

Package ‘BeadArrayUseCases’

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Title Analysing Illumina BeadArray expression data using Bioconductor

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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

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biocViews DifferentialExpression, Microarray

R topics documented: