

# New Paradigm of Identifiable General-response Cognitive Diagnostic Models: Beyond Categorical Data

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## Abstract

Cognitive diagnostic models (CDMs) are a popular family of discrete latent variable models that model students' mastery or deficiency of multiple fine-grained skills. CDMs have been most widely used to model categorical item response data such as binary or polytomous responses. With advances in technology and the emergence of varying test formats in modern educational assessments, new response types, including continuous responses such as response times, and count-valued responses from tests with repetitive tasks or eye-tracking sensors, have also become available. Variants of CDMs have been proposed recently for modeling such responses. However, whether these extended CDMs are identifiable and estimable is entirely unknown. We propose a very general cognitive diagnostic modeling framework for arbitrary types of multivariate responses with minimal assumptions, and establish identifiability in this general setting. Surprisingly, we prove that our general-response CDMs are identifiable under **Q**-matrix-based conditions similar to those for traditional categorical-response CDMs. Our conclusions set up a new paradigm of identifiable general-response CDMs. We propose an EM algorithm to efficiently estimate a broad class of exponential family-based general-response CDMs (ExpCDMs). We conduct simulation studies under various response types. The simulation results not only corroborate our identifiability theory, but also demonstrate the superior empirical performance of our estimation algorithms. We illustrate our methodology by applying it to a TIMSS 2019 response time dataset.

**Keywords:** Cognitive diagnostic model; Diagnostic classification model; EM algorithm; Exponential family; General responses; Generalized linear model; Identifiability; **Q**-matrix.

## 1 Introduction

Cognitive diagnostic models (CDMs, also called diagnostic classification models; see [Rupp et al., 2010](#)) are a popular family of discrete latent variable models in educational and

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psychological measurement. CDMs employ multiple discrete latent variables to model and diagnose students’ mastery or deficiency of a set of fine-grained skills. Popular examples in the literature include the Deterministic Input Noisy output “And” gate model (DINA model; Junker and Sijtsma, 2001), the log-linear CDM (LCDM; Henson et al., 2009), the additive CDM (ACDM; de la Torre, 2011), and general diagnostic models (GDM; von Davier, 2008).

Originally, CDMs were developed mostly to model binary responses and later generalized to polytomous responses, which are both categorical. Recently, with advances in technology and the emergence of varying test formats in modern educational assessments, new response types have become available. In particular, *multivariate continuous* and *count* responses are especially common. Continuous responses arise in the following scenarios: (a) responses that place a mark on a line segment (such as the visual analog scale), (b) assessments that record the probability of each option being correct (probability testing), (c) computer-based tests that record the response time (Minchen et al., 2017). In particular, modeling response times has received great interest for a long time, and many different models have been proposed to this end; see De Boeck and Jeon (2019) for an overview. Another common response type is count responses. They arise in the following scenarios: (a) assessments with repetitive tasks where the number of correct responses is recorded, (b) assessments where examinees read aloud a passage and the number of errors is recorded, (c) modern exams with eye tracking sensors that record students’ visual fixation counts, (d) computer-based tests that record the visit count per item (Man and Harring, 2019; Liu et al., 2022). Rasch (1993) first proposed a Poisson-based item response theory (IRT) model for count data, and many other models have been proposed (Magnus and Thissen, 2017; Man and Harring, 2019, 2022).

Many existing latent variable models for general responses are based on IRT that use continuous latent traits to model the unobserved constructs (Thissen, 1983; van der Linden, 2007; Wang and Xu, 2015). On the other hand, using discrete latent variables as in CDMs can provide students with valuable personalized diagnoses of the mastery/deficiency profiles of the latent skills. Currently, there are a few CDMs proposed for modeling non-categorical responses. For instance, Minchen et al. (2017) proposed a DINA model with a lognormal link to model response time data, whereas Liu et al. (2022, 2023) proposed GDMs (which include the DINA model as a submodel) with a Poisson link and a negative binomial link, respectively, to model visual fixation count data. It is desirable to propose a general modeling

framework of CDMs for flexible types of responses and their associated estimation methods.

When proposing new statistical models, identifiability is a crucial consideration, because it is a fundamental prerequisite for valid statistical estimation and inference. A model is identifiable if the parameters can be uniquely recovered from the observed data distribution. In recent years, many identifiability results have been established for the categorical-response CDMs, including both binary and polytomous cases (e.g., [Chen et al., 2015](#); [Xu and Zhang, 2016](#); [Fang et al., 2019](#); [Gu and Xu, 2019, 2020](#); [Culpepper, 2019, 2023](#)). These studies typically show that CDMs for multivariate categorical data are identifiable under structural constraints on the  $\mathbf{Q}$ -matrix. The  $\mathbf{Q}$ -matrix here is a key component in a CDM that specifies how the observed responses depend on the latent attributes (see its formal definition in [Section 2](#)). However, it is entirely unknown whether or not those extended CDMs for continuous or count data (such as those in [Minchen et al. \(2017\)](#) and [Liu et al. \(2022\)](#)) are identifiable, let alone identifiability of models for more general response types.

This manuscript makes the following contributions. *First*, we propose a very general new framework of  $\mathbf{Q}$ -matrix-based CDMs for modeling rich types of responses. In particular, this framework includes a sub-family of *exponential family-based CDMs* (ExpCDMs), which is a wide class of parametric CDMs using exponential families to model general responses. Our general modeling framework covers existing CDMs for continuous and count responses as special cases ([Minchen et al., 2017](#); [Minchen and de la Torre, 2018](#); [Liu et al., 2022, 2023](#)). *Second*, we provide the crucial identifiability guarantees for the proposed new models. Somewhat surprisingly, we prove that our general model is identifiable under similar structural conditions on the  $\mathbf{Q}$ -matrix, just as those for traditional categorical-response CDMs. This is the first identifiability result for CDMs with non-categorical responses. Our conclusions set up a new paradigm of identifiable general-response CDMs and significantly advances the psychometric theory of diagnostic modeling. *Third*, we propose an EM algorithm to efficiently estimate model parameters. As concrete demonstrations, we consider the DINA and main-effect-based CDMs for general responses and derive explicit updates in the EM algorithms. Our simulation results corroborate the identifiability theory, and also demonstrate the superior empirical performance of the proposed estimation algorithms.

The remainder of this manuscript is organized as follows. [Section 2](#) formally introduces our general-response CDM framework and gives many examples of its parametric submodels.

Section 3 provides conditions for model identifiability. Section 4 proposes a general-purpose EM algorithm to estimate the model parameters. Section 5 presents simulation studies under various response types. Section 6 illustrates our methodology via application to a real-world response time dataset from the Trends in International Mathematics and Science Study (TIMSS) in 2019. Section 7 concludes and discusses future research directions. The Supplementary Material contains the proofs of the theorems and more details of the simulation studies and real data analysis.

## 2 General-response Cognitive Diagnostic Models

### 2.1 General Model Setup

Consider an educational assessment with  $J$  items, and denote an observed response vector by  $\mathbf{Y} = (Y_1, \dots, Y_J) \in \times_{j=1}^J \mathcal{Y}_j$ , where each  $\mathcal{Y}_j$  is the sample space of the random variable  $Y_j$ . We next adopt the measure-theoretic terms to introduce our modeling framework in full generality; these terms follow from classical probability theory textbooks such as [Durrett \(2019\)](#). We consider very general types of responses where the sample spaces  $\mathcal{Y}_j$ 's only need to be separable metric spaces. Let  $\mathcal{F}_j$  and  $m_j$  be the Borel sigma-algebra and a measure on  $\mathcal{Y}_j$ , respectively. In many concrete model examples,  $\mathcal{F}_j$  and  $m_j$  are naturally defined upon specifying the response type of  $Y_j \in \mathcal{Y}_j$ . For instance, when  $\mathcal{Y}_j = \mathbb{R}$  with  $Y_j$  being a continuous response,  $\mathcal{F}_j$  is the collection of Borel sets in  $\mathbb{R}$  and  $m_j$  is the Lebesgue measure on  $\mathbb{R}$ ; when  $\mathcal{Y}_j = \{0, 1, \dots\}$  is the set of nonnegative integers with  $Y_j$  being a count response,  $\mathcal{F}_j$  is the collection of all subsets of  $\mathcal{Y}_j$  and  $m_j$  is the counting measure. Our above general assumption on the response types covers binary, polytomous, continuous, count responses, etc. We also allow the probability triples  $(\mathcal{Y}_j, \mathcal{F}_j, m_j)$  to differ across  $j = 1 \dots, J$ , that is, mixed-type responses (e.g., see [Moustaki and Knott, 2000](#)). We introduce the above abstract measure-theoretic definitions to (a) state the model and identifiability results in the most general setting, and also to (b) deal with discrete and continuous responses in a unified way.

Now we are ready to define the general-response CDMs. We start by specifying the latent part. To serve the diagnostic modeling purposes, we model each student's latent skill profile  $\mathbf{A}$  as a binary  $K$ -dimensional vector  $\mathbf{A} = (A_1, \dots, A_K) \in \{0, 1\}^K$ . Here  $A_k = 1$  or

0 represents the presence or absence of the  $k$ th latent skill. Following most existing studies on CDMs (Chen et al., 2015, 2018), we adopt the saturated model for the latent attributes; that is, for each skill profile  $\alpha \in \{0, 1\}^K$ , define its population proportion parameter as  $p_\alpha = \mathbb{P}(\mathbf{A} = \alpha)$ , which satisfy

$$\sum_{\alpha \in \{0, 1\}^K} p_\alpha = 1, \quad p_\alpha > 0. \quad (1)$$

We collect all proportion parameters in the vector  $\mathbf{p} = (p_\alpha : \alpha \in \{0, 1\}^K)$ .

Next, we define the measurement model, which specifies the conditional distribution of the observed responses  $\mathbf{Y}$  given the latent skills  $\mathbf{A}$ . The main ingredients in this definition are (a) the local independence assumption, and (b) conditional independence constraints induced by the  $\mathbf{Q}$ -matrix (Tatsuoka, 1983). Let  $\mathbb{P}_{j,\alpha}(Y_j | \mathbf{A} = \alpha)$  (or  $\mathbb{P}_{j,\alpha}$  for short) denote the conditional distribution of  $Y_j$  given  $\mathbf{A} = \alpha$ . First, under local independence, the observed  $Y_1, \dots, Y_J$  are conditionally independent given the latent  $A_1, \dots, A_K$ , which implies:

$$\mathbb{P}(\mathbf{Y} \in \times_{j=1}^J S_j | \mathbf{A} = \alpha) = \prod_{j=1}^J \mathbb{P}_{j,\alpha}(Y_j \in S_j | \mathbf{A} = \alpha), \quad \forall S_j \in \mathcal{F}_j, \quad j \in [J] \quad (2)$$

where the notation  $\mathbf{Y} \in \times_{j=1}^J S_j$  means  $Y_j \in S_j$  for all  $j$ , and  $S_j$  is any measurable set in  $\mathcal{F}_j$ .

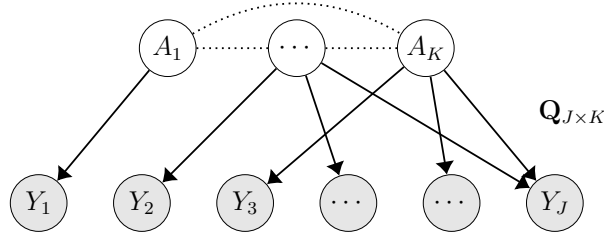


Figure 1: Graphical model of the general-response CDM with  $Y_j \in \mathcal{Y}_j$ . White nodes are latent attributes and grey nodes are observed responses. The directed arrows from the latent to the observed capture the conditional dependence of  $\mathbf{Y}$  given  $\mathbf{A}$ , which is exactly encoded in the  $\mathbf{Q}$ -matrix  $\mathbf{Q}_{J \times K}$ . There is an directed arrow from  $A_k$  to  $Y_j$  if and only if  $q_{j,k} = 1$ .

Next, we describe the constraints on the conditional distributions  $\mathbb{P}_{j,\alpha}$  imposed by the  $\mathbf{Q}$ -matrix. The  $\mathbf{Q}$ -matrix was initially proposed by Tatsuoka (1983) for modeling binary responses using the cognitive diagnostic assumption. The  $\mathbf{Q}$ -matrix  $\mathbf{Q} = (q_{j,k})$  is a  $J \times K$  matrix containing binary entries, where the  $(j, k)$ -th entry  $q_{j,k}$  is equal to 1 if item  $j$  requires or measures the latent attribute  $k$ , and 0 otherwise. Statistically, the entries in the  $\mathbf{Q}$ -matrix

describe how the  $J$  observed item responses depend on the latent attributes. These statistical dependencies can be represented by a directed graph pointing from the latent variables to the observed ones; see Figure 1 for an illustration. Following the convention of probabilistic directed graphical models (Koller and Friedman, 2009), we write  $\text{pa}(j) = \{k \in [K] : q_{j,k} = 1\} = \{k \in [K] : A_k \rightarrow Y_j\}$  as the set of *parent* latent attributes that have a directed arrow towards  $Y_j$  in the graph (or equivalently, the set of attributes required/measured by item  $j$ ). We also write  $\boldsymbol{\alpha}_{\text{pa}(j)}$  as the sub-vector of  $\boldsymbol{\alpha} = (\alpha_1, \dots, \alpha_K)$  that contains those entries indexed by  $k \in \text{pa}(j)$ . To make the  $\mathbf{Q}$ -matrix meaningfully describe the constraints between the latent and the observed variables, we assume that for each  $j \in [J]$ , there exist latent patterns  $\boldsymbol{\alpha}$  and  $\boldsymbol{\alpha}'$  with  $\boldsymbol{\alpha}_{\text{pa}(j)} \neq \boldsymbol{\alpha}'_{\text{pa}(j)}$  such that  $\mathbb{P}_{j,\boldsymbol{\alpha}} \neq \mathbb{P}_{j,\boldsymbol{\alpha}'}$ . According to the graphical model encoded by the  $\mathbf{Q}$ -matrix, the distribution  $\mathbb{P}_{j,\boldsymbol{\alpha}}(Y_j \in S_j \mid \mathbf{A} = \boldsymbol{\alpha}, \mathbf{Q})$  can be equivalently written as

$$\mathbb{P}_{j,\boldsymbol{\alpha}}(Y_j \in S_j \mid \mathbf{A} = \boldsymbol{\alpha}, \mathbf{Q}) = \mathbb{P}_{j,\boldsymbol{\alpha}}(Y_j \in S_j \mid \mathbf{A}_{\text{pa}(j)} = \boldsymbol{\alpha}_{\text{pa}(j)}), \quad \forall S_j \in \mathcal{F}_j; \quad (3)$$

that is, the conditional distribution of  $Y_j$  given all the latent attributes equals the conditional distribution of  $Y_j$  given those parent attributes required by item  $j$  as indicated by the  $\mathbf{Q}$ -matrix. In this work, we consider the confirmatory setting where the  $\mathbf{Q}$ -matrix is known.

