HuExExonProbesetLocation
January 10, 2024

HuExExonProbesetLocation

Probe sequence for microarrays of type HuEx.

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>sequence</th>
<th>character</th>
<th>probe sequence</th>
</tr>
</thead>
<tbody>
<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

* datasets
  HuExExonProbesetLocation, 1

HuExExonProbesetLocation, 1