

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

```
set.seed(7)
d1 <- data.frame(y = rnorm(1000), x1 = c(rep(1, 3), rep(0,
  997)), x2 = c(rep(1, 150), rep(0, 850)), x3 = rnorm(1000),
  cl = as.factor(c(rep(1:10, each = 50), rep(11, 500))))
```

Let us first run a regression of y on x_1 . 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  0.957
```

We can see that the usual robust standard errors (HC1 se) are much smaller than the effective 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 x_2 . There are only 3 treated clusters, so the effective number of observations is again small:

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

```

r1 <- lm(y ~ x2, data = d1)
# Default Imbens-Kolesár method
dfadjustSE(r1, clustervar = d1$c1)
#>
#> Coefficients:
#>           Estimate HC1 se HC2 se Adj. se    df p-value
#> (Intercept) -0.0236 0.0135 0.0169 0.0222 4.94 0.288
#> x2          0.1778 0.0530 0.0621 0.1157 2.43 0.124
# Bell-McCaffrey method
dfadjustSE(r1, clustervar = d1$c1, IK = FALSE)
#>
#> Coefficients:
#>           Estimate HC1 se HC2 se Adj. se    df p-value
#> (Intercept) -0.0236 0.0135 0.0169 0.0316 2.42 0.4547
#> x2          0.1778 0.0530 0.0621 0.1076 2.70 0.0983

```

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:

```

r1 <- lm(y ~ x3 + c1, data = d1)
dfadjustSE(r1, clustervar = d1$c1, ell = c(0, 1, rep(0,
  r1$rank - 2)))
#>
#> Coefficients:
#>           Estimate HC1 se HC2 se Adj. se    df p-value
#> Estimate      0.0261 0.0463 0.0595 0.0928 3.23 0.778
dfadjustSE(r1, clustervar = d1$c1, ell = c(0, 1, rep(0,
  r1$rank - 2)), IK = FALSE)
#>
#> Coefficients:
#>           Estimate HC1 se HC2 se Adj. se    df p-value
#> Estimate      0.0261 0.0463 0.0595 0.0928 3.23 0.778

```

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       3Q      Max
#> -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
#> x2          -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$c1)
#>
#> Coefficients:
#>              Estimate HC1 se  HC2 se Adj. se   df p-value
#> (Intercept) -0.000991 0.00133 0.00205 0.00261 5.50   0.704
#> x2          -0.003590 0.00483 0.00376 0.00554 3.64   0.517
# Bell-McCaffrey method
dfadjustSE(r2, clustervar = d2$c1, IK = FALSE)
#>
#> Coefficients:
#>              Estimate HC1 se  HC2 se Adj. se   df p-value
#> (Intercept) -0.000991 0.00133 0.00205 0.00267 5.10   0.710
#> x2          -0.003590 0.00483 0.00376 0.00554 3.64   0.517

```

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 ,

$$Y_i = X_i' \beta + u_i, \quad E[u_i | X_i] = 0.$$

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

$$E[u_s u_s' | X] = \Omega_s, \quad \text{and} \quad E[u_s u_t' | X] = 0 \quad \text{if } s \neq t.$$

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

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

It has variance

$$V := \text{var}(\ell' \hat{\beta} \mid X) = \ell' (X'X)^{-1} \sum_s X'_s \Omega_s X_s (X'X)^{-1} \ell = \tilde{\ell}' \sum_s Q'_s \Omega_s Q_s \tilde{\ell}.$$

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

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

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

$$\hat{V} = u \sum_s (I - Q Q')'_s a_s a'_s (I - Q Q')_s u,$$

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

If the observations are independent, the vector of leverages $(Q'_1 Q_1, \dots, Q'_n Q_n)$ 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 $a_i = A_i Q'_i \tilde{\ell}$ using vector operations. For the case with clustering, computing the spectral decomposition of $I - Q_s Q'_s$ 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, 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 $A_s = \sum_{i: \lambda_i \neq 1} (1 - \lambda_i)^{-1/2} Q_s r_{is} r'_{is} Q'_s$, satisfies $A_s (I - Q_s Q'_s) A_s = I$.

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:

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

Degrees of freedom correction

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

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

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

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

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 $B \leftarrow a * Q$, and since a_i is a scalar, we have

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

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

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

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

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

where

$$F_{sk} = l'_{n_s} Q_{sk}, \quad D = \text{diag}(a'_s l_{n_s})$$

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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