

# Package ‘EpiMix.data’

March 30, 2023

**Title** Data for the EpiMix package

**Version** 1.0.0

**Description** Supporting data for the EpiMix R package.

It include:

- HM450\_lncRNA\_probes.rda
- HM450\_miRNA\_probes.rda
- EPIC\_lncRNA\_probes.rda
- EPIC\_miRNA\_probes.rda
- EpigenomeMap.rda
- LUAD.sample.annotation
- TCGA\_BatchData
- MET.data
- mRNA.data
- microRNA.data
- lncRNA.data
- Sample\_EpiMixResults\_lncRNA
- Sample\_EpiMixResults\_miRNA
- Sample\_EpiMixResults\_Regular
- lncRNA expression data of tumors from TCGA that are stored in the ExperimentHub.

**License** GPL-3

**Encoding** UTF-8

**Depends** R (>= 4.2.0), ExperimentHub (>= 0.99.6)

**Suggests** rmarkdown, knitr

**biocViews** ExperimentData, ExperimentHub, Genome, RNASeqData,  
ExpressionData

**NeedsCompilation** no

**RoxygenNote** 7.2.1

**VignetteBuilder** knitr

**BugReports** <https://github.com/gevaertlab/EpiMix/issues>

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

**git\_branch** RELEASE\_3\_16

**git\_last\_commit** 6e6eed7

**git\_last\_commit\_date** 2022-11-01

**Date/Publication** 2023-03-30

**Author** Yuanning Zheng [aut, cre]

**Maintainer** Yuanning Zheng <eric2021@stanford.edu>

**R topics documented:**