

# Package ‘prostateCancerGrasso’

December 26, 2024

**Type** Package

**Title** Prostate Cancer Data

**Version** 1.34.0

**Date** 2016-03-24

**Author** Mark Dunning

**Maintainer** Mark Dunning <mark.dunning@cruk.cam.ac.uk>

**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** RELEASE\_3\_20

**git\_last\_commit** 05501d7

**git\_last\_commit\_date** 2024-10-29

**Repository** Bioconductor 3.20

**Date/Publication** 2024-12-26

## Contents

grasso . . . . .	2
<b>Index</b>	<b>3</b>

---

grasso

*The Grasso Cohort from U. Michigan*

---

**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

# Index

\* **datasets**  
    grasso, [2](#)

grasso, [2](#)