```
## Load libraries
library(splines)
library(MASS)
library(swfdr)
## Error in library(swfdr): there is no package called 'swfdr'
library(doParallel) ##to make cluster (on Windows)
## Loading required package: foreach
## Loading required package: iterators
## Loading required package: parallel
library(foreach) ##to use foreach function that does the parallel processing
library(doRNG) ##for reproducible seeds when doing parallel processing
## Loading required package: rngtools
## Warning: package 'rngtools' was built under R version 3.3.2
## Loading required package: pkgmaker
## Warning: package 'pkgmaker' was built under R version 3.3.2
## Loading required package: registry
## Warning: package 'registry' was built under R version 3.3.2
## Attaching package: 'pkgmaker'
## The following object is masked from 'package:base':
##
##
      isNamespaceLoaded
##don't need doRNG here, but easier to keep it in
##Source functions
source("../functions.R")
```

Function to pull out means and variances across simulations:

```
pullMeansVars <- function(pi0EstSim)
{
    ##pull out estimates at lambda=0.8, lambda=0.9, and final estimate
    pi0hat0.8 <- sapply(pi0EstSim, function(x){x[[1]]})
    pi0hat0.9 <- sapply(pi0EstSim, function(x){x[[2]]})
    pi0hatFinal <- sapply(pi0EstSim, function(x){x[[3]]})

    ##get means across simulations
    pi0hatMean0.8 <- rowMeans(pi0hat0.8)
    pi0hatMean0.9 <- rowMeans(pi0hat0.9)
    pi0hatMeanFinal <- rowMeans(pi0hatFinal)</pre>
```

Nothing from alternative distribution, since this is for the global null:

## 1 Probability of being a false positive is 0.9

Perform estimation and save estimates:

```
list=c("tme", "pi0", "pi0MeansVars"))
}

## [1] "alt_beta"

## [1] "alt_chisq_large_3_3"

## [1] "alt_chisq_large"

## [1] "alt_t_large"

## [1] "alt_z_large"
```

## 2 Probability of being a false positive is linear

Perform estimation and save estimates:

## 3 Probability of being a false positive is 1

Perform estimation and save estimates:

```
folder <- "global_null"
load(paste(folder, "simResults_0.RData", sep="/"))
splineMat <- ns(tme, df=3)</pre>
```

## Session info:

```
devtools::session_info()
## Session info ------
## setting value
## version R version 3.3.1 (2016-06-21)
##
  system x86_64, mingw32
## ui
          RTerm
## language (EN)
## collate English_United States.1252
## tz America/New_York
## date
         2018-09-18
## Packages -----
## package * version date
                            source
## codetools 0.2-14 2015-07-15 CRAN (R 3.3.1)
## devtools
            1.12.0 2016-06-24 CRAN (R 3.3.3)
## digest 0.6.12 2017-01-27 CRAN (R 3.3.3)
## doParallel * 1.0.10 2015-10-14 CRAN (R 3.3.1)
## doRNG * 1.6 2014-03-07 CRAN (R 3.3.1)
## evaluate
            0.10 2016-10-11 CRAN (R 3.3.1)
## foreach * 1.4.3 2015-10-13 CRAN (R 3.3.1)
            0.6
                    2016-05-09 CRAN (R 3.3.1)
## highr
## iterators * 1.0.8 2015-10-13 CRAN (R 3.3.0)
## knitr * 1.17 2017-08-10 CRAN (R 3.3.3)
            1.5 2014-11-22 CRAN (R 3.3.1)
## magrittr
## MASS
            * 7.3-45 2016-04-21 CRAN (R 3.3.1)
## memoise
            1.0.0 2016-01-29 CRAN (R 3.3.1)
## pkgmaker * 0.22 2014-05-14 CRAN (R 3.3.2)
## registry * 0.3 2015-07-08 CRAN (R 3.3.2)
```

```
## rngtools * 1.2.4 2014-03-06 CRAN (R 3.3.2)

## stringi 1.1.1 2016-05-27 CRAN (R 3.3.0)

## stringr 1.2.0 2017-02-18 CRAN (R 3.3.3)

## withr 1.0.2 2016-06-20 CRAN (R 3.3.1)

## xtable 1.8-2 2016-02-05 CRAN (R 3.3.1)
```