

Package ‘tinesath1probe’

October 1, 2024

Title Probe sequence data for microarrays of type tinesath1

Version 1.43.0

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 devel

git_last_commit cce5143

git_last_commit_date 2024-04-30

Repository Bioconductor 3.20

Date/Publication 2024-10-01

Contents

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

* **datasets**

 tinesath1probe, [1](#)

tinesath1probe, [1](#)