Supplementary Materials

Semantic integration demands modulate large-scale network interactions in the brain

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Supplementary Methods



Figure S1: *Age distribution of participants.* The density curve (red) represents the estimated probability density function of the age distribution.

Table S1. Demographic and neuropsychological test characteristics

	Mean values of raw scores (SD)
Demographics	
Age (years)	55 (12.77)
Sex (F:M)	15:17
Education (years)	16 (2.44)
Laterality Quotient	91.48 (9)
Neuropsychological Test	
MMSE	29.281 (1.05)
BDI-II	3.594 (3.38)
Digit span forward	6.5 (1.14)
Digit span backward	5.188 (0.99)
TAP (sum score)	193.84

Abbreviations: BDI-II: Beck Depression Inventory; F: Female; M: Male; MMSE: Mini Mental State Examination; SD: Standard Deviation; TAP: Test of Attentional Performance.

Equation S1. Linear mixed-effects model for analysis of reaction times

log (Reaction time) = $\beta 0 + \beta 1$ condition + $\beta 2$ age + $\beta 3$ education + $\beta 4$ condition x age + (1|subject) + (1|stimulus) + ϵ

Equation S2. Generalized linear mixed-effects logistic regression for analysis of accuracy

Error rate = β 0 + β 1condition + β 2age + β 3education + (1|subject) + (1|stimulus) + ϵ

Equation S3. Linear mixed-effects model for analysis of cPPI connectivity effects on reaction times

log (Reaction time) = $\beta 0 + \beta 1$ network measure + $\beta 2$ condition + $\beta 3$ age + βe ducation + $\beta 4$ network measure x condition + (1|subject) + ϵ

Equation S4. Generalized linear mixed-effects logistic regression for analysis of cPPI connectivity effects on accuracy

Error rate = $\beta 0 + \beta 1$ network measure + $\beta 2$ condition + $\beta 3$ age + $\beta 4$ network measure x condition + (1|subject) + ϵ

Note S1. Preprocessing of fMRI-data using fMRIPrep

Results included in this manuscript come from preprocessing performed using *fMRIPrep* 23.0.0 (Esteban, Markiewicz, et al. (2018); Esteban, Blair, et al. (2018); RRID:SCR_016216), which is based on *Nipype* 1.8.5 (K. Gorgolewski et al. (2011); K. J. Gorgolewski et al. (2018); RRID:SCR_002502)

Anatomical data preprocessing

A total of 1 T1-weighted (T1w) images were found within the input BIDS dataset.The T1-weighted (T1w) image was corrected for intensity nonuniformity (INU) with N4BiasFieldCorrection (Tustison et al. 2010), distributed with ANTs 2.3.3 (Avants et al. 2008, RRID:SCR_004757), and used as T1w-reference throughout the workflow. The T1w-reference was then skull-stripped with a *Nipype* implementation of the antsBrainExtraction.sh workflow (from ANTs), using OASIS30ANTs as target template. Brain tissue segmentation of cerebrospinal fluid (CSF), white-matter (WM) and graymatter (GM) was performed on the brain-extracted T1w using fast (FSL 6.0.5.1:57b01774, RRID:SCR_002823, Zhang, Brady, and Smith 2001). Brain surfaces were reconstructed using recon-all (FreeSurfer 7.3.2, RRID:SCR_001847, Dale, Fischl, and Sereno 1999), and the brain mask estimated previously was refined with a custom variation of the method to reconcile ANTs-derived and FreeSurfer-derived segmentations of the cortical graymatter of Mindboggle (RRID:SCR_002438, Klein et al. 2017). Volume-based spatial normalization to two standard spaces (MNI152NLin6Asym, MNI152NLin2009cAsym) was performed through nonlinear registration with antsRegistration (ANTs 2.3.3), using brain-extracted versions of both T1w reference and the T1w template. The following templates were selected for spatial normalization: *FSL's MNI ICBM 152 non-linear 6th Generation Asymmetric Average Brain Stereotaxic Registration Model* [Evans et al. (2012), RRID:SCR_002823; TemplateFlow ID: MNI152NLin2009cAsym]. *ICBM 152 Nonlinear Asymmetrical template version 2009c* [Fonov et al. (2009), RRID:SCR_008796; TemplateFlow ID: MNI152NLin2009cAsym].

