

# Package ‘prostateCancerTaylor’

September 11, 2025

**Type** Package  
**Title** Prostate Cancer Data  
**Version** 1.37.0  
**Date** 2016-03-24  
**Author** Mark Dunning  
**Maintainer** Mark Dunning <mark.dunning@cruk.cam.ac.uk>  
**Description** A Bioconductor data package for the Taylor et al (2010) dataset.  
**License** Artistic-2.0  
**LazyData** TRUE  
**Depends** Biobase, R (>= 3.3)  
**Suggests** GEOquery, org.Hs.eg.db  
**biocViews** ExperimentData, ExpressionData,GEO, MicroarrayData, CancerData, ProstateCancerData  
**NeedsCompilation** no  
**git\_url** <https://git.bioconductor.org/packages/prostateCancerTaylor>  
**git\_branch** devel  
**git\_last\_commit** 190218a  
**git\_last\_commit\_date** 2025-04-15  
**Repository** Bioconductor 3.22  
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taylor

*The MSKCC Cohort from Taylor et al. (2010)*

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**Description**

Object imported using GEOquery and curated to remove variables introduced by GEO. Details in Vignette.

**Usage**

camcap

**Format**

a ExpressionSet instance, 1 row per Affymetrix microarray probe.

**Value**

ExpressionSet object for the dataset

**Author(s)**

Mark Dunning

**Source**

GSE21034

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