A representational similarity analysis of cognitive control during color-word Stroop

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Abstract

Progress in understanding the neural bases of cognitive control has been supported by the paradigmatic color-word Stroop task, in which a target response (color name) must be selected over a more automatic, yet potentially incongruent, distractor response (word). For this paradigm, models have postulated complementary coding schemes: dorsomedial frontal cortex (DMFC) is proposed to evaluate the demand for control via incongruency-related coding, whereas dorsolateral prefrontal cortex (DLPFC) is proposed to implement control via goal and target-related coding. Yet, mapping these theorized schemes to measured neural activity within this task has been challenging. Here, we tested for these coding schemes relatively directly, by decomposing an event-related color-word Stroop task via representational similarity analysis (RSA). Three neural coding models were fit to the similarity structure of multi-voxel patterns of human fMRI activity, acquired from 65 healthy, young-adult males and females. Incongruency coding was predominant in DMFC, whereas both target and incongruency coding were present with indistinguishable strength in DLPFC. In contrast, distractor coding was strongly encoded within early visual cortex. Further, these coding schemes were differentially related to behavior: individuals with stronger DLPFC (and lateral posterior parietal cortex) target coding, but weaker DMFC incongruency coding, exhibited less behavioral Stroop interference. These results highlight the utility of the RSA framework for investigating neural mechanisms of cognitive control and point to several promising directions to extend the Stroop paradigm.

Significant Statement

How the human brain enables cognitive control — the ability to override behavioral habits to pursue internal goals — has been a major focus of neuroscience research. This ability has been frequently investigated by using the Stroop color-word naming task. With the Stroop as a test-bed, many theories have proposed specific neuroanatomical dissociations, in which medial and lateral frontal brain regions underlie cognitive control by encoding distinct types of information. Yet providing a direct confirmation of these claims has been challenging. Here, we demonstrate that representational similarity analysis (RSA), which estimates and models the similarity structure of brain activity patterns, can successfully establish the hypothesized functional dissociations within the Stroop task. RSA may provide a useful approach for investigating cognitive control mechanisms.

1   Introduction

Goals, held in mind, can be used to overcome behavioral habits. Understanding how the human brain enables such *cognitive control* has been a fundamental interest of both basic and translational cognitive neuroscience. Toward this end, the use of *response conflict* tasks has been instrumental (e.g., Botvinick, Braver, Barch, Carter, & Cohen, 2001; Ridderinkhof, Ullsperger, Crone, &

Nieuwenhuis, 2004). These tasks involve trials in which a less-automatic, but goal-relevant course of action, the *target* response, must be selected in the face of a habitual, but goal-irrelevant alternative, the *distractor*. The paradigmatic example is the color-word Stroop task (MacLeod, 1991; Posner & Snyder, 1975; Stroop, 1935): on each trial, the hue of a word must be named, despite the word expressing a potentially conflicting, that is, *incongruent*, color (Figure 1, C). A major goal in this field has been to use measures of neural activity evoked by response conflict tasks, such as Stroop, to test models of cognitive control.

One broad, neurocomputational-level model ascribes particular roles to different frontoparietal regions in overcoming response conflict (Miller & Cohen, 2001; Shenhav, Botvinick, & Cohen, 2013). Central to this view is the type of task information these regions encode. The dorsomedial frontal cortex (DMFC) is proposed to "evaluate" demand for cognitive control, via encoding of *incongruency-related information* (Figure 1, A, bottom row). Such information, according to this view, is used by dorsolateral prefrontal cortex (DLPFC), in concert with lateral posterior parietal cortex (LPPC), to "implement" control, via encoding of goal and *target-related information* (Figure 1, A, top row). Thus, this model predicts key functional dissociations between medial and lateral frontoparietal cortex (Figure 1, B). But, although this view has been influential, directly establishing these dissociations during the performance of standard color-word Stroop tasks has been difficult.

To date, the most traction on this problem has been gained via fMRI designs that temporally dissociate presentation of task-rule and incongruency-related information, in which subjects were instructed prior to each Stroop trial about the task to perform (color-naming, word-reading; Floden, Vallesi, & Stuss, 2010; MacDonald, Cohen, Stenger, & Carter, 2000). But, while these studies generally found supportive evidence for the key claims, results were subject to three notable limitations. First, these studies were likely underpowered for fMRI (e.g., N = 12 in Floden et al., 2010; N = 9 in MacDonald et al., 2000). This fact alone warrants a follow-up study. Second, it is unclear whether the results extend to the more-standard Stroop-task design, in which task rules are not explicitly instructed prior to each trial, but are instead internally maintained. For example, goal-relevant coding in DLPFC may depend upon such explicit rule instruction. Third, the prior results do not speak to functional dissociations *within* a single Stroop trial, during which interference is actually experienced and resolved. It is therefore possible, for instance, that the role of DLPFC (or other frontoparietal regions) in Stroop is primarily preparatory, and is less critical during actual interference resolution.

To address these questions, a neuroanatomically precise technique is needed that does not rely on temporal dissociations, but can instead read out multiple, simultaneously encoded sources of task information from individual brain regions of interest. Multivariate (multivoxel) pattern analysis (MVPA) of fMRI, in popular use for over a decade (Cox & Savoy, 2003; Edelman, Grill-Spector, Kushnir, & Malach, 1998; Haxby, Gobbini, Furey, & Ishai, 2001), accomplishes exactly this purpose.

Surprisingly, however, these methods have not been brought to bear on the question of a functional dissociation between medial and lateral frontoparietal cortex in resolving Stroop conflict.

We fill this gap in the literature by using Representational Similarity Analysis (RSA; Edelman, Grill-Spector, Kushnir, & Malach, 1998); Kriegeskorte 2008), a specific MVPA framework, to test for dissociations in frontoparietal coding during Stroop-task performance (Figure 1, C–E). We conducted a retrospective analysis of data collected as part of the Dual Mechanisms of Cognitive Control project (Braver, Kizhner, Tang, Freund, & Etzel, 2020). Our primary goal was a proof-of-principle: to demonstrate the potential of RSA for testing theorized distinctions in neural coding within cognitive control tasks, such as the Stroop (Freund, Etzel, & Braver, 2021).

Figure 1 *(previous page)*: Schematic of framework and hypotheses. **A**, Conceptual framework. Cognitive theories and computational models of control have decomposed the classic color-word Stroop task into three *task dimensions*: the *target* (the goal-relevant mapping of the stimulus hue to the target response, i.e., color naming), the *distractor* (the prepotent but goal-irrelevant mapping of the stimulus word to the non-target or distractor response, i.e., word reading), and *incongruency* (whether the target and distractor responses match or mismatch). **B**, Hypotheses. Neuroscientific frameworks of cognitive control propose that representation of task dimensions is anatomically dissociated across medial and lateral frontoparietal cortices. Dorsomedial frontal cortex (DMFC), including dorsal anterior cingulate (dACC) and pre-supplementary motor area (pre-SMA), is proposed to "evaluate" demands for control, using information correlated with the *incongruency* dimension (**A**, bottom row), to signal when (and how) the current attentional or action selection policies are suboptimal (Ridderinkhof, Ullsperger, Crone, & Nieuwenhuis, 2004; Shenhav, Botvinick, & Cohen, 2013). Conversely, dorsolateral PFC (DLPFC), in concert with lateral posterior parietal cortex (LPPC), is proposed to guide, or "implement," goal-driven attentional selection and mapping of hue–target-response processes, by way of representing information related to the goal-dependent target dimension (**A**, top row; Miller & Cohen, 2001); Buschman & Miller, 2007). Double dissociations are therefore predicted at multiple levels of analysis. At the group level, incongruency coding (orange univariate distributions) should predominate in DMFC, whereas target coding (green univariate distributions) should predominate in DLPFC. At the individual level, if the strength of target-related coding in DLPFC reflects the robustness of goal-driven selection, then subjects with stronger DLPFC target coding should resolve Stroop interference more efficiently (green bivariate distribution; Kane & Engle, 2002; Braver, 2012). Conversely, if the strength of incongruency-related coding in DMFC indicates a maladaptive selection policy, then subjects with stronger incongruency-related coding should resolve Stroop interference *less* efficiently (orange bivariate distribution; Braver, 2012; MacDonald, Cohen, Stenger, & Carter, 2000). The β notation in axis titles corresponds to that used in **E** (and throughout this manuscript); $\bar{\beta}$ indicates mean over subjects. **C**–**E**, Analytic framework. Participants performed a color-word Stroop task while undergoing an fMRI scan (**C**). To derive neural correlates of the theorized task dimensions (in **A**–**B**), a general linear model estimated the BOLD response evoked by sixteen unique Stroop stimuli (e.g., "WHITE" displayed in blue hue) independently for each voxel. We analyzed a mostly-incongruent trial condition, in order to obtain balanced estimates of each trial type. The Glasser et al. (2016) multi-modal atlas was used to parcellate cortex (**D**, light silver borders). Contiguous sets of parcels that tiled our regions of interest ("superparcels") were defined and treated as analytic units (see Figure 1-1 for list). Within each superparcel, linear correlations among response patterns from the sixteen stimuli were estimated to form an empirical similarity matrix (**E**, left; stimuli that were white are presented here in grey). Through rank regression, these matrices were fit to three representational models (**E**, right), which corresponded to the three hypothesized dimensions of the Stroop task (**A**). The resulting β coefficients summarized the extent to which a parcel emphasized, within its distributed activity patterns, the representation of each unique task dimension. These β coefficients were used as the primary dependent variables in group-level analyses, and primary independent variables in individual-level analyses (e.g., as in **B**). Critically, we verified the specificity of our design and analysis via simulation (Figure 1-2; *cf.*, Cai, Schuck, Pillow, & Niv, 2019).

2 Materials and Methods

We report how we determined our sample size, all data exclusions, all manipulations, and all measures in the study (Simmons, Nelson, & Simonsohn, 2011).

2.1 Code, Data, and Task Accessibility

Code (R Core Team, 2019) and data to reproduce all analyses, in addition to supplementary analysis reports, are publicly available (https://doi.org/10.5281/zenodo.4784067). As part of the planned data release of the Dual Mechanisms of Cognitive Control project, raw and minimally preprocessed fMRI data have been deposited on OpenNeuro (10.18112/openneuro.ds003465.v1.0.3). Additionally, the authors will directly share the specific fMRI data used for this study upon reasonable request. Task scripts are available at the DMC project website (http://pages.wustl.edu/dualmechanisms/tasks). More detailed information regarding all aspects of the project can be found on the Project's OSF page (https://osf.io/xfe32/).

2.2 Participants

Individuals were recruited from the Washington University and surrounding St Louis metropolitan communities for participation in the Dual Mechanisms of Cognitive Control project. The present study began with a subset ($N = 66$; 38 women, 26 men, 1 "prefer not to answer") of these subjects: those with a full set of imaging and behavioral data from the Stroop task during a particular scanning session (the *proactive* session), selected for methodological reasons (see *Selection of Data*). One subject was excluded from all analyses due to a scanner error. We split the remaining sample into two sets of individuals: a *primary analysis set* ($N = 49$; 27 women, 21 men, 1 "prefer not to answer"), which we used in all analyses, and a *validation set* ($N = 16$; 11 women, 5 men), which was only used in the *Model Selection* analysis (see below). This unbalanced partitioning was done to account for the familial structure present within our sample. Specifically, subjects within each set (primary analysis, validation) were all unrelated; however, subjects within the validation set were co-twins of 16 subjects within the primary analysis set. Two of these co-twins were selected for use in the *primary analysis set* as their respective co-twins had atypically high rates of response omission ($>10\%$; $>20\%$ errors of any type); the remaining co-twins were randomly selected. Critically, partitioning the sample in this way ensured that the primary analysis set was a random sample of independent subjects.

The partitioning of the data into two subsets also afforded the opportunity to use the validation subset as held-out data for evaluation of the brain–behavior model within the *Model Selection* analysis. As we performed this sorting of individuals into primary and validation sets only once and did not analyze the validation-set data except to assess predictive accuracy of the final selected model, the validation set provides an unbiased assessment of predictive accuracy, in the sense that no statistical "double-dipping" could have occurred. But because the sets are familially dependent,

it is perhaps more accurate to consider the validation-set analyses as assessing a kind of test–retest reliability (i.e., while eliminating the potential confound of practice effects), rather than providing an estimate of out-of-sample predictive accuracy. To evaluate this matter, follow-up control analyses were conducted in which the co-twins were removed from the primary analysis set.

2.3 Experimental Design and Statistical Analysis

2.3.1 Task

Participants performed the verbal color-word Stroop (1935) task. Names of colors were visually displayed in various hues, and participants were instructed to "say the name of the color, as fast and accurately as possible; do not read the word".

The set of stimuli consisted of two intermixed subsets of color-word stimuli (randomly intermixed during the task): a *mostly incongruent* and an *unbiased* set. Each stimulus set was created by pairing four color words with four corresponding hues in a balanced factorial design, forming 16 unique color-word stimuli within each set. The *mostly incongruent* set consisted of stimuli with hues (and corresponding words) 'blue' (RGB = 0, 0, 255), 'red' (255, 0, 0), 'purple' (128, 0, 128), and 'white' (255, 255, 255); the *unbiased* group, of 'black' (0, 0, 0), 'green' (0, 128, 0), 'pink' (255, 105, 180), and 'yellow' (255, 255, 0). These words were centrally presented in uppercase, bold Courier New font on a grey background (RGB = 191, 191, 191). Of stimuli within the *mostly incongruent* set, incongruent stimuli were presented to subjects more often than congruent stimuli (per block, proportion congruent = 0.25). *Unbiased* stimuli were presented with a balanced frequency (proportion congruent = 0.5). These manipulations of incongruency statistics are standard manipulations to elicit proactive control (Bugg, 2014; e.g., Gonthier, Braver, & Bugg, 2016) and were performed to investigate questions outside the scope of the current study. Thus, as described further below, the unbiased stimulus set was excluded from all analyses.

Each trial (e.g., Figure 1, C) began with a central fixation cross, presented for 300 ms on a grey background (RGB = 191, 191, 191). The color-word stimulus, preceded by a blank screen following fixation offset (100 ms), was centrally presented for a duration of 2000 ms, fixed across trials. The duration of the inter-trial interval (triangle of fixation crosses) was either 900, 2100, or 3300 ms, selected randomly (with uniform probability). Each of two scanning runs consisted of three blocks of 36 trials, intermixed with 4 resting fixation blocks, during which a fixation cross appeared for 30 s. This formed a mixed block-event design (Petersen & Dubis, 2012). Each of the 16 *mostly incongruent* stimuli — that is, each unique colored word (e.g., "BLUE" displayed in red hue) — were presented in both runs. Within each run for each participant, *mostly incongruent* stimuli were presented an equal number of times within each block. Within each block, stimulus order was fully randomized.

