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Christian A. Vossler

1. Introduction

Many laboratory and field experiments in economics involve participants or groups of participants making a sequence of related decisions, usually with feedback, over many choice periods. For instance, this is typical of experimental work on auctions, bargaining, the private provision of public goods, tax compliance, and pollution control instruments. Through repeatedgame play, researchers allow for developments such as learning, strategy refinement, establishment of equilibria, and observances of how decisions or outcomes change in response to experimental design variations. The widespread availability and improving functionality of computer software has made it increasingly common for experiments to be reasonably complex and involve many choice periods.

Experimentalists traditionally have relied on fairly simple and computationally transparent parametric and nonparametric hypothesis tests to evaluate hypotheses (e.g. paired *t*-test, Wilcoxon test), such as those discussed in Davis and Holt (1993). It remains a somewhat common practice to address the time-series dimension superficially by using as the unit of observation the mean outcome across all periods for an individual or group. Time trends may be artificially accounted for by using the average outcome from the last decision period, last few periods, or by separately testing different period groupings. Such analyses rely on the variation in means across individuals or groups and insufficiently accounts for the variation in outcomes across decision periods. These approaches are particularly troublesome for experimental designs that expose the participant to multiple parameter changes.

Particularly in the last several years, experimentalists have relied more on available estimation methods for panel data.¹ These methods include standard random effects (or closely related mixed effects) and fixed effects models, as well as the use of common estimators for cross-section data (e.g. OLS) in tandem with "robust" covariance matrix estimators such as White's (1980) heteroskedasticity-consistent estimator and Beck and Katz's (1995) "panelcorrected standard errors". While these approaches allow analysis of the full data set while accounting for important forms of heterogeneity, they may inadequately address inference issues related to serial correlation. In some instances where serial correlation has been explicitly addressed in experimental analyses, convenient parametric modeling approaches have been employed, such as the inclusion of a lagged dependent variable as an additional covariate or assuming model errors follow an AR(1) process (see, for example, Ashley, Ball, and Eckel 2003; Rassenti, Smith, and Wilson 2003). Alternatively, some recent studies use OLS in tandem with the "cluster-robust" covariance estimator, which – as I discuss in further detail in this study – can lead to valid inferences when within-unit serial correlation is unspecified in the regression model. Surprisingly, many of these papers do not mention serial correlation (e.g. Ashraf, Bonet, and Piankov 2006; Shupp and Williams 2008; Baker, Walker and Williams 2009), and thus the theoretical and empirical properties of this approach may be poorly understood by some.²

Serial correlation is, or at least should be, an important consideration for repeated-game experiments, especially when the number of decision periods is large or when the cross-section and time dimensions are of similar magnitude.³ When serial correlation is left unspecified, the standard errors of common estimators (and sometimes the estimators themselves), and hypothesis tests based on them, are biased. Within a linear regression framework, unlike the case for heteroskedasticity, a consensus has not been reached regarding a covariance estimator for

panel data that is robust to serial correlation of unknown form. This is unfortunate for experimentalists, and indeed many applied researchers who are largely interested in testing hypotheses rather than deciphering the particular structure of the error correlation.

This study endeavors to provide some guidance to those who analyze data from repeatedgame experiments. In particular, I propose the use of heteroskedasticity-autocorrelation consistent (HAC) covariance estimators for panel data, which allows researchers to conduct hypothesis tests without having to place structure on the heteroskedasticity and/or serial correlation likely present in econometric models. Through Monte Carlo experiments I explore the properties of three panel HAC covariance estimators within a linear regression framework, including a new HAC covariance estimator proposed in this study, for a range of cross-section (n) and time (T) dimensions relevant for economics experiments. The new estimator, a randomeffects HAC covariance estimator (hereafter, RE-HAC), is a panel version of the Newey and West (1987) covariance estimator that allows for a unit-specific random effect. The other two HAC covariance estimators investigated, the cluster-robust covariance estimator of Arellano (1987) (hereafter, A-HAC) and the standard panel version of the Newey-West (1987) estimator (hereafter, NW-HAC), are currently available through canned routines in popular econometrics software packages. Overall, the results of the Monte Carlo simulations provide strong support for adding panel HAC covariance estimators to the toolbox of experimentalists.

Most of the previous work on HAC estimation is in the context of time-series data. Although HAC covariance estimators are consistent under reasonable assumptions for $T \rightarrow \infty$ (see Newey and West 1987), results from Monte Carlo experiments suggest that the finite sample properties of HAC covariance estimators can be quite poor. In particular, even with large sample sizes, HAC standard errors tend to be too small in the presence of complicated heteroskedasticity and serial correlation patterns or when the degree of serial correlation is high, leading to gross over-rejection under the null hypothesis (Andrews 1991; Andrews and Monahan 1992; Newey and West 1994; den Haan and Levin 1997; Cushing and McGarvey 1999). Further, unlike the straightforward heteroskedasticity-consistent covariance estimators, at least in the time-series realm, the analyst must choose a kernel (a rule for weighting sample autocovariances) and a bandwidth (the number of autocovariances included). It is also common to use a prewhitening filter, and finite sample performance of HAC covariance estimators can depend greatly on all three choices.⁴

The infrequent use of HAC covariance estimators in the time-series literature is likely a result of unsupportive Monte Carlo evidence. This begs the question: why should we consider using HAC standard errors in a panel data context, in particular for experiment data? There are at least three reasons. First, construction of the panel HAC covariance matrix involves the averaging of autocovariances across cross-section units, and this averaging is likely to lessen the finite sampling variability introduced by the particular kernel and bandwidth chosen by the analyst (see den Hann and Levin 1997; Keifer and Vogelsang 2002, 2005). Thus, for a small or modest *n*, the performance of the HAC covariance estimator is likely to be reasonably insensitive to choice of kernel and bandwidth. Arellano (1987), based on White (1984), proves the $n \rightarrow \infty$ consistency of the A-HAC covariance estimator, which includes all autocovariances and for which all autocovariances are given full weight. In other words, for a large enough cross-section, the analyst is at least theoretically justified setting the bandwidth equal to *T* and foregoing the use of a kernel to weight autocovariances.

The theoretical results of Newey and West (1987) and Arellano (1987) together suggest a second reason to explore HAC covariance estimators in a panel context, namely, that it is

possible to achieve consistent covariance estimation with either large n or T (or both). This suggests that HAC covariance estimators may perform well for data sets with a large crosssection dimension and/or a large time-series dimension. Third, the performance of HAC covariance estimators generally deteriorates when the explanatory variables are themselves serially correlated, and the correlation differs across variables (den Haan and Levin 1997). However, explanatory variables in a regression model for experiment data are typically treatment indicator variables, design-specific variables exogenously determined by the experimentalist, and (time-invariant) participant characteristics.

