

Package: TCGAMethylation450k (via r-universe)

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Type Package

Title 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.

License GPL-2

biocViews Genome, CancerData, MethylationArrayData

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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 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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