# Package 'affycompData'

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Title affycomp data
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<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 OneChannel, Microarray, Preprocessing
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lw.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)

2 rma.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

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 <a href="http://affycomp.jhsph.edu/AFFY2/rafa@jhu.edu/030429.1332/">http://affycomp.jhsph.edu/AFFY2/rafa@jhu.edu/030429.1332/</a>.) 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)
```

rma.sd.assessment 3

### **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.

# **Index**

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