Similar to the time-series literature, much of what we know about panel HAC covariance estimators is based on Monte Carlo experiments, although there have been few such studies. Bertrand, Duflo, and Mullainathan (2004) and Kezdi (2004) investigate the A-HAC covariance estimator within fixed effect frameworks. Similar to these studies, I find that test statistics based on A-HAC have the correct size for panels with a moderate cross-section dimension (e.g. n =50), but with smaller cross-section dimensions (e.g. n = 10) standard errors are biased downward. Driscoll and Kraay (1998) propose a panel HAC covariance estimator that is also robust to spatial correlation, and provide Monte Carlo evidence that their estimator performs better than OLS and seemingly unrelated regression (SUR) when there is spatial correlation. Their estimator is similar to the NW-HAC estimator explored in this study, with the important exception that it is constructed from cross-sectional averages of the autocovariances. This estimator requires that parameters not vary across cross-section units, and unfortunately, this restriction is likely to be violated in the analysis of experiment data (e.g. it would preclude estimation of treatment effects). This study contains further explorations of panel HAC covariance estimators, with a focus on data generating processes (DGPs) and panel dimensions relevant for experimental economics applications. NW-HAC and the proposed RE-HAC estimator have not been previously explored with Monte Carlo methods. In contrast to the existing simulation work on A-HAC in the context of fixed-effects models, I consider estimation in the presence of unobserved heterogeneity in the form of a unit-specific random effect. This is particularly relevant to experimentalists since: (1) unobserved individual or group-specific heterogeneity is unlikely to be correlated with included model covariates; and (2) we are commonly interested in estimating coefficients on time-invariant variables, such as treatment indicator variables and subject-specific characteristics. The simulations further consider serial correlation processes.

The next section presents an overview of HAC covariance estimation in a time-series setting. This background material is useful as two of the HAC covariance estimators are panel extensions of time-series HAC covariance estimators. Section 3 provides some details of the three HAC covariance estimators in the context of panel data. Sections 4 and 5 present Monte Carlo simulation results designed to assess the accuracy of HAC-based hypothesis tests. Section 6 provides some recommendations.

2. HAC Covariance Matrix Estimators for Time-Series Data

This section overviews HAC covariance estimation within the context of analyzing a single time series. Capitalizing on the nice robustness properties of OLS, and beginning with the seminal work of White (1980), researchers in economics and elsewhere have made valid inferences in the presence of unknown heteroskedasticity by estimating coefficients using OLS and using White's heteroskedasticity-consistent covariance estimator in place of the usual OLS covariance matrix. Newey and West (1987) extended consistent covariance estimation by

developing an estimator that is robust to both heteroskedasticity and serial correlation. While the motivation behind the White and Newey-West estimators is the same – to construct a consistent covariance matrix for least squares parameters – controlling for temporal dependence of unknown form is a demanding task.

Consider the least squares regression model:

$$y_t = \mathbf{x}_t' \boldsymbol{\beta} + \varepsilon_t, \tag{1}$$

where $\boldsymbol{\beta}$ and \mathbf{x}_t are $k \times 1$ vectors of estimable parameters and covariates, respectively; ε_t is a mean zero error term (scalar) that is possibly serially correlated and conditionally heteroskedastic, with $E[\varepsilon_t | \mathbf{x}_t] = 0$. With serially correlated and heteroskedastic errors, the asymptotic covariance of the ordinary least squares estimator of $\boldsymbol{\beta}$ is:

Asy. Var
$$[\boldsymbol{b}] = (\mathbf{X}'\mathbf{X})^{-1} \{ \sum_{t=1}^{T} \sum_{j=1}^{T} E[\mathbf{x}_t \varepsilon_t \varepsilon_j \mathbf{x}'_j] \} (\mathbf{X}'\mathbf{X})^{-1},$$
 [2]

where **X** is the full $T \times k$ matrix of covariates. The difficulty lies in suitably estimating the autocovariance matrix, the middle term in equation [2], using the least squares residuals (e_t) as point-wise realizations of the true population disturbances. Newey and West (1987) show that a positive semi-definite, consistent covariance estimator can be constructed by appropriately weighting the sample autocovariances, $\mathbf{x}_t e_t = \mathbf{x}_t (y_t - \mathbf{x}'_t \mathbf{b})$, in such a way that the dependence between observations goes to zero as the distance between observations increases. They suggest using a kernel spectral density estimator evaluated at frequency zero, which requires choosing a kernel function and a bandwidth parameter.

For a given dataset, one can arguably choose among many such kernel/bandwidth pairs to construct a consistent covariance estimator. Since the estimated covariance matrix approaches a constant value as *T* tends towards infinity, HAC-based test statistics typically have a normal (single linear hypotheses) or chi-squared (multiple linear hypotheses) limiting distribution. However, in finite samples, the choice of kernel and bandwidth can severely distort test statistics based on these distributions. That is to say, the choice of kernel and bandwidth introduces finite sampling bias, the extent to which depends on sample size and the underlying DGP. For this reason, Andrews (1991), Newey and West (1994), and others, have developed data-dependent bandwidth selection procedures (taking the kernel as given) under the premise of minimizing the mean-squared error of the HAC covariance matrix. While these selection procedures provide guidance for the analyst and generally perform better than HAC covariance estimators using an arbitrary choice for bandwidth and kernel, Monte Carlo experiments suggest that these data-dependent HAC covariance estimators do not fully resolve the tendency for HAC covariance estimators to over-reject the null hypothesis when it is true (Andrews 1991; Andrews and Monahan 1992; Newey and West 1994; den Haan and Levin 1997; Cushing and McGarvey 1999).

To increase the performance of HAC covariance estimators, Andrews and Monahan (1992) suggest prewhitening the sample autocovariances using a first-order vectorautoregression [VAR(1)] filter. The VAR(1) filter estimates the value of an autoregressive root based on the first-order autocovariance. After filtering this autoregressive root, the autocovariances of the prewhitened residuals may decline more rapidly toward zero, thereby reducing the bias of the kernel-based estimator (Haan and Levin 1997). Andrews and Monahan (1992) show that prewhitening can provide benefits even when the true DGP is not a low-order VAR process. A fairly standard practice involves constructing a HAC covariance estimator by using a prewhitening filter, choosing a kernel, and selecting a bandwidth based on one of the data-dependent selection methods. Alternative approaches include parametric spectral density estimators (see den Haan and Levin 1997) and using the limiting distributions derived by Keifer and Vogelsang (2005) for HAC-based test statistics, which serve as better finite sampling distributions for HAC tests than do the normal or chi-squared distributions.

3. HAC Covariance Estimators for Panel Data

HAC covariance estimation with panel data is not new. In fact, recent versions of the statistical software packages Limdep and Stata include procedures for estimating two panel HAC covariance estimators. The first is a panel extension of the Newey-West estimator (NW-HAC).⁵ The second is the cluster-robust estimator (A-HAC).⁶

Consider the following panel model specification

$$y_{it} = \mathbf{x}'_{it}\boldsymbol{\beta} + u_i + \varepsilon_{it},\tag{3}$$

where the data from the *n* cross-section units are stacked and u_i is a mean-zero, unobserved unitspecific effect. As before, the ε_{it} are possibly serially correlated and conditionally heteroskedastic disturbances. For convenience, and with a slight abuse of notation, the model in [3] can be written as

$$\mathbf{y}_i = \mathbf{X}_i \boldsymbol{\beta} + \mathbf{i} u_i + \boldsymbol{\varepsilon}_i, \tag{4}$$

where \mathbf{y}_i and $\boldsymbol{\varepsilon}_i$ are $T \times 1$ vectors specific to unit *i*, \mathbf{X}_i is a $T \times k$ matrix of covariates for unit *i*, **i** is a $T \times 1$ column of 1s and u_i is defined as before. Assuming away the unit-specific effect for the moment, under the assumption that cross-section units are independent, the asymptotic covariance matrix for the OLS estimator is

Asy. Var
$$[\mathbf{b}] = nT(\sum_{i=1}^{n} \mathbf{X}'_{i} \mathbf{X}_{i})^{-1} \{n^{-1} \sum_{i=1}^{n} \mathbf{V}_{i}\}(\sum_{i=1}^{n} \mathbf{X}'_{i} \mathbf{X})^{-1}.$$
 [5]

HAC covariance estimators differ in how they estimate $\mathbf{V}_i = E[\mathbf{X}'_i \boldsymbol{\varepsilon}_i \boldsymbol{\varepsilon}'_i \mathbf{X}_i] = E[\mathbf{X}'_i \boldsymbol{\Omega}_i \mathbf{X}_i]$.⁷ A-HAC, in contrast to the HAC covariance estimators used for time-series data, uses all autocovariances (i.e. bandwidth equals *T*) and no kernel function to weight them. In particular,

$$\widehat{\mathbf{V}}_{i}^{A-HAC} = T^{-1}\mathbf{X}_{i}'\mathbf{e}_{i}\mathbf{e}_{i}'\mathbf{X}_{i}$$
^[6]

where the \mathbf{e}_i are the OLS residuals. The resulting covariance estimator is consistent for large n and fixed T, but not for fixed n and large T (Arellano 2003). NW-HAC uses the Bartlett kernel, which is computationally simple

$$w_{tj} = 1 - \frac{|t-j|}{m_i}$$
 for $|t-j| < m_i$; $w_{tj} = 0$ for $|t-j| \ge m_i$ [7]

where m_i is the bandwidth parameter and w_{tj} is the weight given to the tj^{th} sample autocovariance. Let $\mathbf{W} = [w_{tj}]$ denote a $T \times T$ matrix of Bartlett kernel weights. NW-HAC estimates \mathbf{V}_i with

$$\widehat{\mathbf{V}}_{i}^{NW-HAC} = T^{-1}\mathbf{X}_{i}'((\mathbf{e}_{i}\mathbf{e}_{i}') \bullet \mathbf{W})\mathbf{X}_{i},$$
[8]

where "•" denotes the Hadamard product operator. For the panel specification (equation [4]), if $E[u_i|\mathbf{X}_i] \neq 0$, in which case consistency of the OLS estimator for $\boldsymbol{\beta}$ generally requires the inclusion of unit-specific fixed effects in **X** (i.e. a fixed-effects model is estimated), A-HAC is consistent in *n* and NW-HAC is consistent in *T*.

Suppose that we seek an alternative to the fixed effects estimator, for instance in a case where we wish to estimate coefficients on time-invariant variables, such as subject characteristics or treatment indicators, and we are comfortable with the assumption of zero correlation between u_i and included covariates. In this case, under standard assumptions we can instead consistently estimate β using OLS (without including the fixed-effects). A-HAC remains consistent in *n*, but NW-HAC is no longer consistent. The intuition behind the inconsistency of NW-HAC is reasonably straightforward. When there is an underlying random effects structure, the OLS residuals are pointwise estimates of $\mathbf{i}u_i + \mathbf{e}_i$, and the variance of the random effect, σ_u^2 , appears (unweighted) in every element of Ω_i (see Greene 2002, pp. 294). Thus, the Bartlett kernel (or any kernel) used for NW-HAC scales the off-diagonal terms of Ω_i and thus underestimates them. In the case of no heteroskedasticity or serial correlation, for example, all Ω_i off-diagonal elements simply equal σ_u^2 but NW-HAC would instead use $w_{tj}\sigma_u^2 < \sigma_u^2$.

I propose an extension of NW-HAC, which I refer to as RE-HAC, which is consistent in *T* while allowing for a unit-specific random effect. To be clear, this is a covariance matrix for the OLS estimator, **b**, and not for the random effects estimator. The extension involves simply incorporating the variance of the random effect, σ_u^2 , into **V**_i. This requires estimates of the population disturbances $\boldsymbol{\varepsilon}$ (to which the kernel should be applied) and an estimate of σ_u^2 (which should appear in every element of $\boldsymbol{\Omega}_i$). The residuals from a fixed effects model are consistent estimates of $\boldsymbol{\varepsilon}$. There are several available consistent estimators for σ_u^2 as discussed in Greene (2002, p. 297-298).

Then, it can be shown that a consistent estimator for V_i under the assumption of random effects is

$$\widehat{\mathbf{V}}_{i}^{RE-HAC} = T^{-1}\mathbf{X}_{i}'((\mathbf{e}_{i}\mathbf{e}_{i}') \bullet (\mathbf{W} + \widehat{\sigma}_{u}^{2}\mathbf{i}\mathbf{i}'))\mathbf{X}_{i}.$$
[9]

The proof of consistency is straightforward; here I just illustrate this through two polar cases. First, if $\sigma_u^2 = 0$ (i.e. there is no unobserved unit effect) then RE-HAC is equivalent to NW-HAC, the properties of which are known. Second, under the assumption of random effects but no serial correlation or conditional heteroskedasticity, the estimator reduces to

$$E[\widehat{\mathbf{V}}_{i}^{RE-HAC}] = T^{-1}\mathbf{X}_{i}'(\sigma_{\varepsilon}^{2}\mathbf{I} + \sigma_{u}^{2}\mathbf{i}\mathbf{i}')\mathbf{X}_{i},$$
^[10]

where **I** is a $T \times T$ identity matrix. Plugging [10] into [5] yields the (consistent) asymptotic covariance matrix for **b** under the standard assumptions for a random effects model.⁸

Important practical considerations for NW-HAC and RE-HAC are, as in the case of timeseries HAC covariance estimators, the choice of bandwidth and prewhitening filter. For the procedures for NW-HAC available in Stata and Limdep the user must specify the bandwidth that is common to all *i* and there is no prewhitening filter option. Alternatively, one might apply one of the data-dependent bandwidth selection methods developed for time-series models (e.g. Andrews 1991; Newey and West 1994). One option would be to apply a data-dependent selection method separately for each cross-section unit (which would result in bandwidths that vary across units). Alternatively, if one desired a single bandwidth, one might rely on averaging (e.g. taking the average unit-specific bandwidths or applying the selection method to averaged residuals). For the purpose of conducting Monte Carlo simulations, I use the popular AR(1) datadependent bandwidth selection procedure of Andrews (1991) applied to each cross-section unit.⁹ No prewhitening filter is used.

4. Monte Carlo Experiment

In this section, I present results from a Monte Carlo experiment in order to help assess the accuracy of the three HAC covariance estimators described above, and to compare them to some familiar estimators. I consider a linear regression with an intercept $\beta_1 = 0.5$ and one intercept shifter $\beta_2 = 5$:

$$y_{it} = 0.5 + 5x_{it} + u_i + \varepsilon_{it},\tag{11}$$

where $x_{it} = 1$ for $t > \frac{1}{2}T$ and equals 0 otherwise. This simple model is intended to correspond with a within-subjects design and two experimental treatment conditions with an equal number of decision periods. The most extensive DGP considered includes a unit-specific AR(2) serial correlation pattern, as well as a unit-specific random effect:

$$\varepsilon_{it} = \rho_1^i \varepsilon_{i,t-1} + \rho_2^i \varepsilon_{i,t-2} + \eta_{it}$$

$$\eta_{it} \sim Normal(0, 1 - r)$$

$$\sigma_u^2 \sim Normal(0, r)$$
[12]

The parameter r determines the relative within versus between unit variation. In particular, $\sigma_u^2 + \sigma_\eta^2 = 1$ and $r = \sigma_u^2/(\sigma_u^2 + \sigma_\eta^2)$. Four values of $r = \{0.0, 0.3, 0.6, 0.9\}$ are explored. By placing restrictions on the DGP above, there are four basic serial correlation cases: (1) no serial correlation $[\rho_1^i = 0, \rho_2^i = 0]$; (2) an AR(1) process that is common to all units $[\rho_1^i = \rho_1; \rho_2^i = 0]$; (3) an AR(2) process that is common to all units $[\rho_1^i = \rho_1; \rho_2^i = \rho_2]$; and (4) an AR(2) process that differs across units. For the AR(1) case, $\rho_1 = \{0.0, 0.3, 0.6, 0.9\}$. For the AR(2) common process case, two sets of values are explored: $\rho_1 = 0.4$, $\rho_2 = 0.2$; and $\rho_1 = 0.5$, $\rho_2 = 0.4$. For the AR(2) heterogeneous process case, draws from a uniform distribution with supports -.2 and .2 are added to the two sets of values: $\rho_1^i = 0.4 + U[-.2, .2]$, $\rho_2^i = 0.2 + U[-.2, .2]$; and $\rho_1^i = 0.2 + U[-.2, .2]$. The interaction of each distinct serial correlation process with the four values of r leads to 32 distinct parameter settings.

Simulation results for four combinations produced with the cross-section and time dimensions $n = \{5, 30\}$ and $T = \{20, 50\}$ are reported. The cross-section dimensions thus capture

a reasonable number of replications for group-level (i.e. n = 5) and individual-level (i.e. n = 30) outcomes, and the time dimensions capture a small and a moderate number of game repetitions. For each $\{n, T\}$ combination, reported results for each of the 32 parameter settings are based on 1,000 simulation repetitions. In the simulations, the coefficients β_1 and β_2 are estimated using OLS along with four sets of standard errors: RE-HAC, A-HAC, NW-HAC, and uncorrected OLS (OLS). Further, the model is estimated using a standard FGLS random effects model (RE) and a random effects estimator that assumes a common AR(1) error process (RE-AR1). Simulations are carried out using Limdep (version 9) software.¹⁰

Tables 1-4 present Monte Carlo experiment results where each table corresponds to a particular $\{n, T\}$ combination. In particular, reported are the empirical probabilities of rejecting the null hypotheses $\beta_1 = 0.5$ and $\beta_2 = 5$ (consistent with the DGP) based on *t*-tests. 5% critical values were used so that the nominal level is 0.05. Thus, rejection probabilities that are close to 0.05 suggest that the test has the correct size, whereas probabilities above (below) 0.05 suggest over-rejection (under-rejection) under the null hypothesis. The coefficient estimators performed well under all scenarios, and common statistics corresponding with the coefficient estimators (e.g. bias, efficiency, mean-squared error) are omitted for brevity.¹¹ Several interesting patterns emerge with respect to the three HAC covariance estimators. First, as conjectured above, hypothesis tests using NW-HAC are severely distorted – in many cases by a factor of 5 or higher – in the presence of unit effects (i.e. r > 0) for all considered sample sizes. The rejection rates have a similar pattern to those involving the usual OLS standard errors, which are of course biased in the presence of a unit-specific effect. In particular, the null hypothesis with respect to β_1 is rejected too often and the rejection rates for β_2 are too low, with less than a 1% rejection

rate for relatively moderate (r = 0.6) or large (r = 0.9) between-unit variation. Performance under both random effects and serial correlation reveals a similar pattern.

The new RE-HAC standard errors are an improvement over NW-HAC, although there are size distortions evident in some simulations.

[Table 1 Here]

For n = 5, T = 20, rejection rates for β_1 are about 2.5 times too large – on average – across scenarios. This size distortion increases with r. For β_2 , the tendency to over-reject the null remains, but is less severe for the no serial correlation DGP. For the serial correlation DGPs, the size distortion of β_2 increases as the degree of serial correlation increases. Across these settings, the average size is about 0.15 or 3 times the significance level. The size of the test statistics improve with increases in n and/or T. With n = 30, rejection rates for β_1 have approximately the correct size.

[Table 2 Here]

Although there is improvement, the tendency to over-reject the null for β_2 remains. Even with n = 30 and T = 50, the rejection rates for β_2 are about 2.5 times too large for moderate values of r and a moderate degree of serial correlation. RE-HAC in particular performs poorly under the AR(2) specification, especially with $\rho_2 = 0.4$, and this is presumably due to the use of the AR(1)-based bandwidth selection procedure. Overall, for β_1 , the RE-HAC rejection rates are similar to tests based on RE-AR1. For β_2 , however, RE-AR1 performs much better when the DGP is random effects with AR(1) serial correlation – which is to be expected since the estimator is fully consistent with the DGP; but, the performance of the two estimators is comparable under AR(2) serial correlation.

The size of tests based on A-HAC is rather promising. For each $\{n, T\}$ combination, there is very little variation in the rejection rates across the DGPs. For n = 5, rejection rates are approximately 3 times too high. For n = 30, however, tests have approximately the correct size: about 0.06 or 6%. There is no evidence of improvement with respect to an increase in *T*.

[Tables 3 and 4 Here]

One interesting observation is that for n = 30 and the AR(1) DGP, the size of A-HAC tests is closer to the nominal 5% level than for RE-AR1. In other words, even though RE-AR1 is fully consistent with the DGP, A-HAC is more accurate for this sample size. In comparison to RE-HAC, A-HAC rejection rates are closer to the nominal level for the serial correlation DGPs for n= 30, and for n = 5 with high degrees of serial correlation. RE-HAC rejection rates are closer to the nominal level for n = 5 for the no serial correlation DGP and the AR(1) DGP with a low or moderate degree of serial correlation.

5. Further Explorations

The results from the Monte Carlo experiment motivate some further explorations. First, the lack of size variation across DGPs for A-HAC, and its improvement for an increase in n, suggests that a degrees of freedom correction, based on n, is justified. In fact, Stata and Limdep both estimate \mathbf{V}_i using $\frac{n}{n-1}\widehat{\mathbf{V}}_i^{A-HAC}$. For n = 5, the size of A-HAC tests using this degrees of freedom adjustment improves from about 0.15 to 0.12. For n = 30 the improvement is from about 0.06 to 0.055. Thus, the degrees of freedom correction appears desirable although for small n the size distortion remains. This same degrees of freedom adjustment does not appear justified for RE-HAC, as size improves for an increase in T as well as n and under no serial correlation and no random effects the size is approximately correct for the sample sizes explored. Second, the superior performance of A-HAC over RE-HAC for n = 30 begs the question of whether it is desirable to set bandwidth equal to T rather than use a bandwidth selection method. To gain some insight, several RE-HAC simulations were conducted where bandwidth was simply set equal to T. The result was that RE-HAC rejection rates are approximately equal to A-HAC. For example, with n = 5, T = 50, r = .6 and $\rho_1 = 0.6$, the rejection rate is .143 for β_1 and .163 for β_2 . For the same sample size, but with r = .9, $\bar{\rho}_1 = 0.2$ and $\bar{\rho}_2 = 0.4$ the rates are .149 and .157, respectively. Similar results are obtained for the other three {n, T} combinations.

Third, given the relative performance of RE-HAC over A-HAC with n = 5 and low to moderate AR(1) serial correlation, I explored the effect of using a prewhitening filter. Using the VAR(1) prewhitening procedure proposed by Andrews and Monahan (1992), I find that test statistics are much closer to the correct size for T = 20 and T = 50. In particular, even for $\rho_1 =$ 0.9, the rejection rates for the AR(1) DGP approximate those from the analogous no serial correlation DGP. In other words, there is no additional distortion from the serial correlation and what is left is the distortion due to the random effect (which is also present for the RE and RE-AR1 estimators). One caveat, however, is that the prewhitening helps very little for the case of AR(2) serial correlation. What is happening is that the filter essentially removes all the firstorder serial correlation and the bandwidth selection procedure, based on an AR(1) serial correlation model, leads to a bandwidth choice that is too small. The small bandwidth fails to capture the second-order serial correlation. When a higher-order process is suspected, the bandwidth selection procedure of Newey and West (1994) is likely preferable, as it is not based solely on an AR(1) process.

6. Recommendations

Experimentalists analyzing panel data, like any analysts, should initially examine their data to discern important properties. One can look at the time-series properties of the data by usual time-series methods. For example, plotting the data against time can be used to examine for trends. And, one can gain insight as to the type of autoregressive and/or moving average process at play by examining the partial autocorrelation function and the autocorrelation function. Wooldridge (2002) proposes a test of AR(1) serial correlation for panel data, which requires minimal assumptions.¹² There are a number of proposed approaches for unit-root testing, and a review of this literature is provided by Baltagi and Kao (2000). If serial correlation is a concern, which is likely for data from repeated-game experiments, this study provides some recommendations on how to proceed using HAC covariance estimators for panel data.

So what is the bottom line? When there is a moderate (or large) number of cross-section units per treatment, which is normal for experiments when data are at the participant-level, the Monte Carlo results suggest that A-HAC (a.k.a. the "cluster-robust" covariance estimator) or the covariance estimator proposed in this study (RE-HAC) with bandwidth equal to *T* are desirable covariance estimators for OLS when there are unobserved unit effects and/or serial correlation of unknown form.^{13,14} Hypothesis tests based these HAC covariance estimators have approximately the correct size. As such, the HAC covariance estimators are as accurate as the RE or RE-AR1 estimator, even when one of the latter estimators are fully consistent with the DGP. When the structure of the serial correlation is misspecified, RE-AR1 or related estimators will lead to biased tests, and A-HAC will be preferred in such instances. Evidence from previous Monte Carlo studies (Bertrand, Duflo, and Mullainathan 2004; Kezdi 2004) provides additional support for A-HAC. On the other hand, if the number of cross-section units per treatment is small, which is more likely when data are at the group-level, such as a case where the experimentalist wishes to analyze measures of market or social efficiency, recommendations are less clear. A-HAC (or RE-HAC with bandwidth equal to T) standard errors tend to be too small. RE-HAC in tandem with a prewhitening filter and data-dependent bandwidth selection appears to have promise, but additional research is warranted. Certainly an analyst who is uncertain about the underlying DGP should look to the RE-HAC estimator, rather than assume a particular structure for the serial correlation as there may be greater size distortion due to misspecification. And it is noted that estimators like RE-AR1 that place structure on the serial correlation, even if approximately correctly specified, produce biased test statistics in small samples (see, for example, Table 1).

On a final note, the HAC covariance estimators investigated are generalizations of the oft-used White's (1980) heteroskedasticity-consistent covariance estimator. As such, although the Monte Carlo simulations here do not consider DGPs with conditional heteroskedasticity, A-HAC and RE-HAC are likewise robust to conditional heteroskedasticity of unknown form. In fact, conditional heteroskedasticity is unlikely to cause any additional size distortions. Evidence in support of these claims for A-HAC can be found in Bertrand, Duflo, and Mullainathan (2004) and Kezdi (2004).

