

Package: BeadArrayUseCases (via r-universe)

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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 MicroarrayData

Config/pak/sysreqs libicu-dev libpng-dev libxml2-dev libssl-dev libx11-dev zlib1g-dev

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