Package 'HelloRangesData'

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Type Package

Title Data for the HelloRanges tutorial vignette

Version 1.34.0

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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.

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Suggests BiocStyle

biocViews ExperimentData, SequencingData

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