2.3.2 Selection of Data

We focused our fMRI pattern analyses solely on trials from the *mostly incongruent* stimulus set within a particular scanning session (the "proactive" session) of our Stroop task. This selection was made purely on the basis of methodological reasoning: these trials were the only set of trials within the larger Dual Mechanisms project in which each unique Stroop stimulus (i.e., one of the 16 color-word combinations) was presented an equal number of times (9) to each participant, constituting a balanced design. Balanced designs ensure that differences in the total number of trials per condition cannot explain any differences observed in pattern correlations among conditions.

2.3.3 Display and Recording Systems

The experiment was programmed in EPrime 2.0 ("E-Prime 2.0," 2013), presented on a Windows 7 Desktop, and back-projected to a screen at the end of the bore for viewing via a mirror head-mount. Verbal responses were recorded for offline transcription and response-time (RT) estimation. The first 45 participants spoke into a MicroOptics MR-compatible electronic microphone (MicroOptics Technologies, Inc.); due to mechanical failure, however, we replaced this microphone with the noise-cancelling FOMRI III (OptoAcoustics, Ltd.), which subsequent participants used. A voice-onset processing script (from the MATLAB Audio Analysis Library) was used to derive response-time estimates on each trial via spectral decomposition (the accuracy of which was verified by manually coding response times from a subsample of subjects and ensuring the two methods gave similar estimates). Code for this algorithm is available within the Dual Mechanisms GitHub repository (https://github.com/ccplabwustl/dualmechanisms/tree/master/preparationsAndConversions/audio).

Importantly, we verified that the change in microphone did not induce confounding between-subject variance in response-time measures of interest. While response time estimates recorded via the Micro-Optics microphone tended to be slower ($b = 102.59, p = 0.01$) and more variable ($\chi^2_9 = 3655, p = 0$), the magnitude of the Stroop effect was not observably impacted by the microphone change ($b = -5.88, p = 0.64$).

2.3.4 Image Acquisition, Preprocessing, and Generalized Linear Model

The fMRI data were acquired with a 3T Siemens Prisma (32 channel head-coil; CMRR multi-band sequence, factor = 4; 2.4 mm isotropic voxel, with 1200 ms TR, no GRAPPA, ipat = 0), and subjected to the minimally pre-processed functional pipeline of the Human Connectome Project (version 3.17.0), outlined in (Glasser et al., 2013). More detailed information regarding acquisition can be found on the Project OSF site (https://osf.io/tbhfg/). All analyses were conducted in volumetric space; surface maps are displayed in figures only for ease of visualization. Note that prior to revision of this manuscript, the data were re-processed with fMRIPrep, using the standard fMRIPrep pipelines (Esteban et al. 2019; 2018; see https://fmriprep.org/en/stable/citing.html#note-for-reviewers-and-editors for more information). At this point, the pre-

processed results with the HCP pipelines were inadvertently removed. Thus, some follow-up control analyses were conducted with the fMRIPrep-preprocessed data (Figure 2-4, Figure 3-2, Figure 4-4). The fMRIPrep pipeline was implemented in a Singularity container (Kurtzer, Sochat, & Bauer, 2017) with additional custom scripts used to implement file management (more detail on the pipeline is available at https://osf.io/6p3en/; container scripts are available at https://hub.docker.com/u/ccplabwustl).

After pre-processing, to estimate activation patterns, we fit a whole-brain voxel-wise generalized linear model (GLM) to blood-oxygen-level dependent (BOLD) time-series in AFNI, version 17.0.00 (R. W. Cox, 1996). To build regressors of primary interest, we convolved with a hemodynamic response function [via AFNI's *BLOCK(1,1)*] 16 boxcar timecourses, each coding for the initial second of presentation of a *mostly incongruent* stimulus that resulted in a correct response. We also included two regressors [similarly created via *BLOCK(1,1)*] to capture signal associated with congruent and incongruent trials of non-interest (*unbiased* stimuli) that prompted correct responses, an "error regressor" coding for any trial in which a response was incorrect or omitted (via *BLOCK*), a sustained regressor coding for task versus rest (via *BLOCK*), a transient regressor coding for task-block onsets [as a set of piecewise linear spline functions via *TENTzero(0,16.8,8)*], six orthogonal motion regressors, five polynomial drift regressors (order set automatically) for each run, and an intercept for each run. These models were created via *3dDeconvolve* and solved via *3dREMLfit*. The data for each subject's model consisted of 2 runs $\times$ 3 blocks $\times$ 36 trials (144 from the *mostly incongruent* stimulus group, 72 from *unbiased*). Frames with FD > 0.9 were censored.

2.3.5   Definition of Regions of Interest

Our primary hypotheses concerned a set of six anatomical regions: DMFC, DLPFC, and LPPC in each hemisphere. Consequently, our primary analyses employed a targeted ROI-based analysis approach. Rather than defining functional regions of interest (ROIs) via a whole-brain searchlight, which has known issues (Etzel, Zacks, & Braver, 2013), we defined ROIs via a cortical parcellation atlas. We selected the MMP atlas (Glasser et al., 2016) for two reasons. (1) The atlas was developed recently via multi-modal imaging measures, and (2) individual MMP parcels are relatively interpretable, as they are heterogeneously sized and have been explicitly connected to a battery of cognitive tasks (Assem, Glasser, Van Essen, & Duncan, 2020), the canonical functional connectivity networks (Ji et al., 2019), and a large body of neuroanatomical research (Glasser et al., 2016). We used a volumetric version, obtained from https://figshare.com/articles/HCP-MMP1_0_projected_on_MNI2009a_GM_volumetric_in_NIfTI_format/3501911?file=5534024 (also available on the project GitHub repository; see *Code, Data, and Task Accessibility*). We then defined a set of six spatially contiguous sets of MMP parcels (three in each hemisphere), which we refer to as "superparcels," that corresponded to each of our ROIs. For full superparcel definitions, see Figure 1-1. "DMFC" was defined as the four parcels covering SMA–pre-SMA and DACC. "DLPFC" was defined as the four parcels that cover middle frontal gyrus (i.e., mid-DLPFC). "LPPC" was

defined as all parcels tiling IPS (both banks and fundus), from the occipital lobe to primary so-matosensory cortex. The overwhelming majority of parcels that met these anatomical criteria were assigned, within a previous report, to the cinguloopercular (most of DMFC), frontoparietal (most of DLPFC), and dorsal-attention (most of LPPC) control networks (Ji et al., 2019). Further, these ROI definitions contain several parcels that correspond to key nodes within the "multiple demand'' network (Assem, Glasser, Van Essen, & Duncan, 2020). To assess the robustness of our results to particular superparcel definitions, we additionally used alternative, more inclusive, superparcel definitions of DMFC and DLPFC (see Figure 1-1). For the brain–behavior model selection analysis (see *Model Selection*), we compiled a larger set of anatomically clustered MMP parcels, covering regions across the cortex (Figure 3-3). Two additional, non-MMP ROIs were included in this set, in order to give better coverage of particular functional brain regions. A mask for ventral somatomotor cortex (the "SomatoMotor–Mouth" network) was obtained from the Gordon atlas (Gordon et al., 2016), as the MMP does not split somatomotor cortex into dorsal and ventral divisions. A mask for left ventral occipito-temporal cortex (encompassing the "visual word-form" area), was obtained using MNI coordinates $-54 < x < -30, -70 < y < -45, -30 < z < -4$, specified in a prior report (Twomey, Kawabata Duncan, Price, & Devlin, 2011). To remove cerebellar voxels from this ROI, we used the Deidrichshen atlas (Diedrichsen, 2006) hosted by AFNI (https://afni.nimh.nih.gov/pub/dist/atlases/SUIT_Cerebellum/SUIT_2.6_1/).

2.3.6   Estimation of Coding Strength β

To estimate the regional strength of target, distractor, and incongruency coding, we used the representational similarity analysis framework (RSA; Kriegeskorte, 2008). The RSA framework consists of modeling the observed similarity structure of activation patterns with a set of theoretically specified model similarity structures (Figure 1, E). For a given subject and cortical region, fMRI GLM coefficient estimates for each of the 16 conditions of interest (4 colors factorially paired with 4 words; e.g., the word "WHITE" presented in blue hue) were assembled into a condition-by-voxel activity pattern matrix $\mathbf{B}$. The observed similarity structure was estimated as the condition-by-condition correlation matrix $\mathbf{R} = \mathrm{Cor}(\mathbf{B})$. Cell R_{ij} of this matrix gives the linear correlation observed between activity patterns evoked by conditions i and j. Model similarity structures were specified in this same correlation matrix form. The target model assumed that conditions (stimuli) with the same hue will evoke identical patterns, regardless of whether the words or congruency match (or mismatch). That is, if the hue of condition $i = j$, this model predicts $R_{ij} = 1$, otherwise 0. (In the 'target' matrix in Figure 1, E, the only cells equal to 1 — i.e., blue cells — are those in which the stimulus hues match.) The distractor model assumed that conditions with the same word will evoke identical patterns, regardless of the hue or congruency. (If the word of condition $i = j$, $R_{ij} = 1$, otherwise 0. In the 'distractor' matrix in Figure 1, E, the only cells equal to 1 are those in which the stimulus words match.) The incongruency model assumed that only conditions that were incongruent would evoke identical patterns, regardless of the hue or word. (If i and j

are both incongruent, $R_{ij} = 1$, otherwise 0. In the 'incongruency' matrix in Figure 1, E, the only cells equal to 1 are those in which both stimuli are incongruent.) As a covariate of non-interest, we also included a model capturing similarity between congruent conditions (if i and j are both congruent, $R_{ij} = 1$, otherwise 0). We additionally examined alternative approaches to RSA modeling of incongruency, to see if our results were robust to this parameterization of incongruency and congruency (see Method section *Alternative RSA Incongruency Models*).

These four models were jointly fitted to the observed similarity structure from each region through multiple regression (ordinary least-squares), separately for each subject. The response vector $\mathbf{y}$ and design matrix of this regression were assembled in a series of steps. (1) The 120 unique off-diagonal elements of of each similarity matrix (one observed and four models) were extracted and unwrapped into vectors. (2) The four model similarity vectors were separately z-scored and assembled into columns. This formed the RSA design matrix. (3) The observed similarity vector was rank-transformed (Nili et al., 2014) then z-score standardized, to form a vector $\mathbf{r}$. (4) The vector $\mathbf{r}$ was prewhitened to remove a specific nuisance component. This component stemmed from the task design: though each *mostly incongruent* stimulus occurred an equal number of times throughout the course of a session, these stimuli were not fully balanced across the two scanning runs. Specifically, half of the stimuli were presented three times in the first run versus six in the second (vice versa for the other half). As each scanning run contains a large amount of run-specific noise (Alink, Walther, Krugliak, Bosch, & Kriegeskorte, 2015; Mumford, Davis, & Poldrack, 2014), this imbalance across runs could lead to a bias in the resulting β coefficients, in which pattern similarity of stimuli that mostly occurred within the same run would be inflated. We formalized this component of bias as another model similarity vector, $\mathbf{v}$, with elements equal to 1 if *the run in which condition i most frequently occurred = the run in which condition j most frequently occurred*, otherwise 0. The magnitude of this bias was estimated as the slope term b_1 in a linear regression $\mathbf{r} = \mathbf{v}b_1 + b_0 + \boldsymbol{\epsilon}$, where b_0 is the intercept coefficient and $\boldsymbol{\epsilon}$ is the residual vector. The model $\mathbf{v}$ was scaled by its magnitude then subtracted from $\mathbf{r}$, forming the RSA response vector $\mathbf{y} = \mathbf{r} - \mathbf{v}b_1$. We additionally used an alternative, downsampling technique, to verify that our primary findings were robust to this issue (see Method section *Downsampling Analysis*).

Thus the RSA regression yielded three β coefficients of interest: β_{target}, $\beta_{\text{distr.}}$, $\beta_{\text{incon.}}$. These coefficients can be understood as a (standardized) contrast on (rank-transformed) correlations of activity patterns, between conditions in which only one task dimension was shared (e.g., the target dimension for β_{target}), versus those in which no dimensions were shared (i.e., different levels of target, distractor, and congruency).

2.3.7 Dimensionality Reduction

We used non-metric multidimensional scaling (Kruskal, 1964), a flexible, non-parametric dimensionality reduction technique, to visualize the structures of activity patterns within selected regions

(Figure 2): ventral somatomotor cortex (corresponding to 'mouth' homunculi), primary visual cortex (V1), and our (left) DMFC superparcel. These parcels were selected to highlight coding of each task dimension. For each selected region, we averaged observed correlation matrices across subjects, then subtracted these values from 1 to obtain a dissimilarity matrix. Before averaging, we z-transformed (inverse hyperbolic tangent, artanh) correlations, and inverted this transform after averaging. (In contrast to the RSA regression above, we did not rank-transform correlation matrices, as non-metric multidimensional scaling incorporates a monotonic regression). Similar to our RSA, we prewhitened each similarity matrix prior to conducting this procedure (see Step 4 in *Estimation of Coding Strength β*). Each mean dissimilarity matrix was submitted to an implementation of Kruskal's non-metric multidimensional scaling, *vegan::metaMDS()* in R, to generate a 2-dimensional configuration (Oksanen et al., 2019).

2.3.8 Group-Level Dissociation Analysis

To test for regional dissociations in coding preferences, we fit a hierarchical linear model on RSA model fits (see Estimation of Coding Strength β) obtained from our three ROIs within each hemisphere, and for our three RSA models. Fixed effects were estimated for the interaction of RSA model, ROI, and hemisphere. Random effects by subject were estimated for the interaction of RSA model and ROI, with a full covariance structure (9×9; Barr, Levy, Scheepers, & Tily, 2013). This model was fit with *lme4::lmer()* in R (Bates, Maechler, Bolker, & Walker, 2014).

Planned contrasts on the fixed effects were performed to test our hypotheses. P-values were estimated using an asymptotic z-test, as implemented by the *multcomp::glht()* function in R (Hothorn, Bretz, & Westfall, 2008). We performed three types of contrasts: (1) to compare coding strengths within-region (e.g., DMFC: incon.$-$target), (2) to compare between regions, within-model (e.g., target: DLPFC$-$DMFC), and (3) to test their interaction [e.g., (target$-$incon.)$\cdot$(DLPFC$-$DMFC)]. These contrasts were performed first by collapsing across hemisphere, then within each hemisphere separately. As we did not have any hypotheses regarding lateralization, p-values from hemisphere-specific contrasts were FDR corrected across hemispheres.

2.3.9 Selection of Behavioral Measures for Individual-Level Analyses

Audio recordings of verbal responses were transcribed and coded for errors offline by two researchers independently. Discrepancies in coding were resolved by a third. Errors were defined as any non-target color word spoken by a subject prior to utterance of the correct response (e.g., including distractor responses, but not disfluencies) or as a response omission. Trials in which responses were present but unintelligible (e.g., due to high scanner noise or poor enunciation) were coded as such.

We fit two hierarchical models on these data, one on errors and one on response times (RT). Several observations were excluded from these models. From the error model, only trials with responses coded as "unintelligible" were excluded (54). From the RT model, several types of trials were excluded: Trials with RTs greater than 3000 ms (1) or less than 250 ms (53; 52 of which were equal

to zero). A cluster of fast and unrealistically invariable RTs from two subjects (23/216, 26/216) that were likely due to an artifact of insufficient voice-onset signal within the recording. Trials with a residual RT that was more extreme than 3 inter-quartile ranges from an initial multi-level model fitted to all subjects data (of the structure of the model equation below; *a la* Baayen & Milin, 2010). All trials with incorrect (137), unintelligible (54), or no response (52). In total, 232 trials were excluded (0–62 per subject), leaving 10352 trials for analysis (154–216 per subject).

RTs for subject s were modeled (following Laird & Ware, 1982 notation):

$$\begin{aligned} \mathrm{RT}_s =& (\mathbf{1}, \mathbf{stroop}_s)\mathbf{b} + (\mathbf{1}, \mathbf{stroop}_s)\mathbf{u}_s + \boldsymbol{\epsilon}_s \\ \mathbf{u}_s \sim& N_2(\mathbf{0}, \mathbf{T}_{2\times 2}^2) \\ \boldsymbol{\epsilon}_s \sim& N(0, c_s\sigma^2) \end{aligned}$$

where RT_s is a column vector of the response times from subject s, $\mathbf{1}$ is an all-ones vector (intercepts), $\mathbf{stroop}_s$ is a vector indicating incongruent trials, and c_s is a subject-specific parameter by which their residual variance was scaled. Critically, $\mathbf{b}$ and $\mathbf{u}_s$ contained coefficients corresponding to the classical Stroop interference effect contrast (incongruent – congruent), for the group ($\mathbf{b}$) and subject ($\mathbf{u}_s$) levels. The scaling parameters c_s relaxed the assumption that each subject had a common residual variance; a well-warranted complexity, given the vastly improved fit of the heterogeneous-variance model ($\chi^2_{48} = 4299, p = 0, \Delta\mathrm{BIC} = -3856, \Delta\mathrm{AIC} = -4203$). To accomplish this, the RT model was fitted with *nlme::lme()* in R (Pinheiro, Bates, DebRoy, Sarkar, & R Core Team, 2019). The behavioral error model had similar predictors, but assumed binomially distributed error, with logit link function (fitted in *lme4*).

This hierarchical modeling framework enabled us to to estimate the amount and internal consistency of individual variability in the Stroop interference effect within both RTs and errors while accounting for trial-level error (e.g., Haines et al., 2020). We used these subject-level estimates to validate that our behavioral measures met prerequisite properties for individual differences analyses. To assess the amount of between-subject variability in the Stroop interference effect, a nested model comparison was conducted, in which the models fitted above were compared to a random-intercept model. Stroop effects differed significantly across subjects in RT ($\chi^2_2 = 87.53, p = 0.00; \Delta\mathrm{BIC} = -69.08; \Delta\mathrm{AIC} = -83.53$), but not in accuracy ($\chi^2_2 = 0.69, p = 0.71; \Delta\mathrm{BIC} = 18; \Delta\mathrm{AIC} = 3$). We therefore did not further analyze accuracy. To assess internal consistency (defined here as cross-run correlation), we fit a model with separate congruency factors (fixed and random) per run, and a full 4×4 covariance structure, which was used to obtain the cross-run correlation in Stroop effect. Individuals' Stroop interference effects in RT were estimated to be highly consistent across scanning runs ($r = 0.95$). We therefore considered our measurements of RT to be adequate for individual

389 differences analysis.

2.3.10 Individual-Level Dissociation Analysis

Similar to our *Group-Level Dissociation Analysis*, we tested our individual-level hypotheses within
a hierarchical modeling framework. Preliminary analyses suggested error measures were inadequate
for individual differences analyses (see *Materials and Methods, Selection of Behavioral Measures for*
Individual-Level Analyses), so we focused solely on RT measures.

We began with the RT model described in the preceding section. However now, for a given RSA
model and region of interest, we incorporated into the fixed effects each subject's estimated coding
strength, β_s, by interacting this coding-strength term with the congruency factor. This formed a
cross-level, continuous-by-categorical interaction, $\beta_s \cdot \mathbf{stroop}_s$. The coefficient on this interaction
term described how the Stroop interference effect (within-subject) varied across subjects as a func-
tion of their coding strength (between-subject). To test our hypotheses, we performed contrasts on
these interaction coefficients, which are outlined within the *Results* describing Figure 3.

2.3.11 Model Selection

To complement our hypothesis-driven brain–behavior analyses, we used a more data-driven model-
selection approach. An expanded set of 24 superparcels (in addition to our six ROIs) was defined
(Figure 3, B; see Figure 3-3 for list). Some superparcels were included as regions of interest,
others were included as negative controls (i.e., regions that were not predicted to be important
for explaining behavioral performance). Subject-level coefficients of the Stroop interference effect
contrast were extracted from the behavior-only RT model and used as the response vector (in the
model equation, the slope elements of $\mathbf{u}_s$; also known as BLUPs or conditional modes). RSA was
conducted on each superparcel (see *Estimation of Coding Strength β*), furnishing three β coefficients
per superparcel. These 72 measures were fitted to the response vector via elastic net regression,
implemented via *glmnet::glmnet()* in R (Friedman, Hastie, & Tibshirani, 2010). Parameter α was
set to 0.5 (balancing lasso and ridge penalties). The parameter λ was tuned via 10-fold cross-
validation (default) using the "1-SE rule": the smallest λ greater than one SE of the minimum
λ across folds was saved (Hastie, Tibshirani, & Friedman, 2009); this routine was repeated 1,000
times and the minimum mode (there were ties) of the saved λs was selected. We selected the
minimum mode because the maximum suppressed all variables from the model.

To assess validation-set accuracy, the selected model coefficients were applied to a design matrix
from validation-set subjects, generating a predicted Stroop effect vector. The linear correlation was
estimated between this predicted Stroop effect and observed Stroop effects (estimated as conditional
modes via a hierarchical model separately fitted to validation-set data). The significance of this
correlation was assessed by randomly permuting the training-set response vector, re-fitting the
model, generating new predicted validation-set values, and re-estimating the predicted–observed
correlation 10,000 times. The p-value was given by the proportion of resamples in which the null

correlation was greater than the observed correlation.

2.3.12 Exploratory Whole-Cortex RSA

The RSA-model fitting procedure, as outlined in *Estimation of Coding Strength β* was separately conducted on each MMP cortical parcel. Inferential statistics followed those suggested by Nili et al. (2014). One-sided signed-rank tests were conducted for significance testing (>0). P-values were FDR-corrected over all 360 parcels, separately for each task dimension (target, distractor, incongruency).

2.3.13 Univariate Activation Analyses

We additionally conducted a standard "univariate activation" analysis on these data. Note that this was not meant to evaluate whether univariate activity was a plausible confounding variable in our analysis, but rather to provide some basis for comparing our data to most extant neuroimaging studies of Stroop. For a given ROI (or MMP parcel), beta coefficients from the first-level fMRI GLM were averaged over voxels by stimuli, then over stimuli by congruency. These mean values were then contrasted, analogous to the behavioral Stroop interference effect (incongruent $-$ congruent). This statistic gives an estimate of the overall (across-voxel) difference in fMRI activity within a give brain region on incongruent versus congruent trials.

2.4 Follow-up Control Analyses

To establish the robustness of our results, we conducted several control analyses that examined a number of confounds and concerns: potential differences in signal-to-noise ratio across prefrontal ROIs, the effects of head motion, different RSA models for incongruency coding, the presence of bias imposed by the experimental design, within-run versus between-run RSA estimation, and the effects of downsampling to account for different trial numbers across runs. These are each described next.

2.4.1 Comparison of Signal-to-Noise Ratios

To test for differences in SNR (signal-to-noise ratios) between DMFC and DLPFC, we estimated "noise ceilings" within each region and contrasted them across regions. Noise ceilings indicate the maximum observable group-level effect size (RSA model fit) given the level of between-subject variability in similarity structure (Nili et al., 2014). Lower (smaller) average noise ceilings indicate poorer SNR for group-level tests. We used the cross-validated "lower-bound" noise ceiling estimator of Nili et al. (2014), as this yields a lower-variance estimate, and therefore more powerful contrast across regions, than the non-cross-validated "upper-bound" (Hastie, Tibshirani, & Friedman, 2009, Section 7.10.1). For a given region, the lower-bound noise ceiling is defined for each subject s in $1, \ldots, N$ as $\mathrm{Cor}(\mathbf{y}_s, \bar{\mathbf{y}}_{-s})$: the linear correlation between a subject s' observed similarity vector, $\mathbf{y}_s$, and the group-mean vector excluding subject s, $\bar{\mathbf{y}}_{-s}$. (See *Estimation of Coding Strength β* for definition of $\mathbf{y}$.) As each of the N estimates are interdependent, we used a percentile bootstrap to contrast noise ceilings across regions. In each resample (of 10,000), noise ceilings were artanh

transformed, contrasted across regions within-subject, then averaged across subjects. A two-sided p-value was provided by computing the proportion of resampled means greater than zero, p, then taking the minimum of $2p$ and $2(1-p)$. Finally, we conducted "two one-sided tests" for equivalence (Lakens, 2017; Schuirmann, 1987) to affirm a null hypothesis of no difference between regions in noise ceiling. This consists of defining a threshold effect size, the minimum effect size of interest, and testing whether the observed difference in noise ceilings is significantly less extreme than the threshold. For an objective threshold, we used the smallest standardized effect size at which our bootstrap procedure was expected to retain 80% power, Cohen's d = 0.35, determined via Monte Carlo simulation.

2.4.2 RSA on Head Motion Estimates

As a negative control analysis for our exploratory whole-cortex RSA, we attempted to decode task variables (target, distractor, incongruency) via RSA from framewise estimates of head motion. The 6 motion regressors that were used in the fMRI GLM as nuisance covariates (corresponding to translation and rotation in 3 dimensions) were regressed upon the design matrix containing predicted BOLD timecourses of our 16 conditions of interest. The coefficient matrix resulting from this regression was then submitted to the RSA procedures described in *Estimation of Coding Strength β* and *Exploratory Whole-Cortex RSA*. To check whether more aggressive movement denoising within the fMRI GLM was warranted (i.e., in addition to the 6 nuisance regressors), we conducted this same movement-based RSA, however using 12 motion regressors (the 6 bases and their temporal derivatives). RSA model fits between the 6 and 12-basis motion-based RSAs were compared via paired-sample signed-rank test.

2.4.3 Alternative RSA Incongruency Models

The RSA incongruency model parameterized the congruent–congruent correlations (i.e., R_{ij} where i and j are both patterns from congruent trials, denoted here simply as CC) with a separate nuisance regressor. That is, these cells were effectively excluded, and the model instead computed the contrast $II - IC$. This exclusion was done because we have no specific hypotheses regarding how congruent trials should be encoded relative to one another. Other parameterizations are possible, however, including models that (a) incorporate CC correlations within the "baseline" or intercept term, by omitting the congruent nuisance regressor [i.e., $II - \text{ave}(IC + CC)$], or (b) that omit IC correlations from the contrast (i.e., $II - CC$). We note, however, that (b) is a suboptimal parameterization as it effectively excludes 40% (48/120) of observations per subject (i.e., all IC cells). To verify that our results were not dependent on the particular modeling approach we chose, we compared the observed RSA model fits to these alternative parameterizations. In brief, subject's model fits were highly similar between the parameterizations (see **Results** sections *Group* and *Exploratory Whole-Cortex*; Figure 2-5). Thus, to help streamline the Results, we only conducted and reported analyses using the original RSA incongruency parameterization (in which the congruency model was a nuisance covariate).

2.4.4 Design Bias

For the current study, a typical within-run form of RSA estimation was implemented, in which correlations were computed among activation patterns estimated within the same scanning run and first-level GLM. Within-run RSA has been criticized because it is susceptible to design biases that occur when trial orders are insufficiently randomized within the experiment (Cai, Schuck, Pillow, & Niv, 2019). A priori, this was not a strong concern in the current design, as trial orders were fully randomized both within and between participants. Nevertheless, we conducted several diagnostic and robustness analyses to validate that our results and conclusions were not impacted by this potential bias. First, we estimated the extent of the potential bias by conducting our RSA technique on data simulated under a "worst-case" scenario, that is, when SNR = 0 (details described within Figure 1-2) and across a wide range of autocorrelation strengths. At worst, the three models were weakly biased (within 0.02 to 0.05 of $\alpha = 0.05$; Figure 1-2). This result indicated that while the bias was present, it was relatively minimal (cf., Cai, Schuck, Pillow, & Niv, 2019). Second, we validated that these simulated estimates were realistic, by using our actual fMRI data to estimate the false positive rate empirically. To do this, we conducted RSA on the first-level GLM coefficients from each subjects' ventricles, as these voxels should contain no brain activity signal but similar noise characteristics as those of interest (a ventricle mask of 2431 voxels was obtained from AFNI servers: https://afni.nimh.nih.gov/pub/dist/tgz/suma_MNI_N27.tgz). By treating the group-level mean and SD of these RSA model fits as the parameters of a non-central null distribution, $X \sim N(\bar{\beta}, SD(\beta))$, we computed the empirical false positive rate as $P(X > z_{\alpha=0.05})$, the proportion of this distribution greater than the customary one-tailed z-criterion. All rates were within 0.03 of $\alpha = 0.05$ (target = 0.07, distractor = 0.04, incongruency = 0.08), confirming the bias was quite minimal.

2.4.5 Between-Run RSA

As an alternative to within-run RSA, various between-run estimation approaches have been proposed which have been shown to be less sensitive to potential design biases. We opted not to use between-run RSA for our primary analyses, both because of the reduced effects of design bias established above, but also because between-run RSA is noted to be considerably more conservative that within-run RSA (Cai, Schuck, Pillow, & Niv, 2019). Moreover, several particulars of the present design are known to further hamper its power. Namely, between-run RSA makes incomplete use of the data, an issue which is exacerbated to the maximum extent possible in the present case, as our design has only the minimum number (two) of cross-validation folds (runs; Diedrichsen, Berlot, Mur, Schütt, & Kriegeskorte, 2020). Additionally, because the image acquisition sequence involved a reversal of phase-encoding direction across the two runs, this effectively effectively adds a strong non-linear component of noise if between-run RSA is used.

