

Simulations

Lauren Quesada

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```
library(tidyverse)

## -- Attaching packages ----- tidyverse 1.3.1 --

## v ggplot2 3.3.5      v purrr 0.3.4
## v tibble 3.1.4       v dplyr 1.0.7
## v tidyr 1.1.3        v stringr 1.4.0
## v readr 2.0.1        v forcats 0.5.1

## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()

library(broom)
LFunky<- function(beta0,beta1,beta2,n_obs,n_data,n_shuffle) {

  set.seed(4747)
  sig<- 0.05

  #If there is no intercept in the fitted regression, d.o.f. = (n-1)
#otherwise, df=n-k-1; k=the number of variables
  dof <- n_obs-1

  #addp_val<-c()

  #intp_val<-c()

  simpp_val<-c()

  simp_t<-c()

  perm_pval<-c()

  for(i in 1:n_data) { # Start of the Monte Carlo loop

    x1<- rnorm(n_obs) # Artificial x series, created just once

    x2<- x1+ rnorm(n_obs) ## Did we figure out why these need to be defined in the loop?
```

```

y<- beta0 + beta1*x1 + beta2*x2 + rnorm(n_obs)    # The DGP includes an intercept

simpfit<- lm(y ~ x1 -1)

#observed p.value for each dataset (n_data)
simpp_val<- c(simpp_val,
              simpfit %>% tidy() %>% filter(term == "x1") %>% select(p.value) %>% pull())

#observed slope for each dataset (n_data)
simp_t<- c(simp_t,
            simpfit %>% tidy() %>% filter(term == "x1") %>% select(statistic) %>% pull())

#obs value for most recent dataset
obsval<- simpfit %>% tidy() %>% filter(term == "x1") %>% select(statistic) %>% pull()

#reassign vector so it doesn't add onto the concatenation each i in 1:n_data loop
simpp_valperm<-c()

  for(j in 1: n_shuffle) {
# A mis-specified model is estimated (unless Beta0 = 0)
#addfit<- lm(y ~ x1+x2 -1)

#intfit<- lm(y ~ x1*x2 -1)

simpfitperm<- lm(y ~ sample(x1) -1)

#addp_val<- c(addp_val, unname(coef(summary(addfit))[, "Pr(>|t|)"]))

#intp_val<- c(intp_val, unname(coef(summary(intfit))[, "Pr(>|t|)"]))

simpp_valperm<- c(simpp_valperm,
                  simpfitperm %>% tidy() %>% filter(term == "sample(x1)") %>% select(statistic) %>%
                  pull())
  }

#p-value for each of my 200 datasets as compared to my permuted distribution of p-values
perm_pval<-c(perm_pval,mean(simpp_valperm>obsval))

} #End of the Monte Carlo Loop

#coefficients, t-test p.values, permutation p.values
data.frame(simp_t, simpp_val, perm_pval)

}

#hist(addp_val, main="Distribution of *ADDITIVE* p-Values", xlab="p-Value", freq=FALSE, border="black")

#hist(intp_val, main="Distribution of *INTERACTIVE* p-Values", xlab="p-Value", freq=FALSE, border="black")

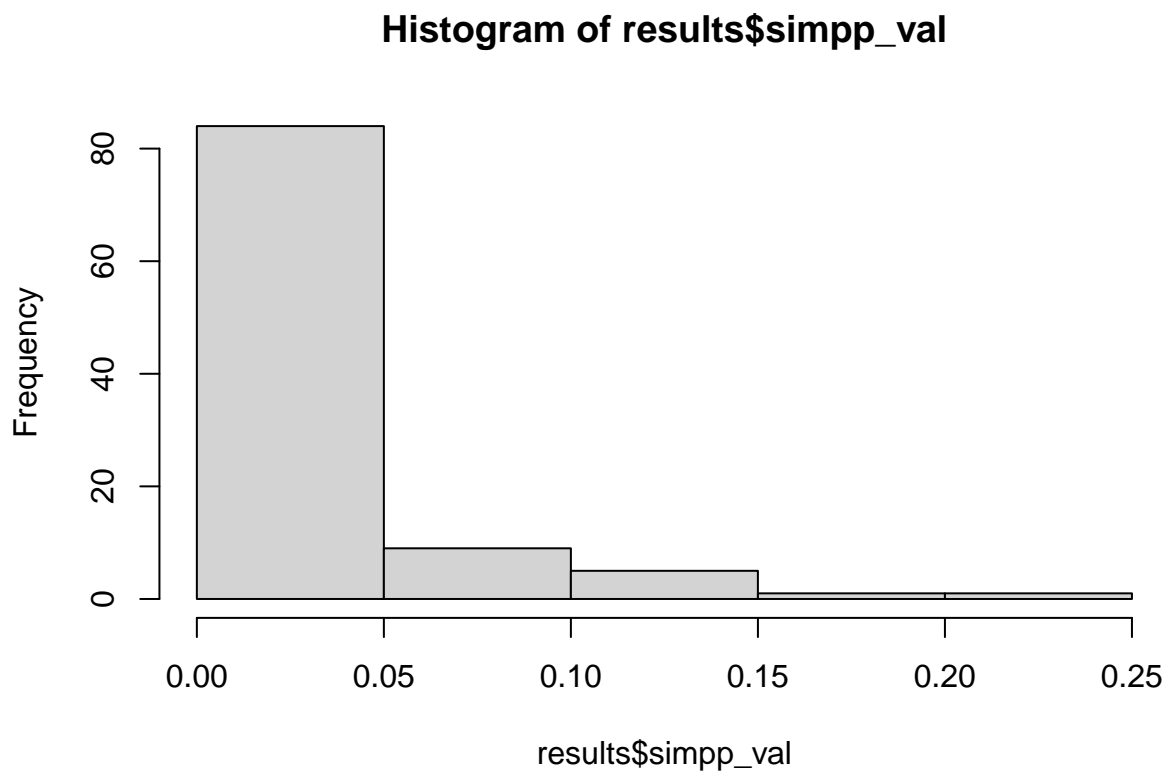
```

```
#hist(simpval, main="Distribution of *SIMPLE* p-Values", xlab="p-Value", freq=FALSE, border="black",
#results %>% ggplot() + geom_hist(aes(x = simpval))
```

First, I'll correctly specify.

1.

```
#beta0,beta1,beta2,n_obs,n_data,n_shuffle
results<-LFunky(0,1,0,15,100,200)
hist(results$simpval)
```



```
hist(results$perm_pval)
```

Histogram of results\$perm_pval