Now, we formally define the general-response CDM based on the above assumptions.

**Definition 1** (General-response CDM). *A general-response CDM with  $K$  binary latent attributes  $\mathbf{A} = (A_1, \dots, A_K)$ ,  $J$  observed responses  $\mathbf{Y} = (Y_1, \dots, Y_J)$ , and model components  $(\mathbf{p}, \{\mathbb{P}_{j,\boldsymbol{\alpha}}\})$  is a statistical model for  $\mathbf{Y}$  that satisfies (1), (2), and (3).*

Under Definition 1, the marginal distribution of  $\mathbf{Y}$  in a general-response CDM is:

$$\mathbb{P}(\mathbf{Y} \in \times_{j=1}^J S_j \mid \mathbf{Q}) = \sum_{\boldsymbol{\alpha} \in \{0,1\}^K} p_{\boldsymbol{\alpha}} \prod_{j=1}^J \mathbb{P}_{j,\boldsymbol{\alpha}}(Y_j \in S_j \mid \mathbf{A}_{\text{pa}(j)} = \boldsymbol{\alpha}_{\text{pa}(j)}), \quad \forall S_j \in \mathcal{F}_j. \quad (4)$$

The proposed general-response CDMs form a very general and flexible framework of cognitive diagnostic models. *First*, this framework includes existing popular CDMs such as the DINA model, Reduced-RUM, LCDM, and GDM for categorical responses as submodels. This is because one can let the conditional distribution  $\mathbb{P}_{j,\boldsymbol{\alpha}}(Y_j \mid \mathbf{A} = \boldsymbol{\alpha})$  be a specific parametric distribution such as the Bernoulli with a certain link function for modeling binary responses

(elaborated in the next paragraph). *Second*, our framework also includes those recently proposed specific CDMs for continuous or count responses (e.g., [Minchen and de la Torre, 2018](#); [Liu et al., 2022](#)) as special cases, and is further able to model other response types.

As a concrete illustration, we next present the special case of our model (4) when the responses are all binary, which is the most widely considered scenario for CDMs. In this case,  $\mathcal{Y}_j = \{0, 1\}$  for all  $j \in [J]$  and  $\mathbb{P}_{j,\alpha}$  is just the Bernoulli distribution. We write the Bernoulli parameter of  $\mathbb{P}_{j,\alpha}$  as  $\theta_{j,\alpha}$ . Note that in the classical binary-response CDM literature,  $\theta_{j,\alpha}$  is often called the positive response probability. The conditional independence condition (3) boils down to the following equality constraints on the parameters  $\theta_{j,\alpha}$ :

$$\theta_{j,\alpha} = \theta_{j,\alpha'}, \quad \forall \alpha, \alpha' \in \{0, 1\}^K \text{ such that } \alpha_{\text{pa}(j)} = \alpha'_{\text{pa}(j)}. \quad (5)$$

The above constraints are satisfied in all existing CDMs for binary responses, and they also appeared as assumptions in the study of identifiability for the related restricted latent class models ([Xu, 2017](#); [Gu and Xu, 2020](#)). As a more concrete example,  $\theta_{j,\alpha}$  under the DINA model with the slipping parameter  $s_j$  and guessing parameter  $g_j$  can be written as

$$\theta_{j,\alpha} = (1 - s_j) \prod_{k=1}^K \alpha^{q_{j,k}} + g_j (1 - \prod_{k=1}^K \alpha^{q_{j,k}}) = (1 - s_j) \prod_{k \in \text{pa}(j)} \alpha_k + g_j (1 - \prod_{k \in \text{pa}(j)} \alpha_k),$$

where  $1 - s_j > g_j$  is typically assumed. This parametrization clearly satisfies (5).

## 2.2 Parametric and Exponential Family-based CDMs (ExpCDMs)

We next define parametric general-response CDMs, and illustrate how our general framework can be specified according to various modeling assumptions and response types. In this section, for notational simplicity, we additionally assume that the response types are the same across the items (i.e.,  $(\mathcal{Y}_j, \mathcal{F}_j, m_j)$  is the same for all  $j \in [J]$ ) and follow the same parametric family  $\mathcal{P} = \{g(\cdot; \boldsymbol{\eta})\}$ . Under this assumption, we can further write  $\mathbb{P}_{j,\alpha}$  as

$$\mathbb{P}_{j,\alpha}(Y_j \in S_j \mid \mathbf{A} = \alpha) = \int_{S_j} g(y; \boldsymbol{\eta}_{j,\alpha}) dm_j(y), \quad (6)$$

Here,  $g(\cdot; \boldsymbol{\eta})$  is the density/mass function of a specific parametric family  $\mathcal{P}$ , and  $\boldsymbol{\eta}_{j,\alpha}$ 's are parameters that can be a scalar or a vector. We next formally define a parametric CDM.

**Definition 2** (Parametric general-response CDM). *Given a parametric family  $\mathcal{P} = \{g(\cdot; \boldsymbol{\eta})\}$ , the parametric general-response CDM with parameters  $(\{\boldsymbol{\eta}_{j,\alpha}\}, \mathbf{p})$  is a general-response CDM whose  $\mathbb{P}_{j,\alpha}$  satisfies (6).*

For parametric general-response CDMs, the  $\mathbf{Q}$ -matrix constraints (3) reduce to the following constraints on the parameters  $\boldsymbol{\eta}$ , which are analogous to (5) in the binary-response case:

$$\boldsymbol{\eta}_{j,\alpha} = \boldsymbol{\eta}_{j,\alpha'}, \quad \forall \alpha, \alpha' \in \{0, 1\}^K \text{ such that } \alpha_{\text{pa}(j)} = \alpha'_{\text{pa}(j)}. \quad (7)$$

In the context of modeling response times in psychometrics, the Gamma (Maris, 1993), Weibull (Loeys et al., 2011), and inverse Gaussian (Lo and Andrews, 2015) distributions have been commonly used in addition to the aforementioned lognormal distribution. In the context of modeling visual fixation counts or correct answer counts, the negative binomial distribution is a popular choice (Man and Harring, 2019; Liu et al., 2023) in addition to the aforementioned Poisson distribution (Liu et al., 2022). In the later sections on estimation methodology and simulation studies, we will consider a variety of distributions for  $\mathcal{P}$ : Normal, transformed-Normal (lognormal, logistic-Normal), Poisson, and negative binomial.

Except for the negative binomial, all parametric distributions mentioned in the previous paragraph are *exponential family distributions* (e.g., see Section 3.4 in Casella and Berger, 2021), for which the probability density/mass function can be written as:

$$g(y_j; \boldsymbol{\eta}) = h(y_j) \exp\{\boldsymbol{\eta}^\top \mathbf{T}(y_j) - A(\boldsymbol{\eta})\}. \quad (8)$$

Following the convention for the exponential family distributions,  $\boldsymbol{\eta}$  collects the natural parameters,  $\mathbf{T}(Y_j)$  collects the sufficient statistics, and  $A(\boldsymbol{\eta})$  is the log-partition function. Both  $\boldsymbol{\eta}$  and  $\mathbf{T}(Y_j)$  are typically multidimensional in our general-response CDM context. Exponential families are a natural and flexible choice to model rich types of responses and are widely used in statistics and psychometrics. This motivates us to define parametric general-response CDMs based on the exponential families; we call these models *ExpCDMs*.

**Definition 3** (ExpCDM). *We define an ExpCDM as a parametric CDM whose parametric*

family  $\mathcal{P} = \{g(\cdot; \boldsymbol{\eta})\}_{\boldsymbol{\eta}}$  is an exponential family distribution.

The broad framework of ExpCDMs encompasses models for various response types including categorical, count, bounded and unbounded continuous data. In the end of this section, we will also briefly mention how to deal with even more general distributions outside of the exponential family, such as the negative binomial.

**Remark 1.** Using exponential family distributions to model general-response multivariate data with latent variables has a long history. [Moustaki and Knott \(2000\)](#) extended the IRT model to general responses, by modeling the natural parameter of the exponential family distribution as the linear combination of the continuous latent traits. This approach extends the generalized linear model (GLM) in [Nelder and Wedderburn \(1972\)](#) to the latent variable setting. In a similar spirit, [Dunson \(2000\)](#) used continuous latent variables and exponential family distributions to model multiple clustered outcomes. In the context of factor analysis, [Skron dal and Rabe-Hesketh \(2004\)](#) extended the linear factor analysis to model general responses by using exponential families and proposed the generalized linear factor model. We emphasize that all these previous models use continuous latent variables, and there do not exist works that adopt a framework as flexible as the exponential family with multidimensional discrete latent variables that serve the diagnostic purposes.

To define a general-response CDM, there are two parts that need to be specified: (a) the type of parametric family that models the response distribution, i.e., the forms of  $\mathcal{P}$  and  $g(\cdot; \boldsymbol{\eta})$ ; and (b) the type of interactions between the latent attributes and the observed responses, i.e., in what way  $\boldsymbol{\eta}_{j,\alpha}$  are subject to the  $\mathbf{Q}$ -matrix constraints, or equivalently, whether the parameters for  $Y_j$  are impacted by the main effects, or interaction effects, or both, of the required attributes. We have already shown that the ExpCDM framework provides a flexible way of specifying part (a). As for part (b), all the existing binary-response CDMs including the conjunctive DINA model, disjunctive DINO model, additive models (ACDM, reduced RUM), and more flexible all-effect models such as GDINA and GDM can be easily extended and incorporated into our ExpCDM framework.

We next show how all the above different types of attribute-item interactions have their ExpCDM counterparts for modeling rich types of data. In the remainder of this section, we describe the exponential family-based DINA (ExpDINA) model and the exponential family-

based additive CDM (ExpACDM). We define these models by further specifying how the natural parameters  $\boldsymbol{\eta}_{j,\boldsymbol{\alpha}}$  satisfy the  $\mathbf{Q}$ -matrix constraints in (7). Specifically, for each model, we present the lognormal distribution and the Poisson distribution as example exponential families to illustrate our general framework. These two distributions are suitable for modeling positive continuous data and nonnegative count data, respectively.

### 2.2.1 ExpDINA for General Responses

The DINA model was proposed by Junker and Sijtsma (2001) for modeling binary responses using a conjunctive assumption. Define the *ideal response* to item  $j$  given attribute pattern  $\mathbf{A} = \boldsymbol{\alpha}$  to be

$$\Gamma_{j,\boldsymbol{\alpha}} = \prod_{k=1}^K \alpha_k^{q_{j,k}} = \prod_{k \in \text{pa}(j)} \alpha_k. \quad (9)$$

In other words,  $\Gamma_{j,\boldsymbol{\alpha}}$  is the binary indicator of whether skill pattern  $\boldsymbol{\alpha}$  masters all of the required skills of item  $j$  and hence is “capable” of item  $j$ . Under the binary-response DINA model, the positive response probability  $\theta_{j,\boldsymbol{\alpha}}$  can be written as

$$\theta_{j,\boldsymbol{\alpha}} = \begin{cases} 1 - s_j =: \eta_{j,(1)}, & \text{if } \Gamma_{j,\boldsymbol{\alpha}} = 1, \\ g_j =: \eta_{j,(0)}, & \text{if } \Gamma_{j,\boldsymbol{\alpha}} = 0, \end{cases} \quad (10)$$

Following the conventional notation in the literature,  $s_j$  is the slipping parameter that gives the probability that a capable subject of item  $j$  provides a incorrect answer, and  $g_j$  is the guessing parameter that gives the probability that an incapable subject provides a correct answer. In (10), we reparametrize the item parameters as  $\eta_{j,(h)}$  for  $h = 1, 0$  with a slight abuse of notation, where  $h$  indicates the value of the ideal response  $\Gamma_{j,\boldsymbol{\alpha}}$ . We are now ready to define the general-response ExpDINA model.

**Definition 4** (ExpDINA). *The ExpDINA model is a submodel of the ExpCDM for which the natural parameters can be written as  $\boldsymbol{\eta}_{j,\boldsymbol{\alpha}} = \boldsymbol{\eta}_{j,(\Gamma_{j,\boldsymbol{\alpha}})}$  for  $j \in [J]$  and  $\boldsymbol{\alpha} \in \{0, 1\}^K$ . The parameters in an ExpDINA include the item parameters  $\{\boldsymbol{\eta}_{j,(0)}, \boldsymbol{\eta}_{j,(1)} : j \in [J]\}$  and proportion parameters  $\mathbf{p}$ .*

**Remark 2.** We remark that for certain distributions there may exist a more conventional parametrization than directly parametrizing the natural parameters to depend on the latent

attributes. For instance, in the ExpDINA with a Normal distribution for  $Y_j \mid \mathbf{A} = \boldsymbol{\alpha}$ , it is more intuitive to parameterize using the mean  $\mu$  and the variance  $\sigma^2$  as depending on  $\boldsymbol{\alpha}$  instead of assuming so for the natural parameters  $\boldsymbol{\eta} = (\mu/\sigma^2, -1/2\sigma^2)^\top$ . Such equivalent re-parametrizations still fit well into our definition of ExpCDMs.

Under an ExpDINA model, each natural parameter for each item  $j$  takes exactly two possible values, for capable ( $\Gamma_{j,\boldsymbol{\alpha}} = 1$ ) and incapable ( $\Gamma_{j,\boldsymbol{\alpha}} = 0$ ) subjects, respectively. Generally, ExpDINA still adopts the conjunctive assumption of required attributes and can be used to model rich types of response data. Under Definition 4, the conditional distribution  $Y_j \mid \mathbf{A} = \boldsymbol{\alpha}$  in (6) is:

$$Y_j \mid \mathbf{A} = \boldsymbol{\alpha} \sim \begin{cases} g(Y_j; \boldsymbol{\eta}_{j,(1)}), & \text{if } \Gamma_{j,\boldsymbol{\alpha}} = 1; \\ g(Y_j; \boldsymbol{\eta}_{j,(0)}), & \text{if } \Gamma_{j,\boldsymbol{\alpha}} = 0. \end{cases} \quad (11)$$

Therefore, ExpDINA essentially models each response  $Y_j$  as a local mixture of two distributions, similar in spirit to the binary-response DINA model. Indeed, ExpDINA covers the binary-response DINA and polytomous-response DINA as special cases. From the definition of  $\Gamma_{j,\boldsymbol{\alpha}}$ , it is clear that the  $\mathbf{Q}$ -matrix constraints (7) are satisfied.

