

1 **The human claustrum is functionally connected to cognitive networks and involved in**
2 **cognitive control**

3

4 Abbreviated title: Claustrum activation during cognition

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29 control, attention, cortex, forebrain

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41

42 **Abstract**

43 The claustrum is among the most highly connected structures in the mammalian brain.
44 However, the function of the claustrum is unknown, which is due to its peculiar anatomical
45 arrangement. Here, we use resting state and task functional magnetic resonance imaging
46 (fMRI) to elucidate claustrum function in human subjects. We first describe a method to reveal
47 claustrum signal with no linear relationship with adjacent regions. We applied this approach to
48 resting state functional connectivity (RSFC) analysis of the claustrum at high resolution (1.5 mm
49 isotropic voxels) using a 7T dataset (n=20) and a separate 3T dataset for replication (n=35). We
50 then assessed claustrum activation during performance of a cognitive task, the multi-source
51 interference task, at 3T (n=33). Extensive functional connectivity was observed between
52 claustrum and cortical regions associated with cognitive control, including anterior cingulate,
53 prefrontal and parietal cortices. Cognitive task performance was associated with widespread
54 activation and deactivation that overlapped with the cortical areas showing functional
55 connectivity to the claustrum. Furthermore, the claustrum was significantly activated at the onset
56 of the difficult condition of the task, but not during the remainder of the difficult condition. These
57 data suggest that the claustrum can be functionally isolated with fMRI, and that it is involved in
58 cognitive control in humans independent of sensorimotor processing.

59 **Highlights**

- 60 • Removing signal from neighboring structures isolates claustrum BOLD signal at 7T and
61 3T field strength
- 62 • Claustrum is extensively functionally connected with cortex, including cognitive networks
- 63 • Claustrum is activated at the onset of a cognitive conflict task
- 64 • Claustrum may be involved in cognition independent of sensorimotor processing

65

66

67 **Introduction**

68 In its mediolateral dimension, the claustrum is thin (submillimeter at certain points), but its
69 rostrocaudal and dorsoventral dimensions are roughly equivalent to that of the striatum.
70 Decades of tract tracing studies in several mammalian species indicate that the claustrum is
71 bidirectionally connected with many cortical areas (Edelstein and Denaro, 2004; Crick and Koch,
72 2005; Mathur et al., 2009; Mathur, 2014; White et al., 2017; Wang et al., 2017) and is estimated
73 by volume to be among the most highly connected structures in the brain (Torgerson et al.,
74 2015). These observations have fueled several hypotheses that the claustrum: 1) binds
75 multimodal sensory information for the generation of conscious perception (Crick and Koch,
76 2005); 2) coordinates somatosensory and motor cortical information (Smith et al., 2012) and; 3)
77 acts as a cortico-cortical relay center supporting attention (Mathur, 2014).

78 Recent comprehensive analyses in a single species of how the cortical mantle connects
79 with the claustrum demonstrate that the claustrum weakly innervates primary sensorimotor
80 cortices, while heavily innervating frontal cortices including the anterior cingulate cortex (ACC)
81 and the prelimbic area of the medial prefrontal cortex (PFC) (White et al., 2017). A strong input
82 from the ACC to the claustrum also exists in rats (Smith and Alloway, 2010; White et al., 2017)
83 and mice, and this input encodes a top-down preparatory signal that is proportional to task
84 difficulty (White et al., 2018). These findings suggest that the claustrum may subserve frontal
85 cortical function, including top-down executive processes (Mathur, 2014; White and Mathur,
86 2018). However, evidence for a role of the human claustrum supporting any of the
87 aforementioned functional hypotheses, including cognitive processing, is particularly lacking.
88 While the anatomical boundaries of the human claustrum can be resolved with relative ease
89 using high-resolution structural magnetic resonance imaging (MRI), functionally resolving this
90 structure for analysis of blood-oxygenation level-dependent (BOLD) signal with functional MRI
91 (fMRI) is challenging, as the signal extracted from the claustrum is heavily mixed with the signal

92 from the neighboring insula and putamen using standard methods. Analysis of BOLD data using
93 standard methods results in similar patterns of functional connectivity (correlation of signal
94 between regions) when comparing claustrum, insula, and putamen. This contrasts with data
95 from multiple tract tracing studies, which instead show unique patterns of anatomical
96 connectivity across these regions (Nakashima et al., 2000; Mathur et al., 2009; Pan et al., 2010;
97 Sato et al., 2013; Wang et al., 2017). Thus, standard fMRI analyses are not capable of
98 functionally resolving the claustrum and may yield inaccurate functional connectivity and
99 activation results.

100 In an effort to elucidate claustrum function, the current study has three goals: 1) devise new
101 fMRI methodology called Small Region Confound Correction (SRCC) to functionally distinguish
102 the claustrum from the insula and putamen, creating a corrected claustrum timeseries; 2)
103 perform resting state functional connectivity analyses with the corrected claustrum timeseries to
104 reveal functional coupling of the claustrum in humans; and 3) test the hypothesis that the
105 claustrum, owing to strong connectivity with frontal cortices and recent data in mice suggesting
106 its involvement in top-down cognitive processing (White et al., 2018), is activated during a
107 cognitive conflict task. Our data indicate that extraction of unique claustrum signal using fMRI is
108 possible and that the claustrum is functionally connected with cognitive networks in the resting
109 state. During task performance, we found the claustrum to be active at the onset of – or switch
110 to – cognitive conflict task engagement.

