

Package ‘tinesath1probe’

April 9, 2019

Title Probe sequence data for microarrays of type tinesath1

Version 1.20.0

Created Mon Aug 7 17:00:54 2006

Author The Bioconductor Project www.bioconductor.org

Description This package was automatically created by package matchprobes version 1.4.0. The probe sequence data was obtained from <http://www.affymetrix.com>.

Maintainer Tine Casneuf <tine@ebi.ac.uk>

Depends R (>= 1.6), AnnotationDbi (>= 1.11.9)

License LGPL

biocViews Arabidopsis_thaliana_Data, SequencingData, MicroarrayData

git_url <https://git.bioconductor.org/packages/tinesath1probe>

git_branch RELEASE_3_8

git_last_commit 12b7e7c

git_last_commit_date 2018-10-30

Date/Publication 2019-04-09

R topics documented:

tinesath1probe 1

Index 3

tinesath1probe	<i>Probe sequence for microarrays of type tinesath1.</i>
----------------	--

Description

This data object was automatically created by the package matchprobes version 1.4.0.

Usage

```
data(tinesath1probe)
```

Format

A data frame with 221802 rows and 6 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 probe sequence data was obtained from <http://www.affymetrix.com>.

Examples

```
data(tinesath1probe)
tinesath1probe
as.data.frame(tinesath1probe[1:3,])
```

Index

*Topic **datasets**

tinesath1probe, [1](#)

tinesath1probe, [1](#)