Robust Standard Errors in Small Samples

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Description

This package implements small-sample degrees of freedom adjustments to robust and cluster-robust standard errors in linear regression, as discussed in Imbens and Kolesár [2016]. The implementation can handle models with fixed effects, and cases with a large number of observations or clusters ¹.

```
library(dfadjust)
```

To give some examples, let us construct an artificial dataset with 11 clusters

Let us first run a regression of y on x1. This is a case in which, in spite of moderate data size, the effective number of observations is small since there are only three treated units:

```
r1 <- lm(y ~ x1, data = d1)

## No clustering

dfadjustSE(r1)

#>

#> Coefficients:

#> Estimate HC1 se HC2 se Adj. se df p-value

#> (Intercept) 0.00266 0.0311 0.031 0.0311 996.00 0.932

#> x1 0.12940 0.8892 1.088 2.3743 2.01 0.916
```

We can see that the usual robust standard errors (HC1 se) are much smaller than the effective

¹We thank Ulrich Müller for suggesting to us the lemma below.

standard errors (Adj. se), which are computed by taking the HC2 standard errors and applying a degrees of freedom adjustment.

Now consider a cluster-robust regression of y on x2. There are only 3 treated clusters, so the effective number of observations is again small:

```
r1 <- lm(y ~ x2, data = d1)

# Default Imbens-Kolesár method
dfadjustSE(r1, clustervar = d1$cl)

#>

#> Coefficients:

#> Estimate HC1 se HC2 se Adj. se df p-value

#> (Intercept) -0.0236 0.0135 0.0169 0.0222 4.94 0.2215

#> x2 0.1778 0.0530 0.0621 0.1157 2.43 0.0826

# Bell-McCaffrey method
dfadjustSE(r1, clustervar = d1$cl, IK = FALSE)

#>

#> Estimate HC1 se HC2 se Adj. se df p-value

#> (Intercept) -0.0236 0.0135 0.0169 0.0316 2.42 0.2766

#> x2 0.1778 0.0530 0.0621 0.1076 2.70 0.0731
```

Now, let us run a regression of y on x3, with fixed effects. Since we're only interested in x3, we specify that we only want inference on the second element:

Finally, an example in which the clusters are large. We have 500,000 observations:

```
d2 <- do.call("rbind", replicate(500, d1, simplify = FALSE))
d2$y <- rnorm(length(d2$y))
r2 <- lm(y ~ x2, data = d2)
summary(r2)
#>
#> Call:
#> lm(formula = y ~ x2, data = d2)
#>
#> Residuals:
```

```
#> Min 1Q Median
                       3 Q
#> -5.073 -0.675 0.000 0.675 4.789
#>
#> Coefficients:
               Estimate Std. Error t value Pr(>|t|)
#> (Intercept) -0.000991
                          0.001535 -0.65
                                             0.52
             -0.003590
                          0.003963 -0.91
                                              0.37
#>
#> Residual standard error: 1 on 499998 degrees of freedom
#> Multiple R-squared: 1.64e-06, Adjusted R-squared: -3.59e-07
\#> F-statistic: 0.821 on 1 and 5e+05 DF, p-value: 0.365
# Default Imbens-Kolesár method
dfadjustSE(r2, clustervar = d2$c1)
#>
#> Coefficients:
#>
               Estimate HC1 se HC2 se Adj. se
#> (Intercept) -0.000991 0.00133 0.00168 0.00294 2.66
                                                      0.603
              -0.003590 0.00483 0.00568 0.00997 2.65
# Bell-McCaffrey method
dfadjustSE(r2, clustervar = d2$c1, IK = FALSE)
#>
#> Coefficients:
              Estimate HC1 se HC2 se Adj. se
                                                 df p-value
#> (Intercept) -0.000991 0.00133 0.00168 0.00315 2.42
                                                      0.607
            -0.003590 0.00483 0.00568 0.00984 2.70
```

Methods

This section describes the implementation of the Imbens and Kolesár [2016] and Bell and McCaffrey [2002] degrees of freedom adjustments.