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	RE-HAC		A-H	IAC	NW-HAC RE-AR1		RE		OLS			
	β ₁	β_2	β_1	β_2	β_1	β_2	β_1	β_2	β ₁	β_2	β_1	β_2
No serial correlation												
<i>r</i> =.0	.041	.065	.136	.155	.057	.051	.045	.072	.035	.047	.046	.046
<i>r</i> =.3	.109	.075	.151	.160	.252	.013	.099	.068	.097	.053	.320	.026
<i>r</i> =.6	.124	.075	.143	.160	.265	.004	.118	.068	.122	.053	.453	.008
<i>r</i> =.9	.133	.075	.142	.160	.192	.000	.135	.068	.133	.053	.561	.001
AR(1) serial correlation, common across units												
<i>r</i> =.0, ρ_1 =.3	.078	.114	.135	.146	.101	.090	.055	.086	.085	.144	.128	.132
<i>r</i> =.3, ρ_1 =.3	.112	.120	.153	.147	.231	.038	.095	.079	.111	.138	.341	.089
<i>r</i> =.6, ρ_1 =.3	.120	.120	.146	.147	.242	.009	.111	.078	.119	.138	.450	.043
<i>r</i> =.9, ρ_1 =.3	.130	.120	.142	.147	.189	.001	.130	.078	.131	.138	.560	.007
<i>r</i> =.0, ρ_1 =.6	.100	.179	.136	.153	.151	.117	.062	.104	.150	.299	.270	.261
<i>r</i> =.3, ρ_1 =.6	.104	.192	.135	.152	.199	.076	.084	.101	.121	.293	.387	.217
<i>r</i> =.6, ρ_1 =.6	.119	.192	.144	.152	.219	.032	.105	.101	.137	.293	.446	.143
<i>r</i> =.9, ρ_1 =.6	.126	.192	.134	.152	.192	.004	.118	.101	.130	.293	.560	.027
<i>r</i> =.0, ρ_1 =.9	.080	.261	.135	.175	.176	.103	.096	.135	.107	.492	.415	.351
<i>r</i> =.3, ρ_1 =.9	.076	.253	.142	.159	.196	.083	.090	.112	.115	.483	.444	.329
<i>r</i> =.6, ρ_1 =.9	086	.253	.134	.159	.184	.056	.096	.112	.107	.483	.457	.277
<i>r</i> =.9, ρ_1 =.9	.116	.253	.142	,159	.188	.016	.126	.112	.131	.483	.528	.141
		A	R(2) set	rial corre	elation, c	common	across t	inits				
<i>r</i> =.0, ρ_1 =.4, ρ_2 =.2	.115	.213	.136	.152	.179	.140	.099	.146	.137	.296	.269	.264
<i>r</i> =.3, ρ_1 =.4, ρ_2 =.2	.109	.218	.134	.150	.232	.079	.097	.150	.123	.287	.391	.202
<i>r</i> =.6, ρ_1 =.4, ρ_2 =.2	.128	.218	.146	.150	.240	.032	.115	.150	.134	.287	.453	.131
<i>r</i> =.9, ρ_1 =.4, ρ_2 =.2	.125	.218	.135	.150	.195	.003	.125	.150	.128	.287	.562	.022
<i>r</i> =.0, ρ_1 =.2, ρ_2 =.4	.125	.232	.135	.153	.198	.158	.120	.206	.130	.275	.251	.231
<i>r</i> =.3, ρ_1 =.2, ρ_2 =.4	.121	.231	.137	.151	.253	.083	.109	.217	.124	.268	.385	.173
<i>r</i> =.6, ρ_1 =.2, ρ_2 =.4	.130	.231	.144	.151	.255	.032	.123	.217	.132	.268	.455	.107
<i>r</i> =.9, ρ_1 =.2, ρ_2 =.4	.128	.231	.133	.151	.197	.002	.125	.217	.129	.268	.560	.018
	r	AR((2) serial	correlat	tion, hete	erogeneo	ous acros	ss units	r			
$r=.0, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.089	.216	.121	.150	.172	.135	.075	.155	.112	.293	.274	.246
$r=.3, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.110	.215	.145	.143	.229	.069	.106	.166	.121	.295	.389	.203
$r=.6, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.135	.215	.146	.143	.241	.029	.128	.166	.145	.295	.485	.128
$r=.9, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.156	.215	.163	.143	.224	.002	.161	.166	.161	.295	.576	.024
$r=.0, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.098	.239	.121	.150	.197	.145	.091	.220	.108	.264	.261	.223
$r=.3, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.113	.220	.146	.143	.247	.066	.118	.220	.117	.261	.380	.181
$r=.6, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.138	.220	.152	.143	.255	.026	.133	.220	.142	.261	.485	.104
$r=.9, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.158	.220	.161	.143	.232	.002	.161	.220	.158	.261	.574	.015

Table 1. Monte Carlo Results: Null Rejection Probabilities for n = 5, T = 20

	RE-I	HAC	A-H	IAC	NW-	HAC	RE-	AR1	R	E	O	LS
	β_1	β_2	β_1	β_2	β_1	β ₂	β_1	β_2	β_1	β ₂	β_1	β_2
No serial correlation												
<i>r</i> =.0	.044	.060	.062	.065	.051	.045	.051	.064	.033	.049	.050	.048
<i>r</i> =.3	.062	.057	.064	.054	.195	.002	.060	.056	.058	.040	.319	.013
<i>r</i> =.6	.067	.057	.066	.054	.151	.000	.073	.056	.066	.040	.435	.002
<i>r</i> =.9	.071	.057	.072	.054	.102	.000	.072	.056	.071	.040	.522	.000
AR(1) serial correlation, common across units												
<i>r</i> =.0, ρ_1 =.3	.063	.089	.055	.052	.086	.061	.034	.053	.074	.112	.131	.103
<i>r</i> =.3, ρ_1 =.3	.070	.089	.068	.059	.178	.012	.066	.052	.072	.120	.347	.058
<i>r</i> =.6, ρ_1 =.3	.076	.089	.071	.059	.149	.000	.069	.052	.078	.120	.472	.020
<i>r</i> =.9, ρ_1 =.3	.068	.089	.072	.059	.110	.000	.069	.052	.069	.120	.548	.000
<i>r</i> =.0, ρ_1 =.6	.056	.146	.053	.053	.114	.077	.053	.080	.100	.271	.257	.239
<i>r</i> =.3, ρ_1 =.6	.066	.142	.062	.053	.132	.024	.061	.078	.089	.270	.388	.175
<i>r</i> =.6, ρ_1 =.6	.071	.142	.065	.053	.121	.005	.069	.078	.077	.270	.460	.090
<i>r</i> =.9, ρ_1 =.6	.073	.142	.069	.053	.113	.000	.071	.078	.076	.270	.541	.004
<i>r</i> =.0, ρ_1 =.9	.036	.154	.068	.057	.115	.035	.064	.083	.054	.439	.396	.291
<i>r</i> =.3, ρ_1 =.9	.040	.162	.065	.050	.107	.031	.071	.088	.059	.446	.395	.261
<i>r</i> =.6, ρ_1 =.9	.039	.162	.062	.050	.111	.017	.070	.088	.053	.446	.439	.209
<i>r</i> =.9, ρ_1 =.9	.051	.162	.060	.050	.095	.001	.058	.088	.058	.446	.497	.057
		A	AR(2) set	rial corre	elation, o	common	across u	inits				
<i>r</i> =.0, ρ_1 =.4, ρ_2 =.2	.060	.168	.050	.054	.139	.087	.069	.113	.094	.267	.267	.230
<i>r</i> =.3, ρ_1 =.4, ρ_2 =.2	.070	.160	.062	.051	.156	.026	.070	.111	.083	.262	.397	.156
<i>r</i> =.6, ρ_1 =.4, ρ_2 =.2	.070	.160	.064	.051	.139	.003	.071	.111	.077	.262	.470	.081
<i>r</i> =.9, ρ_1 =.4, ρ_2 =.2	.074	.160	.067	.051	.116	.000	.072	.111	.077	.262	.550	.002
<i>r</i> =.0, ρ_1 =.2, ρ_2 =.4	.069	.197	.053	.056	.156	.102	.072	.176	.080	.236	.249	.202
<i>r</i> =.3, ρ_1 =.2, ρ_2 =.4	.072	.189	.065	.054	.174	.026	.075	.168	.081	.230	.391	.139
<i>r</i> =.6, ρ_1 =.2, ρ_2 =.4	.073	.189	.065	.054	.156	.003	.075	.168	.074	.230	.468	.065
<i>r</i> =.9, ρ_1 =.2, ρ_2 =.4	.074	.189	.068	.054	.118	.000	.072	.168	.074	.230	.547	.000
	-	AR((2) serial	correlat	tion, het	erogeneo	ous acros	ss units				
$r=.0, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.051	.186	.062	.075	.134	.095	.055	.140	.069	.294	.269	.230
$r=.3, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.060	.195	.071	.078	.167	.044	.059	.142	.074	.312	.380	.201
$r=.6, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.066	.195	.068	.078	.142	.007	.069	.142	.070	.312	.459	.120
$r=.9, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.065	.195	.069	.078	.105	.000	.067	.142	.067	.312	.537	.007
$r=.0, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.057	.211	.062	.073	.145	.103	.061	.195	.066	.260	.260	.211
$r=.3, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.064	.217	.072	.081	.189	.041	.063	.216	.073	.279	.370	.176
$r=.6, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.064	.217	.068	.081	.158	.004	.067	.216	.066	.279	.460	.094
$r=.9, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.065	.217	.070	.081	.106	.000	.066	.216	.066	.279	.530	.004

