

# Package ‘prostateCancerVarambally’

February 13, 2024

**Type** Package

**Title** Prostate Cancer Data

**Version** 1.31.0

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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

**git\_url** <https://git.bioconductor.org/packages/prostateCancerVarambally>

**git\_branch** devel

**git\_last\_commit** 2ac2a23

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**Repository** Bioconductor 3.19

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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

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