Nevertheless, to examine further the extent of design bias in our data, as well as the robustness of our results to the drop in power imposed by between-run RSA, we conducted a follow-up analysis of our

primary results using between-run RSA approaches. We conducted two forms of between-run RSA: the first used the cross-correlation of patterns between scanning runs ("cross-run RSA"; see Alink, Walther, Krugliak, Bosch, & Kriegeskorte, 2015), and the second used a cross-validated form of the Euclidean distance (i.e., on the inner product of pattern contrasts between runs; "cross-validated RSA"; see Walther et al., 2016). We selected these two forms of RSA as they have complementary benefits. Cross-run correlation is most comparable to our original within-run correlation, as they are both linear correlations. However, using this method within our dataset also necessitated employing downsampling (as the number of trials per condition were not perfectly balanced at the run-level; see *Downsampling Analysis*), which increases the variance of resulting estimates due to discarding data. In contrast, cross-validated RSA is insensitive (in terms of expected value) to the issue of trial numbers per condition (Diedrichsen, Berlot, Mur, Schütt, & Kriegeskorte, 2020). Using this method therefore allowed us to conduct the RSA using all the data at once, without downsampling. But, cross-validated RSA tests a more constrained hypothesis than cross-run RSA. Whereas cross-run RSA can be sensitive to non-linear differences between conditions, cross-validated RSA tests linear discriminability between conditions. Thus, when a non-linear boundary separates conditions, cross-validated RSA will fail to detect an effect whereas cross-run RSA could succeed. A non-linear boundary would occur, for example, when one condition (e.g., incongruent stimuli) drives a reliable, common, response while the other conditions (e.g., congruent stimuli) either drive unreliable, or stable but heterogeneous, responses. Nevertheless, to make cross-validated RSA as comparable as possible to our primary RSA, we z-score standardized patterns prior to computation and omitted spatial prewhitening (Walther et al., 2016).

2.4.6 Downsampling Analyses

Lastly, we checked whether the primary results were robust to the prewhitening method of data preprocessing, which was introduced to handle the imbalance of trials across runs (see *Method* section *Estimation of coding strength β*, step 4). In this analysis, we instead handled this issue by performing RSA after equating the number of trials per run*condition, by iteratively downsampling conditions with random subsets of trials. Specifically, we first fitted GLMs on fMRI timeseries, separately for each scanning run, that contained a single regressor per trial (LS-A method of Mumford, Turner, Ashby, & Poldrack, 2012). The minimum number of times we presented each unique Stroop stimulus in a single run was 3; this was the number to which we downsampled all conditions with >3 occurrences. For these conditions, we randomly sampled three trials, and averaged GLM coefficients voxel-wise over these trials. (For 3-trial conditions, we simply averaged all trials that were present.) This formed 16 separate condition-level coefficient vectors (activity patterns) per run. We then averaged these coefficient vectors across run, then estimated condition $\times$ condition correlation matrices from these patterns (averaging across runs was omitted from the downsampled cross-run RSA; see preceding section). We repeated this resampling, averaging, and correlation process 1,000 times, and averaged the resulting correlation matrices across iterations

(after an artanh transform). These correlation matrices were then submitted to the same RSA as outlined in *Estimation of coding strength β*, with the omission of the prewhitening step (4).

3 Results

Influential theories of cognitive control have proposed specific dissociations in the type of task information encoded by human medial and lateral frontoparietal cortex (Figure 1, A–B). But, previous studies have largely approached this question indirectly, by employing tasks designed to recruit these regions differentially in time, then testing for temporal dissociations in regional-mean levels of fMRI activity. Here, we used a more direct approach, by using the similarity structure of neural activity patterns evoked within these regions to estimate their informational content. In particular, through RSA, we compared neural coding of three distinct types of Stroop-task information — target, distractor, and incongruency (Figure 1, D–E) — within each region of interest (ROI) simultaneously, while Stroop interference was being experienced and resolved.

We describe three sets of analyses. First, we examined group-level effects, to test for neuroanatomical dissociations in encoding of task information (Figure 1, B, center). Second, we examined individual-level effects (i.e., individual differences), to test for dissociations in brain–behavior relationships (Figure 1, B, right). These two analyses were ROI-based and primarily focused on dorsomedial frontal and lateral frontoparietal regions (Figure 1, D), but also included sensorimotor ROIs for comparison purposes. The last set of analyses was conducted in whole-cortex exploratory fashion, to provide a more comprehensive picture of the anatomical profile of each task dimension.

3.1 Group

3.1.1 Dorsomedial frontal and dorsolateral prefrontal cortex exhibit distinct coding profiles

Primary group-level results are summarized in Figure 2. Statistical estimates corresponding to results outlined within this section are contained in Table 2.

The DMFC has been strongly associated with the coding of incongruency information in response conflict tasks such as the Stroop. RSA approaches were used to directly test the specificity of that hypothesis. This region was indeed found to encode the incongruency dimension of the Stroop task (left: $\bar{\beta} = 0.11, p = 0.00$, right: $\bar{\beta} = 0.08, p = 0.00$). Further, within left DMFC, a preference for this dimension was observed: incongruency information was encoded more strongly than either target or distractor information (vs. target: $\Delta\bar{\beta} = 0.07, p = 0.05$; vs. distractor: $\Delta\bar{\beta} = 0.10, p = 0.00$; p-values adjusted across hemisphere; Figure 2, A). This DMFC preference was prominent enough that it could be seen in its overall similarity structure within a low-dimensional embedding (Figure 2, B). Further, incongruency coding was neuroanatomically dissociated, as this coding scheme was reflected more strongly in left DMFC than in DLPFC activity patterns ($\Delta\bar{\beta} = 0.06, p = 0.02$; Table

2).

We next focused on lateral frontoparietal regions and the coding of target information. DLPFC indeed encoded target information (left: $\bar{\beta} = 0.03, p = 0.01$ right: $\bar{\beta} = 0.04, p = 0.00$), and more strongly than distractor information ($\Delta\bar{\beta} = 0.04, p = 0.00$). We did not find a preference, however, for target over incongruency information in DLPFC ($\Delta\bar{\beta} = -0.01, p = 0.62$), nor did we observe significantly stronger target coding than in DMFC ($\bar{\beta} = 0.00, p = 0.92$). A qualitatively similar pattern of results was observed in lateral posterior parietal cortex: significant target and incongruency coding, significantly enhanced target versus distractor coding, but no observed preference among target or incongruency nor detectable difference from DMFC target coding (Table 2).

Thus, at the group-level, we observed a single rather than double dissociation between medial and lateral frontoparietal cortex, in the form of enhanced sensitivity to incongruency relative to target coding in left DMFC. In all ROIs, however, these two sources of task information were more strongly encoded than distractor information.

3.1.2 Sensitivity and control analyses.

We next tested a series of hypotheses to scrutinize and extend our results.

First, we conducted a positive control analysis to bolster confidence in the statistical power of RSA methods within the present design. In particular, we sought to determine whether our methods could detect dissociations in task coding that are strongly expected to exist. For this, we focused on primary somatomotor and visual cortical ROIs, the responses of which can be assumed to reflect, relatively selectively, response-related (i.e., motoric) and visual form-related coding. As the distractor (word) defines the visual form of the Stroop stimulus, coding of form-related features should be captured by our distractor model. In parallel, as our analysis included only correct-response trials (i.e., in which the target response was spoken), coding of motoric features should be captured by our target model. Consistent with this logic, within early visual cortex, evidence of preferential distractor coding was observed (Figure 2, C; distractor: $\bar{\beta} = 0.04, p < 0.001$, target: $\bar{\beta} = 0.02, p = 0.17$, distractor vs. target: $\Delta\bar{\beta} = 0.03, p = 0.05$), whereas, within primary ventral somatomotor cortices (encompassing the "mouth" homunculi), a relatively selective pattern of target coding was found (Figure 2, C; distractor: $\bar{\beta} = 0.00, p = 0.9$, target: $\bar{\beta} = 0.13, p < 0.001$, distractor vs. target: $\Delta\bar{\beta} = -0.13, p < 0.001$). Further, coding of these dimensions was strong enough to predominate the overall structure of patterns within a low-dimensional embedding (Figure 2, B). Thus, in primary visual and somatomotor regions, a relatively clear-cut group-level double-dissociation emerged. This suggests that our models were adequately powered, at least in primary sensorimotor cortices, to detect functional distinctions, and that the failure to observe distractor coding in DMFC and DLPFC was not due to a general deficiency in our distractor coding model.

Figure 2: Group-level results. **A**, A single dissociation between dorsolateral versus dorsomedial frontal cortex (DLPFC, DMFC). For simplicity of display, DLPFC estimates are averaged across hemisphere (see Table 2 for per-hemisphere means for each region). Error bars show 95% confidence intervals of between-subject variability (estimated via bias-corrected and accelerated bootstrap). Horizontal grey lines at the top indicate significance of within-subject significance tests (see Table 2 for contrast estimates). While all ROIs encoded target and incongruency information, and did so more strongly than distractor information (coding of which was not detected), incongruency coding was stronger in left DMFC versus DLPFC. These results were generally robust to different analysis decisions and implementations (Figures 2-1, 2-2, 2-4, 2-5). Further, a univariate analysis failed to detect significantly higher activation on incongruent versus congruent trials in these regions (Figure 2-3). **B**, Across-voxel activity patterns from three select regions, embedded within a two-dimensional space (via non-metric multidimensional scaling). These geometries exemplify regional dissocations in coding of Stroop-task dimensions. Letters represent the first letter of the corresponding distractor word (BLUE, PURPLE, RED, WHITE), and hues represent the target color (stimuli that were white are presented here in grey). Connecting lines are drawn to highlight the various structures present. The geometry of (left) dorsomedial frontal cortex (DMFC) is marked by a radial separation of patterns evoked by congruent and incongruent conditions (lines connect congruent and incongruent stimuli of each target color), such that incongruent patterns tend to be located more centrally (i.e., more similar to each other), whereas congruent patterns peripherally diverge (i.e., less similar to each other). This suggests that incongruent trials drove a common component of activation in this region, regardless of target or distractor features. In contrast, within primary ventral somatomotor cortices (vS1/vM1), patterns strongly cluster by target level (color), whereas in primary visual cortex (V1), they tend to cluster by distractor level (word). This pattern formed a double dissociation (**C**), and indicated our distractor model was adequately powered to detect distractor coding in sensory regions (*cf.*, **A**). **C**, Plotting conventions follow those in **A**; horizontal dotted lines indicate $0.1 > p > 0.05$ (see text for estimates).

contrast	$\bar{\beta}$	σ	t	p
DMFC (L) incon.	0.11	0.03	3.65	0.00026
DMFC (R) incon.	0.08	0.03	2.82	0.00485
DMFC (L) target	0.04	0.01	2.61	0.00918
DMFC (R) target	0.05	0.01	3.94	0.00008
DMFC (L) distr.	0.01	0.01	0.78	0.43667
DMFC (R) distr.	0.00	0.01	0.19	0.84907
DLPFC (L) incon.	0.04	0.02	2.04	0.04155
DLPFC (R) incon.	0.05	0.02	2.39	0.01682
DLPFC (L) target	0.03	0.01	2.56	0.01062
DLPFC (R) target	0.04	0.01	3.18	0.00147
DLPFC (L) distr.	-0.01	0.01	-0.98	0.32561
DLPFC (R) distr.	0.00	0.01	0.31	0.75797
LPPC (L) incon.	0.07	0.02	3.07	0.00211
LPPC (R) incon.	0.06	0.02	2.64	0.00818
LPPC (L) target	0.03	0.01	2.74	0.00617
LPPC (R) target	0.05	0.01	4.38	0.00001
LPPC (L) distr.	0.00	0.01	0.28	0.77722
LPPC (R) distr.	0.02	0.01	1.86	0.06359
incon. − target	DMFC (L)	0.07	0.03	2.26	0.04737
incon. − target	DMFC (R)	0.03	0.03	0.91	0.36047
incon. − distr.	DMFC (L)	0.10	0.03	3.22	0.00256
incon. − distr.	DMFC (R)	0.08	0.03	2.64	0.00826
target − distr.	DMFC (L)	0.03	0.02	1.63	0.10230
target − distr.	DMFC (R)	0.05	0.02	3.18	0.00297
incon. − target	DLPFC (L)	0.01	0.02	0.46	0.65809
incon. − target	DLPFC (R)	0.01	0.02	0.44	0.65809
incon. − distr.	DLPFC (L)	0.06	0.02	2.41	0.03212
incon. − distr.	DLPFC (R)	0.05	0.02	2.09	0.03671
target − distr.	DLPFC (L)	0.04	0.02	2.76	0.01145
target − distr.	DLPFC (R)	0.04	0.02	2.33	0.01989
incon. − target	LPPC (L)	0.04	0.03	1.62	0.21226
incon. − target	LPPC (R)	0.01	0.03	0.43	0.66707
incon. − distr.	LPPC (L)	0.07	0.03	2.75	0.01202
incon. − distr.	LPPC (R)	0.04	0.03	1.63	0.10282
target − distr.	LPPC (L)	0.03	0.02	1.87	0.06185
target − distr.	LPPC (R)	0.03	0.02	1.96	0.06185
DMFC (L) − DLPFC	incon.	0.06	0.02	2.38	0.01737
DMFC (L) − DLPFC	target	0.00	0.01	-0.09	0.92467
DMFC (L) − LPPC	incon.	0.04	0.03	1.47	0.14112
DMFC (L) − LPPC	target	-0.01	0.01	-0.56	0.57868
[incon. − target] · [DMFC (L) − DLPFC]	0.06	0.03	2.20	0.02797
[incon. − target] · [LPPC (L) − DLPFC]	0.03	0.03	1.06	0.29066

Table 2 *(previous page)*: Group-level results from RSA in frontoparietal regions of interest. Group-mean RSA model fits and contrasts between RSA models (target, distractor, incongruency) and between ROIs (DMFC, DLPFC, LPPC in each hemisphere). Each row displays the statistical estimates from a given RSA model and ROI, or from a contrast between RSA models or ROIs. Contrasts were conducted either between RSA models (within ROI) or between ROIs (within RSA model). For example, target − distr. | DLPFC (R) indicates the contrast between target and distractor RSA model fits ('coding strengths') within right DLPFC. $\bar{\beta}$ indicates the mean RSA model fit over subjects. Unless noted, contrasts were averaged across hemisphere. For contrasts performed separately in each hemisphere, p-values were FDR-adjusted for two comparisons. Products of bracketed terms indicate interactions between RSA model and ROI. Dark grey rows display contrasts with $p < 0.05$, light grey rows display contrasts with $0.05 < p < 0.1$ Incon., incongruency; distr., distractor.

Second, we conducted sensitivity analyses to assess the robustness of our results to the particular ROI definitions used. In one analysis, we tested more expansive ROI definitions, by using alternatively-defined superparcels (Figure 1-1). These definitions included additional, more rostral PFC parcels (1 in DMFC, 3 in DLPFC), which begin to encroach into ventromedial PFC and frontopolar cortex (e.g., the rostral DMFC parcel was assigned to the Default-Mode network within the Cole-Anticevic divisions). Nevertheless, the previously observed dissociations were robust to these more liberal definitions (Figure 2-1). In the other analysis, we examined whether the overall superparcel coding profiles were representative of individual parcels. While DMFC and DLPFC results generally reflected that of constituent parcels (Figure 2-2, A, B), interestingly, there was substantial heterogeneity within left LPPC (Figure 2-2, C). Similar to left DMFC, a collection of left LPPC regions spanning the length of intraparietal sulcus strongly encoded the incongruency dimension (i.e., IP1, IP2, IPS1, AIP, LIP, MIP).

Third, we examined whether the lack of observed discrimination between target and incongruency coding dimensions within DLPFC could be explained by increased error variance potentially present in fMRI activity patterns within this region. Prior work has suggested that PFC regions might be particularly susceptible to this confound (Bhandari, Gagne, & Badre, 2018). It is possible that we might have observed a dissociation in left DMFC but not DLPFC due to differential levels of statistical power across the two regions. We therefore derived an SNR (signal-to-noise ratio) analysis to determine whether this was a viable explanation (see Method). A paired-sample bootstrap test did not indicate a systematic difference between DLPFC versus left DMFC group-level SNR ($\Delta z = -0.01, p = 0.60$). However, the SNRs in these regions were also not confirmably similar, as indicated by an equivalence test (which provides a confirmatory test of the null hypothesis; $p = 0.10$). Therefore, while we cannot rule out the possibility that the single dissociation observed between left DMFC and DLPFC was driven by better group-level SNR in DMFC, any potential SNR differences between the two regions were not substantial enough for our methods to detect.

Fourth, to provide a basis for comparison to most extant neuroimaging research of the Stroop task, we conducted a "univariate activation" analysis, examining whether these brain regions were generally more active during incongruent versus congruent conditions. No regions were found to respond more strongly overall to incongruent versus congruent conditions (Figure 2-3, A), although the mean contrast in DMFC (L) was positive (i.e., incongruent>congruent). This null result was not surprising, however, due to the high frequency of incongruent trials within the experiment — which is known to reduce both the behavioral and neural univariate Stroop effect (Carter et al., 2000; De Pisapia & Braver, 2006; Logan & Zbrodoff, 1979). While this null result demonstrates the utility of using RSA in this case, it should not, however, be seen as direct evidence for the increased sensitivity of RSA versus univariate methods, as the univariate and RSA-based tests as implemented here are subject to different constraints and are thus incomparable (Allefeld, Görgen, & Haynes, 2016). Finally, the magnitude of the univariate Stroop effects were only weakly correlated incongruency coding model fits (Figure 2-3, B), suggesting these measures were non-redundant.

Finally, we tested whether our pattern of results were robust to alternative RSA techniques, including a downsampling technique to equate trial counts across runs, two "between-run" RSA methods (cross-run RSA and cross-validated RSA; see Method), and alternative parameterizations of the incongruency coding model (see Method). Findings were robust to downsampling and to between-run RSA (Figure 2-4), and were highly similar across different parameterizations (Figure 2-5). Interestingly, however, the detection of incongruency coding in DMFC depended on whether a linear or non-linear RSA method was used. When using an RSA method that tests linear discriminability between conditions (cross-validated RSA), the incongruency coding effect was abolished in DMFC; whereas when using a comparable method that is sensitive to non-linear pattern differences (cross-run RSA), the effect remained quite strong. This pattern of results suggests that incongruency information was encoded non-linearly within DMFC activation patterns. Indeed, this can be seen within the 2-dimensional embedding (Figure 2), as a radial, rather than linear, separation of incongruent (central) and congruent (peripheral) stimuli.

3.2 Individual

Primary individual-level results are summarized in Figure 3. Statistical estimates corresponding to results outlined within this section are contained in Table 3 (see Figure 3-1 for scatter plots of all associations).

3.2.1 Better-performing subjects have stronger lateral frontoparietal target coding.

The fidelity of target-related information in lateral frontoparietal cortex — DLPFC, in particular — is thought to be closely linked to the efficiency with which an individual resolves response conflict in tasks such as Stroop. By using subject-level target-coding estimates (β_{target}) to model behavioral performance (RT), we tested this fundamental prediction relatively directly. Indeed, subjects with stronger target coding in both right DLPFC and right LPPC resolved Stroop interference effects

more quickly (DLPFC: $b = -239.13, p = 0.00, r = -0.44\ [-0.66, -0.15], \rho = -0.39\ [-0.63, -0.08]$; LPPC: $b = -235.34, p = 0.00, r = -0.44\ [-0.63, -0.17], \rho = -0.35\ [-0.59, -0.06]$; p-values corrected across hemisphere; linear correlation, r; rank correlation, ρ; bootstrapped 95% confidence interval, $[lower, upper]$; Figure 3, A; Table 3). Subjects' target coding estimates were moderately correlated ($r = 0.53$) between these two lateral frontoparietal regions, as expected based on their strong neuroanatomical and functional connectivity (e.g., Buschman & Miller, 2007; Petrides & Pandya, 1999). In neither of these lateral regions was incongruency coding significantly related to the Stroop interference effect (DLPFC: $b = 38.42, p = 0.25, r = -0.17\ [-0.14, 0.45], \rho = -0.10\ [-0.19, 0.37]$; LPPC $b = -32.40, p = 0.31, r = -0.15\ [-0.40, 0.13], \rho = -0.18\ [-0.45, 0.12]$; p-values uncorrected), and within right DLPFC, target coding was a considerably stronger explanatory variable than incongruency coding ($\Delta b = -299.21, p = 0.00$; Table 3).

Conversely, incongruency-related responses of DMFC are thought to be positively associated with maladaptive policies of response selection. In line with this notion, subjects with stronger incongruency coding in left DMFC tended to exhibit greater Stroop interference, although this interaction was non-significant ($b = 47.66, p = 0.13, r = -0.26\ [-0.01, 0.49], \rho = -0.17\ [-0.12, 0.44]$; p-value corrected across hemisphere; Figure 3, A). Similarly, incongruency coding was numerically more strongly associated with Stroop interference effects than target coding in this region, though this statistical difference was also non-significant ($\Delta b = -20.92, p = 0.83$; Table 3). While weak, this DMFC–Stroop association notably emerged within the same hemisphere that displayed a group-level preference for incongruency information (Figure 2, B). Importantly, the sign of the correlations we detected within our ROIs were robust to alternative RSA techniques (Figure 3-2; note that general attentuation in effect size is expected with higher variance techniques; see *Downsampling Analysis* and *Between-Run RSA* in Method).

Considering the collective pattern of results, we conclude here that our findings support the hypothesis that target coding in (right) DLPFC and in LPPC reflected a common process of implementing control.

Figure 3 *(previous page)*: Individual-level results. **A**, The coding strength of task variables (β) within right dorsolateral prefrontal cortex (DLPFC) and left dorsomedial frontal cortex (DMFC) accounted for individual differences in the size of the Stroop interference effect (response time, RT) in hypothesized ways. Inset text displays linear correlation coefficients (r) and 95% confidence intervals (estimated via bias-corrected and accelerated bootstrap). Lines indicate a regression line of RT onto Model Fit (β), and confidence bands indicate 95% confidence intervals of predicted values from this regression obtained via percentile bootstrap. The direction of these observed brain–behavior relationships were robust to alternative, more conservative RSA estimation techniques (cross-run RSA and cross-validated RSA; Figure 3-2). For display of relationships between target and incongruency coding in all ROIs (DLPFC, DMFC, LPPC) and Stroop effect, see Figure 3-1. For contrasts on slopes (interaction tests between ROI and RSA model), see Table 3. **B**, A larger set of 24 cortical regions were defined for use in a data-driven model selection analysis. Surface maps display in various colors all left-hemisphere regions (excluding left lateral occipito-temporal cortex); see Figure 2-1 for full list and definitions. Three RSA models (target, distractor, incongruency) were fitted per region, yielding a set of 72 coding-strength estimates. These 72 estimates were used to predict Stroop interference effects within a cross-validated model selection procedure. **C**, Additional brain–behavior associations identified via cross-validated model selection. Left ventral visual incongruency coding (left panel) and left DLPFC distractor coding (right panel) were selected — in addition to right DLPFC target, right LPPC target, and left DMFC incongruency. See caption **A** for description of inset text, lines, and bands. **D**, Estimate of prediction accuracy of the selected model. Within a validation (held-out) set of 17 subjects (monozygotic co-twins of subjects within the analysis sample), observed Stroop effects were moderately correlated (red line) to predictions from the selected model. A permutation test (grey distribution) indicated the selected model significantly explained validation-set variance, suggesting the selected model results are relatively stable.

3.2.2 Model selection affirms a lateral–medial dissociation and identifies unexpected relationships.

Because the preceding hypothesis-driven analysis exclusively focused on a limited set of regions, important brain–behavior relationships may have been missed. A more accurate model may even omit target and incongruency coding from DLPFC, DMFC and LPPC altogether. Consequently, we conducted a more comprehensive test to identify regions and task dimensions that could better account for Stroop performance variability across individuals.

A data-driven model selection analysis was conducted to address this question (see Method). We defined an expanded set of 24 cortical regions (superparcels), including the six defined and used earlier, which covered various areas that may be important for performing the Stroop task (Figure 3, B; Figure 3-3). Conducting RSA on each superparcel furnished three coding estimates (one per coding model) per superparcel. These 72 estimates were then used as features in a cross-validated model selection procedure.

Strikingly, the selected model contained all three hypothesized measures: (right) DLPFC and LPPC

Table 3: Parameter estimates from hierarchical brain–behavior models that explained individual differences in behavioral performance (RT) with variability in the strength of neural coding (β, see Method). Separate models were fit per region and RSA model (target, incongruency) combination. The displayed estimates correspond to the interaction **stroop**$\cdot\beta$, which indicates how subjects' coding strengths relate to the magnitude of their Stroop interference effects. Also displayed are contrasts on **stroop** $\cdot$ β estimates, between RSA models (within ROI) or between ROIs (within RSA model). For example, target $-$ incon. | DLPFC (R) indicates, for right DLPFC, whether target and incongruency coding were differentially related to the size of the Stroop interference effect. Dark grey rows display contrasts with $p < 0.05$, light grey rows display contrasts with $0.05 < p < 0.1$

contrast	b	σ	t	p
DMFC (L) incon.	47.66	25.97	1.84	0.06647
DMFC (R) incon.	-0.10	33.15	0.00	0.99762
DMFC (L) target	48.40	90.63	0.53	0.59330
DMFC (R) target	-24.04	63.75	-0.38	0.70613
DLPFC (L) incon.	14.08	41.17	0.34	0.73227
DLPFC (R) incon.	38.42	33.66	1.14	0.25367
DLPFC (L) target	-105.93	78.34	-1.35	0.17637
DLPFC (R) target	-239.13	71.68	-3.34	0.00085
LPPC (L) incon.	-48.95	38.13	-1.28	0.19921
LPPC (R) incon.	-32.40	32.22	-1.01	0.31467
LPPC (L) target	-1.90	93.65	-0.02	0.98385
LPPC (R) target	-235.34	71.23	-3.30	0.00096
incon. $-$ target	DMFC (L)	20.92	97.23	0.22	0.82962
incon. $-$ target	DLPFC (R)	299.21	80.62	3.71	0.00021
incon. $-$ target	LPPC (R)	204.56	79.94	2.56	0.01050
DMFC (L) $-$ DLPFC (R)	incon.	22.85	52.92	0.43	0.66587
DMFC (L) $-$ DLPFC (R)	target	500.63	139.14	3.60	0.00032
DMFC (L) $-$ LPPC (R)	incon.	134.19	49.85	2.69	0.00710
DMFC (L) $-$ LPPC (R)	target	468.99	137.46	3.41	0.00065

target coding, and (left) DMFC incongruency coding. In addition, two unexpected measures were identified, both with negative slopes: left DLPFC distractor coding ($r = -0.43$ $[-0.66, -0.14]$, $\rho = -0.33$ $[-0.60, -0.04]$) and left ventral visual incongruency coding ($r = -0.35$ $[-0.52, -0.13]\rho = -0.43$ $[-0.62, -0.20]$; bivariate correlations shown in Figure 3, C). We tested the predictive accuracy of the selected model by using a held-out validation set of 17 subjects (co-twins of the primary analysis set). The selected model explained significant variance in Stroop interference effects within the validation set (Figure 3, D). While not fully independent, these validation set data were from a true hold-out and obtained from distinct individuals. This result therefore bolsters claims regarding the stability of the selected model.

Nevertheless, to provide a cursory test of a truly independent validation set (i.e., with no familial dependency to the training set), we excluded all subjects in the training set who were co-twins of with those in the validation set, and reconducted this model selection procedure. This amounted to discarding 16/49 (33%) of training-set observations. The selected model contained only one variable, which was not in our ROIs (but in early visual cortex), and was unable to predict held-out Stroop effects ($r = -0.12$). This is perhaps unsurprising, however, given the substantial reduction in the size of the training dataset for an already high-dimensional model. To reduce the dimensionality, we reconducted this analysis, focusing now instead only on ROIs and coding schemes of interest — target coding in DLPFC and LPPC, and incongruency coding in DMFC (within each hemisphere) — and additionally ensured that all variables were used in prediction (via ridge regression). This model was better able to predict the held-out Stroop effect ($r = 0.20$), in particular, relative to a comparable model that contained theoretically "mismatched" ROI×coding-scheme combinations (incongruency coding in DLPFC and LPPC, target coding in DMFC; $r = -0.17$, bootstrapped $p = 0.088$).

Results from these model selection analyses affirm a functional dissociation across the medial–lateral axis of frontoparietal cortex, and further demonstrate that Stroop-task representations within DLPFC and DMFC hold relatively privileged relationships with behavior.

3.3   Exploratory Whole-Cortex RSA

In a final, exploratory analysis, we estimated RSA models separately for each MMP parcel, to determine more comprehensively how target, distractor, and incongruency coding are distributed across cortex. These three task dimensions were encoded across cortex according to different neuroanatomical profiles. Target coding was widespread (observed in 207/360 parcels), covering substantial portions of the frontal and temporal lobes, including many perisylvian regions (Figure 4, top; Figure 4-1). Notably, the strongest target coding was observed within regions that receive strong sensory and (or) motor-related input (Figure 4-1). Contrastingly, incongruency coding was detected predominantly within prefrontal and intraparietal sulcal parcels — including DMFC, but also left LPPC, bilateral superior frontal gyrus, and left lateral frontopolar cortex (rostral DLPFC)

Figure 4: Cortical distributions of target (top), incongruency (middle) and distractor (bottom) coding, identified via an exploratory, whole-cortex analysis. Color bar indicates test statistic from one-sided sign-ranked test. For all statistical estimates, see Figures 4-1–4-3. Sensitivity and robustness analyses suggested that the core results were robust to various analysis decisions and implementations (handling of head motion, see text in Results; parameterization of incongruency coding, Figure 2-5, B; RSA estimation method, Figure 4-4).

— but additionally within left retrosplenial and right lateral occipital cortex (Figure 4, middle; Figure 4-3). Aside from this occipital area, these incongruency-coding parcels notably belonged to control networks (frontoparietal, cinguloopercular, and dorsal attention; Figure 4-3). In a third, distinct profile, distractor coding was only observed within early visual cortex (left V1 and V2; Figure 4, bottom; Figure 4-2).

As a negative control analysis, we tested whether we could decode these three task variables (target, distractor, incongruency) from framewise head motion estimates, using the same RSA procedures as above. No task variable was significantly encoded within patterns of head movements ($bs < 0.01$, $ps > 0.36$). Using a larger basis set of motion estimates (12 vs 6) did not yield significant decoding ($bs < 0.01$, $ps > 0.17$), nor any increased sensitivity to task variables ($bs < 0.00$, $ps > 0.17$), suggesting that our motion removal procedures (scrubbing and 6 motion regressors) were adequate.

Finally, we repeated this exploratory analysis using alternative RSA methods involving downsampling and between-run RSA (*Downsampling Analysis*, and *Between-Run RSA* sections of Methods; Figure 4-4). Across these analyses, the core results were quite robust: we found incongruency coding in parcels within DMFC (though this depended on the use of non-linear RSA methods, as in Figure 2-4 and Figure 3-2), target coding in mid-DLPFC, and distractor coding in visual cortex. Our findings were therefore not specific to a particular estimation method.

Collectively, these exploratory results confirm and extend our prior findings. (1) As with the

reported brain–behavior associations, target coding was emphasized relative to coding of other task dimensions. (2) Yet despite this emphasis, important and expected dissociations in anatomical profiles were identified across our three coding models, further suggesting that these models were successful in measuring coding of distinct task dimensions (Figure 1, A).

4   Discussion

We analyzed the similarity structure, or representational geometry (Kriegeskorte & Kievit, 2013), of frontoparietal activity patterns associated with cognitive control, during performance of the classic color-word Stroop task. In left DMFC, incongruency coding predominated. While DLPFC and LPPC encoded both target and incongruency-related information, distractor coding was not detected in these regions, but was instead identified in early visual cortex. Further, these neural coding estimates were important and specific indicators of individual differences in magnitude of the behavioral Stroop interference effect. Individuals with stronger target coding in right DLPFC and right LPPC, but weaker incongruency coding in left DMFC, exhibited enhanced cognitive control, in terms of a reduced Stroop effect. Further, in a more comprehensive predictive model that included coding measures from a wide set of cortical regions, coding measures specifically from lateral frontoparietal and dorsomedial frontal regions were privileged in their link to behavior.

On one level, this study is a specific extension of research that has drawn dissociations between control-related functions of DLPFC and DMFC (Floden, Vallesi, & Stuss, 2010; MacDonald, Cohen, Stenger, & Carter, 2000). Most prominently, MacDonald et al. (2000) used a modified Stroop-task design, in which task rules (delivered via pre-cues) randomly alternated between color naming and word reading across trials, to demonstrate that DLPFC and DMFC encoded different types of information during cognitive control engagement. In particular, DLPFC was selectively recruited following cues for the more demanding color-naming task, whereas DMFC was instead driven by incongruent color-naming trials. This pattern of recruitment suggested that DLFPC encodes task-set and rule-related information in a preparatory manner, whereas DMFC encodes incongruency and conflict-related information in a stimulus-evoked manner. In the current study, we leveraged the high spatial dimensionality of fMRI to test whether this functional dissociation can be observed *within* a common time window of response selection, and further with a more traditional Stroop-task design, which does not involve task cues or switches. Our findings reinforce the conclusions of these relatively low-powered studies (N = 12 in Flooden et al., 2010; N = 9 in MacDonald et al., 2000) and indicate that the dissociations were not dependent upon the use of cued task-switching designs. Synthesizing these prior findings with those of the present study hints at a continuity in the putative role of DLPFC during target selection. Rather than exclusively contributing to preparation, DLPFC coding may evolve from proactively representing abstract rule or set-related information, towards more concrete targets and behavioral choices as relevant stimulus information becomes available in the environment. This view accords with work in monkey electrophysiology

(Mante, Sussillo, Shenoy, & Newsome, 2013; Rigotti et al., 2013; Stokes et al., 2013), yet further work is needed to determine whether similar dynamics occur within human DLPFC and how such dynamics may reflect or interact with specific processes of cognitive control in Stroop-like tasks.

More broadly, the results of this study highlight the utility of RSA and the general representational geometric framework for investigating cognitive control. Previous work has used MVPA decoding in the Stroop task to study the impact of control demand on posterior representations (e.g., Banich et al., 2019). Here, we used the RSA framework to explicitly model and decompose control-related frontoparietal representations. In fact, a major motivation of the current study was to assess how well RSA measures of frontoparietal coding map to theorized mechanisms of control. For this purpose, the medial–lateral functional dissociation in frontoparietal cortex was a useful test-bed, as it features in several theoretical accounts (Botvinick, Braver, Barch, Carter, & Cohen, 2001; Miller & Cohen, 2001; Ridderinkhof, Ullsperger, Crone, & Nieuwenhuis, 2004; Shenhav, Botvinick, & Cohen, 2013). Our results were generally in line with these accounts, joining with a growing body of research in suggesting that RSA provides a convenient yet powerful framework from which neural measures can be used to test cognitive control theory (see Freund, Etzel, & Braver, 2021, for a review).

Nevertheless, the current work represents only an initial step in using the RSA framework to investigate cognitive control within Stroop-like tasks. As such, our study raises a number of unaddressed questions. But promisingly, there are ample opportunities for improving and extending the RSA framework highlighted here. For instance, a key limitation of the current study was the finding of widespread coding of the target dimension, suggesting a lack of specificity in the target RSA model. This is perhaps not surprising, however, as the model would capture not only coding of attentional-template and choice-related information, but also hue and response-related information. We mitigated this issue by demonstrating that target coding was selectively related to behavior within DLPFC and LPPC. Yet, this limitation could be addressed more powerfully by experimental design. Adding specific factorial manipulations, such as a task rule manipulation (*a la* MacDonald, et al. 2000; Hall-McMaster, Muhle-Karbe, Myers, & Stokes, 2019), or a response modality manipulation (*a la* Minxha, Adolphs, Fusi, Mamelak, & Rutishauser, 2020; see also Barch et al., 2001), would enable a richer, more precise set of cognitive control-relevant coding variables to be estimated (Figure 5).

Future work could also address some of the complexities revealed by our data that were not entirely accounted for by the theoretical frameworks we used. For one, although predicted coding profiles emerged in some frontoparietal regions of interest, all regions encoded incongruency and target information. This incongruency-coding finding is consistent with prior univariate fMRI research (e.g., Nee, Wager, & Jonides, 2007; Niendam et al., 2012), and a more recent finding that the response of single neurons in human dACC and DLPFC are robustly modulated by conflict (Smith

et al., 2019). With respect to target coding, however, one speculative interpretation is that, during the relatively late phase of response selection and execution, control networks may lose modular structure as the circuitry collectively converges on a behavioral choice. This interpretation accords with the fact that "choice axes" are encoded within multiple key nodes of frontoparietal decision circuitry (e.g., in macaque LIP: Roitman & Shadlen, 2002; in macaque caudal DLPFC: Mante, Sussillo, Shenoy, & Newsome, 2013; in human DACC and pre-SMA: Minxha, Adolphs, Fusi, Mamelak, & Rutishauser, 2020; see also Okazawa, Hatch, Mancoo, Machens, & Kiani, 2021). This account could be addressed using the enriched experimental design described above, identifying when and where choice coding is emphasized over the course of a trial.

Another unexpected finding was a relatively robust negative relationship between the strength of incongruency coding in left ventral visual cortex and the magnitude of the Stroop effect (Figure 3, B, left panel). One interpretation of this finding is provided by the biased competition framework, as an effect of selective visual attention. Prior work has demonstrated that certain ventral visual regions, those which are strongly tuned to target features, activate as a function of Stroop incongruency (Egner & Hirsch, 2005). In our task, mid-ventral stream areas may have received biasing input, selectively on incongruent trials, which enhanced stimulus-related coding and communication with downstream regions. Using the expanded RSA design sketched above, we might expect that such an effect would be limited to color naming conditions, when selective attention processes would be most prominent.

Perhaps the most surprising result was the robust negative correlation observed between left DLPFC distractor coding and the behavioral Stroop effect (Figure 3, B, right panel). At face, accounting for this finding within the framework of top-down biased competition is difficult. But, given the statistics of our task, in which incongruent trials were frequent and congruent were rare, distractor information could have been used to facilitate performance. An association between distractor features and incongruency could have been learned and used to influence response selection — for example, by retrieving and implementing a stimulus-appropriate attentional setting (Bugg & Crump, 2012; Melara & Algom, 2003). In fact, subjects clearly do exploit these associations, as indicated by reduced Stroop effects for stimuli that are "mostly incongruent," also known as "item-specific proportion congruency" effects (ISPC; Bugg, Jacoby, & Chanani, 2011); see also Crump & Milliken, 2009). The prediction that distractor coding might reflect ISPC effects could be tested by varying ISPC levels across different stimuli. For stimuli in which the specific color or word is not predictive of congruency, the relationship between DLPFC distractor coding and improved Stroop performance should not be present.

Finally, the present study sets the stage for using RSA to test the dual-mechanisms framework of cognitive control (Braver, 2012). This framework explains much within and between-individual variability in cognitive control function by the existence of two operational "modes" of cognitive

Figure 5: Expanded experimental designs. The current target coding model (top, green text; see also Figure 1) conflates distinct coding schemes, including those associated with relatively 'early' vision, relatively 'late' motoric commands, or more "central" rule-dependent schemes. By adding a factorial manipulation of task rule (i.e., participants respond to the present set of stimuli, but under both color-naming and word-reading conditions; MacDonald, et al. 2000; light purple arrows and text), the target coding model can be expanded into three more precise models (middle row of matrices), which are identical to our "target coding" model within the top-left quadrant (green square; naming–naming), but are now distinguished among the other quadrants (reading–reading, reading–naming). "Stimulus hue" and "response" models would identify coding that depends upon features of either the stimulus or response, independent of the task rules. Conversely, "target features" would identify a more flexible "attentional template" coding scheme, which depends critically on the current task goals (note that, the lower-right quadrant resembles our "distractor" coding model in Figure 1, as word features become the target in the word-reading condition). Importantly, too, abstract coding of task rule would also now be distinguishable ("rule" matrix on left). For an elegant example of a similar design within cued-task switching, see Hall-McMaster, Muhle-Karbe, Myers, and Stokes (2019). The "response" model could be further elaborated by incorporating another manipulation of response modality (verbal, manual; orange arrows). Now, coding of abstract choice options, independent of effector, could be separated from effector-dependent motoric coding. For an elegant example of a similar manipulation, see Minxha, Adolphs, Fusi, Mamelak, and Rutishauser (2020).

control: proactive and reactive. These modes are proposed to have dissociable signature neural coding schemes. Proactive control should rely heavily on goal-relevant coding schemes that originate in LPFC prior to target-stimulus onset as abstract rule or context coding, but which may morph into target coding post-stimulus onset. In contrast, reactive control should rely on an incongruency-based coding-scheme (including coding of whichever task dimensions are predictive of incongruency), originating post-target onset, with potential loci in DMFC or subcortical structures (Chiu, Jiang, & Egner, 2017; Jiang, Brashier, & Egner, 2015). As suggested here, it may be possible to measure correlates of these neural coding schemes via RSA. Experimental factors that encourage subjects to adopt one mode over another (e.g., strategy training, expectancy of difficulty) should correspondingly shift frontoparietal coding schemes along these proactive and reactive dimensions. Further, their behavioral relevance should predictably change, as well: for example, in task contexts in which a proactive control mode is theoretically maladaptive, subjects with stronger proactive coding should perform worse. Thus, the dual-mechanisms framework suggests a broad range of hypotheses amenable to testing with RSA methods (see, e.g., Hall-McMaster, Muhle-Karbe, Myers, & Stokes, 2019). Such hypotheses can be addressed in the broader Dual Mechanisms of Cognitive Control dataset (Braver, Kizhner, Tang, Freund, & Etzel, 2020), of which the data used here are a small subset.

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 Extended Data

Figure 1-1: Regions of interest, or "superparcels," for primary analyses. Colored parcels were included in these superparcels. For DLPFC and DMFC, we additionally conducted sensitivity analyses with more inclusive definitions (i.e., including parcels with asterisked labels). Network assignments were provided in Ji et al. (2019). Parcel names that are in bold were identified as likely being situated over nodes of the "multiple demand'' network (Assem, et al. 2020), a collection of regions that is commonly recruited by a variety of demanding tasks. Of these, those in grey belonged to the penumbra (recruited somewhat task-selectively) and those in black belonged to the core (robustly recruited regardless of task). DLPFC, dorsolateral prefrontal cortex; DMFC, dorsomedial frontal cortex; LPPC, lateral posterior parietal cortex; MD, multiple demand; FPN, frontoparietal network; CON, cinguloopercular network; DMN, default mode network; DAN, dorsal attention network; SM, somatomotor network; Vis2, 2nd visual network.

Figure 1-2: Simulation to diagnose the presence and extent of design bias. **A**, Estimated false positive rate of each RSA model across a range of autocorrelation strengths. We simulated time-series of pure noise and regressed these onto each subjects' design matrix used in the analysis. Time-series were drawn $\mathbf{Y}_{t \times v} \sim N_t(\mathbf{0}, \mathbf{\Sigma}_{t \times t})$, for t in $1, \ldots, 1080$ TRs and v in $1, \ldots, 100$ voxels. Multiple $\mathbf{\Sigma}_{t \times t}$s were generated to have AR(1) structure across a range of AR coefficients, ϕ in $0, 0.1, 0.2, \ldots, 0.9, 0.99, 0.999$. For each ϕ, 10,000 simulations were conducted. For each simulation, GLM coefficient estimates from these regressions were submitted to RSA as outlined in *Estimation of Coding Strength* β, and the rate of significantly greater-than-zero RSA model fits was tallied as the false positive rate. At worst (low levels of autocorrelation), models were weakly biased. Note that an empirically-derived false positive rate, obtained from fitting RSA models to patterns of "activation" within ventricles, additionally indicated a false positive rate of no worse than than 0.07 (see Method). **B**, the average RSA correlation matrix from the "worst-case" simulation ($\phi = 0$). In the top panel, the color-scale ranges from ~0.3 to 1. As the off-diagonal appears homogeneous, the scale of the bias was quite small. In the bottom panel, the diagonal (all 1s) is masked so the color-scale shows the bias structure. No "representational" structure is readily apparent (*cf.*, Cai et al., 2020).