There are S clusters, and we observe n_s observations in cluster s, for a total of $n = \sum_{s=1}^{S} n_s$ observations. We handle the case with independent observations by letting each observation be in its own cluster, with S = n. Consider the linear regression of a scalar outcome Y_i onto a p-vector of regressors X_i ,

$$Y_i = X_i'\beta + u_i, \qquad E[u_i \mid X_i] = 0.$$

We're interested in inference on $\ell'\beta$ for some fixed vector $\ell \in \mathbb{R}^p$. Let X, u, and Y denote the design matrix, and error and outcome vectors, respectively. For any $n \times k$ matrix M, let M_s denote the $n_s \times k$ block corresponding to cluster s, so that, for instance, Y_s corresponds to the outcome vector in cluster s. For a positive semi-definite matrix M, let $M^{1/2}$ be a matrix satisfying $M^{1/2}M^{1/2} = M$, such as its symmetric square root or its Cholesky decomposition.

Assume that

$$E[u_s u_s' \mid X] = \Omega_s$$
, and $E[u_s u_t' \mid X] = 0$ if $s \neq t$.

Denote the conditional variance matrix of u by Ω , so that Ω_s is the block of Ω corresponding to cluster s. We estimate $\ell'\beta$ using OLS. In R, the OLS estimator is computed via a QR decomposition,

X = QR, where Q'Q = I and R is upper-triangular, so we can write the estimator as

$$\ell'\hat{\beta} = \ell'\left(\sum_s X_s'X_s\right)^{-1} \sum_s X_sY_s = \tilde{\ell}'\sum_s Q_s'Y_s, \qquad \tilde{\ell} = R^{-1'}\ell.$$

It has variance

$$V := \operatorname{var}(\ell'\hat{\beta} \mid X) = \ell' \left(X'X \right)^{-1} \sum_{s} X'_{s} \Omega_{s} X_{s} \left(X'X \right)^{-1} \ell = \tilde{\ell}' \sum_{s} Q'_{s} \Omega_{s} Q_{s} \tilde{\ell}.$$

Variance estimate

We estimate *V* using a variance estimator that generalizes the HC2 variance estimator to clustering. Relative to the LZ2 estimator described in Imbens and Kolesár [2016], we use a slight modification that allows for fixed effects:

$$\hat{V} = \ell'(X'X)^{-1} \sum_{s} X'_{s} A_{s} \hat{u}_{s} \hat{u}'_{s} A'_{s} X_{s} (X'X)^{-1} \ell = \ell' R^{-1} \sum_{s} Q'_{s} A_{s} \hat{u}_{s} \hat{u}'_{s} A'_{s} Q_{s} R'^{-1} \ell = \sum_{s=1}^{S} (\hat{u}'_{s} a_{s})^{2},$$

where

$$\hat{u}_s := Y_s - X_s \hat{\beta} = u_s - Q_s Q' u, \qquad a_s = A'_s Q_s \tilde{\ell},$$

and the matrix A_s is given by the symmetric square root of the inverse of $I - Q_s Q_s'$, or else its pseudo-inverse if it is singular, as is the case, for example, if X contains fixed effects. We do not need to insist on $I - Q_s Q_s'$ to be invertible, since, using the identity

$$\hat{V} = u \sum_{s} (I - QQ')'_{s} a_{s} a'_{s} (I - QQ')_{s} u,$$

one can verify by simple algebra that a sufficient condition for \hat{V} to be unbiased under homoskedasticity is that $Q'_sA_s(I-Q_sQ'_s)A_sQ_s=Q'_sQ_s$ (see, for example, Pustejovsky and Tipton [2018], for details).

If the observations are independent, the vector of leverages $(Q_1'Q_1,\ldots,Q_n'Q_n)$ can be computed directly using the stats::hatvalues function. In this case, use this function to compute $A_i = 1/\sqrt{1-Q_i'Q_i}$ directly, and we then compute $a_i = A_iQ_i'\tilde{\ell}$ using vector operations. For the case with clustering, computing the spectral decomposition of $I-Q_sQ_s'$ can be expensive or even infeasible if the cluster size n_s is large. We therefore use the following result, suggested to us by Ulrich Müller, which allows us to compute a_s by computing a spectral decomposition of a $p \times p$ matrix.

