

1 **Connectivity and functional diversity of different temporo-occipital nodes for**  
2 **action perception**

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11 **ABBREVIATED TITLE:** Temporo-occipital nodes for action perception

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29 **Abstract**

30 The temporo-occipital cortex (TOC) plays a key role in body and action perception, but current  
31 understanding of its functions is still limited. TOC body regions are heterogeneous and their role  
32 in action perception is poorly understood. This study adopted data-driven approaches to region  
33 selectivity and investigated the connectivity of TOC nodes and the functional network sensitivity  
34 for different whole body action videos. In two human 7T fMRI experiments using independent  
35 component analysis, four adjacent body selective nodes were detected within the TOC network  
36 with distinct connectivity profiles and functional roles. Action type independent connectivity was  
37 observed for the posterior-ventral node to the visual cortex, the posterior-dorsal node to the  
38 precuneus and the anterior nodes to the frontal cortex. Action specific connectivity modulations  
39 were found in middle frontal gyrus for the aggressive condition with increased connectivity to  
40 the anterior node and decreased connectivity to the posterior-dorsal node. But for the defensive  
41 condition, node-nonspecific enhancement was found for the TOC-cingulate connectivity. By  
42 addressing the issue of multiple nodes in the temporo-occipital network we show a functional  
43 dissociation of different body selective centres related to the action type and a potential hierarchy  
44 within the TOC body network.

## 45 **1. Introduction**

46 During social interactions, intentions, actions, and emotions are routinely read from nonverbal  
47 communication signals provided by faces, body postures and whole-body movements as well-  
48 documented in studies of humans (Argyle, 1976; de Gelder et al., 2010) and non-human primates  
49 (Vogels, 2022). Despite its importance, the neural basis of whole-body perception is still much  
50 less understood than that of faces (Deen et al., 2023).

51 Recent studies have revealed several brain areas in human and non-human primates. In humans,  
52 the extrastriate body area (EBA) (Downing & Kanwisher, 2001) was the first one to be reported.  
53 It is a region located in the extrastriate cortex that overlaps with other category-selective areas,  
54 such as those dedicated to processing motion (Weiner & Grill-Spector, 2011), tools, and even  
55 action related words (Lingnau & Downing, 2015). Subsequent studies have identified at least  
56 three different body-selective clusters within the extrastriate cortex (Weiner & Grill-Spector,  
57 2011). This anatomical diversity is complemented by functional diversity, as these clusters also  
58 display varying patterns of connectivity (Zimmermann et al., 2018).

59 The actual contribution of these body selective areas is not well understood (de Gelder & Poyo  
60 Solanas, 2021; de Gelder et al., 2010; Vogels, 2022). A number of studies found that these body  
61 selective areas play a role in processing emotional expressions (de Gelder et al., 2004; Grèzes et  
62 al., 2007; Peelen et al., 2007; Pichon et al., 2008), biological movement (Jastorff & Orban, 2009),  
63 specific postural and kinematic features of the body (Marrazzo et al., 2023; Marrazzo et al., 2021;  
64 Poyo Solanas et al., 2020), action recognition (Goldberg et al., 2014; Shmuelof & Zohary, 2005),  
65 motor planning (Zimmermann et al., 2012), as well as social perception (Kret et al., 2011a;  
66 Moreau et al., 2023). Taken together, these findings suggest that body selectivity may be

67 understood not simply as a matter of category selectivity but as resulting from the activity of a  
68 sparsely distributed ensemble of brain areas (de Gelder & Poyo Solanas, 2021; Weiner & Grill-  
69 Spector, 2011). Mapping the broader network's activity may be an important step in uncovering  
70 how these different body-specific nodes collectively contribute to the perception of whole-body  
71 movements at the network level. Several network models have been proposed including the  
72 Action Observation Network (AON) (Caspers et al., 2010), the Default Mode Network (DMN)  
73 (Zhan et al., 2018) or a pathway for social perception (Haak & Beckmann, 2018). However, none  
74 of these directly addresses the specific case of perceiving emotional expressions, intentions, and  
75 actions of conspecifics routinely conveyed by body movements.

76 The goal of this study was to identify a dynamic whole-body network and to investigate how its  
77 network activity and its connectivity with other brain areas supports specific actions such as  
78 defensive or aggressive behaviour. Rather than following the traditional approach of contrast-  
79 based selection of body regions, we approach the question at the network level with data-driven  
80 methods. We used independent component analysis (ICA), which is widely applied in resting-  
81 state and task-based fMRI studies (Du et al., 2017; Jarrahi et al., 2015; Jung et al., 2020), to  
82 identify the body sensitive nodes within the TOC network and further tracked their whole-brain  
83 communications during whole body action processing.

## 84 **2. Method**

85 The study consisted of two experiments: a localizer experiment and the main experiment. First,  
86 we used the localizer experiment to identify the temporo-occipital network associated with body  
87 action perception. This was accomplished through a data-driven strategy based on our previous  
88 study (Li et al., 2023). Next, the data of the main experiment was employed to extract node

89 regions within this network and investigate their connectivity profiles as well as their modulation  
90 by affective body conditions. Nineteen participants took part in the experiment.

## 91 **2.1. Participants**

92 Nineteen healthy participants (mean age = 24.58 years; age range = 19-30 years; 6 males, all  
93 right-handed) took part in the experiment. All participants had a normal or correct-ed-to-normal  
94 vision and no medical history of any psychiatric or neurological disorders. All participants  
95 provided informed written consent before the start of the experiment and received a monetary  
96 reward (vouchers) or course credits for their participation. The experiment was approved by the  
97 Ethical Committee at Maastricht University and was performed in accordance with the  
98 Declaration of Helsinki.

## 99 **2.2. Experiment Design**

### 100 *2.2.1. Network localizer*

101 The functional localizer followed a blocked design with twelve categories of videos com-posed  
102 from three factors: (body / face / object) \* (human / monkey) \* (normal / scramble). Each  
103 category consisted of ten different 1000-ms videos which were presented block-wise in a random  
104 order. Within each block, the ten videos were interleaved by a fixed 500-ms inter-trial interval,  
105 while two consecutive blocks were interleaved by an inter-block interval jittered around 11  
106 seconds. The order of block conditions was randomized for each participant, and each condition  
107 was repeated six times within three runs. Each run contained a catch block where, in one of the  
108 trials, the fixation point changed its shape from a “+” to a “o”. Participants were instructed to  
109 make a button-press response when detecting the fixation shape change. The total length of each

110 run was 735 seconds on average. A detailed description of the localizer stimuli and design can be  
111 found in Li et al. (2023).

