

# Package ‘BeadArrayUseCases’

October 8, 2016

**Version** 1.10.0

**Date** 2012-03-17

**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 <Mike.L.Smith@cancer.org.uk>

**License** GPL-2

**biocViews** MicroarrayData

**NeedsCompilation** no

**R topics documented:**