

# RaExExonProbesetLocation

June 9, 2011

---

RaExExonProbesetLocation

*Probe sequence for microarrays of type RaEx.*

---

## Description

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

## Usage

```
data(RaExExonProbesetLocation)
```

## Format

A data frame with 1065284 rows and 7 columns, as follows.

EPROBESETID	exon-level probeset id
CHR	chroosome
STRAND	chromosome strand
START	start position of the Probe Selection Region on chromosome
END	end position of the Probe Selection Region on chromosome
GPROBESETID	Gene-level probeset id
ANNLEVEL	annotation level

## 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](#)