Functional data preprocessing

For each of the 5 BOLD runs found per subject (across all tasks and sessions), the following preprocessing was performed. First, a reference volume and its skull-stripped version were generated using a custom methodology of fMRIPrep. Head-motion parameters with respect to the BOLD reference (transformation matrices, and six corresponding rotation and translation parameters) are estimated before any spatiotemporal filtering using mcflirt (FSL 6.0.5.1:57b01774, Jenkinson et al. 2002). BOLD runs were slice-time corrected to 0.962s (0.5 of slice acquisition range 0s-1.92s) using 3dTshift from AFNI (Cox and Hyde 1997, RRID:SCR 005927). The BOLD time-series (including slice-timing correction when applied) were resampled onto their original, native space by applying the transforms to correct for head-motion. These resampled BOLD time-series will be referred to as preprocessed BOLD in original space, or just preprocessed BOLD. The BOLD reference was then co-registered to the T1w reference using bbregister (FreeSurfer) which implements boundary-based registration (Greve and Fischl 2009). Co-registration was configured with six degrees of freedom. Several confounding time-series were calculated based on the preprocessed BOLD: framewise displacement (FD), DVARS and three region-wise global signals. FD was computed using two formulations following Power (absolute sum of relative motions, Power et al. (2014)) and Jenkinson (relative root mean square displacement between affines, Jenkinson et al. (2002)). FD and DVARS are calculated for each functional run, both using their im plementations in Nipype (following the definitions by Power et al. 2014). The three global signals are extracted within the CSF, the WM, and the whole-brain masks. Additionally, a set of physiological regressors were extracted to allow for component-based noise correction (CompCor, Behzadi et al. 2007). Principal components are estimated after high-pass filtering the preprocessed BOLD time-series (using a discrete cosine filter with 128s cut-off) for the two CompCor variants: temporal (tCompCor) and anatomical (aCompCor). tCompCor components are then calculated from the top 2% variable voxels within the brain mask. For aCompCor, three probabilistic masks (CSF, WM and combined CSF+WM) are generated in anatomical space. The implementation differs from that of Behzadi et al. in that instead of eroding the masks by 2 pixels on BOLD space, a mask of pixels that likely contain a volume fraction of GM is subtracted from the aCompCor masks. This mask is obtained by dilating a GM mask extracted from the FreeSurfer's aseg segmentation, and it ensures components are not extracted from voxels containing a minimal fraction of GM. Finally, these masks are resampled into BOLD space and binarized by thresholding at 0.99 (as in the original implementation). Components are also calculated separately within the WM and CSF masks. For each CompCor decomposition, the k components with the largest singular values are retained, such that the retained components' time series are sufficient to explain 50 percent of variance across the nuisance mask (CSF, WM, combined, or temporal). The remaining components are dropped from consideration. The head-motion estimates calculated in the correction step were also placed within the corresponding confounds file. The confound time series derived from head motion estimates and global signals were expanded with the inclusion of temporal derivatives and quadratic terms for each (Satterthwaite et al. 2013). Frames that exceeded a threshold of 0.7 mm FD or 1.5 standardized DVARS were annotated as motion outliers. Additional nuisance timeseries are calculated by means of principal components analysis of the signal found within a thin band (crown) of voxels around the edge of the brain, as proposed by (Patriat, Reynolds, and Birn 2017). The BOLD time-series were resampled into standard space, generating a preprocessed BOLD run in MNI152NLin6Asym space. First, a reference volume and its skull-stripped version were generated using a custom methodology of fMRIPrep. All resamplings can be performed with a single interpolation step by composing all the pertinent transformations (i.e. head-motion transform matrices, susceptibility distortion correction when available, and co-registrations to anatomical and output spaces). Gridded (volumetric) resamplings were performed using antsApplyTransforms (ANTs), configured with Lanczos interpolation to minimize the smoothing effects of other kernels (Lanczos 1964). Non-gridded (surface) resamplings were performed using mri_vol2surf (FreeSurfer).

