1 **TITLE:** Precision fMRI reveals densely interdigitated network patches with conserved motifs in
2 the lateral prefrontal cortex
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ABSTRACT:

Dominant models of human lateral prefrontal cortex (LPFC) organization emphasize broad domain-general zones and smooth functional gradients. However, these models rely heavily on group-averaged neuroimaging, which can obscure fine-scale cortical features - especially in highly inter-individually variable regions like the LPFC. To address this limitation, we collected a new precision fMRI dataset from 10 individuals, each with approximately 2 hours of resting-state and 6 hours of task data. We mapped individual-specific LPFC networks using resting-state fMRI and tested network-level functional preferences using task fMRI. We found that individual LPFC organization differed markedly from group-average estimates. Individual maps showed more fragmented and interdigitated networks - especially in anterior LPFC - including novel conserved motifs present across individuals. Task fMRI revealed that distinct but adjacent networks support domain-specific processes (i.e., language, social cognition, episodic projection) versus domain-general control. Sharp functional boundaries were visible at the individual level that could not be observed in group data. These findings uncover previously hidden organizational principles in the LPFC and offer a framework for understanding how the LPFC supports flexible, complex cognition through a finely organized architecture.

INTRODUCTION:

The lateral prefrontal cortex (LPFC) plays a central role in high-level cognitive processes like planning, reasoning, and problem-solving (Fuster, 1989; Stuss & Benson, 1986). Damage to the LPFC disrupts many aspects of goal-directed behavior (Lhermitte et al., 1986; Milner, 1963; Shallice & Burgess, 1991), and abnormalities in LPFC circuitry have been linked to a wide range of neurological and psychiatric disorders (Barbalat et al., 2009; McTeague et al., 2017).

Despite its central role in cognition, there is little consensus on how the LPFC is functionally organized. A major point of debate concerns the degree and topography of functional specialization within the LPFC. Several dominant models portray the LPFC as a broad, multifunctional region with gradual gradients of specialization. For instance, large-scale neuroimaging studies have reported overlapping LPFC activations across diverse tasks (Cabeza & Nyberg, 2000; Duncan & Owen, 2000; Niendam et al., 2012), supporting unitary models of flexible cognitive control (Duncan, 2010; Miller & Cohen, 2001). Others have described smooth topographic gradients, such as a rostral–caudal axis of control-related abstraction (Badre, 2008; Koechlin et al., 2003; Nee & D'Esposito, 2016) and a dorsal–ventral domain shift from spatial to verbal content (Abdallah et al., 2022; Blumenfeld et al., 2013; Rahm et al., 2013). Functional network-based frameworks have identified broad network territories spanning the LPFC (Power et al., 2011; Yeo et al., 2011), dominated by a large frontoparietal network region that is ascribed much of the high-level planning and control functions for which the LPFC is known (Vincent et al., 2008). These networks too have been interpreted along a slowly moving cortex-wide gradient from sensorimotor to default mode regions (Margulies et al., 2016). However, these smooth patterns stand in contrast with some studies based on neuronal recordings or tract tracing of neuroanatomical circuits, which have identified adjacent regions with different cytoarchitectonics and neuronal preferences (Barbas & Pandya, 1989; Funahashi et al., 1993; Petrides, 2005).

We contend that this discrepancy may arise because most neuroimaging studies of human LPFC organization have relied on group-averaged data, aligning individual brains to a common anatomical template and assuming functional regions occur in similar locations across individuals. Recent work has shown that group averaging can obscure fine-scale details and overestimate overlap between distinct functional regions, particularly in areas with high inter-individual variability (Braga & Buckner, 2017; Du et al., 2024; Fedorenko, 2021; Gordon et al., 2017; Petersen et al., 2024). The LPFC has emerged as one of the most inter-individually variable brain regions (Dworetsky et al., 2024; Mueller et al., 2013; Seitzman et al., 2019), raising the possibility that the LPFC may contain fine-scale organization not reflected in current models.

To address the limitations of group-averaged methods, several studies have adopted a "precision neuroimaging" approach—collecting large amounts of data from individual participants to achieve high signal-to-noise at the individual level and analyzing functional organization on a subject-by-subject basis (Fedorenko, 2021; Gordon et al., 2017; Laumann et al., 2015; Michon et al., 2022). In the LPFC, several such studies have identified examples of fine-scale functional specialization not visible in group average models (DiNicola et al., 2020; Fedorenko et al., 2012; Fedorenko & Thompson-Schill, 2014; Michalka et al., 2015; Noyce et al., 2017) and found that functional differentiation often corresponds with distinct functional networks (Blank et al., 2014; Braga et al., 2020; DiNicola et al., 2023; Du et al., 2024; Salvo et al., 2024; Tobyne et al., 2017).

Here, we sought to extend these observations and map the fine-scale organization of the
individual LPFC. To do this, we collected a new high-resolution precision fMRI dataset (PAN:
Precision targeting of Association Networks) from ten individuals (18-33 years; 5F), each with
two hours of resting-state fMRI and 6 hours of task fMRI. The task battery spanned 11 different
cognitive demands known to engage the LPFC including theory of mind, episodic projection,
language processing and eight distinct cognitive control tasks. Using resting-state fMRI, we
derived high-resolution, individual-specific networks and characterized fine-scale features
present in the individual-specific compared to group-average LPFC. Using task fMRI, we
evaluated how individual-specific LPFC networks supported diverse cognitive demands.

We found that individual-specific LPFC network organization systematically differed from group-
average models. While group-average data consisted of large contiguous network regions,
individuals exhibited fragmented and interdigitated networks. In individual-specific data, the
frontoparietal network was systematically smaller and borders between association networks
were denser – especially in anterior LPFC. Despite variability in the exact positioning of
networks, we observed conserved spatial motifs across individuals that could not be seen
viewing group data alone, as well as clearly idiosyncratic features. Using task fMRI, we found
that some LPFC networks supported different domain-specific processes, while others
supported domain-general control. Individual-specific task activations corresponded closely with
individual-specific networks, and revealed sharp functional boundaries that would not have been
visible in group-level data. Together, these findings refine coarse and gradient models of the
LPFC, emphasizing fine-scale architecture and underscoring the importance of individual-level
analysis in the LPFC for research and clinical applications.

RESULTS:

Individual-specific LPFC network and task maps are reliable and unique

Using at least 100 minutes of motion-censored resting state fMRI data per individual, we
generated individual-specific network parcellations for 10 individuals (Figure 1A, whole-cortex in
Supplemental Figure S1, S2) using a state-of-the-art network identification protocol (Lynch et
al., 2024). A group-average estimate was also derived for comparison from a reference set of 37
individuals (Lynch et al., 2024, see Supplemental Figure S3 for comparison with a group
average derived from the 10 individuals in this study). For most analyses reported in this
manuscript, network maps were masked to include only the lateral prefrontal cortex (LPFC). To
establish the reliability and individual specificity of individual-level LPFC network parcellations,
we calculated split-half reliability for two individuals (PAN01 and PAN02) who completed
additional resting-state runs to facilitate reliability analyses (totaling 147 and 159 motion-
censored minutes respectively). As shown in Supplemental Figure S3, their individual-specific
LPFC parcellations were reliable across split halves derived from independent scanning
sessions (mean network Dice overlap = 0.79 ± 0.05 between split halves of the same individual)
and unique (mean network Dice overlap = 0.34 ± 0.13 between split halves of different
individuals). A paired-samples t-test revealed that within-individual similarity was significantly
higher than between-individual similarity (t(6) = 9.8, p = 0.0001, Cohen's d = 3.7).

In addition, we collected ~6 hours of task fMRI data per individual (30-60 minutes per task) and
generated individual-specific z-statistic maps for three cognitive domains of interest (theory of
mind, language processing, and episodic projection) as well as for eight distinct cognitive control
tasks (visual n-back, auditory n-back, spatial memory span, verbal memory span, numeric multi-
source interference, verbal multi-source interference, visual attention, auditory attention). To
validate the reliability of individual-level cognitive control maps, we examined two individuals
(PAN01 and PAN02) who completed approximately twice as many runs per task (50-100
minutes per contrast) as the eight other individuals. Shown in Supplemental Figure S4, their z-

statistic maps were reliable across split halves (r = 0.85 ± 0.06 within individuals) and unique (r
= 0.39 ± 0.09 between individuals). A paired-samples t-test revealed that within-individual
similarity was significantly higher than between-individual similarity across tasks (t(7) = 21.6, p =
0.000001, Cohen's d = 7.6), suggesting that LPFC task responses were reliable and individual
specific.

In the next sections, we characterized the topographical features of LPFC organization visible at
the individual level relative to the group average. Specifically, we: (1) characterized the areal
composition and spatial density of distinct networks in the individual LPFC (Figures 1,2), (2)
described the architecture of conserved network motifs not visible in group-level data (Figures
3,4), and (3) described how this network architecture supports a variety of high-level cognitive
demands (Figures 5-8).

LPFC network organization is systematically different in individuals, especially the
frontoparietal network which is overestimated in group averaged data

Shown in Figure 1A, we observed systematic differences between individual LPFC networks
and the group average. While a similar set of association networks were present in individuals
and the group average (frontoparietal, default A, default B, cingulo-opercular, dorsal attention,
salience/parietal memory, language - see Supplemental Figures S1 and S2 for whole-cortex
maps), both the relative size and spatial patterning of LPFC network regions were different. In
the group average, network regions tended to be large and contiguous, while in individuals,
network regions were often smaller and interdigitated.

This difference was especially stark in the frontoparietal network. In the group averaged data,
the frontoparietal network made up 36% of LPFC surface area and contained a large contiguous
network region (Fig 1A, group average). In individuals, the frontoparietal network was
significantly smaller (25% ± 5.4% of LPFC surface area; one-sample t-test: t(9) = 6.7, p =
0.0001, Cohen's d = 2.1, see Fig 1B,C) and was frequently fragmented into multiple distinct
regions. Only 1/10 individuals had a large and contiguous frontoparietal network as was seen in
the group averaged data (PAN10, see blue arrow in Figure 1A). This result was replicated using
an alternative group atlas which included two frontoparietal networks (Yeo 17 network
parcellation, Supplemental Figure S6). By contrast, default B (t(9) = 3.1, p = 0.013), default A
(t(9) = 4.1, p = 0.003), and visual-stream (t(9) = 4.3, p = 0.0001) networks were larger in
individuals than in the group averaged data (Figure 1B). These findings suggest the LPFC
exhibits a more distributed and fragmented network topography than was previously assumed,
with more cortical surface area dedicated to networks (default B, language) not typically
associated with cognitive control – a topic we will return to later. We further validated network
assignments using seed connectivity for PAN01 and PAN02 to ensure the observed
organization was not an artifact of thresholding or misassignment in the network parcellation
(Supplemental Figure S7).

Figure 1. LPFC network organization differs systematically between individuals and the group average, especially in the frontoparietal network. (A) Individual-specific network parcellations are shown, alongside the group average. While the group average consisted of broad contiguous regions, individual maps were fragmented and interdigitated (see Supplemental Figure S6 for replication with an alternative group average parcellation). This difference was especially pronounced in the frontoparietal network, which was large and contiguous in the group average (blue arrow) but was smaller and fragmented in individuals (except for PAN10, blue arrow). (B,C) On average, frontoparietal network size in the LPFC was overestimated in the group average (t(9) = 6.7, p = 0.00009) while default network B (t(9) = -3.1, p = 0.02), default network A (t(9) = -4.1, p = 0.003), and visual-stream (t(9) = -4.3, p = 0.0008) size were underestimated.

The individual LPFC is densely packed with association network borders, especially in the anterior LPFC

As described above, LPFC organization in individuals exhibited fragmented and interdigitated networks compared with the relatively contiguous networks in the group averaged data. Prior work has speculated that interdigitation of distinct networks could have functional consequences by increasing opportunities for communication at network borders (Assem et al., 2024; Power et al., 2013). To compare interdigitation in the LPFC of individuals relative to the group average, we defined "association network density" as the average number of distinct association

networks within 6-14 mm of each vertex (Figure 2A,B based on "community density" from Power
et al., 2013).

As shown in Figure 2C, association network density was consistently higher in individual-
specific LPFC network maps (mean density = 3.36 +/- 0.22) than in the group average (mean
density = 2.53) (one-sample t-test: t(9) = 11.5, p = 0.000001, Cohen's d = 3.64). In addition to an
overall increase, individuals exhibited a distinct spatial pattern: all individuals showed a hotspot
of high density in anterior LPFC that was not observed in the group average (Figure 2B,D,E).
This was particularly apparent in the difference map between the individual and group-average
density maps (Figure 2E) and in the whole-brain view (Supplemental Figure S8). To ensure that
this difference was not driven by signal-to-noise ratio (SNR) differences within the LPFC, we
examined the spatial SNR pattern across individuals in the LPFC and found that the correlation
between SNR and association network density was near zero (r = 0.02, 95% CI: [–0.002,
0.045]), suggesting no meaningful relationship (Supplemental Figure S9). As with network
composition, these results were replicated using an alternative group network prior (Yeo 17,
Supplemental Figures S10). These findings indicate that interdigitation of networks is
systematically underestimated in LPFC group-average maps and suggest that anterior LPFC
may be a zone of high integrative potential across association networks

225
Figure 2. Association network density is underestimated in LPFC by group averaging,
especially in anterior LPFC. (A) Association network density was computed for each cortical
vertex as the number of unique association networks within a 6–14 mm radius (averaged across
distances of 6, 8, 10, 12, and 14 mm). (B) Individual maps of association network density are
shown with an arrow highlighting the anterior high density region. (C) Mean LPFC association
network density was significantly higher in individuals compared to the group average (One-
sample t-test: t(9) = 11.5, p = 0.000001, Cohen's d = 3.64). (D) On average, individuals
exhibited a distinct spatial pattern, including a region of high density in anterior LPFC. (E) This
anterior pattern was absent in the group average map, suggesting that association network
density is particularly underestimated in anterior LPFC. (See also whole cortex estimates in
Supplemental Figure S7.) These results were replicated using a different group average prior,
suggesting they are not specific to this parcellation (Yeo 17 networks, Supplemental Figure S9).

Individual-specific maps reveal conserved network motifs not present in group average
maps

To better understand the spatial patterns underlying the increased association network density
in the LPFC, we manually examined network organization in each individual. We asked whether
the patterns were completely idiosyncratic or if there were consistent features—conserved
network motifs—that appeared across people. Prior work using precision fMRI has shown that
such motifs can be found across individuals but are too fine-grained or variable in location to
show up in group-average data (Braga & Buckner, 2017; Gordon et al., 2023; Noyce et al.,
2017). In line with this, we identified and validated two spatial motifs in the LPFC that were
shared across most individuals but not visible in the group average. We also identified some
individual-specific spatial features, which we describe later.
249
First, we identified a novel conserved network motif surrounding the rostral cingulo-opercular
network region. In the group averaged data, the rostral cingulo-opercular network region was
bordered by the frontoparietal network posteriorly and by the salience/parietal memory and
default B networks anteriorly. However, in individuals, additional small regions of the language
and dorsal attention networks were present (Figure 3A), forming a three-network motif. Although
the precise shape and extent of these regions varied considerably across individuals, they were
visible in 9/10 and 10/10 individuals, respectively (with PAN01 lacking the language region,
though see evidence of connectivity to the expected location in Figure 3B). These findings were
validated with seed-based functional connectivity analyses (Figure 3B), suggesting they were
real features of functional organization rather than artifacts of our parcellation method. We also
found additional instances of this three-network motif in the caudal strip of the LPFC (Figure
1A), the temporoparietal junction (TPJ; Supplemental Figure S11) and pre-supplementary motor
area (pre-SMA; Supplemental Figure S11), aligning with previous observations that functional
networks often border the same neighbors in multiple cortical regions (Braga & Buckner, 2017).

264
Figure 3. A consistent anterior LPFC motif involving language, cingulo-opercular, and
dorsal attention networks is present across individuals but absent in the group average.
(A) Individual-specific parcellations revealed a consistent spatial motif in 9 out of 10 individuals:
a language network region anterior to rostral cingulo-opercular network and dorsal attention
network region posterior to it. This motif was not observed in the group-average parcellation. (B)
Seed-based connectivity from other frontal regions of language and dorsal attention showed

connectivity with these rostral regions in individuals, supporting their network identity. Weak but
present connectivity was also observed in the group average (blue arrows). The weakness may
be explained by the variability in location/orientation of strong connectivity across people, hence
why these features were not identified in the group-average parcellation.
275
We also note two exceptions to this three-network-motif among the ten individuals. PAN01
exhibited the anterior dorsal attention region but not the anterior language region. However,
their seed connectivity suggested the presence of a putative anterior language region directly
adjacent to CO not captured by the parcellation (Figure 3B, blue arrow). PAN04 displayed two
anterior CO regions. The more posterior CO region followed the motif—language anterior,
dorsal attention posterior—whereas the more anterior region showed the reverse (Figure 3A).
282
Second, we observed that this dorsal attention – cingulo-opercular – language motif was part of
a larger organizational feature in the same location. In 8 out of 10 individuals, the anterior PFC
contained a high-density zone where regions of all seven association networks (dorsal attention,
cingulo-opercular, frontoparietal, salience/parietal memory, default A, default B, language)
appeared in close proximity, with the cingulo-opercular region near its center (Figure 4A). Two
individuals (PAN01, PAN07) did not exhibit a rostral DN-A region, but did show seed-based
connectivity to this location from other DN-A regions (Supplemental Figure S14).
290
This location aligned with the hotspot of association network density in Figure 3C and was the
only region in the LPFC where the salience/parietal memory network consistently appeared
across individuals (Figure 1A). To assess the unique properties of the rostral CO location, we
compared the association network density of the rostral CO region to a cortex-wide rotational
null model. On average across individuals, this region fell in the 99.7th percentile of association
network density relative to a matched randomly rotated null (Figure 4B), suggesting it may have
unique integrative potential. A whole-brain map of association network density further highlights
the prominence of this location, which stands out relative to most other brain regions
(Supplemental Figure S8; note that another high-density region was also observed in the
temporoparietal junction). The gathering of many association networks at specific sites may
have important functional implications, as we later discuss.

Figure 4. An anterior PFC high-density zone including all 7 association networks is
consistently observed across individuals. (A) In the group average, the anterior LPFC
contains a CO network region bordered by FP, DN-B, and SAL/PMN. In individuals, we often
observed regions of all 7 association networks here, underlying the high-density zone observed
in Figure 2. (B) This pattern was observed in 8/10 subjects. PAN01 and PAN07 did not exhibit
rostral DN-A regions but showed seed-based DN-A connectivity to this location (Supplemental
Figure S11). All individuals are masked to show networks within 15mm from the rostral CO
region. (C) The rostral CO network region exhibited significantly higher association network
density versus a matched randomly rotated null distribution (p = 0.003, exceeding 99.7% of
comparisons), suggesting it may have unique integrative qualities.
313
Language, theory of mind, and episodic projection tasks preferentially activate distinct
LPFC networks adjacent to -- but separate from -- control networks.

Up to this point, we have characterized the individual LPFC based on the spatial topography of
large-scale networks defined by resting state functional connectivity. We now turn to task fMRI
to examine how these networks support distinct high-level cognitive demands.

While the LPFC is classically associated with domain-general cognitive control, recent studies
have shown that domain-specific processes —including theory of mind, episodic projection, and
language processing—preferentially engage networks which include LPFC regions: default B,
default A, and language networks (Braga et al., 2020; DiNicola et al., 2020, 2023; Du et al.,
2024; Salvo et al., 2024). Close examinations of task activations with individuals has often
shown these functionally specialized regions lay adjacent to, but not overlapping with, nearby
control-related regions (DiNicola et al., 2023; Fedorenko et al., 2012). Notably, these sharp
anatomical distinctions are obscured in group-averaged data.

Here, we examined individual-specific LPFC responses to theory of mind, episodic projection,
and language processing using well-established localizer tasks (see Methods for task details).

