

```

#~~~~~#
# Code to accompany Jones & Waller: The Normal-Theory #
# and Asymptotic Distribution-Free (ADF) Covariance #
# Matrix of Standardized Regression Coefficients: #
# Theoretical Extensions and Finite Sample Behavior #
#
# This function uses the delta method to construct #
# normal-theory and ADF confidence intervals for #
# standardized regression coefficients. #
#
# Arguments: #
# X           - matrix of predictor scores #
# y           - vector of criterion scores #
# cov.x       - covariance matrix for predictors #
# cov.xy      - vector of covariances between #
#                  predictors and criterion #
# var.y       - criterion variance #
# Nobs        - number of observations #
# alpha       - desired Type I error rate #
#                  default = .05 #
# ADF         - Logical (TRUE/FALSE) to select ADF #
#                  confidence intervals - requires raw X #
#                  and y; default = TRUE #
# digits      - number of significant digits to #
#                  print; default = 3 #
#
# This function accepts either (1) raw data, or (2) #
# second-order moments (covariances) and sample size. #
#
# Output #
# cov.mat    - normal-theory or ADF covariance matrix of #
#                  standardized regression coefficients #
# SEs         - normal-theory or ADF standard errors for #
#                  standardized regression coefficients #
# alpha       - desired Type I error rate #
# CIs         - normal-theory or ADF confidence intervals #
#                  for standardized regression coefficients #
#~~~~~#
seBeta <- function(X = NULL, y = NULL,
                    cov.x = NULL, cov.xy = NULL,
                    var.y = NULL, Nobs = NULL,
                    alpha = .05, ADF = TRUE,
                    digits = 3) {
#~~~~~#Internal
Functions#~~~~~#
# Computes the ADF covariance matrix of covariances
adfCOV <- function(X, y) {

  dev <- scale(cbind(X, y), scale=FALSE)
  nvar <- ncol(dev)
  N <- nrow(dev)

  # number of unique elements in a covariance matrix
  ue <- nvar*(nvar + 1)/2
}

```

```

# container for indices
s <- vector(length=ue, mode="character")
z <- 0
for(i in 1:nvar){
  for(j in i:nvar){
    z <- z + 1
    s[z] <- paste(i, j, sep="")
  }
}

# compute all combinations of elements in s
v <- expand.grid(s, s)

# concatenate index pairs
V <- paste(v[,1], v[,2], sep="")

# separate indices into columns
id.mat <- matrix(0,nrow=ue^2,4)
for(i in 1:4) id.mat[,i] <- as.numeric(sapply(V, substr, i, i))

# fill a matrix, M, with sequence 1:ue^2 by row;
# use M to locate positions of indices in id.mat
M <- matrix(1:ue^2, ue, ue, byrow=TRUE)

# select rows of index pairs
r <- M[lower.tri(M, diag=TRUE) ]
ids <- id.mat[r,]
adfCovMat <- matrix(0,ue,ue)
covs <- matrix(0,nrow(ids),1)

# compute ADF covariance matrix using Browne (1984) Eqn 3.8
for(i in 1:nrow(ids)) {
  w_ij <- cov(dev[,ids[i,1]],dev[,ids[i,2]])*((N-1)/N)
  w_kl <- cov(dev[,ids[i,3]],dev[,ids[i,4]])*((N-1)/N)
  w_ik <- cov(dev[,ids[i,1]],dev[,ids[i,3]])*((N-1)/N)
  w_jl <- cov(dev[,ids[i,2]],dev[,ids[i,4]])*((N-1)/N)

  w_i j k l <- (t(dev[,ids[i,1]]*dev[,ids[i,2]]))%*%
    (dev[,ids[i,3]]*dev[,ids[i,4]])/N

  covs[i] <- (N*(N-1)*(1/((N-2)*(N-3)))*(w_i j k l - w_ij*w_kl) -
    N*(1/((N-2)*(N-3)))*(w_ik*w_jl - (2/(N-1))*w_ij*w_kl))
}

# create ADF Covariance Matrix
adfCovMat[lower.tri(adfCovMat,diag=T)] <- covs
vars <- diag(adfCovMat)
adfCovMat <- adfCovMat + t(adfCovMat) - diag(vars)

adfCovMat
} #end adfCOV

# vech function

