

R code for ordinal effect size

```
#####  
# R code to find confidence intervals in the unrestricted case for a 2xc table #  
# : Modeling and inference for an ordinal effect size measure #  
# : Authors: Euijung Ryu and Alan Agresti #  
# : Statistics in Medicine (2007) #  
#####
```

```
#####  
#### Data: Shoulder-tip pain scores #####
```

```
y1<-c(19, 2, 1, 0, 0)  
y2<-c( 7, 3, 4, 3, 2)
```

```
# number of columns  
c<-length(y1)
```

```
# matrix A (will be used to find confidence intervals)  
zero<-c(rep(0, c))  
one<-c(rep(1, c))  
J<-one%*%t(one)  
J
```

```
DD<-diag(c)  
A<-J-(0.5)*DD
```

```
for(i in 1:c) {  
  for(j in 1:c) {  
    if(j>i) A[i, j]<-0  
  }  
}  
A
```

```
# for 95% confidence intervals  
alpha<-0.05
```

```
#####  
#### Confidence Intervals for the measure theta #####
```

## R code for ordinal effect size

##### Halperin et al. confidence interval #####

```
Halperin.CI <- function(y1, y2){
  crit.c.Norm <- qnorm(1-alpha/2)
  x1 <- y1
  x2 <- y2
  n1 <- sum(x1)
  n2 <- sum(x2)

  pi.hat <- x1/n1
  lambda.hat <- x2/n2
  theta.hat <- c(lambda.hat)%%A%%pi.hat

  A11 <- t(lambda.hat)%%A%%diag(pi.hat)%%t(A)%%lambda.hat
  B11 <- t(pi.hat)%%t(A)%%diag(lambda.hat)%%A%%pi.hat
  A12 <- A13 <- c(rep(0, c-1))

  for(i in 1:(c-1)){
    A12[i] <- pi.hat[i]*((1-lambda.hat[i]) * sum(lambda.hat[(i+1):c]) - (sum(lambda.hat[(i+1):c]))^2)
    A13[i] <- pi.hat[i]*lambda.hat[i]*(1-lambda.hat[i])
  }
  A1 <- A11 - (1/(n2-1))*sum(A12) - (1/(4*(n2-1)))*(sum(A13)+pi.hat[c]*lambda.hat[c]*(1-lambda.hat[c]))
  B12 <- B13 <- c(rep(0, c-1))

  for(j in 2:c){
    B12[j-1] <- lambda.hat[j]*((1-pi.hat[j])*sum(pi.hat[1:(j-1)])) - (sum(pi.hat[1:(j-1)]))^2)
    B13[j-1] <- lambda.hat[j]*pi.hat[j]*(1-pi.hat[j])
  }
  B1 <- B11 - (1/(n1-1))*sum(B12) - (1/(4*(n1-1)))*(sum(B13)+lambda.hat[1]*pi.hat[1]*(1-pi.hat[1]))
  theta.fn <- ((n1*n2-n1-n2+2)*theta.hat-n1*n2*theta.hat^2)/((n1-1)*(n2-1))+A1/(n1-1)+B1/(n2-1)

  epsilon.hat <- 0
  if((abs(A1)>1e-10) | (abs(B1)>1e-10) | abs(theta.hat)>1e-10){
    if((abs(A1-1)>1e-10) | (abs(B1-1)>1e-10) | abs(theta.hat-1)>1e-10){
      epsilon.hat <- ((n1+n2-2)*theta.hat-(n2-1)*A1-(n1-1)*B1)/((n1+n2-2)*theta.fn)
      a.hat <- epsilon.hat
      if(epsilon.hat<0) a.hat <- 0
      if(epsilon.hat>1) a.hat <- 1
      epsilon.hat <- a.hat
    }
  }
}
```

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```
gam<-(n1+n2-1)-(n1+n2-2)*epsi lon.hat
CCC<-gam*cri ti c.Norm^2/(n1*n2)

L.H<-(CCC+2*theta.hat-sqrt(CCC^2+4*CCC*theta.hat*(1-theta.hat)))/(2*(CCC+1))
U.H<-(CCC+2*theta.hat+sqrt(CCC^2+4*CCC*theta.hat*(1-theta.hat)))/(2*(CCC+1))
H.CI<-c(1,1)*c(L.H, U.H)
li st(theta.hat=theta.hat, Hal peri n.CI =H.CI )
}
```

Hal peri n. CI (y1, y2)

##### Newcombe's pseudo-score confidence interval #####

