

# Package ‘prostateCancerVambally’

May 7, 2026

**Type** Package

**Title** Prostate Cancer Data

**Version** 1.41.0

**Date** 2016-03-24

**Author** Mark Dunning

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

**Description** A Bioconductor data package for the Vambally 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/prostateCancerVambally>

**git\_branch** devel

**git\_last\_commit** 45cbbbf

**git\_last\_commit\_date** 2026-04-28

**Repository** Bioconductor 3.24

**Date/Publication** 2026-05-07

## Contents

vambally . . . . .	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](#)