### Preregistration

## My fMRI preregistration

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Working Title	Enter your response here.	
Introduction	Enter your response here.	
Current Study	Enter your response here.  Aims & Hypotheses	
	Essential	Checklist
	Hypotheses include direction of expected results	
	Interactions describe expected shape	or NA
	Maniputated variables include	or NA
	manipulation checks or explain why not	

	Recommended	Checklist
	Figure or table to describe expected	
	results	
	Rationals or frameworks included for	
	why certain hypotheses are being tested	
	Which outcome would be predicted by	or NA
	which theory	
		_
Exisiting Data		
	• Registration prior to creation of da	ca
	• Registration prior to any human ob	servation of the data
	• Registration prior to accessing the	lata
	• Registration prior to analysis of the	data
	• Registration following analysis of the	ne data
Explanation of Exisiting Data	Enter your response here.	
Details of Larger	Is your preregistration part of a large	ger project?
$\mathbf{Study}$	• Yes	
	• No	
	Enter your response here.	
Data Collection	For group comparisons, what vari	ables (if any) were equated across
Procedures	groups?	
	Enter your response here.	
	Study timeline	
	Enter your response here.	

Collection	Checklist
Population	
Recruitment efforts	
Inclusion/Exclusion criteria	
Clinical criteria (if applicable)	
Matching strategy (if applicable)	
Payment for participation	
IRB, consent/assent obtained	
Number of subjects participated and analyzed	
Age	
Sex	
Handedness	

# Sample Size & Stopping Rule

#### Target sample size

Enter your response here.

#### Justification of sample size

Enter your response here.

Power analyses (e.g., Neuropowertools, fmri power)

From Nichols et al., 2016, include:

Effect Size Estimate	Checklist
Effect size used	or NA*
Source of predicted effect size (prior lit, pilot etc.)	or NA*
Significant level	or NA*
Target power	or NA*
Outcome used to calculate	or NA*

#### Stopping rule

Enter your response here.

Contingencies for if your target sample size is not met

Enter your response here.

Measured

Outcome measures

Behavioral Variables

Enter your response here.

Predictor measures

Enter your response here.

Covariate measures

Enter your response here.

How was behavioral task performance measured?

Enter your response here.

Contingency plans for behavioral analysis

Enter your response here.

Additional

Region Specificity

Operational Definitions

Enter your response here.

Any other definitions used across study:

Enter your response here.

Transformations

Enter your response here.

Contingency plans for transformation

Enter your response here.

Code, if applicable: for scoring behavioral data.

Analysis Data

Exclusion

Enter your response here.

### 

Enter your response here.

#### **Design Specifications**

Design	Checklist
Design type (task, rest; event-related, block)	
Conditions & Stimuli (detailed as possible, pictures	
encouraged)	
Number of blocks, trials or experimental units per session	
and/or subject	
Timing and Duration (length of each trial and interval	
between trials)	
Length of experiment (length of full scan and each run)	
Was the design optimized for efficiency, and if so, how?	
Presentation software (name, version, operating system; code	
if possible)	

#### Task Specification

Task	Checklist
Instructions to subjects (what were they asked to do?)	
Stimuli (what were they; how many were there; did it repeat	
across trials?)	
Stimuli presentation & response collection	
Randomization/pseudo-randomized (why/why not done $\&$	
how)	
Run order (of tasks within scanner)	

#### Data acquisition

#### Set up & Acquisitions

Set up & Acquisitions	Checklist
Participant Preparation	
Mock scanning (Report type of mock scanner and protocol;	or NA
i.e. duration, types of sim scans, exper)	

Set up & Acquisitions	Checklist
Specific accommodations (e.g., pediatric, parent present?	or NA
Asleep?)	
Experimental personnel (number of planned personnel to	or NA
interact with subjects)	
MRI System	
Manufacturer, field strength (in Tesla), model name	
MRI acquisition	
Pulse sequence (gradient/spin echo etc.)	
Image type (EPI, spiral, 3D etc.)	
Essential sequence & imaging parameters	
For All	
Echo time (TE)	
Repetition time (TR)	
For multi-shot acquisitions, additionally the time per volume	or NA
Flip angle (FA)	
Acquisition time (duration of acquisition)	
Functional MRI	
Number of volumes	
Inversion Recovery Sequences	
Sparse sampling delay (delay in TR) if used	or NA
B0 Field Maps	
Inversion time (TI) for inversion recovery sequence	or NA
Echo time difference (dTE). Diffusion MRI	or NA
Number of directions	or NA
Direction optimization, if used and type	or NA
b-values	$\_\_\_$ or NA
Number of b=0 images	or NA
Number of averages (if any)	or NA
Single shell, multi-shell (specify equal or unequal spacing)	or NA
Single- or dual-spin-echo, gradient mode (serial or parallel)	or NA
If cardiac gating used	or NA
Imaging Parameters	

Set up & Acquisitions	Checklist
Field of view	or NA
In-plane matrix size, slice thickness and interslice gap, for 2D	or NA
acquisitions	
Slice orientation (Axial, sagittal, coronal or oblique)	or NA
Angulation: If acquistion not aligned with scanner axes,	or NA
specified	
angulation to AC-PC line (see Slice position procedure)	or NA
3D matrix size, for 3D acquisitions	or NA
Additional sequence & imaging parameters	
Phase encoding	or NA
Parallel imaging method & parameters	or NA
Multiband parameters	or NA
Readout parameters	or NA
Fat suppression (for anatomical, state if used)	or NA
Shimming	or NA
Slice order & timing	$\_\_\_$ or NA
Brain coverage (e.g., whole brain, was cerebellum, brain stem	or NA
included)	
Scanner-side preprocessing*	$\_\_\_$ or NA
Scan duration (in seconds)	$\_\_\_$ or NA
Other non-standard procedures	$\_\_\_$ or NA
T1 stabilization (discarded "dummy" scans acquired discarded	or NA
by scanner)	
Diffusion MRI gradient table (Also referred to as the	$\_\_$ or NA
b-matrix, but not to be confused with the 3x3 matrix that	
describes diffusion weighting for a single diffusion weighted	
measurement) scanner-side preprocessing: (e.g., Including:	
Reconstruction matrix size differing from acquisition matrix	
size; Prospective-motion correction (including details of any	
optical tracking, and how motion parameters are used); Signal	
inhomogeneity correction; Distortion-correction.)	

Perfusion: Arterial Spin Labelling MRI	Checklist
ASL Labelling method (e.g. continuous ASL (CASL),	or NA
pseudo-continuous ASL (PCASL), Pulsed ALS	
(PASL), velocity selective ASL (VSASL)	27.4
Use of background suppression pulses and their timing	
label duration	or NA if not
	PCASL or CASL
Label Duration	or NA if not
	PCASL or CASL
Post labeling delay (PLD)	or NA if not
	PCASL or CASL
Location of the labeling plane	or NA
Average labeling gradient	or NA if not
	PCASL
Slice-selective labeling gradient	or NA if not
	PCASL
Flip angle of B1 pulses	or NA if not
	PCASL
Assessment of inversion efficiency	or NA if not
	PCASL
QC used to ensure off-resonance artifacts not	or NA if not
problematic	PCASL
signal obtained over whole brain	or NA if not
	PCASL
Use of a separate labeling coil	or NA if not CASL
Control scan/pulse used	or NA if not CASL
B1 amplitude	or NA if not CASL
ТІ	or NA if not PASL
Labeling slab thickness	or NA if not PASL
Use of QUIPSS pulses and their timing	or NA if not PASL
TI	or NA if not
	VSASL

