```
library(multcomp)
wk1 <- read.csv("data.csv")   ##data.csv is Appendix S1. In column names, litsiz is litter size, sp is species, and ref is references of original data.
resglm <- glm(litsiz ~ sp, data = wk1, family = "poisson")
mltglm <- glht(resglm, linfct = mcp(sp = "Tukey"))
```