111 **Methods**

112 *Overview*

113 Three datasets were analyzed. The first dataset, *7T-Rest*, is publicly available and consisted of
114 scans from 20 healthy humans scanned with 7 Tesla (T) MRI (Gorgolewski et al., 2015). The
115 second dataset, *3T-Rest*, consisted of scans from 35 healthy humans acquired with 3T MRI.

116 The third dataset, *3T-Task*, used the same subjects as *3T-Rest* and consisted of scans from 33
117 healthy humans performing a cognitive interference task acquired with 3T MRI. In *7T-Rest*, we
118 performed seed-based whole-brain resting state functional connectivity (RSFC) analyses using
119 the claustrum and surrounding structures (insula/putamen) as seeds. To examine claustrum
120 functional connectivity while controlling for the influence of insula and putamen signal, we
121 regressed out the timeseries of insula/putamen sub-regions from the claustrum, creating a
122 corrected claustrum signal that we then used to make a claustrum specific RSFC map. We used
123 *3T-Rest* to replicate findings from *7T-Rest*. We applied similar methods used in resting state
124 data to *3T-Task* in order to isolate BOLD signal from the claustrum, and analyzed this signal to
125 determine claustrum activation during cognitive load.

126 *Participants and MRI data*

127 *7T-Rest*: Data from 22 subjects were acquired from a publically available dataset scanned on a
128 7T MR scanner (MAGNETOM 7T, Siemens Healthcare, Erlangen, Germany). We used data
129 from only the first session of this dataset. These data included a T1 weighted structural 3D
130 MP2RAGE image that was used in preprocessing (TR=5000ms, TE=2.45ms, voxels=.7 mm
131 isotropic). *7T-Rest* also included two eyes-open resting state scans using echo planar imaging
132 (EPI) to measure BOLD fMRI while fixating on a plus sign (whole brain coverage, TR=3000ms,
133 TE=17ms, voxels=1.5mm isotropic, slices=70, duration=300 TR). Details of the scans can be
134 found in (Gorgolewski et al., 2015). Two subjects were excluded, one from errors induced by
135 preprocessing and a second from different scanning parameters than the other participants,
136 leaving 20 subjects for analysis (10 women, average age = 25, s.d. = 2).

137 *3T-Rest*. 36 healthy subjects were recruited as a control sample for an ongoing clinical trial and
138 only a baseline scan was used for the following analyses. MRI data were acquired at the
139 University of Maryland, Baltimore Medical Imaging Facility with a Siemens 3T Tim Trio scanner
140 with 32 channel head coil (n=22) or a Siemens 3T Prisma scanner with a 64 channel head coil

141 (n=14) due to a scanner upgrade during data acquisition. We acquired a T1 weighted structural
142 3D MPRAGE scan that was used in preprocessing (whole brain coverage, TR=2300ms,
143 TE=2.98ms, voxels=1.00 mm isotropic). We also acquired eyes-open resting state scans using
144 EPI while subjects fixated on a plus sign (whole brain coverage, TR=2s, TE=28ms, voxels=3.4 x
145 3.4 x 4.0 mm, slices=40, duration=300 TR). One subject was excluded due to poor coverage
146 leaving 35 subjects (31 women, average age=37, s.d. = 13).

147 *3T-Task*: Participants in the *3T-Rest* also performed the multi-source interference task (MSIT;
148 Bush et al., 2003) as a measure of cognitive conflict. Subjects were first trained on the MSIT
149 outside of the scanner. During the MRI session, they performed the MSIT in two runs of about 5
150 minutes each while echo planar imaging with whole brain coverage was acquired (TR=2500ms,
151 TE=30ms, voxels=3.00 mm isotropic, slices=44, duration=121 TR). The task was the same as
152 previously reported (Seminowicz and Davis, 2007, Seminowicz et al., 2011), but is briefly
153 described here. On each trial, the volunteer was presented with an array of three numbers. In
154 each array, two numbers were the same and one number was different. The volunteer was
155 instructed to press a button that corresponded to the number that was different from the two
156 other numbers or characters presented on the screen for that given trial. The control condition
157 was a sequential tapping task in which an asterisk appeared in the same order moving from the
158 left to the right of the screen and the subject pressed a button corresponding to the position of
159 the asterisk. There were two levels of task difficulty (easy, difficult), which were performed in
160 separate 20s blocks (10 stimuli per block). In the easy condition, the different number indicated
161 the position of that number in the array (e.g. “1-2-2”, “3-2-3”, “1-1-3”), and in the difficult
162 condition, the different number did not indicate the position of that number in the array (e.g. “3-
163 2-2”, “3-1-3”, “1-1-2”). Three subjects were excluded because of poor coverage and missing
164 data, leaving a sample size of 33 (29 women, average age=37 s.d.= 12).

165

166 *Initial Data Preprocessing*

167 Data were preprocessed in SPM12 (<http://www.fil.ion.ucl.ac.uk/spm/software/spm12/>), including
168 slice timing correction, realignment (motion correction), coregistration of the T1-weighted
169 structural scan to the mean realigned functional image, segmentation of the structural scan,
170 normalization of the structural and realigned functional images to a standard MNI template, and
171 smoothing with a 6mm full width at half maximum (FWHM) Gaussian kernel for *7T-Rest*, *3T-*
172 *Rest*, and *3T-Task*. We elected to use the same smoothing kernel for 3T-Rest and 7T-Rest to
173 have as comparable processing pipelines as possible. We did not observe obvious qualitative
174 differences for smoothing *7T-Rest* at 6mm FWHM vs a 3mm FWHM kernel (S1).

175 *Analysis of resting state claustrum connectivity*

176 *Resting-state preprocessing.* Whole-brain *7T-Rest* and *3T-Rest* data underwent the following
177 further preprocessing before functional connectivity analyses were performed. Resting-state
178 preprocessing and seed-based analyses were conducted in the Conn toolbox version 17f
179 (<http://www.nitrc.org/projects/conn>). Given continued controversy, we elected not to control for
180 global signal in our analyses (Murphy and Fox, 2017). To control for noise present in white
181 matter and CSF, we used aCompCor (Behzadi et al., 2007; Muschelli et al., 2014) to determine
182 the first five eigenvectors of white matter and CSF. As we did not want to remove global signal,
183 we used a twice eroded CSF mask and a white matter mask with four erosions (that did not
184 include external or extreme capsules), as this level of erosion has been shown to no longer
185 contain global signal (Power et al., 2017). We removed motion related signals estimated from
186 the realignment parameters along with the first order derivatives of these parameters, in addition
187 to the 10 eigenvectors from white matter and CSF. To avoid the reintroduction of noise while
188 removing low frequency artifact, we used linear detrending and also simultaneously high pass
189 filtered our voxelwise and regressor data with a cutoff of 0.008 Hz. No low pass filter was used
190 because signal is present above standard cutoffs (~0.08 Hz, Smith et al., 2013). Despiking was

191 employed after these steps to remove any additional artifact that had not yet been removed
192 (Petel et al., 2014).

193 *Standard analysis*

194 Following artifact removal, ROI analyses were performed by extracting the average timeseries
195 of all voxels within a given set of ROIs using normalized, but not smoothed data. For each
196 subject in *7T-Rest*, left and right claustrum ROIs were hand drawn on the subject's structural
197 image, along with ROIs for the insula and putamen. Fig 1A shows an example tracing for one
198 subject and Fig 1B shows the average ROIs on a group template. In *3T-Rest*, we used mean
199 ROIs obtained from *7T-Rest*, after confirming that these mean images fit well onto the
200 normalized structural data for *3T-Rest*. Functional connectivity was calculated as the Pearson
201 correlation between the timeseries for each ROI and all voxel timeseries across the brain. In the
202 standard analysis for *7T-Rest*, we used the un-corrected claustrum, whole insula, and whole
203 putamen timeseries from individually drawn ROIs to compute RSFC across the whole brain.

204 *Corrected claustrum analysis with Small Region Confound Correction (SRCC)*

205 To remove the possible influence of insula and putamen signals on claustrum timeseries and
206 hence FC, we determined the timeseries of parts of the insula and putamen that 'flanked' the
207 claustrum and treated these as confounding sources for the claustrum. These flanked ROIs
208 were calculated on the individual level for *7T-Rest* by dilating the claustrum 4 functional voxels
209 (6 mm), and determining the overlap between the dilated claustrum and the insula/putamen
210 seeds that were at least 2 functional voxels (3 mm) separated from the original claustrum. This
211 resulted in 'flanking' regions that were continuous with insula and putamen and similar in shape
212 to the claustrum, but still distant enough from the claustrum to ensure that they contained no
213 claustrum data within the seed, as shown in Fig 1C. We call this approach Small Region
214 Confound Correction (SRCC), as it is designed to eliminate partial volume effects that are

215 particularly problematic in small regions of the brain. In *3T-Rest*, this process was repeated, but
216 for the mean ROIs only. We regressed each claustrum timeseries (left and right) on the
217 timeseries from the ipsilateral ‘flanking’ regions, along with previously described artifacts
218 (motion, CSF, etc.), and the residuals from this analysis constituted a corrected claustrum
219 timeseries (e.g. corrected for signal in the flanking insula and putamen sources) that was then
220 used as the seed in whole-brain seed-to-voxel analysis.

221 To determine statistical significance of seed-to-voxel functional connectivity maps, we
222 performed one sample t-tests in SPM12 for each RSFC map (left and right corrected claustrum).
223 These outputs were used for qualitative comparisons between regions. A cluster-forming
224 threshold of $p < 0.001$ was used for all analyses and significant clusters based on FWE
225 correction are reported, as these thresholds have been shown to adequately control for false
226 positive rates (Woo et al., 2014).

227 *Modelling task data*

228 We determined the reaction time (RT) for every trial presented in the MSIT. If the participant did
229 not respond to a particular trial, we designated the RT for that trial as the maximum possible trial
230 duration (i.e. 1500ms). We did not analyze accuracy, as the average was over 90 percent.
231 Typically, the MSIT is modelled in a blocked design, using three types of blocks (tapping, easy,
232 and difficult). However, based on observations of behavioral performance suggesting that the
233 first one to three trials had much poorer performance (see Results) – likely reflecting the
234 cognitive adaptations (or switching) to a new set of rules – we modelled onset and block
235 separately, using an event design to model task onset and a block design to model the
236 remainder of the task.

237 *Preprocessing*

238 For whole brain task analyses, we did not treat insula and putamen signals as a source of noise.

239 For analyses of claustrum activation we used similar approaches as described in *Analysis of*
240 *resting state claustrum connectivity and corrected claustrum analysis with SRCC* (see above)

241 except we did not include despiking nor the first derivative of realignment parameters. This
242 approach created a corrected claustrum timeseries during MSIT that was then used to
243 determine claustrum activation during the task. We then estimated whole brain and corrected
244 claustrum ROI activation patterns.

