

# Package ‘FDRreg’

August 29, 2016

**Type** Package

**Title** False discovery rate regression

**Version** 0.1

**Date** 2014-02-24

**Author** James G. Scott, with contributions from Rob Kass and Jesse Windle

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**Description** Tools for FDR problems, including false discovery rate regression.  
See corresponding paper: “False discovery rate regression: application to neural synchrony detection in primary visual cortex.” James G. Scott, Ryan C. Kelly, Matthew A. Smith, Robert E. Kass.

**License** GPL (>= 3)

**Imports** Rcpp (>= 0.11.0), mosaic (>= 0.8-10)

**Depends** fda (>= 2.4.0), splines (>= 3.0.2)

**LinkingTo** Rcpp, RcppArmadillo

**NeedsCompilation** yes

**Repository** CRAN

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FDRreg-package

*False discovery rate regression*

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## Description

Tools for FDR problems, including false discovery rate regression. Fits models whereby the local false discovery rate may depend upon covariates, either via a linear or additive logistic regression model.

## Details

Package: FDRreg  
Type: Package  
Version: 1.0  
Date: 2014-02-25  
License: GPL (>=3)

The workhouse function is `FDRreg(z,X, ...)`, where `z` is an observed vector of `z` statistics, and `X` is a matrix of covariates. Do not add a column of ones to `X` to get an intercept term; the function does that for you, just like R's base `lm()` and `glm()` functions.

## Author(s)

Author: James G. Scott, with contributions from Rob Kass and Jesse Windle.

Maintainer: James G. Scott <james.scott@mcombs.utexas.edu>

## References

False discovery rate regression: application to neural synchrony detection in primary visual cortex. James G. Scott, Ryan C. Kelly, Matthew A. Smith, Pengcheng Zhou, and Robert E. Kass. arXiv:1307.3495 [stat.ME].

## Examples

```
library(FDRreg)

# Simulated data
P = 2
N = 10000
betatrue = c(-3.5,rep(1/sqrt(P), P))
X = matrix(rnorm(N*P), N,P)
psi = crossprod(t(cbind(1,X)), betatrue)
wsuccess = 1/{1+exp(-psi)}

# Some theta's are signals, most are noise
gammatrue = rbinom(N,1,wsuccess)
```

```

table(gammatrue)

# Density of signals
thetatrue = rnorm(N,3,0.5)
thetatrue[gammatrue==0] = 0
z = rnorm(N, thetatrue, 1)
hist(z, 100, prob=TRUE, col='lightblue', border=NA)
curve(dnorm(x,0,1), add=TRUE, n=1001)

## Not run:
# Fit the model
fdr1 <- FDRreg(z, covars=X, nmc=2500, nburn=100, nmids=120, nulltype='theoretical')

# Show the empirical-Bayes estimate of the mixture density
# and the findings at a specific FDR level
Q = 0.1
plotFDR(fdr1, Q=Q, showfz=TRUE)

# Posterior distribution of the intercept
hist(fdr1$betasave[,1], 20)

# Compare actual versus estimated prior probabilities of being a signal
plot(wsucces, fdr1$priorprob)

# Covariate effects
plot(X[,1], log(fdr1$priorprob/{1-fdr1$priorprob}), ylab='Logit of prior probability')
plot(X[,2], log(fdr1$priorprob/{1-fdr1$priorprob}), ylab='Logit of prior probability')

# Local FDR
plot(z, fdr1$localfdr, ylab='Local false-discovery rate')

# Extract findings at level FDR = Q
myfindings = which(fdr1$FDR <= Q)

## End(Not run)

```

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FDRreg

*False Discovery Rate Regression*


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## Description

Estimate an empirical-Bayes false-discovery rate regression model for test statistics  $z$  and regressors  $X$ .

## Usage

```
FDRreg(z, covars, nulltype = 'empirical', type = 'linear', nmc = 10000, nburn = 500,
nmids = 150, densknots = 10, regknots = 5)
```

**Arguments**

|                        |   |
|------------------------|---|
| <code>z</code>         | An N dimensional vector; $z_i$ is the test statistic for observation $i$ .  |
| <code>covars</code>    | An N x P dimensional design matrix; $x_i$ is the $i$ th row. This is assumed not to have a column of ones representing an intercept; just like in <code>lm()</code> and <code>glm()</code> , this will be added by the fitting algorithm. |
| <code>nulltype</code>  | Choices are 'empirical' for an empirical null using Efron's central-matching method, or 'theoretical' for a standard normal null.   |
| <code>type</code>      | Choices are 'linear' for a standard logistic regression, or 'additive' for an additive logit model, in which case each column of covars is expanded using a b-spline basis.   |
| <code>nmc</code>       | The number of MCMC iterations saved. Defaults to 10,000.  |
| <code>nburn</code>     | The number of initial MCMC iterations discarded as burn-in. Defaults to 500.  |
| <code>nmids</code>     | How many bins should be used in the estimation of the marginal density $f(z)$ ? Defaults to 150.  |
| <code>densknots</code> | How many knots should be used to estimate the marginal density $f(z)$ via spline-based Poisson regression? Defaults to 10; the function will warn you if it looks like you've used too few, using a simple deviance statistic.            |
| <code>regknots</code>  | Used only if <code>type='additive'</code> . How many knots should be used to estimate each partial regression function $f_{-j}(x_{-j})$ ? Defaults to 5.  |

