# Package 'TCGAMethylation450k'

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Title The Cancer Genome Atlas Illumina 450k methylation example data  Version 0.99.3  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  R topics documented:</sdavis2@mail.nih.gov>	
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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

methylumi

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