

Package ‘TCGAMethylation450k’

December 9, 2025

Type Package
Title The Cancer Genome Atlas Illumina 450k methylation example data
Version 1.47.0
Date 2011-10-21
Author Sean Davis
Maintainer Sean Davis <sdavis2@mail.nih.gov>
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
git_last_commit 284c80f
git_last_commit_date 2025-10-29
Repository Bioconductor 3.23
Date/Publication 2025-12-09

Contents

TCGAMethylation450k-package	2
Index	3

TCGAMethylation450k-package

The Cancer Genome Atlas Illumina 450k methylation example data

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)

Sean Davis

Maintainer: Sean Davis <sdavis2@mail.nih.gov>

References

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

See Also

[methylumi](#)

Index

* **package**

TCGAMethylation450k-package, [2](#)

methyllumi, [2](#)

TCGAMethylation450k

(TCGAMethylation450k-package),

[2](#)

TCGAMethylation450k-package, [2](#)