MoExExonProbesetLocation

November 18, 2016

MoExExonProbesetLocation

*ProbesetLocation for microarrays of type MoEx.*

Description

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

Usage

data(MoExExonProbesetLocation)

Format

A data frame with 1256831 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

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

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