

# Package ‘TCGAMethylation450k’

December 18, 2025

**Type** Package  
**Title** The Cancer Genome Atlas Illumina 450k methylation example data  
**Version** 1.47.0  
**Date** 2011-10-21  
**Author** Sean Davis  
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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 methylumiIDAT function.

## Details

Package: TCGAMethylation450k  
Type: Package  
License: GPL-2

## Author(s)

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## References

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

## See Also

[methylumi](#)

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