

# 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,])
```

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