

# Package ‘prebsdata’

May 13, 2025

**Title** Data for 'prebs' package

**Description** This package contains data required to run examples in 'prebs' package.

The data files include:

1) Small sample bam files for demonstration purposes

2) Probe sequence mappings for Cus-

tom CDF (taken from [http://brainarray.mbni.med.umich.edu/brainarray/Database/CustomCDF/genomic\\_curated\\_CDF](http://brainarray.mbni.med.umich.edu/brainarray/Database/CustomCDF/genomic_curated_CDF))

3) Probe sequence mappings for manufacturer's CDF (manually created using bowtie)

**Version** 1.45.0

**Author** Karolis Uziela and Antti Honkela

**Maintainer** Karolis Uziela <karolis.uziela@scilifelab.se>

**Depends** R (>= 2.14.0)

**License** Artistic-2.0

**biocViews** ExperimentData, SequencingData, RNASeqData

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

**git\_branch** devel

**git\_last\_commit** 23fb809

**git\_last\_commit\_date** 2025-04-15

**Repository** Bioconductor 3.22

**Date/Publication** 2025-05-13

## Contents