

# Package ‘tinesath1probe’

September 18, 2025

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

**Version** 1.47.0

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**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>

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|                |                                                          |
|----------------|----------------------------------------------------------|
| tinesath1probe | <i>Probe sequence for microarrays of type tinesath1.</i> |
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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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## \* **datasets**

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