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The relationship between white matter architecture and language lateralisation in the healthy brain

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1 2	The relationship between white matter architecture and language lateralisation in the healthy brain
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42 **Abstract**

Interhemispheric anatomical differences have long been thought to be related to 43 language lateralisation. Previous studies have explored whether asymmetries in the 44 diffusion characteristics of white matter language tracts are consistent with language 45 lateralisation. These studies, typically with smaller cohorts, yielded mixed results. 46 This study investigated whether connectomic analysis of quantitative anisotropy (QA) 47 and shape features of white matter tracts across the whole brain are associated with 48 language lateralisation. We analysed 1040 healthy individuals (562 females) from 49 the Human Connectome Project database. Hemispheric language dominance for 50 each participant was quantified using a laterality quotient (LQ) derived from fMRI 51 activation in regions of interest (ROIs) associated with a language comprehension 52 task compared against a math task. A linear regression model was used to examine 53 the relationship between structural asymmetry and functional lateralisation. 54 Connectometry revealed a significant negative correlation between LQs and QA of 55 corpus callosum tracts, indicating that higher QA in these regions is associated with 56 bilateral and right-hemisphere language representation in frontal and temporal 57 regions, respectively. Left language laterality in temporal lobe was significantly 58 associated with longer right inferior fronto-occipital fasciculus (IFOF) and forceps 59 60 minor tracts. These results suggest that diffusion measures of microstructural architecture as well as geometrical features of reconstructed white matter tracts play 61 a role in language lateralisation. People with increased dependence on right or both 62 frontal hemispheres for language processing may have more developed 63

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64 commissural fibres, which may support more efficient interhemispheric

65 communication.

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Significance statement: The left cerebral hemisphere is dominant for language 68 functions in most people. In some healthy people, language functions are lateralised 69 to the right hemisphere or distributed across both hemispheres. The anatomy 70 underlying patterns of hemispheric language dominance are not well established. 71 Emerging evidence suggests that white matter connectivity and architecture is an 72 73 important feature of cortical functional organisation. In this work, we report that people who have language functions distributed across both hemispheres have 74 greater inter-hemispheric connectivity compared to lateralised people. Our findings 75 provide further insights into the anatomical basis of language function and may have 76 wider clinical implications. 77

78

79 Introduction

It has long been hypothesised that grey matter asymmetries of regions that 80 support language function may be associated with functional lateralisation of 81 language in the human brain (Güntürkün et al., 2020). Associations between cortical 82 asymmetries and hemispheric language dominance (HLD) have been noted in some 83 studies of Wada-tested patients with epilepsy (Dorsaint-Pierre et al., 2006; Foundas 84 et al., 1996, 2002; Keller et al., 2018) and healthy controls who underwent structural 85 and functional MRI (Josse et al., 2009; Keller et al., 2011). However, other studies 86 have reported no association between HLD and structural hemispheric asymmetry in 87 classical language cortical areas (Chiarello et al. 2013; Greve et al. 2013). Attention 88 89 has recently shifted towards the importance of white matter as the basis of lateralised cortical function. Some studies have reported that leftward language 90 lateralisation is associated with a greater volume of the arcuate fasciculus (Propper 91 et al., 2010) and the number of tracts in the corpus callosum (Timocin et al., 2020). 92 93 More recently, HLD has been investigated using microstructural DTI properties, such as fractional anisotropy (FA), in patient cohorts (Tantillo et al., 2016; Barba et al., 94

2020). Some studies have reported relationships between language lateralisation 95 and diffusion characteristics or the size of the corpus callosum (Tantillo et al., 2016), 96 while others did not (Barba et al., 2020). There are limited functional MRI (fMRI)-DTI 97 studies on language lateralisation in healthy individuals. Tractography studies have 98 reported associations between fMRI-determined left HLD and FA of the left arcuate 99 fasciculus (James et al., 2015; Perlaki et al., 2013; Powell et al., 2006; Silva & 100 Citterio, 2017) and corpus callosum (Häberling et al., 2011). However, other studies 101 have not reported relationships between the side or extent of HLD and conventional 102 103 diffusion-based tract characteristics (Karpychev et al., 2022; Vernooij et al., 2007), as well as more sophisticated diffusion MRI measures, such as fibre density cross-104 section (Verhelst et al., 2021). Inconsistencies between studies may be due to 105 methodological differences, including differences in tractography approaches, study 106 designs, patient characteristics, and sample sizes. 107

108 In the present study, we adopted two complementary approaches that potentially overcome some of the methodological shortcomings of previous 109 tractography studies. First, we employed a connectometry approach based on an 110 local analysis of diffusion properties, which uses permutation testing to identify group 111 differences along white matter tracts. This whole-brain approach employs 112 correlational tractography to identify specific subcomponents of white matter tracts 113 that exhibit anisotropy correlated with a predefined variable of interest with superior 114 sensitivity and specificity compared to traditional voxel-based analyses (Yeh et al., 115 2016). Connectometry has recently been used to uncover structural disparities 116 between bilingual and non-bilingual individuals (Rahmani et al., 2017) and to identify 117 structural pathways linked to enhanced language capabilities in individuals with 118 aphasia (Hula et al., 2020; Dresang et al., 2021) and preterm-born children (Barnes-119 Davis et al., 2020, 2022). Second, we employed shape analysis to investigate the 120 geometrical characteristics of white matter tract bundles that comprise the integral 121 components of language networks. This approach captures fundamental shape 122 characteristics, such as volume and surface area, and encompasses advanced 123 morphological properties including white matter bundle curl, elongation, length, span, 124 and diameter (Yeh, 2020). Previous studies have already demonstrated, through the 125 utilization of virtual dissections (Catani et al., 2007) and shape analysis employing 126 127 tractography algorithms (Yeh et al., 2020), that the leftward morphometric

asymmetries of language-associated white matter tracts exist in people without 128

- known hemispheric language dominance (HLD). Whether white matter 129
- interhemispheric asymmetries change in people with atypical HLD remains unclear. 130

The first objective of the present study was to conduct diffusion 131 connectometry analysis in a large cohort of healthy individuals who underwent 132 language fMRI to determine whether microstructural properties of white matter tracts 133 are related to HLD. The second objective was to explore whether interhemispheric 134 135 shape asymmetries of white matter tracts are related to language lateralisation in the Nanus same individuals. 136

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Methods 138

Study data and participants 139

All data were acquired from the Human Connectome Project (HCP) 140 (http://www.humanconnectome.org/) open-access data initiative offering high-guality 141 142 anatomical and functional MRI of the human brain. We used the HCP Young Adults (HCP-YA 1200 Subjects) data release as it contains a large sample of healthy adults 143 for whom both language task fMRI and diffusion MRI sequences were acquired. The 144 dataset comprised 1200 healthy adults, aged 22-35 years. Each participant 145 underwent an identical imaging protocol acquired on the same MRI scanner. 146 Individuals with neuropsychiatric or neurologic disorders, diabetes, high blood 147 pressure, premature birth, and severe symptoms associated with substance use 148 149 were excluded from data collection (Van Essen et al., 2013). The present study focused on language fMRI and diffusion MRI data only. Individuals were only 150 selected for inclusion if they had fMRI data available for the language story task (see 151 below) and had corresponding 3T diffusion MRI data. This resulted in a sample size 152 of 1040 participants (562 females), with a mean age of 28.74 (SD = 3.69) years. 153 According to the Edinburgh Handedness Inventory (Oldfield, 1971), 962 (92%) 154 155 participants preferred their right hand, scoring at least 10 on a scale of -100 (left) to 100 (right). Eighty-five participants preferred left, scoring below -10, and two were 156 157 ambidextrous, scoring zero.

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159 Data acquisition

HCP data were acquired on a Siemens 3T Skyra system, with 32-channel 160 (SC72) head coil. Task fMRI data were collected using gradient-echo echo-planar 161 imaging (EPI) with an isotropic resolution of 2.0 mm (TR = 720ms, TE = 33.1ms, 162 matrix = 104x90, 72 slices, flip angle = 52°, BW = 2290 Hz/Px, FOV = 208 x 180 163 mm, 72 slices, multiband accelerator factor = 8) (Marcus et al., 2013). The HCP 164 dMRI data were acquired using three shells (b=1000, 2000 and 3000 s/mm2) with 90 165 diffusion gradient directions and five b₀ volumes with RL phase encoding direction 166 $(TE = 89.5ms, TR = 5520ms, flip angles = 78/160^\circ, isotropic voxel size = 1.25 mm3,$ 167 multiband factor = 3) (Sotiropoulos et al., 2013). 168

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170 Language paradigm

The language comprehension task used in Human Connectome Project was 171 designed by Binder and colleagues (2011). The task consists of two 3.8-minute runs. 172 Each run has four blocks of story tasks alternating with four blocks of math tasks. 173 The story and math tasks are matched in terms of length, word and phoneme rate, 174 speaking style, and prosodic features. The story blocks present subjects with 5-9 175 auditory sentences, followed by questions about the content of the story. The math 176 177 task requires participants to perform arithmetic operations followed by equals and two choices. Since arithmetic tasks do not engage temporal lobe activity (Baldo & 178 Dronkers, 2007), we decided to use a STORY-MATH contrast, as it effectively 179 isolates regions responsible for language comprehension without "masking" temporal 180 lobe activity. Additionally, temporal lobe is involved in high-level processes of normal 181 consciousness (Spitsyna et al., 2006), thus we avoided passive tasks as a baseline 182 to reduce the risk of masking activities in this region (which is essential for language 183 comprehension). This contrast allowed us to cancel out the regions that are jointly 184 activated in both tasks (such as low-level auditory and phonological input), isolating 185 the regions involved in narrative processing including semantic and non-speech 186 related aspects of language, theory of mind, and inference processing. 187

