

# Package ‘prostateCancerGrasso’

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**Type** Package

**Title** Prostate Cancer Data

**Version** 1.41.0

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**Description** A Bioconductor data package for the Grasso (2012) Prostate Cancer dataset.

**License** Artistic-2.0

**LazyData** TRUE

**Depends** Biobase, R (>= 3.3)

**Suggests** GEOquery

**biocViews** ExperimentData, ExpressionData,GEO, MicroarrayData, CancerData, ProstateCancerData

**NeedsCompilation** no

**git\_url** <https://git.bioconductor.org/packages/prostateCancerGrasso>

**git\_branch** devel

**git\_last\_commit** d568d9a

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**Repository** Bioconductor 3.24

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grasso

*The Grasso Cohort from U. Michigan*

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**Description**

Object imported using GEOquery and curated to remove variables introduced by GEO. See vignette for details

**Usage**

```
grasso
```

**Format**

a ExpressionSet instance, 1 row per Agilent microarray probe.

**Value**

ExpressionSet object for the dataset

**Author(s)**

Mark Dunning

**Source**

GSE35988

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