Perfusion: Arterial Spin Labelling MRI	Checklist
Choice of velocity selection cutoff ("VENC")	or NA if not
	VSASL
Perfusion: Dynamic Susceptibility Contrast	
MRI	
Number of baseline volumes	or NA if not
	Perfusion: Dynamic
	Susceptibility Contras
	MRI
Type, name and manufacturer of intravenous bolus	or NA if not
(e.g. gadobutrol, Gadavist, Bayer)	Perfusion: Dynamic
	Susceptibility Contras
	MRI
Bolus amount and concentration (e.g. 0.1 ml/kg and	or NA if not
0.1 mmol/kg) Injection rate (e.g. 5 ml/s)	Perfusion: Dynamic
	Susceptibility Contras
	MRI
Post-injection of saline (e.g. 20 ml)	or NA if not
	Perfusion: Dynamic
	Susceptibility Contras
	MRI
Injection method (e.g. power injector)	or NA if not
	Perfusion: Dynamic
	Susceptibility Contras
	MRI
Preliminary quality control	
Preliminary quality control	Checklist
Motion monitoring (For functional or diffusion acquisi	tions,
any visual or quantitative checks for severe motion; like	xewise,
for structural images, checks on motion or general images	ıge
quality.)	

Preprocessing

Preliminary quality control	${\bf Checklist}$
Incidental findings (Protocol for review of any incidental	
findings, and how they are handled in particular with respect	
to possible exclusion of a subject's data.)	
Data preprocessing	
For each piece of software used, give the version number number is available, date of last application of updates)	er (or, if no version
If any subjects required different processing operations analysis, those differences should be specified explicitly	or settings in the
Pre-processing: general	Checklist
Specify order of preprocessing operations	
Describe any data quality control measures	
Unwarping of B0 distortions	
Slice timing correction	
Reference slice and type of interpolation used (e.g., "Slice	Fourier phase
timing correction to the first slice as performed, using SPM5's $$	shift
	interpolation")
Motion correction	
Reference scan, image similarity metric, type of interpolation	
used, degrees-of-freedom (if not rigid body) and, ideally,	
optimization method, e.g., "Head motion corrected with FSL's $$	
MCFLIRT by maximizing the correlation ratio between each	
time point and the middle volume, using linear interpolation." $$	
Motion susceptibility correction used	
Smoothing	
Size and type of smoothing kernel (provide justification for	to reduce noise")
size; e.g., for a group study, "12 mm FHWM Gaussian	
smoothing applied to ameliorate differences in intersubject	
localization"; for single subject fMRI "6 mm FWHM Gaussian	
smoothing used	

Intersubject registration	Checklist
Intersubject registration method used	
Illustration of the voxels present in all subjects ("mask	
image") can be helpful, particularly for restricted fields of	
view (to illustrate overlap of slices across all subjects). Better	
still would be an indication of average BOLD sensitivity	
within each voxel in the mask	
Transformation model and optimization	
Transformation model (linear/affine, nonlinear), type of any	
${\it non-linear\ transformations\ (polynomial,\ discrete\ cosine\ basis)},$	
number of parameters (e.g., 12 parameter affine, 3 x 2 x 3	
DCT basis), regularization, image-similarity metric, and	
interpolation method	
Object image information (image used to determine	
transformation to atlas)	
Anatomical MRI? Image properties (see above)	
Co-planar with functional acquisition?	
Functional acquisition co-registered to an atomical? if so, how?	
Segmented gray image?	
Functional image (single or mean)	
Atlas/target information	
Brain image template space, name, modality and resolution	
(e.g., "FSL's MNI Avg152, T1 2 x 2 x 2 mm"; "SPM2's MNI	
gray matter template 2 x 2 x 2 mm")	
Coordinate space	
(Typically MNI, Talairach, or MNI converted to Talairach	
If MNI converted to Talairach, what method? e.g., Brett's	
mni2tal?	
How were anatomical locations (e.g., gyral anatomy,	
Brodmann areas) determined? (e.g., paper atlas, Talairach	
Daemon, manual inspection of individuals' anatomy, etc.)	
Smoothing	

Intersubject registration	Checklist
Size and type of smoothing kernel (provide justification for	to reduce noise")
size; e.g., for a group study, "12 mm FHWM Gaussian	
smoothing applied to ameliorate differences in intersubject	
localization"; for single subject fMRI "6 mm FWHM Gaussian	
smoothing used	

# Statistical modeling

#### Planned comparison

Enter your response here.

