Analysis of BMI GIANT GWAS data - Scott approach

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Load the relevant libraries:

```
library(readr)
library(dplyr)
library(fdrtool)
library(betareg)
library(splines)
library(Hmisc)
library(ggplot2)
library(reshape2)
library(FDRreg)
```

Load the .RData file with the BMI GIANT GWAS meta-analysis data:

```
load("BMI_GIANT_GWAS.RData")

tot <- BMI_GIANT_GWAS</pre>
```

Estimate fraction of true null hypotheses in a regression framework using the Scott approach

Create the design matrix, using natural cubic splines with 5 degrees of freedom to model $\mathbb N$ and 3 discrete categories for the MAFs:

```
X <- model.matrix(~ splines::ns(N,5) + Freq_MAF_Int_Hapmap, data = tot)[,-1]
dim(X)</pre>
```

[1] 2500573 7

```
head(X)
```

```
##
     splines::ns(N, 5)1 splines::ns(N, 5)2 splines::ns(N, 5)3
## 1
           4.414107e-01
                               5.538398e-01
                                                 -0.0017421409
## 2
           0.000000e+00
                               3.954615e-10
                                                 -0.1655612193
## 3
           3.884106e-05
                               9.880678e-01
                                                  0.0099138318
## 4
           3.209714e-04
                               9.891039e-01
                                                  0.0088153851
## 5
           9.327150e-02
                               9.061998e-01
                                                  0.0002901038
## 6
           6.724476e-04
                               9.894590e-01
                                                  0.0082264435
     splines::ns(N, 5)4 splines::ns(N, 5)5 Freq_MAF_Int_Hapmap[0.127,0.302)
##
## 1
           0.0034871548
                             -0.0017450139
                                                                            0
                                                                            0
## 2
           0.3336072837
                              0.8319539352
## 3
           0.0039611703
                              -0.0019816905
                                                                            1
## 4
           0.0035221381
                              -0.0017623939
                                                                            1
## 5
           0.0002264943
                              -0.0001133405
                                                                            1
                                                                            0
## 6
           0.0032867973
                             -0.0016447160
```

```
## Freq_MAF_Int_Hapmap[0.302,0.500]
## 1
## 2
## 3
## 4
## 5
## 6
```

Run code to estimate the fraction of true null hypotheses within a regression framework with the design matrix specified above:

```
##first get z-scores, which are needed for the Scott approach
zScores <- tot$b/tot$se
range(zScores)</pre>
```

[1] -22.41379 26.96667

```
## Warning in doTryCatch(return(expr), name, parentenv, handler): f(z)
## misfit = -0.3. Rerun with increased df.f(z) misfit = -0.7. Rerun with
## increased df.f(z) misfit = 2.3. Rerun with increased df.f(z) misfit = 2.7.
## Rerun with increased df.f(z) misfit = -0.1. Rerun with increased df.f(z)
## misfit = 1. Rerun with increased df.f(z) misfit = 0. Rerun with increased
## df.f(z) misfit = 1. Rerun with increased df.f(z) misfit = 0. Rerun with
## increased df.f(z) misfit = 0. Rerun with increased df.f(z) misfit = 0.
## Rerun with increased df.f(z) misfit = 2. Rerun with increased df.f(z)
## misfit = 1. Rerun with increased df.f(z) misfit = 1. Rerun with increased
## df.f(z) misfit = 0. Rerun with increased df.f(z) misfit = 0. Rerun with
## increased df.f(z) misfit = 0. Rerun with increased df.f(z) misfit = 1.
## Rerun with increased df.f(z) misfit = 0. Rerun with increased df.f(z)
## misfit = 0. Rerun with increased df.f(z) misfit = 2.8. Rerun with increased
## df.f(z) misfit = 36. Rerun with increased df.f(z) misfit = 6.4. Rerun
## with increased df.f(z) misfit = 2.8. Rerun with increased df.f(z) misfit
## = 0.3. Rerun with increased df.f(z) misfit = -2.1. Rerun with increased
## df.f(z) misfit = -2. Rerun with increased df.f(z) misfit = -4.1. Rerun
## with increased df.f(z) misfit = -5.4. Rerun with increased df.f(z) misfit
## = -3.7. Rerun with increased df.f(z) misfit = -6. Rerun with increased
## df.f(z) misfit = -6.3. Rerun with increased df.f(z) misfit = -5.7. Rerun
## with increased df.f(z) misfit = -2.6. Rerun with increased df.f(z) misfit
## = -3.5. Rerun with increased df.f(z) misfit = 0.1. Rerun with increased
## df.f(z) misfit = -0.5. Rerun with increased df.f(z) misfit = 0.6. Rerun
## with increased df.f(z) misfit = 2.7. Rerun with increased df.f(z) misfit
\#\# = 9.8. Rerun with increased df.f(z) misfit = 14.1. Rerun with increased
## df.f(z) misfit = 29.5. Rerun with increased df.f(z) misfit = 29.5. Rerun
## with increased df.f(z) misfit = 39.5. Rerun with increased df.f(z) misfit
## = 50.8. Rerun with increased df.f(z) misfit = 51.5. Rerun with increased
## df.f(z) misfit = 47.8. Rerun with increased df.f(z) misfit = 65.6. Rerun
## with increased df.f(z) misfit = 44.9. Rerun with increased df.f(z) misfit
```

```
## = 34.1. Rerun with increased df.f(z) misfit = 53.9. Rerun with increased
## df.f(z) misfit = 45.8. Rerun with increased df.f(z) misfit = 52.5. Rerun
## with increased df.f(z) misfit = 48.6. Rerun with increased df.f(z) misfit
## = 41.6. Rerun with increased df.f(z) misfit = 35.5. Rerun with increased
## df.f(z) misfit = 25.6. Rerun with increased df.f(z) misfit = 17.1. Rerun
## with increased df.f(z) misfit = -4.2. Rerun with increased df.f(z) misfit
\#\# = -17.6. Rerun with increased df.f(z) misfit = -29.8. Rerun with increased
## df.f(z) misfit = -37.8. Rerun with increased df.f(z) misfit = -30.2. Rerun
## with increased df.f(z) misfit = -16.7. Rerun with increased df.f(z) misfit
## = 4.1. Rerun with increased df.f(z) misfit = 24.4. Rerun with increased
## df.f(z) misfit = 25.1. Rerun with increased df.f(z) misfit = 39.9. Rerun
## with increased df.f(z) misfit = 20. Rerun with increased df.f(z) misfit
\#\#=3.2. Rerun with increased df.f(z) misfit = -12.1. Rerun with increased
## df.f(z) misfit = -39.9. Rerun with increased df.f(z) misfit = -40. Rerun
## with increased df.f(z) misfit = -30.2. Rerun with increased df.f(z) misfit
\#\# = -12.5. Rerun with increased df.f(z) misfit = 5.4. Rerun with increased
## df.f(z) misfit = 27.5. Rerun with increased df.f(z) misfit = 29.9. Rerun
## with increased df.f(z) misfit = 32.5. Rerun with increased df.f(z) misfit
## = 42.2. Rerun with increased df.f(z) misfit = 43.5. Rerun with increased
## df.f(z) misfit = 45.1. Rerun with increased df.f(z) misfit = 41.7. Rerun
## with increased df.f(z) misfit = 38.5. Rerun with increased df.f(z) misfit
## = 36.8. Rerun with increased df.f(z) misfit = 50.4. Rerun with increased
## df.f(z) misfit = 17.9. Rerun with increased df.f(z) misfit = 15.1. Rerun
## with increased df.f(z) misfit = 30. Rerun with increased df.f(z) misfit
\#\# = 42.3. Rerun with increased df.f(z) misfit = 21.6. Rerun with increased
