	<0.001	3.249 (0.334)	<0.001	3.249 (0.334)	<0.001	-0.650 (0.170)	<0.001	-0.934 (0.270)	0.001	-1.068 (0.378)	0.005
Interaction 3 ^c	2.612 (0.316)	<0.001	2.612 (0.316)	<0.001	2.612 (0.316)	<0.001	-0.507 (0.150)	0.001	-0.480 (0.181)	0.008	-0.720 (0.298)	0.016
Interaction 4 ^d	2.946 (0.345)	<0.001	2.946 (0.345)	<0.001	2.946 (0.345)	<0.001	-0.459 (0.128)	<0.001	-0.673 (0.236)	0.004	-0.801 (0.319)	0.012

^a rs252503010 × rs265727287

^b rs246035173 × rs231489766

^c rs231489766 × rs46826545

^d rs51809856 × rs6279141

Table 4.7: Results of the sensitivity analysis for the mice SNPs data when $\sigma_e = 0.82$

Response	Tiebia Length						Bone Condition					
	5%		10%		20%		5%		10%		20%	
	β (S.E.)	p-value										
rs45690064	0.106 (0.349)	0.761	0.106 (0.349)	0.761	0.106 (0.349)	0.761	0.028 (0.096)	0.772	0.069 (0.152)	0.648	0.070 (0.204)	0.734
rs27338905	-0.386 (0.347)	0.266	-0.386 (0.347)	0.266	-0.386 (0.347)	0.266	0.095 (0.101)	0.345	0.213 (0.158)	0.178	0.192 (0.216)	0.374
rs32962338	-0.547 (0.362)	0.131	-0.547 (0.362)	0.131	-0.547 (0.362)	0.131	0.182 (0.095)	0.054	0.258 (0.151)	0.089	0.287 (0.205)	0.161
rs33583459	-0.123 (0.345)	0.722	-0.123 (0.345)	0.722	-0.123 (0.345)	0.722	0.655 (0.109)	<0.001	1.236 (0.237)	<0.001	1.125 (0.383)	0.003
rs224051056	0.086 (0.344)	0.802	0.086 (0.344)	0.802	0.086 (0.344)	0.802	0.033 (0.097)	0.734	0.040 (0.148)	0.785	0.058 (0.201)	0.772
rs33217671	-0.230 (0.358)	0.521	-0.230 (0.358)	0.521	-0.230 (0.358)	0.521	0.241 (0.101)	0.017	0.448 (0.174)	0.010	0.460 (0.243)	0.058
rs38916331	-0.045 (0.340)	0.896	-0.045 (0.340)	0.896	-0.045 (0.340)	0.896	0.013 (0.107)	0.903	0.017 (0.169)	0.919	0.041 (0.224)	0.854
rs47869247	-0.091 (0.354)	0.796	-0.091 (0.354)	0.796	-0.091 (0.354)	0.796	0.035 (0.095)	0.716	0.057 (0.147)	0.699	0.053 (0.198)	0.790
rs217439518	-0.202 (0.345)	0.559	-0.202 (0.345)	0.559	-0.202 (0.345)	0.559	0.062 (0.101)	0.540	0.108 (0.156)	0.490	0.131 (0.209)	0.531
rs29477109	-0.017 (0.355)	0.962	-0.017 (0.355)	0.962	-0.017 (0.355)	0.962	-0.458 (0.107)	<0.001	-0.816 (0.215)	<0.001	-0.812 (0.296)	0.006
rs252503010	-5.065 (1.598)	0.002	-5.065 (1.598)	0.002	-5.065 (1.598)	0.002	1.479 (0.642)	0.021	0.743 (0.464)	0.110	0.969 (0.819)	0.237
rs265727287	8.857 (1.627)	<0.001	8.857 (1.627)	<0.001	8.857 (1.627)	<0.001	-1.879 (0.445)	<0.001	-2.505 (0.549)	<0.001	-2.596 (0.931)	0.005
rs246035173	0.698 (0.481)	0.147	0.698 (0.481)	0.147	0.698 (0.481)	0.147	-0.094 (0.123)	0.445	-0.184 (0.201)	0.360	-0.198 (0.256)	0.438
rs231489766	-0.449 (0.497)	0.366	-0.449 (0.497)	0.366	-0.449 (0.497)	0.366	0.108 (0.137)	0.428	0.088 (0.210)	0.676	0.149 (0.287)	0.604
rs46826545	0.395 (0.411)	0.336	0.395 (0.411)	0.336	0.395 (0.411)	0.336	-0.089 (0.110)	0.419	-0.153 (0.172)	0.375	-0.181 (0.233)	0.437
rs51809856	0.751 (0.455)	0.099	0.751 (0.455)	0.099	0.751 (0.455)	0.099	-0.099 (0.116)	0.390	-0.258 (0.204)	0.205	-0.201 (0.259)	0.436
rs6279141	-0.141 (0.429)	0.743	-0.141 (0.429)	0.743	-0.141 (0.429)	0.743	0.053 (0.106)	0.619	0.041 (0.174)	0.816	0.082 (0.230)	0.722
rs30535702	0.087 (0.357)	0.808	0.087 (0.357)	0.808	0.087 (0.357)	0.808	-0.223 (0.101)	0.028	-0.497 (0.184)	0.007	-0.406 (0.250)	0.104
rs30201629	0.180 (0.381)	0.636	0.180 (0.381)	0.636	0.180 (0.381)	0.636	-0.046 (0.098)	0.641	-0.055 (0.134)	0.682	-0.080 (0.182)	0.662
rs30549753	-0.495 (0.334)	0.138	-0.495 (0.334)	0.138	-0.495 (0.334)	0.138	0.129 (0.090)	0.156	0.100 (0.120)	0.401	0.200 (0.173)	0.247
Interaction 1 ^a	7.178 (0.203)	<0.001	7.178 (0.203)	<0.001	7.178 (0.203)	<0.001	-1.699 (0.598)	0.004	-1.916 (0.352)	<0.001	-1.563 (0.350)	<0.001
Interaction 2 ^b	3.249 (0.334)	<0.001	3.249 (0.334)	<0.001	3.249 (0.334)	<0.001	-0.650 (0.170)	<0.001	-0.956 (0.256)	<0.001	-1.081 (0.383)	0.005
Interaction 3 ^c	2.612 (0.316)	<0.001	2.612 (0.316)	<0.001	2.612 (0.316)	<0.001	-0.507 (0.150)	0.001	-0.540 (0.182)	0.003	-0.746 (0.308)	0.016
Interaction 4 ^d	2.946 (0.345)	<0.001	2.946 (0.345)	<0.001	2.946 (0.345)	<0.001	-0.459 (0.128)	<0.001	-0.678 (0.215)	0.002	-0.804 (0.318)	0.011

^a rs252503010 × rs265727287

^b rs246035173 × rs231489766

^c rs231489766 × rs46826545

^d rs51809856 × rs6279141

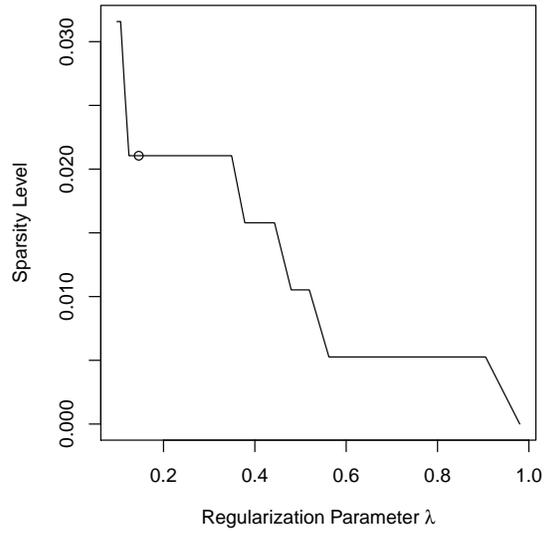
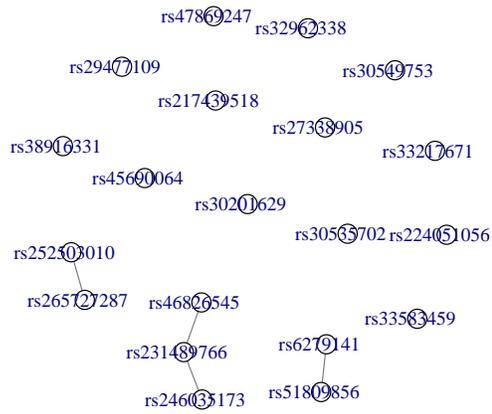


Figure 4.3: *Left panel: the diagram of the covariates structure; Right panel: the solution path sparsity levels*

Chapter 5

Zero-Inflated Poisson Models with Measurement Error in Response

In this chapter, we study the measurement error in the zero-inflated Poisson model. In Section 5.1, we discuss the setup of the response model as well as the measurement error model. In Section 5.2, we examine the effects of measurement error on analyzing count data and develop a method in Bayesian framework to account for the measurement error effects. In Section 5.3, we extend the method to accounting for the effects due to measurement error when validation subsamples are available. In Section 5.4, we illustrate the usage of the method by applying it to the prostate adenocarcinoma genomics data. To evaluate the performance of the method, we conduct simulation studies in Section 5.5.

5.1 Model Setup and Framework

5.1.1 Response Model

For $i = 1, \dots, n$, let Y_i denote the count outcome for subject i taking a non-negative integer value and let X_i denote the associated covariate vector of dimension p_x , where n is the number of subjects in the study. We assume that Y_i and $Y_{i'}$ are independent for any $i \neq i'$. The responses Y_i are sampled from two sources, either from an “at-risk” group where the measurements follow a Poisson distribution, or from a “non-at-risk” group where the measurements are zero. Let A_i be a latent indicator variable showing from which sources Y_i is sampled, where “ $A_i = 1$ ” represents Y_i is sampled from the “at-risk” group, and

“ $A_i = 0$ ” otherwise. For $i = 1, \dots, n$, let $\phi_i = P(A_i = 1|X_i)$ represent the conditional probability of sampling from ‘at-risk’ group, given X_i , and let $\mu_i = E(Y_i|A_i = 1, X_i)$ denote the condition mean of Y_i , given being sampled from the ‘at-risk’ group and the covariate X_i , which are assumed to satisfy $0 < \phi_i < 1$, and $\mu_i > 0$. That is, Y_i is sampled from the “non-at-risk” group with probability $1 - \phi_i$, and sampled from the “at-risk” group with probability ϕ_i , following a Poisson distribution with mean μ_i :

$$\begin{aligned} Y_i &= 0, \text{ with probability } 1 - \phi_i, \\ Y_i &\sim \text{Poisson}(\mu_i), \text{ with probability } \phi_i. \end{aligned} \tag{5.1}$$

Therefore, the zero values of Y_i may come from two sources: either from the “non-at-risk” group or from the “at-risk” group taking a zero count. Consequently, the probability mass function for response Y_i is given by

$$\begin{aligned} P(Y_i = 0|X_i) &= \sum_{k=0}^1 P(Y_i = 0|A_i = k, X_i)P(A_i = k|X_i) \\ &= (1 - \phi_i) + \phi_i e^{-\mu_i}; \\ P(Y_i = y_i|X_i) &= \phi_i \frac{\mu_i^{y_i} e^{-\mu_i}}{y_i!} \quad \text{for } y_i = 1, 2, \dots \end{aligned} \tag{5.2}$$

To facilitate the dependence of ϕ_i and μ_i on covariates X_i , we consider a complementary log-log regression model for ϕ_i and a log linear model for μ_i :

$$\text{cloglog } \phi_i = \beta_{\phi 0} + \beta_{\phi x}^T X_i, \tag{5.3}$$

$$\log \mu_i = \beta_{\mu 0} + \beta_{\mu x}^T X_i, \tag{5.4}$$

where $(\beta_{\phi 0}, \beta_{\phi x}^T)^T$ and $(\beta_{\mu 0}, \beta_{\mu x}^T)^T$ are the coefficients of the binary component and the count component respectively, $\beta = (\beta_{\phi 0}, \beta_{\phi x}^T, \beta_{\mu 0}, \beta_{\mu x}^T)^T$, and $\text{cloglog}(t) = \log\{-\log(1 - t)\}$ refers to the function of complementary log-log link. The complementary log-log link has been frequently used to model zero-inflated Poisson model in the literature (e.g. [Neelon and Chung, 2017](#)), whose interpretation is to be discussed in Section 5.2.1. We comment that although we used the same notation X_i to denote the covariates for ease of notations, the covariates could be different for each component in (5.3) and (5.4) by constraining the corresponding coefficients to be zero.

5.1.2 Measurement Error Model

Due to the measurement error in response Y_i , its precise measurement is not observed for every subject $i \in \{1, \dots, n\}$, but instead, surrogate measurement Y_i^* is observed for $i = 1, \dots, n$.

Measurement error for count data often arises from two distinct scenarios, and we call them the “*add-in*” and “*leave-out*”, respectively. The add-in error generates extra counts that are not supposed to be counted when measuring Y_i , yielding that the surrogate Y_i^* is no smaller than the true value of Y_i . For example, in genomics studies, we are interested in examining the count of copy number variants (CNVs). However, the mapping errors and incorrect sequencing may falsely include some insignificant CNVs, leading to the erroneous count higher than the true value (Xie and Tammi, 2009). On the contrary, the leave-out error may be caused by the loss of counts that should have been counted. In the CNV example, a significant CNV may fail to be identified due to the under-counting from the sequencing error. In the study of COVID-19, the daily reported cases number are often subject to leave-out error due to the limited test capacity and undetected asymptomatic infections as well as unreported cases with a mild symptom.

For measuring Y_i with $i = 1, \dots, n$, let Z_{i+} denote the count due to the add-in error and let Z_{i-} denote the count due to the leave-out error. Here we propose a measurement error model to feature the scenario where both the add-in and leave-out errors may exist; given X_i ,

$$Y_i^* = Y_i + c_+ Z_{i+} - c_- Z_{i-}, \quad (5.5)$$

where Z_{i+} is independent of Y_i and follows the Poisson distribution with mean λ_i , i.e. $\text{Poisson}(\lambda_i)$; Z_{i-} is independent of Z_{i+} but may be dependent on Y_i , and the conditional distribution of Z_{i-} , given $Y_i = y_i$ is the Binomial distribution with the probability π_i , i.e. $\text{Binomial}(y_i, \pi_i)$. Here c_+ and c_- are weights controlling the type of mismeasurements; they may be restricted to take values in $\{0, 1\}$ to facilitate various scenarios. For instance, if both c_+ and c_- are zero, then Y_i^* and Y_i are identical, i.e., no measurement error occurs; if $c_+ = 0$ and $c_- = 1$, then only leave-out error is involved; if $c_+ = 1$ and $c_- = 0$, then only add-in error exists; if $c_+ = c_- = 1$ then both add-in and leave-out errors are equally present. In applications, the background information or researchers’ experience may offer a good sense for specifying suitable values for c_+ and c_- .

Model (5.5) applies to count data and is a form somewhat similar to the widely used classical additive error model for featuring measurement error in continuous covariates. (Stefanski, 2000; Carroll et al., 2006; Yi, 2017, Section 2.6). But two key differences make model (5.5) unique. First, classical additive measurement error models do not differentiate error sources and use a single random variable, say e_i to represent the errors; secondly, the error term e_i is often assumed to be independent of true covariates. In model (5.5), however, the error term is refined by sorting out the errors of different nature. In addition, dependence of the error on the true variables is allowed. Basically, the joint distribution

of Y_i and $\{Z_{i+}, Z_{i-}\}$, is treated as

$$f(y_i, z_{i+}, z_{i-}|x_i) = f(y_i, z_{i-}|x_i)f(z_{i+}|x_i) \quad (5.6)$$

in model (5.5) by allowing the dependence between Y_i and Z_{i-} , where $f(\cdot)$ represents the joint or marginal distribution for the variables indicated by the arguments.

In model (5.5), assuming a Poisson distribution for Z_{i+} reflects its unboundedness yet taking a large value with a small probability. This assumption is feasible in applications where no upper limit is set for an add-in error and assuming errors beyond a certain value is not likely. On the contrary, the leave-out error cannot exceed the value of Y_i itself, so assuming a Binomial distribution for the conditional distribution for Z_{i-} , given Y_i , can be reasonable.

To facilitate different degrees of measurement error, we further model λ_i and π_i via their dependence on predictors, say, W_{i+} and W_{i-} , respectively, where W_{i+} and W_{i-} can be the same or different, and they can be part of covariates X_i or identical to X_i . Let $W_{i+} = (W_{i1+}, \dots, W_{ip+})^T$ and $W_{i-} = (W_{i1-}, \dots, W_{ip-})^T$ denote the covariate vector associated with add-in and leave-out processes, respectively, where p_+ and p_- are the dimension of W_{i+} and W_{i-} , respectively. For ease of exposition, we assume that W_{i+} and W_{i-} are subsets of X_i ; if this is not the case, we can modify our initial definition of X_i to include W_{i+} and W_{i-} as its parts.

The mean parameter λ_i is modeled as

$$\log \lambda_i = \alpha_{+0} + \alpha_{+w}^T W_{i+}, \quad (5.7)$$

and the probability π_i is postulated by a generalized linear model,

$$g(\pi_i) = \alpha_{-0} + \alpha_{-w}^T W_{i-}, \quad (5.8)$$

where $(\alpha_{+0}, \alpha_{+w}^T)^T$ and $(\alpha_{-0}, \alpha_{-w}^T)^T$ are coefficient vectors and $g(\cdot)$ is a link function. Here the link function $g(\cdot)$ can be taken as the logit function $g(t) = \log \frac{t}{1-t}$, the complementary log-log link $g(t) = \log\{-\log(1-t)\}$, or the probit function $g(\cdot) = \Phi^{-1}(\cdot)$, where $\Phi(\cdot)$ is the cumulative distribution function of the standard Gaussian distribution. Let $\alpha = (\alpha_{+0}, \alpha_{+w}^T, \alpha_{-0}, \alpha_{-w}^T)^T$.

5.1.3 Impact of Naive Analysis

In the presence of measurement error in response Y_i , the true response Y_i may not be observed. Instead, its surrogate Y_i^* is available. If we naively replace the response Y_i by

its surrogates Y_i^* in the inference procedure such as the likelihood method, the resulting estimators may not be consistent.

To see how the distribution of Y_i^* is different from Y_i , we consider the conditional distribution of Y_i^* given X_i :

$$\begin{aligned}
f(y_i^*|x_i) &= \sum_{z_{i-}} \sum_{z_{i+}} f_{(Y_i^*, Z_{i+}, Z_{i-})|X_i}(y_i^*, z_{i+}, z_{i-}|x_i) \\
&= \sum_{z_{i-}} \sum_{z_{i+}} f_{(Y_i, Z_{i+}, Z_{i-})|X_i}(y_i - z_{i+} + z_{i-}, z_{i+}, z_{i-}|x_i) \\
&= \sum_{z_{i-}} \sum_{z_{i+}} f_{Y_i|X_i}(y_i - z_{i+} + z_{i-}|x_i) f(z_{i+}|x_i) f(z_{i-}|y_i, x_i), \tag{5.9}
\end{aligned}$$

where the second step is due to (5.5), the third step is due to the independence assumption in (5.6), $f_{Y_i|X_i}(\cdot|x_i)$ is determined by (5.2) together with (5.3) and (5.4), and $f(z_{i+}|x_i)$ and $f(z_{i-}|y_i, x_i)$ are respectively determined by (5.7) and (5.8). Expression (5.9) shows that the conditional distribution for Y_i^* given X_i generally differs from that for Y_i given X_i

However, in some special cases, such as stated in Theorem 5.1 below, the conditional distribution, $f(y_i^*|x_i)$, of Y_i^* given X_i , is closely related to the conditional distribution (5.2) of Y_i given X_i in the structure.

Theorem 5.1 *Suppose Y_i follows the zero-inflated Poisson distribution given by (5.2) and the measurement error model for Y_i is given by (5.5).*

(a) *If $c_+ = 0$ and $c_- = 1$ in (5.5), then Y_i^* also follows a zero-inflated Poisson distribution given by*

$$\begin{aligned}
P(Y_i^* = 0|X_i) &= (1 - \phi_i^*) + \phi_i^* e^{-\mu_i^*}; \\
P(Y_i^* = y_i^*|X_i) &= \phi_i^* \frac{\mu_i^{*y_i^*} e^{-\mu_i^*}}{y_i^{*!}} \quad \text{for } y_i^* = 1, 2, \dots,
\end{aligned}$$

where $\phi_i^* = \phi_i$ and $\mu_i^* = (1 - \pi_i)\mu_i$.

(b) *If $c_+ = 1$ and $c_- = 0$ in (5.5), then Y_i^* follows a mixture distribution of two Poisson distributions, given by*

$$P(Y_i^* = y_i^*|X_i) = (1 - \phi_i) \frac{\lambda_i^{y_i^*} e^{-\lambda_i}}{y_i^{*!}} + \phi_i \frac{(\mu_i + \lambda_i)^{y_i^*}}{y_i^{*!}} e^{-(\mu_i + \lambda_i)} \quad \text{for } y_i^* = 0, 1, 2, \dots$$

(c) *If $c_+ = 1$ and $c_- = 1$ in (5.5), Y_i^* follows a mixture distribution of two Poisson distributions, given by*

$$P(Y_i^* = y_i^*|X_i) = (1 - \phi_i) \frac{\lambda_i^{y_i^*} e^{-\lambda_i}}{y_i^{*!}} + \phi_i \frac{\mu_i^{*y_i^*}}{y_i^{*!}} e^{-\mu_i^*} \quad \text{for } y_i^* = 0, 1, 2, \dots,$$

where $\mu_i^* = (1 - \pi_i)\mu_i + \lambda_i$.

The proof of Theorem 5.1 is presented in Appendix D.1. Theorem 5.1(a) says that if there is no add-in error in the measurement error model (5.5) and the leave-out error follows Binomial(y_i, π_i) then the surrogate variable Y_i^* assumes the same zero-inflated Poisson distribution (5.2) as the true response variable Y_i except for replacing μ_i with $(1 - \pi_i)\mu_i$, where the factor $1 - \pi_i$ reflects the impact of the degree of the leave-out error. We comment that Theorem 5.1 is analogous to the well-known Poisson Process Thinning Theorem (Brown, 1979, Theorem 1) which says that if Y_i follows Poisson(λ_i), then Y_i^* follows Poisson($(1 - \pi_i)\lambda_i$).

Theorem 5.1(b) suggests that if there is no leave-out error in measurement error model (5.5) and the add-in error follows Poisson(λ_i), then the distribution of the surrogate variable Y_i^* is determined by two Poisson distributions, given by

$$\begin{aligned} Y_i^* &\sim \text{Poisson}(\lambda_i), \text{ with probability } 1 - \phi_i, \\ Y_i^* &\sim \text{Poisson}(\mu_i + \lambda_i), \text{ with probability } \phi_i. \end{aligned} \quad (5.10)$$

Theorem 5.1(c) may be viewed as a combined result from Theorem 5.1(a) and (b), saying that when both the add-in error and the leave-out error are present, the distribution of the surrogate variable Y_i^* assumes the same form as (5.10) except that μ_i is replaced by $(1 - \pi_i)\mu_i$.

Next, we discuss possible biases of the naive analysis which disregards the difference between Y_i and Y_i^* . That is, we naively assume that Y_i^* follows the same distribution form as Y_i , then we replace Y_i in (5.2) with Y_i^* and let ϕ_i^* and μ_i^* denote the resulting quantities corresponding to ϕ_i and μ_i in (5.2), respectively; in addition, the same model forms as (5.3) and (5.4) are assumed for ϕ_i^* and μ_i^* :

$$\text{cloglog } \phi_i^* = \beta_{\phi 0}^* + \beta_{\phi x}^{*\text{T}} X_i, \quad (5.11)$$

$$\log \mu_i^* = \beta_{\mu 0}^* + \beta_{\mu x}^{*\text{T}} X_i, \quad (5.12)$$

where $\beta^* \triangleq (\beta_{\phi 0}^*, \beta_{\phi x}^{*\text{T}}, \beta_{\mu 0}^*, \beta_{\mu x}^{*\text{T}})^{\text{T}}$ are the associated parameters which may differ from the corresponding parameters in the models (5.3) and (5.4). Without adding any constraint on the measurement error (5.5), it is generally expected that β^* differs from β . Even with certain conditions for the measurement error model (5.5), such as those discussed in Theorem 5.1(b)(c), Y_i^* does not follow a zero-inflated Poisson distribution, and thus $\beta^* \neq \beta$. However, for the case considered in Theorem 5.1(a), the following theorem describes the relationship between β^* and β , which shows a scenario where conducting the naive analysis can still yield consistent estimators for some parameters.

Theorem 5.2 *If the conditions in Theorem 5.1(a) holds, then we have*

- (i) $\beta_{\phi 0}^* = \beta_{\phi 0}$ and $\beta_{\phi x}^* = \beta_{\phi x}$,
- (ii) $\beta_{\mu 0}^* = \beta_{\mu 0} + \log(1 - \pi_i)$,
- (iii) $\beta_{\mu x}^* = \beta_{\mu x}$.

The proof of Theorem 5.2 is presented in Appendix D.2. Theorem 5.2 says that when there is only leave-out error in model (5.5), within the frequentist framework, point estimators of the parameters for the response models (5.3) and (5.4) except for the intercept in (5.4) are still consistent if using the naive method by disregarding measurement error. Furthermore, Theorem 5.2(ii) implies that the estimator for $\beta_{\mu 0}$ obtained from the naive method can be adjusted by subtracting $\log(1 - \pi_i)$ to produce a consistent estimator. On the other hand, Theorem 5.2 shows that if π_i is unknown, nonidentifiability arises because $\beta_{\mu 0}$ and π_i cannot be separated when using the surrogate measurements Y_i^* together with the covariates X_i . However, this nonidentifiability issue can be circumvented if we conduct inferences in the Bayesian framework with a weakly informative prior imposed.

5.2 Bayesian Analysis Methodology

5.2.1 Bayesian Inference and Data Augmentation

Here we propose a Bayesian method for conducting inference about β by using the surrogate measurements Y_i^* , together with the covariates, where the effects of measurement error are accommodated.

Let $\beta = (\beta_{\phi 0}, \beta_{\phi x}^T, \beta_{\mu 0}, \beta_{\mu x}^T)^T$ and let $\theta = (\beta^T, \alpha^T)^T$. Inference about the parameter θ is based on the posterior distribution of θ given Y_i^* and X_i , given by

$$f(\theta|y_i^*, x_i) = \frac{f(y_i^*, \theta|x_i)}{f(y_i^*|x_i)} \propto f(y_i^*|x_i; \theta)\pi(\theta), \quad (5.13)$$

where $f(y_i^*, \theta|x_i)$ represents the joint distribution of Y_i and θ , $\pi(\theta)$ is the prior distribution of parameter θ , $f(y_i^*|x_i; \theta)$ is given by (5.9), and $f(y_i^*|x_i) = \int f(y_i^*|x_i; \theta)\pi(\theta)d\theta$. Then, the Bayes estimator of the parameters are given by the posterior mean $\hat{\theta} = E(\theta|Y_i^*, X_i)$.

The basic idea of implementing Bayesian estimation is to sample a sequence of parameters from their posterior distribution given by (5.13). Then the Bayes point estimator

$\widehat{\theta}$ is given by taking the sample mean of the sampled parameter sequence, and the $\gamma\%$ -credibility interval is given by $(q_{1-\gamma\%}, q_{\gamma\%})$, where $0 < \gamma < 1$, $q_{\gamma\%}$ is the $\gamma\%$ quantile of the sampled parameter sequence.

To this end, one may employ a sampling algorithm such as the Gibbs sampling method to sample a sequence of values from the posteriors distribution (5.13), which, however, can be challenging due to the complex structure of the probability mass function of Y_i in (5.2). To circumvent this, we consider an alternative way to express the distribution of Y_i by using two latent variables, say U_{i1} and U_{i2} , which are conditionally independent given X_i , and each follows a Poisson distribution. Rather than directly characterizing the distribution of Y_i by using (5.2) together with (5.3) and (5.4), we separately describe (5.3) and (5.4) each using U_{i1} and U_{i2} , respectively, to gain the flexibility in modeling of the distribution of Y_i .

To be specific, we assume that given X_i , U_{i1} and U_{i2} are conditionally independent, and that given X_i the conditional distributions of U_{i1} and U_{i2} are given by

$$\begin{aligned} U_{i1}|X_i &\sim \text{Poisson}(\mu_{i1}), \\ U_{i2}|X_i &\sim \text{Poisson}(\mu_i), \end{aligned}$$

where $\mu_{i1} = \exp(\beta_{\phi 0} + \beta_{\phi x}^T X_i)$ and $\mu_i = \exp(\beta_{\mu 0} + \beta_{\mu x}^T X_i)$ with $\beta_{\phi 0}$, $\beta_{\phi x}$, $\beta_{\mu 0}$ and $\beta_{\mu x}$ being the parameters in (5.3) and (5.4). Then (5.3) is equivalently written as $\phi_i = 1 - \exp(-\mu_{i1})$, which can be viewed as the probability $P(U_{i1} > 0|X_i)$. Therefore, the initial definition (5.1) for Y_i is equivalently expressed as

$$\begin{aligned} Y_i &= 0, \text{ with probability } P(U_{i1} = 0|X_i), \\ Y_i &= U_{i2}, \text{ with probability } P(U_{i1} > 0|X_i). \end{aligned} \tag{5.14}$$

In other words, the values of Y_i may be viewed by the distributions of U_{i1} and U_{i2} in such a way:

$$\begin{aligned} &\text{if } U_{i1} = 0, \text{ then we set } Y_i = 0; \\ &\text{if } U_{i1} > 0, \text{ then we set } Y_i = U_{i2}; \end{aligned}$$

and thus, we write $Y_i = 0 \cdot I(U_{i1} = 0) + U_{i2} \cdot I(U_{i1} > 0)$, which is

$$Y_i = U_{i2} I(U_{i1} > 0), \tag{5.15}$$

where $I(\cdot)$ is the indicator function.

Consequently, the original distribution (5.2) of Y_i together with (5.3) and (5.4) can now be equivalently described by using U_{i1} and U_{i2} via (5.15). Thereby, using the idea of

data augmentation (van Dyk and Meng, 2001), U_{i1} and U_{i2} can be used to ease sampling procedures directly based on (5.13), which is complicated to realize. In particular, rather than using $f(y_i^*|x_i; \theta)$ in (5.13) directly, we use (5.9) with Y_i replaced by (5.15) in its derivation, and sampling parameter values from (5.13) can be equivalently re-expressed as follows.

To see the idea, we consider the case with $c_+ = c_- = 1$ when using (5.5). First, fixing the initial parameter θ , we treat U_{i1} , U_{i2} , Z_{i+} and Z_{i-} as “missing data” and calculate their posterior distribution, $f(u_{i1}, u_{i2}, z_{i+}, z_{i-}|y_i^*, x_i; \theta)$, given $\{Y_i^*, X_i\}$ and θ , which is given by

$$\begin{aligned} & f(u_{i1}, u_{i2}, z_{i+}, z_{i-}|y_i^*, x_i; \theta) \\ &= P(U_{i1} = u_{i1}, U_{i2} = u_{i2}, Z_{i+} = y_i^* - I(u_{i1} > 0)u_{i2} + z_{i-}, Z_{i-} = z_{i-}|x_i; \theta) \\ &= P(U_{i1} = u_{i1}|x_i) P(U_{i2} = u_{i2}|x_i) P(Z_{i-} = z_{i-}|U_{i1} = u_{i1}, U_{i2} = u_{i2}, x_i) \\ & \quad \times P(Z_{i+} = y_i^* - I(u_{i1} > 0)u_{i2} + z_{i-}|x_i), \end{aligned} \quad (5.16)$$

where the first equality is due to (5.5) and (5.15), and in the second equality we use the conditional independence between U_{i1} and U_{i2} given X_i as well as (5.6).

Next, we re-express the posterior distribution (5.13) of θ by replacing Y_i with U_{i1} and U_{i2} and using the measurement error model (5.5):

$$\begin{aligned} & f(\theta|y_i^*, x_i, u_{i1}, u_{i2}, z_{i+}, z_{i-}) \\ & \propto f(u_{i1}, u_{i2}, z_{i+}, z_{i-}|y_i^*, x_i; \theta) \pi(\theta) \\ &= P(U_{i1} = u_{i1}|x_i; \beta_{\phi 0}, \beta_{\phi x}) P(U_{i2} = u_{i2}|x_i; \beta_{\mu 0}, \beta_{\mu x}) P(Z_{i-} = z_{i-}|u_{i1}, u_{i2}, x_i; \alpha_{-0}, \alpha_{-w}) \\ & \quad \times P(Z_{i+} = y_i^* - I(u_{i1} > 0)u_{i2} + z_{i-}|x_i; \alpha_{+0}, \alpha_{+w}) \pi(\theta), \end{aligned} \quad (5.17)$$

where the second step is due to (5.16) with the dependence on the parameters spelled out explicitly. The advantage of (5.17) lies in its separation of the components of θ by using distributions for different random variables, i.e., U_{i1} , U_{i2} , Z_{i-} and Z_{i+} . For example, given the rest parameters, the posterior distribution of parameters $\beta_{\phi 0}$ and $\beta_{\phi x}$ are simplified by (5.17) as

$$\begin{aligned} & f(\beta_{\phi 0}, \beta_{\phi x}|y_i^*, x_i, u_{i1}, u_{i2}, z_{i+}, z_{i-}; \beta_{\mu 0}, \beta_{\mu x}, \alpha_{-0}, \alpha_{-w}, \alpha_{+0}, \alpha_{+w}) \\ & \propto f(u_{i1}|x_i; \beta_{\phi 0}, \beta_{\phi x}) \pi(\theta). \end{aligned}$$

Therefore, sampling values of θ from (5.17) can be easily realized by sampling values for $(\beta_{\phi 0}, \beta_{\phi x}^T)^T$, $(\beta_{\mu 0}, \beta_{\mu x}^T)^T$, $(\alpha_{+0}, \alpha_{+w}^T)^T$ and $(\alpha_{-0}, \alpha_{-w}^T)^T$, separately from their posterior distribution $f(\beta_{\phi 0}, \beta_{\phi x}|u_{i1}, x_i)$, $f(\beta_{\mu 0}, \beta_{\mu x}|u_{i2}, x_i)$, $f(\alpha_{+0}, \alpha_{+w}|z_{i+}, x_i)$ and $f(\alpha_{-0}, \alpha_{-w}|z_{i-}, u_{i1}, u_{i2}, x_i)$.

5.2.2 Implementation: Monte Carlo Markov Chain Method with Data Augmentation

In this subsection, we describe the details of implementing the data augmentation idea described in Section 5.2.1 by using an MCMC algorithm, which is summarized as follows.

Step 1: (Data Augmentation) Generate U_{i1} , U_{i2} , Z_{i+} and Z_{i-} : For $i = 1, \dots, n$ and given Y_i^* , X_i and θ , generate U_{i1} , U_{i2} , Z_{i+} and Z_{i-} jointly from the distribution (5.16), which can be realized by the inversion sampling algorithm described in Appendix D.3.

Step 2: Update α_{+j} using (5.7): For $j = 0, \dots, p_+$, let α_{+j} be the j th element of α_{+w} , let $W_{i0+} = 1$ and let W_{ij+} be the j th element of W_{i+} . Given Z_{i+} obtained from Step 1, we generate α_{+j} from the posterior distribution

$$\begin{aligned} \prod_{i=1}^n f(\alpha_{+j}|z_{i+}) &\propto \prod_{i=1}^n f(z_{i+}|\alpha_{+j})\pi(\alpha_{+}) \\ &\propto \exp\left(\sum_{i=1}^n z_{i+}w_{ij+}\alpha_{+j}\right) \exp\left\{-\sum_{i=1}^n \exp(\alpha_{+}^T w_{i+})\right\} \pi(\alpha_{+}), \end{aligned} \quad (5.18)$$

where the second step comes from (5.7), and $\pi(\alpha_{+})$ is the prior distribution of α_{+} which may, for example, take a log-Gamma or a normal distribution with hyperparameters whose values are specified. When W_{ij+} is binary, then we take $\pi(\alpha_{+})$ to be a conjugate log-Gamma(c, d) prior with $\pi(\alpha_{+j}) \propto \exp\{-c \exp(\alpha_{+j})\} \exp(d\alpha_{+j})$, so that the posterior distribution (5.18) becomes

$$\begin{aligned} &\prod_{i=1}^n f(\alpha_{+j}|z_{i+}) \\ &\propto \exp\left(\sum_{i=1}^n z_{i+}w_{ij+}\alpha_{+j}\right) \exp\left\{-\sum_{i=1}^n \exp(\alpha_{+}^T w_{i+})\right\} \exp\{-c \exp(\alpha_{+j})\} \exp(d\alpha_{+j}) \\ &= \exp\left\{\left(d + \sum_{i=1}^n z_{i+}w_{ij+}\right)\alpha_{+j}\right\} \exp\left[-\left\{c + \sum_{i=1}^n \exp\left(\sum_{j^* \neq j} w_{ij^*+} \alpha_{+j^*}\right) w_{ij+}\right\} \exp(\alpha_{+j})\right], \end{aligned}$$

which is the log-Gamma distribution, log-Gamma($c + \sum_{i=1}^n \exp(\sum_{j^* \neq j} W_{ij^*+} \alpha_{+j^*}) W_{ij+}$, $d + \sum_{i=1}^n z_{i+} w_{ij+}$).

Step 3: Update α_{-j} : For $j = 0, \dots, p_-$, let α_{-j} be the j th element of α_{-w} , let $W_{i0-} = 1$, and let that W_{ij-} is the j th element of W_{i-} . We generate α_{-j} from the posterior distribution

$$\begin{aligned} \prod_{i=1}^n f(\alpha_{-j}|y_i, z_{i-}) &\propto \prod_{i=1}^n f(y_i, z_{i-}|\alpha_{-j})\pi(\alpha_{-j}) \\ &\propto \prod_{i=1}^n \binom{y_i}{z_{i-}} \{g^{-1}(\alpha_{-}^T w_{i-})\}^{z_{i-}} \{1 - g^{-1}(\alpha_{-}^T w_{i-})\}^{y_i - z_{i-}} \pi(\alpha_{-j}), \end{aligned}$$

where the second step comes from (5.8) and $\pi(\alpha_{-j})$ the probability density function of the prior of α_{-j} , which can be taken as a normal distribution.

Step 4: Update β : Since both U_{i1} and U_{i2} follow a conditional Poisson distribution, given X_i , we update them in the same way as in Step 2. Let $\beta_{\phi j}$ and $\beta_{\mu j}$ respectively be the j th element of β_{ϕ} and β_{μ} . Let $X_{i0} = 1$ and X_{ij} be the j th element of X_i . For $j = 0, \dots, p_x$, update $\beta_{\phi j}$ by sampling it from

$$\prod_{i=1}^n f(\beta_{\phi j}|u_{i1}) \propto \exp\left(\sum_{i=1}^n u_{i1} x_{ij} \beta_{\phi j}\right) \exp\left\{-\sum_{i=1}^n \exp(\beta_{\phi}^T x_i)\right\} \pi(\beta_{\phi}),$$

and update $\beta_{\mu j}$ by sampling it from

$$\prod_{i=1}^n f(\beta_{\mu j}|u_{i2}) \propto \exp\left(\sum_{i=1}^n u_{i2} x_{ij} \beta_{\mu j}\right) \exp\left\{-\sum_{i=1}^n \exp(\beta_{\mu}^T x_i)\right\} \pi(\beta_{\mu}).$$

where $\pi(\beta_{\phi})$ and $\pi(\beta_{\mu})$ are prior distributions for β_{ϕ} and β_{μ} , respectively. For instance, if the covariates X_i are binary, we may take the conjugate log-Gamma prior for β_{ϕ} and β_{μ} .

5.3 Extension to the Main/Validation Studies

The Bayesian inference circumvents the traditional identifiability issue in the frequentist framework (e.g. [Gelman et al., 2013](#), Page 412) by using weakly informative priors. In some applications, however, even weakly informative priors are not available or cannot be precisely set. In this circumstance, the study design can provide extra information regarding the measurement error process through validation data.

Let \mathcal{M} denote the index set of the subjects in the main study, where $\{(y_i^*, x_i) : i \in \mathcal{M}\}$ is available. Let \mathcal{V} represent the index set of the subjects in the validation data. For internal validation, the validation data contain $\{(y_i^*, y_i, x_i) : i \in \mathcal{V}\}$ with $\mathcal{V} \subset \mathcal{M}$; for external validation, the validation data contain $\{(y_i^*, y_i, w_{i+}, w_{i-}) : i \in \mathcal{V}\}$ with $\mathcal{M} \cap \mathcal{V} = \emptyset$. Let m denote the size of the validation subsample \mathcal{V} and let n be the size of \mathcal{M} as used in Sections 5.1–5.2.

5.3.1 Main/External Validation Study

With external validation data, we write the posterior function of θ combining the main and validation data:

$$\begin{aligned} & \left\{ \prod_{i \in \mathcal{M}} f(\theta | y_i^*, x_i) \right\} \left\{ \prod_{i \in \mathcal{V}} f(\theta | y_i^*, y_i, w_{i+}, w_{i-}) \right\} \\ & \propto \pi(\theta) \left\{ \prod_{i \in \mathcal{M}} f(y_i^* | x_i; \theta) \right\} \left\{ \prod_{i \in \mathcal{V}} f(y_i^* | y_i, w_{i+}, w_{i-}; \alpha) \right\}, \end{aligned} \quad (5.19)$$

where $f(y_i^* | x_i; \theta)$ comes from (5.9), $f(y_i^* | y_i, w_{i+}, w_{i-}; \alpha)$ is modeled by (5.5), and $\pi(\theta)$ is a prior function.

Similar to the development of Section 5.2, instead of directly using (5.19) for sampling values of the parameters, we apply the following sampling procedures:

Step 1: (Data Augmentation) Generate U_{i1} , U_{i2} , Z_{i+} and Z_{i-} . For $i \in \mathcal{M}$, we generate augmented data in the same way as in Section 5.2.2. For $i \in \mathcal{V}$, we generate Z_{i+} and Z_{i-} from their joint posterior distribution

$$f(z_{i+}, z_{i-} | Y_i^* = y_i^*, Y_i = y_i; \alpha, \beta) = f(Z_{i+} = y_i^* - y_i + z_{i-}, Z_{i-} = z_{i-}; \alpha, \beta), \quad (5.20)$$

which is determined by (5.5).

Steps 2-3: Update α_{+j} and α_{-j} . These two steps are similar to Steps 2–3 in Section 5.2.2 except for replacing the summation $\sum_{i=1}^n$ with $\sum_{i \in \mathcal{M} \cup \mathcal{V}}$. For example, we update α_{+j} by sampling it from the posterior

$$\prod_{i \in \mathcal{M} \cup \mathcal{V}} f(\alpha_{+j} | z_{i+}) \propto \exp \left(\sum_{i \in \mathcal{M} \cup \mathcal{V}} z_{i+} w_{ij} \alpha_{+j} \right) \exp \left\{ - \sum_{i \in \mathcal{M} \cup \mathcal{V}} \exp(\alpha_{+j}^T w_{i+}) \right\} \pi(\alpha_{+j}),$$

Step 4: This is identical to Step 4 in Section 5.2.2.

5.3.2 Main/Internal Validation Study

When internal validation data is available, the posterior function of parameter θ now becomes

$$\begin{aligned} & \left\{ \prod_{i \in \mathcal{M} \setminus \mathcal{V}} f(\theta | y_i^*, x_i) \right\} \left\{ \prod_{i \in \mathcal{V}} f(\theta | y_i^*, y_i, x_i) \right\} \\ & \propto \pi(\theta) \left\{ \prod_{i \in \mathcal{M} \setminus \mathcal{V}} f(y_i^* | x_i; \theta) \right\} \left[\prod_{i \in \mathcal{V}} \{f(y_i^* | y_i; \alpha) f(y_i | x_i; \beta)\} \right], \end{aligned} \quad (5.21)$$

where $f(y_i^* | x_i; \theta)$ comes from (5.9), $f(y_i | x_i; \beta)$ is from (5.2), $f(y_i^* | y_i; \alpha)$ is from (5.5), and $\pi(\theta)$ is the prior function of parameters θ . Similar to the development of Section 5.2, instead of directly using (5.21) for sampling values of the parameters, we apply the following sampling procedures:

Step 1: (Data Augmentation) Generate U_{i1} , U_{i2} , Z_{i+} and Z_{i-} . For $i \in \mathcal{M} \setminus \mathcal{V}$, we generate the augmented data in a way similar to that in Section 5.2.2. For $i \in \mathcal{V}$, we generate the variables in the following steps iteratively.

1. We update the latent data augmentation variable U_{i1} according to the value of U_{i2} and Y_i , which includes three circumstances according to (5.15):
 - (i). Case 1 with $Y_i = 0$ and $U_{i2} = 0$: By (5.15), $Y_i = 0$ if and only if $U_{i1} = 0$ or $U_{i2} = 0$. That is, there is no restrictions on the U_{i1} when $U_{i2} = 0$, so in this case U_{i1} is generated from $\text{Poisson}(\mu_{i1})$.
 - (ii). Case 2 with $Y_i = 0$ and $U_{i2} \neq 0$: (5.15) says that U_{i1} is surely be 0. Hence, we set $U_{i1} = 0$.
 - (iii). Case 3 with $Y_i > 0$: (5.15) implies that $U_{i1} > 0$. Hence, we update U_{i1} by a truncated $\text{Poisson}(\mu_{i1})$ at 0.
2. Given U_{i1} obtained in part 1, we update the latent variable U_{i2} , which includes two cases:
 - (i). Case 1 with $U_{i1} > 0$: we generate U_{i2} by setting it equal to Y_{i1} by (5.15).
 - (ii). Case 2 with $U_{i1} = 0$: (5.15) shows that $Y_{ij} = 0$, and thus, there are no constraints on the variable U_{i2} and we generate U_{i2} from $\text{Poisson}(\mu_i)$.
3. Generate Z_{i+} and Z_{i-} in the same way as in Step 1 in Section 5.3.1.

Steps 2–4: The steps are the same as Steps 2–4 in Section 5.2.2.

5.4 Application to Prostate Adenocarcinoma Genomics Data

5.4.1 Study Background

Here we apply the proposed methods to a multi-center molecular prostate cancer study. We are interested in predicting whether or not cancer-related pathways are activated during the prostate cancer progression and how the number of genes with copy number variations (CNVs) within each pathway is associated with the risk factors. The data contain two datasets that are linked by the genes in The Cancer Genome Atlas (TCGA) data that are annotated in the Kyoto Encyclopedia of Genes and Genomes (KEGG) pathways data through website cBioPortal. The first part includes the pathway information arising from the KEGG pathways data, and the second part is the putative CNV data with 465 subjects collected from two sources with 185 subjects from Broad Institute ([Banerji et al., 2012](#)) and 280 subjects from Memorial Sloan-Kettering Cancer Center (MSKCC) ([Taylor et al., 2010](#)) for prostate adenocarcinoma.

In this analysis, similar to [Neelon and Chung \(2017\)](#), we consider four pathways: mitogen-activated protein kinase (MAPK) signaling, cytokine-cytokine receptor (CCR) interaction, endocytosis (EC), and P53. Genes in the MAPK pathway are related to various cellular functions, such as cell proliferation, differentiation, and migration; genes in the CCR interaction pathway are associated with inflammatory host defenses, cell growth, differentiation and death, and the restoration of homeostasis; the genes in the EC pathway are related to the mechanisms of cells transporting ligands, nutrients, proteins, and lipids from the cell surface to the cell interior; and the p53 pathway is induced by a number of stress signals, including DNA damage, oxidative stress, and activated oncogenes ([Alberts et al., 2002](#)).

In our study here, we conduct four marginal analysis separately for each pathway, where the response for each individual (Y_i) is defined as the count of genes with significant CNVs (with reading valued as either -2 or 2), which reflects the level of mutation in the individual. We implement the Vuong tests ([Vuong, 1989](#)) to assess whether or not zero-inflation exists in the response. With a p-value smaller than 0.001 for all four pathways, the test result shows a strong sign of zero inflation. We investigate two different risk factors that may be associated with the CNVs counts in two separate studies, which are reported in Sections [5.4.2](#) and [5.4.3](#), respectively. In the first study, the covariate is denoted as X_{i1} , which is taken as the tumor stage, which is given by an indicator variable, taking value 0 or 1, according to the T2 or T3+ tumor stage for subject i ; and in the second study, the

covariate is denoted as X_{i2} , which represents the cancer recurrence, with $X_i = 1$ if cancer recurrence occurs and $X_i = 0$ otherwise.

We are interested in understanding the relationship between Y_i and a covariate in each study. However, due to the potential sequencing error, the CNV reading of insignificant gene can be falsely measured as significant, whereas the gene with significant CNVs can be missed to be counted, and thus, the observed count number (denoted as Y_i^*) may considerably differ from the true value of Y_i . To feature this difference, we consider the measurement error model (5.5) with $c_+ = c_- = 1$.

5.4.2 Association of Tumor Stage and CNVs

We conduct analysis for each pathway separately using the zero-inflated model (5.2), (5.3) and (5.4) to feature the dependence of Y_i on the covariate X_{i1} . The dataset is combined from multiple sources, and the data quality and genetic sequencing protocols can be different. Thus, in this study, we perceive that the measurement error process is associated with the data source and use the measurement error models (5.7) and (5.8), where the covariate W_i is a binary indicator for the data source, with $W_i = 1$ if the subject i is from the broad institute, and 0 otherwise.

In implementing the Bayesian procedures described in Section 5.2, we consider an uninformative prior, log-Gamma(1000, 0.001), for the parameters of models (5.3), (5.4) and (5.7). For the parameter α_{-0} in the model (5.8), we consider the prior, Normal(-2, 10), where the negative mean reflects our expectation of a negative value for α_{-0} , and a large variance shows a flat prior. We use the Gelman-Rubin method (Gelman et al., 1992) to diagnose the convergence of Monte Carlo Markov chains, and the results show that MCMC series for all the parameters well converge after running 250,000 iterations of sampling steps and discarding the first 5000 as burn-in.

Our first interest lies in whether CNVs counts change as the tumor progresses for patients who have activated the pathway. We implement the proposed method described in Section 5.2.1, and for comparison, we also implement the naive method based on Neelon and Chung (2017) where the difference of Y_i and Y_i^* is neglected. The analysis results of parameter estimation are presented in Figure 5.1. Both the naive method and the proposed method find that the $\beta_{\mu x}$ for all pathways are not significantly different from zero, suggesting that patients in tumor stage T3+ do not have different mutations than those in T2 stage. The estimates of the intercept of the count model (5.4), $\beta_{\mu 0}$, for the MAPK, CCR and EC pathways are higher than those of $\beta_{\mu 0}$ in the P53 pathway, showing

that prostate cancer patients have more mutations in the genes involved in the pathways of MAPK, CCR, and EC than the P53 pathway.

Our second interest is the probability of activation for patients as the tumor grows, which is reflected by the estimation of parameters associated with ϕ_i . It is clear that the proposed method with the measurement error effects accounted for yields results different from the naive method. In Figure 5.2, we report credible intervals for the probability of pathway activation using (5.3) together with estimates of $\beta_{\phi 0}$ and $\beta_{\phi x}$. The proposed method indicates that the difference in the probability of pathway activation is close to zero for patients in tumor stage T2 versus the patients in T3+. However, the naive method suggests that the difference is very large.

5.4.3 Association of Cancer Recurrence and CNVs

We consider the endpoint to be, alternatively, the recurrence status of a prostate cancer patient after being cured. We are interested in: 1) whether the status of cancer recurrence is associated with activation of the pathway; and 2) for the subjects who have activated the pathway, whether the status of cancer recurrence is associated with the CNVs counts. We conduct an analysis for each pathway separately using (5.2), (5.3) and (5.4) to feature the dependence of Y_i on the covariate X_{i2} . Since the covariate of cancer recurrence information is only available in the MSKCC study, we focus on the analysis of 280 subjects in the MSKCC study and consider measurement error models (5.7) and (5.8) with constant parameters α_{0+} and α_{0-} only, where W_i is no longer included in the models.

In implementing the Bayesian procedures described in Section 5.2, we consider the same priors for parameters as in Section 5.4.2. We run 250,000 iterations of the sampling steps and discard the first 5000 as burn-in. The resultant Monte Carlo Markov chains converge according to the Gelman-Rubin method (Gelman et al., 1992).

The results are exhibited in Figures 5.3–5.4. First, using the proposed methods in Section 5.2 in contrast to the naive method as described in Neelon and Chung (2017), we study the association between the status of recurrence of prostate cancer and the number of CNVs for patients with activated pathways. In Figure 5.3, the proposed method suggests that for all the pathways, under the significance level of 0.05, the number of CNVs is not significantly associated with a higher risk of cancer recurrence, where the estimate of $\beta_{\mu x}$ is, respectively, 0.007 and the credible interval $(-0.393, 0.401)$ for MAPK, 0.294 and the credible interval $(-0.098, 0.614)$ for CCR, 0.144 and the credible interval $(-0.346, 0.655)$ for EC, and 0.001 and the credible interval $(-1.152, 1.116)$ for P53. On the other hand, the naive method shows that the patients with higher CNVs in the CCR pathway has a

higher risk of recurrence, with the estimate of $\beta_{\mu x}$ being 0.454 and the credible interval (0.106, 0.802).

Secondly, we study the association between the activation of pathways and the risk of cancer recurrence. We observe that the patients with the MAPK, CCR, or EC pathway activated tend to have a higher risk of the prostate cancer recurrence, where the estimate of $\beta_{\phi x}$ is, respectively, 0.906 and the credible interval (0.061, 1.737), 1.021 and the credible interval (0.145, 1.999), and 1.205 and the credible interval (0.311, 2.103) for each pathway. On the other hand, the naive method indicates that the activation of the pathway is associated with a lower cancer risk because the estimates of $\beta_{\phi x}$ are negative. We estimate the probability of pathway activation using (5.3) and present the results in Figure 5.4. The proposed method generally suggests that the cancer patients have low probabilities of activation of the pathway, while the naive method indicates opposite findings.

5.5 Simulation Studies

In this section, we conduct simulation studies to evaluate the performance of the proposed method. For the sake of comparison, we also implement the naive method where no action is taken to deal with the measurement error in response.

We conduct two simulation studies. In the first simulation study, we evaluate the performance of the proposed method under different settings of the parameters, leading to different percentages of zeros in the responses. We conduct sensitivity analyses by exploring different settings of the prior distribution of the parameters. In the second simulation study, we evaluate the performance of the methods under different degrees of measurement error.

For both simulation studies, we run 1000 simulations for each setting. The sample size is taken as $n = 5000$, and we consider model (5.3) with covariates X_{i1} and (5.4) with covariates X_{i2} , where covariate X_{i1} is generated from Binomial(0.5) and X_{i2} is independently generated from Uniform[0, 1]. The true response Y_i is generated from (5.2), where $\phi_i = 1 - \exp\{-\exp(\beta_{\phi 0} + \beta_{\phi x} X_{i1})\}$ and $\mu_i = \beta_{\mu 0} + \beta_{\mu x} X_{i2}$.

To generate surrogate measurements Y_i^* of Y_i , we consider the measurement error models (5.7) and (5.8) each associated with a covariate W_{i1} and W_{i2} , respectively, where W_{i1} is independently generated from $U[0, 1]$, and W_{i2} is independently generated from $U[0, 2]$. Furthermore, we generate Z_{i+} from exponential($\alpha_{+0} + \alpha_{+w} W_{i1}$) and Z_{i-} from Binomial($Y_i, \frac{\exp(\alpha_{-0} + \alpha_{-w} W_{i2})}{1 + \exp(\alpha_{-0} + \alpha_{-w} W_{i2})}$). As a result, Y_i^* is determined by $Y_i^* = Y_i + Z_{i+} - Z_{i-}$.

To summarize the simulation results, we report biases (denoted ‘‘Bias’’) by calculating $\frac{1}{N} \sum_{i=1}^N \hat{\theta}_k - \theta_k$, model-based standard errors (denoted ‘‘SEM’’), empirical standard error

of the point estimates (denoted “SEE”), and the coverage rate (in percent) of 95% credible intervals for a parameter, say θ_k , (denoted “CR”), defined as

$$\frac{1}{N} \sum_{i=1}^N I(\hat{\theta}_k^{(L)} < \theta_k < \hat{\theta}_k^{(U)}),$$

where N is the number of simulations, $\hat{\theta}_k^{(L)}$ and $\hat{\theta}_k^{(U)}$ are respectively the 2.5% and 97.5% quantile for the sampled parameter values.

5.5.1 Simulation 1: Performance of the Proposed Method with Different Zero Percentages and Hyperparameters

Two parameter settings are considered. In Setting 1, we consider $(\beta_{\phi 0}, \beta_{\phi x}, \beta_{\mu 0}, \beta_{\mu x})^T = (-0.7, 0.7, 1, -0.5)^T$, yielding about 60% zeros; in Setting 2, we consider $(\beta_{\phi 0}, \beta_{\phi x}, \beta_{\mu 0}, \beta_{\mu x})^T = (-0.2, 0.7, 1, 0.5)^T$, yielding about 30% zeros. In both settings, we consider the parameters of the measurement error model (5.5) with $\alpha_{+0} = \alpha_{-0} = 0$, and $(\alpha_{+w}, \alpha_{-w})^T = (0.5, -2.3)^T$. For each setting, we study the sensitivity of results with respect to different priors when implementing the proposed method and the naive method which disregards the difference in Y_i and Y_i^* . In the first set of priors, we consider uninformative priors log-Gamma(1000, 0.001) for $\beta_{\phi 0}$, $\beta_{\phi x}$, $\beta_{\mu 0}$, $\beta_{\mu x}$, α_{+w} , and Normal(0, 1000²) for α_{-w} . In the second set of priors, we choose log-Gamma(1, 1) for $\beta_{\phi 0}$, $\beta_{\phi x}$, $\beta_{\mu 0}$, $\beta_{\mu x}$, α_{+w} , and Normal(-2, 2²) for α_{-w} .

Table 5.1 shows that without accounting for measurement error in response, the naive model produces biased estimates of the parameters and meaningless coverage rates of 95% credible intervals. On the other hand, the proposed method considerably reduces the biases resulting from the measurement error effects and provides reasonable standard errors. The performance of the proposed method is satisfactory for different settings, regardless of the specification of the prior distribution.

5.5.2 Simulation 2: Performance of Method with Different Degrees of Measurement Error

In this subsection, we evaluate how the performance of the proposed method may be affected by different degrees of measurement error resulting from different parameters in the add-in process and the leave-out process. For the add-in process, we set the parameters

$(\alpha_{+0}, \alpha_{+w})^T$ in (5.7) to be $(-1, 0.6)^T$ or $(2, -1.2)^T$, leading to the mean of Z_{i+} to be 0.5 (small) or 5.0 (substantial). For the leave-out process, we take the parameters $(\alpha_{-0}, \alpha_{-w})^T$ in (5.8) to be $(-1, -1.2)^T$ or $(-0.8, -1)^T$, respectively, leading to 5% (low) or 10% (high) counts being neglected.

To create a dataset for a main/external validation study, we randomly choose 2500 subjects and only keep the variables of Y_i^* , Y_i , W_{i1} and W_{i2} . For the case of a main/internal validation study, we randomly select 2500 subjects and keep their variables of Y_i^* , Y_i , X_{i1} , X_{i2} , W_{i1} and W_{i2} to serve as the validation data.

We implement the methods described in Sections 5.2 and 5.3, as opposed to the naive method by replacing Y_i with Y_i^* in the analysis. When implementing the methods, we take log-Gamma(1, 1) as the prior for $\beta_{\phi 0}$, $\beta_{\phi x}$, $\beta_{\mu 0}$, $\beta_{\mu x}$, α_{+0} , and Normal($-2, 2^2$) for α_{-0} and α_{-w} .

The results are displayed in Table 5.2. Our proposed methods outperform the naive method, regardless of the parameter settings for the measurement error model. The proposed methods yield small finite sample biases for the point estimates and reasonable coverage rates for 95% credible intervals.

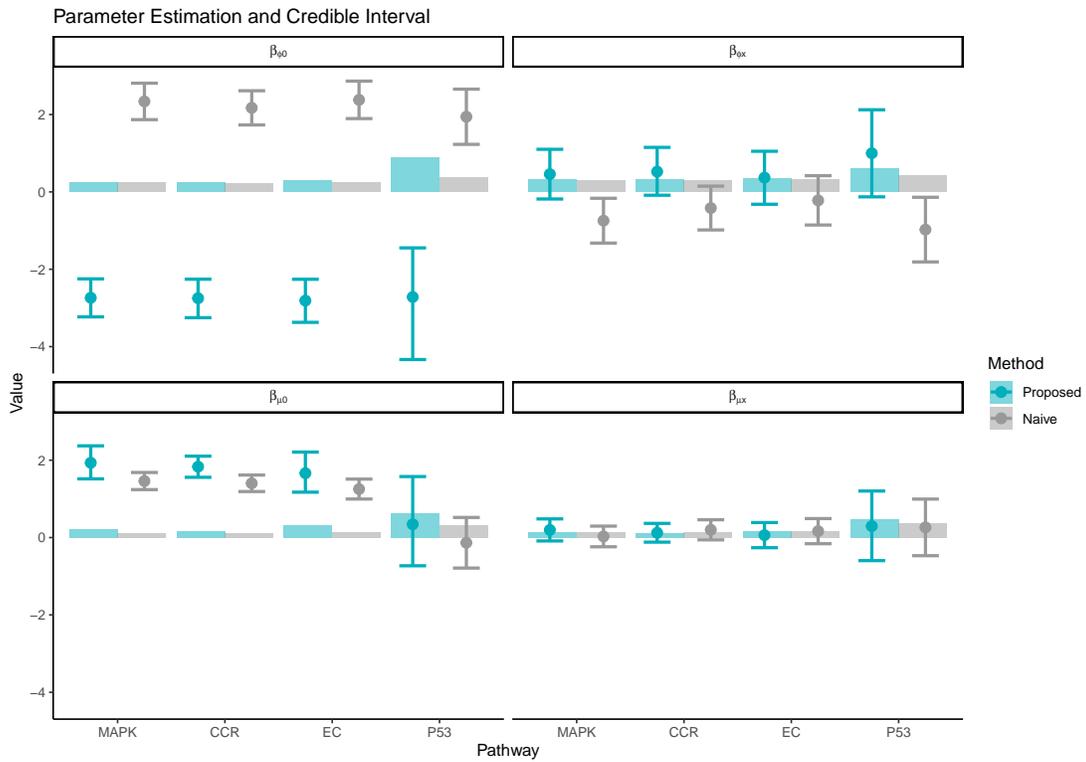


Figure 5.1: The plot of parameter estimation of the zero-inflated Poisson models for the association between tumor stage and CNVs. The point indicates the point estimation and the line segments represent the 95% credible interval. The barplot is the associated standard error for the estimation of the corresponding parameter.

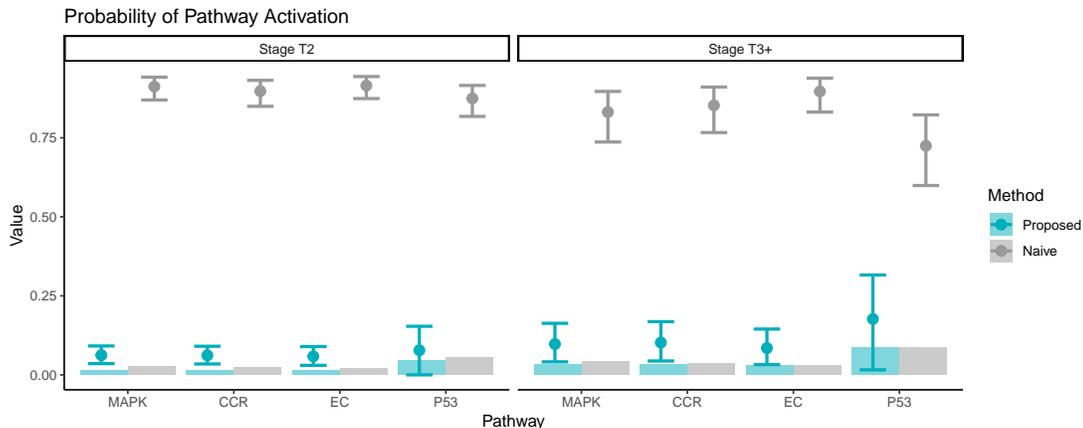


Figure 5.2: The plot of probability of pathway activation in the study of the association between tumor stage and CNVs. The point indicates the point estimation and the line segments represent the 95% credible interval. The barplot is the associated standard error for the estimation of the corresponding parameter.

Table 5.1: Results for Simulation 1 with different zero percentage and prior parameters

Parameter	Prior	Naive Method				Proposed Method			
		Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%
Setting 1: zero-percentage 60%									
$\beta_{\phi 0}$	Uninformative	1.506	0.030	0.029	0.0	0.004	0.089	0.086	93.3
$\beta_{\phi x}$		0.340	0.051	0.049	0.0	0.004	0.082	0.079	94.1
$\beta_{\mu 0}$		0.122	0.020	0.017	0.0	0.030	0.069	0.068	91.9
$\beta_{\mu x}$		0.816	0.032	0.028	0.0	0.004	0.082	0.081	94.5
α_{+w}		-	-	-	-	0.001	0.044	0.045	94.7
α_{-w}		-	-	-	-	0.115	0.562	0.560	90.5
$\beta_{\phi 0}$	Informative	1.506	0.030	0.029	0.0	0.007	0.088	0.085	93.4
$\beta_{\phi x}$		0.340	0.051	0.049	0.0	0.005	0.081	0.078	94.2
$\beta_{\mu 0}$		0.122	0.020	0.017	0.0	0.002	0.070	0.064	92.7
$\beta_{\mu x}$		0.817	0.032	0.028	0.0	0.004	0.082	0.080	94.3
α_{+w}		-	-	-	-	0.005	0.044	0.045	95.2
α_{-w}		-	-	-	-	0.243	0.733	0.648	91.5
Setting 2: zero-percentage 30%									
$\beta_{\phi 0}$	Uninformative	1.001	0.030	0.029	0.0	0.001	0.049	0.048	93.9
$\beta_{\phi x}$		0.332	0.052	0.049	0.2	0.002	0.056	0.055	94.5
$\beta_{\mu 0}$		0.124	0.020	0.017	0.0	0.021	0.046	0.052	93.7
$\beta_{\mu x}$		0.185	0.033	0.028	0.0	0.004	0.047	0.047	94.5
α_{+w}		-	-	-	-	0.004	0.053	0.053	94.7
α_{-w}		-	-	-	-	0.100	0.370	0.430	94.0
$\beta_{\phi 0}$	Informative	1.000	0.030	0.029	0.0	0.002	0.049	0.048	93.5
$\beta_{\phi x}$		0.332	0.051	0.049	0.0	0.003	0.056	0.055	95.7
$\beta_{\mu 0}$		0.124	0.020	0.017	0.0	0.012	0.043	0.052	96.6
$\beta_{\mu x}$		0.184	0.033	0.028	0.0	0.003	0.047	0.047	94.2
α_{+w}		-	-	-	-	0.006	0.053	0.053	94.5
α_{-w}		-	-	-	-	0.017	0.389	0.456	96.0

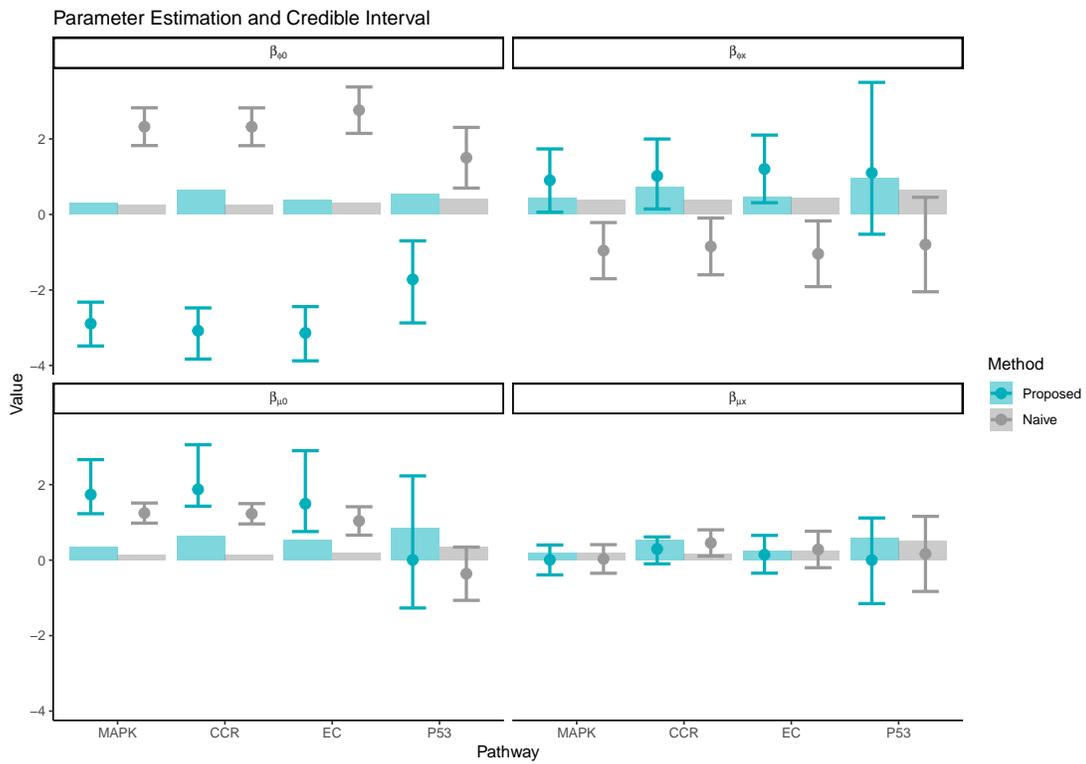


Figure 5.3: The plot of parameter estimation of the zero-inflated Poisson models for the association between cancer recurrence and CNVs. The point indicates the point estimation and the line segments represent the 95% credible interval. The barplot is the associated standard error for the estimation of the corresponding parameter.

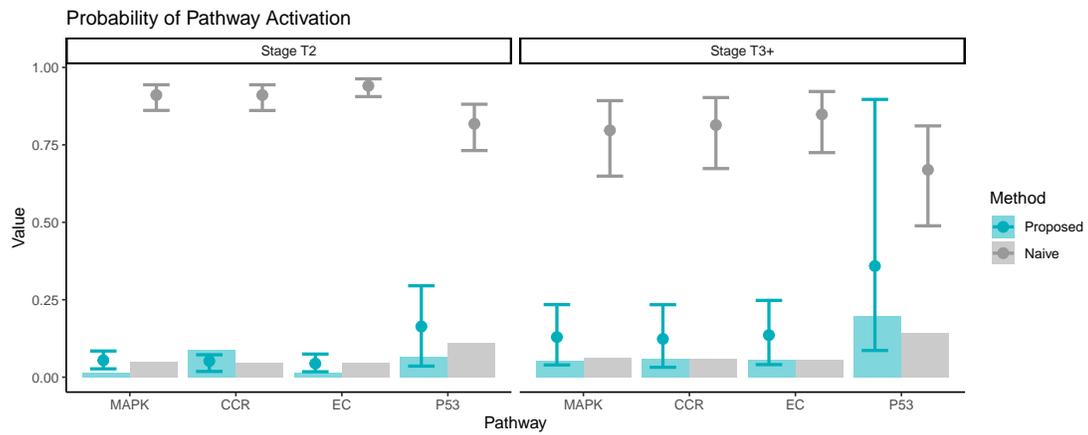


Figure 5.4: *The plot of probability of pathway activation in the study of the association between cancer recurrence and CNVs. The point indicates the point estimation and the line segments represent the 95% credible interval. The barplot is the associated standard error for the estimation of the corresponding parameter.*

Table 5.2: Results for Simulation 2 with different degrees of measurement error

Parameter	Naive Method						Proposed Method													
	No Validation			External Validation			No Validation			External Validation										
	α_{+0}	α_{+w}	α_{-0}	α_{-w}	Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%	Bias	SEE	SEM	CR%				
β_{ϕ_0}					0.523	0.027	0.027	0.0	0.005	0.043	0.043	93.9	0.001	0.051	0.051	93.8	0.001	0.030	0.031	96.0
β_{ϕ_x}					0.201	0.041	0.041	0.2	0.003	0.047	0.048	95.2	0.009	0.064	0.067	94.9	0.005	0.039	0.043	96.0
β_{μ_0}					0.054	0.022	0.020	23.3	0.028	0.033	0.041	94.4	0.003	0.036	0.037	95.6	0.000	0.023	0.023	94.2
β_{μ_x}					0.081	0.036	0.032	30.4	0.005	0.040	0.041	95.2	0.010	0.055	0.057	95.6	0.002	0.036	0.036	94.3
α_{+0}	-1.0	0.6	-1.0	-1.2	-	-	-	-	0.016	0.114	0.117	94.7	0.002	0.058	0.062	95.7	0.002	0.057	0.062	95.4
α_{+w}					-	-	-	-	0.011	0.157	0.161	94.5	0.004	0.092	0.097	95.9	0.004	0.093	0.097	95.7
α_{-0}					-	-	-	-	0.087	0.178	0.191	95.6	0.004	0.079	0.079	95.1	0.004	0.078	0.079	94.2
α_{-w}					-	-	-	-	0.466	0.292	0.481	94.9	0.005	0.091	0.092	95.1	0.005	0.092	0.091	95.3
β_{ϕ_0}					0.528	0.027	0.027	0.0	0.005	0.044	0.044	94.0	0.002	0.052	0.052	94.1	0.001	0.030	0.032	95.9
β_{ϕ_x}					0.200	0.041	0.042	0.2	0.003	0.049	0.048	94.6	0.010	0.064	0.068	95.3	0.005	0.039	0.043	96.8
β_{μ_0}					0.091	0.022	0.020	0.5	0.049	0.036	0.045	86.9	0.002	0.037	0.039	95.3	0.000	0.023	0.023	94.7
β_{μ_x}					0.084	0.037	0.033	29.2	0.005	0.042	0.042	95.0	0.009	0.058	0.059	94.4	0.002	0.037	0.037	94.1
α_{+0}	-1.0	0.6	-0.8	-1.0	-	-	-	-	0.012	0.120	0.120	93.4	0.000	0.060	0.063	95.9	0.000	0.060	0.063	95.6
α_{+w}					-	-	-	-	0.010	0.161	0.161	94.0	0.002	0.096	0.098	95.8	0.002	0.096	0.099	96.0
α_{-0}					-	-	-	-	0.126	0.158	0.175	91.9	0.004	0.070	0.072	94.6	0.002	0.070	0.071	94.6
α_{-w}					-	-	-	-	0.486	0.259	0.414	88.8	0.004	0.074	0.076	95.3	0.004	0.074	0.076	95.1
β_{ϕ_0}					1.676	0.044	0.044	0.0	0.054	0.090	0.097	94.1	0.007	0.081	0.084	94.9	0.001	0.033	0.036	96.7
β_{ϕ_x}					0.494	0.101	0.086	2.3	0.004	0.075	0.073	93.3	0.016	0.093	0.098	95.1	0.006	0.045	0.048	95.1
β_{μ_0}					0.785	0.014	0.011	0.0	0.053	0.056	0.072	97.3	0.001	0.055	0.059	95.9	0.001	0.026	0.026	95.1
β_{μ_x}					0.334	0.025	0.019	0.0	0.001	0.064	0.068	96.3	0.021	0.078	0.083	95.5	0.003	0.041	0.041	95.1
α_{+0}	2.0	-1.2	-1.0	-1.2	-	-	-	-	0.010	0.021	0.022	93.2	0.001	0.013	0.014	96.4	0.000	0.013	0.014	96.5
α_{+w}					-	-	-	-	0.021	0.044	0.047	93.7	0.001	0.028	0.030	95.6	0.001	0.028	0.029	95.1
α_{-0}					-	-	-	-	0.110	0.280	0.309	96.4	0.008	0.149	0.151	94.7	0.006	0.149	0.149	95.3
α_{-w}					-	-	-	-	0.699	0.458	0.740	96.4	0.046	0.207	0.208	93.3	0.038	0.206	0.205	94.6
β_{ϕ_0}					1.671	0.043	0.044	0.0	0.056	0.096	0.104	94.9	0.011	0.083	0.087	95.7	0.001	0.034	0.036	96.2
β_{ϕ_x}					0.490	0.100	0.085	2.8	0.004	0.077	0.076	94.4	0.018	0.096	0.101	95.1	0.006	0.046	0.048	95.4
β_{μ_0}					0.772	0.014	0.011	0.0	0.076	0.058	0.079	95.9	0.000	0.056	0.062	96.3	0.001	0.026	0.026	94.6
β_{μ_x}					0.338	0.025	0.019	0.0	0.002	0.067	0.071	96.2	0.020	0.082	0.086	95.0	0.003	0.042	0.041	95.2
α_{+0}	2.0	-1.2	-0.8	-1.0	-	-	-	-	0.010	0.020	0.022	93.9	0.001	0.014	0.014	95.9	0.000	0.013	0.014	96.3
α_{+w}					-	-	-	-	0.020	0.044	0.048	93.8	0.000	0.028	0.030	95.6	0.001	0.028	0.030	95.5
α_{-0}					-	-	-	-	0.149	0.261	0.287	94.7	0.008	0.132	0.132	94.4	0.004	0.130	0.131	94.7
α_{-w}					-	-	-	-	0.690	0.367	0.608	92.8	0.026	0.155	0.157	94.5	0.022	0.155	0.155	94.2

Chapter 6

Autoregressive Models with Data Subject to Measurement Error

In this chapter, we discuss error-contaminated time series data. The notation and the setup for the autoregressive time series model and the proposed measurement error models are introduced in Section 6.1. In Section 6.2, we present the theoretical results for characterizing the impact of measurement error on the analysis of time series data. In Section 6.3, we develop an estimating equation approach to adjust for the biases due to measurement error. In Section 6.4, we implement the proposed method to analyze the COVID-19 data in four provinces in Canada.

6.1 Model Setup and Framework

6.1.1 Time Series Model

Consider a $T \times 1$ vector of time series, $X^{(T)} = (X_1, X_2, \dots, X_T)^\top$. We are interested in modeling the dependence of X_t on its previous observations $X^{(t-1)}$ and we consider it to be postulated by an autoregressive model with lag p

$$X_t = \phi_0 + \sum_{j=1}^p \phi_j X_{t-j} + \epsilon_t, \quad (6.1)$$

where p is an integer smaller than T , $\epsilon^{(t)} = (\epsilon_1, \dots, \epsilon_t)^\top$ is independent of $X^{(t)} = (X_1, \dots, X_t)^\top$ with each ϵ_t having zero mean and variance σ_ϵ^2 , ϕ_0 is a constant drift, and $\phi = (\phi_1, \dots, \phi_p)^\top$ is the regression coefficient.

The additive form in (6.1) and the zero mean assumption of ϵ_t show that ϕ_0 and ϕ are constrained by

$$\phi_0 = E(X_t) - \{E(\tilde{X}_{t-1})\}^T \phi, \quad (6.2)$$

where $\tilde{X}_{t-1} = (X_{t-1}, \dots, X_{t-p})^T$. To make the process of X_t stationary, ϕ_1, \dots, ϕ_p are further constrained such that all the roots of the equation in z

$$z^p - \phi_1 z^{p-1} - \dots - \phi_p = 0$$

have absolute values smaller than 1 (Brockwell and Davis, 2002, Section 3.1.). For example, a stationary AR(1) process requires that $|\phi_1| < 1$, and a stationary AR(2) process needs that $(\phi_1 + \phi_2) < 1$, $(\phi_2 - \phi_1) < 1$ and $|\phi_2| < 1$. Here we are interested in the estimation of parameters, ϕ and ϕ_0 . Let μ denote the mean $E(X_t)$ of the time series, which equals $\frac{\phi_0}{1 - \phi_1 - \dots - \phi_p}$ if X_t is (weakly) stationary. When $p = 1$, the stationarity of a time series implies $\text{Var}(X_t) = \frac{\sigma_\epsilon^2}{1 - \phi^2}$ for $t = 1, \dots, T$.

6.1.2 Estimation of Model Parameters

The estimation of the parameters in the AR(p) time series model (6.1) can be carried out by the least squares method. To see this, we first focus on estimation of $\phi = (\phi_1, \dots, \phi_p)^T$. Let $S(\phi) = \sum_{t=p+1}^T \{X_t - (\phi_0 + \sum_{j=1}^p \phi_j X_{t-j})\}^2$ be the sum of the squared difference between X_t and its linearly combined history with lag p . Then applying the constraint (6.2) gives $S(\phi) = \sum_{t=p+1}^T \left[\{X_t - E(X_t)\} - \{\tilde{X}_{t-1} - E(\tilde{X}_{t-1})\}^T \phi \right]^2$.

To minimize $S(\phi)$ with respect to ϕ , we solve $\frac{\partial S(\phi)}{\partial \phi} = 0$ for ϕ and obtain the solution

$$\hat{\phi}^{(\text{LS})} = \left(\sum_{t=p+1}^T \left\{ \tilde{X}_{t-1} - E(\tilde{X}_{t-1}) \right\} \left\{ \tilde{X}_{t-1} - E(\tilde{X}_{t-1}) \right\}^T \right)^{-1} \sum_{t=p+1}^T \left\{ \tilde{X}_{t-1} - E(\tilde{X}_{t-1}) \right\} \{X_t - E(X_t)\}, \quad (6.3)$$

where for $t = 1, \dots, T$, $E(X_t)$ can be estimated by $\frac{1}{T} \sum_{t=1}^T X_t$, which is denoted as $\hat{\mu}$.

Next, by the constraint (6.2), replacing $E(X_t)$ by $\hat{\mu}$ gives an estimator of ϕ_0 :

$$\hat{\phi}_0^{(\text{LS})} = \hat{\mu} - \hat{\mu} \cdot \sum_{j=1}^p \hat{\phi}_j. \quad (6.4)$$

Re-expressing (6.1) as $\epsilon_t = X_t - (\phi_0 + \sum_{j=1}^p \phi_j X_{t-j})$ and by the definition of $S(\phi)$, we may estimate $\text{Var}(\epsilon_t) = \sigma_\epsilon^2$ by

$$\begin{aligned}\hat{\sigma}_\epsilon^{2(\text{LS})} &= \frac{1}{T-p} S(\hat{\phi}) \\ &= \frac{1}{T-p} \sum_{t=p+1}^T \{X_t - E(X_t)\}^2 - \frac{2}{T-p} \sum_{t=p+1}^T \{X_t - E(X_t)\} \{\tilde{X}_{t-1} - E(\tilde{X}_{t-1})\}^T \hat{\phi} \\ &\quad + \frac{1}{T-p} \sum_{t=p+1}^T \hat{\phi}^T \{\tilde{X}_{t-1} - E(\tilde{X}_{t-1})\} \{\tilde{X}_{t-1} - E(\tilde{X}_{t-1})\}^T \hat{\phi}\end{aligned}\quad (6.5)$$

with $E(X_t)$ estimated by $\hat{\mu}$.

Estimators (6.3)–(6.5) can be derived in an alternative way. First, by the stationarity of the X_t , for $k = 0, \dots, p$ and $p \leq t$, $\text{Cov}(X_t, X_{t-k})$ is time-independent and let γ_k denote it; it is clear that γ_0 represents $\text{Var}(X_t)$ for any t . Let Γ be the autocovariance matrix

$$\Gamma = \begin{pmatrix} \gamma_0 & \cdots & \gamma_{p-1} \\ \vdots & \ddots & \vdots \\ \gamma_{p-1} & \cdots & \gamma_0 \end{pmatrix}.$$

Let $\hat{\gamma} = (\hat{\gamma}_1, \dots, \hat{\gamma}_p)^T$ with $\hat{\gamma}_k = \frac{1}{T-k} \sum_{t=k+1}^T (X_t - \hat{\mu})(X_{t-k} - \hat{\mu})$ being an estimator of γ_k for $k = 0, \dots, p$, and let $\hat{\Gamma}$ be the estimator of Γ with γ_k replaced by $\hat{\gamma}_k$ for $k = 0, \dots, p-1$.

Next, we examine the summation terms in (6.3) and (6.5) by using the fact that as $T \rightarrow \infty$, $\frac{1}{T-p} \sum_{t=p+1}^T \{X_t - E(X_t)\}^2 \xrightarrow{p} \gamma_0$, $\frac{1}{T-p} \sum_{t=p+1}^T \{X_t - E(X_t)\} \{\tilde{X}_{t-1} - E(\tilde{X}_{t-1})\}^T \xrightarrow{p} \gamma$, and $\frac{1}{T-p} \sum_{t=p+1}^T \{\tilde{X}_{t-1} - E(\tilde{X}_{t-1})\} \{\tilde{X}_{t-1} - E(\tilde{X}_{t-1})\}^T \xrightarrow{p} \Gamma$. Then, (6.3)–(6.5) motivate an alternative method of finding estimators for ϕ , ϕ_0 , and σ_ϵ^2 , by solving the estimating equations:

$$\begin{aligned}\phi &= \hat{\Gamma}^{-1} \hat{\gamma}; \\ \phi_0 &= \left(1 - \sum_{i=1}^p \phi_i\right) \hat{\mu}; \\ \sigma_\epsilon^2 &= \hat{\gamma}_0 - 2\phi^T \hat{\gamma} + \phi^T \hat{\Gamma} \phi,\end{aligned}\quad (6.6)$$

for ϕ , ϕ_0 , and σ_ϵ^2 . Let $\hat{\phi}$, $\hat{\phi}_0$ and $\hat{\sigma}_\epsilon^2$ denote the resultant estimators of ϕ , ϕ_0 , and σ_ϵ^2 , respectively. These estimators are asymptotically equivalent to the least squares estimators

$\widehat{\phi}^{(\text{LS})}$, $\widehat{\phi}_0^{(\text{LS})}$, and $\widehat{\sigma}_\epsilon^{2(\text{LS})}$ in a sense that $\widehat{\phi} - \widehat{\phi}^{(\text{LS})} \xrightarrow{p} 0$, $\widehat{\phi}_0 - \widehat{\phi}_0^{(\text{LS})} \xrightarrow{p} 0$ and $\widehat{\sigma}_\epsilon^2 - \widehat{\sigma}_\epsilon^{2(\text{LS})} \xrightarrow{p} 0$, as $T \rightarrow \infty$, and hence, they are consistent (Box et al., 2015, Section A.7.4).

Estimating equations (6.6) offer a unified estimation framework in its connections with not only the least squares estimation but also the maximum likelihood method under the assumption of Gaussian error as well as the Yule-Walker method. Similar to the least squares method, finding estimators using one of those approaches is asymptotically equivalent to solving (6.6) for ϕ , ϕ_0 and σ_ϵ^2 (Box et al., 2015, Section A.7.4).

6.2 Measurement Error and Impact

6.2.1 Measurement Error Models

Suppose that for $t = 1, \dots, T$, the observation of X_t is subject to measurement error and the precise measurement of X_t may not be observed, but its surrogate measurement X_t^* is available. We consider two measurement error models.

The first measurement error model takes an additive form

$$X_t^* = \alpha_0 + \alpha_1 X_t + e_t \quad (6.7)$$

for $t = 1, \dots, T$, where the error term e_t is independent of X_t with mean 0 and time-independent variance σ_e^2 and is assumed to be mutually independent for $t = 1, \dots, T$, and $\alpha = (\alpha_0, \alpha_1)^\top$ is the parameter vector. Here, α_0 represents the systematic error and α_1 represents the constant inflation (or shrinkage) due to the measurement error. For instance, if $\alpha_0 = 0$, then setting $\alpha_1 < 1$ (or $\alpha_1 > 1$) features the scenario where X_t^* tends to be smaller (or larger) than X_t if the noise term is ignored. This model generalizes the classical additive model considered by Staudenmayer and Buonaccorsi (2005) who considered the case with $\alpha_0 = 0$ and $\alpha_1 = 1$.

By the stationarity of the X_t , we note that model (6.7) yields $E(X_t^*) = \alpha_0 + \alpha_1 \mu$ and

$$\text{Var}(X_t^*) = \alpha_1^2 \gamma_0 + \sigma_e^2; \quad (6.8)$$

the variability of the X_t^* can be greater or smaller than that of the X_t , depending on the value of α_1 .

The second measurement error model assumes a multiplicative form:

$$X_t^* = \beta_0 u_t X_t, \quad (6.9)$$

for $t = 1, \dots, T$, where β_0 is a positive scaling parameter, and the u_t are the error terms which are independent of each other as well as of the X_t , and have mean one and time-independent variance σ_u^2 . Depending on the distribution of the error term u_t , (6.9) can feature different types of discrepancy between X_t and X_t^* .

The stationarity of the X_t together with model (6.9) implies $E(X_t^*) = \beta_0\mu$, and

$$\begin{aligned}
\text{Var}(X_t^*) &= \text{Var}(\beta_0 X_t u_t) \\
&= \beta_0^2 \{E(X_t^2 u_t^2) - E^2(X_t u_t)\} \\
&= \beta_0^2 \{E(X_t^2)E(u_t^2) - E^2(X_t)E^2(u_t)\} \\
&= \beta_0^2 \{(\text{Var}(X_t) + E^2(X_t))(\sigma_u^2 + 1) - E^2(X_t)\} \\
&= \beta_0^2 \{(\sigma_u^2 + 1)\gamma_0 + \sigma_u^2 \mu^2\}, \tag{6.10}
\end{aligned}$$

where the third step is because of the independence of X_t and u_t .

Since $E(X_t^*)$ is time-independent for both (6.7) and (6.9), in the following discussion, we let μ^* denote $E(X_t^*)$ for $t = 1, \dots, T$. The modeling of the measurement error process by (6.7) or (6.9) introduces extra parameters $\{\alpha_0, \alpha_1, \sigma_e^2\}$ or $\{\beta_0, \sigma_u^2\}$, where the variance of the error term is bounded by the variability of X_t^* together with others. Clearly, (6.8) shows that $\sigma_e^2 < \text{Var}(X_t^*)$ and (6.10) implies that $\sigma_u^2 < \frac{\text{Var}(X_t^*)}{\beta_0^2 \mu^2}$.

6.2.2 Naive Estimation and Bias for AR(1) Model

Estimating equations (6.6) are useful when measurements of X_t are available. However, due to the measurement error, X_t is not observed so (6.6) cannot be directly used for estimation of the parameters for model (6.1). As the surrogate X_t^* for X_t is available, one may attempt to employ the naive analysis to model (6.1) with X_t replaced by X_t^* . Here we study the impact of measurement error on the naive analysis disregarding the difference between X_t and X_t^* . We start with the AR(1) model, i.e., model (6.1) with $p = 1$.

If we naively replace X_t in (6.1) by X_t^* , then the time series model (6.1) becomes

$$X_t^* = \phi_0^* + \phi_1^* X_{t-1}^* + \epsilon_t^*, \tag{6.11}$$

where $(\phi_0^*, \phi_1^*)^T$ and ϵ_t^* show possible differences from the corresponding quantity in the model (6.1). To estimate ϕ_0^* and ϕ_1^* , we may employ the ordinary least squares (OLS) method. Specifically, we minimize $S(\phi_0^*, \phi_1^*) = \sum_{t=2}^T (X_t^* - \phi_0^* - \phi_1^* X_{t-1}^*)^2$ with respective

to ϕ_0^* and ϕ_1^* , yielding the OLS estimators of ϕ_1^* and ϕ_0^* :

$$\widehat{\phi}_1^* = \frac{\sum_{t=2}^T (X_{t-1}^* - \bar{X}_{(-1)}^*)(X_t^* - \bar{X}^*)}{\sum_{t=2}^T (X_{t-1}^* - \bar{X}_{(-1)}^*)^2},$$

and

$$\widehat{\phi}_0^* = \bar{X}_t^* - \widehat{\phi}_1^* \bar{X}^*, \quad (6.12)$$

where $\bar{X}_{(-1)}^* = \frac{1}{T-1} \sum_{t=2}^T X_{t-1}^*$ and $\bar{X}^* = \frac{1}{T-1} \sum_{t=2}^T X_t^*$.

Theorem 6.1 *Let $\omega_1 = \frac{\alpha_1^2 \sigma_\epsilon^2}{\alpha_1^2 \sigma_\epsilon^2 + \sigma_\epsilon^2 (1 - \phi_1^*)}$, $\phi_1^* = \phi_1 \omega_1$, and $\phi_0^* = \left(\alpha_0 + \frac{\alpha_1 \phi_0}{1 - \phi_1} \right) (1 - \phi_1 \omega_1)$. Assume the stationarity of the times series. If the measurement error process satisfies (6.7), then*

- (1) $\widehat{\phi}_1^* \xrightarrow{p} \phi_1^*$ and $\widehat{\phi}_0^* \xrightarrow{p} \phi_0^*$ as $T \rightarrow \infty$,
- (2) $\epsilon_t^* = \alpha_0 (1 - \phi_1^*) + \alpha_1 \phi_0 - \phi_0^* + \alpha_1 (\phi_1 - \phi_1^*) X_{t-1} + (1 - \phi_1^*) e_t + \alpha_1 \epsilon_t$ for $t = 1, \dots, T$,
and hence $\text{Var}(\epsilon_t^*) = \phi_1^2 \alpha_1^2 (1 - \omega_1)^2 \left(\frac{\sigma_\epsilon^2}{1 - \phi_1^2} \right) + (1 - \omega_1 \phi_1)^2 \sigma_\epsilon^2 + \alpha_1^2 \sigma_\epsilon^2$.

The proof of the theorem is included in Appendix E.2. This theorem essentially implies that the naive estimator under the additive form in (6.7) is inconsistent because $\phi_1^* \neq \phi_1$ and $\phi_0^* \neq \phi_0$. The naive estimator $\widehat{\phi}_1^*$ attenuates and the attenuation factor ω_1 depends on the parameters α_1 and σ_ϵ^2 of the measurement error model (6.7) as well as ϕ_1 and σ_ϵ^2 in the time series model (6.1). The coefficient α_1 in the measurement error model (6.7) affects the estimation of the both naive estimators $\widehat{\phi}_1^*$ and $\widehat{\phi}_0^*$, while the intercept α_0 influences the estimation of ϕ_0^* only, but not ϕ_1^* or $\text{Var}(\epsilon^*)$.

Theorem 6.2 *Let $\omega_2 = \left\{ 1 + \sigma_u^2 + \frac{(1 + \phi_1) \sigma_u^2 \phi_0^2}{(1 - \phi_1) \sigma_\epsilon^2} \right\}^{-1}$, $\phi_1^* = \phi_1 \omega_2$, and $\phi_0^* = \frac{\beta_0 \phi_0}{1 - \phi_1} (1 - \omega_2 \phi_1)$. If the times series is stationary and the measurement error process satisfies (6.9), then*

- (1) $\widehat{\phi}_1^* \xrightarrow{p} \phi_1^*$ and $\widehat{\phi}_0^* \xrightarrow{p} \phi_0^*$ as $T \rightarrow \infty$,
- (2) $\epsilon_t^* = \beta_0 \phi_0 u_t - \phi_0^* + \beta_0 X_{t-1} (\phi_1 u_t - \omega_2 \phi_1 u_{t-1}) + \beta_0 u_t \epsilon_t$ for $t = 1, \dots, T$,
and hence $\text{Var}(\epsilon_t^*) = \beta_0^2 \{ \sigma_u^2 \phi_0^2 + (1 + \sigma_u^2) \sigma_\epsilon^2 \} + \beta_0^2 \phi_1^2 \frac{(1 + \omega_2^2)}{\omega_2} \frac{\sigma_\epsilon^2}{(1 - \phi_1^2)}$.

The proof of the theorem is included in Appendix E.3. This theorem says the attenuation effect resulting from the measurement error on estimation of ϕ_1 . The constant scaling parameter β_0 in the measurement error model (6.9) does not influence the estimation of ϕ_1 but affects the estimation of ϕ_0 and σ_ϵ^2 . The attenuation factor ω_2 is determined by the magnitude σ_u^2 of measurement error as well as the values of ϕ_0 , ϕ_1 , and σ_ϵ^2 of the time series model (6.1).

6.2.3 Naive Estimation and Bias for AR(p) Model with $p \geq 2$

We now extend the discussion in Section 6.2.2 to the AR(p) model with $p \geq 2$. Replacing X_t with X_t^* in (6.1) gives the working model

$$X_t^* = \phi_0^* + \sum_{j=1}^p \phi_j^* X_{t-j}^* + \epsilon_t^*, \quad (6.13)$$

where $\phi^* = (\phi_1^*, \dots, \phi_p^*)^\top$ and ϵ_t^* may differ from the corresponding symbol in (6.1). If mimicking the procedure of using (6.6) with X_t replaced by X_t^* to estimate ϕ^* , ϕ_0^* and σ_ϵ^{2*} in (6.13), then we let $\hat{\phi}^* = (\hat{\phi}_1^*, \dots, \hat{\phi}_p^*)^\top$, $\hat{\phi}_0^*$ and $\hat{\sigma}_\epsilon^{2*}$ denote the resultant estimators. Similar to $\hat{\gamma}_k$ and $\hat{\mu}$, we define $\hat{\mu}^* = \frac{1}{T} \sum_{t=1}^T X_t^*$ and $\hat{\gamma}_k^* = \frac{1}{T-k} \sum_{t=1}^{T-k} (X_t^* - \hat{\mu}^*)(X_{t+k}^* - \hat{\mu}^*)$ for $k = 1, \dots, p$. Let $\hat{\gamma}^* = (\hat{\gamma}_1^*, \dots, \hat{\gamma}_p^*)^\top$ and $\hat{\gamma}_0^* = \frac{1}{T} \sum_{t=1}^T (X_t^* - \hat{\mu}^*)(X_t^* - \hat{\mu}^*)$.

We now discuss the asymptotic results of the naive estimators under different measurement error models.

Theorem 6.3 *Let $\mathbf{1}_p$ be the $p \times 1$ unit and let I_p be the $p \times p$ identity matrix. Define $\gamma^* = \alpha_1^2 \gamma$, $\gamma_0^* = \alpha_1^2 \gamma_0 + \sigma_e^2$, $\phi^* = \alpha_1^2 (\alpha_1^2 \Gamma + \sigma_e^2 I_p)^{-1} \gamma$, $\phi_0^* = (1 - \phi^* \cdot \mathbf{1}_p) (\alpha_0 + \alpha_1 \mu)$ and $\sigma_\epsilon^{2*} = \alpha_1^2 \gamma_0 + \sigma_e^2 - \alpha_1^4 \gamma^\top (\alpha_1^2 \Gamma + \sigma_e^2 I_p)^{-1} \gamma$. Under regularity conditions, if the time series is stationary and the measurement error process satisfies (6.7), then*

$$(1) \hat{\gamma}^* \xrightarrow{P} \gamma^* \text{ and } \hat{\gamma}_0^* \xrightarrow{P} \gamma_0^* \text{ as } T \rightarrow \infty.$$

$$(2) \hat{\phi}^* \xrightarrow{P} \phi^*, \hat{\phi}_0^* \xrightarrow{P} \phi_0^*, \text{ and } \hat{\sigma}_\epsilon^{2*} \xrightarrow{P} \sigma_\epsilon^{2*} \text{ as } T \rightarrow \infty.$$

(3) *Let Q_1 denote the $(p+1) \times (p+1)$ asymptotic covariance matrix of $\sqrt{T} \{(\hat{\gamma}_0^*, \hat{\gamma}^{*\top})^\top - (\gamma_0^*, \gamma^{*\top})^\top\}$ as $T \rightarrow \infty$. Then the elements of Q_1 are given by*

$$\begin{aligned} q_{100}^* &= \alpha_1^4 q_{00} + 4\alpha_1^2 \gamma_0 \sigma_e^2 + E(e_t^4) - \sigma_e^4; \\ q_{10p}^* &= \alpha_1^4 q_{0p} + 4\alpha_1^2 \gamma_p \sigma_e^2; \\ q_{1pr}^* &= \alpha_1^4 q_{pr} + 2\alpha_1^2 \sigma_e^2 (\gamma_{|p-r|} + \gamma_{p+r}) \text{ for } r \neq 0, r \neq p; \\ q_{1pp}^* &= \alpha_1^4 q_{pp} + 2\alpha_1^2 \sigma_e^2 (\gamma_0 + \gamma_{2p}) + \sigma_e^4; \end{aligned}$$

for $p \geq 1$, where q_{jk} is the (j, k) element of the asymptotic covariance matrix of $(\hat{\gamma}_0, \hat{\gamma}^\top)^\top$, given by (Brockwell et al., 1991, Section 7.3)

$$q_{jk} = (\eta - 3) \gamma_j \gamma_k + \sum_{i=-\infty}^{\infty} (\gamma_i \gamma_{i-j+k} + \gamma_{i+k} \gamma_{i-j}) \quad (6.14)$$

for $(j, k) = (0, 0), (0, p), (p, p)$ and (p, r) with $r \neq 0$ and $r \neq p$, with $\eta = E(\epsilon_t^4)/\sigma_\epsilon^4$.

The proof of Theorem 6.3 is presented in the Appendix E.4. Similar to the results in Theorem 6.1, the intercept α_0 only influences ϕ_0 and does not influence ϕ .

Theorem 6.4 *Let $\gamma^* = \beta_0^2 \gamma$, $\gamma_0^* = \beta_0^2 \{(\sigma_u^2 + 1)\gamma_0 + \sigma_u^2 \mu^2\}$, $\phi^* = \{\Gamma + \sigma_u^2(\gamma_0 + \mu^2)I_p\}^{-1} \gamma$, $\phi_0^* = \beta_0(1 - \phi^{*\top} \cdot \mathbf{1}_p) \mu$, and $\sigma_\epsilon^{2*} = \beta_0^2(\sigma_u^2 + 1)\gamma_0 + \beta_0^2 \sigma_u^2 \mu^2 - \beta_0^2 \gamma^\top \{\Gamma + \sigma_u^2(\gamma_0 + \mu^2)I_p\}^{-1} \gamma$. Under regularity conditions, if the time series are stationary and the measurement error process satisfy (6.9), then*

$$(1) \hat{\gamma}^* \xrightarrow{p} \gamma^* \text{ and } \hat{\gamma}_0^* \xrightarrow{p} \gamma_0^* \quad \text{as } T \rightarrow \infty.$$

$$(2) \hat{\phi}^* \xrightarrow{p} \phi^*, \hat{\phi}_0^* \xrightarrow{p} \phi_0^*, \text{ and } \hat{\sigma}_\epsilon^{2*} \xrightarrow{p} \sigma_\epsilon^{2*} \quad \text{as } T \rightarrow \infty.$$

(3) *Let Q_2 denote the $(p + 1) \times (p + 1)$ asymptotic covariance matrix of $\sqrt{T} \{(\hat{\gamma}_0^*, \hat{\gamma}^{*\top})^\top - (\gamma_0^*, \gamma^{*\top})^\top\}$ as $T \rightarrow \infty$. Then the elements of Q_2 are given by*

$$\begin{aligned} q_{200}^* &= \beta_0^4(\sigma_u^2 + 1)^2 q_{00} + \beta_0^4 \{E(u_t^4) - (\sigma_u^2 + 1)^2\} E(X_t - \mu)^4 \\ &\quad + 4\mu\beta_0^4 \sigma_u^2 (\sigma_u^2 + 1) v_0 + 4\mu\beta_0^4 \{E(u_t^4) - E(u_t^3) - \sigma_u^2(\sigma_u^2 + 1)\} E(X_t - \mu)^3 \\ &\quad + 2\mu^2 \beta_0^4 \{E(u_t^4) - 2E(u_t^3) + 1 - \sigma_u^4\} \gamma_0 \\ &\quad + 4\mu^2 \beta_0^4 \left[\sigma_u^4 \sum_{h=-\infty}^{\infty} \gamma_h + \{E(u_t^4) - 2E(u_t^3) + \sigma_u^2 + 1 - \sigma_u^4\} \gamma_0 \right] + \mu^4 \beta_0^4 [E\{(u_t - 1)^4\} - \sigma_u^4]; \\ q_{20p}^* &= \beta_0^4 q_p(\sigma_u^2 + 1) + \beta_0^4 \{E(u_t^3) - (\sigma_u^2 + 1)\} [E\{(X_t - \mu)^3(X_{t+p} - \mu)\} + E\{(X_t - \mu)^3(X_{t-p} - \mu)\}] \\ &\quad + 2\mu\beta_0^4 \sigma_u^2 v_{0p} + \mu\beta_0^4 E\{3u_t^3 - 3u_t^2 - 2\sigma_u^2\} [E\{(X_t - \mu)^2(X_{t-p} - \mu)\} + E\{(X_t - \mu)^2(X_{t+p} - \mu)\}] \\ &\quad + 6\mu^2 \beta_0^4 E(u_t - 1)^3 \gamma_p + 4\mu^2 \beta_0^4 \sigma_u^2 \gamma_p; \\ q_{2pr}^* &= \beta_0^4 q_{pr} + \beta_0^4 \sigma_u^2 [E\{(X_t - \mu)^2(X_{t+p} - \mu)(X_{t+r} - \mu)\} + E\{(X_t - \mu)(X_{t+p} - \mu)^2(X_{t+p+r} - \mu)\}] \\ &\quad + E\{(X_{t-r} - \mu)(X_t - \mu)^2(X_{t+p} - \mu)\} + E\{(X_t - \mu)(X_{t+p-r} - \mu)(X_{t+p} - \mu)^2\}] \\ &\quad + \mu\beta_0^4 \sigma_u^2 [E\{(X_t - \mu)(X_{t+p} - \mu)(X_{t+r} - \mu)\} + E\{(X_t - \mu)(X_{t+p} - \mu)(X_{t+p+r} - \mu)\}] \\ &\quad + E\{(X_{t-r} - \mu)(X_t - \mu)(X_{t+p} - \mu)\} + E\{(X_t - \mu)(X_{t+p-r} - \mu)(X_{t+p} - \mu)\}] \\ &\quad + 2\mu^2 \beta_0^4 \sigma_u^2 (\gamma_{|p-r|} + \gamma_{p+r}) \text{ for } r \neq p, r \neq 0; \\ q_{2pp}^* &= \beta_0^4 q_{pp} + \beta_0^4 (\sigma_u^4 + 2\sigma_u^2) \text{Var}\{(X_t - \mu)(X_{t+p} - \mu)\} + 2\beta_0^4 E\{(X_t - \mu)(X_{t+p} - \mu)^2(X_{t+2p} - \mu)\} \\ &\quad + \mu\beta_0^4 \sigma_u^2 [E\{(X_t - \mu)(X_{t+p} - \mu)^2\} + 2E\{(X_t - \mu)(X_{t+p} - \mu)(X_{t+2p} - \mu)\}] \\ &\quad + E\{(X_t - \mu)^2(X_{t+p} - \mu)\}] + 2\mu^2 \beta_0^4 \sigma_u^4 \gamma_p + 2\mu^2 \beta_0^4 \sigma_u^2 (\gamma_0 + \gamma_{2p}) + \mu^4 \beta_0^4 \sigma_u^4, \end{aligned}$$

where the q_{jk} are given by (6.14), for $(j, k) = (0, 0), (0, p), (p, p)$ and (p, r) with $r \neq 0$ and $r \neq p$, and $v_p = \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T E\{(X_t - \mu)(X_{t+p} - \mu)(X_s - \mu)\}$.

The proof of the theorem is presented in Appendix E.5. The multiplicative measurement error u_t contributes to the biasedness of the parameter estimation for ϕ , while the scaling parameter β_0 has no effects on the naive estimator $\hat{\phi}^*$.

6.3 Methodology of Correcting Measurement Error Effects

6.3.1 Estimation of Model Parameters

In the presence of measurement error, measurements of the X_t are not always available but surrogate measurements X_t^* are available. It may be tempting to conduct a naive analysis by implementing (6.6) with the X_t replaced by the X_t^* , or equivalently with $\hat{\mu}$ and $\hat{\gamma}_k$ replaced by $\hat{\mu}^*$ and the $\hat{\gamma}_k^*$, respectively, to find estimators of ϕ , ϕ_0 and σ_ϵ^2 . However, by Theorems 6.3–6.4, such a procedure typically yields biased estimators. In this section, we develop new estimators accounting for the measurement error effects described by either the additive model (6.7) or the multiplicative model (6.9).

Our idea is still to employ (6.6) to find consistent estimators of ϕ , ϕ_0 and σ_ϵ^2 , but instead of replacing $\hat{\mu}$ and the $\hat{\gamma}_k$ with $\hat{\mu}^*$ and the $\hat{\gamma}_k^*$ as in the naive analysis, we replace $\hat{\mu}$ and the $\hat{\gamma}_k$ in (6.6) with new functions of the X_t^* , denoted as $\tilde{\mu}$ and the $\tilde{\gamma}_k$, which adjust for the measurement error effects. Specifically, if we can find $\tilde{\mu}$ and the $\tilde{\gamma}_k$ such that they resemble $\hat{\mu}$ and the $\hat{\gamma}_k$ in the sense that as $T \rightarrow \infty$,

$$\begin{aligned} &\tilde{\mu} \text{ and } \hat{\mu} \text{ have the same limit in probability,} \\ \text{and } &\tilde{\gamma}_k \text{ and } \hat{\gamma}_k \text{ have the same limit in probability for } k = 0, \dots, p, \end{aligned} \quad (6.15)$$

then substituting $\hat{\mu}$ and the $\hat{\gamma}_k$ with $\tilde{\mu}$ and the $\tilde{\gamma}_k$ in (6.6) yields consistent estimators of ϕ , ϕ_0 and σ_ϵ^2 .

With the availability of the $\tilde{\gamma}_k$ satisfying (6.15), let $\tilde{\Gamma}$ denote Γ with the γ_k replaced by the $\tilde{\gamma}_k$. Then provided regularity conditions, consistent estimators of ϕ , ϕ_0 and σ_ϵ^2 can be obtained by solving the estimating equations for ϕ , ϕ_0 , and σ_ϵ^2 :

$$\begin{aligned} \phi &= \tilde{\Gamma}^{-1} \tilde{\gamma}, \\ \phi_0 &= \left(1 - \sum_{i=1}^p \phi_i \right) \tilde{\mu}, \\ \sigma_\epsilon^2 &= \tilde{\gamma}_0 - 2\phi^T \tilde{\gamma} + \phi^T \tilde{\Gamma} \phi. \end{aligned} \quad (6.16)$$

It is immediate to obtain the following result.

Theorem 6.5 *Assume regularity conditions hold and the time series are stationary. If $\tilde{\mu}$ and the $\tilde{\gamma}_k$ are functions of the X_t^* with $t = 1, \dots, T$ and they satisfy (6.15), and let ϕ , ϕ_0 ,*

and $\tilde{\sigma}_\epsilon^2$ denote the estimators for ϕ , ϕ_0 and σ_ϵ^2 , respectively, obtained by solving (6.16). Then, as $T \rightarrow \infty$

(1) $\tilde{\phi} \xrightarrow{p} \phi$, $\tilde{\phi}_0 \xrightarrow{p} \phi_0$, and $\tilde{\sigma}_\epsilon^2 \xrightarrow{p} \sigma_\epsilon^2$;

(2) $\sqrt{n}(\tilde{\phi} - \phi) \xrightarrow{d} N(0, GQG^T)$,

where G is the matrix of derivatives of $\tilde{\phi}$ with respect to the components of $(\hat{\gamma}_0^*, \hat{\gamma}^{*T})^T$. Here $Q = Q_1$, the matrix in Theorem 6.3, if measurement error follows the model (6.7); and $Q = Q_2$, the matrix in Theorem 6.4, if measurement error follows the model (6.9).

Now we discuss explicitly how to determine $\tilde{\mu}$ and the $\tilde{\gamma}_k$ under the measurement error model (6.7) or (6.9). With (6.7), take $\tilde{\mu} = \frac{\hat{\mu}^*}{\alpha_1} - \alpha_0$, $\tilde{\gamma}_0 = \frac{1}{\alpha_1^2}(\hat{\gamma}_0^* - \sigma_\epsilon^2)$, and $\tilde{\gamma}_k = \frac{\hat{\gamma}_k^*}{\alpha_1^2}$ for $k = 1, \dots, p$. With (6.9), take $\tilde{\mu} = \frac{\hat{\mu}^*}{\beta_0}$, $\tilde{\gamma}_0 = \frac{\hat{\gamma}_0^*}{(1+\sigma_u^2)\beta_0^2} - \frac{\sigma_u^2\mu^2}{\sigma_u^2+1}$, and $\tilde{\gamma}_k = \frac{\hat{\gamma}_k^*}{\beta_0^2}$ for $k = 1, \dots, p$. By the results in Theorem 6.3(1) and Theorem 6.4(1), it can be easily verified that these $\tilde{\mu}$ and the $\tilde{\gamma}_k$ satisfy (6.15).

We conclude this section with a procedure of estimating the asymptotic covariance matrix for the estimator $\tilde{\phi}$. While Theorem 6.5 presents the sandwich form of the asymptotic covariance matrix of $\tilde{\phi}$, its evaluation involves lengthy calculations. We may alternatively employ the block bootstrap algorithm (Lahiri, 1999) to obtain variance estimates for $\tilde{\phi}$ using the following steps. Firstly, we set a positive integer, say N , as the number for the bootstrap sampling; N can be set as a large number such as 1000. Next, we repeat through the following five steps:

Step 1: At iteration $n \in \{1, \dots, N\}$, we initialize a null time series $X^{(n,0)}$ of dimension 0 and specify a block length, say b , which is an integer between 0 and T . Initialize $m=1$.

Step 2: Sample an index, say i , from $\{0, \dots, T - b\}$, and then define $X_{\text{add}}^{(m-1)} = \{X_{i+1}, \dots, X_{i+b}\}$.

Step 3: Update the previous time series $X^{(n,m-1)}$ by appending $X_{\text{add}}^{(m-1)}$ to it, and let $X^{(n,m)}$ denote the new time series.

Step 4: If the dimension $X^{(n,m)}$ is smaller than T then return to Steps 2 and 3; otherwise drop the elements in the time series with the index greater than T to ensure the dimension of $X^{(n,m)}$ is identical to T and then go to Step 5.

Step 5: Obtain an estimate $\tilde{\phi}^{(n)}$ of parameter ϕ by applying the times series $X^{(n,m)}$ to (6.16). If $n < N$, then set n to be $n + 1$ and go back to Step 1 to repeat; otherwise stop.

Let $\tilde{\phi} = \frac{1}{N} \sum_{n=1}^N \tilde{\phi}^{(n)}$ be the sample mean. The bootstrap variance of $\tilde{\phi}$ is then given by,

$$\text{Var}_{\text{boot}}(\tilde{\phi}) = \frac{1}{N} \sum_{n=1}^N (\tilde{\phi}^{(n)} - \tilde{\phi})^2.$$

6.3.2 Forecasting and Prediction Error

Forecasting is an important application of the autoregressive models. Specifically, in forecasting based on the observed time series $X_{(T)} = \{x_1, \dots, x_T\}$, we are interested in the predictions of $\{X_{T+1}, \dots, X_{T+H}\}$ for a positive integer H , which is done one by one starting from the nearest time point $T + 1$ to the farthest time point $T + H$. To this end, let $h = 1, \dots, H$, the h -step forecasting of X_{T+h} is based on its history of lag- p , $\{X_{T+h-1}, \dots, X_{T+h-p}\}$, by using the conditional expectation $E(X_{T+h} | x_{T+h-1}, \dots, x_{T+h-p})$, denoted \hat{X}_{T+h} , where for $j = T + h - 1, \dots, T + h - p$, x_j is the observe value of X_j if $j \leq T$; and x_j is the predicted value of X_j , \hat{X}_j , if $j > T$. This prediction minimizes the squared prediction error $E(\hat{X}_{T+h} - X_{T+h})^2$ (e.g., Box et al., 2015, Page 131).

If no measurement error is involved, due to the zero mean of the random error term ϵ_t in the AR(p) model (6.1), for $h = 1, \dots, H$, the conditional expectation can be calculated by

$$\hat{X}_{T+h} = \phi_0 + \phi_1 x_{T+h-1} + \dots + \phi_p x_{T+h-p}. \quad (6.17)$$

When measurement error appears, the observe values x_j for $j = T, \dots, T - p + 1$ in (6.17) are no longer available but their surrogates X_j^* are available. We now provide a sensible estimate of X_j by using the measurement error model for characterizing the relationship of X_j and X_j^* . If measurement error follows (6.7), we “estimate” X_j by

$$\hat{X}_j = \frac{1}{\alpha_1} (X_j^* - \alpha_0) \quad \text{for } j = t, \dots, t - p + 1; \quad (6.18)$$

if the measurement error follows (6.9), then \hat{X}_j is “estimated” by

$$\hat{X}_j = \frac{X_j^*}{\beta_0} \quad \text{for } j = t, \dots, t - p + 1. \quad (6.19)$$

These “estimates” are unbiased in the sense that $E(\widehat{X}_j) = X_j$ for $j = t, \dots, t - p + 1$. Consequently, for $h = 1, \dots, H$, X_{T+h} is predicted as

$$\widehat{X}_{T+h} = \phi_0 + \phi_1 \widehat{X}_{T+h-1} + \dots + \phi_p \widehat{X}_{T+h-p}. \quad (6.20)$$

In contrast to the observed values $\{x_T, \dots, x_{T-p+1}\}$, also referred to as the initial values of the forecasting of X_{T+1}, \dots, X_{T+H} , the estimates determined by (6.18) or (6.19) introduce additional prediction error which should be characterized. Without the loss of generality, we consider $p = 1$ to illustrate the recursive calculation of the prediction error; the prediction error with a higher order of autoregressive process can be derived recursively in a similar way but with more complex expressions.

If the measurement error follows (6.7), the mean squared prediction error of the 1-step prediction is given by

$$\begin{aligned} P_e^{(1)} &= E(\widehat{X}_{T+1} - X_{T+1})^2 \\ &= E\{(\phi_0 + \phi_1 \widehat{X}_T) - (\phi_0 + \phi_1 X_T + \epsilon_{T+1})\}^2 \\ &= E\left\{\phi_1 \left(X_T + \frac{e_T}{\alpha_1}\right) - \phi_1 X_T - \epsilon_{T+1}\right\}^2 \\ &= \frac{\phi_1^2 \sigma_e^2}{\alpha_1^2} + \sigma_\epsilon^2, \end{aligned}$$

where the last step is due to the independence between e_t and ϵ_{t+1} , as well as $E(e_t^2) = \sigma_e^2$ and $E(\epsilon_t^2) = \sigma_\epsilon^2$.

Then, the h -step prediction error is given by

$$\begin{aligned} P_e^{(h)} &= E(\widehat{X}_{T+h} - X_{T+h})^2 \\ &= E(\phi_0 + \phi_1 \widehat{X}_{T+h-1} - \phi_0 - \phi_1 X_{T+h-1} - \epsilon_{T+1})^2 \\ &= E\left\{\phi_1 \left(\widehat{X}_{T+h-1} - X_{T+h-1}\right) - \epsilon_{T+1}\right\}^2 \\ &= \phi_1^2 P_e^{(h-1)} + \sigma_\epsilon^2 \\ &= \frac{\phi_1^{2h} \sigma_e^2}{\alpha_1^2} + \sum_{i=0}^{h-1} \phi_1^{2i} \sigma_\epsilon^2, \end{aligned} \quad (6.21)$$

where the last step comes from the recursive evaluation of $P_e^{(h-1)}$.

Similarly, if the measurement error follows (6.9), the mean squared prediction error is given by

$$\begin{aligned}
P_e^{(1)} &= E(\widehat{X}_{T+1} - X_{T+1})^2 \\
&= E(\{\phi_0 + \phi_1 \widehat{X}_T\} - \{\phi_0 + \phi_1 X_T + \epsilon_{T+1}\})^2 \\
&= E\{\phi_1 X_T (u_T - 1) - \epsilon_{T+1}\}^2 \\
&= E\{\phi_1 X_T (u_T - 1)\}^2 + E(\epsilon_{T+1}^2) \\
&= \phi_1^2 E\{X_T^2 (u_T - 1)^2\} + E(\epsilon_{T+1}^2) \\
&= \phi_1^2 E\{X_T^2\} E(u_T^2 - 2u_T + 1) + E(\epsilon_{T+1}^2) \\
&= \phi_1^2 \{\text{Var}(X_T) + E^2(X_T)\} \{E(u_T^2) - 2E(u_T) + 1\} + \sigma_\epsilon^2 \\
&= \phi_1^2 \{\text{Var}(X_T) + E^2(X_T)\} \{\text{Var}(u_T) + E^2(u_T) - 2E(u_T) + 1\} + \sigma_\epsilon^2 \\
&= \phi_1^2 \{\text{Var}(X_T) + E^2(X_T)\} \text{Var}(u_T) + \sigma_\epsilon^2 \\
&= \phi_1^2 \left\{ \frac{\sigma_\epsilon^2}{1 - \phi_1^2} + \mu^2 \right\} \sigma_u^2 + \sigma_\epsilon^2,
\end{aligned}$$

where the fourth step is due to the independence of ϵ_{t+1} , u_t and X_t , the sixth step is due to the independence of u_t and X_t , the second last step is due to $E(u_t) = 1$, and the last step is because $\text{Var}(X_t) = \frac{\sigma_\epsilon^2}{1 - \phi_1^2}$ due to the stationary AR(1) process. Hence,

$$\begin{aligned}
P_e^{(h)} &= E(\widehat{X}_{T+h} - X_{T+h})^2 \\
&= E\{(\phi_0 + \phi_1 \widehat{X}_{T+h-1}) - (\phi_0 + \phi_1 X_{T+h-1} + \epsilon_{T+1})\}^2 \\
&= E\left\{\phi_1 \left(\widehat{X}_{T+h-1} - X_{T+h-1}\right) - \epsilon_{T+1}\right\}^2 \\
&= \phi_1^2 P_e^{(h-1)} + \sigma_\epsilon^2 \\
&= \phi_1^{2h-2} P_e^{(1)} + \sum_{i=0}^{h-2} \phi_1^{2i} \sigma_\epsilon^2 \\
&= \phi_1^{2h} \left\{ \frac{\sigma_\epsilon^2}{1 - \phi_1^2} + \mu^2 \right\} \sigma_u^2 + \sum_{i=0}^{h-1} \phi_1^{2i} \sigma_\epsilon^2. \tag{6.22}
\end{aligned}$$

The evaluation of the mean squared prediction error $P_e^{(h)}$ is carried out by replacing the parameters with their estimators. We comment that the common second term in (6.21) and (6.22), $\sum_{i=0}^{h-1} \phi_1^{2i} \sigma_\epsilon^2$, is the mean squared prediction error for the AR(1) model for error-free settings (e.g. Box et al., 2015, Page 152), which equals $\frac{1 - \phi_1^{2h}}{1 - \phi_1^2} \sigma_\epsilon^2$.

For an α with $0 < \alpha < 1$, then h -step $(1 - \alpha)$ -prediction interval is constructed as

$$\left[\widehat{X}_{T+h} - q_{\frac{\alpha}{2}} P_e^{(h)}, \widehat{X}_{T+h} + q_{\frac{\alpha}{2}} P_e^{(h)} \right],$$

where $q_{\frac{\alpha}{2}}$ the α -level quantile of the distribution of $\widehat{X}_{T+h} - X_{T+h}$. In practice, under normal assumption of ϵ_t and e_t , one can take $q_{\frac{\alpha}{2}}$ to be the α -level quantile of the standard normal distribution (Brockwell and Davis, 2002, Page 108).

6.4 Analysis of COVID-19 Death Rates

6.4.1 Study Objective

Using Canadian provincial COVID-19 data containing the daily confirmed cases and deaths from April 3, 2020 to May 4, 2020, we compare the times series of death rates for British Columbia, Ontario, Quebec, and Alberta, the four provinces in Canada which experience severe situations. The daily confirmed cases and fatalities are taken from “1Point3Acres.com” (<https://coronavirus.1point3acres.com/>).

In epidemiology, the mortality rate, defined as the proportion of cumulative deaths of the disease in the total number of people diagnosed with the disease (Kanchan et al., 2015), is often used to measure the severeness of an infectious disease. For COVID-19, determining the mortality rate is not trivial due to the difficulty in precisely determining the number of infected cases. Due to the limited test capacity, individuals with light symptoms are not being tested. Asymptomatic infections and the incubation period make it difficult to acquire an accurate number of infections. To circumvent this, we explore different definitions of death rates. *Definition 1* is from Baud et al. (2020) who estimated mortality rates by dividing the number of deaths on a given day by the number of patients with confirmed COVID-19 infection 14 days before, with the consideration of the maximum incubation time to be 14 days. On the other hand, the median time from symptom onset to intensive care unit admission is about 10 days ([3] in Baud et al., 2020), so we consider *Definition 2* which is the number of deaths of COVID-19 on day t divided by the number of confirmed cases at day $(t - 10)$. In comparison, we also consider *Definition 3* by calculating the death rate on day t as the ratio of the number of deaths on day t to the number of confirmed cases on the day t .

While the first two ways may help more reasonably estimate mortality rates than the third definition, these calculated rates still differ from the true mortality rates because

of under-reported cases which are primarily due to limited test capacity and undetected asymptomatic infections. To reflect the discrepancy between the *reported* and the *true* mortality rates for each province, for each definition of the mortality rate, we let $X_{1,t}$, $X_{2,t}$, $X_{3,t}$, and $X_{4,t}$, represent the *true* mortality rate on day t for British Columbia, Ontario, Quebec and Alberta, respectively; and let $X_{1,t}^*$, $X_{2,t}^*$, $X_{3,t}^*$ and $X_{4,t}^*$ denote the *reported* mortality rate on day t in British Columbia, Ontario, Quebec and Alberta, respectively. The objective is to use the reported mortality rates $\{X_{it}^* : t = 1, \dots, 31\}$ to infer the true mortality rates $X_{i,t}$ which are modeled by (6.1) separately for $i = 1, \dots, 4$. In addition, we want to forecast the true mortality rate of COVID-19 for a future time period. Due to the undetected asymptomatic cases and untested cases for light symptoms, the reported mortality rates $X_{i,t}^*$ are typically overestimated (i.e., $X_{i,t}^* \geq X_{i,t}$) for $i = 1, \dots, 4$. As there is no exact information to guide us how to characterize the relationship between X_{it}^* and X_{it} , here we conduct sensitivity studies by considering measurement error model (6.7) or (6.9). We use the observed data $X_{i,t}^*$ from April 3, 2020 to May 4, 2020, i.e., $\{X_{i,t}^* : t = 1, \dots, T_i\}$ with $T_1 = T_2 = 31$, to estimate the model parameters in (6.1) with measurement error effects accounted for, and then forecast the mortality rate of COVID-19, from May 5, 2020 to May 9, 2020, in British Columbia, Ontario, Quebec and Alberta, Canada.

6.4.2 Models Building

Figure 6.1 displays the trajectory of the mortality rates of COVID-19 in the four provinces that are obtained from the three definitions. To assess the stationarity of the X_{it}^* , we conduct the augmented Dickey–Fuller (ADF) tests (Cheung and Lai, 1995) to times series $\{X_{i,t}^* : t = 1, \dots, T\}$, or its differencing transformation $\{X_{i,(t+1)}^* - X_{i,t}^* : t = 1, \dots, T\}$ for $i = 1, \dots, 4$ in each definition. Table 6.1 presents the test statistics and p -value of the ADF test for each time series, where “TSV” represents a test statistics value.

To determine the lag value p for the autoregression model (6.1) used for the time series $\{X_{i,t} : t = 1, \dots, T_i\}$ with $T_1 = T_2 = 31$ for $i = 1, \dots, 4$, we fit the naive model (6.13) with ϵ_t^* assumed to follow a normal distribution $N(0, \sigma_\epsilon^{*2})$, and use the AIC criterion by minimizing

$$-2 \sum_{t=p}^T \log f(x_t^* | x_{t-1}^*, \dots, x_{t-p}^*) + 2p, \quad (6.23)$$

where $f(x_t^* | x_{t-1}^*, \dots, x_{t-p}^*)$ is the conditional probability of X_t^* given $X_{t-1}^*, \dots, X_{t-p}^*$. The results are summarized in Table 6.2, where no-differencing or 1-differencing is applied, the entries with “-” indicate that the corresponding model is not applicable due to the ADF test results.

We take those lag values for an AR(p) model to feature the true mortality rate $X_{i,t}$ for each definition and $i = 1, \dots, 4$. To be specific, for the British Columbia data, with Definition 1 we consider two models: AR(1) model for the time series with 1-order differencing and AR(2) model for the time series with no-differencing; with Definitions 2 and 3, we consider AR(2) and AR(1) models, respectively, for the time series with 1-order differencing. For the Ontario data, we consider AR(1) and AR(4) for the time series with 1-order differencing in Definitions 1 and 3, respectively, and AR(2) for Definition 2 with no transformation. For the Quebec data, we consider AR(1) and AR(2) models for the times series with 1-order differencing in Definitions 1 and 2, respectively. For Alberta data, we consider the AR(1) model for the times series with 1-order differencing for both Definitions 1 and 2.

6.4.3 Sensitivity Analyses

As there are no additional data available for estimating the parameters for the model (6.7) or (6.9), we conduct sensitivity analyses using the findings in the literature. Different studies showed different estimates of the asymptomatic infection rates, changing from 17.9% to 78.3% (Kimball, 2020; Day, 2020). To accommodate the heterogeneity of different studies, He et al. (2020) carried out a meta-analysis and obtained an estimate of the asymptomatic infection rate to be 46%. If under-reported confirmed cases are only caused from undetected asymptomatic cases, then $X_t = (1 - \tau_A)X_t^*$, or equivalently,

$$X_t^* = \frac{1}{1 - \tau_A} X_t, \quad (6.24)$$

where τ_A represents the rate of asymptomatic infections.

Now we use (6.24) as a starting point to conduct sensitivity analyses. In the multiplicative model (6.9), we take $\beta_0 u_t = \frac{1}{1 - \tau_A}$. With $E(u_t) = 1$, we set $\beta_0 = \frac{1}{1 - \tau_A}$ by setting $\tau_A = 46\%$, the value from the meta-analysis of He et al. (2020). To see different degrees of error, we consider σ_u^2 to take a small value, say σ_{u1}^2 , and a large value, say, σ_{u2}^2 , which is alternatively reflected by the change of the coefficient of variation, $CV = \frac{\sigma_u}{E(u_t)}$, of the error term u_t from $\sigma_{u1} \times 100\%$ to $\sigma_{u2} \times 100\%$.

When using the additive model (6.7) to characterize the measurement error process, motivated by (6.24), we set $\alpha_0 = 0$ and $\alpha_1 = \frac{1}{1 - 46\%}$, and let σ_e^2 take a small value, say σ_{e1}^2 , and a large value, say σ_{e2}^2 , to feature an increasing degree of measurement error. Due to the constraints for the parameters discussed for (6.8) and (6.10), we set the values for σ_{u1} , σ_{u2} , σ_{e1} , and σ_{e2} case by case for each definition and for each province, which are recorded in Table 6.3.

The model-fitting results are reported in Tables 6.4–6.6 for the three definitions of mortality rates, where the point estimates (EST), the associated standard errors (SE), and the p-values for the model parameters are included. Table 6.4 shows that with Definition 1, the estimates of ϕ_0 in the absolute value from the proposed method are smaller than those of naive method, while the estimates of ϕ_1 produced from the proposed and naive methods exhibit an opposite direction. As expected, the standard errors for the proposed method are generally larger than those of the naive method. However, both methods find no evidence to support that ϕ_0 and ϕ_1 are different from zero for the data of British Columbia and Ontario, suggesting that the mortality rates of these two provinces remain statistically unchanged. At the significance level 0.1, the naive method and the proposed method show different evidence for the data of Quebec and Alberta. The naive method suggests a likely downward trend with p-value 0.071 and 0.061 for testing of ϕ_0 for Quebec and Alberta, respectively. The proposed method, on the other hand, shows that ϕ_0 is insignificant for these two provinces.

Table 6.5 displays the results for Definition 2. For the British Columbia data, the estimates of the three parameters ϕ_1 , ϕ_2 and ϕ_3 produced from the proposed method are smaller than those yielded from the naive method, whereas the standard errors output from the proposed method are larger than those from the naive method. However, at the significance level 0.05, both methods find no evidence to show the significance of ϕ_0 , ϕ_1 and ϕ_2 , suggesting that the mortality rate of British Columbia remain unchanged with time. Similar findings are revealed for the Alberta data except that the parameter estimates output from the proposed method are larger than those produced from the naive method. For the Ontario and Quebec data, the revealings from the two methods are quite different. For Ontario, both methods show that ϕ_0 is insignificant and ϕ_1 is significant. The evidence of ϕ_2 , however, depends on the nature of measurement error. On the contrary, the findings for Quebec do not tend to show a definite direction, and they vary with the model form or degree of the measurement error process.

Table 6.6 shows the results for Definition 3. For the British Columbia data, the estimates produced by the proposed method are smaller than those yielded from the naive method. The standard errors output from the proposed method inflate as the degree of measurement error increases. The naive and proposed methods reveal different evidence for the significance of ϕ_0 and ϕ_1 , and the degree of measurement error affects the findings too. For the Ontario data, both methods uncover the same type of evidence for all the parameters at the significance level 0.05, except for the case with the large error under the multiplicative model.

6.4.4 Forecasting

With the fitted model for each time series in Section 6.4.3, we forecast the true mortality rate for the subsequent five days (May 5 – May 9) using the method described in Section 6.3.2. Specifically, since the true mortality rates are not observable, we “estimate” them using (6.18) and (6.19), respectively, for the measurement error models (6.7) and (6.9), and then we forecast the values of $X_{i,32}$, $X_{i,33}$, $X_{i,34}$, $X_{i,35}$, and $X_{i,36}$ using (6.20).

To quantify the forecasting performance, we calculate $P_e^{(h)}$ for $h = 1, \dots, H$ for each specified model of the mortality rates $X_{i,t}$, and we report the results, together with the total $\sum_{h=1}^H P_e^{(h)}$ in Tables 6.7–6.9, where H is set as 5. For $h = 1, \dots, H$, we report the observed prediction error $(X_{T+h} - \hat{X}_{T+h})^2$, and the expected prediction error defined in (6.21) and (6.22).

Forecasting results based on the three definitions of mortality rates are reported in Figures 6.2–6.8 for the four provinces, where the prediction results after May 4 are marked in blue and red for the measurement error models (6.7) and (6.9), respectively, together with prediction areas marked in shaded parts, as well as the prediction results obtained from the naive method by using (6.20) with naive estimates of ϕ (marked in dark yellow). In comparison, we display the reported mortality rate (in black) from Apr 3, 2020 to May 9, 2020 as well as the adjusted mortality rates obtained from (6.24) (in green); in addition, we report the fitted values using (6.17) in blue points. To compare the forecasting results in the presence of different degrees of measurement error. We report the results derived from a mild degree of measurement error in top subfigures and place those obtained from a large degree of measurement error in bottom subfigures.

The results for British Columbia are presented in Figures 6.2–6.5. With Definition 1, the methods with measurement error effects accommodated suggest that the mortality rate in the past and its forecasting values are around 4%, whereas the results obtained from the method without accounting for measurement error effects indicate that the mortality rates over time are higher than 6%. With Definition 2, the methods with or without accounting for measurement error effects reveal that the mortality rates over time are, respectively, below 3.5% and above 5%. With Definition 3, the methods with or without accounting for measurement error effects indicate that the mortality rates over time are around 3% and above 4%, respectively.

The results for Ontario are presented in Figures 6.6–6.8. With Definition 1, the methods with measurement error effects accommodated suggest that the mortality rate over time is around 7% over time, while the reported mortality rate over time is about 12.5%. With Definition 2, the methods with and without incorporating the feature of measurement error

indicate the mortality rate in the past and its forecasting values are, respectively, below 6% and around 10%. With Definition 3, the mortality rate increases over time substantially. The methods with measurement error effects accommodated suggest that the mortality rate increases from 2% to above 4% whereas the reported mortality rate shows that rate increases from below 4% to above 8%.

The results for Quebec are presented in Figures 6.9–6.10. With Definition 1 the methods with measurement error effects accommodated show that the mortality rate is around 6.5% over time, whereas the method without considering measurement error indicates the mortality rate is over 10%. With Definition 2, the methods with or without addressing the measurement error effects show that the mortality rates over time are, respectively, below 6% and above 7.5%.

The results for Alberta are presented in Figures 6.11–6.12. With Definition 1 the methods with and without measurement error accommodated suggest that the mortality rates are, respectively, around 2% and 4% over time. With Definition 2, the methods with or without addressing the measurement error effects show that the historical mortality rate and its predictions are, respectively, below 2% and above 2%.

6.4.5 Model Assessment

The specification of lag p for model (6.1) of the true mortality rates $\{X_{i,t} : t = 1, \dots, T\}$ is based on (6.23) which is derived from the reported mortality rates $\{X_{i,t}^* : t = 1, \dots, T\}$, but not from $\{X_{i,t} : t = 1, \dots, T\}$ itself. This discrepancy introduces the possibility of model misspecification when featuring the series $X_{i,t}$ using (6.1). To investigate this, we conduct a sensitivity analysis by considering the AR(p) with a different value of p for the $X_{i,t}$ from Definition 1. As Table 6.2 indicates the feasibility of using AR(1) for all four provinces, here we further employ the AR(2) model to do forecasting for the period from May 5 to May 9.

In Table 6.10, we report the observed and expected prediction errors of the forecasting using AR(2) models in comparison with AR(1) models. Comparing different lag orders of the autoregressive models, we find that in terms of the observed prediction error, the selected AR(1) models have better performance than the AR(2) models for the data of Ontario and Alberta, and the results for British Columbia and Quebec are fairly similar. It is noticed that both the observed prediction error and the expected prediction error associated with the proposed method tend to become small when the degree of measurement error increases for British Columbia, Ontario, and Quebec.

Table 6.1: The results of the augmented Dickey-Fuller test

Definition	Transformation	British Columbia		Ontario		Quebec		Alberta	
		TSV	p-value	TSV	p-value	TSV	p-value	TSV	p-value
Definition 1	X_t	-8.346	<0.01	-1.527	0.755	-1.813	0.645	-2.850	0.245
	$X_{t+1} - X_t$	-6.974	<0.01	-5.522	<0.01	-3.880	0.027	-3.516	0.059
Definition 2	X_t	-1.208	0.878	-4.294	<0.01	-2.018	0.566	-1.768	0.662
	$X_{t+1} - X_t$	-3.336	0.084	-2.599	0.342	-3.340	0.084	-3.296	0.090
Definition 3	X_t	-1.325	0.833	-2.264	0.471	0.098	0.999	-2.688	0.307
	$X_{t+1} - X_t$	-3.590	0.048	-4.584	<0.01	-2.209	0.492	-2.008	0.569

Table 6.2: The results of the augmented Dickey-Fuller test

Definition	British Columbia		Ontario		Quebec		Alberta	
	Differencing	lag p	Differencing	lag p	Differencing	lag p	Differencing	lag p
Definition 1	1 degree	1	1 degree	1	1 degree	1	1 degree	1
	no differencing	2	-	-	-	-	-	-
Definition 2	1 degree	2	no differencing	2	1 degree	2	1 degree	1
Definition 3	1 degree	1	1 degree	4	-	-	-	-

Table 6.3: The parameter values of σ_e^2 or σ_u^2 for the measurement error model (6.7) or (6.9) that are used for sensitivity analyses

Definition	Error Model	British Columbia	Ontario	Quebec	Alberta
Definition 1		AR(1)	AR(1)	AR(1)	AR(1)
	Additive (σ_e^2)	0.1	0.5	0.5	0.1
	Multiplicative (σ_u^2)	0.3	0.5	0.5	0.4
		AR(2)*	-	-	-
Definition 2		AR(2)	AR(2)*	AR(2)	AR(1)
	Additive (σ_e^2)	0.1	-	-	-
	Multiplicative (σ_u^2)	0.01	-	-	-
		0.02	-	-	-
Definition 3		AR(2)	AR(2)	AR(2)	AR(1)
	Additive (σ_e^2)	0.05	0.05	0.1	0.05
	Multiplicative (σ_u^2)	0.2	0.005	0.01	0.3
		0.5	0.005	0.01	0.6
Definition 4		AR(2)	AR(4)	-	-
	Additive (σ_e^2)	0.03	0.02	0.05	-
	Multiplicative (σ_u^2)	0.3	0.1	0.2	-
		0.6	0.2	-	-

* The time series with no differencing

Table 6.4: Definition 1: The parameter estimation under different measurement error models: the AR(1) model with “order-1 differencing” is used to fit the data of British Columbia, Ontario, Quebec and Alberta

Method	Error Degree	Parameter	British Columbia			Ontario			Quebec			Alberta		
			EST	SE	p-value	EST	SE	p-value	EST	SE	p-value	EST	SE	p-value
Naive	-	ϕ_0	-0.050	0.043	0.272	-0.215	0.243	0.384	-0.340	0.180	0.071	-0.031	0.016	0.061
		ϕ_1	0.138	0.214	0.533	0.215	0.157	0.183	0.012	0.124	0.923	0.052	0.144	0.721
The Proposed Method with Additive Error	Small ($\sigma_{\varepsilon_1}^2$)	ϕ_0	-0.027	0.025	0.313	-0.113	0.134	0.406	-0.183	0.111	0.112	-0.017	0.009	0.088
		ϕ_1	0.146	0.532	0.788	0.237	0.280	0.406	0.014	1.566	0.993	0.056	0.185	0.764
	Large ($\sigma_{\varepsilon_2}^2$)	ϕ_0	-0.027	0.025	0.298	-0.097	0.263	0.715	-0.181	0.100	0.083	-0.014	0.073	0.845
		ϕ_1	0.146	0.468	0.760	0.345	0.939	0.717	0.027	0.323	0.934	0.183	1.596	0.909
The Proposed Method with Multiplicative Error	Small ($\sigma_{\varepsilon_1}^2$)	ϕ_0	-0.027	0.024	0.286	-0.107	0.152	0.488	-0.183	0.099	0.078	-0.017	0.009	0.080
		ϕ_1	0.151	0.236	0.535	0.275	0.238	0.260	0.016	0.166	0.923	0.060	0.180	0.740
	Large ($\sigma_{\varepsilon_2}^2$)	ϕ_0	-0.025	0.024	0.308	-0.078	1.690	0.964	-0.180	0.127	0.170	-0.016	0.015	0.299
		ϕ_1	0.192	0.300	0.535	0.476	3.955	0.905	0.031	1.327	0.981	0.087	0.360	0.812

Table 6.5: Definition 2: The parameter estimation under different measurement error models: the AR(2) model with “no differencing” is used to fit the data of Ontario, the AR(1) model with “order-1 differencing” is used to fit the data of Alberta, and the AR(2) model with “order-1 differencing” is used to fit the data of British Columbia and Quebec

Method	Error Degree	Parameter	British Columbia			Ontario			Quebec			Alberta		
			EST	SE	p-value	EST	SE	p-value	EST	SE	p-value	EST	SE	p-value
Naive		ϕ_0	0.062	0.034	0.097	2.126	1.388	0.138	0.225	0.058	0.001	-0.013	0.022	0.561
	-	ϕ_1	-0.415	0.186	0.046	1.167	0.209	<0.001	-0.122	0.136	0.380	-0.124	0.172	0.477
		ϕ_2	-0.254	0.185	0.195	-0.370	0.140	0.014	-0.309	0.092	0.003	-	-	-
The Proposed Method with Additive Error	Small (σ_{e1}^2)	ϕ_0	0.034	0.020	0.114	1.146	0.759	0.144	0.174	0.042	0.000	-0.007	0.012	0.567
		ϕ_1	-0.432	0.201	0.053	1.173	0.216	<0.001	0.124	0.032	0.001	-0.131	0.185	0.486
		ϕ_2	-0.268	0.205	0.215	-0.375	0.141	0.014	-0.130	0.165	0.435	-	-	-
	Large (σ_{e2}^2)	ϕ_0	0.036	0.024	0.164	1.138	0.747	0.140	-0.327	0.096	0.002	-0.007	0.012	0.554
		ϕ_1	-0.497	0.265	0.085	1.189	0.239	<0.001	0.162	0.044	0.001	-0.158	0.247	0.529
		ϕ_2	-0.320	0.354	0.384	-0.390	0.172	0.032	0.132	0.041	0.004	-	-	-
The Proposed Method with Multiplicative Error	Small (σ_{a1}^2)	ϕ_0	0.034	0.020	0.115	1.139	0.748	0.141	-0.164	0.229	0.480	-0.007	0.012	0.564
		ϕ_1	-0.439	0.205	0.053	1.188	0.231	<0.001	-0.394	0.199	0.059	-0.144	0.205	0.487
		ϕ_2	-0.273	0.204	0.205	-0.389	0.162	0.024	0.128	0.042	0.006	-	-	-
	Large (σ_{a2}^2)	ϕ_0	0.039	0.032	0.236	1.112	0.747	0.149	0.127	0.036	0.002	-0.008	0.012	0.546
		ϕ_1	-0.584	0.339	0.111	1.255	0.503	0.020	-0.143	0.194	0.467	-0.205	0.317	0.524
		ϕ_2	-0.393	0.322	0.245	-0.451	0.510	0.384	-0.353	0.111	0.004	-	-	-

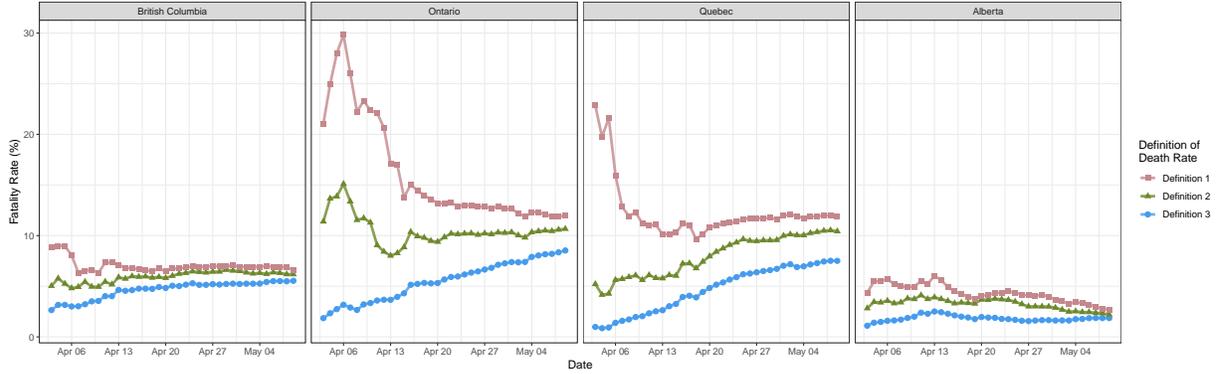


Figure 6.1: *The time series plots of the death rate with different definitions*

Table 6.6: Definition 3: The parameter estimation under different measurement error models: the AR(1) model with “order-1 differencing” is used to fit the data of British Columbia and the AR(4) model with “order-1 differencing” is used to fit the data of Ontario

Method	Error Degree	Parameter	British Columbia			Ontario		
			EST	SE	p-value	EST	SE	p-value
Naive	-	ϕ_0	0.105	0.038	0.018	0.379	0.057	<0.001
		ϕ_1	-0.207	0.077	0.020	-0.086	0.099	0.391
		ϕ_2	-	-	-	-0.287	0.106	0.012
		ϕ_3	-	-	-	-0.301	0.094	0.004
		ϕ_4	-	-	-	-0.284	0.078	0.001
The Proposed Method with Additive Error	Small (σ_{e1}^2)	ϕ_0	0.057	0.021	0.021	0.206	0.031	<0.001
		ϕ_1	-0.213	0.086	0.029	-0.088	0.100	0.383
		ϕ_2	-	-	-	-0.290	0.109	0.014
		ϕ_3	-	-	-	-0.303	0.094	0.003
		ϕ_4	-	-	-	-0.287	0.081	0.002
	Large (σ_{e2}^2)	ϕ_0	0.058	0.021	0.017	0.212	0.036	<0.001
		ϕ_1	-0.234	0.147	0.137	-0.102	0.123	0.417
		ϕ_2	-	-	-	-0.306	0.139	0.037
		ϕ_3	-	-	-	-0.318	0.107	0.006
		ϕ_4	-	-	-	-0.308	0.093	0.003
The Proposed Method with Multiplicative Error	Small (σ_{u1}^2)	ϕ_0	0.058	0.023	0.027	0.210	0.033	<0.001
		ϕ_1	-0.244	0.090	0.019	-0.097	0.107	0.375
		ϕ_2	-	-	-	-0.300	0.117	0.016
		ϕ_3	-	-	-	-0.312	0.098	0.004
		ϕ_4	-	-	-	-0.300	0.087	0.002
	Large (σ_{u2}^2)	ϕ_0	0.066	0.035	0.087	0.230	0.058	0.001
		ϕ_1	-0.401	0.219	0.092	-0.139	0.183	0.454
		ϕ_2	-	-	-	-0.347	0.213	0.116
		ϕ_3	-	-	-	-0.354	0.159	0.035
		ϕ_4	-	-	-	-0.361	0.149	0.023

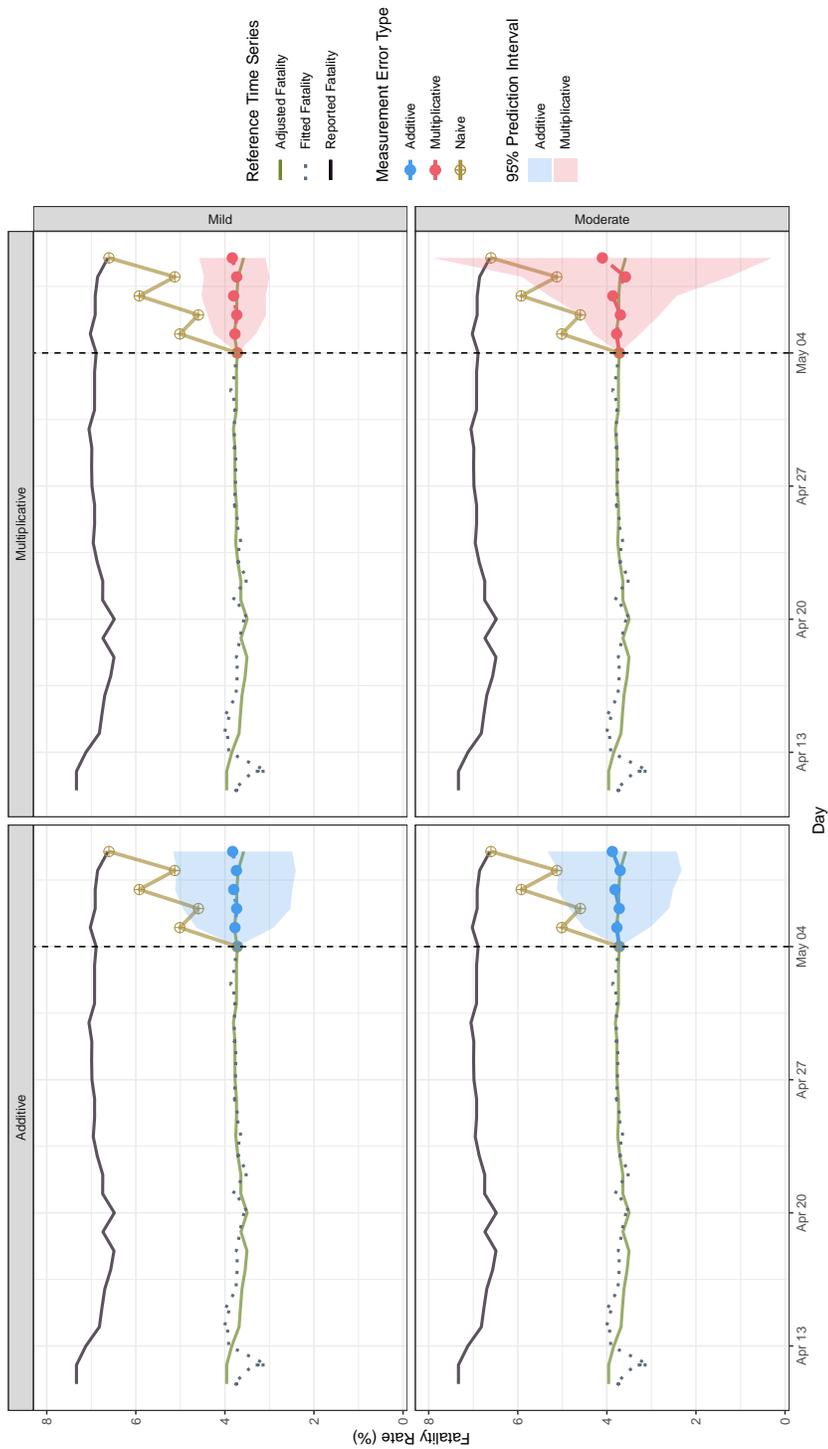


Figure 6.2: British Columbia by Definition 1 ($AR(2)$, no differencing): A 5-day forecasting of the true mortality rate (May 5 - May 9) based on the additive (in blue) or multiplicative (in red) versus the naive model (in dark yellow); the reported mortality rates (in black) and the adjusted true mortality rate accounting for the asymptomatic cases (in green).

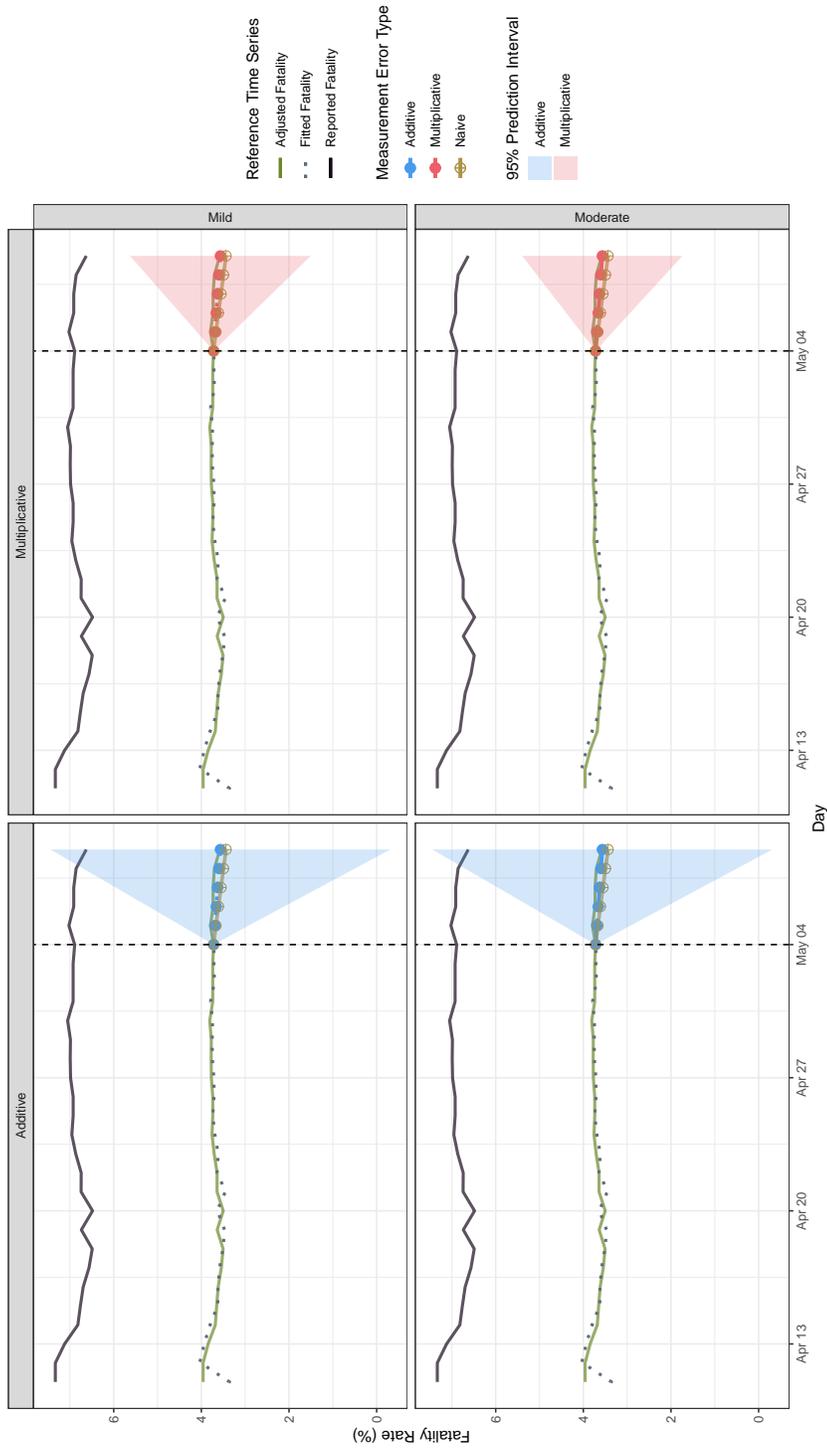


Figure 6.3: British Columbia by Definition 1 (AR(1), order-1 differencing): A 5-day forecasting of the true mortality rate (May 5 - May 9) based on the additive (in blue) or multiplicative (in red) versus the naive model (in dark yellow); the reported mortality rates (in black) and the adjusted true mortality rate accounting for the asymptomatic cases (in green).

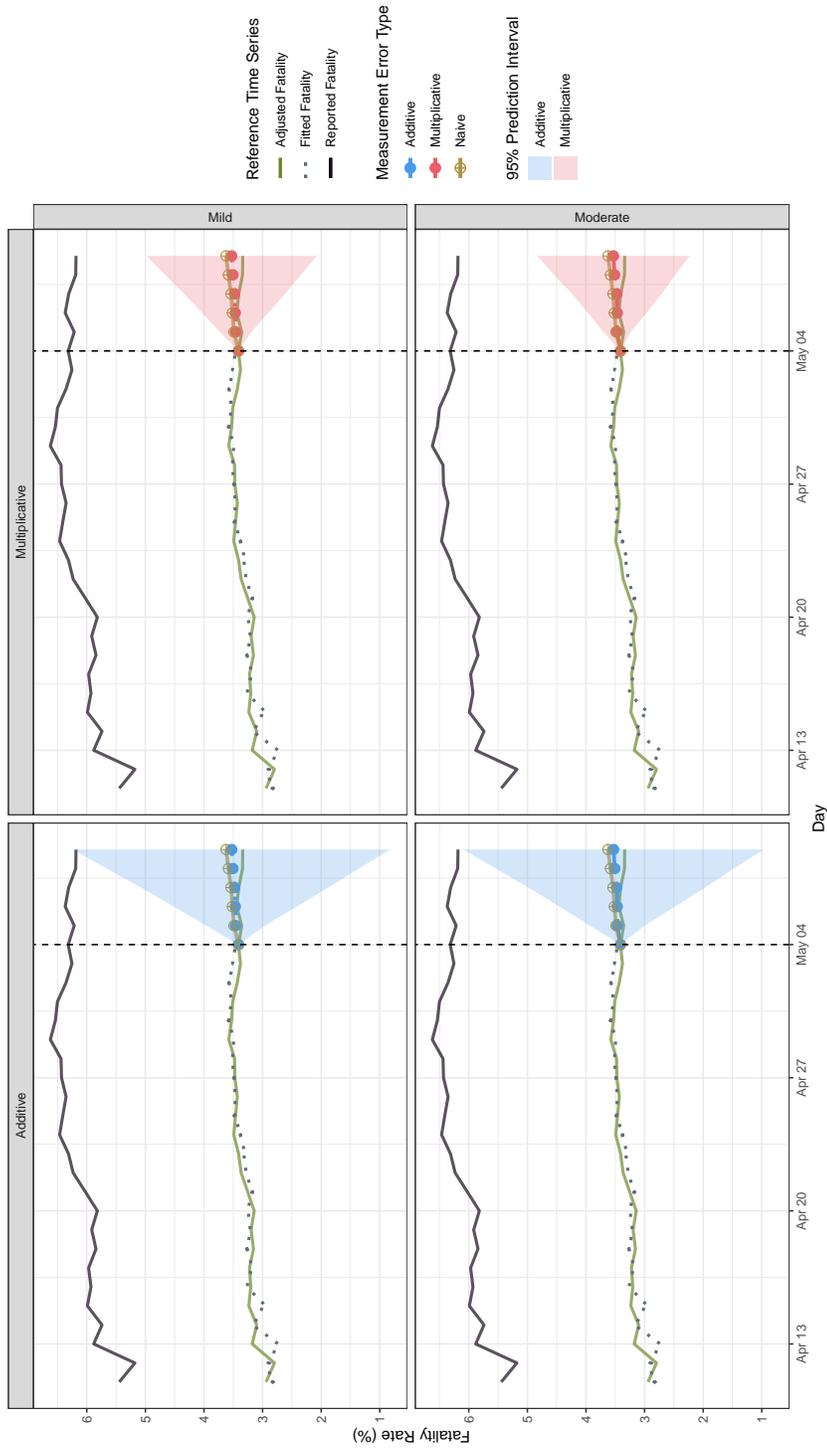


Figure 6.4: British Columbia by Definition 2 ($AR(3)$, order-1 differencing): A 5-day forecasting of the true mortality rate (May 5 - May 9) based on the additive (in blue) or multiplicative (in red) versus the naive model (in dark yellow); the reported mortality rates (in black) and the adjusted true mortality rate accounting for the asymptomatic cases (in green).