```
Newcombe.CI <-function(y1, y2){
  cri ti c.Norm<-qnorm(1-alpha/2)
  x1<-y1
  x2<-y2
  n1<-sum(x1)
  n2<-sum(x2)

  pi .hat<-x1/n1
  l am b d a .hat<-x2/n2
  theta.hat<-c(l am b d a .hat)%*%A*%pi .hat

  aa<-(n1+n2)/2-1
  bb<-(cri ti c.Norm^2)/(n1*n2)
  cc<-theta.hat

  c1<- -(1+bb)-2*aa*bb
  c2<-(1+2*bb+2*cc)+4*aa*bb
  c3<-2+2*bb-(bb+2*cc)-cc^2-aa*bb
  c4<-cc^2-2*bb-4*cc-aa*bb
  c5<-2*cc^2

  i ter<-0
  di ff<-1
  rel tol =1e-6
  maxi ter=200
  theta0<-0.01

  whi l e((di ff>rel tol) & (i ter<-i ter+1)<=maxi ter) {
    theta1<-theta0
```

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```

f<-c1*theta1^4+c2*theta1^3+c3*theta1^2+c4*theta1+c5
der. f<-4*c1*theta1^3+3*c2*theta1^2+2*c3*theta1+c4
theta2<-theta1-(1/der. f)*f
di ff<-(theta2-theta1)^2
theta0<-theta2
theta0
di ff
}
left<-theta0

i ter<-0

di ff<-1
rel tol <-1e-6
maxi ter<-200
theta0<-. 99

whi le((di ff>rel tol) & (i ter<=maxi ter)) {
theta1<-theta0
f<-c1*theta1^4+c2*theta1^3+c3*theta1^2+c4*theta1+c5
der. f<-4*c1*theta1^3+3*c2*theta1^2+2*c3*theta1+c4
theta2<-theta1-(1/der. f)*f
di ff<-(theta2-theta1)^2
theta0<-theta2
theta0
di ff
i ter<-i ter+1
}
ri ght<-theta0

CI<-c(1, 1)*c(left, ri ght)
li st(Newcombe. CI =CI)
}
Newcombe. CI (y1, y2)

```

##### Unrestricted Wald-type confidence intervals #####

```

Wald. CI s<-function(y1, y2){
cri ti c. Norm<-qnorm(1-alpha/2)

```

R code for ordinal effect size

```

x1<-y1
x2<-y2
n1<-sum(x1)
n2<-sum(x2)

pi . hat<-x1/n1
l ambda. hat<-x2/n2
theta. hat<-c(l ambda. hat)%**A%**pi . hat

CC<-t(l ambda. hat)%**A%**di ag(pi . hat)%**t(A)%**l ambda. hat
DD<-t(pi . hat)%**t(A)%**di ag(l ambda. hat)%**A%**pi . hat

V. 1<-((n2-1)*(CC-theta. hat^2)+(n1-1)*(DD-theta. hat^2)+theta. hat*(1-theta. hat)-.25*t(pi . hat)%**l ambda. hat)/(n1*n2)
V. 1<-abs(V. 1)
W. 1<-theta. hat+c(-1, 1)*cri ti c. Norm*sqrt(V. 1)

Logi t. 1<-Logi t. 2<-c(0, 1)
i f( abs(theta. hat)>1e-8 & abs(theta. hat-1)>1e-8 ){
  l ogi t<-l og(theta. hat/(1-theta. hat))

  L. 1<-l ogi t-cri ti c. Norm*sqrt(V. 1)/(theta. hat*(1-theta. hat))
  U. 1<-l ogi t+cri ti c. Norm*sqrt(V. 1)/(theta. hat*(1-theta. hat))

  L. theta. W. Logi t. 1<-exp(L. 1)/(1+exp(L. 1))
  U. theta. W. Logi t. 1<-exp(U. 1)/(1+exp(U. 1))

  Logi t. 1<-c(L. theta. W. Logi t. 1, U. theta. W. Logi t. 1)
}

l i st(MLE. theta=theta. hat, Wal d. CI =W. 1, Logi t. Wal d. CI =Logi t. 1)
}

Wal d. CI s(y1, y2)

```

##### Unrestricted LRT, Score, and Pseudo-score confidence intervals #####

```

## To use the following function, we need to get "mph. fi t" functi on
## from Dr. Joseph Lang (j oseph-l ang@ui owa. edu)
## It takes some time to find these confidence intervals
## We can change "starti ng. boundari es" i nsi de the follo wi ng functi on i f you want