Figure 2-1: Parameter estimates from a sensitivity test of the group-level RSA, which used alternative (more inclusive) definitions of DLPFC and DMFC (see Figure 1-1). This table contains statistics analogous to Table 2. $\bar{\beta}$ indicates mean RSA model fit over subjects. Unless noted, contrasts were collapsed across hemisphere. For contrasts performed separately in each hemisphere, p-values were FDR-adjusted for multiple comparisons. Incon., incongruency; distr., distractor.

contrast	$\bar{\beta}$	σ	t	p
DMFC (L) incon.	0.10	0.02	4.16	0.00003
DMFC (R) incon.	0.08	0.02	3.16	0.00157
DMFC (L) target	0.03	0.01	2.38	0.01729
DMFC (R) target	0.05	0.01	3.65	0.00027
DMFC (L) distr.	0.01	0.01	0.82	0.41106
DMFC (R) distr.	0.00	0.01	0.05	0.95851
DLPFC (L) incon.	0.05	0.02	2.34	0.01913
DLPFC (R) incon.	0.04	0.02	2.18	0.02927
DLPFC (L) target	0.03	0.01	2.08	0.03738
DLPFC (R) target	0.05	0.01	3.17	0.00153
DLPFC (L) distr.	-0.01	0.01	-0.70	0.48529
DLPFC (R) distr.	0.00	0.01	0.27	0.78954
incon. − target	DMFC (L)	0.07	0.03	2.64	0.01636
incon. − target	DMFC (R)	0.03	0.03	0.97	0.33044
incon. − distr.	DMFC (L)	0.09	0.02	3.67	0.00049
incon. − distr.	DMFC (R)	0.08	0.02	3.11	0.00188
target − distr.	DMFC (L)	0.02	0.02	1.33	0.18277
target − distr.	DMFC (R)	0.05	0.02	2.99	0.00551
incon. − target	DLPFC (L)	0.02	0.03	0.68	0.95934
incon. − target	DLPFC (R)	0.00	0.03	-0.05	0.95934
incon. − distr.	DLPFC (L)	0.06	0.02	2.28	0.04530
incon. − distr.	DLPFC (R)	0.04	0.02	1.63	0.10409
target − distr.	DLPFC (L)	0.04	0.02	2.24	0.02501
target − distr.	DLPFC (R)	0.04	0.02	2.39	0.02501
DMFC (L) − DLPFC	incon.	0.06	0.02	3.41	0.00065
DMFC (L) − DLPFC	target	0.00	0.02	-0.17	0.86124
DMFC (L) − LPPC	incon.	0.03	0.02	2.01	0.04441
DMFC (L) − LPPC	target	-0.01	0.02	-0.51	0.61058
[incon. − target] · [DMFC (L) − DLPFC]	0.06	0.02	2.84	0.00457
[incon. − target] · [LPPC (L) − DLPFC]	0.03	0.02	1.57	0.11590

Figure 2-2: Mean RSA model fits in each MMP parcel within frontoparietal ROIs. (**A**) DLPFC, (**B**) DMFC, (**C**) LPPC. Hemispheres displayed separately. Error-bars represent 95% confidence intervals on between subject variability (estimated via bias-corrected and accelerated bootstrap, 10,000 resamples).

Figure 2-3: Univariate analysis. **A**, Across-voxel mean levels of activity in response to incongruent versus congruent conditons ("univariate Stroop contrast") within each frontoparietal ROI. Error-bars represent 95% confidence intervals of between-subject variability (estimated via bootstrap). **B**, Subject-level correlations between the univariate Stroop contrast and RSA incongruency model fit.

Figure 2-4: Group-level results from alternative RSA techniques. "Within-run downsampled" columns correspond to results from the downsampling analysis, while "Cross-run downsampled" and "Cross-validated" columns correspond to results from between-run RSA (see relevant Method sections). Following Nili et. al (2014), one-sample contrasts are sign-rank tested against a one-tailed hypothesis, while model comparisons are two-tailed. P-values uncorrected.

contrast	Within-run downsampled			Cross-run downsampled			Cross-validated		
	$\bar{\beta}$	σ	p	$\bar{\beta}$	σ	p	$\bar{\beta}$	σ	p
DMFC (L) incon.	0.095	0.187	0.001	0.064	0.139	0.001	0.010	0.169	0.318
DMFC (R) incon.	0.062	0.173	0.012	0.047	0.134	0.006	-0.007	0.154	0.660
DMFC (L) target	0.044	0.081	0.000	0.015	0.047	0.012	0.020	0.095	0.066
DMFC (R) target	0.057	0.091	0.000	0.025	0.059	0.003	0.041	0.101	0.009
DMFC (L) distr.	0.007	0.064	0.311	-0.002	0.044	0.441	0.006	0.079	0.293
DMFC (R) distr.	0.003	0.057	0.486	0.000	0.039	0.522	-0.009	0.067	0.805
DLPFC (L) incon.	0.035	0.156	0.117	0.020	0.134	0.209	0.037	0.136	0.046
DLPFC (R) incon.	0.070	0.159	0.003	0.026	0.145	0.153	0.037	0.134	0.059
DLPFC (L) target	0.025	0.072	0.011	0.012	0.047	0.026	0.013	0.063	0.052
DLPFC (R) target	0.049	0.085	0.000	0.004	0.055	0.311	0.018	0.080	0.081
DLPFC (L) distr.	-0.012	0.065	0.960	-0.006	0.043	0.954	-0.007	0.062	0.870
DLPFC (R) distr.	-0.005	0.067	0.724	-0.008	0.046	0.850	0.001	0.064	0.502
LPPC (L) incon.	0.125	0.205	0.000	0.095	0.163	0.000	0.027	0.181	0.233
LPPC (R) incon.	0.078	0.209	0.008	0.059	0.166	0.016	0.040	0.152	0.035
LPPC (L) target	0.021	0.063	0.023	0.004	0.041	0.243	0.009	0.080	0.143
LPPC (R) target	0.051	0.080	0.000	0.022	0.044	0.001	0.030	0.080	0.005
LPPC (L) distr.	0.022	0.063	0.048	0.006	0.043	0.165	0.016	0.071	0.080
LPPC (R) distr.	0.026	0.072	0.018	0.010	0.057	0.186	0.015	0.074	0.052
incon. − target | DMFC (L)	0.051	0.196	0.092	0.049	0.150	0.043	-0.010	0.186	0.907
incon. − target | DMFC (R)	0.005	0.188	0.907	0.021	0.149	0.351	-0.048	0.194	0.080
incon. − distr. | DMFC (L)	0.088	0.190	0.003	0.066	0.142	0.001	0.004	0.174	0.733
incon. − distr. | DMFC (R)	0.059	0.181	0.038	0.047	0.140	0.024	0.001	0.183	0.859
target − distr. | DMFC (L)	0.037	0.083	0.002	0.017	0.053	0.086	0.014	0.107	0.436
target − distr. | DMFC (R)	0.054	0.094	0.000	0.025	0.062	0.016	0.049	0.111	0.006
incon. − target | DLPFC (L)	0.010	0.165	0.875	0.008	0.142	0.939	0.024	0.148	0.346
incon. − target | DLPFC (R)	0.022	0.183	0.593	0.022	0.160	0.362	0.019	0.155	0.492
incon. − distr. | DLPFC (L)	0.046	0.166	0.078	0.026	0.148	0.357	0.044	0.143	0.094
incon. − distr. | DLPFC (R)	0.076	0.169	0.006	0.035	0.150	0.092	0.036	0.152	0.182
target − distr. | DLPFC (L)	0.037	0.082	0.005	0.018	0.050	0.015	0.020	0.067	0.061
target − distr. | DLPFC (R)	0.054	0.093	0.000	0.012	0.066	0.273	0.018	0.090	0.182
incon. − target | LPPC (L)	0.104	0.210	0.004	0.091	0.167	0.001	0.018	0.184	0.600
incon. − target | LPPC (R)	0.028	0.218	0.711	0.037	0.181	0.282	0.009	0.165	0.764
incon. − distr. | LPPC (L)	0.104	0.222	0.007	0.089	0.166	0.001	0.011	0.192	0.749
incon. − distr. | LPPC (R)	0.052	0.219	0.239	0.049	0.180	0.157	0.025	0.149	0.373
target − distr. | LPPC (L)	0.000	0.069	0.796	-0.002	0.046	0.788	-0.007	0.075	0.572
target − distr. | LPPC (R)	0.024	0.094	0.104	0.012	0.066	0.230	0.016	0.101	0.512
DMFC (L) − DLPFC | incon.	0.058	0.208	0.116	0.056	0.141	0.007	-0.006	0.179	0.772
DMFC (L) − DLPFC | target	0.007	0.076	0.418	0.006	0.054	0.310	0.004	0.093	0.644
DMFC (L) − LPPC | incon.	-0.007	0.178	0.891	-0.013	0.173	0.559	-0.023	0.196	0.608
DMFC (L) − LPPC | target	0.008	0.075	0.566	0.001	0.046	0.843	0.000	0.085	0.666
[incon. − target] · [DMFC (L) − DLPFC]	0.051	0.196	0.092	0.049	0.150	0.043	-0.010	0.186	0.907
[incon. − target] · [LPPC (L) − DLPFC]	0.050	0.224	0.176	0.054	0.153	0.020	-0.006	0.189	0.875

Figure 2-5: Comparison of results from alternative RSA incongruency models to original parameterization. Two alternative incongruency models were fitted, and model fits (β) were compared to our original parameterization of incongruency coding. See *Alternative RSA Incongruency Models* section of Methods for description of alternative models. Note that the $II - CC$ parameterization is expected to yield higher variance fits than the other two, due to excluding substantially more data per subject. **A**, Incongruency model fits for each parameterization in each of our ROIs. Error bars show bootstrapped 95% CI. Each line is a subject. Model fits were highly similar across parameterizations. **B**, Incongruency model fits for each parameterization in each MMP parcel (each point is a parcel). Model fits were highly similar across parameterizations.

Figure 3-1: Associations between subjects' coding strength (β) and Stroop interference effect (response time) separately displayed within each hemisphere, ROI, and coding model (target, incongruency). Green, target coding model; orange; incongruency coding model. Confidence bands indicate 95% confidence interval from percentile bootstrap (10,000 resamples).

Figure 3-2: Robustness check of individual-level correlations found in our primary analysis (see Figure 3 and Table 3) to alternative RSA techniques. "Within-run downsampled" corresponds to results from the downsampling analysis, while "Cross-run downsampled" and "Cross-validated" corresponds to the between-run RSA (see relevant Method sections). While attenuation of effect sizes is generally expected when moving to higher variance methods (e.g., downsampling, between-run RSA), critically, the signs of the effects are robust to technique. Notably, however, the large reduction in DMFC incongruency coding correlation with cross-validated RSA, but preservation of the effect with cross-run RSA, matches what is seen at the group level (Figure 2-4, Figure 4-4), and suggests that incongruency information was encoded non-linearly within DMFC activation patterns.

Figure 3-3: Superparcels defined for model-selection analysis. All superparcels were lateralized (one in each hemisphere) except for relatively early sensory and motor regions ('Aud,' 'V1–V3,' 'VSM'). Ventral somatomotor cortex was defined using the Gordon et al. (2016) atlas 'somato-motor–mouth network,' as MMP's somatomotor parcels encompass all of somatomotor strip (i.e., dorsal and ventral areas). Left ventral occipitotemporal cortex ("visual word-form area") was defined using coordinates from Twomey et al. (2011). DPM, dorsal premotor cortex; VPM, ventral premotor cortex; VVis, ventral visual cortex; AIns, anterior (frontal) insular cortex; IFC, inferior frontal cortex; OFC, orbitofrontal cortex; FPC, frontoparietal cortex; Aud, early auditory cortex; V1–V3, visual cortex, VSM, ventral somatomotor cortex. LVOT, left visual occipitotemporal cortex.

superparcel	constituent MMP parcels	# voxels (L, R)
DPM	6a, FEF, 6ma	508, 495
VPM	6v, PEF, 6r, 55b, IFJp	629, 572
VVis	FFC, VVC, V8, VMV3	648, 721
AIns	FOP4, FOP5, FOP3, AVI, MI	527, 536
IFC	IFJa, 44, 45, IFSa, p47r, IFSp, a47r, 47l	1065, 1263
OFC	47s, 47m, 11l, 13l, OFC, 10pp, 10v, 10r, s32, p32, a24, 25, pOFC	1927, 1810
FPC	46, 9-46d, a9-46v, a10p, p10p, a47r, 9a, 9p, 10d	1657, 1769
Aud	LBelt, A1, MBelt, PBelt	473
V1–V3	V1, V2, V3	3904
VSM	none; Gordon et al. (2016)	1114
LVOT	none; Twomey et al. (2011)	939

Figure 4-1: MMP parcels with significant target model fits. P-values are FDR-corrected across all 360 cortical parcels. $\bar{\beta}$, Across-subject mean of RSA model fits. v, Wilcoxon sum of signed-rank test statistic. network, Cole-Anticevic network assignemnt. md, Multiple-Demand assignment (Assem, et al. 2020).