We first assessed network-level functional preferences within the LPFC, aiming to replicate prior
findings that theory of mind, episodic projection, and language processing preferentially engage
the default B, default A, and language networks, respectively. Using paired t-tests, we compared
the mean z-values of LPFC regions in each target network to those of all six other association
networks. The results replicated prior network-level dissociations: theory of mind preferentially
activated LPFC default B regions (corrected $p < 0.01$ for all comparisons), episodic projection
preferentially activated LPFC default A regions (corrected $p < 0.00005$), and language
processing preferentially activated LPFC language network regions (corrected $p < 0.001$; Figure
5A). Full statistical details—including corrected and uncorrected p-values, t-statistics, and
Cohen's d—are provided in Supplemental Table S1.

Figure 5. Theory of mind, episodic projection, and language demands preferentially
recruit distinct individual-specific LPFC networks. Average z-statistics were computed for
LPFC regions of each individual-specific network for theory of mind, episodic projection, and
language processing tasks. Paired t-tests comparing the target network to each of the other
LPFC networks revealed that: (A) theory of mind preferentially activated default B (all
comparisons, corrected $p < 0.01$), (B) episodic projection preferentially activated default A (all
comparisons, corrected $p < 0.00005$), and (C) language processing preferentially activated the
language network (all comparisons, corrected $p < 0.001$).

We next assessed how individual-specific LPFC task activations corresponded with individual-
specific versus group average network boundaries. As described earlier, individual-specific
LPFC networks differed systematically from group-average maps—they were more fragmented,
interdigitated, and featured a smaller frontoparietal region. Figure 6A shows three example
individuals, with the top 25% of activated LPFC vertices for each task overlaid on both
individual-specific and group network boundaries (see Supplemental Figure S13 for all
individual-specific maps). We observed that task activations were patchy and idiosyncratic, and
aligned more closely with individual-specific than group-averaged network boundaries.

To quantify this, we calculated the average Dice overlap between thresholded task activations
(top 15%, 20%, 25% of LPFC vertices by z-statistic) and network boundaries for individual-
specific, group-average, and individual-non-specific networks. As shown in Figure 6B, task
activations overlapped more with individual-specific networks than with either group-average or
individual-non-specific networks across all three tasks (corrected $p < 0.05$ for all comparisons;
see Supplemental Table S2 for full statistics). This replicated prior findings that task activations
align with individual-specific networks using other task contrasts (Gordon et al., 2017; Salvo et
al., 2021; Tavor et al., 2016). Notably, although this pattern was evident across cortex, the effect
was significantly stronger in the LPFC than in other regions (Figure 6C), consistent with the idea
that the LPFC exhibits particularly high individual variability and contains fine-scale organization
not captured in group-averaged data.

Figure 6. LPFC non-control task activations are patchy and correspond with individual-specific network boundaries. (A) The top 25% most active LPFC vertices (ranked by z-statistic) are shown for three exemplar individuals for theory of mind, episodic projection and language demands, overlaid with both individual-specific (left column) and group-averaged

(right column) network boundaries. All individuals are shown in Supplemental Figure S13. (B)
Average dice overlap was calculated between thresholded (top 15%, 20%, 25%) LPFC task
activation maps and individual-specific, group-average, and non-specific individual parcellations.
Task activations showed significantly greater overlap with individual-specific LPFC networks
than with either group-average networks or non-specific individual networks (corrected $p < 0.05$
for all comparisons). (C) This effect was larger in the LPFC than in non-LPFC cortical regions (p
< 0.0001).

Finally, we observed that individual-specific activations for these three domains closely matched
individual network boundaries even when these networks appeared in unexpected locations.
For example, a subset of individuals displayed extensive, unexpected segments of the language
and default B networks interdigitated within discrete FP territories in the mid-LPFC (Figure 7A,
Supplemental Figure S14) far from their expected group-average locations.
384
Language network regions were found within the canonical group-average frontoparietal
territory of the LPFC in 4 out of 10 individuals (Figure 7A). No comparable regions could be
observed in the other 6 individuals, suggesting this is not a conserved motif. In all 4 cases, the
variant regions were adjacent to individual-specific FP regions (Figure 7B) and showed strong
seed-based functional connectivity with other LPFC language network territories (Figure 7C),
indicating they were not artifacts of the parcellation. Functionally, these variant language
network regions responded robustly to language processing demands (Figure 7D) and did not
respond to spatial working memory demands. By contrast, nearby individual FP regions showed
the opposite pattern—activation during spatial working memory but not language processing
(Figure 7E). In other individuals, ROIs matched to the variant locations did not show stronger
responses to language than to spatial working memory, further supporting the individual-specific
nature of these regions (Figure 7F). Similar effects were seen for default network B variants in
2/10 individuals (Supplemental Figure S14).

These findings provide functional evidence supporting the highly interdigitated organization of
the LPFC and challenge models suggesting gradual, continuous functional gradients. These
cases also vividly illustrate how averaging across individuals can produce misleading
impressions of blurred boundaries between functionally specialized and domain-general control
regions due to their variable location across individuals.

Figure 7. Language regions are embedded unexpectedly within mid-LPFC canonical frontoparietal territory in a subset of individuals.

(A) Language network parcellations for 10 individuals and the group average are shown,

overlaid with the group-average frontoparietal network. Four individuals (PAN01, PAN02,
PAN06, PAN09) showed large language regions in mid-LPFC, well outside typical language
territory. (B) These variant language regions were interdigitated with fragmented individual-
specific FP regions. (C) Variant regions showed strong seed-based connectivity with other
LPFC language network regions, supporting their network identity. (D) Language processing
task activations showed positive responses in the variant language regions but not in adjacent
FP regions. (E) Spatial working memory activations showed the opposite pattern: activation in
adjacent FP regions but not in the variant Language regions. (F) Language > spatial working
memory z-values are shown for the variant individuals and for the same ROI locations averaged
across the six non-variant individuals. Variant-region ROIs in non-variant individuals did not
show language > spatial working memory activity, validating the individual-specificity of variants.
418
Cognitive control demands recruit the dorsal attention, frontoparietal, and cingulo-
opercular networks with strong activations at their borders

Having observed that some LPFC networks were preferentially activated by domain-specific
non-control processes, we then examined how individual-specific LPFC networks were engaged
by cognitive control demands in a wide variety of contexts. We used eight well-validated tasks
spanning multiple stimulus modalities, domains, and control demands (N-back: Visual and
Auditory, Memory Span: Spatial and Verbal, Multi-Source Interference: Numeric and Verbal,
Attention: Visual and Auditory). Each task map was calculated as the difference between high-
demand and low-demand conditions (N-back: 2-back versus 0-back, Memory Span: 8 items
versus 4 items, Multi-Source Interference: conflict versus no-conflict, Attention: sustained
attention versus sensorimotor control).

As shown in Figure 8A, the eight cognitive control tasks frequently engaged three networks—
the frontoparietal, dorsal attention, and cingulo-opercular networks (corrected p < 0.05 dorsal
attention: 8/8 tasks; frontoparietal: 5/8 tasks; cingulo-opercular: 4/8 tasks). Other networks, such
as salience/parietal memory (2/8 tasks) and language (3/8 tasks), were engaged in specific task
contexts (see Supplemental Table S3 for full statistics). For example, the language network was
activated during verbal working memory, auditory working memory, and auditory attention tasks,
all of which involved additional reading or listening in the high- versus low-demand conditions.

As was observed for domain-specific tasks, cognitive control activation maps were both
individual-specific and patchy. Quantified in Figure 8B and visible for three individuals in Figure
8C, LPFC activation maps were unique across individuals for the same task (cross-individual r =
0.27 ± 0.03) but strikingly consistent within individuals across the 8 different control tasks (r =
0.58 ± 0.03, two-sample t-test contrasting across task vs. across individual similarity: $t(16)$ =
6.94, p = 0.00001, Cohen's d = 1.68). This is visible for three exemplar individuals in Figure 8C,
where a consistent set of distributed regions tended to be active across all 8 control tasks,
despite large task-specific differences in stimuli and control demands (see Supplemental
Figures S15 and S16 for all individual cognitive control activation maps). Unlike LPFC
activations for theory of mind, episodic projection, which tended to fill the borders of a single
network, these activations often appeared to sit at the borders between FP, DAN and CO
(activations highlighted with black outlined circles in Figure 8C). To quantify this tendency, we
measured the geodesic distance between active regions and the nearest other control network
(e.g., if an FP vertex, the distance to the nearest CO or DAN vertex). We found that high-
activation vertices (z>2) were significantly closer to other control network borders than were
lower-activation vertices (z<2). This effect was replicated across multiple z-thresholds
(Supplemental Figure S17). Discussed later, this observation suggests that borders between
networks may be particularly important regions for cognitive control and cross-network

interactions.

456
Figure 8. Cognitive control tasks strongly recruit a set of distributed, individual-specific
regions concentrated along the borders of frontoparietal, cingulo-opercular, and dorsal
attention networks in the LPFC.
(A) Mean network-level LPFC activations (z value) are shown for all eight cognitive control
tasks. Bars are grouped by network and colored accordingly. Statistically significant activations

(corrected p < 0.05) are marked with asterisks. The dorsal attention (DAN), frontoparietal (FP),
and cingulo-opercular (CO) networks were the most consistently engaged across tasks (8/8,
5/8, and 4/8 tasks, respectively). Language and salience/parietal memory networks showed
more selective engagement (3/8 and 2/8 tasks). (B) Vertex-wise activation patterns were
individual-specific (r = 0.27 ± 0.03 between individuals for the same task) and relatively more
consistent within individuals across tasks (r = 0.58 ± 0.03), indicating a stable individual-specific
activation motif across domains. (C) The top 25% most active LPFC vertices are shown for each
task in three exemplar individuals (same as Figure 6, all individuals shown in Supplemental
Figures S15 and S16), overlaid with their FP, DAN, and CO network borders. Task responses
consistently involved a distributed set of regions often located near the borders between the FP,
DAN and CO networks which are highlighted here with black circles. (D) To quantify this border
proximity effect, we measured the geodesic distance from each LPFC vertex in FP, DAN, or CO
to the nearest vertex belonging to a different control network. Highly active vertices (z > 2) were
significantly closer to other control networks than less active vertices (z < 2) in all eight tasks (all
p < 0.001). This was replicated at two other thresholds (Supplemental Figure S17).

