

# Small sample methods for cluster-robust variance estimation and hypothesis testing in fixed-effect models

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## **Abstract**

The text of your abstract. 200 or fewer words.

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\*The authors thank Dan Knopf for helpful discussions about the linear algebra behind the cluster-robust variance estimator. Coady Wing,...

# 1 INTRODUCTION

Fixed-effect models are an important tool for applied economic analysis. Controlling for unobserved confounding factors. Leading cases: panel models for repeated measurements on a set of individuals, organizations, or other aggregate units; block-randomized experiments (or analogous observational studies). Bertrand et al. highlight the need to use cluster-robust variance estimation.

Problems with standard CRVE.

Recent solutions.

## 1.1 Econometric framework

We consider a generic fixed effects model in which

$$\mathbf{y}_j = \mathbf{R}_j\boldsymbol{\beta} + \mathbf{S}_j\boldsymbol{\gamma} + \mathbf{T}_j\boldsymbol{\delta} + \boldsymbol{\epsilon}_j, \quad (1)$$

where  $\mathbf{R}_j$  is an  $n_j \times r$  matrix of covariates,  $\mathbf{S}_j$  is an  $n_j \times s$  matrix describing fixed effects that vary across clusters, and  $\mathbf{T}_j$  is an  $n_j \times t$  matrix describing fixed effects that are identified only within clusters. For example, in a balanced state-by-year panel model where the variance is estimated by clustering on states,  $\mathbf{T}_j$  would consist of an indicator for state  $j$ ,  $\mathbf{S}_j$  would include indicators for each time period, and  $\mathbf{R}_j$  would include a policy indicator or set of indicators.

We assume that  $E(\boldsymbol{\epsilon}_j | \mathbf{R}_j, \mathbf{S}_j, \mathbf{T}_j) = \mathbf{0}$  and  $\text{Var}(\boldsymbol{\epsilon}_j | \mathbf{R}_j, \mathbf{S}_j, \mathbf{T}_j) = \boldsymbol{\Sigma}_j$ , for  $j = 1, \dots, m$ , where the form of  $\boldsymbol{\Sigma}_1, \dots, \boldsymbol{\Sigma}_m$  may be unknown but the errors are independent across clusters. For notational convenience, let  $\mathbf{U}_j = [\mathbf{R}_j \ \mathbf{S}_j]$  denote the set of predictors that vary across clusters,  $\mathbf{X}_j = [\mathbf{U}_j \ \mathbf{T}_j]$  denote the full set of predictors,  $\boldsymbol{\alpha} = (\boldsymbol{\beta}', \boldsymbol{\gamma}', \boldsymbol{\delta}')'$ , and  $x = r + s + t$ . Denote the total number of individual observations by  $N = \sum_{j=1}^m n_j$ . Let  $\mathbf{y}$ ,  $\mathbf{R}$ ,  $\mathbf{S}$ ,  $\mathbf{T}$ ,  $\mathbf{U}$ , and  $\mathbf{X}$  denote the matrices obtained by stacking their corresponding components, as in  $\mathbf{R} = (\mathbf{R}'_1 \ \mathbf{R}'_2 \ \dots \ \mathbf{R}'_m)'$ .

In this model, inferential interest is confined to  $\boldsymbol{\beta}$  and the fixed effects  $\boldsymbol{\gamma}$  and  $\boldsymbol{\delta}$  are treated as nuisance parameters. The distinction between the covariates  $\mathbf{R}_j$  versus the fixed effects  $[\mathbf{S}_j \ \mathbf{T}_j]$  thus depends on context and the analyst's inferential goals. However, the distinction between the two fixed effect matrices  $\mathbf{S}_j$  and  $\mathbf{T}_j$  is unambiguous, in that

the within-cluster fixed effects satisfy  $\mathbf{T}_j \mathbf{T}'_k = \mathbf{0}$  for  $j \neq k$ . We further assume that  $(\mathbf{U}'\mathbf{U} - \mathbf{U}'_j\mathbf{U}_j)$  is of full rank for  $j = 1, \dots, m$ .

We shall consider weighted least-squares (WLS) estimation of  $\beta$ . For each cluster  $j$ , let  $\mathbf{W}_j$  be a symmetric,  $n_j \times n_j$  weighting matrix of full rank. The WLS framework includes the unweighted case (where  $\mathbf{W}_j = \mathbf{I}_j$ , an identity matrix), as well as feasible GLS.<sup>1</sup> In the latter case, it is assumed that  $\text{Var}(\mathbf{e}_j | \mathbf{X}_j) = \Phi_j$ , where  $\Phi_j$  is a known function of a low-dimensional parameter. For example, an auto-regressive error structure might be posited to describe repeated measures on an individual over time. The weighting matrices are then taken to be  $\mathbf{W}_j = \hat{\Phi}_j^{-1}$ , where the  $\hat{\Phi}_j$  are constructed from estimates of the variance parameter. Finally, for analysis of data from complex survey designs, WLS may be used with sampling weights in order to account for unequal selection probabilities.

Several approaches computing the WLS estimator are possible. One possibility is to calculate WLS estimates of the full parameter vector  $\alpha$  directly. However, this method can be computationally intensive and numerically inaccurate if the fixed effects specification is large (i.e.,  $s + t$  large). An alternative is to first absorb the fixed effect specification. We shall describe the latter approach because it is more efficient and numerically accurate.

Denote the full block-diagonal weighting matrix as  $\mathbf{W} = \text{diag}(\mathbf{W}_1, \dots, \mathbf{W}_m)$ . Let  $\mathbf{K}$  be the  $x \times r$  matrix that selects the covariates of interest, so that  $\mathbf{X}\mathbf{K} = \mathbf{R}$  and  $\mathbf{K}'\alpha = \beta$ . For a generic matrix  $\mathbf{Z}$  of full column rank, let  $\mathbf{M}_Z = (\mathbf{Z}'\mathbf{W}\mathbf{Z})^{-1}$  and  $\mathbf{H}_Z = \mathbf{Z}\mathbf{M}_Z\mathbf{Z}'\mathbf{W}$ .