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189 fMRI preprocessing and analysis

The preprocessed task fMRI data were retrieved from HCP database 190 (https://db.humanconnectome.org). The HCP pre-processing included fMRIVolume 191 and *fMRISurface* pipelines, which were primarily built using tools from FSL 192 (Jenkinson et al., 2012; http://www.fmrib.ox.ac.uk/fsl), Freesurfer (Fischl, 2012) and 193 the HCP Workbench (Marcus et al., 2013). Details of the pre-processing steps have 194 been described previously (Glasser et al., 2013). The goal of the first fMRIVolume 195 pipeline was to generate 4D whole-brain timeseries. This was accomplished by (1) 196 removing spatial distortions by gradient non-linearity distortion correction, (2) 197 198 realigning volumes using rigid-body motion correction using a single-band reference image as the target, and (3) estimating (using FSL toolbox "topup") and correcting 199 field map-based EPI distortions. The resulting EPI data was (4) registered to T1-200 weighted scan, then (5) non-linearly (FNIRT) into Montreal Neurological Institute 201 (MNI) space, and (6) blood-oxygen-level-dependent (BOLD) signal intensity was 202 normalized by the average. This process resulted in individual subjects being 203 mapped with a notable degree of left-right symmetry (Elam et al, 2021), which aligns 204 with laterality research recommendations (Vingerhoets et al., 2023). 205

The goal of *fMRISurface* pipeline was to transform the resulting 4D timeseries 206 to Connectivity Informatics Technology Initiative (CIFTI) grayordinate space, 207 encompassing cortical, subcortical, and cerebellar grey matter collectively (Pham et 208 al, 2022). This was accomplished by mapping fMRI data within cortical grey matter 209 ribbon onto the native cortical surface, registering it into CIFTI grayordinate space 210 (surface representation with 32,492 vertices on each hemisphere), and mapping the 211 set of subcortical grey matter voxels from each subcortical parcel in each individual 212 to a standard set of voxels in each atlas parcel, resulting in 2mm average surface 213 vertex and subcortical volume voxel spacing. Finally, grayordinate space data was 214 smoothed using Gaussian kernel. 215

We used a fully processed task-based STORY-MATH fMRI activation
 Contrast Of Parameter Estimates (COPE) map, which was generated by FSL FEAT
 and is readily available on https://db.humanconnectome.org as part of the "S1200
 Subjects" dataset. Considering the spatial heterogeneity of the individual brain
 scans, the MSM-All (Multimodal Surface Matching) registered dataset was used,
 which uses information on areal features derived from the resting state network,
 myelin maps, and alignment of folding. The motivation for using MSM-All over MSM-

Sulc (cortical folding-based registration) came from previous studies that
demonstrated a weaker correlation between sulcal depth and local curvature with
regions responsible for higher cognitive functions, including Broca's area (Fischl et
al., 2008; Van Essen, 2005), compared to the MSM-All registration, which showed
improved cross-subject alignment of independent task fMRI datasets (Robinson et
al., 2018).

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230 Language Comprehension Laterality Quotient

Grayordinates localised regions-of-interest (ROIs) on the "inflated" brain 231 surface (Van Essen & Glasser, 2016). A laterality quotient (LQ) was calculated to 232 assess HLD for each participant's task fMRI activation using the CIFTI toolbox in 233 MATLAB in ROIs associated with language comprehension. The analyses were 234 conducted separately for frontal and temporal regions, given the well-documented 235 phenomenon of crossed language dominance, where a participant may exhibit 236 dominance in one hemisphere for frontal regions and the opposite hemisphere for 237 temporal regions (Seghier, 2008). For the frontal ROIs, Brodmann areas 44 and 45 238 were selected due to their established high reliability in determining language 239 dominance during semantic tasks (Sabbah et al., 2003; Seghier et al., 2008). In our 240 temporal lobe laterality analyses, ROIs were chosen within the anterior temporal lobe 241 242 (TGd, TGv, TE1a, TE2a, STGa, STSva, STSda) because these areas have been shown to be heavily involved in language comprehension (Binder et al., 2011) 243 (Figure 1). 244

To work with CIFTI files, we generated dscalar files for each ROI using 245 wb_command, imported them into MATLAB, and extracted z-values from ROIs using 246 the CIFTI toolbox. The z-values were thresholded for each participant by including 247 only grayordinates with values greater than the median in each ROI (Dietz et al., 248 2016). To account for the unequal number of grayordinates between the left and 249 right hemisphere (approximately 100 more grayordinates on the left than the right), 250 we corrected for these regional differences to ensure that comparisons between 251 hemispheres were not skewed by differences in their sizes. This adjustment involved 252 dividing the total sum of thresholded z-values by the number of grayordinates in 253 each hemisphere for both frontal and temporal ROIs separately. The laterality 254

quotient (LQ) was then computed for each participant's normalised z-values usingthe equations below:

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$$LQ = \frac{(L-R)}{\max(L,R)}$$

259

Where L represents the normalised z-values in the left, and R in the right ROI. 260 We chose to employ an innovative LQ formula based on the Jaccard-Tanimoto index 261 to provide a more sophisticated approach in evaluating and classifying language 262 lateralisation (Seghier, 2019). This revised formula defines LQ as a metric of 263 distance that adheres to a consistent distribution pattern, thus enhancing its 264 sensitivity towards hemisphere activity differences, accentuating the distinctions 265 between the two hemispheres. The values above +1/3 indicate left language 266 dominance (LLD), values below -1/3 indicate right language dominance (RLD), and 267 values between -1/3 and +1/3 indicate bilateral language representation (BLR), 268 269 ensuring an equal cumulative probability in each dominance category.

270

271 Diffusion processing

Diffusion data was downloaded from the HCP S1200 Young Adult Data 272 Release and preprocessed using the HCP Diffusion preprocessing pipeline using 273 FMRIB diffusion toolbox in FSL. Briefly, the pipeline included b₀ image intensity 274 normalisation, removing EPI susceptibility-induced field distortions with FSL's "topup" 275 algorithm (Andersson et al, 2016), correcting for eddy current distortions, head 276 277 movements, and gradient nonlinearities (Glasser et al., 2013). Quality control of the preprocessed diffusion MRI data was performed using DSI studio software 278 (http://dsi-studio.labsolver.org). An automatic quality control routine then checked the 279 280 b-table to ensure its accuracy (Schilling et al., 2019). The diffusion data were coregistered in MNI space using q-space diffeomorphic reconstruction (Yeh & Tseng, 281 2011) to obtain the spin distribution function (SDF) with a recommended length ratio 282 of 1.25, as specified in the original study (Yeh et al., 2010) (Figure 2). 283

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285 Connectometry analysis

We applied a whole-brain group connectometry analysis using DSI Studio as 286 287 described in previous applications (Barnes-Davis et al., 2020, 2022; Dresang et al., 2021; Rahmani et al., 2017) to study the relationship between regional white matter 288 quantitative anisotropy (QA) and language lateralisation measures derived from LQs 289 (Figure 2). The connectometry approach derives the QA measure from the SDF in 290 each fibre orientation, which defines the number of anisotropic spins along that 291 direction in each streamline (Yeh et al., 2010, 2013). The anisotropy in each section 292 293 of a white matter tract is then correlated with the study variable (Yeh et al., 2016). Unlike a voxel-based FA metric, which attributes identical anisotropy values to all 294 295 fibre orientations within a voxel, QA demonstrates a discerning capability by identifying specific axonal orientations in each peak orientation of the SDF (Yeh et al, 296 2013). 297

Our connectometry analyses were conducted in two phases: the initial phase 298 focused on examining the lateralisation of frontal regions during language 299 comprehension, and the subsequent phase investigated temporal regions. Initially, 300 301 connectometry analyses were performed on all participant groups concurrently, followed by post-hoc analyses on three distinct groups separately to aid in 302 interpretation and capture varying effects related to different degrees of laterality. 303 304 Specifically, the first post-hoc analysis included participants with LLD and BLR, the second consisted of participants with RLD and BLR, and the third included 305 306 individuals with both LLD and RLD. The linear effect of handedness, sex, and age was mitigated using a partial linear correlation. A nonparametric Spearman partial 307 correlation was used to derive the continuous segments correlating with a LQ (Yeh 308 309 et al., 2016). Each reconstructed white matter tract within a voxel was tracked to extract a QA map for each participant (Yeh et al., 2013). A T-score threshold was 310 assigned to the highest level of three to reduce the possibility of false positive results 311 (Ashraf-Ganjouei et al., 2019). The tracks were filtered by topology-informed pruning 312 with sixteen iterations to remove implausible spurious connections (Yeh et al., 2019). 313 Given the large sample size in our study, and to prevent false positives, a 314 conservative false discovery rate (FDR) correction for multiple comparisons was 315

employed with a threshold of 0.01 to select tracks showing significant associations
between LQ and QA. To estimate the false discovery rate, 5000 randomized
permutations were applied to the group label to obtain the null distribution of the
track length. After the correlational results were obtained, additional categorical
analyses were performed at the group level (LLD/RLD, LLD/BLR, RLD/BLR). Short
tracts (<20mm) were removed for easier interpretation of our results.

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323 Shape analysis

The SDF maps generated from the connectometry analysis were used for 324 tract shape analysis and automatic fibre tractography was performed using a 325 deterministic fibre tracking algorithm utilising DSI studio software (Yeh, 2020). 326 Eleven white matter tract bundles that are part of language comprehension networks 327 (Friederici et al., 2007; Harvey et al., 2013; Ivanova et al., 2021; Rollans & Cummine, 328 2018; Shin et al., 2019; Zhong et al., 2022; Forkel et al, 2022) were then 329 automatically tracked and recognised based on the HCP-842 tractography atlas (Yeh 330 et al., 2018) (Figure 3). These include the arcuate fasciculus (AF), corpus callosum 331 body, corpus callosum forceps major (splenium), corpus callosum forceps minor 332 (genu), inferior fronto-occipital fasciculus (IFOF), frontal aslant tract (FAT), inferior 333 longitudinal fasciculus (ILF), the three branches of the superior longitudinal 334 fasciculus (dorsal SLF1, middle SLF2 and ventral SLF3), and the uncinate 335 fasciculus. All white matter bundles were independently tracked within the left and 336 right hemispheres, while the corpus callosum bundles were tracked as a whole. The 337 diffusion sampling length ratio was set at 1.25 and the output resolution was 338 resampled to 2 mm isotropic. To remove false connections, topology-informed 339 pruning was applied with 32 iterations (Yeh et al., 2019). We decided to exclude 340 participants for whom we could not reconstruct at least one of their ROI bundles. As 341 a result, 290 participants were excluded, leaving us with a final sample size of 750 342 participants. Finally, after identifying all white matter tracts of interest, the following 343 shape metrics were extracted: tract length, span, curl, elongation, diameter, volume, 344 and surface area were extracted (Figure 4). 345

To evaluate the statistical significance of differences among various laterality groups (LLD, RLD, BLR), we conducted an analysis of variance (ANOVA). This

- analysis utilized the same laterality groupings based on frontal and temporal ROIs
- and included covariates consistent with those used in the connectometry analyses.

All computations were performed using R (version 4.4.1).

351

352 **Results**

353 fMRI

Activations in frontal ROIs revealed a weak leftward lateralisation on the 354 group level (LQ=0.33±0.31), while BOLD activations in anterior temporal lobe ROIs 355 showed a more bilateral pattern (LQ=0.17±0.2). Based on the frontal ROIs of the 356 fMRI language comprehension task, 581 participants were classified as left-357 hemisphere dominant (56%; LQ=0.53±0.16), 426 as bilateral (41%; LQ=0.13±0.17), 358 and the remaining 33 as right-hemisphere dominant (3%; LQ=-0.64±0.17). For 359 temporal ROIs, only 193 participants (19%; LQ=0.41±0.09) were left-hemisphere 360 dominant, while 80% (n=833) were classified as bilateral (LQ=0.13±0.14). Only 14 361 participants (1%; LQ=-0.58±0.28) were right lateralised. In both laterality groups, 362 strong neural activity is observed in brain areas associated with semantic processing 363 i.e. anterior, and posterior temporal lobes as well as in the left inferior frontal gyrus 364 (Jackson, 2021) (Figure 5). In line with previous reports that have used similar 365 story/narrative materials, also the ventral angular gyrus bordering the 366 temporoparietal junction was engaged during language comprehension (Lerner et al, 367 2011; Branzi et al, 2020; Branzi et al., 2021; Humbphreys & Lambon Ralph, 2015). 368

369

370 Connectometry

Two connectometry analyses were conducted. Firstly, in the investigation of 371 language lateralisation in frontal regions, a total of 2028 white matter tracts exhibiting 372 a significant negative correlation with LQ (n=1040; p<0.01, FDR corrected) were 373 identified. Notably, the majority of these tracts were commissural, constituting 82% of 374 the identified tracts, including the forceps minor (63%) and corpus callosum body 375 (19%). A smaller portion of these tracts were positioned in bilateral fornix (18%). 376 377 Conversely, 218 tracts were linked to higher frontal LQ values, all of which were situated in bilateral cingulum (n=1040; p<0.01, FDR corrected). Subsequently, 378

categorical post-hoc analyses were conducted to further illustrate the laterality 379 groups influencing significant differences, with a specific focus on the disparities 380 between LLD and BLR, RLD and BLR, and RLD and LLD. Within the cohort of 381 bilateral and left lateralised individuals (n=1007), it was observed that individuals with 382 left lateralisation exhibited higher QA in 2126 tracts, predominantly in the forceps 383 minor (48%), corpus callosum body (38%), and bilateral fornix (8%), implying that the 384 negative correlations in the primary findings were mainly driven by individuals with 385 bilateral language dominance (p<0.01, FDR corrected). The remaining streamlines 386 387 consisted of the right IFOF, right AF, and middle cerebellar peduncle (MCP). No tracts were found to be associated with left lateralised individuals. In the analysis 388 encompassing individuals with bilateral and right hemisphere dominance (n=459), 389 only 20 streamlines in the forceps minor exhibited higher QA in individuals with right 390 hemisphere dominance compared to those with bilateral dominance (p<0.01, FDR 391 corrected). No significant differences were observed between LLD and RLD (n = 392 614; p<0.01, FDR corrected). 393

The second connectometry analysis revealed a significantly negative 394 correlation between anterior temporal lobe LQ and QA in 2,408 tracts (n=1040; 395 p<0.01, FDR corrected). These tracts were predominantly located in the corpus 396 callosum body (67%), with additional distributions in the left corticospinal tract (9%), 397 left cingulum (7%), left medial lemniscus (5%), right dentarubrothalamic tract (3%), 398 and bilateral AF (3%). Additionally, 118 tracts showed a positive correlation between 399 QA and higher LQ in the anterior temporal regions, all of which were located in the 400 forceps minor (n=1040; p<0.01, FDR corrected). Categorical post-hoc analyses 401 found no significant differences between LLD and BLR (n=1029; p<0.01, FDR 402 403 corrected) or between RLD and BLR (n=847; p<0.01, FDR corrected). However, LLD and RLD comparison (n=210) identified 391 streamlines with higher QA in RLD 404 compared to LLD. These streamlines were distributed in the forceps minor (29%), 405 corpus callosum tapetum (27%), bilateral arcuate fasciculus (26%), and right IFOF 406 (8%). 407

408

409 Shape analysis

ANOVA analysis for language lateralisation in anterior temporal lobe showed 410 that mean length of right IFOF was significantly different between three laterality 411 groups ($F_2 = 9.8$; p = 0.005, FDR corrected). Tukey post-hoc tests showed that left 412 lateralised individuals had longer right IFOF compared to people with BLR (p=0.01, 413 FWE corrected). Another tract that showed significant difference was forceps minor, 414 with different mean lengths between laterality groups (F₂=10.1; p=0.005, FDR 415 corrected). Similarly, Tukey post-hoc tests showed that left lateralised people had 416 longer forceps minor compared to bilateral individuals (p=0.01, FWE corrected) 417 418 (Figure 7). No significant differences between frontal laterality groups were found in Manu 419 relation to shape metrics.

420

Discussion 421

Biological implications 422

We report significantly increased QA of the corpus callosum in individuals who 423 had lower LQ in both frontal and temporal ROIs. Post-hoc analyses revealed that this 424 result was driven by individuals with both BLR and RLD when laterality quotient was 425 calculated using frontal regions, and by RLD when laterality quotient was based on 426 the anterior temporal lobe. This observation aligns with prior research in patient 427 cohorts (Tantillo et al., 2016) and supports the hypothesis that individuals who rely 428 more on both cerebral hemispheres for language processing may have more 429 developed commissural fibres. These fibres could potentially serve as a 430 compensatory mechanism for the heightened metabolic energy requirements 431 associated with information transfer and the reduction of transmission times 432 (Laughlin & Sejnowski, 2003). An alternative hypothesis suggests that using both 433 hemispheres for language processing is more natural than previously thought 434 (Newport et al., 2022). This implies that the shift of language function towards the left 435 hemisphere might result from less developed commissural tracts. Well-developed 436 commissural tracts may facilitate bilateral language processing in frontal regions by 437 allowing the right hemisphere to participate fully in language tasks (Newport et al., 438 2017). A higher forceps minor anisotropy in right lateralised people compared to 439 people with LLD and BLR (Figures 6-1, 7-1) suggests that commissural tracts may 440

441 not necessarily enhance the function of both hemispheres but rather maintain right442 hemisphere involvement in language comprehension tasks.

Other imaging research has also suggested the importance of the corpus 443 callosum for language lateralisation, although the link between structural measures 444 of the corpus callosum and HLD is unclear. One study reported a greater FA of the 445 whole corpus callosum in people with atypical language lateralisation (defined as 446 RLD and BLR together) and, consistent with our results of anterior temporal lobe 447 lateralisation, greatest anisotropy in people with RLD (Häberling et al., 2011). We 448 found a positive correlation between a small segment of the left forceps minor and 449 450 temporal LQ, suggesting a more complex role of the forceps minor than previously understood (Figure 6). This finding was supported by shape analysis, which revealed 451 452 a longer forceps minor in individuals with left temporal language lateralisation. This aligns with a recent study using fixel-based analysis, which reported greater left fibre 453 bundle cross-section asymmetry of the forceps minor in individuals with left language 454 dominance (LLD) and higher asymmetry of the right forceps minor in those with RLD 455 (Verhelst et al., 2021). Thus, while the body of the corpus callosum is associated 456 with atypical (both bilateral and right) lateralisation for language comprehension in 457 our study, the posterior and anterior parts of the corpus callosum are mainly 458 associated with lateralised language, whether right or left, consistent with previous 459 studies (Karpychev et al., 2022; Westerhausen et al., 2006). Although commissural 460 fibres have shown relationships with language lateralisation in both previous and 461 current studies, there is inconsistency regarding which individuals exhibit greater 462 microstructural differences inferred by diffusion scalar metrics. This inconsistency 463 may stem from methodological factors, such as the inclusion or exclusion of 464 individuals with BLR and the type of language lateralisation assessed. 465

Our study reported a significant association between asymmetry in frontal 466 activation and QA in the fornix bilaterally, particularly in individuals exhibiting an LQ 467 closer to zero, indicating a more bilateral hemispheric representation. While there is 468 469 limited existing literature on the relationship between HLD and diffusion measures in these tracts, some have highlighted the involvement of them in language processing 470 471 (Hula et al., 2020; Sihvonen et al., 2021). Our findings related to temporal lobe laterality revealed a small number of streamlines in the AF bilaterally exhibiting 472 higher anisotropy in individuals with RLD compared to LLD, but not in those with 473

BLR compared to LLD. This suggests that increased QA in these regions may be
linked to right-hemisphere language lateralisation. Previous research has also found
associations between AF asymmetry and language lateralisation in several cohorts
of healthy individuals (Ocklenburg et al., 2013; Propper et al., 2010), although the
exact relationship between AF asymmetry and HLD remains unclear (Gerrits et al.,
2022; Verhelst et al., 2021). Further studies are needed to better understand these
tracts' importance for language lateralisation.

481

482 Methodological considerations

To investigate the relationship between language lateralisation and white 483 matter characteristics, we used both correlational methods and traditional categorical 484 classifications (left, right, bilateral). Analysing raw LQ values for the entire sample 485 helped reduce the subjectivity inherent in categorical groupings (Wegrzyn et al., 486 2019; Westerhausen et al., 2006). Instead of relying solely on global BOLD 487 activation patterns to assess language lateralisation, we conducted two distinct 488 studies focusing on the frontal and anterior temporal lobes. This approach allowed 489 us to obtain more detailed insights. For instance, whilst lower LQs in both frontal and 490 temporal regions were associated with increased anisotropy in the corpus callosum, 491 frontal regions engaged a larger portion of the anterior corpus callosum body and the 492 forceps minor. Interestingly, lower LQs in the temporal lobe were linked to increased 493 494 anisotropy in the bilateral AF, which suggested a right hemisphere dominance, as indicated by post-hoc analyses—an effect not observed in the frontal lobe. 495 496 Furthermore, while frontal LQs were positively correlated with fractional anisotropy in the bilateral cingulum, temporal LQs were associated with increased QA in the left 497 forceps minor. Analysing language lateralisation separately for the frontal and 498 499 temporal regions thus provided richer and more nuanced insights into the 500 relationship between language comprehension and white matter characteristics.

501 The number of individuals with right hemisphere dominance, especially for 502 anterior temporal region, is small in the present study compared to previous studies 503 (Chang et al., 2011). This disparity is primarily attributed to the task used to assess 504 HLD, which entailed semantic processing, resulting in more bilateral fMRI activations 505 (Binder et al., 2011; Metoki et al., 2022; Walenski et al., 2019). This might be

because comprehension tasks rely less on the left-lateralised dorsal pathway, 506 strongly associated with cognitive operations crucial for language production (e.g., 507 retrieval and production of speech sounds) (Hickok & Poeppel 2007). Other non-508 mutually exclusive explanations include differences in the complexity of the linguistic 509 stimuli (single words typically used in fluency tasks versus sentences typically used 510 in comprehension tasks) and the demands that these tasks typically involve (see 511 Peelle, 2012). For example, the verbal stimuli used for the stories likely portrav 512 social concepts (e.g., theory of mind, intention, emotion, morality) and/or they may 513 514 contain different amounts of metaphor, idiom, or implied meaning. All these aspects have been associated with the recruitment of fronto-temporal and parietal regions in 515 the right hemisphere (Miller et al. 1997; Olson et al. 2007; Yang, 2014; Schmidt et 516 517 al., 2007).

The fact that we employed a language comprehension task might explain why 518 519 only a small subset of participants demonstrated strong leftward lateralisation, while the majority displayed mild lateralisation. A recent systematic review comparing 520 different language tasks has highlighted that language production tasks may be 521 522 more robust in accurately assessing language laterality than language comprehension tasks (Bradshaw et al., 2017). For instance, there is evidence that 523 between 6 and 24 years of age, there is an increase in frontal asymmetry during 524 tasks involving the articulation of words. However, this asymmetry is not present 525 during story listening. This suggests partly different maturational mechanisms 526 between language comprehension and production (Lidzba et al., 2011; Berl et al., 527 2010). Future studies will have to use tasks that typically generate strongly left 528 lateralised neural responses such as verbal fluency tasks to corroborate our findings. 529

530 Our methodology offers several practical benefits. Firstly, despite the limited proportion of right hemisphere dominant individuals identified through the functional 531 532 language task, the sizeable sample size (N= 1040) employed in this study is the largest ever utilised in such investigations, thus potentially decreasing the likelihood 533 534 of false positives. Secondly, we employed two complementary methodologies, namely connectometry and tractography, to delineate the precise white matter 535 536 characteristics associated with HLD. In doing so, we have provided new insights into the anatomical basis of language lateralisation insomuch that white matter tract 537 geometrical features (shape analysis) are unrelated to HLD at the group-level. 538

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540 Conclusion

- The findings of our study suggest that measures of diffusion-based 541 542 microstructural architecture of reconstructed white matter tracts, are linked to language lateralisation. Specifically, individuals who exhibit a greater reliance on 543 both cerebral hemispheres for language comprehension may possess more highly 544 developed CC body fibres, thereby promoting more efficient interhemispheric 545 communication. The involvement of anterior and posterior parts of the CC in 546 asymmetrical temporal lobe activity (i.e., either left or right) for language 547 comprehension unveils a more complex and nuanced role of the forceps minor. 548 Future research should further investigate these relationships, employing tasks that 549 550 typically generate strongly left-lateralized neural responses, to validate and expand
- 551 upon our findings.

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- 925 **Figure 2.** Flowchart of the methods pipeline. The pre-processed diffusion MRI data was
- 926 reconstructed in an MNI space. The outputs of the reconstruction and SDFs were calculated
- to obtain the fibre orientations using DSI studio. Then, two different approaches were used
- to examine the white matter tracts associated with language laterality. The connectometry
- approach involved obtaining a local connectome matrix and finding out its association with
 LQ. Shape analysis involved the recognition of the WM tracts using HCP atlas and mapping
- eleven WM fibre bundles important for language function. The measures of key shape
- 932 features, such as curl and volume, were extracted and linear regression analyses were used
- 933 to look at the associations between shape metrics and LQ.
- Figure 3. Eleven white matter tracts were reconstructed for shape analysis based on the
 HCP842 atlas computed on 1065 healthy people (Yeh et al, 2018).
- **Figure 4.** Schematic illustration of the shape analysis of the white matter tracts. (a) The area metrics used in the included surface area (mm). (b) The length metrics used in the study included mean tract length (mm) as well as span bundle (mm) and diameter (mm) of the bundle. (c) The volume metrics used in the study included branch volume (mm3) (blue
- dotted line), trunk volume (mm3) (right dotted line), and total bundle volume (mm3) (black
- 941 dotted line) (d) The shape metrics used in the study included curl and elongation.
- Figure 5. Cohen's d maps of language comprehension task. The colour bar indicates Z
 scores; L left hemisphere; R right hemisphere.
- Figure 6. Connectometry results for the language lateralisation in frontal lobe (n=1040). a)
 Tract sections negatively correlated with LQ (p<0.01, FDR corrected). b) Tract sections
 positively correlated with LQ (p<0.01, FDR corrected). Abbreviations: CC, corpus callosum.
 Colour bar represents t-statistic. See Extended Figure 6-1 for the group level posthoc
 analyses.
- Figure 7. Connectometry results for the language lateralisation in temporal lobe (n=1040). a)
 Tract sections negatively correlated with LQ (p<0.01, FDR corrected). b) Tract sections
 positively correlated with LQ (p<0.01, FDR corrected). Abbreviations: AF -arcuate
 fasciculus; CC, corpus callosum; CS corticospinal; DRT dentatorubrothalamic. Colour bar
- 953 represents t-statistic. See Extended Figure 7-1 for the group level posthoc analyses.
- **Figure 8.** Violin plots (right) illustrating the mean length distribution in (a) the right IFOF and (b) the forceps minor (illustrated in left), across different temporal language lateralisation groups. Group differences are analysed using Tukey post-hoc tests, with adjustments for
- 957 FDR. Significant differences are indicated by p-values less than 0.05 (FDR corrected), while
- non-significant differences are denoted by p-values greater than 0.05 (FDR corrected).
- 959

960 **Table 1**. A list of abbreviations used in this article.

SDF

Spin Distribution Function

Abbreviation Definition AF Arcuate Fasciculus ANOVA Analysis of Variance BLR Bilateral Language Representation BOLD Blood-Oxygen-Level-Dependent Signal CIFTI Connectivity Informatics Technology Initiative Grayordinate Space COPE Contrast Of Parameter Estimates (COPE) Map FDR False Discovery Rate FAT Frontal Aslant Tract a "medium resolution" non-linear registration method FNIRT FSL The FMRIB Software Library FWE Family-Wise Error fMRI Functional Magnetic Resonance Imaging HCP Human Connectome Project HLD Hemispheric Language Dominance Inferior Fronto-Occipital Fasciculus IFOF ILF Inferior Longitudinal Fasciculus Left Language Dominance LLD LQ Laterality Quotient MNI Montreal Neurological Institute Space MSM-ALL Multimodal Surface Matching Registration MSM-SULC **Cortical Folding-Based Registration** QA Quantitative Anisotropy ROI Region Of Interest RLD **Right Language Dominance**

SLF Superior Longitudinal Fasciculus

STGa Superior Temporal Gyrus anterior

STSda Superior Temporal Sulcus dorsal anterior

STSva Superior Temporal Sulcus ventral anterior

Methosciaccepted Manuscrip TE1a Temporal Area 1 anterior