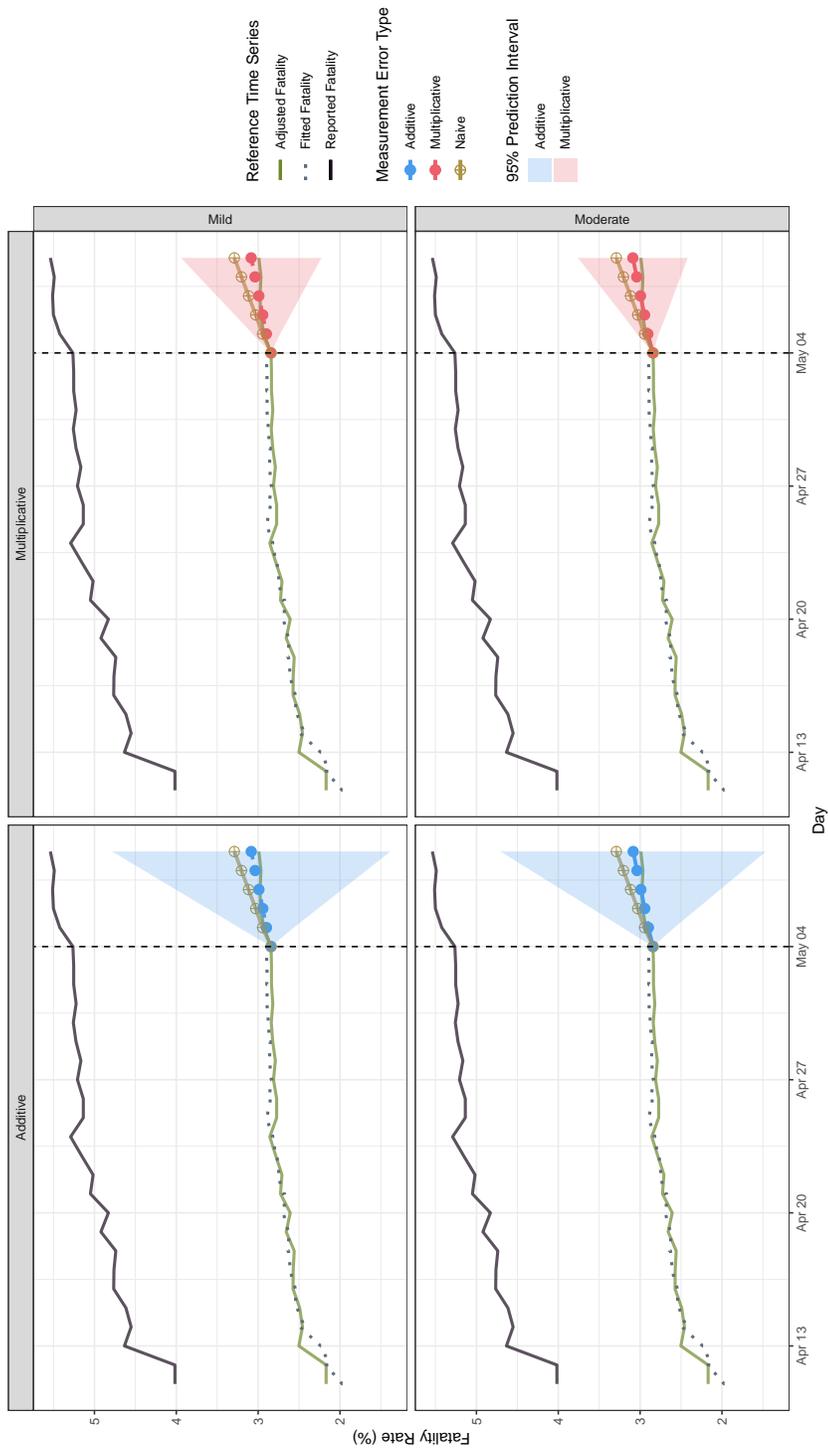


Figure 6.5: British Columbia by Definition 3 (AR(1), order-1 differencing): A 5-day forecasting of the true mortality rate (May 5 - May 9) based on the additive (in blue) or multiplicative (in red) versus the naive model (in dark yellow); the reported mortality rates (in black) and the adjusted true mortality rate accounting for the asymptomatic cases (in green).

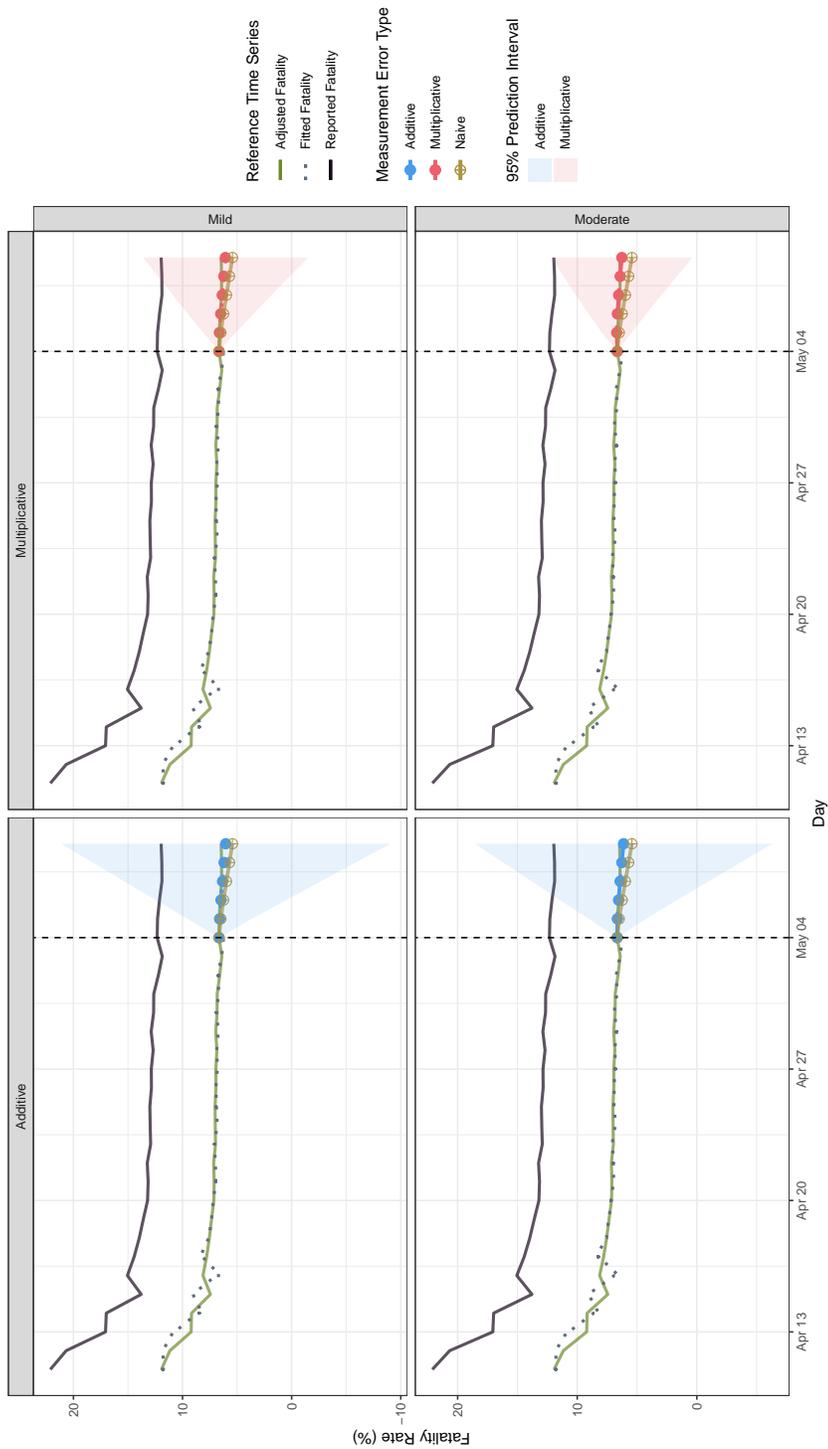


Figure 6.6: Ontario by Definition 1 (AR(1), order-1 differencing): A 5-day forecasting of the true mortality rate (May 5 - May 9) based on the additive (in blue) or multiplicative (in red) versus the naive model (in dark yellow); the reported mortality rates (in black) and the adjusted true mortality rate accounting for the asymptomatic cases (in green).

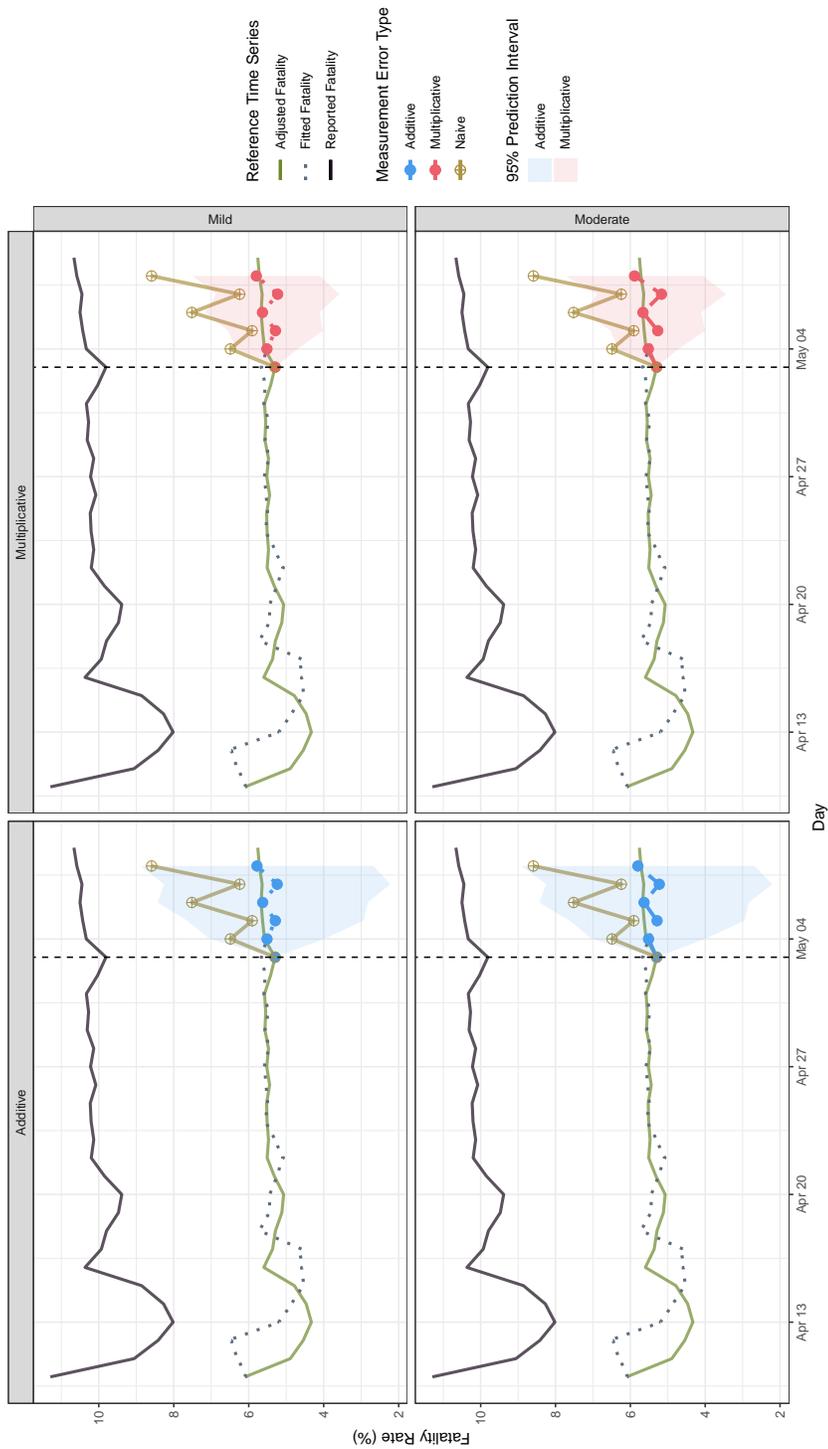


Figure 6.7: Ontario by Definition 2 ($AR(1)$, no differencing): A 5-day forecasting of the true mortality rate (May 5 - May 9) based on the additive (in blue) or multiplicative (in red) versus the naive model (in dark yellow); the reported mortality rates (in black) and the adjusted true mortality rate accounting for the asymptomatic cases (in green).

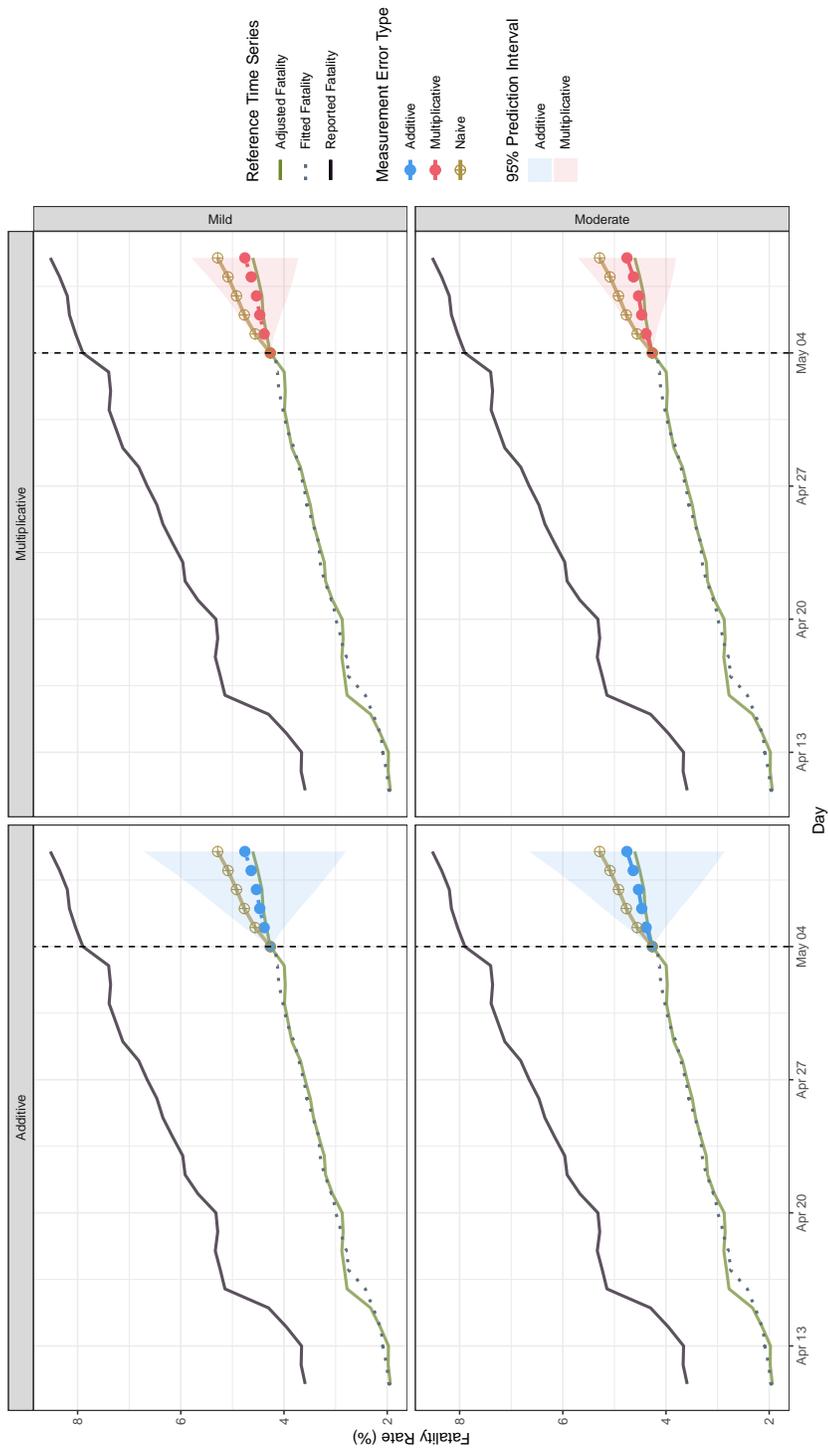


Figure 6.8: Ontario by Definition 3 ($AR(4)$, order-1 differencing): A 5-day forecasting of the true mortality rate (May 5 - May 9) based on the additive (in blue) or multiplicative (in red) versus the naive model (in dark yellow); the reported mortality rates (in black) and the adjusted true mortality rate accounting for the asymptomatic cases (in green).

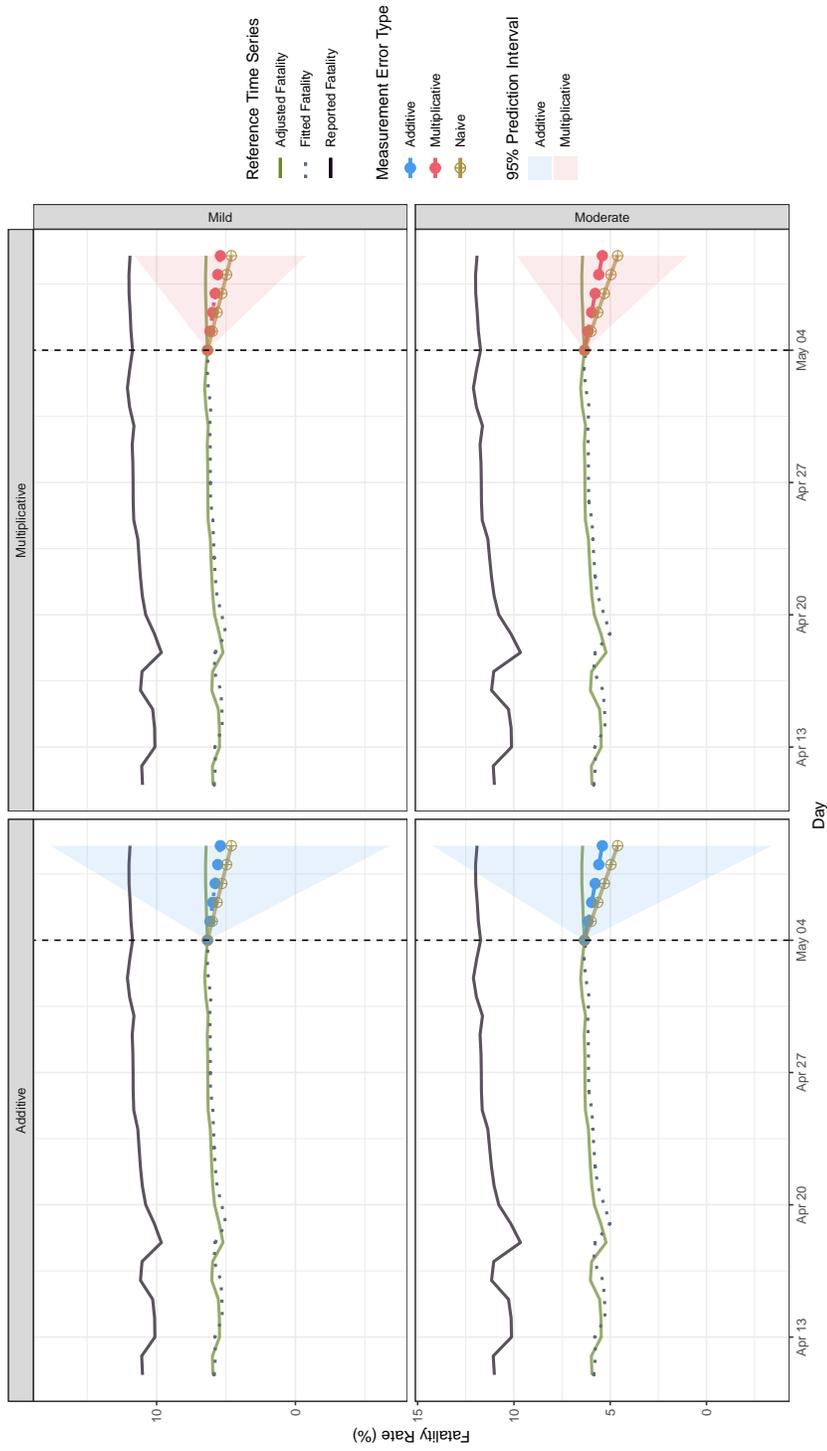


Figure 6.9: Quebec by Definition 1 ($AR(1)$, order-1 differencing): A 5-day forecasting of the true mortality rate (May 5 - May 9) based on the additive (in blue) or multiplicative (in red) versus the naive model (in dark yellow); the reported mortality rates (in black) and the adjusted true mortality rate accounting for the asymptomatic cases (in green).

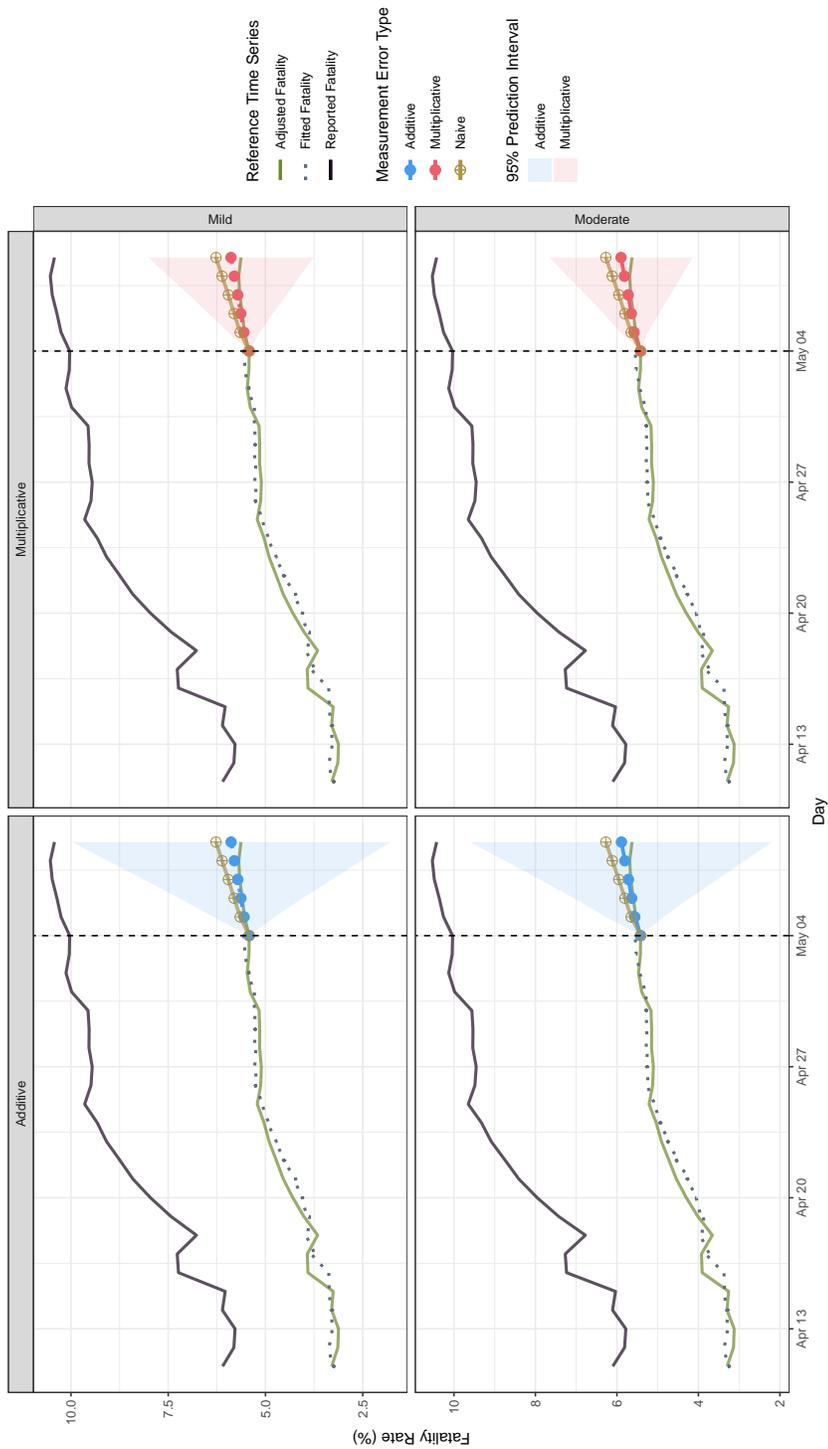


Figure 6.10: Quebec by Definition 2 (AR(2), order-1 differencing): A 5-day forecasting of the true mortality rate (May 5 - May 9) based on the additive (in blue) or multiplicative (in red) versus the naive model (in dark yellow); the reported mortality rates (in black) and the adjusted true mortality rate accounting for the asymptomatic cases (in green).

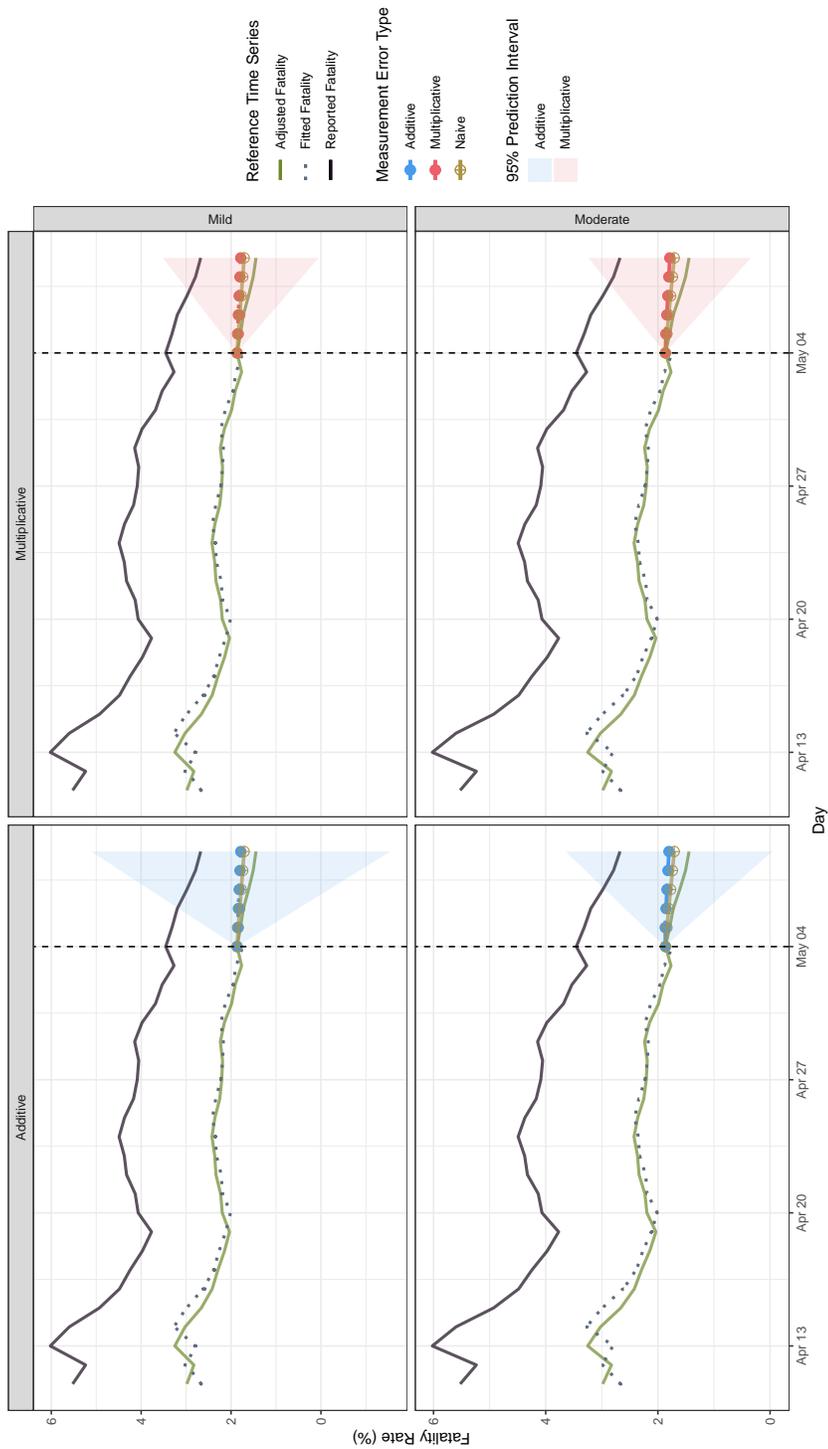


Figure 6.11: Alberta by Definition 1 (AR(1), order-1 differencing): A 5-day forecasting of the true mortality rate (May 5 - May 9) based on the additive (in blue) or multiplicative (in red) versus the naive model (in dark yellow); the reported mortality rates (in black) and the adjusted true mortality rate accounting for the asymptomatic cases (in green).

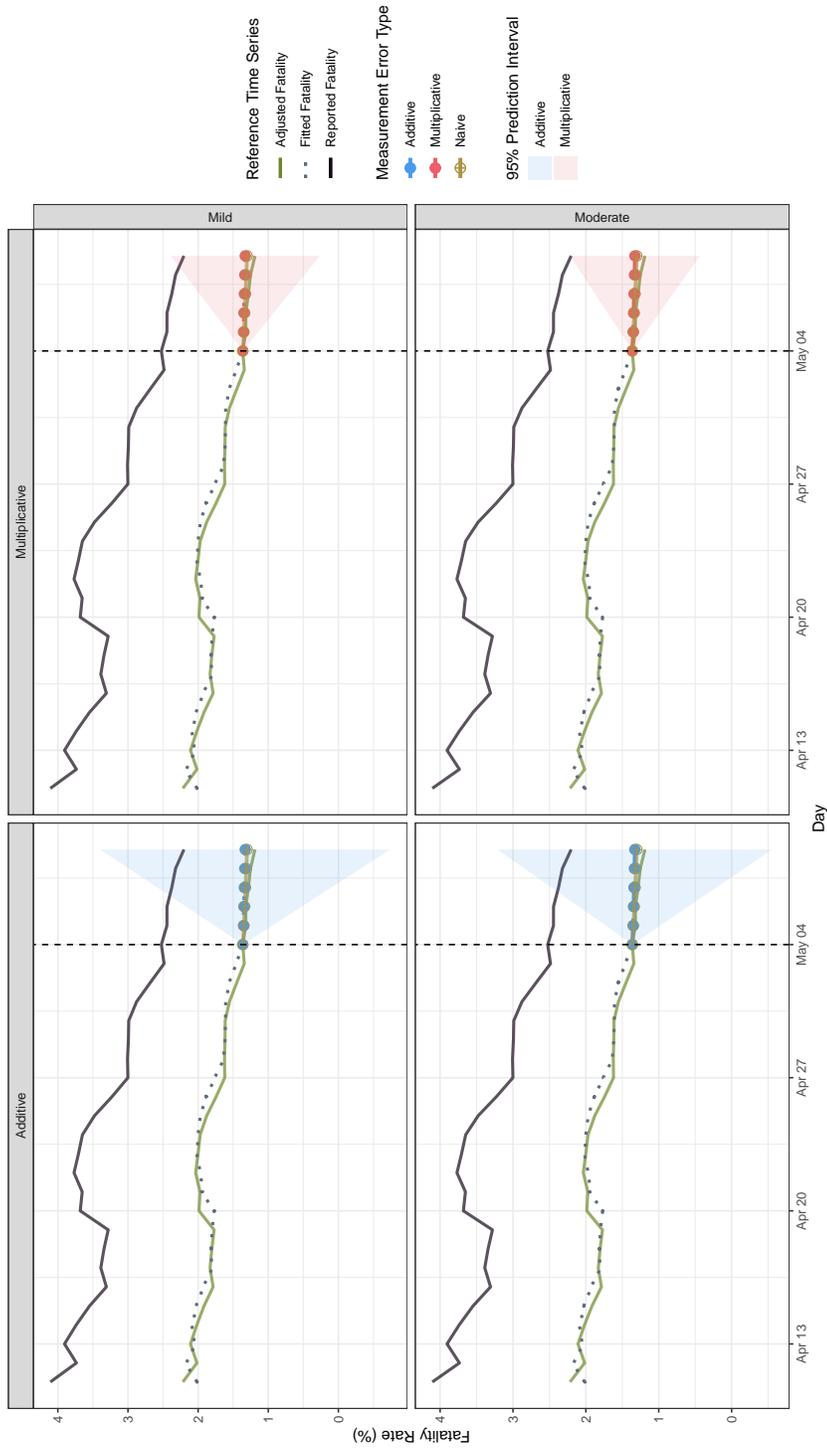


Figure 6.12: Alberta by Definition 2 ($AR(1)$, order-1 differencing): A 5-day forecasting of the true mortality rate (May 5 - May 9) based on the additive (in blue) or multiplicative (in red) versus the naive model (in dark yellow); the reported mortality rates (in black) and the adjusted true mortality rate accounting for the asymptomatic cases (in green).

Table 6.7: Definition 1: The observed prediction error and expected prediction error for different definition of death rates

Method	σ_e^2 (or σ_u^2)	Observed Prediction Error					Expected Prediction Error						
		Day 1	Day 2	Day 3	Day 4	Day 5	$\sum_{h=1}^H$ OPE(h)	Day 1	Day 2	Day 3	Day 4	Day 5	$\sum_{h=1}^H$ EPE(h)
Definition 1													
British Columbia													
Naive	-	0.017	0.006	0.017	0.058	0.081	0.178	0.069	0.081	0.081	0.082	0.083	0.396
Additive	Mild	0.011	0.001	0.005	0.027	0.035	0.078	0.066	0.078	0.078	0.080	0.080	0.382
	Moderate	0.012	0.001	0.005	0.027	0.036	0.080	0.057	0.070	0.070	0.072	0.073	0.342
Multiplicative	Mild	0.011	0.001	0.005	0.027	0.035	0.078	0.020	0.023	0.023	0.023	0.023	0.111
	Moderate	0.013	0.001	0.004	0.029	0.037	0.083	0.015	0.019	0.018	0.019	0.019	0.090
Ontario													
Naive	-	0.830	0.077	3.409	0.360	8.264	12.940	0.612	1.446	2.048	2.372	2.514	8.991
Additive	Mild	0.004	0.116	0.002	0.161	0.004	0.288	0.607	1.440	2.046	2.373	2.517	8.983
	Moderate	0.004	0.119	0.001	0.172	0.007	0.304	0.591	1.422	2.040	2.378	2.531	8.963
Multiplicative	Mild	0.004	0.119	0.001	0.171	0.007	0.302	0.176	0.420	0.602	0.704	0.754	2.655
	Moderate	0.003	0.132	0.000	0.225	0.029	0.389	0.169	0.418	0.630	0.775	0.888	2.879
Quebec													
Naive	-	0.163	0.607	1.357	2.289	3.294	7.709	1.811	1.811	1.811	1.811	1.811	9.057
Additive	Mild	0.061	0.216	0.479	0.778	1.053	2.587	1.561	1.561	1.561	1.561	1.561	7.807
	Moderate	0.060	0.215	0.478	0.776	1.051	2.580	0.811	0.811	0.811	0.811	0.811	4.057
Multiplicative	Mild	0.061	0.216	0.479	0.778	1.053	2.586	0.399	0.399	0.399	0.399	0.399	1.995
	Moderate	0.060	0.215	0.477	0.776	1.050	2.578	0.205	0.205	0.205	0.205	0.205	1.025
Alberta													
Naive	-	0.002	0.007	0.027	0.055	0.070	0.160	0.125	0.125	0.125	0.125	0.125	0.627
Additive	Mild	0.004	0.012	0.044	0.087	0.115	0.262	0.115	0.115	0.115	0.115	0.115	0.577
	Moderate	0.006	0.017	0.052	0.098	0.129	0.302	0.035	0.035	0.035	0.035	0.035	0.177
Multiplicative	Mild	0.004	0.012	0.044	0.087	0.115	0.263	0.031	0.031	0.031	0.031	0.031	0.157
	Moderate	0.005	0.013	0.045	0.089	0.118	0.270	0.022	0.022	0.022	0.022	0.022	0.109

Table 6.8: Definition 2: The observed prediction error and expected prediction error for different definition of death rates

Method	σ_e^2 (or σ_u^2)	Observed Prediction Error					Expected Prediction Error						
		Day 1	Day 2	Day 3	Day 4	Day 5	Day 1	Day 2	Day 3	Day 4	Day 5	$\sum_{h=1}^H$ EPE(h)	
Definition 2													
British Columbia													
Naive	-	0.015	0.015	0.032	0.043	0.020	0.126	0.167	0.167	0.167	0.167	0.167	0.834
Additive	Mild	0.010	0.005	0.011	0.011	0.000	0.037	0.157	0.157	0.157	0.157	0.157	0.783
	Moderate	0.010	0.005	0.011	0.011	0.000	0.037	0.154	0.157	0.157	0.157	0.157	0.784
Multiplicative	Mild	0.010	0.005	0.011	0.011	0.000	0.037	0.044	0.044	0.044	0.044	0.044	0.222
	Moderate	0.010	0.005	0.011	0.011	0.000	0.037	0.034	0.035	0.035	0.035	0.035	0.174
Ontario													
Naive	-	0.020	0.087	0.196	0.521	1.059	1.884	2.643	2.649	2.649	2.649	2.649	13.117
Additive	Mild	0.001	0.004	0.007	0.056	0.175	0.243	2.264	2.391	2.399	2.399	2.399	11.853
	Moderate	0.000	0.000	0.000	0.023	0.110	0.134	1.453	1.626	1.646	1.649	1.649	8.023
Multiplicative	Mild	0.000	0.002	0.003	0.044	0.152	0.201	0.558	0.599	0.603	0.603	0.603	2.965
	Moderate	0.004	0.010	0.014	0.000	0.035	0.063	0.270	0.331	0.345	0.348	0.348	1.642
Quebec													
Naive	-	0.013	0.044	0.086	0.183	0.413	0.739	0.174	0.176	0.191	0.192	0.193	0.926
Additive	Mild	0.000	0.001	0.003	0.013	0.065	0.081	0.163	0.165	0.181	0.182	0.183	0.874
	Moderate	0.000	0.002	0.003	0.014	0.068	0.087	0.130	0.133	0.149	0.151	0.153	0.716
Multiplicative	Mild	0.000	0.002	0.003	0.013	0.066	0.084	0.044	0.045	0.049	0.049	0.050	0.236
	Moderate	0.002	0.003	0.004	0.017	0.073	0.098	0.030	0.030	0.033	0.034	0.034	0.162
Alberta													
Naive	-	0.001	0.000	0.002	0.003	0.012	0.019	0.047	0.047	0.047	0.047	0.047	0.236
Additive	Mild	0.001	0.001	0.003	0.007	0.019	0.031	0.044	0.045	0.045	0.045	0.045	0.223
	Moderate	0.001	0.001	0.003	0.006	0.019	0.031	0.036	0.037	0.037	0.037	0.037	0.185
Multiplicative	Mild	0.001	0.001	0.003	0.006	0.019	0.031	0.012	0.012	0.012	0.012	0.012	0.059
	Moderate	0.001	0.001	0.003	0.006	0.019	0.030	0.008	0.008	0.008	0.008	0.008	0.042

Table 6.9: Definition 3: The observed prediction error and expected prediction error for different definition of death rates

Method	σ_e^2 (or σ_u^2)	Observed Prediction Error					Expected Prediction Error					
		Day 1	Day 2	Day 3	Day 4	Day 5	Day 1	Day 2	Day 3	Day 4	Day 5	$\sum_{h=1}^H$ EPE(h)
Definition 3												
British Columbia												
Naive	-	0.000	0.003	0.020	0.057	0.090	0.170	0.030	0.031	0.031	0.031	0.155
Additive	Mild	0.001	0.001	0.000	0.005	0.009	0.016	0.029	0.030	0.030	0.030	0.151
	Moderate	0.001	0.001	0.000	0.005	0.009	0.016	0.026	0.028	0.028	0.028	0.137
Multiplicative	Mild	0.001	0.001	0.000	0.005	0.009	0.016	0.007	0.008	0.008	0.008	0.038
	Moderate	0.001	0.001	0.000	0.006	0.010	0.017	0.005	0.005	0.005	0.005	0.023
Ontario												
Naive	-	0.048	0.132	0.243	0.333	0.464	1.219	0.039	0.039	0.039	0.042	0.202
Additive	Mild	0.002	0.004	0.011	0.017	0.024	0.058	0.039	0.039	0.039	0.042	0.200
	Moderate	0.002	0.004	0.011	0.016	0.023	0.057	0.036	0.036	0.036	0.039	0.187
Multiplicative	Mild	0.002	0.004	0.011	0.016	0.023	0.057	0.011	0.011	0.011	0.012	0.056
	Moderate	0.002	0.004	0.011	0.015	0.023	0.055	0.009	0.009	0.009	0.010	0.048

Table 6.10: The observed prediction error and expected prediction error for different lag order of autoregressive models

Method	σ_e^2 (or σ_u^2)	Model	Observed Prediction Error					Expected Prediction Error						
			Day 1	Day 2	Day 3	Day 4	Day 5	Day 1	Day 2	Day 3	Day 4	Day 5	$\sum_{h=1}^H$ EPPE(h)	
British Columbia														
Naive	-		0.015	0.015	0.032	0.043	0.020	0.126	0.167	0.167	0.167	0.167	0.167	0.834
Mild			0.010	0.005	0.011	0.011	0.000	0.037	0.157	0.157	0.157	0.157	0.783	
Additive	Moderate	AR(1)^a	0.010	0.005	0.011	0.011	0.000	0.037	0.154	0.157	0.157	0.157	0.784	
Mild			0.010	0.005	0.011	0.011	0.000	0.037	0.044	0.044	0.044	0.044	0.222	
Multiplicative	Moderate		0.010	0.005	0.011	0.011	0.000	0.037	0.034	0.035	0.035	0.035	0.174	
Naive	-		0.016	0.014	0.031	0.042	0.019	0.122	0.161	0.165	0.167	0.167	0.828	
Mild			0.010	0.005	0.010	0.010	0.000	0.035	0.151	0.155	0.157	0.157	0.777	
Additive	Moderate	AR(2)	0.010	0.005	0.010	0.010	0.000	0.035	0.151	0.155	0.157	0.157	0.778	
Mild			0.010	0.005	0.010	0.010	0.000	0.035	0.043	0.044	0.044	0.044	0.220	
Multiplicative	Moderate		0.010	0.005	0.010	0.010	0.000	0.034	0.034	0.035	0.035	0.035	0.173	
Ontario														
Naive	-		0.020	0.087	0.196	0.521	1.059	1.884	2.527	2.643	2.649	2.649	2.649	13.117
Mild			0.001	0.004	0.007	0.056	0.175	0.243	2.264	2.391	2.399	2.399	11.853	
Additive	Moderate	AR(1)^a	0.000	0.000	0.000	0.023	0.110	0.134	1.453	1.626	1.646	1.649	8.023	
Mild			0.000	0.002	0.003	0.044	0.152	0.201	0.558	0.599	0.603	0.603	2.965	
Multiplicative	Moderate		0.004	0.010	0.014	0.000	0.035	0.063	0.270	0.331	0.345	0.348	1.642	
Naive	-		0.073	0.107	0.240	0.550	1.111	2.081	2.517	2.648	2.649	2.649	13.111	
Mild			0.029	0.014	0.026	0.083	0.227	0.379	2.256	2.398	2.399	2.399	11.851	
Additive	Moderate	AR(2)	0.045	0.008	0.031	0.063	0.221	0.368	1.470	1.658	1.649	1.649	8.072	
Mild			0.034	0.012	0.027	0.076	0.222	0.370	0.571	0.606	0.603	0.603	2.986	
Multiplicative	Moderate		0.085	0.001	0.071	0.024	0.310	0.491	0.454	0.469	0.469	0.469	2.103	
Quebec														
Naive	-		0.163	0.607	1.357	2.289	3.294	7.709	1.811	1.811	1.811	1.811	9.057	
Mild			0.061	0.216	0.479	0.778	1.053	2.587	1.561	1.561	1.561	1.561	7.807	
Additive	Moderate	AR(1)^a	0.060	0.215	0.478	0.776	1.051	2.580	0.811	0.811	0.811	0.811	4.057	
Mild			0.061	0.216	0.479	0.778	1.053	2.586	0.399	0.399	0.399	0.399	1.995	
Multiplicative	Moderate		0.060	0.215	0.477	0.776	1.050	2.578	0.205	0.205	0.205	0.205	1.025	
Naive	-		0.129	0.524	1.226	2.115	3.085	7.079	1.746	1.746	1.809	1.811	8.921	
Mild			0.052	0.195	0.446	0.734	1.002	2.429	1.375	1.375	1.447	1.451	7.096	
Additive	Moderate	AR(2)	0.032	0.109	0.247	0.396	0.519	1.303	0.413	0.413	0.407	0.402	2.043	
Mild			0.051	0.190	0.438	0.723	0.988	2.390	0.345	0.345	0.356	0.357	1.760	
Multiplicative	Moderate		0.038	0.141	0.333	0.560	0.774	1.847	0.332	0.332	0.234	0.234	1.319	
Alberta														
Naive	-		0.002	0.007	0.027	0.055	0.070	0.160	0.125	0.125	0.125	0.125	0.627	
Mild			0.004	0.012	0.044	0.087	0.115	0.262	0.115	0.115	0.115	0.115	0.577	
Additive	Moderate	AR(1)^a	0.006	0.017	0.052	0.098	0.129	0.302	0.035	0.035	0.035	0.035	0.177	
Mild			0.004	0.012	0.044	0.087	0.115	0.263	0.031	0.031	0.031	0.031	0.157	
Multiplicative	Moderate		0.005	0.013	0.045	0.089	0.118	0.270	0.022	0.022	0.022	0.022	0.109	
Naive	-		0.003	0.010	0.033	0.064	0.081	0.191	0.122	0.122	0.125	0.125	0.621	
Mild			0.005	0.016	0.051	0.097	0.127	0.296	0.112	0.112	0.115	0.115	0.570	
Additive	Moderate	AR(2)	0.006	0.018	0.056	0.104	0.136	0.320	0.081	0.081	0.085	0.085	0.419	
Mild			0.005	0.016	0.052	0.099	0.129	0.301	0.030	0.031	0.031	0.031	0.155	
Multiplicative	Moderate		0.006	0.019	0.059	0.109	0.141	0.334	0.022	0.022	0.022	0.022	0.109	

^a The selected model

Chapter 7

Summary and Discussion

In this thesis, we investigate several important research problems concerning correlated responses with measurement error or misclassification. The results in this thesis have been or will be prepared as papers for dissemination. The research in Chapter 2 has been prepared as a paper, [Zhang and Yi \(2020b\)](#), and has been accepted by *Statistics in Medicine*; the results in Chapter 3 have been written up as a paper, [Zhang and Yi \(2020c\)](#), which has been invited by *Statistical Methods in Medical Research* for revision; the results in Chapter 4 have been included in the paper, [Zhang and Yi \(2020a\)](#), which has been submitted for publication; the results in Chapter 5 are being prepared as the paper, [Zhang and Yi \(2020e\)](#), which is to be submitted for publication soon; the results in Chapter 6 have already been wrapped up as the paper, [Zhang and Yi \(2020d\)](#), and submitted for publication. Below we present a summary for each chapter with discussions.

Chapter 2

When jointly modeling the mixed type of continuous and binary responses, we often encounter responses that are subject to measurement error and misclassification. To remove the bias resulting from the mismeasurement, it is necessary to address both measurement error and misclassification simultaneously. In this chapter, we develop two inference approaches to account for the effects due to mismeasurement in responses under latent variable models. The induced likelihood method can be easily implemented by R function `optim()` and the EM algorithm has the advantage of dealing with associated integrals by employing a complete likelihood formulation.

Although measurement error and misclassification is an inevitable issue in practice, such features are often ignored in genetic association studies. Even in the statistical literature,

available work mainly focuses on a single type of mismeasurement in responses, either measurement error or misclassification but not both. In this chapter, we propose two valid methods to account for measurement error and misclassification in mixed continuous and discrete responses. Our methods can be applied to handle error-contaminated data arising from genomewide-association studies for which mixed responses with a continuous variable and a binary variable may be subjected to mismeasurement.

Our development is carried out for the generalized linear mixed model (2.1) where a common random effect u_i is introduced to feature the mean structure of the two response components. More generally, one may use different random effects, say u_{i1} and u_{i2} , to describe the mean structure of Y_{i1} and Y_{i2} , respectively. That is, we write model (2.1) as

$$\begin{bmatrix} g_1(\mu_{i1}) \\ g_2(\mu_{i2}) \end{bmatrix} = \begin{bmatrix} X_i^T \beta_1 \\ X_i^T \beta_2 \end{bmatrix} + \begin{bmatrix} u_{i1} \\ u_{i2} \end{bmatrix}, \quad (7.1)$$

where u_{i1} and u_{i2} are random effects, and other symbols are defined in the same way as for (2.1). Then we set the random effects vector u to be $\{(u_{i1}, u_{i2}) : i = 1, \dots, n\}$ and modify the development accordingly.

Finally, in the development here, random effects u are assumed to have the covariance structure $\sigma_g^2 R$ with a pre-specified matrix R and an unknown σ_g^2 . Letting R be pre-specified allows us to incorporate a priori information of the study. In circumstances where R is impossible to be feasibly prespecified, we write the covariance matrix of random effects to be a single matrix, say \tilde{R} , which may contain multiple parameters rather than a single parameter σ_g^2 considered here. Then we carry out the inferential procedures similar to the development here by replacing σ_g^2 in the parameter vector θ in Section 2.2.1 with the parameters in \tilde{R} .

Chapter 3

Error-contaminated mixed responses with a continuous and a binary variable present a new challenge in joint modeling and analysis of multiple responses. In this chapter, we develop a generalized estimating equation approach to incorporate the dependence among responses and develop an insertion strategy to adjust for the effects of mismeasurement in responses. We propose valid estimators that apply when either internal validation data or external validation data are available. Our methods are robust to model misspecification and produce small finite sample biases. We develop a weighted estimator to improve the efficiency of parameter estimation in the presence of internal validation data.

The generalized estimation equation is robust to model misspecification at the price of the efficiency loss. To overcome this disadvantage, in addition to the weighted estimators

proposed in Section 3.3.3, other strategies may also be considered. For example, [Hall and Severini \(1998\)](#) proposed the extended generalized estimating equation (EGEE) based on the idea of extended Quasi-Likelihood. The EGEE has better efficiency than the original GEE approach ([Prentice and Zhao, 1991](#)) in some scenarios. The EGEE approach can be easily adapted in our method with minor modifications in the estimation part.

Measurement error and misclassification are inevitable in many cases. In this chapter, we propose several methods to address response mismeasurement in different types of study designs. We have shown that under certain regularity conditions, the proposed estimators are asymptotically normal and consistent. The methods are fast to implement and can apply for various settings.

Chapter 4

Identifying interactions among genetic variants is important in the analysis of gene networks. In this chapter, we develop a generalized network model to facilitate the relationship between genetic variants with a complex structure and the mixed responses via a two-step procedure. We further extend the development to handle data with measurement error and misclassification in responses. Theoretical justifications are provided to ensure the validity of the proposed method, and numerical studies demonstrate satisfactory finite sample performance of the proposed method.

In the development here, we consider continuous covariates that are featured by the Gaussian graphical model. It is interesting to generalize our method to accommodate discrete covariates or mixed covariates with both discrete and continuous components.

Our methods focus on addressing the effects due to mismeasurement in mixed bivariate responses, where covariates are assumed to be precisely measured. It is interesting to extend our work here to handle data which contain error-contaminated covariates, in addition to having mismeasured responses. In such a circumstance, adjusting the effects of measurement error in covariates is necessary for the first step for identifying the network structure for the true covariates. This research warrants exploration in depth.

Chapter 5

Zero-inflated Poisson models are useful in cancer genomics studies, which are, however, challenged by the presence of measurement error. While this problem is important, not much work has been available. We provide a general strategy in dealing with error-contaminated count data and proposed a flexible modeling scheme for measurement error in count data. We introduce a mixture model to facilitate an add-in process and a leave-out process for characterizing different types of measurement error associated count data.

