

# Package ‘genomationData’

April 13, 2022

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

**Version** 1.26.0

**Date** 2013-10-25

**Author** Altuna Akalin, Vedran Franke

**Maintainer** Vedran Franke <vedran.franke@gmail.com>

**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\_14

**git\_last\_commit** 58bb495

**git\_last\_commit\_date** 2021-10-26

**Date/Publication** 2022-04-13

**R topics documented:**