Many internal operations of *fMRIPrep* use *Nilearn* 0.9.1 (Abraham et al. 2014, RRID:SCR_001362), mostly within the functional processing workflow. For more details of the pipeline, see the section corresponding to workflows in *fMRIPrep*'s documentation.

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Errors

No errors to report!



Figure S2. Conjunction of the contrasts "unexpected sentence endings > expected sentence endings \cap anomalous sentence endings > expected sentence endings". Results are FWE-corrected at p < 0.05 at the cluster level.

Table S2. Coordinates of seed regions for gPPI analysis

ROI	Region	Side	MNI	MNI Coordinates (in mm)						
			х	У	Z					
1	Inferior frontal gyrus (pars triangularis)	L	-54	36	3					
2	Presupplementary motor area	L	-6	12	51					
3	Cerebellum VI	R	27	-66	-27					
4	Frontal pole / orbital gyrus	L	36	36	-15					
5	Frontal pole / inferior frontal gyrus	R	48	36	15					

Note: Seed regions were obtained from the conjunction of the contrasts "anomalous > expected" and "unexpected > expected" sentence endings. **Abbreviation:** ROI: Region of Interest.

Supplementary Results

	Lo	Error rat	e			
Predictors	Estimate	CI	Р	Odds	CI	Р
	5			Ratios		
(Intercept)	6.94	6.90 - 6.97	<0.001	0.11	0.08 - 0.17	<0.001
condition [unexpected]	0.09	0.06 - 0.12	<0.001			
condition [anomalous]	0.24	0.21 - 0.27	<0.001			
condition [pseudoword]	0.26	0.24 – 0.29	<0.001			
age (centered)	0.00	-0.00 - 0.00	0.222	1.02	1.01 - 1.04	0.002
education (centered)	0.00	-0.01 - 0.02	0.736	0.98	0.90 - 1.07	0.645
condition [unexpected] ×	0.00	0.00 - 0.00	0.002			
age (centered)						
condition [anomalous] ×	0.00	0.00 - 0.00	0.002			
age (centered)						
condition [pseudoword] ×	-0.00	-0.00 - 0.00	0.843			
age (centered)						
condition [expected]				0.09	0.05 – 0.17	<0.001
condition [pseudoword]				0.35	0.22 – 0.56	<0.001
condition [unexpected]				0.13	0.07 – 0.24	<0.001
Random Effects						
σ²	0.02			3.29		
τ ₀₀	0.01 stimulus			1.72 _{stim}	ulus	
	0.01 subject			0.27 _{subj}	ect	
ICC	0.46			0.38		
Ν	32 subject			32 subject		
	300 stimulus			300 stimu	llus	
Observations	8406			9565		
Marginal R ² / Conditional R ²	0.255 / 0.595			0.132 /	0.460	

Note: Significant effects are marked in bold. Contrasts are treatment coded. P-values were obtained via likelihood ratio tests. **Abbreviation:** CI: Confidence interval.



Figure S3. Effect of age on behavioral data.



Figure S4. Group-level results from univariate fMRI-analysis. Results are FWE-corrected at p < 0.05 at the cluster level. No significant results were found for the contrast "pseudoword > anomalous".

Region	Side	MN	Coordinates (in	mm)	t	k
J. J		x	y `	z		
Anomalous > expected			_			
Inferior frontal gyrus (p. op.)	L	-57	21	24	9.15	1387
Presupplementary motor area	L	-6	21	45	9.06	659
Frontal orbital cortex / insula	R	33	27	0	7.96	820
Cerebellum VI	L	-27	-66	-27	7.78	265
Superior temporal gyrus	L	-60	-30	6	6.59	208
Cerebellum crus II	R	12	-81	-30	6.07	219
Anomalous > pseudoword						
Frontal pole	L	-39	42	-15	6.16	58
Angular gyrus	L	-48	-54	24	5.88	473
Superior frontal gyrus	L	-9	27	63	5.32	295
Cerebellum Crus I	R	24	48	-24	5.08	88
Superior frontal gyrus	L	-9	-60	33	4.60	50
Anomalous > unexpected						
Insula cortex	R	33	24	-3	5.74	96
Presupplementary motor area	R	6	21	45	5.50	130
Inferior parietal lobe	R	51	-42	45	5.00	82
Inferior frontal gyrus	R	45	12	21	4.96	87
Precentral gyrus	L	-39	3	33	4.88	57
Frontal operculum	L	-42	21	6	4.82	66
Expected > anomalous						
Precuneous cortex	R	9	-57	24	5.78	189
Inferior parietal lobe	R	42	-54	24	4.95	75
Expected > pseudoword						
Inferior parietal lobe	L	-42	-78	33	8.20	3387
Middle temporal gyrus	R	60	-9	-18	6.86	187
Middle frontal gyrus	L	-39	21	51	6.65	357
Temporal pole	L	-51	6	-24	6.09	196
Frontal pole	L	-12	63	21	5.94	414
Middle frontal gyrus	R	21	57	39	5.75	124
Expected > unexpected						
Angular gyrus	R	63	-48	36	6.26	286
Supramarginal gyrus	L	-54	-48	45	4.72	45
Pseudoword > expected			-			
Inferior frontal gyrus (p. op.)	L	-42	6	24	9.21	1163
Frontal orbital cortex	R	33	27	3	7.52	193
Superior temporal gyrus	ĸ	63	-3	-3	7.41	89
Presupplementary motor area	L	-6	12	54	6.65	414
wilddie frontal gyrus	К	51	36	24	6.55	311
Pseudoword > unexpected	D	20	20	0	F 10	C A
Providal orbital cortex	К D	59	30	40	5.1ð 4.70	04
Faracinguidle gyrus	ĸ	D	24	42	4.72	44
Planum tomporala		62	71	E	E 00	70
Frantal operation	D	-03	-21	2	5.89 77	62
riontai operculuin	n	50	27	Э	4.//	02

Table S4. Group-level activation peaks from univariate fMRI-analysis

Unexpected > anomalous						
Inferior parietal lobe	L	-54	-63	18	7.19	311
Lateral occipital cortex	R	51	-60	21	5.91	150
Precuneous cortex	L	-6	-57	21	5.69	197
Frontal medial cortex	R	3	45	-15	5.26	130
Unexpected > expected						
Inferior frontal gyrus (p. op.)	L	-39	6	24	7.65	1309
Presupplementary motor area	L	-3	12	48	5.88	237
Precentral gyrus	R	48	9	27	5.16	89
Frontal orbital cortex	R	42	33	-12	5.07	76
Cerebellum VI	R	30	-66	-24	5.04	73
Inferior temporal gyrus	L	-54	-57	-12	4.11	42
Unexpected > pseudoword						
Inferior parietal lobe	L	-45	-60	18	8.70	2424
Frontal pole	L	-15	48	39	8.34	1211
Middle temporal gyrus	L	-57	-9	-21	6.63	502
Middle temporal gyrus	R	57	-9	-15	6.25	186
Inferior parietal lobe	R	57	-60	18	5.84	456
Middle temporal gyrus	L	-63	-42	-3	5.63	113
Parahippocampal gyrus	R	30	-33	-18	5.23	107
Word > pseudoword						
Frontal pole	L	-15	48	39	11.13	1098
Inferior parietal lobe	L	-45	-63	21	9.41	3410
Middle temporal gyrus	L	-57	-9	-21	7.40	335
Parahippocampal gyrus	L	-30	-27	-25	6.35	122
Middle temporal gyrus	R	57	0	-24	6.15	215

Note: Cluster corrected at FWE p < 0.05 with a voxel-wise threshold at p < 0.001. Clusters represent local maxima. Cluster size (k) is given in mm³. No significant cluster were found for the contrast "sentences with pseudoword endings > anomalous endings". **Abbreviation:** P. op.: Pars opercularis.

