# Package 'PRECISIONDATA'

July 7, 2016

Title Datasets for the PaiREd miCrorna sImulation on Study desIgn for

mOlecular classificatioN (PRECISION) Package

Type Package

Version 0.1	
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Author Rebecca Huang, Li-Xuan Qin	
Maintainer Rebecca Huang <huangh4@mskcc.org></huangh4@mskcc.org>	
<b>Description</b> Different versions of a unique pair of Agilent microRNA microarray datasets that can be used for the PRECISION package.	
License GPL (>= 2)	
<b>Depends</b> R (>= $3.0.2$ )	
Suggests rmarkdown, knitr	
LazyData TRUE	
RoxygenNote 5.0.1	
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noba.nuhdata.log2

The non-uniformly-handled probe-level dataset, before preprocessing

## **Description**

The non-uniformly-handled probe-level dataset, with control probes removed. The expressions are on a log2 scale without background adjustment. The sample IDs (the column names) ending with "E" or "V" are used to indicate whether a sample is endometrial or ovarian tumor sample. There are 96 endometrial and 96 ovarian tumor samples.

#### Usage

noba.nuhdata.log2

#### **Format**

A data matrix with 53880 rows (probes) and 192 columns (samples).

#### References

http://clincancerres.aacrjournals.org/content/20/13/3371.long

noba.nuhdata.log2.p10 *The non-uniformly-handled probe-level dataset, 10 probes for each unique probe* 

## **Description**

The non-uniformly-handled probe-level dataset with 10 probes for each unique probe. It is truncated from the non-uniformly-handled probe-level dataset (noba.nuhdata.log2.p10), by keeping only the first 10 probes for each unique probe.

#### Usage

noba.nuhdata.log2.p10

#### **Format**

A data matrix with 35230 rows (probes) and 192 columns (samples).

noba.nuhdata.log2.p10.psl

The non-uniformly-handled probe-set-level dataset

## **Description**

The non-uniformly-handled probe-set-level dataset. It is probe-set summarized from the non-uniformly-handled probe-level dataset with 10 probes for each unique probe (noba.nuhdata.log2.p10), using median.

## Usage

```
noba.nuhdata.log2.p10.psl
```

#### **Format**

A data matrix with 3523 rows (probes) and 192 columns (samples).

noba.uhdata.log2

The uniformly-handled probe-level dataset, before preprocessing

## Description

The uniformly-handled probe-level dataset, with control probes removed. The expressions are on a log2 scale without background adjustment. The sample IDs (the column names) ending with "E" or "V" are used to indicate whether a sample is endometrial or ovarian tumor sample. There are 96 endometrial and 96 ovarian tumor samples.

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noba.uhdata.log2
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 ${\it noba.uhdata.log2.p10} \quad \textit{The uniformly-handled probe-level dataset, 10 probes for each unique probe}$ 

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## Usage

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noba.uhdata.log2.p10
```

#### **Format**

A data matrix with 35230 rows (probes) and 192 columns (samples).

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noba.uhdata.log2.p10.psl
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The uniformly-handled probe-set-level dataset

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noba.uhdata.log2.p10.psl
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#### **Format**

A data matrix with 3523 rows (probes) and 192 columns (samples).

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