

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.

License GPL (>= 2)

NeedsCompilation no

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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	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 parameter***Description**

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

Usage

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

Arguments

x	beta value
a	a parameter (first)
b	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 intensities corresponding to 30 specimens and 30 CpGs.

Usage

exampleMethBS

Format

30 x 30 matrix (CpGs x Specimens)

`exampleMethOxBs`

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

Description

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

Usage

```
exampleUnmethOxBs
```

Format

30 x 30 matrix (CpGs x Specimens)

fitOneOxBs*Fit one OxyBS result*

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
betaOxBs	beta value from oxy-bisulfite conversion
signalBS	total signal from conventional bisulfite conversion
signalOxBs	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

[fitOxBs](#)

fitOxBs*Fit OxyBS for one specimen***Description**

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

Usage

```
fitOxBs(betaBS, betaOxBs, signalBS, signalOxBs, eps=1E-5)
```

Arguments

<code>betaBS</code>	beta value from conventional bisulfite conversion
<code>betaOxBs</code>	beta value from oxy-bisulfite conversion
<code>signalBS</code>	total signal from conventional bisulfite conversion
<code>signalOxBs</code>	total signal from oxy-bisulfite conversion
<code>eps</code>	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

# Create container for results
```

```

MethOxy <- array(NA,dim=c(nCpGs,nSpecimens,3))
dimnames(MethOxy) <- list(
  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,] <- fitOxBs(betaBS[,i],betaOxBs[,i],signalBS[,i],signalOxBs[,i])
}

# 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)

```

likeOxBs

*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
betaOxBs	beta value from oxy-bisulfite conversion
signalBS	total signal from conventional bisulfite conversion
signalOxBs	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.

Value

likelihood for C/5mC/5hmC likelihood

Author(s)

E. Andres Houseman

See Also

[fitOne0xBS](#)

score0xBS

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

Description

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

Usage

`score0xBS(theta, betaBS, beta0xBS, signalBS, signal0xBS)`

Arguments

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

[fitOne0xBS](#)

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