

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

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NeedsCompilation no

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correctedMarkers	<i>A correctedMarkers Function</i>
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Description

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

Usage

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

Arguments

<code>dat</code>	Data matrix.
<code>fac</code>	Index of the group of samples.
<code>con</code>	Index of correction variable.
<code>comp</code>	Index (range) of measured compounds.

Examples

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

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