

Package ‘HelloRangesData’

September 16, 2021

Type Package

Title Data for the HelloRanges tutorial vignette

Version 1.18.0

Author Michael Lawrence

Maintainer Michael Lawrence <michaf1a@gene.com>

Description Provides the data that were used in the bedtools tutorial by Aaron Quinlan (<http://quinlanlab.org/tutorials/bedtools/bedtools.html>). Includes a subset of the DnaseI hypersensitivity data from ``Maurano et al. Systematic Localization of Common Disease-Associated Variation in Regulatory DNA. Science. 2012. Vol. 337 no. 6099 pp. 1190-1195." The rest of the tracks were originally downloaded from the UCSC table browser. See the HelloRanges vignette for a port of the bedtools tutorial to R.

License GPL (>= 2)

Suggests BiocStyle

biocViews ExperimentData, SequencingData

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

git_branch RELEASE_3_13

git_last_commit 3657376

git_last_commit_date 2021-05-19

Date/Publication 2021-09-16

R topics documented: