

# Package ‘prostateCancerTaylor’

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**Type** Package

**Title** Prostate Cancer Data

**Version** 1.39.0

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**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** b7b0d5d

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**Repository** Bioconductor 3.23

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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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