# Package 'affycompData'

September 19, 2024

september 19, 2021
<b>Version</b> 1.42.0
Title affycomp data
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<pre>URL https://bioconductor.org/packages/affycompData</pre>
<pre>BugReports https://github.com/rafalab/affyCompData/issues</pre>
<b>Depends</b> R ( $>= 2.13.0$ ), methods, Biobase ( $>= 2.3.3$ ), affycomp
<b>Description</b> Data needed by the affycomp package.
License GPL (>= 2)
biocViews MicroarrayData
git_url https://git.bioconductor.org/packages/affycompData
git_branch RELEASE_3_19
git_last_commit 59d20c9
git_last_commit_date 2024-04-30
Repository Bioconductor 3.19
Date/Publication 2024-09-19
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lw.sd.assessment

An example of the result of an SD assessment

# Description

The Dilution files were processed with the dChip package (using PM-only), and then the function assessSD from the affycomp package was applied.

#### Usage

```
data(lw.sd.assessment)
```

#### **Format**

A list.

mas5.assessment

Examples of the result of assessments

# Description

The Dilution and both (HGU95 and HGU133) types of Spike-in data were processed with Affymetrix MAS 5.0 software, yielding three "MAS 5.0" ExpressionSet's. (These are available, in csv-format, at http://affycomp.jhsph.edu/AFFY2/rafa@jhu.edu/030424.1033/.) Then various assessment functions from the affycomp package (most recently, version 1.28.0) were applied. mas5.assessment resulted from assessAll on Dilution and HGU95; mas5.assessment.133 from assessSpikeIn on HGU133; mas5.assessment2 from assessSpikeIn2 on HGU95; and mas5.assessment2.133 from assessSpikeIn2 on HGU133.

#### Usage

```
data(mas5.assessment)
data(mas5.assessment.133)
data(mas5.assessment2)
data(mas5.assessment2.133)
```

#### **Format**

A list of list.

rma.assessment 3

rma.assessment

Examples of the result of assessments

#### Description

The Dilution and both (HGU95 and HGU133) types of Spike-in data were processed with the (version 1.0) function rma, yielding three "RMA" ExpressionSet's. (These are available, in csv-format, at http://affycomp.jhsph.edu/AFFY2/rafa@jhu.edu/030429.1332/.) Then various assessment functions from the affycomp package (most recently, version 1.28.0) were applied. rma.assessment resulted from assessAll on Dilution and HGU95; rma.assessment.133 from assessSpikeIn on HGU133; rma.assessment2 from assessSpikeIn2 on HGU95; and rma.assessment2.133 from assessSpikeIn2 on HGU133.

## Usage

```
data(rma.assessment)
data(rma.assessment.133)
data(rma.assessment2)
data(rma.assessment2.133)
```

#### **Format**

A list of list.

rma.sd.assessment

An example of the result of an SD assessment

## **Description**

The Dilution files were processed with the affy version 1.0 package rma add-on function, and then the function assessSD from the affycomp package was applied.

# Usage

```
data(rma.sd.assessment)
```

#### **Format**

A list.

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