Analysis of BMI GIANT GWAS data

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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(swfdr)
```

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

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
                                                 -0.1655612193
                               3.954615e-10
## 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
## 2
                                                                            0
           0.3336072837
                              0.8319539352
## 3
           0.0039611703
                              -0.0019816905
                                                                            1
## 4
           0.0035221381
                              -0.0017623939
                                                                            1
## 5
           0.0002264943
                              -0.0001133405
                                                                            1
## 6
                                                                            0
           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:

```
pi0Est <- lm_pi0(pValues=tot$p, X=X, smooth.df=3)</pre>
```

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```

##caution: this should take about an hour!

Save results:

```
fitted0.8 <- pi0Est$pi0.lambda[,pi0Est$lambda==0.8]
fitted0.9 <- pi0Est$pi0.lambda[,round(pi0Est$lambda,2)==0.9]
fitted.final.smooth <- pi0Est$pi0
save(fitted0.8, fitted0.9, fitted.final.smooth, file="BMI_GIANT_GWAS_results.RData")</pre>
```

Session Information

```
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
           America/New_York
## tz
           2016-10-27
## date
## Packages ------
## package
               * version
                           date
                                    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)
## betareg
               * 3.0-5
                           2014-09-25 CRAN (R 3.3.1)
## BiocStyle
               * 2.0.2
                           2016-05-16 Bioconductor
                2.3 - 47
                           2015-06-24 CRAN (R 3.3.1)
## chron
##
   cluster
                2.0.4
                           2016-04-18 CRAN (R 3.3.1)
## codetools
               0.2-14
                          2015-07-15 CRAN (R 3.3.1)
               1.2-6
1.9.6
0.4-1
## colorspace
                          2015-03-11 CRAN (R 3.3.1)
## data.table
                           2015-09-19 CRAN (R 3.3.1)
## DBI
                           2016-05-08 CRAN (R 3.3.1)
## devtools
               1.12.0
                          2016-06-24 CRAN (R 3.3.1)
## digest
                0.6.9
                          2016-01-08 CRAN (R 3.3.1)
                           2016-06-24 CRAN (R 3.3.1)
## dplyr
               * 0.5.0
## evaluate
                0.9
                           2016-04-29 CRAN (R 3.3.1)
## fdrtool
               * 1.2.15
                          2015-07-08 CRAN (R 3.3.0)
## flexmix
                2.3-13
                          2015-01-17 CRAN (R 3.3.1)
## foreign
                0.8-66
                           2015-08-19 CRAN (R 3.3.1)
## formatR
                1.4
                           2016-05-09 CRAN (R 3.3.1)
## Formula
               * 1.2-1
                           2015-04-07 CRAN (R 3.3.0)
## 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
                           2016-02-26 CRAN (R 3.3.1)
                0.2.0
## Hmisc
               * 3.17-4
                           2016-05-02 CRAN (R 3.3.1)
## htmltools
                0.3.5
                           2016-03-21 CRAN (R 3.3.1)
               1.13
## knitr
                           2016-05-09 CRAN (R 3.3.1)
## lattice
               * 0.20-33
                          2015-07-14 CRAN (R 3.3.1)
```

```
## latticeExtra
                  0.6-28
                              2016-02-09 CRAN (R 3.3.1)
                              2015-06-06 CRAN (R 3.3.1)
##
   lmtest
                   0.9 - 34
##
                   1.5
                              2014-11-22 CRAN (R 3.3.1)
   magrittr
## 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)
##
   munsell
                   0.4.3
                              2016-02-13 CRAN (R 3.3.1)
   nnet
                              2016-02-02 CRAN (R 3.3.1)
##
                  7.3-12
##
   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)
## Rcpp
                   0.12.6
                              2016-07-19 CRAN (R 3.3.1)
## readr
                 * 0.2.2
                              2015-10-22 CRAN (R 3.3.1)
## reshape2
                              2014-12-06 CRAN (R 3.3.1)
                 * 1.4.1
## rmarkdown
                   1.0
                              2016-07-08 CRAN (R 3.3.1)
   rpart
                              2015-06-29 CRAN (R 3.3.1)
##
                   4.1-10
##
   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
                              2015-04-30 CRAN (R 3.3.1)
                   1.0.0
##
   stringr
                              2016-05-11 CRAN (R 3.3.1)
## survival
                 * 2.39-4
## swfdr
                 * 0.0.0.9000 2016-10-21 local
## tibble
                   1.1
                              2016-07-04 CRAN (R 3.3.1)
                              2016-06-20 CRAN (R 3.3.1)
##
   withr
                   1.0.2
##
                              2014-06-12 CRAN (R 3.3.1)
   yaml
                  2.1.13
## zoo
                  1.7-13
                              2016-05-03 CRAN (R 3.3.1)
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