477
DISCUSSION:

Neuroimaging studies have frequently reported overlapping activations across large portions of
the LPFC, supporting the idea that it is a multifunctional, flexible region (Cabeza & Nyberg,
2000; Duncan & Owen, 2000; Niendam et al., 2012). However, these reports may have missed
fine-scale organizational structure, as group-averaging methods can blur functionally distinct
regions across individuals—particularly in the highly inter-individually variable LPFC (Dworetsky
et al., 2024; Mueller et al., 2013; Seitzman et al., 2019). In this study, we collected a new
precision fMRI dataset (PAN) with both resting-state and task data to examine the fine-scale
organization of LPFC networks in 10 deeply sampled individuals. We found that LPFC network
architecture is systematically different at the individual level compared to group averages: while
group-level data show large, contiguous swaths of networks, individual data reveal dense
interdigitation of distinct networks. This difference was most striking in anterior LPFC where a
high-density zone of association networks was consistently observed across individuals but was
not present in group-level data. We also validated that distinct LPFC network regions participate
in distinct cognitive processes using task fMRI. This replicated prior findings that some LPFC
networks are functionally specialized for domain-specific processes (e.g., LANG, DN-B, DN-A),
while others support cognitive control across diverse task contexts (e.g., FP, DAN, CO). Below,
we explore the implications of this organizational structure for prior and future work.

Within-individual analysis reveals interdigitated functionally specialized and domain-
general control networks in the LPFC

Our findings build on a growing body of work that has identified functionally specialized regions
in the LPFC using methods that preserve individual-specific organization (Assem et al., 2025;
Braga et al., 2020; DiNicola et al., 2020; Fedorenko et al., 2011, 2012; Michalka et al., 2015). In
several cases, these studies showed—as we do here—that functionally specialized regions are
distinct from, but often adjacent to, domain-general control regions (Assem et al., 2025; DiNicola
et al., 2023; Fedorenko et al., 2012; Noyce et al., 2017). Here, we extend this work with a more
comprehensive description of the patchy and interdigitated nature of LPFC organization,
delineating fine-scale individual networks and dissociating their functional roles across domain-
specific and domain-general control demands. We propose that fine-scale interdigitation may
reflect a general organizational principle of the LPFC and suggest that further high-resolution,

within-individual studies (e.g., Allen et al., 2022; Braga et al., 2020) will uncover additional
examples of functional specialization.

Conceptually, this interdigitated organization contrasts with influential unitary and gradient-
based accounts of LPFC organization. Such accounts emphasize smooth functional transitions
along axes of control-related abstraction (Badre, 2008; Koechlin et al., 2003; Nee & D'Esposito,
2016), informational domain (Abdallah et al., 2022; Blumenfeld et al., 2013; Rahm et al., 2013)
and a principal gradient of representational abstraction (Margulies et al., 2016). In contrast, we
observed sharp boundaries in both task activation and functional connectivity throughout the
individual LPFC, often with multiple noncontiguous patches of the same network appearing in
different LPFC subregions. This was observed most dramatically in a subset of individuals
where language and default B regions were idiosyncratically embedded within the canonical
frontoparietal mid-LPFC region (Figure 7, Supplemental Figure S14). In these cases, functional
preference shifted abruptly and in individual-specific ways. These cases highlight how group
averaging can obscure sharp functional boundaries and give the illusion of broad
multifunctionality, or smooth gradients. Consistent with recent theoretical arguments, (Petersen
et al., 2024), we suggest that while gradients may emerge when data are smoothed—either
analytically or through group averaging—they are substantially reduced in high-resolution,
within-individual analyses. We propose that future work examine how the distinct, interdigitated
networks visible in individuals may nevertheless participate along hierarchical, informational, or
representational axes.

Functional implications of network interdigitation

As discussed above, individual-level analyses revealed a densely interdigitated organization in
the LPFC, with high association network density, especially in the anterior LPFC (Figure 1). This
organization was not visible in group-averaged data and has not been comprehensively
described in canonical network models (e.g., Power et al., 2011; Yeo et al., 2011). Here, we
discuss a potential functional implication of this architecture: interdigitation may increase
opportunities for information integration at the borders between networks.

Several influential theories propose that cognitive control emerges from coordination between
multiple large-scale networks (Assem et al., 2024; Cole et al., 2013; Dosenbach et al., 2008;
Petersen & Sporns, 2015). Consistent with these ideas, we found that LPFC regions belonging
to the dorsal attention, frontoparietal, and cingulo-opercular networks were frequently recruited
across a range of cognitive control tasks (Figure 8A). Further, using within-individual analysis, it
was possible to observe that these networks frequently bordered one another in individual-
specific topographies (Figure 1A) and that cognitive control task activations frequently peaked
near the borders between them (Figure 8D). This result echoes a recent report showing that
cognitive control-related activations tend to be localized near the borders between "core" and
"non-core" multiple demand regions (Assem et al., 2024) with activations shifting depending on
task-specific demands. It also aligns with spatial descriptions of "connector hubs" — regions
exhibiting strong functional connectivity to multiple distinct networks — which are often found
near network borders or "articulation points." (Power et al., 2013, Figure 7; Gratton et al., 2018
Figure 3).
549
Importantly, while prior group-level findings could have been driven by individual variability and
smoothing, the current findings provide corroboration of the importance of border regions within
individuals. Further, we note that anatomical tracing studies in nonhuman primates have shown
strong short-range connectivity across borders between distinct cytoarchitectonic areas in visual

cortex (Kennedy & Bullier, 1985; Rockland & Lund, 1983), and LPFC (Barbas & Pandya, 1989;
Petrides & Pandya, 2002), providing a plausible structural basis for cross-network interactions.
556
557   **Anterior LPFC density and relationship to its special LPFC properties**
558
From an integration perspective, regions where many distinct networks converge may play a
unique role in cognition. In this study, we identified a consistent high-density zone in the anterior
LPFC where many association networks were frequently represented. We speculate that such
regions could have unique functions – perhaps acting as "diverse club" hubs that facilitate
communication across many association networks (Bertolero et al., 2017, 2018) or supporting
conjunctive coding - integrating stimulus, context, and response at the intersection of multiple
networks (Badre, 2025; Fusi et al., 2016; Rigotti et al., 2013). Such ideas are especially
compelling given early studies identifying the anterior LPFC at the apex of a cognitive control
hierarchy, specialized for highly abstract or integrative functions (Badre, 2008; Koechlin et al.,
2003). It is possible that highly abstract or complex cognitive functions may require more cross-
network interactions. In this way, we speculate that studies of network density could reconcile
with gradient frameworks.
571
Further, the anterior LPFC has been identified as the effective site for transcranial magnetic
stimulation (TMS) in major depression (Siddiqi et al., 2020). One possible consequence of
applying stimulation here is the potential to influence multiple networks simultaneously. While
most studies of the LPFC TMS target have focused on its connectivity with the subgenual
cingulate (Fox et al., 2012), we propose that future studies may examine how the uniquely
dense topography may contribute to treatment effects. Dense interdigitation may also help
explain the broad cognitive deficits observed following LPFC lesions, which are often
nonspecific and span multiple domains (Luria, 1966; Roca et al., 2010; Shallice & Burgess,
1991; though see Stuss, 2011). Lesions in this densely packed area may more easily cross
network boundaries, resulting in less selective functional deficits.
582
These findings offer a new framework for considering the LPFC's functional role in comparison
to other brain regions. The LPFC has been historically considered a special area with strong
integrative qualities distinct from posterior brain regions (Miller, 2000). However, later network
analyses revealed that the LPFC contains the same association networks also present in the
parietal, temporal, and cingulate cortices (Power et al., 2011; Yeo et al., 2011), raising the
possibility that unique qualities ascribed to the LPFC may be better attributed to specific highly
integrative networks (Cole & Schneider, 2007). Here, we propose a unifying hypothesis: the
LPFC is not unique in network identity but rather in network density.

Functional implications of distributed patchy networks

Another possibility is that the distributed patchy network structure in LPFC reflects functional
differentiation between different patches of the same network. Over the past decade, imaging
neuroscience has shifted from a localist to a more distributed perspective, where cognitive
processes are thought to emerge from coordinated activity across widely distributed networks
rather than being localized to single regions (Bassett & Sporns, 2017; Petersen & Sporns, 2015;
Sporns, 2013). An open question remains as to what unique roles are played by individual
nodes within these highly connected networks.

Although functional imaging reveals highly coordinated activity across these regions within the
same network, evidence from lesion and neurodegeneration patients suggests differentiation of

distinct network regions. For example, classic models of language processing propose distinct
roles for different subcomponents of the language system—such as comprehension in posterior
STG and production in inferior frontal gyrus (Geschwind, 1970; Hickok & Poeppel, 2007).
Similarly, a seminal account of the spatial attention network identified specialized roles for
posterior parietal cortex (spatial location), frontal eye fields (top-down control of saccades), and
cingulate cortex (motivational relevance) (Mesulam, 1981, 1990). It could likewise be the case
that the multiple patches of a single network in LPFC reflect distinct functional roles. While we
focused here on network-level activity, future work could aim to dissociate the roles of individual
LPFC nodes within the same network.

Precision fMRI reveals common cortical features obscured by group averaging

This study uncovered fine-scale features of LPFC network organization that were consistent
across individuals but not visible in group-averaged data. Notably, we identified a reproducible
DAN–CO–LANG motif in anterior LPFC (Figure 3) which was absent from group-level data. In
contrast, a prominent feature of the group-average map—a continuous frontoparietal region in
dorsolateral PFC (Figure 1)—was present in only 1 of 10 individuals. This highlights a key
limitation of group averaging: it can produce features that do not reflect the organization of any
individual participant.

