

Package ‘BeadArrayUseCases’

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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 <Mike.L.Smith@cancer.org.uk>

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biocViews MicroarrayData

NeedsCompilation no

R topics documented: