

# Package ‘tinesath1probe’

October 17, 2019

**Title** Probe sequence data for microarrays of type tinesath1

**Version** 1.22.0

**Created** Mon Aug 7 17:00:54 2006

**Author** The Bioconductor Project [www.bioconductor.org](http://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>.

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**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\_9

**git\_last\_commit** 3ae4818

**git\_last\_commit\_date** 2019-05-02

**Date/Publication** 2019-10-17

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tinesath1probe      *Probe sequence for microarrays of type tinesath1.*

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### 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,])
```

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