

# HuExExonProbesetLocation

March 18, 2021

---

HuExExonProbesetLocation

*Probe sequence for microarrays of type HuEx.*

---

## Description

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

## Usage

```
data(HuExExonProbesetLocation)
```

## Format

A data frame with 1432143 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

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

# Index

## \* datasets

HuExExonProbesetLocation, [1](#)

HuExExonProbesetLocation, [1](#)