

# Package ‘tinesath1probe’

July 9, 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](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>.

**Maintainer** Tine Casneuf <[tine@ebi.ac.uk](mailto: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** 2024-07-09

## Contents

tinesath1probe . . . . .	1
<b>Index</b>	<b>3</b>

---

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](#)