	IC04	IC11	IC12	IC13	IC16	IC17	IC18	IC21	IC25	IC33	IC35
ContA	0.002	0.024	0.050	0.031	0.011	0.075	0.019	0.001	0.241	0.020	0.002
ContB	0.001	0.022	0.033	0.012	0.006	0.159	0.004	0.005	0.064	0.033	0.050
ConC	0.010	0.075	0.033	0.025	0.000	0.087	0.000	0.019	0.000	0.001	0.003
DefaultA	0.011	0.225	0.005	0.015	0.000	0.082	0.013	0.007	0.006	0.015	0.142
DefaultB	0.026	0.023	0.034	0.014	0.005	0.048	0.004	0.007	0.081	0.078	0.273
DefaultC	0.000	0.059	0.000	0.017	0.014	0.002	0.016	0.045	0.000	0.002	0.003
DorsAttnA	0.000	0.056	0.083	0.040	0.035	0.013	0.112	0.051	0.064	0.000	0.002
DorsAttnB	0.004	0.018	0.154	0.079	0.010	0.005	0.016	0.001	0.013	0.001	0.002
LimbicA	0.001	0.000	0.000	0.001	0.001	0.001	0.000	0.001	0.000	0.002	0.004
LimbicB	0.000	0.020	0.000	0.000	0.000	0.004	0.003	0.000	0.001	0.002	0.006
SalVentAttnA	0.087	0.040	0.063	0.240	0.000	0.006	0.017	0.001	0.011	0.022	0.007
SalVentAttnB	0.008	0.004	0.009	0.067	0.000	0.037	0.007	0.007	0.029	0.118	0.020
SomMotA	0.041	0.038	0.274	0.023	0.014	0.002	0.001	0.000	0.000	0.000	0.030
SomMotB	0.288	0.041	0.003	0.080	0.001	0.000	0.017	0.002	0.010	0.001	0.017
TempPar	0.156	0.015	0.000	0.017	0.003	0.020	0.094	0.000	0.043	0.028	0.024
VisCent	0.002	0.004	0.001	0.005	0.027	0.000	0.197	0.185	0.002	0.018	0.012
VisPeri	0.000	0.051	0.000	0.019	0.001	0.004	0.001	0.286	0.000	0.122	0.003

Table S5. Jaccard indices for independent components and cognitive networks (Yeo et al., 2011)

Note: The selected network labels for the respective independent components (ICs) are shown in bold, whereas all cognitive networks that showed a higher Similarity Coefficient than J = 0.15 are shown in italics. No Similarity Coefficient over J = 0.15 was found for IC16 and for IC33.

	expected	expected vs. rest unexpected vs. rest		anomalou	s vs. rest	pseudowo	pseudoword vs. rest		unexpected vs.		anomalous vs.		pseudoword vs.	
									expe	expected		ected	expected	
	t-values	p-values	t-values	p-values	t-values	p-values	t-values	p-values	t-values	p-values	t-values	p-values	t-values	p-values
IC04	8.548	0.000	8.498	0.000	8.006	0.000	8.565	0.000	-0.891	0.576	0.655	0.790	2.126	0.114
IC11	-4.832	0.000	-4.719	0.000	-5.127	0.000	-6.961	0.000	-0.348	0.803	-1.718	0.296	-3.597	0.004
IC12	3.998	0.001	4.213	0.001	3.712	0.002	3.174	0.009	0.823	0.576	-0.464	0.790	-1.176	0.431
IC13	0.756	0.556	1.671	0.192	0.761	0.453	1.320	0.216	-1.377	0.490	-0.078	0.998	1.059	0.431
IC16	1.752	0.226	1.240	0.287	2.704	0.024	1.749	0.142	1.153	0.567	1.537	0.296	0.173	0.863
IC17	-1.338	0.262	-3.972	0.001	-1.503	0.197	-1.875	0.129	3.023	0.055	-0.553	0.790	-1.025	0.431
IC18	-1.584	0.226	-1.187	0.287	-2.227	0.061	-1.583	0.151	-0.819	0.576	-1.570	0.296	-0.191	0.863
IC21	0.469	0.706	0.491	0.627	0.846	0.444	0.169	0.867	-0.077	0.939	0.632	0.790	-0.584	0.689
IC25	1.353	0.262	3.265	0.006	4.662	0.000	4.626	0.000	-2.419	0.119	5.084	0.000	4.037	0.004
IC33	0.096	0.924	1.146	0.287	1.110	0.337	1.667	0.145	-1.766	0.320	1.826	0.296	3.615	0.004
IC35	-1.656	0.226	-1.350	0.287	-1.510	0.197	-2.929	0.014	-0.602	0.674	-0.002	0.998	-1.976	0.126

Table S6. Statistical results for domain-specific network activity

Note: P-values are FDR-corrected at *p* < 0.05. Bold values indicate significant comparisons in the two-sided t-tests.

Α



В

Conta Conta Conta SomMota Defaulta Defaulta Cerebellum Defaulta



pseudoword vs. expected





Figure S5. Functional coupling between task-relevant networks. A. Chord diagrams show (additional) significant results of functional coupling between ICA-derived networks for the contrasts "words vs. pseudowords" and "unexpected vs. anomalous". Connectivity values are partial correlations. The color

intensity and width of a connection indicate its correlational strength. Higher correlation values (r) indicate positive coupling and negative values indicate decoupling between networks. B. Heatmaps show correlation matrices of functional coupling between ICA-derived networks. The color intensity indicates correlational strength between two networks. Significant correlations are highlighted by black boxes. Higher values indicate positive coupling and negative values indicate decoupling between networks. Abbreviations: ContA/B: Control network A/B; DefaultA/B: Default mode network A/B; SomMotA/B: Somatomotor network A/B.

Table S7. CPPI connectivity effects on response time contrasting unexpected vs. expected sentence endings

	Log (Reacti	ion times)	
Predictors	Estimates	CI	р
(Intercept)	6.72	6.41 - 7.03	<0.001
DefaultASomMotA	-0.03	-0.07 - 0.02	0.217
condition [unexpected]	-0.09	-0.100.07	<0.001
age	0.00	-0.00 - 0.00	0.426
education	0.00	-0.02 - 0.02	0.790
DefaultASomMotA ×	0.02	0.01 - 0.03	0.001
condition [unexpected]			
Random Effects			
σ²	0.03		
τ _{00 subject}	0.01		
ICC	0.31		
N subject	32		
Observations	3673		
Marginal R ² / Conditional R ²	0.064 / 0.3	57	

Note: Significant effects are marked in bold. Contrasts are treatment coded. P-values were obtained via likelihood ratio tests. **Abbreviations:** CI: Confidence interval; DefaultA: Default mode network A; SomMotA: Somatomotor network A.

	Log	(Reaction time	s)	Log	(Reaction time	s)		
Predictors	Estimates	CI	р	Estimates	CI	р		
(Intercept)	6.83	6.51 – 7.14	<0.001	6.81	6.48 – 7.13	<0.001		
SomMotBContA	-0.00	-0.04 - 0.04	0.999					
condition [anomalous]	-0.23	-0.240.22	<0.001	-0.23	-0.24 – -0.22	<0.001		
age	0.00	-0.00 - 0.00	0.370	0.00	-0.00 - 0.00	0.412		
education	0.00	-0.02 - 0.02	0.993	0.00	-0.02 - 0.02	0.863		
SomMotBContA ×	-0.02	-0.030.01	0.002					
condition [anomalous]								
SomMotBDefaultB				-0.01	-0.06 - 0.04	0.696		
SomMotB DefaultB ×				0.02	0.01 - 0.03	0.001		
condition [anomalous]								
Random Effects								
σ ²	0.03			0.03				
τ ₀₀	0.01 subject			0.01 subject				
ICC	0.31			0.31				
Ν	32 subject			32 subject				
Observations	3654			3654				
Marginal R ² / Conditional R ²	0.232 / 0.4	73		0.233 / 0.473				

Table S8. CPPI connectivity effects on response time contrasting anomalous vs. expected sentence endings

Note: Significant effects are marked in bold. Contrasts are treatment coded. P-values were obtained via likelihood ratio tests. **Abbreviations:** CI: Confidence interval; ContA: Control network A; DefaultB: Default mode network B; SomMotB: Somatomotor network B.

Table S9. CPPI connectivity effects on response time contrasting pseudoword vs. expected sentence endings

	Log	g (Reaction times)	1	Log	g (Reaction times)		Log	(Reaction times)		Log ((Reaction times)		Log	(Reaction times)	1			Log (Reaction times)
Predictors	Estimates	CI	р	Estimates	CI	р	Estimates	CI	р	Estimates	CI	р	Estimates	CI	р	Estimates	CI	p
(Intercept)	6.78	6.48 - 7.08	<0.001	6.78	6.50 - 7.07	<0.001	6.81	6.52 - 7.09	<0.001	6.76	6.45 - 7.06	<0.001	6.79	6.50 - 7.07	<0.001	6.86	6.58 - 7.13	<0.001
DefaultAContA	0.00	-0.04 - 0.04	0.848															
condition [pseudoword]	0.27	0.26 - 0.28	<0.001	0.27	0.26 - 0.28	<0.001	0.27	0.26-0.28	<0.001	0.27	0.26-0.28	<0.001	0.27	0.26-0.28	<0.001	0.27	0.26-0.28	<0.001
age	0.00	-0.00 - 0.00	0.447	0.00	-0.00 - 0.00	0.595	0.00	-0.00 - 0.00	0.498	0.00	-0.00 - 0.00	0.404	0.00	-0.00 - 0.00	0.579	0.00	-0.00 - 0.00	0.904
education	0.01	-0.01 - 0.02	0.541	0.01	-0.01 - 0.02	0.449	0.00	-0.01 - 0.02	0.633	0.01	-0.01 - 0.02	0.493	0.01	-0.01 - 0.02	0.486	0.00	-0.01 - 0.02	0.633
DefaultAContA ×	0.03	0.02 - 0.04	<0.001															
condition [pseudoword]																		
SomMotACerebellum				-0.03	-0.07 - 0.01	0.145												
SomMotACerebellum ×				-0.02	-0.030.01	<0.001												
condition [pseudoword]																		
ContBDefaultB							0.03	-0.01 - 0.07	0.106									
ContBDefaultB ×							-0.02	-0.030.01	0.001									
condition [pseudoword]																		
SomMotBContB										-0.02	-0.06 - 0.03	0.442						
SomMotBContB ×										0.01	0.01 - 0.02	0.003						
condition [pseudoword]																		
SomMotBDefaultA													-0.03	-0.07 - 0.01	0.118			
SomMotBDefaultA ×													0.01	0.00 - 0.02	0.007			
condition [pseudoword]																		
SomMotADefaultB																0.05	0.01 - 0.09	0.009
SomMotADefaultB ×																-0.01	-0.020.00	0.021
condition [pseudoword]																		
Random Effects																		
σ²	0.03			0.03			0.03			0.03			0.03			0.03		
τ	0.01 subject			0.01 subject			0.01 subject			0.01 subject			0.01 subject			0.01 subject		
ICC	0.31			0.29			0.30			0.31			0.30			0.27		
N	32 subject			32 subject			32 subject			32 subject			32 subject			32 subject		
Observations	5494			5494			5494			5494			5494			5494		
Marginal R ² / Conditional R ²	0.279 / 0.505	5		0.293 / 0.502	L		0.290 / 0.501			0.278 / 0.502			0.290 / 0.501			0.314 / 0.499		

Note: Significant effects are marked in bold. Contrasts are treatment coded. P-values were obtained via likelihood ratio tests. **Abbreviations:** CI: Confidence interval; ContA/B: Control network A/B; DefaultA/B: DefaultA/B: Default mode network A/B; SomMotA/B: Somatomotor network A/B.