#### A CONDITIONAL-HETEROSKEDASTICITY-ROBUST CONFIDENCE INTERVAL FOR THE AUTOREGRESSIVE PARAMETER

By

#### **Donald W.K. Andrews and Patrik Guggenberger**

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# A Conditional-Heteroskedasticity-Robust Confidence Interval for the Autoregressive Parameter

Donald W. K. Andrews<sup>\*</sup> Cowles Foundation for Research in Economics Yale University

> Patrik Guggenberger Department of Economics UCSD

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#### Abstract

This paper introduces a new confidence interval (CI) for the autoregressive parameter (AR) in an AR(1) model that allows for conditional heteroskedasticity of general form and AR parameters that are less than or equal to unity. The CI is a modification of Mikusheva's (2007a) modification of Stock's (1991) CI that employs the least squares estimator and a heteroskedasticity-robust variance estimator. The CI is shown to have correct asymptotic size and to be asymptotically similar (in a uniform sense). It does not require any tuning parameters. No existing procedures have these properties. Monte Carlo simulations show that the CI performs well in finite samples in terms of coverage probability and average length, for innovations with and without conditional heteroskedasticity.

*Keywords*: Asymptotically similar, asymptotic size, autoregressive model, conditional heteroskedasticity, confidence interval, hybrid test, subsampling test, unit root.

JEL Classification Numbers: C12, C15, C22.

#### 1 Introduction

We consider confidence intervals (CI's) for the autoregressive parameter (AR)  $\rho$  in a conditionally heteroskedastic AR(1) model in which  $\rho$  may be close to, or equal to, one. The observed time series  $\{Y_i : i = 0, ..., n\}$  is based on a latent no-intercept AR(1) time series  $\{Y_i^* : i = 0, ..., n\}$ :

$$Y_{i} = \mu + Y_{i}^{*},$$
  

$$Y_{i}^{*} = \rho Y_{i-1}^{*} + U_{i} \text{ for } i = 1, ..., n,$$
(1.1)

where  $\rho \in [-1 + \varepsilon, 1]$  for some  $0 < \varepsilon < 2$ ,  $\{U_i : i = ..., 0, 1, ...\}$  are stationary and ergodic under the distribution F, with conditional mean 0 given a  $\sigma$ -field  $\mathcal{G}_{i-1}$  for which  $U_j \in \mathcal{G}_i$  for all  $j \leq i$ , conditional variance  $\sigma_i^2 = E_F(U_i^2|\mathcal{G}_{i-1})$ , and unconditional variance  $\sigma_U^2 \in (0, \infty)$ . The distribution of  $Y_0^*$  is the distribution that yields strict stationarity for  $\{Y_i^* : i \leq n\}$  when  $\rho < 1$ . That is,  $Y_0^* = \sum_{j=0}^{\infty} \rho^j U_{-j}$  when  $\rho < 1$ . When  $\rho = 1$ ,  $Y_0^*$  is arbitrary.

Models of this sort are applicable to exchange rate and commodity and stock prices, e.g., see Kim and Schmidt (1993). Simulations in Mikusheva (2007b, Table II) show that CI's not designed to handle conditional heteroskedasticity may perform poorly in terms of coverage probabilities when conditional heteroskedasticity is present. In fact, most have incorrect asymptotic size in this case.<sup>1</sup>

For the case of conditional homoskedasticity, several CI's with correct asymptotic size have been introduced, including those in Stock (1991), Andrews (1993), Andrews and Chen (1994), Nankervitz and Savin (1996), Hansen (1999), Elliot and Stock (2001), Romano and Wolf (2001), Chen and Deo (2007), and Mikusheva (2007a).<sup>2</sup> Of these CI's the only one that has correct asymptotic size in the presence of conditional heteroskedasticity is the symmetric two-sided subsampling CI of Romano and Wolf (2001).<sup>3</sup> The latter CI has the disadvantages that it is not asymptotically similar, requires a tuning parameter (the subsample size), and is far from being equal-tailed when  $\rho$  is near one.<sup>4</sup>

The first CI's that were shown to have correct asymptotic size under conditional heteroskedasticity and an AR parameter that may be close to, or equal to, unity were introduced in Andrews and Guggenberger (2009) (AG09).<sup>5</sup> These CI's are based on inverting a t statistic constructed using a feasible quasi-generalized least squares (FQGLS) estimator of  $\rho$ . AG09 shows that equal-tailed and symmetric two-sided CI's based on hybrid (fixed/subsampling) critical values have correct asymptotic size.<sup>6</sup> These CI's are robust to misspecification of the form of the conditional heteroskedasticity. However, they are not asymptotically similar and require the specification of a tuning parameter—the subsample size.

The contribution of this paper is to introduce a CI that (i) has correct asymptotic size for a parameter space that allows for general forms of conditional heteroskedasticity and for an AR parameter close to, or equal to, unity, (ii) is asymptotically similar, and (iii) does not require any tuning parameters.

The CI is constructed by inverting tests constructed using a t statistic based on the LS estimator of  $\rho$  and a heteroskedasticity-consistent (HC) variance matrix estimator. For the latter, we use a variant of the HC3 version defined in MacKinnon and White (1985), which we call HC5. It employs an adjustment that improves the finite-sample coverage probabilities. This t statistic is asymptotically nuisance parameter-free under the null hypothesis under drifting sequences of null parameters  $\rho$ , whether or not these parameters are local to unity. In consequence, critical values can be obtained by matching the given null value of  $\rho$  and sample size n with a local-tounity parameter  $h = n(1 - \rho)$ . Then, one uses the quantile(s) from the corresponding local-to-unity asymptotic distribution which depends on h. This method is employed by Stock (1991), Andrews and Chen (1994, Sec. 4), and Mikusheva (2007a) (in her modification of Stock's CI).<sup>7</sup> The resulting CI is the same as Mikusheva's (2007a) modification of Stock's (1991) CI applied to the LS estimator of  $\rho$ , except that we use the HC5 variance estimator in place of the homoskedastic variance estimator and we use a stationary initial condition rather than a zero initial condition.<sup>8,9</sup> We refer to the new CI as the CHR CI (which abbreviates "conditional-heteroskedasticity-robust").

The use of the LS estimator, rather than the FQGLS estimator, is important because the latter has an asymptotic distribution in the local-to-unity case that is a convex combination of a random variable with a unit-root distribution and an independent standard normal random variable with coefficients that depend on the strength of the conditional heteroskedasticity, see Seo (1999), Guo and Phillips (2001), and Andrews and Guggenberger (2012). Hence, a nuisance parameter appears in the asymptotic distribution of the FQGLS estimator that does not appear with the LS estimator. This yields a trade-off when constructing a CI between using a more efficient estimator (FQGLS) combined with critical values that do not lead to an asymptotically similar CI and using a less efficient estimator (LS) with critical values that yield an asymptotically similar CI.

The use of an HC variance matrix estimator with the new CHR CI is important to obtain a (nuisance-parameter free) standard normal asymptotic distribution of the t statistic when the sequence of true  $\rho$  parameters converges to a value less than one as  $n \to \infty$  and conditional heteroskedasticity is present. It is not needed to yield a nuisance parameter-free asymptotic distribution when  $\rho$  converges to unity (either at a  $O(n^{-1})$  rate or more slowly).<sup>10</sup> This follows from results in Seo (1999). Also see Guo and Phillips (2001), Cavaliere and Taylor (2009), and Andrews and Guggenberger (2012).

Simulations indicate that the CHR CI has good finite-sample coverage probabilities and has shorter average lengths—often noticeably shorter—than the hybrid CI of AG09 (based on the FQGLS estimator) for a variety of GARCH(1,1) processes whose parametrizations are empirically relevant. When no conditional heterskedasticity is present, the CHR CI performs very well in finite samples relative to CI's that are designed for the i.i.d. innovation case. Hence, there is little cost to achieving robustness to conditional heteroskedasticity. The asymptotic size and similarity results for the new CI are obtained by employing the asymptotic results of Andrews and Guggenberger (2012) for FQGLS estimators under drifting sequence of distributions, which include LS estimators as a special case, combined with the generic uniformity results in Andrews, Cheng, and Guggenberger (2009).

The CHR CI yields a unit root test that is robust to conditional heteroskedasticity. One rejects a unit root if the CI does not include unity. Seo (1999), Guo and Phillips (2001), and Cavaliere and Taylor (2009) also provide unit root tests that are robust to conditional heteroskedasticity.

The CHR CI for  $\rho$  can be extended to give a CI for the sum of the AR coefficients in an AR(k) model when all but one root is bounded away from the unit circle, e.g., as in Andrews and Chen (1994, Sec. 4) and Mikusheva (2007a). In this case, the asymptotic distributions (and hence the CHR critical values) are unchanged. See the end of Section 2 for more details. The CHR CI for  $\rho$  also can be extended to models with a linear time trend.<sup>11</sup> In this case, the asymptotic distributions are given in (7.7) of Andrews and Guggenberger (2009) with  $h_{2,7} = 1$ . Extending the proof of Theorem 1 below for these two cases requires additional detailed analysis, e.g., as in Mikusheva (2007a, Sec. 7). For brevity, we do not provide such proofs here.

The paper is structured as follows. Section 2 defines the new CI and establishes its large sample properties. Section 3 provides tables of critical values. Section 4 contains a Monte Carlo study. An Appendix provides: (i) the local asymptotic false coverage probabilities of the CHR CI, (ii) asymptotic and finite-sample assessments of the price the CHR CI pays in the i.i.d. case for obtaining robustness to conditional heteroskedasticity, (iii) probabilities of obtaining disconnected CHR CI's, (iv) definitions, tables of critical values, and simulation results for symmetric two-sided CHR CI's, (v) details concerning the simulations, (vi) description of a recursive residualbased wild bootstrap version of the CHR CI, (vii) proofs of the asymptotic results for the CHR CI, and (viii) a proof that the symmetric two-sided subsampling CI of Romano and Wolf (2001) has correct asymptotic size under conditional heteroskedasticity.

#### 2 The CHR CI for the AR parameter

For the exposition of the theory, we focus on equal-tailed two-sided CI's for  $\rho$ .<sup>12,13</sup> The CI is obtained by inverting a test of the null hypothesis that the true value is  $\rho$ . The model (1.1) can be rewritten as  $Y_i = \tilde{\mu} + \rho Y_{i-1} + U_i$ , where  $\tilde{\mu} = \mu(1-\rho)$  for i = 1, ..., n. We use the t statistic

$$T_n(\rho) = \frac{n^{1/2}(\widehat{\rho}_n - \rho)}{\widehat{\sigma}_n},$$
(2.1)

where  $\hat{\rho}_n$  is the LS estimator from the regression of  $Y_i$  on  $Y_{i-1}$  and 1 and  $\hat{\sigma}_n^2$  is the (1, 1) element of the HC5 heteroskedasticity-robust variance estimator, defined below,

for the LS estimator in the preceding regression. More explicitly, let  $Y, U, X_1$ , and  $X_2$  be *n*-vectors with *i*th elements given by  $Y_i, U_i, Y_{i-1}$ , and 1, respectively. Let  $X = [X_1 : X_2], P_X = X(X'X)^{-1}X'$ , and  $M_X = I_n - P_X$ . Let  $\hat{U}_i$  denote the *i*th element of the residual vector  $M_X Y$ . Let  $p_{ii}$  denote the *i*th diagonal element of  $P_X$ . Let  $p_{ii}^* = \min\{p_{ii}, n^{-1/2}\}$ . Let  $\Delta$  be the diagonal  $n \times n$  matrix with *i*th diagonal element given by  $\hat{U}_i/(1 - p_{ii}^*)$ .<sup>14</sup> Then, the LS estimator of  $\rho$  and the HC5 estimator  $\hat{\sigma}_n^2$  are

$$\widehat{\rho}_{n} = (X_{1}'M_{X_{2}}X_{1})^{-1}X_{1}'M_{X_{2}}Y, \text{ and}$$

$$\widehat{\sigma}_{n}^{2} = \left(n^{-1}X_{1}'M_{X_{2}}X_{1}\right)^{-1}\left(n^{-1}X_{1}'M_{X_{2}}\Delta^{2}M_{X_{2}}X_{1}\right)\left(n^{-1}X_{1}'M_{X_{2}}X_{1}\right)^{-1}.$$
(2.2)

Equivalently,  $\hat{\sigma}_n^2$  is the (1,1) element of  $n (X'X)^{-1} X' \Delta^2 X (X'X)^{-1}$ . The parameter space for  $(\rho, F)$  is given by

$$\Lambda = \{\lambda = (\rho, F) : \rho \in [-1 + \varepsilon, 1], \{U_i : i = 0, \pm 1, \pm 2, ...\} \text{ are stationary}$$
and strong mixing under  $F$  with  $E_F(U_i | \mathcal{G}_{i-1}) = 0$  a.s.,  $E_F(U_i^2 | \mathcal{G}_{i-1}) = \sigma_i^2$  a.s., where  $\mathcal{G}_i$  is some non-decreasing sequence of  $\sigma$ -fields for  $i = ..., 1, 2, ...$  for which  $U_j \in \mathcal{G}_i$  for all  $j \leq i$ , the strong-mixing numbers  $\{\alpha_F(m) : m \geq 1\}$  satisfy  $\alpha_F(m) \leq Cm^{-3\zeta/(\zeta-3)} \ \forall m \geq 1, \ \sup_{i,s,t,u,v,A} E_F |\prod_{a \in A} a|^{\zeta} \leq M, \ \text{where}$  $0 \leq i, s, t, u, v < \infty, \ \text{and } A \ \text{is any non-empty subset of} \ \{U_{i-s}, U_{i-t}, U_{i+1}^2, U_{-u}, U_{-v}, U_1^2\}, \ \text{and} \ E_F U_1^2 \geq \delta\},$ (2.3)

for some constants  $0 < \varepsilon < 2$ ,  $\zeta > 3$ ,  $C < \infty$ , and  $\delta > 0$ .

Next, we define the critical values used in the construction of the CI. They are based on the asymptotic distributions of the test statistic under drifting sequences  $\{\lambda_n = (\rho_n, F_n) : n \ge 1\}$  of AR parameters  $\rho_n$  and distributions  $F_n$ , when  $n(1 - \rho_n) \rightarrow h \in [0, \infty)$ . When  $F_n$  depends on n,  $\{U_i : i \le n\}$  for  $n \ge 1$  form a triangular array of random variables and  $U_i = U_{n,i}$ . To describe the asymptotic distribution, let  $W(\cdot)$  be a standard Brownian motion on [0, 1]. Let  $Z_1$  be a standard normal random variable that is independent of  $W(\cdot)$ . Define

$$I_{h}(r) = \int_{0}^{r} \exp(-(r-s)h) dW(s), \ I_{h}^{*}(r) = I_{h}(r) + \frac{1}{\sqrt{2h}} \exp(-hr) Z_{1} \text{ for } h > 0,$$
  

$$I_{h}^{*}(r) = W(r) \text{ for } h = 0, \text{ and } I_{D,h}^{*}(r) = I_{h}^{*}(r) - \int_{0}^{1} I_{h}^{*}(s) ds.$$
(2.4)

Andrews and Guggenberger (2012, Theorem 1) (with a minor adjustment for the  $p_{ii}^*$  term in  $\Delta$ ) shows that, under any sequence  $\lambda_n = (\rho_n, F_n) \in \Lambda$  such that  $n(1 - \rho_n) \to h \in [0, \infty],$ 

$$T_n(\rho_n) \to_d J_h, \tag{2.5}$$

where  $J_h$  is defined as follows. For  $h = \infty$ ,  $J_h$  is the N(0, 1) distribution, and for  $h \in [0, \infty)$ ,  $J_h$  is the distribution of

$$\int_{0}^{1} I_{D,h}^{*}(r) dW(r) / \left( \int_{0}^{1} I_{D,h}^{*}(r)^{2} dr \right)^{1/2}.$$
(2.6)

For  $\alpha \in (0, 1)$ , let  $c_h(1 - \alpha)$  denote the  $(1 - \alpha)$ -quantile of  $J_h$ . The second component of  $I_h^*(r)$  in (2.4) is due to the stationary start-up of the AR(1) process when  $\rho < 1$ , as in Elliott (1999), Elliott and Stock (2001), Müller and Elliott (2003), and Andrews and Guggenberger (2009, 2012). Giraitis and Phillips (2006) and Phillips and Magdalinos (2007) provide similar results for the LS estimator for the case  $h = \infty$ under assumptions that do not allow for conditional heteroskedasticity.

The new nominal  $1 - \alpha$  equal-tailed two-sided CHR CI for  $\rho$  is

$$CI_{CHR,n} = \{ \rho \in [-1 + \varepsilon, 1] : c_h(\alpha/2) \le T_n(\rho) \le c_h(1 - \alpha/2) \text{ for } h = n(1 - \rho) \}.$$
 (2.7)

The CI  $CI_{CHR,n}$  can be calculated by taking a fine grid of values  $\rho \in [-1 + \varepsilon, 1]$  and comparing  $T_n(\rho)$  to  $c_h(\alpha/2)$  and  $c_h(1 - \alpha/2)$ , where  $h = n(1 - \rho)$ . Tables of values of  $c_h(\alpha/2)$  and  $c_h(1 - \alpha/2)$  are given in Section 3 below. Given these values, calculation of  $CI_{CHR,n}$  is simple and fast.<sup>15</sup>

One could replace the asymptotic quantiles  $c_h(\alpha/2)$  and  $c_h(1 - \alpha/2)$  in (2.7) by recursive residual-based wild bootstrap quantiles and the CI would still have correct asymptotic size. (For brevity, we do not prove this claim.) The resulting CI is a grid bootstrap, as in Hansen (1999), but is designed to allow for conditional heteroskedasticity. Note that the bootstrap needs to be defined carefully. See the Appendix for its definition.<sup>16</sup> The bootstrap version of the CI is much less convenient computationally because one cannot use tables of critical values. Rather, one has to compute bootstrap critical values for each value of  $\rho$  to determine whether  $\rho$  is in the CI.

The main theoretical result of this paper shows that  $CI_{CHR,n}$  has correct asymptotic size for the parameter space  $\Lambda$  and is asymptotically similar. Let  $P_{\lambda}$  denote probability under  $\lambda = (\rho, F) \in \Lambda$ .

**Theorem 1** Let  $\alpha \in (0, 1)$ . For the parameter space  $\Lambda$ , the nominal  $1 - \alpha$  confidence interval  $CI_{CHR,n}$  for the AR parameter  $\rho$  satisfies

$$AsySz \equiv \liminf_{n \to \infty} \inf_{\lambda = (\rho, F) \in \Lambda} P_{\lambda}(\rho \in CI_{CHR, n}) = 1 - \alpha.$$

Furthermore,  $CI_{CHR,n}$  is asymptotically similar, that is,

$$\liminf_{n \to \infty} \inf_{\lambda = (\rho, F) \in \Lambda} P_{\lambda}(\rho \in CI_{CHR, n}) = \limsup_{n \to \infty} \sup_{\lambda = (\rho, F) \in \Lambda} P_{\lambda}(\rho \in CI_{CHR, n}).$$

Theorem 2 in the Appendix establishes the local asymptotic false coverage probabilities of the CHR CI, which are directly related to their length. As noted in the Introduction, the CHR CI for  $\rho$  can be extended to give a CI for the sum of the AR coefficients in an AR(k) model when all but one root is bounded away from the unit circle. The AR(k) model written in augmented Dickey-Fuller form is

$$Y_i = \mu + Y_i^*$$
 and  $Y_i^* = \rho Y_{i-1}^* + \sum_{j=1}^{k-1} \psi_j \delta Y_{i-j}^* + U_i$  for  $i = 1, ..., n,$  (2.8)

where  $\delta Y_{i-j}^* = Y_{i-j}^* - Y_{i-j-1}^*$ . Here  $\rho$  equals the sum of the k AR coefficients. For this model, the estimator  $\hat{\rho}_n$  of  $\rho$  that we consider is the LS estimator from the regression of  $Y_i$  on  $Y_{i-1}, \delta Y_{i-1}, ..., \delta Y_{i-k+1}$ , and 1, where  $\delta Y_{i-j} = Y_{i-j} - Y_{i-j-1}$ . The estimator  $\hat{\sigma}_n^2$  that we consider is the (1,1) element of  $n (X'X)^{-1} X' \Delta^2 X (X'X)^{-1}$ , where  $X = [X_1 : X_2 : \cdots : X_{k+1}], X_1, X_2, ..., X_k, X_{k+1}$  are the *n*-vectors with *i*th elements equal to  $Y_{i-1}, \delta Y_{i-1}, ..., \delta Y_{i-k+1}, 1$ , respectively, and  $\Delta$  is defined as in the paragraph containing (2.1) but with X defined as immediately above, rather than as in the paragraph containing (2.1). The CHR CI for  $\rho$  in (2.8) is defined exactly as the CHR CI for  $\rho$  in (1.1) is defined, but with the definitions of  $\hat{\rho}_n$  and  $\hat{\sigma}_n^2$  given immediately above rather than just below (2.1).

#### 3 Tables of Critical Values

Table 1 reports the quantiles  $c_h(.025)$  and  $c_h(.975)$  (for a broad range of values of h) which are used to calculate 95% equal-tailed CHR CI's. Table 2 reports analogous quantiles used to calculate 90% equal-tailed CHR CI's. These tables also can be used for 97.5% and 95% lower and upper one-sided CI's. Section 9 in the Appendix provides critical values for symmetric two-sided CHR CI's.

For given  $\alpha$ ,  $c_h(\alpha)$  (the  $\alpha$ -quantile of  $J_h$  in (2.5)) is computed by simulating the asymptotic distribution  $J_h$ . To do so, 300,000 independent AR(1) sequences are generated from the model in (1.1) with innovations  $U_i \sim iid N(0,1)$ ,  $\mu = 0$ , stationary startup, n = 25,000, and  $\rho_h = 1 - h/n$ . For each sequence, the test statistic  $T_n(\rho_h)$ (defined in (2.1) but using the homoskedastic variance estimator) is calculated. Then, the simulated estimate of  $c_h(\alpha)$  is the  $\alpha$ -quantile of the empirical distribution of the 300,000 realizations of the test statistic.

In Table 1, the critical values do not reach the  $h = \infty$  values of -1.96 and 1.96 for h = 500. Larger values of h, which would be needed only in very large samples, yield the following:  $c_{1,000}(.025) = -2.02$ ,  $c_{5,000}(.025) = -1.98$ ,  $c_{10,000}(.025) = -1.97$ ,  $c_{1,000}(.975) = 1.90$ ,  $c_{5,000}(.975) = 1.93$ , and  $c_{10,000}(.975) = 1.94$ .

h	0	.2	.4	.6	.8	1.0	1.4	1.8	2.2	2.6	3.0	3.4	3.8
$c_h$	-3.13	-3.09	-3.06	-3.03	-3.00	-2.98	-2.93	-2.89	-2.85	-2.83	-2.80	-2.77	-2.75
											13		
$c_h$	-2.73	-2.71	-2.69	-2.65	-2.62	-2.59	-2.56	-2.54	-2.52	-2.50	-2.48	-2.47	-2.45
											200		
$c_h$	-2.39	-2.35	-2.32	-2.28	-2.24	-2.23	-2.21	-2.19	-2.18	-2.17	-2.11	-2.08	-2.05

Table 1(a). Values of  $c_h(.025)$ , the .025 Quantile of  $J_h$ , for Use with 95% Equal-Tailed Two-Sided Confidence Intervals

Table 1(b). Values of  $c_h(.975)$ , the .975 Quantile of  $J_h$ , for Use with 95% Equal-Tailed Two-Sided Confidence Intervals

h	0	.2	.4	.6	.8	1.0	1.4	1.8	2.2	2.6	3.0	3.4	3.8
$c_h(.975)$	.24	.31	.36	.41	.45	.50	.57	.64	.69	.74	.79	.84	.88
h	4.2	4.6	5.0	6.0	7.0	8.0	9.0	10	11	12	13	14	15
$c_h(.975)$	.92	.95	.99	1.06	1.12	1.17	1.21	1.25	1.29	1.32	1.34	1.37	1.39
h	20	25	30	40	50	60	70	80	90	100	200	300	500
$c_h(.975)$	1.47	1.51	1.55	1.61	1.65	1.67	1.69	1.71	1.73	1.74	1.81	1.83	1.86

Table 2(a). Values of  $c_h(.05)$ , the .05 Quantile of  $J_h$ , for Use with 90% Equal-Tailed Two-Sided Confidence Intervals

h	0	.2	.4	.6	.8	1.0	1.4	1.8	2.2	2.6	3.0	3.4	3.8
$c_h$	-2.87	-2.83	-2.79	-2.76	-2.73	-2.70	-2.65	-2.61	-2.57	-2.54	-2.51	-2.48	-2.46
h	4.2	4.6	5.0	6.0	7.0	8.0	9.0	10	11	12	13	14	15
$c_h$	-2.44	-2.42	-2.39	-2.35	-2.32	-2.29	-2.26	-2.23	-2.21	-2.19	-2.18	-2.16	-2.14
h	20	25	30	40	50	60	70	80	90	100	200	300	500
$c_h$	-2.09	-2.05	-2.01	-1.97	-1.93	-1.91	-1.89	-1.87	-1.86	-1.85	-1.79	-1.76	-1.74

Table 2(b). Values of  $c_h(.95)$ , the .95 Quantile of  $J_h$ , for Use with 90% Equal-Tailed Two-Sided Confidence Intervals

h	0	.2	.4	.6	.8	1.0	1.4	1.8	2.2	2.6	3.0	3.4	3.8
$c_h(.95)$	07	02	.04	.08	.13	.17	.25	.31	.37	.43	.48	.52	.57
h	4.2	4.6	5.0	6.0	7.0	8.0	9.0	10	11	12	13	14	15
$c_h(.95)$	.61	.64	.68	.75	.81	.87	.91	.95	.98	1.01	1.03	1.05	1.08
h	20	25	30	40	50	60	70	80	90	100	200	300	500
$c_h(.95)$	1.15	1.20	1.24	1.30	1.34	1.36	1.39	1.40	1.42	1.43	1.49	1.52	1.55

#### 4 Finite-Sample Simulation Results

Here we compare the finite-sample coverage probabilities (CP's) and average lengths of the new CHR CI and the hybrid CI of AG09.<sup>17,18</sup> For brevity, we focus on nominal 95% equal-tailed two-sided CI's. Results for symmetric CI's, including the symmetric subsampling CI of Romano and Wolf (2001), are provided in the Appendix.

We consider a wide range of  $\rho$  values: .99, .9, .5, .0, -.9. The innovations are of the form  $U_i = \sigma_i \varepsilon_i$ , where  $\{\varepsilon_i : i \ge 1\}$  are i.i.d. standard normal and  $\sigma_i$  is the multiplicative conditional heteroskedasticity. Let GARCH- $(ma, ar; \psi)$  denote a GARCH(1, 1) process with MA, AR, and intercept parameters  $(ma, ar; \psi)$  and let ARCH- $(ar_1, ..., ar_4; \psi)$  denote an ARCH(4) process with AR parameters  $(ar_1, ..., ar_4)$ and intercept  $\psi$ . We consider five specifications for the conditional heteroskedasticity of the innovations: (i) GARCH-(.05, .9; .001), (ii) GARCH-(.15, .8; .2), (iii) i.i.d., (iv) GARCH-(.25, .7; .2), and (v) ARCH-(.3, .2, .2, .2; .2). Specifications (i)-(iii) are the most relevant ones empirically.<sup>19</sup> Specifications (iv) and (v) are included for purposes of robustness. They exhibit stronger conditional heteroskedasticity than in cases (i)-(iii). In cases (i)-(iv), the hybrid CI has an unfair advantage over the CHR CI, because it uses a GARCH(1, 1) model which is correctly specified in these cases. The results are invariant to the choice of  $\mu$ .

We consider a sample size of n = 130. The hybrid CI is based on a GARCH(1, 1) specification.<sup>20</sup> The hybrid critical values use subsamples of size b = 12, as in AG09.

We report average lengths of "CP-corrected" CI's. A CP-corrected CI equals the actual nominal 95% CI if its CP is at least .95 (for the given data-generating process), but otherwise equals the CI implemented at a nominal CP that makes the finite-sample CP equal to  $.95.^{21}$  All simulation results are based on 30,000 simulation repetitions.

Table 3 reports the results. CHR denotes the CI in (2.7). Hyb denotes the hybrid CI of AG09. The new CHR CI has very good finite-sample coverage probabilities. Specifically, its CP's (×100) are in the range [94.1, 94.8] for all values of  $\rho$  in cases (i)-(iii). For cases (iv) and (v), the range is [93.2, 94.5]. The hybrid CI has CP's in the range [94.2, 98.5] for cases (i)-(iii) and [93.9, 98.5] for cases (iv) and (v). These CP's reflect the fact that the hybrid CI is not asymptotically similar due to its reliance on subsampling.

The average length results of Table 3 (CP-corrected) show that the CHR CI is shorter than the hybrid CI for all values of  $\rho$  in cases (i)-(iv). The greatest length reductions are for  $\rho = .5, .0$ , where the CHR CI is from .69 to .83 times the length of the hybrid CI in cases (i)-(iii). For  $\rho = .99, .9$ , it is from .86 to .91 times the length of the hybrid CI in cases (i)-(iii). In cases (iv) and (v), the CHR and hybrid CI's have similar lengths for  $\rho = .99, .9$ . In cases (iv) and (v), for  $\rho = .5, .0$ , the CHR CI is from .82 to .98 times the length of the hybrid CI. In conclusion, in an overall sense, the CHR CI out-performs the hybrid CI in terms of average length by a noticeable margin in the cases considered.<sup>22</sup> Simulations for the symmetric two-sided subsampling CI of Romano and Wolf (2001) given in the Appendix show that the latter CI under-covers substantially in some cases (e.g., its CP's (×100) are 88.9, 88.3, 86.7 for b = 8, 12, 16 in case (ii) with  $\rho = .0$ ). It is longer than the symmetric and equal-tailed CHR CI's when  $\rho = .99$  in cases (i)-(v) and has similar average length (CP-corrected) in other cases. Hence, the CHR CI's out-perform the Romano and Wolf (2001) CI in the finite-sample cases considered.

Results reported in the Appendix compare the CHR CI in the i.i.d. case with the analogous CI that employs the homoskedastic variance estimator.<sup>23</sup> The use of the HC5 variance matrix estimator increases the deviations of the CP's (×100) from 95.0 compared to the homoskedastic variance estimator somewhat, but even so, the deviations for the equal-tailed CI's are only .3 on average over the five  $\rho$  values. It has no impact on the average lengths except when  $\rho = .99$ , in which case the impact is very small (8.3 for the equal-tailed CHR CI versus 8.1 for the equal-tailed homoskedastic variance CI). Hence, the CHR CI pays a very small price in the i.i.d. case for its robustness to conditional heteroskedasticity.

Table 3. Coverage Probabilities and (C.	P-Corrected) Average Lengths of Nominal
95% Equal-Tailed Two-Sided CI's: CHR a	nd Hybrid

								Ave	rage	Leng	gths (	×100)
			Cove	rage P	robabi	lities (	$\times 100)$		(CP)	-Cor	rected	1)
Innovations	$\operatorname{CI}$	$\rho$ :	.99	.9	.5	.0	9	.99	.9	.5	.0	9
(i) $GARCH(1, 1)$ -	CHR		94.2	94.7	94.8	94.5	94.4	8.5	19	33	37	17
(.05, .9; .001)	Hyb		98.5	98.3	96.5	95.2	95.6	9.6	21	46	47	19
(ii) $GARCH(1, 1)$ -	CHR		94.2	94.6	94.7	94.1	94.2	8.8	20	37	43	18
(.15, .8; .2)	Hyb		98.0	97.9	96.0	94.3	95.0	9.8	22	49	52	21
(iii) I.i.d.	CHR		94.5	94.7	94.8	94.7	94.6	8.3	18	31	35	16
	Hyb		97.7	97.6	95.7	94.2	94.8	9.6	21	45	48	19
(iv) $GARCH(1, 1)$ -	CHR		94.3	94.5	94.4	93.7	94.1	9.2	21	42	49	20
(.25,.7;.2)	Hyb		98.4	98.3	95.9	94.8	95.1	9.5	22	51	54	21
	-											
(v) $ARCH(4)$ -	CHR		94.5	94.3	93.9	93.2	94.0	9.6	23	48	56	22
(.3,.2,.2,.2;.2)	Hyb		98.5	98.2	95.9	94.2	95.4	9.1	22	53	57	21

### Footnotes

<sup>1</sup> Throughout this paper we use the term "asymptotic size" to mean the limit as  $n \to \infty$  of the finite-sample size. Uniformity in the asymptotics is built into this definition because finite-sample size is a uniform concept. By the definition of asymptotic size, the infimum of the coverage probability over different values of  $\rho$  and different innovation distributions is taken before the limit as  $n \to \infty$  is taken.

 $^{2}$  The CI of Stock (1991) needs to be modified as in Mikusheva (2007a) to have correct asymptotic size.

<sup>3</sup> The correct asymptotic size of this CI is established in the Appendix. The equaltailed subsampling CI of Romano and Wolf (2001) does not have correct asymptotic size under homoskedasticity or heteroskedasticity, see Mikusheva (2007a) and AG09.

<sup>4</sup> Lack of asymptotic similarity implies that the CI over-covers asymptotically for some sequences of  $\rho$  values. This may yield a longer CI than is possible.

<sup>5</sup> Gonçalves and Kilian (2004, 2007) also consider inference in autoregressive models with conditional heteroskedasticity using bootstrap methods. Their results do not allow for unit roots or roots near unity. Kuersteiner (2001) provides some related results for stationary models with conditional heteroskedasticity. Cavaliere and Taylor (2009) provide unit root tests in models with conditional heteroskedasticity using a recursive wild bootstrap. Inoue and Kilian (2002) consider bootstrap methods for autoregressions with unit roots but their results only apply to non-unit root parameters and they do not allow for conditional heteroskedasticity.

<sup>6</sup> AG09 also introduces several other CI's that have correct asymptotic size under conditional heteroskedasticity using size-corrected fixed critical values and sizecorrected subsampling critical values (for equal-tailed CI's). The performance of these CI's is not as good as that of the FQGLS-based hybrid CI, so we do not discuss these CI's further here.

<sup>7</sup> As in Mikusheva's (2007a) modification of Stock's CI, we invert the t statistic that is designed for a given value of  $\rho$ , not the t statistic for testing  $H_0: \rho = 1$  which is employed in Stock (1991). This is necessary to obtain correct asymptotic coverage when  $\rho$  is not  $O(n^{-1})$  local to unity.

<sup>8</sup> Mikusheva's (2007a) results do not cover the new CI because (i) she does not consider innovations that have conditional heteroskedasticity and (ii) even in the i.i.d. innovation case the t statistic considered here does not lie in the class of test statistics that she considers.

<sup>9</sup> The use of a stationary initial condition when  $\rho < 1$ , rather than a zero initial condition, is not crucial to obtaining robustness to conditional heteroskedasticity. Our results also apply to the case of a zero initial condition, in which case the second component of  $I_h^*(r)$  in (2.4) below is deleted.

<sup>10</sup> That is, when  $\rho$  converges to unity, one obtains the same asymptotic distribution whether an HC or a homoskedastic variance estimator is employed.

<sup>11</sup> Note that there is a one-to-one mapping between the sum of the AR coefficients and the cumulative impulse response. Hence, a CI for the former yields a CI for the latter. See Andrews and Chen (1994) for a discussion of the advantages of the sum of the AR coefficients over the largest AR root as a measure of long run dynamics of an AR(k) process.

<sup>12</sup> Symmetric two-sided and one-sided CI's can be handled in a similar fashion, see the Appendix for details.

<sup>13</sup> We prefer equal-tailed CI's over symmetric CI's in the AR(1) context because the latter can have quite unequal coverage probabilities for missing the true value above and below when  $\rho$  is near unity, which is a form of biasedness, due to the lack of symmetry of the near-unit root distributions.

<sup>14</sup> The quantity  $p_{ii}^*$  used in HC5 is a finite-sample adjustment to the standard HC variance estimator. In contrast, the HC3 variance estimator uses  $p_{ii}$  in the definition of  $\Delta$ . The use of  $p_{ii}^*$  guarantees that the finite-sample adjustment does not affect the asymptotics. When  $n(1 - \rho_n) \rightarrow h < \infty$ , it is straightforward to show that the use of  $p_{ii}$  is valid asymptotically. In other cases, it is more difficult to do so. However, the finite-sample results reported below are essentially the same whether  $p_{ii}^*$  or  $p_{ii}$  is used. Users may find it more convenient to use the HC3 version because it is computable in STATA using the linear regression option vce(hc3). Note that the asymptotic results given in the paper hold if one sets  $p_{ii}^* = 0$ , which yields the standard HC variance estimator.

<sup>15</sup> Note that it is possible for  $CI_{CHR,n}$  to consist of two disconnected intervals of the form  $[a, b] \cup [c, 1]$ , where  $-1 + \varepsilon \leq a < b < c \leq 1$ . This occurs with very low probability in most cases, and low probability in all cases, see the Appendix for details.

<sup>16</sup> For example, the fixed-design wild bootstrap and the pairs bootstrap considered in Gonçalves and Kilian (2004, 2007) for stationary observations are not suitable, see the Appendix. A suitable bootstrap is similar to, but different from, the recursive wild bootstrap considered by Cavaliere and Taylor (2009), which is designed for unit root tests, and the recursive wild bootstrap considered by Gonçalves and Kilian (2004), which is designed for stationary observations, see the Appendix.

<sup>17</sup> See MacKinnon and White (1985) and Long and Ervin (2000) for simulation results concerning the properties of the HC3 estimator in the standard linear regression model with i.i.d. observations.

<sup>18</sup> The hybrid CI is defined as in AG09 using the standard HC variance estimator with  $p_{ii}^* = 0$ , not the HC5 estimator.

<sup>19</sup> For example, see Bollerslev (1987), Engle, Ng, and Rothschild (1990), and, for more references, Ma, Nelson, and Startz (2007).

 $^{20}$  See Section 10 of the Appendix for more details concerning the definition and computation of the hybrid CI. Note that the hybrid CI has correct asymptotic size whether or not the GARCH(1, 1) specification is correct.

<sup>21</sup> When calculating the average length of a CI, we restrict the CI to the interval

[-1, 1]. The search to find the nominal significance level such that the actual finitesample CP (×100) equals 95.0 is done with stepsize .025. In the case of a disconnected CI, the "gap" in the CI is not included in its length.

<sup>22</sup> The CHR CI also out-performs the hybrid CI based on the infeasible QGLS estimator, see the Appendix. The CP (×100) results of Table 3 using  $p_{ii}$ , rather than  $p_{ii}^*$ , are the same in all cases except case (i)  $\rho = .99$ , case (iv)  $\rho = .5, .0$ , and case (v)  $\rho = .99$ , where the differences are .1% (e.g., 94.2% versus 94.3%), and case (v)  $\rho = .5, .0$ , where the differences are .2% and .3%, respectively. There are no differences in the average lengths. For the symmetric two-sided CHR CI, the CP results and the average length results compared to the hybrid CI are similar to those in Table 3, although slightly better in both dimensions, see the Appendix.

<sup>23</sup> The latter CI is Mikusheva's (2007a) modification of Stock's (1991) CI applied to the LS estimator of  $\rho$ , but with a stationary initial condition when  $\rho < 1$ , rather than a zero initial condition.

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# Appendix

to

A Conditional-Heteroskedasticity-Robust Confidence Interval for the Autoregressive Parameter

Donald W. K. Andrews Cowles Foundation for Research in Economics Yale University

> Patrik Guggenberger Department of Economics UCSD

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### 5 Outline

This Appendix is organized as follows: Section 6 establishes the asymptotic false coverage probabilities (FCP's) of the equal-tailed CHR CI under local alternatives to the true values. Section 7 assesses the asymptotic and finite-sample price the CHR CI pays in the i.i.d. innovation case for its robustness to conditional heteroskedasticity. Section 8 provides the probabilities of obtaining disconnected CHR CI's. Section 9 defines symmetric two-sided CHR CI's and gives tables of critical values for them. It also provides simulation results for a variety symmetric two-sided confidence intervals, including the CHR CI, the hybrid and subsampling CI's of Andrews and Guggenberger (2009a) (AG09a), and the subsampling CI of Romano and Wolf (2001). Section 10 provides details regarding the implementation of the Monte Carlo simulations in the paper and the Appendix. Section 11 defines a recursive residual-based wild bootstrap procedure which could be used to obtain the critical values for the CHR CI. Section 12 provides the proof of Theorem 1 of the paper. Section 13 gives the proof of the asymptotic FCP result that is stated in Section 6. Section 14 provides a proof of the correct asymptotic size in the presence of conditional heteroskedasticity of Romano and Wolf's (2001) symmetric two-sided subsampling CI, which is based on a least squares (LS) based t statistic with a homoskedastic variance estimator (designed for the i.i.d. innovations case).

# 6 Local Asymptotic False Coverage Probabilities of the CHR CI

In this section, we determine the asymptotic FCP's of the CHR CI for sequences of local alternatives  $\{\rho_n^* : n \ge 1\}$  to the true parameters  $\{\rho_n : n \le 1\}$ . We provide results for the full spectrum of cases in which  $n(1 - \rho_n) \to h$  for (i)  $0 \le h < \infty$ , (ii)  $h = \infty$  and  $\rho_n \to 1$ , and (iii)  $h = \infty$  and  $\rho_n \to \rho_\infty < 1$ . Asymptotic results of this sort are not available currently for any of the CI's in the literature under conditional homoskedasticity or conditional heteroskedasticity.

Theorem 1 of Andrews and Guggenberger (2012) (AG12) shows that under sequences  $\{(\rho_n, F_n) \in \Lambda : n \geq 1\}$  for which  $n(1-\rho_n) \to h \in [0, \infty]$  the LS estimator  $\hat{\rho}_n$ , the heteroskedasticity consistent HC5 variance estimator  $\hat{\sigma}_n^2$ , and the corresponding t statistic  $T_n(\rho) = n^{1/2}(\hat{\rho}_n - \rho)/\hat{\sigma}_n$  satisfy:

$$(n^{1/2}d_n(\widehat{\rho}_n - \rho_n), \ d_n\widehat{\sigma}_n, \ T_n(\rho_n))' \to_d (N_h, S_h, T_h).$$

$$(6.1)$$

The result of AG12 actually is for the HC variance estimator with 0 in place of  $p_{ii}^*$ . Because  $\max_{i \leq n} p_{ii}^* \leq n^{-1/2} \to 0$ , the proofs go through with  $\hat{\sigma}_n^2$  being the HC5 variance estimator.

In (6.1), (i)  $\{d_n : n \ge 1\}$  is a sequence of constants defined below, (ii)  $T_h = N_h/S_h$ ,

(iii) when  $h = \infty$ ,  $N_h \sim N(0, 1)$  and  $S_h = 1$ , and (iv) when  $0 \le h < \infty$ ,

$$N_{h} = \int_{0}^{1} I_{D,h}^{*}(r) dW(r) / \int_{0}^{1} I_{D,h}^{*}(r)^{2} dr \text{ and } S_{h} = \left(\int_{0}^{1} I_{D,h}^{*}(r)^{2} dr\right)^{-1/2}, \quad (6.2)$$

where  $W(\cdot)$  and  $I^*_{D,h}(\cdot)$  are defined in (2.4) of the paper.<sup>1</sup>

The constants  $\{d_n : n \ge 1\}$  depend on  $(\rho_n, F_n)$ , although their order of magnitude only depends on  $\rho_n$ . When  $0 \le h < \infty$ ,  $d_n = n^{1/2}$ . When  $h = \infty$ ,

$$d_n = E_{F_n} Y_{n,i-1}^{*2} / (E_{F_n} Y_{n,i-1}^{*2} U_{n,i}^2)^{1/2},$$
(6.3)

where by stationarity  $d_n$  does not depend on *i*.

The result in (6.1) shows that the local alternatives for which the CHR CI has non-trivial asymptotic FCP's (i.e., asymptotic FCP's less than  $1 - \alpha$ ) are of the form:

$$\rho_n^* = \rho_n - \frac{\delta_n}{n^{1/2} d_n} \tag{6.4}$$

for any sequence of constants  $\{\delta_n : n \ge 1\}$  such that  $\delta_n \to \delta \in \mathbb{R}$  and  $h + \delta \ge 0$  if  $0 \le h < \infty$ .

If  $0 \le h < \infty$ , then the local alternatives  $\{\rho_n^* : n \ge 1\}$  are  $n^{-1}$ -local alternatives from the true values  $\{\rho_n : n \ge 1\}$  because  $n^{1/2}d_n = n$ .

If  $h = \infty$ ,  $\rho_n \to \rho_\infty < 1$ , and  $\lim_{n\to\infty} d_n = d_\infty \in (0,\infty)$ , then

$$\rho_n^* = \rho_n - (\delta d_\infty^{-1} / n^{1/2})(1 + o(1)).$$
(6.5)

In this case,  $\{\rho_n^* : n \ge 1\}$  are  $n^{-1/2}$ -local alternatives from  $\{\rho_n : n \ge 1\}$ . Note that the condition on  $d_n$  is not stringent. For example, it holds if  $(\rho_n, F_n)$  does not depend on n and  $E_F(Y_i^* - E_F Y_i^*)^2 U_i^2 \in (0, \infty)$ .

If  $h = \infty$  and  $\rho_n \to 1$ , then

$$d_n = (1 - \rho_n^2)^{-1/2} (1 + o(1)) \tag{6.6}$$

by equation (11) of AG12 and the first two results of Lemma 6 in AG12. The  $n^{1/2}(1-\rho_n^2)^{-1/2}$  rate of convergence of the LS estimator in this case was first obtained by Giraitis and Phillips (2006) and Phillips and Magdalinos (2007), but under assumptions that rule out conditional heteroskedasticity in the innovations. In the present case, the local alternatives are of the form

$$\rho_n^* = \rho_n - \frac{\delta(1 - \rho_n^2)^{1/2}}{n^{1/2}} (1 + o(1)), \tag{6.7}$$

which constitute deviations that are smaller than  $O(n^{-1/2})$  and larger than  $O(n^{-1})$ .

By definition, let  $h + \delta = \infty$  if  $h = \infty$  and  $\delta \in R$ .

We have the following asymptotic FCP result.

<sup>&</sup>lt;sup>1</sup>When  $\hat{\rho}_n$  is the LS estimator, then (in the notation of AG12)  $\hat{\phi}_{n,i} = \phi_{n,i} = 1 \quad \forall i \leq n$  and the constants in Thm. 1 of AG12 simplify:  $h_{2,1} = h_{2,2} = \lim_{n \to \infty} E_{F_n} U_i^2$  and  $h_{2,5} = h_{2,7} = 1$ . This yields the form of the asymptotic distributions in (6.1) and (6.2).

**Theorem 2** Let  $\alpha \in (0,1)$ . Let  $\{(\rho_n, F_n) \in \Lambda : n \geq 1\}$  be any sequence of true parameters and distributions for which  $n(1 - \rho_n) \to h \in [0, \infty], \rho_n \to \rho_\infty \in (-1, 1],$ and  $n^{1/2}d_n \to \infty$  if  $\rho_\infty < 1$ . Let  $\{\rho_n^* : n \geq 1\}$  be any sequence of alternative parameters that satisfies  $\rho_n^* = \rho_n - \delta_n/(n^{1/2}d_n)$ , where  $\delta_n \to \delta \in R$  and  $h + \delta \geq 0$  if  $0 \leq h < \infty$ . Then, the equal-tailed nominal  $1 - \alpha$  confidence interval  $CI_{CHR,n}$  for the AR parameter  $\rho$  satisfies

$$\lim_{n \to \infty} P_{\lambda_n}(\rho_n^* \in CI_{CHR,n}) = P(c_{h+\delta}(\alpha/2) \le T_h + \delta/S_h \le c_{h+\delta}(1-\alpha/2)).$$

**Comments. 1.** If  $h = \infty$ , then  $h + \delta = \infty$ ,  $c_{h+\delta}(\alpha/2) = -z_{1-\alpha/2}$ ,  $c_{h+\delta}(1-\alpha/2) = z_{1-\alpha/2}$ ,  $T_h \sim N(0,1)$ ,  $S_h = 1$ , and the limit FCP in the result of Theorem 2 equals  $P(|T_h + \delta| \le z_{1-\alpha/2})$ , which is less than  $1 - \alpha$  for all  $\delta \ne 0$ . This result holds even if  $\rho_n \rightarrow 1$  provided  $n(1-\rho_n) \rightarrow \infty$ . In this case, the distance of the local alternatives from the true values depends on how fast  $\rho_n$  goes to one via  $1 - \rho_n^2$ , as in (6.7), but the form of the asymptotic FCP is the same as when  $\rho_n \rightarrow \rho_\infty < 1$ . If  $0 \le h < \infty$ , then the limit FCP in the result of Theorem 2 is not a standard quantity, but it can be simulated quite easily.

2. Using the results of AG12, one could establish analogous results to those in Theorem 2 for the hybrid CI of AG09a and the symmetric subsampling CI of Romano and Wolf (2001). For brevity, we do not do so here.

#### 7 Price for Robustness of the CHR CI

This section assesses the price the CHR CI pays, in terms of asymptotic FCP's and finite-sample average lengths, in the i.i.d. innovation case to obtain robustness to conditional heteroskedasticity.

First, we show that the asymptotic price (to first order) is zero. Consider the CHR CI and the same CI but constructed with the homoskedastic variance estimator in place of the heteroskedasticity-consistent variance estimator. The latter CI is referred to as the MSS CI, because it is the same as Mikusheva's (2007) modification of Stock's (1991) CI applied to the LS estimator of  $\rho$ , but with a stationary initial condition when  $\rho < 1$ , rather than a zero initial condition. The latter affects the asymptotic distribution of the t statistic and hence the definition of the critical values. Under i.i.d. innovations, these two CI's have the same asymptotic FCP's (and coverage probabilities (CP's)).<sup>2</sup> The asymptotic FCP's of the CHR CI are given in Theorem 2. Identical results for the MSS CI under i.i.d. innovations hold because the two test statistics have the same asymptotic distributions in this case both when  $\rho_n \to 1$  and

<sup>&</sup>lt;sup>2</sup>Note that the MSS CI does not have correct asymptotic size when conditionally heteroskedastic innovation distributions are included in the parameter space. This is because the MSS t statistic has an asymptotic distribution that depends on the form of the conditional heteroskedasticity and is not standard normal when  $\rho_n \to \rho_\infty < 1$ , but the critical value is taken from the i.i.d. innovation case which is a standard normal quantile when  $\rho_n \to \rho_\infty < 1$ .

when  $\rho_n \to \rho_{\infty} < 1$ . When  $\rho_n \to 1$  this holds by the proof given in Section 14 below of the correct asymptotic size of Romano and Wolf's (2001) symmetric subsampling CI. When  $\rho_n \to \rho_{\infty} < 1$  and the innovations are i.i.d., it holds by standard results because the two variance estimators have the same probability limit.

Next, because the asymptotic price is zero, we use simulations to assess the finitesample price. The data generating process considered is the i.i.d. standard normal innovation case (i.e., case (iii) in the paper). The sample size is n = 130.

Table A-1 reports CP's ( $\times 100$ ) and (CP-corrected) average lengths for the CHR and MSS CI's with nominal size 95.0%. Results are given for both equal-tailed and symmetric two-sided CI's.

Table A-1 shows that the use of the HC5 variance matrix estimator increases the deviations of the CP's (×100) from 95.0 slightly compared to the homoskedastic variance estimator. It has no impact on average length except when  $\rho = .99$ , in which case the impact is very small. Thus, the CHR CI pays a very small price in the i.i.d. case for its robustness to conditional heteroskedasticity.

Note that CI's analogous to the CHR CI's (equal-tailed, symmetric, and onesided) based on the HC5 variance estimator can be defined with other versions of the HC variance matrix, such as the HC, HC1, HC2, and HC3 estimators defined in MacKinnon and White (1985), but we find that the HC5 variance estimator gives the best finite-sample CP's and the choice has very little effect on the (CP-corrected) average lengths.<sup>3</sup> As noted in a footnote in Section 2 of the paper, the HC5 estimator and the HC3 (without the (n - 1)/n term) estimator have almost the same finitesample properties for the cases in Table 3 of the paper.

Next, we briefly discuss simulation comparisons between the (equal-tailed) CHR CI and the infeasible hybrid CI which is based on the infeasible QGLS estimator. By definition, the infeasible QGLS estimator takes the GARCH(1, 1) (or ARCH(4)) specification as known and its parameter values as known. Simulations for the hybrid CI based on the infeasible QGLS estimator (not reported) show that it over-covers in many cases and its CP's exceed those of the FQGLS hybrid CI in almost all cases. In consequence, its average lengths are the same or slightly longer than those of the FQGLS hybrid CI in cases (i)-(iv) and only slightly shorter in case (v), reported in Table 3 of the paper. Hence, the CHR CI out-performs the infeasible QGLS hybrid CI in the cases considered.

 $<sup>^{3}</sup>$ We denote the HC variance estimator introduced in this paper as HC5 because Cribari-Neto (2004) uses HC4 to denote a different HC variance estimator.

