

Package ‘BeadArrayUseCases’

December 19, 2024

Version 1.44.1

Date 2024-11-13

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 RELEASE_3_20

git_last_commit ea0832f

git_last_commit_date 2024-11-13

Repository Bioconductor 3.20

Date/Publication 2024-12-19

Contents