

Package ‘BeadArrayUseCases’

June 25, 2024

Version 1.42.0

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 ($\geq 2.3.18$), 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_19

git_last_commit 05c10b5

git_last_commit_date 2024-04-30

Repository Bioconductor 3.19

Date/Publication 2024-06-25

Contents