

# Package ‘genomationData’

October 17, 2019

**Title** Experimental data for showing functionalities of the genomation package

**Version** 1.16.0

**Date** 2013-10-25

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**Description** The package contains Chip Seq, Methylation and Cage data, downloaded from Encode

**License** GPL-3

**VignetteBuilder** knitr

**biocViews** ExperimentData, Genome, StemCell, SequencingData, ChIPSeqData, ChipOnChipData, ENCODE

**Suggests** knitr

**git\_url** <https://git.bioconductor.org/packages/genomationData>

**git\_branch** RELEASE\_3\_9

**git\_last\_commit** ed76790

**git\_last\_commit\_date** 2019-05-02

**Date/Publication** 2019-10-17

**R topics documented:**