

1 **Supplementary Information to**

2 **“Serum Neurofilament Light Concentrations are Associated with**
3 **Cortical Thinning in Anorexia Nervosa”**

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1. Materials and Methods

1.1 Participants

Healthy control participants (HC) were recruited through advertisement among middle school, high school and university students. Normal weight in HC was defined by a body mass index (BMI) between 18.5 kg/m² and 30 kg/m² (or between the 10th and the 97th age percentile, in participants younger than 18 years). Additional exclusion criteria for HC were a lifetime BMI below 17.5 kg/m² (or below the 10th age percentile when younger than 18 years) and substantial weight loss in the four weeks preceding study participation. Participants with anorexia nervosa (AN) were included if no substantial weight gain was reported in the four weeks preceding the first study point. In addition to the exclusion criteria mentioned in the main manuscript, participants in both groups were excluded if they had any history of bulimia nervosa or binge eating disorder, and any medical or neurological condition that might affect appetite, eating behavior, or body weight, any history of organic brain syndrome, dementia, schizophrenia, psychosis, or bipolar disorder. Further exclusion criteria for all participants were an IQ below 85, current substance abuse, current inflammatory, neurologic, or metabolic illness, clinically relevant anemia, pregnancy, and breast feeding. Information relevant to inclusion/exclusion criteria, including lifetime neurological disorders and possible confounding variables was obtained using the SIAB-EX (Fichter & Quadflieg, 1999), our own semi-structured interview, and medical records. Information about comorbid diagnoses in AN participants was obtained from medical records and confirmed by an expert clinician. Of the AN participants, 51 identified as European and one as Asian. In the HC group, 144 individuals identified as European, two as African and one as Asian.

Serum marker values of the complete AN sample and of a subgroup of the HC sample have been analyzed before (Doose et al., 2021; Hellerhoff et al., 2021) and the current structural magnetic resonance imaging (sMRI) sample partially overlaps with the sMRI data analyzed by Bernardoni et al. and Bahnsen et al. (Bahnsen et al., 2022; Bernardoni et al., 2016). The adequacy of the sample size for the analysis of cortical thickness (CT) was assured by consulting previous studies from our and other groups (Bernardoni et al., 2016; Cascino et al., 2020; King et al., 2015). Due to missing effect sizes

1 regarding neurofilament light (NF-L), tau protein, and glial fibrillary acidic protein (GFAP) levels in
2 AN, studies on other neurological or psychiatric disorders were consulted to assure the adequacy of the
3 sample size (Evered, Silbert, Scott, Zetterberg, & Blennow, 2018; Rohrer et al., 2016; Weston et al.,
4 2017). Study data were managed using the secure, web-based electronic data capture tool REDCap
5 (Harris et al., 2009).

6

7 **1.2 Clinical measures**

8 IQ was estimated using full or short versions of age-appropriate German versions of the Wechsler
9 Intelligence Scales for adults or children (HAWIK-IV [Petermann & Petermann, 2007] in $n = 97$; WISC-
10 IV [Petermann & Petermann, 2011] in $n = 1$; WISC-V [Petermann, 2017] in $n = 2$; WIE [von Aster et
11 al., 2008] in $n = 88$; WAIS-IV [Petermann, 2012] in $n = 6$). Short versions were used in 151 participants.
12 In five cases, IQ was missing. Missing clinical measures were not imputed.

13

14 **1.3 Blood sampling and simoa analysis**

15 Venous blood samples were collected into vacutainer tubes between 7 and 9 a.m. after an overnight fast.
16 To yield blood serum, blood samples were left to clot for 30 min at 6–8°C and then centrifuged (800 x
17 g for 15 min) in a pre-cooled (5°C) centrifuge. The samples were then aliquoted into pre-cooled
18 Eppendorf Tubes® and stored at -80° C. Determination of NF-L, tau protein, and GFAP levels was
19 carried out using the digital Simoa™ Human Neurology 4-Plex A assay in combination with the
20 Simoa™ HD-1 Analyzer (both Quanterix, Lexington, MA, USA) following the manufacturer's
21 instructions. All samples were measured in duplicates, and the mean of the two measurements was used
22 for all statistical analyses (except in one case, where only a single measurement could be obtained). The
23 mean coefficient of variation between duplicates was 6.6% (NF-L), 7.9% (tau protein), and 3.4%
24 (GFAP), respectively.

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1 **1.4 MRI acquisition and processing**

2 1.4.1 MRI acquisition

3 As in our previous studies (Bahnsen et al., 2022; Bernardoni et al., 2016), high-resolution three-
4 dimensional T1-weighted structural scans were acquired on a 3T scanner (Magnetom Trio, Siemens,
5 Erlangen, Germany) using a rapid acquisition gradient echo (MP-RAGE) sequence with the following
6 parameters: 176 sagittal slices (1 mm thickness, no gap), TR = 1900 ms; TE = 2.26 ms; flip angle = 9°;
7 voxel size = $1.0 \times 1.0 \times 1.0\text{mm}^3$, FoV = $256 \times 224\text{mm}^2$, bandwidth of 200 Hz/pixel).

8

9 1.4.2 Cortical thickness estimation

10 As in Bernardoni et al. (2016), we used standard FreeSurfer procedures to measure CT (Dale, Fischl, &
11 Sereno, 1999; Desikan et al., 2006; Fischl & Dale, 2000; Fischl et al., 2002; Fischl, Sereno, & Dale,
12 1999; Fischl et al., 2004). Surface reconstruction was first performed for each hemisphere. The quality
13 of the surface reconstruction and segmentation was assured by visual inspection by a trained examiner
14 with the support of quality assurance tools implemented in FreeSurfer
15 (<https://surfer.nmr.mgh.harvard.edu/fswiki/QATools>). All images underwent further longitudinal
16 preprocessing with the FreeSurfer longitudinal stream. Specifically, the longitudinal pipeline first
17 creates an unbiased within-subject template space and image (Reuter & Fischl, 2011) using robust,
18 inverse consistent registration (Reuter, Rosas, & Fischl, 2010). Subsequently, several processing
19 procedures are implemented including skull stripping, Talairach transformation and atlas registration in
20 addition to generation of spherical surface maps and parcellations with common information from the
21 within-subject template. The thickness at each vertex is computed as the average of two distances (Fischl
22 & Dale, 2000; Greve & Fischl, 2018): the first is the distance from each white surface vertex to their
23 corresponding closest point on the pial surface (not necessarily at a pial vertex); the second is the
24 distance from the corresponding pial vertex to the closest point on the white surface (again, not
25 necessarily at a vertex; Greve & Fischl, 2018). CT was smoothed using a Gaussian kernel with a full-
26 width-at-half-maximum (FWHM) of 10 mm.

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1 **1.5 Statistical analyses**

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3 1.5.1 Outlier exclusion

4 Extreme outliers (> 3 standard deviations (SD) from the mean of the respective diagnostic group and
 5 time point after logarithmization) were excluded from all analyses resulting in the exclusion of four NF-
 6 L concentration values ($n = 1$ acAN-TP1, $n = 1$ acAN-TP2, and $n = 2$ HC), four tau protein concentration
 7 values ($n = 1$ acAN-TP1, $n = 1$ acAN-TP2, and $n = 2$ HC), and two GFAP concentration values ($n = 1$
 8 acAN-TP2, and $n = 1$ HC).

9

10 1.5.2 Supplementary linear mixed effects model (LME)

11 To locate areas with an effect of study group (acAN vs. HC) on CT, we built a model including both
 12 acAN and HC participants. In this model, CT was estimated as

$$13 \quad CT = A + \Delta_{acAN} + B_{acAN}(b_t - b_{TP1}) + C \textit{ age}$$

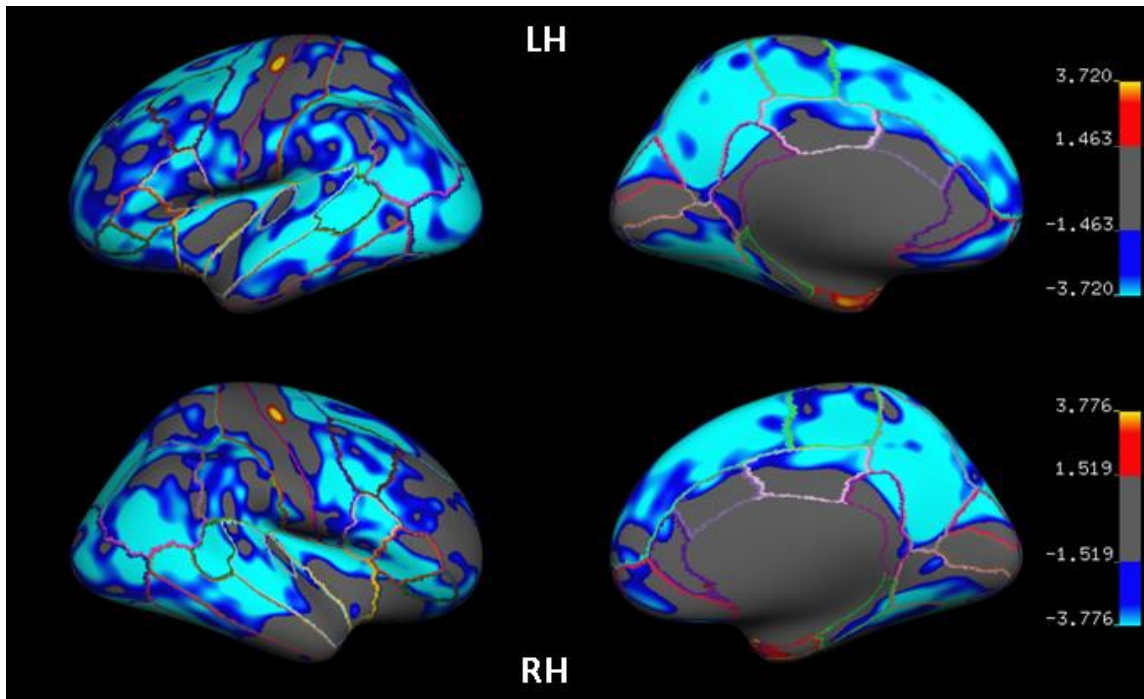
14 A in this model is the CT in HC at the mean age (of the complete sample), \textit{age} is the mean-subtracted
 15 age of the subject at the time of the scan, and C is the rate of age-related changes. b_t is BMI-SDS at time
 16 t . B_{acAN} is thus the rate of CT change associated with change in BMI-SDS. Δ_{acAN} is the difference in CT
 17 at mean age between acAN-TP1 and HC (because $(b_t - b_{TP1}) = 0$ when $t = TP1$).

1 **2. Results**

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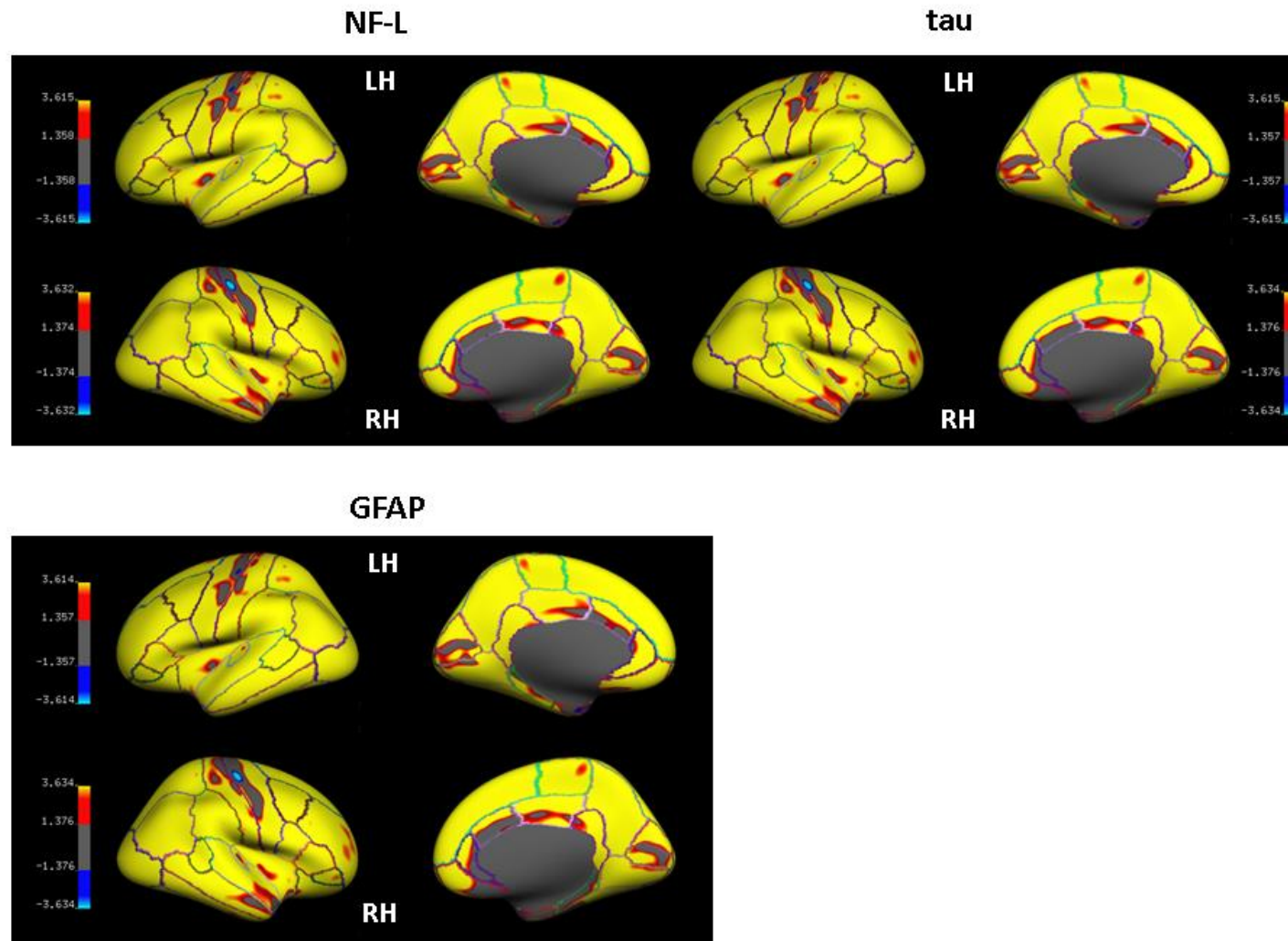
3 **2.1 Supplementary linear mixed-effect (LME) Model**

4 In the supplementary LME model including all participants, CT was strongly and mainly positively
5 associated with change in BMI-SDS in both hemispheres. Age was negatively associated with CT in
6 widespread clusters of both hemispheres, with exceptions in some small positive clusters in entorhinal,
7 temporal, postcentral, and occipital regions. Study group was negatively associated with CT in
8 widespread clusters of both hemispheres (thinner cortex in acAN compared to HC), with exceptions in
9 some small positive clusters in entorhinal/temporal, precentral and anterior cingulate regions
10 (Supplementary Figure S1).

