

## Overall Effect Size Calculation

```
library(meta)
library(metafor)
# Loading
library("readxl")
# xls files
data <- read_excel("Revised_database_for_R.xlsx")

# Random-effects meta-analysis with pre-calculated estimates (SMD) and using
DerSimonian and Laird method

m.hksj <- metagen(data$SMD,
                  data$se,
                  data=data,
                  studlab=paste(data$Study),
                  comb.fixed = FALSE,
                  comb.random = TRUE,
                  method.tau = "DL",
                  hakn = TRUE,
                  prediction=TRUE,
                  sm="SMD")
```

## Subgroup Analyses

```
## Example for Households vs. Individuals
x.sub1 <- subset(data, data$Households==1)
x.sub2 <- subset(data, data$Households==0)

m.sub1 <- metagen(x.sub1$SMD,
                  x.sub1$se,
                  data=x.sub1,
                  studlab=paste(x.sub1$Study),
                  comb.fixed = FALSE,
                  comb.random = TRUE,
                  method.tau = "DL",
                  hakn = TRUE,
                  prediction=TRUE,
                  sm="SMD")

m.sub2 <- metagen(x.sub2$SMD,
                  x.sub2$se,
                  data=x.sub2,
                  studlab=paste(x.sub2$Study),
                  comb.fixed = FALSE,
                  comb.random = TRUE,
                  method.tau = "DL",
                  hakn = TRUE,
                  prediction=FALSE,
                  sm="SMD")
```

## Funnel Plot

```
## Small-study bias
funnel(m.hksj, xlab="Effect estimate",
```

```

                                RCodes
col ="bl ack", cex=1. 5, hl i nes="red", bg=' bl ack' , y l i m=(c(1. 1, 0)),
  contour = c(. 95, . 975, . 99), col . contour=c("darkgray", "gray", "l i ghtgray"))+
  legend(1. 4, 0, c("Studi es", "< 0. 01", "p<0. 025", "p < 0. 05", "p>0. 05" ), bty = "o",
    pch=c(16, NA, NA, NA, NA),
    fill=c("whi te", "l i ghtgray", "gray", "darkgray", "whi te"), bg=' whi te' , border =
"bl ack")

```

## Cumul ative Meta-anal ysi s

```

#Fit a simple random-effects meta-analytic model
random = rma(yi=data$SMD, vi=data$se, method="DL",
test="t", slab=paste(data$Study), tau2="TRUE")
tmp <- cumul (random, order=order(data$Year), transf=TRUE)
forest(tmp, xli m=c(-4, 2), at=(c(-2, -1, -0. 5, -. 1, 0)),
  di gi ts=c(2, 2), cex=0. 5, yaxt="n" )

```