

# Package ‘allenpvc’

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**Title** GEO accession data GSE71585 as a SingleCellExperiment

**Version** 1.0.0

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**Description** Celular taxonomy of the primary visual cortex in adult mice based on single cell RNA-sequencing from a study performed by the Allen Institute for Brain Science. In said study 49 transcriptomic cell types are identified.

**License** CC0

**NeedsCompilation** no

**Depends** R (>= 3.5.0), AnnotationHub, ExperimentHub (>= 1.7.0), SingleCellExperiment

**Suggests** BiocStyle, knitr, rmarkdown

**VignetteBuilder** knitr

**Encoding** UTF-8

**biocViews** ExperimentData, ExpressionData, SingleCellData, RNASeqData

**RoxygenNote** 6.0.1

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

**git\_branch** RELEASE\_3\_8

**git\_last\_commit** b7bed86

**git\_last\_commit\_date** 2018-10-30

**Date/Publication** 2019-04-11

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allenpvc

*Adult mouse cortical cell taxonomy by single cell transcriptomics*

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### **Description**

Cellular taxonomy of the primary visual cortex in adult mice based on single cell RNA-sequencing from a study performed by the Allen Institute for Brain Science. In said study 49 transcriptomic cell types are identified. This data set is the supplementary data from GEO accession [GSE71585](#) encapsulated in a [SingleCellExperiment](#).

### **Format**

The data is encapsulated in a [SingleCellExperiment](#) object available through [ExperimentHub](#)

### **Details**

See the vignette for examples of using these data in differential gene expression analysis.

```
browseVignettes("allenpvc")
```

Details of how this data was created are in the `inst/scripts/` directory of the source package.

### **Source**

<https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE71585>

### **References**

Tasic, Bosiljka, et al. Adult mouse cortical cell taxonomy revealed by single cell transcriptomics. *Nature neuroscience* 19.2 (2016): 335.

### **Examples**

```
allenpvc()
```

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