

# Package ‘LiebermanAidenHiC2009’

April 11, 2019

**Title** Selected data from the HiC paper of E. Lieberman-Aiden et al. in  
Science (2009)

**Version** 0.20.0

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**Description** This package provides data that were presented in  
the article ``Comprehensive mapping of long-range interactions reveals  
folding principles of the human genome'', Science 2009 Oct 9;326(5950):289-  
93. PMID: 19815776

**biocViews** ExperimentData, Genome, SequencingData, DNaseqData,  
ChIPSeqData, miRNAData, SmallRNAData, ChipOnChipData, GEO, NCI,  
ENCODE

**License** LGPL

**Depends** KernSmooth, IRanges

**Suggests**

**git\_url** <https://git.bioconductor.org/packages/LiebermanAidenHiC2009>

**git\_branch** RELEASE\_3\_8

**git\_last\_commit** 0c4ea54

**git\_last\_commit\_date** 2018-10-30

**Date/Publication** 2019-04-11

## R topics documented: