

Package ‘tinesath1probe’

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Title Probe sequence data for microarrays of type tinesath1

Version 1.18.0

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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>.

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

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biocViews Arabidopsis_thaliana_Data, SequencingData, MicroarrayData

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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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*Topic **datasets**

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