These results parallel recent precision fMRI studies that have revealed similarly fine-scale
motifs obscured in group data, including parallel default networks (DN-A, DN-B) (Braga &
Buckner, 2017; DiNicola et al., 2020), a novel somato-cognitive-action network (Gordon et al.,
2023), and a unified salience/parietal memory network (Kwon et al., 2025) among others.

Although the three-network motif identified here has not been the central focus of prior work,
converging evidence supports its existence. Assem et al., 2024 reported strong connectivity
between the dorsal attention network and a rostral LPFC region in group-average data and
hypothesized the existence of a dorsal attention region in that location (their Supplemental
Figure 4). Braga et al., 2020 identified an anterior LPFC region assigned to the language
network across individuals, not captured in group-level atlases. Du et al., 2024 observed a
similar motif to the one we observed in multiple individuals (Figures 2, 5, 14–16), though it was
not a focal point of their study and was detected using a different parcellation method (MS-
HBM). We also found this motif repeated in other cortical zones—including pre-SMA, TPJ, and
dorsal and ventral caudal LPFC—consistent with prior work showing that networks often border
the same neighbors in multiple locations (Braga & Buckner, 2017; DiNicola et al., 2020). This
repeating spatial structure aligns with classic proposals of parallel, functionally specialized
processing streams (Goldman-Rakic, 1988).

Identifying common fine-scale features is essential for grounding theoretical models of brain
organization. In the present study, detecting small anterior regions of the dorsal attention and
language networks in individuals was critical for revealing the high association network density
region in the anterior LPFC and for localizing the CO region as its spatial anchor (Figure 4).
Further, while influential models propose a single coarse hierarchical arrangement of LPFC—
from dorsal attention to frontoparietal to cingulo-opercular and salience networks—they do not
account for the presence of multiple, spatially distinct patches of each network. Future models
can incorporate the patchy, interdigitated organization revealed by precision fMRI.

Precision fMRI can identify individual-specific cortical features

In addition to common cortical features, precision fMRI enables the identification of idiosyncratic
network regions that do not conform to a shared cortical template and are present in only a
subset, or even a single individual (Dworetsky et al., 2024; Glasser et al., 2016; Gordon et al.,
2017; Laumann et al., 2015; Seitzman et al., 2019). In this study, we characterized several such
regions of the language and default B networks embedded in the mid-LPFC frontoparietal
region in a subset of individuals. We validated these regions using task activations and
observed they were particularly strong examples of adjacent specialized and domain-general
control activity in the LPFC.

Prior work has differentiated distinct types of idiosyncratic features: "ectopic intrusions" --
islands of altered connectivity and function, and "border shifts"– local expansions or contractions
of commonly shared network regions. It was found that ectopic intrusions are most prevalent in
the LPFC while border shifts are most prevalent in the TPJ (Seitzman et al., 2019, Dworetsky et
al., 2024). In this study, we characterized both ectopic (both DN-B variants, LANG variants
PAN02 and PAN09) and border shifts (LANG variants PAN01 and PAN06). Important questions
remain about why ectopic intrusions are so common in the LPFC and how they develop. Prior
work showed that these regions are less heritable than border shifts and so may relate more to
experience-dependent processes. The LPFC undergoes particularly protracted development
(Gogtay et al., 2004), which may render it especially plastic to experience-dependent processes
(Johnson, 2001; Petanjek et al., 2011). Future work may investigate how ectopic intrusions in
LPFC arise and vary across individuals, potentially offering new insight into how life experience
shapes large-scale cortical organization.

Limitations
666
This study has several methodological limitations, which we have attempted to address where
possible. First, in deriving individual-specific networks, we chose one specific parcellation
strategy (vertex-level Infomap with a template matching manual consensus procedure following
Lynch et al., 2024) based on an 18-network group prior. We selected this prior because it
allowed us to test a priori hypotheses about the functional preferences of specific networks (i.e.,
language, default A/B) and selected vertex-level resolution to identify small network regions.
However, this raises the possibility that our findings may be specific to the parcellation method
or group prior. For example, other parcellations have suggested there are two frontoparietal
networks, though only one is present in our 18-network group prior. To mitigate this risk, we
validated the parcellation using seed maps (Supplemental Figure S7) and replicated network
composition and interdigitation findings using an alternative group prior with two frontoparietal
networks (Yeo 17-network atlas; Supplemental Figures S6, S10).
679
Second, to examine the relationship between individual task activation maps and individual-
specific networks, we required highly reliable individual-level task data. We therefore used block
designs across all tasks, as these have been shown to produce more robust and reliable
individual z-statistic maps than mixed or event-related designs (Mumford et al., 2014; Noble et
al., 2021). However, block designs limit the ability to resolve temporally distinct cognitive
processes within tasks—an important caveat in cognitive control paradigms, where different
networks may operate at different timescales (Dosenbach et al., 2008). Future work may be
able to produce temporally sensitive, individual-specific maps of these control-related dynamics.
688
Third, our analyses of variant LPFC regions (Figure 7, Supplemental Figure S14) were
constrained by our relatively small sample size (N = 10). While this is typical for precision fMRI
studies (e.g., Allen et al., 2022; Braga & Buckner, 2017; DiNicola et al., 2023; Du et al., 2024;
Gordon et al., 2017; Kwon et al., 2025; Salvo et al., 2024), it limits our ability to assess the

prevalence of variants in the broader population or to detect variants not represented in our
sample. We present these findings as illustrative examples of individual variability, with the
assumption that additional variants likely exist. Larger-sample studies (e.g. Seitzman et al.,
2019) have shown that network variants can reflect meaningful subtypes and relate to behavior;
future work could assess how prevalent specific archetypal variants are and how they are
associated with behavior.
699
Finally, all estimates of network topography and task activation are limited by the inherent
smoothness of fMRI data. In particular, the border activations observed between control-related
networks (Figure 8D) may partly reflect signal spillover between adjacent networks. To reduce
this risk, we used small voxel sizes (2.5 mm³) and minimal smoothing (1 mm Gaussian kernel).
This high-resolution approach enabled detection of small, spatially reliable network regions
critical for identifying LPFC motifs which were not visible in group averaged data (Figure 3, see
PAN05 dorsal attention network), while still preserving reliable measures within individuals.
Additionally, we observed that some control network regions showed minimal or no activation
during control tasks, inconsistent with the idea that widespread activation plus smoothing alone
explains the border peaks. Still, replication with higher-resolution methods (e.g., 7T fMRI) will be
important to confirm these findings and rule out residual smoothing artifacts.

Conclusion

In this study, we collected a new precision fMRI dataset (PAN) to map the fine-scale network
organization of the lateral prefrontal cortex (LPFC) in 10 deeply sampled individuals. While the
LPFC is often portrayed as a broad, multifunctional region, our data revealed a patchy,
interdigitated architecture in which distinct networks support domain-general control and
domain-specific processing. We identified both conserved motifs—such as a consistent high-
density zone in anterior LPFC—and individual-specific variants. These features were validated
using both task and resting-state data.

Together, these findings offer a more detailed picture of LPFC organization. They highlight the
value of individual-specific approaches for revealing fine-scale functional architecture and
suggest that spatial patterns of network interdigitation may play a role in how the LPFC supports
flexible and complex cognition.

METHODS:
724
Participants
Ten paid participants (ages 18-33 years, mean age = 26.4, SD = 4.8, 5F) completed one
behavioral training session and 7-10 fMRI sessions. Each fMRI session lasted ~1.5 hours,
comprising 8-11 task runs (5-10 minutes each) and 4-5 resting-state runs (5 minutes each).
Participants were required to have corrected-to-normal vision, be native/fluent English speakers,
be right-handed, and have no history of neurological disorders. All participants provided written
informed consent, and the study was approved by the Northwestern University and Florida State
Institutional Review Boards.
733
MRI Acquisition
Data for one individual (PAN01) was acquired at the Northwestern University Center for
Translational Imaging (CTI) on a 3T Siemens Prisma MRI scanner with a 64-channel head coil
(Siemens Healthcare, Erlangen, Germany). Data for the remaining 9 individuals was acquired at
the Florida State University MRI Facility, also on a 3T Siemens Prisma MRI scanner with a 64-

