Distributional Treatment Effect with Latent Rank Invariance*

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September 30, 2025

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Abstract

Treatment effect heterogeneity is of a great concern when evaluating policy impact: "is the treatment Pareto-improving?", "what is the proportion of people who are better off under the treatment?", etc. However, even in the simple case of a binary random treatment, existing analysis has been mostly limited to an average treatment effect or a quantile treatment effect, due to the fundamental limitation that we cannot simultaneously observe both treated potential outcome and untreated potential outcome for a given unit. This paper assumes a conditional independence assumption that the two potential outcomes are independent of each other given a scalar latent variable. With a specific example of strictly increasing conditional expectation, I label the latent variable as 'latent rank' and motivate the identifying assumption as 'latent rank invariance.' In implementation, I assume a finite support on the latent variable and propose an estimation strategy based on a nonnegative matrix factorization. A limiting distribution is derived for the distributional treatment effect estimator, using Neyman orthogonality.

Keywords: distributional treatment effect, proximal inference, finite mixture, nonnegative matrix factorization, *U*-statistic, Neyman orthogonality.

JEL classification codes: C13

*I thank Stéphane Bonhomme, Bernard Salanie, Myunghwan Seo and Martin Weidner for their valuable comments. I acknowledge the support from the European Research Council through the grant ERC-2018-CoG-819086-PANEDA. Any and all errors are my own.

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1 Introduction

The fundamental limitation that we cannot simultaneously observe the two potential outcomes—treated potential outcome and untreated potential outcome—for a given unit makes the task of identifying the distribution of treatment effect particularly complicated. Thus, instead of estimating the entire distribution of treatment effect, researchers often estimate some summary measures of the treatment effect distribution, such as the average treatment effect (ATE) or the quantile treatment effect (QTE). These summary measures provide insights into the treatment effect distribution and thus help researchers with policy recommendations. However, there still remain a lot of questions that can only be answered with the distribution of the treatment effect: e.g., is the treatment Pareto improving?; how heterogeneous is the treatment effect at the unit level? Also, the distribution is important in empirical contexts where participation cannot be mandated. To anticipate the participation rate, we need to identify the share of people who are better off under the treatment regime and thus would select into treatment.

Consider a potential outcome setup with a binary treatment:

$$Y = D \cdot Y(1) + (1 - D) \cdot Y(0).$$

Y(1) is the treated potential outcome, Y(0) is the untreated potential outcome, and $D \in \{0,1\}$ is the binary treatment variable. The questions above correspond to testing $H_0: F_{Y(1)-Y(0)}(0) = 0$ and estimating Var(Y(1)-Y(0)). Note that these quantities, $F_{Y(1)-Y(0)}(0)$ and Var(Y(1)-Y(0)), all come from the distribution of individual-level treatment effect Y(1)-Y(0). To answer questions that relate to the distributional concerns in policy recommendation more broadly, I focus on the following two parameters of interest:

$$F_{Y(1),Y(0)}(y_1,y_0) = \Pr\{Y(1) \le y_1, Y(0) \le y_0\}$$
 for some (y_1,y_0) ,
 $F_{Y(1)-Y(0)}(\delta) = \Pr\{Y(1) - Y(0) \le \delta\}$ for some δ .

The first parameter is the joint distribution of the two potential outcomes and the second parameter is the marginal distribution of the treatment effect. For the rest of the paper, I refer to these quantities as the distributional treatment effect (DTE) parameters.¹

¹Some previous works in the literature use the terminology 'distributional effect' to discuss parameters that are a functional of the marginal distributions of the potential outcomes; e.g., Firpo and Pinto (2016). To avoid confusion, I will reserve the expression 'distributional' to only when the object involves the joint distribution of the two potential

When we believe that there is no dependence between the two potential outcomes, meaning that a realized value of the treated potential outcome has no information on the individual-level heterogeneity and thus has no predictive power for the untreated potential outcome and vice versa, identification of the joint distribution of the two potential outcomes becomes trivial. Once we identify the marginal distributions of the two potential outcomes, the joint distribution becomes their product. However, this assumption is extremely restrictive. Thus, I instead assume *conditional* independence, by assuming a scalar latent variable that captures the individual-level heterogeneity in terms of the dependence between the two potential outcomes. For illustration, consider a simple additive model: the two potential outcomes are constructed with a unit-level latent variable $U \in \mathbb{R}$ and two treatment-status-specific random shocks $\varepsilon(1)$ and $\varepsilon(0)$:

$$Y(1) = \mu_1(U) + \varepsilon(1),\tag{1}$$

$$Y(0) = \mu_0(U) + \varepsilon(0). \tag{2}$$

When

$$\varepsilon(1) \perp \!\!\!\perp \varepsilon(0) \mid U,$$
 (3)

we can characterize the joint distribution of the two potential outcome as follows:

$$\Pr\{Y(1) \le y_1, Y(0) \le y_0\} = \mathbf{E} \left[\Pr\{Y(1) \le y_1 | U\} \cdot \Pr\{Y(0) \le y_0 | U\} \right].$$

Thus, the task of identifying the joint distribution of the two potential outcomes becomes that of identifying the conditional distribution of $\varepsilon(1)$ given U, the conditional distribution of $\varepsilon(0)$ given U, and the marginal distribution of U.

To identify the conditional distribution of $\varepsilon(d)$ given U and the marginal distribution of U, I assume that there are two additional proxy variables X, Z that are conditionally independent of each other and the potential outcomes, given U. This identification strategy is drawn from the nonclassical measurement error literature and the proximal inference literature: see Hu and Schennach (2008); Miao et al. (2018); Deaner (2023); Kedagni (2023); Nagasawa (2022) and more. In the simple example (1)-(2), the proxy variables X, Z will shift $\mu(U)$ independently of $(\varepsilon(1), \varepsilon(0))$, allowing us to decompose the variation of Y(d) into the variation of U and the variation of $\varepsilon(d)$.

outcomes.

Additionally, since I do not adopt the 'measurement error' interpretation on the proxy variables as in the nonclassical measurement error literature, I assume that there exists a functional of the conditional distribution of the potential outcomes given the latent variable, which strictly increases in the latent variable U. An example of such a functional is conditional expectation. Suppose that the two conditional expectations $\mathbf{E}[Y(1)|U=u]$ and $\mathbf{E}[Y(0)|U=u]$ are strictly increasing in u. In this example, the latent variable U can be thought of as the rank of the conditional expectations $\mathbf{E}[Y(1)|U]$ and $\mathbf{E}[Y(0)|U]$; hence 'latent rank invariance.' The conditional independence assumption and the latent rank invariance assumption are the key assumptions in identification.

In developing estimators for the distributional treatment effect parameters, I additionally assume a finite support on U. The finite support assumptions has several merits. Firstly, it motivates a simple estimation method based on a nonnegative matrix factorization algorithm. Secondly, the conditional independence assumption can be interpreted as finite mixture whose properties are well-studied in the literature. Lastly, under the finite support assumption, the identification of the DTE parameters reduces down to a GMM model with quadratic moments, giving us some insights on how the DTE parameters are identified. Though I assume that U is finitely discrete in the estimation, the identification result does not require such a restriction and I develop an alternative estimation method based on sieve maximum likelihood for a setup with continuous U, in the Appendix subsection A.2.

The estimation procedure is two-step. In the first step, I estimate the conditional probability $\Pr\{U=u|Z=z\}$, using the nonnegative matrix factorization. In the second step, I identify a DTE parameter with a moment condition involving probabilities $\Pr\{Y \leq y|D=d, Z=z\}$ and $\Pr\{U=u|Z=z\}$. The former probability is directly observed from the dataset and the latter is estimated in the first step. Thus, the estimator can be thought of as a plug-in GMM estimator, where nuisance parameters are estimated in the first-step nonnegative matrix factorization. Asymptotic normality of the distributional treatment effect parameters is established. In deriving asymptotic normality, I construct a moment condition that satisfies Neyman orthogonality to be robust to the first-step estimation error from the nonnegative matrix factorization.

This paper makes contribution to the distributional treatment effect literature by proposing a framework where the joint distribution of the potential outcomes and thus the marginal distribution of treatment effect are point identified, without imposing any functional form assumptions. This is in contrast to the partial identification results in the literature: Fan and Park (2010); Fan et al. (2014); Firpo and Ridder (2019); Frandsen and Lefgren (2021); Kaji and Cao (2023) and

more. There exist several notable point identification results: Heckman et al. (1997); Carneiro et al. (2003). These point identification results either assume independence on potential outcomes conditioning on observables only, or assume structural assumptions on treatment and/or potential outcomes: e.g. a Roy model for treatment and a factor structure for potential outcomes. In terms of estimation, Wu and Perloff (2006); Noh (2023) also develop DTE estimators; unlike this paper, they both build on the point identification result without latent conditioning variable and develop a deconvolution-based estimator. Bedoya et al. (2018) provides an insightful overview on recent developments on both identification and estimation in program evaluation literature regarding distributional concerns.

This paper also makes contribution to the nonclassical measurement error/proximal inference literature and the finite mixture literature: Hu and Schennach (2008); Henry et al. (2014); Miao et al. (2018); Deaner (2023); Kedagni (2023); Nagasawa (2022) and more. In terms of identification, the latent rank invariance assumption provides an alternative assumption in labeling the latent variable that uses the information from the outcome variable Y, as opposed to the "measure of location" assumption suggested in Hu and Schennach (2008) that uses the information from the proxy variable X. Alternatively, this paper can be thought of as adding an additional identifying assumption—conditional independence between Y(d) and X—to narrow down the identified set of Henry et al. (2014) to a singleton. In terms of the asymptotic theory on the estimator, this paper is in a similar setup as Hu (2008), assuming a finite support. Unlike existing estimators based on the principal component analysis, the estimation strategy based on the nonnegative matrix factorization as proposed in this paper has guarantee that the estimated conditional distributions are indeed nonnegative and sum-to-one. The \sqrt{n} -consistency is proven in this paper so that future works may build upon the nonnegative matrix factorization estimator.

The rest of the paper is organized as follows. Section 2 discusses the identification result for the joint distribution of the two potential outcomes. Section 3 explains the estimation method for the two DTE parameters and develops asymptotic theory for the estimators. Section 4 contains Monte Carlo simulation restuls and Section 5 applies the estimation procedure to an empirical dataset from Jones et al. (2019).

2 Identification

An econometrican observes a dataset $\{Y_i, D_i, X_i, Z_i\}_{i=1}^n$ where $Y_i, X_i, Z_i \in \mathbb{R}$ and $D_i \in \{0, 1\}$. Y_i is an outcome variable, D_i is a binary treatment variable and X_i, Z_i are two proxy variables. The outcome Y_i is constructed with two potential outcomes.

$$Y_i = D_i \cdot Y_i(1) + (1 - D_i) \cdot Y_i(0). \tag{4}$$

In addition to $(Y_i(1), Y_i(0), D_i, X_i, Z_i)$, there is a latent variable $U_i \in \mathcal{U} \subset \mathbb{R}$. U_i plays a key role in putting restrictions on the joint distribution of $Y_i(1)$ and $Y_i(0)$ and overcoming the fundamental limitation that we observe only one potential outcome for a given unit. The dataset comes from random sampling: $(Y_i(1), Y_i(0), D_i, X_i, Z_i, U_i) \stackrel{iid}{\sim} \mathcal{F}$.

Firstly, I assume conditional random assignment on the treatment D_i and exclusion restriction on the proxy variable Z_i .

Assumption 1. (assignment/exclusion restriction) $(Y_i(1), Y_i(0), X_i) \perp (D_i, Z_i) \mid U_i$.

Assumption 1 assumes that the treatment is as good as random with regard to the potential outcomes and X_i after conditioning on the latent variable U_i . In this sense, Assumption 1 is a restriction on treatment endogeneity. In addition, Assumption 1 assumes that the proxy variable Z_i does not have any additional information on the potential outcomes after conditioning on the latent variable U_i , satisfying exclusion restriction. Note that Assumption 1 does not impose any restriction on the dependence between Z_i and D_i . The proxy variable Z_i may still depend on treatment. This assumption imposes nontrivial restriction on treatment endogeneity in a non-experimental context. Thus, for the rest of the paper, I focus on a randomly assigned treatment, limiting my attention to randomized controlled trials. The following condition is a sufficient condition for Assumption 1.

Remark. A sufficient condition for Assumption 1 is

$$(Y_i(1), Y_i(0), X_i, U_i) \perp D_i$$
 and $(Y_i(1), Y_i(0), X_i) \perp Z_i \mid (D_i, U_i)$.

Thus, in a randomized controlled trial setup, Assumption 1 is satisfied when there are one proxy variable that is independent of the treatment D_i and another proxy variable that only depends the latent heterogeneity U_i and the treatment D_i .

When U_i is observed, Assumption 1 identifies numerous treatment effect parameters such as

average treatment effect (ATE), quantile treatment effect (QTE) and more. However, even when U_i is observed, we still cannot identify the distribution of treatment effect from Assumption 1 since Assumption 1 does not tell us anything about the dependence between $Y_i(1)$ and $Y_i(0)$.

To impose restrictions on the joint distribution of $Y_i(1)$ and $Y_i(0)$ and have more identifying power, I assume that the latent variable U_i captures all of the dependence between the two potential outcomes and the proxy variable X_i .

Assumption 2. (conditional independence) $Y_i(1)$, $Y_i(0)$ and X_i are all mutually independent given U_i .

Assumption 2 assumes that the two potential outcomes $Y_i(1)$ and $Y_i(0)$ and the proxy variable X_i are mutually independent of each other conditioning on U_i . Given U_i , the proxy variable X_i does not give us additional information on the distribution of the potential outcomes. Note that the latent variable U_i lies in \mathbb{R} as do $Y_i(1)$ and $Y_i(0)$. This excludes a non-binding case where $U_i = (Y_i(1), Y_i(0))$.

When U_i is observed, Assumptions 1-2 identify the joint distribution of the two potential outcomes and various distributional treatment effect parameters. Examples include the variance of the treatment effect $Var(Y_i(1) - Y_i(0))$ and the marginal distribution of the treatment effect $Pr\{Y_i(1) - Y_i(0) \le \delta\}$. Since U_i is not observed, identifying the conditional densities of $Y_i(1), Y_i(0)$ given U_i and the marginal density of U_i will be the main challenge in the identification.

Assumptions 1-2 play a key role in the identification result. Below I present three examples of econometric frameworks that motivate Assumptions 1-2. The first example is rank invariance, which is widely used in the quantile treatment effect literature and the quantile IV literature: see Chernozhukov and Hansen (2005, 2006); Athey and Imbens (2006); Vuong and Xu (2017); Callaway and Li (2019); Han and Xu (2023) and more.

Example 1. (rank invariance) The econometrician observes $\{Y_i, D_i\}_{i=1}^n$ and

$$Y_i = D_i \cdot Y_i(1) + (1 - D_i) \cdot Y_i(0).$$

The treatment D_i is randomly assigned and the potential outcomes $Y_i(1)$ and $Y_i(0)$ have the same rank:

$$\Pr\{F_{Y(1)}(Y_i(1)) = F_{Y(0)}(Y_i(0))\} = 1.$$

Then, Assumptions 1-2 are satisfied with $U_i = X_i = Z_i = F_{Y(1)}(Y_i(1)) = F_{Y(0)}(Y_i(0))$.

The usage of this rank invariance assumption is mostly limited to the quantile treatment effect and not applied to the distributional treatment effect, due to the fact that it imposes excessive restriction on the joint distribution of the two potential outcomes. Under the rank invariance, $\operatorname{Var}(Y_i(1)|Y_i(0)) = 0$ and vice versa. Thus, the treatment effect $Y_i(1) - Y_i(0)$ is also a deterministic function of $Y_i(1)$ and of $Y_i(0)$.

In this paper, I relax this deterministic relationship among the potential outcomes and the latent variable, by assuming the rank invariance not on the potential outcomes directly, but on some functional of the conditional distribution of the potential outcome given U_i . In this sense, the econometric framework of this paper is a relaxation of the rank invariance assumption in the quantile treatment effect literature and the quantile IV literature.

The second example is a panel data model with a latent state variable that is first-order Markovian, which is often referred to as a hidden Markov model. The hidden Markov model is widely discussed in the dynamic panel data model literature, especially in the context of the dynamic discrete choice and conditional choice probability estimation: see Kasahara and Shimotsu (2009); Arcidiacono and Miller (2011); Hu and Shum (2012); Hu and Sasaki (2018) for more.

Example 2. (panel with a latent state variable) The econometrician observes $\{\{Y_{it}\}_{t=1}^3, D_i\}_{i=1}^n$ where

$$Y_{it}(d) = g_d(V_{it}, \varepsilon_{it}(d))$$

for t = 1, 2, 3 and d = 0, 1 and

$$Y_{it} = \begin{cases} Y_{i1}(0) & \text{if } t = 1 \\ D_i \cdot Y_{i2}(1) + (1 - D_i) \cdot Y_{i2}(0) & \text{if } t = 2 \cdot \\ Y_{i3}(1) & \text{if } t = 3 \end{cases}$$

 $\{V_{it}\}_{t=1}^3$ is first-order Markovian given D_i and $(\{V_{it}\}_{t=1}^3, D_i)$, $\varepsilon_{i1}, \varepsilon_{i2}(1), \varepsilon_{i2}(0)$ and ε_{i3} are mutually independent. D_i is randomly assigned at time t=2: $\{V_{it}\}_{t=1}^2 \perp \!\!\!\perp D_i$. Then, Assumptions 1-2 are

$$D_{i2} = \mathbf{1} \{ \mathbf{E} \left[Y_{i2}(1) - Y_{i2}(0) | V_{i2} \right] \ge \eta_i \}$$

The assignment model above assumes that at the timing of selection, individuals are only aware of their common

²Even when the treatment D_i is not random, Assumption 1 may not be too restrictive an assumption in the context of Example 2. Suppose that the common shock V_{it} is drawn first and then the treatment-status-specific shocks $\varepsilon_{it}(1)$ and $\varepsilon_{it}(0)$ are drawn subsequently and that at time t=2, individuals select into treatment by comparing their expected gain from being treated with their costs η_i before the treatment-status-specific shocks are realized:

satisfied with $Y_i = Y_{i2}, X_i = Y_{i1}, Z_i = Y_{i3}$ and $U_i = V_{i2}$. Similarly, Assumptions 1-2 are also satisfied with $Y_i = Y_{i2}, X_i = Y_{i1}, Z_i = Y_{i3}$ and $U_i = V_{i2}$ when $Y_{i3} = D_i \cdot Y_{i3}(1) + (1 - D_i) \cdot Y_{i3}(0)$.

In this nonlinear panel data model, the potential outcome $Y_{it}(d)$ is a function of a latent variable V_{it} and an error term $\varepsilon_{it}(d)$. Note that V_{it} appears in the model twice; for $Y_{it}(1)$ and for $Y_{it}(0)$. In this sense, V_{it} is a common shock to the potential outcomes where $\varepsilon_{it}(d)$ is a treatment-status-specific shock. The key elements of Example 2 are that the common shock process $\{V_{it}\}_{t=1}^3$ and the treatment-status-specific shocks $\varepsilon_{i1}(1), \ldots, \varepsilon_{i3}(0)$ are all mutually independent and that dependence within $\{V_{it}\}_{t=1}^3$ themselves is restricted to be first-order Markovian given D_i . Thus, V_{i2} has sufficient information on the dependence between $Y_{i2}(1)$ and $Y_{i2}(0)$ and the past and the future outcomes Y_{i1} and Y_{i3} can be used as proxies for V_{i2} .

The hidden Markov model is mostly applied to a single observed outcome setup. In this paper, I extend the hidden Markov model to a potential outcome setup so that there are two idiosyncratic error terms $\varepsilon_{it}(0)$ and $\varepsilon_{it}(1)$, specific to each treatment status. Moreover, I add one more conditional independence to the hidden Markov model by assuming that the two error terms are independent across the treatment status conditioning on U_i .

The second example closely relates to Section 5 of this paper. In Section 5, I revisit Jones et al. (2019) and estimate the full distribution of treatment effect, in the empirical context of workplace wellness program as a treatment and monthly medical spending as an outcome. The dataset used in Jones et al. (2019) contains short panel data on monthly medical spending, with one pretreatment time period. Thus, by assuming that the monthly medical spending is a function of two different types of random shocks—a transitory, idiosyncratic shock and a systemic health shock that is first-order Markovian—, the model described in Example 2 can be applied to the dataset and we can use the pretreatment medical spending and the post-treatment medical spending as the two proxy variables.

The third example is where we have economic interpretation on the latent variable U_i and therefore can find measurements on the latent variable. There are several notable papers in labor economics that adopts this approach: see Carneiro et al. (2003); Cunha and Heckman (2008); Cunha et al. (2010) and more.

shock V_{i2} and thus their (conditionally) expected gain $\mathbf{E}[Y_{i2}(1) - Y_{i2}(0)|V_{i2}]$, but not the realized gain $Y_i(1) - Y_i(0)$. When η_i , the idiosyncratic shock in the assignment model, is independent of the shocks in the outcome model, Assumption 1 is satisfied.

Example 3. (repeated measurements) The econometrician observes $\{Y_i, D_i, X_i, Z_i\}_{i=1}^n$ and

$$Y_i = D_i \cdot Y_i(1) + (1 - D_i) \cdot Y_i(0).$$

 Y_i is earning, D_i is treatment, X_i, Z_i are test scores, and $U_i = (U_{X,i}, U_{Z,i})$ is innate ability.

$$Y_i(d) = \frac{1}{\theta_d} \log \left(\rho_d U_{X,i}^{\theta_d} + (1 - \rho) U_{Z,i}^{\theta_d} \right) + \varepsilon_i(d) \quad \text{for } d = 0, 1$$

$$X_i = g_X(U_{X,i}) + \varepsilon_{X,i},$$

$$Z_i = g_Z(U_{Z,i}) + \varepsilon_{Z,i}.$$

Conditioning on U_i , D_i , $\varepsilon_i(0)$, $\varepsilon_i(1)$, $\varepsilon_{X,i}$ and $\varepsilon_{Z,i}$ are mutually independent.

The above model is a simplified version of the framework in Cunha et al. (2010), applied to a potential outcome setup. In this example, an economic model gives us an interpretation on the latent variable U_i and helps us find measurements on the latent variable. For example, in Cunha et al. (2010), $U_{X,i}$ and $U_{Z,i}$ are assumed to be cognitive skill and noncognotive skill. Then, various measures on cognitive ability, temperament, motor and social developments and such are used as proxy variables. In this paper, I consider a more flexible outcome function than the CES function, at the cost of assuming a univariate U_i and a strong independence assumption on the error terms.³ In this sense, this paper can also be thought of as nonparametric version of the Carneiro et al. (2003)'s framework.

