Package 'OxyBS'

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Type Package
Title Processing of Oxy-Bisulfite Microarray Data
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Depends R (>= $3.2.2$)
Description Provides utilities for processing of Oxy-Bisulfite microarray data (e.g. via the Illumina Infinium platform, http://www.illumina.com) with tandem arrays, one using conventional bisulfite conversion, the other using oxy-bisulfite conversion.
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diffBeta1

First derivative of beta minus-log-pdf with respect to first parameter

Description

First derivative of -log(beta pdf) wrt a (first) parameter

Usage

```
diffBeta1(x,a,b)
```

Arguments

x beta value

a a parameter (first)

b parameter (second)

Details

First derivative of beta minus-log-pdf with respect to first parameter; used for maximum likelihood estimation, not typically called by user.

Value

first derivative with respect to a (first) parameter

Author(s)

E. Andres Houseman

See Also

diffBeta2,score0xBS

diffBeta2	First derivative of beta minus-log-pdf with respect to second parame-
	ter

Description

First derivative of -log(beta pdf) wrt b (second) parameter

Usage

```
diffBeta2(x,a,b)
```

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Arguments

X	beta val	ne

a a parameter (first)

b parameter (second)

Details

First derivative of beta minus-log-pdf with respect to second parameter; used for maximum likelihood estimation, not typically called by user.

Value

first derivative with respect to b (second) parameter

Author(s)

E. Andres Houseman

See Also

diffBeta1,score0xBS

exampleMethBS

Sample Data: Methylation (red) signals from conventional bisulfite conversion.

Description

Matrix of signal intensitities corresponding to 30 specimens and 30 CpGs.

Usage

exampleMethBS

Format

30 x 30 matrix (CpGs x Specimens)

exampleMeth0xBS

Sample Data: Methylation (red) signals from oxy-bisulfite conversion.

Description

Matrix of signal intensitities corresponding to 30 specimens and 30 CpGs.

Usage

exampleMethOxBS

Format

30 x 30 matrix (CpGs x Specimens)

exampleUnmethBS

Sample Data: Unmethylated (green) signals from conventional bisulfite conversion.

Description

Matrix of signal intensitities corresponding to 30 specimens and 30 CpGs.

Usage

exampleUnmethBS

Format

30 x 30 matrix (CpGs x Specimens)

exampleUnmethOxBS

Sample Data: Unmethylated (green) signals from oxy-bisulfite conversion.

Description

Matrix of signal intensitities corresponding to 30 specimens and 30 CpGs.

Usage

exampleUnmethOxBS

Format

30 x 30 matrix (CpGs x Specimens)

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fitOneOxBS	Fit one OxyBS result	
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Description

Uses maximum likelihood to estimate (C,5mC,5hmC) for one CpG and one specimen

Usage

```
fitOneOxBS(betaBS, betaOxBS, signalBS, signalOxBS, eps=1E-5)
```

Arguments

betaBS	beta value from conventional bisulfite conversion
beta0xBS	beta value from oxy-bisulfite conversion
signalBS	total signal from conventional bisulfite conversion
signal0xBS	total signal from oxy-bisulfite conversion
eps	small positive value representing numerical zero

Details

Uses maximum likelihood to estimate (C,5mC,5hmC) for one CpG and one specimen; not typically called by user.

Value

(C,5mC,5hmC) for one CpG and one specimen.

Author(s)

E. Andres Houseman

See Also

fit0xBS

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fit0xBS	Fit OxyBS for one specimen
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Description

Uses maximum likelihood to estimate (C,5mC,5hmC) vectors for one specimen

Usage

```
fit0xBS(betaBS, beta0xBS, signalBS, signal0xBS, eps=1E-5)
```

Arguments

betaBS	beta value from conventional bisulfite conversion
beta0xBS	beta value from oxy-bisulfite conversion
signalBS	total signal from conventional bisulfite conversion
signal0xBS	total signal from oxy-bisulfite conversion
eps	small positive value representing numerical zero

Details

Uses maximum likelihood to estimate (C,5mC,5hmC) one specimen (many CpGs).

Value

matrix of (C,5mC,5hmC) values (each row corresponds to a separate CpG).

Author(s)

E. Andres Houseman

Examples

```
## Not run:
data(OxyBSSampleData)

nSpecimens <- 30
nCpGs <- 30

# Calculate Total Signals
signalBS <- exampleMethBS+exampleUnmethBS
signalOxBS <- exampleMethOxBS+exampleUnmethOxBS

# Calculate Beta Values
betaBS <- exampleMethBS/signalBS
betaOxBS <- exampleMethOxBS/signalOxBS</pre>
# Create container for results
```

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```
MethOxy <- array(NA,dim=c(nCpGs,nSpecimens,3))</pre>
dimnames(MethOxy) <- list(</pre>
  rownames(exampleMethBS)[1:nCpGs],
  colnames(exampleMethBS)[1:nSpecimens],
  c("C", "5mC", "5hmC"))
# Process results (one array at a time)
for(i in 1:nSpecimens){
  MethOxy[,i,] <- fit0xBS(betaBS[,i],beta0xBS[,i],signalBS[,i],signal0xBS[,i])</pre>
# Check that results sum to one
table(apply(MethOxy,1:2,sum))
# First specimen
MethOxy[,1,]
# Ranges
range(MethOxy[,,1])
range(MethOxy[,,2])
range(MethOxy[,,3])
## End(Not run)
```

like0xBS

Likelihood function for C/5mC/5hmC likelihood estimator

Description

Likelihood function for C/5mC/5hmC likelihood estimator

Usage

```
likeOxBS(theta, betaBS, betaOxBS, signalBS, signalOxBS)
```

Arguments

theta	2-element parameter vector
betaBS	beta value from conventional bisulfite conversion
beta0xBS	beta value from oxy-bisulfite conversion
signalBS	total signal from conventional bisulfite conversion
signal0xBS	total signal from oxy-bisulfite conversion

Details

Likelihood function for C/5mC/5hmC likelihood estimator; used for maximum likelihood estimation, not typically called by user.

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Value

likelihood for C/5mC/5hmC likelihood

Author(s)

E. Andres Houseman

See Also

fitOneOxBS

score0xBS

Score function for C/5mC/5hmC likelihood estimator

Description

Score function for C/5mC/5hmC likelihood estimator

Usage

```
scoreOxBS(theta, betaBS, betaOxBS, signalBS, signalOxBS)
```

Arguments

theta 2-element parameter vector

betaBS beta value from conventional bisulfite conversion

beta0xBS beta value from oxy-bisulfite conversion

signalBS total signal from conventional bisulfite conversion

signalOxBS total signal from oxy-bisulfite conversion

Details

Score function for C/5mC/5hmC likelihood estimator; used for maximum likelihood estimation, not typically called by user.

Value

score vector for C/5mC/5hmC likelihood

Author(s)

E. Andres Houseman

See Also

fitOneOxBS

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