channel head coil. A multi-echo multiband sequence was used for all functional runs (voxel size:
isotropic 2.4mm, TR = 1.355s, TE = 12.0ms, 32.4ms, 52.0ms, 71.6ms, 91.2ms, flip angle = 59°,
multi-band factor = 6, in-plane acceleration factor = 2, AP phase encoding direction,
FOV=216mm, Slices = 54, see Lynch et al., 2020). Two structural (T1) scans were acquired per
individual on separate days [voxel size: isotropic 1mm, TR = 2.3s, TE = 1.86ms, 3.78ms, TI =
1180ms, flip angle = 7°, FOV = 256mm, Slices = 208]. A gradient-echo fieldmap with the same
geometric parameters as the functional sequence was acquired in each fMRI session. Motion
was monitored in real-time using Framewise Integrated Real-time MRI Monitoring (Dosenbach
et al., 2017) and individual's eyes were video monitored for signs of sleepiness.
748
fMRI Processing
All fMRI data was processed using fMRIprep 23.0.2 – a standardized open-source pipeline for
minimal fMRI preprocessing. This included the creation of a reference T1w, skull-stripping, head
motion correction (rigid body correction using 6 parameters), segmentation of white matter, gray
matter, and CSF, susceptibility distortion correction, optimal combination of echoes, registration
to T1w, spatial normalization to MNI152NLin6Asym 2mm isotropic resolution template space,
and generation of a cortical surface with FreeSurfer (Dale et al., 1999; Esteban et al.,
2019). The resulting surfaces were registered into fs_LR_32k surface space as described in
Glasser et al., 2013. For resting-state fixation runs, after running fMRIprep 23.0.2, data was
further processed as in Power et al., 2014 to reduce the effect of artifacts on functional
connectivity estimation. Data was demeaned and detrended, nuisance signals (motion
parameters, white matter, gray matter, CSF, global signal) were regressed, high motion frames
(fFD > 0.1mm, as in Gratton et al., 2020) were censored and their data interpolated, the residual
data were band-pass filtered (0.08-0.009 Hz). After preprocessing in the volume, cortical
functional data were registered to the fslr32k surface. Cortical surface and volumetric
subcortical and cerebellar data were combined into CIFTI format using Connectome Workbench
(Marcus et al., 2011) and data were (minimally) smoothed (Gaussian kernel, sigma = 1mm)
using 2D geodesic smoothing on the surface and 3-D Euclidean smoothing for subcortical
volumetric data.
768
fMRI Data Quality Control
For all runs, motion-contaminated frames were identified by filtered framewise displacement
(fFD) (Gratton et al., 2020). Frames with fFD > 0.1 were flagged as high motion frames. For task
data, runs were excluded if > 20% of frames were flagged as high motion frames or if task
performance was below chance. In total, 791 task runs were completed in this study, and 11
runs were excluded. PAN02 had 1 Auditory Attention run excluded for task performance, PAN05
had one Auditory Attention run excluded for task performance, PAN06 had one Episodic
Projection run excluded for motion, PAN07 had two runs (Auditory Attention, Theory of Mind -
False Belief) excluded for task performance, and one run (Visual Language) removed for
motion. PAN08 had two runs (Auditory Attention, Theory of Mind) excluded for task
performance, and one run (Episodic Projection) excluded for motion. PAN09 had one run
(Episodic Projection) excluded for motion. PAN10 had one run (Visual Working Memory)
excluded for task performance. For functional connectivity analyses using resting state data, all
runs were included but high motion frames were censored to minimize bias from motion-related
artifacts (Gratton et al., 2020; Power et al., 2014b). On average across individuals, 95.5% +/-
2.1% of frames were retained, leaving a minimum of 101 motion-censored minutes of data per
individual (range = [101 minutes – 159 minutes], average = 118 minutes). Prior work suggests
at least 40 minutes of data are necessary for precision mapping of functional networks, though
possibly less with multi-echo data (Gordon et al., 2017; Laumann et al., 2015; Lynch et al.,
2020).
789

Precision Functional Network Mapping

Individual-specific functional networks were derived using the InfoMap community detection algorithm (Rosvall & Bergstrom, 2008), following the approach described in Lynch et al., 2024. For each participant, we built a functional connectivity matrix by correlating time series between all cortical and subcortical vertices across all resting-state runs and sessions. To reduce the influence of spatially local correlations, connections between nodes ≤ 20 mm apart were set to zero—using geodesic distance for cortical-cortical connections and Euclidean distance for cortical-subcortical ones. These matrices were thresholded at a range of densities (0.01%, 0.02%, 0.05%, 0.1%, 0.2%, 0.5%, 1%, 2%, and 5%) to retain only the strongest edges per vertex. Each thresholded matrix was input to InfoMap to identify community structure. We focused on the 0.1% threshold, as prior work has shown it produces networks with the highest size-weighted homogeneity compared to randomly rotated null models (Gordon et al., 2020; Lynch et al., 2024). At this threshold, individuals showed an average of 62.2 ± 5.0 functional communities.

Network identities were assigned to each community by matching them to a reference set of 18 independently derived functional networks (default B, default A, visual-lateral, visual-stream, visual-V1, visual-V5, frontoparietal, dorsal attention, premotor, language, salience, parietal memory, cingulo-opercular, auditory, somatomotor-hand, somatomotor-face, somatomotor-foot, somato-cognitive-action), based on both functional connectivity and spatial location. These priors were adapted (removing two default-related networks from their prior in order to test specific hypotheses about DN-A and DN-B) from a precision mapping dataset of 37 healthy adults (Lynch et al., 2024). A confidence score was automatically generated for each match, equally weighting functional connectivity and spatial topography. Communities with low confidence (≤ 1) were manually reviewed, and in ambiguous cases, their labels were adjusted by Z.L and verified by K.K. When a community didn't resemble any known network—often in low-signal regions—it was labeled "noise." On average, 3.3% ± 0.7% of cortical vertices were reassigned to a different network after review, and 1.7% ± 1.0% were labeled as noise. Following Gordon et al. (2017), we removed contiguous network patches smaller than 50 mm² and filled them in by dilating neighboring labels. Although networks were estimated for subcortical structures, only cortical assignments were used in subsequent analyses. Finally, based on recent work suggesting that the salience and parietal memory networks are part of a single large-scale system (Du et al., 2024; Kwon et al., 2025) and consistent with our own seed-based maps, we combined them into one network. To ensure our results didn't depend on a single parcellation approach, we repeated the network assignment step using the Yeo 17-network atlas (Yeo et al., 2011). Because this alternative prior lacked connectivity templates, assignments were based on spatial overlap alone.

Group Average Network Parcellation

Comparisons were made between individual network parcellations and a group-average parcellation. The primary group parcellation was defined as the mode network assignment across the 37 healthy controls used as priors here and in Lynch et al. (2024). For comparison, we also generated a group-average parcellation from the 10 individuals included in the main analyses. This was done by averaging their dense connectivity matrices and running them through the same precision functional mapping procedure. The resulting parcellation was highly similar to the Lynch et al. (2024) group average (Supplemental Figure S3), confirming its validity for our sample. As described above, individual network maps were also generated using the Yeo et al. (2011) 17-network parcellation as a spatial prior (Supplemental Figure S6). These were compared to the publicly available group-average 17 network parcellation (https://github.com/ThomasYeoLab/CBIG/tree/master/stable_projects/brain_parcellation/Yeo2011_fcMRI_clustering). For more straightforward comparison, LPFC networks in the Yeo 17-

841   network parcellation were manually renamed to match the networks used in the current study
842   where possible (their control A = frontoparietal A, control B = frontoparietal B, control C =
843   parietal memory, temporal parietal = language, default C = default A, default A = default B1,
844   default B = default B2, salience/ventral attention A = cingulo-opercular A, salience/ventral
845   attention B = cingulo-opercular B).
846

Lateral Prefrontal Cortex Mask
An average LPFC mask was created and used across all individuals. This mask was generated
by taking the union of Freesurfer aparc.a2009s parcellations from all 10 individuals, including
the following parcels: L_G_and_S_frontomargin, L_G_and_S_transv_frontopol, L_G_front_inf-
Opercular, L_G_front_inf-Triangul, L_G_front_middle, L_Lat_Fis-ant-Vertical, L_S_front_inf,
L_S_front_middle, L_S_front_sup, L_S_orbital_lateral, L_S_precentral-inf-part, and
L_S_precentral-sup-part.
854

Split Half Functional Network Reliability
For two individuals (PAN01 and PAN02), approximately 2.5 hours of motion-censored resting-
state data were collected (160 minutes and 146 minutes, respectively) in order to enable
reliability testing. These individuals were used as pilot cases to assess the reliability of the
precision functional mapping approach. Their data were split into odd and even sessions
(always collected on different days), and precision functional mapping was run independently on
each half, starting from the construction of functional connectivity matrices as described in
"Precision Functional Mapping."
863

The reliability of individual LPFC networks (Supplemental Figure S4) was assessed by
computing the Dice overlap for each LPFC network between the two halves. To quantify
whether within-individual reliability was greater than between-individual similarity, the average
dice overlap value per network was calculated over the two individuals, and a one-sided paired
t-test was used to compare the overlap of networks within individuals across split halves to the
overlap of networks across individuals within the same split half.
870

871   **Surface Area Analyses**
872   For each individual, the percent of total surface area for each network was calculated using the
873   final network estimates, masked to include only the LPFC. Surface area per network was
874   defined as the sum of the surface area of each vertex assigned to that network (Figure 1).
875   Vertex-wise surface areas were calculated using the Conte69 32k fs_LR template surface.
876

Association Network Density
For each individual, association network density was calculated at each vertex as the number of
unique association networks present within a fixed-radius neighborhood (Figure 4). This
approach was adapted from Power et al., 2013, where it was termed "community density" and
was calculated on all communities across different thresholds. Here, we refer to it as association
network density, as the calculation was based on the final labeled network identities and limited
to the 7 association networks (dorsal attention, frontoparietal, default A, default B,
salience/parietal memory, cingulo-opercular, language). For the Yeo 17-networks parcellation,
10 association networks were included (dorsal attention B, cingulo-opercular A, cingulo-
opercular B, frontoparietal A, frontoparietal B, default A, default B1, default B2, parietal memory,
language).
888

A geodesic distance matrix was computed using the Conte69 32k fs_LR group template surface
and applied uniformly across individuals. Density was calculated at five radii (6 mm, 8 mm, 10
mm, 12 mm, 14 mm), and we reported the average value across these radii per vertex.

Additionally, canonically low signal regions (the inferior temporal cortex and orbitofrontalcortex)
were excluded from density calculations via a custom mask. This is visible in the whole brain
view (see Supplementary Figure S8). The same mask was used for all subjects and was based
on mode 1000 BOLD signal values from an independent set of 120 subjects from a previous
study (Power et al., 2011).
897
Density Rotation Analysis
In each individual, the left hemisphere rostral cingulo-opercular network region was defined as a
region of interest (ROI) using Connectome Workbench's cifti-cluster function, followed by
manual selection of the rostral subregion from the resulting network clusters. Association
network density was computed by averaging the density values of all vertices within this ROI. To
assess whether the observed density was higher than expected by chance, a surface-based
rotation analysis was performed. The ROI was randomly rotated across the individual's cortical
surface 10000 times, preserving its shape and size. Rotated instances were retained only if all
ROI vertices remained within a single association network in the individual-specific parcellation.
For each valid rotation, the mean association network density was calculated. These values
were aggregated across individuals to generate a null distribution, against which the true ROI
value was compared (Figure 4C).
910
Task Paradigms
All individuals completed at least 30 minutes per task across 11 different cognitive tasks
targeting language, theory of mind, episodic projection, and cognitive control demands. Task
paradigms were selected from the literature based on their ability to elicit reliable LPFC activity
within single individuals and their suitability for repeat testing (i.e., being resilient to habituation
or allowing for easy generation of novel stimuli). Tasks were adapted with minimal modifications
from the original publications, with any relevant changes described below. All tasks were
implemented using Psychtoolbox-3.
919
920 **Language**
921 Two language tasks were administered: a visual language task and an auditory language task
922 (described below). A high-level language processing map was generated by averaging the
923 activation maps from both tasks.
924
<u>Visual Language Task:</u>
The visual language task was adapted from Fedorenko et al., 2011 using publicly available code
and stimuli (https://www.evlab.mit.edu/resources-all/download-localizer-tasks). Individuals
passively read either real sentences (Sentences condition) or pronounceable nonword
sequences (Nonword condition), presented one (non)word at a time. Each experimental block
lasted 18 seconds and included three trials. Each trial lasted 6 seconds, beginning and ending
with 100 ms of blank screen. During the trial, 12 (non)words were presented every 450 ms,
followed by a 400 ms response cue prompting participants to press a button to mark the end of
the trial.
934
Individuals completed 16 experimental blocks, intermixed with five 14-second fixation blocks
placed at the start of each run and after every four experimental blocks. Each person completed
at least six runs of the visual language task (5 minutes 58 seconds per run; 35 minutes 48
seconds total). Condition order was counterbalanced across runs and held constant across
individuals. Evidence suggests that contrasting Sentence > Nonwords blocks reliably identifies
high-level language processing regions, including in the LPFC (Fedorenko et al., 2011, 2012).
941