**Example 1** (Lognormal-DINA, Logistic-Normal-DINA, and transformed-Normal-DINA). As an illustrative example, consider using the lognormal distribution to model positive continuous response. The lognormal is very commonly used to model response times (van der Linden, 2006, 2007; Minchen et al., 2017). The lognormal density with mean parameter  $\mu$  and variance parameter  $\sigma^2$  takes the form of

$$g^{\text{lognormal}}(y; \mu, \sigma^2) = \frac{1}{y\sqrt{2\pi\sigma^2}} \exp\left[-\frac{(\log y - \mu)^2}{2\sigma^2}\right], \quad y \in (0, \infty).$$

By specifying the mean  $\mu_{j,h}$  and variance  $\sigma_{j,h}^2$  for  $h = 1, 0$ , our lognormal-DINA model becomes the so-called C-DINA model proposed by Minchen et al. (2017):

$$Y_j \mid \mathbf{A} = \boldsymbol{\alpha} \sim \begin{cases} \text{lognormal}(\mu_{j,1}, \sigma_{j,1}^2), & \text{if } \Gamma_{j,\boldsymbol{\alpha}} = 1, \\ \text{lognormal}(\mu_{j,0}, \sigma_{j,0}^2), & \text{if } \Gamma_{j,\boldsymbol{\alpha}} = 0. \end{cases} \quad (12)$$

Note that under the same parameter values  $(\mu, \sigma^2)$ , the lognormal distribution is just an exponential transformation of the Normal distribution. Hence, the lognormal-DINA and normal-DINA models are equivalent up to an invertible transform of  $Y$ . Importantly, ExpCDMs based on *any monotone-increasing transformation* of the Normal are equivalent to the *Normal-CDMs*, that is, when  $Y_j \mid \mathbf{A} = \boldsymbol{\alpha}$  follows a Normal distribution. This observation can be used to build ExpCDMs for continuous responses that take values in a restricted range. For example, in addition to lognormal-CDM, we can define logistic-Normal-CDM by applying a logit transform of a Normal random variable to model bounded responses ranging in the interval  $(0, 1)$ . Here, the density function of the logistic-Normal( $\mu, \sigma^2$ ) is

$$g^{\text{logistic-Normal}}(y; \mu, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}y(1-y)} \exp \left[ -\frac{(\log(y/(1-y)) - \mu)^2}{2\sigma^2} \right], \quad y \in (0, 1).$$

The *logistic-Normal-DINA* model can be defined as:

$$Y_j \mid \mathbf{A} = \boldsymbol{\alpha} \sim \text{logistic-Normal}(\mu_{j,h}, \sigma_{j,h}^2), \quad h = \Gamma_{j,\boldsymbol{\alpha}}.$$

This newly proposed logistic-Normal-DINA model may be useful for analyzing continuous bounded responses in psychological or educational assessments.

**Example 2** (Poisson-DINA). As another example, we consider using the Poisson distribution to model count responses. The Poisson distribution is a canonical choice for modeling count responses, and was recently used in [Liu et al. \(2022\)](#) to model the number of correct answers in repetitive tasks under a diagnostic model. Poisson distribution with mean parameter  $\lambda$  has the probability mass function

$$g(y; \lambda) = \frac{e^{-\lambda} \lambda^y}{y!}, \quad y \in \mathbb{N},$$

where  $\mathbb{N}$  denotes the collection of all nonnegative integers. By defining the mean for  $h = 0, 1$  as  $\lambda_{j,h}$  (or equivalently defining  $\boldsymbol{\eta}_{j,(h)} = \log \lambda_{j,h}$ ), the distribution  $Y_j \mid \mathbf{A}$  can be written as

$$Y_j \mid \mathbf{A} = \boldsymbol{\alpha} \sim \begin{cases} \text{Poisson}(\lambda_{j,1}), & \text{if } \Gamma_{j,\boldsymbol{\alpha}} = 1, \\ \text{Poisson}(\lambda_{j,0}), & \text{if } \Gamma_{j,\boldsymbol{\alpha}} = 0. \end{cases} \quad (13)$$

This can be viewed as a reparameterization of the PDCM-DINA model in [Liu et al. \(2022\)](#).

### 2.2.2 ExpACDM for General Responses

We next define the *exponential family-based Additive CDMs* (ExpACDM) for general responses. The word *additive* refers to the assumption that the required skills of item  $j$  enter the conditional distribution of  $Y_j$  through an additive linear combination of the latent attributes  $A_1, \dots, A_K$ . Variants of such additive diagnostic models are very popular in the binary-response CDM literature. Examples include the Reduced Reparameterized Unified Model (R-RUM; [DiBello et al., 1995](#)), the Additive Cognitive Diagnostic Model (ACDM; [de la Torre, 2011](#)), and the Logistic Linear Model (LLM; [Maris, 1999](#)).

We continue to work in the general framework of general-response CDMs in (6). Under the additive assumption, we write the parameters  $\boldsymbol{\eta}_{j,\alpha}$  as

$$\boldsymbol{\eta}_{j,\alpha} = \mathbf{h} \left( \beta_{j,0} + \sum_{k=1}^K \beta_{j,k} q_{j,k} \alpha_k, \gamma_j \right) = \mathbf{h} \left( \beta_{j,0} + \sum_{k \in \text{pa}(j)} \beta_{j,k} \alpha_k, \gamma_j \right). \quad (14)$$

We explain the notations in the above expression in turn. First, similar to binary-response additive CDMs, the corresponding  $\beta_{j,k}$  are not needed to define the model when  $q_{j,k} = 0$ . So without the loss of generality, we assume  $\beta_{j,k} = 0$  if  $q_{j,k} = 0$ . If  $q_{j,k} = 1$ , then  $\beta_{j,k}$  is the main-effect coefficient for the  $k$ th latent attribute. One can see that the linear combination  $\beta_{j,0} + \sum_{k=1}^K \beta_{j,k} q_{j,k} \alpha_k$  depends only on those required attributes of item  $j$ , so  $\boldsymbol{\eta}_{j,\alpha}$  defined by (14) satisfies the  $\mathbf{Q}$ -matrix constraints (7). The  $\gamma_j$  in (14) denotes potential additional model parameters that do not depend on the latent attributes (e.g.,  $\gamma_j$  maybe be the dispersion parameter if an exponential family distribution is used), and  $\mathbf{h}$  in (14) is a link function that plays a very similar role as the link function in the generalized linear models (e.g., [Nelder and Wedderburn, 1972](#)). Here,  $\mathbf{h}$  is introduced to map the linear combination of the required attributes to the natural parameters  $\boldsymbol{\eta}$ , as the natural parameter space may be different from the space of the linear combinations. Note that  $\boldsymbol{\eta}$  may be multidimensional, and hence  $\mathbf{h}$  can be a mapping between multidimensional spaces.

With these notations at hand, we next formally define ExpACDMs.

**Definition 5** (ExpACDM). *An ExpACDM is a submodel of the ExpCDM for which the*

natural parameters  $\boldsymbol{\eta}_{j,\boldsymbol{\alpha}}$  satisfy (7). The parameters in an ExpACDM with a link function  $\mathbf{h}$  include  $\{\beta_{j,0} : j \in [J]\}$ ,  $\{\beta_{j,k} : q_{j,k} = 1\}$ ,  $\{\gamma_j : j \in [J]\}$  and proportion parameters  $\mathbf{p}$ .

Note that for the ExpDINA model defined earlier in Section 2.2.1, we have parameterized the item parameters as  $(\boldsymbol{\eta}_{j,(0)}, \boldsymbol{\eta}_{j,(1)})$ , because there are only two possible values of  $\{\boldsymbol{\eta}_{j,\boldsymbol{\alpha}} : \boldsymbol{\alpha} \in \{0,1\}^K\}$  due to the conjunctive modeling assumption. In contrast, the ExpACDM has more possible values of  $\{\boldsymbol{\eta}_{j,\boldsymbol{\alpha}}\}$  when the attribute profile  $\boldsymbol{\alpha}$  ranges in  $\{0,1\}^K$ . So in Definition 5, we have parameterized ExpACDMs using the  $\beta$ - and  $\gamma$ -coefficients.

**Example 3** (lognormal-ACDM). Continuing from Example 1, we consider the lognormal distribution for modeling the responses but now with an additive structure of the attributes. For the lognormal parameters  $(\mu_{j,\boldsymbol{\alpha}}, \sigma_{j,\boldsymbol{\alpha}}^2)$ , one can choose to model  $\mu_{j,\boldsymbol{\alpha}}$  as additive in  $\alpha_k$ 's and  $\sigma_{j,\boldsymbol{\alpha}}^2$  to not depend on  $\boldsymbol{\alpha}$ :

$$\mu_{j,\boldsymbol{\alpha}} = \beta_{j,0} + \sum_{k=1}^K \beta_{j,k} q_{j,k} \alpha_k, \quad \sigma_{j,\boldsymbol{\alpha}}^2 = \gamma_j.$$

Here, we can view the link function  $\mathbf{h}$  as  $\mathbf{h}(\mu, \sigma^2) = (\mu/\sigma^2, -1/2\sigma^2)$  that maps the mean and variance to the natural parameters under the lognormal distribution. By plugging this parametrization in (6),  $Y_j \mid \mathbf{A}$  can be written as

$$Y_j \mid \mathbf{A} = \boldsymbol{\alpha} \sim \text{lognormal} \left( \beta_{j,0} + \sum_{k=1}^K \beta_{j,k} q_{j,k} \alpha_k, \gamma_j \right). \quad (15)$$

Note that there are multiple modeling choices regarding how to specify the dependence of  $\mu_{j,\boldsymbol{\alpha}}$  and  $\sigma_{j,\boldsymbol{\alpha}}^2$  on the linear (i.e., additive) combinations of  $\alpha_k$ 's. One could model a additive mean and a constant variance after the log transformation as done in the above (15). Alternatively, one could even model the variance also as additive in  $\alpha_k$ 's.

**Example 4** (Poisson-ACDM). As an counterpart of the Poisson-DINA model in Example 2, we define an ExpACDM that uses the Poisson distribution to model count data. Since Poisson only has one rate parameter, there is no need to introduce  $\gamma_j$  and we only need to express the rate parameter as the linear combination of those required attributes:

$$\lambda_{j,\boldsymbol{\alpha}} = \beta_{j,0} + \sum_{k=1}^K \beta_{j,k} q_{j,k} \alpha_k.$$

Recall that the natural parameter under Poisson is  $\boldsymbol{\eta} = \log \lambda$ , so we define the link function  $\mathbf{h}$  as  $\mathbf{h}(\lambda) = \log \lambda$ . We additionally assume  $\beta_{j,k} \geq 0$  to ensure that the rate parameters  $\lambda_{j,\alpha}$  are nonnegative. By plugging the above expression in (6),  $Y_j \mid \mathbf{A}$  can be written as

$$Y_j \mid \mathbf{A} = \boldsymbol{\alpha} \sim \text{Poisson} \left( \beta_{j,0} + \sum_{k=1}^K \beta_{j,k} q_{j,k} \alpha_k \right).$$

Alternatively, one could let  $\mathbf{h}$  to be the identity link  $\mathbf{h}(\lambda) = \lambda$  without assuming  $\beta_{j,k} \geq 0$ .

In Supplementary Material S.1, we define the exponential family-based general diagnostic model (ExpGDM) for general responses, and also discuss how to define the general-response DINA and ACDM for distributions outside of the exponential family, such as the negative binomial distribution. In this section, we have focused on cases where the parametric families  $\mathcal{P}$  do not change across different items  $j \in [J]$ . But it is actually straightforward to use our framework to model different distribution families for different items.

## 3 Identifiability of General-response CDMs

### 3.1 Strict Identifiability of General-response CDMs

Model identifiability is a fundamental prerequisite for valid statistical estimation and inference. We will prove that our general-response CDMs are identifiable under transparent conditions on the  $\mathbf{Q}$ -matrix. In this subsection, we first define strict identifiability for general-response CDMs, and then propose conditions for strict identifiability.

Recall that  $(\mathbf{p}, \{\mathbb{P}_{j,\alpha} : j \in [J], \alpha \in \{0,1\}^K\})$  can be viewed as the model parameters of a general-response CDM. We will say that  $(\mathbf{p}, \{\mathbb{P}_{j,\alpha}\})$  is equal to  $(\bar{\mathbf{p}}, \{\bar{\mathbb{P}}_{j,\alpha}\})$  up to a sign flip for each coordinate of  $\boldsymbol{\alpha}$ , if there exist permutation maps  $\tau_k : \{0,1\} \rightarrow \{0,1\}$  such that  $p_{\alpha} = \bar{p}_{(\tau_1(\alpha_1), \dots, \tau_K(\alpha_K))}$  and  $\mathbb{P}_{j,\alpha} = \bar{\mathbb{P}}_{j,(\tau_1(\alpha_1), \dots, \tau_K(\alpha_K))}$  for any  $j \in [J]$  and  $\boldsymbol{\alpha} \in \{0,1\}^K$ . The sign flipping issue of the binary latent variables in Definition 6 is inevitable for general-response CDMs under the minimal assumptions that we make on the response types. Nevertheless, sign flipping is a trivial ambiguity and can be easily fixed in parametric submodels including most ExpCDMs. For example, in traditional CDMs for binary responses, the sign of latent variables is fixed by making the monotonicity assumption that students who possess all

required skills of an item are more likely to correctly answer it compared to those who lack some required skills. Similarly, to resolve the sign flipping issue, in the lognormal-DINA model for continuous responses, one can assume that the students who possess all skills required by an item have a larger mean parameter in the lognormal distribution than those who do not:  $\mu_{j,1} > \mu_{j,0}$ . More generally, for ExpACDMs, assuming  $\beta_{j,k} > 0$  when  $q_{j,k} = 1$  resolves the sign flipping issue. This claim is formally presented in the later Proposition 1. However, it is less convincing to assume monotonicity for every response type, as not all responses increase with more skill mastery, and moreover, an ordering may not be defined on the sample space of the response  $\mathcal{Y}_j$  if it is not a subset of the real line  $\mathbb{R}$ . We next formally define strict identifiability up to the sign flipping.

**Definition 6** (Strict identifiability). *Consider a general-response CDM with a known  $\mathbf{Q}$ -matrix. The model is strictly identifiable if for any parameters  $(\mathbf{p}, \{\mathbb{P}_{j,\alpha}\})$  and  $(\bar{\mathbf{p}}, \{\bar{\mathbb{P}}_{j,\alpha}\})$  satisfying (3), the following equations hold if and only if  $(\mathbf{p}, \{\mathbb{P}_{j,\alpha}\})$  is equal to  $(\bar{\mathbf{p}}, \{\bar{\mathbb{P}}_{j,\alpha}\})$  up to a sign flip for each coordinate of  $\alpha$ :*

$$\mathbb{P}(\mathbf{Y} \in \times_{j=1}^J S_j \mid \bar{\mathbf{p}}, \bar{\mathbb{P}}, \mathbf{Q}) = \mathbb{P}(\mathbf{Y} \in \times_{j=1}^J S_j \mid \mathbf{p}, \mathbb{P}, \mathbf{Q}), \quad \forall S_j \in \mathcal{F}_j. \quad (16)$$

Now we are ready to state the main theorem on strict identifiability.

**Theorem 1.** *Under the general-response CDM, the model components  $(\mathbf{p}, \{\mathbb{P}_{j,\alpha}\})$  are strictly identifiable if the following conditions hold.*

- A. *The true  $\mathbf{Q}$ -matrix contains two identity submatrices  $\mathbf{I}_K$  after row swapping, i.e.,  $\mathbf{Q}$  can be written as*

$$\mathbf{Q} = [\mathbf{I}_K, \mathbf{I}_K, \mathbf{Q}^{*\top}]^\top.$$

- B. *Suppose that the  $\mathbf{Q}$ -matrix is written as in A. For any  $\alpha \neq \alpha'$ , there exists  $j > 2K$  such that  $\mathbb{P}_{j,\alpha} \neq \mathbb{P}_{j,\alpha'}$ .*

The conditions in Theorem 1 are similar in spirit to existing identifiability conditions for CDMs with binary or polytomous responses, such as Xu (2017) and Culpepper (2019). When considering unidimensional responses  $Y_j$ 's for each item, we can additionally make a monotonicity assumption to avoid the sign flipping issue. The following proposition estab-

lishes the strongest possible notion of identifiability under a monotonicity assumption (17). Here,  $\mathbf{0}_K$  and  $\mathbf{1}_K$  denote the all-zero and all-one vectors of length  $K$ , respectively.

**Proposition 1.** *Suppose that a general-response CDM satisfies conditions A and B. Without loss of generality, suppose the first  $K$  rows in  $\mathbf{Q}$  form an identity matrix  $\mathbf{I}_K$ . Additionally, suppose that  $\mathcal{Y}_j \subseteq \mathbb{R}$ , and the conditional distributions  $\{\mathbb{P}_{j,\alpha}\}$  satisfy the following monotonicity assumption:*

$$\mathbb{E}(Y_j \mid \mathbf{A} = \mathbf{0}_K) < \mathbb{E}(Y_j \mid \mathbf{A} = \mathbf{1}_K), \quad (17)$$

for  $1 \leq j \leq K$ . Then, the model components  $(\mathbf{p}, \{\mathbb{P}_{j,\alpha}\})$  are strictly identifiable with no sign flipping issues. In other words, (16) implies  $(\mathbf{p}, \{\mathbb{P}_{j,\alpha}\}) = (\bar{\mathbf{p}}, \{\bar{\mathbb{P}}_{j,\alpha}\})$ .

The proofs of all theoretical results are deferred to the Supplementary Material. We next provide the high-level intuition of our proof argument. The main idea is to discretize all responses by first constructing a partition of each sample space  $\mathcal{Y}_j$  into a finite number of categories, and then defining surrogate categorical variables based on the partitions. In this case, our model in (4) implies a tensor decomposition for the probability mass function of the surrogate categorical variables. We then leverage Kruskal’s Theorem (Kruskal, 1977) to establish uniqueness of this tensor decomposition under the  $\mathbf{Q}$ -matrix conditions. Finally, we link such uniqueness results back to identifiability of parameters based on the original distribution of the general responses  $Y_j \in \mathcal{Y}_j$ . Our proof strategy is motivated by Theorem 8 in Allman et al. (2009) which established identifiability of mixtures of products of nonparametric probability densities. However, Allman et al. (2009)’s result cannot be directly applied to our setup as we consider more general response types and our conditional distributions are subject to  $\mathbf{Q}$ -matrix constraints.

Theorem 1 and Proposition 1 hold true under minimal assumptions, without assuming a specific measurement model such as DINA or ACDM. For binary responses, there is an extensive literature on model identifiability under various assumptions (e.g. Chen et al., 2015; Xu, 2017; Xu and Shang, 2018; Chen et al., 2020; Gu and Xu, 2021). It is worth noting that when not assuming a specific measurement model, our results in Theorem 1 and Proposition 1 nearly match existing weakest identifiability conditions for categorical-response CDMs. For example, Theorem 1 in Xu (2017) stated that the parameters  $(\mathbf{p}, \boldsymbol{\Theta})$  in a restricted latent class model are strictly identifiable under condition A and the following condition B1:

B1. For any  $k \in [K]$ , there exists one  $j > 2K$  such that  $\theta_{j,\mathbf{e}_k} \neq \theta_{j,\mathbf{0}_K}$ .

Here,  $\Theta = (\theta_{j,\alpha})_{j \in [J], \alpha \in \{0,1\}^K}$  is the positive response probability matrix with  $\theta_{j,\alpha} = \mathbb{P}_{j,\alpha}(Y_j = 1 \mid \mathbf{A} = \alpha)$ , and this determines the  $\mathbb{P}_{j,\alpha}$ 's, as mentioned in Section 2. Also,  $\mathbf{e}_k$  is the standard basis vector whose  $k$ th entry is 1 and other entries are zeros. For exploratory CDMs with polytomous responses, Theorem 2 in Culpepper (2019) proved that the model parameters are identifiable under conditions A, B, and an additional condition C:

C. Each item  $j \in [J]$  has distinct item response functions for at least two latent classes.

For general-response CDMs, the above additional assumption is equivalent to “for all  $j \in [J]$ , there exist skill patterns  $\alpha \neq \alpha'$  such that  $\mathbb{P}_{j,\alpha} \neq \mathbb{P}_{j,\alpha'}$ ”. Our Theorem 1 does not impose such an assumption for all items, but only implicitly imposes it for  $j = 1, \dots, 2K$  (because the first  $2K$  rows in  $\mathbf{Q}$  form two copies of the identity matrix  $\mathbf{I}_K$ ). As for  $j > 2K$ , we do not require there exist  $\alpha \neq \alpha'$  such that  $\mathbb{P}_{j,\alpha} \neq \mathbb{P}_{j,\alpha'}$  for each  $j$ . In fact, Theorem 1 allows some  $Y_j$  for  $j > 2K$  to not depend on any attributes; or equivalently, the corresponding row vector  $\mathbf{q}_j$  in the  $\mathbf{Q}$ -matrix is an all-zero row vector. Such all-zero row vectors can be absorbed into the submatrix  $\mathbf{Q}^*$  in condition A of Theorem 1.

Finally, in the following propositions, we propose variations of Theorem 1 by modifying condition B to more easily checkable ones. In Proposition 2, we replace condition B in Theorem 1 by a stronger, but more intuitive condition. This condition has been previously proposed to establish identifiability of exploratory diagnostic models for categorical responses (Fang et al., 2019; Gu and Dunson, 2023).

**Proposition 2.** *Condition B in Theorem 1 holds when  $\mathbf{Q}^*$  contains an identity submatrix  $\mathbf{I}_K$ . Hence, the general-response CDM is strictly identifiable when it vertically stacks three identity submatrices  $\mathbf{I}_K$  after some row swapping.*

In the next proposition, we consider the ExpACDM with parameters  $(\mathbf{p}, \boldsymbol{\beta}, \gamma, \mathbf{Q})$ , instead of the general-response CDM, and show how condition B can be simplified.

**Proposition 3.** *For the ExpACDM, condition B in Theorem 1 reduces to:*

B2. *For any  $\alpha \neq \alpha' \in \{0,1\}^K$ , there exists  $j > 2K$  such that  $\sum_{k=1}^K q_{j,k} \beta_{j,k} (\alpha_k - \alpha'_k) \neq 0$ .*

*Hence, the ExpACDM is strictly identifiable when the  $\mathbf{Q}$ -matrix satisfies condition A, and the main-effect coefficients  $\boldsymbol{\beta}$  satisfies condition B2.*

### 3.2 Generic Identifiability of ExpACDMs

Although strict identifiability is the strongest possible identifiability notion, in practice it may impose too stringent conditions on the  $\mathbf{Q}$ -matrix. For instance, the  $\mathbf{Q}$ -matrix for the TIMSS 2019 response time dataset analyzed in Section 6 requires one content skill and one cognitive skill for each item (see Table 1 and related discussion there). This  $\mathbf{Q}$ -matrix does not contain any  $K$ -dimensional standard basis vectors as row vectors, and hence does not contain an identity submatrix  $\mathbf{I}_K$ . In this subsection, we study generic identifiability, which is a slightly weaker notion than strict identifiability proposed by Allman et al. (2009). Generic identifiability requires that the model parameters are identifiable almost everywhere in the parameter space, in the sense that they are identifiable except in a Lebesgue measure-zero subset of the parameter space. Existing studies such as Gu and Xu (2020) and Chen et al. (2020) proposed practical generic identifiability conditions for binary-response CDMs that contain main effects of latent attributes. Those conditions are much weaker than the strict identifiability conditions.

To study generic identifiability, we next focus on parametric forms of  $\mathbb{P}_{j,\alpha}$  and especially consider ExpACDMs that model the main effects of latent attributes. For notational simplicity, we let  $\boldsymbol{\beta}_{\text{main}} = \{\beta_{j,k}\}_{k \in [J], k \in [K]}$  denote all main-effect coefficients, and  $\boldsymbol{\beta} = \boldsymbol{\beta}_{\text{main}} \cup \{\beta_{j,0}\}_{j \in [J]}$  denote the collection of all intercepts and main-effect coefficients. Before defining the concept of generic identifiability, we define the true parameter space  $\Omega(\boldsymbol{\beta}_{\text{main}}; \mathbf{Q})$  for  $\boldsymbol{\beta}_{\text{main}}$  as follows:

$$\Omega(\boldsymbol{\beta}_{\text{main}}; \mathbf{Q}) = \{\boldsymbol{\beta}_{\text{main}} : \beta_{j,k} = 0 \quad \text{if} \quad q_{j,k} = 0\}. \quad (18)$$

Here, we are using the previous assumption that  $\beta_{j,k} = 0$  if  $q_{j,k} = 0$  (see the text following (14)). Recall that in an ExpACDM,  $\gamma$  denotes the dispersion parameters of the exponential family and  $\mathbf{p}$  denotes the proportion parameters of the latent attribute patterns. We define the joint parameter space of all parameters  $\mathbf{p}$ ,  $\boldsymbol{\beta}$ , and  $\gamma$  by

$$\Omega(\mathbf{p}, \boldsymbol{\beta}, \gamma; \mathbf{Q}) = \{(\mathbf{p}, \boldsymbol{\beta}, \gamma) : \sum_{\alpha \in \{0,1\}^K} p_{\alpha} = 1, \quad p_{\alpha} \geq 0, \quad \boldsymbol{\beta}_{\text{main}} \in \Omega(\boldsymbol{\beta}_{\text{main}}; \mathbf{Q})\}. \quad (19)$$

In order to define generic identifiability without the sign flipping issue, we next adopt

the monotonicity assumption in (17). In particular, we assume that  $\beta_{j,k} > 0$  when  $q_{j,k} = 1$ , which means that the main effects of the required attributes are all positive. Here, the actual sign of  $\beta_{j,k}$  is not important and one may assume  $\beta_{j,k} < 0$  for all  $j$  and  $k$  instead to achieve identifiability. Now we are ready to define generic identifiability for ExpACDMs.

**Definition 7.** *An ExpACDM with parameters  $(\mathbf{p}, \boldsymbol{\beta}, \gamma)$  is generically identifiable if the following set has measure zero with respect to the Lebesgue measure on  $\Omega(\mathbf{p}, \boldsymbol{\beta}, \gamma; \mathbf{Q})$ :*

$$\{(\mathbf{p}, \boldsymbol{\beta}, \gamma) \in \Omega(\mathbf{p}, \boldsymbol{\beta}, \gamma; \mathbf{Q}) : \text{there exist alternative parameters } (\bar{\mathbf{p}}, \bar{\boldsymbol{\beta}}, \bar{\gamma}) \neq (\mathbf{p}, \boldsymbol{\beta}, \gamma) \text{ in } \Omega(\mathbf{p}, \boldsymbol{\beta}, \gamma; \mathbf{Q}) \text{ such that } \mathbb{P}(\mathbf{Y} \mid \mathbf{p}, \boldsymbol{\beta}, \gamma) = \mathbb{P}(\mathbf{Y} \mid \bar{\mathbf{p}}, \bar{\boldsymbol{\beta}}, \bar{\gamma})\}.$$

We next propose generic identifiability conditions for ExpACDMs that are substantially weaker than the strict identifiability conditions in Section 3.1.

**Theorem 2.** *Consider an ExpACDM. Assume that  $\mathbf{h}$  is an analytic function, and that the true parameters  $(\mathbf{p}, \boldsymbol{\beta}, \gamma)$  lie in  $\Omega(\mathbf{p}, \boldsymbol{\beta}, \gamma; \mathbf{Q})$ . Then, the model parameters  $(\mathbf{p}, \boldsymbol{\beta}, \gamma)$  are generically identifiable if the following conditions on the  $\mathbf{Q}$ -matrix hold.*

*A\*. The true  $\mathbf{Q}$ -matrix can be written as  $\mathbf{Q} = [\mathbf{Q}_1^\top, \mathbf{Q}_2^\top, \mathbf{Q}^{*\top}]^\top$  after some row permutation, where  $\mathbf{Q}_1, \mathbf{Q}_2$ 's are  $K \times K$  matrices such that all diagonal entries are equal to one. In other words, we can write*

$$\mathbf{Q}_i = \begin{pmatrix} 1 & * & \cdots & * \\ * & 1 & \cdots & * \\ \vdots & \vdots & \ddots & \vdots \\ * & * & \cdots & 1 \end{pmatrix}$$

*for  $i = 1, 2$ . Here,  $*$  indicates an arbitrary value in  $\{0, 1\}$ .*

*B\*. Suppose that the  $\mathbf{Q}$ -matrix is written as in A\*, and  $\mathbf{Q}^*$  does not have any all-zero columns. In other words, for any  $k$ , there exists  $j > 2K$  such that  $q_{j,k} = 1$ .*

Conditions in Theorem 2 only depend on the  $\mathbf{Q}$ -matrix but not on other model components. Theorem 2 is motivated by Gu and Xu (2020) and Chen et al. (2020), which are the first studies that considered generic identifiability for CDMs. Theorem 4.3 in Gu and

Xu (2020) and Theorem 1 in Chen et al. (2020) showed that binary-response CDMs are generically identifiable under conditions  $A^*$  and  $B^*$ .

Note that Theorem 2 holds without assuming (1) that  $p_\alpha > 0$  for all  $\alpha \in \{0, 1\}^K$ . This is because  $\{\mathbf{p} : \exists \alpha \text{ such that } p_\alpha = 0\}$  is a measure-zero subset of  $\{\mathbf{p} : \sum_\alpha p_\alpha = 1, p_\alpha \geq 0\}$ , and hence relaxing the condition from  $p_\alpha > 0$  to  $p_\alpha \geq 0$  does not violate generic identifiability. In addition, we point out that the exponential family assumption for  $\mathcal{P}$  is not crucial for generic identifiability to hold. In fact,  $\mathcal{P}$  can be any parametric family as long as the model parameters  $\boldsymbol{\eta}_{j,\alpha}$  satisfy the additive assumption in (14). Hence, the conditions in Theorem 2 also guarantee the generic identifiability of ACDMs with non-exponential family distributions; see more discussions in Remark S.1 in the Supplementary Material S.2.

Theorem 2 may be extended to the setting of ExpGDM and other all-effect CDMs for general responses. This is because these models include the main effects of the latent attributes, which are the key components underlying generic identifiability. We leave the rigorous statement of generic identifiability of ExpGDMs as future work. On the other hand, Theorem 2 does not apply to the DINA model, because the DINA model is conjunctive and does not include the main effects of the latent attributes. Under the DINA model for general responses, we conjecture that another identifiability notion similar to the  $\mathbf{p}$ -partial identifiability in Gu and Xu (2020) for conventional categorical-response CDMs can be studied.

## 4 Universal EM Algorithms for Estimating ExpCDMs

Our identifiability result in Theorem 1 has a nice consequence of statistical consistency of the maximum likelihood estimator (MLE). We next formally establish this result. Consider a sample with  $N$  independent and identically distributed response vectors  $\mathbf{Y}_{1:N} = \{\mathbf{Y}_1, \dots, \mathbf{Y}_N\}$  from an ExpCDM in (6) with true parameters  $(\boldsymbol{\eta}_0, \mathbf{p}_0)$ . Given the  $\mathbf{Q}$ -matrix, the marginal log-likelihood can be written as

$$\ell(\boldsymbol{\eta}, \mathbf{p} \mid \mathbf{Y}_{1:N}, \mathbf{Q}) = \sum_{i=1}^N \log \mathbb{P}(\mathbf{Y} = \mathbf{Y}_i \mid \boldsymbol{\eta}, \mathbf{p}) = \sum_{i=1}^N \log \left( \sum_{\alpha \in \{0,1\}^K} p_\alpha \prod_{j=1}^J g(Y_{i,j}; \boldsymbol{\eta}_{j,\alpha}) \right), \quad (20)$$

where  $\boldsymbol{\eta}$  is the collection of all  $\boldsymbol{\eta}_{j,\alpha}$ 's that are subject to the  $\mathbf{Q}$ -matrix constraints under the considered ExpCDM. Define the MLE of model parameters as

$$(\hat{\boldsymbol{\eta}}, \hat{\mathbf{p}}) = \underset{\boldsymbol{\eta}, \mathbf{p}}{\operatorname{argmax}} \ell(\boldsymbol{\eta}, \mathbf{p} \mid \mathbf{Y}_{1:N}).$$

Following from the identifiability conclusions, we have the following proposition on parameter estimation consistency.

**Proposition 4** (Parameter Estimation Consistency). *Consider an ExpCDM in the form (6) where the true parameters  $(\boldsymbol{\eta}_0, \mathbf{p}_0)$  lie in the interior of the parameter space. Suppose the identifiability conditions A and B in Theorem 1 hold. Then  $(\hat{\boldsymbol{\eta}}, \hat{\mathbf{p}})$  converge to  $(\boldsymbol{\eta}_0, \mathbf{p}_0)$  almost surely as  $N \rightarrow \infty$ .*

Next we propose an expectation-maximization (EM, [Dempster et al., 1977](#)) algorithm for parameter estimation. For the purpose of developing an EM algorithm, consider the log-likelihood for the complete data  $(\mathbf{Y}_{1:N}, \mathbf{A}_{1:N}) = (\mathbf{Y}_i, \mathbf{A}_i)_{i=1, \dots, N}$ :

$$\begin{aligned} \ell_c(\boldsymbol{\eta}, \mathbf{p} \mid \mathbf{Y}_{1:N}, \mathbf{A}_{1:N}, \mathbf{Q}) &= \log \left[ \prod_{i=1}^N \mathbb{P}(\mathbf{A}_i) \mathbb{P}(\mathbf{Y}_i \mid \mathbf{A}_i, \mathbf{Q}) \right] \\ &= \sum_{\boldsymbol{\alpha} \in \{0,1\}^K} \sum_{i=1}^N \mathbb{1}(\mathbf{A}_i = \boldsymbol{\alpha}) \left( \log p_{\boldsymbol{\alpha}} + \sum_{j=1}^J \log g(Y_{i,j}; \boldsymbol{\eta}_{j,\boldsymbol{\alpha}}) \right), \end{aligned} \quad (21)$$

where  $\mathbf{A}_i \in \{0,1\}^K$  denotes the latent attribute profile of the  $i$ th subject in the sample. In the following Section 4.1 and Section 4.2, we present details of the EM algorithms for the general ExpDINA and ExpACDM models.

## 4.1 EM Algorithm for the ExpDINA

Consider the ExpDINA model defined in Section 2.2.1, which is parametrized by item parameters  $\{\boldsymbol{\eta}_{j,(h)}\}_{j \in [J], h=0,1}$  and  $\mathbf{p}$ . In this subsection, for notational simplicity, we write  $\boldsymbol{\eta} = \{\boldsymbol{\eta}_{j,(h)}\}_{j \in [J], h=0,1}$  to denote all item parameters. For the ExpDINA model, the complete data log-likelihood (21) can be rewritten as:

$$\ell_c^{\text{ExpDINA}}(\boldsymbol{\eta}, \mathbf{p} \mid \mathbf{Y}_{1:N}, \mathbf{A}_{1:N}) = \sum_{\boldsymbol{\alpha} \in \{0,1\}^K} \sum_{i=1}^N \mathbb{1}(\mathbf{A}_i = \boldsymbol{\alpha}) \log p_{\boldsymbol{\alpha}} \quad (22)$$

$$\begin{aligned}
& + \sum_{\alpha \in \{0,1\}^K} \sum_{i=1}^N \mathbb{1}(\mathbf{A}_i = \alpha) \sum_{j=1}^J \left[ (\boldsymbol{\eta}_{j,(0)}^\top \mathbf{T}(Y_{i,j}) - A(\boldsymbol{\eta}_{j,(0)}) + \log h(Y_{i,j})) (1 - \Gamma_{j,\alpha}) \right. \\
& \left. + (\boldsymbol{\eta}_{j,(1)}^\top \mathbf{T}(Y_{i,j}) - A(\boldsymbol{\eta}_{j,(1)}) + \log h(Y_{i,j})) \Gamma_{j,\alpha} \right].
\end{aligned}$$

Recall that  $\Gamma_{j,\alpha}$  is the binary ideal response, which equals one if and only if the latent skill pattern  $\alpha$  possesses all skills required by item  $j$ . We next present details of the E step and M step in the EM algorithm.

**E step for ExpDINA.** In the E step of the  $t$ -th EM iteration, we calculate the conditional expectation of  $\ell_c^{\text{ExpDINA}}$  in (22) given the current parameter values  $\boldsymbol{\eta}^{(t)}, \mathbf{p}^{(t)}$ . As  $\mathbf{A}_i$  is discrete and ranges in  $\{0,1\}^K$ , it suffices to evaluate  $\varphi_{i,\alpha}^{(t+1)} := \mathbb{P}(\mathbf{A}_i = \alpha \mid \boldsymbol{\eta}^{(t)}, \mathbf{p}^{(t)})$  for all  $\alpha \in \{0,1\}^K$ . This can be calculated by noting that

$$\varphi_{i,\alpha}^{(t+1)} = p_\alpha \prod_{j=1}^J \mathbb{P}(Y_{i,j} \mid \boldsymbol{\eta}^{(t)}, \mathbf{p}^{(t)}, \mathbf{A}_i = \alpha) / \left( \sum_{\alpha' \in \{0,1\}^K} p_{\alpha'} \prod_{j=1}^J \mathbb{P}(Y_{i,j} \mid \boldsymbol{\eta}^{(t)}, \mathbf{p}^{(t)}, \mathbf{A}_i = \alpha') \right).$$

The exact form of  $\varphi_{i,\alpha}^{(t+1)}$  is given in Algorithm 1. Now using  $\varphi_{i,\alpha}^{(t+1)}$ , we can calculate the conditional expectation of  $\ell_c^{\text{ExpDINA}}$  as follows and denote it as  $Q(\boldsymbol{\eta}, \mathbf{p} \mid \boldsymbol{\eta}^{(t)}, \mathbf{p}^{(t)})$ :

$$\begin{aligned}
Q(\boldsymbol{\eta}, \mathbf{p} \mid \boldsymbol{\eta}^{(t)}, \mathbf{p}^{(t)}) &= \mathbb{E} \left[ \ell_c(\boldsymbol{\eta}, \mathbf{p} \mid \mathbf{Y}, \mathbf{A}) \mid \boldsymbol{\eta}^{(t)}, \mathbf{p}^{(t)} \right] \\
&= \sum_{\alpha \in \{0,1\}^K} \sum_{i=1}^N \varphi_{i,\alpha}^{(t+1)} \log p_\alpha \\
&\quad + \sum_{\alpha \in \{0,1\}^K} \sum_{i=1}^N \varphi_{i,\alpha}^{(t+1)} \sum_{j=1}^J \left[ (\boldsymbol{\eta}_{j,(0)}^\top \mathbf{T}(Y_{i,j}) - A(\boldsymbol{\eta}_{j,(0)}) + \log h(Y_{i,j})) (1 - \Gamma_{j,\alpha}) \right. \\
&\quad \left. + (\boldsymbol{\eta}_{j,(1)}^\top \mathbf{T}(Y_{i,j}) - A(\boldsymbol{\eta}_{j,(1)}) + \log h(Y_{i,j})) \Gamma_{j,\alpha} \right].
\end{aligned}$$

**M step for ExpDINA.** Next, in the M step in the  $t$ th EM iteration, we maximize  $Q(\boldsymbol{\eta}, \mathbf{p} \mid \boldsymbol{\eta}^{(t)}, \mathbf{p}^{(t)})$  with respect to  $(\boldsymbol{\eta}, \mathbf{p})$  and update the model parameters as follows:

$$(\boldsymbol{\eta}^{(t+1)}, \mathbf{p}^{(t+1)}) = \underset{\boldsymbol{\eta}, \mathbf{p}}{\operatorname{argmax}} Q(\boldsymbol{\eta}, \mathbf{p} \mid \boldsymbol{\eta}^{(t)}, \mathbf{p}^{(t)}).$$

Since  $\boldsymbol{\eta}_{j,(0)}, \boldsymbol{\eta}_{j,(1)}, \mathbf{p}$  are continuous parameters, we can set their partial gradients to zero and update parameters as the solutions to the gradient equations. Specifically, we will solve

$$\operatorname{argmax}_{\mathbf{p}} \sum_{\boldsymbol{\alpha} \in \{0,1\}^K} \sum_{i=1}^N \varphi_{i,\boldsymbol{\alpha}}^{(t+1)} \log p_{\boldsymbol{\alpha}}, \quad \text{subject to} \quad \sum_{\boldsymbol{\alpha} \in \{0,1\}^K} p_{\boldsymbol{\alpha}} = 1, \quad (23)$$

$$\operatorname{argmax}_{\boldsymbol{\eta}_{j,(0)}} \sum_{\boldsymbol{\alpha} \in \{0,1\}^K} \sum_{i=1}^N \varphi_{i,\boldsymbol{\alpha}}^{(t+1)} \sum_{j=1}^J (\boldsymbol{\eta}_{j,(0)}^\top \mathbf{T}(Y_{i,j}) - A(\boldsymbol{\eta}_{j,(0)})) (1 - \Gamma_{j,\boldsymbol{\alpha}}), \quad \forall j \in [J], \quad (24)$$

$$\operatorname{argmax}_{\boldsymbol{\eta}_{j,(1)}} \sum_{\boldsymbol{\alpha} \in \{0,1\}^K} \sum_{i=1}^N \varphi_{i,\boldsymbol{\alpha}}^{(t+1)} \sum_{j=1}^J (\boldsymbol{\eta}_{j,(1)}^\top \mathbf{T}(Y_{i,j}) - A(\boldsymbol{\eta}_{j,(1)})) \Gamma_{j,\boldsymbol{\alpha}}, \quad \forall j \in [J]. \quad (25)$$

The first optimization problem in (23) has a closed-form solution for  $\mathbf{p}$ :

$$p_{\boldsymbol{\alpha}} = \frac{\sum_{i=1}^N \varphi_{i,\boldsymbol{\alpha}}^{(t+1)}}{\sum_{\boldsymbol{\alpha} \in \{0,1\}^K} \sum_{i=1}^N \varphi_{i,\boldsymbol{\alpha}'}^{(t+1)}}, \quad \forall \boldsymbol{\alpha} \in \{0,1\}^K.$$

For optimization problems in (24) and (25) for the item parameters, there also often exist closed-form updates for many distributions. Consider  $\boldsymbol{\eta}_{j,(1)}$  in (25) for some item  $j \in [J]$ . Setting the partial gradient of (25) with respect to  $\boldsymbol{\eta}_{j,(1)}$  to zero, we obtain:

$$\sum_{\boldsymbol{\alpha}} \sum_i \varphi_{i,\boldsymbol{\alpha}}^{(t+1)} \Gamma_{j,\boldsymbol{\alpha}} (\mathbf{T}(Y_{i,j}) - \nabla_{\boldsymbol{\eta}} A(\boldsymbol{\eta})) = 0, \implies \nabla_{\boldsymbol{\eta}_{j,(1)}} A(\boldsymbol{\eta}) = \frac{\sum_{i,\boldsymbol{\alpha}} \mathbf{T}(Y_{i,j}) \varphi_{i,\boldsymbol{\alpha}}^{(t+1)} \Gamma_{j,\boldsymbol{\alpha}}}{\sum_{i,\boldsymbol{\alpha}} \varphi_{i,\boldsymbol{\alpha}}^{(t+1)} \Gamma_{j,\boldsymbol{\alpha}}}. \quad (26)$$