The remainder of this section outlines the identification argument. For illustration purposes only, let Y_i, X_i, Z_i, U_i be discrete: $Y_i \in \{y^1, \dots, y^{M_Y}\}, X_i \in \{x^1, \dots, x^{M_X}\}, Z_i \in \{z^1, \dots, z^{M_Z}\}$ and $U_i \in \{u^1, \dots, u^K\}$. With $M = M_Y \cdot M_X$, we can construct a $M \times M_Z$ matrix \mathbf{H}_d of conditional probabilities as follows:

$$\mathbf{H}_{d} = \begin{bmatrix} \Pr\left\{ (Y_{i}, X_{i}) = \left(y^{1}, x^{1}\right) \mid (D_{i}, Z_{i}) = \left(d, z^{1}\right) \right\} & \cdots & \Pr\left\{ (Y_{i}, X_{i}) = \left(y^{1}, x^{1}\right) \mid (D_{i}, Z_{i}) = \left(d, z^{M_{Z}}\right) \right\} \\ \vdots & \ddots & \vdots \\ \Pr\left\{ (Y_{i}, X_{i}) = \left(y^{M_{Y}}, x^{M_{X}}\right) \mid (D_{i}, Z_{i}) = \left(d, z^{1}\right) \right\} & \cdots & \Pr\left\{ (Y_{i}, X_{i}) = \left(y^{M_{Y}}, x^{M_{X}}\right) \mid (D_{i}, Z_{i}) = \left(d, z^{M_{Z}}\right) \right\} \end{bmatrix}$$

³The main focus of Cunha et al. (2010) is less on the outcome, but more on the skill formation. In the full framework of Cunha et al. (2010), there exists time dimension and the skills vector U_{it} is modeled with a dynamic process and the paper nonparametrically identifies the skill evolution process.

for each d = 0, 1. \mathbf{H}_0 is the conditional probability of (Y_i, X_i) given Z_i in the untreated subsample and \mathbf{H}_1 is the conditional probability in the treated subsample. From Assumptions 1-2, both \mathbf{H}_0 and \mathbf{H}_1 decompose into a multiplication of two matrices: for each d=0,1,

$$\mathbf{H}_d = \Gamma_d \cdot \Lambda_d \tag{5}$$

where

$$\Gamma_{d} = \begin{pmatrix}
\Pr\{(Y_{i}(d), X_{i}) = (y^{1}, x^{1}) | U_{i} = u^{1}\} & \cdots & \Pr\{(Y_{i}(d), X_{i}) = (y^{1}, x^{1}) | U_{i} = u^{K}\} \\
\vdots & \ddots & \vdots \\
\Pr\{(Y_{i}(d), X_{i}) = (y^{M_{Y}}, x^{M_{X}}) | U_{i} = u^{1}\} & \cdots & \Pr\{(Y_{i}(d), X_{i}) = (y^{M_{Y}}, x^{M_{X}}) | U_{i} = u^{K}\} \end{pmatrix},$$

$$\Lambda_{d} = \begin{pmatrix}
\Pr\{(U_{i} = u^{1} | (D_{i}, Z_{i}) = (d, z^{1})\} & \cdots & \Pr\{(U_{i} = u^{1} | (D_{i}, Z_{i}) = (d, z^{M_{Z}})\}\} \\
\vdots & \ddots & \vdots \\
\Pr\{(U_{i} = u^{K} | (D_{i}, Z_{i}) = (d, z^{1})\} & \cdots & \Pr\{(U_{i} = u^{K} | (D_{i}, Z_{i}) = (d, z^{M_{Z}})\}\} \end{pmatrix}.$$
(6)

Note that the discreteness of Y_i, X_i, Z_i is nonbinding; we can use partitioning on \mathbb{R} when they are continuous. 4 The remaining discretization on U_{i} is imposed only for the expositional brevity; the identification argument does not hinge on the discreteness of U_i . The continuous U_i version of the identification follows the same argument and uses one additional assumption to find a labeling on the infinite number of functions: Assumption 5. I present more discussion on Assumption 5 later in this section and a full identification argument for continuous U_i is provided in Subsection A.1 of Appendix.

The equation $\mathbf{H}_d = \Gamma_d \cdot \Lambda_d$ shows us that the conditional density model in (16) is indeed a mixture model. For each subpopulation $\{i:(D_i,Z_i)=(d,z)\}$, there is a column in the matrix Λ_d which denotes the subpopulation-specific distribution of U_i . Then, the density of (Y_i, X_i) in that subpopulation admits a mixture model with the aforementioned columns of Λ_d as mixture weights

$$\left\{ \mathcal{Y}^m = \left(y^{m-1}, y^m \right) \right\}_{m=1}^{M_Y}, \quad \left\{ \mathcal{X}^m = \left(x^{m-1}, x^m \right) \right\}_{m=1}^{M_X}, \quad \left\{ \mathcal{Z}^m = \left(z^{m-1}, z^m \right) \right\}_{m=1}^{M_Z}$$

 $\left\{\mathcal{Y}^m = \left(y^{m-1}, y^m\right]\right\}_{m=1}^{M_Y}, \quad \left\{\mathcal{X}^m = \left(x^{m-1}, x^m\right]\right\}_{m=1}^{M_X}, \quad \left\{\mathcal{Z}^m = \left(z^{m-1}, z^m\right]\right\}_{m=1}^{M_Z}$ where $y^0 = x^0 = z^0 = -\infty$ and $y^{M_Y} = x^{M_X} = z^{M_Z} = \infty$. Let $\mathcal{W}^1 = \mathcal{Y}^1 \times \mathcal{X}^1, \mathcal{W}^2 = \mathcal{Y}^2 \times \mathcal{X}^1, \cdots, \mathcal{W}^M = \mathcal{Y}^{M_Y} \cdot \mathcal{X}^{M_X}$. $\left\{\mathcal{W}^m\right\}_{m=1}^{M}$ is a partition on \mathbb{R}^2 . Then, \mathbf{H}_d becomes

$$\mathbf{H}_{d} = \begin{pmatrix} \Pr\left\{ (Y_{i}, X_{i}) \in \mathcal{W}^{1} | D_{i} = d, Z_{i} \in \mathcal{Z}^{1} \right\} & \cdots & \Pr\left\{ (Y_{i}, X_{i}) \in \mathcal{W}^{1} | D_{i} = d, Z_{i} \in \mathcal{Z}^{M_{Z}} \right\} \\ \vdots & \vdots & \vdots \\ \Pr\left\{ (Y_{i}, X_{i}) \in \mathcal{W}^{M} | D_{i} = d, Z_{i} \in \mathcal{Z}^{1} \right\} & \cdots & \Pr\left\{ (Y_{i}, X_{i}) \in \mathcal{W}^{M} | D_{i} = d, Z_{i} \in \mathcal{Z}^{M_{Z}} \right\} \end{pmatrix}$$

for each d=0,1. Γ_d and Λ_d are similarly constructed with partitioned Y_i,X_i and Z_i .

⁴Consider partitions on \mathbb{R} such that

and the conditional density of $(Y_i(d), X_i)$ given U_i as mixture component densities. The equation $\mathbf{H}_d = \Gamma_d \cdot \Lambda_d$ aggregates the finite mixture formulations across the subpopulations.

Note that from Assumption 2, the joint distribution of $Y_i(1)$ and $Y_i(0)$ is identified if the conditional distribution of $Y_i(1)$ given U_i , the conditional distribution of $Y_i(0)$ given U_i , and the marginal distribution of U_i are identified. The first two distributions correspond to Γ_1 and Γ_0 in the discretization. The last distribution is a function of Λ_1, Λ_0 and the distribution of (D_i, Z_i) , which is observed. Thus, to identify of the distributional treatment effect parameter is to identify $\Gamma_1, \Gamma_0, \Lambda_1$ and Λ_0 .

To decompose \mathbf{H}_d into Γ_d and Λ_d , first fix $y \in \{y^1, \dots, y^{M_Y}\}$ and extract rows of \mathbf{H}_d and Γ_d that correspond to $(y, x^1), \dots, (y, x^{M_X})$:

$$\mathbf{H}_{d}(y) = \left(\Pr \left\{ (Y_{i}, X_{i}) = (y, x^{j}) \mid (D_{i}, Z_{i}) = (d, z^{k}) \right\} \right)_{1 \leq j \leq M_{X}, 1 \leq k \leq M_{Z}}$$

$$\Gamma_{d}(y) = \left(\Pr \left\{ (Y_{i}(d), X_{i}) = (y, x^{j}) \mid U_{i} = u^{k} \right\} \right)_{1 \leq j \leq M_{X}, 1 \leq k \leq K}$$

for d = 0, 1. From Assumption 2, the mixture component density matrix $\Gamma_d(y)$ can be further decomposed:

$$\Gamma_{d}(y) = \begin{pmatrix} \Pr\left\{X_{i} = x^{1} \middle| U_{i} = u^{1}\right\} & \cdots & \Pr\left\{X_{i} = x^{1} \middle| U_{i} = u^{K}\right\} \\ \vdots & \ddots & \vdots \\ \Pr\left\{X_{i} = x^{M_{X}} \middle| U_{i} = u^{1}\right\} & \cdots & \Pr\left\{X_{i} = x^{M_{X}} \middle| U_{i} = u^{K}\right\} \end{pmatrix}$$

$$\cdot \operatorname{diag}\left(\Pr\left\{Y_{i}(d) = y \middle| U_{i} = u^{1}\right\}, \cdots, \Pr\left\{Y_{i}(d) = y \middle| U_{i} = u^{K}\right\}\right)$$

$$=: \Gamma_{X} \cdot \Delta_{d}(y).$$

Now, sum $\mathbf{H}_d(y)$ across y^1, \dots, y^{M_Y} :

$$\sum_{y} \mathbf{H}_{d}(y) = \Gamma_{X} \cdot \sum_{y} \Delta_{d}(y) \cdot \Lambda_{d} = \Gamma_{X} \cdot \Lambda_{d}.$$

Find that when $M_X = M_Z = K$ and both Γ_X and Λ_d have full rank,

$$\mathbf{H}_{d}(y) \left(\sum_{y} \mathbf{H}_{d}(y) \right)^{-1} = \Gamma_{X} \cdot \Delta_{d}(y) \cdot \Lambda_{d} \left(\Gamma_{X} \cdot \Lambda_{d} \right)^{-1}$$
$$= \Gamma_{X} \cdot \Delta_{d}(y) \cdot \Gamma_{X}^{-1}.$$

Given a no repeated eigenvalue condition that for any $u \neq u'$ there exist some (y, d) such that $\Pr\{Y_i(d) = y | U_i = u\} \neq \Pr\{Y_i(d) = y | U_i = u'\}$, diagonalization of $\mathbf{H}_d(y) \left(\sum_y \mathbf{H}_d(y)\right)^{-1}$ across different y and d identifies Γ_X and $\{\Delta_d(y)\}_{y^1 \leq y \leq y^{M_Y}}$. Once Γ_X is identified, the identification of Λ_0, Λ_1 follows from Γ_X having full rank. When M_X or M_Z is bigger than K, we may stack some of the rows or the columns of $\sum_y \mathbf{H}_d(y)$ to make it into a square matrix.

Assumption 3 formally states the full rank condition and the no repeated eigenvalue condition for discrete U_i .

Assumption 3.

- **a.** (finitely discrete U_i) $\mathcal{U} = \{u^1, \dots, u^K\}$.
- **b.** (full rank) Λ_0 , Λ_1 and Γ_X have rank K.
- **c.** (no repeated eigenvalue) For any $k \neq k'$, there exist some $y, y' \in \{y^1, \dots, y^{M_Y}\}$ such that

$$\Pr \left\{ Y_i(0) = y | U_i = u^k \right\} \neq \Pr \left\{ Y_i(0) = y | U_i = u^{k'} \right\},$$

$$\Pr \left\{ Y_i(1) = y' | U_i = u^k \right\} \neq \Pr \left\{ Y_i(1) = y' | U_i = u^{k'} \right\}.$$

Assumption 3.b implicitly assumes that $M_X, M_Z \geq K$. The restriction that $M_X, M_Z \geq K$ is sensible since I use the variation in the conditional density of X_i given $Z_i = z$ across z to capture the variation in the latent variable U_i . The support for the two proxy variables has to be at least as rich as the support of the latent variable. Assumption 3.c assumes that the eigenvalue decomposition does not have repeated eigenvalues.

Assumption 4 reiterates Assumption 3 for a setup where U_i are continuous. Let $f_{Y(d)|U}$ denote the conditional density of $Y_i(d)$ given U_i , $f_{X|U}$ denote the conditional density of X_i given U_i , and $f_{U|D=d,Z}$ denote the conditional density of U_i given $D_i = d$ and Z_i , for d = 0, 1. Define integral operators $L_{X|U}$ and $L_{U|D=d,Z}$ that map a function in $\mathcal{L}^1(\mathbb{R})$ to a function in $\mathcal{L}^1(\mathbb{R})$: for d = 0, 1,

$$\begin{split} \left[L_{X|U}g\right](x) &= \int_{\mathbb{R}} f_{X|U}(x|u)g(u)du, \\ \left[L_{U|D=d,Z}g\right](u) &= \int_{\mathbb{R}} f_{U|D=d,Z}(u|z)g(z)dz. \end{split}$$

Assumption 4. Assume

⁵Eigenvalue decomposition on its own is not unique but we have sufficiently many constraints on Γ_X for unqueness; Γ_X , the eigenvector matrix, is nonnegative and its column-wise sums are one since they are conditional probabilities. See Hu and Schennach (2008) for more.

- **a.** (continuous U_i) $\mathcal{U} = [0, 1]$.
- **b.** (bounded density) The conditional densities $f_{Y(1)|U}$, $f_{Y(0)|U}$, $f_{X|U}$, $f_{U|D=1,Z}$ and $f_{U|D=0,Z}$ and the marginal densities f_U , $f_{Z|D=1}$ and $f_{Z|D=0}$ are bounded.
- **c.** (completeness) The integral operators $L_{X|U}, L_{X|D=1,Z}$ and $L_{X|D=0,Z}$ are injective on $\mathcal{L}^1(\mathbb{R})$.
- **d.** (no repeated eigenvalue) For any $u \neq u'$,

$$\Pr \{ f_{Y(d)|U}(Y_i|u) \neq f_{Y(d)|U}(Y_i|u') | D_i = d \} > 0$$

for each d=0,1.

Assumption 4.c corresponds to Assumption 3.b and Assumption 4.d to Assumption 3.c.

When U_i is continuous, we need an additional assumption for the identification. This is because when U_i is discrete and finite, a bijection between u and $\Pr\{X_i = \cdot | U_i = u\}$ needs not be specified. However, when U_i is continuous, we need an ordering on the infinite collection $\{f_{X|U}(\cdot|u)\}_u$ to connect u to $f_{X|U}(\cdot|u)$.

Assumption 5. (latent rank) There exists a functional M defined on $\mathcal{L}^1(\mathbb{R})$ such that either

$$h(u) = M f_{Y(1)|U}(\cdot|u) \qquad or \qquad h(u) = M f_{Y(0)|U}(\cdot|u)$$

defined on \mathcal{U} is strictly increasing and continuously differentiable.

The functional M provides us an ordering on the infinite collection $\{f_{X|U}(\cdot|u)\}_u$, by applying the functional to $\{f_{Y(1)|U}(\cdot|u), f_{Y(0)|U}(\cdot|u)\}_u$. A simple example where Assumption 5 fails is when $\mathcal{U} = [-1, 1]$ and $Y_i(d)|U_i = u \sim \mathcal{N}(u^2 + d, \sigma^2)$. Neither $f_{Y(1)|U}$ nor $f_{Y(0)|U}$ helps us find an ordering between $f_{X|U}(\cdot|u)$ and $f_{X|U}(\cdot|u)$.

Along with Assumptions 1-2, Assumption 5 is a key identifying assumption in the case of continuous U_i . As hinted by its label, Assumption 5 draws the inspiration from the rank invariance assumption in Example 1. Suppose that Assumption 5 holds true for $\mathbf{E}[Y_i(1)|U_i=u]$ and $\mathbf{E}[Y_i(0)|U_i=u]$. Then, the two potential outcomes of a given unit have the same 'latent rank' in the sense that their expected values $\mathbf{E}[Y_i(1)|U_i]$ and $\mathbf{E}[Y_i(0)|U_i]$ have the same rank in their respective distributions. A similar assumption can be made with other summary measures such as median or mode. Recall that Assumptions 1-2 is a relaxation of the rank invariance assumption.

Assumption 5 allows us to retain the rank interpretation on the latent variable U_i . Conditioning on the latent variable U_i , some summary measure applied to the conditional distributions of the potential outcomes has the same rank.

Theorem 1 formally states the identification result.

Theorem 1. Either Assumptions 1-3 or Assumptions 1-2, 4-5 hold. Then, the joint density of $(Y_i(1), Y_i(0), D_i, X_i, Z_i)$ is identified.

Proof. See Appendix.
$$\Box$$

The result of Theorem 1 can be understood as applying the identification result of Hu and Schennach (2008) twice, once to the treated population and again to the untreated population, and then connecting the two identification results. Also, when U_i is finite, the result of Theorem 1 can be understood as a point identification adaptation of the partial identification result from Henry et al. (2014); the additional identifying power comes from the conditional independence between $Y_i(d)$ and X_i given U_i .

It directly follows Theorem 1 that any functional of the joint distribution of $Y_i(1)$ and $Y_i(0)$ is identified: e.g., $Var(Y_i(1) - Y_i(0))$, $Pr(Y_i(1) \ge Y_i(0))$, $Pr(Y_i(1) \ge Y_i(0)|Y_i(0))$ and etc. The rest of the section discusses the restrictions on the joint distribution of $Y_i(1)$ and $Y_i(0)$ implied by the identifying assumptions and a testable implication of the identifying assumptions which proposes a falsification test.

2.1 Restriction on the joint distribution

Assumption 2 assumes that there exists a latent variable U_i which contains sufficient information on the dependence between a treated potential outcome and an untreated potential outcome. Assumption 3.b and Assumption 4.c assume that the proxy variable X_i and Z_i create sufficient variation in the latent variable U_i . By assuming X_i , Z_i and U_i are scalar variables, I exclude the trivial case of $U_i = (Y_i(1), Y_i(0))$ and impose implicit restrictions on the joint distribution of $Y_i(1)$ and $Y_i(0)$.

To discuss the implicit restrictions imposed by the identifying assumptions, let us consider a simple quantity of $\mathbf{E}[Y_i(1)Y_i(0)]$. $\mathbf{E}[Y_i(1)Y_i(0)]$ is a key ingredient in identifying $\text{Var}(Y_i(1) - Y_i(0))$, a measure of the treatment effect heterogeneity. In most econometric frameworks that identify ATE or QTE, $\mathbf{E}[Y_i(1)Y_i(0)]$ still remains unidentified. In this paper, using Assumption 3.b or Assumption 4.c, the conditional density of $Y_i(1)$ given $Y_i(0)$ is identified as a weighted average

of the conditional densities of Y_i given $(D_i = 1, Z_i)$, identifying $\mathbf{E}[Y_i(1)Y_i(0)]$. The core idea in constructing the weights is that the conditional density $f_{Y(1)|U}$ is identified as a weighted average of $\{f_{Y|D=1,Z}(\cdot|z)\}_z$, from the completeness of $L_{U|D=1,Z}$. With $w(\cdot,\cdot)$ denoting the weighting function,

$$f_{Y(1)|Y(0)}(\cdot|y) = \int_{\mathbb{R}} \frac{w(y,z)}{f_{Y(0)}(y)} \cdot f_{Y|D=1,Z}(\cdot|z)dz$$

and thus

$$\mathbf{E}[Y_i(1)|Y_i(0) = y] = \int_{\mathbb{R}} \frac{w(y,z)}{f_{Y(0)}(y)} \cdot \mathbf{E}[Y_i|D_i = 1, Z_i = z]dz.$$

 $\mathbf{E}[Y_i(1)Y_i(0)]$ is identified as

$$\begin{aligned} \mathbf{E}[Y_i(1)Y_i(0)] &= \mathbf{E}[\mathbf{E}[Y_i(1)|Y_i(0)] \cdot Y_i(0)] = \int_{\mathbb{R}} w(y,z) \cdot \mathbf{E}[Y_i|D_i = 1, Z_i = z] y dy dz \\ &= \mathbf{E}\left[\frac{w(Y_i(0), Z_i)}{f_{Y_i(0), Z}(Y_i(0), Z_i)} \cdot \mathbf{E}[Y_i|D_i = 1, Z_i] Y_i(0)\right]. \end{aligned}$$

Note that $\mathbf{E}[Y_i(1)Y_i(0)]$ is identified as an expected product of two random variables $Y_i(0)$ and $\mathbf{E}[Y_i|D_i=1,Z_i]$, reweighted with $\frac{w}{f_{Y(0),Z}}$. Even though we do not observe $Y_i(1)$ and $Y_i(0)$ simultaneously, the result above shows us that we can instead use $\mathbf{E}[Y_i|D_i=1,Z_i]$, a random variable that is observed for every untreated unit, in place of $Y_i(1)$ and reweight the joint density of $Y_i(0)$ and Z_i with $w(\cdot,\cdot)$. Thus, the implicit restriction in identifying $\mathbf{E}[Y_i(1)Y_i(0)]$ is that the conditional expectation $\mathbf{E}[Y_i(1)|Y_i(0)=y]$ must be spanned by the observed conditional expectations $\{\mathbf{E}[Y_i|D_i=1,Z_i=z]\}_z$. Since the above identification argument can be rewritten with $\mathbf{E}[Y_i(0)|Y_i(1)=y]$, another implicit restriction is that the conditional expectation $\mathbf{E}[Y_i(0)|Y_i(1)=y]$ must be spanned by the observed conditional expectations $\{\mathbf{E}[Y_i|D_i=0,Z_i=z]\}_z$. The identification argument can also be extended to conditional densities, instead of conditional expectations; thus, more generally, the implicit restriction imposed on the joint distribution of $Y_i(1)$ and $Y_i(0)$ is that the conditional distribution of $Y_i(1)$ given $Y_i(0)$ must be spanned by the conditional distribution of Y_i given $(D_i=1,Z_i)$ and vice versa.

2.2 Testable implication

When we extend Assumption 5 so that both $u \mapsto Mf_{Y(1)|U}(\cdot|u)$ and $u \mapsto Mf_{Y(0)|U}(\cdot|u)$ are strictly increasing and continuously differentiable, we have a testable implication of Assumptions 1-2 and 4-5, from over-identification. Suppose that $\mathbf{E}[Y_i(1)|U_i=u]$ and $\mathbf{E}[Y_i(0)|U_i=u]$ are strictly increasing in u. Then, the conditional densities $(f_{Y(1)|U}, f_{X|U}, f_{U|D=1,Z})$ are identified in the treated

subsample and the conditional densities $(f_{Y(0)|U}, f_{X|U}, f_{U|D=0,Z})$ are identified in the untreated subsample. Let $f_{X|D=1,U}$ denote the conditional density of X_i given U_i , identified from the treated subsample and likewise for $f_{X|D=0,U}$. Then, Assumption 1 imposes that

$$\min_{\tilde{g}:\text{monotone}} \mathbf{E} \left[\int_{\mathbb{R}} \left(f_{X|D=1,U}(x|U_i) - f_{X|D=0,U}(x|\tilde{g}(U_i)) \right)^2 dx \middle| D_i = 1 \right] = 0$$
 (7)

since $f_{X|D=1,U} = f_{X|D=0,U}$. In (7), a monotone function \tilde{g} is used to connect the identification result from the treated subpopulation to the untreated subpopulation, now that $f_{X|U}$ is not used to connect the two identification results. A test that uses (7) as a null can be used as a falsification test on the framework proposed in this paper.

What does a test on the null (7) exactly test? The mixture model on the conditional density $f_{Y,X|D=d,Z}$ assumes that conditioning on U_i , the potential outcome $Y_i(d)$ and the proxy variable X_i are independent of each other. Recall that in Example 2, the proxy variable X_i is a past outcome. Thus, in the panel context, we can understand the falsification test as testing whether we can find a latent variable U_i conditioning on which the outcomes are intertemporally independent. Note that the key identifying assumption is that the potential outcomes independent across the treatment status. While the conditional independence assumption across the treatment status remains untestable due to the limitation that we only observe either a treated potential outcome or a untreated potential outcome for a given unit, the falsification test in Example 2 tests if the outcomes are intertemporally independent, conditioning on some latent variable.

In the case of discrete U_i , Assumption 5 was not used in the identification. In fact, without introducing any further assumptions, we have a testable implication:

$$\sum_{k=1}^{K} \min_{k'} \sum_{j=1}^{M_X} \left(\Pr\left\{ X_i = x^j \middle| (D_i, U_i) = (1, u^k) \right\} - \Pr\left\{ X_i = x^j \middle| (D_i, U_i) = (0, u^k) \right\} \right)^2 = 0.$$
 (8)

I develop an asymptotic theory in the next section under the finite support assumption on U_i , formally proposing a falsification test.

3 Implementation

Based on the identification result for discrete U_i , I estimate the conditional density of $Y_i(1)$ and $Y_i(0)$ given U_i , by assuming a finite support for U_i and solving a nonnegative matrix factorization (NMF) problem. The focus on the case of discrete U_i has several reasons. Firstly, a dicretization

is often used in econometric models with latent heterogeneity as an approximation to a continuous latent heterogeneity space: see Bonhomme et al. (2022) for more. Secondly, with parametrization, the estimation of infinite-dimensional objects such as conditional densities $f_{U|D=0,Z}$ and $f_{U|D=1,Z}$ becomes an estimation of finite-dimensional objects Λ_0 and Λ_1 , giving us \sqrt{n} rate. The \sqrt{n} rate becomes helpful in deriving an asymptotic distribution for the distributional treatment effect estimators. Lastly, the linearity induced from discretization reduces the computational burden substantially. This does not mean that there is no feasible estimation method for continuous U_i . For a continuous latent variable case, we can construct a sieve maximum likelihood estimator, as suggested in the nonclassical measurement error literature. The specifics are discussed in the appendix subsection A.2.

The parameters of interest in this paper are the joint distribution of the potential outcomes $Y_i(1)$ and $Y_i(0)$ and the marginal distribution of the treatment effect $Y_i(1) - Y_i(0)$. To estimate these distributional treatment effect (DTE) parameters, I first estimate the conditional probabilites of U_i given Z_i , namely the mixture weight matrices Λ_0 and Λ_1 in the finite mixture interpretation, by solving a nonnegative matrix factorization problem. Given the first step estimators on Λ_0 and Λ_1 , I characterize the the joint distribution of $Y_i(1)$ and $Y_i(0)$ and the marginal distribution of $Y_i(1) - Y_i(0)$ as quadratic moments and estimate the distributions by plugging in the first step estimates to the induced U-statistics. In doing so, to account for the estimation error from the first step, I orthogonalize the score function. Neyman orthogonality makes the plug-in estimator robust to the first step estimation error and helps derive a limiting distribution for the estimator.

All of the discussions in this section assume that K is known. In practice, we often do not have a priori choice of K. Thus, in Appendix B, I discuss how we could utilize the existing econometric methods such as the eigenvalue ratio estimator from Ahn and Horenstein (2013) and the rank test from Kleibergen and Paap (2006) for guidance on the choice of K.

3.1 Nonnegative matrix factorization

To estimate the mixture weight matrices Λ_0 and Λ_1 from (6), I first let $M_Z = K$ by using a partition on \mathbb{R} when the support of Z_i has more than K points and construct sample analogues of

the conditional probability matrices \mathbf{H}_0 and \mathbf{H}_1 defined in the previous section: for d=0,1, let

$$\mathbb{H}_{d} = \begin{pmatrix} \frac{\sum_{i=1}^{n} \mathbf{1} \left\{ (Y_{i}, D_{i}, X_{i}, Z_{i}) = \left(y^{1}, d, x^{1}, z^{1}\right) \right\}}{\sum_{i=1}^{n} \mathbf{1} \left\{ (D_{i}, Z_{i}) = (d, z^{1}) \right\}} & \cdots & \frac{\sum_{i=1}^{n} \mathbf{1} \left\{ (Y_{i}, D_{i}, X_{i}, Z_{i}) = \left(y^{1}, d, x^{1}, z^{K}\right) \right\}}{\sum_{i=1}^{n} \mathbf{1} \left\{ (D_{i}, Z_{i}) = (d, z^{K}) \right\}} \\ & \vdots & \ddots & \vdots \\ \frac{\sum_{i=1}^{n} \mathbf{1} \left\{ (Y_{i}, D_{i}, X_{i}, Z_{i}) = \left(y^{MY}, d, x^{MX}, z^{1}\right) \right\}}{\sum_{i=1}^{n} \mathbf{1} \left\{ (D_{i}, Z_{i}) = (d, z^{1}) \right\}} & \cdots & \frac{\sum_{i=1}^{n} \mathbf{1} \left\{ (Y_{i}, D_{i}, X_{i}, Z_{i}) = \left(y^{MY}, d, x^{MX}, z^{K}\right) \right\}}{\sum_{i=1}^{n} \mathbf{1} \left\{ (D_{i}, Z_{i}) = (d, z^{K}) \right\}} \end{pmatrix}.$$

Each column of \mathbb{H}_0 is a conditional empirical distribution function of (Y_i, X_i) given $(D_i = 0, Z_i)$ and each column of \mathbb{H}_1 is a conditional empirical distribution function of (Y_i, X_i) given $(D_i = 1, Z_i)$. As discussed in Section 2, I use partitioning on \mathbb{R} in constructing \mathbb{H}_0 and \mathbb{H}_1 when any of Y_i, X_i and Z_i is continuous.

To estimate Λ_0 and Λ_1 , I formulate a nonnegative matrix factorization problem. Let ι_x be a x-dimensional column vector of ones. Then, the nonnegative matrix factorization problem is constructed as follows:

$$\min_{\Lambda_0, \Lambda_1, \Gamma_0, \Gamma_1} \| \mathbb{H}_0 - \Gamma_0 \Lambda_0 \|_F^2 + \| \mathbb{H}_1 - \Gamma_1 \Lambda_1 \|_F^2$$
(9)

subject to linear constraints that

$$\Lambda_0 \in \mathbb{R}_+^{K \times K}, \quad \Lambda_1 \in \mathbb{R}_+^{K \times K}, \quad \Gamma_0 \in \mathbb{R}_+^{M \times K}, \quad \Gamma_1 \in \mathbb{R}_+^{M \times K},$$

$$\iota_K^{\mathsf{T}} \Lambda_0 = \iota_K^{\mathsf{T}}, \quad \iota_K^{\mathsf{T}} \Lambda_1 = \iota_K^{\mathsf{T}}, \quad \iota_M^{\mathsf{T}} \Gamma_0 = \iota_K^{\mathsf{T}}, \quad \iota_M^{\mathsf{T}} \Gamma_1 = \iota_K^{\mathsf{T}}$$

and quadratic constraints that

$$\Pr\left\{ (Y_i(d), X_i) = (y, x) | U_i = u^k \right\}$$

$$= \left(\sum_{k=1}^{M_X} \Pr\left\{ \left(Y_i(d), X_i \right) = \left(y, x^k \right) | U_i = u^k \right\} \right) \cdot \left(\sum_{j=1}^{M_Y} \Pr\left\{ \left(Y_i(d), X_i \right) = \left(y^j, x \right) | U_i = u^k \right\} \right)$$
(10)

for each (y,x). The linear constraints are probabilities being nonnegative and summing to one. The quadratic constraints are X_i satisfying the exclusion restriction from Assumption 2. When \mathbb{H}_0 and \mathbb{H}_1 are sufficiently close to \mathbf{H}_0 and \mathbf{H}_1 , the identification result discussed in the previous section says that there is a unique decomposition of \mathbb{H}_0 and \mathbb{H}_1 which satisfies the linear and the quadratic constraints.

Note that the objective function in (9) is quadratic when we fix either (Λ_0, Λ_1) or (Γ_0, Γ_1) . Moreover, Γ_0 and Γ_1 can be further decomposed into three matrices $\Gamma_X, \Gamma_{Y(0)}, \Gamma_{Y(1)}$, each of which correponds to the conditional probabilities of X_i given U_i , $Y_i(0)$ given U_i , and $Y_i(1)$ given U_i , respectively. Let $\Gamma_d(\cdot,\cdot)$ denote how Γ_X and $\Gamma_{Y(d)}$ recover Γ_d : $\Gamma_d = \Gamma_d \left(\Gamma_X, \Gamma_{Y(d)}\right)$. The quadratic constraints are trivially imposed by optimizing over $\Gamma_X, \Gamma_{Y(0)}$ and $\Gamma_{Y(1)}$. Using these, I propose an iterative algorithm to solve the minimization problem.

- 1. Initialize $\Gamma_0^{(0)}, \Gamma_1^{(0)}$.
- **2.** (Update Λ) Given $\left(\Gamma_0^{(s)}, \Gamma_1^{(s)}\right)$, solve the following quadratic program:

$$\left(\Lambda_{0}^{(s+1)}, \Lambda_{1}^{(s+1)}\right) = \arg\min_{\Lambda_{0}, \Lambda_{1}} \left\| \mathbb{H}_{0} - \Gamma_{0}^{(s)} \Lambda_{0} \right\|_{F}^{2} + \left\| \mathbb{H}_{1} - \Gamma_{1}^{(s)} \Lambda_{1} \right\|_{F}^{2}$$

subject to $\Lambda_0 \in \mathbb{R}_+{}^{K \times K}, \Lambda_1 \in \mathbb{R}_+{}^{K \times K}, \iota_K{}^\intercal \Lambda_0 = \iota_K{}^\intercal$ and $\iota_K{}^\intercal \Lambda_1 = \iota_K{}^\intercal$.

3. $(Update\ \Gamma_X)$ Given $\left(\Lambda_0^{(s+1)}, \Lambda_1^{(s+1)}, \Gamma_{Y(0)}^{(s)}, \Gamma_{Y(1)}^{(s)}\right)$, solve the following quadratic program:

$$\left(\Gamma_{X}^{(s+1)}\right) = \arg\min_{\Gamma_{Y}} \left\| \mathbb{H}_{0} - \Gamma_{0}\left(\Gamma_{X}, \Gamma_{Y(0)}^{(s)}\right) \Lambda_{0}^{(s+1)} \right\|_{F}^{2} + \left\| \mathbb{H}_{1} - \Gamma_{1}\left(\Gamma_{X}, \Gamma_{Y(1)}^{(s)}\right) \Lambda_{1}^{(s+1)} \right\|_{F}^{2}$$

subject to $\Gamma_X \in \mathbb{R}_+^{M_X \times K}, \iota_{M_X}^{\mathsf{T}} \Gamma_X = \iota_K^{\mathsf{T}}.$

4. (Update Γ_Y) Given $\left(\Lambda_0^{(s+1)}, \Lambda_1^{(s+1)}, \Gamma_X^{(s+1)}\right)$, solve the following quadratic program:

$$\begin{split} & \left(\Gamma_{Y(0)}^{(s+1)}, \Gamma_{Y(0)}^{(s+1)}\right) \\ &= \arg\min_{\Gamma_{Y(0)}, \Gamma_{Y(1)}} \left\| \mathbb{H}_0 - \Gamma_0 \left(\Gamma_X^{(s+1)}, \Gamma_{Y(0)}\right) \Lambda_0^{(s+1)} \right\|_F^2 + \left\| \mathbb{H}_1 - \Gamma_1 \left(\Gamma_X^{(s+1)}, \Gamma_{Y(1)}\right) \Lambda_1^{(s+1)} \right\|_F^2 \end{split}$$

$$\text{subject to } \Gamma_{Y(0)} \in \mathbb{R_+}^{M_Y \times K}, \Gamma_{Y(1)} \in \mathbb{R_+}^{M_Y \times K}, \iota_{M_Y} {}^{\mathsf{T}} \Gamma_{Y(0)} = \iota_K {}^{\mathsf{T}}, \iota_{M_Y} {}^{\mathsf{T}} \Gamma_{Y(1)} = \iota_K {}^{\mathsf{T}}.$$

5. Repeat **2-4** until convergence.

Each step of the iteration is a quadratic programming with linear constraints, which can be solved with a built-in optimization tool in most statistical softwares. The stepwise optimization assures a convergence to a local minimum. To find the global minimum, I consider various initial values $\left(\Gamma_0^{(0)}, \Gamma_1^{(0)}\right)$.

Let $\widehat{\Lambda}_0$, $\widehat{\Lambda}_1$, $\widehat{\Gamma}_0$ and $\widehat{\Gamma}_1$ denote the solution to the minimization problem. Note that when Y_i and X_i are discrete, the estimates $\widehat{\Gamma}_0$ and $\widehat{\Gamma}_1$ directly estimate the conditional distribution of $Y_i(1)$ and

⁶To initialize $\Gamma_0^{(0)}$, $\Gamma_1^{(1)}$, I consider columns from \mathbb{H}_d and weighted sums of columns of \mathbb{H}_d with randomly drawn K sets of weights that sum to one as initial values. Alternatively, we can select the eigenvectors associated with the first K largest eigenvalues of $\mathbb{H}_d^{\mathsf{T}}\mathbb{H}_d$ as an initial value.

 $Y_i(0)$ given U_i . When Y_i are X_i are continuous and therefore partitioning was used in constructing $\mathbf{H}_0, \mathbf{H}_1$, we use $\widehat{\Lambda}_0$ and $\widehat{\Lambda}_1$ to estimate the distribution of $Y_i(1)$ and $Y_i(0)$ given U_i .

3.2 Distributional treatment effect estimators and falsification test

Given the estimates of the two mixture weights matrices Λ_0 and Λ_1 , I construct an estimator for the joint distribution of $Y_i(1)$ and $Y_i(0)$ and the marginal distribution of $Y_i(1) - Y_i(0)$. Firstly, find that for any $y \in \mathbb{R}$,

$$\left(F_{Y|D=d,Z}(y|z^1) \quad \cdots \quad F_{Y|D=d,Z}(y|z^K)\right) = \left(F_{Y(d)|U}(y|u^1) \quad \cdots \quad F_{Y(d)|U}(y|u^K)\right)\Lambda_d$$

Since Λ_d is full rank, we have

$$\left(F_{Y(d)|U}(y|u^1) \quad \cdots \quad F_{Y(d)|U}(y|u^K)\right) = \left(F_{Y|D=d,Z}(y|z^1) \quad \cdots \quad F_{Y|D=d,Z}(y|z^K)\right) (\Lambda_d)^{-1}.$$

The conditional distribution of $F_{Y(d)|U}(\cdot|u)$ is identified as a linear combination of the observed distributions $\{F_{Y|D=d,Z}(\cdot|z)\}_{z=1}^{K}$. Building on this, let

$$\tilde{\Lambda}_d = (\Lambda_d)^{-1}$$

for d = 0, 1. Let $\tilde{\lambda}_{jk,d}$ denote the j-th row and k-th column component of $\tilde{\Lambda}_d$. $\left(\tilde{\lambda}_{1k,d}, \cdots, \tilde{\lambda}_{Kk,d}\right)^{\mathsf{T}}$, the k-th column of $\tilde{\Lambda}_d$, is a set of linear coefficients on $\{F_{Y|D=d,Z}(\cdot|z)\}_{z=1}^K$ to retrieve the conditional distribution of $Y_i(d)$ given $U_i = u^k$. Using the estimators on Λ_0, Λ_1 from the nonnegative matrix factorization, we estimate the linear coefficients as follows:

$$\widehat{\widetilde{\Lambda}}_d = \left(\widehat{\Lambda}_d\right)^{-1}$$

for d = 0, 1.

Secondly, the distribution of U_i is also identified from Λ_0 and Λ_1 :

$$\begin{pmatrix}
\Pr\{U_{i} = u^{1}\} \\
\vdots \\
\Pr\{U_{i} = u^{K}\}
\end{pmatrix} = \Lambda_{0} \begin{pmatrix}
\Pr\{D_{i} = 0, Z_{i} = z^{1}\} \\
\vdots \\
\Pr\{D_{i} = 0, Z_{i} = z^{K}\}
\end{pmatrix} + \Lambda_{1} \begin{pmatrix}
\Pr\{D_{i} = 1, Z_{i} = z^{1}\} \\
\vdots \\
\Pr\{D_{i} = 1, Z_{i} = z^{K}\}
\end{pmatrix}.$$
(11)

Let $p_U(k)$ denote $\Pr\{U_i=u^k\}$ for $k=1,\cdots,K$ and let $p_{D,Z}(d,j)$ denote $\Pr\{D_i=d,Z_i=z^j\}$ for

d=0,1 and $j=1,\cdots,K$. Then, I estimate p_U and $p_{D,Z}$ with

$$\hat{p}_{D,Z}(d,j) = \frac{1}{n} \sum_{i=1}^{n} \mathbf{1} \{ D_i = d, Z_i = z^j \}$$

and

$$\hat{p}_{U} = \begin{pmatrix} \hat{p}_{U}(1) \\ \vdots \\ \hat{p}_{U}(K) \end{pmatrix} = \hat{\Lambda}_{0} \begin{pmatrix} \frac{1}{n} \sum_{i=1}^{n} \mathbf{1} \{D_{i} = 0, Z_{i} = z^{1} \} \\ \vdots \\ \frac{1}{n} \sum_{i=1}^{n} \mathbf{1} \{D_{i} = 0, Z_{i} = z^{K} \} \end{pmatrix} + \hat{\Lambda}_{1} \begin{pmatrix} \frac{1}{n} \sum_{i=1}^{n} \mathbf{1} \{D_{i} = 1, Z_{i} = z^{1} \} \\ \vdots \\ \frac{1}{n} \sum_{i=1}^{n} \mathbf{1} \{D_{i} = 1, Z_{i} = z^{K} \} \end{pmatrix}.$$

By combining the two results, we get

$$F_{Y(0),Y(1)}(y,y') = \sum_{k=1}^{K} p_{U}(k) F_{Y(0)}(y) F_{Y(1)}(y')$$

$$= \sum_{k=1}^{K} p_{U}(k) \left(\sum_{j=1}^{K} \tilde{\lambda}_{jk,0} F_{Y|D=0,Z}(y|z^{j}) \right) \cdot \left(\sum_{j'=1}^{K} \tilde{\lambda}_{j'k,1} F_{Y|D=1,Z}(y'|z^{j'}) \right)$$

$$= \sum_{j=1}^{K} \sum_{j'=1}^{K} \left(\sum_{k=1}^{K} p_{U}(k) \tilde{\lambda}_{jk,0} \tilde{\lambda}_{j'k,1} \right) F_{Y|D=0,Z}(y|z^{j}) \cdot F_{Y|D=1,Z}(y'|z^{j'}).$$

Using this characterization, I estimate the joint distribution of $Y_i(1)$ and $Y_i(0)$ as a linear combination of $\{F_{Y|D=0,Z}(y|z^j)\cdot F_{Y|D=1,Z}(y'|z^{j'})\}_{j,j'}$ where the weights are computed with $\widehat{\Lambda}_0, \widehat{\Lambda}_1$ and $\{\widehat{p}_{D,Z}(d,j)\}_{d,j}$. We can derive a similar result for the marginal distribution of $Y_i(1) - Y_i(0)$: for any $\delta \in \mathbb{R}$,

$$F_{Y(1)-Y(0)|U}(\delta|u) = \int_{\mathbb{R}} F_{Y(1)|U}(y+\delta|u) \cdot f_{Y(0)|U}(y|u) dy,$$

$$F_{Y(1)-Y(0)}(\delta) = \sum_{j=1}^{K} \sum_{j'=1}^{K} \left(\sum_{k=1}^{K} p_{U}(k) \tilde{\lambda}_{jk,0} \tilde{\lambda}_{j'k,1} \right) \int_{\mathbb{R}} F_{Y|D=1,U}(y+\delta|z^{j}) \cdot f_{Y|D=0,U}(y|z^{j'}) dy.$$

Both parameters of interest are identified as a weighted sum of quantities that are indexed by pairs of subpopulations $\{i: D_i = 0, Z_i = z^j\}$ and $\{i: D_i = 1, Z_i = z^{j'}\}$. As shown above, weights are estimated from the first step nonnegative matrix factorization and empirical measures of the subpopulations.

Likewise, we can find a similar characterization of the conditional distributions used in the

falsification test suggested in Subsection 2.2: for d = 0, 1 and $k = 1, \dots, K$,

$$F_{X|U}(x|u^k) = \sum_{j=1}^K \tilde{\lambda}_{jk,d} F_{X|D=d,Z}(x|z^j).$$

Again, the conditional distributions are identified as a weighted sum of quantities that are indexed by subpopulations $\{i: D_i = d, Z_i = z^j\}$. Since the distributions are conditional on U_i , the weights are estimated only from the first step nonnegative matrix factorization.

Now, I show that the characterization of the two DTE parameters and the conditional distribution of X_i given $(D_i = d, U_i)$ for the falsification test can be transformed into moment conditions with nuisance parameters $\tilde{\lambda}_0, \tilde{\Lambda}_1, p_U, p_{D,Z}$. Then, to be robust to the first step estimation error in the nuisance parameters, I orthogonalize the moment. I will discuss the moment condition and the orthogonalization for the marginal distribution of $Y_i(1) - Y_i(0)$; the case for the joint distribution of $Y_i(1)$ and $Y_i(0)$ and that for the conditional probability of X_i given $(D_i = d, U_i)$ follow naturally.

For some δ , let

$$\theta = F_{Y(1)-Y(0)}(\delta).$$

Firstly, find that θ is a summation over K treated subpopulations and K untreated subpopulations. Fix j, j' and let

$$\theta_{jj'} := \left(\sum_{k=1}^{K} p_U(k) \tilde{\lambda}_{jk,0} \tilde{\lambda}_{j'k,1} \right) \int_{\mathbb{R}} F_{Y|D=1,U}(y+\delta|z^j) \cdot f_{Y|D=0,U}(y|z^{j'}) dy.$$

Find that

$$\int_{\mathbb{R}} F_{Y|D=1,U}(y+\delta|z^{j}) \cdot f_{Y|D=0,U}(y|z^{j'}) dy = \frac{\mathbf{E}\left[\mathbf{1}\{Y_{i'} \leq Y_{i} + \delta, D_{i} = 0, Z_{i} = z^{j}, D_{i} = 1, Z_{i'} = z^{j'}\}\right]}{\mathbf{E}\left[\mathbf{1}\{D_{i} = 0, Z_{i} = z^{j}, D_{i'} = 1, Z_{i'} = z^{j'}\}\right]}$$

with $(Y_i, D_i, Z_i) \perp \!\!\! \perp (Y_{i'}, D_{i'}, Z_{i'})$. Thus, $\theta_{jj'}$ is identified from a quadratic moment

$$\mathbf{E}\left[m_{jj'}\left(W_i,W_{i'};\theta_{jj'},\tilde{\Lambda}_0,\tilde{\Lambda}_1,\{p_U(k)\}_k,\{p_{D,Z}(d,j)\}_{d,j}\right)\right]=0$$

where $W_i = (Y_i, D_i, X_i, Z_i)$ and

$$\begin{split} m_{jj'}\left(W_{i},W_{i'};\theta_{jj'},\tilde{\Lambda}_{0},\tilde{\Lambda}_{1},\{p_{U}(k)\}_{k},\{p_{D,Z}(d,j)\}_{d,j}\right) \\ &= \frac{\sum_{k=1}^{K}p_{U}(k)\tilde{\lambda}_{jk,0}\tilde{\lambda}_{j'k,1}}{p_{D,Z}(0,j)\cdot p_{D,Z}(1,j')}\cdot \left(\frac{1}{2}\mathbf{1}\{Y_{i'}\leq Y_{i}+\delta,D_{i}=0,Z_{i}=z^{j},D_{i}=1,Z_{i'}=z^{j'}\}\right. \\ &\qquad \qquad + \frac{1}{2}\mathbf{1}\{Y_{i}\leq Y_{i'}+\delta,D_{i}=1,Z_{i}=z^{j'},D_{i}=0,Z_{i'}=z^{j}\}\right) - \theta_{jj'}. \end{split}$$

By summing over j and j', we can construct a moment function $m = \sum_{j=1}^K \sum_{j'=1}^K m_{jj'}$ such that

$$\mathbf{E}\left[m\left(W_i, W_{i'}; \theta, \tilde{\Lambda}_0, \tilde{\Lambda}_1, \{p_U(k)\}_k, \{p_{D,Z}(d,j)\}_{d,j}\right)\right] = 0$$

identifies θ .

If the nuisance parameters $\tilde{\Lambda}_0, \tilde{\Lambda}_1, p_U, p_{D,Z}$ were known, the standard asymptotic theory of U statistic would apply to the GMM estimator of θ using $\mathbf{E}[m(W_i, W_{i'}; \theta)] = 0$ as the moment condition. However, in practice, we use first step estimates for the nuisance parameters. Thus, to account for the first step estimation error, we orthogonalize the moment function. Even though the NMF estimators $(\hat{\Lambda}_0, \hat{\Lambda}_1)$ and the induced estimators $(\hat{\Lambda}_0, \hat{\Lambda}_1)$ are complex nonlinear functions of the data matrix \mathbb{H}_0 and \mathbb{H}_1 , $(\tilde{\Lambda}_0, \tilde{\Lambda}_1)$ satisfy the following equations at their true values: for all (y, d, x, k),

$$\sum_{j=1}^{K} \tilde{\lambda}_{jk,d} \Pr \left\{ Y_i = y, X_i = x | D_i = d, Z_i = z^j \right\}$$

$$= \sum_{j=1}^{K} \tilde{\lambda}_{jk,d} \Pr \left\{ Y_i = y | D_i = d, Z_i = z^j \right\} \cdot \sum_{j=1}^{K} \tilde{\lambda}_{jk,d} \Pr \left\{ X_i = x | D_i = d, Z_i = z^j \right\}$$

$$\Pr \left\{ X_i = x \right\} = \sum_{k=1}^{K} p_U(k) \sum_{j=1}^{K} \tilde{\lambda}_{jk,d} \Pr \left\{ X_i = x | D_i = d, Z_i = z^j \right\}.$$
(13)

Equation (12) corresponds to the conditional independence assumption that

$$\Pr\{Y_i(d) = y, X_i = x | U_i = u\} = \Pr\{Y_i(d) = y | U_i = u\} \cdot \Pr\{X_i = x | U_i = u\}.$$

and Equation (13) corresponds to the law of iterated expectation that

$$\Pr\{X_i = x\} = \sum_{k=1}^K p_U(k) \Pr\{X_i = x | U_i = u^k\}.$$

Given $\{p_{D,Z}(d,j)\}_{d,j}$, Equation (12) can be written as a quadratic moment condition and Equation (13) as a linear moment condition. I use these additional moments in orthogonalizing the moment m so that the Neyman orthogonality holds.

Let $\tilde{\lambda}$ and p denote vectorizations of $(\tilde{\Lambda}_0, \tilde{\Lambda}_1)$ and $(\{p_U(k)\}_k, \{p_{D,Z}(d,j)\}_{d,j})$. The orthogonalized score is constructed with the additional moment function

 ϕ simply collects the quadratic moments from (12) across (y, d, x, k), the linear moments from (13) across (d, x), and the linear moment

$$p_{D,Z}(d,j) = \mathbf{E}[\mathbf{1}\{D_i = d, Z_i = z^j\}]$$
 (14)

across (d, j). To complete the orthogonalization, I show that the Jacobian matrix of ϕ has full rank.

Lemma 1. Assumptions 1-3 hold. Then,

$$\begin{pmatrix}
\mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} \phi(W_i, W_{i'}; \tilde{\lambda}, p) \right] \\
\mathbf{E} \left[\frac{\partial}{\partial p} \phi(W_i, W_{i'}; \tilde{\lambda}, p) \right]
\end{pmatrix}$$

has a full rank.

Proof. See Appendix. \Box

Then, with an additional nuisance parameter

$$\mu = \begin{pmatrix} \mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} \phi(W_i, W_{i'}; \tilde{\lambda}, p) \right] \\ \mathbf{E} \left[\frac{\partial}{\partial p} \phi(W_i, W_{i'}; \tilde{\lambda}, p) \right] \end{pmatrix}^{\mathsf{T}} \\ \cdot \begin{pmatrix} \left[\mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} \phi(W_i, W_{i'}; \tilde{\lambda}, p) \right] \\ \mathbf{E} \left[\frac{\partial}{\partial p} \phi(W_i, W_{i'}; \tilde{\lambda}, p) \right] \end{pmatrix} \begin{pmatrix} \mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} \phi(W_i, W_{i'}; \tilde{\lambda}, p) \right] \\ \mathbf{E} \left[\frac{\partial}{\partial p} \phi(W_i, W_{i'}; \tilde{\lambda}, p) \right] \end{pmatrix}^{\mathsf{T}} \end{pmatrix}^{\mathsf{T}} \\ \cdot \begin{pmatrix} \mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} m(W_i, W_{i'}; \tilde{\lambda}, p) \right] \\ \mathbf{E} \left[\frac{\partial}{\partial p} m(W_i, W_{i'}; \tilde{\lambda}, p) \right] \end{pmatrix},$$

the orthogonalized score

$$\psi(W_i, W_{i'}; \theta, \tilde{\lambda}, p, \mu) = m(W_i, W_{i'}; \theta, \tilde{\lambda}, p) - \mu^{\mathsf{T}} \phi(W_i, W_{i'}; \tilde{\lambda}, p)$$

satisfies the Neyman orthogonality. μ is estimated by taking a sample analogue of the expression above. Given estimators $(\hat{\lambda}, \hat{p}, \hat{\mu})$, I estimate θ with

$$\binom{n}{2}^{-1} \sum_{i < i'} \psi\left(W_i, W_{i'}; \hat{\theta}, \hat{\tilde{\lambda}}, \hat{p}, \hat{\mu}\right) = 0.$$

The same procedure applies to $F_{Y(0),Y(1)}(y,y')$.

In the case of the conditional distribution of X_i given $(D_i = d, U_i)$, there are a few deviations. Firstly, I solve the nonnegative matrix factorization problem without imposing the exclusion restriction, by optimizing over $\Gamma_{X,D=1}$, $\Gamma_{X,D=0}$. Then, the moment condition

$$\mathbf{E}\left[\sum_{j=1}^{K} \frac{\tilde{\lambda}_{jk}}{\mu_{D,Z}(d,j)} \mathbf{1}\{D_i = d, X_i \le x, Z_i = z^j\} - F_{X|U}(x|u^k)\right] = 0$$

is orthogonalized with (12) and (14). The auxiliary moment (13) is not used here since the nuisance parameter p_U is not used. Once I have estimators from each subpopulation, I extend Assumption 5 to find a labeling on U_i across treated subpopulation and untreated subpopulation. For example, if I assume Assumption 5 holds with conditional expectation both for $f_{Y(1)|U}$ and $f_{Y(0)|U}$, I estimate $\{\mathbf{E}[Y_i(d)|U_i=u^k]\}_{k=1}^K$ and relabel $\{f_{X|D=d,U}(\cdot|u^k)\}_{k=1}^K$ based on the estimates.

 $\widehat{F}_{Y(0),Y(1)}$ and $\widehat{F}_{Y(1)-Y(0)}$ denote the distributional treatment effect estimators and $\widehat{F}_{X|D=d,U}$ denotes the conditional distribution estimator.

3.3 Asymptotic properties

Theorem 2 establishes the consistency of the mixture weight estimators $\hat{\Lambda}_0$ and $\hat{\Lambda}_1$.

Theorem 2. Assumptions 1-3 hold. Up to some permutation on $\{u^1, \dots, u^K\}$,

$$\left\| \widehat{\Lambda}_0 - \Lambda_0 \right\|_F = O_p \left(\frac{1}{\sqrt{n}} \right) \quad and \quad \left\| \widehat{\Lambda}_1 - \Lambda_1 \right\|_F = O_p \left(\frac{1}{\sqrt{n}} \right)$$

as $n \to \infty$.

Proof. See Appendix. \Box

A direct corollary of Theorem 2 is that $\widehat{\tilde{\Lambda}}_0$, $\widehat{\tilde{\Lambda}}_1$ are consistent for $\tilde{\Lambda}_0$ and $\tilde{\Lambda}_1$ at the rate of $\frac{1}{\sqrt{n}}$. Theorem 3 establishes the asymptotic normality of the distributional treatment effect estimators.

Theorem 3. Assumptions 1-3 hold. Then, for any $(y, y') \in \mathbb{R}^2$ and $\delta \in \mathbb{R}$,

$$\sqrt{n}\left(\widehat{F}_{Y(0),Y(1)}(y,y') - F_{Y(0),Y(1)}(y,y')\right) \xrightarrow{d} \mathcal{N}\left(0,\sigma(y,y')^{2}\right)$$

$$\sqrt{n}\left(\widehat{F}_{Y(1)-Y(0)}(\delta) - F_{Y(1)-Y(0)}(\delta)\right) \xrightarrow{d} \mathcal{N}\left(0,\sigma(\delta)^{2}\right)$$

as $n \to \infty$. Additionally, Assumption 5 holds with both $f_{Y(1)|U}$ and $f_{Y(0)|U}$ in a strict sense that

$$\mathbf{E}[Y_i(d)|U_i = u^k] < \mathbf{E}[Y_i(d)|U_i = u^{k+1}]$$

for every d=0,1 and $k=1,\cdots,K-1$. Then, for any $(x^1,\cdots,x^{\tilde{M}})\in\mathbb{R}$,

$$\sqrt{n} \begin{pmatrix} \widehat{F}_{X|D=0,U}(x^{1}|u^{1}) \\ \widehat{F}_{X|D=1,U}(x^{1}|u^{1}) \\ \vdots \\ \widehat{F}_{X|D=1,U}(x^{\tilde{M}}|u^{K}) \end{pmatrix} - \begin{pmatrix} F_{X|U}(x^{1}|u^{1}) \\ F_{X|U}(x^{1}|u^{1}) \\ \vdots \\ F_{X|U}(x^{\tilde{M}}|u^{K}) \end{pmatrix} \xrightarrow{d} \mathcal{N} \left(\boldsymbol{o}_{2\tilde{M}K}, \Sigma \right)$$

Proof. See Appendix.

The asymptotic variances are computed from a projection of the orthogonal scores:

$$\tilde{\psi}(w) = \mathbf{E}\left[\psi(W_i, w)\right]$$
 and $\sigma^2 = \mathbf{E}\left[\tilde{\psi}(W_i)^2\right]$.

In Sections 4-5, the standard error is obtained with a plug-in estimator for the asymptotic variance.

A direct corollary of Theorem 3 is the asymptotic validity of the falsification test. Firstly, fix some $x^1 < \cdots < x^{\tilde{M}}$ such that $F_{X|U}(x^1|u) > 0$ and $F_{X|U}(x^{\tilde{M}}|u) < 1$ for every u. The test statistic for the falsification test is

$$T_{n} = n \begin{pmatrix} \widehat{F}_{X|D=1,U}(x^{1}|u^{1}) - \widehat{F}_{X|D=0,U}(x^{1}|u^{1}) \\ \vdots \\ 1 - \widehat{F}_{X|D=1,U}(x^{\tilde{M}}|u^{K}) - \left(1 - \widehat{F}_{X|D=0,U}(x^{\tilde{M}}|u^{K})\right) \end{pmatrix}^{\mathsf{T}} \widetilde{\Sigma}^{-1} \\ \cdot \begin{pmatrix} \widehat{F}_{X|D=1,U}(x^{1}|u^{1}) - \widehat{F}_{X|D=0,U}(x^{1}|u^{1}) \\ \vdots \\ 1 - \widehat{F}_{X|D=1,U}(x^{\tilde{M}}|u^{K}) - \left(1 - \widehat{F}_{X|D=0,U}(x^{\tilde{M}}|u^{K})\right) \end{pmatrix}$$
(15)

where $\tilde{\Sigma} = A \Sigma A^{\dagger}$ with

$$A = \begin{pmatrix} -1 & 1 & 0 & 0 & \cdots & 0 & 0 \\ 1 & -1 & -1 & 1 & \cdots & 0 & 0 \\ \vdots & \vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & 0 & 0 & 0 & \cdots & 1 & -1 \end{pmatrix}.$$

 $\tilde{\Sigma}$ can be consistently estimated.

Corollary 1. Assumptions 1-3 hold and Assumption 5 holds with both $f_{Y(1)|U}$ and $f_{Y(0)|U}$ in a strict sense that

$$\mathbf{E}[Y_i(d)|U_i = u^k] < \mathbf{E}[Y_i(d)|U_i = u^{k+1}]$$

for every d = 0, 1 and $k = 1, \dots, K - 1$. Then,

$$T_n \stackrel{d}{\to} \chi^2(K\tilde{M})$$

as $n \to \infty$.

The test statistic is constructed with $K(\tilde{M}+1)$ -dimensional vector but the degree of freedom is $K\tilde{M}$ since the conditional probabilities sum to one.

4 Simulation

In this section, I discuss Monte Carlo simulation results. I generated B = 200 random samples from DGPs with discrete $Y_i(1), Y_i(0), X_i, Z_i$ and U_i where $M_Y = 3, M_X = 6, M_Z = 3$ and K = 3: $Y_i \in \{1, 2, 3\}, X_i \in \{1, 2, 3, 4, 5, 6\}$ and $Z_i \in \{1, 2, 3\}$. The treatment D_i was drawn randomly, independent of $Y_i(1), Y_i(0), X_i, Z_i$. In the first step nonnegative matrix factorization, I collapsed the support of X_i so that the effective number of points in the support of X_i is three. Thus, the conditional probability matrix \mathbb{H}_0 and \mathbb{H}_1 were 9×3 matrices. Across difference DGPs, I varied Λ , the conditional probability of U_i given Z_i which is shared across treated and untreated subpopulation, to vary the informativeness of the proxy variable Z_i with regard to the latent variable U_i .

Table 1 contains the bias and the root mean squared error (rMSE) of the distributional treatment effect estimators $\widehat{F}_{Y(1)-Y(0)}$. As Λ becomes less informative about the distribution of U_i , i.e. the smallest singular value $\sigma_{\min}(\Lambda)$ decreases, the rMSE goes up. This suggests that the first step nonnegative matrix factorization estimation quality depends on how informative the proxy variables X_i and Z_i are for the latent variable U_i . Additionally, Table 2 contains the coverage probability

$$\Gamma_X = \begin{pmatrix} 0.778 & 0.028 & 0.022 \\ 0.067 & 0.050 & 0.033 \\ 0.056 & 0.422 & 0.044 \\ 0.044 & 0.422 & 0.056 \\ 0.033 & 0.050 & 0.067 \\ 0.022 & 0.028 & 0.778 \end{pmatrix}, \quad \Gamma_{Y(1)} = \begin{pmatrix} 0.656 & 0.022 & 0.000 \\ 0.117 & 0.706 & 0.117 \\ 0.228 & 0.272 & 0.883 \end{pmatrix}, \quad \Gamma_{Y(0)} = \begin{pmatrix} 0.756 & 0.122 & 0.078 \\ 0.167 & 0.756 & 0.167 \\ 0.078 & 0.122 & 0.756 \end{pmatrix},$$

and Λ s in the order of decreasing smallest singular value are

$$\Lambda = \begin{pmatrix} 0.840 & 0.091 & 0.040 \\ 0.077 & 0.772 & 0.056 \\ 0.083 & 0.137 & 0.905 \end{pmatrix}, \quad \begin{pmatrix} 0.722 & 0.134 & 0.078 \\ 0.124 & 0.665 & 0.095 \\ 0.154 & 0.201 & 0.827 \end{pmatrix}, \quad \begin{pmatrix} 0.611 & 0.175 & 0.120 \\ 0.168 & 0.563 & 0.137 \\ 0.221 & 0.262 & 0.744 \end{pmatrix}$$

$\widehat{F}_{Y(1)-Y(0)}$									
	$\sigma_{\min}(\Lambda) = 0.701$		$\sigma_{\min}(\Lambda)$	= 0.501	$\sigma_{\min}(\Lambda)$	$\sigma_{\min}(\Lambda) = 0.310$			
δ	bias	rMSE	bias	rMSE	bias	rMSE			
-2	0.000	0.006	0.001	0.010	0.001	0.025			
-1	-0.000	0.017	0.000	0.025	-0.002	0.052			
0	-0.007	0.028	-0.012	0.040	-0.014	0.076			
1	-0.009	0.025	-0.014	0.040	-0.015	0.084			

Table 1: Bias and rMSE of DTE estimator, B = 200.

⁷The specifics of the DGPs are as follows: $p_U = (0.286, 0.286, 0.438)$,

		$\widehat{F}_{Y(1)-Y(0)}$	
	$\sigma_{\min}(\Lambda) = 0.701$	$\sigma_{\min}(\Lambda) = 0.501$	$\sigma_{\min}(\Lambda) = 0.310$
$\Pr\left\{F_{Y(1)-Y(0)}(-2) \in \widehat{CI}\right\}$	0.968	0.970	0.990
$\Pr\left\{F_{Y(1)-Y(0)}(-1) \in \widehat{CI}\right\}$	0.978	0.960	0.970
$\Pr\left\{F_{Y(1)-Y(0)}(0) \in \widehat{CI}\right\}$	0.960	0.975	0.990
$\Pr\left\{F_{Y(1)-Y(0)}(1) \in \widehat{CI}\right\}$	0.970	0.970	0.980
$\Pr\left\{\text{reject } F_{X D=1,U} = F_{X D=0,U}\right\}$	0.070	0.063	0.049

Table 2: Coverage of CI and type I error of falsification test, B = 200.

of the confidence interval constructed with the asymptotic standard error and the type I error of the falsification test proposed in Subsection 2.2. The 95% confidence interval shows mostly correct coverage, sometimes slightly too conservative, and the falsification test is valid.

5 Empirical illustration

In this section, we revisit Jones et al. (2019) and estimate the distributional treatment effect of workplace wellness program on medical spending. Jointly with the Campus Well-being Services at the University of Illinois Urbana-Champaign, the authors of Jones et al. (2019) conducted a large-scale randomized controlled trials. The experiment started in July 2016, by inviting 12,459 eligible university employees to participate in an online survey. Of 4,834 employees who completed the survey, 3,300 employees were randomly selected into treatment, being offered to participate in a workplace wellness program names iThrive. The participation itself was not enforced; the treated individuals were merely financially incentivized to participate by being offered monetary reward for completing each step of the wellness program. Thus, the main treatment effect parameter of Jones et al. (2019) is the 'intent-to-treat' effect. The workplace wellness program consisted of various activities such as chronic disease management, weight management, and etc. The treated individuals were offered to participate in the wellness program starting the fall semester of 2016, until the spring semester of 2018.

One of the main outcome variables that Jones et al. (2019) studied is the monthly medical spending. Since the authors had access to the university-sponsored health insurance data, they had detailed information on the medical spending behaviors of the participants. Taking advantage of the randomness in assigning eligibility to the participants, Jones et al. (2019) estimated the intent-to-treat type ATE of the workplace wellness program on the monthly medical spending.

The ATE estimate on the first-year monthly medical spending, from August 2016 to July 2017, showed that the eligibility for the wellness program raised the monthly medical spending by \$10.8, with p-value of 0.937, finding no significant intent-to-treat effect.

In Jones et al. (2019), the authors acknowledge that the null effect on the mean does not necessarily mean null effect everywhere, though they themselves do not explore the treatment effect heterogeneity in the paper.⁸ On page 1890, Jones et al. (2019) state "there may exist subpopulations who did benefit from the intervention or who would have benefitted hard they participated." I build onto this observation and estimate the distributional treatment effect of the randomly assigned eligibility for the wellness program. By looking at the distribution, I find the proportion of the subpopulation among treated population that benefitted from the treatment.

The dataset built by the authors of Jones et al. (2019) fits the context of the short panel model in Example 2. For each individual, the dataset contains monthly medical spending records for the following three time durations: July 2015-July 2016, August 2016-July 2017 and August 2017-January 2019. Since the experiment started in the summer of 2016 and the treated individuals were offered to participate in the wellness program starting the fall semester of 2016, the monthly medical spending record for July 2015-July 2016 could be thought of as a 'pretreatment' outcome variable. Thus, we could use the information from the distribution of the pretreatment outcome variable to connect the treated subsample and the untreated subsample. The followings are the variables taken from the dataset.

 Y_i : monthly medical spending for August 2016-July 2017

 D_i : a binary variable for whether eligible to participate in the wellness program

 X_i : monthly medical spending for July 2015-July 2016

 \mathbb{Z}_i : monthly medical spending for August 2017-January 2019

In this specific empirical context, the common shock V_{it} could be thought of as underlying health status and the treatment-status-specific shocks $(\varepsilon_{it}(1), \varepsilon_{it}(0))$ could be thought of as additional random shocks such as susceptibility to the workplace wellness program or transient health shock

 $^{^{8}}$ In the original dataset used in Jones et al. (2019), the authors had connected the medical spending variables to additional survey variables such as age, health behavior, salary, etc. They did not explore how the treatment effect interacts with the additional characteristics, but they did add these additional control variables through double Lasso. Adding the control variables increased the point estimate for the ATE (\$34.9) but the estimate still remained insignificant, with p-value being 0.859.

⁹Damon Jones, David Molitor, and Julian Reif, "What do workplace wellness programs do? Evidence from the Illinois workplace wellness study," *The Quarterly Journal of Economics*, vol. 134, no.4 (2019): 1747-1791.

which does not persist over time. The first-order Markovian assumption in Example 2 is consistent with the health economics literature and broader economics literature of modeling household choices regarding health expenditure: Grossman (1972); Wagstaff (1993); Jacobson (2000); Yogo (2016) and more. Applying assumptions in Example 2, the treatment is allowed to affect the underlying health status in the post-treatment period of August 2017-January 2019, but is assumed to be independent of the underlying health status in July 2015-July 2017.

To choose, K, the size of the support for the latent variable U_i , I applied the eigenvalue ratio estimator and the Kleibergen-Paap rank test to a 12×10 matrix of $\Pr\left\{X_i \in \mathcal{X}^l | D_i = d, Z_i \in \mathcal{Z}^j\right\}$. Under Assumptions 1-3, the matrix is at most rank K. Given a matrix, the eigenvalue ratio estimator from Ahn and Horenstein (2013) estimates K by choosing the value of K maximizes eigenvalue/growth ratio and the Kleibergen-Paap test statistic tests the null hypothesis of rank K against the alternative of rank being bigger than K. For detailed discussion, see Appendix B.