#### General issues

Enter your response here.

#### First level (fx) modeling

First level (fx) modeling	Checklist
Event related design predictors	
Modeled duration, if other than zero	
Parametric modulation	
Block Design predictors(Note whether baseline was explicitly	
modeled)	
HRF basis, typically one of:	
Canonical only	
Canonical plus temporal derivative	
Canonical plus temporal and dispersion derivative. Smooth	
basis (e.g. SPM "informed" or Fourier basis; FSL's FLOBS)	
Finite Impulse Response model	
Drift regressors (e.g. DCT basis in SPM, with specified	
cut-off)	
Movement regressors; specify if squares and/or temporal	
derivative used	
Any other nuisance regressors, and whether they were entered	
as interactions (e.g. with a task effect in 1st level fMRI, or	
with group effect)	

First level (fx) modeling	Checklist
Any orthogonalization of regressors, and set of other	
regressors used to orthogonalize against	
Contrast construction (Exactly what terms are subtracted	
from what? Define these in terms of task or stimulus	
conditions (e.g., using abstract names such as AUDSTIM,	
VISSTIM) instead of underlying psychological concepts	
Autocorrelation model type (e.g., $AR(1)$ , $AR(1) + WN$ , or	
arbitrary autocorrelation function), and whether global or	
local (e.g., for SPM2/SPM5, 'Approximate $AR(1)$	
autocorrelation model estimated at omnibus F-significant	
voxels (P $< 0.001$ ), used globally over the whole brain'; for	
FSL, 'Autocorrelation function estimated locally at each	
voxel, tapered and regularized in space.')	

### Second level (group) modeling

Second level (group) modeling	Checklist
Statistical model and estimation method, inference type	
(mixed/random effects or fixed), e.g., "Mixed effects inference	
with one sample t-test on summary statistic" (SPM2/SPM5),	
e.g., "Mixed effects inference with Bayesian 2-level model with	
fast approximation to posterior probability of activation."	
(FSL)	
If fixed effects inference used, justify	
If more than 2-levels, describe the levels and assumptions of	
the model (e.g., are variances assumed equal between groups)	
Repeated measures?	

Second level (group) modeling	Checklist
If multiple measurements per subject, list method to account	
for within subject correlation, exact assumptions made about	
correlation/variance e.g., SPM: "Within-subject correlation	
estimated at F-significant voxels (P $<$ 0.001), then used	
globally over whole brain"; or, if variances for each measure	
are allowed to vary, "Within-subject correlation and relative	
variance estimated"	
For group model with repeated measures, specify:	
How condition effects are modeled (e.g. as factors, or as linear	
trends)	
Whether subject effects are modeled (i.e. as regressors, as	
opposed to	
with a covariance structure)	
For group effects: clearly state whether or not covariates are	
split by group (i.e. fit as a group-by-covariate interaction)	
Model type (Some suggested terms include:	
"Mass Univariate"	
"Multivariate" (e.g. ICA on whole brain data)	
"Mass Multivariate" (e.g. MANOVA on diffusion or	
morphometry tensor data)	
"Local Multivariate" (e.g. "searchlight")	
"Multivariate, intra-subject predictive" (e.g. classify individual	
trials in	
event-related fMRI)	
"Multivariate inter-subject predictive" (e.g. classify subjects as	
patient vs.	
control)	
"Representational Similarity Analysis")	
Model settings (The essential details of the model For mass	
univariate, first level fMRI, these include:	
Drift model, if not already specified as a dependent variable	
(e.g. locally linear detrending of data & regressors, as in FSL)	

