

# Package ‘BeadArrayUseCases’

January 9, 2025

**Version** 1.44.1

**Date** 2024-11-13

**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

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

**git\_branch** RELEASE\_3\_20

**git\_last\_commit** ea0832f

**git\_last\_commit\_date** 2024-11-13

**Repository** Bioconductor 3.20

**Date/Publication** 2025-01-09

## Contents