### 112 2.2.2. *Main experiment*

113 For the main experiment, we applied a mixed block/event-related design (Visscher et al., 2003)  
114 consisting of five conditions of videos: three human body conditions (aggressive, defensive, and  
115 neutral), one neutral human face condition and one neutral object condition (**Figure 1**). Each  
116 condition consisted of ten different 1000-ms videos. The body and face videos were chosen from  
117 the same stimulus set as described in Kret et al. (2011b). The object videos were selected from  
118 the same set as used in the localizer experiment but differed from the ones used in that  
119 experiment.

120 During the experiment, each condition was presented block-wise with a jittered inter-trial  
121 interval of around 3 seconds and an inter-block interval of 12 seconds. For each trial, the video  
122 was centered and presented on a uniform gray background. The sizes of stimuli were 3.5\*7.5  
123 degrees of visual angle for bodies and objects, and 3.5\*3.5 degrees of visual angle for faces. The  
124 order of block conditions was randomized for each participant, and each condition was repeated  
125 ten times within five runs. Two extra blocks with a catch trial were inserted in each run where  
126 participants were instructed to detect the fixation shape changes as described in the localizer  
127 experiment. The total length of each run was 480 seconds.

128 Both the main experiment and the localizer experiment were programmed using the  
129 Psychtoolbox (<https://www.psychtoolbox.net>) implemented in Matlab 2018b  
130 (<https://www.mathworks.com>). Stimuli were projected onto a screen at the end of the scanner  
131 bore with a Panasonic PT-EZ57OEL projector (screen size = 30 \* 18 cm, resolution = 1920 \*

132 1200 pixel). Participants viewed the stimuli through a mirror attached to the head coil (screen-to-  
133 eye distance = 99 cm, visual angle =  $17.23 \times 10.38$  degrees).

### 134 2.2.3. *fMRI data acquisition*

135 All images were acquired with a 7T MAGNETOM scanner at the Maastricht Brain Imaging  
136 Centre (MBIC) of Maastricht University, the Netherlands. Functional images were collected  
137 using the T2\*-weighted multi-band accelerated EPI 2D BOLD sequence (TR/TE = 1000/20 ms,  
138 multiband acceleration factor = 3, in-plane isotropic resolution = 1.6 mm, number of slices per  
139 volume = 68, matrix size =  $128 \times 128$ , volume number = 735 for the network localizer and 480  
140 for the main experiment). T1-weighted anatomical images were obtained using the 3D-  
141 MP2RAGE sequence (TR/TE = 5000/2.47 ms, Inverse time T11/I2 = 900/2750 ms, flip angle  
142 FA1/FA2 =  $5/3^\circ$ , in-plane isotropic resolution = 0.7 mm, matrix size =  $320 \times 320$ , slice number =  
143 240). Physiological parameters were recorded via pulse oximetry on the index finger of the left  
144 hand and with a respiratory belt.

### 145 2.2.4. *fMRI image preprocessing*

146 Anatomical and functional images were preprocessed using the Brainvoyager 22 (Goebel, 2012),  
147 and the Neuroelf toolbox in Matlab (<https://neuroelf.net/>). For anatomical images, brain  
148 extraction was conducted with INV2 images to correct for MP2RAGE background noise. The  
149 resolution was then downsampled to 0.8 mm for better alignment to the 1.6 mm resolution of  
150 functional images. For functional images, the preprocessing steps included EPI distortion  
151 correction (Bremner et al., 2020), slice scan time correction, 3D head-motion correction, and  
152 high-pass temporal filtering (GLM with Fourier basis set of 3 cycles, including linear trend).  
153 Coregistration was first conducted between the anatomical image and its most adjacent

154 functional run using a boundary-based registration (BBR) algorithm (Greve & Fischl, 2009), and  
155 all the other functional runs were coregistered to the aligned run. Individual images were  
156 normalized to Talairach space (Collins, Neelin, Peters, & Evans, 1994) with 3 mm Gaussian  
157 spatial smoothing. Trilinear/sinc interpolation was used in the motion correction step, and sinc  
158 interpolation was used in all the other steps.

159 Physiological parameters were collected as confound factors for the functional imaging data. The  
160 physiological data were preprocessed using the RETROspective Image CORrection  
161 (RETROICOR; Glover et al., 2000; Harvey et al., 2008) pipeline, which uses Fourier expansions  
162 of different orders for the phase of cardiac pulsation (3rd order), respiration (4th order) and  
163 cardio-respiratory interaction (1st order). Eighteen physiological confound factors were finally  
164 created for each participant.

165 The anatomical labeling of the brain areas reported in this study was performed according to the  
166 Talairach Daemon (<http://www.talairach.org/daemon.html>) in combination with the Multilevel  
167 Human Brain Atlas (<https://ebrains.eu/service/human-brain-atlas>).

## 168 **2.3. Data analysis**

### 169 *2.3.1. Body network extraction*

170 The Infomax algorithm implemented in the Group ICA of fMRI Toolbox (GIFT; Calhoun et al.,  
171 2001) was used to identify body selective networks within the localizer experiment. This resulted  
172 in 75 spatial independent components. Individual ICs were back-reconstructed using the GIG-  
173 ICA algorithm from the aggregated group ICs (Du & Fan, 2013). The stability of group ICA was  
174 assessed by the ICASSO module implemented in the GIFT, which repeated the Infomax



175 decomposition 20 times and resulted in an index of stability (Iq) for each IC (Himberg et al.,  
176 2004). Prior to the group-ICA, physiological and motion confounds were regressed out from the  
177 preprocessed functional images. The resulting time courses were then transformed into  
178 percentages of signal change to enhance the ICA stability (Allen et al., 2011). Components  
179 showing large white matter / cerebrospinal fluid coverage were excluded from further analysis.

180 To identify body selective networks, we conducted a GLM on each reconstructed subject-level  
181 IC time course, which estimated the IC response for each condition. In the design matrix, each  
182 condition predictor was modeled as a boxcar function with the same duration of the block and  
183 convolved with the canonical HRF. The estimated betas were first averaged across all runs for  
184 each participant and were then used to calculate the contrast of [2 \* human body (normal -  
185 scramble) – (human face (normal - scramble) + human object (normal - scramble))]. Right-tailed  
186 t-tests and Benjamini-Hochberg multiple comparison corrections were conducted at the group  
187 level to find significant body sensitivity.

188 To define the group-level coverage of the IC networks, the individual IC maps were normalized  
189 to z-scores and averaged across all runs for each participant. A group t-test against zero was  
190 computed using the z-scored maps of each subject and corrected using a cluster-threshold  
191 statistical procedure based on Monte-Carlo simulation (initial  $p < 0.005$ , alpha level = 0.05,  
192 iteration = 5000). The group-level coverage of the network was then used as the initial mask for  
193 the body node extraction in the main experiment.