The absorption technique involves obtaining the residuals from the regression of  $\mathbf{y}$  on  $\mathbf{T}$  and from the multivariate regressions of  $\mathbf{U} = [\mathbf{R} \ \mathbf{S}]$  on  $\mathbf{T}$ . The  $\mathbf{y}$  residuals and  $\mathbf{R}$  residuals are then regressed on the  $\mathbf{S}$  residuals. Finally, these twice-regressed  $\mathbf{y}$  residuals are regressed on the twice-regressed  $\mathbf{R}$  residuals to obtain the WLS estimates of  $\beta$ . Let  $\check{\mathbf{S}} = (\mathbf{I} - \mathbf{H}_T) \mathbf{S}$ ,  $\check{\mathbf{R}} = (\mathbf{I} - \mathbf{H}_{\check{\mathbf{S}}}) (\mathbf{I} - \mathbf{H}_T) \mathbf{R}$ , and  $\check{\mathbf{y}} = (\mathbf{I} - \mathbf{H}_{\check{\mathbf{S}}}) (\mathbf{I} - \mathbf{H}_T) \mathbf{y}$ . In what follows, subscripts on  $\check{\mathbf{R}}$ ,  $\check{\mathbf{S}}$ ,  $\check{\mathbf{U}}$ , and  $\check{\mathbf{y}}$  refer to the rows of these matrices corresponding to a specific

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<sup>1</sup>The WLS estimator also encompasses the estimator proposed by Ibragimov and Müller (2010) for clustered data. Assuming that  $\mathbf{X}_j$  has rank  $p$  for  $j = 1, \dots, m$ , their proposed approach involves estimating  $\beta$  separately within each cluster and taking the simple average of these estimates. The resulting average is equivalent to the WLS estimator with weights  $\mathbf{W}_j = \mathbf{X}_j (\mathbf{X}'_j \mathbf{X}_j)^{-2} \mathbf{X}_j$ .

cluster. The WLS estimator of  $\beta$  can then be written as

$$\hat{\beta} = \mathbf{M}_{\ddot{\mathbf{R}}} \sum_{j=1}^m \ddot{\mathbf{R}}_j' \mathbf{W}_j \ddot{\mathbf{y}}_j. \quad (2)$$

This estimator is algebraically identical to the direct WLS estimator based on the full set of predictors,

$$\hat{\beta} = \mathbf{K}' \mathbf{M}_{\mathbf{X}} \sum_{j=1}^m \mathbf{X}_j' \mathbf{W}_j \mathbf{y}_j,$$

but avoids the need to solve a system of  $x$  linear equations.

The variance of the WLS estimator is

$$\text{Var}(\hat{\beta}) = \mathbf{M}_{\ddot{\mathbf{R}}} \left( \sum_{j=1}^m \ddot{\mathbf{R}}_j' \mathbf{W}_j \boldsymbol{\Sigma}_j \mathbf{W}_j \ddot{\mathbf{R}}_j \right) \mathbf{M}_{\ddot{\mathbf{R}}}, \quad (3)$$

which depends upon the unknown variance matrices  $\boldsymbol{\Sigma}_j$ . One approach to estimating this variance is based on a parametric model for the error structure. If this approach is used, each  $\boldsymbol{\Sigma}_j$  is substituted with an estimate  $\hat{\boldsymbol{\Phi}}_j$ , producing the model-based variance estimator

$$\mathbf{V}^M = \mathbf{M}_{\ddot{\mathbf{R}}} \left( \sum_{j=1}^m \ddot{\mathbf{R}}_j' \mathbf{W}_j \hat{\boldsymbol{\Phi}}_j \mathbf{W}_j \ddot{\mathbf{R}}_j \right) \mathbf{M}_{\ddot{\mathbf{R}}}. \quad (4)$$

However, if the working model is mis-specified, the model-based variance estimator will be inconsistent and inferences based upon it will be invalid.

Do we even need to write out the formula for  $\mathbf{V}^M$ ?

## 1.2 Standard CRVE

Cluster-robust variance estimators provide a means of estimating  $\text{Var}(\hat{\beta})$  and testing hypotheses regarding  $\beta$  in the absence of a valid parametric model for the error structure. They are thus a generalization of heteroskedasticity-consistent (HC) variance estimators. Like the HC estimators, several different variants have been proposed, with different rationales and different finite-sample properties. Each of these are of the form

$$\mathbf{V}^{CR} = \mathbf{M}_{\ddot{\mathbf{R}}} \left( \sum_{j=1}^m \ddot{\mathbf{R}}_j' \mathbf{W}_j \mathbf{A}_j \mathbf{e}_j \mathbf{e}_j' \mathbf{A}_j' \mathbf{W}_j \ddot{\mathbf{R}}_j \right) \mathbf{M}_{\ddot{\mathbf{R}}}, \quad (5)$$

for some  $n_j$  by  $n_j$  adjustment matrix  $\mathbf{A}_j$ . The form of these adjustments parallels those of the heteroscedasticity-consistent (HC) variance estimators proposed by MacKinnon and

Citations on originators of CRVE? Would be a long list...

White (1985). Setting  $\mathbf{A}_j = \mathbf{I}_j$ , an  $n_j \times n_j$  identity matrix, results in the most basic form, described by Liang and Zeger (1986). Following Cameron and Miller (2015), we refer to this estimator as  $\mathbf{V}^{CR0}$ . Setting  $\mathbf{A}_j = c\mathbf{I}_j$ , where  $c = \sqrt{(m/(m-1))(N/(N-p))}$ , results in a slightly larger estimator, denoted  $\mathbf{V}^{CR1}$ . Note that when  $N \gg p$ ,  $c \approx \sqrt{m/(m-1)}$ , and some software uses the latter approximation. Both the CR0 and CR1 estimators rely on asymptotic properties of the residuals in order to consistently estimate  $\boldsymbol{\Sigma}_j$ . The correction constant used in the CR1 estimator does not depend on  $\mathbf{X}_j$ , and so cannot account for features of the covariates that might cause the cross-product of the residuals to better or worse estimates of the true variance.