245 *Analysis*

246 To determine significant task activation for whole brain analyses, we performed one sample t-
247 tests for contrast maps with a cluster-forming threshold of $p < 0.001$ and a FWE cluster
248 correction. Contrast maps for difficult and easy blocks and onsets were calculated based on a
249 tapping baseline (e.g. [difficult > tapping] and [easy > tapping]). To determine claustrum
250 activation, we performed a one-sample t-test on extracted activation estimates for each contrast
251 extracted using MarsBaR v 0.44 (<http://marsbar.sourceforge.net/>). To examine the overlap
252 between claustrum connectivity and task-related activation during MSIT, we created an overlap
253 map between areas of significant deactivation/activation and areas with significant claustrum
254 functional connectivity. We then used this overlap map to calculate the percentage of task-
255 responding voxels that functionally connected with the claustrum. Average RT for each task
256 condition (tapping, easy, difficult) was calculated for each subject, and these subject-averaged
257 RTs were compared between each task type using a one-way repeated measures ANOVA with
258 Greenhouse-Geisser correction. Post-hoc t-tests were used to identify specific task type
259 differences with Tukey's correction for multiple comparisons. We also performed a one-way
260 ANOVA with Greenhouse-Geisser correction and used a Dunnett's test to compare the first trial
261 to all other trials. Whole brain contrast of difficult vs easy tasks gave task-positive
262 (difficult>easy, EMN) and task-negative (easy>difficult, DMN) networks. For event-related plots

263 in Fig 4C, we extracted timeseries
264 data for the claustrum and for EMN
265 and DMN. These timeseries were
266 each averaged for the task of
267 interest.

268

269 Results

270 *Resting state connectivity of the*
271 *claustrum at 7T: methodological*
272 *approach*

273 We used high spatial resolution fMRI
274 data in *7T-Rest* and hand drawn
275 ROIs of claustrum, insula, and
276 putamen to analyze whole brain
277 functional connectivity of these ROIs.
278 Despite excellent resolution, we
279 found that functional connectivity
280 between the claustrum and the
281 insula/putamen in this standard
282 analysis was high (average FC of L/R
283 claustrum with insula/putamen
284 ranged from 0.32 to 0.5; Fig 1D). This
285 contrasts with known unique
286 connectivity of the claustrum relative

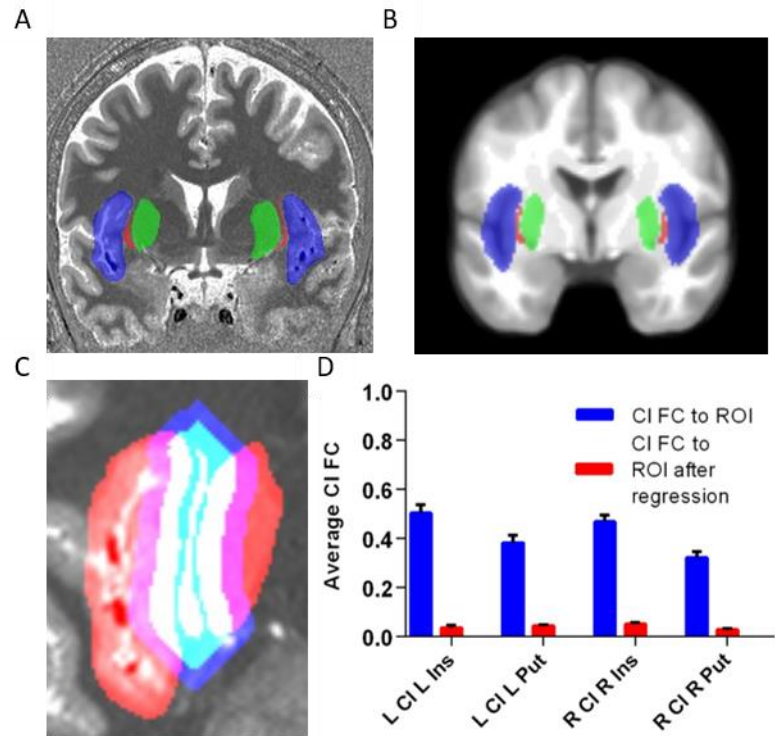


Fig 1. The claustrum is a thin sheet of gray matter and its fMRI signal is normally confounded by neighboring structures. A) Structural MRI image with insula (blue), claustrum (red), and putamen (green) for one subject. These regions were hand-drawn for each subject on their native anatomical image. B) Average insula (blue), claustrum (red), and putamen (green) averaged over 20 hand-drawn images displayed on an average template. C) To remove signal from neighboring structures, the claustrum was dilated, adding 6 mm around the claustrum, while maintaining its shape (blue). This dilated claustrum was overlapped with the neighboring insula and putamen (red). Voxels that contained insula/putamen and dilated claustrum separated by at least 3mm from the original claustrum were categorized as flanking regions. Flanking regions, shown in purple, were regressed out of the claustrum signal to create a corrected claustrum timeseries. D) Correlation (FC) of claustrum timeseries with L/R insula/putamen before and after regressing out flanking regions (i.e. following Small Region Confound Correction (SRCC)). The average connectivity between claustrum and its neighbors approaches zero following this approach. Error bars show standard error of the mean. Cl=claustrum, Ins=insula, Put=putamen, ROI=region of interest, FC=functional connectivity.

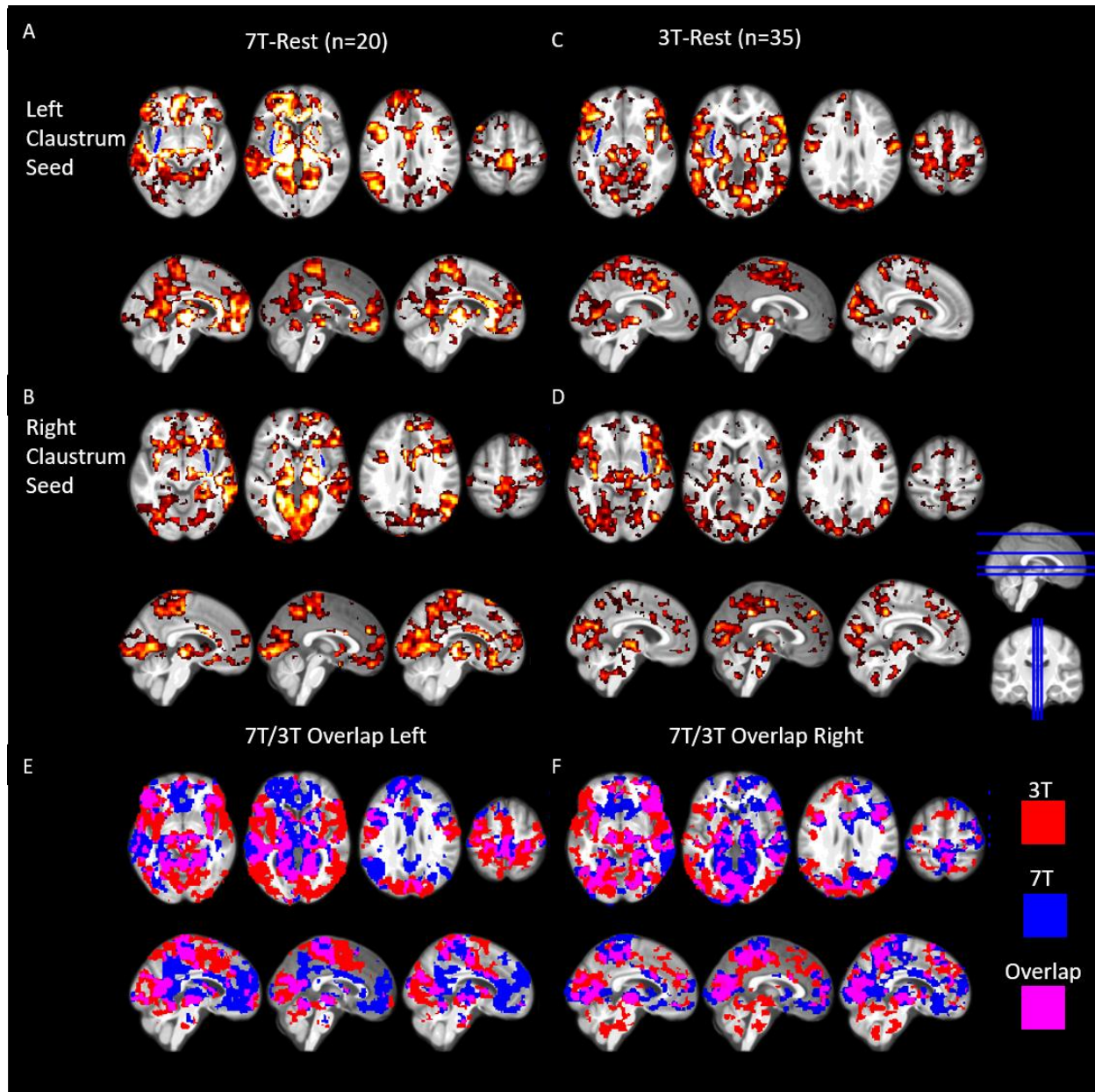


Fig 2. Resting state connectivity of the left and right claustrum at 7T and 3T showing widespread connectivity to cortical and subcortical regions. A) RSFC of the left claustrum in 7T-Rest. B) RSFC of the right claustrum in 7T-Rest. C) RSFC of the left claustrum in 3T-Rest. D) RSFC of the right claustrum in 3T-Data. E) and F) show the overlap of these thresholded RSFC maps. Data were voxelwisethresholded at $p < 0.001$ followed by FWE cluster correction. Blue ROI in A-D represents an average claustrum dilated by 1.5 mm (for visualization only). Cl=claustrum.

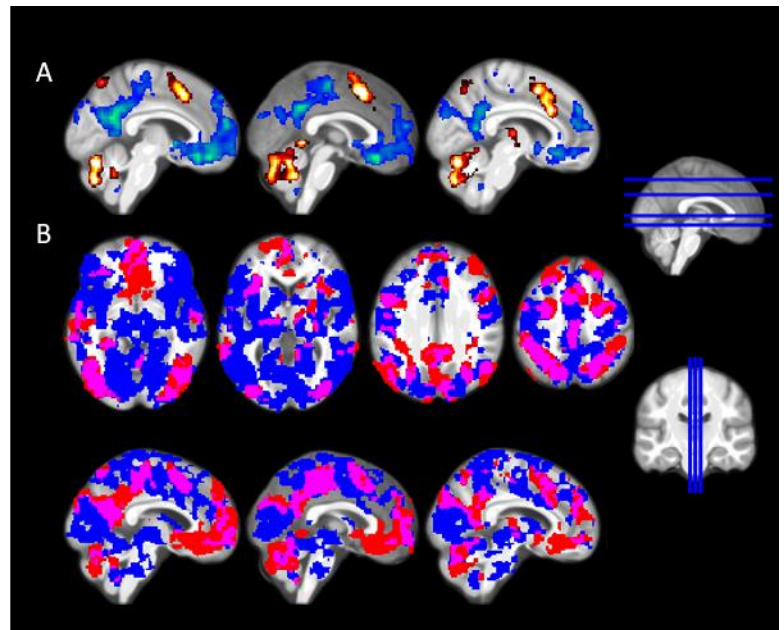
287 to insula and putamen (Nakashima et al., 2000; Mathur et al., 2009; Pan et al., 2010; Sato et al.,
288 2013; Wang et al., 2017), and suggests that even with high spatial resolution and well identified
289 ROIs that insula and putamen signals are sampled in claustrum voxels. This confounds the

290 interpretation of small structure studies
291 in fMRI even when the regions can be
292 identified with relative ease in structural
293 scans.

294 *Creating a corrected claustrum signal* 295 *with SRCC*

296 To mitigate the effect of insula/putamen
297 sampling in claustrum voxels, we
298 generated a corrected claustrum
299 timeseries using SRCC by regressing
300 out signal from insula and putamen
301 regions separated by two voxels from
302 claustrum (3 mm) as sources of noise,
303 identical to how nuisance white matter
304 (WM) and cerebro-spinal fluid (CSF)
305 signals are treated (see Fig 1C). This
306 approach eliminated the linear
307 relationship between claustrum and the
308 surrounding structures, and the resulting
309 average correlation between the
310 corrected claustrum timeseries and
311 neighboring regions reduced effectively
312 to zero (Fig 1D).

313 *Resting state connectivity of the corrected claustrum at 7T and 3T*



C

Fig 3. Claustrum functional connectivity substantially overlaps with DMN and EMN. A) Contrasting the difficult block and the easy block of MSIT reveals DMN (easy>difficult, shown in blue) and EMN (difficult>easy, shown in red). B) Areas functionally connected with L/R claustrum in 3T-Rest (blue) and the DMN and EMN shown above (red) show a high degree of overlap (purple) C) Quantification of DMN/EMN voxels from panel A and the percent that also have FC with claustrum. Claustrum FC maps are the same as shown in Figure 2B. Task contrasts maps used a $p < 0.001$ threshold with FWE cluster correction. Easy=easy condition of MSIT, Diff=Difficult condition of MSIT, CI=claustrum, DMN=default mode network, EMN=extrinsic mode network.

314 We estimated functional connectivity of the corrected claustrum timeseries across the whole
315 brain in two datasets. Given previous findings in rats (White et al., 2017), we anticipated
316 claustrum RSFC to be widespread and to feature connectivity with cingulate cortex, PFC, visual
317 cortex, and intraparietal sulcus (IPS), among other regions. In *7T-Rest*, this analysis identified
318 connectivity with the: thalamus, particularly the pulvinar; nucleus accumbens; visual cortex; both
319 anterior and posterior cingulate cortex; PFC (including the dorsal lateral PFC, medial PFC, and
320 ventral lateral PFC); precuneus; angular gyrus; sensorimotor cortex; parahippocampal gyrus;
321 superior and inferior temporal gyri; and IPS (Fig 2A and B and Table S1 and S2).

322 We found a more extensive pattern of claustrum functional connectivity in *3T-Rest*, however, the
323 regions displaying FC with claustrum were largely similar to *7T-Rest*. In *3T-Rest* we identified
324 connectivity with: the thalamus, mainly the pulvinar; nucleus accumbens; visual cortex; both
325 anterior and posterior cingulate cortex; PFC, precuneus; angular gyrus; sensorimotor cortex;
326 parahippocampal gyrus; temporal gyri; and IPS (Fig 2C and D and Table S3 and S4). The
327 similar pattern of claustrum functional connectivity across *7T-Rest* and *3T-Rest*, and the
328 bilateral nature of claustrum functional connectivity (i.e. left claustrum functionally connects with
329 right) suggest our confound-corrected claustrum timeseries is not artifactual.

330 *Extrinsic and Default Mode Network overlap with claustrum functional connectivity*

331 Given the high degree of claustrum functional connectivity with regions involved in cognitive
332 control (e.g. ACC) and prior literature showing claustrum involvement in top-down cognitive
333 processing (White et al., 2018), we sought to quantify the overlap between resting-state
334 claustrum connectivity, the cognitive conflict task positive network (i.e. extrinsic mode network;
335 EMN) (Hugdahl., 2015) and the task negative network (i.e. default mode network; DMN)
336 (Raichle et al., 2001) evoked by the multi-source interference task (MSIT; Bush et al., 2003).
337 MSIT features a difficult condition where there is conflict between the position and identity of a
338 number that must be selected and an easy condition, where there is no conflict. The neural

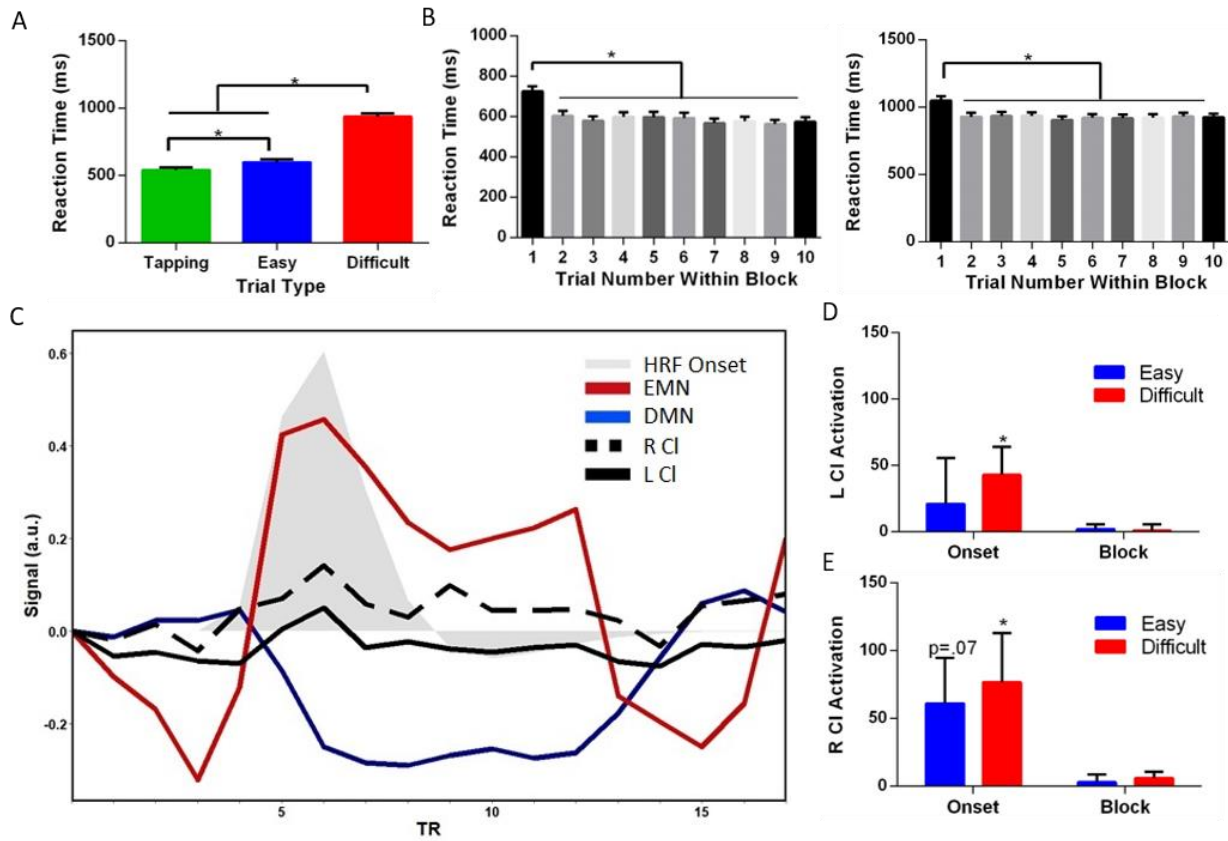


Fig 4. Claustrum activates when cognitive load peaks. A) Reaction time performance in MSIT varies based on the condition, with the difficult blocks having the slowest reaction times. B) The first trial of the easy and difficult conditions is behaviorally unique from the other trials in the block. As a result, we modelled the onset separately from the remainder of the block for difficult and easy conditions in an adaptive model to capture individual variability. C) Time course of claustrum shows a response during the difficult task, but exclusively during the onset of that task. The contrast map localizing DMN and the contrast map localizing EMN (Fig 3A) were used as regions of interest. We extracted the signal from these areas along with the left and right claustrum averaged over every difficult condition for all subjects. The task starts at the third TR and ends at the 11th. TRs 1:2 and 12:15 show the control condition. We under laid a canonical hemodynamic response function (HRF) convolved with an onset of duration 3.2 second (two trials in duration). D) Left and E) right claustrum activates at the onset – but not block – of the difficult MSIT condition. Error bars show standard error of the mean. * represents a p value of <0.05. HRF=Hemodynamic Response Function, CI=claustrum, DMN=default mode network, EMN=extrinsic mode network.

339 activation associated with these conditions are contrasted to a motor and visual control
 340 condition. Examining the difficult-block greater than easy-block contrast identified the standard
 341 EMN, including fronto-parietal network (FPN), dorsal-attention network (DAN), and dorsal ACC
 342 (Fox et al., 2005; see Fig 3A and Table S5). The easy-block greater than difficult-block contrast
 343 identified the DMN. We found left and right claustrum functional connectivity overlapped with

344 more than half of the regions in both the extrinsic mode and default mode networks (Fig 3C).
345 Additionally, every major region with significant task activation was also functionally connected
346 to the claustrum (i.e. at least partially overlapped).

347 *MSIT performance*

348 Consistent with past literature, we found a main effect of reaction time (RT; $F_{1.76,56.38}=374.2$,
349 $p<0.05$), with RT for tapping<easy<difficult trials within the MSIT, and all trial types differed from
350 one another in terms of RT (Fig 4A).

351 When analyzing RT at the group level, we observed that RT was significantly higher for the first
352 trial of easy and difficult blocks than the remaining 9 trials, a finding observed in many cognitive
353 tasks (Blais et al., 2014) (Fig 4B) (Difficult: $F_{6.27,200.7}=44.17$, $p<0.05$; Easy: $F_{4.33,138.5}=25.43$,
354 $p<0.05$). These data suggest that after the first trial participants transition into a stable level of
355 performance and therefore the beginning or onset of difficult and easy conditions should be
356 modelled separately from the block (creating 5 conditions to be modelled: tapping, easy onset,
357 easy block, difficult onset, difficult block). While at the group level a transition from high initial RT
358 for the first trial to a near-mean RT by the second trial occurred, we observed variability within
359 subjects and found that subjects sometimes took longer to transition into this stable level of
360 performance where RT was close to the block mean. That is, for a given block of stimuli some
361 subjects had clearly slower RTs for both the first and second/third trials relative to the RT mean.
362 In order to capture this inter- and intra- individual variation as well as possible, we allowed the
363 duration of the easy onset and difficult onset to vary based on RT of the subject. For start time
364 of onset events, we used the beginning of the task block. For duration, we created an adaptive
365 program that defined the duration of the easy onset and difficult onset events as either the first
366 trial RT, the second trial RT (+ trial one duration of 1.6 s), or the third trial RT (+ trial one and
367 two duration of 3.2 s). The choice of which trial RT to use as duration (one, two, or three) was
368 based on the z scores for each of these trials using the mean calculated for each subject of the