## df.f(z) misfit = 16.8. Rerun with increased df.f(z) misfit = 25. Rerun
## with increased df.f(z) misfit = 8. Rerun with increased df.f(z) misfit
## = 2. Rerun with increased df.f(z) misfit = 3.4. Rerun with increased
## df.f(z) misfit = -0.7. Rerun with increased df.f(z) misfit = -3.5. Rerun
## with increased df.f(z) misfit = -6. Rerun with increased df.f(z) misfit
## = -7.3. Rerun with increased df.f(z) misfit = -7.5. Rerun with increased
## df.f(z) misfit = -6.3. Rerun with increased df.f(z) misfit = -5. Rerun
## with increased df.f(z) misfit = -4.2. Rerun with increased df.f(z) misfit
\#\# = -0.2. Rerun with increased df.f(z) misfit = 1.7. Rerun with increased
## df.f(z) misfit = 1.7. Rerun with increased df.f(z) misfit = 8.2. Rerun
## with increased df.f(z) misfit = 1.9. Rerun with increased df.f(z) misfit
\#\# = 3. Rerun with increased df.f(z) misfit = 2. Rerun with increased df.f(z)
## misfit = 4. Rerun with increased df.f(z) misfit = 5. Rerun with increased
## df.f(z) misfit = 11. Rerun with increased df.f(z) misfit = 4. Rerun with
## increased df.f(z) misfit = 3. Rerun with increased df.f(z) misfit = 0.
## Rerun with increased df.f(z) misfit = 0. Rerun with increased df.f(z)
## misfit = 0. Rerun with increased df.f(z) misfit = 0. Rerun with increased
## df.f(z) misfit = 0. Rerun with increased df.f(z) misfit = 0. Rerun with
## increased df.f(z) misfit = 0. Rerun with increased df.f(z) misfit = 0.
## Rerun with increased df.f(z) misfit = 0. Rerun with increased df.f(z)
## misfit = 0. Rerun with increased df.f(z) misfit = 0. Rerun with increased
## df.f(z) misfit = 0. Rerun with increased df.f(z) misfit = 0. Rerun with
## increased df.f(z) misfit = 2. Rerun with increased df.f(z) misfit = 1.
## Rerun with increased df.f(z) misfit = 0. Rerun with increased df.f(z)
## misfit = 0.9. Rerun with increased df.f(z) misfit = -0.2. Rerun with
## increased df.f(z) misfit = 0.4. Rerun with increased df.f(z) misfit = 0.
## Rerun with increased df.f(z) misfit = -1.1. Rerun with increased df.f(z)
## misfit = 0.1. Rerun with increased df.f(z) misfit = -1.7. Rerun with
## increased df.f(z) misfit = -1.5. Rerun with increased df.f(z) misfit =
```

```
## -0.7. Rerun with increased df.f(z) misfit = -0.2. Rerun with increased
## df.f(z) misfit = 1.4. Rerun with increased df.f(z) misfit = 1.4. Rerun with
## increased df.f(z) misfit = 0.9. Rerun with increased df.f(z) misfit = -0.4.
## Rerun with increased df.f(z) misfit = -0.8. Rerun with increased df.f(z)
## misfit = 0.2. Rerun with increased df.
##get prior probabilities
pi0EstScott <- 1-fdr$priorprob</pre>
Save results:
save(x=pi0EstScott, file="BMI_GIANT_GWAS_results_Scott.RData")
Session Information
devtools::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
           2016-10-28
## Packages -----
               * version date
## package
                                  source
## acepack
                 1.3-3.3 2014-11-24 CRAN (R 3.3.0)
## assertthat
                0.1 2013-12-06 CRAN (R 3.3.1)
## BayesLogit * 0.5.1 2014-07-21 CRAN (R 3.3.0)
               * 3.0-5 2014-09-25 CRAN (R 3.3.1)
## betareg
## BiocStyle
               * 2.0.2
                        2016-05-16 Bioconductor
## chron
                2.3-47 2015-06-24 CRAN (R 3.3.1)
## cluster
                2.0.4
                        2016-04-18 CRAN (R 3.3.1)
## codetools
                0.2-14 2015-07-15 CRAN (R 3.3.1)
## colorspace
                1.2-6
                        2015-03-11 CRAN (R 3.3.1)
## data.table
               1.9.6
                        2015-09-19 CRAN (R 3.3.1)
## DBI
                0.4-1
                        2016-05-08 CRAN (R 3.3.1)
## devtools
                1.12.0 2016-06-24 CRAN (R 3.3.1)
                0.6.9
                        2016-01-08 CRAN (R 3.3.1)
## digest
## dplyr
               * 0.5.0
                        2016-06-24 CRAN (R 3.3.1)
## evaluate
                0.9
                        2016-04-29 CRAN (R 3.3.1)
               * 2.4.4
## fda
                        2014-12-16 CRAN (R 3.3.1)
                        2016-08-30 Github (jgscott/FDRreg@8025d1a)
## FDRreg
               * 0.2-1
```

* 1.2.15 2015-07-08 CRAN (R 3.3.0) 2.3-13 2015-01-17 CRAN (R 3.3.1)

fdrtool

flexmix

```
0.8-66 2015-08-19 CRAN (R 3.3.1)
    foreign
##
    formatR
                   1.4
                           2016-05-09 CRAN (R 3.3.1)
                           2015-04-07 CRAN (R 3.3.0)
##
    Formula
                 * 1.2-1
                           2016-04-27 CRAN (R 3.3.1)
##
    ggdendro
                   0.1-20
    ggplot2
                 * 2.1.0
                           2016-03-01 CRAN (R 3.3.1)
##
    gridExtra
                   2.2.1
                           2016-02-29 CRAN (R 3.3.1)
##
    gtable
                   0.2.0
                           2016-02-26 CRAN (R 3.3.1)
##
    Hmisc
                 * 3.17-4
                           2016-05-02 CRAN (R 3.3.1)
##
    htmltools
                   0.3.5
                            2016-03-21 CRAN (R 3.3.1)
##
    knitr
                   1.13
                           2016-05-09 CRAN (R 3.3.1)
##
    lattice
                 * 0.20-33 2015-07-14 CRAN (R 3.3.1)
##
                   0.6-28 2016-02-09 CRAN (R 3.3.1)
    latticeExtra
                   0.2.0
                           2016-06-12 CRAN (R 3.3.1)
    lazyeval
##
    lmtest
                   0.9 - 34
                           2015-06-06 CRAN (R 3.3.1)
##
    magrittr
                   1.5
                            2014-11-22 CRAN (R 3.3.1)
##
    MASS
                   7.3-45
                           2016-04-21 CRAN (R 3.3.1)
##
    Matrix
                 * 1.2-6
                           2016-05-02 CRAN (R 3.3.1)
                           2016-01-29 CRAN (R 3.3.1)
##
    memoise
                   1.0.0
##
    modeltools
                   0.2-21
                           2013-09-02 CRAN (R 3.3.0)
                           2016-07-29 CRAN (R 3.3.1)
##
    mosaic
                   0.14.4
##
    mosaicData
                   0.14.0
                           2016-06-17 CRAN (R 3.3.1)
##
    munsell
                   0.4.3
                           2016-02-13 CRAN (R 3.3.1)
##
                 * 1.0-5
                           2016-02-02 CRAN (R 3.3.0)
    mvtnorm
##
    nnet
                   7.3-12
                           2016-02-02 CRAN (R 3.3.1)
##
    plyr
                   1.8.4
                           2016-06-08 CRAN (R 3.3.1)
   R6
                   2.1.2
                           2016-01-26 CRAN (R 3.3.1)
##
    RColorBrewer
                   1.1-2
                           2014-12-07 CRAN (R 3.3.0)
##
                   0.12.6
                           2016-07-19 CRAN (R 3.3.1)
    Rcpp
##
                           2015-10-22 CRAN (R 3.3.1)
    readr
                 * 0.2.2
   reshape2
                 * 1.4.1
                           2014-12-06 CRAN (R 3.3.1)
                           2016-07-08 CRAN (R 3.3.1)
##
    rmarkdown
                   1.0
##
    rpart
                   4.1-10
                           2015-06-29 CRAN (R 3.3.1)
##
    sandwich
                   2.3 - 4
                           2015-09-24 CRAN (R 3.3.1)
##
    scales
                   0.4.0
                           2016-02-26 CRAN (R 3.3.1)
                           2016-05-27 CRAN (R 3.3.0)
##
    stringi
                   1.1.1
   stringr
##
                   1.0.0
                           2015-04-30 CRAN (R 3.3.1)
##
    survival
                 * 2.39-4
                           2016-05-11 CRAN (R 3.3.1)
## tibble
                   1.1
                           2016-07-04 CRAN (R 3.3.1)
                           2016-06-14 CRAN (R 3.3.1)
##
   tidyr
                   0.5.1
##
                           2016-06-20 CRAN (R 3.3.1)
    withr
                   1.0.2
    yaml
                   2.1.13
                           2014-06-12 CRAN (R 3.3.1)
                   1.7-13 2016-05-03 CRAN (R 3.3.1)
##
    zoo
```