# 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 <grimbough@gmail.com>

**License** GPL-2

**biocViews** MicroarrayData

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