parcel	$\bar{\beta}$	v	p	network	md
3b_L	0.08	1108	0.00001	Somatomotor	
4_L	0.07	1097	0.00001	Somatomotor	
FEF_R	0.05	1063	0.00010	Cingulo-Opercular	
4_R	0.06	1049	0.00016	Somatomotor	
3b_R	0.06	1047	0.00016	Somatomotor	
OP4_L	0.06	1033	0.00031	Somatomotor	
3a_L	0.07	1009	0.00088	Somatomotor	
PFop_L	0.05	1008	0.00088	Cingulo-Opercular	
11l_L	0.04	1004	0.00096	Frontoparietal	penumbra
11l_R	0.04	1000	0.00106	Frontoparietal	penumbra
3a_R	0.04	993	0.00135	Somatomotor	
PFop_R	0.04	989	0.00139	Cingulo-Opercular	
STSda_R	0.04	989	0.00139	Default	
a47r_L	0.04	985	0.00156	Frontoparietal	penumbra
V8_L	0.04	982	0.00164	Visual2	
V6_R	0.04	981	0.00164	Visual2	
LBelt_L	0.04	978	0.00165	Auditory	
p10p_L	0.05	978	0.00165	Frontoparietal	penumbra
5mv_R	0.05	977	0.00165	Cingulo-Opercular	
55b_L	0.05	976	0.00165	Language	
8C_R	0.04	974	0.00172	Frontoparietal	core
p9-46v_R	0.05	971	0.00188	Frontoparietal	core
POS2_L	0.05	970	0.00188	Frontoparietal	penumbra
6v_R	0.05	969	0.00188	Somatomotor	
PFcm_L	0.04	966	0.00206	Cingulo-Opercular	
55b_R	0.04	965	0.00207	Language	
43_R	0.04	963	0.00209	Cingulo-Opercular	
STGa_L	0.04	963	0.00209	Language	
7PC_R	0.04	957	0.00252	Somatomotor	
MI_R	0.04	957	0.00252	Cingulo-Opercular	
SCEF_L	0.03	956	0.00255	Cingulo-Opercular	core
23c_L	0.05	953	0.00280	Cingulo-Opercular	
FOP5_L	0.05	952	0.00283	Cingulo-Opercular	penumbra
TGv_R	0.04	951	0.00286	Language	
a32pr_R	0.04	950	0.00289	Cingulo-Opercular	penumbra
6d_L	0.04	949	0.00293	Somatomotor	
7PC_L	0.04	947	0.00310	Somatomotor	
A4_R	0.04	945	0.00319	Auditory	
AIP_R	0.05	945	0.00319	Dorsal-Attention	penumbra
IP0_R	0.04	943	0.00328	Dorsal-Attention	
MBelt_R	0.04	943	0.00328	Auditory	
STSda_L	0.05	942	0.00334	Language	
MI_L	0.03	941	0.00339	Cingulo-Opercular	
FOP4_R	0.05	938	0.00371	Cingulo-Opercular	
SCEF_R	0.04	937	0.00371	Cingulo-Opercular	core
TGd_R	0.04	937	0.00371	Default	
PBelt_R	0.05	936	0.00377	Auditory	
47l_L	0.03	935	0.00377	Default	
VVC_L	0.04	935	0.00377	Visual2	
5m_L	0.04	934	0.00384	Somatomotor	
6ma_R	0.04	933	0.00384	Cingulo-Opercular	
V3B_L	0.04	933	0.00384	Visual2	
6r_R	0.04	932	0.00391	Cingulo-Opercular	penumbra
6v_L	0.04	931	0.00399	Somatomotor	
6a_R	0.03	930	0.00407	Dorsal-Attention	
a10p_R	0.03	928	0.00424	Frontoparietal	penumbra
POS2_R	0.04	928	0.00424	Frontoparietal	penumbra
2_L	0.04	927	0.00433	Somatomotor	
DVT_R	0.04	923	0.00494	Visual1	
PGi_L	0.03	920	0.00543	Default	
9p_L	0.04	919	0.00554	Default	
p32pr_R	0.04	917	0.00587	Cingulo-Opercular	
FOP4_L	0.03	916	0.00599	Cingulo-Opercular	
IP1_R	0.04	915	0.00611	Frontoparietal	core
8BM_R	0.05	912	0.00660	Frontoparietal	core
PFm_R	0.03	912	0.00660	Frontoparietal	core
OP2-3_L	0.03	910	0.00698	Somatomotor	
6mp_L	0.04	909	0.00713	Somatomotor	
FOP3_R	0.04	908	0.00717	Cingulo-Opercular	
V4t_L	0.03	908	0.00717	Visual2	
OP4_R	0.04	907	0.00733	Somatomotor	
d32_R	0.04	906	0.00738	Frontoparietal	penumbra
VVC_R	0.04	906	0.00738	Visual2	
43_L	0.03	905	0.00744	Cingulo-Opercular	

Figure 4-1 continued

parcel	$\bar{\beta}$	v	p	network	md
TGd_L	0.04	905	0.00744	Default	
IFSp_R	0.04	904	0.00760	Frontoparietal	
47s_L	0.04	903	0.00776	Default	
PBelt_L	0.04	902	0.00793	Auditory	
45_L	0.04	899	0.00847	Language	
TA2_L	0.03	899	0.00847	Auditory	
V6A_R	0.03	899	0.00847	Visual2	
a24pr_R	0.03	897	0.00895	Cingulo-Opercular	
d32_L	0.03	896	0.00915	Default	penumbra
44_R	0.03	895	0.00935	Frontoparietal	
FEF_L	0.04	893	0.00976	Cingulo-Opercular	
IFJa_R	0.04	893	0.00976	Language	
VIP_L	0.03	892	0.00997	Visual2	
24dd_L	0.04	891	0.01008	Somatomotor	
47m_R	0.03	891	0.01008	Default	
AVI_R	0.03	890	0.01030	Frontoparietal	core
31pv_R	0.04	889	0.01041	Default	
FOP5_R	0.05	889	0.01041	Cingulo-Opercular	penumbra
TE2a_L	0.03	886	0.01136	Default	
FOP1_R	0.03	885	0.01136	Cingulo-Opercular	
FOP3_L	0.04	885	0.01136	Cingulo-Opercular	
TPOJ1_L	0.03	885	0.01136	Language	
24dv_R	0.03	884	0.01162	Somatomotor	
AVI_L	0.03	883	0.01166	Frontoparietal	core
PoI2_L	0.04	883	0.01166	Cingulo-Opercular	
FOP1_L	0.03	882	0.01166	Cingulo-Opercular	
TA2_R	0.02	882	0.01166	Auditory	
TE1a_R	0.04	882	0.01166	Default	
TE2p_L	0.03	882	0.01166	Dorsal-Attention	
V7_L	0.03	880	0.01231	Visual2	
10d_L	0.03	879	0.01235	Default	
8Ad_R	0.03	879	0.01235	Default	
AIP_L	0.03	879	0.01235	Dorsal-Attention	penumbra
FST_L	0.03	878	0.01251	Visual2	
SFL_L	0.04	878	0.01251	Language	
a9-46v_R	0.04	877	0.01279	Frontoparietal	core
STV_L	0.02	876	0.01308	Language	
IP2_R	0.04	875	0.01325	Frontoparietal	core
IPS1_R	0.02	875	0.01325	Visual2	
10pp_L	0.03	874	0.01343	Default	
8Ad_L	0.03	874	0.01343	Default	
FFC_L	0.03	872	0.01393	Visual2	
FST_R	0.04	872	0.01393	Visual2	
PeEc_L	0.04	872	0.01393	Ventral-Multimodal	
OFC_R	0.03	871	0.01424	Frontoparietal	
RI_L	0.04	870	0.01456	Auditory	
PHT_L	0.03	869	0.01476	Dorsal-Attention	
TE1a_L	0.03	869	0.01476	Default	
a10p_L	0.03	868	0.01509	Frontoparietal	penumbra
PGi_R	0.02	867	0.01543	Default	
IFJp_R	0.03	865	0.01625	Frontoparietal	core
8Av_R	0.03	864	0.01635	Default	
i6-8_L	0.03	864	0.01635	Frontoparietal	core
p47r_L	0.03	864	0.01635	Frontoparietal	penumbra
5mv_L	0.03	862	0.01709	Cingulo-Opercular	
FFC_R	0.02	862	0.01709	Visual2	
8BL_R	0.03	858	0.01892	Default	
PI_R	0.03	858	0.01892	Cingulo-Opercular	
PEF_L	0.03	857	0.01904	Dorsal-Attention	
STV_R	0.03	857	0.01904	Posterior-Multimodal	
V2_L	0.03	857	0.01904	Visual2	
OFC_L	0.02	856	0.01904	Default	
PSL_R	0.03	856	0.01904	Cingulo-Opercular	
STSdp_R	0.03	856	0.01904	Language	
V3_L	0.03	856	0.01904	Visual2	
24dd_R	0.03	855	0.01919	Somatomotor	
46_L	0.03	855	0.01919	Cingulo-Opercular	
7AL_R	0.03	855	0.01919	Somatomotor	
IFSa_L	0.02	854	0.01934	Frontoparietal	
V2_R	0.02	854	0.01934	Visual2	
V6A_L	0.03	854	0.01934	Visual2	
1_L	0.03	853	0.01976	Somatomotor	
A5_L	0.02	851	0.02049	Language	
OP1_L	0.03	851	0.02049	Somatomotor	
OP1_R	0.02	851	0.02049	Somatomotor	
10r_R	0.03	850	0.02067	Default	
6r_L	0.03	850	0.02067	Cingulo-Opercular	penumbra
PFcm_R	0.02	850	0.02067	Cingulo-Opercular	
PH_L	0.03	848	0.02172	Visual2	
1_R	0.03	847	0.02190	Somatomotor	
A5_R	0.03	847	0.02190	Language	
p9-46v_L	0.04	847	0.02190	Frontoparietal	core
Ig_L	0.03	845	0.02272	Somatomotor	

Figure 4-1 continued

parcel	$\bar{\beta}$	v	p	network	md
PCV_R	0.03	845	0.02272	Posterior-Multimodal	
VMV3_R	0.03	845	0.02272	Visual2	
LO2_L	0.02	844	0.02320	Visual2	
MT_R	0.02	842	0.02421	Visual2	
TF_R	0.02	842	0.02421	Ventral-Multimodal	
6mp_R	0.02	840	0.02525	Somatomotor	
TGv_L	0.03	840	0.02525	Language	
OP2-3_R	0.03	837	0.02721	Somatomotor	
LIPv_L	0.02	836	0.02745	Visual2	
LO3_R	0.03	836	0.02745	Visual2	
PEF_R	0.03	836	0.02745	Cingulo-Opercular	
LIPv_R	0.03	835	0.02802	Visual2	
a24_L	0.03	834	0.02827	Default	
DVT_L	0.03	834	0.02827	Visual1	
IFJa_L	0.03	834	0.02827	Language	
a47r_R	0.03	832	0.02963	Frontoparietal	penumbra
6d_R	0.03	831	0.02990	Somatomotor	
A4_L	0.03	831	0.02990	Auditory	
TE1m_L	0.02	831	0.02990	Default	penumbra
SFL_R	0.03	830	0.03051	Language	
8Av_L	0.02	829	0.03114	Default	
8C_L	0.03	827	0.03260	Frontoparietal	core
TPOJ1_R	0.03	826	0.03327	Language	
24dv_L	0.03	824	0.03481	Somatomotor	
9a_R	0.02	823	0.03531	Default	
V8_R	0.03	823	0.03531	Visual2	
9m_R	0.03	822	0.03582	Default	
ProS_R	0.03	822	0.03582	Visual1	
8BM_L	0.03	821	0.03634	Frontoparietal	core
H_R	0.02	821	0.03634	Default	
AAIC_R	0.02	820	0.03707	Orbito-Affective	
10v_R	0.02	818	0.03834	Default	
44_L	0.02	818	0.03834	Language	
p32pr_L	0.02	818	0.03834	Cingulo-Opercular	
a24pr_L	0.03	817	0.03910	Cingulo-Opercular	
a24_R	0.02	815	0.04064	Default	
PHA3_L	0.03	815	0.04064	Dorsal-Attention	
FOP2_L	0.02	814	0.04143	Somatomotor	
IFSp_L	0.02	811	0.04432	Language	
10v_L	0.02	810	0.04471	Default	
23c_R	0.02	810	0.04471	Cingulo-Opercular	
7PL_R	0.03	810	0.04471	Dorsal-Attention	
9a_L	0.02	808	0.04643	Default	
STSvp_R	0.03	808	0.04643	Default	
9-46d_R	0.03	806	0.04750	Cingulo-Opercular	
IFJp_L	0.02	806	0.04750	Frontoparietal	core
IP0_L	0.02	806	0.04750	Dorsal-Attention	
PGp_L	0.02	806	0.04750	Dorsal-Attention	
POS1_R	0.03	806	0.04750	Default	
s6-8_R	0.03	805	0.04839	Frontoparietal	penumbra

Figure 4-2: MMP parcels with significant distractor model fits. P-values are FDR-corrected across all 360 cortical parcels. $\bar{\beta}$, Across-subject mean of RSA model fits. v, Wilcoxon sum of signed-rank test statistic. network, Cole-Anticevic network assignemnt. md, Multiple-Demand assignment (Assem, et al. 2020).

parcel	$\bar{\beta}$	v	p	network	md
V2_L	0.05	1029	0.00144	Visual2	
V1_L	0.05	1025	0.00144	Visual1	

Figure 4-3: MMP parcels with significant incongruency model fits. P-values are FDR-corrected across all 360 cortical parcels. $\bar{\beta}$, Across-subject mean of RSA model fits. v, Wilcoxon sum of signed-rank test statistic. network, Cole-Anticevic network assignemnt. md, Multiple-Demand assignment (Assem, et al. 2020).

parcel	$\bar{\beta}$	v	p	network	md
LIPd_L	0.12	978	0.01973	Dorsal-Attention	penumbra
MIP_L	0.13	967	0.01973	Dorsal-Attention	penumbra
s6-8_R	0.09	961	0.01973	Frontoparietal	penumbra
IP1_L	0.14	956	0.01973	Frontoparietal	core
6ma_L	0.09	937	0.02743	Cingulo-Opercular	
RSC_L	0.09	937	0.02743	Frontoparietal	
9-46d_L	0.07	934	0.02743	Cingulo-Opercular	
IP0_R	0.10	920	0.03821	Dorsal-Attention	
6ma_R	0.08	918	0.03821	Cingulo-Opercular	
LO2_R	0.09	914	0.03821	Visual2	
8BM_R	0.09	913	0.03821	Frontoparietal	core
p32pr_R	0.08	907	0.04334	Cingulo-Opercular	

Figure 4-4: Robustness check of results from exploratory whole-cortex RSA (see Figure 4) to alternative RSA techniques. "Within-run downsampled" corresponds to results from the downsampling analysis, while "Cross-run downsampled" and "Cross-validated" corresponds to between-run RSA (see relevant Method sections). Maps are threhsolded for model fits greater than 0 with significance at FDR-corrected (across all 360 parcels) p-value of 0.05. Note that, in the present design, between-run RSA methods are expected to strongly attenuate effect sizes (see Method). Nevertheless, our core results were robust: in both downsampled and between-run RSA methods, we identified incongruency coding in parcels within DMFC, target coding in mid-DLPFC, and distractor coding in visual cortex. Notably, the reduction in DMFC incongruency coding with cross-validated RSA, but preservation of the effect with cross-run RSA, matches what is seen at the individual-difference level (Figure 3-2), and suggests that incongruency information was encoded non-linearly within DMFC activation patterns (see also Figure 2, B). Relative to the prewhitening approach we used to handle imbalance of trials across runs, the downsampling approach appears to have increased sensitivity to incongruency coding, in terms of coverage (cf., Figure 4). This could be due to the prewhitening approach (in a separate, preceeding regression) being a somewhat more aggressive denoising strategy. Critically, however, this did not impact our main conclusions: we still observed stronger incongruency coding in DMFC than in DLPFC (see Figure 2-4).