Table 2. Monte Carlo Results: Null Rejection Probabilities for n = 30, T = 20

	RE-I	HAC	A-H	IAC	NW-	HAC	RE-	AR1	R	E	O	LS
	β_1	β_2	β_1	β_2	β_1	β_2	β_1	β_2	β_1	β_2	β_1	β_2
				No s	erial cor	relation						
<i>r</i> =.0	.042	.056	.155	.167	.059	.049	.042	.062	.037	.052	.055	.049
<i>r</i> =.3	.142	.057	.146	.165	.375	.007	.138	.059	.141	.050	.501	.025
<i>r</i> =.6	.144	.057	.143	.165	.330	.002	.146	.059	.144	.050	.637	.006
<i>r</i> =.9	.134	.057	.136	.165	.251	.000	.133	.059	.134	.050	.723	.001
AR(1) serial correlation, common across units												
<i>r</i> =.0, ρ_1 =.3	.066	.115	.156	.171	.101	.099	.044	.069	.082	.155	.152	.150
<i>r</i> =.3, ρ_1 =.3	.138	.109	.151	.164	.298	.021	.135	.065	.139	.154	.511	.104
<i>r</i> =.6, ρ_1 =.3	.142	.109	.144	.164	.297	.006	.146	.065	.146	.154	.628	.047
<i>r</i> =.9, ρ_1 =.3	.139	.109	.138	.164	.229	.001	.141	.065	.140	.154	.714	.003
<i>r</i> =.0, ρ_1 =.6	.083	.154	.151	.172	.135	.123	.050	.088	.159	.337	.318	.324
<i>r</i> =.3, ρ_1 =.6	.120	.156	.139	.163	.210	.049	.111	.079	.151	.307	.510	.251
<i>r</i> =.6, ρ_1 =.6	.138	.156	.150	.163	.244	.013	.139	.079	.146	.307	.609	.184
<i>r</i> =.9, ρ_1 =.6	.145	.156	.144	.163	.215	.001	.150	.079	.149	.307	.701	.046
<i>r</i> =.0, ρ_1 =.9	.084	.228	.148	.150	.174	.121	.081	.105	.183	.599	.565	.530
<i>r</i> =.3, ρ_1 =.9	.100	.232	.145	.158	.180	.104	.095	.105	.166	.585	.581	.506
<i>r</i> =.6, ρ_1 =.9	.115	.232	.146	.158	.188	.068	.113	.105	.157	.585	.619	.474
<i>r</i> =.9, ρ_1 =.9	.137	.232	.145	.158	.210	.018	.143	.105	.155	.585	.684	.333
		A	R(2) set	rial corre	elation, c	common	across u	inits				
<i>r</i> =.0, ρ_1 =.4, ρ_2 =.2	.103	.192	.148	.170	.179	.153	.085	.144	.168	.348	.339	.333
<i>r</i> =.3, ρ_1 =.4, ρ_2 =.2	.125	.186	.140	.161	.312	.101	.130	.136	.152	.326	.529	.263
<i>r</i> =.6, ρ_1 =.4, ρ_2 =.2	.141	.186	.151	.161	.408	.041	.143	.135	.149	.326	.617	.182
<i>r</i> =.9, ρ_1 =.4, ρ_2 =.2	.146	.186	.145	.161	.478	.003	.149	.135	.151	.326	.703	.043
<i>r</i> =.0, ρ_1 =.2, ρ_2 =.4	.121	.232	.146	.164	.231	.202	.118	.227	.163	.343	.344	.327
<i>r</i> =.3, ρ_1 =.2, ρ_2 =.4	.136	.228	.140	.162	.389	.137	.137	.224	.151	.326	.533	.256
<i>r</i> =.6, ρ_1 =.2, ρ_2 =.4	.145	.228	.151	.162	.487	.067	.147	.224	.148	.326	.622	.175
<i>r</i> =.9, ρ_1 =.2, ρ_2 =.4	.147	.228	.145	.162	.551	.006	.149	.224	.149	.326	.703	.042
	1	AR((2) serial	correlat	tion, hete	erogeneo	ous acros	ss units	1	1		
$r=.0, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.091	.192	.157	.150	.181	.136	.084	.167	.174	.399	.405	.380
$r=.3, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.133	.198	.160	.140	.325	.096	.142	.168	.164	.400	.556	.336
$r=.6, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.142	.198	.154	.140	.420	.049	.143	.168	.156	.400	.673	.255
$r=.9, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.149	.198	.150	.140	.475	.008	.155	.168	.153	.400	.742	.074
$r=.0, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.111	.244	.156	.150	.232	.168	.131	.251	.167	.397	.399	.373
$r=.3, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.141	.248	.160	.140	.398	.142	.147	.252	.160	.389	.554	.329
$r=.6, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.148	.248	.155	.140	.490	.065	.150	.252	.157	.389	.670	.243
$r=.9, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.149	.248	.151	.140	.549	.010	.152	.252	.152	.389	.740	.064

