HuExExonProbesetLocationHg18

February 28, 2017

HuExExonProbesetLocation

Probe sequence for microarrays of type HuEx.

Description

This data object was automatically created by the package AnnotationDbi version 1.8.1.

Usage

data(HuExExonProbesetLocation)

Format

A data frame with 1425756 rows and 7 columns, as follows.

<table>
<thead>
<tr>
<th></th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>EPROBESETID</td>
<td>exon-level probeset id</td>
</tr>
<tr>
<td>CHR</td>
<td>chromosome</td>
</tr>
<tr>
<td>STRAND</td>
<td>chromosome strand</td>
</tr>
<tr>
<td>START</td>
<td>start position of the Probe Selection Region on chromosome</td>
</tr>
<tr>
<td>END</td>
<td>end position of the Probe Selection Region on chromosome</td>
</tr>
<tr>
<td>GPROBESETID</td>
<td>Gene-level probeset id</td>
</tr>
<tr>
<td>ANNLEVEL</td>
<td>annotation level</td>
</tr>
</tbody>
</table>

Source

The exon-level probeset genome location was retrieved from Netaffx using AffyCompatible.

Examples

library(HuExExonProbesetLocationHg18)
data(HuExExonProbesetLocation)
as.data.frame(HuExExonProbesetLocation[1:3,])
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