Several further small-sample corrections for CRVE do account for features of the covariates. The BRL approach (described in the next section) is an extension of the HC2 estimator for regressions with heteroskedastic but uncorrelated errors; we therefore refer to it as CR2. A further alternative is CR3, which uses adjustment matrices given by  $\mathbf{A}_j = \left(\mathbf{I} - \ddot{\mathbf{R}}_j \mathbf{M}_{\ddot{\mathbf{R}}} \ddot{\mathbf{R}}_j' \mathbf{W}_j\right)^{-1}$ . The CR3 estimator closely approximates the jackknife re-sampling variance estimator.

## 2 BIAS REDUCED LINEARIZATION

The BRL approach chooses adjustment matrices so that the variance estimator is exactly unbiased under a specific working model for the data. It is therefore directly analogous to the HC2 heteroskedasticity-robust estimator, which is exactly unbiased under homoskedasticity. Bell and McCaffrey (2002) developed the BRL estimator for linear regression models in which the errors have an unknown dependence structure within clusters. However, their implementation is not applicable to many fixed effect models, where the adjustment matrices may be undefined. For instance, Angrist and Pischke (2009) pointed out that Bell and McCaffrey’s approach cannot be applied in balanced state-by-year panels with fixed effects for states and for years because the adjustment matrices involve inverses of matrices that are not of full rank. The form of the Bell and McCaffrey matrices also varies depending on whether fixed effects are absorbed or estimated directly by WLS, which is undesirable. Our implementation of BRL addresses both of these issues and can be implemented in models with quite general fixed effects specifications. It reduces to Bell and McCaffrey’s

implementation for models without fixed effects.

Let  $\Phi_j$  be a working model for the covariance of the errors in cluster  $j$ , and denote  $\Phi = \text{diag}(\Phi_1, \dots, \Phi_m)$ . Consider adjustment matrices satisfying the following criterion:

$$\ddot{\mathbf{R}}_j' \mathbf{W}_j \mathbf{A}_j (\mathbf{I} - \mathbf{H}_{\mathbf{X}})_j \Phi (\mathbf{I} - \mathbf{H}_{\mathbf{X}})_j' \mathbf{A}_j' \mathbf{W}_j \ddot{\mathbf{R}}_j = \ddot{\mathbf{R}}_j' \mathbf{W}_j \Phi_j \mathbf{W}_j \ddot{\mathbf{R}}_j, \quad (6)$$

where  $(\mathbf{I} - \mathbf{H}_{\mathbf{X}})_j$  denotes the rows of  $\mathbf{I} - \mathbf{H}_{\mathbf{X}}$  corresponding to cluster  $j$ . A variance estimator that uses such adjustment matrices will be exactly unbiased when the working model is correctly specified.<sup>2</sup> When the working model deviates from the true covariance  $\Sigma_j$ , the variance estimator remains biased. However, Bell and McCaffrey (2002) showed that the CR2 estimator still greatly reduces the bias compared to the more basic CR0 and CR1 estimators (thus the name “bias reduced linearization”). Extensive simulation results indicate that the remaining bias is typically minimal, even for large deviations from the assumed structure (CITE). Furthermore, as the number of clusters increases, the reliance on the working model diminishes. In a sense, CR2 provides necessary scaffolding in the small sample case, which falls away when there is sufficient data.

Criterion (6) does not uniquely define  $\mathbf{A}_j$ . Following McCaffrey, Bell and Botts (2001), we propose to use a symmetric solution in which

$$\mathbf{A}_j = \mathbf{D}_j' \mathbf{B}_j^{+1/2} \mathbf{D}_j, \quad (7)$$

where  $\mathbf{D}_j$  is the upper-right triangular Cholesky factorization of  $\hat{\Phi}_j$ ,

$$\mathbf{B}_j = \mathbf{D}_j (\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{R}}})_j (\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{S}}}) (\mathbf{I} - \mathbf{H}_{\mathbf{T}}) \Phi (\mathbf{I} - \mathbf{H}_{\mathbf{T}})' (\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{S}}})' (\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{R}}})_j' \mathbf{D}_j', \quad (8)$$

and  $\mathbf{B}_j^{+1/2}$  is the symmetric square root of the Moore-Penrose inverse of  $\mathbf{B}_j$ . The Moore-Penrose inverse is well-defined and unique even when  $\mathbf{B}_j$  is not of full rank (Banerjee and Roy, 2014, Thm. 9.18). Theorem 1 in Appendix A shows that the adjustment matrices given by (7) and (8) satisfy criterion (6). Furthermore, because the adjustment matrices are defined in terms of all three components of the predictors ( $\mathbf{R}$ ,  $\mathbf{S}$ , and  $\mathbf{T}$ ), they are invariant

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<sup>2</sup>Note that this criterion differs from the criterion used by Bell and McCaffrey (2002) in that it pre- and post-multiplies both sides by  $\mathbf{W}_j \ddot{\mathbf{R}}_j$ . As will be seen, this modification justifies the use of generalized matrix inverses in calculating the adjustment matrices, thus avoiding rank-deficiency problems that would otherwise leave them undefined.

to whether the model is estimated by direct WLS estimation or after absorbing some or all of the fixed effects.

In many applications, it will make sense to choose weighting matrices that are the inverses of the working covariance model, so that  $\mathbf{W}_j = \mathbf{\Phi}_j^{-1}$ . In this case, the adjustment matrices can be calculated using  $\tilde{\mathbf{B}}_j$  in place of  $\mathbf{B}_j$ , where

$$\tilde{\mathbf{B}}_j = \mathbf{D}_j (\mathbf{I} - \mathbf{H}_{\tilde{\mathbf{R}}})_j (\mathbf{I} - \mathbf{H}_{\tilde{\mathbf{S}}}) \mathbf{\Phi} (\mathbf{I} - \mathbf{H}_{\tilde{\mathbf{S}}})' (\mathbf{I} - \mathbf{H}_{\tilde{\mathbf{R}}})_j' \mathbf{D}_j'. \quad (9)$$

Theorem 2 in Appendix A demonstrates that using  $\tilde{\mathbf{B}}_j$  rather than  $\mathbf{B}_j$  leads to algebraically identical adjustment matrices; the form of  $\tilde{\mathbf{B}}_j$  is simply more convenient for computation. In the simple case of ordinary (unweighted) least squares, in which the working variance model posits that the errors are all independent and homoskedastic and  $\mathbf{W} = \mathbf{\Phi} = \mathbf{I}$ , the adjustment matrices simplify further to

$$\mathbf{A}_j = \left( \mathbf{I}_j - \ddot{\mathbf{U}}_j \left( \ddot{\mathbf{U}}' \ddot{\mathbf{U}} \right)^{-1} \ddot{\mathbf{U}}_j' \right)^{+1/2},$$

where  $\ddot{\mathbf{U}} = (\mathbf{I} - \mathbf{H}_{\mathbf{T}}) \mathbf{U}$ .

The remainder of the paper considers hypothesis testing procedures based on the BRL variance estimator,  $\mathbf{V}^{CR2}$ .

### 3 HYPOTHESIS TESTING

Wald-type test statistics based on CRVEs are often used to test hypotheses regarding the coefficients in the regression specification. Such procedures are justified based on the asymptotic behavior of robust Wald statistics as the number of clusters grows large (i.e., as  $m \rightarrow \infty$ ). However, evidence from a wide variety of contexts indicates that the asymptotic limiting distribution can be a very poor approximation when the number of clusters is small, even when small-sample corrections such as CR2 are employed (Bell and McCaffrey, 2002; Bertrand, Duflo and Mullainathan, 2004; Cameron, Gelbach and Miller, 2008). Furthermore, the accuracy of asymptotic approximations depends on design features such as the degree of imbalance in the covariates, skewness of the covariates, and similarity of cluster sizes (McCaffrey et al., 2001; Tipton and Pustejovsky, forthcoming; Webb and

MacKinnon, 2013). Consequently, no simple rule-of-thumb exists for what constitutes an adequate sample size to trust the asymptotic test.

We will consider linear constraints on  $\beta$ , where the null hypothesis has the form  $H_0 : \mathbf{C}\beta = \mathbf{d}$  for fixed  $q \times r$  matrix  $\mathbf{C}$  and  $q \times 1$  vector  $\mathbf{d}$ . For a general CRVE estimator, the Wald statistic is then

$$Q = (\mathbf{C}\hat{\beta} - \mathbf{d})' (\mathbf{C}\mathbf{V}^{CR}\mathbf{C}')^{-1} (\mathbf{C}\hat{\beta} - \mathbf{d}). \quad (10)$$

The asymptotically valid Wald test rejects  $H_0$  at level  $\alpha$  if  $Q$  exceeds  $\chi^2(\alpha; q)$ , the  $\alpha$  critical value from a chi-squared distribution with  $q$  degrees of freedom. When the number of clusters is small, it is common to instead use the test statistic  $F = Q/q$ , compared to the  $F(q, m - 1)$  reference distribution.

Citations to evidence that asymptotic test is way too liberal?

Is this really standard?

Small-sample adjustments to hypothesis tests based on CRVE have largely focused on tests for single coefficients. The following subsection reviews approaches for one-dimensional hypothesis tests, with special attention to the Satterthwaite approximation approach proposed by Bell and McCaffrey (2002). We then propose a method for testing more general,  $q$ -dimensional linear hypotheses regarding  $\beta$ . Our approach is similar to a Satterthwaite approximation, in that it involves approximating the distribution of  $Q$  using an  $F$  distribution with estimated degrees of freedom.

### 3.1 Small-sample corrections for t-tests

Consider testing the hypothesis  $H_0 : \mathbf{c}'\beta = 0$  for some fixed  $r \times 1$  contrast vector. For this one-dimensional constraint, an equivalent to the Wald F test is to use the test statistic  $Z = \mathbf{c}'\hat{\beta} / \sqrt{\mathbf{c}'\mathbf{V}^{CR}\mathbf{c}}$ , which follows a standard normal in large samples. In small samples, it is common to instead approximate the distribution of  $Z$  by a  $t(m - 1)$  distribution. Hansen (2007) provided one justification for the use of a  $t(m - 1)$  reference distribution by identifying conditions under which  $Z$  converges in distribution to  $t(m - 1)$  as the within-cluster sample sizes grow large, with  $m$  fixed (see also Donald and Lang, 2007). Ibragimov and Müller (2010) proposed a weighting technique derived so that that  $t(m - 1)$  critical values would be conservative (leading to rejection rates less than or equal to  $\alpha$ ). However, both of these arguments require that  $\mathbf{c}'\beta$  be separately identified within each cluster. Outside of



these circumstances, using  $t(m-1)$  critical values can still lead to over-rejection (Cameron and Miller, 2015). Furthermore, this correction does not take into account that the distribution of  $\mathbf{V}^{CR}$  is affected by the structure of the covariate matrix. An alternative, proposed by Bell and McCaffrey (2002), is to approximate the distribution of  $Z$  by a  $t$  distribution with degrees of freedom determined by a Satterthwaite approximation, under the working covariance model.

The  $t$ -test developed by Bell and McCaffrey (2002) involves using a  $t(\nu)$  reference distribution with degrees of freedom estimated by a Satterthwaite approximation. The Satterthwaite approximation (Satterthwaite, 1946) entails using degrees of freedom that are a function of the first two moments of the sampling distribution of  $\mathbf{c}'\mathbf{V}^{CR}\mathbf{c}$ . Theoretically, these degrees of freedom should be

$$\nu = \frac{2 [\mathbf{E} (\mathbf{c}'\mathbf{V}^{CR2}\mathbf{c})]^2}{\text{Var} (\mathbf{c}'\mathbf{V}^{CR2}\mathbf{c})}. \quad (11)$$

Expressions for the first two moments of  $\mathbf{c}'\mathbf{V}^{CR2}\mathbf{c}$  can be derived under the assumption that the errors  $\boldsymbol{\epsilon}_1, \dots, \boldsymbol{\epsilon}_m$  are normally distributed; see Appendix B.

