

Main Analysis

Giovani Genesi, MSc / Phillips Lab (UPenn)

2024-06-17

```
# stroop_data <- read.table("https://raw.githubusercontent.com/Lakens/Stroop/master/stroop.txt",  
# sep = "\t", header = TRUE)  
#  
# write.table(stroop_data, file = "stroop.csv", quote = F, row.names = F)  
  
stroop_data <- read.csv("stroop.csv", sep = " ", header = TRUE)
```