HuExExonProbesetLocation
April 26, 2017

HuExExonProbesetLocation

*Probe sequence for microarrays of type HuEx.*

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**Description**

This data object was automatically created by the package AnnotationForge version 1.7.17.

**Usage**

data(HuExExonProbesetLocation)

**Format**

A data frame with 1432143 rows and 7 columns, as follows.

<table>
<thead>
<tr>
<th>Column</th>
<th>Type</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>sequence</td>
<td>character</td>
<td>probe sequence</td>
</tr>
<tr>
<td>x</td>
<td>integer</td>
<td>x-coordinate on the array</td>
</tr>
<tr>
<td>y</td>
<td>integer</td>
<td>y-coordinate on the array</td>
</tr>
<tr>
<td>Probe.Set.Name</td>
<td>character</td>
<td>Affymetrix Probe Set Name</td>
</tr>
<tr>
<td>Probe.Interrogation.Position</td>
<td>integer</td>
<td>Probe Interrogation Position</td>
</tr>
<tr>
<td>Target.Strandedness</td>
<td>factor</td>
<td>Target Strandedness</td>
</tr>
</tbody>
</table>

**Source**

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

**Examples**

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

«Topic **datasets**
   HuExonProbesetLocation, 1

HuExonProbesetLocation, 1