

# HuExExonProbesetLocationHg18

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HuExExonProbesetLocation

*Probe sequence for microarrays of type HuEx.*

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## Description

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

## Usage

```
data(HuExExonProbesetLocation)
```

## Format

A data frame with 1425756 rows and 7 columns, as follows.

|             |  |
|-------------|--|
| EPROBESETID | exon-level probeset id                                     |
| CHR         | chromosome   |
| 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

```
library(HuExExonProbesetLocationHg18)
data(HuExExonProbesetLocation)
as.data.frame(HuExExonProbesetLocation[1:3,])
```

# Index

## \* datasets

HuExExonProbesetLocation, [1](#)

HuExExonProbesetLocation, [1](#)