Auditory Language Task: The auditory language task was adapted from Scott et al., 2017
using publicly available code and stimuli (https://www.evlab.mit.edu/resources-all/download-
localizer-tasks), along with additional stimuli created in-house. Individuals listened to 18-second
clips of either intact speech (Intact condition) or acoustically degraded speech (Degraded
condition). Degraded clips retained auditory properties of speech but were not intelligible. To
expand the stimulus set beyond the 32 publicly available clips, we created 64 additional stimuli
by extracting 18-second segments from episodes of *The Moth* podcast and degrading them
using the procedure described below. Each run included 16 experimental blocks (one 18-
second audio clip per block) and five 14-second fixation blocks, placed at the beginning of each
run and after every four experimental blocks. All individuals completed six runs of the auditory
language task (5 minutes 58 seconds per run; 35 minutes 48 seconds total). Condition order
was counterbalanced across runs and held constant across individuals. Evidence suggests that
contrasting Intact > Degraded blocks reliably identifies high-level language-processing regions,
including regions in the LPFC (Salvo et al., 2024; Scott et al., 2017).
956
Auditory Degradation: To degrade novel audio clips, we followed the procedure outlined in Scott
et al. (2017) using their publicly available code. Briefly, a low-pass version of the intact clip was
created with a passband cutoff at 500 Hz. A noise track was generated by temporally
scrambling the intact audio and modulating it by the original amplitude envelope to preserve
naturalistic intensity variation. This noise signal was then bandpass filtered between 8000–
10,000 Hz and added to the low-pass version of the clip. All degraded clips were manually
checked to ensure they were unintelligible while preserving general acoustic similarity to the
original speech.
965
Episodic Projection
The episodic projection task was adapted from Andrews-Hanna et al., 2010 and DiNicola et al.,
2020. Code and stimuli for the task were generously provided by DiNicola and colleagues. This
task probes remembering and prospection demands by asking participants to respond to
scenarios regarding past, future, or present events. While participants were presented with
scenarios in six possible categories (Past Self, Present Self, Future Self, Past Non-Self, Present
Non-Self, Future Non-Self), only self-related scenarios were used in this manuscript. As in
DiNicola et al., 2020, a general episodic projection contrast was derived by averaging together
contrasts targeting Retrospection (Past Self > Present Self) and Prospection (Future Self >
Present Self). Episodic projection runs were 624 seconds long, consisting of 30 20-second
experimental blocks (5 per category) and 12-second fixation blocks beginning and ending the
run. Each experimental block consisted of a 5-second fixation period, a 10-second trial period,
and another 5-second fixation period. During the trial period, participants viewed a scenario and
related question and were asked to select from three possible responses with a button press.
The order of scenarios was randomized and kept consistent across participants. All participants
completed six runs of the episodic projection task (10 minutes 24 seconds each; 62 minutes 24
seconds total). Evidence suggests that the general episodic projection contrast dissociates
regions of default A from regions of default B, including regions in the LPFC (DiNicola et al.,
2020).
985
Theory of Mind
Two theory of mind tasks were administered: False Belief and Emotional/Physical Pain, both
replicated from DiNicola et al. (2020). A high-level theory of mind processing contrast was
created by averaging the contrasts from these two tasks (Belief > Photo and Emotional Pain >
Physical Pain). Evidence suggests that this contrast dissociates regions of default Network B
from regions of default Network A, including regions in the LPFC (DiNicola et al., 2020).
992

<u>False Belief/Photo:</u> The False Belief task was generously shared by the Braga Lab, adapted
from publicly available materials developed by Dodell-Feder et al., 2011
(https://saxelab.mit.edu/use-our-efficient-false-belief-localizer/). In this task, participants read
short stories in which either a character held a potentially false belief (False Belief condition) or
an object (e.g., a photograph or map) contained potentially false information (False Photo
condition). They then answered a true/false question via button press. Each run was 324
seconds long, with ten 30-second trials (five per condition) and 12-second fixation periods at the
beginning and end. Each trial included a 10-second story, 5-second response, and 15-second
fixation. Story order was randomized and consistent across participants. All individuals
completed four runs (5 minutes 24 seconds each; 21 minutes 36 seconds total).

<u>Emotional/Physical Pain:</u> The Emotional/Physical Pain task was generously shared by the
Braga Lab, adapted from publicly available materials developed by Jacoby et al., 2016
(https://saxelab.mit.edu/theory-mind-and-pain-matrix-localizer-narratives/). In this task,
participants read stories describing a character experiencing either emotional or physical pain
and rated the character's pain on a 1–4 scale via button press. Each run lasted 324 seconds,
with ten 30-second trials (five per condition) and 12-second fixation periods at the beginning and
end. Each trial included a 10-second story, 5-second response, and 15-second fixation. Story
order was randomized and consistent across participants. All individuals completed four runs (5
minutes 24 seconds each; 21 minutes 36 seconds total).

Cognitive Control
We report on eight contrasts derived from six distinct cognitive control tasks. Four tasks (MSIT,
VMSIT, Spatial Memory Span, Verbal Memory Span) were adapted from Fedorenko et al.
(2013), which identified "domain-general control" regions based on greater activation for "hard
" versus "easy" conditions across diverse tasks. Two additional tasks—Visual vs. Auditory
Attention and Visual vs. Auditory N-back—were adapted from Noyce et al., 2017 and Michalka
et al., 2015. We hypothesized these control tasks would show distinct recruitment from the
language, episodic projection, and theory of mind contrasts described above.

<u>Visual and Auditory Attention:</u> The visual and auditory sustained spatial attention task was
adapted from Michalka et al., 2015 with minimal modification. Participants were simultaneously
presented with four rapid serial streams of stimuli (two auditory, two visual), each containing
either numeric or alphabetical characters. They were instructed to monitor one stream for target
digits (1–4) while ignoring distractor letters in that stream ('A', 'F', 'G', 'H', 'J', 'K', 'L', 'M', 'N', 'P',
'R', 'X', 'Y') and the other three streams, which contained only digits (1–9, excluding 7). Each
visual stream was flanked by three additional digit-only distractor streams. Participants
responded via button press (1–4) to target digits in the attended stream, which occurred three
times per block. In sensorimotor control ("passive") blocks, participants viewed the same stream
layout, but all streams contained only digits (1–9), and participants pressed each response
button once at a relaxed pace. Fixation control blocks showed a white cross on a black screen.
Each run lasted 363.2 seconds and included an 8-second initial and 12-second final fixation, ten
26-second experimental blocks (two per stream type and two passive), two 26-second fixation
blocks, and 2.6-second audiovisual cue periods announcing upcoming block types. Each
experimental block contained 40 stimuli, presented audiovisually for 300 ms with a 350 ms inter-
stimulus interval. Block order was counterbalanced across runs and consistent across
individuals. A Visual Attention contrast was defined as Visual > Passive; an Auditory Attention
contrast was defined as Auditory > Passive. All individuals completed at least six runs (7
minutes 3.2 seconds each; 36 minutes 19 seconds total).

