

# Package ‘diffloopdata’

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**Type** Package

**Title** Example ChIA-PET Datasets for the diffloop Package

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**Description** ChIA-PET example datasets and additional data for use with the diffloop package.

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**LazyData** TRUE

**biocViews** ExperimentData, SequencingData

**Suggests** knitr, rmarkdown

**VignetteBuilder** knitr

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

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**R topics documented:**