

R codes of analysis serum (mice / rat) (pos)

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1 File: serum_pos.R

```
## this script run source of serum_neg stat, to identically
## stat serum pos data
source("~/operation/superstat.R")
file <- "~/operation/batch_serum_pos/batch_serum_pos.csv"
## run script of serum_neg stat
source("~/outline/serum_neg/serum_neg_stat.R")
## the `export` is the stat results
## -----
source("~/outline/serum_neg/serum_neg_stat_metabo.R")
## -----
source("~/outline/serum_neg/serum_neg_stat_metabo_collate.R")
## -----
```