Package ‘FlowSorted.DLPFC.450k’

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Version 1.40.0

Title Illumina HumanMethylation data on sorted frontal cortex cell populations

Description Raw data objects for the Illumina 450k DNA methylation microarrays.

Author Andrew E Jaffe, Zachary A. Kaminsky

Maintainer Andrew E Jaffe <andrew.jaffe@libd.org>

License Artistic-2.0

Depends R (>= 2.13.0), minfi (>= 1.21.2)

LazyData yes

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**FlowSorted.DLPFC.450k**  
*Illumina Human Methylation data from 450k on sorted frontal cortex cell populations*

**Description**

This RGChannelSet contains Illumina 450k DNA methylation measurements on 58 flow-sorted dorsolateral prefrontal cortex samples from non-psychiatric controls from Guintivano et al. 2013. These samples were separated into neuronal (NeuN+) and non-neuronal (NeuN-) cell types. These data can be used by the minfi package to estimate cellular composition from bulk frontal cortex samples. This data may also be useful to individuals as example Illumina 450k data for trying preprocessing methods across a variety of Bioconductor packages.

**Usage**

```r
data(FlowSorted.DLPFC.450k)
```

**Format**

An object of class RGChannelSet.

**Details**

The `FlowSorted.DLPFC.450k` objects is based on samples assayed as part of Guintivano et al (2013). Please cite this paper, if the data is used. If you’re using this data together with the minfi package, please see the package vignette for details on how to cite that package.

**References**


**Examples**

```r
data(FlowSorted.DLPFC.450k)
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