

huex.1.0.st.v2frmavecs

September 24, 2013

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Vectors used by fRMA for huex.1.0.st.v2.

Description

Frozen parameter vectors for use with the `frma` and `GNUSE` functions.

Usage

```
data(huex.1.0.st.v2frmavecs)
```

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 when summarizing to core transcripts
<code>probeVecExt</code>	probe effect vector when summarizing to extended transcripts
<code>probeVecFull</code>	probe effect vector when summarizing to full transcripts
<code>mapCore</code>	mapping between exons and core transcripts
<code>mapExt</code>	mapping between exons and extended transcripts
<code>mapFull</code>	mapping between exons and full transcripts

Examples

```
data(huex.1.0.st.v2frmavecs)  
str(huex.1.0.st.v2frmavecs)
```

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