Table 3. Monte Carlo Results: Null Rejection Probabilities for n = 5, T = 50

	RE-	HAC	A-H	IAC	NW-	HAC	RE-	AR1	RE		OLS	
	β_1	β_2	β_1	β_2	β_1	β_2	β_1	β_2	β_1	β_2	β_1	β_2
				No s	erial cor	relation						
<i>r</i> =.0	.032	.057	.057	.069	.050	.052	.037	.062	.031	.055	.048	.055
<i>r</i> =.3	.057	.057	.057	.064	.282	.002	.058	.060	.056	.056	.499	.023
<i>r</i> =.6	.061	.057	.062	.064	.213	.000	.062	.060	.061	.056	.633	.006
<i>r</i> =.9	.063	.057	.064	.064	.124	.000	.063	.060	.063	.056	.696	.000
AR(1) serial correlation, common across units												
<i>r</i> =.0, ρ_1 =.3	.061	.105	.056	.066	.092	.087	.037	.072	.084	.159	.139	.156
<i>r</i> =.3, ρ_1 =.3	.058	.095	.059	.064	.210	.010	.057	.063	.062	.140	.489	.091
<i>r</i> =.6, ρ_1 =.3	.061	.095	.062	.064	.182	.000	.060	.063	.062	.140	.619	.027
<i>r</i> =.9, ρ_1 =.3	.063	.095	.063	.064	.124	.000	.059	.063	.063	.140	.696	.000
<i>r</i> =.0, ρ_1 =.6	.058	.122	.061	.068	.103	.095	.043	.084	.128	.319	.316	.308
<i>r</i> =.3, ρ_1 =.6	.053	.124	.055	.060	.150	.019	.056	.072	.074	.314	.498	.250
<i>r</i> =.6, ρ_1 =.6	.055	.124	.059	.060	.144	.001	.056	.072	.063	.314	.598	.156
<i>r</i> =.9, ρ_1 =.6	.063	.124	.062	.060	.102	.000	.063	.072	.064	.314	.685	.013
<i>r</i> =.0, ρ_1 =.9	.048	.163	.067	.060	.099	.077	.056	.072	.100	.579	.565	.502
<i>r</i> =.3, ρ_1 =.9	.046	.173	.060	.064	.123	.069	.058	.079	.082	.600	.556	.499
<i>r</i> =.6, ρ_1 =.9	.054	.173	.061	.064	.117	.034	.055	.079	.076	.600	.581	.456
<i>r</i> =.9, ρ_1 =.9	.051	.173	.058	.064	.097	.002	.058	.079	.063	.600	.652	.289
	r	A	R(2) set	rial corre	elation, c	common	across u	inits				
<i>r</i> =.0, ρ_1 =.4, ρ_2 =.2	.064	.156	.060	.069	.133	.121	.071	.137	.131	.339	.326	.320
<i>r</i> =.3, ρ_1 =.4, ρ_2 =.2	.059	.153	.055	.060	.251	.060	.059	.128	.073	.333	.510	.256
<i>r</i> =.6, ρ_1 =.4, ρ_2 =.2	.059	.153	.060	.060	.342	.013	.058	.128	.062	.333	.609	.157
<i>r</i> =.9, ρ_1 =.4, ρ_2 =.2	.064	.153	.063	.060	.389	.000	.065	.128	.064	.333	.688	.013
<i>r</i> =.0, ρ_1 =.2, ρ_2 =.4	.085	.203	.060	.067	.194	.163	.095	.204	.125	.339	.336	.324
<i>r</i> =.3, ρ_1 =.2, ρ_2 =.4	.061	.196	.057	.059	.344	.105	.067	.197	.072	.338	.512	.246
<i>r</i> =.6, ρ_1 =.2, ρ_2 =.4	.061	.196	.060	.059	.426	.024	.058	.197	.062	.338	.609	.151
<i>r</i> =.9, ρ_1 =.2, ρ_2 =.4	.064	.196	.064	.059	.483	.000	.065	.197	.064	.338	.688	.012
	1	AR((2) serial	correlat	tion, het	erogeneo	ous acros	ss units				
$r=.0, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.050	.162	.044	.070	.122	.100	.064	.160	.099	.386	.395	.357
$r=.3, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.061	.157	.058	.055	.223	.063	.063	.149	.075	.403	.524	.309
$r=.6, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.057	.157	.056	.055	.301	.018	.060	.149	.065	.403	.604	.220
$r=.9, \overline{\rho}_1=.4, \overline{\rho}_2=.2$.055	.157	.057	.055	.355	.000	.056	.149	.057	.403	.689	.033
$r=.0, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.064	.200	.047	.071	.163	.133	.082	.241	.097	.373	.382	.351
$r=.3, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.065	.187	.058	.057	.293	.091	.068	.219	.071	.392	.522	.286
$r=.6, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.061	.187	.057	.057	.378	.030	.064	.219	.066	.392	.607	.197
$r=.9, \overline{\rho}_1=.2, \overline{\rho}_2=.4$.056	.187	.057	.057	.426	.000	.056	.219	.057	.392	.689	.025

Table 4. Monte Carlo Results: Null Rejection Probabilities for n = 30, T = 50

¹ Note that the idea for this paper originated in 2002, when panel data analysis was the exception rather than the rule for drawing inferences from experimental data. Although panel data models are increasing used for analyzing repeated-game experiment data, and indeed more and more researchers have relied on using cluster-robust standard errors, it remains common for journal referees to request simple statistical tests even when their validity is questionable.

² Indeed, on numerous occasions I have reviewed papers that simply mention heteroskedasticity when justifying the use of cluster-robust standard errors. Further, it is fairly common for some to use the standard random effects estimator in tandem with cluster-robust standard errors. This approach is internally inconsistent, as the random effects estimator assumes a specific form of within-unit serial correlation but the use of cluster-robust standard errors suggests that the assumed form of serial correlation is incorrect.

³ Data sets of this sort tend to be labeled as "time-series cross-section" or TSCS data.

⁴ A prewhitening filter attempts to remove some correlation in the residuals from a regression model, which has been shown to improve the performance of HAC-based techniques (see Andrews and Monahan 1992).

⁵ Stata's *newey* command (with option *force*) produces standard OLS coefficients (without any adjustment for a fixed or random-effects structure) along with the NW-HAC estimator. Limdep estimates an equivalent NW-HAC estimator, but with a fixed-effects estimator for model coefficients.

⁶ This covariance estimator is produced when one specifies the *cluster* option for Stata or Limdep's *regress* command. To estimate a fixed-effects model with A-HAC errors in Stata, one can jointly use the *cluster* and *fe* options for Stata's *xtreg* command.

⁷ For purpose of identification, the HAC covariance estimators considered here require that $n \ge k$.

⁸ Of course, this estimator would be inefficient, and the standard FGLS random effects estimator for β is preferable. ⁹ As suggested by Andrews (1991), to calculate the bandwidths I use a weight of 0 for the autoregressive parameter associated with the model intercept and a weight of 1 on other autoregressive parameters. See Andrews (1991) for details on using this procedure.

¹⁰ OLS and random effects models are estimated using canned procedures in Limdep. RE-AR1 uses an estimate of ρ from a fixed effects model. To construct the RE-HAC covariance estimator I use the estimate of σ_u^2 generated by Limdep's random effects estimator.

¹¹ These statistics are available upon request.

¹² There is a user-written program available to run this test in Stata (Drukker, 2003).

¹⁴ If a fixed-effects structure is preferred or assumed, then a fixed effects coefficient estimator with A-HAC is recommended.

¹³ Since there are canned procedures in Stata and Limdep for A-HAC (with the degrees of freedom correction discussed above), it is likely preferable from the practitioner's viewpoint.