

# Package ‘EmpericalBrownsMethod’

September 4, 2015

**Title** Uses Brown's method to combine p-values from dependent tests.

**Version** 0.1

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**Description** Combining P-values from multiple statistical tests is common in bioinformatics. However, this procedure is non-trivial for dependent P-values. This package implements an empirical adaptation of Brown's Method (an extension of Fisher's Method) for combining dependent P-values which is appropriate for highly correlated data sets found in high-throughput biological experiments.

**Depends** R (>= 3.2.0)

**Suggests** BiocStyle, testthat, knitr

**License** Apache License, Version 2.0

**VignetteBuilder** knitr

**URL** <https://github.com/IlyaLab/CombiningDependentPvaluesUsingEBM.git>

**LazyData** true

**Encoding** UTF-8

**NeedsCompilation** no

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`allPvals`*Data used in tests*

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**Description**

This data is used in the unit tests and shows a usage example.

**Usage**`testData`**Format**`data.frame`**Source**`GEO`

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`dat`*Data used in tests*

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**Description**

This data is used in the unit tests and shows a usage example.

**Usage**`testData`**Format**`data.frame`**Source**`GEO`

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empiricalBrownsMethod *EmpiricalBrownsMethod*


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**Description**

Combine p-values

**Usage**

```
empiricalBrownsMethod(data_matrix, p_values, extra_info)
```

**Arguments**

```
data_matrix    a
p_values       b
extra_info     c
```

**Examples**

```
## restore the saved values to the current environment
data(testData)
# glypGenes <- pathways$gene[pathways$pathway == "GLYPICAN 3 NETWORK"];
# glypPvals <- allPvals$pvalue.with.CHD4[allPvals$gene %in% glypGenes];
# glypDat   <- dat[dat$V1 %in% glypGenes, 2:ncol(dat)];
# print(empiricalBrownsMethod(data_matrix=glypDat, p_values=glypPvals, extra_info=T));
```

---

pathways *Data used in tests*


---

**Description**

This data is used in the unit tests and shows a usage example.

**Usage**

```
testData
```

**Format**

```
data.frame
```

**Source**

```
GEO
```

---

randData	<i>Data used in tests</i>
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---

**Description**

This data is used in the unit tests and shows a usage example.

**Usage**

testData

**Format**

data.frame

**Source**

GEO

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testData	<i>Data used in tests</i>
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**Description**

This data is used in the unit tests and shows a usage example.

**Usage**

testData

**Format**

Four items

**Source**

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