							Avera	age I	lengt	hs (>	(100)	
			Cove	rage P	robabi	lities (	(	CP-	Corre	ected	)	
CI-Type	$\operatorname{CI}$	$\rho$ :	.99	.9	.5	.0	9	.99	.9	.5	.0	9
Equal-tailed	CHR		94.5	94.7	94.8	94.7	94.6	8.3	18	31	35	16
	MSS		95.0	94.9	95.1	94.9	94.6	8.1	18	31	35	16
Symmetric	CHR		94.5	95.2	95.1	94.8	95.6	10.6	19	31	35	16
	MSS		95.1	95.2	95.4	95.2	95.4	10.4	19	31	35	16

Table A-1. Coverage Probabilities and (CP-Corrected) Average Lengths of Nominal 95% Two-Sided CI's with I.i.d. Innovations: CHR and MSS

#### 8 Disconnected CHR Confidence Intervals

It is possible for CHR CI's to consist of two disjoint intervals of the form  $[a, b] \cup [c, 1]$ for  $-1 + \varepsilon \leq a < b < c \leq 1.^4$  To see why, consider the nominal 95% equal-tailed CHR CI. The critical values  $c_h(.025)$  and  $c_h(.975)$  are increasing and concave as functions of h, see Tables 1 and 2 of the paper. Thus,  $c_{n(1-\rho)}(.025)$  and  $c_{n(1-\rho)}(.975)$  are decreasing concave functions of  $\rho$ . Viewed as a function of  $\rho$  on [-1, 1], the critical value functions are essentially flat and take values close to -1.96 and 1.96, respectively, for most of [-1, 1] and dip down to the values -3.13 and .24 as  $\rho$  approaches one.

The test statistic is a linear function of  $\rho$  with negative slope. The test statistic takes the value 0 for the value of  $\rho$  equal to  $\hat{\rho}_n$ . The CI consists of all values  $\rho$  where the linear test statistic function lies between the mostly horizontal critical value curves. Drawing the corresponding picture, one can see that disconnected CI's can occur if the linear test statistic line cuts across the lower critical value curve as it dips near one and intersects with it at two places. This only occurs for a very small range of values of  $n^{1/2}$ ,  $\hat{\rho}_n$ , and  $\hat{\sigma}_n$ .

No such disconnected CI feature occurs if the test statistic line intersects the upper critical value curve in two places because  $\rho$  values between the two intersection points are in the CI, not excluded from it, in this case.

Table A-2 provides simulated values of the probability that the CHR equal-tailed and symmetric CI's are disconnected for the five cases considered in Table 3 of the paper.<sup>5</sup> The sample size is n = 130. In cases (i)-(iii) except for  $\rho = .9$  and in cases (iv) and (v) except for  $\rho = .9, .5$ , the probability of a disconnected CI is essentially zero. For  $\rho = .9$  in cases (i)-(iii) the probability is still quite small ( $\leq 5/1000$ ). For  $\rho = .9, .5$ in cases (iv) and (v), which are the stronger and less realistic forms of conditional heteroskedasticity, the probabilities are larger, but still small ( $\leq 19/1000$ ).

<sup>&</sup>lt;sup>4</sup>The same is true of Mikusheva's (2007) modification of Stock's (1991) CI.

<sup>&</sup>lt;sup>5</sup>These results are based on 30,000 simulation repetitions with the asymptotic critical values computed using 100,000 repetitions and n = 30,000.

			$\Pr$	obability	v of	
			Disconi	nected (	CHR CI	
$\operatorname{CI}$	$\rho$ :	.99	.9	.5	.0	9
eq-tail		.0001	.0008	.0000	.0000	.0000
$\operatorname{sym}$		.0001	.0028	.0000	.0000	.0000
eq-tail		.0001	.0023	.0020	.0000	.0000
$\operatorname{sym}$		.0002	.0050	.0011	.0000	.0000
eq-tail		.0000	.0008	.0000	.0000	.0000
sym		.0002	.0014	.0000	.0000	.0000
eq-tail		.0001	.0035	.0089	.0002	.0000
sym		.0010	.0074	.0054	.0002	.0000
eq-tail		.0003	.0054	.0190	.0020	.0000
sym		.0008	.0102	.0170	.0016	.0000
	eq-tail sym eq-tail sym eq-tail sym eq-tail sym eq-tail	eq-tail sym eq-tail sym eq-tail sym eq-tail sym eq-tail	CI $\rho$ :         .99           eq-tail         .0001           sym         .0001           eq-tail         .0001           sym         .0001           eq-tail         .0001           sym         .0002           eq-tail         .0000           sym         .0002           eq-tail         .0000           sym         .0001           eq-tail         .0001           sym         .0010           eq-tail         .0003	$\begin{array}{c c c c c c c c c c c c c c c c c c c $	$\begin{array}{c c c c c c c c c c c c c c c c c c c $	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$

Table A-2. Probabilities of Obtaining Disconnected CHR CI's

#### 9 Symmetric Two-Sided Confidence Intervals

Let  $CI_{CHR,n}^{sym}$  denote the symmetric two-sided nominal  $1 - \alpha$  CHR CI that is analogous to the equal-tailed CHR CI introduced in the paper. It is defined as follows:

$$CI_{CHR,n}^{sym} = \{ \rho \in [-1 + \varepsilon, 1] : |T_n(\rho)| \le c_h^{sym}(1 - \alpha) \text{ for } h = n(1 - \rho) \},$$
(9.8)

where  $c_h^{sym}(1-\alpha)$  is the  $1-\alpha$  quantile of the asymptotic distribution  $|J_h|$  of  $|T_n(\rho)|$ .

Tables A-3 to A-5 report  $c_h^{sym}(1-\alpha)$  for  $\alpha = .05, .01, .1$ , respectively, and a range of h values, which are used to calculate 95%, 99%, and 90% symmetric two-sided CHR CI's.

Table A-3. Values of  $c_h^{sym}(.95)$ , the .95 Quantile of  $|J_h|$ , for Use with 95% Symmetric Two-Sided Confidence Intervals

h	0	.2	.4	.6	.8	1.0	1.4	1.8	2.2	2.6	3.0	3.4	3.8
$c_h^{sym}(.95)$	2.87	2.83	2.79	2.76	2.73	2.70	2.65	2.61	2.57	2.54	2.51	2.49	2.46
	4.2												
$c_h^{sym}(.95)$	2.44	2.42	2.40	2.36	2.32	2.30	2.27	2.25	2.23	2.21	2.20	2.19	2.17
$c_h^{sym}(.95)$	2.13	2.10	2.08	2.05	2.03	2.02	2.01	2.01	2.00	2.00	1.98	1.97	1.96

Table A-4. Values of  $c_h^{sym}(.99)$ , the .99 Quantile of  $|J_h|$ , for Use with 99% Symmetric Two-Sided Confidence Intervals

h	0	.2	.4	.6	.8	1.0	1.4	1.8	2.2	2.6	3.0	3.4	3.8
$c_h^{sym}(.99)$	3.44	3.40	3.38	3.35	3.32	3.30	3.26	3.22	3.19	3.16	3.14	3.11	3.09
h	4.2	4.6	5.0	6.0	7.0	8.0	9.0	10	11	12	13	14	15
$c_h^{sym}(.99)$	3.07	3.05	3.03	3.00	2.97	2.94	2.92	2.90	2.88	2.87	2.85	2.84	2.82
h	20	25	30	40	50	60	70	80	90	100	200	300	500
$c_h^{sym}(.99)$	2.77	2.74	2.72	2.68	2.66	2.65	2.64	2.63	2.62	2.61	2.59	2.58	2.57

Table A-5. Values of  $c_h^{sym}(.90)$ , the .90 Quantile of  $|J_h|$ , for Use with 90% Symmetric Two-Sided Confidence Intervals

h	0	.2	.4	.6	.8	1.0	1.4	1.8	2.2	2.6	3.0	3.4	3.8
$c_h^{sym}(.9)$	2.57	2.53	2.49	2.45	2.42	2.39	2.34	2.30	2.26	2.22	2.19	2.16	2.13
h	4.2	4.6	5.0	6.0	7.0	8.0	9.0	10	11	12	13	14	15
$c_h^{sym}(.9)$	2.11	2.09	2.07	2.02	1.98	1.95	1.93	1.91	1.89	1.87	1.86	1.85	1.83
h	20	25	30	40	50	60	70	80	90	100	200	300	500
$c_h^{sym}(.9)$	1.79	1.77	1.75	1.72	1.70	1.69	1.69	1.68	1.68	1.67	1.66	1.65	1.65

Under the conditions of Theorem 1, the symmetric two-sided nominal  $1 - \alpha$  CHR CI has asymptotic size equal to  $1 - \alpha$  and is asymptotically similar. Under the conditions of Theorem 2, the FCP's of the symmetric two-sided CHR CI satisfy

$$\lim_{n \to \infty} P_{\lambda_n}(\rho_n^* \in CI_{CHR,n}) = P(|T_h + \delta/S_h| \le c_{h+\delta}^{sym}(1-\alpha)).$$
(9.9)

(The proofs of these results are analogous to those given for the equal-tailed CHR CI in Sections 12 and 13 below.)

Table A-6 reports simulation results analogous to those in Table 3 of the paper except for symmetric two-sided CI's, rather than equal-tailed CI's. It reports results for the symmetric CHR, hybrid, and FQGLS subsampling (Sub<sub>GLS</sub>) CI's. The hybrid and Sub<sub>GLS</sub> CI's are proposed in AG09a. They are based on the FQGLS estimator with standard heteroskedasticity-consistent variance estimator (as in Table 3 of the paper), coupled with hybrid (fixed/subsampling) and subsampling critical values, respectively, see Section 10 for more details. The sample size is n = 130 and the subsample size is b = 12.

The results in Table A-6 are similar to those in Table 3 of the paper, but the CHR CI performs slightly better in terms CP's and in terms of average length compared to the hybrid CI. The hybrid and  $\text{Sub}_{GLS}$  CI's have very similar finite-sample properties (which is expected because they have the same asymptotic properties in terms of CP's and FCP's).

				, e	· · · ·		0.20					
								Ave	rage	Leng	gths (	×100)
			Cove	rage P	robabi	lities (	$\times 100)$		(CP)	-Cor	recteo	d)
Innovations	$\operatorname{CI}$	$\rho$ :	.99	.9	.5	.0	9	.99	.9	.5	.0	9
(i) $GARCH(1, 1)$ -	CHR		94.6	95.2	95.1	94.8	95.4	11	20	33	37	16
(.05, .9; .001)	Hyb		95.9	98.0	97.2	96.1	95.8	15	28	46	45	19
	$\operatorname{Sub}_{GLS}$		95.9	98.0	97.0	95.6	95.0	15	28	45	44	18
(ii) GARCH(1, 1)-	CHR		94.9	95.3	95.0	94.5	95.3	11	21	37	42	18
(.15,.8;.2)	Hyb		96.0	97.8	97.0	96.0	95.7	15	29	49	49	20
	$\operatorname{Sub}_{GLS}$		96.0	97.8	96.8	95.4	95.1	15	29	49	48	20
(iii) I.i.d.	CHR		94.5	95.2	95.1	94.8	95.6	11	19	31	35	16
	Hyb		95.3	97.4	96.8	95.6	95.2	15	27	45	45	19
	$\operatorname{Sub}_{GLS}$		95.3	97.4	96.6	95.1	94.5	15	27	45	44	19
(iv) $GARCH(1, 1)$ -	CHR		95.3	95.4	94.8	94.1	95.2	11	23	41	48	19
(.25,.7;.2)	Hyb		96.5	98.0	97.2	96.2	95.9	15	30	51	51	20
· · · ·	$\operatorname{Sub}_{GLS}$		96.5	98.0	97.0	95.6	95.4	15	30	51	50	20
(v) ARCH $(4)$ -	CHR		95.8	95.5	94.4	93.6	95.0	12	24	47	55	21
(.3,.2,.2,.2;.2)	Hyb		96.9	98.1	97.2	96.0	96.4	15	30	53	53	21
	$\operatorname{Sub}_{GLS}$		96.9	98.1	96.9	95.5	95.9	15	30	53	52	21

Table A-6. Coverage Probabilities and (CP-Corrected) Average Lengths of Nominal 95% Symmetric Two-Sided CI's: CHR, Hybrid, and  $Sub_{GLS}$ 

Next, Table A-7 reports results analogous to those of Table 3 of the paper but for the symmetric two-sided CHR and symmetric two-sided subsampling CI of Romano and Wolf (2001) (Sub<sub>*RW*</sub>). The Sub<sub>*RW*</sub> CI is based on the LS estimator and the homoskedastic variance matrix estimator. For the Sub<sub>*RW*</sub> CI, we compute results for subsample sizes b = 8, 12, 16, 20. In Table A-7, we report results for b = 8, 12, because they provide the best results in terms of CP's and average lengths. We discuss the results for Sub<sub>*RW*</sub> CI with b = 8 because they are better than those for b = 12 in terms of CP's. Table A-7 shows that the Sub<sub>*RW*</sub> CI exhibits problems with under-coverage in some cases. For example, when  $\rho = .0$ , its CP's (×100) in the five cases considered lie in the interval [82.5, 92.8], whereas those of the CHR CI lie in [93.6, 94.8]. When  $\rho = .5$ , the CP's (×100) of the Sub<sub>*RW*</sub> CI in the five cases lie in [87.4, 95.8], whereas those of the CHR CI lie in [94.4, 95.1]. The average (CP-corrected) lengths of the Sub<sub>*RW*</sub> CI's are noticeably longer than those of the CHR CI for  $\rho = .99$  (for all five specifications of the conditional heteroskedasticity), but similar for most other parameter values.

									Average Lengths $(\times 100)$					
			Coverage Probabilities $(\times 100)$						(CP-Corrected)					
Innovations	$\operatorname{CI}$	$\rho$ :	.99	.9	.5	.0	9	.9	9	.9	.5	.0	9	
(i) $GARCH(1, 1)$ -	CHR		94.6	95.2	95.1	94.8	95.4	1	1	20	33	37	16	
(.05, .9; .001)	$\operatorname{Sub}_{RW} b \!=\! 8$		95.3	97.3	95.2	91.8	93.4	1	4	23	34	39	14	
	$\operatorname{Sub}_{RW} b = 12$		93.4	96.1	93.6	90.8	91.8	1	5	22	35	40	15	
(ii) $GARCH(1, 1)$ -	$\operatorname{CHR}$		94.9	95.3	95.0	94.5	95.3	1	1	21	37	42	18	
(.15, .8; .2)	$\operatorname{Sub}_{RW} b \!=\! 8$		94.7	96.5	93.0	88.9	92.2	1	5	23	38	45	15	
	$\operatorname{Sub}_{RW} b = 12$		93.0	95.2	91.5	88.3	90.6	1	5	22	40	45	17	
(iii) I.i.d.	CHR		94.5	95.2	95.1	94.8	95.6	1	1	19	31	35	16	
	$\operatorname{Sub}_{RW} b = 8$		95.6	97.6	95.8	92.8	93.8	1	1	23	34	36	13	
	$\operatorname{Sub}_{RW} b = 12$		93.7	96.4	94.2	91.6	92.3	1	5	22	34	37	15	
(iv) $GARCH(1, 1)$ -	CHR		95.3	95.4	94.8	94.1	95.2	1	1	23	41	48	19	
(.25,.7;.2)	$\operatorname{Sub}_{RW} b = 8$		94.2	95.4	90.5	85.9	91.3	1	5	23	43	52	16	
	$\operatorname{Sub}_{RW} b = 12$		92.6	94.1	89.2	85.6	90.0	1	5	23	45	47	17	
(v) $ARCH(4)$ -	CHR		95.8	95.5	94.4	93.6	95.0	12	2	24	47	55	21	
(.3,.2,.2,.2;.2)	$\operatorname{Sub}_{RW} b = 8$		93.8	93.8	87.4	82.5	90.5	1	3	25	52	54	17	
	$\operatorname{Sub}_{RW} b = 12$		92.4	92.5	86.2	82.9	89.5	1'	7	26	46	49	18	

Table A-7. Coverage Probabilities and (CP-Corrected) Average Lengths of Nominal 95% Symmetric Two-Sided CI's: CHR and the Romano and Wolf (2001) Subsampling CI (Sub<sub>RW</sub>) with Subsample Sizes b = 8 and b = 12

In sum, the symmetric two-sided subsampling CI of Romano and Wolf (2001) does not perform as well as the symmetric CHR CI due its noticeable under-coverage in some cases.

Comparing the results of the  $\text{Sub}_{GLS}$  and  $\text{Sub}_{RW}$  CI's in Tables A-6 and A-7, it is clear that the use of the feasible FQGLS estimator of  $\rho$  combined with a HC variance matrix estimator, compared to the LS estimator of  $\rho$  combined with the homoskedastic variance estimator, improves the finite-sample coverage probabilities of the subsampling CI's noticeably.

Additional simulations show that much of this difference is due to the use of the HC variance matrix estimator (even though the latter is not necessary to obtain correct asymptotic size). Specifically, we computed CP's for the symmetric subsampling CI based on the LS estimator combined with the standard heteroskedasticity-consistent variance estimator, denoted  $\text{Sub}_{LS,Het}$ , which differs from  $\text{Sub}_{RW}$  only in the choice of the variance matrix estimator. For i.i.d. innovations, n = 130, b = 12, and

 $\rho = .99, .9, .5, .0, -.9$ , the Sub<sub>RW</sub> and Sub<sub>LS,Het</sub> CI's have CP's (×100): (93.7, 98.7), (96.4, 99.1), (94.2, 98.4), (91.6, 97.3), (92.3, 96.6). Hence, even in the i.i.d. innovations case, there are substantial differences between the n = 130 finite-sample CP's of the Sub<sub>RW</sub> and Sub<sub>LS,Het</sub> CI's.

#### 10 Monte Carlo Details

The hybrid and  $\operatorname{Sub}_{GLS}$  CI's reported in Table 3 of the paper and Table A-6 are based on a t statistic constructed using a FQGLS estimator of  $\rho$  that employs estimators  $\{\widehat{\phi}_{n,i}^2 : i \leq n\}$  of the conditional variances  $\{\sigma_i^2 : i \leq n\}$ . The studentized t statistic is  $T_{GLS,n}(\rho) = n^{1/2}(\widehat{\rho}_{GLS,n} - \rho)/\widehat{\sigma}_{GLS,n}$ , where  $\widehat{\rho}_{GLS,n}$  is the LS estimator from the regression of  $Y_i/\widehat{\phi}_i$  on  $Y_{i-1}/\widehat{\phi}_i$  and  $1/\widehat{\phi}_i$  for i = 1, ..., n and  $\widehat{\sigma}_{GLS,n}^2$  is the (1, 1) element of the standard heteroskedasticity-robust variance estimator for the LS estimator in the preceding regression (which does not employ the HC5 adjustment factor  $1/(1 - p_{ii}^*)$ ).

The estimators  $\{\widehat{\phi}_{n,i}^2 : i \leq n\}$  are based on a GARCH(1, 1) parametric specification of the conditional heteroskedasticity. The GARCH(1, 1) model is estimated using the closed-form estimator of Kristensen and Linton (2006) applied to the LS residuals. This estimator is employed in the simulations because it is very quick to compute. More precisely, we use two Newton-Raphson iterations (see Kristensen and Linton's (2006) equation (17)), and we initialize the iteration using their closed-form estimator (see their equation (10) on p. 326) implemented with  $w_1 = w_2 = w_3 = 1/3$  and with their  $\widehat{\phi}$  Winsorized to the interval [.001, .999]. In each iteration step, we initialize the  $\widehat{\sigma}_{k,t}^2$  (see their p. 329, line 5 from the bottom) by setting it equal to the squared first data observation.<sup>6</sup>

In the simulations using an ARCH(4) data generating process, the GARCH(1, 1) specification is incorrect. Nevertheless, the hybrid and  $Sub_{GLS}$  CI's still have correct asymptotic size, see AG09a.

The asymptotic distribution of the FQGLS estimator in the  $n^{-1}$ -local to unity case depends on the parameter  $h_{2,7} = Corr(U_i, U_i/\phi_i^2)$ , where  $\phi_i^2$  is the conditional variance of the innovations based on the GARCH(1, 1) specification (which may or may not be correctly specified), with GARCH(1, 1) parameter values evaluated at the probability limit of the GARCH parameter estimators, see AG12. For the five processes considered in the simulations,  $h_{2,7}$  equals .98, .86, 1.00, .74, and .54, respectively.

For the equal-tailed two-sided nominal  $1 - \alpha$  hybrid CI, the upper critical value is the maximum of the subsampling critical value for nominal size  $1 - \alpha/2$  and the standard normal quantile  $z_{1-\alpha/2}$ . The lower critical value is the minimum of the

<sup>&</sup>lt;sup>6</sup>For simplicity, this estimator is not discretized and the GARCH(1, 1) process is not truncated to conform to the theoretical results given in the Section 3.4 of AG12 for the asymptotic equivalence of feasible and infeasible QGLS statistics.

subsampling critical value for nominal size  $\alpha/2$  and  $z_{\alpha/2}$ . For the symmetric twosided nominal  $1 - \alpha$  subsampling CI, i.e.,  $\operatorname{Sub}_{GLS}$ , the test statistic is the absolute value of  $T_{GLS,n}(\rho)$  and the critical value is the  $1 - \alpha$  sample quantile of the absolute values of the subsample t statistics. For the symmetric hybrid CI, the test statistic is the same, but the critical value is the maximum of the latter subsampling critical value and  $z_{1-\alpha/2}$ .

The subsample FQGLS t statistics use the full-sample estimator of the conditional heteroskedasticity { $\hat{\phi}_{n,i} : i \leq n$ }, which is justified because the feasible QGLS and infeasible QGLS t statistics are asymptotically equivalent in the full sample and in subsamples. In addition, the subsample FQGLS t statistics are defined with the fullsample FQGLS estimators  $\hat{\rho}_{GLS,n}$  in place of the null value  $\rho$  in the expression for  $T_{GLS,n}(\rho)$ . That is, the subsample t statistic is of the form:  $b^{1/2}(\hat{\rho}_{GLS,b} - \hat{\rho}_{GLS,n})/\hat{\sigma}_{GLS,b}$ , where  $\hat{\rho}_{GLS,b}$  and  $\hat{\sigma}_{GLS,b}$  are the estimators based on the subsample of size b.

The Sub<sub>*RW*</sub> CI reported in Table A-7 is based on the LS estimator and the homoskedastic variance estimator, as in Romano and Wolf (2001). The Sub<sub>*RW*</sub> critical values are based on subsample statistics that are defined analogously to those for Sub<sub>*GLS*</sub> except that  $\hat{\rho}_{GLS,n}$ ,  $\hat{\rho}_{GLS,b}$ , and  $\hat{\sigma}_{GLS,b}$  are replaced by the full-sample and subsample LS estimators and the subsample homoskedastic standard error estimator, respectively.

The sample size, subsample size, and number of subsamples (for the subsampling and hybrid CI's) employed are 130, 12, and 119. For the  $\text{Sub}_{RW}$  CI we also consider results for subsample sizes b = 8, 16, 20 (with n - b + 1 = 131 - b subsamples in each case).

To mitigate the effect of the initialization on the (G)ARCH processes, we simulate time series of innovations of length 1130 and eliminate the first 1000 observations.

The CHR CP and (CP-corrected) average length results in Tables 4, A-1, A-6, and A-7 are computed in two steps. First, we simulate the asymptotic critical values using 30,000 repetitions, n = 25,000, and standard normal innovations. Then, using these critical values, we simulate the CP's and (CP-corrected) average lengths using 30,000 repetitions and n = 130. To compute CP's, all we need to consider are the true  $\rho$  values of interest: .99, .9, .5, .0, -.9 and one or two quantiles, such as  $c_h(\alpha/2)$  and  $c_h(1-\alpha/2)$ , where  $h=n(1-\rho)$ , for equal-tailed CHR CI's. However, to compute the average lengths we need to determine which values of  $\rho$  are in the CI. To do this, we consider 401 equally spaced grid points for  $\rho$  in [-1, 1] and we determine whether each of these points is in the CI or not. This requires computing the appropriate quantiles for each of the 401  $\rho$  values, such as  $c_h(\alpha/2)$  and  $c_h(1-\alpha/2)$  for  $h = n(1-\rho)$  and n = 130. Furthermore, to carry out CP-correction of the average lengths, we need to determine the value  $\alpha'$  such that the nominal  $1 - \alpha'$  CI has finite-sample CP equal to the desired value  $1 - \alpha$  for the data generating process being considered. To do this, we need to compute the asymptotic critical values not only for one or two quantiles, but rather, for a broad range of potential values of  $1 - \alpha'$ . Hence, when computing the asymptotic critical values in the first step, we consider a grid of 3,200 values of  $100 \times \beta$  (taking values in [20, 99.999] with a step size of .025) and 401 values of  $\rho$  and we compute  $c_h(\beta)$  for  $h = n(1 - \rho)$  for all of these values. Given this 3200 by 401 dimensional matrix of  $c_h(\beta)$  values, we compute the CP's and (CP-corrected) average lengths of the CHR CI in the second step.

For the subsampling and hybrid CI's, we use the same grid of 401  $\rho$  values and 3,200 values of  $100 \times \beta$  when computing the CP-corrected average lengths.

#### 11 Recursive Residual-Based Wild Bootstrap

The recursive residual-based wild bootstrap referred to in Section 2 of the paper is a variant of the grid bootstrap of Hansen (1999) to allow for conditional heterskedasticity. It is defined as follows. For given  $\rho$  (and corresponding  $h = n(1 - \rho)$ ), we desire bootstrap analogues of the quantiles  $c_h(\alpha/2)$  and  $c_h(1 - \alpha/2)$ . Given  $\rho$ , a bootstrap sample of observations indexed by i = 0, ..., n is constructed recursively via  $Y_i^b = \rho Y_{i-1}^b + U_i^b$  with (i)  $U_i^b = \hat{U}_i V_i$ , where  $\hat{U}_i = Y_i - \hat{\rho}_n Y_{i-1}$ ,  $V_i$  has mean 0, variance 1, and finite moments of all orders (e.g.,  $V_i \sim N(0, 1)$ ),  $V_i$  is i.i.d. across  $i, V_i$ is independent of all other variables, and  $\hat{\rho}_n$  is the LS estimator of  $\rho$  based on the original sample, and (ii)  $Y_0^b \sim N(0, \hat{\sigma}_n^2/(1 - \rho^2))$ , where  $\hat{\sigma}_n^2 = n^{-1} \sum_{i=1}^n (Y_i - \overline{Y}_n)^2$ ,  $\overline{Y}_n = n^{-1} \sum_{i=1}^n Y_i$ , and  $Y_0^b$  is independent of all other variables.<sup>7,8</sup>

The bootstrap t statistic is defined just as  $T_n(\rho)$  is defined in (2.1) of the paper, but using the bootstrap sample in place of the original sample. The desired bootstrap quantiles are the  $\alpha/2$  and  $1 - \alpha/2$  quantiles of the distribution of the bootstrap t statistic. These quantiles are computed by using the sample quantiles of the bootstrap t statistics from a large number of bootstrap samples.

The t statistic is asymptotically nuisance parameter free under the null hypothesis and the bootstrap just specified should provide consistent estimators of the quantities that arise in the second term of the Edgeworth expansion of the t statistic. In consequence, we conjecture that this bootstrap version of the CHR CI provides higherorder refinements compared to the CHR CI based on the asymptotic critical values in stationary, unit root, and near unit root scenarios (i.e., the errors in coverage

<sup>&</sup>lt;sup>7</sup>The "wild" nature of this bootstrap is not actually needed for the bootstrap to have correct asymptotic size, but may have better finite sample properties than the "non-wild" version. The "non-wild" version of the bootstrap uses  $U_i^b = V_i$  and  $Y_0^b \sim N(0, 1/(1 - \rho^2))$ .

<sup>&</sup>lt;sup>8</sup>When one is interested in a CI for the sum of the AR coefficients in an AR(k) model, the bootstrap sample is constructed recursively using the augmented Dickey-Fuller (ADF) representation of the model with the specified value  $\rho$  as the sum of the AR coefficients (i.e., the coefficient on  $Y_{i-1}^b$  in the ADF representation) and with the coefficients on the lagged differenced variables in the ADF representation set equal to their LS estimators from the original sample, call them  $\hat{\psi}_1, ..., \hat{\psi}_{k-1}$ . That is,  $\Delta Y_i^b = \rho Y_{i-1}^b + \hat{\psi}_1 \Delta Y_{i-1}^b + ... + \hat{\psi}_{k-1} \Delta Y_{i-k+2}^b + U_i^b$  for i = 1, ..., n, where  $\Delta Y_i^b = Y_i^b - Y_{i-1}^b, Y_0^b$  is as in the AR(1) model,  $\Delta Y_0^b = ... = \Delta Y_{-k+2}^b = 0$  for  $k \ge 2$ , and  $U_i^b$  is as in the AR(1) model, but with  $\hat{U}_i$  given by the *i*th residual from the LS regression of the ADF model.

probabilities of the bootstrap version of the CI converge to zero at a faster rate as  $n \to \infty$ ). A proof of this conjecture is beyond the scope of this paper.

Note that the fixed-design wild bootstrap considered in Gonçalves and Kilian (2004, 2007) in a stationary setting is not appropriate here (i.e., will not provide correct asymptotic size) for two reasons. First, the fixed design does not properly capture the time series properties of the series in the unit root and near unit root contexts. Second, the use of an estimated value  $\rho$  to construct the bootstrap sample does not work properly in the unit root and near unit root contexts. The recursive bootstrap considered in Gonçalves and Kilian (2004) in a stationary setting is not appropriate here for the second reason. The pairs bootstrap considered in Gonçalves and Kilian (2004) in a stationary setting is not appropriate here for the first reason.

When  $\rho = 1$ , the bootstrap procedure outlined above is similar to the bootstrap procedure of Cavaliere and Taylor (2009). It differs in that it employs a stationary start-up, see the definition of  $Y_0^*$  which depends on  $\rho$ . Also note that Cavaliere and Taylor's (2009) unit root test statistic uses a homoskedastic variance estimator which is not appropriate in the context of this paper because a heteroskedasticity-consistent variance matrix estimator is required if the observations are stationary.

#### 12 Proof of Theorem 1

The proof of Theorem 1 relies on Theorem 1 of AG12, which provides the asymptotic distribution of the t statistic under certain drifting sequences of distributions, as specified in (2.5) of the paper. As noted above, the proofs in AG12 need to be adjusted slightly because of the  $p_{ii}^*$  term in the HC5 variance estimator, which does not appear in the variance estimator in AG12. Because  $\max_{i\leq n} p_{ii}^* \leq n^{-1/2} \to 0$ , the adjustment is simple. Theorem 1 of AG12 applies because the restrictions imposed in the definition of  $\Lambda$  include those imposed in Assumption INNOV in AG12 simplified to the case where  $\phi_{n,i} = 1$  in that paper.<sup>9</sup>

The asymptotic results in (2.5) are sufficient to determine the asymptotic size of the CHR CI and to show that it is asymptotically similar using Theorem 2.1 of Andrews, Cheng, and Guggenberger (2009) (ACG).

To describe the result in that paper, using general terminology, let  $\{CS_n : n \ge 1\}$ be a sequence of confidence sets for a parameter  $r(\lambda)$ , where  $\lambda$  indexes the true distribution of the observations. The parameter space for  $\lambda$  is denoted by  $\Lambda$ . Let  $CP_n(\lambda)$  denote the coverage probability of  $CS_n$  under  $\lambda$ . The asymptotic size of  $CS_n$ 

<sup>&</sup>lt;sup>9</sup>Note that Assumption INNOV(v) in AG12 is only needed for the asymptotic results in the case where  $\rho_n \to 1$ . Assumption INNOV(v) in the case where  $\phi_{n,i} = 1$  reduces to the smaller eigenvalue of the 2 × 2 matrix with diagonal elements  $E_{F_n}Y_0^{*2}U_1^2$  and  $E_{F_n}U_1^2$  and off-diagonal element equal to  $E_{F_n}Y_0^*U_1^2$  being larger than  $\delta$  for all n sufficiently large. By Lemma 6 in AG12  $E_{F_n}Y_0^{*2}U_1^2 \to \infty$ and  $E_{F_n}Y_0^*U_1^2 = O(1)$  when  $\rho_n \to 1$ . Assuming  $E_{F_n}U_1^2 \ge \delta$  then ensures that the smaller eigenvalue of the matrix above is not smaller than  $\delta$  for all large enough n. This can be seen by straightforward calculations using l'Hôpital's rule.

is defined as

$$AsySz = \liminf_{n \to \infty} \inf_{\lambda \in \Lambda} CP_n(\lambda).$$
(12.10)

We say a sequence  $\{CS_n : n \ge 1\}$  is asymptotically similar (in a uniform sense) if

$$\liminf_{n \to \infty} \inf_{\lambda \in \Lambda} CP_n(\lambda) = \limsup_{n \to \infty} \sup_{\lambda \in \Lambda} CP_n(\lambda).$$
(12.11)

Corollary 2.1(c) of ACG shows that under Assumptions B1<sup>\*</sup> and B2, stated below, { $CS_n : n \ge 1$ } is asymptotically similar and satisfies AsySz = CP. Let { $h_n(\lambda) : n \ge 1$ } be a sequence of functions on  $\Lambda$ , where  $h_n(\lambda) = (h_{n,1}(\lambda), ..., h_{n,J}(\lambda), h_{n,J+1}(\lambda))'$ ,  $h_{n,j}(\lambda) \in R \ \forall j \le J$ , and  $h_{n,J+1}(\lambda) \in \mathcal{T}$  for some compact pseudo-metric space  $\mathcal{T}$ .

**Assumption B1**<sup>\*</sup>. For any sequence  $\{\lambda_n \in \Lambda : n \geq 1\}$  for which  $h_n(\lambda_n) \to h \in H$ ,  $CP_n(\lambda_n) \to CP$  for some constant  $CP \in [0, 1]$  and some index set H.

Assumption B2. For any subsequence  $\{p_n\}$  of  $\{n\}$  and any sequence  $\{\lambda_{p_n} \in \Lambda : n \geq 1\}$  for which  $h_{p_n}(\lambda_{p_n}) \to h \in H$ , there exists a sequence  $\{\lambda_n^* \in \Lambda : n \geq 1\}$  such that  $h_n(\lambda_n^*) \to h \in H$  and  $\lambda_{p_n}^* = \lambda_{p_n} \ \forall n \geq 1$ .

To prove Theorem 1, it is sufficient to verify Assumptions B1\* and B2 for  $CS_n = CI_{CHR,n}$ . In the present case  $\lambda = (\rho, F)$ ,  $r(\lambda) = \rho$ ,  $h_n(\lambda) = n(1-\rho) \in R$ ,  $H = [0, \infty]$ , and the parameter space  $\Lambda$  is defined in (2.3). Thus, J = 1 and there is no (J + 1)-st component in  $h_n(\lambda)$ . For Assumption B1\*, consider a sequence  $\{\lambda_n = (\rho_n, F_n) \in \Lambda : n \geq 1\}$  for which  $h_n(\lambda_n) \to h \in H$ , i.e.,  $\rho_n = 1 - h_n/n$  and  $h_n \to h \in [0, \infty]$ . We have  $CP_n(\lambda_n) = P_{\lambda_n}(\rho_n \in CI_{CHR,n}) = P_{\lambda_n}(c_{h_n}(\alpha/2) \leq T_n(\rho_n) \leq c_{h_n}(1-\alpha/2))$ . By (2.5) of the paper, we have  $T_n(\rho_n) \to_d J_h$  under  $\{\lambda_n \in \Lambda : n \geq 1\}$ . In addition,  $c_{h_n}(\beta) \to c_h(\beta)$  for  $\beta = \alpha/2$  and  $1-\alpha/2$ . (The latter is proved as follows: Because  $J_h$  is strictly increasing at its  $\beta$ -quantile, for any  $\varepsilon > 0$ ,  $L_n(c_h(\beta) - \varepsilon) \to J_h(c_h(\beta) - \varepsilon) < \beta$  and  $L_n(c_h(\beta) + \varepsilon) \to J_h(c_h(\beta) + \varepsilon) > \beta$ , where  $L_n(x)$  denotes the df of  $T_n(\rho_n)$  at x. This and the definition  $c_{h_n}(\beta) = \inf\{x \in R : L_n(x) \geq \beta\}$  yield  $1\{c_h(\beta) - \varepsilon \leq c_{h_n}(\beta) \leq c_h(\beta) + \varepsilon\} \to 1$  as  $n \to \infty$  for any  $\varepsilon > 0$ .) By the definition of convergence in distribution and continuity of  $J_h$ , it follows that  $CP_n(\lambda_n) \to 1 - \alpha$ . Assumption B1\* therefore holds with  $CP = 1 - \alpha$  for all  $h \in H$ .

For Assumption B2, assume we are given  $\{\lambda_{p_n} \in \Lambda : n \geq 1\}$  for a subsequence  $\{p_n\}$  of  $\{n\}$  such that  $h_{p_n}(\lambda_{p_n}) \to h \in H$ . Define  $\{\lambda_n^* : n \geq 1\}$  by (i)  $\lambda_{p_n}^* = \lambda_{p_n}$  $\forall n \geq 1$ , (ii) when  $h < \infty$  and  $m \neq p_n$ , define  $\lambda_m^* = (1 - h/m, F^*)$ , and (iii) when  $h = \infty$  and  $m \neq p_n$ , define  $\lambda_m^* = (0, F^*)$ , where  $F^*$  is the distribution such that  $\{U_i : i = 0, \pm 1, \pm 2, ...\}$  are i.i.d., standard normal. Then,  $\lambda_n^* \in \Lambda$  for all  $n \geq 1$  and by construction  $h_n(\lambda_n^*) \to h \in H$ . This verifies Assumption B2 and completes the proof.  $\Box$ 

#### 13 Proof of Theorem 2

By the result of Theorem 1 of AG12 stated in (6.1), we have

$$T_n(\rho_n^*) = T_n(\rho_n) + \delta_n / (d_n \widehat{\sigma}_n) \to_d T_h + \delta / S_h$$
(13.12)

under sequences  $\{(\rho_n, F_n) \in \Lambda : n \ge 1\}$  for which  $n(1 - \rho_n) \to h$ .

We show now that  $\{\rho_n^* : n \ge 1\}$  satisfies

$$n(1 - \rho_n^*) \to h + \delta \in [0, \infty].$$
(13.13)

First, suppose  $0 \leq h < \infty$ . Then,  $d_n = n^{1/2}$ ,  $\rho_n^* = \rho_n - \delta_n/n$ , and  $n(1 - \rho_n^*) = n(1 - \rho_n) + \delta_n \to h + \delta \in [0, \infty)$ . Second, suppose  $h = \infty$  and  $\rho_n \to \rho_\infty < 1$ . Then,  $\rho_n^* = \rho_n - \delta_n/(n^{1/2}d_n) \to \rho_\infty < 1$  and  $n(1 - \rho_n^*) \to \infty = h + \delta$ . Third, suppose  $h = \infty$  and  $\rho_n \to 1$ . Then,  $d_n = (1 - \rho_n^2)^{-1/2}(1 + o(1))$  by (6.6). Note that  $n(1 - \rho_n^2) = (1 + \rho_n)n(1 - \rho_n) = 2n(1 - \rho_n)(1 + o(1))$ . Hence, we have

$$n(1 - \rho_n^*) = n(1 - \rho_n) + n^{1/2} \delta_n / d_n$$
  
=  $n(1 - \rho_n) + \delta(n(1 - \rho_n^2))^{1/2} (1 + o(1))$   
=  $n(1 - \rho_n) + \delta 2^{1/2} (n(1 - \rho_n))^{1/2} (1 + o(1))$   
=  $n(1 - \rho_n) [1 + \delta 2^{1/2} (n(1 - \rho_n))^{-1/2} (1 + o(1))]$   
 $\rightarrow \infty = h + \delta.$  (13.14)

Hence, (13.13) is established.