**Details**

This model assumes that a z-statistic  $z$  arises from

$$f(z_i) = w_i f^1(z) + (1 - w_i) f^0(z),$$

where  $f^1(z)$  and  $f^0(z)$  are the densities/marginal likelihoods under the alternative and null hypotheses, respectively, and where  $w_i$  is the prior probability that  $z_i$  is a signal (non-null case). Efron's method is used to estimate  $f(z)$  nonparametrically;  $f^0(z)$  may either be the theoretical (standard normal) null, or an empirical null which can be estimated using the middle 25 percent of the data. The prior probabilities  $w_i$  are estimated via logistic regression against covariates, using the Poly-Gamma Gibbs sampler of Polson, Scott, and Windle (JASA, 2013).

**Value**

|                       |  |
|-----------------------|--|
| <code>z</code>        | The test statistics provided as the argument <code>z</code> .  |
| <code>localfdr</code> | The corresponding vector of local false discovery rates (lfdr) for the elements of <code>z</code> . <code>localfdr[i]</code> is simply 1 minus the fitted posterior probability that <code>z[i]</code> comes from the non-null (signal) population. It is important to remember that <code>localfdr</code> is not necessarily monotonic in <code>z</code> , because the regression model allows the prior probability that <code>z[i]</code> is a signal to change with covariates <code>x[i]</code> . |
| <code>FDR</code>      | The corresponding vector of cut-level false discovery rates (FDR) for the elements of <code>z</code> . Used for extracting findings at a given FDR level. <code>FDR[i]</code> is the estimated false discovery rate for the cohort of test statistics whose local <code>lfdr</code> 's are at least as small as <code>localfdr[i]</code> — that is, the <code>z[j]</code> 's such that <code>localfdr[j] &lt;= localfdr[i]</code> .  |

|                             |  |
|-----------------------------|--|
| <code>X</code>              | The design matrix used in the regression. This will include an added column for an intercept, along with the spline basis expansion if <code>type='additive'</code> .  |
| <code>grid</code>           | Length <code>nmids</code> : equally-spaced midpoints of the histogram bins used to estimate $f(z)$ via Poisson spline regression.  |
| <code>breaks</code>         | Length <code>nmids</code> : the breakpoints of the histogram used to estimate $f(z)$ via Poisson spline regression.  |
| <code>grid.fz</code>        | Length <code>nmids</code> : the estimated value of $f(z)$ at the histogram midpoints.  |
| <code>grid.f0z</code>       | Length <code>nmids</code> : the estimated value of $f^0(z)$ , the assumed (either theoretical or empirical) null density at the histogram midpoints.   |
| <code>grid.zcounts</code>   | Length <code>nmids</code> : The number of z-scores that fell into each histogram bin.  |
| <code>dnull</code>          | The estimated (or assumed) null density at each of the observed z scores; <code>dnull[i]</code> corresponds to $z[i]$ .  |
| <code>dmix</code>           | The estimated marginal density $f(z)$ at each point $z[i]$ . This should look like a good, smooth fit to the histogram of $z$ .  |
| <code>empirical.null</code> | A list with two members <code>mu0</code> and <code>sig0</code> , representing the mean and standard deviation of the empirical null estimated using Efron's central-matching method. Always returned, but only used if <code>nulltype='empirical'</code> .   |
| <code>betasave</code>       | A matrix of posterior draws. Each row is a single posterior draw of the vector of regression coefficients corresponding to the columns of the returned <code>X</code> .  |
| <code>priorprob</code>      | The estimated prior probability of being a signal for each observation $z_i$ . Here <code>priorprob[i] = P(z_i is non-null)</code> .   |
| <code>postprob</code>       | The estimated posterior probabilities of being a signal each observation $z_i$ : <code>postprob[i] = P(z_i is non-null   data)</code> , and <code>localfdr[i] = 1 - postprob[i]</code> .   |
| <code>fjindex</code>        | A list of indices of length <code>ncol(covars)</code> , where <code>covars</code> is the matrix of covariates you fed in. Mainly useful if <code>type='additive'</code> , in which case <code>fjind[[j]]</code> gives you a vector of indices telling you which columns of the returned <code>X</code> and <code>betasave</code> correspond to the basis expansion of the original design matrix <code>covars[,j]</code> . |

## References

- J.G. Scott, R. Kelly, M.A. Smith, P. Zhou, and R.E. Kass (2013). False discovery rate regression: application to neural synchrony detection in primary visual cortex. arXiv:1307.3495 [stat.ME].
- N.G. Polson, J.G. Scott, and J. Windle (2013). Bayesian inference for logistic models using Poly-Gamma latent variables. *Journal of the American Statistical Association (Theory and Methods)* 108(504): 1339-49 (2013). arXiv:1205.0310 [stat.ME].
- Efron (2004). Large-scale simultaneous hypothesis testing: the choice of a null hypothesis. *J. Amer. Statist. Assoc.* 99, 96-104.
- Efron (2005). Local false discovery rates. Preprint, Dept. of Statistics, Stanford University.

## Examples

```
library(FDRreg)

# Simulated data
```

```

P = 2
N = 10000
betatrue = c(-3.5,rep(1/sqrt(P), P))
X = matrix(rnorm(N*P), N,P)
psi = crossprod(t(cbind(1,X)), betatrue)
wsuccess = 1/{1+exp(-psi)}

# Some theta's are signals, most are noise
gammatrue = rbinom(N,1,wsuccess)
table(gammatrue)

# Density of signals
thetatrue = rnorm(N,3,0.5)
thetatrue[gammatrue==0] = 0
z = rnorm(N, thetatrue, 1)
hist(z, 100, prob=TRUE, col='lightblue', border=NA)
curve(dnorm(x,0,1), add=TRUE, n=1001)

## Not run:
# Fit the model
fdr1 <- FDRreg(z, covars=X, nmc=2500, nburn=100, nmids=120, nulltype='theoretical')

# Show the empirical-Bayes estimate of the mixture density
# and the findings at a specific FDR level
Q = 0.1
plotFDR(fdr1, Q=Q, showfz=TRUE)

# Posterior distribution of the intercept
hist(fdr1$betasave[,1], 20)

# Compare actual versus estimated prior probabilities of being a signal
plot(wsuccess, fdr1$priorprob)

# Covariate effects
plot(X[,1], log(fdr1$priorprob/{1-fdr1$priorprob}), ylab='Logit of prior probability')
plot(X[,2], log(fdr1$priorprob/{1-fdr1$priorprob}), ylab='Logit of prior probability')

# Local FDR
plot(z, fdr1$localfdr, ylab='Local false-discovery rate')

# Extract findings at level FDR = Q
myfindings = which(fdr1$FDR <= Q)

## End(Not run)

```

**Description**

Plots the results of a fitted FDR regression model from FDRreg.

**Usage**

```
plotFDR(fdr, Q=0.1, showrug=TRUE, showfz=TRUE, showsub=TRUE)
```

**Arguments**

|         |  |
|---------|--|
| fdr     | A fitted model object from FDRreg.   |
| Q       | The desired level at which FDR should be controlled. Defaults to 0.1, or 10 percent.   |
| showrug | Logical flag indicating whether the findings at the specified FDR level should be displayed in a rug plot beneath the histogram. Defaults to TRUE. |
| showfz  | Logical flag indicating the fitted marginal density $f(z)$ should be plotted. Defaults to TRUE.  |
| showsub | Logical flag indicating whether a subtitle should be displayed describing features of the plot. Defaults to TRUE.                                  |

**Details**

It is important to remember that `localfdr` (and therefore global FDR) is not necessarily monotonic in  $z$ , because the regression model allows the prior probability that  $z[i]$  is a signal to change with covariates  $x[i]$ .

**Value**

No return value.

**Examples**

```
library(FDRreg)

# Simulated data
P = 2
N = 10000
betatrue = c(-3.5, rep(1/sqrt(P), P))
X = matrix(rnorm(N*P), N, P)
psi = crossprod(t(cbind(1, X)), betatrue)
wsuccess = 1/{1+exp(-psi)}

# Some theta's are signals, most are noise
gammatrue = rbinom(N, 1, wsuccess)
table(gammatrue)

# Density of signals
thetatrue = rnorm(N, 3, 0.5)
thetatrue[gammatrue==0] = 0
```

```
z = rnorm(N, thetatrue, 1)
hist(z, 100, prob=TRUE, col='lightblue', border=NA)
curve(dnorm(x,0,1), add=TRUE, n=1001)

## Not run:
# Fit the model
fdr1 <- FDRreg(z, covars=X, nmc=2500, nburn=100, nmids=120, nulltype='theoretical')
# Show the empirical-Bayes estimate of the mixture density
# and the findings at a specific FDR level
Q = 0.1
plotFDR(fdr1, Q=Q, showfz=TRUE)

## End(Not run)
```



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