### 194 2.3.2. *Connectivity of the TOC body nodes*

195 The analysis for the main experiment is illustrated in **Figure 1**. First, a fixed-effects GLM was  
196 conducted on each participant's functional images with each different video as a separate

197 predictor. Fifty betas were estimated on each voxel for each participant. Next, a group ICA was  
198 conducted on the estimated betas within the TOC body network, following its definition with the  
199 network localizer. Fifteen ICs were extracted along with their video-wise betas. The component  
200 betas were then averaged for each condition and entered a group-level t-test for the contrast of  
201 [(aggressive body + defensive body + neutral body) > (face + object)] to select the body nodes.

202 To track the connectivity between the body nodes and the rest of the brain, a whole brain GLM  
203 was conducted with the IC responses as predictors. An event-related design matrix was first  
204 constructed by convolving each stimulus duration with the HRF and binned for each condition.  
205 Next, since the IC betas were extracted video-wise, the time-courses of each IC could be  
206 reconstructed by convolving the betas with a canonical HRF according to the on/offsets of the  
207 corresponding videos. These reconstructed time courses were then incorporated into the GLM  
208 design matrix to reflect their connectivity. Moreover, the IC betas were normalized within each  
209 condition so that the reconstructed time-courses reflected the item-to-item variance while  
210 omitting the categorical baseline modulations. The IC time-courses were modeled separately for  
211 each IC and each condition. The resulting betas were then entered in a voxel-wise ANOVA with  
212 factors Body conditions (aggressive / defensive / neutral) and Seeds nodes to assess the action  
213 type modulated connectivity. Statistical maps of significant main and interaction effects were  
214 corrected with the Monte Carlo cluster-threshold (initial  $p < 0.005$ , alpha level = 0.05, iteration =  
215 3000). Further multiple comparisons and simple effect tests were conducted at the ROI level for  
216 each significant cluster.

### 217 3. Results

218 Among the nineteen participants, two of them were excluded from both experiments and an extra  
219 one was excluded from the main experiment analysis due to large distortion of the functional or  
220 anatomical images.

221 In the localizer experiment, 75 independent components (ICs) were extracted from each subject's  
222 preprocessed functional images. Noise-induced ICs were identified excluded according to the  
223 spatial overlap with the white matter / cerebrospinal fluid mask, the mean response, and the  $r^2$  of  
224 the general linear modelling (GLM) fitting on the IC time-course. The details of the criteria are  
225 described in Li et al. (2023). By conducting GLM on the IC time-courses, the body-selective  
226 occipital network was identified by the analysis of the contrast [2 \* human body (normal -  
227 scramble) > human face (normal - scramble) + human object (normal - scramble)]. It exhibited a  
228 significant preference for human bodies over objects (**Figure 2a**,  $t(16) = 4.24$ , Benjamini-  
229 Hochberg False Discovery Rate corrected  $q = 0.006$ , right-tailed) in the bilateral lateral occipital  
230 cortex (LOC) and also included bilateral fusiform cortex, superior parietal lobe (SPL), posterior  
231 superior temporal sulcus (pSTS) / temporoparietal junction (TPJ), pulvinar and amygdala.

232 Following the identification of body network defined above (referred to as TOC network for  
233 simplicity), we examined their connectivity patterns and how they were influenced by the  
234 affective conditions. This analysis utilized the data from the main experiment, where 50 betas  
235 were extracted for each subject. A GIG-ICA procedure was then conducted within the predefined  
236 TOC network on each subject's 50 betas, resulting in 15 ICs along with their video-wise betas.  
237 The component betas were then averaged by condition and entered a group-level t-test for the  
238 contrast of [(aggressive body + defensive body + neutral body) > (face + object)] to select body

239 nodes. After multiple comparison correction, four adjacent nodes showed significant body  
240 selectivity (C04,  $t(15) = 8.08$ , corrected  $q < 0.001$ ; C06,  $t(15) = 4.21$ , corrected  $q = 0.002$ ; C07,  
241  $t(15) = 2.66$ , corrected  $q = 0.033$ ; C09,  $t(15) = 4.90$ , corrected  $q < 0.001$ ; all right-tailed; **Figure**  
242 **2b, c**). The decomposed beta values are shown in **Figure 2b**. Since the data was demeaned  
243 before entering the ICA, the zero-point in the plots indicate the averaged beta value across all  
244 masked voxels. The nodes C04 and C09 were distributed bilaterally covering the inferior LOC  
245 (**Figure 2c**, green component, LOCi for abbreviation) and the superior LOC (**Figure 2c**, purple  
246 component, LOCs for abbreviation) regions, respectively. The C06 and C07 had a unilateral  
247 distribution and covered the posterior middle temporal gyrus on the left (**Figure 2c**, red  
248 component, lpMTG) and right (**Figure 2c**, blue component, rpMTG) hemispheres, respectively.  
249 Consistent with a previous study on subdivisions of EBA (Weiner & Grill-Spector, 2011), these  
250 body nodes partially overlapped with and surrounded the hMT in each hemisphere (**Figure 2d**).

251 By comparing the spatial maps of the four independent components, we further investigated the  
252 voxels which were specifically dominant in each of the nodes. As shown in **Figure 3a**, non-  
253 overlapping clusters were identified where the component weights exhibited significantly larger  
254 values for one of the components than for the other three. The composition of each cluster was  
255 then labelled following the nomenclature of the Human Connectome Project multi-modal  
256 parcellation (HCP-MMP; Glasser et al., 2016). This showed that the dominance of the LOCi  
257 node was mainly found in V4t, PH, LO2, and FST (**Figure 3b**). The dominance of the pMTG  
258 nodes included PHT and TPOJ2 on the left side (**Figure 3b**), and further extended to TPOJ1 on  
259 the right side (**Figure 3b**). For the LOCs node, dominance was found especially in TPOJ3  
260 (**Figure 3b**).

261 To track the connectivity between the body nodes and the rest of the brain, a whole brain GLM  
262 was conducted with the IC responses as predictors. Since the IC betas were extracted item-wise,  
263 the time-courses of each IC can be reconstructed by convolving the betas with a canonical  
264 hemodynamic response function (HRF) according to the on/offsets of the corresponding videos.  
265 The IC time-courses were modeled separately for each IC and each condition, resulting in 5  
266 conditions \* 4 nodes (LOCi, LOCs, lpMTG, & rpMTG) = 20 connectivity terms added to the  
267 GLM. Next, to assess connectivity modulated by the action type, the resulting betas were entered  
268 in a voxel-wise ANOVA with the 3 body conditions (aggressive / defensive / neutral) \* 4 nodes.  
269 A significant main effect of node was found in widely distributed clusters, showing that different  
270 TOC body nodes have differentiated connectivity profiles (**Figure 4; Table 1**). The LOCi and  
271 LOCs nodes showed stronger connectivity to visual cortex and posterior cingulate cortex (PCC)  
272 while the two pMTG nodes were connected more dominantly to the middle & posterior insula,  
273 supramarginal gyrus (SMG) and frontal regions (**Figure 4; Table 1**).

274 Within the areas showing a significant main effect of body condition, affective modulations were  
275 found in anterior / posterior cingulate cortex (ACC / PCC) and caudate. Subsequent post hoc  
276 tests revealed a significant enhancement of overall node connectivity specifically for the  
277 defensive body condition (**Figure 5a; Table 1**). On the other hand, significant interaction effects  
278 between body condition and node node were found in middle frontal gyrus / frontal eye field and  
279 in cerebellum. In both clusters, the aggressive body condition significantly increased  
280 connectivity to the lpMTG node, while it decreased the connectivity to the LOCs node (**Figure**  
281 **5b; Table 1**).

## 282 **4. Discussion**

283 We investigated dynamic whole body processing using data driven methods and a network  
284 approach to address three fundamental questions about body and action representation in the  
285 brain: (1) is there a network in the brain showing body sensitivity; (2) what is the connectivity  
286 profile of the different nodes within this body selective network; and (3) are these nodes  
287 differently modulated by action specific information. Our results demonstrated four different  
288 nodes involved in body representation, each showing clear anatomical and functional distinctions.  
289 Furthermore, each of these nodes had a unique connectivity profile sustaining the processing of  
290 different affective whole-body action videos. These data-driven findings on multiple body nodes  
291 extend previous proposals based on traditional category localizer methods and provide novel  
292 insights into their functional connectivity sustaining action processing.

293

### 294 **4.1. *Four different body selective network nodes***

295 The current study identified four body-selective nodes within the TOC body network (Li et al.,  
296 2023). The largest node was the bilateral LOC<sub>i</sub> located in the inferior division of LOC and the  
297 lateral occipital sulcus (LOS). Another node, the bilateral LOC<sub>s</sub>, was defined in the superior  
298 division of LOC. Anterior to the LOC<sub>i</sub> and LOC<sub>s</sub>, two unilateral nodes were found in the  
299 posterior middle temporal gyrus (MTG), one on the left hemisphere and one on the right (lpMTG  
300 and rpMTG). Overall, our findings provide evidence for a distributed network of body  
301 representation consistent with an early proposal by Weiner and Grill-Spector (2010). The authors  
302 not only identified a single body category, but they also discovered three limb-selective areas

303 organized in a crescent shape around, yet not overlapping with, hMT+. Furthermore, other areas  
304 were also involved in limb representation including ITG and MTG. Our results confirm their  
305 findings on the anterior-posterior separation of body areas (Weiner & Grill-Spector, 2011). Yet  
306 importantly, we extend these findings by demonstrating a distinction between the dorsal and  
307 ventral LOC in terms of body processing, challenging the conventional definition of EBA as a  
308 homogeneous cluster in previous studies.

309 To understand differences among the four nodes, we investigated the composition of voxels  
310 exhibiting a significantly higher weight for each node compared to the other three. Voxels  
311 showing LOC<sub>i</sub> dominance were found in shape selective areas, such as LO2 and PH (Kolster et  
312 al., 2010), as well as in motion and body part selective areas including V4t and FST (Glasser et  
313 al., 2016; Kolster et al., 2010). In agreement with our findings, previous studies have suggested  
314 that the LOC<sub>i</sub> region may serve both as an entry of higher-level visual streams as well as a node  
315 for integrating the dorsal and ventral stream feedback (Grill-Spector & Malach, 2004; Larsson &  
316 Heeger, 2006; Sayres & Grill-Spector, 2008). Recent studies have also found involvement of the  
317 LOC<sub>i</sub> in body kinematics (Marrazzo et al., 2023) and motion feature processing (Robert et al.,  
318 2023). Thus, this node may be related to an early integration of the low-/mid-level visual  
319 information.

320 In contrast, LOC<sub>s</sub> dominance was mainly associated with TOPJ3, a region believed to be  
321 modulated by higher-level functions such as theory of mind (ToM) and working memory  
322 (Glasser et al., 2016). This once again underscores the functional divergence between the ventral  
323 and dorsal LOC. Similarly, both the lpMTG and rpMTG nodes involved multifunctional areas  
324 such as PHT and TOPJ2. Both TPOJ3 and PHT/TPOJ2 have shown selectivity for body and face  
325 stimuli (Glasser et al., 2016). Interestingly, stronger face form selectivity has been documented

326 for TOPJ3 compared to PHT/TPOJ2, whereas PHT/TPOJ2 has been associated with stronger  
327 body form selectivity (Glasser et al., 2016). Thus, the pMTG nodes and the LOCs node may be  
328 sensitive to different kinds of visual features associated with specific features of social  
329 information.

#### 330 **4.2. Different connectivity profiles**

331 An ANOVA with seed nodes and affective conditions as factors was used to compare the nodes'  
332 connectivity profiles. Distinct global connectivity profiles were revealed by the significant main  
333 effect of the seed node. The only cluster showing stronger connectivity to the LOCi node was  
334 EVA/V2. Regarding to the LOCs node, the strongest connectivity was found with default mode  
335 network and dorsal attention network nodes including PCC, precuneus, and superior frontal  
336 gyrus (SFG). This is consistent with other studies reporting ventral stream connectivity to the  
337 LOC (overlap with the LOCi) and dorsal stream connectivity to the EBA (overlap with the LOCs)  
338 (Zimmermann et al., 2018). Compared to the LOCi and LOCs, more widespread connectivity  
339 was found for the two anterior nodes, lpMTG and rpMTG, which suggested the pMTG nodes  
340 may serve as network hubs connecting the TOC network to the global-level computation. Both  
341 pMTG nodes had the highest connectivity to SMG and insula. However, in the case of the  
342 rpMTG, this connectivity extended further to encompass the MFG, angular gyrus and SFG. Also,  
343 different from the LOCs, the anterior nodes were linked to nodes of ventral attention / salience  
344 networks (VAN & SN). The asymmetric results are consistent with the right-lateralized  
345 distribution of the VAN (Vossel et al., 2014). Thus, these two nodes could contribute to a  
346 processing pathway for evaluating both valence as well as relevance in the observer.



### 347 **4.3. *Affective modulation for defensive vs. aggressive body movements***

348 The most significant and novel finding derived from our analysis is that for defensive/fearful  
349 body images we observed enhanced connectivity between the TOC and the PCC / precuneus as  
350 well as the caudate. The former are part of Posteromedial cortex (PMC), known for involvement  
351 in episodic memory, including autobiographical memory (Bubb et al., 2017; Leech & Sharp,  
352 2014), and are also part of the default mode network. The PCC has also been shown to exhibit  
353 increased activity when attention is directed towards a target of high motivational value (Leech  
354 & Sharp, 2014). Furthermore, recent evidence indicates that Anterior Precuneus is causally  
355 linked to the bodily self, or the brain resources involved with the observers' body schema (Lyu et  
356 al., 2023). Seen against this background, witnessing fear/defensive actions may engage brain  
357 activity intricately associated with the bodily self, in the sense of increasing the involvement of  
358 the observer. These results suggest that observing fearful expressions is associated with  
359 concomitant neural activity preparing the brain for defensive actions (de Gelder et al., 2004). In  
360 support, several brain areas identified in studies of body schema (Berlucchi & Aglioti, 2010),  
361 sense of self and its deficits in pathological conditions (Dary et al., 2023) overlap with the  
362 network described here.

363 In contrast, the connectivity modulation for aggressive actions showed a different pattern and  
364 varied across different nodes. As revealed by the ANOVA analysis, a significant interaction effect  
365 was observed for FEF and cerebellum regions. When viewing the aggressive videos, the two  
366 regions showed increased connectivity to the lpMTG node, and decreased connectivity to the  
367 LOCs node. As mentioned above, both lpMTG and LOCs were composed of voxels from social  
368 perception regions such as TPOJ and PHT and may be related to the processing of different types

369 of social features. And compared to LOCs, the lpMTG node exhibited a higher level of global  
370 connectivity. Moreover, the FEF region is known for its involvement in shaping attention  
371 (Veniero et al., 2021) and as such the FEF may be part of decision making (Krajbich et al., 2021).  
372 Thus, the opposite FEF connectivity pattern to the lpMTG and LOCs may suggest the  
373 reorientation of salient features and the distributed computation during the presence of  
374 aggressive stimuli.

## 375 **5. Conclusions**

376 We showed that body representation is implemented in the brain in several different body  
377 sensitive hubs that each have their connectivity network. These hubs and connectivity networks  
378 have a relatively specific functional role in body and action perception. This supports the notion  
379 that these are body sensitive hubs that are not so much defined by abstract body category  
380 selectivity than by their different functional roles in action understanding.

381

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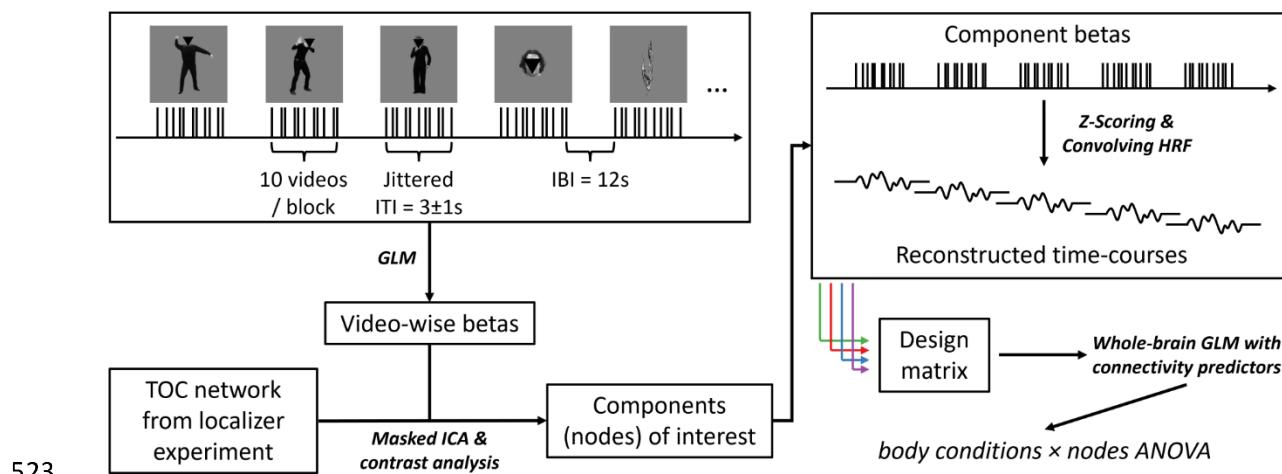
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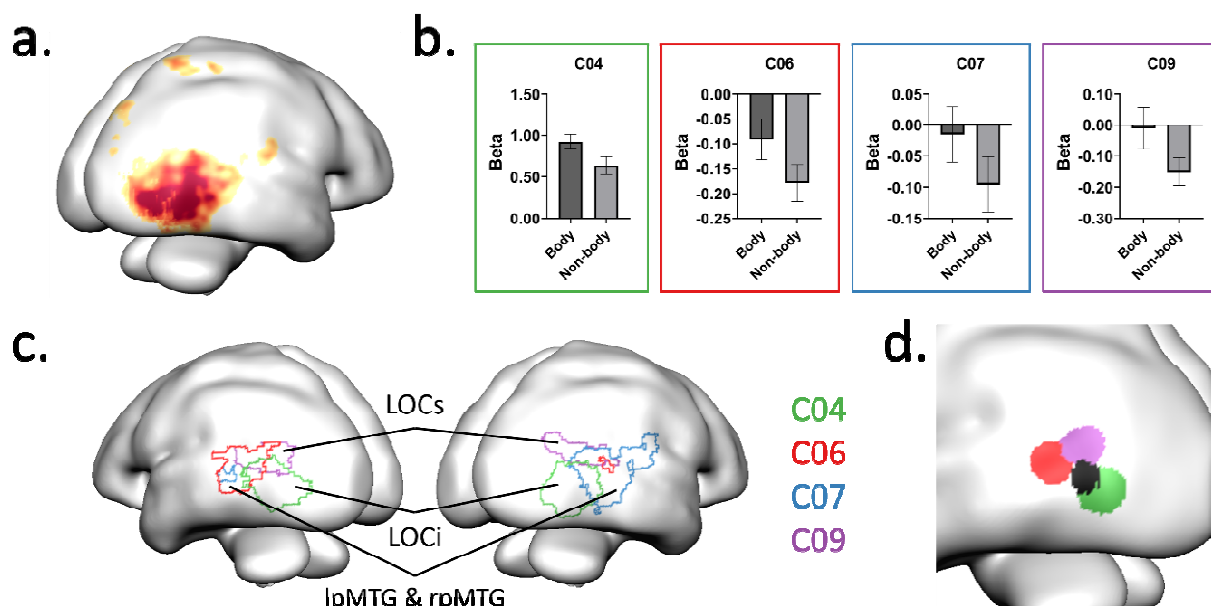
521

522 **Figures & tables**



524 Figure 1. Illustration of the main experiment design and analysis. Five video conditions  
525 (aggressive / defensive / neutral face blurred body, neutral face, and object; face region overed  
526 here for privacy) were presented in a mixed block/event-related design, in which the stimuli were  
527 blocked for each condition while with jittered inter-trial-interval around 3s. For each condition,  
528 ten different videos were included and were repeated ten times across five runs. GLM was  
529 conducted to estimate the response for each different video, resulting in 50 betas extracted for  
530 each participant. The video-wise betas were then entered to an ICA procedure within the body  
531 sensitive TOC network identified by the localizer experiment. Body selective network nodes  
532 were defined by higher component responses for body videos than for non-body videos. To track  
533 the whole-brain connectivity of each selected node, the video-wise betas were z-scored and  
534 convolved with the hemodynamic response function within each condition, resulting in five  
535 reconstructed time-courses for each component. The reconstructed time-courses for all selected  
536 components were then added to a whole-brain GLM design matrix as the predictors for seed-  
537 based connectivity. Finally, two-factor ANOVA was conducted with connectivity betas across all  
538 participants to test their modulations from the body conditions or the node components.





540

541 Figure 2. (a). The coverage of TOC network as defined in the network localizer experiment. (b).

542 Beta plots of the four body-nodes from the main experiment. Zero-point indicates the mean beta

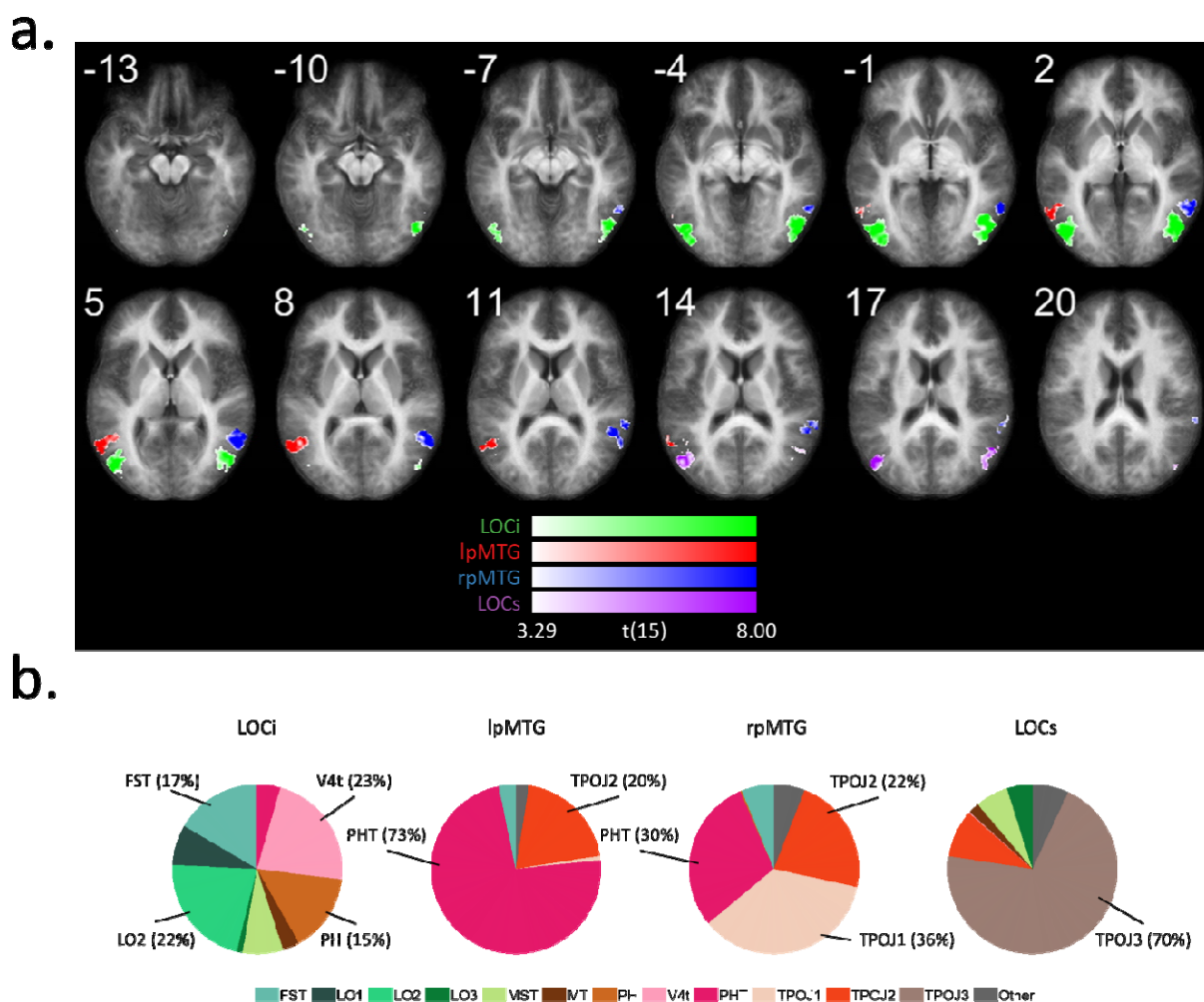
543 value across all masked voxels. Colors indicate the component indexes. (c). The spatial

544 distribution of the four body nodes. (d). The relative position of the centers of body nodes and

545 the hMT (black) on the left hemisphere.

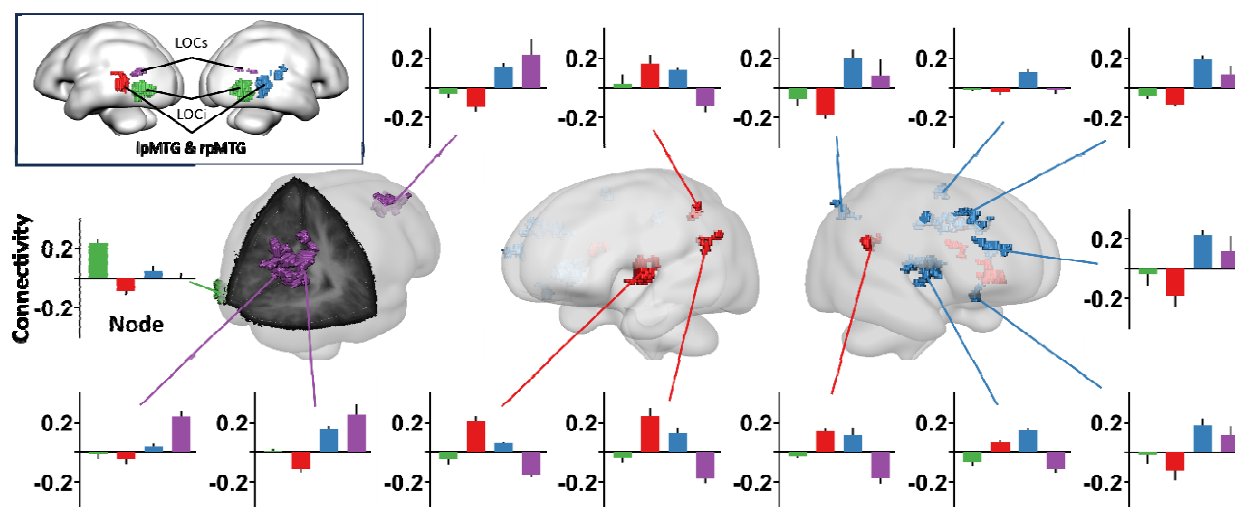
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547  
 548 Figure 3. (a). Map of voxels with significantly higher contribution from each of the nodes. The  
 549 voxel-wise IC weight from each node was compared to the other three nodes and entered a  
 550 group-level t-test against zero (two-tailed). The resulting map was corrected by a cluster-  
 551 threshold statistical procedure based on Monte-Carlo simulation (initial  $p < 0.005$ , alpha level =  
 552 0.05, iteration = 5000). Slice numbers indicate the Z coordinates in the Talairach space. (b). The  
 553 voxel composition for each cluster in (a.), labeled by the HCP-MMP atlas (Glasser et al., 2016).

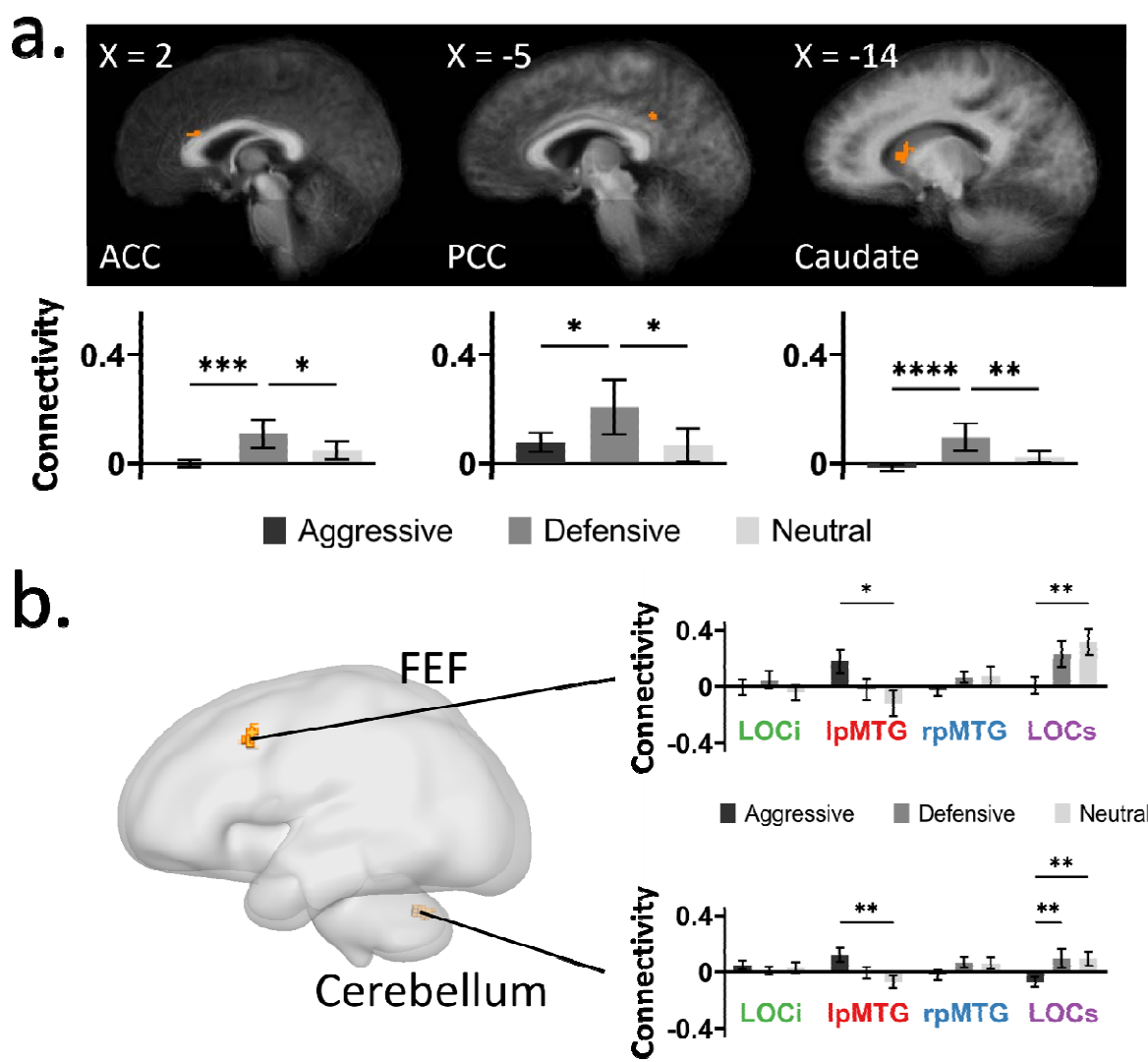
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555

556 Figure 4. Clusters with significant main effect of the seed node on connectivity. Cluster colors  
557 indicate the seed nodes with the highest connectivity to the corresponding clusters. Bar plots  
558 show the detailed connectivity profile of each cluster, with bar colors indicating the seed nodes  
559 as shown in the upper-left framed panel.

560



561

562 Figure 5. (a) Clusters with significant main effect of the body condition on TOC connectivity

563 (averaged across nodes). (b) Clusters with significant interaction between the body condition and

564 the seed node. Bar plots show the detailed connectivity profile of each cluster. Asterisks indicate

565 the significant pairwise comparisons after Bonferroni correction (\*:  $p < 0.05$ ; \*\*:  $p < 0.01$ ; \*\*\*:

566  $p < 0.001$ ; \*\*\*\*:  $p < 0.0001$ ).