In practice, both moments involve the variance structure  $\boldsymbol{\Sigma}$ , which is unknown. Bell and McCaffrey (2002) proposed to estimate the moments based on the same working model as used to derive the adjustment matrices. This “model-based” estimate of the degrees of freedom is then calculated as

$$\nu_M = \frac{\left( \sum_{j=1}^m \mathbf{p}_j' \hat{\boldsymbol{\Phi}} \mathbf{p}_j \right)^2}{\sum_{i=1}^m \sum_{j=1}^m \left( \mathbf{p}_i' \hat{\boldsymbol{\Phi}} \mathbf{p}_j \right)^2}, \quad (12)$$

where  $\mathbf{p}_j = (\mathbf{I} - \mathbf{H}_\mathbf{X})'_j \mathbf{A}_j \mathbf{W}_j \ddot{\mathbf{R}}_j \mathbf{M}_{\ddot{\mathbf{R}}} \mathbf{c}$ . Alternately, for any of the CRVEs one could instead use an “empirical” estimate of the degrees of freedom, constructed by substituting  $\mathbf{e}_j \mathbf{e}_j'$  in place of  $\boldsymbol{\Sigma}_j$ . However, Bell and McCaffrey (2002) found using simulation that this plug-in degrees of freedom estimate led to very conservative rejection rates.

Can we use  $\mathbf{H}_{\hat{\mathbf{U}}}$  here instead?

The Bell and McCaffrey (2002) approach has been shown to perform well in a variety of conditions (CITE simulation studies). These studies encompass a variety of data generation processes and covariate types. A key finding is that the degrees of freedom depend not only on the number of clusters  $m$ , but also on features of the covariates. When the covariate is balanced across clusters—as occurs in balanced panels with a dichotomous covariate with

the same proportion of ones in each cluster—the degrees of freedom are  $m - 1$  even in small samples. However, when the covariate exhibits large imbalances—as occurs when the panel is not balanced or if the proportion of ones varies from cluster to cluster—the degrees of freedom can be considerably smaller. Similarly, covariates with large leverage points will tend to exhibit lower degrees of freedom. Because the degrees of freedom are covariate-dependent, it is not possible to assess whether a small-sample correction is needed based solely on the total number of clusters in the data. Consequently, we recommend that t-tests based on CRVE should routinely use the CR2 variance estimator and the Satterthwaite degrees of freedom, regardless even when  $m$  appears to be large.

### 3.2 Small-sample corrections for F-tests

Little extant research has considered small-sample corrections for multiple-constraint hypothesis tests based on robust Wald statistics. A simple correction, analogous to the CR1 for t-tests, would be to compare  $Q/q$  to an  $F(q, m - 1)$  reference distribution. As we will show in our simulation study, like the t-test case, this test tends to be overly liberal. The ideal adjustment, therefore, would be to determine empirically the degrees of freedom of the  $F$  distribution using an approach similar to that for the BRL t-test. In the broad literature, several small-sample corrections for multiple-constraint Wald tests of this form have been proposed. While this broader literature includes methods based on spectral decomposition (CITE), as well as several methods based on the Wishart distribution (which we focus in on here), we ultimately focus here on the development of a single test that performs well under a vareity of conditions (see Tipton Pustejovsky 2015).

We already said this.

Why ideal? Doesn't matter if it's an F, as long as it works, no?

Following the approach of Pan and Wall (2002), who developed a similar method in the context of CRVE for generalized estimating equations, the method we propose involves approximating the distribution of  $\mathbf{CV}^{CR2}\mathbf{C}'$  by a multiple of a Wishart distribution. From this it follows that  $Q$  approximately follows a multiple of an F distribution. Specifically, if  $\eta\mathbf{CV}^{CR2}\mathbf{C}'$  approximately follows a Wishart distribution with  $\eta$  degrees of freedom and scale matrix  $\mathbf{CVar}\left(\mathbf{C}\hat{\beta}\right)\mathbf{C}'$ , then

$$\left(\frac{\eta - q + 1}{\eta q}\right) Q \sim F(q, \eta - q + 1). \quad (13)$$

We will refer to this as the approximate Hotelling's  $T^2$  (AHT) test, and the remainder of this section will develop this test in greater detail.

Just as in the t-test case, our goal is to develop a strategy to estimate the degrees of freedom of this F-test (through the parameter  $\eta$ ). To do so, we estimate the degrees of freedom of the Wishart distribution so that they match the mean and variance of  $\mathbf{C}\mathbf{V}^{CR}\mathbf{C}'$ . A problem that arises in doing so is that when  $q > 1$  it is not possible to exactly match both moments. In developing the test, we therefore borrow strategies from the literature on CRVE found more broadly. One approach, developed by Pan and Wall (2002), is to use as degrees of freedom the value that minimizes the squared differences between the covariances among the entries of  $\eta\mathbf{C}\mathbf{V}^{CR}\mathbf{C}'$  and the covariances of the Wishart distribution with  $\eta$  degrees of freedom and scale matrix  $\mathbf{C}\mathbf{V}^{CR}\mathbf{C}'$ . Another approach, developed by Zhang (2012a,1,1) in the context of heteroskedastic and multivariate analysis of variance models, is to instead match the mean and total variance of  $\mathbf{C}\mathbf{V}^{CR}\mathbf{C}'$  (i.e., the sum of the variances of its entries), which avoids the need to calculate any covariances. In what follows we focus on this latter approach, which we find performs best in practice (see Tipton Pustejovsky 2015).

Let  $\mathbf{c}_1, \dots, \mathbf{c}_q$  denote the  $p \times 1$  row-vectors of  $\mathbf{C}$ . Let  $\mathbf{p}_{sh} = (\mathbf{I} - \mathbf{H})'_h \mathbf{A}'_h \mathbf{W}_h \mathbf{X}_h \mathbf{M} \mathbf{c}_s$  for  $s = 1, \dots, q$  and  $h = 1, \dots, m$ . The degrees of freedom are then estimated under the working model as

$$\eta_M = \frac{\sum_{s,t=1}^q \sum_{h,i=1}^m b_{st} \mathbf{p}'_{sh} \hat{\Omega} \mathbf{p}_{th} \mathbf{p}'_{si} \hat{\Omega} \mathbf{p}_{ti}}{\sum_{s,t=1}^q \sum_{h,i=1}^m \mathbf{p}'_{sh} \hat{\Omega} \mathbf{p}_{ti} \mathbf{p}'_{sh} \hat{\Omega} \mathbf{p}_{ti} + \mathbf{p}'_{sh} \hat{\Omega} \mathbf{p}_{si} \mathbf{p}'_{th} \hat{\Omega} \mathbf{p}_{ti}}, \quad (14)$$

where  $b_{st} = 1 + (s = t)$  for  $s, t = 1, \dots, q$ . Note that  $\eta_M$  reduces to  $\nu_M$  if  $q = 1$ .

