

# Package ‘GSE13015’

October 19, 2021

**Title** GEO accession data GSE13015\_GPL6106 as a SummarizedExperiment

**Description** Microarray expression matrix platform GPL6106 and clinical data for 67 septicemic patients and made them available as GEO accession [GSE13015](<https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE13015>). GSE13015 data have been parsed into SummarizedExperiment object available in ExperimentHub. This data data could be used as an example supporting BloodGen3Module R package.

**Version** 1.1.0

**biocViews** ExperimentData, MicroarrayData, ExperimentHub

**Depends** Biobase, GEOquery

**Imports** preprocessCore, SummarizedExperiment, GEOquery, Biobase

**Suggests** ExperimentHub (>= 0.99.6), knitr, BiocStyle, rmarkdown

**License** MIT License

**VignetteBuilder** knitr

**NeedsCompilation** no

**RoxygenNote** 7.1.0

**git\_url** <https://git.bioconductor.org/packages/GSE13015>

**git\_branch** master

**git\_last\_commit** 11ea831

**git\_last\_commit\_date** 2021-05-19

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**R topics documented:**