```

```

vech <- function(x) t(x[!upper.tri(x)])  

  

# Transition or Duplicator Matrix
Dn <- function(x){
  mat <- diag(x)
  index <- seq(x * (x+1) / 2)
  mat[lower.tri(mat, TRUE)] <- index
  mat[upper.tri(mat)] <- t(mat)[upper.tri(mat)]
  outer(c(mat), index, function(x, y) ifelse(x == y, 1, 0))
}
#~~~~~End Define Internal Functions~~~~~#
#~~~~~ Error Checking ~~~~~#
if(is.null(X) & !is.null(y))
  stop("\n y is not defined\n Need to specify both X and y\n")
if(!is.null(X) & is.null(y))
  stop("\n X is not defined\n Need to specify both X and y\n")
if(is.null(X) & is.null(y)) {
  if(is.null(cov.x) | is.null(cov.xy) | is.null(var.y) | is.null(Nobs))
    stop("\nYou need to specify covariances and sample size\n")
  scov <- rbind(cbind(cov.x, cov.xy), c(cov.xy, var.y))
  N <- Nobs
  p <- nrow(cov.x)
}  

} else {
  scov <- cov(cbind(X, y))
  N <- length(y)
  p <- ncol(X)
}  

  

if(ADF) {
  cov.cov <- adfCOV(X, y)
} else {
# create normal-theory covariance matrix of covariances
# See Browne (1984) Eqn 4.6
  Kp.lft <- solve(t(Dn(p + 1)) %*% Dn(p + 1)) %*% t(Dn(p + 1))
  cov.cov <- 2 * Kp.lft %*% (scov %x% scov) %*% t(Kp.lft)
}  

  

param <- c(vech(scov))
ncovs <- length(param)  

  

# find vector element numbers for variances of X
v.x.pl <- c(1, rep(0, p - 1))
for(i in 2:p) v.x.pl[i] <- v.x.pl[i - 1] + p - (i - 2)  

  

# store covariances and variances
cx <- scov[1:p, 1:p]
cxy <- scov[1:p, p+1]
vy <- scov[p+1, p+1]
sx <- sqrt(diag(cx))
sy <- sqrt(vy)
bu <- solve(cx) %*% cxy
ncx <- length(vech(cx))

```

```

# compute derivatives of standardized regression
# coefficients using Yuan and Chan (2011) Equation 13
db <- matrix(0, p, ncovs)
V <- matrix(0, p, ncx)
V[as.matrix(cbind(1:p, v.x.pl))] <- 1

db[, 1:ncx] <- (diag(c(solve(diag(2 * sx * sy)) %*% bu)) %*% V -
                  diag(sx / sy) %*% (t(bu) %*% solve(cx)) %*% Dn(p))

db[, (ncx+1):(ncx+p)] <- diag(sx / sy) %*% solve(cx)
db[,ncovs] <- -diag(sx / (2 * sy^3)) %*% bu

# re-order derivatives
cx.nms <- matrix(0, p, p)
cxy.nms <- c(rep(0, p), "var_y")

for(i in 1:p) for(j in 1:p) cx.nms[i, j] <- paste("cov_x", i, "x", j,
sep=' ')
for(i in 1:p) cxy.nms[i] <- paste("cov_x", i, "y", sep=' ')

old.ord <- c(vech(cx.nms), cxy.nms)
new.ord <- vech(rbind(cbind(cx.nms, cxy.nms[1:p]), c(cxy.nms)))

db <- db[, match(new.ord, old.ord)]

# compute covariance matrix of standardized
# regression coefficients using the Delta Method
DEL.cmat <- db %*% cov.cov %*% t(db) / N
b.nms <- NULL

for(i in 1:p) b.nms[i] <- paste("beta_", i, sep=' ')
rownames(DEL.cmat) <- colnames(DEL.cmat) <- b.nms

# compute standard errors and confidence intervals
DELse <- sqrt(diag(DEL.cmat))
CIs <- as.data.frame(matrix(0, p, 3))
colnames(CIs) <- c("lbound", "estimate", "ubound")
for(i in 1:p) rownames(CIs)[i] <- paste("beta_", i, sep=' ')

tc <- qt(alpha / 2, N - p - 1, lower = F)
beta <- diag(sx) %*% bu * sy^-1
for(i in 1:p) {
  CIs[i,] <- c(beta[i] - tc * DELse[i], beta[i], beta[i] + tc *
  DELse[i])
}
cat("\n", 100 * (1 - alpha),
    "% CIs for Standardized Regression Coefficients:\n\n", sep=' ')
print(round(CIs,digits))

invisible(list(cov.mat=DEL.cmat,SEs=DELse,alpha=alpha,CIs=CIs))
}

```