

# Package ‘BeadArrayUseCases’

October 18, 2022

**Version** 1.34.0

**Date** 2021-07-26

**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

**git\_url** <https://git.bioconductor.org/packages/BeadArrayUseCases>

**git\_branch** RELEASE\_3\_15

**git\_last\_commit** 64c0a11

**git\_last\_commit\_date** 2022-04-26

**Date/Publication** 2022-10-18

**R topics documented:**