Package 'dam'

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Title Data Analysis Metabolomics

Version 0.0.1

Description A collection of functions which aim to assist common computational workflow for analysis of matabolomic data..

Depends R (>= 3.2.3)

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Encoding UTF-8

LazyData true

RoxygenNote 5.0.1.9000

NeedsCompilation no

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R topics documented:

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correctedMarkers *A correctedMarkers Function*

Description

This function allows you to perform several multiple regression on metabolomic data.

Usage

```
correctedMarkers(dat, fac, con, comp)
```

Arguments

`dat`	Data matrix.
`fac`	Index of the group of samples.
`con`	Index of correction variable.
`comp`	Index (range) of measured compounds.

Examples

```
correctedMarkers(iris, 5, 1, 2:4)
```

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