## nature portfolio

Corresponding author(s):	Thilo van Eimeren
Last updated by author(s):	12.09.2023

## **Reporting Summary**

Data will be made available upon reasonable request.

Nature Portfolio wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Portfolio policies, see our <u>Editorial Policies</u> and the <u>Editorial Policy Checklist</u>.

Statistics					
For all statistical a	nalyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.				
n/a Confirmed					
The exact	e exact sample size $(n)$ for each experimental group/condition, given as a discrete number and unit of measurement				
A statem	A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly				
The statis	The statistical test(s) used AND whether they are one- or two-sided Only common tests should be described solely by name; describe more complex techniques in the Methods section.				
A descrip	A description of all covariates tested				
A descrip	A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons				
A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)					
For null hypothesis testing, the test statistic (e.g. <i>F</i> , <i>t</i> , <i>r</i> ) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted Give <i>P</i> values as exact values whenever suitable.					
For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings					
For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes					
$\square$ Estimates of effect sizes (e.g. Cohen's $d$ , Pearson's $r$ ), indicating how they were calculated					
	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.				
Software an	d code				
Policy information	about <u>availability of computer code</u>				
Data collection	No commercial , open source or custom code was used to collect data in this study.				
Data analysis	There was no commercial, open source or custom code generated for data analysis. SPM12 and SPSS Version 28.0.1.0 was used.				
· ·	g custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and encourage code deposition in a community repository (e.g. GitHub). See the Nature Portfolio <u>guidelines for submitting code &amp; software</u> for further information.				
Data					
All manuscripts n - Accession code - A description c	about <u>availability of data</u> nust include a <u>data availability statement</u> . This statement should provide the following information, where applicable: es, unique identifiers, or web links for publicly available datasets of any restrictions on data availability assets or third party data, please ensure that the statement adheres to our policy				

Research involving human participants, their data, or biological material			
Policy information about studies with			

## Magnetic resonance imaging

## Experimental design

Design type N/A

Design specifications	N/A			
Behavioral performance measure	es N/A			
Acquisition				
Imaging type(s)	Structural			
Field strength	ЗТ			
Sequence & imaging parameters	T1-weighted MPRAGE acquisition parameters were as follows: repetition time = $2300  \text{ms}$ , echo time = $2.32  \text{ms}$ , flip angle = $8$ , field of view = $230  \text{mm}$ , slice thickness = $0.9  \text{mm}$ , voxel size = $0.9  \text{x}  0.9  \text{x}  0.9  \text{mm}$ , number of slices = $192  \text{ms}$ .			
Area of acquisition	Whole Brain			
Diffusion MRI Used	Not used     Not used			
Preprocessing				
Preprocessing software	SPM 12, CAT 12			
	T1 images are normalized to a template space and segmented into gray matter (GM), white matter (WM), and cerebrospinal fluid (CSF).			
Normalization template	TPM template of SPM			
Noise and artifact removal	N/A			
Volume censoring	N/A			
Statistical modeling & infere	nce			
Model type and settings	Multiple linear regression			
	Define precise effect in terms of the task or stimulus conditions instead of psychological concepts and indicate whether ANOVA or factorial designs were used.			
Specify type of analysis: Whole brain ROI-based Both				
Statistic type for inference	cluster wise inference p<0.05			
(See Eklund et al. 2016)				
Correction	FWE			
Models & analysis				
n/a   Involved in the study				
Functional and/or effective connectivity				
Graph analysis				
Multivariate modeling or pr	redictive analysis			