Second level (group) modeling	Checklist
Autocorrelation model (e.g. global approximate AR(1) in	
${\rm SPM; \ locally \ regularized \ autocorrelation \ function \ in \ FSL)}$	
$For \ mass \ univariate \ second \ level \ fMRI \ these \ include:$	
Fixed effects (all subjects' data in one model)	
Random or mixed-effects model, implemented with:	
Ordinary least squares (OLS, aka unweighted summary	
statistics approach; SPM default, FSL FEAT's "Simple OLS")	
weighted least squares (i.e. FSL FEAT's "FLAME 1"), using	
voxel-wise estimate of between subject variance	
Global weighted least squares (i.e. SPM's MFX)	
With any group (multi-subject) model, indicate any specific	
variance structure, e.g.	
Un-equal variance between groups (and if globally pooled, as	
in SPM)	
If repeated measures, the specific covariance structure	
assumed (e.g.	
compound symmetric, or arbitrary; if globally pooled)	
For local-multivariate report:	
The number of voxels in the local model	
Local model used (e.g. Canonical Correlation Analysis) with	
any	
constraints (e.g. positive weights only)	

#### ROI analysis

ROI analysis	Checklist
How were ROIs defined (e.g., functional, anatomical, parcel	
localizer)?	
How was signal extracted within ROI? (e.g., average	
parameter estimates, FIR deconvolution?)	
If percent signal change reported, how was scaling factor	
determined (e.g., height of block regressor or height of	
isolated event regressor)?	

ROI analysis	Checklist
Is change relative to voxel-mean, or whole-brain mean?	
Justify definition of ROI and analysis conducted with it: (e.g., $$	
if your ROI is defined based on the cluster; how will you	
ensure your ROI analyses are not circular?)	

Enter your response here.

# Statistical inference

#### Statistical inference

Statistical inference	Checklist
Search region (Type of search region for analysis, and the	or NA
volume in voxels or CC)	
If not whole brain, state how region was determined; method	or NA
for constructing region should be independent of present	
statistic image	
Whole brain or "small volume"; carefully describe any small	or NA
volume correction used for each contrast	
If a small-volume correction mask is defined anatomically,	or NA
provide named anatomical regions from a publicly available	
ROI atlas	
If small-volume correction mask is functionally defined, clearly	or NA
describe the functional task and identify any risk of circularity	
All small-volume corrections should be fully described in the	or NA
methods section, not just mentioned in passing in the results	
Statistical type (Typically one of:	
Voxel-wise (aka peak-wise in SPM)	or NA
Cluster-wise	or NA
Cluster size	or NA
Cluster mass	or NA
Threshold-free Cluster Enhancement (TFCE)	or NA
For cluster size or mass, report:	
Cluster-forming threshold	or NA
~	

Statistical inference	Checklist
For all cluster-wise methods, report:	
Neighborhood size used to form clusters (e.g. 6, 18 or 26)	or NA
For TFCE, report:	
Use of non-default TFCE parameters.)	or NA
P value computation (Report if anything but standard	
parametric inference used to obtain (uncorrected) P-values. If	
nonparametric method was used, report method	
(e.g. permutation or bootstrap) and number of	
permutations/samples used.)	
Multiple test correction (For mass-univariate, specify the type	or NA
of correction and how it is obtained, especially if not the	
typical usage.)	
Usually one of:	
Familywise Error	
Random Field Theory (typical)	or NA
Permutation	or NA
Monte Carlo	or NA
Bonferroni	or NA
False Discovery Rate	
Benjamini & Hochberg FDR (typical)	or NA
Positive FDR	or NA
Local FDR	or NA
Cluster-level FDR	or NA
None/Uncorrected	or NA
If permutation or Monte Carlo, report the number of	or NA
permutations/samples. If Monte Carlo, note the brain mask	
and smoothness used, and how smoothness was estimated	
If correction is limited to a small volume, the method for	or NA
selecting the region should be stated explicitly	
If no formal multiple comparisons method is used, the	or NA
inference must be explicitly labeled "uncorrected."	

