

# RaExExonProbesetLocation

October 7, 2014

---

RaExExonProbesetLocation

*Probe sequence for microarrays of type RaEx.*

---

## Description

This data object was automatically created by the package AnnotationForge version 1.5.21.

## Usage

```
data(RaExExonProbesetLocation)
```

## Format

A data frame with 1064901 rows and 7 columns, as follows.

sequence	character	probe sequence
x	integer	x-coordinate on the array
y	integer	y-coordinate on the array
Probe.Set.Name	character	Affymetrix Probe Set Name
Probe.Interrogation.Position	integer	Probe Interrogation Position
Target.Strandedness	factor	Target Strandedness

## Source

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

## Examples

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

# Index

\*Topic **datasets**

RaExExonProbesetLocation, [1](#)

RaExExonProbesetLocation, [1](#)