```

## R code for ordinal effect size

```

LRT. Score. Pseudo. Score. CI <- function(y1, y2){
  starting.boundaries <- Newcombe.CI(y1, y2)$Newcombe.CI
  theta.set <- seq(starting.boundaries[1]-0.02, starting.boundaries[2]+0.02, length=1000)

  x1 <- y1
  x2 <- y2
  n1 <- sum(x1)
  n2 <- sum(x2)

  pi.hat <- x1/n1
  lambda.hat <- x2/n2
  theta.hat <- c(lambda.hat)%%A%%pi.hat

  L.cal1 <- c(rep(0, c))
  L.cal2 <- c(rep(0, c))
  for(j in 1:c){
    if(x1[j]>0) L.cal1[j] <- x1[j]*log(pi.hat[j])
    if(x2[j]>0) L.cal2[j] <- x2[j]*log(lambda.hat[j])
  }

  L.HA <- sum(L.cal1)+sum(L.cal2)

  yy <- matrix(c(y1, y2), 2*c, 1)
  Z <- ZF <- kronecker(diag(2), matrix(1, c, 1))

  critical.ChiSq <- qchiSq(1-alpha, 1)
  store.LRT <- store.Score <- store.Pseudo.Score <- c(rep(0, length(theta.set)))
  LRT.interval <- Score.interval <- Pseudo.Score.interval <- c(rep(1e+5, length(theta.set)))

  for(k in 1:length(theta.set)){
    theta.null <- theta.set[k]
    cat("index=", k, "theta.value=", theta.null, "\n")

    h.fct <- function(m){
      p <- diag(c(1/(Z%%t(Z)%%m)))%%m
      t(p[(c+1):(2*c)])%%A%%p[1:c]-theta.null
    }

    if( (1-theta.hat)>.1 & (theta.hat >.1) ) {
      a <- mph.fit(yy, Z, ZF, h.fct=h.fct, norm.diff.conv=10, maxiter=200, norm.score.conv=1e-5)
    }

    if( (1-theta.hat)>.1 & (theta.hat <=.1) ) {
      a <- mph.fit(yy, Z, ZF, h.fct=h.fct, norm.diff.conv=10, maxiter=200, norm.score.conv=1e-5, y.eps=.1)
    }
  }
}

```

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```

if( (1-theta.hat)<= .1 & ( theta.hat > .1)) {
  a<-mph. fit(yy, Z, ZF, h. fct=h. fct, norm. di ff. conv=10, maxi ter=200, norm. score. conv=1e-5, y. eps=. 1)
}

if( (1-theta.hat)<= .1 & ( theta.hat <= .1 )){
  a<-mph. fit(yy, Z, ZF, h. fct=h. fct, norm. di ff. conv=10, maxi ter=200, norm. score. conv=1e-5, y. eps=. 1)
}

lang<-a$p
pi. fi nal <-lang[1: c]
lambda. fi nal <-lang[(c+1): (2*c)]
theta. fi nal <-t(lambda. fi nal )%%A%%pi. fi nal

l rt<-score<-pseudo. score<-1e10

if(abs(theta. fi nal - theta. nul l) <1e-5){
  L. cal 10<-c(rep(0, c))
  L. cal 20<-c(rep(0, c))
  for(j in 1:c){
    if(x1[j ]>0) L. cal 10[j ]<-x1[j ]*l og(pi. fi nal [j ])
    if(x2[j ]>0) L. cal 20[j ]<-x2[j ]*l og(lambda. fi nal [j ])
  }

  L. H0<-sum(L. cal 10)+sum(L. cal 20)
  l rt<- -2*(L. H0-L. HA)
  score<-a$Xsq

  CC<-t(lambda. fi nal )%%A%%di ag(pi. fi nal )%%t(A)%%lambda. fi nal
  DD<-t(pi. fi nal )%%t(A)%%di ag(lambda. fi nal )%%A%%pi. fi nal

var. theta. fi nal <-((n2-1)*(CC-theta. nul l ^2)+(n1-1)*(DD-theta. nul l ^2)+theta. nul l *(1-theta. nul l )-. 25*t(pi. fi nal )%%lambda. fi nal )/(n1
*n2)

  pseudo. score<-(theta. hat-theta. nul l )^2/var. theta. fi nal
}

cat("LRT. stat=", l rt, "Score. stat=", score, "Pseudo. Score. stat=", pseudo. score, "\n")

store. LRT[k]<-l rt
store. Score[k]<-score
store. Pseudo. Score[k]<-pseudo. score

if(l rt<cri ti c. Chi sq) {LRT. i nterval [k]<-theta. nul l }
if(score<cri ti c. Chi sq) {Score. i nterval [k]<-theta. nul l }
if(pseudo. score<cri ti c. Chi sq) {Pseudo. Score. i nterval [k]<-theta. nul l }
}

```

R code for ordinal effect size

```
LRT.set<-LRT.interval [LRT.interval <1e+5]  
LRT.CI<-cbind(mi n(LRT.set), max(LRT.set))
```

```
Score.set<-Score.interval [Score.interval <1e+5]  
Score.CI<-cbind(mi n(Score.set), max(Score.set))
```

```
Pseudo.Score.set<-Pseudo.Score.interval [Pseudo.Score.interval <1e+5]  
Pseudo.Score.CI<-cbind(mi n(Pseudo.Score.set), max(Pseudo.Score.set))
```

```
l i st(LRT.CI=LRT.CI, Score.CI=Score.CI, Pseudo.Score.CI=Pseudo.Score.CI)
```

```
}
```

```
LRT.Score.Pseudo.Score.CI s(y1, y2)
```