1 **2.2 Supplementary Figures**

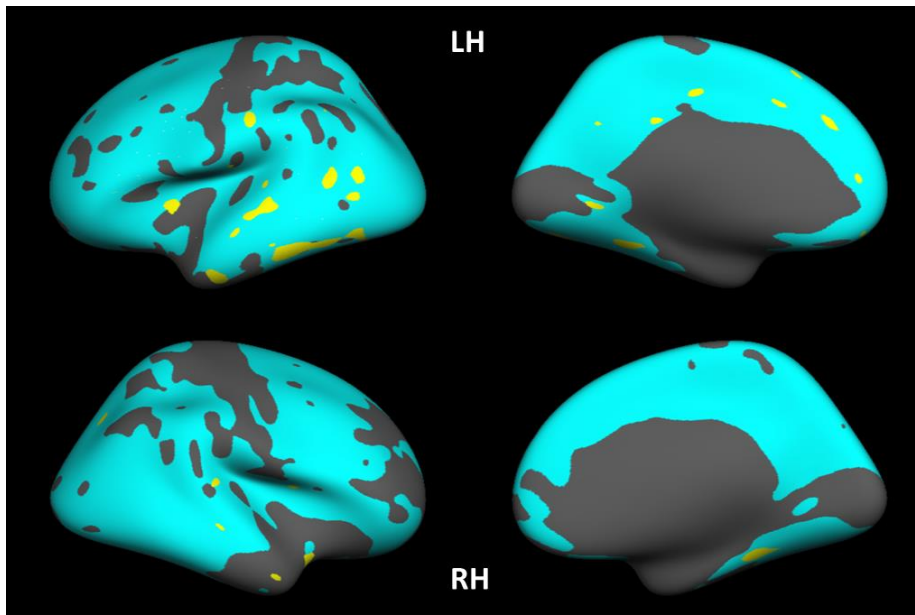
2

3 **Supplementary Figure S1.** Results of vertex-wise analysis of cortical thickness (CT) in acAN-TP1 patients
4 relative to healthy controls (HC). FDR-corrected statistical maps ($q < .05$) displaying regions in which study
5 group is associated with CT plotted on the inflated surface of a standard average subject. The color scale shows
6 p-values expressed as $-\log_{10}(p)$. Warm colors indicate an increased CT in patients with acute AN, cool colors
7 indicate a decreased CT in patients with acute AN. Abbreviations: LH, left hemisphere; RH, right hemisphere.
8 Colored outlines correspond to anatomical labels of the Desikan-Killiany atlas (Desikan et al., 2006).

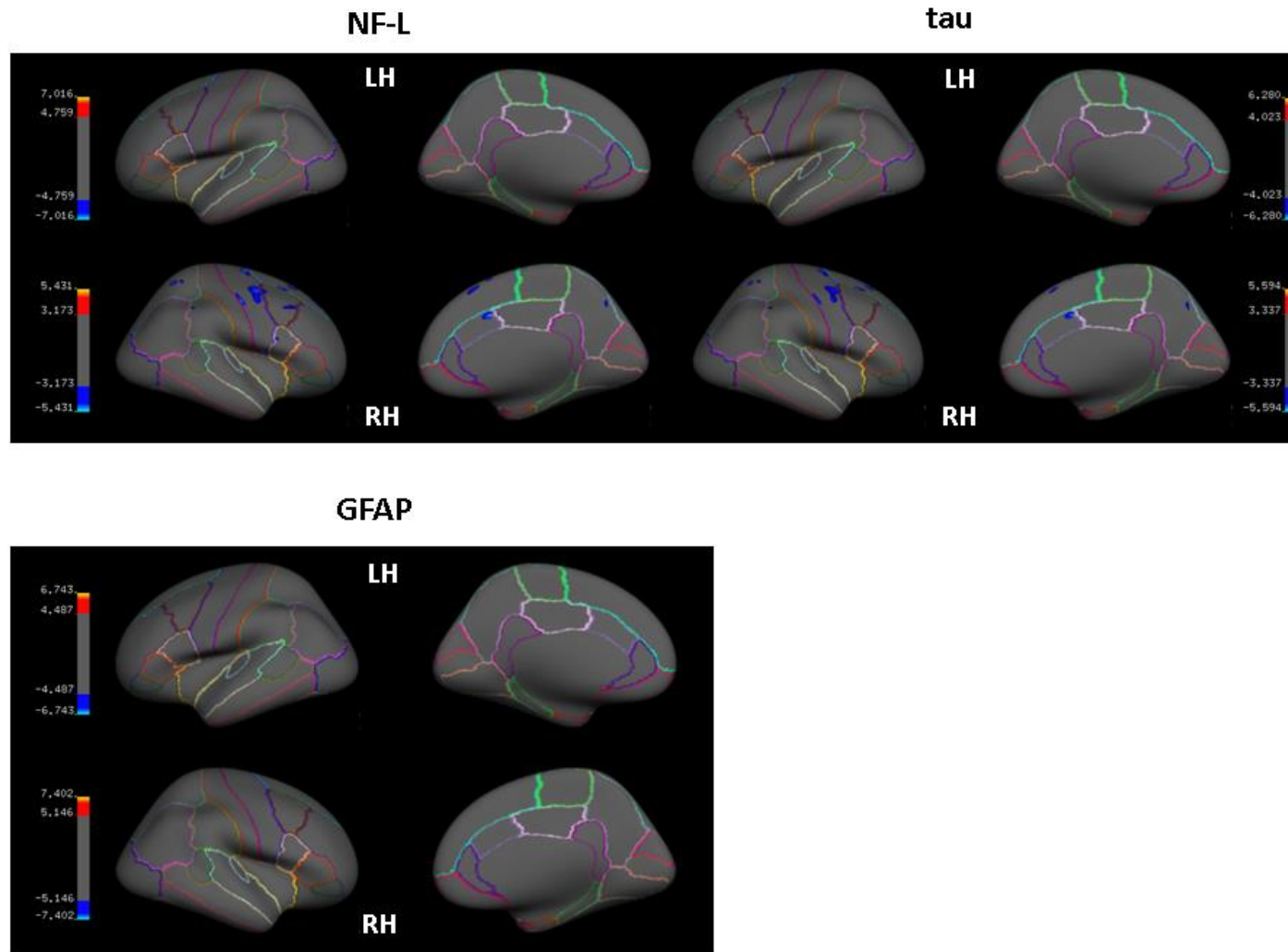


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2 **Supplementary Figure S2.** Association of change in Body Mass Index (BMI-SDS) with cortical thickness (CT) in patients with anorexia nervosa (AN) after partial weight
 3 restoration. FDR-corrected statistical maps ($q < .05$) displaying regions in which change in BMI-SDS is associated with CT plotted on the inflated surface of a standard average
 4 subject. The color scale shows p-values expressed as $-\log_{10}(p)$. Warm colors indicate an increased CT as a function of greater change in BMI-SDS. Abbreviations: LH, left
 5 hemisphere; RH, right hemisphere. Colored outlines correspond to anatomical labels of the Desikan-Killiany atlas (Desikan et al., 2006).



1 **Supplementary Figure S3.** Overlap between areas, in which cortical thickness (CT) in the supplementary
2 analyses was reduced in patients with acute anorexia nervosa (acAN-TP1) (turquoise) and areas, in which CT in
3 participants with anorexia nervosa was associated with neurofilament light levels in the main analyses (yellow).
4 For better readability, the small clusters of increased CT in acAN-TP1 are not displayed in this image.
5 Abbreviations: LH, left hemisphere; RH, right hemisphere.



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2 **Supplementary Figure S4.** Association of age with cortical thickness (CT) in patients with anorexia nervosa (AN). FDR-corrected statistical maps ($q < .05$) displaying regions in
 3 which (mean-subtracted) age is associated with CT plotted on the inflated surface of a standard average subject. The color scale shows p-values expressed as $-\log_{10}(p)$. Cool
 4 colors indicate a decrease in CT as a function of higher age. Abbreviations: LH, left hemisphere; RH, right hemisphere. Colored outlines correspond to anatomical labels of the
 5 Desikan-Killiany atlas (Desikan et al., 2006).

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