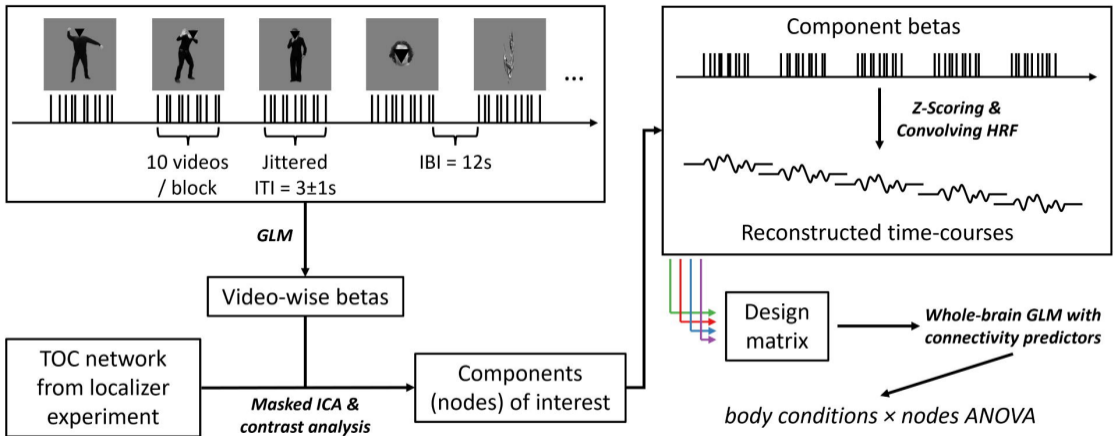
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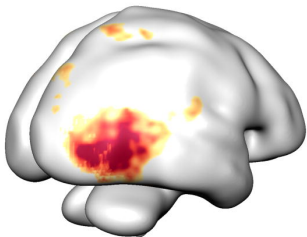
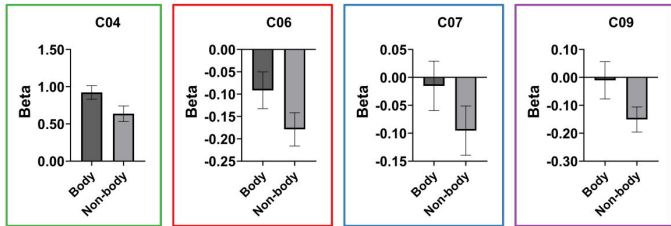
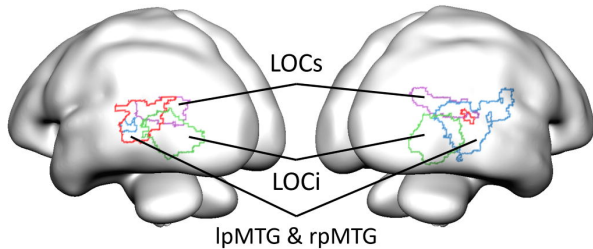
Table 1.

Results from the 3 (body condition)  $\times$  4 (node) ANOVA on the connectivity betas.

	Talairach label	Brodmann label	Talairach coordinates			ROI statistics
			x	y	z	
Mean effect: Node						F(3,45)
	Middle Occipital Gyrus	BA 18	-18	-95	9	6.01
	Posterior Cingulate	BA 30	-9	-62	14	5.76
		BA 31	12	-54	25	6.57
	Angular Gyrus	BA 39	39	-58	39	5.54
	Inferior Parietal Lobule	BA 40	-60	-36	26	5.60
	Postcentral Gyrus	BA 2	-50	-25	38	5.80
		BA 40	62	-22	22	6.12
	Medial Frontal Gyrus	BA 6	10	-4	57	6.57
	Insula	BA 13	-39	1	1	6.23
	Precentral Gyrus	BA 44	49	4	12	6.11
	Superior Frontal Gyrus	BA 8	23	28	46	6.19
		BA 10	41	49	20	5.97
	Inferior Frontal Gyrus	BA 47	30	30	-9	5.97
	Middle Frontal Gyrus	BA 9	30	33	30	5.90
Mean effect: Body condition						F(2,30)
	Cingulate Gyrus	BA 31	-4	-44	33	7.53
	Caudate		-12	15	9	7.21
	Anterior Cingulate	BA 33	4	22	22	6.75
Interaction effect: Node * Body						F(6,90)
	Middle Frontal Gyrus	BA 8	-44	10	41	3.82
	Cerebellum		1	-60	-30	3.94





**a.****b.****c.**

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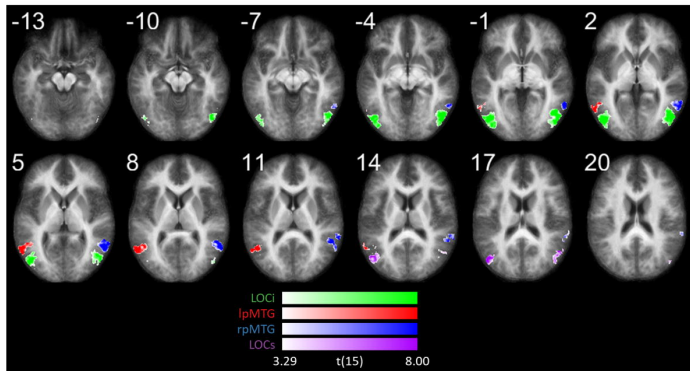
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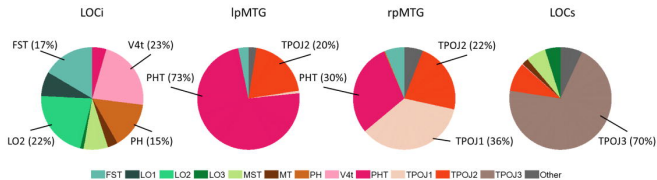
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**d.**

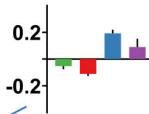
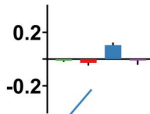
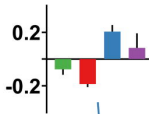
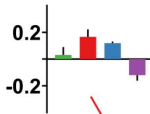
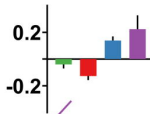
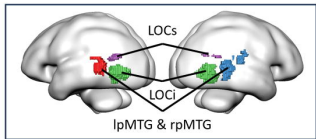
a.



b.







Connectivity

Node

