

# Package ‘TCGAMethylation450k’

May 28, 2026

**Type** Package

**Title** The Cancer Genome Atlas Illumina 450k methylation example data

**Version** 1.49.0

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**Description** The Cancer Genome Atlas (TCGA) is applying genomics technologies to over 20 different types of cancer. This package contains a small set of 450k array data in idat format.

**License** GPL-2

**biocViews** Genome, CancerData, MethylationArrayData

**git\_url** <https://git.bioconductor.org/packages/TCGAMethylation450k>

**git\_branch** devel

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TCGAMethylation450k-package  
*The Cancer Genome Atlas Illumina 450k methylation example data*

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## Description

The Cancer Genome Atlas (TCGA) is applying genomics technologies to over 20 different types of cancer. This package contains a small set of 450k array data in idat format.

Data are included in this package in extdata/idat. A typical way of accessing these data would be to use the methylumi package methylumIDAT function.

**Details**

Package: TCGAMethylation450k  
Type: Package  
License: GPL-2

**Author(s)**

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**References**

<http://cancergenome.nih.gov/>

**See Also**

[methyumi](#)

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