

# Package ‘prostateCancerStockholm’

July 2, 2024

**Type** Package

**Title** Prostate Cancer Data

**Version** 1.33.0

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**Description** A Bioconductor data package for the Stockholm 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/prostateCancerStockholm>

**git\_branch** devel

**git\_last\_commit** bcb7da2

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**Repository** Bioconductor 3.20

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stockholm

*The Stockholm Cohort from Ross-Adams et al. (2015)*

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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 Illumina HuamnV4 probe.

**Value**

ExpressionSet object for the dataset

**Author(s)**

Mark Dunning

**Source**

GSE70769

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