K	1	2	3	4	5	6	7	8
eigenvalue ratio	3.505	3.991	4.029	2.721	1.653	1.863	1.418	3.309
growth ratio	0.964	1.135	1.472	1.353	0.893	0.956	0.580	1.035

Table 3: Eigenvalue ratios and growth ratios

K	1	2	3	4	5	6
test statistic	884.82	116.23	35.75	20.08	13.80	7.94
<i>p</i> -value	0.000	0.001	0.984	0.998	0.995	0.992

Table 4: Kleibergen-Paap rank test statistics for H_0 : rank = K and their p-values

Both the eigenvalue ratio estimator and the Kleibergen-Paap rank test suggest K=3. To be conservative, I used K=5 for the main estimation specification. Thus, when constructing \mathbb{H}_0 and \mathbb{H}_1 to be used in the first step nonnegative matrix factorization, I used the quintiles of the marginal distributions: $(-\infty, F_Y^{-1}(0.2), F_Y^{-1}(0.4), F_Y^{-1}(0.6), F_Y^{-1}(0.8), \infty)$ and similarly for X_i . The matrices \mathbb{H}_0 and \mathbb{H}_1 are 25×5 -dimensional.

With K=5, the conditional independence framework of the paper passes the falsification test. T_n from (15) is computed with $(x^1, \dots, x^4) = (F_X^{-1}(0.2), F_X^{-1}(0.4), F_X^{-1}(0.6), F_X^{-1}(0.8))$ and

The partition for X_i is constructed with $F_X^{-1}(1/12), \dots, F_X^{-1}(11/12)$ and the partition for Z_i is constructed with $F_Z^{-1}(1/5), \dots, F_Z^{-1}(4/5)$.

therefore

$$T_n \stackrel{d}{\to} \chi^2(20)$$

as $n \to \infty$. T_n is 16.435 and its *p*-value is 0.689, passing the falsification test.

Figure 1 contains the estimated joint distribution of the two potential outcomes from the non-negative matrix factorization algorithm with K=5. For visibility, I first partitioned the potential outcome variable with quantiles $F_Y^{-1}(1/7), \dots, F_Y^{-1}(6/7)$ and plotted the joint distribution of partitioned potential outcomes. Since the treated potential outcomes are plotted on the vertical axis, higher mass on the left-upper triangle means that the treatment reduces the medical spending. Overall, there is no definitive pattern. One notable observation is that the joint density is higher where $F_Y(Y_i(1)) \approx F_Y(Y_i(0)) \approx 0$ and $F_Y(Y_i(1)) \approx F_Y(Y_i(0)) \approx 1$. This is intuitive since on the two ends of the underlying health status spectrum, the effectiveness of the workplace wellness program must be limited.

Figure 2 contains the estimated marginal distribution of the treatment effect and its 95% pointwise confidence interval. Note that the point estimates are mostly upward-sloping and lie between zero and one. Though the quadratic moment representation used in the DTE estimators does not impose any monotonicity or nonnegativity restrictions, the estimated marginal distribution violates these constraints only on a small subset of the range [-1000, 1000]. Overall, it is unclear if more than half of the people would be better off from the treatment; the confidence interval for

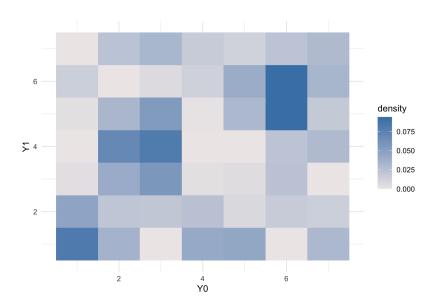


Figure 1: Joint density of $F_Y(Y_i(1))$ and $F_Y(Y_i(0))$, K = 5.

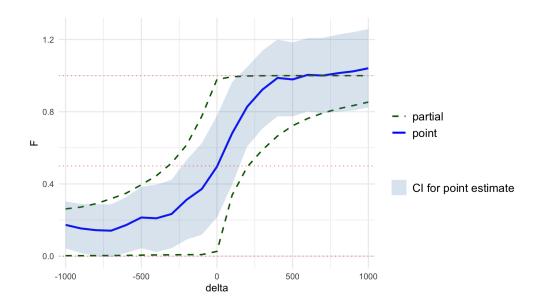


Figure 2: Marginal distribution of $Y_i(1) - Y_i(0)$, K = 5.

 $\Pr\{Y_i(1) - Y_i(0) \ge 0\}$ contains 0.5, not being able to reject the null $\Pr\{Y_i(1) - Y_i(0) \ge 0\} \le 0.5$.

As comparison, estimates for the upper bound and the lower bound from Makarov (1982); Fan and Park (2010) are also provided in Figure 2, as green dotted lines. The point estimates are consistent with the partial identification result, lying between the lower bound and the upper bound. The comparison highlights the gain of the point identification result, at the cost of assuming stronger identifying assumptions. For $\delta \in [-500, 600]$, the 95% confidence interval is included in the partially identified set, giving us much bigger power in inference.

Lastly, the point identification helps us analyze the pattern of the treatment heterogeneity. Recall that the ATE estimate was inconclusive about the effectiveness of the treatment. However, the DTE estimates on $\Pr\{Y_i(1) - Y_i(0) \le \delta\}$ for $\delta \le -600$ and the DTE estimates on $\Pr\{Y_i(1) - Y_i(0) \le \delta\}$ for $\delta \ge 400$ shows us interesting treatment effect heterogeneity patterns, in favor of implementing the treatment. The negative impact of the treatment, i.e. how much more money you spend under the treatment, is capped at \$400: $\widehat{F}_{Y(1)-Y(0)}(400) \approx 1$. On the other hand, the left tail of the treatment effect distribution is thicker, implying that some people are greatly benefitted from participating in the program: $\widehat{F}_{Y(1)-Y(0)}(-600) \approx 0.15$.

6 Conclusion

This paper presents an identification result for the joint distribution of treated potential outcome and untreated potential outcome, given conditionally random binary treatment. The key assumptions in the identification are that there exists a latent variable that captures the dependence between the two potential outcomes and that there exist two proxy variables for the latent variable. By assuming strict monotonicity for some functional of the conditional distribution of potential outcomes given the latent variable, I interpret the latent variable as 'latent rank' and strict monotonicity as 'latent rank invariance.' In implementation, I propose a first step nonnegative matrix factorization and a second step plug-in GMM. \sqrt{n} -consistency of the first-step estimator and the asymptotic normality of the second step GMM estimator are established. Lastly, I apply the estimation method to revisit Jones et al. (2019) and find that the potential medical spendings are positively correlated at the two ends of the support and the marginal distribution of the treatment effect has thicker left tail.

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APPENDIX

A Discussion on a continuous latent variable

A.1 Identification

Assumptions 1-2 are powerful enough for us to apply the known spectral decomposition results with proxy variables (see Hu (2008); Hu and Schennach (2008) and more) to each of the treated subsample and the untreated subsample. Let $f_{Y=y,X|D=d,Z}(x|z)$ denote the conditional density of (Y_i, X_i) given (D_i, Z_i) evaluated at $Y_i = y$ and $D_i = d$; the density has only two arguments x and z. Likewise, let $f_{U|D=d,Z}$ denote the conditional density of U_i given (D_i, Z_i) evaluated at $D_i = d$. From Assumptions 1-2, we obtain the following integral representation: for $x, z \in \mathbb{R}$,

$$f_{Y=y,X|D=d,Z}(x|z) = \int_{\mathcal{U}} f_{Y(d),X|D=d,Z,U}(y,x|z,u) \cdot f_{U|D=d,Z}(u|z) du$$

$$= \int_{\mathcal{U}} f_{Y(d),X|U}(y,x|u) \cdot f_{U|D=d,Z}(u|z) du \quad \therefore \text{ Assumption 1}$$

$$= \int_{\mathcal{U}} f_{Y(d)|U}(y|u) \cdot f_{X|U}(x|u) \cdot f_{U|D=d,Z}(u|z) du \quad \therefore \text{ Assumption 2}$$

$$f_{X|D=d,Z}(x|z) = \int f_{X|U}(x|u) \cdot f_{U|D=d,Z}(u|z) du.$$

$$(16)$$

To discuss the spectral decomposition result of Hu and Schennach (2008), let us construct integral operators $L_{X|U}$, $L_{U|D=d,Z}$ and a diagonal operator $\Delta_{Y(d)=y|U}$ which map a function in $\mathcal{L}^1(\mathbb{R})$ to a function in $\mathcal{L}^1(\mathbb{R})$:

$$\begin{aligned} \left[L_{X|U}g\right](x) &= \int_{\mathbb{R}} f_{X|U}(x|u)g(u)du, \\ \left[L_{U|D=d,Z}g\right](u) &= \int_{\mathbb{R}} f_{U|D=d,Z}(u|z)g(z)dz, \\ \left[\Delta_{Y(1)=y|U}g\right](u) &= f_{Y(1)|U}(y|u)g(u). \end{aligned}$$

For example, when g is a density, $L_{X|U}$ takes the density g as a marginal density of U_i and maps it to a marginal density of X_i , implied by $f_{X|U}$ and g. Define $L_{Y=y,X|D=d,Z}$ and $L_{X|D=d,Z}$ similarly,

with the conditional density $f_{Y=y,X|D=d,X}$ and $f_{X|D=d,Z}$. Then,

$$L_{Y=y,X|D=d,Z} = L_{X|U} \cdot \Delta_{Y(d)|U} \cdot L_{U|D=d,Z},$$

$$L_{X|D=d,Z} = L_{X|U} \cdot L_{U|D=d,Z}.$$

To get to a spectral decomposition result, we additionally assume that the conditional density $f_{X|D=d,Z}$ is complete. The completeness assumption imposes restriction on the proxy variables X_i and Z_i ; the conditional density of U_i given Z_i , within each subsample, should preserve the variation in the conditional density of X_i given U_i . With completeness condition on the conditional density $f_{X|D=d,Z}$, we can define an inverse of the integral operator $L_{X|D=d,Z}$ and therefore obtain a spectral decomposition:

$$L_{Y=y,X|D=d,Z} \cdot (L_{X|D=d,Z})^{-1} = L_{X|U} \cdot \Delta_{Y(d)=y|U} \cdot (L_{X|U})^{-1}$$
.

The RHS of the equation above admits a spectral decomposition with $\{f_{X|U}(\cdot|u)\}_u$ as eigenfunctions and $\{f_{Y(d)|U}(y|u)\}_u$ as eigenvalues.

However, the individual spectral decomposition results on the two subsamples by themselves are not enough to identify the joint distribution of the potential outcomes. To connect the two spectral decomposition results, we resort to Assumption 1. Under Assumption 1, the conditional density of X_i given U_i is identical across the two subsamples. Thus, the two decomposition results should admit the same density functions $\{f_{X|U}(\cdot|u)\}_u$ as eigenfunctions. Using this, we connect the eigenvalues of the two decompositions; we identify $\{f_{Y(1)|U}(\cdot|u) \cdot f_{Y(0)|U}(\cdot|u)\}_u$.

Lastly, to find the marginal distribution of U_i , we fully invoke the latent rank interpretation and assume that there is some functional M defined on $\mathcal{L}^1(\mathbb{R}^2)$ such that $Mf_{Y(d)|U}(\cdot|u)$ is strictly increasing in u, with some d=0,1. An example of such a functional is expectation:

$$Mf = \int_{\mathbb{R}} y f(y) dy.$$

When $Mf_{Y(1)|U}(\cdot|u)$ and $Mf_{Y(0)|U}(\cdot|u)$ are both strictly increasing in u, the latent rank invariance holds in a truer sense that U_i determines the rank of $\mathbf{E}[Y_i(1)|U_i]$ and the rank of $\mathbf{E}[Y_i(0)|U_i]$ and that the two ranks coincide. The latent rank assumption finds on ordering on the eigenfunctions $\{f_{X|U}(\cdot|u)\}_u$ using information from $\{f_{Y(1)|U}(\cdot|u)\}_u$ or $\{f_{Y(0)|U}(\cdot|u)\}_u$ and allows us to use a transformation on U_i without precisely locating U_i .

A.2 Sieve maximum likelihood

To estimate the conditional densities of interest, i.e. $f_{Y(1)|U}$, $f_{Y(0)|U}$, $f_{X|U}$, $f_{U|D=1,Z}$, $f_{U|D=0,Z}$, we again utilize the decomposition given in (5). Especially, with U_i being a continuous random variable, the decomposition can be rewritten as an integration:

$$f_{Y,X|D,Z}(y,x|d,z) = \int_{\mathcal{U}} f_{Y(d)|U}(y|u) \cdot f_{X|U}(x|u) \cdot f_{U|D=d,Z}(u|z) du.$$

Given some sieves to approximate the conditional densities, characterized with finite-dimensional parameters $\theta = (\theta_1, \theta_0, \theta_X, \theta_{1Z}, \theta_{0Z})$, the sieve ML estimator is:

$$\hat{\theta} = \arg\max_{\theta \in \Theta_n} \sum_{i=1}^n \log f_{Y,X|D,Z,n}(Y_i, X_i|D_i, Z_i; \theta)$$

$$= \arg\max_{\theta \in \Theta_n} \sum_{i=1}^n \left(D_i \log \int_{\mathcal{U}} f_{Y(1)|U,n}(Y_i|u; \theta_1) \cdot f_{X|U,n}(X_i|u; \theta_X) \cdot f_{U|D=1,Z,n}(u|Z_i; \theta_{1Z}) du \right)$$

$$(1 - D_i) \log \int_{\mathcal{U}} f_{Y(0)|U,n}(Y_i|u; \theta_0) \cdot f_{X|U,n}(X_i|u; \theta_X) \cdot f_{U|D=0,Z,n}(u|Z_i; \theta_{0Z}) du$$

$$(1 - D_i) \log \int_{\mathcal{U}} f_{Y(0)|U,n}(Y_i|u; \theta_0) \cdot f_{X|U,n}(X_i|u; \theta_X) \cdot f_{U|D=0,Z,n}(u|Z_i; \theta_{0Z}) du$$

In particular, we propose tensor product spaces of Bernstein polynomials as sieves $\{\Theta_n\}_{n=1}^{\infty}$. For example, the conditional density $f_{Y(1)|U}$ approximated to a tensor product space with a given dimension of $(p^y + 1, p^u + 1)$ is as follows: with y normalized to be on [0, 1],

$$f_{Y(1)|U,n}(y|u;\theta_1) = \sum_{j=0}^{p^y} \sum_{k=0}^{p^u} \theta_{jk,1} \binom{p^y}{j} y^j (1-y)^{p^y-j} \cdot \binom{p^u}{k} u^k (1-u)^{p^u-k}$$

and $\theta_1 = \{\theta_{jk,1}\}_{0 \leq j \leq p^j, 0 \leq k \leq p^u}$. The tensor product construction and the properties of Bernstein polynomials make it remarkably straightforward to impose that the approximated functions are densities. Using properties of Berstein polynomials, we can impose that $f_{Y(1)|U,n}(y|u;\theta_1)$ is

¹¹The degree of Bernstein polynomial does not need to be uniform across different conditional densities; for example p^y for $f_{Y(1)|U,n}$ may differ from p^y for $f_{Y(0)|U,n}$. However, p^u being uniform across all five conditional densities facilitates computation.

nonnegative and integrate to one, by imposing that

$$\theta_{jk,1} \ge 0 \quad \forall j, k$$
 (nonnegative)

$$\sum_{i=0}^{p^y} \frac{\theta_{j0,1}}{p^y + 1} = 1$$
 (sum-to-one)

$$\sum_{l=0}^{k} \sum_{j=0}^{p^{y}} \frac{1}{p^{y}+1} (-1)^{k-l} \binom{p^{u}}{k} \binom{k}{l} \theta_{jl,1} = 0 \quad \forall k = 1, \cdots, p^{u} \qquad (sum\text{-}to\text{-}one)$$

Moreover, when the latent rank interpretation from Assumption 5 is assumed with average, the monotonicity condition can be easily imposed as linear constraints. For example, $\mathbf{E}[Y_i(1)|U_i=u]$ being monotone increasing in u translates to

$$\sum_{j=0}^{p^y} w_j \theta_{jk,1} \le \sum_{j=0}^{p^y} w_j \theta_{jk+1,1} \quad \forall k = 0, \dots, p^u - 1$$
 (monotonicity)

Below are the details on the linear constraints that correspond to nonnegativity, sum-to-one and monotonicity. Use the same example from before— $f_{Y(1)|U,n}$ —and find that we can rearrange the approximated function as a univariate Bernstein polynomial of degree p^u by fixing u:

$$f_{Y(1)|U,n}(y|u;\theta_1) = \sum_{j=0}^{p^y} \left(\sum_{k=0}^{p^u} \theta_{jk,1} \binom{p^u}{k} u^k (1-u)^{p^u-k} \right) \binom{p^y}{j} y^j (1-y)^{p^y-j}.$$

 $f_{Y(1)|U,n}(y|u;\theta_1)$ is nonnegative if and only if

$$\sum_{k=0}^{p^{u}} \theta_{jk,1} \binom{p^{u}}{k} u^{k} (1-u)^{p^{u}-k} \ge 0$$

for every $j = 0, \dots, p^y$ at the fixed u. Since $f_{Y(1)|U,n}(y|u;\theta_1)$ needs to be a nonnegative function at any value of u, this translates to $\sum_{k=0}^{p^u} \theta_{jk,1} \binom{p^u}{k} u^k (1-u)^{p^u-k}$, which is a Bernstein polynomial itself, being a nonnegative function. Thus, the nonnegativity constraints become

$$\theta_{jk,1} \ge 0 \quad \forall j, k.$$

Also, find that

$$\int_{0}^{1} f_{Y(1)|U,n}(y|u;\theta_{1}) dy = \sum_{k=0}^{p^{u}} \left(\sum_{j=0}^{p^{y}} \theta_{jk,1} \int_{0}^{1} \sum_{j=0}^{p^{y}} {p^{y} \choose j} y^{j} (1-y)^{p^{y}-j} dy \right) {p^{u} \choose k} u^{k} (1-u)^{p^{u}-k}$$

$$= \sum_{k=0}^{p^{u}} \sum_{j=0}^{p^{y}} \frac{\theta_{jk,1}}{p^{y}+1} {p^{u} \choose k} u^{k} (1-u)^{p^{u}-k}.$$

For $\int_0^1 f_{Y(1)|U,n}(y|u;\theta_1)dy=1$ to hold uniformly over u, $\sum_{k=0}^{p^u} \sum_{j=0}^{p^y} \frac{\theta_{jk,1}}{p^y+1} {p^u \choose k} u^k (1-u)^{p^u-k}$ must be constant in u and equal to one. Again, $\sum_{k=0}^{p^u} \sum_{j=0}^{p^y} \frac{\theta_{jk,1}}{p^y+1} {p^u \choose k} u^k (1-u)^{p^u-k}$ is a Bernstein polynomial itself and can be transformed to a sum of monomials:

Thus, the sum-to-one constraints are

$$\sum_{j=0}^{p^y} \frac{\theta_{j0,1}}{p^y + 1} = 1,$$

$$\sum_{l=0}^k \sum_{j=0}^{p^y} \frac{1}{p^y + 1} (-1)^{k-l} \binom{p^u}{k} \binom{k}{l} \theta_{jl,1} = 0 \quad \forall k = 1, \dots, p^u.$$

Lastly, for the monotonicity constraint, find that

$$\int_{0}^{1} y f_{Y(1)|U,n}(y|u;\theta_{1}) dy = \sum_{k=0}^{p^{u}} \underbrace{\left(\sum_{j=0}^{p^{y}} \theta_{jk,1} \int_{0}^{1} \binom{p^{y}}{j} y^{j+1} (1-y)^{p^{y}-j} dy\right)}_{=:\theta_{\cdot k,1}} \binom{p^{u}}{k} u^{k} (1-u)^{p^{u}-k}$$

Again, the conditional expectation is also a Berstein polynomial and it is monotone increasing if and only if $\theta_{\cdot k,1} \leq \theta_{\cdot k+1,1}$ for $k = 0, \dots, p^u - 1$. By applying the monomial transformation again,

we get

$$\binom{p^y}{j} y^{j+1} (1-y)^{p^y-j} = \binom{p^y}{j} \binom{p^y+1}{j+1}^{-1} \sum_{l=j+1}^{p^y+1} (-1)^{l-j-l} \binom{p^y+1}{j+1} \binom{j+1}{l} u^l,$$

$$\int_0^1 \binom{p^y}{j} y^{j+1} (1-y)^{p^y-j} dy = \frac{j+1}{p^y+1} \sum_{l=j+1}^{p^y+1} (-1)^{l-j-l} \binom{p^y+1}{j+1} \binom{j+1}{l} \frac{1}{l+1} =: w_j.$$

The monotonicity constraints are

$$\sum_{j=0}^{p^{y}} w_{j} \theta_{jk,1} \leq \sum_{j=0}^{p^{y}} w_{j} \theta_{jk+1,1} \quad \forall k = 0, \dots, p^{u} - 1.$$

Now, we discuss how to estimate the distributional treatment effect parameters. Unlike the nonnegative matrix factorization estimator, the sieve ML estimator fully estimates the five conditional densities. Thus, an estimator on the joint distribution of the potential outcomes and the marginal distribution of treatment effect can be directly constructed from $\hat{\theta}$. For example, the joint density estimator can be constructed as follows: for any (y, y'),

$$\widehat{F}_{Y(1),Y(0)}(y,y') = \frac{1}{n} \sum_{i=1}^{n} \int_{\mathcal{U}} \int_{-\infty}^{y} \int_{-\infty}^{y'} f_{Y(1)|U,n}(w|u;\hat{\theta}_{1}) \cdot f_{Y(0)|U}(w'|u;\hat{\theta}_{0}) dw dw'$$

$$\cdot \left(D_{i} f_{U|D=1,Z,n}(u|Z_{i};\hat{\theta}_{1Z}) + (1-D_{i}) f_{U|D=0,Z,n}(u|Z_{i};\hat{\theta}_{0Z}) \right) du.$$

Likewise, the marginal treatment effect distribution estimator can be constructed as follows: for any δ ,

$$\widehat{F}_{Y(1)-Y(0)}(\delta) = \frac{1}{n} \sum_{i=1}^{n} \int_{\mathcal{U}} \int_{\mathbb{R}} \int_{-\infty}^{y+\delta} f_{Y(1)|U}(y'|u; \hat{\theta}_{1}) \cdot f_{Y(0)|U}(y|u; \hat{\theta}_{0}) dy' dy$$

$$\cdot \left(D_{i} f_{U|D=1,Z,n} \left(u|Z_{i}; \hat{\theta}_{1Z} \right) + (1 - D_{i}) f_{U|D=0,Z,n} \left(u|Z_{i}; \hat{\theta}_{0Z} \right) \right) du.$$

In constructing induced estimators, the conditional densities $f_{U|D=1,Z}$ and $f_{U|D=0,Z}$ are used to obtain the marginal density of U_i , taking advantage of the following equivalence:

$$\mathbf{E}\left[g(U_i)\right] = \mathbf{E}\left[\mathbf{E}\left[g(U_i)|D_i,Z_i\right]\right].$$

B Choice of K for a discrete latent variable

The finite support assumption in Assumption 3 has definite merits such as significantly alleviating the computational burden compared to Assumption 4. However, the finite support assumption requires researcher to commit to a specific value of K in the estimation, which is a nontrivial task. In the framework of this paper, choosing K to be used in estimation is equivalent to estimating the rank of the matrix \mathbf{H}_0 and \mathbf{H}_1 . Thus, in this section, I introduce existing rank estimation and inference methods in the literature and connect them to the choice of K in the estimation.

As for the low-rank matrix of rank K to discuss choice of K with, I consider a $M_X \times 2M_Z$ matrix \mathbf{H}_X :

$$\mathbf{H}_{X} = \begin{pmatrix} \Pr\left\{X_{i} = x^{1} | (D_{i}, Z_{i}) = (0, z^{1})\right\} & \cdots & \Pr\left\{X_{i} = x^{1} | (D_{i}, Z_{i}) = (1, z^{M_{Z}})\right\} \\ \vdots & \ddots & \vdots \\ \Pr\left\{X_{i} = x^{M_{X}} | (D_{i}, Z_{i}) = (0, z^{1})\right\} & \cdots & \Pr\left\{X_{i} = x^{M_{X}} | (D_{i}, Z_{i}) = (1, z^{M_{Z}})\right\} \end{pmatrix}.$$

Again, when X_i and Z_i are continuous random variables, we may use partitioning on \mathbb{R} to construct such \mathbf{H}_X . The construction of \mathbf{H}_X pools information from the treated subpopulation and the untreated subpopulation; the columnwise concatenation of \mathbf{H}_0 and \mathbf{H}_1 is in general not rank K since the conditional distribution of $Y_i(1)$ given U_i is likely different from that of $Y_i(0)$ given U_i . On the other hand, since the conditional distribution of X_i given U_i is the same across the two subpopulations from Assumption 1, the finite support assumption in Assumption 3.a implies that $\operatorname{rank}(\mathbf{H}_X) \leq K$. In fact, Assumption 3.b implies that $M_X, M_Z \geq K$ and $\operatorname{rank}(\mathbf{H}_X)$ is exactly K. I estimate \mathbf{H}_X with

$$\mathbb{H}_{X} = \begin{pmatrix} \frac{\sum_{i=1}^{n} \mathbf{1} \left\{ (D_{i}, X_{i}, Z_{i}) = (0, x^{1}, z^{1}) \right\}}{\sum_{i=1}^{n} \mathbf{1} \left\{ (D_{i}, Z_{i}) = (0, z^{1}) \right\}} & \cdots & \frac{\sum_{i=1}^{n} \mathbf{1} \left\{ (D_{i}, X_{i}, Z_{i}) = (1, x^{1}, z^{M}Z) \right\}}{\sum_{i=1}^{n} \mathbf{1} \left\{ (D_{i}, Z_{i}) = (1, z^{M}Z) \right\}} \\ & \vdots & \ddots & \vdots \\ \frac{\sum_{i=1}^{n} \mathbf{1} \left\{ (D_{i}, X_{i}, Z_{i}) = (0, x^{M}X, z^{1}) \right\}}{\sum_{i=1}^{n} \mathbf{1} \left\{ (D_{i}, X_{i}, Z_{i}) = (1, x^{M}X, z^{M}Z) \right\}} \\ & \frac{\sum_{i=1}^{n} \mathbf{1} \left\{ (D_{i}, X_{i}, Z_{i}) = (1, z^{M}Z, z^{M}Z) \right\}}{\sum_{i=1}^{n} \mathbf{1} \left\{ (D_{i}, Z_{i}) = (1, z^{M}Z, z^{M}Z) \right\}} \end{pmatrix}.$$

Firstly, to estimate rank(\mathbf{H}_X), I propose the eigenvalue ratio estimator from Ahn and Horenstein (2013). The eigenvalue ratio estimator is developed for a setup where a low-rank matrix of growing dimension is estimated, which is different from the setup of this paper; as n increases, the dimension of \mathbf{H}_X stays the same while the estimation error of \mathbb{H}_X decreases. Higgins (2025) discusses a similar setup to mine where a factor model is assumed for short T panel data and the dimension of the factor model is estimated with a reduced/collapsed data matrix whose dimension is fixed. Following

Higgins (2025)'s treatment, I apply the eigenvalue ratio estimator of Ahn and Horenstein (2013) to \mathbb{H}_X . Let $\nu_k(\mathbf{H})$ denote the k-th largest eigenvalue of $\mathbf{H}\mathbf{H}^{\dagger}$ when $M_X \leq 2M_Z$ and that of $\mathbf{H}^{\dagger}\mathbf{H}$ when $M_X \geq 2M_Z$. The eigenvalue ratio estimator is

$$\widehat{K}_{ER} = \max_{K \le K_{\text{max}}} \frac{\nu_K(\mathbb{H}_X)}{\mu_{K+1}(\mathbb{H}_X)}$$

with $K_{\text{max}} = \min\{M_X, 2M_Z\} - 1$. Similarly, Ahn and Horenstein (2013) also proposes an estimator based on the growth rate of eigenvalues:

$$\widehat{K}_{GR} = \max_{K \le K_{\text{max}}} \frac{\log \left(1 + \frac{\nu_K(\mathbb{H}_X)}{\sum_{k=K+1}^{M_X} \nu_k(\mathbb{H}_X)} \right)}{\log \left(1 + \frac{\nu_{K+1}(\mathbb{H}_X)}{\sum_{k=K+2}^{M_X} \nu_k(\mathbb{H}_X)} \right)}$$

with $K_{\text{max}} = \min\{M_X, 2M_Z\} - 2$. Both estimators are consistent for true K as $n \to \infty$.

Secondly, to infer on rank(\mathbf{H}_X), I use the Kleibergen-Paap (KP) rank test: Kleibergen and Paap (2006). For a given K, the KP rank test tests the null hypothesis H_0 : rank(\mathbf{H}_X) = K against the alternative hypothesis H_1 : rank(\mathbf{H}_X) $\geq K+1$. The KP rank test is developed for a general setup where a low-dimensional, low-rank matrix is estimated with estimators that are asymptotically normal. Since $\text{vec}(\mathbb{H}_X)$ is asymptotically normal around $\text{vec}(\mathbf{H}_X)$, we can directly apply the KP rank test. The construction of the KP rank test statistic is notationally complex but is implemented easily with singular value decomposition. The key element of the KP test statistic is the contruction of a $(M_X - K) \times (2M_Z - K)$ matrix that is a linear transformation of $\text{vec}(\mathbb{H}_X)$ and therefore asymptotically normal. The matrix is all zeros when $\text{rank}(\mathbf{H}_X) = K$ and contains nonzero element when $\text{rank}(\mathbf{H}_X) \geq K+1$. Kleibergen and Paap (2006) provides an explicit formula for the matrix through singular value decomposition. With the KP rank test, I test if there is significant evidence against K chosen to be used in the estimation.

C Proofs

C.1 Proof for Theorem 1

This subsection completes the proof for Theorem 1 under Assumptions 1-2, 4-5, by extending the spectral decomposition result of Hu and Schennach (2008).¹² For the proof of the spectral decom-

¹²The identification under Assumptions 1-3 is straightforward from the discussion in the main text.

position results, refer to Hu and Schennach (2008). By applying assumptions of Hu and Schennach (2008), except their Assumption 5, we have a collection of $\{f_{Y(1)|U}(\cdot|u), f_{Y(0)|U}(\cdot|u), f_{X|U}(\cdot|u)\}_{u\in\mathcal{U}}$, without labeling on u; we have separated the triads of conditional densities for each value of u, but we have not labeled each triad with their respective values of u. To find an ordering on the infinite number of triads, WLOG let $\tilde{U}_i = h(U_i) := M f_{Y(0)|U}(\cdot|U_i)$ and $\tilde{U} = h(\mathcal{U})$. Now, we have labeled each triad with $\tilde{u} = h(u)$ and therefore identified $f_{Y(1)|\tilde{U}}(\cdot|\cdot), f_{Y(0)|\tilde{U}}(\cdot|\cdot)$ and $f_{X|\tilde{U}}(\cdot|\cdot)$. The remainder of the proof constructs conditional densities and a marginal density in terms of the new latent variable \tilde{U}_i as ingredients in identifying the joint density of $Y_i(1)$ and $Y_i(0)$ and shows that the strict monotonicity of h allows us to identify the joint distribution of $Y_i(1)$ and $Y_i(0)$ using \tilde{U}_i instead of U_i .

Firstly, let us establish the injectivity of the integral operator based on the conditional density of X_i given \tilde{U}_i . Find that

$$\begin{split} f_{X|\tilde{U}}(x|\tilde{u}) &= f_{X|U}\left(x|h^{-1}(u)\right) \\ \left[L_{X|\tilde{U}}g\right](x) &= \int_{\tilde{\mathcal{U}}} f_{X|\tilde{U}}(x|\tilde{u})g(\tilde{u})d\tilde{u} = \int_{\tilde{\mathcal{U}}} f_{X|U}\left(x|h^{-1}(\tilde{u})\right)g(\tilde{u})d\tilde{u} \\ &= \int_{\tilde{\mathcal{U}}} f_{X|U}\left(x|h^{-1}(\tilde{u})\right)g\left(h\left(h^{-1}\left(\tilde{u}\right)\right)\right)d\tilde{u} \\ &= \int_{\mathcal{U}} f_{X|U}(x|u)g\left(h(u)\right)h'(u)du, \quad \text{by letting } \tilde{u} = h(u). \end{split}$$

From the completeness of $f_{X|U}$, $L_{X|\tilde{U}}g=0$ implies that g(h(u))h'(u)=0 for almost everywhere on \mathcal{U} . Since h is strictly increasing, h'(u)>0. Thus, we have $g(\tilde{u})=0$ almost everywhere on $\tilde{\mathcal{U}}$: the completeness of $f_{X|\tilde{U}}$ follows. Using the completeness, we identify $f_{\tilde{U}|D=d,Z}$ from

$$f_{X|D=d,Z} = \int_{\mathbb{R}} f_{X|\tilde{U}}(x|\tilde{u}) f_{\tilde{U}|D=d,Z}(\tilde{u}|z) d\tilde{u}.$$

Since the conditional density of Z_i given $D_i = d$ is directly observed, the marginal density of \tilde{U}_i is also identified.

Secondly, it remains to show that the arbitrary choice of \tilde{U}_i does not matter. Under the conditional independence of $Y_i(1)$ and $Y_i(0)$ given U_i , the joint distribution of $Y_i(1)$ and $Y_i(0)$ is a function of three distributions: the conditional distribution of $Y_i(1)$ given U_i , the conditional

distribution of $Y_i(0)$ given U_i and the marginal distribution of U_i . For each $(y_1, y_0) \in \mathbb{R}^2$,

$$\begin{split} f_{Y(1),Y(0)}(y_1,y_0) &= \int_{\mathcal{U}} f_{Y(1)|U}(y_1|u) f_{Y(0)|U}(y_0|u) f_U(u) du \\ &= \int_{\mathcal{U}} f_{Y(1)|\tilde{U}}(y_1|h(u)) f_{Y(0)|\tilde{U}}(y_0|h(u)) f_U(u) du \\ &= \int_{\tilde{U}} f_{Y(1)|\tilde{U}}(y_1|\tilde{u}) f_{Y(0)|\tilde{U}}(y_0|\tilde{u}) \frac{f_U\left(h^{-1}(\tilde{u})\right)}{h'\left(h^{-1}(\tilde{u})\right)} d\tilde{u}, \quad \text{by letting } u = h^{-1}(\tilde{u}) \\ &= \int_{\tilde{U}} f_{Y(1)|\tilde{U}}(y_1|\tilde{u}) f_{Y(0)|\tilde{U}}(y_0|\tilde{u}) f_{\tilde{U}}(\tilde{u}) d\tilde{u}, \quad \text{since } F_U\left(h^{-1}(\tilde{u})\right) = F_{\tilde{U}}(\tilde{u}). \end{split}$$

The last two equalities are from the inverse function theorem: $(h^{-1}(\tilde{u}))' = 1/h'(h^{-1}(\tilde{u}))$. The joint distribution of $Y_i(1)$ and $Y_i(0)$ is identified. The expansion to include (D_i, X_i, Z_i) follows the same argument.

C.2 Proof for Lemma 1

Let us consider three different parts of ϕ : ϕ_A, ϕ_B, ϕ_C . Firstly, ϕ_A is the part of ϕ that corresponds to the quadratic constraints (12). Fix some (y, d, x, k) and let

$$\phi_{A}(W_{i}, W_{i'}; \tilde{\lambda}, p)$$

$$= \sum_{j} \frac{\tilde{\lambda}_{jk,d}}{p_{D,Z}(d,j)} \cdot \frac{\mathbf{1}\{Y_{i} = y, D_{i} = d, X_{i} = x, Z_{i} = z^{j}\} + \mathbf{1}\{Y_{i'} = y, D_{i'} = d, X_{i'} = x, Z_{i'} = z^{j}\}}{2}$$

$$- \sum_{j,j'} \frac{\tilde{\lambda}_{jk,d} \tilde{\lambda}_{j'k,d}}{p_{D,Z}(d,j) \cdot p_{D,Z}(d,j')} \cdot \frac{1}{2} \left(\mathbf{1}\{Y_{i} = y, D_{i} = d, Z_{i} = z^{j}, X_{i'} = x, D_{i'} = d, Z_{i'} = z^{j'}\} \right)$$

$$+ \mathbf{1}\{X_{i} = x, D_{i} = d, Z_{i} = z^{j'}, Y_{i'} = y, D_{i'} = d, Z_{i'} = z^{j}\} \right).$$

Then,

$$\mathbf{E}\left[\frac{\partial}{\partial \tilde{\lambda}_{jk,d}}\phi_{A}(W_{i},W_{i'};\tilde{\lambda},p)\right]$$

$$=\Pr\{Y_{i}=y,X_{i}=x|D_{i}=d,Z_{i}=z^{j}\}-\Pr\{Y_{i}=y|D_{i}=d,Z=z^{j}\}\cdot\Pr\{X_{i}=x|U_{i}=u^{k}\}$$

$$-\Pr\{X_{i}=x|D_{i}=d,Z=z^{j}\}\cdot\Pr\{Y_{i}(d)=y|U_{i}=u^{k}\}$$

and $\mathbf{E}\left[\frac{\partial}{\partial \tilde{\lambda}_{jk',d'}}\phi_A(W_i,W_{i'};\tilde{\lambda},p)\right]$ is zero when $k'\neq k$ or $d'\neq d$. $\mathbf{E}\left[\frac{\partial}{\partial p_U(k)}\phi_A(W_i,W_{i'};\tilde{\lambda},p)\right]=0$ for every k. Lastly,

$$\begin{split} \mathbf{E} \left[\frac{\partial}{\partial p_{D,Z}(d,j)} \phi_A \big(W_i, W_{i'}; \tilde{\lambda}, p \big) \right] \\ &= -\frac{\tilde{\lambda}_{jk,d}}{p_{D,Z}(d,j)} \cdot \Pr\{Y_i = y, X_i = x | D_i = d, Z_i = z^j \} \\ &+ \frac{\tilde{\lambda}_{jk,d}}{p_{D,Z}(d,j)} \cdot \Pr\{Y_i = y | D_i = d, Z_i = z^j \} \cdot \Pr\{X_i = x | U_i = u^k \} \\ &+ \frac{\tilde{\lambda}_{jk,d}}{p_{D,Z}(d,j)} \cdot \Pr\{X_i = x | D_i = d, Z_i = z^j \} \cdot \Pr\{Y_i(d) = y | U_i = u^k \} \end{split}$$

and $\mathbf{E}\left[\frac{\partial}{\partial p_{D,Z}(d',j)}\phi_A(W_i,W_{i'};\tilde{\lambda},p)\right]$ is zero when $d' \neq d$.

Secondly, ϕ_B is the part of ϕ that corresponds to the linear constraints (13). Fix some (d, x) and let

$$\phi_B(W_i, W_{i'}; \tilde{\lambda}, p)$$

$$= \frac{\mathbf{1}\{X_i = x\} + \mathbf{1}\{X_{i'} = x\}}{2}$$

$$- \sum_k p_U(k) \sum_j \frac{\tilde{\lambda}_{jk,d}}{p_{D,Z}(d,j)} \cdot \frac{\mathbf{1}\{D_i = d, X_i = x, Z_i = z^j\} + \mathbf{1}\{D_{i'} = d, X_{i'} = x, Z_{i'} = z^j\}}{2}.$$

Then,

$$\mathbf{E}\left[\frac{\partial}{\partial \tilde{\lambda}_{jk,d}} \phi_B(W_i, W_{i'}; \tilde{\lambda}, p)\right] = -p_U(k) \cdot \Pr\{X_i = x | D_i = d, Z_i = z^j\}$$

and $\mathbf{E}\left[\frac{\partial}{\partial \tilde{\lambda}_{ik,d'}}\phi_B(W_i, W_{i'}; \tilde{\lambda}, p)\right]$ is zero when $d' \neq d$. Also,

$$\mathbf{E}\left[\frac{\partial}{\partial p_{U}(k)}\phi_{B}(W_{i},W_{i'};\tilde{\lambda},p)\right] = -\Pr\{X_{i} = x|U_{i} = u^{k}\}$$

$$\mathbf{E}\left[\frac{\partial}{\partial p_{D,Z}(d,j)}\phi_{B}(W_{i},W_{i'};\tilde{\lambda},p)\right] = \sum_{k=1}^{K} \frac{p_{U}(k)\tilde{\lambda}_{jk,d}}{p_{D,Z}(d,j)} \cdot \Pr\{X_{i} = x|D_{i} = d, Z_{i} = z^{j}\}$$

and $\mathbf{E}\left[\frac{\partial}{\partial p_{D,Z}(d',j)}\phi_B(W_i,W_{i'};\tilde{\lambda},p)\right]$ is zero when $d'\neq d$.

Thirdly, ϕ_C is the moment condition for $p_{D,Z}$. Fix some (d,j) and let

$$\phi_C(W_i, W_{i'}; \tilde{\lambda}, p) = \frac{\mathbf{1}\{D_i = d, Z_i = z^j\} + \mathbf{1}\{D_{i'} = d, Z_{i'} = z^j\}}{2} - p_{D,Z}(d, j).$$

Then, $\mathbf{E}\left[\frac{\partial}{\partial \tilde{\lambda}_{ik.d'}}\phi_{C}\left(W_{i},W_{i'};\tilde{\lambda},p\right)\right]$ and $\mathbf{E}\left[\frac{\partial}{\partial p_{U}(k)}\phi_{C}\left(W_{i},W_{i'};\tilde{\lambda},p\right)\right]$ are zero for every (d',j,k). Also,

$$\mathbf{E}\Big[\frac{\partial}{\partial p_{D,Z}(d,j)}\phi_C(W_i,W_{i'};\tilde{\lambda},p)\Big] = -1$$

and $\mathbf{E}\left[\frac{\partial}{\partial p_{D,Z}(d',j')}\phi_C(W_i,W_{i'};\tilde{\lambda},p)\right]$ is zero when $d'\neq d$ or $j'\neq j$.

The order of ϕ_A , ϕ_B and ϕ_C across different values of (y, x, d, j, k) in ϕ is as follows. Firstly, stack ϕ_A across every value of (y, x) for (d = 0, k = 1) and then for (d = 1, k = 1). Then, repeat this for $k = 2, \dots, K$. These will be the first 2MK components of ϕ . Secondly, stack ϕ_B across every value of x for d = 0 and then for d = 1. These will be the second $2M_X$ components of ϕ . Then, stack ϕ_C across every value of j for d = 0 and then for d = 1. These will be the last 2K components of ϕ .

Also, we need to decide on the order of $\tilde{\lambda}_{jk,d}$ in vectorized $\tilde{\lambda}$ and similarly for p. In a similar manner to ϕ , collect $\tilde{\lambda}_{jk,d}$ across j for (d=0,k=1) and then for (d=1,k=1). Then, repeat this for $k=2,\cdots,K$. These will be the $2K^2$ -dimensional vector $\tilde{\lambda}$. For p, first collect $p_U(k)$ across k, collect $p_{D,Z}(0,j)$ across j, and then collect $p_{D,Z}(1,j)$ across j.

With this order of stacking/vectorization, the Jacobian matrix becomes

$$\begin{pmatrix}
\mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} \phi \left(W_{i}, W_{i'}; \tilde{\lambda}, p \right) \right] \\
\mathbf{E} \left[\frac{\partial}{\partial p} \phi \left(W_{i}, W_{i'}; \tilde{\lambda}, p \right) \right]
\end{pmatrix}$$

$$= \begin{pmatrix}
\mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} \phi_{A} \left(W_{i}, W_{i'}; \tilde{\lambda}, p \right) \right] & \mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} \phi_{B} \left(W_{i}, W_{i'}; \tilde{\lambda}, p \right) \right] & \mathbf{O}_{2K^{2} \times 2K} \\
\mathbf{O}_{K \times 2MK} & \mathbf{E} \left[\frac{\partial}{\partial p_{D}} \phi_{B} \left(W_{i}, W_{i'}; \tilde{\lambda}, p \right) \right] & \mathbf{O}_{K \times 2K} \\
\mathbf{E} \left[\frac{\partial}{\partial p_{D,Z}} \phi_{A} \left(W_{i}, W_{i'}; \tilde{\lambda}, p \right) \right] & \mathbf{E} \left[\frac{\partial}{\partial p_{D,Z}} \phi_{B} \left(W_{i}, W_{i'}; \tilde{\lambda}, p \right) \right] & -\mathbf{I}_{2K \times 2K}
\end{pmatrix}.$$

It suffices to show that the submatrix

$$\begin{pmatrix}
\mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} \phi_A \left(W_i, W_{i'}; \tilde{\lambda}, p \right) \right] & \mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} \phi_B \left(W_i, W_{i'}; \tilde{\lambda}, p \right) \right] \\
\mathbf{O}_{K \times 2MK} & \mathbf{E} \left[\frac{\partial}{\partial p_U} \phi_B \left(W_i, W_{i'}; \tilde{\lambda}, p \right) \right]
\end{pmatrix}.$$
(18)

is full rank. Assume to the contrary that the rows of the submatrix from (18) are linearly dependent:

with linear coefficients $\alpha = (\alpha_{A,1}, \dots, \alpha_{A,2K^2}, \alpha_{B,1}, \dots, \alpha_{B,2K})^{\mathsf{T}}$,

$$\alpha^{\mathsf{T}} \begin{pmatrix} \mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} \phi_A \left(W_i, W_{i'}; \tilde{\lambda}, p \right) \right] & \mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} \phi_B \left(W_i, W_{i'}; \tilde{\lambda}, p \right) \right] \\ \mathbf{O}_{K \times 2MK} & \mathbf{E} \left[\frac{\partial}{\partial p_U} \phi_B \left(W_i, W_{i'}; \tilde{\lambda}, p \right) \right] \end{pmatrix} = \mathbf{0}.$$

Note that $\mathbf{E}\left[\frac{\partial}{\partial \tilde{\lambda}}\phi_A\left(W_i,W_{i'};\tilde{\lambda},p\right)\right]$ is a diagonal block matrix, consisting of 2K block matrices, each of which is a $K\times M$ matrix. For example, the first block matrix is

$$\Lambda_0^{\mathsf{T}} \Gamma_0^{\mathsf{T}} - (\Lambda_0^{\mathsf{T}} \Gamma_X^{\mathsf{T}}) \otimes \left(\Pr\left\{ Y_i(0) = y^1 | U_i = u^1 \right\} \right) \cdots \quad \Pr\left\{ Y_i(0) = y^{M_Y} | U_i = u^1 \right\} \right)$$
$$- \left(\Pr\left\{ X_i = x^1 | U_i = u^1 \right\} \right) \cdots \quad \Pr\left\{ X_i = x^{M_X} | U_i = u^1 \right\} \right) \otimes \Lambda_0^{\mathsf{T}} \Gamma_{Y(0)}^{\mathsf{T}}$$

where \otimes is the Kronecker product. From Assumption 3.b-c, the rows of the block matrices are linearly independent. Thus, the first $2K^2$ components of α are zeroes. Then, it must satisfy that

$$\alpha_B{}^\intercal \mathbf{E} \left[\frac{\partial}{\partial p_U} \phi_B \left(W_i, W_{i'}; \tilde{\lambda}, p \right) \right] = \alpha_B{}^\intercal \Gamma_X{}^\intercal = \mathbf{0}.$$

From Assumption 3.b, α_B must be a zero vector. The Jacobian matrix has full rank.

C.3 Proof for Theorem 2

All of the following proof is for $K \geq 2$.

$$\textbf{Step 1.} \ \left\| \Gamma_0 \Lambda_0 - \widehat{\Gamma}_0 \widehat{\Lambda}_0 \right\|_F^{\ 2} = O_p \left(\tfrac{1}{\sqrt{n}} \right) \ \text{and} \ \left\| \Gamma_1 \Lambda_1 - \widehat{\Gamma}_1 \widehat{\Lambda}_1 \right\|_F^{\ 2} = O_p \left(\tfrac{1}{\sqrt{n}} \right).$$

From iid-ness of observations, we have

$$\|\mathbb{H}_0 - \mathbf{H}_0\|_F = O_p\left(\frac{1}{\sqrt{n}}\right)$$
 and $\|\mathbb{H}_1 - \mathbf{H}_1\|_F = O_p\left(\frac{1}{\sqrt{n}}\right)$.

From the definition of $\widehat{\Lambda}_0$ and $\widehat{\Lambda}_1$, we have

$$\begin{split} \left\| \mathbb{H}_{0} - \widehat{\Gamma}_{0} \widehat{\Lambda}_{0} \right\|_{F}^{2} + \left\| \mathbb{H}_{1} - \widehat{\Gamma}_{1} \widehat{\Lambda}_{1} \right\|_{F}^{2} &\leq \left\| \mathbb{H}_{0} - \Gamma_{0} \Lambda_{0} \right\|_{F}^{2} + \left\| \mathbb{H}_{1} - \Gamma_{1} \Lambda_{1} \right\|_{F}^{2} \\ &= \left\| \mathbb{H}_{0} - \mathbf{H}_{0} \right\|_{F}^{2} + \left\| \mathbb{H}_{1} - \mathbf{H}_{1} \right\|_{F}^{2} = O_{p} \left(\frac{1}{n} \right). \end{split}$$

Then,

$$\left\| \Gamma_0 \Lambda_0 - \widehat{\Gamma}_0 \widehat{\Lambda}_0 \right\|_F^2 = \left\| \mathbf{H}_0 - \widehat{\Gamma}_0 \widehat{\Lambda}_0 \right\|_F^2 \le \left(\left\| \mathbf{H}_0 - \mathbb{H}_0 \right\|_F + \left\| \mathbb{H}_0 - \widehat{\Gamma}_0 \widehat{\Lambda}_1 \right\|_F \right)^2 = O_p \left(\frac{1}{n} \right)$$

and likewise for $\|\Gamma_1\Lambda_1 - \widehat{\Gamma}_1\widehat{\Lambda}_1\|_F = \|\mathbf{H}_1 - \widehat{\Gamma}_1\widehat{\Lambda}_1\|_F$. From the submultiplicavity of $\|\cdot\|_F$, we also get $\|P\Gamma_1\Lambda_1 - P\widehat{\Gamma}_1\widehat{\Lambda}_1\|_F = \|P\Gamma_0\Lambda_1 - P\widehat{\Gamma}_0\widehat{\Lambda}_1\|_F = O_p\left(\frac{1}{\sqrt{n}}\right)$.

To avoid repetition, we will only prove the consistency of $\widehat{\Lambda}_0$; the same argument applies to $\widehat{\Lambda}_1$.

Step 2. $\|\widehat{\Gamma}_0 - \Gamma_0 A\|_F = O_p\left(\frac{1}{\sqrt{n}}\right)$ with some $K \times K$ matrix A.

Firstly, I show that $\widehat{\Lambda}_0^{-1}$ exists with probability going to one. Find that

$$\left\| \Gamma_0 \mathsf{^T} \widehat{\Gamma}_0 \widehat{\Lambda}_0 - \Gamma_0 \mathsf{^T} \Gamma_0 \Lambda_0 \right\|_F \le \| \Gamma_0 \|_F \cdot \left\| \Gamma_0 \Lambda_0 - \widehat{\Gamma}_0 \widehat{\Lambda}_0 \right\|_F = O_p \left(\frac{1}{\sqrt{n}} \right).$$

The determinant of $\Gamma_0^{\dagger}\widehat{\Gamma}_0\widehat{\Lambda}_0$ converges in probability to the determinant of $\Gamma_0^{\dagger}\Gamma_0\Lambda_0$, which is nonzero. Thus, with probability converging to one, both $\Gamma_0^{\dagger}\widehat{\Gamma}_0$ and $\widehat{\Lambda}_0$ have full rank and $\left(\Gamma_0^{\dagger}\widehat{\Gamma}_0\right)^{-1}$ and $\widehat{\Lambda}_0^{-1}$ exist.

Let

$$A = \begin{cases} \Lambda_0 \left(\widehat{\Lambda}_0 \right)^{-1}, & \text{if } \Gamma_0^{\mathsf{T}} \widehat{\Gamma}_0 \widehat{\Lambda}_0 \text{ is invertible} \\ \mathbf{I}_K, & \text{if } \Gamma_0^{\mathsf{T}} \widehat{\Gamma}_0 \widehat{\Lambda}_0 \text{ is not invertible} \end{cases}$$

with \mathbf{I}_K being the $K \times K$ identity matrix. When $\Gamma_0^{\mathsf{T}} \widehat{\Gamma}_0 \widehat{\Lambda}_0$ is invertible,

$$\begin{split} \left\| \widehat{\Gamma}_0 - \Gamma_0 A \right\|_F &= \left\| \left(\widehat{\Gamma}_0 \widehat{\Lambda}_0 - \Gamma_0 \Lambda_0 \right) \widehat{\Lambda}_0^{-1} \right\|_F \\ &\leq \left\| \widehat{\Gamma}_0 \widehat{\Lambda}_0 - \Gamma_0 \Lambda_0 \right\|_F \left\| \left(\Gamma_0^{\mathsf{T}} \widehat{\Gamma}_0 \widehat{\Lambda}_0 \right)^{-1} \right\|_F \left\| \Gamma_0^{\mathsf{T}} \widehat{\Gamma}_0 \right\|_F. \end{split}$$

There is some $\delta > 0$ such that $\left\| \Gamma_0 {}^\intercal \widehat{\Gamma}_0 \widehat{\Lambda}_0 - \Gamma_0 {}^\intercal \Gamma_0 \Lambda_0 \right\|_F \le \delta$ implies the invertibility of $\Gamma_0 {}^\intercal \widehat{\Gamma}_0 \widehat{\Lambda}_0$ and

$$C = \left\{ \left\| \left(\Gamma_0^{\mathsf{T}} \widehat{\Gamma}_0 \widehat{\Lambda}_0 \right)^{-1} \right\|_F : \left\| \Gamma_0^{\mathsf{T}} \widehat{\Gamma}_0 \widehat{\Lambda}_0 - \Gamma_0^{\mathsf{T}} \Gamma_0 \Lambda_0 \right\|_F \le \delta \right\} < \infty$$

since $\|\left(\Gamma_0^{\dagger}\widehat{\Gamma}_0\widehat{\Lambda}_0\right)^{-1}\|_F$ is a continuous function of $\Gamma_0^{\dagger}\widehat{\Gamma}_0\widehat{\Lambda}_0$ and

$$\left\{\Gamma_0{}^{\intercal}\widehat{\Gamma}_0\widehat{\Lambda}_0: \left\|\Gamma_0{}^{\intercal}\widehat{\Gamma}_0\widehat{\Lambda}_0 - \Gamma_0{}^{\intercal}\Gamma_0\Lambda_0\right\|_F \leq \delta\right\}$$

is closed and bounded. Then,

$$\Pr\left\{ \left(\left\| \Gamma_0^{\mathsf{T}} \widehat{\Gamma}_0 \widehat{\Lambda}_0 \right)^{-1} \right\|_F \ge C, \Gamma_0^{\mathsf{T}} \widehat{\Gamma}_0 \widehat{\Lambda}_0 \text{ is invertible} \right\} = o(1)$$

Also, $\left\|\Gamma_0^{\mathsf{T}}\widehat{\Gamma}_0\right\|_F$ is bounded by K^2 . Thus,

$$\begin{split} & \Pr\left\{\sqrt{n} \left\| \widehat{\Gamma}_0 - \Gamma_0 A \right\|_F \ge \varepsilon \right\} \\ & \leq \Pr\left\{\sqrt{n} \left\| \widehat{\Gamma}_0 \widehat{\Lambda}_0 - \Gamma_0 \Lambda_0 \right\|_F \left\| \left(\Gamma_0^{\mathsf{T}} \widehat{\Gamma}_0 \widehat{\Lambda}_0\right)^{-1} \right\|_F \left\| \Gamma_0^{\mathsf{T}} \widehat{\Gamma}_0 \right\|_F \ge \varepsilon, \Gamma_0^{\mathsf{T}} \widehat{\Gamma}_0 \widehat{\Lambda}_0 \text{ is invertible} \right\} + o(1) \\ & \leq \Pr\left\{\sqrt{n} \left\| \widehat{\Gamma}_0 \widehat{\Lambda}_0 - \Gamma_0 \Lambda_0 \right\|_F \ge \frac{\varepsilon}{CK^2} \right\} + o(1) \end{split}$$

Therefore, we have

$$\left\| \widehat{\Gamma}_0 - \Gamma_0 A \right\|_F = O_p \left(\frac{1}{\sqrt{n}} \right).$$

A is a $K \times K$ matrix that reorders the columns of Γ_0 so that it resembles $\widehat{\Gamma}_0$. Let a_{jk} denote the j-th row and k-th column element of A and $a_{\cdot k}$ denote the k-th column of A. In this sense, $a_{\cdot k}$ is a set of weights on the columns of $\widehat{\Gamma}_0$ so that we get the k-th column in Γ_0 .

Step 3. Each column of A converges to an elementary vector at the rate of $n^{-\frac{1}{2}}$. Firstly, the columns of A sum to one. To see this, compute column-wise sums of

$$\widehat{\Gamma}_0 = \Gamma_0 A + \left(\widehat{\Gamma}_0 \widehat{\Lambda}_0 - \Gamma_0 \Lambda_0\right) \widehat{\Lambda}_0^{-1}$$

when $\Gamma_0{}^{\intercal}\widehat{\Gamma}_0\widehat{\Lambda}_0$ is invertible:

$$\begin{split} \iota_{M}{}^{\mathsf{T}}\widehat{\Gamma}_{0} &= \iota_{M}{}^{\mathsf{T}}\Gamma_{0}A + \iota_{M}{}^{\mathsf{T}}\left(\widehat{\Gamma}_{0}\widehat{\Lambda}_{0} - \Gamma_{0}\Lambda_{0}\right)\widehat{\Lambda}_{0}^{-1} \\ \iota_{K}{}^{\mathsf{T}} &= \iota_{K}{}^{\mathsf{T}}A + \left(\iota_{K}{}^{\mathsf{T}}\widehat{\Lambda}_{0} - \iota_{K}{}^{\mathsf{T}}\Lambda_{0}\right)\widehat{\Lambda}_{0}^{-1} \\ \iota_{K}{}^{\mathsf{T}} &= \iota_{K}{}^{\mathsf{T}}A + \left(\iota_{K}{}^{\mathsf{T}} - \iota_{K}{}^{\mathsf{T}}\right)\widehat{\Lambda}_{0}^{-1} \\ \iota_{K}{}^{\mathsf{T}} &= \iota_{K}{}^{\mathsf{T}}A. \end{split}$$

Secondly, with probability going to one, the columns of A are bounded with $\|\cdot\|_{\infty}$. To see this, let $\Gamma_{0,k}$ be the k-the column of Γ_0 and let $\Gamma_{0,-k}$ be the rest of the K-1 columns formed into a $M \times (K-1)$ matrix. Let

$$\delta^* := \min_{k} \|\Gamma_{0,k} - \Gamma_{0,-k} (\Gamma_{0,-k}^{\mathsf{T}} \Gamma_{0,-k})^{-1} \Gamma_{0,-k}^{\mathsf{T}} \Gamma_{0,k} \|.$$

 $\delta^* > 0$ from Assumption 3.b. Then, for any linear combination of $\Gamma_{0,-k}$,

$$\|\Gamma_{0,k} - \Gamma_{0,-k}\alpha\|_{\infty} \ge \frac{\delta^*}{2\sqrt{M}}.$$

Since each column of A sum to one, a k-th column element of $\Gamma_0 A$ can be written as follows:

$$\begin{split} &\sum_{j=1}^{K} \Pr\{Y_i(0) = y, X_i = x | U_i = u^j\} a_{jk} \\ &= \Pr\{Y_i(0) = y, X_i = x | U_i = u^1\} \\ &+ (1 - a_{1k}) \left(\sum_{j=2}^{K} \Pr\{Y_i(0) = y, X_i = x | U_i = u^j\} \cdot \frac{a_{jk}}{\sum_{j=2}^{K} a_{jk}} - \Pr\{Y_i(0) = y, X_i = x | U_i = u^1\} \right) \end{split}$$

For any given $\{a_{jk}\}_{j=2}^K$, we know from the construction of δ^* that there must be a row in $\Gamma_0 A$ such that

$$\left| \Pr\{Y_i(0) = y, X_i = x | U_i = u^1\} - \sum_{j=2}^K \Pr\{Y_i(0) = y, X_i = x | U_i = u^j\} \cdot \frac{a_{jk}}{\sum_{j=2}^K a_{jk}} \right| \ge \frac{\delta^*}{2\sqrt{M}}.$$

Thus, $\sum_{j=1}^{K} \Pr\{Y_i(0) = y, X_i = x | U_i = u^j\} a_{jk}$ lies outside of

$$\Pr\{Y_i(0) = y, X_i = x | U_i = u^1\} + \left[-\frac{|1 - a_{1k}|\delta^*}{2\sqrt{M}}, \frac{|1 - a_{1k}|\delta^*}{2\sqrt{M}} \right]$$

and

$$\Pr\left\{|1 - a_{1k}| \ge \frac{4\sqrt{M}}{\delta^*}\right\} \le \Pr\left\{\left\|\widehat{\Gamma}_0 - \Gamma_0 A\right\|_F \ge 1\right\} = o(1).$$

The inequality holds since $\widehat{\Gamma}_0$ is a well-defined probability matrix and therefore its elements all lie between 0 and 1. We can repeat this for every a_{jk} and we have $\Pr\left\{\|a_{\cdot k}\|_{\infty} \geq \frac{4\sqrt{M}}{\delta^*} + 1\right\} = o(1)$ for every k.

Using these two observations, now I show that each column of A converges to an elementary vector at the rate of $\frac{1}{\sqrt{n}}$: with e_k being the k-th elementary vector whose k-th element is one and the rest are zeros and some $\varepsilon > 0$,

$$\Pr\left\{\sqrt{n} \cdot \min_{k} \|a_{\cdot 1} - e_{k}\| \ge \varepsilon\right\} = o(1).$$

To put a bound on the probability, I first show that $\sqrt{n} \cdot \min_k \|a_{\cdot 1} - e_k\| \ge \varepsilon$ implies that there is at least one j such that $|a_{j1}| \ge \frac{1}{K}$ and another $j' \ne j$ such that $|a_{j'1}| \ge \frac{\varepsilon}{2\sqrt{n}K}$. The existence of such j is trivial from $\sum_{k=1}^K a_{k1} = 1$. Assume to the contrary that there exists only one j such that $|a_{j1}| \ge \frac{\varepsilon}{2\sqrt{n}K}$. Then, for the rest of K-1 elements, it must be that $|a_{k1}| \le \frac{\varepsilon}{2\sqrt{n}K}$, which leads to $a_{j1} \in [1 - \frac{\varepsilon}{2\sqrt{n}}, 1 + \frac{\varepsilon}{2\sqrt{n}}]$. Then,

$$||a_{\cdot 1} - e_j|| \le \left(\frac{\varepsilon^2}{4n} \cdot \frac{K - 1}{K^2} + \frac{\varepsilon^2}{4n}\right)^{\frac{1}{2}} \le \frac{\varepsilon}{\sqrt{2n}} < \min_k ||a_{\cdot 1} - e_k||,$$

which leads to a contradiction. Thus, we have

$$\Pr\left\{\sqrt{n} \cdot \min_{k} \|a_{\cdot 1} - e_{k}\| \ge \varepsilon\right\} \le \Pr\left\{\exists j, j' \text{ such that } j \ne j', |a_{j1}| \ge \frac{1}{K}, |a_{j'1}| \ge \frac{\varepsilon}{2\sqrt{n}K}\right\}.$$

Two elements of $a_{\cdot 1}$ being away from zero creates a contradiction to $\|\widehat{\Gamma}_0 - \Gamma_0 A\|_F = O_p\left(\frac{1}{\sqrt{n}}\right)$ since the convergence says that each column of $\Gamma_0 A$ can be well-approximated by a column in $\widehat{\Gamma}_0$, which satisfies the quadratic constraints (10). To see this, let $\widetilde{\Gamma}_{0,k}$ be a $M_X \times M_Y$ matrix whose m-th row and m'-th column element is

$$\Pr\left\{Y_i(0) = y^{m'}, X_i = x^m | U_i = u^k\right\}.$$

 $\tilde{\Gamma}_{0,k}$ takes the k-th column of Γ_0 and makes it into a $M_X \times M_Y$ matrix. Note that $\tilde{\Gamma}_{0,k} = p_k q_{0k}^{\mathsf{T}}$, with

$$p_{k} = \left(\Pr \left\{ X_{i} = x^{1} | U_{i} = u^{k} \right\} \right) \cdots \Pr \left\{ X_{i} = x^{M_{X}} | U_{i} = u^{k} \right\} \right)^{\mathsf{T}},$$

$$q_{dk} = \left(\Pr \left\{ Y_{i}(d) = y^{1} | U_{i} = u^{k} \right\} \right) \cdots \Pr \left\{ Y_{i}(d) = y^{M_{Y}} | U_{i} = u^{k} \right\} \right)^{\mathsf{T}} \quad \forall k = 1, \dots, K.$$

Then, $\min_{p,q} \left\| \sum_{k=1}^K \tilde{\Gamma}_{0,k} a_{k1} - pq^{\mathsf{T}} \right\|_F = O_p\left(\frac{1}{\sqrt{n}}\right)$ since

$$\min_{p \in \mathbb{R}^{M_X}, q \in \mathbb{R}^{M_Y}} \left\| \sum_{k=1}^K \widetilde{\Gamma}_{0,k} a_{k1} - pq^\mathsf{T} \right\|_F \leq \left\| \sum_{k=1}^K \widetilde{\Gamma}_{0,k} a_{k1} - \widehat{\widetilde{\Gamma}}_{0,1} \right\|_F \leq \left\| \widehat{\Gamma}_0 - \Gamma_0 A \right\|_F$$

with $\widehat{\Gamma}_{0,k}$ constructed from $\widehat{\Gamma}_0$ in the same manner as $\widetilde{\Gamma}_{0,k}$. The first inequality holds from the construction of the estimator $\widehat{\Gamma}_0$; the estimated mixture component distribution satisfies the exclusion restriction of $Y_i(0)$ and X_i given U_i and thus $\widehat{\widetilde{\Gamma}}_{0,1}$ is a rank one matrix. The second inequality

holds since $\sum_{k=1}^K \tilde{\Gamma}_{0,k} a_{k1}$ corresponds to the first column of $\Gamma_0 A$ and $\hat{\Gamma}_{0,1}$ corresponds to the first column of $\hat{\Gamma}_0$. However, since two elements of $a_{\cdot 1}$ are away from zero, the matrix $\sum_{k=1}^K \tilde{\Gamma}_{0,k} a_{k1}$ cannot be well-approximated by a rank one matrix as implied by $\|\hat{\Gamma}_0 - \Gamma_0 A\|_F = O_p\left(\frac{1}{\sqrt{n}}\right)$, giving us a contradiction.

The rest of the step completes the argument. Assume that there exist some j, j' such that $j \neq j', |a_{j1}| \geq \frac{1}{K}, |a_{j'1}| \geq \frac{\varepsilon}{2\sqrt{n}K}$. Let $p_k(x) = \Pr\{X_i = x | U_i = u^k\}, q_{dk}(y) = \Pr\{Y_i(d) = y | U_i = u^k\}$ for $k = 1, \dots, K$ and let

$$w(y) = \begin{pmatrix} a_{11}q_{01}(y) & \cdots & a_{K1}q_{0K}(y) \end{pmatrix}^{\mathsf{T}}.$$

Then,

$$\sum_{k=1}^{K} \tilde{\Gamma}_{0,k} a_{k1} = \sum_{k=1}^{K} a_{k1} p_k q_{0k}^{\mathsf{T}} = \Gamma_X \left(w(y^1) \quad \cdots \quad w(y^{M_Y}) \right).$$

From Assumption 3.c,

$$c^* := \min_{k \neq k'} \left\{ \max_{y} \left(q_{0k}(y) - q_{0k'}(y) \right) \right\} > 0.$$

WLOG let y^1 and y^2 satisfy that

$$q_{0i}(y^1) - q_{0i'}(y^1) \ge c^*$$
 and $q_{0i'}(y^2) - q_{0i}(y^2) \ge c^*$.

Then, since $(q_{0j}(y^1)q_{0j'}(y^2) - q_{0j'}(y^1)q_{0j}(y^2)) \ge c^{*2}$,

$$\left| w_j(y^1) w_{j'}(y^2) - w_{j'}(y^1) w_j(y^2) \right| = \left| a_{j1} a_{j'1} \right| \left(q_{0j}(y^1) q_{0j'}(y^2) - q_{0j'}(y^1) q_{0j}(y^2) \right) \ge \frac{\varepsilon c^{*2}}{2\sqrt{n}K^2}.$$

With the columns corresponding to (y^1, y^2) , the submatrix of $\sum_{k=1}^K \tilde{\Gamma}_{0,k} a_{k1}$ is

$$\tilde{A} = \Gamma_X \left(w(y^1) \quad w(y^2) \right).$$

Then,

$$\min_{p,q} \Big\| \sum_{k=1}^K \tilde{\Gamma}_{0,k} a_{k1} - pq^\intercal \Big\|_F \ge \min_{p \in \mathbb{R}^{M_X}, q \in \mathbb{R}^2} \Big\| \tilde{A} - pq^\intercal \Big\|_F = \text{ the smallest singular value of } \tilde{A}.$$

The equality is from the Echkart-Young theorem. The smallest singular value of Γ_X is bounded away from zero from Assumption 3.b. To show that the smallest singular value of $\begin{pmatrix} w(y^1) & w(y^2) \end{pmatrix}$ is bounded away from zero with a lower bound proportional to $\frac{1}{\sqrt{n}}$, I use the following result:

Theorem 1 Hong and Pan (1992) Let $A \in \mathbb{R}^{\rho \times \rho}$. Then, singular values of A are bounded from below by

$$\left(\frac{\rho-1}{\rho}\right)^{\frac{\rho-1}{2}} |det(A)| \max \left\{ \frac{\min_{r} ||A_{r\cdot}||_{2}}{\prod_{r=1}^{\rho} ||A_{r\cdot}||_{2}}, \frac{\min_{s} ||A_{\cdot s}||_{2}}{\prod_{s=1}^{\rho} ||A_{\cdot s}||_{2}} \right\}$$

where A_r is the r-th row of A and A_{s} is the s-th column of A.