We explore the effects of different measurement error models on the analysis. Numerical studies demonstrate satisfactory performance of the proposed method.

The development in this chapter can be modified to address the measurement error of count data in other models. For example, besides the zero-inflated model, the hurdle model (Mullahy, 1986) is also frequently used to account for excessive zeros in count data. Our Bayesian method can be adapted to suit the hurdle model. Sometimes, it is interesting to consider the overdispersion in count data. Our method can be further extended to deal with zero-inflated Negative Binomial models (Yau et al., 2003).

Chapter 6

We investigate the impact of measurement error on time series analysis under autoregressive models and establish analytic results under the additive and multiplicative measurement error models. We propose an estimating equation method to correct for the biases induced from the naive analysis which disregards the differences between the true measurements and their surrogate measurements. We rigorously establish the theoretical results for the proposed method. As a genuine application, we apply to the proposed method to analyze the mortality rates of COVID-19 data in four provinces, British Columbia, Ontario, Quebec, and Alberta, which have the most severe virus outbreaks in Canada. The real data analysis clearly demonstrates that incorporating measurement error in the analysis can uncover various different results.

Our method has the flexibility or robustness in that distribution assumptions are required to describe the measurement error process as well as the time series autoregressive process. While our research is motivated by the faulty nature of COVID-19 data, the proposed method can be applied to handle other problems related to error-contaminated time series. Our development here is directed to using autoregressive models to delineate time series data. The same principles can be applied to other model forms such as moving average models or autoregressive moving average models which may be used to handle error-prone time series data, where technical details can be more notationally involved.

When checking the stationarity of time series, we apply the ADF test to the observed time series X_t^* , which is mainly driven by the unavailability of the true values of X_t , as well as the fact that the weakly stationarity of observed time series implies the weakly stationarity of the true time series if measurement error is featured with (6.7) or (6.9). It is interesting to rigorously develop a formal test similar to the ADF test to handle time series subject to measurement error.

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APPENDICES

In this part, we include supplementary materials associated with Chapters 2–6, including regularity conditions, the proofs of the theoretical results, and additional calculations or discussions.

Appendix A

Supplement Materials for Chapter 2

A.1 Identifiability Issue

Model (2.8) may incur identifiability issues in some circumstances. For example, consider that $g_1(x) = x$ for (2.1) and e_i follows a normal distribution for (2.8). Then, the first component of (2.1) is equivalent to

$$Y_{i1} = \beta_{10} + \beta_{11}^T X_i + \epsilon_i, \quad (\text{A.1})$$

where β_{10} is the first element of β_1 in (2.1) and β_{11}^T is the remaining vector, and ϵ_i is independent of X_i and follows normal distribution with zero mean and variance σ^2 .

Plugging (A.1) into the additive measurement error model (2.8) gives

$$Y_{i1}^* = \beta_0^* + \beta_1^{*T} X_i + \gamma_2 f(Y_{i2}) + \gamma_3 Z_i + e_i^*,$$

where $e_i^* = e_i + \epsilon_i$ is independent of $\{X_i, Y_{i2}, Z_i\}$ and follows $N(0, \sigma^2 + \sigma_e^2)$, and

$$\beta_0^* = \gamma_0 + \gamma_1 \beta_{10}; \quad \beta_1^{*T} = \gamma_1 \beta_1^T. \quad (\text{A.2})$$

(A.2) shows that based on the observed data, we are not able to separate γ_0 from $\gamma_1 \beta_{10}$, γ_1 from β_1 , and σ^2 from σ_e^2 .

To overcome model nonidentifiability, we may add extra constraints on each group of parameters as commented by Yi (2017, Page 52). For example, we may specify $\gamma_0 = -\frac{1}{2}\gamma_2$, $\gamma_1 = 1$ and $\sigma^2 = \sigma_e^2$, which is equivalent to specifying $f(t) = 2t - 1$, $\gamma_0 = 0$, $\gamma_1 = 1$ and $\sigma^2 = \sigma_e^2$.

A.2 Gaussian Quadrature Approximation of the Expectation

In this section, we illustrate how to approximate

$$E_{u_i, Y_{i1}, Y_{i2}} \{g(Y_{i1}^*, Y_{i2}^*, Y_{i1}, Y_{i2}, u_i; \theta)\}, \quad (\text{A.3})$$

using Gaussian-Hermite Quadrature.

Define the notation

$$\begin{aligned} L_{y2}(y_{i1}, u_i, y_{i1}^*, y_{i2}^*) &= \int_{y_{i2}} g(y_{i1}^*, y_{i2}^*, y_{i1}, y_{i2}, u_i; \theta) f(y_{i2}|u_i, x_i) dy_{i2}, \\ L_{y1}(u_i, y_{i1}^*, y_{i2}^*) &= \int_{y_{i1}} L_{y2}(y_{i1}, u_i, y_{i1}^*, y_{i2}^*) f(y_{i1}|u_i, x_i) dy_{i1}, \\ L_u(y_{i1}^*, y_{i2}^*) &= \int_{u_i} L_{y1}(u_i, y_{i1}^*, y_{i2}^*) f(u_i|x_i) du_i. \end{aligned}$$

Since Y_{i2} is a binary variable following Bernoulli distribution, we can compute the exact expectation,

$$\begin{aligned} L_{y2}(y_{i1}, u_i, y_{i1}^*, y_{i2}^*) &= g(y_{i1}^*, y_{i2}^*, y_{i1}, y_{i2} = 1, u_i; \theta) f(y_{i2} = 1|u_i) \\ &\quad + g(y_{i1}^*, y_{i2}^*, y_{i1}, y_{i2} = 0, u_i; \theta) f(y_{i2} = 0|u_i). \end{aligned}$$

Consider the case where R is an diagonal matrix, using Gaussian Quadrature, we can approximate

$$\begin{aligned} L_{y1}(u_i, y_{i1}^*, y_{i2}^*) &\approx \frac{1}{\sqrt{\pi}} \sum_{j=1}^{S(y)} w_j^{(y)} L_{y2}(y_{i1} = \sqrt{2}\sigma v_j^{(y)} + \beta_1^T x_i + u_i, u_i, y_{i1}^*, y_{i2}^*), \\ L_u(y_{i1}^*, y_{i2}^*) &\approx \frac{1}{\sqrt{\pi}} \sum_{j=1}^{S(u)} w_j^{(u)} L_{y1}(u_i = \sqrt{2R_{ii}}\sigma_g v_j^{(u)}, y_{i1}^*, y_{i2}^*). \end{aligned}$$

where $v_j^{(y)}, v_j^{(u)}$ are the roots of the Hermite polynomial $H_n(x)$ for $j = 1, 2, \dots, n$ ([Abramowitz and Stegun, 1972](#), Page 890), and $w_j^{(y)}, w_j^{(u)}$ are the associated weights given by

$$w_j^{(y)} = \frac{2^{n-1}n!\sqrt{\pi}}{n^2[H_{n-1}(v_j^{(y)})]^2} \quad \text{and} \quad w_j^{(u)} = \frac{2^{n-1}n!\sqrt{\pi}}{n^2[H_{n-1}(v_j^{(u)})]^2}.$$

Based on the derivation above, the expectation (A.3) can be approximated as

$$\frac{1}{\pi} \sum_{j=1}^{S^{(u)}} \sum_{k=1}^{S^{(y)}} w_j^{(u)} w_k^{(y)} \left\{ g(y_{i1}^*, y_{i2}^*, y_{i1}, y_{i2} = 1, u_i; \theta) f(y_{i2} = 1 | u_i) \right. \\ \left. + g(y_{i1}^*, y_{i2}^*, y_{i1}, y_{i2} = 0, u_i; \theta) f(y_{i2} = 0 | u_i) \right\} \Bigg|_{\substack{y_{i1} = \sqrt{2} \sigma v_k^{(y)} + \beta_1^T x_i + u_i, \\ u_i = \sqrt{2 R_{ii}} \sigma_g v_j^{(u)}}}. \quad .$$

Appendix B

Proofs of the Results in Chapter 3

B.1 Proof of $E\{U_i^{**}(\theta)|Y_{i1}, Y_{i2}, X_i\} = U_i(\theta)$

Step 1: First, we show that

$$E(Y_{i2}^{**}|Y_{i1}, Y_{i2}, X_i) = Y_{i2}. \quad (\text{B.1})$$

Indeed, by the definition of Y_{i2}^{**} ,

$$\begin{aligned} E(Y_{i2}^{**}|Y_{i1}, Y_{i2} = j, X_i) &= E\left(\frac{Y_{i2}^* - \pi_{i0}}{1 - \pi_{i0} - \pi_{i1}} \middle| Y_{i1}, Y_{i2} = j, X_i\right) \\ &= \frac{E(Y_{i2}^*|Y_{i1}, Y_{i2} = j, X_i) - \pi_{i0}}{1 - \pi_{i0} - \pi_{i1}} \\ &= \begin{cases} \frac{1 \times \pi_{i0} + 0 \times (1 - \pi_{i0}) - \pi_{i0}}{1 - \pi_{i0} - \pi_{i1}}, & \text{if } j = 0 \\ \frac{0 \times \pi_{i1} + 1 \times (1 - \pi_{i1}) - \pi_{i0}}{1 - \pi_{i0} - \pi_{i1}}, & \text{if } j = 1 \end{cases} \\ &= \begin{cases} 0, & \text{if } j = 0 \\ 1, & \text{if } j = 1 \end{cases} \\ &= j. \end{aligned}$$

Thus, (B.1) holds.

Step 2: Next, we show that

$$E(Y_{i1}^{**}|Y_{i1}, Y_{i2}, X_i) = Y_{i1}. \quad (\text{B.2})$$

By the definition of Y_{i1}^{**} , we obtain that

$$\begin{aligned}
E(Y_{i1}^{**}|Y_{i1}, Y_{i2}, X_i) &= E\left(\frac{Y_{i1}^* - \gamma_0 - \gamma_2 Y_{i2}^{**} - \gamma_3^T X_i}{\gamma_1} \middle| Y_{i1}, Y_{i2}, X_i\right) \\
&= E\left(\frac{Y_{i1}^* - \gamma_0 - \gamma_3^T X_i}{\gamma_1} \middle| Y_{i1}, Y_{i2}, X_i\right) - E\left(\frac{\gamma_2}{\gamma_1} Y_{i2}^{**} \middle| Y_{i1}, Y_{i2}, X_i\right) \\
&= \frac{1}{\gamma_1} E(Y_{i1}^* | Y_{i1}, Y_{i2}, X_i) - \frac{\gamma_0}{\gamma_1} - \frac{\gamma_2}{\gamma_1} Y_{i2} - \frac{\gamma_3^T X_i}{\gamma_1} \\
&= Y_{i1},
\end{aligned}$$

where the third step is due to (B.1), and the last step comes from measurement error model (3.9) together with $E(e_i | Y_{i1}, Y_{i2}, X_i) = 0$.

Step 3: Since e_i in (3.9) is independent of Y_{i1} , Y_{i2} and Y_{i2}^{**} , we have that

$$E(Y_{i1} e_i | Y_{i1}, Y_{i2}, X_i) = 0 \quad (\text{B.3})$$

and

$$E\{e_i(Y_{i2} - Y_{i2}^{**}) | Y_{i1}, Y_{i2}, X_i\} = 0. \quad (\text{B.4})$$

Now we prove that

$$E(\Delta_i | Y_{i1}, Y_{i2}, X_i) = E\{(Y_{i2} - Y_{i2}^{**})^2 | Y_{i1}, Y_{i2}, X_i\}. \quad (\text{B.5})$$

Indeed, by the definition of Δ_i ,

$$\begin{aligned}
E(\Delta_i | Y_{i1}, Y_{i1}, Y_{i2} = j, X_i) &= E\left(\frac{\Delta_{i0}^{1-Y_{i2}^*} \Delta_{i1}^{Y_{i2}^*} - \Delta_{i0} \pi_{i1} - \Delta_{i1} \pi_{i0}}{1 - \pi_{i1} - \pi_{i0}} \middle| Y_{i1}, Y_{i2} = j, X_i\right) \\
&= \begin{cases} \pi_{i0} \left(\frac{\Delta_{i1} - \Delta_{i0} \pi_{i1} - \Delta_{i1} \pi_{i0}}{1 - \pi_{i1} - \pi_{i0}}\right) + (1 - \pi_{i0}) \left(\frac{\Delta_{i0} - \Delta_{i0} \pi_{i1} - \Delta_{i1} \pi_{i0}}{1 - \pi_{i1} - \pi_{i0}}\right), & \text{if } j = 0 \\ (1 - \pi_{i1}) \left(\frac{\Delta_{i1} - \Delta_{i0} \pi_{i1} - \Delta_{i1} \pi_{i0}}{1 - \pi_{i1} - \pi_{i0}}\right) + \pi_{i1} \left(\frac{\Delta_{i0} - \Delta_{i0} \pi_{i1} - \Delta_{i1} \pi_{i0}}{1 - \pi_{i1} - \pi_{i0}}\right), & \text{if } j = 1 \end{cases} \\
&= \begin{cases} \Delta_{i0}, & \text{if } j = 0 \\ \Delta_{i1}, & \text{if } j = 1 \end{cases} \\
&= \Delta_{ij}.
\end{aligned}$$

On the other hand,

$$\begin{aligned}
E \left\{ (Y_{i2} - Y_{i2}^{**})^2 \mid Y_{i1}, Y_{i2} = j, X_i \right\} &= E \left\{ \left(Y_{i2} - \frac{Y_{i2}^* - \pi_{i0}}{1 - \pi_{i0} - \pi_{i1}} \right)^2 \mid Y_{i1}, Y_{i2} = j, X_i \right\} \\
&= \begin{cases} \pi_{i0} \left(-\frac{1 - \pi_{i0}}{1 - \pi_{i0} - \pi_{i1}} \right)^2 + (1 - \pi_{i0}) \left(\frac{\pi_{i0}}{1 - \pi_{i0} - \pi_{i1}} \right)^2, & \text{if } j = 0 \\ (1 - \pi_{i1}) \left(1 - \frac{1 - \pi_{i0}}{1 - \pi_{i0} - \pi_{i1}} \right)^2 + \pi_{i1} \left(1 + \frac{\pi_{i0}}{1 - \pi_{i0} - \pi_{i1}} \right)^2, & \text{if } j = 1 \end{cases} \\
&= \begin{cases} \Delta_{i0}, & \text{if } j = 0 \\ \Delta_{i1}, & \text{if } j = 1 \end{cases} \\
&= \Delta_{ij}.
\end{aligned}$$

Hence, (B.5) is proved.

Step 4: We show that

$$E(Y_{i1}^{**} \mid Y_{i1}, Y_{i2}, X_i) = Y_{i1}^2. \quad (\text{B.6})$$

By the definition of Y_{i1}^{**} ,

$$\begin{aligned}
E(Y_{i1}^{**} \mid Y_{i1}, Y_{i2}, X_i) &= E \left(Y_{i1}^{**2} - \frac{\sigma_e^2}{\gamma_1^2} - \frac{\gamma_2^2}{\gamma_1^2} \Delta_i \mid Y_{i1}, Y_{i2}, X_i \right) \\
&= E \left[\left\{ \frac{\gamma_1 Y_{i1} + e_i + \gamma_2 (Y_{i2} - Y_{i2}^{**})}{\gamma_1} \right\}^2 - \frac{\sigma_e^2}{\gamma_1^2} - \frac{\gamma_2^2}{\gamma_1^2} \Delta_i \mid Y_{i1}, Y_{i2}, X_i \right] \\
&= E \left\{ Y_{i1}^2 + \frac{\gamma_2^2}{\gamma_1^2} (Y_{i2} - Y_{i2}^{**})^2 - \frac{\gamma_2^2}{\gamma_1^2} \Delta_i \right. \\
&\quad \left. + \frac{2Y_{i1}e_i}{\gamma_1} + \frac{2\gamma_2 e_i (Y_{i2} - Y_{i2}^{**})}{\gamma_1^2} + \frac{2\gamma_2 Y_{i1} (Y_{i2} - Y_{i2}^{**})}{\gamma_1} \mid Y_{i1}, Y_{i2}, X_i \right\} \\
&= Y_{i1}^2 + \frac{\gamma_2^2}{\gamma_1^2} E \left\{ (Y_{i2} - Y_{i2}^{**})^2 - \Delta_i \mid Y_{i1}, Y_{i2}, X_i \right\} \\
&= Y_{i1}^2,
\end{aligned}$$

where the second step comes from the definition of Y_{i1}^{**} and the model (3.9), the fourth step comes from (B.3), (B.4) and $E \{ Y_{i1} (Y_{i2} - Y_{i2}^{**}) \mid Y_{i1}, Y_{i2}, X_i \} = 0$ which is due to (B.1), and the last step is due to (B.5).

Step 5: We show that

$$E(Y_{i12}^{**}|Y_{i1}, Y_{i2}, X_i) = Y_{i1}Y_{i2}. \quad (\text{B.7})$$

Similarly, by definition, we obtain that

$$\begin{aligned}
& E(Y_{i12}^{**}|Y_{i1}, Y_{i2}, X_i) \\
&= E\left(Y_{i1}^{**}Y_{i2}^{**} + \frac{\gamma_2}{\gamma_1}\Delta_i|Y_{i1}, Y_{i2}, X_i\right) \\
&= E\left(\frac{Y_{i1}^* - \gamma_0 - \gamma_2 Y_{i2}^{**} - \gamma_3^T X_i}{\gamma_1} Y_{i2}^{**} + \frac{\gamma_2}{\gamma_1}\Delta_i|Y_{i1}, Y_{i2}, X_i\right) \\
&= E\left\{\frac{Y_{i1}^* - \gamma_0 - \gamma_2 Y_{i2} - \gamma_3^T X_i + \gamma_2(Y_{i2} - Y_{i2}^{**})}{\gamma_1} Y_{i2}^{**} + \frac{\gamma_2}{\gamma_1}\Delta_i|Y_{i1}, Y_{i2}, X_i\right\} \\
&= E\left(\frac{Y_{i1}^* - \gamma_0 - \gamma_2 Y_{i2} - \gamma_3^T X_i}{\gamma_1} Y_{i2}^{**}|Y_{i1}, Y_{i2}, X_i\right) + \frac{\gamma_2}{\gamma_1} E\left\{(Y_{i2} - Y_{i2}^{**})Y_{i2}^{**} + \Delta_i|Y_{i1}, Y_{i2}, X_i\right\} \\
&= E\left(\frac{Y_{i1}^* - \gamma_0 - \gamma_2 Y_{i2} - \gamma_3^T X_i}{\gamma_1}|Y_{i1}, Y_{i2}, X_i\right) E\left(Y_{i2}^{**}|Y_{i1}, Y_{i2}, X_i\right) \\
&\quad + \frac{\gamma_2}{\gamma_1} E\left\{(Y_{i2} - Y_{i2}^{**})Y_{i2}^{**}|Y_{i1}, Y_{i2}, X_i\right\} + \frac{\gamma_2}{\gamma_1} E\left(\Delta_i|Y_{i1}, Y_{i2}, X_i\right) \\
&= E\left(Y_{i1} + \frac{e_i}{\gamma_1}|Y_{i1}, Y_{i2}, X_i\right) Y_{i2} + \frac{\gamma_2}{\gamma_1} E\left\{(Y_{i2} - Y_{i2}^{**})Y_{i2}^{**}|Y_{i1}, Y_{i2}, X_i\right\} + \frac{\gamma_2}{\gamma_1} E\left(\Delta_i|Y_{i1}, Y_{i2}, X_i\right) \\
&= Y_{i1}Y_{i2} + \frac{\gamma_2}{\gamma_1} E\left\{(Y_{i2} - Y_{i2}^{**})Y_{i2}^{**}|Y_{i1}, Y_{i2}, X_i\right\} + \frac{\gamma_2}{\gamma_1} E\left(\Delta_i|Y_{i1}, Y_{i2}, X_i\right), \quad (\text{B.8})
\end{aligned}$$

where the fifth step is due to the conditional independence assumption for Y_{i1}^* and Y_{i2}^* given by (2.4), and the sixth step comes from (3.9) and (B.1), and last step comes from $E(e_i|Y_{i1}, Y_{i2}, X_i) = 0$.

By (B.5), we obtain that

$$\begin{aligned}
& E(\Delta_i|Y_{i1}, Y_{i2}, X_i) = E\left\{(Y_{i2} - Y_{i2}^{**})^2|Y_{i1}, Y_{i2}, X_i\right\} \\
&= E\left\{(Y_{i2} - Y_{i2}^{**})Y_{i2}|Y_{i1}, Y_{i2}, X_i\right\} - E\left\{(Y_{i2} - Y_{i2}^{**})Y_{i2}^{**}|Y_{i1}, Y_{i2}, X_i\right\} \\
&= -E\left\{(Y_{i2} - Y_{i2}^{**})Y_{i2}^{**}|Y_{i1}, Y_{i2}, X_i\right\},
\end{aligned}$$

where the last step is due to (B.1). Consequently, (B.8) gives (B.7).

Step 6: In (3.10) we replace Y_{i1} , Y_{i1}^2 and $Y_{i1}Y_{i2}$, respectively, with Y_{i1}^{**} , Y_{i11}^{**} , and Y_{i12}^{**} , and

then we obtain $U_i^{**}(\theta) = (U_{i1}^{**\text{T}}(\theta), U_{i2}^{**\text{T}}(\theta))^{\text{T}}$ where

$$\begin{aligned} U_{i1}^{**}(\theta) &= D_{1i}^{\text{T}} V_{i1}^{-1} \begin{pmatrix} Y_{i1}^{**} - \mu_{i1} \\ Y_{i2}^{**} - \mu_{i2} \end{pmatrix}, \\ U_{i2}^{**}(\theta) &= D_{2i}^{\text{T}} V_{i2}^{-1} \begin{pmatrix} Y_{i11}^{**} - 2\mu_{i1} Y_{i1}^{**} + \mu_{i1}^2 - \xi_{i1} \\ Y_{i12}^{**} - Y_{i1}^{**} \mu_{i2} - Y_{i2}^{**} \mu_{i1} + \mu_{i1} \mu_{i2} - \xi_{i2} \\ Y_{i2}^{**} - 2\mu_{i2} Y_{i2}^{**} + \mu_{i2}^2 - \xi_{i3} \end{pmatrix}. \end{aligned}$$

Then applying (B.1), (B.2), (B.6), and (B.7) gives that

$$E\{U_i^{**}(\theta) | Y_{i1}, Y_{i2}, X_i\} = U_i(\theta).$$

B.2 The Consistency and Normality of the Proposed Estimator with External Validation Data

Assume that subjects are randomly assigned to the validation or nonvalidation sample. Let $\delta_i = I(i \in \mathcal{V})$, where $I(\cdot)$ is the indicator function. Define $H_{i1}(\theta, \eta) = (1 - \delta_i)U_{i1}^{**}(\theta, \eta)$, $H_{i2}(\theta, \eta) = (1 - \delta_i)U_{i2}^{**}(\theta, \eta)$, $H_{i3}(\eta) = \delta_i S_i(\eta)$, and $H_i(\theta, \eta) = \{H_{i1}^{\text{T}}(\theta, \eta), H_{i2}^{\text{T}}(\theta, \eta), H_{i3}^{\text{T}}(\eta)\}^{\text{T}}$. Then, (3.13) is equivalent to

$$U^{(E)}(\theta, \eta) = \sum_{i \in \mathcal{M} \cup \mathcal{V}} H_i(\theta, \eta).$$

Since $H_i(\theta, \eta)$ is an unbiased estimating function, i.e., $E\{H_i(\theta, \eta)\} = 0$, then by estimating function theory (Godambe, 1991; Newey and McFadden, 1994; Heyde, 1997, Ch.12) we conclude that under regularity conditions, solving $\sum_{i \in \mathcal{M} \cup \mathcal{V}} H_i(\theta, \eta) = 0$ gives a consistent estimator, $(\hat{\theta}_E^{\text{T}}, \hat{\eta}_E^{\text{T}})^{\text{T}}$, of $(\theta^{\text{T}}, \eta^{\text{T}})^{\text{T}}$.

Applying the Taylor series expansion to $\sum_{i \in \mathcal{M} \cup \mathcal{V}} H_i(\hat{\theta}_E, \hat{\eta}_E) = 0$, we obtain

$$\sum_{i \in \mathcal{M} \cup \mathcal{V}} H_i(\theta, \eta) + \sum_{i \in \mathcal{M} \cup \mathcal{V}} \left(\frac{\partial H_i(\theta, \eta)}{\partial \theta^{\text{T}}} \quad \frac{\partial H_i(\theta, \eta)}{\partial \eta^{\text{T}}} \right) \left\{ \begin{pmatrix} \hat{\theta}_E \\ \hat{\eta}_E \end{pmatrix} - \begin{pmatrix} \theta \\ \eta \end{pmatrix} \right\} + o_p(1) = 0,$$

which leads to

$$\begin{aligned} & \sqrt{1 + \frac{m}{n}} \left\{ -\frac{1}{n+m} \sum_{i \in \mathcal{M} \cup \mathcal{V}} \left(\frac{\partial H_i(\theta, \eta)}{\partial \theta^{\text{T}}} \quad \frac{\partial H_i(\theta, \eta)}{\partial \eta^{\text{T}}} \right) \right\} \sqrt{n} \left\{ \begin{pmatrix} \hat{\theta}_E \\ \hat{\eta}_E \end{pmatrix} - \begin{pmatrix} \theta \\ \eta \end{pmatrix} \right\} \\ &= \frac{1}{\sqrt{n+m}} \sum_{i \in \mathcal{M} \cup \mathcal{V}} H_i(\theta, \eta) + o_p(1). \end{aligned} \tag{B.9}$$

Let

$$\Gamma_E = \lim_{n \rightarrow \infty} \left\{ -\frac{1}{n+m} \sum_{i \in \mathcal{M} \cup \mathcal{V}} E \left(\begin{array}{cc} \frac{\partial H_i(\theta, \eta)}{\partial \theta^T} & \frac{\partial H_i(\theta, \eta)}{\partial \eta^T} \end{array} \right) \right\}$$

and

$$\Sigma_E = \lim_{n \rightarrow \infty} \text{Var} \left(\frac{1}{\sqrt{n+m}} \sum_{i \in \mathcal{M} \cup \mathcal{V}} H_i(\theta, \eta) \right).$$

Then applying the central limit theorem to (B.9) gives that

$$\sqrt{n} \left\{ (\hat{\theta}_E^T, \hat{\eta}_E^T)^T - (\theta^T, \eta^T)^T \right\} \xrightarrow{d} N \left(0, \frac{1}{1+\rho} \Gamma_E^{-1} \Sigma_E (\Gamma_E^{-1})^T \right) \quad \text{as } n \rightarrow \infty.$$

Now it remains to show that Γ_E and Σ_E are identical to (3.14). By the definitions of $H_i(\theta, \eta)$, we derive that

$$\begin{aligned} \Gamma_E &= \lim_{n \rightarrow \infty} \left\{ -\frac{1}{n+m} E \left(\begin{array}{cc} \sum_{i \in \mathcal{M}} \frac{\partial U_{i1}^{**}(\theta, \eta)}{\partial \theta^T} & \sum_{i \in \mathcal{M}} \frac{\partial U_{i1}^{**}(\theta, \eta)}{\partial \eta^T} \\ \sum_{i \in \mathcal{M}} \frac{\partial U_{i2}^{**}(\theta, \eta)}{\partial \theta^T} & \sum_{i \in \mathcal{M}} \frac{\partial U_{i2}^{**}(\theta, \eta)}{\partial \eta^T} \\ 0 & 0 \end{array} \right) - \frac{1}{n+m} E \left(\begin{array}{cc} 0 & 0 \\ 0 & 0 \\ 0 & \sum_{i \in \mathcal{V}} \frac{\partial S_i(\eta)}{\partial \eta^T} \end{array} \right) \right\} \\ &= \lim_{n \rightarrow \infty} \left\{ -\frac{n}{n+m} E \left(\begin{array}{cc} \frac{\partial U_{i1}^{**}(\theta, \eta)}{\partial \theta^T} & \frac{\partial U_{i1}^{**}(\theta, \eta)}{\partial \eta^T} \\ \frac{\partial U_{i2}^{**}(\theta, \eta)}{\partial \theta^T} & \frac{\partial U_{i2}^{**}(\theta, \eta)}{\partial \eta^T} \\ 0 & 0 \end{array} \right) - \frac{m}{n+m} E \left(\begin{array}{cc} 0 & 0 \\ 0 & 0 \\ 0 & \frac{\partial S_i}{\partial \eta^T} \end{array} \right) \right\} \\ &= -\frac{1}{1+\rho} \left(\begin{array}{cc} E \left(\frac{\partial U_{i1}^{**}(\theta, \eta)}{\partial \theta^T} \right) & E \left(\frac{\partial U_{i1}^{**}(\theta, \eta)}{\partial \eta^T} \right) \\ E \left(\frac{\partial U_{i2}^{**}(\theta, \eta)}{\partial \theta^T} \right) & E \left(\frac{\partial U_{i2}^{**}(\theta, \eta)}{\partial \eta^T} \right) \\ 0 & 0 \end{array} \right) - \frac{\rho}{1+\rho} \left(\begin{array}{cc} 0 & 0 \\ 0 & 0 \\ 0 & E \left(\frac{\partial S_i(\eta)}{\partial \eta^T} \right) \end{array} \right). \end{aligned}$$

Similarly, we have that

$$\begin{aligned}
\Sigma_E &= \lim_{n \rightarrow \infty} \frac{1}{n+m} \sum_{i \in \mathcal{M} \cup \mathcal{V}} \text{Var}\{H_i(\theta, \eta)\} \\
&= \lim_{n \rightarrow \infty} \frac{1}{n+m} \sum_{i \in \mathcal{M} \cup \mathcal{V}} E\{H_i(\theta, \eta)H_i^T(\theta, \eta)\} \\
&= \lim_{n \rightarrow \infty} \frac{1}{n+m} \sum_{i \in \mathcal{M} \cup \mathcal{V}} E \begin{pmatrix} (1-\delta_i)^2 U_{i1}^{**}(\theta, \eta) U_{i1}^{**T}(\theta, \eta) & (1-\delta_i)^2 U_{i1}^{**}(\theta, \eta) U_{i2}^{**T}(\theta, \eta) & 0 \\ (1-\delta_i)^2 U_{i2}^{**}(\theta, \eta) U_{i1}^{**T}(\theta, \eta) & (1-\delta_i)^2 U_{i2}^{**}(\theta, \eta) U_{i2}^{**T}(\theta, \eta) & 0 \\ 0 & 0 & \delta_i^2 S_i(\eta) S_i^T(\eta) \end{pmatrix} \\
&= \lim_{n \rightarrow \infty} \frac{n}{n+m} \frac{1}{n} \sum_{i \in \mathcal{M}} \begin{pmatrix} E\{U_{i1}^{**}(\theta, \eta) U_{i1}^{**T}(\theta, \eta)\} & E\{U_{i1}^{**}(\theta, \eta) U_{i2}^{**T}(\theta, \eta)\} & 0 \\ E\{U_{i2}^{**}(\theta, \eta) U_{i1}^{**T}(\theta, \eta)\} & E\{U_{i2}^{**}(\theta, \eta) U_{i2}^{**T}(\theta, \eta)\} & 0 \\ 0 & 0 & 0 \end{pmatrix} \\
&\quad + \lim_{n \rightarrow \infty} \frac{m}{n+m} \frac{1}{m} \sum_{i \in \mathcal{V}} \begin{pmatrix} 0 & 0 & 0 \\ 0 & 0 & 0 \\ 0 & 0 & E\{S_i(\eta) S_i^T(\eta)\} \end{pmatrix} \\
&= \frac{1}{1+\rho} \begin{pmatrix} E\{U_{i1}^{**}(\theta, \eta) U_{i1}^{**T}(\theta, \eta)\} & E\{U_{i1}^{**}(\theta, \eta) U_{i2}^{**T}(\theta, \eta)\} & 0 \\ E\{U_{i2}^{**}(\theta, \eta) U_{i1}^{**T}(\theta, \eta)\} & E\{U_{i2}^{**}(\theta, \eta) U_{i2}^{**T}(\theta, \eta)\} & 0 \\ 0 & 0 & 0 \end{pmatrix} + \frac{\rho}{1+\rho} \begin{pmatrix} 0 & 0 & 0 \\ 0 & 0 & 0 \\ 0 & 0 & E\{S_i(\eta) S_i^T(\eta)\} \end{pmatrix},
\end{aligned}$$

where the third step is due to $(1-\delta_i)^2 = (1-\delta_i)$ and $\delta_i^2 = \delta_i$.

B.3 The Consistency and Normality of the Proposed Estimator with Internal Validation Data

Similar to Section B.2, we assume that subjects are randomly assigned to the validation or nonvalidation sample. We define $H_{i1}(\theta, \eta) = (1-\delta_i)U_{i1}^{**}(\theta, \eta) + \delta_i U_{i1}(\theta, \eta)$, $H_{i2}(\theta, \eta) = (1-\delta_i)U_{i2}^{**}(\theta, \eta) + \delta_i U_{i2}(\theta, \eta)$, $H_{i3}(\theta, \eta) = \delta_i S_i(\eta)$, and $H_i(\theta, \eta) = \{H_{i1}^T(\theta, \eta), H_{i2}^T(\theta, \eta), H_{i3}^T(\eta)\}^T$. Then, (3.15) is equivalent to

$$U^{(I)}(\theta, \eta) = \sum_{i \in \mathcal{M}} H_i(\theta, \eta).$$

Similar to Section B.2, we conclude that under regularity conditions, solving $\sum_{i \in \mathcal{M}} H_i(\theta, \eta) = 0$ gives a consistent estimator, $(\hat{\theta}_I^T, \hat{\eta}_I^T)^T$, of $(\theta^T, \eta^T)^T$.

Applying the Taylor series expansion to $\sum_{i \in \mathcal{M}} H_i(\hat{\theta}_I^T, \hat{\eta}_I^T) = 0$, we obtain

$$\sum_{i \in \mathcal{M}} H_i(\theta, \eta) + \sum_{i \in \mathcal{M}} \begin{pmatrix} \frac{\partial H_i(\theta, \eta)}{\partial \theta^T} & \frac{\partial H_i(\theta, \eta)}{\partial \eta^T} \end{pmatrix} \left\{ \begin{pmatrix} \hat{\theta}_I \\ \hat{\eta}_I \end{pmatrix} - \begin{pmatrix} \theta \\ \eta \end{pmatrix} \right\} + o_p(1) = 0,$$

which leads to

$$\left\{ -\frac{1}{n} \sum_{i \in \mathcal{M}} \begin{pmatrix} \frac{\partial H_i(\theta, \eta)}{\partial \theta^T} & \frac{\partial H_i(\theta, \eta)}{\partial \eta^T} \end{pmatrix} \right\} \sqrt{n} \left\{ \begin{pmatrix} \hat{\theta}_I \\ \hat{\eta}_I \end{pmatrix} - \begin{pmatrix} \theta \\ \eta \end{pmatrix} \right\} = \frac{1}{\sqrt{n}} \sum_{i \in \mathcal{M}} H_i(\theta, \eta) + o_p(1). \quad (\text{B.10})$$

Let

$$\Gamma_I = \lim_{n \rightarrow \infty} \left\{ -\frac{1}{n} \sum_{i \in \mathcal{M}} E \begin{pmatrix} \frac{\partial H_i(\theta, \eta)}{\partial \theta^T} & \frac{\partial H_i(\theta, \eta)}{\partial \eta^T} \end{pmatrix} \right\}$$

and

$$\Sigma_I = \lim_{n \rightarrow \infty} \text{Var} \left(\frac{1}{\sqrt{n}} \sum_{i \in \mathcal{M}} H_i(\theta, \eta) \right).$$

Then applying the central limit theorem to (B.10) gives that

$$\sqrt{n} \left\{ (\hat{\theta}_I^T, \hat{\eta}_I^T)^T - (\theta^T, \eta^T)^T \right\} \xrightarrow{d} N(0, \Gamma_I^{-1} \Sigma_I (\Gamma_I^{-1})^T) \quad \text{as} \quad n \rightarrow \infty.$$

Now it remains to show that Γ_I and Σ_I are identical to (3.17). By definition of $H_i(\theta, \eta)$, Γ_I equals

$$\begin{aligned} & \lim_{n \rightarrow \infty} \left\{ -\frac{1}{n} E \begin{pmatrix} \sum_{i \in \mathcal{M} \setminus \mathcal{V}} \frac{\partial U_{i1}^{**}(\theta, \eta)}{\partial \theta^T} & \sum_{i \in \mathcal{M} \setminus \mathcal{V}} \frac{\partial U_{i1}^{**}(\theta, \eta)}{\partial \eta^T} \\ \sum_{i \in \mathcal{M} \setminus \mathcal{V}} \frac{\partial U_{i2}^{**}(\theta, \eta)}{\partial \theta^T} & \sum_{i \in \mathcal{M} \setminus \mathcal{V}} \frac{\partial U_{i2}^{**}(\theta, \eta)}{\partial \eta^T} \\ 0 & 0 \end{pmatrix} - \frac{1}{n} E \begin{pmatrix} \sum_{i \in \mathcal{V}} \frac{\partial U_{i1}(\theta, \eta)}{\partial \theta^T} & 0 \\ \sum_{i \in \mathcal{V}} \frac{\partial U_{i2}(\theta, \eta)}{\partial \theta^T} & 0 \\ 0 & \sum_{i \in \mathcal{V}} \frac{\partial S_i(\eta)}{\partial \eta^T} \end{pmatrix} \right\} \\ &= \lim_{n \rightarrow \infty} \left\{ -\frac{n-m}{n} E \begin{pmatrix} \frac{\partial U_{i1}^{**}(\theta, \eta)}{\partial \theta^T} & \frac{\partial U_{i1}^{**}(\theta, \eta)}{\partial \eta^T} \\ \frac{\partial U_{i2}^{**}(\theta, \eta)}{\partial \theta^T} & \frac{\partial U_{i2}^{**}(\theta, \eta)}{\partial \eta^T} \\ 0 & 0 \end{pmatrix} - \frac{m}{n} E \begin{pmatrix} \frac{\partial U_{i1}(\theta, \eta)}{\partial \theta^T} & 0 \\ \frac{\partial U_{i2}(\theta, \eta)}{\partial \theta^T} & 0 \\ 0 & \frac{\partial S_i(\eta)}{\partial \eta^T} \end{pmatrix} \right\} \\ &= -(1-\rho) \begin{pmatrix} E \left(\frac{\partial U_{i1}^{**}(\theta, \eta)}{\partial \theta^T} \right) & E \left(\frac{\partial U_{i1}^{**}(\theta, \eta)}{\partial \eta^T} \right) \\ E \left(\frac{\partial U_{i2}^{**}(\theta, \eta)}{\partial \theta^T} \right) & E \left(\frac{\partial U_{i2}^{**}(\theta, \eta)}{\partial \theta^T} \right) \\ 0 & 0 \end{pmatrix} - \rho \begin{pmatrix} E \left(\frac{\partial U_{i1}(\theta, \eta)}{\partial \theta^T} \right) & 0 \\ E \left(\frac{\partial U_{i2}(\theta, \eta)}{\partial \theta^T} \right) & 0 \\ 0 & E \left(\frac{\partial S_i(\eta)}{\partial \eta^T} \right) \end{pmatrix}. \end{aligned}$$

Similarly, we have that

$$\begin{aligned}
\Sigma_I &= \lim_{n \rightarrow \infty} \frac{1}{n} \sum_{i \in \mathcal{M}} \text{Var} \{H_i(\theta, \eta)\} \\
&= \lim_{n \rightarrow \infty} \frac{1}{n} \sum_{i \in \mathcal{M}} E\{H_i(\theta, \eta)H_i^\top(\theta, \eta)\} \\
&= \lim_{n \rightarrow \infty} \frac{1}{n} \sum_{i \in \mathcal{M}} E \begin{pmatrix} (1 - \delta_i)^2 U_{i1}^{**}(\theta, \eta) U_{i1}^{**\top}(\theta, \eta) & (1 - \delta_i)^2 U_{i1}^{**}(\theta, \eta) U_{i2}^{**\top}(\theta, \eta) & 0 \\ (1 - \delta_i)^2 U_{i2}^{**}(\theta, \eta) U_{i1}^{**\top}(\theta, \eta) & (1 - \delta_i)^2 U_{i2}^{**}(\theta, \eta) U_{i2}^{**\top}(\theta, \eta) & 0 \\ 0 & 0 & 0 \end{pmatrix} \\
&\quad + \lim_{n \rightarrow \infty} \frac{1}{n} \sum_{i \in \mathcal{M}} E \begin{pmatrix} \delta_i^2 U_{i1}(\theta, \eta) U_{i1}^\top(\theta, \eta) & \delta_i^2 U_{i1}(\theta, \eta) U_{i2}^\top(\theta, \eta) & \delta_i^2 U_{i1}(\theta, \eta) S_i^\top(\eta) \\ \delta_i^2 U_{i2}(\theta, \eta) U_{i1}^\top(\theta, \eta) & \delta_i^2 U_{i1}(\theta, \eta) U_{i1}^\top(\theta, \eta) & \delta_i^2 U_{i1}(\theta, \eta) S_i^\top(\eta) \\ \delta_i^2 U_{i1}(\theta, \eta) S_i^\top(\eta) & \delta_i^2 U_{i2}(\theta, \eta) S_i^\top(\eta) & \delta_i^2 S_i(\eta) S_i^\top(\eta) \end{pmatrix} \\
&= \lim_{n \rightarrow \infty} \frac{n - m}{n} \frac{1}{n - m} \sum_{i \in \mathcal{M} \setminus \mathcal{V}} \begin{pmatrix} E\{U_{i1}^{**}(\theta, \eta) U_{i1}^{**\top}(\theta, \eta)\} & E\{U_{i1}^{**}(\theta, \eta) U_{i2}^{**\top}(\theta, \eta)\} & 0 \\ E\{U_{i2}^{**}(\theta, \eta) U_{i1}^{**\top}(\theta, \eta)\} & E\{U_{i2}^{**}(\theta, \eta) U_{i2}^{**\top}(\theta, \eta)\} & 0 \\ 0 & 0 & 0 \end{pmatrix} \\
&\quad + \lim_{n \rightarrow \infty} \frac{m}{n} \frac{1}{m} \sum_{i \in \mathcal{V}} \begin{pmatrix} E\{U_{i1}(\theta, \eta) U_{i1}^\top(\theta, \eta)\} & E\{U_{i1}(\theta, \eta) U_{i2}^\top(\theta, \eta)\} & E\{U_{i1}(\theta, \eta) S_i^\top(\theta, \eta)\} \\ E\{U_{i2}(\theta, \eta) U_{i1}^\top(\theta, \eta)\} & E\{U_{i1}(\theta, \eta) U_{i1}^\top(\theta, \eta)\} & E\{U_{i1}(\theta, \eta) S_i^\top(\theta, \eta)\} \\ E\{U_{i1}(\theta, \eta) S_i^\top(\theta, \eta)\} & E\{U_{i2}(\theta, \eta) S_i^\top(\theta, \eta)\} & E\{S_i(\theta, \eta) S_i^\top(\theta, \eta)\} \end{pmatrix} \\
&= (1 - \rho) \begin{pmatrix} E\{U_{i1}^{**}(\theta, \eta) U_{i1}^{**\top}(\theta, \eta)\} & E\{U_{i1}^{**}(\theta, \eta) U_{i2}^{**\top}(\theta, \eta)\} & 0 \\ E\{U_{i2}^{**}(\theta, \eta) U_{i1}^{**\top}(\theta, \eta)\} & E\{U_{i2}^{**}(\theta, \eta) U_{i2}^{**\top}(\theta, \eta)\} & 0 \\ 0 & 0 & 0 \end{pmatrix} \\
&\quad + \rho \begin{pmatrix} E\{U_{i1}(\theta, \eta) U_{i1}^\top(\theta, \eta)\} & E\{U_{i1}(\theta, \eta) U_{i2}^\top(\theta, \eta)\} & E\{U_{i1}(\theta, \eta) S_i^\top(\eta)\} \\ E\{U_{i2}(\theta, \eta) U_{i1}^\top(\theta, \eta)\} & E\{U_{i1}(\theta, \eta) U_{i1}^\top(\theta, \eta)\} & E\{U_{i1}(\theta, \eta) S_i^\top(\eta)\} \\ E\{U_{i1}(\theta, \eta) S_i^\top(\eta)\} & E\{U_{i2}(\theta, \eta) S_i^\top(\eta)\} & E\{S_i(\eta) S_i^\top(\eta)\} \end{pmatrix},
\end{aligned}$$

where the third step is due to $\delta_i(1 - \delta_i) = 0$ and the fourth step comes from $(1 - \delta_i)^2 = (1 - \delta_i)$ and $\delta_i^2 = \delta_i$.

B.4 Determination the values of $\widehat{\text{Var}}(\widehat{\theta}_{I_r}^{(1)})$, $\widehat{\text{Var}}(\widehat{\theta}_{I_r}^{(0)})$ and $\widehat{\text{Cov}}(\widehat{\theta}_{I_r}^{(0)}, \widehat{\theta}_{I_r}^{(1)})$

To study the covariance between $\widehat{\theta}_{I_r}^{(0)}$ and $\widehat{\theta}_{I_r}^{(1)}$, we jointly combine the estimating procedure by stacking the estimating functions from validation data and nonvalidation data. However, this procedure makes the resulting dimension of estimating functions be greater than the dimension of parameter θ . To overcome this problem, following the spirit of (Shu and Yi, 2017), we enlarge the original parameter space by using different symbols, say, $\theta^{(0)}$ and $\theta^{(1)}$, respectively, to represent the parameter θ in the estimating function of $\widehat{\theta}_{I_r}^{(0)}$ and $\widehat{\theta}_{I_r}^{(1)}$, where the true value of $\theta^{(0)}$ and $\theta^{(1)}$ are identical to that of θ . Specifically, consider the estimating functions

$$\Psi_i(\theta^{(0)\text{T}}, \eta^{\text{T}}, \theta^{(1)\text{T}}) = \begin{pmatrix} I(i \in \mathcal{V}) \cdot U_{i1}(\theta^{(0)}, \eta) \\ I(i \in \mathcal{V}) \cdot U_{i2}(\theta^{(0)}, \eta) \\ I(i \in \mathcal{V}) \cdot S_i(\eta) \\ I(i \in \mathcal{M} \setminus \mathcal{V}) \cdot U_{i1}^{**}(\theta^{(1)}, \eta) \\ I(i \in \mathcal{M} \setminus \mathcal{V}) \cdot U_{i2}^{**}(\theta^{(1)}, \eta) \end{pmatrix}. \quad (\text{B.11})$$

Solving the estimating function $\sum_{i=1}^n \Psi_i(\theta^{(0)\text{T}}, \eta^{\text{T}}, \theta^{(1)\text{T}}) = 0$, we obtain an estimator, $(\widehat{\theta}^{(0)\text{T}}, \widehat{\eta}^{\text{T}}, \widehat{\theta}^{(1)\text{T}})^{\text{T}}$, of $(\theta^{(0)\text{T}}, \eta^{\text{T}}, \theta^{(1)\text{T}})^{\text{T}}$. By estimating function theory (e.g. Godambe, 1991; Newey and McFadden, 1994; Yi, 2017, Section 1.3.2), the variance of $(\widehat{\theta}^{(0)\text{T}}, \widehat{\eta}^{\text{T}}, \widehat{\theta}^{(1)\text{T}})^{\text{T}}$ can be estimated by the empirical sandwich estimator

$$\widehat{\text{Var}} \left\{ (\widehat{\theta}^{(0)\text{T}}, \widehat{\eta}^{\text{T}}, \widehat{\theta}^{(1)\text{T}})^{\text{T}} \right\} = \frac{1}{n} \Gamma_{\Psi}^{-1} \Sigma_{\Psi} \Gamma_{\Psi}^{-1\text{T}}, \quad (\text{B.12})$$

where $\Gamma_{\Psi} = \frac{1}{n} \sum_{i=1}^n \left\{ -\frac{\partial}{\partial(\theta^{(0)\text{T}}, \eta^{\text{T}}, \theta^{(1)\text{T}})^{\text{T}}} \Psi_i(\widehat{\theta}^{(0)}, \widehat{\eta}, \widehat{\theta}^{(1)}) \right\}$ and $\Sigma_{\Psi} = \frac{1}{n} \sum_{i=1}^n \left\{ \Psi_i(\widehat{\theta}^{(0)}, \widehat{\eta}, \widehat{\theta}^{(1)}) \Psi_i^{\text{T}}(\widehat{\theta}^{(0)}, \widehat{\eta}, \widehat{\theta}^{(1)}) \right\}$.

Therefore, $\widehat{\text{Var}}(\widehat{\theta}_{I_r}^{(0)})$, $\widehat{\text{Var}}(\widehat{\theta}_{I_r}^{(1)})$ and $\widehat{\text{Cov}}(\widehat{\theta}_{I_r}^{(0)}, \widehat{\theta}_{I_r}^{(1)})$ are, respectively, the covariance matrix, $\widehat{\text{Var}} \left\{ (\widehat{\theta}^{(0)\text{T}}, \widehat{\eta}^{\text{T}}, \widehat{\theta}^{(1)\text{T}})^{\text{T}} \right\}$, corresponding to elements (r, r) , $(r+q, r+q)$ and $(r, r+q)$ where $q = p_{\theta} + p_{\eta}$.

Appendix C

Conditions and Proofs of the Results in Chapter 4

C.1 Regularity Conditions

(R1) The dimension p of the covariates is of a polynomial order of the sample size n . That is, $p = O(n^\gamma)$ for a constant $\gamma > 0$.

(R2) There exists $0 \leq \kappa < 1$ so that

$$\max_{s \in V} |\mathcal{N}(s)| = O(n^\kappa).$$

(R3) There exists some $m < \infty$ so that

$$\max_{s \in V, t \in \mathcal{N}(s)} |\mathcal{N}(s) \cap \mathcal{N}(t)| \leq m.$$

(R4) There exists a constant $\delta > 0$ and ξ with $\kappa < \xi \leq 1$ for κ in condition (R2), such that for every edge $(s, t) \in E$,

$$|\pi_{st}| \geq \delta n^{-\frac{1-\xi}{2}},$$

where π_{st} is the partial correlation between X_{is} and X_{it} after eliminating the linear effects from all remaining variables $\{X_{ik} : k \in V \setminus \{s, t\}\}$.

(R5) The covariance matrix of X_i is non-singular.

- (R6) The parameter space \mathcal{B} of $(\beta_M^T, \beta_I^T, \phi^T)^T$ is compact.
- (R7) Given the data, the estimating functions $\sum_{i=1}^n U_i(\beta_M, \beta_I, \phi)$ are continuous in $(\beta_M^T, \beta_I^T, \phi^T)^T$ everywhere, and satisfy the condition (12.5) in Theorem 12.1 of [Heyde \(1997\)](#).
- (R8) Given the data, the estimating functions $\sum_{i=1}^n U_i(\beta_M, \beta_I, \phi)$ are continuously differentiable in a neighbourhood of $(\beta_M^T, \beta_I^T, \phi^T)^T$.
- (R9) For $U_i(w; \beta_M, \beta_I, \phi)$ defined in (4.8) with $w = (y, x^T)^T$, there exists a function $h(w)$ with $E\{h(W_i)\} < \infty$, such that $|U_i(w, \beta_M, \beta_I, \phi)| < h(w)$ for all β_M, β_I , and ϕ , where $W_i = (Y_i, X_i^T)^T$.
- (R10) The equation $E_{\text{vix}}\{U_i(\beta_{M(0)}, \beta_{I(0)}, \phi_0)\} = 0$ has an unique solution.

Conditions (R1)–(R5) are the regularity conditions discussed by [Meinshausen and Bühlmann \(2006\)](#). Condition (R1) allows for a high dimension of the covariates and regulates the dimension of covariates on a scale relative to the sample size. Conditions (R2) and (R3) basically regulate the sparsity of the graph and the maximum possible growth rate of the size of neighborhoods. Condition (R4) provides a lower bound of the magnitude of partial correlations to ensure the consistency of variable selection of edge set E . Condition (R5) requires the existence of the precision matrix. Conditions (R6)–(R8) are the regularity conditions for estimating functions discussed by [Heyde \(1997\)](#). Condition (R9) is the condition for Theorem 2 in [Jennrich \(1969\)](#). Condition (R10) is used to show the consistency in Theorem 2, which was also assumed by [Yi and Reid \(2010\)](#), among others.

C.2 Proof of Theorem 4.1

For κ and ξ defined in Conditions (R3) and (R4), consider a tuning parameter λ in (4.6) satisfying $\lambda \sim dn^{-(1-\epsilon)/2}$ with $\kappa < \epsilon < \xi$ and a scaling constant $d > 0$. Then according to [Meinshausen and Bühlmann \(2006, Page 1445\)](#), with regularity conditions assumed, there exists $c > 0$ such that

$$P(\widehat{E} = E) = 1 - O(\exp(-cn^\epsilon)), \tag{C.1}$$

yielding, by the definition of $\widehat{\beta}_{\text{II}}$, that

$$P(\widehat{\beta}_{\text{II}} = \beta_{\text{II}}) \geq 1 - O(\exp(-cn^\epsilon)).$$

Thus, the conclusion follows from that $0 < \epsilon < \xi \leq 1$.

C.3 Proof of Theorem 4.2

Proof of Theorem 4.2 (i)

Step 1: We introduce basic notation first

Let $(\beta_M, \beta_I, \beta_{II}, \phi)$ be the generic symbol of parameters from parameter space \mathcal{B} , and let $(\beta_0^T, \phi_0^T)^T = (\beta_{M(0)}^T, \beta_{I(0)}^T, \beta_{II(0)}^T, \phi_0^T)^T$ denote the true value of the parameters $\beta = (\beta_M^T, \beta_I^T, \beta_{II}^T, \phi^T)^T$. Let $\mathcal{E} = \{E^a : E^a \subseteq \tilde{E}\}$ denote the collection of all possible subsets of \tilde{E} . For an estimated edge set \hat{E} for E , let $\beta^{*(\hat{E})} = \{\beta_{st} : (s, t) \in \hat{E}\}$ denote the subvector of $(\beta_I^T, \beta_{II}^T)^T$ with the indexes included in \hat{E} , and let $\beta^{**(\hat{E})} = \{\beta_{st} : (s, t) \notin \hat{E}\}$ be the complement of $\beta^{*(\hat{E})}$, i.e., the subvector of $(\beta_I^T, \beta_{II}^T)^T$ with the indexes not included in \hat{E} . The introduction of $\beta^{*(\hat{E})}$ and $\beta^{**(\hat{E})}$ offers a new way to partition the vector $(\beta_I^T, \beta_{II}^T)^T$, or E , according to the estimated set \hat{E} .

For $U_i(\beta_M, \beta_I, \phi)$ in (4.8) and a generic element E^a in \mathcal{E} , let $U_i^\dagger(\beta_M, \beta^{*(E^a)}, \phi)$ denote the estimating function $U_i(\beta_M, \beta_I, \phi)$ with β_I replaced by $\beta^{*(E^a)}$, and define

$$H(\beta_M, \beta_I, \beta_{II}, \phi) = \begin{pmatrix} E_{Y|X}\{U_i(\beta_M, \beta_I, \phi)\} \\ \beta_{II} \end{pmatrix},$$

$$H_n^\dagger(\beta_M, \beta_I, \beta_{II}, \phi) = \sum_{E^a \in \mathcal{E}} \begin{pmatrix} \frac{1}{n} \sum_{i=1}^n U_i^\dagger(\beta_M, \beta^{*(E^a)}, \phi) \\ \beta^{**(\hat{E}^a)} \end{pmatrix} \cdot I(E^a = \hat{E}), \quad (\text{C.2})$$

and

$$H^\dagger(\beta_M, \beta_I, \beta_{II}, \phi) = E_{Y|X}\{H_n^\dagger(\beta_M, \beta_I, \beta_{II}, \phi)\}, \quad (\text{C.3})$$

the expectation is taken with respect to Y_i given X_i , and $H^\dagger(\beta_M, \beta_I, \beta_{II}, \phi)$ can be written as $\sum_{E^a \in \mathcal{E}} \begin{pmatrix} E_{Y|X}\{U_i^\dagger(\beta_M, \beta^{*(E^a)}, \phi)\} \\ \beta^{**(\hat{E}^a)} \end{pmatrix} \cdot I(E^a = \hat{E})$.

Step 2: To show the consistency of $(\hat{\beta}_M^T, \hat{\beta}_I^T, \hat{\phi}^T)^T$, we apply Theorem 5.9 of [Van der Vaart \(2000, Page 46\)](#) by varying the required conditions. That is, it suffices to show that

$$\text{Claim 1: } \inf_{(\beta^T, \phi^T)^T \in \mathcal{B}(\eta)} \|H(\beta, \phi)\| > 0 = \|H(\beta_0, \phi_0)\|.$$

$$\text{Claim 2: } \sup_{\beta, \phi} \|H_n^\dagger(\beta, \phi) - H(\beta, \phi)\| \xrightarrow{p} 0 \text{ as } n \rightarrow \infty.$$

Step 3: Show Claim 1.

By Condition (R10) and the definition of β_{II} , $(\beta_{\text{M}(0)}^{\text{T}}, \beta_{\text{I}(0)}^{\text{T}}, \beta_{\text{II}(0)}^{\text{T}}, \phi_0^{\text{T}})^{\text{T}}$ is the unique solution to the equation $H(\beta_{\text{M}}, \beta_{\text{I}}, \beta_{\text{II}}, \phi) = 0$.

Given the data, by Conditions (R6) and (R7), applying by the Heine-Cantor Theorem (Rudin, 1976, Theorem 4.19), $(U_i^{\text{T}}(\beta_{\text{M}}, \beta_{\text{I}}, \phi), \beta_{\text{II}}^{\text{T}})^{\text{T}}$ is uniformly continuous in β and ϕ , implying that $\|H(\beta, \phi)\|$ is continuous, where $\|\cdot\|$ is the Euclidean norm. Since the set $\mathcal{B}(\eta) = \{(\beta^{\text{T}}, \phi^{\text{T}})^{\text{T}} \in \mathcal{B} : \|(\beta^{\text{T}}, \phi^{\text{T}})^{\text{T}} - (\beta_0^{\text{T}}, \phi_0^{\text{T}})^{\text{T}}\| \geq \eta\}$ is a compact subset of \mathcal{B} for any $\eta > 0$, there exists $(\beta_1^{\text{T}}, \phi_1^{\text{T}})^{\text{T}} \in \mathcal{B}(\eta)$ such that

$$\inf_{(\beta^{\text{T}}, \phi^{\text{T}})^{\text{T}} \in \mathcal{B}(\eta)} \|H(\beta, \phi)\| = \|H(\beta_1, \phi_1)\|.$$

Since $(\beta_0^{\text{T}}, \phi_0^{\text{T}})^{\text{T}}$ is the unique solution of $H(\beta, \phi) = 0$, then for any $(\beta_1^{\text{T}}, \phi_1^{\text{T}})^{\text{T}} \neq (\beta_0^{\text{T}}, \phi_0^{\text{T}})^{\text{T}}$, we have that $\|H(\beta_1, \phi_1)\| > 0$. That is,

$$\inf_{(\beta^{\text{T}}, \phi^{\text{T}})^{\text{T}} \in \mathcal{B}(\eta)} \|H(\beta, \phi)\| > 0 = \|H(\beta_0, \phi_0)\|.$$

Step 4: Show Claim 2.

Noting that

$$\sup_{\beta, \phi} \|H_n^{\dagger}(\beta, \phi) - H(\beta, \phi)\| \leq \sup_{\beta, \phi} \|H_n^{\dagger}(\beta, \phi) - H^{\dagger}(\beta, \phi)\| + \sup_{\beta, \phi} \|H^{\dagger}(\beta, \phi) - H(\beta, \phi)\|, \quad (\text{C.4})$$

we examine the two terms on the right-hand side of (C.4) separately.

1°. For the first term on the right-hand side of (C.4), by (C.2) and (C.3),

$$\begin{aligned} & \sup_{\beta, \phi} \|H_n^{\dagger}(\beta, \phi) - H^{\dagger}(\beta, \phi)\| \\ &= \sup_{\beta, \phi} \left\| \sum_{E^a \in \mathcal{E}} \left(\frac{1}{n} \sum_{i=1}^n U_i^{\dagger}(\beta_{\text{M}}, \beta^{*(E^a)}, \phi) - E\{U_i^{\dagger}(\beta_{\text{M}}, \beta^{*(E^a)}, \phi)\} \right) \cdot I(E^a = \widehat{E}) \right\|, \\ &= \sup_{\beta, \phi} \left\| \sum_{E^a \in \mathcal{E}} \left[\frac{1}{n} \sum_{i=1}^n U_i^{\dagger}(\beta_{\text{M}}, \beta^{*(E^a)}, \phi) - E\{U_i^{\dagger}(\beta_{\text{M}}, \beta^{*(E^a)}, \phi)\} \right] \cdot I(E^a = \widehat{E}) \right\|, \\ &= \sup_{\beta, \phi} \left\| \frac{1}{n} \sum_{i=1}^n U_i^{\dagger}(\beta_{\text{M}}, \beta^{*(\widehat{E})}, \phi) - E\{U_i^{\dagger}(\beta_{\text{M}}, \beta^{*(\widehat{E})}, \phi)\} \right\|. \end{aligned} \quad (\text{C.5})$$

Now we show the convergence of $\frac{1}{n} \sum_{i=1}^n U_i^\dagger(\beta_M, \beta^{*(\widehat{E})}, \phi)$ for a given estimated graph \widehat{E} . By Condition (R9) and the uniform weak law of large numbers (Newey and McFadden, 1994, Lemma 2.4), we have that

$$\sup_{\beta, \phi} \left\| \frac{1}{n} \sum_{i=1}^n U_i^\dagger(\beta_M, \beta^{*(\widehat{E})}, \phi) - E\{U_i^\dagger(\beta_M, \beta^{*(\widehat{E})}, \phi)\} \right\| \xrightarrow{p} 0,$$

and thus by (C.5),

$$\sup_{\beta, \phi} \|H_n^\dagger(\beta, \phi) - H^\dagger(\beta, \phi)\| \xrightarrow{p} 0 \quad \text{as } n \rightarrow \infty. \quad (\text{C.6})$$

2°. Next, for the second term on the right-hand side of (C.4), we have that

$$\begin{aligned} & \lim_{n \rightarrow \infty} P \left(\sup_{\beta, \phi} \|H^\dagger(\beta, \phi) - H(\beta, \phi)\| > \epsilon \right) \\ &= \lim_{n \rightarrow \infty} P \left(\sup_{\beta, \phi} \|H^\dagger(\beta, \phi) - H(\beta, \phi)\| > \epsilon \mid \widehat{E} = E \right) P(\widehat{E} = E) \\ & \quad + \lim_{n \rightarrow \infty} P \left(\sup_{\beta, \phi} \|H^\dagger(\beta, \phi) - H(\beta, \phi)\| > \epsilon \mid \widehat{E} \neq E \right) P(\widehat{E} \neq E) \\ &= \lim_{n \rightarrow \infty} P \left(\sup_{\beta, \phi} \|H^\dagger(\beta, \phi) - H(\beta, \phi)\| > \epsilon \mid \widehat{E} = E \right) \\ &= \lim_{n \rightarrow \infty} P \left(\sup_{\beta, \phi} \left\| \sum_{E^a \in \mathcal{E}} \left(E\{U_i^\dagger(\beta_M, \beta^{*(E^a)}, \phi)\} \right) \cdot I(E^a = \widehat{E}) - H(\beta, \phi) \right\| > \epsilon \mid \widehat{E} = E \right) \\ &= \lim_{n \rightarrow \infty} P \left(\sup_{\beta, \phi} \left\| \sum_{E^a \in \mathcal{E}} \left(E\{U_i^\dagger(\beta_M, \beta^{*(E^a)}, \phi)\} \right) \cdot I(E^a = E) - H(\beta, \phi) \right\| > \epsilon \right) \\ &= \lim_{n \rightarrow \infty} P \left(\sup_{\beta, \phi} \left\| \left(E\{U_i^\dagger(\beta_M, \beta^{*(E)}, \phi)\} \right) - H(\beta, \phi) \right\| > \epsilon \right) = 0, \end{aligned}$$

where the second step is because of $\lim_{n \rightarrow \infty} P(\widehat{E} \neq E) = 0$, the third step is because of (C.3) and (C.2), and last step is by the definition $H(\beta, \phi)$.

Therefore,

$$\sup_{\beta, \phi} \|H^\dagger(\beta, \phi) - H(\beta, \phi)\| \xrightarrow{p} 0 \text{ as } n \rightarrow \infty. \quad (\text{C.7})$$

Combining (C.4), (C.6) and (C.7) shows Claim 2.

Proof of Theorem 4.2 (ii)

By (C.1),

$$P(\widehat{E} = E) \rightarrow 0 \text{ as } n \rightarrow \infty,$$

which implies that as $n \rightarrow \infty$

$$P \left\{ \sum_{i=1}^n U_i^\dagger(\beta_M, \beta^{*(\widehat{E})}, \phi) = \sum_{i=1}^n U_i(\beta_M, \beta_I, \phi) \right\} \rightarrow 0,$$

and

$$P \left\{ \sum_{i=1}^n \left(\frac{\partial U_i^\dagger(\beta_M, \beta^{*(\widehat{E})}, \phi)}{\partial \beta_M^T} \quad \frac{\partial U_i(\beta_M, \beta^{*(\widehat{E})}, \phi)}{\partial \beta_I^T} \quad \frac{\partial U_i(\beta_M, \beta^{*(\widehat{E})}, \phi)}{\partial \phi^T} \right) = \sum_{i=1}^n \left(\frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \beta_M^T} \quad \frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \beta_I^T} \quad \frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \phi^T} \right) \right\} \rightarrow 0,$$

because by definition of $U_i^\dagger(\beta_M, \beta^{*(\widehat{E})}, \phi)$, we have that $U_i^\dagger(\beta_M, \beta^{*(\widehat{E})}, \phi) = U_i(\beta_M, \beta_I, \phi)$ if $\widehat{E} = E$.

Hence, we have that

$$\sum_{i=1}^n U_i^\dagger(\beta_M, \beta^{*(\widehat{E})}, \phi) = \sum_{i=1}^n U_i(\beta_M, \beta_I, \phi) + o_p(1) \quad (\text{C.8})$$

and

$$\begin{aligned} & \sum_{i=1}^n \left(\frac{\partial U_i^\dagger(\beta_M, \beta^{*(\widehat{E})}, \phi)}{\partial \beta_M^T} \quad \frac{\partial U_i(\beta_M, \beta^{*(\widehat{E})}, \phi)}{\partial \beta_I^T} \quad \frac{\partial U_i(\beta_M, \beta^{*(\widehat{E})}, \phi)}{\partial \phi^T} \right) \\ &= \sum_{i=1}^n \left(\frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \beta_M^T} \quad \frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \beta_I^T} \quad \frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \phi^T} \right) + o_p(1). \end{aligned}$$

Applying the Taylor series expansion to $\sum_{i=1}^n U_i^\dagger(\widehat{\beta}_M, \widehat{\beta}^{*(\widehat{E})}, \widehat{\phi}) = 0$ around $(\beta_{M(0)}^T, \beta_{I(0)}^T, \phi_0^T)^T$, we obtain

$$\begin{aligned} & \sum_{i=1}^n U_i(\beta_{M(0)}, \beta_{I(0)}, \phi) + \sum_{i=1}^n \left(\frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \beta_M^T} \quad \frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \beta_I^T} \quad \frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \phi^T} \right) \Bigg|_{\substack{\beta_M = \beta_{M(0)} \\ \beta_I = \beta_{I(0)} \\ \phi = \phi_0}} \\ & \quad \times \left\{ \begin{pmatrix} \widehat{\beta}_M \\ \widehat{\beta}_I \\ \widehat{\phi} \end{pmatrix} - \begin{pmatrix} \beta_{M(0)} \\ \beta_{I(0)} \\ \phi_0 \end{pmatrix} \right\} + o_p(1) = 0, \end{aligned}$$

yielding

$$\begin{aligned} & \left\{ -\frac{1}{n} \sum_{i=1}^n \left(\frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \beta_M^T} \quad \frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \beta_I^T} \quad \frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \phi^T} \right) \right\} \Bigg|_{\substack{\beta_M = \beta_{M(0)} \\ \beta_I = \beta_{I(0)} \\ \phi = \phi_0}} \times \sqrt{n} \left\{ \begin{pmatrix} \widehat{\beta}_M \\ \widehat{\beta}_I \\ \widehat{\phi} \end{pmatrix} - \begin{pmatrix} \beta_{M(0)} \\ \beta_{I(0)} \\ \phi_0 \end{pmatrix} \right\} \\ &= \frac{1}{\sqrt{n}} \sum_{i=1}^n U_i(\beta_{M(0)}, \beta_{I(0)}, \phi_0) + o_p(1). \end{aligned} \quad (\text{C.9})$$

Let

$$\Gamma_0 = \lim_{n \rightarrow \infty} E_{(X, Y)} \left\{ \frac{1}{n} \sum_{i=1}^n \left(\frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \beta_M^T} \quad \frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \beta_I^T} \quad \frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \phi^T} \right) \right\} \Bigg|_{\substack{\beta_M = \beta_{M(0)} \\ \beta_I = \beta_{I(0)} \\ \phi = \phi_0}}$$

and

$$\Sigma_0 = \lim_{n \rightarrow \infty} \text{Var} \left\{ \frac{1}{\sqrt{n}} \sum_{i=1}^n U_i(\beta_{M(0)}, \beta_{I(0)}, \phi_0) \right\}.$$

Then applying the central limit theorem to (C.9) gives that

$$\sqrt{n} \left\{ (\widehat{\beta}_M^T, \widehat{\beta}_I^T, \widehat{\phi}^T)^T - (\beta_{M(0)}^T, \beta_{I(0)}^T, \phi_0^T)^T \right\} \xrightarrow{d} N(0, \Gamma_0^{-1} \Sigma_0 (\Gamma_0^{-1})^T) \quad \text{as } n \rightarrow \infty,$$

where the expectation and variance are taken with respect to the joint distribution of Y_i and X_i with the model parameters taken as their true value.

Now it remains to show that Γ_0 and Σ_0 are identical to those specified as in (4.9). By the assumption that the variables are independent and identically are distributed, it is immediate that

$$\Gamma_0 = \left[E \left\{ \frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \beta_M^T} \right\} \quad E \left\{ \frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \beta_I^T} \right\} \quad E \left\{ \frac{\partial U_i(\beta_M, \beta_I, \phi)}{\partial \phi^T} \right\} \right] \Bigg|_{\substack{\beta_M = \beta_{M(0)} \\ \beta_I = \beta_{I(0)} \\ \phi = \phi_0}}$$

and

$$\begin{aligned} \Sigma_0 &= \lim_{n \rightarrow \infty} \text{Var} \left\{ \frac{1}{\sqrt{n}} \sum_{i=1}^n U_i(\beta_{M(0)}, \beta_{I(0)}, \phi_0) \right\} \\ &= \lim_{n \rightarrow \infty} \frac{1}{n} \sum_{i=1}^n \text{Var} \{ U_i(\beta_{M(0)}, \beta_{I(0)}, \phi_0) \} \\ &= E \{ U_i^T(\beta_{M(0)}, \beta_{I(0)}, \phi_0) U_i(\beta_{M(0)}, \beta_{I(0)}, \phi_0) \} - E^T \{ U_i(\beta_{M(0)}, \beta_{I(0)}, \phi_0) \} E \{ U_i(\beta_{M(0)}, \beta_{I(0)}, \phi_0) \} \\ &= E \{ U_i^T(\beta_{M(0)}, \beta_{I(0)}, \phi_0) U_i(\beta_{M(0)}, \beta_{I(0)}, \phi_0) \}. \end{aligned}$$

C.4 Proof of Theorem 4.4

To reflect the randomness introduced by the selection of the edge set, we use $U_i^{*\dagger}(\beta_M, \beta^{*(\widehat{E})}, \phi, \eta)$ to denote the estimating function $U_i^*(\beta_M, \beta_I, \phi, \eta)$ in (4.14) with the edge set \widehat{E} , estimate by \widehat{E} , where $\beta^{*(\widehat{E})} = \{\beta_{st} : (s, t) \in \widehat{E}\}$ represents the subvector of $(\beta_I^T, \beta_{II}^T)^T$ with the indexes included in \widehat{E} .

Assume the external validation sample is randomly formed. To simplify the notation, let $\beta = (\beta_M^T, \beta_I^T, \phi^T)^T$ and $\beta^\dagger = (\beta_M^T, \beta^{*(\widehat{E})}, \phi^T)^T$. Let $\delta_i = I(i \in \mathcal{V})$, where $I(\cdot)$ is the indicator function. Define $F_{i1}(\beta, \eta) = (1 - \delta_i)U_i^*(\beta, \eta)$, $F_{i2}(\eta) = \delta_i S_i(\eta)$, and $F_i(\beta, \eta) = \{F_{i1}^T(\beta, \eta), F_{i2}^T(\eta)\}^T$, and define $F_{i1}^\dagger(\beta^\dagger, \eta) = (1 - \delta_i)U_i^{*\dagger}(\beta^\dagger, \eta)$, and $F_i^\dagger(\beta^\dagger, \eta) = \{F_{i1}^{\dagger T}(\beta^\dagger, \eta), F_{i2}^T(\eta)\}^T$. Then, (4.14) is equivalent to solving

$$U^{(\text{EV})}(\beta, \eta) = \sum_{i \in \mathcal{M} \cup \mathcal{V}} F_i^\dagger(\beta^\dagger, \eta).$$

Similar to the proof of Theorem 4.2(i), with $U_i(\beta_M, \beta_I, \phi)$ replaced by $F_i(\beta, \eta)$ and $U_i^\dagger(\beta_M, \beta^{*(\widehat{E})}, \phi)$ replaced by $F_i^\dagger(\beta^\dagger, \eta)$, we can show that solving $\sum_{i \in \mathcal{M} \cup \mathcal{V}} F_i^\dagger(\beta^\dagger, \eta) = 0$ gives a consistent estimator, $(\widehat{\beta}^{(\text{EV})T}, \widehat{\eta}^{(\text{EV})T})^T$, of $(\beta^T, \eta^T)^T$.

Similar to the derivation of (C.8) in the proof of Theorem 4.2(ii), we have that

$$\sum_{i=1}^n U_i^{*\dagger}(\beta_M, \beta^{*(E)}, \phi, \eta) = \sum_{i=1}^n U_i^*(\beta_M, \beta_I, \phi, \eta) + o_p(1)$$

and hence

$$\sum_{i \in \mathcal{M} \cup \mathcal{V}} F_i^\dagger(\beta^\dagger, \eta) = \sum_{i \in \mathcal{M} \cup \mathcal{V}} F_i(\beta, \eta) + o_p(1).$$

Applying the Taylor series expansion to $\sum_{i \in \mathcal{M} \cup \mathcal{V}} F_i^\dagger(\widehat{\beta}^{(\text{EV})}, \widehat{\eta}^{(\text{EV})}) = 0$, we obtain

$$\sum_{i \in \mathcal{M} \cup \mathcal{V}} F_i(\beta_0, \eta_0) + \sum_{i \in \mathcal{M} \cup \mathcal{V}} \left(\frac{\partial F_i(\beta, \eta)}{\partial \beta^T} \quad \frac{\partial F_i(\beta, \eta)}{\partial \eta^T} \right) \Bigg|_{\substack{\beta=\beta_0 \\ \eta=\eta_0}} \times \left\{ \begin{pmatrix} \widehat{\beta} \\ \widehat{\eta} \end{pmatrix} - \begin{pmatrix} \beta_0 \\ \eta_0 \end{pmatrix} \right\} + o_p(1) = 0,$$

which leads to

$$\begin{aligned} & \sqrt{1 + \frac{m}{n}} \left\{ \frac{1}{n+m} \sum_{i \in \mathcal{M} \cup \mathcal{V}} \left(\frac{\partial F_i(\beta, \eta)}{\partial \beta^T} \quad \frac{\partial F_i(\beta, \eta)}{\partial \eta^T} \right) \right\} \Bigg|_{\substack{\beta=\beta_0 \\ \eta=\eta_0}} \times \sqrt{n} \left\{ \begin{pmatrix} \widehat{\beta} \\ \widehat{\eta} \end{pmatrix} - \begin{pmatrix} \beta_0 \\ \eta_0 \end{pmatrix} \right\} \\ &= \frac{1}{\sqrt{n+m}} \sum_{i \in \mathcal{M} \cup \mathcal{V}} F_i(\beta_0, \eta_0) + o_p(1). \end{aligned} \tag{C.10}$$

Let

$$\Gamma_{(\text{EV})} = \lim_{n \rightarrow \infty} \left\{ \frac{1}{n+m} \sum_{i \in \mathcal{M} \cup \mathcal{V}} E \left(\begin{array}{cc} \frac{\partial F_i(\beta, \eta)}{\partial \beta^{\text{T}}} & \frac{\partial F_i(\beta, \eta)}{\partial \eta^{\text{T}}} \end{array} \right) \right\} \Bigg|_{\substack{\beta = \beta_0 \\ \eta = \eta_0}}$$

and

$$\Sigma_{(\text{EV})} = \lim_{n \rightarrow \infty} \text{Var} \left(\frac{1}{\sqrt{n+m}} \sum_{i \in \mathcal{M} \cup \mathcal{V}} F_i(\beta_0, \eta_0) \right).$$

Then applying the central limit theorem to (C.10) gives that

$$\sqrt{n} \left\{ (\widehat{\beta}^{(\text{EV})\text{T}}, \widehat{\eta}^{(\text{EV})\text{T}})^{\text{T}} - (\beta_0^{\text{T}}, \eta_0^{\text{T}})^{\text{T}} \right\} \xrightarrow{d} N \left(0, \frac{1}{1+\rho} \Gamma_{(\text{EV})}^{-1} \Sigma_{(\text{EV})} (\Gamma_{(\text{EV})}^{-1})^{\text{T}} \right) \quad \text{as } n \rightarrow \infty.$$

Now it remains to show that $\Gamma_{(\text{EV})}$ and $\Sigma_{(\text{EV})}$ are identical to (4.15). By the definitions of $F_i(\beta, \eta)$, we derive that

$$\begin{aligned} \Gamma_{(\text{EV})} &= \lim_{n \rightarrow \infty} \left\{ \frac{1}{n+m} E \left(\begin{array}{cc} \sum_{i \in \mathcal{M}} \frac{\partial U_i^*(\beta, \eta)}{\partial \beta^{\text{T}}} & \sum_{i \in \mathcal{M}} \frac{\partial U_i^*(\beta, \eta)}{\partial \eta^{\text{T}}} \\ 0 & 0 \end{array} \right) + \frac{1}{n+m} E \left(\begin{array}{cc} 0 & 0 \\ 0 & \sum_{i \in \mathcal{V}} \frac{\partial S_i(\eta)}{\partial \eta^{\text{T}}} \end{array} \right) \right\} \\ &= \lim_{n \rightarrow \infty} \left\{ \frac{n}{n+m} E \left(\begin{array}{cc} \frac{\partial U_i^*(\beta, \eta)}{\partial \beta^{\text{T}}} & \frac{\partial U_i^*(\beta, \eta)}{\partial \eta^{\text{T}}} \\ 0 & 0 \end{array} \right) + \frac{m}{n+m} E \left(\begin{array}{cc} 0 & 0 \\ 0 & \frac{\partial S_i(\eta)}{\partial \eta^{\text{T}}} \end{array} \right) \right\} \\ &= \frac{1}{1+\rho} \left(E \left(\begin{array}{cc} \frac{\partial U_i^*(\beta, \eta)}{\partial \beta^{\text{T}}} & \frac{\partial U_i^*(\beta, \eta)}{\partial \eta^{\text{T}}} \\ 0 & 0 \end{array} \right) + \frac{\rho}{1+\rho} \left(\begin{array}{cc} 0 & 0 \\ 0 & E \left(\frac{\partial S_i(\eta)}{\partial \eta^{\text{T}}} \right) \end{array} \right) \right). \end{aligned}$$

For $\Sigma_{(\text{EV})}$, we have that

$$\begin{aligned}
\Sigma_{(\text{EV})} &= \lim_{n \rightarrow \infty} \frac{1}{n+m} \sum_{i \in \mathcal{M} \cup \mathcal{V}} \text{Var}\{F_i(\beta, \eta)\} \\
&= \lim_{n \rightarrow \infty} \frac{1}{n+m} \sum_{i \in \mathcal{M} \cup \mathcal{V}} E\{F_i(\beta, \eta)F_i^{\text{T}}(\beta, \eta)\} \\
&= \lim_{n \rightarrow \infty} \frac{1}{n+m} \sum_{i \in \mathcal{M} \cup \mathcal{V}} E \begin{pmatrix} (1-\delta_i)^2 U_i^*(\beta, \eta) U_i^{*\text{T}}(\beta, \eta) & 0 \\ 0 & \delta_i^2 S_i(\eta) S_i^{\text{T}}(\eta) \end{pmatrix} \\
&= \lim_{n \rightarrow \infty} \frac{n}{n+m} \frac{1}{n} \sum_{i \in \mathcal{M}} \begin{pmatrix} E\{U_i^*(\beta, \eta) U_i^{*\text{T}}(\beta, \eta)\} & 0 \\ 0 & 0 \end{pmatrix} \\
&\quad + \lim_{n \rightarrow \infty} \frac{m}{n+m} \frac{1}{m} \sum_{i \in \mathcal{V}} \begin{pmatrix} 0 & 0 \\ 0 & E\{S_i(\eta) S_i^{\text{T}}(\eta)\} \end{pmatrix} \\
&= \frac{1}{1+\rho} \begin{pmatrix} E\{U_i^*(\beta, \eta) U_i^{*\text{T}}(\beta, \eta)\} & 0 \\ 0 & 0 \end{pmatrix} + \frac{\rho}{1+\rho} \begin{pmatrix} 0 & 0 \\ 0 & E\{S_i(\eta) S_i^{\text{T}}(\eta)\} \end{pmatrix},
\end{aligned}$$

where the fourth step is due to $(1 - \delta_i)^2 = (1 - \delta_i)$ and $\delta_i^2 = \delta_i$.

C.5 Proof of Theorem 4.5

Similar to Section C.4, we assume the internal validation subsample is randomly formed. We define $F_{i1}(\beta, \eta) = (1 - \delta_i)U_{i1}^*(\beta, \eta) + \delta_i U_{i1}(\beta, \eta)$, $F_{i2}(\beta, \eta) = \delta_i S_i(\eta)$, and $F_i(\beta, \eta) = \{F_{i1}^{\text{T}}(\beta, \eta), F_{i2}^{\text{T}}(\eta)\}^{\text{T}}$, and define $F_{i1}^{\dagger}(\beta^{\dagger}, \eta) = (1 - \delta_i)U_{i1}^*(\beta^{\dagger}, \eta) + \delta_i U_{i1}(\beta^{\dagger}, \eta)$, and $F_i^{\dagger}(\beta^{\dagger}, \eta) = \{F_{i1}^{\dagger\text{T}}(\beta^{\dagger}, \eta), F_{i2}^{\text{T}}(\eta)\}^{\text{T}}$. Then, (4.17) is equivalent to

$$U^{(\text{iv})}(\beta, \eta) = \sum_{i \in \mathcal{M}} F_i^{\dagger}(\beta^{\dagger}, \eta).$$

Similar to the proof of Theorem 4.2(i), with $U_i(\beta_{\text{M}}, \beta_{\text{I}}, \phi)$ replaced by $F_i(\beta, \eta)$ and $U_i^{\dagger}(\beta_{\text{M}}, \beta^*(\hat{E}), \phi)$ replaced by $F_i^{\dagger}(\beta^{\dagger}, \eta)$, we conclude that under regularity conditions, solving $\sum_{i \in \mathcal{M}} F_i^{\dagger}(\beta^{\dagger}, \eta) = 0$ gives a consistent estimator, $(\hat{\beta}^{(\text{iv})\text{T}}, \hat{\eta}^{(\text{iv})\text{T}})^{\text{T}}$, of $(\beta_0, \eta_0)^{\text{T}}$.

Applying the Taylor series expansion to $\sum_{i \in \mathcal{M}} F_i(\hat{\beta}^{(\text{iv})\text{T}}, \hat{\eta}^{(\text{iv})\text{T}}) = 0$, we obtain

$$\sum_{i \in \mathcal{M}} F_i(\beta_0, \eta_0) + \sum_{i \in \mathcal{M}} \left(\frac{\partial F_i(\beta, \eta)}{\partial \beta^{\text{T}}} \quad \frac{\partial F_i(\beta, \eta)}{\partial \eta^{\text{T}}} \right) \Bigg|_{\substack{\beta = \beta_0 \\ \eta = \eta_0}} \left\{ \begin{pmatrix} \hat{\beta} \\ \hat{\eta} \end{pmatrix} - \begin{pmatrix} \beta_0 \\ \eta_0 \end{pmatrix} \right\} + o_p(1) = 0,$$

which leads to

$$\left\{ -\frac{1}{n} \sum_{i \in \mathcal{M}} \begin{pmatrix} \frac{\partial F_i(\beta, \eta)}{\partial \beta^T} & \frac{\partial F_i(\beta, \eta)}{\partial \eta^T} \end{pmatrix} \right\} \Big|_{\substack{\beta = \beta_0 \\ \eta = \eta_0}} \times \sqrt{n} \left\{ \begin{pmatrix} \hat{\beta} \\ \hat{\eta} \end{pmatrix} - \begin{pmatrix} \beta_0 \\ \eta_0 \end{pmatrix} \right\} = \frac{1}{\sqrt{n}} \sum_{i \in \mathcal{M}} F_i(\beta_0, \eta_0) + o_p(1). \quad (\text{C.11})$$

Let

$$\Gamma_{(\text{iv})} = \lim_{n \rightarrow \infty} \left\{ \frac{1}{n} \sum_{i \in \mathcal{M}} E \begin{pmatrix} \frac{\partial F_i(\beta, \eta)}{\partial \beta^T} & \frac{\partial F_i(\beta, \eta)}{\partial \eta^T} \end{pmatrix} \right\} \Big|_{\substack{\beta = \beta_0 \\ \eta = \eta_0}}$$

and

$$\Sigma_{(\text{iv})} = \lim_{n \rightarrow \infty} \text{Var} \left(\frac{1}{\sqrt{n}} \sum_{i \in \mathcal{M}} F_i(\beta_0, \eta_0) \right).$$

Then applying the central limit theorem to (C.11) gives that

$$\sqrt{n} \left\{ \begin{pmatrix} \hat{\beta}^{(\text{iv})T} & \hat{\eta}^{(\text{iv})T} \end{pmatrix}^T - \begin{pmatrix} \beta_0^T & \eta_0^T \end{pmatrix}^T \right\} \xrightarrow{d} N \left(0, \Gamma_{(\text{iv})}^{-1} \Sigma_{(\text{iv})} (\Gamma_{(\text{iv})}^{-1})^T \right) \quad \text{as} \quad n \rightarrow \infty.$$

Now it remains to show that $\Gamma_{(\text{iv})}$ and $\Sigma_{(\text{iv})}$ are identical to (4.19). By definition of $F_i(\beta, \eta)$, $\Gamma_{(\text{iv})}$ equals

$$\begin{aligned} & \lim_{n \rightarrow \infty} \left\{ \frac{1}{n} E \begin{pmatrix} \sum_{i \in \mathcal{M} \setminus \mathcal{V}} \frac{\partial U_i^*(\beta, \eta)}{\partial \beta^T} & \sum_{i \in \mathcal{M} \setminus \mathcal{V}} \frac{\partial U_i^*(\beta, \eta)}{\partial \eta^T} \\ 0 & 0 \end{pmatrix} + \frac{1}{n} E \begin{pmatrix} \sum_{i \in \mathcal{V}} \frac{\partial U_i(\beta, \eta)}{\partial \beta^T} & 0 \\ 0 & \sum_{i \in \mathcal{V}} \frac{\partial S_i(\eta)}{\partial \eta^T} \end{pmatrix} \right\} \\ &= \lim_{n \rightarrow \infty} \left\{ \frac{n-m}{n} E \begin{pmatrix} \frac{\partial U_i^*(\beta, \eta)}{\partial \beta^T} & \frac{\partial U_i^*(\beta, \eta)}{\partial \eta^T} \\ 0 & 0 \end{pmatrix} + \frac{m}{n} E \begin{pmatrix} \frac{\partial U_i(\beta, \eta)}{\partial \beta^T} & 0 \\ 0 & \frac{\partial S_i(\eta)}{\partial \eta^T} \end{pmatrix} \right\} \\ &= (1-\rho) \begin{pmatrix} E \left(\frac{\partial U_i^*(\beta, \eta)}{\partial \beta^T} \right) & E \left(\frac{\partial U_i^*(\beta, \eta)}{\partial \eta^T} \right) \\ 0 & 0 \end{pmatrix} + \rho \begin{pmatrix} E \left(\frac{\partial U_i(\beta, \eta)}{\partial \beta^T} \right) & 0 \\ 0 & E \left(\frac{\partial S_i(\eta)}{\partial \eta^T} \right) \end{pmatrix}. \end{aligned}$$

Similarly, we have that

$$\begin{aligned}
\Sigma_{(iv)} &= \lim_{n \rightarrow \infty} \frac{1}{n} \sum_{i \in \mathcal{M}} \text{Var} \{F_i(\beta, \eta)\} \\
&= \lim_{n \rightarrow \infty} \frac{1}{n} \sum_{i \in \mathcal{M}} E\{F_i(\beta, \eta)F_i^T(\beta, \eta)\} \\
&= \lim_{n \rightarrow \infty} \frac{1}{n} \sum_{i \in \mathcal{M}} E \begin{pmatrix} (1 - \delta_i)^2 U_i^*(\beta, \eta)U_i^{*\text{T}}(\beta, \eta) & 0 \\ 0 & 0 \end{pmatrix} \\
&\quad + \lim_{n \rightarrow \infty} \frac{1}{n} \sum_{i \in \mathcal{M}} E \begin{pmatrix} \delta_i^2 U_i(\beta, \eta)U_i^T(\beta, \eta) & \delta_i^2 U_i(\beta, \eta)S_i^T(\eta) \\ \delta_i^2 U_i(\beta, \eta)S_i^T(\eta) & \delta_i^2 S_i(\eta)S_i^T(\eta) \end{pmatrix} \\
&= \lim_{n \rightarrow \infty} \frac{n - m}{n} \frac{1}{n - m} \sum_{i \in \mathcal{M} \setminus \mathcal{V}} \begin{pmatrix} E\{U_i^*(\beta, \eta)U_i^{*\text{T}}(\beta, \eta)\} & 0 \\ 0 & 0 \end{pmatrix} \\
&\quad + \lim_{n \rightarrow \infty} \frac{m}{n} \frac{1}{m} \sum_{i \in \mathcal{V}} \begin{pmatrix} E\{U_i(\beta, \eta)U_i^T(\beta, \eta)\} & E\{U_i(\beta, \eta)S_i^T(\beta, \eta)\} \\ E\{U_i(\beta, \eta)S_i^T(\beta, \eta)\} & E\{S_i(\beta, \eta)S_i^T(\beta, \eta)\} \end{pmatrix} \\
&= (1 - \rho) \begin{pmatrix} E\{U_i^*(\beta, \eta)U_i^{*\text{T}}(\beta, \eta)\} & 0 \\ 0 & 0 \end{pmatrix} \\
&\quad + \rho \begin{pmatrix} E\{U_i(\beta, \eta)U_i^T(\beta, \eta)\} & E\{U_i(\beta, \eta)S_i^T(\eta)\} \\ E\{U_i(\beta, \eta)S_i^T(\eta)\} & E\{S_i(\eta)S_i^T(\eta)\} \end{pmatrix},
\end{aligned}$$

where the third step is due to $\delta_i(1 - \delta_i) = 0$ and the fourth step comes from $(1 - \delta_i)^2 = (1 - \delta_i)$ and $\delta_i^2 = \delta_i$.

C.6 Proof of Theorem 4.6

By (4.12) and (4.20), showing (4.23) is equivalent to showing

$$\{(1 - \rho)\Gamma + \rho\Gamma_0\}^{-1}\{(1 - \rho)\Sigma + \rho\Sigma_0 - \rho\Delta\}\{(1 - \rho)\Gamma + \rho\Gamma_0\}^{-1\text{T}} \leq \Gamma^{-1}\Sigma\Gamma^{-1\text{T}}. \quad (\text{C.12})$$

Left multiplying $(1 - \rho)\Gamma + \rho\Gamma_0$ and right multiplying $\{(1 - \rho)\Gamma + \rho\Gamma_0\}^T$ on both sides of (C.12), we obtain

$$(1 - \rho)\Sigma + \rho\Sigma_0 - \rho\Delta \leq (1 - \rho)^2\Sigma + \rho(1 - \rho)\Gamma_0\Gamma^{-1}\Sigma + (1 - \rho)\rho\Sigma\Gamma^{-1\text{T}}\Gamma_0^T + \rho^2\Gamma_0\Gamma^{-1}\Sigma\Gamma^{-1\text{T}}\Gamma_0^T. \quad (\text{C.13})$$

Left multiplying Γ_0^{-1} and right multiplying Γ_0^{-1T} on the both sides of (C.13) gives

$$\begin{aligned} & (1 - \rho)\Gamma_0^{-1}\Sigma\Gamma_0^{-1T} + \rho\Gamma_0^{-1}\Sigma_0\Gamma_0^{-1T} - \rho\Gamma_0^{-1}\Delta\Gamma_0^{-1T} \\ & \leq (1 - \rho)^2\Gamma_0^{-1}\Sigma\Gamma_0^{-1T} + \rho(1 - \rho)\Gamma^{-1}\Sigma\Gamma_0^{-1T} + (1 - \rho)\rho\Gamma_0^{-1}\Sigma\Gamma^{-1T} + \rho^2\Gamma^{-1}\Sigma\Gamma^{-1T}. \end{aligned}$$

Then combining the terms with $\Gamma_0^{-1}\Sigma\Gamma_0^{-1T}$ and dividing the both sides by ρ , we get

$$\begin{aligned} (1 - \rho)\Gamma_0^{-1}\Sigma\Gamma_0^{-1T} + \Gamma_0^{-1}\Sigma_0\Gamma_0^{-1T} - \Gamma_0^{-1}\Delta\Gamma_0^{-1T} & \leq (1 - \rho)\Gamma^{-1}\Sigma\Gamma_0^{-1T} + (1 - \rho)\Gamma_0^{-1}\Sigma\Gamma^{-1T} \\ & + \rho\Gamma^{-1}\Sigma\Gamma^{-1T}. \end{aligned} \quad (\text{C.14})$$

It now suffices to show that (C.14) is true when the conditions (4.21) and (4.22) are satisfied. For the case with $\rho = 1$, the inequality (C.14) is equivalent to

$$\Gamma_0^{-1}\Sigma_0\Gamma_0^{-1T} - \Gamma_0^{-1}\Delta\Gamma_0^{-1T} \leq \Gamma^{-1}\Sigma\Gamma^{-1T},$$

which is true by the condition (4.21) together with the fact that $\Gamma_0^{-1}\Delta\Gamma_0^{-1T}$ is non-negative definite.

For the case with $\rho < 1$, dividing $(1 - \rho)$ on both sides, (C.14) is equivalent to

$$\Gamma_0^{-1}\Sigma\Gamma_0^{-1T} + \frac{1}{1 - \rho}\Gamma_0^{-1}\Sigma_0\Gamma_0^{-1T} - \frac{1}{1 - \rho}\Gamma_0^{-1}\Delta\Gamma_0^{-1T} \leq \Gamma^{-1}\Sigma\Gamma_0^{-1T} + \Gamma_0^{-1}\Sigma\Gamma^{-1T} + \frac{\rho}{1 - \rho}\Gamma^{-1}\Sigma\Gamma^{-1T}, \quad (\text{C.15})$$

which is true, because

$$\begin{aligned} & \text{the left hand side of (C.15)} \\ & \leq \Gamma_0^{-1}\Sigma\Gamma_0^{-1T} + \frac{1}{1 - \rho}\Gamma_0^{-1}\Sigma_0\Gamma_0^{-1T} - \Gamma_0^{-1}\Delta\Gamma_0^{-1T} \\ & = \Gamma_0^{-1}\Sigma\Gamma_0^{-1T} + \Gamma_0^{-1}\Sigma_0\Gamma_0^{-1T} + \frac{\rho}{1 - \rho}\Gamma_0^{-1}\Sigma_0\Gamma_0^{-1T} - \Gamma_0^{-1}\Delta\Gamma_0^{-1T} \\ & \leq \Gamma_0^{-1}\Sigma\Gamma_0^{-1T} + \Gamma_0^{-1}\Sigma_0\Gamma_0^{-1T} + \frac{\rho}{1 - \rho}\Gamma^{-1}\Sigma\Gamma^{-1T} - \Gamma_0^{-1}\Delta\Gamma_0^{-1T} \\ & \leq \Gamma^{-1}\Sigma\Gamma_0^{-1T} + \Gamma_0^{-1}\Sigma\Gamma^{-1T} + \frac{\rho}{1 - \rho}\Gamma^{-1}\Sigma\Gamma^{-1T} \\ & = \text{the right hand side of (C.15)}, \end{aligned}$$

where the first step is because $\rho \in (0, 1)$, the third step is due to the condition (4.21) that $\Gamma_0^{-1}\Sigma_0\Gamma_0^{-1T} \leq \Gamma^{-1}\Sigma\Gamma^{-1T}$, and the fourth step is due to the condition (4.22).

Appendix D

Proofs of the Results in Chapter 5

D.1 Proof of Theorem 5.1

Proof of Theorem 5.1(a):

First, consider

$$\begin{aligned} P(Y_i^* = 0|X_i) &= \sum_{k=0}^{\infty} P(Y_i^* = 0|Y_i = k, X_i)P(Y_i = k|X_i) \\ &= \sum_{k=0}^{\infty} P(Z_{i-} = k|Y_i = k, X_i)P(Y_i = k|X_i) \\ &= P(Y_i = 0|X_i)P(Z_{i-} = 0|Y_i = 0, X_i) \\ &\quad + \sum_{k=1}^{\infty} P(Z_{i-} = k|Y_i = k, X_i)P(Y_i = k|X_i) \\ &= P(Y_i = 0|X_i) + \sum_{k=1}^{\infty} \binom{k}{k} \pi_i^k (1 - \pi_i)^0 \times \frac{\mu_i^k}{k!} e^{-\mu_i} \\ &= P(Y_i = 0|X_i) + \phi_i \left(\sum_{k=1}^{\infty} \frac{(\pi_i \mu_i)^k}{k!} e^{-\mu_i \pi_i} \right) e^{-\mu_i + \mu_i \pi_i} \\ &= P(Y_i = 0|X_i) + \phi_i (1 - e^{-\mu_i \pi_i}) e^{-\mu_i (1 - \pi_i)} \\ &= 1 - \phi_i + \phi_i e^{-\mu_i (1 - \pi_i)}, \end{aligned} \tag{D.1}$$

where the second step comes from (5.5) with $c_+ = 0$ and $c_- = 1$, the fourth step comes from (5.2) and the distributional assumption of Z_{i-} , the fifth step is because $\sum_{k=0}^{\infty} \frac{(\pi_i \mu_i)^k}{k!} e^{-\mu_i \pi_i} = 1$, and the last step is due to (5.2).

Next, for $Y_i^* = k \geq 1$, we calculate

$$\begin{aligned}
P(Y_i^* = k | X_i) &= \sum_{r=0}^{\infty} P(Z_{i-} = r | Y = k + r, X_i) P(Y = k + r | X_i) \\
&= \phi_i \sum_{r=0}^{\infty} \binom{k+r}{r} \pi_i^r (1 - \pi_i)^k \times \frac{\mu_i^{k+r}}{(k+r)!} e^{-\mu_i} \\
&= \phi_i \sum_{r=0}^{\infty} \frac{1}{r! k!} (1 - \pi_i)^k \mu_i^k (\mu_i \pi_i)^r e^{-\mu_i} \\
&= \phi_i \frac{\{\mu_i(1 - \pi_i)\}^k}{k!} e^{-\mu_i(1 - \pi_i)}, \tag{D.2}
\end{aligned}$$

where the first step is due to (5.5) with $c_+ = 0$ and $c_- = 1$ and the second step is because of (5.2) and the distributional assumption of Z_{i-} .

Therefore, comparing (D.1) and (D.2) to (5.2), we conclude that conditional on X_i , Y_i^* follows a zero-inflated Poisson distribution with mean $\mu_i(1 - \pi_i)$ and the probability ϕ_i .

Proof of Theorem 5.1(b):

First, we consider the case with $Y_i^* = 0$. Under (5.5) with $c_+ = 1$ and $c_- = 0$, we note that $Y_i^* = 0$ if and only if $Y_i = 0$ and $Z_{i+} = 0$. Then,

$$\begin{aligned}
P(Y_i^* = 0 | X_i) &= P(Y_i = 0, Z_{i+} = 0 | X_i) \\
&= (1 - \phi_i + \phi_i e^{-\mu_i}) e^{-\lambda_i} \\
&= (1 - \phi_i) e^{-\lambda_i} + \phi_i e^{-(\mu_i + \lambda_i)},
\end{aligned}$$

where the second step is due to the conditional independence assumption between Y_i and Z_{i+} given X_i , as well as (5.2) and the distributional assumption of Z_{i+} .

Next, for $k \geq 1$, we have that

$$\begin{aligned}
f(Y_i^* = k|X_i) &= P(Y_i = 0, Z_{i+} = k|X_i) + \sum_{t=1}^k P(Y_i = t, Z_{i+} = k - t|X_i) \\
&= (1 - \phi_i + \phi_i e^{-\mu_i}) \frac{\lambda_i^k e^{-\lambda_i}}{k!} + \sum_{t=1}^k \phi_i \frac{\mu_i^t e^{-\mu_i}}{t!} \times \frac{\lambda_i^{k-t}}{(k-t)!} e^{-\lambda_i} \\
&= (1 - \phi_i) \frac{\lambda_i^k e^{-\lambda_i}}{k!} + \sum_{t=0}^k \phi_i \frac{\mu_i^t e^{-\mu_i}}{t!} \times \frac{\lambda_i^{k-t}}{(k-t)!} e^{-\lambda_i} \\
&= (1 - \phi_i) \frac{\lambda_i^k e^{-\lambda_i}}{k!} + \phi_i e^{-\mu_i} e^{-\lambda_i} \sum_{t=0}^k \frac{\mu_i^t \lambda_i^{k-t}}{t!(k-t)!} \\
&= (1 - \phi_i) \frac{\lambda_i^k e^{-\lambda_i}}{k!} + \phi_i \frac{(\mu_i + \lambda_i)^k}{k!} e^{-(\mu_i + \lambda_i)} \sum_{t=0}^k \frac{k!}{t!(k-t)!} \left(\frac{\mu_i}{\mu_i + \lambda_i} \right)^t \left(\frac{\lambda_i}{\mu_i + \lambda_i} \right)^{k-t} \\
&= (1 - \phi_i) \frac{\lambda_i^k e^{-\lambda_i}}{k!} + \phi_i \frac{(\mu_i + \lambda_i)^k}{k!} e^{-(\mu_i + \lambda_i)},
\end{aligned}$$

where the second step is due to the conditional independence assumption between Y_i and Z_{i+} given X_i , as well as (5.2) and the distributional assumption of Z_{i+} , and the last step is due to the Binomial theorem. Thus, the conclusion follows.

Proof of Theorem 5.1(c):

Model (5.5) with $c_+ = c_- = 1$ can be viewed as $Y_i^* = (Y_i - Z_{i-}) + Z_{i+}$, where by Theorem 5.1(a), the first term $(Y_i - Z_{i-})$ follows a zero-inflated Poisson distribution with parameters ϕ_i and $\mu_i^* = (1 - \pi_i)\mu_i$. Then applying Theorem 5.1(b) to Y_i^* , we conclude that

$$P(Y_i^* = y_i^*|X_i) = (1 - \phi_i) \frac{\lambda_i^{y_i^*} e^{-\lambda_i}}{y_i^{*!}} + \phi_i \frac{\mu_i^{y_i^*}}{y_i^{*!}} e^{-\mu_i^*} \quad \text{for } y_i^* = 0, 1, 2, \dots,$$

where $\mu_i^* = (1 - \pi_i)\mu_i + \lambda_i$.

D.2 Proof of Theorem 5.2

Proof:

By Theorem 1(a), we have that if $c_+ = 0$ and $c_- = 1$, Y_i^* follows zero-inflated Poisson distribution with parameter ϕ_i^* and μ_i^* , where $\phi_i^* = \phi_i$ and $\mu_i^* = \mu_i(1 - \pi)$. Thus, by (5.4),

$$\begin{aligned}\log \mu_i^* &= \log(1 - \pi_i) + \log \mu_i \\ &= \log(1 - \pi_i) + \beta_{\mu 0} + \beta_{\mu x}^T X_i \\ &= \beta_{\mu 0}^* + \beta_{\mu x}^{*T} X_i.\end{aligned}\tag{D.3}$$

Comparing (D.3) to (5.12), we conclude that $\beta_{\mu 0}^* = \beta_{\mu 0} + \log(1 - \pi_i)$ and $\beta_{\mu x}^* = \beta_{\mu x}$.

D.3 Inverse Sampling of Multivariate Discrete Variables

To execute inverse sampling for multivariate discrete variables (Loukas and Kemp, 1983), we evaluate the joint distribution of U_{i1} , U_{i2} , Z_{i-} and Z_{i+} , expressed in (5.16). Noting that although U_{i1} and U_{i2} are unbounded (and Z_{i-} is bounded by U_{i2}), the probability (5.16) is extremely small for sufficiently large values. We focus only those values of U_{i1} and U_{i2} bounded by sufficiently large values, say, J and K , respectively, and hence Z_{i-} is also bounded by K . Specifically, for $j = 0, \dots, J$, $k = 0, \dots, K$, and $l = 0, \dots, K$, we evaluate (5.16)

$$\begin{aligned}p_{jkl} &= P(U_{i1} = j, U_{i2} = k, Z_{i-} = l, Z_{i+} = y_i^* - I(j > 0)k + l | x_i) \\ &= P(Z_{i+} = y_i^* - I(j > 0)k + l | x_i) P(Z_{i-} = l | U_{i1} = j, U_{i2} = k | x_i) P(U_{i1} = j | x_i) P(U_{i2} = k | x_i),\end{aligned}$$

where each probability is computed based on the distribution assumed for the associated random variables.

To ensure legitimate probabilities induced from the imposition of bounds to U_{i1} and U_{i2} , we normalize the p_{ijk} by calculating $p_{jkl}^* = \frac{p_{jkl}}{\sum_{j',k',l'} p_{j'k'l'}}$. Let Φ be the $(J+1)(K+1)^2$ -dimensional column vector consisting of the p_{jkl}^* with $j = 0, \dots, J$; $k = 0, \dots, K$; and $l = 0, \dots, K$, and let Φ_t denote the t th element of Φ for $t = 1, \dots, (J+1)(K+1)^2$.

Generate a random value V from Uniform[0,1] and find the smallest x such that $\sum_{t=1}^x \Phi_t \geq V$. Examining Φ_x , we identify j_0 , k_0 and l_0 such that $\Phi_x = p_{j_0 k_0 l_0}^*$. Then we set j_0 , k_0 and l_0 to be the values for U_{i1} , U_{i2} , and Z_{i-} , respectively, and take Z_{i+} as $Y_i^* - I(U_{i1} > 0)U_{i2} + Z_{i-} = y_i^* - I(j > 0)k + l$.

Appendix E

Conditions and Proofs of the Results in Chapter 6

E.1 Regularity Conditions

- (R1) The time series $\{X_t : t = 1, \dots, T\}$ is stationary.
- (R2) The observed error-prone time series $\{X_t^* : t = 1, \dots, T\}$ is stationary.
- (R3) For any $t \in \{1, \dots, T\}$, $\frac{1}{T} \sum_{s=1}^T \gamma_{|s-t|} \rightarrow 0$ as $T \rightarrow \infty$.
- (R4) For any p , $\frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T E\{(X_t - \mu)(X_{t+p} - \mu)(X_s - \mu)\} < \infty$.

While the two process $\{X_t : t = 1, \dots, T\}$ and $\{X_t^* : t = 1, \dots, T\}$ are constrained by the measurement error model (6.7) or (6.9), they can both be assumed to be stationary without inducing conflicting requirements on the associated processes. Obviously, the weak stationarity of $\{X_t : t = 1, \dots, T\}$ implies the weak stationarity of $\{X_t^* : t = 1, \dots, T\}$ if they are linked by (6.7) or (6.9). Condition (R3) says that as the time series goes long enough, the average of the covariances between any paired variables is negligible. Condition (R4) requires the summation of the third moment of X_t is $O(T)$, which is needed in Theorem 6.4 when $\phi_0 \neq 0$; this condition can be satisfied if $E(\epsilon_t^3) = 0$, for example.

E.2 The proof of Theorem 6.1

Applying the weak law of large numbers to $\widehat{\phi}_1^*$ given by (6.12), we obtain that the estimator $\widehat{\phi}_1^*$ converges in probability to $\frac{\text{Cov}(X_t^*, X_{t-1}^*)}{\text{Var}(X_{t-1}^*)}$, which is denoted as ϕ_1^* . Now we further examine ϕ_1^* by using the AR(1) model (6.1) and the measurement error model (6.7):

$$\begin{aligned}
 \phi_1^* &= \frac{\text{Cov}(X_t^*, X_{t-1}^*)}{\text{Var}(X_{t-1}^*)} \\
 &= \frac{\text{Cov}(\alpha_0 + \alpha_1 X_t + e_t, \alpha_0 + \alpha_1 X_{t-1} + e_{t-1})}{\text{Var}(\alpha_0 + \alpha_1 X_t + e_t)} \\
 &= \frac{\alpha_1^2 \text{Cov}(X_t, X_{t-1})}{\alpha_1^2 \text{Var}(X_t) + \text{Var}(e_t)} \\
 &= \frac{\alpha_1^2 \text{Cov}(\phi_0 + \phi_1 X_{t-1} + \epsilon_t, X_{t-1})}{\alpha_1^2 \text{Var}(X_t) + \text{Var}(e_t)} \\
 &= \phi_1 \cdot \frac{\alpha_1^2 \text{Var}(X_{t-1})}{\alpha_1^2 \text{Var}(X_t) + \text{Var}(e_t)},
 \end{aligned}$$

where the second step is due to (6.7), the third step is because of the independence among the X_t and the e_t , and the fourth step is because of (6.1). Since the time series $\{X_t\}$ is stationary, it follows that $\text{Var}(X_t) = \text{Var}(X_{t-1}) = \frac{\sigma_\epsilon^2}{1-\phi_1^2}$, and hence

$$\phi_1^* = \phi_1 \cdot \frac{\alpha_1^2 \sigma_\epsilon^2}{\alpha_1^2 \sigma_\epsilon^2 + \sigma_\epsilon^2 (1 - \phi_1^2)} = \phi_1 \omega_1. \tag{E.1}$$

Next, applying the Slutsky's theorem to (6.12), we have that as $T \rightarrow \infty$,

$$\widehat{\phi}_0^* \xrightarrow{p} E(X_t^*) - \phi_1^* E(X_t^*),$$

where the limit equals $\left(\alpha_0 + \frac{\alpha_1 \phi_0}{1-\phi_1}\right) (1-\phi_1 \omega_1)$ by (E.1) and the fact that $E(X_t^*) = \alpha_0 + \frac{\alpha_1 \phi_0}{1-\phi_1}$.

Finally, plugging the AR(1) model (6.1) into the measurement error model (6.11), we obtain that

$$X_t^* = \alpha_0 + \alpha_1 (\phi_0 + \phi_1 X_{t-1} + \epsilon_t) + e_t. \tag{E.2}$$

On the other hand, plugging the measurement error model (6.7) into the working model (6.11), we obtain that

$$X_t^* = \phi_0^* + \phi_1^* (\alpha_0 + \alpha_1 X_{t-1} + e_t) + \epsilon_t^*. \tag{E.3}$$

Then equating (E.2) and (E.3) that

$$\epsilon^* = \alpha_0(1 - \phi_1^*) + \alpha_1\phi_0 - \phi_0^* + \alpha_1(\phi_1 - \phi_1^*)X_{t-1} + (1 - \phi_1^*)e_t + \alpha_1\epsilon_t.$$

Consequently, by the independence assumption for X_{t-1} , e_t and ϵ_t , we obtain that

$$\begin{aligned} \text{Var}(\epsilon_t^*) &= \phi_1^2\alpha_1^2(1 - \omega_1)^2\text{Var}(X_{t-1}) + (1 - \omega_1\phi_1)^2\text{Var}(e_t) + \alpha_1^2\text{Var}(\epsilon_t) \\ &= \phi_1^2\alpha_1^2(1 - \omega_1)^2 \left(\frac{\sigma_\epsilon^2}{1 - \phi_1^2} \right) + (1 - \omega_1\phi_1)^2\sigma_e^2 + \alpha_1^2\sigma_\epsilon^2. \end{aligned}$$

E.3 The proof of Theorem 6.2

As noted in the beginning of E.2, as $T \rightarrow \infty$, $\hat{\phi}_1^* \xrightarrow{p} \phi_1^*$ where

$$\hat{\phi}_1^* = \frac{\text{Cov}(X_t^*, X_{t-1}^*)}{\text{Var}(X_{t-1}^*)}.$$

Now we further examine ϕ_1^* by using the AR(1) model (6.1) and the measurement error model (6.9):

$$\begin{aligned} \phi_1^* &= \frac{\text{Cov}(X_t^*, X_{t-1}^*)}{\text{Var}(X_{t-1}^*)} \\ &= \frac{\text{Cov}(\beta_0 u_t X_t, \beta_0 u_{t-1} X_{t-1})}{\text{Var}(\beta_0 u_{t-1} X_{t-1})} \\ &= \frac{\beta_0^2 \text{Cov}(u_t X_t, u_{t-1} X_{t-1})}{\beta_0^2 \text{Var}(u_{t-1} X_{t-1})} \\ &= \frac{\text{Cov}\{u_t(\phi_0 + \phi_1 X_{t-1} + \epsilon_t), u_{t-1} X_{t-1}\}}{\text{Var}(X_{t-1} u_{t-1})} \\ &= \phi_1 \frac{\text{Cov}(u_t X_{t-1}, u_{t-1} X_{t-1})}{\text{Var}(u_{t-1} X_{t-1})} \\ &= \phi_1 \frac{E(u_t u_{t-1} X_{t-1}^2) - E(u_t X_{t-1})E(u_{t-1} X_{t-1})}{E(u_{t-1}^2 X_{t-1}^2) - E^2(u_{t-1} X_{t-1})} \\ &= \phi_1 \frac{E(u_t)E(u_{t-1})E(X_{t-1}^2) - E(u_t)E(u_{t-1})E^2(X_{t-1})}{E(u_{t-1}^2)E(X_{t-1}^2) - E^2(u_{t-1} X_{t-1})}, \end{aligned}$$

where the second step is due to measurement error model (6.9).

Then, because u_t , u_{t-1} and X_{t-1} are mutually independent, we further have that

$$\begin{aligned}
\phi_1^* &= \phi_1 \frac{E(u_t)E(u_{t-1})\text{Var}(X_{t-1})}{\{\text{Var}(u_{t-1}) + E^2(u_{t-1})\}\{\text{Var}(X_{t-1}) + E^2(X_{t-1})\} - E^2(u_{t-1})E^2(X_{t-1})} \\
&= \phi_1 \frac{\text{Var}(X_{t-1})}{\{\text{Var}(u_{t-1}) + 1\}\{\text{Var}(X_{t-1}) + E^2(X_{t-1})\} - E^2(X_{t-1})} \\
&= \phi_1 \frac{\text{Var}(X_{t-1})}{\text{Var}(u_{t-1})\text{Var}(X_{t-1}) + \text{Var}(u_{t-1})E^2(X_{t-1}) + \text{Var}(X_{t-1})}, \tag{E.4}
\end{aligned}$$

where the second last step is due to $E(u_t) = 1$. Since the time series $\{X_t\}$ is stationary, it follows that $E(X_t) = E(X_{t-1}) = \frac{\phi_0}{1-\phi_1}$ and $\text{Var}(X_t) = \text{Var}(X_{t-1}) = \frac{\sigma_\epsilon^2}{1-\phi_1^2}$. Hence (E.4) becomes

$$\begin{aligned}
\phi_1^* &= \phi_1 \frac{\text{Var}(X_{t-1})}{\text{Var}(u_{t-1})\text{Var}(X_{t-1}) + \text{Var}(u_{t-1})E^2(X_{t-1}) + \text{Var}(X_{t-1})} \\
&= \phi_1 \frac{\frac{\sigma_\epsilon^2}{1-\phi_1^2}}{\sigma_u^2 \frac{\sigma_\epsilon^2}{1-\phi_1^2} + \sigma_u^2 \left(\frac{\phi_0}{1-\phi_1}\right)^2 + \frac{\sigma_\epsilon^2}{1-\phi_1^2}} \\
&= \phi_1 \frac{\sigma_\epsilon^2}{\sigma_\epsilon^2 \sigma_u^2 + \sigma_\epsilon^2 + \sigma_u^2 \phi_0^2 \frac{1+\phi_1}{1-\phi_1}} = \phi_1 \omega_2. \tag{E.5}
\end{aligned}$$

Next, applying the Slutsky's Theorem to (6.12) gives that as $T \rightarrow \infty$,

$$\widehat{\phi}_0^* \xrightarrow{p} \left(\frac{\beta_0 \phi_0}{1-\phi_1}\right) (1 - \phi_1 \omega_2)$$

by (E.5) as well as $E(X_t^*) = \frac{\beta_0 \phi_0}{1-\phi_1}$.

Finally plugging the AR(1) model (6.1) into the measurement error model (6.9), we obtain that

$$X_t^* = \beta_0(\phi_0 + \phi_1 X_{t-1} + \epsilon_t)u_t. \tag{E.6}$$

On the other hand, plugging the measurement error model (6.9) into the working model (6.11), we obtain that

$$X_t^* = \phi_0^* + \phi_1^*(\beta_0 X_{t-1} u_{t-1}) + \epsilon_t^*. \tag{E.7}$$

Then equating (E.6) and (E.7) gives that

$$\epsilon^* = \beta_0 \phi_0 u_t - \phi_0^* + \beta_0 X_{t-1} (\phi_1 u_t - \omega_2 \phi_1 u_{t-1}) + \beta_0 u_t \epsilon_t.$$

yielding that

$$\begin{aligned}
\text{Var}(\epsilon_t^*) &= \phi_0^2 \beta_0^2 \text{Var}(u_t) + \beta_0^2 \phi_1^2 \text{Var}(X_{t-1} u_t) + \beta_0^2 \omega_2^2 \phi_1^2 \text{Var}(X_{t-1} u_{t-1}) + \beta_0^2 \text{Var}(u_t \epsilon_t) \\
&= \phi_0^2 \beta_0^2 \sigma_u^2 + (\beta_0^2 \phi_1^2 + \beta_0^2 \omega_2^2 \phi_1^2) \{E(X_{t-1}^2 u_{t-1}^2) - E^2(X_{t-1}) E^2(u_{t-1})\} \\
&\quad + \beta_0^2 \{E(u_t^2) E(\epsilon_t^2) - E^2(u_t) E^2(\epsilon_t)\} \\
&= \phi_0^2 \beta_0^2 \sigma_u^2 + (\beta_0^2 \phi_1^2 + \beta_0^2 \omega_2^2 \phi_1^2) \{E(X_{t-1}^2) E(u_{t-1}^2) - E^2(X_{t-1}) E^2(u_{t-1})\} + \beta_0^2 (\sigma_u^2 + 1) \sigma_\epsilon^2 \\
&= \beta_0^2 \{\sigma_u^2 \phi_0^2 + (1 + \sigma_u^2) \sigma_\epsilon^2\} \\
&\quad + \beta_0^2 \phi_1^2 (1 + \omega_2^2) [\{\text{Var}(u_{t-1}) + E^2(u_{t-1})\} \{\text{Var}(X_{t-1}) + E^2(X_{t-1})\} - E^2(X_{t-1})] \\
&= \beta_0^2 \{\sigma_u^2 \phi_0^2 + (1 + \sigma_u^2) \sigma_\epsilon^2\} \\
&\quad + \beta_0^2 \phi_1^2 (1 + \omega_2^2) [\{\text{Var}(u_{t-1}) + 1\} \{\text{Var}(X_{t-1}) + E^2(X_{t-1})\} - E^2(X_{t-1})] \\
&= \beta_0^2 \{\sigma_u^2 \phi_0^2 + (1 + \sigma_u^2) \sigma_\epsilon^2\} \\
&\quad + \beta_0^2 \phi_1^2 (1 + \omega_2^2) \{\text{Var}(u_{t-1}) \text{Var}(X_{t-1}) + \text{Var}(u_{t-1}) E^2(X_{t-1}) + \text{Var}(X_{t-1})\} \\
&= \beta_0^2 \{\sigma_u^2 \phi_0^2 + (1 + \sigma_u^2) \sigma_\epsilon^2\} + \beta_0^2 \phi_1^2 (1 + \omega_2^2) \frac{\text{Var}(X_{t-1})}{\omega_2} \\
&= \beta_0^2 \{\sigma_u^2 \phi_0^2 + (1 + \sigma_u^2) \sigma_\epsilon^2\} + \beta_0^2 \phi_1^2 \frac{1 + \omega_2^2}{\omega_2} \frac{\sigma_\epsilon^2}{1 - \phi_1^2},
\end{aligned}$$

where the second step is because of the independence assumption as well as $E(u_{t-1}^2) = E(u_t^2)$ and $E(u_{t-1}) = E(u_t)$ such that $\text{Var}(X_{t-1} u_t) = \text{Var}(X_{t-1} u_{t-1})$, and the second last step is due to $\omega_2 = \frac{\text{Var}(X_{t-1})}{\text{Var}(u_{t-1}) \text{Var}(X_{t-1}) + \text{Var}(u_{t-1}) E^2(X_{t-1}) + \text{Var}(X_{t-1})}$ in (E.5).

E.4 The proof of Theorem 6.3

Proof of Theorem 6.3(1):

For $k = 1, \dots, p$, applying the weak law of large numbers to $\hat{\gamma}_k^*$, we obtain that as $T \rightarrow \infty$, the estimator $\hat{\gamma}_k^*$ converges in probability to $\text{Cov}(X_t^*, X_{t-k}^*)$, denoted γ_k^* .

Next, we examine γ_k . By the form of measurement error model (6.7), we have that for $0 < k < t$,

$$\begin{aligned}
\text{Cov}(X_t^*, X_{t-k}^*) &= \text{Cov}(\alpha_0 + \alpha_1 X_t + e_t, \alpha_0 + \alpha_1 X_{t-k} + e_{t-k}) \\
&= \alpha_1^2 \text{Cov}(X_t, X_{t-k}) = \alpha_1^2 \gamma_k,
\end{aligned}$$

and by (6.8), $\text{Var}(X_t^*) = \alpha_1^2 \gamma_0 + \sigma_e^2$, which is denoted as γ_0^* .

Thus, Theorem 6.3(1) follows.

Proof of Theorem 6.3(2):

First, by Theorem 6.3(1), we write

$$\widehat{\gamma}^* = \alpha_1^2 \gamma + o_p(1) \quad (\text{E.8})$$

and

$$\widehat{\Gamma}^* = \alpha_1^2 \Gamma + \sigma_e^2 I_p + o_p(1),$$

where $\widehat{\Gamma}^* = \begin{pmatrix} \widehat{\gamma}_0^* & \cdots & \widehat{\gamma}_{p-1}^* \\ \vdots & \ddots & \vdots \\ \widehat{\gamma}_{p-1}^* & \cdots & \widehat{\gamma}_0^* \end{pmatrix}$. Then the naive estimator $\widehat{\phi}^*$ is obtained by replacing $\widehat{\gamma}_k$ in (6.6) with $\widehat{\gamma}_k^*$,

$$\widehat{\phi}^* = \{\alpha_1^2 \Gamma + \sigma_e^2 I_p + o_p(1)\}^{-1} \{\alpha_1^2 \gamma + o_p(1)\} = \alpha_1^2 (\alpha_1^2 \Gamma + \sigma_e^2 I_p)^{-1} \gamma + o_p(1), \quad (\text{E.9})$$

and hence $\phi^* = \alpha_1^2 (\alpha_1^2 \Gamma + \sigma_e^2 I_p)^{-1} \gamma$ such that $\widehat{\phi}^* \xrightarrow{p} \phi^*$ as $T \rightarrow \infty$.

Again, replacing $\widehat{\gamma}_k$ in (6.6) with $\widehat{\gamma}_k^*$ gives the naive estimator $\widehat{\phi}_0^*$

$$\begin{aligned} \widehat{\phi}_0^* &= \frac{1}{T-p} \sum_{t=p}^T X_t^* - \left(\sum_{k=1}^p \widehat{\phi}_k^* \right) \left(\frac{1}{T-p} \sum_{t=p}^T X_{t-k}^* \right) \\ &= E(X_t^*) - E(X_t^*) \sum_{k=1}^p \widehat{\phi}_k^* + o_p(1) \\ &= \alpha_0 + \alpha_1 E(X_t) - \{\alpha_0 + \alpha_1 E(X_t)\} \sum_{k=1}^p \{\phi_k^* + o_p(1)\} + o_p(1) \\ &= (1 - \phi^{*\text{T}} \cdot \mathbf{1}_p) (\alpha_0 + \alpha_1 \mu) + o_p(1), \end{aligned}$$

where $\widehat{\phi}_k$ and ϕ_k are respectively the k th element of $\widehat{\phi}$ and ϕ , the third step is because $\widehat{\phi}_k = \phi_k + o_p(1)$ by (E.9) as well as the model form (6.7), and the last step is due to the stationarity of the time series $\{X_t\}$ such that $E(X_t) = \mu$.

Finally, noting that the native estimator $\hat{\sigma}_\epsilon^{2*}$ is given by $\hat{\sigma}_\epsilon^{2*} = \hat{\gamma}_0^* - 2\hat{\phi}^{*\top}\hat{\gamma}^* + \hat{\phi}^{*\top}\hat{\Gamma}^*\hat{\phi}^*$ by applying a version similar to (6.6), we obtain that

$$\begin{aligned}\hat{\sigma}_\epsilon^{2*} &= \hat{\gamma}_0^* - 2\hat{\phi}^{*\top}\hat{\gamma}^* + \hat{\phi}^{*\top}\hat{\Gamma}^*\hat{\phi}^* \\ &= (\alpha_1^2\gamma_0^2 + \sigma_e^2) - 2\alpha_1^4\gamma^\top(\alpha_1^2\Gamma + \sigma_e^2I_p)^{-1}\gamma \\ &\quad + \alpha_1^4\gamma^\top(\alpha_1^2\Gamma + \sigma_e^2I_p)^{-1}(\alpha_1^2\Gamma + \sigma_e^2I_p)(\alpha_1^2\Gamma + \sigma_e^2I_p)^{-1}\gamma + o_p(1) \\ &= \alpha_1^2\gamma_0 + \sigma_e^2 - \alpha_1^4\gamma^\top(\alpha_1^2\Gamma + \sigma_e^2I_p)^{-1}\gamma + o_p(1),\end{aligned}$$

where the second step is due to (6.8), (E.8) and (E.9).

Proof of Theorem 6.3(3):

Step 1: We show certain identities before proving Theorem 6.3(3).

1. By model (6.7), we have that

$$\begin{aligned}X_t^* - \hat{\mu}^* &= \alpha_0 + \alpha_1 X_t + e_t - \frac{1}{T} \sum_{t=1}^T (\alpha_0 + \alpha_1 X_t + e_t) \\ &= \alpha_1 \left(X_t - \frac{1}{T} \sum_{t=1}^T X_t \right) + \left(e_t - \frac{1}{T} \sum_{t=1}^T e_t \right) \\ &= \alpha_1 (X_t - \hat{\mu}) + (e_t - \bar{e}),\end{aligned}\tag{E.10}$$

where the first step is because $\hat{\mu}^* = \frac{1}{T} \sum_{t=1}^T X_t^*$ and in the last step $\bar{e} = \frac{1}{T} \sum_{t=1}^T e_t$.

2. For any t and s , we have that

$$\begin{aligned}&\text{Cov} \{ (X_t - \hat{\mu})^2, (X_s - \hat{\mu})(e_s - \bar{e}) \} \\ &= E \{ (X_t - \hat{\mu})^2 (X_s - \hat{\mu})(e_s - \bar{e}) \} - \{ E(X_t - \hat{\mu})^2 \} E \{ (X_s - \hat{\mu})(e_s - \bar{e}) \} \\ &= E \{ (X_t - \hat{\mu})^2 (X_s - \hat{\mu}) \} E(e_s - \bar{e}) - \{ E(X_t - \hat{\mu})^2 \} E(X_s - \hat{\mu}) E(e_s - \bar{e}) \\ &= 0,\end{aligned}\tag{E.11}$$

where the second step is due to the independence of e_t and X_t , and the last step is by $E(e_s - \bar{e}) = 0$.

3. By the independence of e_t and e_s for $t \neq s$, we have that

$$\begin{aligned}
& \text{Cov} \{ (X_t - \hat{\mu})(e_t - \bar{e}), (X_s - \hat{\mu})(e_s - \bar{e}) \} \\
&= E \{ (X_t - \hat{\mu})(e_t - \bar{e})(X_s - \hat{\mu})(e_s - \bar{e}) \} - E \{ (X_t - \hat{\mu})(e_t - \bar{e}) \} E \{ (X_s - \hat{\mu})(e_s - \bar{e}) \} \\
&= E \{ (X_t - \hat{\mu})(X_s - \hat{\mu}) \} E \{ (e_t - \bar{e}) \} E \{ (e_s - \bar{e}) \} \\
&\quad - E \{ (X_t - \hat{\mu}) \} E \{ (e_t - \bar{e}) \} E \{ (X_s - \hat{\mu}) \} E \{ (e_s - \bar{e}) \} \\
&= 0,
\end{aligned} \tag{E.12}$$

where the second step is due to the independence of the e_t and the X_t , and the last step is by $E(e_s - \bar{e}) = 0$.

4. For any t , we have that

$$\begin{aligned}
& \text{Var} \{ (X_t - \hat{\mu})(e_t - \bar{e}) \} \\
&= E \{ (X_t - \hat{\mu})^2 (e_t - \bar{e})^2 \} - E^2 \{ (X_t - \hat{\mu})(e_t - \bar{e}) \} \\
&= E \{ (X_t - \hat{\mu})^2 \} E \{ (e_t - \bar{e})^2 \} - E^2 \{ (X_t - \hat{\mu}) \} E^2 \{ (e_t - \bar{e}) \} \\
&= E \{ (X_t - \hat{\mu})^2 \} E \{ (e_t - \bar{e})^2 \}.
\end{aligned} \tag{E.13}$$

5. For any t , we have

$$\begin{aligned}
& \lim_{T \rightarrow \infty} E \{ (X_t - \hat{\mu})^2 \} \\
&= \lim_{T \rightarrow \infty} E \{ (X_t - \mu)^2 + (\mu - \hat{\mu})^2 + 2(X_t - \mu)(\mu - \hat{\mu}) \} \\
&= \gamma_0 + \lim_{T \rightarrow \infty} E \{ (\hat{\mu} - \mu)^2 \} + 2 \lim_{T \rightarrow \infty} E \{ (X_t - \mu)(\mu - \hat{\mu}) \} \\
&= \gamma_0 + \lim_{T \rightarrow \infty} E \{ (\hat{\mu} - \mu)^2 \} - 2 \lim_{T \rightarrow \infty} E \left[(X_t - \mu) \left\{ \frac{1}{T} \sum_{s=1}^T (X_s - \mu) \right\} \right] \\
&= \gamma_0 + \lim_{T \rightarrow \infty} \text{Var}(\hat{\mu}) - 2 \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{s=1}^T E \{ (X_t - \mu)(X_s - \mu) \} \\
&= \gamma_0 + 0 - 2 \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{s=1}^T \gamma_{|s-t|} \\
&= \gamma_0,
\end{aligned} \tag{E.14}$$

where the third step is due to $\hat{\mu} - \mu = \frac{1}{T} \sum_{s=1}^T (X_s - \mu)$, and the fourth step is because $E(\hat{\mu} - \mu) = 0$ by stationarity of the time series, the second last step is due to $\lim_{T \rightarrow \infty} \text{Var}(\hat{\mu}) = 0$ (Brockwell et al., 1991, Theorem 7.1.1.), and the last step due to Condition (R3).

6. Similar to (E.14), we have that

$$\begin{aligned}
& \lim_{T \rightarrow \infty} E\{(X_t - \hat{\mu})(X_{t-p} - \hat{\mu})\} \\
&= \lim_{T \rightarrow \infty} E\{(X_t - \mu + \mu - \hat{\mu})(X_{t-p} - \mu + \mu - \hat{\mu})\} \\
&= \lim_{T \rightarrow \infty} [E\{(X_t - \mu)(X_{t-p} - \mu)\} + E\{(\mu - \hat{\mu})(X_{t-p} - \mu)\} \\
&\quad + E\{(\mu - \hat{\mu})(X_t - \mu)\} + E\{(\mu - \hat{\mu})(\mu - \hat{\mu})\}] \tag{E.15}
\end{aligned}$$

$$\begin{aligned}
&= \gamma_p + \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{s=1}^T (\gamma_{|t-s|} + \gamma_{|t-s-p|}) + \lim_{T \rightarrow \infty} \text{Var}(\hat{\mu}) \\
&= \gamma_p, \tag{E.16}
\end{aligned}$$

where the last step is due to Condition (R3) and $\lim_{T \rightarrow \infty} \text{Var}(\hat{\mu}) = 0$ (Brockwell et al., 1991, Theorem 7.1.1).

7. For any t , we have

$$\begin{aligned}
& E\{(e_t - \bar{e})^2\} \\
&= E\{e_t^2 - 2e_t\bar{e} + \bar{e}^2\} \\
&= \left\{ E(e_t^2) - \frac{2}{T} \sum_{s=1}^T E(e_t e_s) + \frac{1}{T^2} \sum_{t=1}^T \sum_{s=1}^T E(e_t e_s) \right\} \\
&= E(e_t^2) + \left\{ -\frac{2}{T} E(e_t e_t) + \frac{1}{T^2} \sum_{t=1}^T E(e_t^2) \right\} \\
&= \frac{T-1}{T} E(e_t^2) = \frac{T-1}{T} \sigma_e^2, \tag{E.17}
\end{aligned}$$

so $\lim_{T \rightarrow \infty} E\{(e_t - \bar{e})^2\} = \sigma_e^2$.

8. By the independence of e_t and X_t , for any s and t , we have that

$$\begin{aligned}
& \text{Cov}\{(X_t - \hat{\mu})(e_t - \bar{e}), (e_s - \bar{e})^2\} \\
&= E\{(X_t - \hat{\mu})(e_t - \bar{e})(e_s - \bar{e})^2\} - E\{(X_t - \hat{\mu})(e_t - \bar{e})\} E\{(e_s - \bar{e})^2\} \\
&= E(X_t - \hat{\mu}) E\{(e_t - \bar{e})(e_s - \bar{e})^2\} - E(X_t - \hat{\mu}) E(e_t - \bar{e}) E(e_s - \bar{e})^2 \\
&= 0, \tag{E.18}
\end{aligned}$$

where the last step is due to $E(X_t - \hat{\mu}) = 0$ and $E(e_t - \bar{e}) = 0$.

9. For any $t \neq s$, $\text{Cov}\{(e_t - \bar{e})^2, (e_s - \bar{e})^2\} = 0$; and for $t = s$,

$$\begin{aligned}
& \text{Var}\{(e_t - \bar{e})^2\} \\
&= E\{(e_t - \bar{e})^4\} - E^2\{(e_t - \bar{e})^2\} \\
&= E(e_t^4) - 4E(e_t^3\bar{e}) + 6E(e_t^2\bar{e}^2) - 4E(e_t\bar{e}^3) + E(\bar{e}^4) - \{E(e_t^2) - 2E(e_t\bar{e}) + E(\bar{e}^2)\}^2 \\
&= E(e_t^4) - \frac{4}{T}E(e_t^4) + \left[\frac{6(T-1)}{T^2}\{E(e_t^2)\}^2 + \frac{6}{T^2}E(e_t^4) \right] - \frac{4}{T^3}E(e_t^4) \\
&\quad + \left[\frac{1}{T^3}E(e_t^4) + \frac{3(T-1)}{T^3}\{E(e_t^2)\}^2 \right] - \left\{ E(e_t^2) - \frac{2}{T}E(e_t^2) + \frac{1}{T}E(e_t^2) \right\}^2, \quad (\text{E.19})
\end{aligned}$$

so $\lim_{T \rightarrow \infty} \text{Var}\{(e_t - \bar{e})^2\} = E(e_t^4) - \{E(e_t^2)\}^2 = E(e_t^4) - \sigma_e^4$.

10. Similar to the derivation in (E.19), we can show $\text{Cov}\{(e_t - \bar{e})^2, (e_s - \bar{e})(e_{s+p} - \bar{e})\} = 0$ for $s \neq t$ and $s \neq t - p$. For a given t ,

$$\begin{aligned}
& \text{Cov}\{(e_t - \bar{e})^2, (e_t - \bar{e})(e_{t+p} - \bar{e})\} \\
&= E\{(e_t - \bar{e})^3(e_{t+p} - \bar{e})\} - E\{(e_t - \bar{e})^2\}E\{(e_t - \bar{e})(e_{t+p} - \bar{e})\}, \quad (\text{E.20})
\end{aligned}$$

which can be derived analogously to the (E.19) that $\lim_{T \rightarrow \infty} E\{(e_t - \bar{e})^3(e_{t+p} - \bar{e})\} - E\{(e_t - \bar{e})^2\}E\{(e_t - \bar{e})(e_{t+p} - \bar{e})\} = E\{e_t^3 e_{t+p}\} - E\{e_t^2\}E\{e_t e_{t+p}\} = 0$ and similarly $\lim_{T \rightarrow \infty} \text{Cov}\{(e_t - \bar{e})^2, (e_{t-p} - \bar{e})(e_t - \bar{e})\} = 0$.

11. For any t ,

$$\begin{aligned}
& \text{Cov}\{(X_t - \hat{\mu})(e_{t+p} - \bar{e}), (X_{t+p-r} - \hat{\mu})(e_{t+p} - \bar{e})\} \\
&= [E\{(X_t - \hat{\mu})(X_{t+p-r} - \hat{\mu})(e_{t+p} - \bar{e})^2\} - E(X_t - \hat{\mu})E(X_{t+p-r} - \hat{\mu})E^2(e_{t+p} - \bar{e})] \\
&= E\{(X_t - \hat{\mu})(X_{t+p-r} - \hat{\mu})\}E\{(e_{t+p} - \bar{e})^2\} \\
&= \gamma_{|p-r|} \left(\frac{T-1}{T} \right) \sigma_e^2, \quad (\text{E.21})
\end{aligned}$$

where the second step is because of $E(X_t - \hat{\mu}) = 0$ and the independence of X_t and e_t , the third step is due to (E.17) and (E.15). Hence,

$$\lim_{T \rightarrow \infty} \text{Cov}\{(X_t - \hat{\mu})(e_{t+p} - \bar{e}), (X_{t+p-r} - \hat{\mu})(e_{t+p} - \bar{e})\} = \gamma_{|p-r|} \sigma_e^2.$$

Similarly,

$$\lim_{T \rightarrow \infty} \text{Cov}\{(X_{t+p} - \hat{\mu})(e_t - \bar{e}), (X_{t-r} - \hat{\mu})(e_t - \bar{e})\} = \gamma_{|p-r|} \sigma_e^2.$$

Similarly,

$$\begin{aligned}
& \text{Cov} \{ (X_t - \hat{\mu})(e_{t+p} - \bar{e}), (X_{t+p+r} - \hat{\mu})(e_{t+p} - \bar{e}) \} \\
&= [E \{ (X_t - \hat{\mu})(X_{t+p+r} - \hat{\mu})(e_{t+p} - \bar{e})^2 \} \\
&\quad - E(X_t - \hat{\mu})E(X_{t+p+r} - \hat{\mu})E^2(e_{t+p} - \bar{e})] \\
&= E \{ (X_t - \hat{\mu})(X_{t+p+r} - \hat{\mu}) \} E \{ (e_t - \bar{e})^2 \} \\
&= \gamma_{p+r} \left(\frac{T-1}{T} \right) \sigma_e^2, \tag{E.22}
\end{aligned}$$

and hence

$$\lim_{T \rightarrow \infty} \text{Cov} \{ (X_t - \hat{\mu})(e_{t+p} - \bar{e}), (X_{t+p+r} - \hat{\mu})(e_{t+p} - \bar{e}) \} = \gamma_{p+r} \sigma_e^2.$$

Similarly,

$$\lim_{T \rightarrow \infty} \text{Cov} \{ (X_{t+p} - \hat{\mu})(e_t - \bar{e}), (X_{t+r} - \hat{\mu})(e_t - \bar{e}) \} = \gamma_{p+r} \sigma_e^2.$$

12. By independence assumption between $\{e_t\}$, if $t \neq s$ or $p \neq r$, we have that

$$\text{Cov} \{ (e_t - \bar{e})(e_{t+p} - \bar{e}), (e_s - \bar{e})(e_{s+r} - \bar{e}) \} = 0. \tag{E.23}$$

In addition, by (E.17), we have that

$$\begin{aligned}
& \text{Var} \{ (e_t - \bar{e})(e_{t+p} - \bar{e}) \} \\
&= E \{ (e_t - \bar{e})^2 (e_{t+p} - \bar{e})^2 \} \\
&= E \{ (e_t - \bar{e})^2 \} E \{ (e_{t+p} - \bar{e})^2 \}, \\
&= \left(\frac{T-1}{T} \right)^2 \sigma_e^4, \tag{E.24}
\end{aligned}$$

so $\lim_{T \rightarrow \infty} \text{Var} \{ (e_t - \bar{e})(e_{t+p} - \bar{e}) \} = \sigma_e^4$.

Step 2: Now we prove the results in (3).

1°. We first show the derivation of q_{100}^* as follows:

$$\begin{aligned}
q_{100}^* &= \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T} \sum_{t=1}^T (X_t^* - \hat{\mu}^*)^2, \frac{1}{T} \sum_{s=1}^T (X_s^* - \hat{\mu}^*)^2 \right\} \\
&= \lim_{T \rightarrow \infty} TCov \left[\frac{1}{T} \sum_{t=1}^T \{ \alpha_1^2 (X_t - \hat{\mu})^2 + 2\alpha_1 (X_t - \hat{\mu})(e_t - \bar{e}) + (e_t - \bar{e})^2 \}, \right. \\
&\quad \left. \frac{1}{T} \sum_{s=1}^T \alpha_1^2 (X_s - \hat{\mu})^2 + 2\alpha_1 (X_s - \hat{\mu})(e_s - \bar{e}) + (e_s - \bar{e})^2 \right] \\
&= \alpha_1^4 \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T} \sum_{t=1}^T (X_t - \hat{\mu})^2, \frac{1}{T} \sum_{s=1}^T (X_s - \hat{\mu})^2 \right\} \\
&\quad + \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T} \sum_{t=1}^T 2\alpha_1 (X_t - \hat{\mu})(e_t - \bar{e}), \frac{1}{T} \sum_{s=1}^T 2\alpha_1 (X_s - \hat{\mu})(e_s - \bar{e}) \right\} \\
&\quad + \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T} \sum_{t=1}^T (e_t - \bar{e})^2, \frac{1}{T} \sum_{s=1}^T (e_s - \bar{e})^2 \right\} \\
&= \alpha_1^4 q_{00} + \lim_{T \rightarrow \infty} \frac{4\alpha_1^2}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ (X_t - \hat{\mu})(e_t - \bar{e}), (X_s - \hat{\mu})(e_s - \bar{e}) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ (e_t - \bar{e})^2, (e_s - \bar{e})^2 \} \\
&= \alpha_1^4 q_{00} + \lim_{T \rightarrow \infty} \frac{4\alpha_1^2}{T} \sum_{t=1}^T Cov \{ (X_t - \hat{\mu})(e_t - \bar{e}), (X_t - \hat{\mu})(e_t - \bar{e}) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T Cov \{ (e_t - \bar{e})^2, (e_t - \bar{e})^2 \} \\
&= \alpha_1^4 q_{00} + 4\alpha_1^2 E \{ (X_t - \hat{\mu})^2 (e_t - \bar{e})^2 \} + E(e_t^4) - \{ E(e_t^2) \}^2 \\
&= \alpha_1^4 q_{00} + 4\alpha_1^2 \gamma_0 \sigma_e^2 + E(e_t^4) - \sigma_e^4,
\end{aligned}$$

where the second step is due to (E.10), the third step is because of (E.11), (E.18), and the definition $q_{00} = \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T} \sum_{t=1}^T (X_t - \hat{\mu})^2, \frac{1}{T} \sum_{s=1}^T (X_s - \hat{\mu})^2 \right\}$, the fifth step is due to (E.12) and (E.19), and the sixth step is because (E.13) and (E.19), and the last step is because (E.17) and (E.18).

2°. We derive the value of q_{10p}^* :

$$\begin{aligned}
q_{10p}^* &= \lim_{T \rightarrow \infty} \text{TCov} \left\{ \frac{1}{T} \sum_{t=1}^T (X_t^* - \hat{\mu}^*)^2, \frac{1}{T-p} \sum_{s=1}^{T-p} (X_s^* - \hat{\mu}^*)(X_{s+p}^* - \hat{\mu}^*) \right\} \\
&= \lim_{T \rightarrow \infty} \text{TCov} \left[\frac{1}{T} \sum_{t=1}^T \{ \alpha_1^2 (X_t - \hat{\mu})^2 + 2\alpha_1 (X_t - \hat{\mu})(e_t - \bar{e}) + (e_t - \bar{e})^2 \}, \right. \\
&\quad \left. \frac{1}{T-p} \sum_{s=1}^{T-p} \alpha_1^2 (X_s - \hat{\mu})(X_{s+p} - \hat{\mu}) + \alpha_1 (X_s - \hat{\mu})(e_{s+p} - \bar{e}) \right. \\
&\quad \left. + \alpha_1 (X_{s+p} - \hat{\mu})(e_s - \bar{e}) + (e_s - \bar{e})(e_{s+p} - \bar{e}) \right] \\
&= \alpha_1^4 \lim_{T \rightarrow \infty} \text{TCov} \left\{ \frac{1}{T} \sum_{t=1}^T (X_t - \hat{\mu})^2, \frac{1}{T-p} \sum_{s=1}^{T-p} (X_s - \hat{\mu})(X_{s+p} - \hat{\mu}) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \text{TCov} \left\{ \frac{1}{T} \sum_{t=1}^T 2\alpha_1 (X_t - \hat{\mu})(e_t - \bar{e}), \frac{1}{T-p} \sum_{s=1}^{T-p} \alpha_1 (X_s - \hat{\mu})(e_{s+p} - \bar{e}) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \text{TCov} \left\{ \frac{1}{T} \sum_{t=1}^T 2\alpha_1 (X_t - \hat{\mu})(e_t - \bar{e}), \frac{1}{T-p} \sum_{s=1}^{T-p} \alpha_1 (X_{s+p} - \hat{\mu})(e_s - \bar{e}) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \text{TCov} \left\{ \frac{1}{T} \sum_{t=1}^T (e_t - \bar{e})^2, \frac{1}{T-p} \sum_{s=1}^{T-p} (e_s - \bar{e})(e_{s+p} - \bar{e}) \right\} \\
&= \alpha_1^4 q_{0p} + \lim_{T \rightarrow \infty} \frac{2\alpha_1^2}{T-p} \sum_{t=1}^T \sum_{s=1}^{T-p} \text{Cov} \{ (X_t - \hat{\mu})(e_t - \bar{e}), (X_s - \hat{\mu})(e_{s+p} - \bar{e}_s) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{2\alpha_1^2}{T-p} \sum_{t=1}^T \sum_{s=1}^{T-p} \text{Cov} \{ (X_t - \hat{\mu})(e_t - \bar{e}), (X_{s+p} - \hat{\mu})(e_s - \bar{e}) \} \\
&= \alpha_1^4 q_{0p} + \lim_{T \rightarrow \infty} \frac{2\alpha_1^2}{T-p} \sum_{\substack{t=p \\ (s=t-p)}}^T \text{Cov} \{ (X_t - \hat{\mu})(e_t - \bar{e}), (X_{t-p} - \hat{\mu})(e_t - \bar{e}) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{2\alpha_1^2}{T-p} \sum_{\substack{t=1 \\ (s=t)}}^{T-p} \text{Cov} \{ (X_t - \hat{\mu})(e_t - \bar{e}), (X_{t+p} - \hat{\mu})(e_t - \bar{e}) \} \\
&= \alpha_1^4 q_{0p} + 2\alpha_1^2 E \{ (X_t - \hat{\mu})(X_{t-p} - \hat{\mu})(e_t - \bar{e})^2 \} + 2\alpha_1^2 E \{ (X_t - \hat{\mu})(X_{t+p} - \hat{\mu})(e_t - \bar{e})^2 \} \\
&= \alpha_1^4 q_{0p} + 4\alpha_1^2 \gamma_p \sigma_e^2,
\end{aligned}$$

where the second step is due to (E.10), the third step is because of (E.11) and (E.18), the fourth step is by definition that $q_{0p} = \lim_{T \rightarrow \infty} \text{TCov} \left\{ \frac{1}{T} \sum_{t=1}^T (X_t - \hat{\mu})^2, \frac{1}{T-p} \sum_{s=1}^{T-p} (X_s - \hat{\mu})(X_{s+p} - \hat{\mu}) \right\}$ and (E.20), the fifth step is due to (E.12), and the last step is result from (E.17) and (E.15).

3°. We derive q_{1pr}^* for $p > 0$, $r > 0$ and $p \neq r$:

$$\begin{aligned}
q_{1pr}^* &= \lim_{T \rightarrow \infty} \text{TCov} \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} (X_t^* - \hat{\mu}^*)(X_{t+p}^* - \hat{\mu}^*), \frac{1}{T-r} \sum_{s=1}^{T-r} (X_s^* - \hat{\mu}^*)(X_{s+r}^* - \hat{\mu}^*) \right\} \\
&= \lim_{T \rightarrow \infty} \text{TCov} \left[\frac{1}{T-p} \sum_{t=1}^{T-p} \left\{ \alpha_1^2 (X_t - \hat{\mu})(X_{t+p} - \hat{\mu}) + \alpha_1 (X_t - \hat{\mu})(e_{t+p} - \bar{e}) \right. \right. \\
&\quad \left. \left. + \alpha_1 (X_{t+p} - \hat{\mu})(e_t - \bar{e}) + (e_t - \bar{e})(e_{t+p} - \bar{e}) \right\}, \right. \\
&\quad \left. \frac{1}{T-r} \sum_{s=1}^{T-r} \left\{ \alpha_1^2 (X_s - \hat{\mu})(X_{s+r} - \hat{\mu}) + \alpha_1 (X_s - \hat{\mu})(e_{s+r} - \bar{e}) \right. \right. \\
&\quad \left. \left. + \alpha_1 (X_{s+r} - \hat{\mu})(e_s - \bar{e}) + (e_s - \bar{e})(e_{s+r} - \bar{e}) \right\} \right] \\
&= \alpha_1^4 \lim_{T \rightarrow \infty} \text{TCov} \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} (X_t - \hat{\mu})(X_{t+p} - \hat{\mu}), \frac{1}{T-r} \sum_{s=1}^{T-r} (X_s - \hat{\mu})(X_{s+r} - \hat{\mu}) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \text{TCov} \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} \alpha_1 (X_t - \hat{\mu})(e_{t+p} - \bar{e}), \frac{1}{T-r} \sum_{s=1}^{T-r} \alpha_1 (X_s - \hat{\mu})(e_{s+r} - \bar{e}) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \text{TCov} \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} \alpha_1 (X_t - \hat{\mu})(e_{t+p} - \bar{e}), \frac{1}{T-r} \sum_{s=1}^{T-r} \alpha_1 (X_{s+r} - \hat{\mu})(e_s - \bar{e}) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \text{TCov} \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} \alpha_1 (X_{t+p} - \hat{\mu})(e_t - \bar{e}), \frac{1}{T-r} \sum_{s=1}^{T-r} \alpha_1 (X_s - \hat{\mu})(e_{s+r} - \bar{e}) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \text{TCov} \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} \alpha_1 (X_{t+p} - \hat{\mu})(e_t - \bar{e}), \frac{1}{T-r} \sum_{s=1}^{T-r} \alpha_1 (X_{s+r} - \hat{\mu})(e_s - \bar{e}) \right\} \\
&= \alpha_1^4 q_{pr} + \alpha_1^2 \lim_{T \rightarrow \infty} \frac{T}{(T-p)(T-r)} \sum_{\substack{t=\max(1, r-p+1) \\ (s=t+p-r)}}^{T-p} \text{Cov} \{ (X_t - \hat{\mu})(e_{t+p} - \bar{e}), (X_{t+p-r} - \hat{\mu})(e_{t+p} - \bar{e}) \} \\
&\quad + \alpha_1^2 \lim_{T \rightarrow \infty} \frac{T}{(T-p)(T-r)} \sum_{\substack{t=1 \\ (s=t+p)}}^{T-p-r} \text{Cov} \{ (X_t - \hat{\mu})(e_{t+p} - \bar{e}), (X_{t+p+r} - \hat{\mu})(e_{t+p} - \bar{e}) \} \\
&\quad + \alpha_1^2 \lim_{T \rightarrow \infty} \frac{T}{(T-p)(T-r)} \sum_{\substack{t=r+1 \\ (s=t-r)}}^{T-p} \text{Cov} \{ (X_{t+p} - \hat{\mu})(e_t - \bar{e}), (X_{t-r} - \hat{\mu})(e_t - \bar{e}) \} \\
&\quad + \alpha_1^2 \lim_{T \rightarrow \infty} \frac{T}{(T-p)(T-r)} \sum_{\substack{t=1 \\ (s=t)}}^{T-\max(p, r)} \text{Cov} \{ (X_{t+p} - \hat{\mu})(e_t - \bar{e}), (X_{t+r} - \hat{\mu})(e_t - \bar{e}) \} \\
&= \alpha_1^4 q_{pr} + 2\alpha_1^2 \sigma_e^2 (\gamma_{|p-r|} + \gamma_{p+r}). \tag{E.25}
\end{aligned}$$

where the second step is due to (E.10), the third step is because of (E.11) and a similar version to (E.18), the fourth step is because (E.23) and by the definition that

$q_{pr} = \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} (X_t - \hat{\mu})(X_{t+p} - \hat{\mu}), \frac{1}{T-r} \sum_{s=1}^{T-r} (X_s - \hat{\mu})(X_{s+r} - \hat{\mu}) \right\}$, and the last step is from (E.21) and (E.22).

4°. Finally, we present the derivation of q_{1pp}^* for $p \neq 0$,

$$\begin{aligned}
q_{1pp}^* &= \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} (X_t^* - \hat{\mu}^*)(X_{t+p}^* - \hat{\mu}^*), \frac{1}{T-p} \sum_{s=1}^{T-p} (X_s^* - \hat{\mu}^*)(X_{s+p}^* - \hat{\mu}^*) \right\} \\
&= \lim_{T \rightarrow \infty} TCov \left[\frac{1}{T-p} \sum_{t=1}^{T-p} \left\{ \alpha_1^2 (X_t - \hat{\mu})(X_{t+p} - \hat{\mu}) + \alpha_1 (X_t - \hat{\mu})(e_{t+p} - \bar{e}) \right. \right. \\
&\quad \left. \left. + \alpha_1 (X_{t+p} - \hat{\mu})(e_t - \bar{e}) + (e_t - \bar{e})(e_{t+p} - \bar{e}) \right\}, \right. \\
&\quad \left. \frac{1}{T-p} \sum_{s=1}^{T-p} \left\{ \alpha_1^2 (X_s - \hat{\mu})(X_{s+p} - \hat{\mu}) + \alpha_1 (X_s - \hat{\mu})(e_{s+p} - \bar{e}) \right. \right. \\
&\quad \left. \left. + \alpha_1 (X_{s+p} - \hat{\mu})(e_s - \bar{e}) + (e_s - \bar{e})(e_{s+p} - \bar{e}) \right\} \right] \\
&= \alpha_1^4 \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} (X_t - \hat{\mu})(X_{t+p} - \hat{\mu}), \frac{1}{T-p} \sum_{s=1}^{T-p} (X_s - \hat{\mu})(X_{s+p} - \hat{\mu}) \right\} \\
&\quad + \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} \alpha_1 (X_t - \hat{\mu})(e_{t+p} - \bar{e}), \frac{1}{T-p} \sum_{s=1}^{T-p} \alpha_1 (X_s - \hat{\mu})(e_{s+p} - \bar{e}) \right\} \\
&\quad + \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} \alpha_1 (X_t - \hat{\mu})(e_{t+p} - \bar{e}), \frac{1}{T-p} \sum_{s=1}^{T-p} \alpha_1 (X_{s+p} - \hat{\mu})(e_s - \bar{e}) \right\} \\
&\quad + \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} \alpha_1 (X_{t+p} - \hat{\mu})(e_t - \bar{e}), \frac{1}{T-p} \sum_{s=1}^{T-p} \alpha_1 (X_s - \hat{\mu})(e_{s+p} - \bar{e}) \right\} \\
&\quad + \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} \alpha_1 (X_{t+p} - \hat{\mu})(e_t - \bar{e}), \frac{1}{T-p} \sum_{s=1}^{T-p} \alpha_1 (X_{s+p} - \hat{\mu})(e_s - \bar{e}) \right\} \\
&\quad + \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} (e_t - \bar{e})(e_{t+p} - \bar{e}), \frac{1}{T-p} \sum_{s=1}^{T-p} (e_s - \bar{e})(e_{s+p} - \bar{e}) \right\},
\end{aligned}$$

where the second step is due to (E.10), the third step is because of (E.11) and a similar version to (E.18),

Because (E.23) and by the definition that

$$q_{pp} = \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T-p} \sum_{t=1}^{T-p} (X_t - \hat{\mu})(X_{t+p} - \hat{\mu}), \frac{1}{T-p} \sum_{s=1}^{T-p} (X_s - \hat{\mu})(X_{s+p} - \hat{\mu}) \right\},$$

we have that

$$\begin{aligned}
q_{1pp}^* &= \alpha_1^4 q_{pp} + \alpha_1^2 \lim_{T \rightarrow \infty} \frac{T}{(T-p)^2} \sum_{\substack{t=1 \\ s=t}}^{T-p} \text{Cov} \{ (X_t - \hat{\mu})(e_{t+p} - \bar{e}), (X_t - \hat{\mu})(e_{t+p} - \bar{e}) \} \\
&+ \alpha_1^2 \lim_{T \rightarrow \infty} \frac{T}{(T-p)^2} \sum_{\substack{t=1 \\ s=t+p}}^{T-2p} \text{Cov} \{ (X_t - \hat{\mu})(e_{t+p} - \bar{e}), (X_{t+2p} - \hat{\mu})(e_{t+p} - \bar{e}) \} \\
&+ \alpha_1^2 \lim_{T \rightarrow \infty} \frac{T}{(T-p)^2} \sum_{\substack{t=1+p \\ s=t-p}}^{T-p} \text{Cov} \{ (X_{t+p} - \hat{\mu})(e_t - \bar{e}), (X_{t-p} - \hat{\mu})(e_t - \bar{e}) \} \\
&+ \alpha_1^2 \lim_{T \rightarrow \infty} \frac{T}{(T-p)^2} \sum_{\substack{t=1 \\ s=t}}^T \text{Cov} \{ (X_{t+p} - \hat{\mu})(e_t - \bar{e}), (X_{t+p} - \hat{\mu})(e_t - \bar{e}) \} \\
&+ \alpha_1^2 \lim_{T \rightarrow \infty} \frac{T}{(T-p)^2} \text{Var} \{ (e_t - \bar{e})(e_{t+p} - \bar{e}) \} \\
&= \alpha_1^4 q_{pp} + 2\alpha_1^2 \sigma_e^2 (\gamma_0 + \gamma_{2p}) + \sigma_e^4, \tag{E.26}
\end{aligned}$$

where the last step is because of (E.24), and (E.21) and (E.22) with $q = p$.

E.5 The proof of Theorem 6.4

Proof of Theorem 6.4(1):

For $k = 1, \dots, p$, applying the weak law of large numbers to $\hat{\gamma}_k^*$, we obtain that as $T \rightarrow \infty$, the estimator $\hat{\gamma}_k^*$ converges in probability to $\text{Cov}(X_t^*, X_{t-k}^*)$, which is denoted as γ_k^* .

Next, we examine γ_k . By the form of measurement error model (6.9), we have that for $0 < k < t$,

$$\begin{aligned}
&\text{Cov}(X_t^*, X_{t-k}^*) \\
&= \text{Cov}(\beta_0 X_t u_t, \beta_0 X_{t-k} u_{t-k}) \\
&= \beta_0^2 \{ E(X_t u_t X_{t-k} u_{t-k}) - E(X_t u_t) E(X_{t-k} u_{t-k}) \} \\
&= \beta_0^2 \{ E(u_t) E(u_{t-k}) \text{Cov}(X_t, X_{t-k}) \} \\
&= \beta_0^2 \{ \text{Cov}(X_t, X_{t-k}) \} = \beta_0^2 \gamma_k,
\end{aligned}$$

and by (6.10), $\text{Var}(X_t^*) = \beta_0^2 \{(\sigma_u^2 + 1)\gamma_0 + \sigma_u^2\mu^2\}$, which is denoted as γ_0^* . Thus, Theorem 6.4(1) follows.

Proof of Theorem 6.4(2):

First, by Theorem 6.4(1), we write

$$\widehat{\gamma}^* = \beta_0^2 \gamma + o_p(1)$$

and

$$\begin{aligned} \widehat{\Gamma}^* &= \begin{pmatrix} \beta_0^2(\sigma_u^2 + 1)\gamma_0 + \beta_0\sigma_u^2\mu^2 & \beta_0^2\gamma_1 & \cdots & \beta_0^2\gamma_{p-1} \\ \vdots & \ddots & & \vdots \\ \beta_0^2\gamma_{p-1} & \beta_0^2\gamma_{p-2} & \cdots & \beta_0^2(\sigma_u^2 + 1)\gamma_0 + \beta_0\sigma_u^2\mu^2 \end{pmatrix} + o_p(1) \\ &= \beta_0^2 \{ \Gamma + \sigma_u^2(\gamma_0 + \mu^2)I_p \} + o_p(1), \end{aligned}$$

where $\widehat{\Gamma}^* = \begin{pmatrix} \widehat{\gamma}_0^* & \cdots & \widehat{\gamma}_{p-1}^* \\ \vdots & \ddots & \vdots \\ \widehat{\gamma}_{p-1}^* & \cdots & \widehat{\gamma}_0^* \end{pmatrix}$. Then the naive estimator $\widehat{\phi}^*$ is obtained by replacing $\widehat{\gamma}_k$ in (6.6) with $\widehat{\gamma}_k^*$,

$$\widehat{\phi}^* = [\beta_0^2 \{ \Gamma + \sigma_u^2(\gamma_0 + \mu^2)I_p \} + o_p(1)]^{-1} \{ \beta_0^2 \gamma + o_p(1) \} = \{ \Gamma + \sigma_u^2(\gamma_0 + \mu^2)I_p \}^{-1} \gamma + o_p(1), \quad (\text{E.27})$$

and hence $\phi^* = \{ \Gamma + \sigma_u^2(\gamma_0 + \mu^2)I_p \}^{-1} \gamma$ such that $\widehat{\phi}^* \xrightarrow{p} \phi^*$ as $T \rightarrow \infty$.

Again, by replacing $\widehat{\gamma}_k$ in (6.6) with $\widehat{\gamma}_k^*$ gives the naive estimator $\widehat{\phi}_0^*$

$$\begin{aligned} \widehat{\phi}_0^* &= \frac{1}{T-p} \sum_{t=p}^T X_t^* - \left(\sum_{k=1}^p \widehat{\phi}_k^* \right) \left(\frac{1}{T-p} \sum_{t=p}^T X_{t-k}^* \right) \\ &= E(X_t^*) - E(X_t^*) \sum_{k=1}^p \widehat{\phi}_k^* + o_p(1) \\ &= \beta_0 E(X_t) - \beta_0 E(X_t) \sum_{k=1}^p \{ \phi_k^* + o_p(1) \} + o_p(1), \\ &= \beta_0 (1 - \phi^{*\top} \cdot \mathbf{1}_p) \mu + o_p(1), \end{aligned}$$

where $\widehat{\phi}_k$ and ϕ_k are respectively the k th element of $\widehat{\phi}$ and ϕ , the third step is because $\widehat{\phi}_k = \phi_k + o_p(1)$ by (E.27) as well as the model form (6.9), and the last step is due to the stationarity of the time series $\{X_t\}$ such that $E(X_t) = \mu$.

Finally, noting that the native estimator $\widehat{\sigma}_\epsilon^{*2}$ is given by $\widehat{\sigma}_\epsilon^{*2} = \widehat{\gamma}_0^* - 2\widehat{\phi}^{*\top}\widehat{\gamma}^* + \widehat{\phi}^{*\top}\widehat{\Gamma}^*\widehat{\phi}^*$ by applying a version similar to (6.6), we obtain that

$$\begin{aligned}\widehat{\sigma}_\epsilon^{*2} &= \widehat{\gamma}_0^* - 2\widehat{\phi}^{*\top}\widehat{\gamma}^* + \widehat{\phi}^{*\top}\widehat{\Gamma}^*\widehat{\phi}^* \\ &= \beta_0^2 \{(\sigma_u^2 + 1)\gamma_0 + \sigma_u^2\mu^2\} - 2\beta_0^2\gamma^\top \{\Gamma + \sigma_u^2(\gamma_0 + \mu^2)I\}^{-1}\gamma \\ &\quad + \beta_0^2\gamma^\top \{\Gamma + \sigma_u^2(\gamma_0 + \mu^2)I\}^{-1} \{\Gamma + \sigma_u^2(\gamma_0 + \mu^2)I\} \{\Gamma + \sigma_u^2(\gamma_0 + \mu^2)I\}^{-1}\gamma + o_p(1) \\ &= \beta_0^2 \{(\sigma_u^2 + 1)\gamma_0 + \sigma_u^2\mu^2\} - \beta_0^2\gamma^\top \{\Gamma + \sigma_u^2(\gamma_0 + \mu^2)I\}^{-1}\gamma + o_p(1).\end{aligned}$$

Proof of Theorem 6.4(3):

Step 1: We show that as $T \rightarrow \infty$,

$$\sqrt{T} \left(\frac{1}{T} \sum_{t=1}^T (X_t^* - \mu^*)(X_{t+p}^* - \mu^*) - \frac{1}{T-p} \sum_{t=1}^{T-p} (X_t^* - \widehat{\mu}^*)(X_{t+p}^* - \widehat{\mu}^*) \right) = o_p(1). \quad (\text{E.28})$$

With some simple algebra,

$$\begin{aligned}& \sqrt{T} \left\{ \frac{1}{T} \sum_{t=1}^T (X_t^* - \mu^*)(X_{t+p}^* - \mu^*) - \frac{1}{T-p} \sum_{t=1}^{T-p} (X_t^* - \widehat{\mu}^*)(X_{t+p}^* - \widehat{\mu}^*) \right\} \\ &= \sqrt{T} \left\{ \frac{1}{T} \sum_{t=1}^T (X_t^* - \mu^*)(X_{t+p}^* - \mu^*) - \frac{1}{T-p} \sum_{t=1}^{T-p} (X_t^* - \mu^* + \mu^* - \widehat{\mu}^*)(X_{t+p}^* - \mu^* + \mu^* - \widehat{\mu}^*) \right\} \\ &= \sqrt{T} \left\{ \frac{1}{T} \sum_{t=1}^T (X_t^* - \mu^*)(X_{t+p}^* - \mu^*) - \frac{1}{T-p} \sum_{t=1}^{T-p} (X_t^* - \mu^*)(X_{t+p}^* - \mu^*) \right. \\ &\quad \left. - \frac{1}{T-p} \sum_{t=1}^{T-p} (X_t^* - \mu^*)(\mu^* - \widehat{\mu}^*) - \frac{1}{T-p} \sum_{t=1}^{T-p} (X_{t+p}^* - \mu^*)(\mu^* - \widehat{\mu}^*) - \frac{1}{T-p} \sum_{t=1}^{T-p} (\mu^* - \widehat{\mu}^*)^2 \right\} \\ &= \sqrt{T} \left(\frac{T-p}{T} - 1 \right) \frac{1}{T-p} \sum_{t=1}^{T-p} (X_t^* - \mu^*)(X_{t+p}^* - \mu^*) + \frac{1}{\sqrt{T}} \sum_{t=T-p+1}^T (X_t^* - \mu^*)(X_{t+p}^* - \mu^*) \\ &\quad + \sqrt{T}(\widehat{\mu}^* - \mu^*) \left(\frac{1}{T-p} \sum_{t=1}^{T-p} X_t^* + \frac{1}{T-p} \sum_{t=1}^{T-p} X_{t+p}^* - \widehat{\mu}^* - \mu^* \right) \end{aligned} \quad (\text{E.29})$$

$$\triangleq I_1 + I_2 + I_3.$$

Now we examine each term in (E.29) as $T \rightarrow \infty$ separately. First,

$$\begin{aligned} I_1 &= -\frac{p}{\sqrt{T}} \frac{1}{T-p} \sum_{t=1}^{T-p} (X_t^* - \mu^*)(X_{t+p}^* - \mu^*) \\ &= -\frac{p}{\sqrt{T}} \{\gamma_p^* + o_p(1)\} = o_p(1) \quad \text{as } T \rightarrow \infty. \end{aligned} \quad (\text{E.30})$$

Next, we examine the second term I_2 in (E.29). Since $T^{-\frac{1}{2}} E[\sum_{t=T-p+1}^T (X_t^* - \mu^*)(X_{t+p}^* - \mu^*)] \leq T^{-\frac{1}{2}} p \text{Var}(X_t)$ (Brockwell et al., 1991, Page 230) and $T^{-\frac{1}{2}} p \text{Var}(X_t) \rightarrow 0$ as $T \rightarrow \infty$, we have that

$$I_2 = \frac{1}{\sqrt{T}} \sum_{t=T-p+1}^T (X_t^* - \mu^*)(X_{t+p}^* - \mu^*) = o_p(1). \quad (\text{E.31})$$

Finally, we examine I_3 in (E.29).

$$\begin{aligned} & \frac{1}{T-p} \sum_{t=1}^{T-p} X_{t+p}^* - \hat{\mu}^* \\ &= \frac{1}{T-p} \sum_{t=1}^{T-p} X_{t+p}^* - \frac{1}{T} \sum_{t=1}^p X_t^* - \frac{1}{T} \sum_{t=p+1}^T X_t^* \\ &= \frac{1}{T-p} \sum_{t=1}^{T-p} X_{t+p}^* - \frac{1}{T} \sum_{t=1}^p X_t^* - \frac{1}{T} \sum_{t=1}^{T-p} X_{t+p}^* \\ &= \left(\frac{1}{T-p} - \frac{1}{T} \right) \sum_{t=1}^{T-p} X_{t+p}^* - \frac{1}{T} \sum_{t=1}^p X_t^* \\ &= o_p(1) \quad \text{as } T \rightarrow \infty, \end{aligned} \quad (\text{E.32})$$

where $\hat{\mu}^* = \frac{1}{T} \sum_{t=1}^T X_t^*$, and $\frac{1}{T} \sum_{t=1}^p X_t^* = o_p(1)$ because $E(\frac{1}{T} \sum_{t=1}^p X_t^*) = \frac{1}{T} p E(X_t) \rightarrow 0$ as $T \rightarrow \infty$. In addition, by the weak law of large numbers,

$$\frac{1}{T-p} \sum_{t=1}^{T-p} X_t^* - \mu^* \xrightarrow{p} 0 \quad \text{as } T \rightarrow \infty. \quad (\text{E.33})$$

By condition (R2) and the central limit theorem for strictly stationary p -dependent sequences (Brockwell et al., 1991, Theorem 6.4.2), we have

$$\sqrt{T}(\hat{\mu}^* - \mu^*) = O_p(1). \quad (\text{E.34})$$

Therefore, applying (E.30), (E.31), (E.32), (E.33) and (E.34) yields (E.28).

Step 2: We now show that as $T \rightarrow \infty$, the asymptotic covariance matrix of $\sqrt{T} \{(\hat{\gamma}_0^*, \hat{\gamma}^{*\top})^\top - (\gamma_0^*, \gamma^{*\top})^\top\}$ equals

$$\lim_{T \rightarrow \infty} \text{Cov} \left\{ \frac{1}{\sqrt{T}} \sum_{t=1}^T (X_t^* - \mu^*)(X_{t+r}^* - \mu^*), \frac{1}{\sqrt{T}} \sum_{s=1}^T (X_s^* - \mu^*)(X_{s+q}^* - \mu^*) \right\}.$$

For $k \leq p$

$$\begin{aligned} & \sqrt{T}(\hat{\gamma}_k - \gamma_k) \\ &= \sqrt{T} \left\{ \frac{1}{T-k} \sum_{t=1}^{T-k} (X_t^* - \hat{\mu}^*)(X_{t+k}^* - \hat{\mu}^*) - \gamma_k \right\} \\ &= \sqrt{T} \left\{ \frac{1}{T} \sum_{t=1}^T (X_t^* - \mu^*)(X_{t+k}^* - \mu^*) - \gamma_k \right\} \\ & \quad + \sqrt{T} \left\{ \frac{1}{T-k} \sum_{t=1}^{T-k} (X_t^* - \hat{\mu}^*)(X_{t+k}^* - \hat{\mu}^*) - \frac{1}{T} \sum_{t=1}^T (X_t^* - \mu^*)(X_{t+k}^* - \mu^*) \right\} \\ &= \left\{ \frac{1}{\sqrt{T}} \sum_{t=1}^T (X_t^* - \mu^*)(X_{t+k}^* - \mu^*) - \gamma_k \right\} + o_p(1), \end{aligned}$$

where the last step is due to (E.28).

Hence, the (r, q) element of matrix $\lim_{T \rightarrow \infty} \text{Var} \left(\sqrt{T} \{(\hat{\gamma}_0^*, \hat{\gamma}^{*\top})^\top - (\gamma_0^*, \gamma^{*\top})^\top\} \right)$ is given by

$$\lim_{T \rightarrow \infty} \text{Cov} \left\{ \frac{1}{\sqrt{T}} \sum_{t=1}^T (X_t^* - \mu^*)(X_{t+r}^* - \mu^*), \frac{1}{\sqrt{T}} \sum_{s=1}^T (X_s^* - \mu^*)(X_{s+q}^* - \mu^*) \right\}.$$

Step 3: We show certain identities to be used for proving Theorem 6.4(3):

1. By model (6.9), we have that

$$\begin{aligned} X_t^* - \mu^* &= \beta_0 X_t u_t - \beta_0 \mu \\ &= \beta_0 X_t u_t - \beta_0 u_t \mu + \beta_0 u_t \mu - \beta_0 \mu \\ &= \beta_0 \{u_t (X_t - \mu) + \mu (u_t - 1)\}, \end{aligned} \tag{E.35}$$

where the first step is because $\mu^* = E(\beta_0 X_t u_t) = \beta_0 E(X_t)E(u_t) = \beta_0 \mu$.

2. We have that

$$\begin{aligned}
& \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov} \{u_t^2 (X_t - \mu)^2, u_s^2 (X_s - \mu)^2\} \\
&= \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T [E\{u_t^2 u_s^2 (X_t - \mu)^2 (X_s - \mu)^2\} - E(u_t^2)E(u_s^2)E\{(X_t - \mu)^2\}E\{(X_s - \mu)^2\}], \\
&= \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{\substack{s=1 \\ s \neq t}}^T [E(u_t^2 u_s^2)E\{(X_t - \mu)^2 (X_s - \mu)^2\} - E(u_t^2)E(u_s^2)E\{(X_t - \mu)^2\}E\{(X_s - \mu)^2\}] \\
&\quad + \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{\substack{t=1 \\ s=t}}^T [E(u_t^4)E\{(X_t - \mu)^4\} - E^2(u_t^2)E^2\{(X_t - \mu)^2\}], \\
&= \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{\substack{s=1 \\ s \neq t}}^T [E(u_t^2)E(u_s^2)\text{Cov}\{(X_t - \mu)^2, (X_s - \mu)^2\}] + \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{\substack{t=1 \\ s=t}}^T E^2(u_t^2)\text{Var}\{(X_t - \mu)^2\} \\
&\quad + \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{\substack{t=1 \\ s=t}}^T \{E(u_t^4) - E^2(u_t^2)\} E\{(X_t - \mu)^4\} \\
&= \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T [E(u_t^2)E(u_s^2)\text{Cov}\{(X_t - \mu)^2, (X_s - \mu)^2\}] \tag{E.36}
\end{aligned}$$

$$\begin{aligned}
& + \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \{E(u_t^4) - E^2(u_t^2)\} E\{(X_t - \mu)^4\} \\
&= (\sigma_u^2 + 1)^2 q_{00} + \{E(u_t^4) - (\sigma_u^2 + 1)^2\} E\{(X_t - \mu)^4\}, \tag{E.37}
\end{aligned}$$

where the second and third step is due to the independence between u_t and X_t . In the last step, we use the definition $q_{00} = \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov}\{(X_t - \mu)^2, (X_s - \mu)^2\}$, $E(u_t^2) = \sigma_u^2 + 1$, and the fact that $E(u_t^4)$ and $E\{(X_t - \mu)^4\}$ are time-independent which are derived from Conditions (R1) and (R2) together with independence between u_t and X_t .

3. Similar to the derivation in (E.36), now we derive the summation of $\text{Cov}\{\beta_0^2 u_t^2 (X_t - \mu)^2, \beta_0^2 u_s u_{s+p} (X_s - \mu)(X_{s+p} - \mu)\}$ for $p > 0$,

$$\begin{aligned}
& \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov}\{\beta_0^2 u_t^2 (X_t - \mu)^2, \beta_0^2 u_s u_{s+p} (X_s - \mu)(X_{s+p} - \mu)\} \\
&= \lim_{T \rightarrow \infty} \frac{\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T [E(u_t^2 u_s u_{s+p}) E\{(X_t - \mu)^2 (X_s - \mu)(X_{s+p} - \mu)\} \\
&\quad - E(u_t^2) E(u_s) E(u_{s+p}) E(X_t - \mu)^2 E\{(X_s - \mu)(X_{s+p} - \mu)\}] \\
&= \lim_{T \rightarrow \infty} \frac{\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T E(u_t^2) E(u_s) E(u_{s+p}) \text{Cov}\{(X_t - \mu)^2, (X_s - \mu)(X_{s+p} - \mu)\} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\beta_0^4}{T} \sum_{\substack{t=1 \\ s=t}}^T \{E(u_t^3) E(u_{t+p}) - E(u_t^2) E(u_t) E(u_{t+p})\} E\{(X_t - \mu)^3 (X_{t+p} - \mu)\} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\beta_0^4}{T} \sum_{\substack{t=1 \\ s=t-p}}^T \{E(u_t^3) E(u_{t-p}) - E(u_t^2) E(u_t) E(u_{t-p})\} E\{(X_t - \mu)^3 (X_{t-p} - \mu)\} \\
&= \lim_{T \rightarrow \infty} \frac{\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T (\sigma_u^2 + 1) \text{Cov}\{(X_t - \mu)^2, (X_s - \mu)(X_{s+p} - \mu)\} \\
&\quad + \beta_0^4 \{E(u_t^3) - E(u_t^2)\} E\{(X_t - \mu)^3 (X_{t+p} - \mu)\} \\
&\quad + \beta_0^4 \{E(u_t^3) - E(u_t^2)\} E\{(X_t - \mu)^3 (X_{t-p} - \mu)\}, \\
&= \beta_0^4 \{E(u_t^3) - (\sigma_u^2 + 1)\} [E\{(X_t - \mu)^3 (X_{t+p} - \mu)\} + E\{(X_t - \mu)^3 (X_{t-p} - \mu)\}] \\
&\quad + \beta_0^4 q_{0p} (\sigma_u^2 + 1), \tag{E.38}
\end{aligned}$$

where the first step is because X_t and u_t are independent, and the second last step is due to $E(u_t^2) = \text{Var}(u_t) + E(u_t^2) = \sigma_u^2 + 1$ and is derived similar to the second and third step in (E.36), and the last step is because of the definition that $q_{0p} = \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov}\{(X_t - \mu)^2, (X_s - \mu)(X_{s+p} - \mu)\}$ and the fact that $E\{(X_t - \mu)^3 (X_{t+p} - \mu)\}$, $E\{(X_t - \mu)^3 (X_{t-p} - \mu)\}$ and $E(u_t^3)$ are time-independent, derived from Conditions (R1) and (R2) together with the independence between u_t and X_t .

4. Analogous to the derivation in (E.36) and (E.38), we derive the summation of $\text{Cov}\{u_t u_{t+p}(X_t - \mu)(X_{t+p} - \mu), u_s u_{s+r}(X_s - \mu)(X_{s+r} - \mu)\}$ for $p > 0$, $r > 0$ and $p \neq r$,

$$\begin{aligned}
& \beta_0^4 \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov}\{u_t u_{t+p}(X_t - \mu)(X_{t+p} - \mu), u_s u_{s+r}(X_s - \mu)(X_{s+r} - \mu)\} \\
&= \beta_0^4 \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T E(u_t u_{t+p} u_s u_{s+r}) \text{Cov}\{(X_t - \mu)(X_{t+p} - \mu), (X_s - \mu)(X_{s+r} - \mu)\} \\
&\quad + \beta_0^4 \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{\substack{t=1 \\ s=t}}^T \{E(u_t^2) E(u_{t+p}) E(u_{t+r}) - 1\} E\{(X_t - \mu)^2 (X_{t+p} - \mu)(X_{t+r} - \mu)\} \\
&\quad + \beta_0^4 \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{\substack{t=1 \\ s=t+p}}^T \{E(u_{t+p}^2) E(u_t) E(u_{t+p+r}) - 1\} E\{(X_t - \mu)(X_{t+p} - \mu)^2 (X_{t+p+r} - \mu)\} \\
&\quad + \beta_0^4 \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{\substack{t=1 \\ s=t-r}}^T \{E(u_t^2) E(u_{t+p}) E(u_{t-r}) - 1\} E\{(X_{t-r} - \mu)(X_t - \mu)^2 (X_{t+p} - \mu)\} \\
&\quad + \beta_0^4 \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{\substack{t=1 \\ s=t+p-r}}^T \{E(u_{t+p}^2) E(u_t) E(u_{t+p-r}) - 1\} E\{(X_t - \mu)(X_{t+p-r} - \mu)(X_{t+p} - \mu)^2\} \\
&= \beta_0^4 q_{pr} + \beta_0^4 \sigma_u^2 E\{(X_t - \mu)^2 (X_{t+p} - \mu)(X_{t+r} - \mu)\} \\
&\quad + \beta_0^4 \sigma_u^2 E\{(X_t - \mu)(X_{t+p} - \mu)^2 (X_{t+p+r} - \mu)\} \\
&\quad + \beta_0^4 \sigma_u^2 E\{(X_{t-r} - \mu)(X_t - \mu)^2 (X_{t+p} - \mu)\} \\
&\quad + \beta_0^4 \sigma_u^2 E\{(X_t - \mu)(X_{t+p-r} - \mu)(X_{t+p} - \mu)^2\}, \tag{E.39}
\end{aligned}$$

where the third step is derived analogously to the second step of (E.38), and $E(u_t u_{t+p} u_s u_{s+r}) = 1$, and the last step is due to the definition $q_{pr} = \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov}\{(X_t - \mu)(X_{t+p} - \mu), (X_s - \mu)(X_{s+r} - \mu)\}$ and the fact that $E\{(X_t - \mu)^2 (X_{t+p} - \mu)(X_{t+r} - \mu)\}$, $E\{(X_t - \mu)(X_{t+p} - \mu)^2 (X_{t+p+r} - \mu)\}$, $E\{(X_{t-r} - \mu)(X_t - \mu)^2 (X_{t+p} - \mu)\}$, and $E\{(X_t - \mu)(X_{t+p} - \mu)^2 (X_{t+p-r} - \mu)\}$ are time-independent derived from Conditions (R1) and (R2).

5. Similar to the derivation in (E.36), (E.38), and (E.39), we derive the summation of $\text{Cov}\{u_t u_{t+p}(X_t - \mu)(X_{t+p} - \mu), u_s u_{s+p}(X_s - \mu)(X_{s+p} - \mu)\}$ for $p > 0$,

$$\begin{aligned}
& \beta_0^4 \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov}\{u_t u_{t+p}(X_t - \mu)(X_{t+p} - \mu), u_s u_{s+p}(X_s - \mu)(X_{s+p} - \mu)\} \\
&= \beta_0^4 \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T E(u_t)E(u_{t+p})E(u_s)E(u_{s+p}) \text{Cov}\{(X_t - \mu)(X_{t+p} - \mu), (X_s - \mu)(X_{s+p} - \mu)\} \\
&\quad + \beta_0^4 \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{\substack{t=1 \\ s=t}}^T \{E(u_t^2)E(u_{t+p}^2) - 1\} \text{Var}\{(X_t - \mu)(X_{t+p} - \mu)\} \\
&\quad + \beta_0^4 \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{\substack{t=1 \\ s=t+p}}^T \{E(u_{t+p}^2)E(u_t)E(u_{t+2p}) - 1\} E\{(X_t - \mu)(X_{t+p} - \mu)^2(X_{t+2p} - \mu)\} \\
&\quad + \beta_0^4 \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{\substack{t=1 \\ s=t-p}}^T \{E(u_t^2)E(u_{t-p})E(u_{t+p}) - 1\} E\{(X_{t-p} - \mu)(X_t - \mu)^2(X_{t+p} - \mu)\} \\
&= \beta_0^4 q_{pp} + \beta_0^4 (\sigma_u^4 + 2\sigma_u^2) \text{Var}\{(X_t - \mu)(X_{t+p} - \mu)\} + 2\beta_0^4 E\{(X_t - \mu)(X_{t+p} - \mu)^2(X_{t+2p} - \mu)\}, \tag{E.40}
\end{aligned}$$

where the last step is by the definition $q_{pp} = \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov}\{(X_t - \mu)(X_{t+p} - \mu), (X_s - \mu)(X_{s+p} - \mu)\}$ and $E\{(X_t - \mu)(X_{t+p} - \mu)^2(X_{t+2p} - \mu)\} = E\{(X_{t-p} - \mu)(X_t - \mu)^2(X_{t+p} - \mu)\}$ due to the stationarity of the time series and the fact that $\text{Var}\{(X_t - \mu)(X_{t+p} - \mu)\}$ and $E\{(X_t - \mu)(X_{t+p} - \mu)^2(X_{t+2p} - \mu)\}$ are time-independent, resulted from the Conditions (R1) and (R2).

6. For any t, s and p , we have that

$$\begin{aligned}
& \text{Cov}\{(X_t - \mu)(X_{t-p} - \mu), (X_s - \mu)\} \\
&= E\{(X_t - \mu)(X_{t-p} - \mu)(X_s - \mu)\} - E\{(X_t - \mu)(X_{t-p} - \mu)\}E(X_s - \mu) \\
&= E\{(X_t - \mu)(X_{t-p} - \mu)(X_s - \mu)\}, \tag{E.41}
\end{aligned}$$

where the last step is because $E(X_s - \mu) = 0$.

7. For any t and s , we have that

$$\begin{aligned}
& \text{Cov}\{u_t(u_t - 1)(X_t - \mu), u_s(u_s - 1)(X_s - \mu)\} \\
&= E\{u_t(u_t - 1)(X_t - \mu)u_s(u_s - 1)(X_s - \mu)\} - E\{u_t(u_t - 1)(X_t - \mu)\}E\{u_s(u_s - 1)(X_s - \mu)\} \\
&= E\{u_t(u_t - 1)(X_t - \mu)u_s(u_s - 1)(X_s - \mu)\} \\
&= E\{u_t(u_t - 1)u_s(u_s - 1)\}E\{(X_t - \mu)(X_s - \mu)\}, \tag{E.42}
\end{aligned}$$

where the second step is because of the independence between u_t and X_t and that $E(X_t - \mu) = 0$. Then, $E\{u_t(u_t - 1)u_s(u_s - 1)\} = \sigma_u^4$ for $t \neq s$ and $E\{u_t^2(u_t - 1)^2\} = E(u_t^4) - 2E(u_t^3) + \sigma_u^2 + 1$ for any t .

By (E.42), we have that

$$\begin{aligned}
& \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov}\{u_t(u_t - 1)(X_t - \mu), u_s(u_s - 1)(X_s - \mu)\} \\
&= \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T E\{u_t(u_t - 1)u_s(u_s - 1)\}E\{(X_t - \mu)(X_s - \mu)\} \\
&= \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T \sigma_u^4 E\{(X_t - \mu)(X_s - \mu)\} \\
&\quad + \lim_{T \rightarrow \infty} \frac{1}{T} \sum_{\substack{t=1 \\ s=t}}^T \{E(u_t^4) - 2E(u_t^3) + \sigma_u^2 + 1 - \sigma_u^4\} E\{(X_t - \mu)^2\} \\
&= \sigma_u^4 \sum_{h=-\infty}^{\infty} \gamma_h + \{E(u_t^4) - 2E(u_t^3) + \sigma_u^2 + 1 - \sigma_u^4\} \gamma_0, \tag{E.43}
\end{aligned}$$

where the last is because $\lim_{T \rightarrow \infty} \frac{1}{T} \sum_{t=1}^T \sum_{s=1}^T E\{(X_t - \mu)(X_s - \mu)\} = \sum_{h=-\infty}^{\infty} \gamma_h$ (Brockwell et al., 1991, Theorem 7.1.1).

8. For any t , s and $p > 0$, we have that

$$\begin{aligned}
& \text{Cov}\{u_t(u_t - 1)(X_t - \mu), u_{s+p}(u_s - 1)(X_{s+p} - \mu)\} \\
&= E\{u_t(u_t - 1)(X_t - \mu)u_{s+p}(u_s - 1)(X_{s+p} - \mu)\} \\
&\quad - E\{u_t(u_t - 1)(X_t - \mu)\}E\{u_{s+p}(u_s - 1)(X_{s+p} - \mu)\} \\
&= E\{u_t(u_t - 1)u_{s+p}(u_s - 1)\}E\{(X_t - \mu)(X_{s+p} - \mu)\} \\
&= E\{u_t(u_t - 1)u_{s+p}(u_s - 1)\}\gamma_{|s+p-t|}, \tag{E.44}
\end{aligned}$$

where the second step is because of the independence between u_t and X_t and that $E(X_t - \mu) = 0$. Then, $E\{u_t(u_t - 1)u_{s+p}(u_s - 1)\} = 0$ for $t \neq s$ and $E\{u_t(u_t - 1)^2u_{t+p}\} = E\{u_t(u_t - 1)^2\} = E\{(u_t - 1)^3\} + \sigma_u^2$ for any $s = t$.

9. By independence of u_t and u_s , for $t \neq s$, we have that

$$\text{Cov}\{u_t^2(X_t - \mu)^2, (u_s - 1)^2\} = 0, \quad (\text{E.45})$$

and for any t ,

$$\begin{aligned} & \text{Cov}\{u_t^2(X_t - \mu)^2, (u_t - 1)^2\} \\ &= E\{u_t^2(u_t - 1)^2(X_t - \mu)^2\} - E\{u_t^2(X_t - \mu)^2\}E\{(u_t - 1)^2\} \\ &= [E\{u_t^2(u_t - 1)^2\} - E(u_t^2)E(u_t - 1)^2] E\{(X_t - \mu)^2\} \\ &= \{E(u_t^4) - 2E(u_t^3) + \sigma_u^2 + 1 - \sigma_u^4 - \sigma_u^2\} \gamma_0 \\ &= \{E(u_t^4) - 2E(u_t^3) + 1 - \sigma_u^4\} \gamma_0. \end{aligned} \quad (\text{E.46})$$

10. By independence of u_t and u_s , for $s \neq t$, $s \neq t + p$ and any p , we have that

$$\text{Cov}\{u_t u_{t+p}(X_t - \mu)(X_{t+p} - \mu), (u_s - 1)^2\} = 0. \quad (\text{E.47})$$

For any t and $p > 0$,

$$\begin{aligned} & \text{Cov}\{u_t u_{t+p}(X_t - \mu)(X_{t+p} - \mu), (u_t - 1)^2\} \\ &= E\{u_t u_{t+p}(u_t - 1)^2(X_t - \mu)(X_{t+p} - \mu)\} - E\{u_t u_{t+p}(X_t - \mu)(X_{t+p} - \mu)\}E\{(u_t - 1)^2\}, \\ &= [E\{u_t u_{t+p}(u_t - 1)^2\} - E(u_t u_{t+p})E\{(u_t - 1)^2\}] E\{(X_t - \mu)(X_{t+p} - \mu)\} \\ &= E\{(u_t - 1)^3\} \gamma_p, \end{aligned} \quad (\text{E.48})$$

and

$$\text{Cov}\{u_t u_{t-p}(X_t - \mu)(X_{t-p} - \mu), (u_t - 1)^2\} = E\{(u_t - 1)^3\} \gamma_p.$$

11. For any t and s , and $r \neq p$ and $r > 0$, we have that

$$\text{Cov}\{u_t u_{t+p}(X_t - \mu)(X_{t+p} - \mu), (u_s - 1)(u_{s+r} - 1)\} = 0. \quad (\text{E.49})$$

By independence of u_t and u_s , for $t \neq s$ and any p , we have that

$$\text{Cov}\{u_t u_{t+p}(X_t - \mu)(X_{t+p} - \mu), (u_s - 1)(u_{s+p} - 1)\} = 0, \quad (\text{E.50})$$

and for any t and $p > 0$,

$$\begin{aligned} & \text{Cov}\{u_t u_{t+p}(X_t - \mu)(X_{t+p} - \mu), (u_t - 1)(u_{t+p} - 1)\} \\ &= E\{u_t u_{t+p}(u_t - 1)(u_{t+p} - 1)(X_t - \mu)(X_{t+p} - \mu)\} \end{aligned} \quad (\text{E.51})$$

$$\begin{aligned} & - E\{u_t u_{t+p}(X_t - \mu)(X_{t+p} - \mu)\} E\{(u_t - 1)(u_{t+p} - 1)\} \\ &= E\{u_t(u_t - 1)\} E\{u_{t+p}(u_{t+p} - 1)\} E\{(X_t - \mu)(X_{t+p} - \mu)\} \\ &= \sigma_u^4 \gamma_p. \end{aligned} \quad (\text{E.52})$$

12. For any t , we have that

$$\begin{aligned} & \text{Cov}\{u_t(u_t - 1)(X_t - \mu), (u_s - 1)^2\} \\ &= E\{u_t(u_t - 1)(X_t - \mu)(u_s - 1)^2\} - E\{u_t(u_t - 1)(X_t - \mu)\} E\{(u_s - 1)^2\} \\ &= [E\{u_t(u_t - 1)(u_s - 1)^2\} - E\{u_t(u_t - 1)\} E\{(u_s - 1)^2\}] E(X_t - \mu) = 0, \end{aligned} \quad (\text{E.53})$$

where the last step is because $E(X_t - \mu) = 0$.

13. By independence assumption between $\{u_t\}$, if $t \neq s$ or $p \neq r$, we have that

$$\text{Cov}\{(u_t - 1)(u_{t+p} - 1), (u_s - 1)(u_{s+r} - 1)\} = 0. \quad (\text{E.54})$$

In addition, for any t and p we have that

$$\begin{aligned} & \text{Var}\{(u_t - 1)(u_{t+p} - 1)\} \\ &= E\{(u_t - 1)^2(u_{t+p} - 1)^2\} \\ &= E\{(u_t - 1)^2\} E\{(u_{t+p} - 1)^2\} \\ &= \sigma_u^4, \end{aligned} \quad (\text{E.55})$$

and for any t , we have that

$$\begin{aligned} & \text{Var}(u_t - 1)^2 \\ &= E\{(u_t - 1)^4\} - E^2\{(u_t - 1)^2\} \\ &= E\{(u_t - 1)^4\} - \sigma_u^4. \end{aligned} \quad (\text{E.56})$$

Step 4: Now we prove the results in (3).

1°. We first show the derivation of q_{200}^* as follows:

$$\begin{aligned}
q_{200}^* &= \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T} \sum_{t=1}^T (X_t^* - \mu^*)^2, \frac{1}{T} \sum_{s=1}^T (X_s^* - \mu^*)^2 \right\} \\
&= \lim_{T \rightarrow \infty} \frac{\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ u_t^2 (X_t - \mu)^2 + 2\mu u_t (u_t - 1)(X_t - \mu) + \mu^2 (u_t - 1)^2, \right. \\
&\quad \left. u_s^2 (X_s - \mu)^2 + 2\mu u_s (u_s - 1)(X_s - \mu) + \mu^2 (u_s - 1)^2 \right\} \\
&= \lim_{T \rightarrow \infty} \frac{\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ u_t^2 (X_t - \mu)^2, u_s^2 (X_s - \mu)^2 \right\} \\
&\quad + \lim_{T \rightarrow \infty} \frac{4\mu\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ u_t^2 (X_t - \mu)^2, u_s (u_s - 1)(X_s - \mu) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \frac{2\mu^2\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ u_t^2 (X_t - \mu)^2, (u_s - 1)^2 \right\} \\
&\quad + \lim_{T \rightarrow \infty} \frac{4\mu^2\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ u_t (u_t - 1)(X_t - \mu), u_s (u_s - 1)(X_s - \mu) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\mu^4\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ (u_t - 1)^2, (u_s - 1)^2 \right\}, \\
&= \beta_0^4 (\sigma_u^2 + 1)^2 q_0 + \beta_0^4 \{ E(u_t^4) - (\sigma_u^2 + 1)^2 \} E\{(X_t - \mu)^4\} \\
&\quad + 4\mu\beta_0^4 \sigma_u^2 (\sigma_u^2 + 1) v_{00} + 4\mu\beta_0^4 \{ E(u_t^4) - E(u_t^3) - \sigma_u^2 (\sigma_u^2 + 1) \} E\{(X_t - \mu)^3\} \\
&\quad + 2\mu^2\beta_0^4 \{ E(u_t^4) - 2E(u_t^3) + 1 - \sigma_u^4 \} \gamma_0 \\
&\quad + 4\mu^2\beta_0^4 \left[\sigma_u^4 \sum_{h=-\infty}^{\infty} \gamma_h + \{ E(u_t^4) - 2E(u_t^3) + \sigma_u^2 + 1 - \sigma_u^4 \} \gamma_0 \right] \\
&\quad + \mu^4\beta_0^4 [E\{(u_t - 1)^4\} - \sigma_u^4],
\end{aligned}$$

where the second step is due to (E.35), the third step is because of (E.53), the last step is by (E.36), (E.41), (E.43), (E.45), (E.46), and (E.56).

2°. Then we derive the value of q_{20p}^* :

$$\begin{aligned}
q_{20p}^* &= \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T} \sum_{t=1}^T (X_t^* - \mu^*)^2, \frac{1}{T} \sum_{s=1}^T (X_s^* - \mu^*)(X_{s+p}^* - \mu^*) \right\} \\
&= \lim_{T \rightarrow \infty} \frac{\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ u_t^2 (X_t - \mu)^2 + 2\mu u_t (u_t - 1)(X_t - \mu) + \mu^2 (u_t - 1)^2, \right. \\
&\quad \left. u_s u_{s+p} (X_s - \mu)(X_{s+p} - \mu) + \mu u_s (u_{s+p} - 1)(X_s - \mu) \right. \\
&\quad \left. + \mu u_{s+p} (u_s - 1)(X_{s+p} - \mu) + \mu^2 (u_s - 1)(u_{s+p} - 1) \right\} \\
&= \lim_{T \rightarrow \infty} \frac{\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ u_t^2 (X_t - \mu)^2, u_s u_{s+p} (X_s - \mu)(X_{s+p} - \mu) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\mu \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ u_t^2 (X_t - \mu)^2, u_s (u_{s+p} - 1)(X_s - \mu) + u_{s+p} (u_s - 1)(X_{s+p} - \mu) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \frac{2\mu \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ u_s u_{s+p} (X_s - \mu)(X_{s+p} - \mu), u_t (u_t - 1)(X_t - \mu) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\mu^2 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T [Cov \left\{ u_t^2 (X_t - \mu)^2, (u_s - 1)(u_{s+p} - 1) \right\} \\
&\quad + Cov \left\{ (u_t - 1)^2, u_s u_{s+p} (X_s - \mu)(X_{s+p} - \mu) \right\}] \\
&\quad + \lim_{T \rightarrow \infty} \frac{2\mu^2 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ u_t (u_t - 1)(X_t - \mu), u_s (u_{s+p} - 1)(X_s - \mu) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \frac{2\mu^2 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ u_t (u_t - 1)(X_t - \mu), u_{s+p} (u_s - 1)(X_{s+p} - \mu) \right\} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\mu^4 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ (u_t - 1)^2, (u_s - 1)(u_{s+p} - 1) \right\}, \\
&= \beta_0^4 q_{0p} (\sigma_u^2 + 1) + \beta_0^4 \{ E(u_t^3) - (\sigma_u^2 + 1) \} [E\{(X_t - \mu)^3 (X_{t+p} - \mu)\} + E\{(X_t - \mu)^3 (X_{t-p} - \mu)\}] \\
&\quad + \mu \beta_0^4 E\{u_t^3 - u_t^2\} [E\{(X_t - \mu)^2 (X_{t-p} - \mu)\} + E\{(X_t - \mu)^2 (X_{t+p} - \mu)\}] \\
&\quad + 2\mu \beta_0^4 \sigma_u^2 v_{0p} + 2\mu \beta_0^4 E\{u_t^3 - u_t^2 - \sigma_u^2\} [E\{(X_t - \mu)^2 (X_{t-p} - \mu)\} + E\{(X_t - \mu)^2 (X_{t+p} - \mu)\}] \\
&\quad + 2\mu^2 \beta_0^4 E(u_t - 1)^3 \gamma_p + 4\mu^2 \beta_0^4 \{ E(u_t - 1)^3 + \sigma_u^2 \} \gamma_p + \mu^4 \beta_0^4 \sigma_u^4 \\
&= \beta_0^4 q_p (\sigma_u^2 + 1) + \beta_0^4 \{ E(u_t^3) - (\sigma_u^2 + 1) \} [E\{(X_t - \mu)^3 (X_{t+p} - \mu)\} + E\{(X_t - \mu)^3 (X_{t-p} - \mu)\}] \\
&\quad + 2\mu \beta_0^4 \sigma_u^2 v_p + \mu \beta_0^4 E\{3u_t^3 - 3u_t^2 - 2\sigma_u^2\} [E\{(X_t - \mu)^2 (X_{t-p} - \mu)\} + E\{(X_t - \mu)^2 (X_{t+p} - \mu)\}] \\
&\quad + 6\mu^2 \beta_0^4 E(u_t - 1)^3 \gamma_p + 4\mu^2 \beta_0^4 \sigma_u^2 \gamma_p,
\end{aligned}$$

where the second step is by (E.35), the third step is because (E.41) and (E.53), and the second last step is because (E.38), (E.49), (E.48), (E.44), and (E.54).

3°. Then we derive the value of q_{2pr}^* for $r \neq p$

$$\begin{aligned}
q_{2pr}^* &= \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T} \sum_{t=1}^T (X_t^* - \mu^*)(X_{t+p}^* - \mu^*), \frac{1}{T} \sum_{s=1}^T (X_s^* - \mu^*)(X_{s+r}^* - \mu^*) \right\} \\
&= \lim_{T \rightarrow \infty} \frac{\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ u_t u_{t+p} (X_t - \mu)(X_{t+p} - \mu) + \mu u_t (u_{t+p} - 1)(X_t - \mu) \\
&\quad + \mu u_{t+p} (u_t - 1)(X_{t+p} - \mu) + \mu^2 (u_t - 1)(u_{t+p} - 1), \\
&\quad u_s u_{s+r} (X_s - \mu)(X_{s+r} - \mu) + \mu u_s (u_{s+r} - 1)(X_s - \mu) + \mu u_{s+r} (u_s - 1)(X_{s+r} - \mu) + \mu^2 (u_s - 1)(u_{s+r} - 1) \} \\
&= \lim_{T \rightarrow \infty} \frac{\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ u_t u_{t+p} (X_t - \mu)(X_{t+p} - \mu), u_s u_{s+r} (X_s - \mu)(X_{s+r} - \mu) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\mu \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ u_s u_{s+r} (X_s - \mu)(X_{s+r} - \mu), u_t (u_{t+p} - 1)(X_t - \mu) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\mu \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ u_s u_{s+r} (X_s - \mu)(X_{s+r} - \mu), u_{t+p} (u_t - 1)(X_{t+p} - \mu) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\mu \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ u_t u_{t+p} (X_t - \mu)(X_{t+p} - \mu), u_s (u_{s+r} - 1)(X_s - \mu) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\mu \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ u_t u_{t+p} (X_t - \mu)(X_{t+p} - \mu), u_{s+r} (u_s - 1)(X_{s+r} - \mu) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{2\mu^2 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ u_t u_{t+p} (X_t - \mu)(X_{t+p} - \mu), (u_s - 1)(u_{s+r} - 1) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\mu^2 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ u_t (u_{t+p} - 1)(X_t - \mu), u_s (u_{s+r} - 1)(X_s - \mu) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\mu^2 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ u_t (u_{t+p} - 1)(X_t - \mu), u_{s+r} (u_s - 1)(X_{s+r} - \mu) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\mu^2 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ u_{t+p} (u_t - 1)(X_{t+p} - \mu), u_s (u_{s+r} - 1)(X_s - \mu) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\mu^2 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ u_{t+p} (u_t - 1)(X_{t+p} - \mu), u_{s+r} (u_s - 1)(X_{s+r} - \mu) \} \\
&\quad + \lim_{T \rightarrow \infty} \frac{\mu^4 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \{ (u_t - 1)(u_{t+p} - 1), (u_s - 1)(u_{s+r} - 1) \},
\end{aligned}$$

where the second step is by (E.35), the third step is because (E.41) and (E.53). Then because (E.38), (E.49), (E.44), and (E.54), we have that

$$\begin{aligned}
q_{2pr}^* &= \beta_0^4 q_{pr} + \beta_0^4 \sigma_u^2 \left[E\{(X_t - \mu)^2 (X_{t+p} - \mu)(X_{t+r} - \mu)\} \right. \\
&\quad + E\{(X_t - \mu)(X_{t+p} - \mu)^2 (X_{t+p+r} - \mu)\} \\
&\quad + E\{(X_{t-r} - \mu)(X_t - \mu)^2 (X_{t+p} - \mu)\} \\
&\quad + E\{(X_t - \mu)(X_{t+p-r} - \mu)(X_{t+p} - \mu)^2\} \left. \right] \\
&\quad + \mu \beta_0^4 \sigma_u^2 \left[E\{(X_t - \mu)(X_{t+p} - \mu)(X_{t+r} - \mu)\} \right. \\
&\quad + E\{(X_t - \mu)(X_{t+p} - \mu)(X_{t+p+r} - \mu)\} \\
&\quad + E\{(X_{t-r} - \mu)(X_t - \mu)(X_{t+p} - \mu)\} \\
&\quad + E\{(X_t - \mu)(X_{t+p-r} - \mu)(X_{t+p} - \mu)\} \left. \right] \\
&\quad + 2\mu^2 \beta_0^4 \sigma_u^2 (\gamma_{|p-r|} + \gamma_{p+r}). \tag{E.57}
\end{aligned}$$

4°. Finally, similar to the derivation of q_{2pq}^* , now we derive the value of q_{2pp}^* . By (E.35),

$$\begin{aligned}
q_{2pp}^* &= \lim_{T \rightarrow \infty} TCov \left\{ \frac{1}{T} \sum_{t=1}^T (X_t^* - \mu^*)(X_{t+p}^* - \mu^*), \frac{1}{T} \sum_{s=1}^T (X_s^* - \mu^*)(X_{s+p}^* - \mu^*) \right\} \\
&= \lim_{T \rightarrow \infty} \frac{\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T Cov \left\{ u_t u_{t+p} (X_t - \mu)(X_{t+p} - \mu) + \mu u_t (u_{t+p} - 1)(X_t - \mu) \right. \\
&\quad + \mu u_{t+p} (u_t - 1)(X_{t+p} - \mu) + \mu^2 (u_t - 1)(u_{t+p} - 1), \\
&\quad u_s u_{s+r} (X_s - \mu)(X_{s+p} - \mu) + \mu u_s (u_{s+p} - 1)(X_s - \mu) \\
&\quad \left. + \mu u_{s+p} (u_s - 1)(X_{s+p} - \mu) + \mu^2 (u_s - 1)(u_{s+p} - 1) \right\}.
\end{aligned}$$

Then, because (E.41) and (E.53), we have that,

$$\begin{aligned}
q_{2pp}^* &= \lim_{T \rightarrow \infty} \frac{\beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov} \{u_t u_{t+p} (X_t - \mu)(X_{t+p} - \mu), u_s u_{s+p} (X_s - \mu)(X_{s+p} - \mu)\} \\
&+ \lim_{T \rightarrow \infty} \frac{2\mu^2 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov} \{u_t u_{t+p} (X_t - \mu)(X_{t+p} - \mu), (u_s - 1)(u_{s+p} - 1)\} \\
&+ \lim_{T \rightarrow \infty} \frac{\mu \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov} \{u_s u_{s+p} (X_s - \mu)(X_{s+p} - \mu), u_t (u_{t+p} - 1)(X_t - \mu)\} \\
&+ \lim_{T \rightarrow \infty} \frac{\mu \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov} \{u_s u_{s+p} (X_s - \mu)(X_{s+p} - \mu), u_{t+p} (u_t - 1)(X_{t+p} - \mu)\} \\
&+ \lim_{T \rightarrow \infty} \frac{\mu \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov} \{u_t u_{t+p} (X_t - \mu)(X_{t+p} - \mu), u_s (u_{s+p} - 1)(X_s - \mu)\} \\
&+ \lim_{T \rightarrow \infty} \frac{\mu \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov} \{u_t u_{t+p} (X_t - \mu)(X_{t+p} - \mu), u_{s+p} (u_s - 1)(X_{s+p} - \mu)\} \\
&+ \lim_{T \rightarrow \infty} \frac{\mu^2 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov} \{u_t (u_{t+p} - 1)(X_t - \mu), u_s (u_{s+p} - 1)(X_s - \mu)\} \\
&+ \lim_{T \rightarrow \infty} \frac{\mu^2 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov} \{u_t (u_{t+p} - 1)(X_t - \mu), u_{s+p} (u_s - 1)(X_{s+p} - \mu)\} \\
&+ \lim_{T \rightarrow \infty} \frac{\mu^2 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov} \{u_{t+p} (u_t - 1)(X_{t+p} - \mu), u_s (u_{s+p} - 1)(X_s - \mu)\} \\
&+ \lim_{T \rightarrow \infty} \frac{\mu^2 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov} \{u_{t+p} (u_t - 1)(X_{t+p} - \mu), u_{s+p} (u_s - 1)(X_{s+p} - \mu)\} \\
&+ \lim_{T \rightarrow \infty} \frac{\mu^4 \beta_0^4}{T} \sum_{t=1}^T \sum_{s=1}^T \text{Cov} \{(u_t - 1)(u_{t+p} - 1), (u_s - 1)(u_{s+p} - 1)\}.
\end{aligned}$$

Then, because (E.40), (E.50), (E.51) and (E.55), we have that,

$$\begin{aligned}
q_{2pp}^* &= \beta_0^4 q_{pp} + \beta_0^4 (\sigma_u^4 + 2\sigma_u^2) \text{Var}\{(X_t - \mu)(X_{t+p} - \mu)\} \\
&+ 2\beta_0^4 E\{(X_t - \mu)(X_{t+p} - \mu)^2(X_{t+2p} - \mu)\} \\
&+ \mu \beta_0^4 \sigma_u^2 \left[E\{(X_t - \mu)(X_{t+p} - \mu)^2\} + 2E\{(X_t - \mu)(X_{t+p} - \mu)(X_{t+2p} - \mu)\} \right. \\
&+ \left. E\{(X_t - \mu)^2(X_{t+p} - \mu)\} \right] \\
&+ 2\mu^2 \beta_0^4 \sigma_u^4 \gamma_p + 2\mu^2 \beta_0^4 \sigma_u^2 (\gamma_0 + \gamma_{2p}) + \mu^4 \beta_0^4 \sigma_u^4. \tag{E.58}
\end{aligned}$$