

# Package ‘prostateCancerVarambally’

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**Type** Package

**Title** Prostate Cancer Data

**Version** 1.0.0

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**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

## R topics documented:

varambally . . . . . 1

**Index** 3

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

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## 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

\*Topic **datasets**  
varambally, 1

varambally, 1