

Package ‘BeadArrayUseCases’

January 23, 2022

Version 1.33.0

Date 2021-07-26

Title Analysing Illumina BeadArray expression data using Bioconductor

Author Mark Dunning, Wei Shi, Andy Lynch, Mike Smith, Matt Ritchie

Description Example data files and use cases for processing Illumina BeadArray expression data using Bioconductor

Imports beadarray (>= 2.3.18), limma, GEOquery

Suggests Biostrings, GenomicRanges, illuminaHumanv1.db, illuminaHumanv2.db, illuminaHumanv3.db

Maintainer Mike Smith <grimbough@gmail.com>

License GPL-2

biocViews MicroarrayData

git_url <https://git.bioconductor.org/packages/BeadArrayUseCases>

git_branch master

git_last_commit b15b092

git_last_commit_date 2021-10-26

Date/Publication 2022-01-23

R topics documented: