

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 DifferentialExpression, Microarray

R topics documented: