

Package: ggsegChen (via r-universe)

August 22, 2024

Title Chen datasets for the ggseg-plotting tool

Version 1.0.01

Description This is a support package for the ggseg, and ggseg3d packages. It contains the Chen atlases to plot using functions from those two packages.

License MIT + file LICENSE

Encoding UTF-8

RoxygenNote 7.1.1

Depends R (>= 2.10)

LazyData true

Suggests ggseg, ggseg3d, ggplot2, tidyverse, knitr, rmarkdown, covr, testthat (>= 2.1.0), devtools

VignetteBuilder knitr

Remotes LCBC-UiO/ggseg, LCBC-UiO/ggseg3d

URL <https://github.com/LCBC-UiO/ggsegChen>

BugReports <https://github.com/LCBC-UiO/ggsegChen/issues>

Repository <https://ggseg.r-universe.dev>

RemoteUrl <https://github.com/ggseg/ggsegChen>

RemoteRef HEAD

RemoteSha af5c130078cea024d43b25fdf90cab857ca6f14e

Contents

| | |
|--------|---|
| chenAr | 2 |
| chenTh | 3 |

Index

4

chenAr

Genetic topography of brain region morphology

Description

Genetic topography of brain region morphology

Usage

chenAr

Format

A data.frame with 11341 observations and 11 variables

long coordinates for the x-axis

lat coordinates for the y-axis

region name of region

hemi dummy name of the hemisphere

side which side to view (sagittal)

label label name from Freesurfer

atlas name of the atlas

References

Chen et al. (2013) PNAS, 110 (42) 17089-17094; ([PubMed](#))

See Also

Other ggseg_atlases: [chenTh](#)

Examples

```
data(chenAr)
```

chenTh

Genetic topography of brain thickness morphology

Description

Genetic topography of brain thickness morphology

Usage

chenTh

Format

A data.frame with 11341 observations and 11 variables

long coordinates for the x-axis
lat coordinates for the y-axis
region name of region
hemi dummy name of the hemisphere
side which side to view (sagittal)
label label name from Freesurfer
atlas name of the atlas

References

Chen et al. (2013) PNAS, 110 (42) 17089-17094; ([PubMed](#))

See Also

Other ggseg_atlases: [chenAr](#)

Examples

```
data(chenTh)
```

Index

- * **datasets**
 - chenAr, [2](#)
 - chenTh, [3](#)
- * **ggseg_atlases**
 - chenAr, [2](#)
 - chenTh, [3](#)
- chenAr, [2, 3](#)
- chenTh, [2, 3](#)