

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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1 INTRODUCTION

2 STANDARD CLUSTER-ROBUST VARIANCE ESTIMATION

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.

2.1 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 might include indicators for each time period, and \mathbf{R}_j might include a set of policy 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 $\mathbf{U}_j = [\mathbf{R}_j \ \mathbf{S}_j]$, $\mathbf{X}_j = [\mathbf{U}_j \ \mathbf{T}_j]$, $\boldsymbol{\alpha} = (\boldsymbol{\beta}', \boldsymbol{\gamma}', \boldsymbol{\delta}')'$, 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. The distinction between the two fixed effect matrices \mathbf{S}_j and \mathbf{T}_j is less ambiguous, 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 $\boldsymbol{\beta}$. 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.¹ In the latter case, it is assumed that $\text{Var}(\mathbf{e}_j | \mathbf{X}_j) = \boldsymbol{\Phi}_j$, where $\boldsymbol{\Phi}_j$ 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 $\mathbf{W}_j = \hat{\boldsymbol{\Phi}}_j^{-1}$, where the $\hat{\boldsymbol{\Phi}}_j$ 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 recover design-unbiased estimates of population parameters.

Several approaches computing the WLS estimator are possible. One possibility is to calculate WLS estimates of the full parameter vector $\boldsymbol{\alpha}$ 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 $\mathbf{K}'\boldsymbol{\alpha} = \boldsymbol{\beta}$. 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 $\boldsymbol{\beta}$. Let

¹The 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 $\boldsymbol{\beta}$ separately within each cluster and taking the simple average of these estimates. The resulting average is equivalent to the WLS estimator with weights $\mathbf{W}_j = \mathbf{X}_j (\mathbf{X}_j' \mathbf{X}_j)^{-2} \mathbf{X}_j$.

$\ddot{\mathbf{S}} = (\mathbf{I} - \mathbf{H}_{\mathbf{T}}) \mathbf{S}$, $\ddot{\mathbf{R}} = (\mathbf{I} - \mathbf{H}_{\mathbf{S}}) (\mathbf{I} - \mathbf{H}_{\mathbf{T}}) \mathbf{R}$, and $\ddot{\mathbf{y}} = (\mathbf{I} - \mathbf{H}_{\mathbf{S}}) (\mathbf{I} - \mathbf{H}_{\mathbf{T}}) \mathbf{y}$. In what follows, subscripts on $\ddot{\mathbf{R}}$, $\ddot{\mathbf{S}}$, $\ddot{\mathbf{U}}$, and $\ddot{\mathbf{y}}$ refer to the rows of these matrices corresponding to a specific cluster. The WLS estimator of $\boldsymbol{\beta}$ can then be written as

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

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

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

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

The variance of the WLS estimator is

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

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.

2.2 Standard CRVE

Cluster-robust variance estimators provide a means of estimating $\text{Var}(\hat{\boldsymbol{\beta}})$ and testing hypotheses regarding $\boldsymbol{\beta}$ 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

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

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

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 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 *Longitudinal data analysis using generalized linear models* (1986), among others. 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 Σ_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 will be described in the next section. Because it is an extension of the HC2 estimator for regressions with heteroskedastic but uncorrelated errors, we refer to it as CR2. A further alternative is to use a jack-knife resampling estimator, or an approximation to it. The CR3 estimator closely approximates the jack-knife approach, by taking $\mathbf{A}_j = \left(\mathbf{I} - \ddot{\mathbf{R}}_j \mathbf{M}_{\ddot{\mathbf{R}}} \ddot{\mathbf{R}}_j' \mathbf{W}_j \right)^{-1}$.

3 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 follow 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 example, 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 Φ_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.² When the working model deviates from the true covariance Σ_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. One way to understand this approach is that it 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 even when \mathbf{B}_j is not of full rank. [Theorem 1 in Appendix](#)

Add citation.

²Note 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 permits the use of generalized matrix inverses in calculating the adjustment matrices, thus avoiding rank-deficiency problems that would otherwise leave them undefined.

A shows that the adjustment matrices given by (7) and (8) satisfy criterion (6) and are invariant 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 $\ddot{\mathbf{U}} = (\mathbf{I} - \mathbf{H}_{\mathbf{T}}) \mathbf{U}$.

In the remainder of this paper, we will focus on this BRL approach, using the \mathbf{V}^{CR2} estimator throughout.

4 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., $m \rightarrow \infty$). However, evidence from a wide variety of contexts indicates that the asymptotic results 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 β , 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 α if Q exceeds $\chi^2(\alpha; q)$, the α critical value from a chi-squared distribution with q degrees of freedom. When samples are small, in standard practice instead the test $F = Q/q$ is often used with the CR1 estimator and the reference distribution $F(q, m - 1)$.

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, i.e., $H_0 : \beta_i = 0$. For one-dimensional constraints, an equivalent to the Wald F test is to use the test statistic $Z = \hat{\beta}_i / \sqrt{V_i^{CR}}$, the distribution of which is approximately standard normal in large samples. In small samples, one can instead approximate the distribution of Z by a $t(m - 1)$ distribution. However, Bell and McCaffrey (2002) showed that, even when the CR2 estimator is used, Z does not follow a $t(m - 1)$ distribution in small samples. They then proposed 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 following subsection reviews the Bell and McCaffrey (2002) approach for one-dimensional hypothesis tests, which applies directly in fixed-effect models. We then propose a method for testing more general, q -dimensional linear hypotheses regarding β , by approximating the distribution of Q using an F distribution with estimated degrees of freedom.

4.1 Small-sample corrections for t-tests

Our approach to developing a better small-sample F-test parallels that the t-test developed by Bell and McCaffrey (2002). In this section, we therefore review this approach. As noted, the standard test implemented in software uses the CR1 correction with the $t(m - 1)$ reference distribution. The first and surely most common approach is to compare $|Z|$ to the

appropriate critical value from a t distribution with $m-1$ degrees of freedom. 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 α). However, both of these arguments require that $\mathbf{c}'\boldsymbol{\beta}$ 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.

In contrast, the approach developed by McCaffrey et al. (2001) is to instead estimate the degrees of freedom of the t-test using a Satterthwaite approximation (Satterthwaite, 1946). This approach compares Z to a t reference distribution, with degrees of freedom ν that are estimated from the data. Theoretically, the 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. McCaffrey et al. (2001) proposed to estimate the moments based on the same working model as used to derive the adjustment matrices. A “model-based” estimate of the degrees of freedom is then calculated as

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

where $\mathbf{s}_j = (\mathbf{I} - \mathbf{H})'_j \mathbf{A}'_j \mathbf{W}_j \mathbf{X}_j \mathbf{M} \mathbf{c}$. 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 the plug-in degrees of freedom estimate produced very conservative rejection rates.

The McCaffrey et al. (2001) 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. Importantly, 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 – 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 considerably from cluster to cluster – these degrees of freedom can be tremendously smaller. Similarly, covariates with large leverage points can exhibit similar losses in terms of degrees of freedom. The result is that the small-sample corrections are required even when the number of clusters seems large, suggesting that this CR2 t-test be applied as a default in all CRVE based analyses.

4.2 Small-sample corrections for F-tests

Compared to single-constraint tests, fewer approaches to small-sample correction are available for multiple-constraint tests. 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).

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 η degrees of freedom and

scale matrix $\mathbf{CVar}(\mathbf{C}\hat{\beta})\mathbf{C}'$, 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 η). To do so, we estimate the degrees of freedom of the Wishart distribution so that they match the mean and variance of $\mathbf{CV}^{CR}\mathbf{C}'$. 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 $\eta\mathbf{CV}^{CR}\mathbf{C}'$ and the covariances of the Wishart distribution with η degrees of freedom and scale matrix $\mathbf{CV}^{CR}\mathbf{C}'$. 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 $\mathbf{CV}^{CR}\mathbf{C}'$ (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 $\mathbf{t}_{sh} = (\mathbf{I} - \mathbf{H})'_h \mathbf{A}'_h \mathbf{W}_h \mathbf{X}_h \mathbf{M} \mathbf{c}_s$ for $s = 1, \dots, q$ and $h = 1, \dots, m$. The degrees of freedom are then estimated under the working model as

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

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

5 SIMULATION EVIDENCE

6 EXAMPLES

6.1 Tennessee STAR class-size experiment.

6.2 Heterogeneous treatment impacts

6.3 Robust Hausmann test

7 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 Φ 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 orthogonal 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 the CR2 estimator can be written in the form $\mathbf{V}^{CR2} = \sum_{j=1}^M \mathbf{T}_j \mathbf{e}_j \mathbf{e}_j' \mathbf{T}_j'$ for $p \times n_j$ matrices $\mathbf{T}_j = \mathbf{M} \mathbf{X}_j' \mathbf{W}_j \mathbf{A}_j$. 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} :

$$\mathbf{c}_1' \mathbf{V}^{CR2} \mathbf{c}_2 = \mathbf{Y}' \left(\sum_{j=1}^m \mathbf{t}_{2j} \mathbf{t}_{1j}' \right) \mathbf{Y},$$

for $N \times 1$ vectors $\mathbf{t}_{sh} = (\mathbf{I} - \mathbf{H})_h' \mathbf{T}_h' \mathbf{c}_s$, $s = 1, \dots, 4$, and $h = 1, \dots, m$.

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

$$\mathbb{E} (\mathbf{c}_1' \mathbf{V}^{CR2} \mathbf{c}_2) = \sum_{j=1}^m \mathbf{t}_{1j}' \boldsymbol{\Sigma} \mathbf{t}_{2j} \quad (18)$$

$$\text{Var} (\mathbf{c}_1' \mathbf{V}^{CR2} \mathbf{c}_2) = \sum_{i=1}^m \sum_{j=1}^m (\mathbf{t}_{1i}' \boldsymbol{\Sigma} \mathbf{t}_{2j})^2 + \mathbf{t}_{1i}' \boldsymbol{\Sigma} \mathbf{t}_{1j} \mathbf{t}_{2i}' \boldsymbol{\Sigma} \mathbf{t}_{2j} \quad (19)$$

$$\text{Cov} (\mathbf{c}_1' \mathbf{V}^{CR2} \mathbf{c}_2, \mathbf{c}_3' \mathbf{V}^{CR} \mathbf{c}_4) = \sum_{i=1}^m \sum_{j=1}^m \mathbf{t}_{1i}' \boldsymbol{\Sigma} \mathbf{t}_{4j} \mathbf{t}_{2i}' \boldsymbol{\Sigma} \mathbf{t}_{3j} + \mathbf{t}_{1i}' \boldsymbol{\Sigma} \mathbf{t}_{3j} \mathbf{t}_{2i}' \boldsymbol{\Sigma} \mathbf{t}_{4j}. \quad (20)$$

Furthermore, the distribution of $\mathbf{c}_1' \mathbf{V}^{CR2} \mathbf{c}_2$ can be expressed as a weighted sum of χ_1^2 distributions, with weights given by the eigen-values of the $m \times m$ matrix with $(i, j)^{th}$ entry $\mathbf{t}_{1i}' \boldsymbol{\Sigma} \mathbf{t}_{2j}$, $i, j = 1, \dots, m$.

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