

# Package ‘BeadArrayUseCases’

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**Version** 1.0.8

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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:**