Now note that an exponential family distribution has a nice property that  $A(\boldsymbol{\eta})$  is always a convex function of  $\boldsymbol{\eta}$  (Casella and Berger, 2021). This is because the Hessian matrix of  $A(\boldsymbol{\eta})$  equals the covariance matrix of the sufficient statistics  $\mathbf{T}$  and hence is always positive definite. This property implies that  $\boldsymbol{\eta} \rightarrow \nabla_{\boldsymbol{\eta}} A(\boldsymbol{\eta})$  is always an invertible map of  $\boldsymbol{\eta}$  and there exists a unique solution to the gradient equation in (26). In fact, for many widely used exponential family distributions, (26) has an explicit solution and hence the corresponding ExpDINA has explicit M step updates. For example, under the *Lognormal-DINA* model, maximizing (24)–(25) gives the following updates of the mean and variance parameters:

$$\mu_{j,1}^{(t+1)} = \frac{\sum_{i,\boldsymbol{\alpha}} \varphi_{i,\boldsymbol{\alpha}} \Gamma_{j,\boldsymbol{\alpha}} \log Y_{i,j}}{\sum_{i,\boldsymbol{\alpha}} \varphi_{i,\boldsymbol{\alpha}} \Gamma_{j,\boldsymbol{\alpha}}}, \quad (\sigma_{j,1}^2)^{(t+1)} = \frac{\sum_{i,\boldsymbol{\alpha}} \varphi_{i,\boldsymbol{\alpha}} \Gamma_{j,\boldsymbol{\alpha}} (\log Y_{i,j} - \mu_{j,1}^{(t+1)})^2}{\sum_{i,\boldsymbol{\alpha}} \varphi_{i,\boldsymbol{\alpha}} \Gamma_{j,\boldsymbol{\alpha}}};$$

$$\mu_{j,0}^{(t+1)} = \frac{\sum_{i,\alpha} \varphi_{i,\alpha} (1 - \Gamma_{j,\alpha}) \log Y_{i,j}}{\sum_{i,\alpha} \varphi_{i,\alpha} (1 - \Gamma_{j,\alpha})}, \quad (\sigma_{j,0}^2)^{(t+1)} = \frac{\sum_{i,\alpha} \varphi_{i,\alpha} (1 - \Gamma_{j,\alpha}) (\log Y_{i,j} - \mu_{j,1}^{(t+1)})^2}{\sum_{i,\alpha} \varphi_{i,\alpha} (1 - \Gamma_{j,\alpha})}.$$

Updates for ExpDINA for other transformed-Normal distributions (e.g., logistic-Normal) can be obtained by simply replacing the  $\log Y_{i,j}$  term in the above display with the corresponding transform. As another example, under the *Poisson-DINA* model, maximizing (24)–(25) gives the following updates for the Poisson rate parameters:

$$\lambda_{j,1}^{(t+1)} = \frac{\sum_{i,\alpha} \phi_{i,\alpha} \Gamma_{j,\alpha} Y_{i,j}}{\sum_{i,\alpha} \phi_{i,\alpha} \Gamma_{j,\alpha}}, \quad \lambda_{j,0}^{(t+1)} = \frac{\sum_{i,\alpha} \phi_{i,\alpha} (1 - \Gamma_{j,\alpha}) Y_{i,j}}{\sum_{i,\alpha} \phi_{i,\alpha} (1 - \Gamma_{j,\alpha})}.$$

## 4.2 EM Algorithm for the ExpACDM

In this subsection we propose an EM algorithm to estimate the model parameters under the ExpACDM defined in Section 2.2.2. Recall that the ExpACDM is parametrized by  $\boldsymbol{\eta}_{j,\alpha} = \mathbf{h}(\beta_{j,0} + \sum_{k=1}^K \beta_{j,k} q_{j,k} \alpha_k, \gamma_j)$  with all parameters collected in  $(\boldsymbol{\beta}, \gamma, \mathbf{p})$ . The complete data log-likelihood (21) under an ExpACDM can be written as:

$$\begin{aligned} \ell_c^{\text{ExpACDM}}(\boldsymbol{\beta}, \gamma, \mathbf{p} \mid \mathbf{Y}, \mathbf{A}) &= \sum_{\boldsymbol{\alpha} \in \{0,1\}^K} \sum_{i=1}^N \mathbb{1}(\mathbf{A}_i = \boldsymbol{\alpha}) \log p_{\boldsymbol{\alpha}} \\ &+ \sum_{\boldsymbol{\alpha} \in \{0,1\}^K} \sum_{i=1}^N \mathbb{1}(\mathbf{A}_i = \boldsymbol{\alpha}) \sum_{j=1}^J \left\{ \mathbf{h}(\beta_{j,0} + \sum_{k=1}^K \beta_{j,k} q_{j,k} \alpha_k, \gamma_j)^\top \mathbf{T}(Y_{i,j}) \right. \\ &\quad \left. - A(\mathbf{h}(\beta_{j,0} + \sum_{k=1}^K \beta_{j,k} q_{j,k} \alpha_k, \gamma_j)) + \log h(Y_{i,j}) \right\}. \end{aligned} \quad (27)$$

Based on the above expression, we can derive an EM algorithm to estimate parameters in an ExpACDM. The detailed steps are summarized in Algorithm 2.

An interesting fact is that in Algorithm 2, maximizing  $(\boldsymbol{\beta}_j, \gamma_j)$  in each M step is similar to obtaining the MLE of the regression coefficients in a *generalized linear model*, but with the observed covariates replaced with the latent attributes evaluated in the E step. In particular, in the special case of the transformed-Normal distributions, this maximization is similar to linear regression and has a closed form solution. For example, when  $\mathcal{P}$  is the

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**Algorithm 1:** EM Algorithm for the General ExpDINA Model

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**Data:** Responses  $\mathbf{Y} = (Y_{ij})_{N \times J}$ ,  $\mathbf{Q}$ -matrix  $\mathbf{Q} = (q_{jk})_{J \times K}$ .

Initialize parameters  $(\boldsymbol{\eta}, \mathbf{p})$ .

**while** *log-likelihood has not converged* **do**

    In the  $(t + 1)$ th iteration,

    // E step

**for**  $(i, \boldsymbol{\alpha}) \in [N] \times \{0, 1\}^K$  **do**

$$\begin{aligned} \varphi_{i,\boldsymbol{\alpha}}^{(t+1)} &= \mathbb{P}(\mathbf{A}_i = \boldsymbol{\alpha} \mid \mathbf{Y}, \boldsymbol{\eta}^{(t)}, \mathbf{p}^{(t)}) \\ &= \frac{p_{\boldsymbol{\alpha}}^{(t)} \exp \left( \sum_j [\boldsymbol{\eta}_{j,(0)}^{(t)\top} \mathbf{T}(Y_{i,j}) - A(\boldsymbol{\eta}_{j,(0)}^{(t)})](1 - \Gamma_{j,\boldsymbol{\alpha}}) + [\boldsymbol{\eta}_{j,(1)}^{(t)\top} \mathbf{T}(Y_{i,j}) - A(\boldsymbol{\eta}_{j,(1)}^{(t)})]\Gamma_{j,\boldsymbol{\alpha}} \right)}{\sum_{\boldsymbol{\alpha}'} p_{\boldsymbol{\alpha}'}^{(t)} \exp \left( \sum_j [\boldsymbol{\eta}_{j,(0)}^{(t)\top} \mathbf{T}(Y_{i,j}) - A(\boldsymbol{\eta}_{j,(0)}^{(t)})](1 - \Gamma_{j,\boldsymbol{\alpha}'}) + [\boldsymbol{\eta}_{j,(1)}^{(t)\top} \mathbf{T}(Y_{i,j}) - A(\boldsymbol{\eta}_{j,(1)}^{(t)})]\Gamma_{j,\boldsymbol{\alpha}'} \right)}; \end{aligned}$$

    // M step

**for**  $\boldsymbol{\alpha} \in \{0, 1\}^K$  **do**

$$p_{\boldsymbol{\alpha}}^{(t+1)} = \frac{\sum_i \varphi_{i,\boldsymbol{\alpha}}^{(t+1)}}{\sum_{i,\boldsymbol{\alpha}'} \varphi_{i,\boldsymbol{\alpha}'}^{(t+1)}};$$

**for**  $j \in [J], h \in \{0, 1\}$  **do**

$$\boldsymbol{\eta}_{j,(h)}^{(t+1)} = \underset{\boldsymbol{\eta}}{\operatorname{argmax}} \sum_{i,\boldsymbol{\alpha}} \left[ (1 - \Gamma_{j,\boldsymbol{\alpha}})^{1-h} \Gamma_{j,\boldsymbol{\alpha}}^h (\boldsymbol{\eta}^\top \mathbf{T}(Y_{i,j}) - A(\boldsymbol{\eta})) \varphi_{i,\boldsymbol{\alpha}}^{(t+1)} \right];$$

    Or equivalently, solve the following gradient equations for  $(\boldsymbol{\eta}_{j,(0)}^{(t+1)}, \boldsymbol{\eta}_{j,(1)}^{(t+1)})$ :

$$\begin{aligned} (\nabla_{\boldsymbol{\eta}} A)(\boldsymbol{\eta}_{j,(1)}^{(t+1)}) &= \frac{\sum_{i,\boldsymbol{\alpha}} \mathbf{T}(Y_{i,j}) \varphi_{i,\boldsymbol{\alpha}}^{(t+1)} \Gamma_{j,\boldsymbol{\alpha}}}{\sum_{i,\boldsymbol{\alpha}} \varphi_{i,\boldsymbol{\alpha}}^{(t+1)} \Gamma_{j,\boldsymbol{\alpha}}}, \\ (\nabla_{\boldsymbol{\eta}} A)(\boldsymbol{\eta}_{j,(0)}^{(t+1)}) &= \frac{\sum_{i,\boldsymbol{\alpha}} \mathbf{T}(Y_{i,j}) \varphi_{i,\boldsymbol{\alpha}}^{(t+1)} (1 - \Gamma_{j,\boldsymbol{\alpha}})}{\sum_{i,\boldsymbol{\alpha}} \varphi_{i,\boldsymbol{\alpha}}^{(t+1)} (1 - \Gamma_{j,\boldsymbol{\alpha}})}. \end{aligned}$$

**Output:** Estimated parameters  $(\boldsymbol{\eta}, \mathbf{p})$ .

---

Normal distribution, the maximization for  $(\boldsymbol{\beta}_j, \gamma_j)$  in the M step can be written as:

$$(\boldsymbol{\beta}_j, \gamma_j)^{(t+1)} = \underset{\boldsymbol{\beta}_j, \gamma_j}{\operatorname{argmax}} \sum_{i,\boldsymbol{\alpha}} \left( -\frac{(Y_{i,j} - \beta_{j,0} - \sum_{k=1}^K \beta_{j,k} q_{j,k} \alpha_k)^2}{2\gamma_j^2} - \frac{1}{2} \log \gamma_j^2 \right) \varphi_{i,\boldsymbol{\alpha}}^{(t+1)}$$

---

**Algorithm 2:** EM Algorithm for the General ExpACDM

---

**Data:** Responses  $\mathbf{Y} = (Y_{ij})_{N \times J}$ ,  $\mathbf{Q}$ -matrix  $\mathbf{Q} = (q_{jk})_{J \times K}$

Initialize parameters  $\beta_j, \gamma_j, p_\alpha$ 's.

**while** *log-likelihood has not converged* **do**

    In the  $(t+1)$ th iteration,

    // E step

**for**  $j \in [J], \alpha \in \{0, 1\}^K$  **do**

$\boldsymbol{\eta}_{j,\alpha}^{(t)} = \mathbf{h}(\beta_{j,0} + \sum_k \beta_{j,k} \alpha_k, \gamma_j^{(t)})$

**for**  $(i, \alpha) \in [N] \times \{0, 1\}^K$  **do**

$$\varphi_{i,\alpha}^{(t+1)} = \mathbb{P}(\mathbf{A}_i = \alpha \mid \mathbf{Y}, \beta_j^{(t)}, \gamma_j^{(t)}, p_\alpha^{(t)}) = \frac{p_\alpha \exp(\sum_j [\boldsymbol{\eta}_{j,\alpha}^{(t)\top} \mathbf{T}(Y_{i,j}) - A(\boldsymbol{\eta}_{j,\alpha}^{(t)})])}{\sum_{\alpha'} p_{\alpha'} \exp(\sum_j [\boldsymbol{\eta}_{j,\alpha'}^{(t)\top} \mathbf{T}(Y_{i,j}) - A(\boldsymbol{\eta}_{j,\alpha'}^{(t)})])};$$

    // M step

**for**  $\alpha \in \{0, 1\}^K$  **do**

$$p_\alpha^{(t+1)} = \frac{\sum_i \varphi_{i,\alpha}^{(t+1)}}{\sum_{i,\alpha'} \varphi_{i,\alpha'}^{(t+1)}}$$

**for**  $j \in [J]$  **do**

$$\begin{aligned} (\beta_j, \gamma_j)^{(t+1)} = \operatorname{argmax}_{\beta_j, \gamma_j} \sum_{i,\alpha} & \left[ \mathbf{h}(\beta_{j,0} + \sum_{k=1}^K \beta_{j,k} q_{j,k} \alpha_k, \gamma_j)^\top \mathbf{T}(Y_{i,j}) \varphi_{i,\alpha}^{(t+1)} \right. \\ & \left. - A(\mathbf{h}(\beta_{j,0} + \sum_{k=1}^K \beta_{j,k} q_{j,k} \alpha_k, \gamma_j)) \varphi_{i,\alpha}^{(t+1)} \right] \end{aligned}$$

**Output:**  $\beta_j, \gamma_j, p_\alpha$ 's.

---

for any  $j \in [J]$ . So for those  $q_{j,k} = 1$ , we can get the updates

$$\hat{\beta}_{j,k}^{(t+1)} = \frac{\sum_{i,\alpha} Y_{i,j} \alpha_k \varphi_{i,\alpha}^{(t+1)}}{\sum_{i,\alpha} \alpha_k \varphi_{i,\alpha}^{(t+1)}}, \quad \hat{\gamma}_j^{(t+1)} = \sqrt{\frac{\sum_{i,\alpha} (Y_{i,j} - \beta_{j,0} - \sum_{k=1}^K \beta_{j,k} q_{j,k} \alpha_k)^2 \varphi_{i,\alpha}^{(t+1)}}{N}}.$$

The updates under the transformed-Normal distributions can be obtained similarly by applying the corresponding transform of  $Y_{i,j}$ . For other members in the exponential family beyond the transformed-Normal, maximizing  $(\beta_j, \gamma_j)$  may not have closed forms and one may use available optimization software to find  $(\hat{\beta}_j, \hat{\gamma}_j)$  in the M step.

Although we have focused on describing EM algorithms for ExpDINA and ExpACDM

in this section, we remark that an EM algorithm with closed-form M step can be similarly developed for the ExpGDM (ExpGDINA). Furthermore, the exponential family assumption in Algorithms 1 and 2 is not essential to our EM procedures. In Supplementary Material S.4, we demonstrate how our algorithms can be modified to estimate the negative-binomial-based DINA and negative-binomial-based ACDM.

## 5 Simulation Studies

We conduct simulation studies under various models in the proposed family. We have two goals in the simulations: (a) to empirically verify the theoretical results of identifiability and consistency; and (b) to assess the computational performance of the proposed EM algorithms. Under the ExpDINA and the ExpACDM, we consider the Normal and transformed Normal distributions (i.e., lognormal and logistic normal) for continuous responses in Section 5.1, and the Poisson and negative binomial distributions for count responses in Section 5.2. We remark that in addition to the distributions considered in this section, it is also easy to use our framework and estimation procedures for other exponential family distributions such as Gamma and Beta to model continuous positive, and continuous bounded data, respectively.

### 5.1 Simulations under the Normal- and transformed Normal-CDMs

We first describe the true parameter settings used in the simulations. In all simulations, we set the  $\mathbf{Q}$ -matrix and proportion parameter  $\mathbf{p}$  as follows. Consider  $K = 5$  latent attributes and  $J = 20$  items. The  $\mathbf{Q}$ -matrix takes the form

$$\mathbf{Q} = \begin{pmatrix} \mathbf{I}_K \\ \mathbf{I}_K \\ \mathbf{I}_K \\ \mathbf{Q}_1 \end{pmatrix}, \quad \text{where } \mathbf{Q}_1 = \begin{pmatrix} 1 & 1 & & 0 \\ 1 & \ddots & \ddots & \\ & \ddots & \ddots & 1 \\ 0 & & 1 & 1 \end{pmatrix}_{K \times K}. \quad (28)$$

The above  $\mathbf{Q}$ -matrix satisfies our strict identifiability conditions in Theorem 1 and Proposition 2. The proportion parameters are set to be uniform with  $p_\alpha = 1/2^K$  for all  $\alpha \in \{0, 1\}^K$ . We consider varying sample sizes  $N = 500, 1000, 1500, 2000$ . In each setting, 100 indepen-

dent simulation replicates are performed. Under the Normal-DINA model, we set the true item parameters as

$$\mu_{j,0} = -1, \quad \mu_{j,1} = 2, \quad \sigma_{j,0} = 1, \quad \sigma_{j,1} = 1.$$

Under the Normal-ACDM, we set the coefficients  $\beta_{j,k}$  by

$$\beta_{j,0} = -1; \quad \beta_{j,k} = \frac{3}{\sum_{k'=1}^K q_{j,k'}} \mathbb{1}(q_{j,k} = 1), \quad \forall j \in [J], k \in [K].$$

The variance parameter  $\gamma_j = \sigma_j^2$  is fixed to be 1 for all  $j$ .

Recall that in Example 1, we have mentioned that general-response CDMs based on any transformed Normal distributions (such as lognormal and logistic Normal) are equivalent to the Normal-based CDM. This fact implies that the estimation procedures for these models are identical up to an invertible transformation in the M step. In preliminary simulations, we have tried estimating the Normal-based and lognormal-based CDMs independently, and the estimation accuracies for the two models are exactly the same. Hence, we only report the estimation accuracy for the Normal-based CDM in this section.

In each of the  $C = 100$  independent replicates, we generate data using the above parameter settings and fit our EM Algorithm 1 or 2 with a random initialization. We calculate the root average mean squared error (RMSE) of the proportion parameters and item parameters based on the simulation replicates. Figure 2 displays the average RMSEs for the Normal-DINA and Normal-ACDM. In each simulation setting, the RMSE is defined as

$$\sqrt{\frac{1}{C} \sum_{c=1}^C \frac{\|\hat{\boldsymbol{\theta}}^{(c)} - \boldsymbol{\theta}_0\|_2^2}{\dim(\boldsymbol{\theta}_0)}},$$

where  $\boldsymbol{\theta}_0$  is the true parameter vector,  $\hat{\boldsymbol{\theta}}^{(c)}$  is the estimator in the  $c$ -th simulation replication, and  $\dim(\boldsymbol{\theta}_0)$  denotes the dimension of the vector  $\boldsymbol{\theta}_0$ . The exact RMSE values are included in Tables S.1, S.2, and S.3 in the Supplementary Material.

Figure 2 clearly shows that the RMSE decreases as the sample size  $N$  increases. Furthermore, in Figure 3, we plot the average RMSE of  $\boldsymbol{p}$  under the Normal-DINA model versus  $1/\sqrt{N}$ , and it is clear that the RMSE is linear with respect to  $1/\sqrt{N}$ . This observation

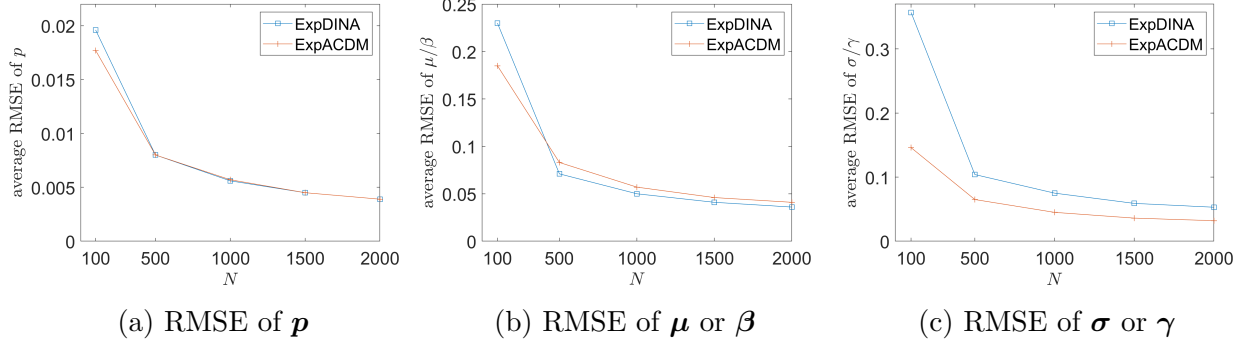


Figure 2: RMSEs of parameters under the Normal-DINA (red) and Normal-ACDM (blue).

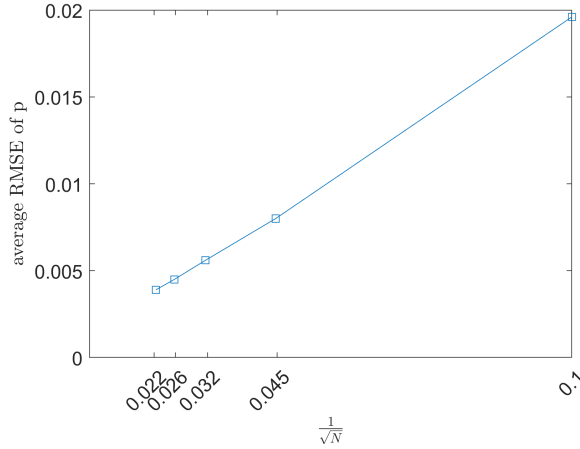


Figure 3: RMSEs of  $p$  with respect to  $\frac{1}{\sqrt{N}}$  under the Normal-DINA model.

empirically validates that our estimation procedure is statistically consistent and converges at the usual parametric rate  $1/\sqrt{N}$ . In addition, the above simulation results also show that our EM algorithms have good computational performance and can efficiently find the MLE.

## 5.2 Simulations under the Poisson- and negative binomial-CDMs

We also conduct simulation studies for general-response CDMs for multivariate count data. Here, we consider Poisson and negative binomial distributions to model the count responses. The  $\mathbf{Q}$ -matrix and the proportion parameters  $\mathbf{p}$  are set to be the same as described in the previous subsection. The distribution-specific item parameters are set as follows. For the Poisson-DINA model, we set the item parameters as  $\lambda_{j,0} = 1$  and  $\lambda_{j,1} = 3$ . For the Poisson-

ACDM, we set the true model parameters as

$$\beta_{j,0} = 1, \quad \beta_{j,k} = \frac{2}{\sum_{k'=1}^K q_{j,k'}} \mathbb{1}(q_{j,k} = 1), \quad \forall j \in [J], \quad k \in [K].$$

For the negative binomial-DINA model (NegBin-DINA, defined in Supplementary Material S.1), we set the item parameters as

$$r_{j,0} = 1, \quad r_{j,1} = 3, \quad \pi_{j,0} = 0.5, \quad \pi_{j,1} = 0.5, \quad \forall j \in [J],$$

where  $r_{j,0}, r_{j,1}$  are the number of successes and  $\pi_{j,0}, \pi_{j,1}$  are the success probability in a negative binomial distribution. For the negative binomial-ACDM (NegBin-ACDM, defined in Supplementary Material S.1), we set the true model parameters as

$$\beta_{j,0} = 1, \quad \beta_{j,k} = \frac{2}{\sum_{k'=1}^K q_{j,k'}} \mathbb{1}(q_{j,k} = 1), \quad \gamma_j = \pi_j = 0.5, \quad \forall j \in [J], \quad k \in [K].$$

Figures 4 and 5 report the RMSEs of the estimated parameters obtained from the replicated simulations. Similar to the Normal-based CDMs, the RMSEs here also decrease as  $N$  increases, at the typical  $1/\sqrt{N}$  rate.

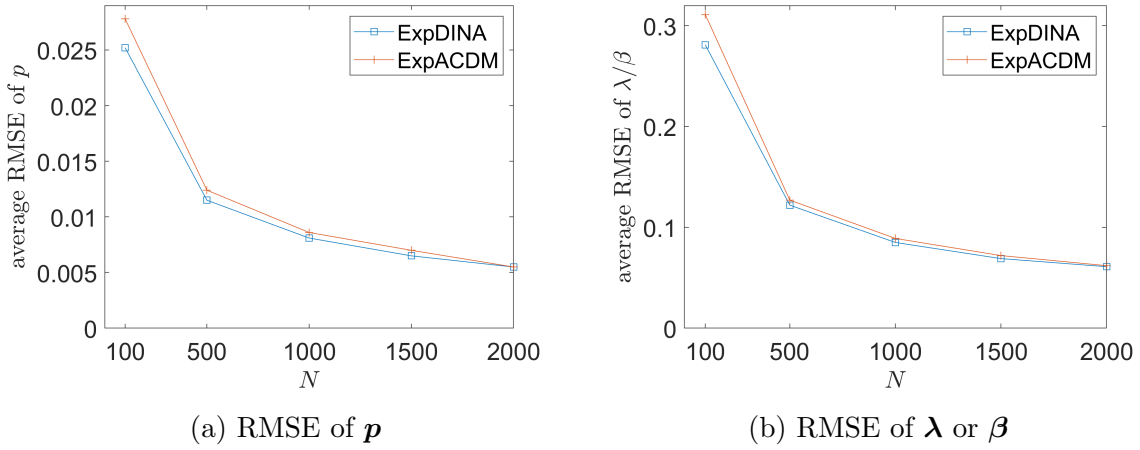


Figure 4: RMSEs of parameters under the Poisson-DINA (red) and Poisson-ACDM (blue).

Finally, we remark that our estimation methods are computationally quite efficient. In the simulation settings considered in this section, the computation time taken by our EM algorithm is less than one minute on average even for sample size as large as  $N = 2000$ .

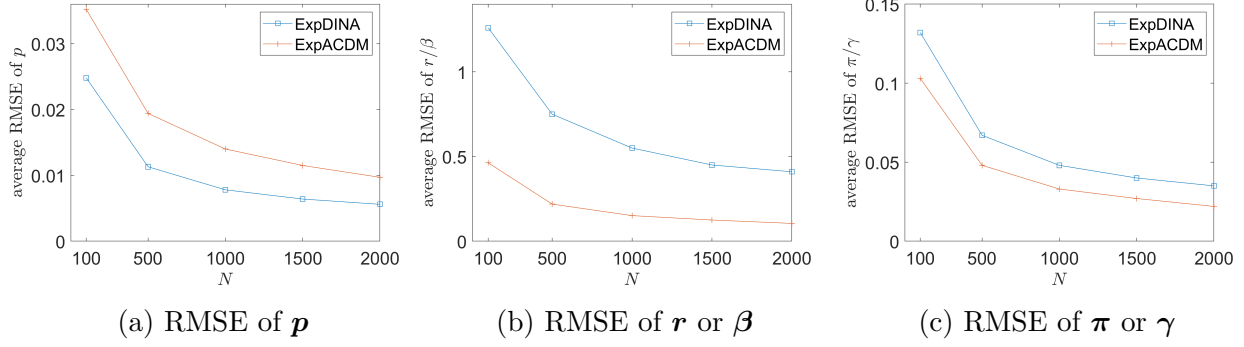


Figure 5: RMSEs of parameters under the NegBin-DINA (red) and NegBin-ACDM (blue).

We present the average number of EM iterations and computation time in Supplementary Material S.4.

## 6 Application to the TIMSS 2019 Response Time Data

We demonstrate the proposed general-response CDM by applying it to a response time dataset extracted from the TIMSS 2019 assessment (Fishbein et al., 2021). This dataset is from the mathematics assessment data of eighth-grade students in the United States. It consists of each student’s time spent on each item screen (in seconds). If a question has sub-questions that share the same screen, all the sub-questions combined are regarded as a single question, and the overall response time is recorded. We focus on the students who received booklet number 14. After data preprocessing (see details in Supplementary Material S.5), the dataset consists of  $N = 620$  students’ response times to  $J = 29$  items.

The TIMSS 2019 mathematics assessment aims to measure four content skills (Number, Algebra, Geometry, and Data and Probability) and three cognitive skills (Knowing, Applying, and Reasoning). The TIMSS 2019 database specifies how each of the  $J = 29$  items measures one content skill and one cognitive skill. Based on this information, we construct a  $29 \times 7$   $\mathbf{Q}$ -matrix with  $K = 7$  skill attributes:  $A_1$ : Number,  $A_2$ : Algebra,  $A_3$ : Geometry,  $A_4$ : Data and Probability,  $A_5$ : Knowing,  $A_6$ : Applying,  $A_7$ : Reasoning. Each row of this  $\mathbf{Q}$ -matrix has exactly two nonzero entries in one content skill and one cognitive skill. We provide the details of this  $\mathbf{Q}$ -matrix in Table 1. It is not hard to verify that this  $\mathbf{Q}$ -matrix satisfies our generic identifiability conditions in Theorem 2 under any general-response Ex-

pACDMs, and we choose to use the lognormal-ACDM to analyze this dataset. The TIMSS database also provides additional item information, including a brief description of the item type (whether it is a multiple choice item or a constructed response item), and the correct response percentage among the U.S. students. We present this information in Table 2.