This F-test shares features with the t-test developed by Bell and McCaffrey. Like the t-test, the degrees of freedom of this F-test depend non only on the number of clusters, but also on features of the covariates being tested. Again, these degrees of freedom can be much smaller than  $m - 1$ , and are particularly smaller when the covariates being tested exhibit high imbalances or leverage. Unlike the t-test case, however, in multi-parameter case, it is often more difficult to diagnose the cause of these small degrees of freedom. In some situations, however, these are straightforward extensions to the findings in t-tests. For example, if the goal is to test if there are differences across a four-arm treatment study, the degrees of freedom are largest (and close to  $m - 1$ ) when the treatment is allocated equally

across the four groups within each cluster. When the proportion varies across clusters, these degrees of freedom fall, often leading to degrees of freedom in the "small sample" territory even when the number of clusters is large. In the next section, we will illustrate these principles in a simulation study.

## 4 SIMULATION EVIDENCE

## 5 EXAMPLES

### 5.1 Tennessee STAR class-size experiment.

### 5.2 Heterogeneous treatment impacts

### 5.3 Robust Hausmann test

## 6 DISCUSSION

## A BRL adjustment matrices

This appendix states and provides proof of two theorems regarding the BRL adjustment matrices.

**Theorem 1.** *Let  $\mathbf{L} = (\ddot{\mathbf{U}}'\ddot{\mathbf{U}} - \ddot{\mathbf{U}}_j'\ddot{\mathbf{U}}_j)$  and assume that  $\mathbf{L}$  has full rank  $r + s$ , so that its inverse exists. Then the adjustment matrices  $\mathbf{A}_j$  defined in (7) and (8) satisfy criterion (6) and  $\mathbf{V}^{CR2}$  is exactly unbiased when the working covariance model  $\Phi$  is correctly specified.*

*Proof.* The Moore-Penrose inverse of  $\mathbf{B}_j$  can be computed from its eigen-decomposition. Let  $b \leq n_j$  denote the rank of  $\mathbf{B}_j$ . Let  $\mathbf{\Lambda}$  be the  $b \times b$  diagonal matrix of the positive eigenvalues of  $\mathbf{B}_j$  and  $\mathbf{V}$  be the  $n_j \times b$  matrix of corresponding eigen-vectors, so that  $\mathbf{B}_j = \mathbf{V}\mathbf{\Lambda}\mathbf{V}'$ . Then  $\mathbf{B}_j^+ = \mathbf{V}\mathbf{\Lambda}^{-1}\mathbf{V}'$  and  $\mathbf{B}_j^{+1/2} = \mathbf{V}\mathbf{\Lambda}^{-1/2}\mathbf{V}'$ .

Now, observe that  $(\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{R}}})_j (\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{S}}}) (\mathbf{I} - \mathbf{H}_{\mathbf{T}}) = (\mathbf{I} - \mathbf{H}_{\mathbf{X}})_j$ . Thus,

$$\begin{aligned} \ddot{\mathbf{R}}'_j \mathbf{W}_j \mathbf{A}_j (\mathbf{I} - \mathbf{H}_{\mathbf{X}})_j \boldsymbol{\Phi} (\mathbf{I} - \mathbf{H}_{\mathbf{X}})'_j \mathbf{A}'_j \mathbf{W}_j \ddot{\mathbf{R}}_j &= \ddot{\mathbf{R}}'_j \mathbf{W}_j \mathbf{D}_j \mathbf{B}_j^{+1/2} \mathbf{B}_j \mathbf{B}_j^{+1/2} \mathbf{D}'_j \mathbf{W}_j \ddot{\mathbf{R}}_j \\ &= \ddot{\mathbf{R}}'_j \mathbf{W}_j \mathbf{D}_j \mathbf{V} \mathbf{V}' \mathbf{D}'_j \mathbf{W}_j \ddot{\mathbf{R}}_j. \end{aligned} \quad (15)$$

Because  $\mathbf{D}_j$ , and  $\boldsymbol{\Phi}$  are positive definite and  $\mathbf{B}_j$  is symmetric, the eigenvectors  $\mathbf{V}$  define an orthonormal basis for the column span of  $(\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{U}}})_j$ . We now show that  $\ddot{\mathbf{U}}_j$  is in the column space of  $(\mathbf{I} - \mathbf{H}_{\mathbf{X}})_j$ . Let  $\mathbf{Z}_j$  be an  $n_j \times (r + s)$  matrix of zeros. Let  $\mathbf{Z}_k = -\ddot{\mathbf{U}}_k \mathbf{L}^{-1} \mathbf{M}_{\ddot{\mathbf{U}}}^{-1}$ , for  $k \neq j$  and take  $\mathbf{Z} = (\mathbf{Z}'_1, \dots, \mathbf{Z}'_m)'$ . Now observe that  $(\mathbf{I} - \mathbf{H}_{\mathbf{T}}) \mathbf{Z} = \mathbf{Z}$ . It follows that

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$$\begin{aligned} (\mathbf{I} - \mathbf{H}_{\mathbf{X}})_j \mathbf{Z} &= (\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{U}}})_j (\mathbf{I} - \mathbf{H}_{\mathbf{T}}) \mathbf{Z} = (\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{U}}})_j \mathbf{Z} \\ &= \mathbf{Z}_j - \ddot{\mathbf{U}}_j \mathbf{M}_{\ddot{\mathbf{U}}} \sum_{k=1}^m \ddot{\mathbf{U}}'_k \mathbf{W}_k \mathbf{Z}_k = \ddot{\mathbf{U}}_j \mathbf{M}_{\ddot{\mathbf{U}}} \left( \sum_{k \neq j} \ddot{\mathbf{U}}'_k \mathbf{W}_k \ddot{\mathbf{U}} \right) \mathbf{L}^{-1} \mathbf{M}_{\ddot{\mathbf{U}}}^{-1} \\ &= \ddot{\mathbf{U}}_j. \end{aligned}$$

Thus, there exists an  $N \times (r + s)$  matrix  $\mathbf{Z}$  such that  $(\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{U}}})_j \mathbf{Z} = \ddot{\mathbf{U}}_j$ , i.e.,  $\ddot{\mathbf{U}}_j$  is in the column span of  $(\mathbf{I} - \mathbf{H}_{\mathbf{X}})_j$ . Because  $\mathbf{D}_j \mathbf{W}_j$  is positive definite and  $\ddot{\mathbf{R}}_j$  is a sub-matrix of  $\ddot{\mathbf{U}}_j$ ,  $\mathbf{D}_j \mathbf{W}_j \ddot{\mathbf{R}}_j$  is also in the column span of  $(\mathbf{I} - \mathbf{H}_{\mathbf{X}})_j$ . It follows that

$$\ddot{\mathbf{R}}'_j \mathbf{W}_j \mathbf{D}_j \mathbf{V} \mathbf{V}' \mathbf{D}'_j \mathbf{W}_j \ddot{\mathbf{R}}_j = \ddot{\mathbf{R}}'_j \mathbf{W}_j \boldsymbol{\Phi}_j \mathbf{W}_j \ddot{\mathbf{R}}_j. \quad (16)$$

Substituting (16) into (15) demonstrates that  $\mathbf{A}_j$  satisfies criterion (6).

Under the working model, the residuals from cluster  $j$  have mean  $\mathbf{0}$  and variance

$$\text{Var}(\ddot{\mathbf{e}}_j) = (\mathbf{I} - \mathbf{H}_{\mathbf{X}})_j \boldsymbol{\Phi} (\mathbf{I} - \mathbf{H}_{\mathbf{X}})'_j,$$

It follows that

$$\begin{aligned} \text{E}(\mathbf{V}^{CR2}) &= \mathbf{M}_{\ddot{\mathbf{R}}} \left[ \sum_{j=1}^m \ddot{\mathbf{R}}'_j \mathbf{W}_j \mathbf{A}_j (\mathbf{I} - \mathbf{H}_{\mathbf{X}})_j \boldsymbol{\Phi} (\mathbf{I} - \mathbf{H}_{\mathbf{X}})'_j \mathbf{A}_j \mathbf{W}_j \ddot{\mathbf{R}}_j \right] \mathbf{M}_{\ddot{\mathbf{R}}} \\ &= \mathbf{M}_{\ddot{\mathbf{R}}} \left[ \sum_{j=1}^m \ddot{\mathbf{R}}'_j \mathbf{W}_j \boldsymbol{\Phi}_j \mathbf{W}_j \ddot{\mathbf{R}}_j \right] \mathbf{M}_{\ddot{\mathbf{R}}} \\ &= \text{Var}(\hat{\boldsymbol{\beta}}) \end{aligned}$$

□

**Theorem 2.** Let  $\tilde{\mathbf{A}}_j = \mathbf{D}'_j \tilde{\mathbf{B}}_j^{+1/2} \mathbf{D}_j$ , where  $\tilde{\mathbf{B}}_j$  is given in (9). If  $\mathbf{T}_j \mathbf{T}'_k = \mathbf{0}$  for  $j \neq k$  and  $\mathbf{W} = \Phi^{-1}$ , then  $\mathbf{A}_j = \tilde{\mathbf{A}}_j$ .

*Proof.* From the fact that  $\ddot{\mathbf{U}}'_j \mathbf{W}_j \mathbf{T}_j = \mathbf{0}$  for  $j = 1, \dots, m$ , it follows that

$$\begin{aligned} \mathbf{B}_j &= \mathbf{D}_j (\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{U}}})_j (\mathbf{I} - \mathbf{H}_{\mathbf{T}}) \hat{\Phi} (\mathbf{I} - \mathbf{H}_{\mathbf{T}})' (\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{U}}})'_j \mathbf{D}'_j \\ &= \mathbf{D}_j (\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{U}}} - \mathbf{H}_{\mathbf{T}})_j \hat{\Phi} (\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{U}}} - \mathbf{H}_{\mathbf{T}})'_j \mathbf{D}'_j \\ &= \mathbf{D}_j \left( \Phi_j - \ddot{\mathbf{U}}_j \mathbf{M}_{\ddot{\mathbf{U}}} \ddot{\mathbf{U}}'_j - \mathbf{T}_j \mathbf{M}_{\mathbf{T}} \mathbf{T}'_j \right) \mathbf{D}'_j \end{aligned}$$

and

$$\mathbf{B}_j^+ = (\mathbf{D}'_j)^{-1} \left( \Phi_j - \ddot{\mathbf{U}}_j \mathbf{M}_{\ddot{\mathbf{U}}} \ddot{\mathbf{U}}'_j - \mathbf{T}_j \mathbf{M}_{\mathbf{T}} \mathbf{T}'_j \right)^+ \mathbf{D}_j^{-1}. \quad (17)$$

Let  $\Omega_j = \left( \Phi_j - \ddot{\mathbf{U}}_j \mathbf{M}_{\ddot{\mathbf{U}}} \ddot{\mathbf{U}}'_j \right)^+$ . Using a generalized Woodbury identity (Henderson and Searle, 1981),

$$\Omega_j = \mathbf{W}_j + \mathbf{W}_j \ddot{\mathbf{U}}_j \mathbf{M}_{\ddot{\mathbf{U}}} \left( \mathbf{M}_{\ddot{\mathbf{U}}} - \mathbf{M}_{\ddot{\mathbf{U}}} \ddot{\mathbf{U}}'_j \mathbf{W}_j \ddot{\mathbf{U}}_j \mathbf{M}_{\ddot{\mathbf{U}}} \right)^+ \mathbf{M}_{\ddot{\mathbf{U}}} \ddot{\mathbf{U}}'_j \mathbf{W}_j.$$

