

# Package ‘MACSdata’

October 16, 2021

**Title** Test datasets for the MACSr package

**Version** 1.0.0

**Description** Test datasets from the MACS3 test examples are used in the examples of the ‘MACSr’ package. All 9 datasets are uploaded to the ‘ExperimentHub’. The original data can be found at: <https://github.com/macs3-project/MACS/>.

**License** file LICENSE

**Encoding** UTF-8

**LazyData** true

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.1.1

**biocViews** ExperimentHub, ExperimentData, ChIPSeqData

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

**git\_branch** RELEASE\_3\_13

**git\_last\_commit** 218fc1a

**git\_last\_commit\_date** 2021-05-19

**Date/Publication** 2021-10-16

**Author** Qiang Hu [aut, cre]

**Maintainer** Qiang Hu <[Qiang.Hu@roswellpark.org](mailto:Qiang.Hu@roswellpark.org)>

## R topics documented:

MACSdata . . . . . 1

**Index** . . . . . 2

---

MACSdata	<i>MACSdata</i>
----------	-----------------

---

## Description

This is an ExperimentHub Package for files that used in MACS unit test. Details can be found at: <https://macs3-project.github.io/macs/>.

# Index

MACSdata, [1](#)