

# Package ‘FDRreg’

February 19, 2015

**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

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

**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(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)

```

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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 fdr'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 <code>z[i]</code> .
<code>dmix</code>	The estimated marginal density $f(z)$ at each point <code>z[i]</code> . This should look like a good, smooth fit to the histogram of <code>z</code> .
<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 <code>z_i</code> . Here <code>priorprob[i] = P(z_i is non-null)</code> .
<code>postprob</code>	The estimated posterior probabilities of being a signal each observation <code>z_i</code> : <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)
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# Density of signals
thetatrue = rnorm(N, 3, 0.5)
thetatrue[gammatrue==0] = 0
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```
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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