It follows that  $\Omega_j \mathbf{T}_j = \mathbf{W}_j \mathbf{T}_j$ . Another application of the generalized Woodbury identity gives

$$\begin{aligned} \left( \Phi_j - \ddot{\mathbf{U}}_j \mathbf{M}_{\ddot{\mathbf{U}}} \ddot{\mathbf{U}}'_j - \mathbf{T}_j \mathbf{M}_{\mathbf{T}} \mathbf{T}'_j \right)^+ &= \Omega_j + \Omega_j \mathbf{T}_j \mathbf{M}_{\mathbf{T}} (\mathbf{M}_{\mathbf{T}} - \mathbf{M}_{\mathbf{T}} \mathbf{T}'_j \Omega_j \mathbf{T}_j \mathbf{M}_{\mathbf{T}})^+ \mathbf{M}_{\mathbf{T}} \mathbf{T}'_j \Omega_j \\ &= \Omega_j + \mathbf{W}_j \mathbf{T}_j \mathbf{M}_{\mathbf{T}} (\mathbf{M}_{\mathbf{T}} - \mathbf{M}_{\mathbf{T}} \mathbf{T}'_j \mathbf{W}_j \mathbf{T}_j \mathbf{M}_{\mathbf{T}})^+ \mathbf{M}_{\mathbf{T}} \mathbf{T}'_j \mathbf{W}_j \\ &= \Omega_j. \end{aligned}$$

The last equality follows from the fact that  $\mathbf{T}_j \mathbf{M}_{\mathbf{T}} (\mathbf{M}_{\mathbf{T}} - \mathbf{M}_{\mathbf{T}} \mathbf{T}'_j \mathbf{W}_j \mathbf{T}_j \mathbf{M}_{\mathbf{T}})^- \mathbf{M}_{\mathbf{T}} \mathbf{T}'_j = \mathbf{0}$  because the fixed effects are nested within clusters. Substituting into (17), we then have that  $\mathbf{B}_j^+ = (\mathbf{D}'_j)^{-1} \Omega_j \mathbf{D}_j^{-1}$ . But

$$\tilde{\mathbf{B}}_j = \mathbf{D}_j (\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{U}}})_j \Phi (\mathbf{I} - \mathbf{H}_{\ddot{\mathbf{U}}})'_j \mathbf{D}'_j = \mathbf{D}_j \left( \Phi_j - \ddot{\mathbf{U}}_j \mathbf{M}_{\ddot{\mathbf{U}}} \ddot{\mathbf{U}}'_j \right) \mathbf{D}'_j = \mathbf{D}_j \Omega_j^+ \mathbf{D}'_j,$$

and so  $\mathbf{B}_j^+ = \tilde{\mathbf{B}}_j^+$ . It follows that  $\mathbf{A}_j = \tilde{\mathbf{A}}_j$  for  $j = 1, \dots, m$ .  $\square$

## B DISTRIBUTION THEORY FOR $\mathbf{V}^{CR}$

The small-sample approximations for t-tests and F-tests both involve the distribution of the entries of  $\mathbf{V}^{CR2}$ . This appendix explains the relevant distribution theory.

First, note that any of the CR estimators can be written in the form  $\mathbf{V}^{CR2} = \sum_{j=1}^M \mathbf{P}_j \mathbf{e}_j \mathbf{e}_j' \mathbf{P}_j'$  for  $r \times n_j$  matrices  $\mathbf{P}_j = \mathbf{M}_{\ddot{\mathbf{R}}} \ddot{\mathbf{R}}_j' \mathbf{W}_j \mathbf{A}_j$ . Let  $\mathbf{c}_1, \mathbf{c}_2, \mathbf{c}_3, \mathbf{c}_4$  be fixed,  $p \times 1$  vectors and consider the linear combination  $\mathbf{c}_1' \mathbf{V}^{CR2} \mathbf{c}_2$ . Bell and McCaffrey (2002, Theorem 4) show that the linear combination is a quadratic form in  $\mathbf{y}$ :

$$\mathbf{c}_1' \mathbf{V}^{CR2} \mathbf{c}_2 = \mathbf{y}' \left( \sum_{j=1}^m \mathbf{p}_{2j} \mathbf{p}_{1j}' \right) \mathbf{y},$$

for  $N \times 1$  vectors  $\mathbf{p}_{sh} = (\mathbf{I} - \mathbf{H}_{\mathbf{X}})'_h \mathbf{P}'_h \mathbf{c}_s$ ,  $s = 1, \dots, 4$ , and  $h = 1, \dots, m$ .

Standard results regarding quadratic forms can be used to derive the moments of the linear combination (e.g., Searle, 2006, Sec. 13.5). We now assume that  $\boldsymbol{\epsilon}_1, \dots, \boldsymbol{\epsilon}_m$  are multivariate normal with zero mean and variance  $\boldsymbol{\Sigma}$ . It follows that

$$\mathbb{E}(\mathbf{c}_1' \mathbf{V}^{CR2} \mathbf{c}_2) = \sum_{j=1}^m \mathbf{p}_{1j}' \boldsymbol{\Sigma} \mathbf{p}_{2j} \quad (18)$$

$$\text{Var}(\mathbf{c}_1' \mathbf{V}^{CR2} \mathbf{c}_2) = \sum_{i=1}^m \sum_{j=1}^m (\mathbf{p}_{1i}' \boldsymbol{\Sigma} \mathbf{p}_{2j})^2 + \mathbf{p}_{1i}' \boldsymbol{\Sigma} \mathbf{p}_{1j} \mathbf{p}_{2i}' \boldsymbol{\Sigma} \mathbf{p}_{2j} \quad (19)$$

$$\text{Cov}(\mathbf{c}_1' \mathbf{V}^{CR2} \mathbf{c}_2, \mathbf{c}_3' \mathbf{V}^{CR} \mathbf{c}_4) = \sum_{i=1}^m \sum_{j=1}^m \mathbf{p}_{1i}' \boldsymbol{\Sigma} \mathbf{p}_{4j} \mathbf{p}_{2i}' \boldsymbol{\Sigma} \mathbf{p}_{3j} + \mathbf{p}_{1i}' \boldsymbol{\Sigma} \mathbf{p}_{3j} \mathbf{p}_{2i}' \boldsymbol{\Sigma} \mathbf{p}_{4j}. \quad (20)$$

Furthermore, the distribution of  $\mathbf{c}_1' \mathbf{V}^{CR2} \mathbf{c}_2$  can be expressed as a weighted sum of  $\chi_1^2$  distributions (Mathai and Provost, 1992), with weights given by the eigen-values of the  $m \times m$  matrix with  $(i, j)^{th}$  entry  $\mathbf{p}_{1i}' \boldsymbol{\Sigma} \mathbf{p}_{2j}$ ,  $i, j = 1, \dots, m$ .

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