

Package ‘genomationData’

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Title Experimental data for showing functionalities of the genomation package

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

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VignetteBuilder knitr

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

Suggests knitr

git_url <https://git.bioconductor.org/packages/genomationData>

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R topics documented: