

Asymmetry of large-scale functional organization: heritable and phylogenetic profiles

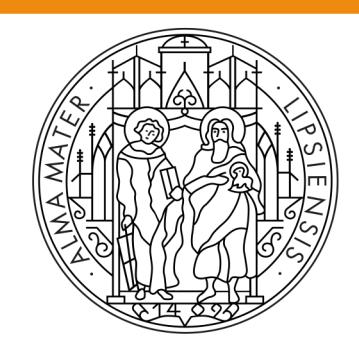
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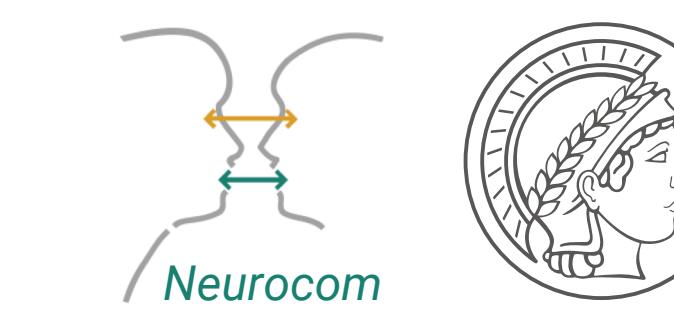
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Introduction

- Asymmetry is a key organisational feature of the human brain and related to the hemispheric preference of functional processes, for example language. However, to what extent the functional organisation in the cerebral cortex is asymmetric, and whether this asymmetry is heritable is not known to date.
- The present study aims to investigate asymmetry of functional organization in healthy humans, study its heritability using a twin-sample, and analyse the asymmetry in non-human primates (i.e., macaque monkeys) to map phylogenetic conservation.

Methods

- Human data: HCP S1200 (n=1014) (www.humanconnectome.org/)
- Macaque data: PRIMATE-DE (n=19). (Milham et al., 2018)
- Functional connectivity (FC) included intra-hemispheric (left-left and right-right) and inter-hemispheric (left-right and right-left) patterns.
- Gradients of each FC pattern were computed using BrainSpace (Vos de Wael et al., 2020) and aligned to the template gradients (gradients of group mean left-left FC).
- Asymmetry index: left minus right hemisphere in the homologous regions.
- We computed heritability by means of family structure of HCP using SOLAR eclipse 8.4.1b (<http://solar-eclipse-genetics.org/>). Heritability $h^2 = \sigma_g^2$ (total additive genetic variance) / σ_p^2 (phenotypic variance).
- Functions along the asymmetry axis was computed using projection of meta-analytical task-based activity generated by NeuroSynth (Chang et al., 2013) (<https://neurosynth.org/>).

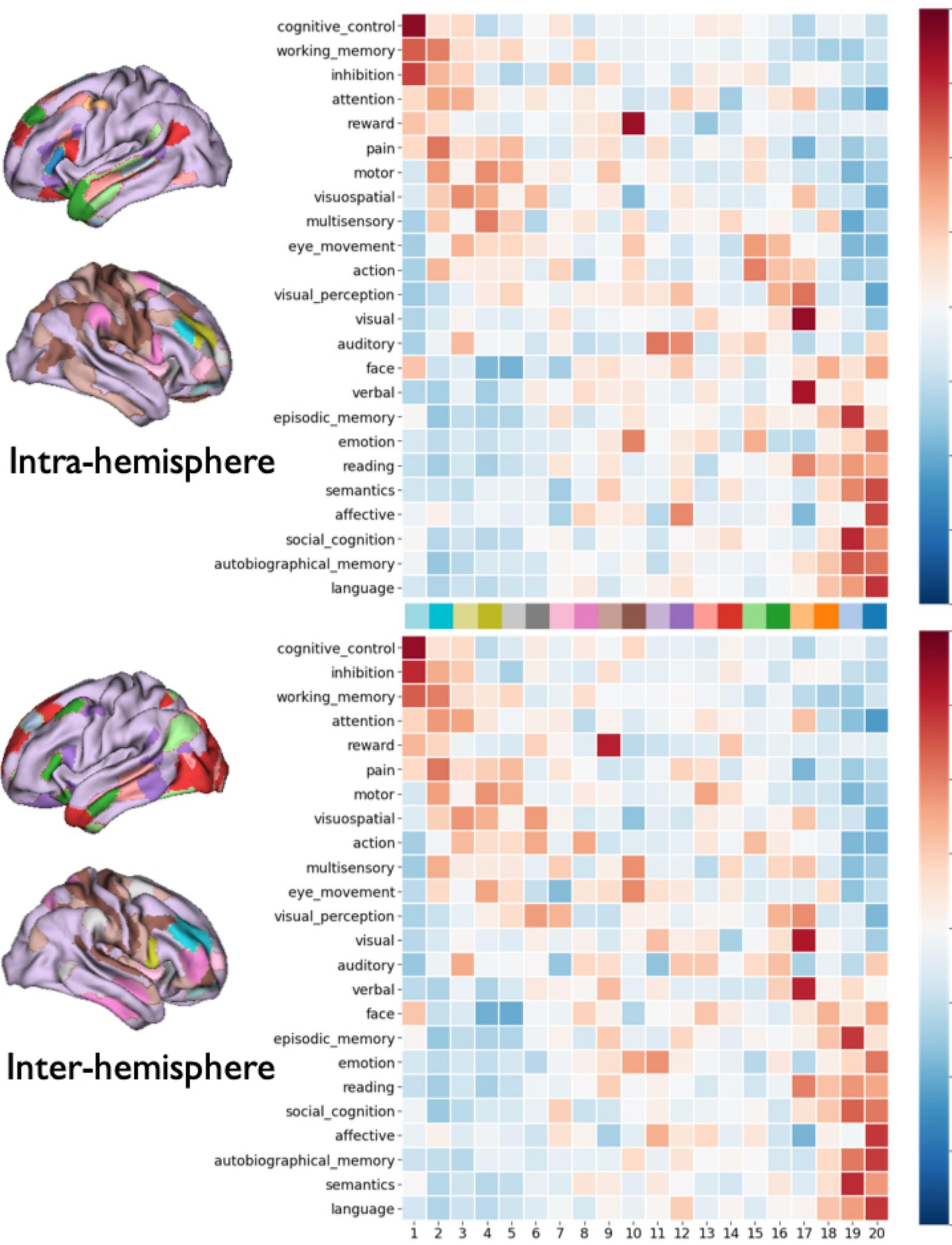


Fig. 4. Projection of meta-analytical task-based function along asymmetric effects of G1. Right panel is the functional decoding of the intra-hemispheric pattern and the left panel is the inter-hemispheric pattern.

Results

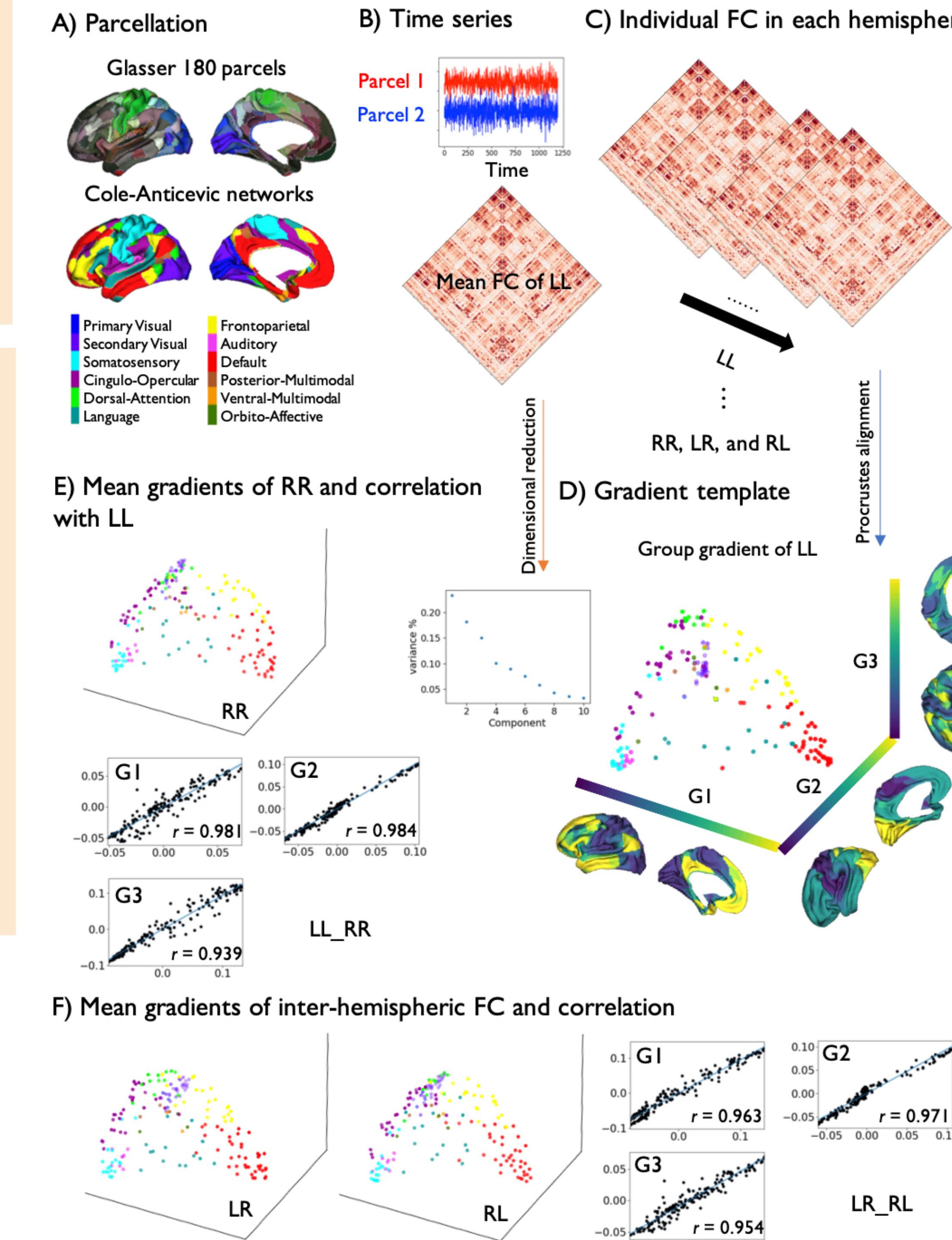


Fig 1. Processing of functional gradients. A) Parcellation using Glasser 180 atlas in each hemisphere and Cole-Anticevic networks. B) Time series of two parcels and mean functional connectivity (FC) matrix between left and left hemisphere (LL). C) Individual FC in each hemispheric pattern, i.e., (LL), right-right (RR), left-right (LR), and right-left (RL). D) Gradient template using group level gradient LL. E) Mean gradients of RR and correlation with LL F) Mean gradients of inter-hemispheric FC and correlation. Color of dots in 3D plotting means Cole-Anticevic networks.

Discussion

- For the intra-hemispheric pattern, transmodal systems showed great asymmetry (e.g., language, DMN, FPN), but unimodal systems showed small or no asymmetry (e.g., visual). For the inter-hemispheric pattern, we did not observe strong patterns of asymmetry (particularly in primary cortex). It indicates that neurons involved in higher order cognition are asymmetrically organized within hemisphere, but the asymmetric organization decreases when it needs corpus callosum to be connected.
- Excluding visual cortex, asymmetry of both intra and inter-hemispheric asymmetry was significantly heritable, indicating that asymmetry of functional gradients may be under genetic control. As noted, heritability of asymmetry of Wernicke's area peaked among the cortex, indicating

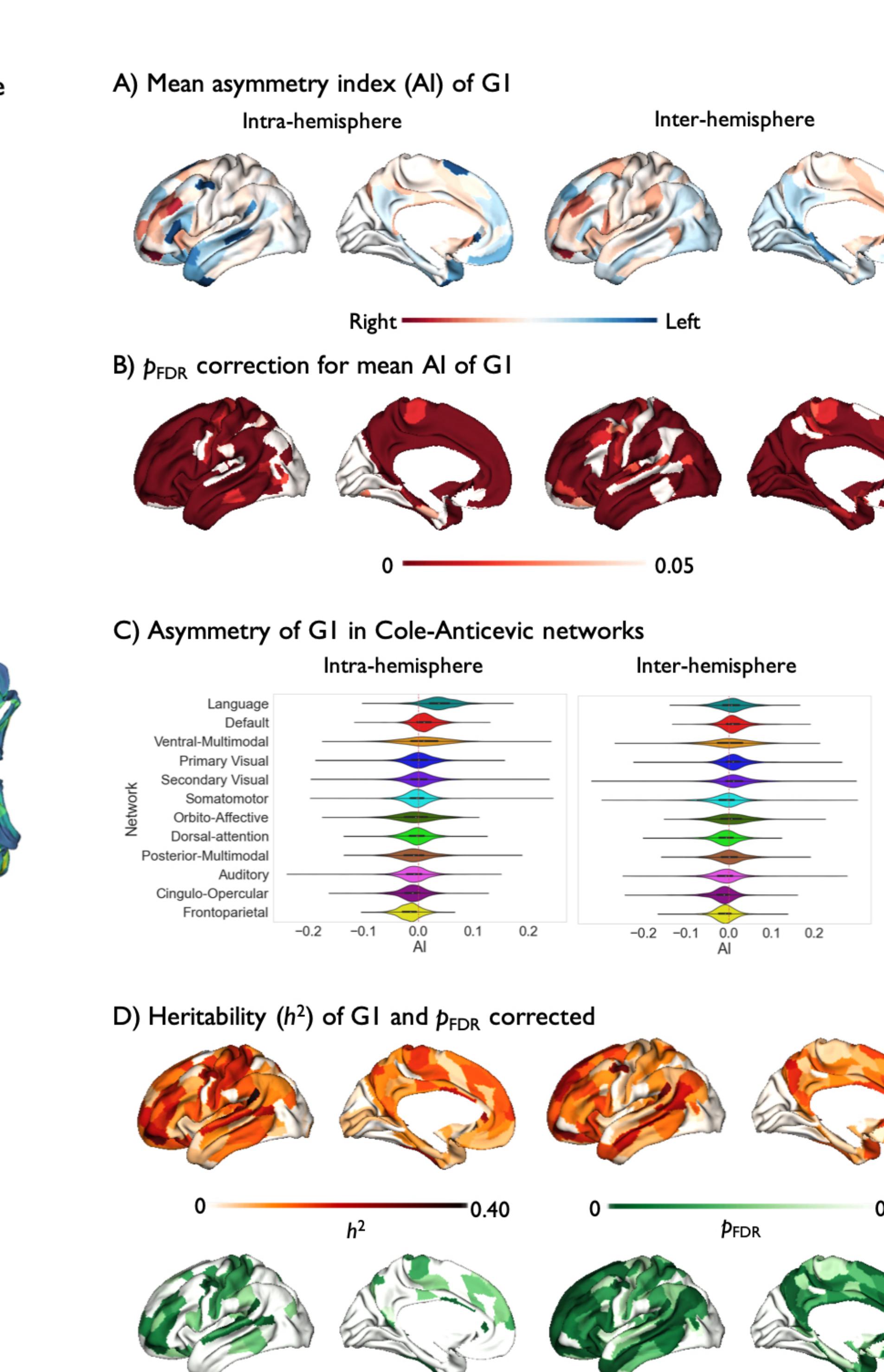


Fig 2. Asymmetry of functional gradients in humans. A) Mean asymmetric index (AI) of intra-hemispheric pattern and inter-hemispheric pattern in humans. B) FDR correction for the p value. C) Asymmetries of the first gradient in Cole-Anticevic networks. D) Heritability (h^2) of G1 and FDR correction for p value.

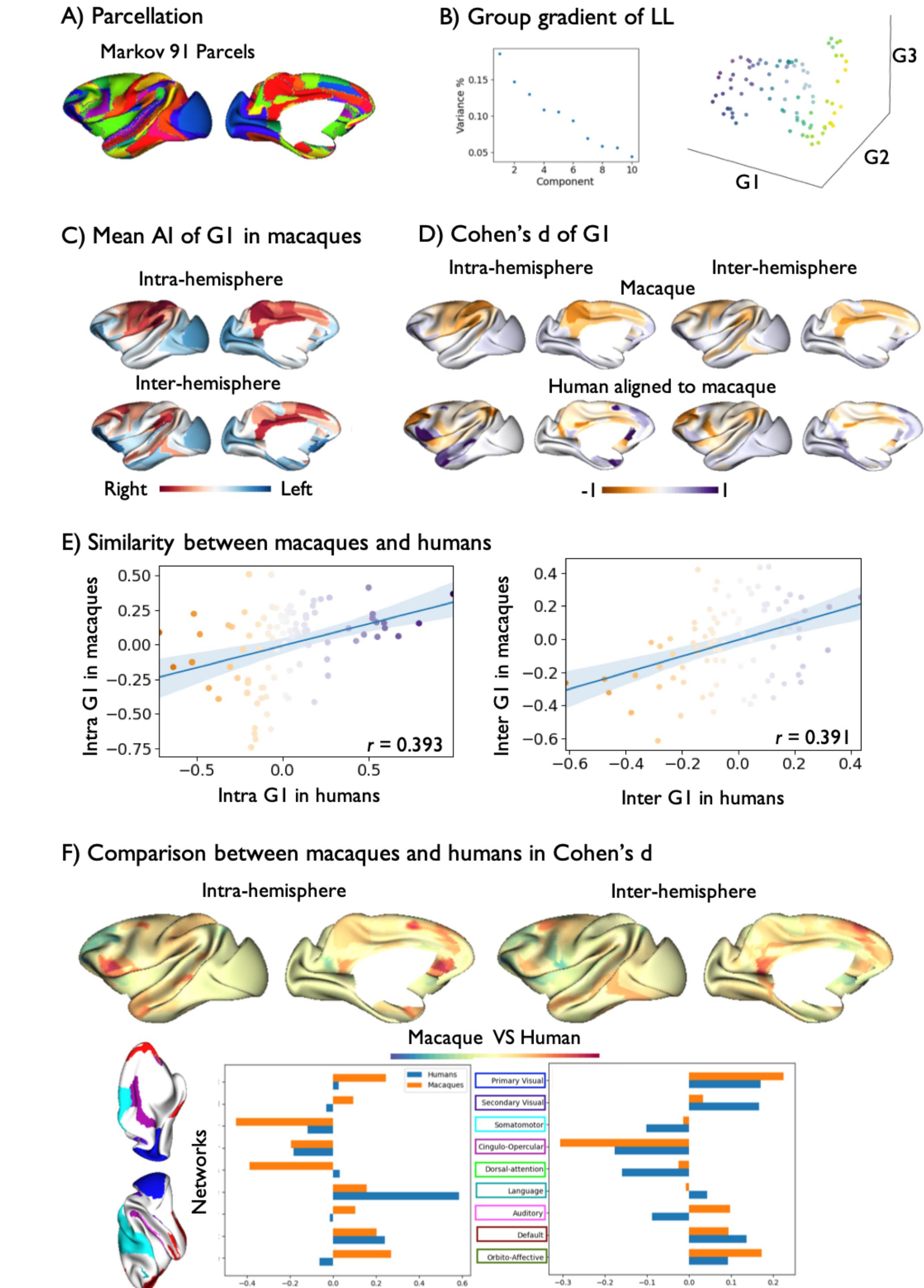


Fig. 3. Asymmetry of functional gradients in macaques. A) Parcellation used Markov 91 atlas in macaques. B) Template gradients. C) Mean asymmetry index of G1 in macaques. D) Asymmetric effects (Cohen's d) of the G1 in macaques and humans aligned to macaques. E) Similarity of G1 in humans with G1 in macaques. F) Comparison between macaques and humans in Cohen's d of G1.

References

- Buckner, R. L., & Krienen, F. M. (2013). The evolution of distributed association networks in the human brain. *Trends in Cognitive Sciences*, 17(12), 648–665.
 Chang, L. J., Yarkoni, T., Khaw, M. W., & Sanfey, A. G. (2013). Decoding the role of the insula in human cognition: functional parcellation and large-scale reverse inference. *Cerebral Cortex*, 23(3), 739–749.
 Ge, T., Holmes, A. J., Buckner, R. L., Smoller, J. W., & Sabuncu, M. R. (2017). Heritability analysis with repeat measurements and its application to resting-state functional connectivity. *Proceedings of the National Academy of Sciences of the United States of America*, 114(11), 5521–5526.
 Glasser, M. F., Coalson, T. S., Robinson, E. C., Hacker, C. D., Harwell, J., Yacoub, E., Ugurbil, K., Andersson, J., Beckmann, C. F., Jenkinson, M., Smith, S. M., & Van Essen, D. C. (2016). A multimodal parcellation of human cerebral cortex. *Nature*, 536(7615), 171–178.
 Karolis, V. R., Corbetta, M., & Thiebaut de Schotten, M. (2019). The architecture of functional lateralisation and its relationship to callosal connectivity in the human brain. *Nature Communications*, 10(1).
 Margulies, D. S., Ghosh, S. S., Goulas, A., Falkiewicz, M., Huntenburg, J. M., Langs, G., Bezgin, G., Eickhoff, S. B., Castellanos, F. X., Petrides, M., Jefferies, E., & Smallwood, J. (2016). Situating the default-mode network along a principal gradient of macroscale cortical organization. *Proceedings of the National Academy of Sciences of the United States of America*, 113(44), 12574–12579.
 Markov, N. T., Ercsey-Ravasz, M. M., Ribeiro Gomes, A. R., Lamy, C., Magrou, L., Vezoli, J., Misery, P., Faucher, A., Quillardet, R., Gariel, M. A., Sallet, J., Garamut, R., Huissoud, C., Clavagnier, S., Giroud, P., Sappéy-Marinier, D., Barone, P., Dehay, C., Toroczkai, Z., ... Kennedy, H. (2014). A Weighted and Directed Interareal Connectivity Matrix for Macaque Cerebral Cortex. *Cerebral Cortex*, 24(1), 17–36.
 Milham, M. P., Al, L., Koo, B., Xu, T., Amiez, C., Balezeau, F., Baxter, M. G., Bleizer, E. L., Brochier, T., Chen, A., Croxson, P. L., Damatac, C. G., Dehaene, S., Everling, S., Fair, D. A., Fleysher, L., Freiwald, W., Froudrist-Walsh, S., Griffiths, T. D., ... Schroeder, C. E. (2018). An Open Resource for Non-human Primate Imaging. *Neuron*, 100(1), 61–74.e2.
 Vos de Wael, R., Benkarim, O., Paquola, C., Larivière, S., Royer, J., Tavakol, S., Xu, T., Hong, S.-J., Langs, G., Valk, S., Misic, B., Milham, M., Margulies, D., Smallwood, J., & Bernhardt, B. C. (2020). BrainSpace: a toolbox for the analysis of macroscale gradients in neuroimaging and connectomics datasets. *Communications Biology*, 3(1), 103.
 Xu, T., Nenning, K. H., Schwartz, E., Hong, S. J., Vogelstein, J. T., Goula, A., Fair, D. A., Schroeder, C. E., Margulies, D. S., Smallwood, J., Milham, M. P., & Langs, G. (2020). Cross-species functional alignment reveals evolutionary hierarchy within the connectome. *NeuroImage*, 223, 117346.