<u>Visual and Auditory N-Back:</u> The visual and auditory working memory task was adapted from
Noyce et al., 2017. Participants performed either a 2-back working memory task or a 0-back
sensorimotor control task on visual (faces) or auditory (animal sounds) stimuli, presented in
separate blocks. Visual stimuli were black-and-white photographs of young adult faces from the
Chicago Face Database (Ma et al., 2015) with male and female faces presented in separate
blocks to increase difficulty. Auditory stimuli consisted of cat and dog sounds, also presented in
separate blocks. Each run lasted 372 seconds and included an initial 8-second fixation period, a
mid-run 8-second fixation break (after four blocks), and a final 12-second fixation. Runs included
eight 43-second blocks: one per combination of stimulus category (Male Faces, Female Faces,
Cat Sounds, Dog Sounds) and demand level (2-back or sensorimotor). Each block began with a
3-second cue indicating the upcoming category, followed by a 40-second trial period. Visual
stimuli were displayed for 1000 ms with a 250 ms ISI; auditory stimuli lasted ~500 ms and were
spaced 1250 ms apart to match timing. In 2-back blocks, participants responded to each
stimulus as "new" or a "2-back repeat" (25% of trials). In sensorimotor blocks, no repeats
occurred, and participants made random button presses to each stimulus. Block order was
counterbalanced across runs and consistent across individuals. A Visual n-back contrast was
defined as Visual 2-back > Visual Sensorimotor; an Auditory n-back contrast as Auditory 2-back
> Auditory Sensorimotor. All participants completed at least six runs (6 minutes 12 seconds
each; 37 minutes 12 seconds total).
1062
<u>Spatial Memory Span:</u> The spatial memory span task was adapted from Fedorenko et al., 2011.
Participants viewed a 3×4 grid and were sequentially shown spatial locations to remember, with
either four or eight total locations depending on difficulty (easy vs. hard conditions). At the end
of each trial, they selected the correct grid from two options via button press; the incorrect grid
contained one or two wrong locations. In easy blocks, locations were presented one at a time
(four total); in hard blocks, locations were shown in pairs (eight total). Each run lasted 436
seconds and included ten 34-second experimental blocks (five easy, five hard) and six 16-
second fixation blocks. Each experimental block contained four trials, with each trial consisting
of a 0.5-second fixation, four 1-second stimulus presentations, and a 4-second response
window. The order of easy and hard blocks was counterbalanced across runs and kept
consistent across individuals. All participants completed at least four runs (7 minutes 52
seconds each; 31 minutes 28 seconds total).
1075
<u>Verbal Memory Span:</u> The verbal memory span task was adapted from Fedorenko et al., 2011.
Participants were sequentially presented with spoken digit names and asked to remember either
four or eight digits depending on condition difficulty (easy vs. hard). At the end of each trial, they
selected the correct sequence from two options via button press; incorrect options contained
one or two wrong digits. In hard blocks, digit names were presented in pairs (eight total); in easy
blocks, they were presented one at a time (four total). Each run lasted 436 seconds and
included ten 34-second experimental blocks (five easy, five hard) and six 16-second fixation
blocks. Each block contained four trials, with each trial consisting of a 0.5-second fixation, four
1-second digit presentations, and a 4-second response window. Block order was
counterbalanced across runs and consistent across individuals. All participants completed at
least four runs (7 minutes 52 seconds each; 31 minutes 28 seconds total).
1087
<u>Numeric Multi-Source Interference Task (MSIT):</u> The numeric Multi-Source Interference Task
was adapted from Fedorenko et al., 2011. which was originally adapted from Bush et al., 2003.
Participants were shown digit triplets (e.g., "123") and asked to respond based on the identity—
rather than the position—of the unique (non-repeated) digit. In the easy condition, the identity of
the target digit matched its position, and the distractors were not valid response options (e.g.,
"003"). In the hard condition, the identity of the target digit did not match its position, and the

distractors were valid response options (e.g., "131"). Each run lasted 396 seconds and included
two 30-second fixation blocks (at the beginning and end) and eight 42-second experimental
blocks. Each experimental block contained 24 trials with a 1.5-second stimulus presentation and
a 0.25-second interstimulus interval (ISI). The order of hard and easy blocks was
counterbalanced across runs and consistent across participants. All individuals completed at
least four runs (6 minutes 36 seconds each; 26 minutes 24 seconds total).
1100
Verbal Multi-Source Interference Task (VMSIT): The Verbal Multi-Source Interference Task
(VMSIT) was adapted from Fedorenko et al., 2011 which was originally adapted from Bush et
al., 2003. Participants were shown triplets of words (e.g., "none," "left," "middle," "right") and
asked to respond based on the meaning of the unique (non-repeated) word, rather than its
spatial position. In the easy condition, the position of the target word matched its meaning, and
the distractor words were not valid response options (e.g., "none none right"). In the hard
condition, the target word's position conflicted with its meaning, and the distractor word(s) were
valid response options (e.g., "left right left"). Each run lasted 396 seconds and included two 30-
second fixation blocks (at the beginning and end) and eight 42-second experimental blocks.
Each block contained 24 trials, with 1.5-second stimulus presentation and a 0.25-second
interstimulus interval (ISI). The order of easy and hard blocks was counterbalanced across runs
and consistent across individuals. All participants completed at least four runs (6 minutes 36
seconds each; 26 minutes 24 seconds total).
1114
Task Analysis
All tasks were analyzed using the general linear model (GLM) implemented via the
3dDeconvolve function in AFNI. All task conditions were included in the block design model.
Surface-projected data were analyzed within individuals on a per-run basis. For each contrast, t-
statistic maps were generated and converted to z-statistic maps in AFNI, then averaged across
runs to produce a mean z-statistic map for each individual and contrast.
1121
Within Individual Task fMRI Reliability
While there is growing literature on the amount of resting-state fMRI data needed for reliable
functional connectivity measures (Birn et al., 2013; Elliott et al., 2020; Gordon et al., 2017;
Laumann et al., 2015; Noble et al., 2017), it is less clear how much data is required for reliable
within-individual task fMRI. The answer likely depends on the task contrast and the design of
the paradigm (e.g., block, event-related, or mixed). To ensure the robustness of our study, we
first collected task data in two pilot individuals (PAN01 and PAN02), acquiring approximately
twice as much data as in previous studies. This allowed us to assess test-retest reliability for
each task. As shown in Supplemental Figure S5, we found that most tasks demonstrated strong
test-retest reliability with 30 minutes of data per split half. We evaluated this based on the
correlation of LPFC z-statistic maps. Based on these results, we proceeded with collecting at
least 30 minutes of data per task per individual in the main study.
1134
Network Task Activity Comparison
To compare across cognitive domains, contrasts were grouped as described in the Task
Paradigms section. For the language processing map, z-statistic maps from the [Sentence –
Nonword] and [Intact Speech – Degraded Speech] contrasts were averaged. For the theory of
mind map, [Emotional Pain – Physical Pain] and [False Belief – Photo] maps were averaged.
For the episodic projection map, [Past Self – Present Self] and [Future Self – Present Self]
maps were averaged. For each of 8 cognitive control tasks, a z-statistic map was created from
the relevant high demand – low demand contrast (N-back: 2-back versus 0-back, Memory Span:
8 items versus 4 items, Multi-Source Interference: conflict versus no-conflict, Attention:
sustained attention versus sensorimotor control).

1145
1146 To compare the activity of LPFC network regions, all individual network maps were masked
1147 using the LPFC mask described above. Mean z-values for each network were extracted from
1148 each of the contrast maps per individual, using their own LPFC individual-specific network
1149 maps, resulting in one z-value per network per individual per task.
1150
For the specialized cognitive domains (Language Processing, Theory of Mind, Episodic
Projection), one-tailed paired t-tests were used to compare the activation of the hypothesized
target LPFC network—language, default B, and default A, respectively—against each of the
other six LPFC association networks (language, default A, default B, frontoparietal, dorsal
attention, cingulo-opercular). To correct for multiple comparisons, p-values were Bonferroni
corrected for six comparisons (p × 6).
1157
1158 For the cognitive control tasks, one-tailed one-sample t-tests were used to identify LPFC
1159 association networks with significantly greater-than-zero activation. These p-values were
1160 Bonferroni corrected for seven comparisons (p × 7). All statistical tests were conducted using the
1161 *ttest* function in MATLAB.
1162
1163 **Task Visualization**
1164 The activation maps for all tasks were visualized relative to the boundaries of the relevant LPFC
1165 networks. In all figures, activity was thresholded to display the top 25% of LPFC vertices per
1166 hemisphere, based on z-statistic values, excluding any vertices with $z < 0$.
1167
Task/Network Overlap
The spatial overlap between thresholded task activation maps and different network
parcellations—individual-specific, group-average, and other-individuals' maps—was quantified
using the Dice similarity coefficient (Dice, 1945). Individual-specific task activation maps were
thresholded at three levels (top 15%, 20%, and 25% of LPFC vertices) and compared to each
LPFC network parcellation. For each individual and task, the average Dice overlap values
(across thresholds) were compared between individual networks and (1) group-average
networks or (2) other-individual networks using one-sided paired t-tests. P-values were
corrected for 2 comparisons (p × 2). To test for regional specificity, the difference in Dice overlap
between individual-specific and other-individual networks was compared for LPFC versus non-
LPFC regions. The same procedure described above was repeated for non-LPFC task
activations and networks. LPFC and non-LPFC values were compared using one-sided paired t-
tests for each task and a composite score averaged across the three tasks.
1181
Distance to Control Network Analysis
It was observed that cognitive control activations appeared near the borders of multiple control
networks (frontoparietal, dorsal attention, cingulo-opercular). This was quantified by comparing
the mean distance of activated versus non-activated FP, DAN, and CO network vertices (at $z >$
1, $z > 2$, $z > 3$) to the nearest vertex of either of the other two control networks (for FP, the
nearest DAN or CO vertex, for example). Minimum distances were measured from each vertex
to each network were measured on the Conte32k_fsLR template surface which was used
across all individuals.
1190
1191 **Task Activation Similarity Analysis**
1192 To compare the similarity of activation patterns for each of the 8 cognitive control demands
1193 within and across individuals (Figure 8B), unthresholded task activation maps masked to the
1194 LPFC were correlated within and across individuals.
1195

Seed Connectivity Analyses

Seed-based connectivity was used to validate network parcellations throughout the manuscript (Figures 3 and 7, Supplemental Figures S7, S12, S14). In each case, the seed was manually selected from within network boundaries to best represent the network's characteristic connectivity pattern.

Variant ROI Task Analyses

Language and default B variant regions described in the manuscript (Figure 7 and Supplemental Figure S14) were defined as ROIs by clustering the relevant network using Connectome Workbench's cifti-cluster function and manually selecting the cluster corresponding to the variant. In cases where the variant partially overlapped with the canonical group-average network, the overlapping portion of the group-average network cluster was subtracted from the manually defined ROI. Task activation z-statistics were averaged across all vertices within each ROI for both the variant individual and all individuals without a variant in that location. A variant task activation metric was computed as the difference between the relevant specialized demand (language processing or theory of mind) and the expected cognitive control demand (spatial memory). This metric was then compared between variant individuals and non-variant individuals.

Data and Software Availability

Raw MRI data, preprocessed cifti resting-state and task timecourses, task z-statistic maps, and individual-specific networks, have been deposited in the Openfmri data repository (https://openfmri.org/) under the label "Ladwig 2025 - LPFC". Code to perform preprocessing and analysis is available at https://github.com/GrattonLab/Ladwig-LPFC.

Declaration of generative AI and AI-assisted technologies in the writing process

During the preparation of this work the author(s) used chatGPT in order to improve the readability of the text. After using this tool/service, the author(s) reviewed and edited the content as needed and take(s) full responsibility for the content of the publication.

Declaration of competing interests

The authors declare no competing interests.

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