Package ‘TCGAMethylation450k’

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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
NeedsCompilation no

R topics documented:

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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/

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