

# Package ‘BeadArrayUseCases’

May 19, 2026

**Version** 1.51.0

**Date** 2024-11-13

**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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**License** GPL-2

**biocViews** MicroarrayData

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

**git\_branch** devel

**git\_last\_commit** 3bc582d

**git\_last\_commit\_date** 2026-04-28

**Repository** Bioconductor 3.24

**Date/Publication** 2026-05-19

## Contents