

# Package ‘prostateCancerVambally’

October 18, 2017

**Type** Package  
**Title** Prostate Cancer Data  
**Version** 1.4.0  
**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

## R topics documented:

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| vambally | <i>The Vambally Cohort from U. Michigan et al. (2005)</i> |
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## Description

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

## Usage

```
vambally
```

## Format

a ExpressionSet instance, 1 row per microarray probe.

**Value**

ExpressionSet object for the dataset

**Author(s)**

Mark Dunning

**Source**

GSE3325

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