Find that

the smallest eigenvalue of
$$\left(w(y^1) \quad w(y^2)\right) = \min_{p \in \mathbb{R}^{M_X}, q \in \mathbb{R}^2} \left\| \left(w(y^1) \quad w(y^2)\right) - pq^\intercal \right\|_F$$

$$\geq \min_{p,q \in \mathbb{R}^2} \left\| \left(w_j(y^1) \quad w_j(y^2) \atop w_{j'}(y^1) \quad w_{j'}(y^2)\right) - pq^\intercal \right\|_F$$

$$= \text{ the smallest eigenvalue of } \left(w_j(y^1) \quad w_j(y^2) \atop w_{j'}(y^1) \quad w_{j'}(y^2)\right).$$

We have shown that

$$\det \begin{pmatrix} w_{j}(y^{1}) & w_{j}(y^{2}) \\ w_{j'}(y^{1}) & w_{j'}(y^{2}) \end{pmatrix} \ge \frac{\varepsilon c^{*2}}{2\sqrt{n}K^{2}}.$$

With probability going to one, $w(y^1)$ and $w(y^2)$ is bounded by $\frac{4\sqrt{M}}{\delta^*} + 1$ and therefore

$$(w_j(y^1)^2 + w_j(y^1)^2)^{-\frac{1}{2}} \le \left(\frac{4\sqrt{2M}}{\delta^*} + \sqrt{2}\right)^{-1} > 0.$$

Thus, with probability going to one,

the smallest eigenvalue of
$$\left(w(y^1) \quad w(y^2)\right) \ge \frac{1}{\sqrt{n}} \cdot \frac{\varepsilon c^{*2}}{2K^2} \cdot \left(\frac{4\sqrt{2M}}{\delta^*} + \sqrt{2}\right)^{-1}$$

Consequently, with some constant $C^* > 0$ which does not depend on ε ,

$$\Pr\left\{\sqrt{n} \cdot \min_{k} \|a_{\cdot 1} - e_{k}\| \ge \varepsilon\right\}$$

$$\leq \Pr\left\{\exists j, j' \text{ such that } j \neq j', |a_{j1}| \ge \frac{1}{K}, |a_{j'1}| \ge \frac{\varepsilon}{2\sqrt{n}K}\right\}$$

$$\leq \Pr\left\{\left\|\widehat{\Gamma}_{0} - \Gamma_{0}A\right\|_{F} \ge \frac{C^{*}\varepsilon}{\sqrt{n}}\right\} + \Pr\left\{\exists y \text{ s.t. } \|w(y)\|_{\infty} \ge \frac{4\sqrt{M}}{\delta^{*}} + 1\right\} = o(1).$$

We repeat this for every column of $A: a_{\cdot 2}, \dots, a_{\cdot K}$.

Step 4. No two columns of A converge to the same elementary vector.

It remains to show that A is indeed a permutation; each of the elementary vector e_1, \dots, e_K has to show up once and only once, across the columns of A. To see this, let

$$\delta^{**} = \min_{1 \le k \le K} \max_{1 \le j \le K} \Pr\{U_i = u^k | D_i = 0, Z_i = z^j\} > 0.$$

 δ^{**} finds row-wise maximums of Λ_0 and then finds the minimum among the maximum values. $\delta^{**} > 0$ since there cannot be a zero row in Λ_0 , due to Assumption 3.b. From the result of Step 3, we have

$$\sum_{k=1}^{K} \Pr\left\{ \min_{k'} \|a_{\cdot k} - e_{k'}\| \ge \frac{\delta^{**}}{K} \right\} = o(1).$$

If $\min_{k'} \|a_{\cdot k} - e_{k'}\| \le \frac{\delta^{**}}{K}$ for every k, there is a bijection between the columns of A and $\{e_1, \dots, e_K\}$. Firstly, see that $\|a_{\cdot 1} - e_k\| \le \frac{\delta^{**}}{K}$ means that

$$||a_{\cdot 1} - e_{k'}|| \ge 1 - \frac{\delta^{**}}{K} > \frac{\delta^{**}}{K} \quad \forall k' \ne k$$

since $\delta^* * < 1$ and $K \geq 2$. Thus, $\pi(k) = \arg\min_{k'} \|a_{\cdot k} - e_{k'}\|$ is a well-defined function when $\min_{k'} \|a_{\cdot k} - e_{k'}\| \leq \frac{\delta^{**}}{K}$ for every k. Secondly, assume to the contrary that there is some j such that $j \neq \pi(k)$ for every k. Then, the j-th row of A lies in $\left[-\frac{\delta^{**}}{K}, \frac{\delta^{**}}{K}\right]$. Since the columns of $\tilde{\Lambda}_0$ sum to one, the j-th row of $\Lambda_0 = A\hat{\Lambda}_0$ lies in $\left[-\frac{\delta^{**}}{K}, \frac{\delta^{**}}{K}\right]$, leading to a contradiction. Thus, π is a bijection.

Thus, with some permutation on the rows of $\widehat{\Lambda}_0$,

$$\Pr\left\{\sqrt{n} \|A - \mathbf{I}_K\|_F \ge \varepsilon\right\}$$

$$\leq \Pr\left\{\sqrt{n} \|A - \mathbf{I}_K\|_F \ge \varepsilon, \min_{k'} \|a_{\cdot k} - e_{k'}\| \le \frac{\delta^{**}}{K} \text{ for every } k\right\} + o(1)$$

$$\leq \sum_{k=1}^K \Pr\left\{\sqrt{n} \cdot \min_{k'} \|a_{\cdot k} - e_{k'}\| \ge \frac{\varepsilon}{\sqrt{K}}\right\} + o(1) = o(1).$$

Step 5. Lastly,
$$\|\widehat{\Lambda}_0 - \Lambda_0\|_F = O_p\left(\frac{1}{\sqrt{n}}\right)$$
.

Find that

$$\begin{split} & \left\| \Lambda_{0} - \widehat{\Lambda}_{0} \right\|_{F} \\ & \leq \left\| \Lambda_{0} - (\Gamma_{0}^{\mathsf{T}} \Gamma_{0})^{-1} \Gamma_{0}^{\mathsf{T}} \widehat{\Gamma}_{0} \widehat{\Lambda}_{0} \right\|_{F} + \left\| (\Gamma_{0}^{\mathsf{T}} \Gamma_{0})^{-1} \Gamma_{0}^{\mathsf{T}} \widehat{\Gamma}_{0} \widehat{\Lambda}_{0} - \widehat{\Lambda}_{0} \right\|_{F} \\ & \leq \left\| (\Gamma_{0}^{\mathsf{T}} \Gamma_{0})^{-1} \Gamma_{0}^{\mathsf{T}} \right\|_{F} \cdot \left\| \Gamma_{0} \Lambda_{0} - \widehat{\Gamma}_{0} \widehat{\Lambda}_{0} \right\|_{F} + \left\| (\Gamma_{0}^{\mathsf{T}} \Gamma_{0})^{-1} \Gamma_{0}^{\mathsf{T}} \widehat{\Gamma}_{0} - \mathbf{I}_{K} \right\|_{F} \cdot \left\| \widehat{\Lambda}_{0} \right\|_{F} \\ & \leq \left\| (\Gamma_{0}^{\mathsf{T}} \Gamma_{0})^{-1} \Gamma_{0}^{\mathsf{T}} \right\|_{F} \cdot \left\| \Gamma_{0} \Lambda_{0} - \widehat{\Gamma}_{0} \widehat{\Lambda}_{0} \right\|_{F} \\ & + \left\| \widehat{\Lambda}_{0} \right\|_{F} \cdot \left(\left\| (\Gamma_{0}^{\mathsf{T}} \Gamma_{0})^{-1} \Gamma_{0}^{\mathsf{T}} \left(\widehat{\Gamma}_{0} - \Gamma_{0} A \right) \right\|_{F} + \left\| (\Gamma_{0}^{\mathsf{T}} \Gamma_{0})^{-1} \Gamma_{0}^{\mathsf{T}} \Gamma_{0} (A - \mathbf{I}_{K}) \right\|_{F} \right) \\ & = \left(\left\| (\Gamma_{0}^{\mathsf{T}} \Gamma_{0})^{-1} \Gamma_{0}^{\mathsf{T}} \right\|_{F} + \left\| \widehat{\Lambda}_{0} \right\|_{F} \cdot \left\| (\Gamma_{0}^{\mathsf{T}} \Gamma_{0})^{-1} \Gamma_{0}^{\mathsf{T}} \right\|_{F} + \left\| \widehat{\Lambda}_{0} \right\|_{F} \right) \cdot O_{p} \left(\frac{1}{\sqrt{n}} \right). \end{split}$$

C.4 Proof for Theorem 3

Step 1.
$$\|\hat{\tilde{\Lambda}}_d - \tilde{\Lambda}_d\|_F = O_p\left(\frac{1}{\sqrt{n}}\right)$$
.

Find that

$$\begin{aligned} \left\| \widehat{\tilde{\Lambda}}_0 - \widetilde{\Lambda}_0 \right\|_F &= \left\| \widehat{\Lambda}_0^{-1} \left(\Lambda_0 - \widehat{\Lambda}_0 \right) \Lambda_0^{-1} \right\|_F \\ &\leq \left\| \widehat{\Lambda}_0^{-1} \right\|_F \cdot \left\| \Lambda_0 - \widehat{\Lambda}_0 \right\|_F \cdot \left\| \Lambda_0^{-1} \right\|_F \end{aligned}$$

and $\|\widehat{\Lambda}_0^{-1}\|_F = O_p(1)$.

Step 2.
$$\|\hat{p} - p\| = O_p\left(\frac{1}{\sqrt{n}}\right)$$
 and $\|\hat{\mu} - \mu\| = O_p\left(\frac{1}{\sqrt{n}}\right)$ as $n \to \infty$. Firstly,

$$\hat{p}_{D,Z} = \begin{pmatrix} \frac{1}{n} \sum_{i=1}^{n} \mathbf{1} \{ D_i = 0, Z_i = z^1 \} \\ \vdots \\ \frac{1}{n} \sum_{i=1}^{n} \mathbf{1} \{ D_i = 1, Z_i = z^K \} \end{pmatrix}$$

is $O_p\left(\frac{1}{\sqrt{n}}\right)$ from the central limit theorem. Thus,

$$\hat{p}_{U} = \widehat{\Lambda}_{0} \begin{pmatrix} \frac{1}{n} \sum_{i=1}^{n} \mathbf{1} \{ D_{i} = 0, Z_{i} = z^{1} \} \\ \vdots \\ \frac{1}{n} \sum_{i=1}^{n} \mathbf{1} \{ D_{i} = 0, Z_{i} = z^{K} \} \end{pmatrix} + \widehat{\Lambda}_{1} \begin{pmatrix} \frac{1}{n} \sum_{i=1}^{n} \mathbf{1} \{ D_{i} = 1, Z_{i} = z^{1} \} \\ \vdots \\ \frac{1}{n} \sum_{i=1}^{n} \mathbf{1} \{ D_{i} = 1, Z_{i} = z^{K} \} \end{pmatrix}$$

is also $O_p\left(\frac{1}{\sqrt{n}}\right)$.

Secondly, let

$$\partial \phi = \begin{pmatrix} \mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} \phi(W_i, W_{i'}; \tilde{\lambda}, p) \right] \\ \mathbf{E} \left[\frac{\partial}{\partial p} \phi(W_i, W_{i'}; \tilde{\lambda}, p) \right] \end{pmatrix},$$

$$\partial m = \begin{pmatrix} \mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} m(W_i, W_{i'}; \tilde{\lambda}, p) \right] \\ \mathbf{E} \left[\frac{\partial}{\partial p} m(W_i, W_{i'}; \tilde{\lambda}, p) \right] \end{pmatrix}.$$

and let $\widehat{\partial \phi}$ and $\widehat{\partial m}$ be the estimators of $\partial \phi$ and ∂m by taking their sample analogues, plugging in \widehat{p} and $\widehat{\lambda}$. μ is estimated with

$$\hat{\mu} = \widehat{\partial \phi}^{\mathsf{T}} \left(\widehat{\partial \phi} \widehat{\partial \phi}^{\mathsf{T}} \right)^{-1} \widehat{\partial m}.$$

 $\widehat{\partial \phi}$ and $\widehat{\partial m}$ converge to $\partial \phi$ and ∂m at the rate of $\frac{1}{\sqrt{n}}$ in $\|\cdot\|_F$ since each element of $\widehat{\partial \phi}$ and $\widehat{\partial m}$ is a ratio of a product of \sqrt{n} -consistent estimators over a product of \sqrt{n} -consistent estimators which converge to a nonzero constant. For example,

$$\mathbf{E}\left[\frac{\partial}{\partial \tilde{\lambda}_{jk,d}}\phi_{A}(W_{i},W_{i'};\tilde{\lambda},p)\right]$$

$$=\Pr\{Y_{i}=y,X_{i}=x|D_{i}=d,Z_{i}=z^{j}\}-\Pr\{Y_{i}=y|D_{i}=d,Z=z^{j}\}\cdot\Pr\{X_{i}=x|U_{i}=u^{k}\}$$

$$-\Pr\{X_{i}=x|D_{i}=d,Z=z^{j}\}\cdot\Pr\{Y_{i}(d)=y|U_{i}=u^{k}\}$$

is estimated with

$$\binom{n}{2}^{-1} \sum_{i \neq i'} \frac{\frac{1}{2} \mathbf{1} \{ Y_i = y, D_i = d, X_i = x, Z_i = z^j \}}{\hat{p}_{D,Z}(d,j)}$$

$$- \binom{n}{2}^{-1} \sum_{i \neq i'} \frac{\frac{1}{2} \mathbf{1} \{ Y_i = y, D_i = d, Z_i = z^j \} \cdot \sum_{j'=1}^K \hat{\lambda}_{j'k,d} \mathbf{1} \{ D_{i'} = d, X_{i'} = x, Z_{i'} = z^{j'} \}}{\hat{p}_{D,Z}(d,j) \cdot \hat{p}_{U}(k)}$$

$$- \binom{n}{2}^{-1} \sum_{i \neq i'} \frac{\frac{1}{2} \mathbf{1} \{ D_i = d, X_i = x, Z_i = z^j \} \cdot \sum_{j'=1}^K \hat{\lambda}_{j'k,d} \mathbf{1} \{ Y_{i'} = y, D_{i'} = d, Z_{i'} = z^{j'} \}}{\hat{p}_{D,Z}(d,j) \cdot \hat{p}_{U}(k)}.$$

The exact expression of $\partial \phi$ and ∂m is given in the proof for Lemma 1. Then,

$$\begin{split} \widehat{\mu} - \mu &= \widehat{\partial \phi}^{\mathsf{T}} \left(\widehat{\partial \phi} \widehat{\partial \phi}^{\mathsf{T}} \right)^{-1} \widehat{\partial m} - \partial \phi^{\mathsf{T}} \left(\partial \phi \partial \phi^{\mathsf{T}} \right)^{-1} \partial m \\ &\leq \left\| \widehat{\partial \phi}^{\mathsf{T}} \left(\widehat{\partial \phi} \widehat{\partial \phi}^{\mathsf{T}} \right)^{-1} \right\|_{F} \cdot \left\| \widehat{\partial m} - \partial m \right\|_{F} + \left\| \widehat{\partial \phi}^{\mathsf{T}} - \partial \phi^{\mathsf{T}} \right\|_{F} \cdot \left\| \left(\partial \phi \partial \phi^{\mathsf{T}} \right)^{-1} \partial m \right\|_{F} \\ &+ \left\| \widehat{\partial \phi}^{\mathsf{T}} \left(\partial \phi \partial \phi^{\mathsf{T}} \right)^{-1} \right\|_{F} \cdot \left\| \partial \phi \partial \phi^{\mathsf{T}} - \widehat{\partial \phi} \widehat{\partial \phi}^{\mathsf{T}} \right\|_{F} \cdot \left\| \left(\widehat{\partial \phi} \widehat{\partial \phi}^{\mathsf{T}} \right)^{-1} \partial m \right\|_{F} \\ &= O_{p} \left(\frac{1}{\sqrt{n}} \right). \end{split}$$

Step 3. Find that

$$\begin{split} &\psi\left(W_{i},W_{i'};\hat{\theta},\hat{\tilde{\lambda}},\hat{p},\hat{\mu}\right) \\ &=\psi\left(W_{i},W_{i'};\theta,\tilde{\lambda},p,\mu\right) \\ &+\frac{\partial}{\partial\theta}\psi(W_{i},W_{i'};\bar{\theta},\bar{\lambda},\bar{p},\bar{\mu})\cdot\left(\hat{\theta}-\theta\right)+\frac{\partial}{\partial\tilde{\lambda}}\psi(W_{i},W_{i'};\bar{\theta},\bar{\lambda},\bar{p},\bar{\mu})^{\mathsf{T}}\cdot\left(\hat{\tilde{\lambda}}-\tilde{\lambda}\right) \\ &+\frac{\partial}{\partial p}\psi(W_{i},W_{i'};\bar{\theta},\bar{\lambda},\bar{p},\bar{\mu})^{\mathsf{T}}\cdot(\hat{p}-p)+\frac{\partial}{\partial\mu}\psi(W_{i},W_{i'};\bar{\theta},\bar{\lambda},\bar{p},\bar{\mu})^{\mathsf{T}}\cdot(\hat{\mu}-\mu) \\ &=\psi\left(W_{i},W_{i'};\theta,\tilde{\lambda},p,\mu\right) \\ &-\left(\hat{\theta}-\theta\right)+\frac{\partial}{\partial\tilde{\lambda}}m(W_{i},W_{i'};\bar{\theta},\bar{\lambda},\bar{p})^{\mathsf{T}}\cdot\left(\hat{\tilde{\lambda}}-\tilde{\lambda}\right)-\mu^{\mathsf{T}}\frac{\partial}{\partial\tilde{\lambda}}\phi(W_{i},W_{i'};\bar{\lambda},\bar{p})^{\mathsf{T}}\cdot\left(\hat{\tilde{\lambda}}-\tilde{\lambda}\right) \\ &+\frac{\partial}{\partial p}m(W_{i},W_{i'};\bar{\theta},\bar{\lambda},\bar{p})^{\mathsf{T}}\cdot(\hat{p}-p)-\mu^{\mathsf{T}}\frac{\partial}{\partial p}\phi(W_{i},W_{i'};\bar{\lambda},\bar{p})^{\mathsf{T}}\cdot(\hat{p}-p) \\ &+\phi(W_{i},W_{i'};\bar{\lambda},\bar{p})^{\mathsf{T}}(\hat{\mu}-\mu) \end{split}$$

with $(\bar{\theta}, \bar{\lambda}, \bar{p}, \bar{\mu})$ being the intermediate values between $(\theta, \tilde{\lambda}, p, \mu)$ and $(\hat{\theta}, \hat{\bar{\lambda}}, \hat{p}, \hat{\mu})$. Therefore,

$$\begin{split} &\sqrt{n}\left(\hat{\theta}-\theta\right) \\ &= \sqrt{n}\binom{n}{2}^{-1} \sum_{i < i'} \psi\left(W_{i}, W_{i'}; \theta, \tilde{\lambda}, p, \mu\right) \\ &+ \binom{n}{2}^{-1} \sum_{i < i'} \left(\frac{\partial}{\partial \tilde{\lambda}} m(W_{i}, W_{i'}; \bar{\theta}, \bar{\lambda}, \bar{p})^{\mathsf{T}} - \mu^{\mathsf{T}} \frac{\partial}{\partial \tilde{\lambda}} \phi(W_{i}, W_{i'}; \bar{\lambda}, \bar{p})^{\mathsf{T}}\right) \cdot \sqrt{n} \left(\hat{\tilde{\lambda}} - \tilde{\lambda}\right) \\ &+ \binom{n}{2}^{-1} \sum_{i < i'} \left(\frac{\partial}{\partial p} m(W_{i}, W_{i'}; \bar{\theta}, \bar{\lambda}, \bar{p})^{\mathsf{T}} - \mu^{\mathsf{T}} \frac{\partial}{\partial p} \phi(W_{i}, W_{i'}; \bar{\lambda}, \bar{p})^{\mathsf{T}}\right) \cdot \sqrt{n} \left(\hat{p} - p\right) \\ &+ \binom{n}{2}^{-1} \sum_{i < i'} \phi(W_{i}, W_{i'}; \bar{\lambda}, \bar{p})^{\mathsf{T}} \cdot \sqrt{n} \left(\hat{\mu} - \mu\right). \end{split}$$

The intermediate values $(\bar{\theta}, \bar{\lambda}, \bar{p}, \bar{\mu})$ depend on $(W_i, W_{i'})$. From the construction of the Neyman orthogonal score and the consistency of the nuisance parameter estimators,

$$\binom{n}{2}^{-1} \sum_{i < i'} \left(\frac{\partial}{\partial \tilde{\lambda}} m(W_i, W_{i'}; \bar{\theta}, \bar{\lambda}, \bar{p}) - \frac{\partial}{\partial \tilde{\lambda}} \phi(W_i, W_{i'}; \bar{\lambda}, \bar{p}) \mu \right)$$

$$\stackrel{p}{\to} \mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} m(W_i, W_{i'}; \bar{\theta}, \bar{\lambda}, \bar{p}) \right] - \mathbf{E} \left[\frac{\partial}{\partial \tilde{\lambda}} \phi(W_i, W_{i'}; \bar{\lambda}, \bar{p}) \right] \mu = \mathbf{0}_{2K^2}$$

and similarly for $\binom{n}{2}^{-1} \sum_{i < i'} \left(\frac{\partial}{\partial p} m(W_i, W_{i'}; \bar{\theta}, \bar{\lambda}, \bar{p}) - \frac{\partial}{\partial p} \phi(W_i, W_{i'}; \bar{\lambda}, \bar{p}) \mu \right)$. From $\sqrt{n} \left(\hat{\lambda} - \tilde{\lambda} \right) = O_p(1)$, $\sqrt{n} \left(\hat{p} - p \right) = O_p(1)$, $\sqrt{n} \left(\hat{\mu} - \mu \right) = O_p(1)$ and the asymptotic theory for U statistics, we get

$$\sqrt{n}\left(\hat{\theta} - \theta\right) = \frac{1}{\sqrt{n}} \sum_{i=1}^{n} \tilde{\psi}\left(W_i; \theta, \tilde{\lambda}, p, \mu\right) + o_p(1)$$

where

$$\tilde{\psi}\left(w;\theta,\tilde{\lambda},p,\mu\right) = \mathbf{E}\left[\psi(w,W_i;\theta,\tilde{\lambda},p,\mu)\right].$$

The asymptotic normality of $\widehat{F}_{X|D=d,U}$ follows the same argument.