

# Package ‘prostateCancerVarambally’

April 16, 2024

**Type** Package  
**Title** Prostate Cancer Data  
**Version** 1.30.0  
**Date** 2016-03-24  
**Author** Mark Dunning  
**Maintainer** Mark Dunning <mark.dunning@cruk.cam.ac.uk>  
**Description** A Bioconductor data package for the Varambally 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/prostateCancerVarambally>  
**git\_branch** RELEASE\_3\_18  
**git\_last\_commit** 3c16ae4  
**git\_last\_commit\_date** 2023-10-24  
**Repository** Bioconductor 3.18  
**Date/Publication** 2024-04-16

## R topics documented:

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

---

varambally

*The Varambally Cohort from U. Michigan et al. (2005)*

---

**Description**

Object imported using GEOquery and curated to remove variables introduced by GEO

**Usage**

varambally

**Format**

a ExpressionSet instance, 1 row per microarray probe.

**Value**

ExpressionSet object for the dataset

**Author(s)**

Mark Dunning

**Source**

GSE3325

# Index

\* **datasets**

varambally, [2](#)

varambally, [2](#)