Item ID.	Number	Algebra	Geometry	Data and Prob.	Knowing	Applying	Reasoning
1	1	0	0	0	1	0	0
2	1	0	0	0	0	1	0
3	1	0	0	0	0	1	0
4	1	0	0	0	1	0	0
5	0	1	0	0	0	1	0
6	0	1	0	0	1	0	0
7	0	1	0	0	0	0	1
8	0	1	0	0	0	1	0
9	0	0	1	0	0	0	1
10	0	0	1	0	0	1	0
11	0	0	1	0	0	1	0
12	0	0	0	1	0	1	0
13	0	0	0	1	0	1	0
14	1	0	0	0	1	0	0
15	1	0	0	0	1	0	0
16	1	0	0	0	0	0	1
17	1	0	0	0	0	1	0
18	0	1	0	0	1	0	0
19	0	1	0	0	1	0	0
20	0	1	0	0	0	1	0
21	0	1	0	0	1	0	0
22	0	1	0	0	0	1	0
23	0	0	1	0	1	0	0
24	0	0	1	0	0	0	1
25	0	0	1	0	0	0	1
26	0	0	1	0	0	1	0
27	0	0	0	1	0	1	0
28	0	0	0	1	0	1	0
29	0	0	0	1	0	0	1

Table 1: **Q**-matrix for the TIMSS 2019 math assessment booklet 14.

Given the **Q**-matrix in Table 1, we fit the lognormal-ACDM using Algorithm 2. In addition to the identifiability consideration, our rationale for adopting the additive model assumption is that in order to solve each problem, students need to perform operations (i.e., attributes) that are specified by the **Q**-matrix additively. We assume that each operation is carried out separately, and the total log-time is the sum of the log-times for each operation. For instance, the question “Value of  $X$  in  $10/15 = X/18$ ” (this is the 15th question in our dataset; see Tables 1 and 2 for more details) measures the content skill “Number” and

Item	Description	Item Type	% Correct Response
1	Octagon with equivalent shading	MC	55
2	Percentage of laps finished	CR	32*
3	Multiples of 3	MC	42
4	Convert decimal to a fraction	CR	42
5	Expression for area of rectangle	MC	51
6	Ratio of boys	MC	56
7	Rule for number of matches	CR	11*
8	Graph of $y = 2x$	MC	41
9	Rotation and reflection	MC	34
10	Surface area of the prism	MC	41
11	Value of angle $x$ outside triangle	MC	44
12	Number of balls in a bag	MC	69
13	Liv's smartphone use	MC	35*
14	Number with most factors	MC	54
15	Value of $X$ in $10/15 = X/18$	CR	50
16	Make two fractions with largest product	CR	41
17	Ratio of boys to girls - eighth grade	CR	53*
18	Expression for money in savings account	MC	73
19	Value of $k$ when $(k + 7) / 3 = 6$	CR	60
20	Formula for students' overall scores in course	CR	54
21	Expression equivalent to $x + 2y + 6$	MC	42
22	Pair of equations for ticket price (Derived)	MC	30
23	Shape reflected over dotted line (Derived)	MC	58
24	Value of angle $x$ with parallel lines $a$ and $b$	CR	11
25	Areas of triangles $ABC$ and $ABD$	CR	8
26	Surface area of rectangular prism	CR	19
27	Bar graph of Internet users in world	MC	60
28	Graph that best shows favorite fruit	MC	65
29	Mean heights of basketball team	CR	30

Table 2: Additional item information for items in TIMSS 2019 math assessment booklet 14. Starred (\*) items are items that are composed of sub-questions; for these items, we display the smallest correct response percentage among all sub-questions.

cognitive skill “Knowing”, and we assume that the log-time is the sum of:

$\beta_{j,0}$ : how long it takes to read the problem and click/type the answer;

$\beta_{j,K}A_K$ : how long it takes to re-formulate the problem as “ $10 \times 18 \div 15 = X$ ”, regarding whether a student “knows” the necessary concepts, i.e. whether the student possesses the cognitive skill “Knowing”;

$\beta_{j,N}A_N$ : how long it takes to compute and find the answer  $X = 12$ , regarding whether the student possesses the content skill “Number”.

Summing up the above components and adding a normal error gives the following response time distribution, which corresponds to the lognormal-ACDM defined in Example 3:

$$\log Y_{i,j} \mid \mathbf{A} = \underset{\text{intercept}}{\beta_{j,0}} + \underset{\text{cognitive skill}}{\beta_{j,K}A_K} + \underset{\text{content skill}}{\beta_{j,N}A_N} + \epsilon_{i,j}, \text{ where } \epsilon_{i,j} \stackrel{\text{i.i.d.}}{\sim} N(0, \gamma_j).$$

Decomposing the response time into several components has a long history. [Sternberg \(1969\)](#) and [Sternberg \(1980\)](#) considered a linear regression model based on a sequence of hypothetical processes that students go through to solve a problem. However, these studies do not introduce individual-level latent variables. [Maris \(1993\)](#) considered a latent variable model with Gamma distribution for the response time, with a similar additive combination of the latent variables. These models are also referred to as exploratory response time models in the literature (e.g. [De Boeck and Jeon, 2019](#)). The recent papers [Minchen et al. \(2017\)](#) and [Minchen and de la Torre \(2018\)](#) incorporate this line of thinking into a CDM framework with the lognormal distribution for modeling the response times. The empirical results in these papers show that students possessing more required attributes take a longer time to respond, so we anticipate that our main-effect coefficients  $\beta_{j,k}$  for the attributes are nonnegative.

We run our EM Algorithm 2 with 20 random parameter initializations with  $\beta_{j,k} > 0$ . We do not impose any strict constraints on the sign of  $\beta_{j,k}$ , and allow the algorithm to update parameters in the unconstrained space. Among the resulting 20 parameter estimates, we select the estimate with the largest log-likelihood. Figure 6 presents the estimated main-effect coefficients. This figure shows that students spend the most time reading the question and clicking/typing the answer (represented by the intercepts in the first column), compared to the time it takes to re-formulate and solve the problem (represented by the main-effect coefficients from the second column to the last column). The intercept values vary a lot across different questions, ranging from 1.0 to 3.1. This indicates that the length and the abstractness of the questions vary substantially across items. One interesting observation is the relation between the magnitude of the estimated intercept  $\beta_{j,0}$  and the type of each item. Recall that Table 2 provides information about whether each item is a multiple choice item or a constructed response item. Nonetheless, Figure 6 reveals that, among the first ten items, the three items that ask students to construct the response (items 2, 4, 7) have the largest intercept values. This is clearly visible by looking at the darker-colored entries (or

the numbers shown in white) in the first column of Figure 6. This result can be interpreted as that students spend more time typing and checking the answer when solving constructed response items. Note that the item type information is *not used* in our estimation procedure, but our method can automatically distinguish the multiple choice items from the constructed response items via the estimated parameters.

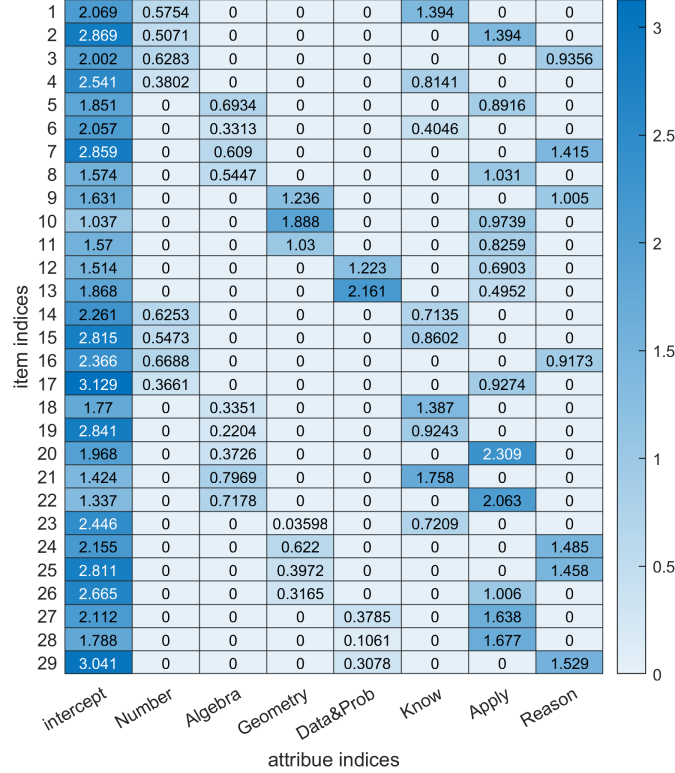


Figure 6: Heatmap of the estimated  $\beta$ -parameters under the lognormal-ACDM for the TIMSS 2019 response time dataset.

As for the main-effect coefficients  $\beta_{j,k}$  for  $k \geq 1$ , the computation time represented by the content skills' coefficients and formulation time represented by the cognitive skills' coefficients also differ a lot across the items. These coefficients range from 0.03 to 2.3. Even though we do not constrain  $\beta_{j,k}$  to be positive, all estimated main-effect coefficients turn out to be positive, which is consistent with empirical findings in previous studies of response time modeling. This observation also indicates that our additive model assumption is indeed plausible and yields interpretable parameter estimates.

We also provide the correlation plot of the estimated latent attributes in Figure 7. The

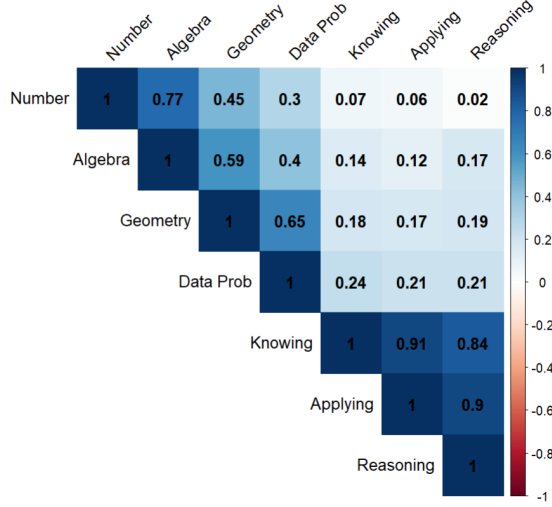


Figure 7: Correlation plot of the latent attributes under the estimated lognormal-ACDM for the TIMSS 2019 response time dataset.

EM algorithm does not estimate the individual latent attributes directly, but instead computes the conditional expectations  $\hat{\varphi}_{i,\alpha} = \mathbb{P}(\mathbf{A}_i = \alpha \mid \mathbf{Y}, \hat{\boldsymbol{\eta}}, \hat{\mathbf{p}})$  for all  $i \in [N]$  and  $\alpha \in \{0, 1\}^K$ . Therefore, we estimate the latent attribute profiles  $\mathbf{A}_i$  by

$$\hat{\mathbf{A}}_i = \underset{\alpha \in \{0,1\}^K}{\operatorname{argmax}} \hat{\varphi}_{i,\alpha},$$

and use these estimated  $\hat{\mathbf{A}}_1, \dots, \hat{\mathbf{A}}_N$  to compute the sample correlation between the attributes. Figure 7 reveals interesting observations about the intrinsic dependence among the attributes. There are higher correlations among the four content skills and also higher correlations among the three cognitive skills, compared to the correlations between one content skill and one cognitive skill. This phenomenon not only supports that our model is a reasonable choice as it outputs interpretable correlation structures among latent attributes, but also implies that it may be plausible to model high-order latent traits behind the seven fine-grained attributes (de la Torre and Douglas, 2004; Zhan et al., 2018a). Future studies are warranted to explore the identifiability and interpretability of such high-order extensions of our general-response CDMs.

## 7 Discussion

In this paper, we have proposed a flexible new framework of cognitive diagnostic models for multivariate general responses beyond the traditional binary or polytomous responses. Our modeling framework incorporates the  $\mathbf{Q}$ -matrix constraints in a unified way and covers popular existing CDMs as submodels. An important contribution of this work is to provide the crucial identifiability theory for all these general-response CDMs. Interestingly and somewhat surprisingly, we have shown that the general-response CDMs are identifiable under similar conditions on the  $\mathbf{Q}$ -matrix as the binary-response CDMs. Our identifiability theory has the nice implication of consistent parameter estimation via the maximum likelihood. For computation, we have proposed an efficient EM algorithm for parameter estimation under various response types. Extensive simulation studies not only corroborate the identifiability conclusions, but also demonstrate the favorable computational performance of our algorithms. We have analyzed a response time dataset from the TIMSS 2019 assessment using the proposed lognormal-ACDM and obtained interpretable results.

The proposed new paradigm of identifiable general-response cognitive diagnostic models open up a number of interesting possibilities for future research. *First*, our current identifiability results and estimation procedure assume that the  $\mathbf{Q}$ -matrix is known. In practice, many modern assessment datasets may not come with a readily available  $\mathbf{Q}$ -matrix. In these exploratory settings, it would be interesting to directly identify and estimate the  $\mathbf{Q}$ -matrix along with other model parameters for general-response CDMs. In terms of identifiability, we conjecture that our identifiability proof technique could be generalized to also handle the identifiability of the  $\mathbf{Q}$ -matrix. In terms of estimation, it may be possible to extend existing methods for estimating the  $\mathbf{Q}$ -matrix such as penalized likelihood methods in [Chen et al. \(2015\)](#) and [Ma et al. \(2023\)](#), or Bayesian approaches in [Chen et al. \(2018\)](#) and [Liu et al. \(2020\)](#) from the binary-response CDMs to general-response CDMs. We plan to pursue these directions of exploratory general-response CDMs in the future.

*Second*, it would be interesting to explore possibilities of strengthening our identifiability results by developing weaker identifiability conditions for special submodels of general-response CDMs. Our main theorems provide transparent sufficient conditions for identifiability in a very general setting. But for binary-response CDMs under some special modeling

assumptions, existing studies show that there exist weaker conditions that can guarantee identifiability. For example, for the DINA model with binary responses, [Gu and Xu \(2019\)](#) proposed weaker conditions that are necessary and sufficient for strict identifiability. Establishing necessary identifiability conditions in the most general setting with arbitrary response types is a nontrivial but interesting future direction.

*Third*, many educational assessments naturally contain multiple types of responses for each item. For instance, it is common to record the response accuracy and visual fixation counts/visit counts in addition to the response time for each item ([Zhan et al., 2022](#); [Fishbein et al., 2021](#)). Jointly modeling multiple types of responses has received great attention in the measurement literature ([van der Linden, 2007](#); [Molenaar et al., 2015](#); [Zhan et al., 2018b](#); [Wang et al., 2018](#); [Man and Harring, 2022](#); [Kang et al., 2023](#)). Among these modeling approaches, CDM-based methods such as [Zhan et al. \(2018b\)](#) and [Zhan et al. \(2022\)](#) use binary latent variables to model the response accuracy only (i.e., binary responses of correct or wrong answers to items), and use an IRT-based model with continuous latent variables to model the response time and fixation counts. In fact, as suggested by our analysis of the TIMSS response time data, using binary latent skills to model the response time can also yield interpretable results. Therefore, it would be interesting in the future to extend our current work and propose identifiable general-response CDMs for joint responses.

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