Given (13.13), we have  $c_{n(1-\rho_n^*)}(\beta) \to c_{h+\delta}(\beta)$  for all  $\beta \in (0,1)$  by the same argument as in the proof of Theorem 1 with  $(\rho_n^*, h+\delta)$  in place of  $(\rho_n, h)$ . Using these results and the definition of the CHR CI, we obtain

$$P_{\lambda_n}(\rho_n^* \in CI_{CHR,n})$$

$$= P_{\lambda_n}(c_{n(1-\rho_n^*)}(\alpha/2) \leq T_n(\rho_n^*) \leq c_{n(1-\rho_n^*)}(1-\alpha/2))$$

$$\rightarrow P(c_{h+\delta}(\alpha/2) \leq T_h + \delta/S_h \leq c_{h+\delta}(1-\alpha/2)), \qquad (13.15)$$

where the convergence holds by the definition of convergence in distribution and the continuity of the distribution of  $T_h + \delta/S_h$ .  $\Box$ 

# 14 Asymptotic Validity of Romano and Wolf's (2001) Symmetric Subsampling CI

In this section, we show that the symmetric two-sided subsampling CI of Romano and Wolf (2001) (RW), denoted  $\operatorname{Sub}_{RW}$  above, has asymptotic size equal to its nominal size for the parameter space  $\Lambda$  defined in (2.3), which allows for conditional heteroskedasticity. The derivations below also imply that the lower one-sided version of this CI has correct asymptotic size. The CI in RW is based on a t statistic that employs the LS estimator of  $\rho$ , a homoskedastic standard error estimator, and subsampling critical values.

RW demonstrate that this CI is pointwise asymptotically valid, while Andrews and Guggenberger (2007, Sections 9, 15) show that it has correct asymptotic size for a parameter space that imposes conditional homoskedasticity. (However, the equaltailed two-sided and upper one-sided versions of this CI do not have asymptotically correct size under homoskedasticity or conditional heteroskedasticity, see Mikusheva (2007) and Andrews and Guggenberger (2007).)

Note that AG09a also analyzes a CI based on a t statistic and subsampling critical values, denoted by  $\operatorname{Sub}_{GLS}$  above. They consider a different test statistic than RW. Specifically, they consider a t statistic based on a FQGLS estimator that employs estimators  $\{\widehat{\phi}_{n,i}^2 : i \leq n\}$  of the conditional variances  $\{\sigma_i^2 : i \leq n\}$ , combined with a heteroskedasticity-consistent standard error estimator, see Section 10 above for more details. AG09a proves that the resulting symmetric two-sided subsampling CI has asymptotic size equal to its nominal size for a parameter space that is comparable to  $\Lambda$  but with some additional restrictions on the quantities  $\{\phi_{n,i}^2 : i \leq n\}$  that  $\{\widehat{\phi}_{n,i}^2 : i \leq n\}$  estimate.

The CI in RW is based on a studentized t statistic

$$|T_{Hom,n}(\rho)| = |\frac{n^{1/2}(\widehat{\rho}_n - \rho)}{\widehat{\sigma}_{Hom,n}}|, \qquad (14.16)$$

where  $\hat{\rho}_n$  is the LS estimator defined in (2.2) and  $\hat{\sigma}_{Hom,n}^2$  is the (1,1) element of the standard variance estimator for the LS estimator under the assumption of homoskedasticity. More explicitly,

$$\widehat{\sigma}_{Hom,n}^2 = \left(n^{-1} X_1' M_{X_2} X_1\right)^{-1} \left(n^{-1} Y' M_X Y\right).$$
(14.17)

RW use subsampling critical values, denoted here by  $c_{n,b}(1-\alpha)$ , where *b* denotes the subsample size that satisfies  $b \to \infty$  and  $b/n \to 0$ , and  $1-\alpha$  is the nominal size. The critical value is the  $(1-\alpha)$ -quantile of the empirical distribution of the subsample test statistics over the q = n - b + 1 subsamples of data consisting of *b* consecutive observations from the original data set. The subsample test statistics  $|T_{Hom,n,b,s}(\hat{\rho}_n)|$  for s = 1, ..., q are defined in the same way as the full-sample statistic  $|T_{Hom,n}(\rho)|$  except that only the *b* observations in the *s*-th subsample are used and the hypothesized parameter  $\rho$  is replaced by the full-sample LS estimator  $\hat{\rho}_n$ .

The symmetric two-sided CI in RW is given by the collection of all  $\rho$  ( $\in [-1 + \varepsilon, 1]$  for some  $\varepsilon > 0$ ) that satisfy

$$|T_{Hom,n}(\rho)| \le c_{n,b}(1-\alpha).$$
 (14.18)

Equivalently, the RW CI can be written as

$$[\widehat{\rho}_n - n^{-1/2} c_{n,b} (1-\alpha) \widehat{\sigma}_{Hom,n}, \widehat{\rho}_n + n^{-1/2} c_{n,b} (1-\alpha) \widehat{\sigma}_{Hom,n}].$$

$$(14.19)$$

We now show that this subsampling CI has correct asymptotic size. The proof is quite similar to that for the symmetric subsampling CI based on the FQGLS estimator in Sec. 7 of AG09a and Sec. S10 of Andrews and Guggenberger (2009b) (AG09b). A special case of the FQGLS estimator obtained by taking  $\widehat{\phi}_{n,i} = 1 \quad \forall i \leq n$  is the LS estimator. In this case, the only difference between the test statistics considered in RW and AG09a,b is that the former uses the homoskedasticity variance estimator, whereas the latter uses the standard heteroskedasticity-consistent variance estimator.

The asymptotic size calculations given in AG09a,b depend on the limit as  $n \to \infty$ of CP's of the CI under sequences  $\{\lambda_n = (\rho_n, F_n) \in \Lambda : n \ge 1\}$  for which  $\rho_n = 1 - h_n/n, h_n \to h \in [0, \infty], \rho_n \to \rho_\infty$  for some  $-1 + \varepsilon \le \rho_\infty \le 1, E_{F_n} U_{n,i}^2 \to \sigma_{U,\infty}^2 > 0$ , and  $b_n(1 - \rho_n) \to g \in [0, \infty]$  for  $g \le h$  (where  $b = b_n$  is the subsample size).<sup>10</sup> Provided we show that the limit of the CP's of the RW symmetric two-sided CI is greater than or equal to the nominal size  $1 - \alpha$  for all such sequences, the remainder of the proof of the correct asymptotic size for the RW CI is almost the same as that given in AG09a,b.

When  $\rho_n \to \rho_\infty < 1$ , the subsample and the full-sample t statistic  $|T_{Hom,n}(\rho_n)|$ have the same limiting distribution, a zero mean normal with a sandwich variance expression, and no asymptotic discontinuity arises. Hence, by standard arguments, e.g., see AG09b, the limit of the CP of the RW CI in this case equals the nominal size  $1 - \alpha$ .

Below we show: when  $\rho_n \to 1$  the asymptotic distribution of the RW statistic  $|T_{Hom,n}(\rho_n)|$  is the same as the asymptotic distribution of the AG09a,b t statistic  $|T_n(\rho_n)| = |n^{1/2}(\hat{\rho}_n - \rho_n)/\hat{\sigma}_n|$  (defined in (2.1) of the paper) based on the LS estimator and a heteroskedasticity-consistent variance estimator. Given this, by the arguments in AG09b, the limit of the CP's of the RW CI when  $\rho_n \to 1$  equals that of the AG09a,b CI, which is greater than or equal to the nominal size  $1 - \alpha$ . Hence, the RW CI has correct asymptotic size.

It remains to show the result stated in the previous paragraph. Without loss of generality, we can assume that  $\mu = 0$ , because both  $\hat{\rho}_n - \rho_n$  and  $\hat{\sigma}_{Hom,n}$  are invariant to the choice of  $\mu$ . All limits below are taken as  $n \to \infty$ .

First, suppose  $\rho_n \to 1$  and  $h < \infty$ . By Theorem 1 of AG12, we have

$$n(\hat{\rho}_n - \rho_n) \to_d \frac{\int_0^1 I_{D,h}^*(r) dW(r)}{\int_0^1 I_{D,h}^*(r)^2 dr},$$
(14.20)

where the right-hand side expression uses the fact that the quantities  $h_{2,1}, h_{2,2}, h_{2,5}$ , and  $h_{2,7}$  in AG12 equal  $h_{2,1} = h_{2,2} = \sigma_{U,\infty}^2$  (=  $\lim_{n\to\infty} E_{F_n} U_i^2$ ),  $h_{2,5} = 1$ , and  $h_{2,7} = 1$ when  $\hat{\rho}_n$  is the LS estimator (which corresponds to  $\hat{\phi}_{n,i} = \phi_{n,i} = 1$ ). From eqn. (28) in AG12, it follows that jointly with (14.20) we have

$$n^{-2}X_1'M_{X_2}X_1 \to_d \sigma_{U,\infty}^2 \left(\int_0^1 I_h^*(r)^2 dr - \left(\int_0^1 I_h^*(r) dr\right)^2\right) = \sigma_{U,\infty}^2 \int_0^1 I_{D,h}^*(r)^2 dr.$$
(14.21)

<sup>&</sup>lt;sup>10</sup>In AG09a,b, h and g are denoted by  $h_1$  and  $g_1$ .

Next, Lemma 5(c)-(d) and 5(f)-(h) in AG12 implies that when  $0 < h < \infty$ ,

$$\frac{n^{-1}Y'M_XY}{\sigma_{U,\infty}^2} = \frac{n^{-1}U'M_XU}{\sigma_{U,\infty}^2} = \frac{n^{-1}U'U}{\sigma_{U,\infty}^2} - \frac{n^{-1}U'P_XU}{\sigma_{U,\infty}^2} \to_p 1.$$
 (14.22)

When h = 0, the same result holds by Lemma 5(1) and the arguments in (35) and (36) of AG12 by writing the projection matrix  $P_X$  equivalently as the projection matrix  $P_{X^*}$ , where  $X^* = [X_1 - \overline{Y}_{-1,n}^* \mathbf{1}_n : \mathbf{1}_n]$ , where  $\mathbf{1}_n = (1, ..., 1)' \in \mathbb{R}^n$  and  $\overline{Y}_{-1,n}^* = n^{-1} \sum_{i=1}^n Y_{n,i-1}^* (= n^{-1} \mathbf{1}'_n X_1)$ .

Combining (14.20), (14.21), and (14.22), it follows that the asymptotic distribution of  $|T_{Hom,n}(\rho_n)|$  is

$$\int_{0}^{1} I_{D,h}^{*}(r) dW(r) | / (\int_{0}^{1} I_{D,h}^{*}(r)^{2} dr)^{1/2}, \qquad (14.23)$$

which is the same as that of  $|T_n(\rho_n)|$ , see Theorem 1(a) in AG12 with  $h_{2,7} = 1$  (or (2.6) of the present paper).

Next, suppose  $\rho_n \to 1$  and  $h = \infty$ . By Theorem 1(b) and the definition of  $a_n$  in eqn. (11) in AG12, we have

$$n^{1/2} \frac{E_{F_n} Y_{n,0}^{*2}}{(E_{F_n} Y_{n,0}^{*2} U_{n,1}^2)^{1/2}} (\widehat{\rho}_n - \rho_n) \to_d N(0,1).$$
(14.24)

By (40) of AG12,

$$\frac{n^{-1}X_1'M_{X_2}X_1}{E_{F_n}Y_{n,0}^{*2}} \to_p 1.$$
(14.25)

We also have

$$\frac{n^{-1}Y'M_XY}{\sigma_{U,\infty}^2} = \frac{n^{-1}U'M_XU}{\sigma_{U,\infty}^2} = \frac{n^{-1}U'U}{\sigma_{U,\infty}^2} - \frac{n^{-1}U'P_XU}{\sigma_{U,\infty}^2} \to_p 1$$
(14.26)

by a law of large numbers and

$$n^{-1}U'P_XU \to_p 0. \tag{14.27}$$

When  $\rho_n \to 1$  and  $h = \infty$ , (14.27) holds by the following calculations:

$$n^{-1}U'P_XU = U'X(X'X)^{-1}(n^{-1}X'X)(X'X)^{-1}X'U = v'_n(n^{-1}X'X)v_n = o_p(1),$$
(14.28)

where  $v_n = (X'X)^{-1}X'U = (O_p((1-\rho_n)^{1/2}n^{-1/2}), O_p(n^{-1/2}))'$  by Lemma 8(d) in AG12, the first equality holds because  $P_X = P_X P_X$ , the second equality holds by the definition of  $v_n$ , and the third equality holds by the properties of  $v_n$  and the result that the (1,1), (1,2), and (2,2) components of  $n^{-1}X'X$  are  $O_p((1-\rho_n)^{-1})$ ,

 $O_p((1-\rho_n)^{-1/2})$ , and O(1), respectively, by the first result of Lemma 6 and Lemma 8(a) and (b) in AG12. Hence, (14.26) and (14.27) both hold.

Note that when  $\rho_n \to 1$  and  $h_n \to \infty$ , then from the first two results in Lemma 6 of AG12, we have

$$\frac{E_{F_n}Y_{n,0}^{*2}U_{n,1}^2}{E_{F_n}Y_{n,0}^{*2}E_{F_n}U_{n,1}^2} = \frac{(1-\rho_n^2)^{-1}(E_{F_n}U_{n,1}^2)^2 + O(1)}{((1-\rho_n^2)^{-1}(E_{F_n}U_{n,1}^2) + O(1))(E_{F_n}U_{n,1}^2)} \\
= \frac{(E_{F_n}U_{n,1}^2)^2 + O(1)}{((E_{F_n}U_{n,1}^2) + O(1))(E_{F_n}U_{n,1}^2)} \\
= 1 + o(1).$$
(14.29)

Combining (14.24), (14.25), (14.26), and (14.29), it follows that the asymptotic distribution |N(0,1)| of  $|T_{Hom,n}(\rho_n)|$  is the same as that of  $|T_n(\rho_n)|$ , see Theorem 1(b) in AG12. This completes the proof.

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