

# R codes of analysis kidney (mice / rat) (neg)

## Contents

1 File: kidney\_neg.R

1

## 1 File: kidney\_neg.R

```
## this script run source of serum_neg stat, to identically  
## stat kidney neg data  
source("~/operation/superstat.R")  
file <- "~/operation/batch_kidney_neg/batch_kidney_neg.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")  
## -----
```