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install.packages("metaSEM")
library(metaSEM)
setwd("E:/R Practice")

My.R<-readLowTriMat("Data-Eleanor.dat", no.var=10)

is.pd(My.R)

Myn <- c(107, 436, 23, 128, 92, 132, 285, 226, 862, 85, 66, 146, 133, 78, 302, 590, 216, 178, 175,
416, 101, 129, 219, 82, 354, 180, 1308, 153, 76, 702, 401, 126, 194, 105, 965, 98, 47, 1002, 25,
220, 497, 168, 253, 41, 34, 527, 70, 351, 227, 209, 208, 82, 145, 126, 102, 59, 71, 136, 180, 145,
38, 277, 138, 170)

pattern.n(My.R, Myn)
pattern.na(My.R, show.na=FALSE)

fixed1 <- tssem1(My.R, Myn, method = "FEM")
summary(fixed1)

fixed2<-tssem1(My.R, Myn, method = "REM", RE.type = "Diag", acov = "weighted")
summary(fixed2)

fixed3 <- tssem1(My.R, Myn, method = "REM", RE.type="Zero")
summary(fixed3)

random1<-tssem1(My.R, Myn, method = "REM")
summary(random1)

random2<-tssem1 (My.R, Myn, method = "REM", RE.type = "Diag", acov = "weighted")
summary(random2)

```