• Let $Q_s'Q_s = \sum_{i=1}^p \lambda_{is}r_{is}r_{is}'$ be the spectral decomposition of $Q_s'Q_s$. Then $A_s = \sum_{i: \lambda_i \neq 1} (1 - \lambda_i)^{-1/2}Q_sr_{is}r_{is}'Q_s'$, satisfies $A_s(I - Q_sQ_s')A_s = I$.

This follows from the fact that $I - Q_s Q_s'$ has eigenvalues $1 - \lambda_{is}$ and eigenvectors $Q_s r_{is}$, and hence its pseudoinverse is $\sum_{i: \lambda_i \neq 1} (1 - \lambda_i)^{-1} Q_s r_{is} r_{is}' Q_s'$.

Using the lemma, we can compute a_s efficiently as:

$$a_s = \sum_{i: \lambda_i \neq 1} (1 - \lambda_i)^{-1/2} Q_s r_{is} r'_{is} Q'_s Q_s \tilde{\ell} = Q_s D_s \tilde{\ell}, \qquad D_s = \sum_{i: \lambda_i \neq 1} \lambda_i (1 - \lambda_i)^{-1/2} r_{is} r'_{is}.$$

Degrees of freedom correction

Let *G* be an $n \times S$ matrix with columns $(I - QQ')'_s a_s$. Then the Bell and McCaffrey [2002] adjustment sets the degrees of freedom to

$$f_{\rm BM} = \frac{\operatorname{tr}(G'G)^2}{\operatorname{tr}((G'G)^2)}.$$

Since $(G'G)_{st} = a'_s(I - QQ')_s(I - QQ)'_t a_t = a_s(\mathbb{1}\{s = t\} - Q_sQ'_t)a_t$, the matrix G'G can be efficiently computed as

$$G'G = \operatorname{diag}(a'_s a_s) - BB'$$
 $B_{sk} = a'_s Q_{sk}$.

Note that *B* is an $S \times p$ matrix, so that computing the degrees of freedom adjustment only involves $p \times p$ matrices:

$$f_{\text{BM}} = \frac{(\sum_{s} a'_{s} a_{s} - \sum_{s,k} B_{sk}^{2})^{2}}{\sum_{s} (a'_{s} a_{s})^{2} - 2\sum_{s,k} (a'_{s} a_{s}) B_{sk}^{2} + \sum_{s,t} (B'_{s} B_{t})^{2}}.$$

If the observations are independent, we compute B directly as B < -a*Q, and since a_i is a scalar, we have

$$f_{\text{BM}} = \frac{(\sum_{i} a_i^2 - \sum_{sk} B_{sk}^2)^2}{\sum_{i} a_i^4 - 2\sum_{i} a_i^2 B_i' B_i + \sum_{i,j} (B_i' B_j)^2}.$$

The Imbens and Kolesár [2016] degrees of freedom adjustment instead sets

$$f_{IK} = \frac{\operatorname{tr}(G'\hat{\Omega}G)^2}{\operatorname{tr}((G'\hat{\Omega}G)^2)},$$

where $\hat{\Omega}$ is an estimate of the Moulton [1986] model of the covariance matrix, under which $\Omega_s = \sigma_{\epsilon}^2 I_{n_s} + \rho \iota_{n_s} \iota'_{n_{\epsilon}}$. Using simple algebra, one can show that in this case,

$$G'\Omega G = \sigma_{\epsilon}^2 \operatorname{diag}(a_s'a_s) - \sigma_{\epsilon}^2 BB' + \rho(D - BF')(D - BF')',$$

where

$$F_{sk} = \iota'_{n_s} Q_{sk}, \qquad D = \operatorname{diag}(a'_s \iota_{n_s})$$

which can again be computed even if the clusters are large. The estimate $\hat{\Omega}$ replaces σ_{ϵ}^2 and ρ with analog estimates.

References

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