

mogene.1.0.st.v1frmavecs

May 15, 2019

mogene.1.0.st.v1barcodevecs

Vectors used by barcode function for mogene.1.0.st.v1.

Description

Parameters of the background distribution for use with the barcode function. These correspond to version 3.0 of the Gene Expression Barcode website.

Usage

`data(mogene.1.0.st.v1barcodevecs)`

Format

A list with 3 elements.

<code>mu</code>	background means
<code>tau</code>	background standard deviations
<code>entropy</code>	observed gene entropy

Examples

```
data(mogene.1.0.st.v1barcodevecs)
str(mogene.1.0.st.v1barcodevecs)
```

mogene.1.0.st.v1frmavecs

Vectors used by fRMA for mogene.1.0.st.v1.

Description

Frozen parameter vectors for use with the fRMA and GNUMSE functions.

Usage

`data(mogene.1.0.st.v1frmavecs)`

Format

A list with 6 elements.

<code>normVec</code>	normalization vector
<code>probeVec</code>	probe effect vector
<code>probeVarWithin</code>	within batch probe variance
<code>probeVarBetween</code>	between batch probe variance
<code>probesetSD</code>	within probeset standard deviation
<code>medianSE</code>	median standard error for gene expression estimates
<code>probeVecCore</code>	exon effect vector for summarizing to core transcripts
<code>mapCore</code>	mapping between exons and core transcripts

Examples

```
data(mogene.1.0.st.v1frmavecs)
str(mogene.1.0.st.v1frmavecs)
```

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