Statistical inference	Checklist
If FWE found by random field theory list the smoothness in	or NA
mm FWHM and the RESEL count	
If FWE found by simulation (e.g., AFNI AlphaSim), provide	or NA
details of parameters for simulation	
If not a standard method, specify the method for finding	or NA
significance (e.g., "Custom in-lab software was used to	
construct statistic maps and thresholded at FDR < $0.05$	
(Benjamini and Hochberg, 1995)"	
If cluster-wise significance, state cluster-defining threshold	or NA
(e.g., P = 0.001)	
False negative discussion	
Any discussion of failure to reject the null hypothesis (e.g.,	or NA
lack of activation in a particular region) should be	
accompanied by SNR or effect size of the actually observed	
effect (allows reader to infer power to estimate an effect)	

#### Functional Connectivity

Functional Connectivity	Checklist
Confound adjustment & filtering Report:	
Method for detecting movement artifacts, movement-related	or NA
variation, and remediation (e.g. 'scrubbing', 'despiking', etc)	
Use of global signal regression, exact type of global signal	
used and how it	
was computed	or NA
Whether a high or lowpass temporal filtering is applied to	or NA
data, and at which point in the analysis pipeline. Note, any	
temporal regression model using filtered data should have its	
regressors likewise filtered	
Multivariate method: Independent Component Analysis	
Report:	
Algorithm to estimate components	or NA

Functional Connectivity	Checklist
Number of components (if fixed), or algorithm for estimating	
number of	
components	or NA
If used, method to synthesize multiple runs	or NA
Sorting method of IC's, if any	or NA
Detailed description of how components were chosen for	or NA
further analysis	
Dependent variable definition	
For seed-based analyses report:	
Definition of the seed region(s)	or NA
Rationale for choosing these regions	or NA
For region-based analyses report:	
Number of ROIs	or NA
How the ROI's are defined (e.g. citable anatomical atlas;	
auxiliary fMRI	
experiments); note if ROIs overlap	or NA
Assignment of signals to regions (i.e. how a time series is	
obtained from	
each region, e.g. averaging or first singular vector)	or NA
Note if considering only bilateral (L+R) merged regions	or NA
Note if considering only interhemispheric homotopic	or NA
connectivity	
Functional connectivity measure/model Report:	
Measure of dependence used, e.g. Pearson's (full) correlation,	
partial correlation, mutual information, etc; also specify:	
Use of Fisher's Z-transform (Yes/No) and, if standardised,	or NA
effective N is used to compute standard error (to account for	
any filtering operations on the data)	
Estimator used for partial correlation	or NA
Estimator used for mutual information	or NA
Regression model used to remove confounding effects (Pearson	or NA
or partial correlation)	

	Functional Connectivity	Checklist
	Effectivity connectivity Report:	_
	Model	or NA
	Algorithm used to fit model	or NA
	If per subject model, method used to generalize inferences to	or NA
	population. Itemize models considered, and method used for	
	model comparison	
	Graph analysis	
	Report the 'dependent variable' and 'functional connectivity	
	measure' used (see above). Specify either:	
	Weighted graph analysis or	or NA
	Binarized graph analysis is used, clarifying the method used	or NA
	for thresholding (e.g. a $10\%$ density threshold, or a	
	statistically -defined threshold); consider the sensitivity of	
	your findings to the particular choice of threshold used	
	Itemise the graph summaries used (e.g. clustering coefficient,	or NA
	efficiency, etc), whether these are global or per-node/per-edge	
	summaries. In particular with fMRI or EEG, clarify if	
	measures applied to individual subject networks or group	
	networks	
Follow-up	Enter your regnered have	
_	Enter your response here.	
Analyses		
Exploratory	Enter your response here.	
Analyses		
	References	
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