369 same task type (difficult or easy) for the fourth to tenth trials over all runs. Specifically, trial one
370 RT was the onset duration if trial two had a z-score \leq to 0.5; trail two was used if trail one RT
371 had a z score $>$ 0.5 and if trial three had a z score \leq to 0.5; trial three was used if these prior
372 stipulations were not met. Hence, the onset event duration was adapted based on individual
373 level behavioral data, which ensured that for each subject we captured the period of greatest
374 cognitive conflict, reflective of the switch to a new task. The block of the task type was modelled
375 as the remainder of the block not included in onset. We used this hybrid event-block design for
376 all task analysis of MSIT and found activations and deactivations for the onset and block for
377 both easy and difficult conditions across the brain (Table S5, S6, S7, and S8).

378 *Clastrum responds to the switch to task onset*

379 The onset of tasks involved switching from performing a tapping visuomotor control task to
380 performing either a no-cognitive conflict easy task or a cognitive conflict difficult task. When
381 analyzing claustrum activation independent of surrounding insula and putamen using this RT
382 adaptive model, we found that claustrum significantly responded to difficult onset (Left: $t_{32}=2.1$,
383 $p<0.05$; Right: $t_{32}=2.2$, $p<0.05$), though the same effect was not found for easy onset (Left:
384 $t_{32}=0.6$, $p>.2$; Right: $t_{32}=1.8$, $p=0.07$, Fig 4D-E). However, we found no evidence that claustrum
385 responded to the difficult block nor to the easy block and found no evidence that claustrum
386 responded to difficult onset more than to easy onset (all $p>0.2$). This strongly suggests that
387 claustrum is involved in the switch to active cognitive control, rather than maintaining
388 performance during high cognitive demand.

389 **Discussion**

390 In this study, we provide a novel approach we term Small Region Confound Correction to detect
391 the activity and functional connectivity of the human claustrum. In doing so, we find that the
392 claustrum is strongly functionally connected to cingulate and prefrontal cortices at rest and that

393 there is considerable overlap between claustrum connectivity maps and cognitive task-related
394 networks. Supporting a role of the claustrum in cognition, we show that claustrum is activated at
395 the onset of a demanding cognitive task, which is also associated with the onset of EMN
396 engagement.

397 We isolated a distinct claustrum signal by regressing putamen and insula signals from the
398 claustrum. This is a conservative approach, assuming that any linear relationship between
399 claustrum and insula/putamen is a result of a partial volume effect, defined as signal from
400 outside structures being erroneously incorporated into an ROI by virtue of the ROI only
401 occupying a portion of a measured volume (Dukart and Bertolino, 2014; Du et al., 2014). Thus,
402 while our approach mitigates the partial volume effect, a degree of true claustrum signal may be
403 dampened. However, our resting state analyses showing similar RSFC across left/right
404 claustrum, similar RSFC over two datasets, bilateral claustrum RSFC, and task response of the
405 claustrum, argue that the signal that we do derive is sufficiently robust. This approach offers a
406 generalizable method to assay the function of the claustrum, or other small or oddly shaped
407 neural structures, independent of partial volume effects from surrounding structures. SRCC
408 serves as a platform for studying a host of regions across the brain, such as the habenula
409 (Shelton et al., 2012; Hetu et al., 2016) and other thalamic association nuclei.

410 Our claustrum functional connectivity data reveals co-activation of claustrum with executive
411 cortical regions including ACC and medial PFC. This is in line with decades of neuronal tract
412 tracing studies from rodent to monkeys (for review see Mathur, 2014), and particularly in line
413 with reports in rat indicating dense claustrum connections with ACC and prelimbic PFC (Smith
414 and Alloway, 2010; White et al., 2017) and in with area 24 in the common marmoset (Reser et
415 al., 2017). The present work also indicates that claustrum is functionally connected to posterior
416 cingulate cortex, precuneus, angular gyrus, cuneus, visual cortex, and sensorimotor cortex,
417 which is in line with connections from claustrum to parietal association cortex and sensorimotor

418 cortices in rats (White et al., 2017), connections from claustrum to visual cortex in cats (LeVay
419 and Sherk, 1981) and projections from claustrum to parietal cortex in monkeys (Gamberini et
420 al., 2017). The more extensive pattern of claustrum functional connectivity observed in *3T-Rest*
421 may be a result of the larger sample size compared to *7T-Rest* (35 for 3T vs. 20 for 7T). The
422 additional functional connectivity seen in *3T-Rest* with midbrain and nucleus accumbens in
423 particular, is likely to reflect indirect connectivity as direct connections between these regions
424 are not currently strongly supported in the tract tracing literature. Alternatively, functional
425 connectivity of claustrum with these structures may reflect connections that are unique to the
426 human brain.

427 Our data also suggest involvement of the claustrum at the start of a difficult condition when new
428 rules come into play that require a change of cognitive strategy and task set instantiation
429 (Dosenbach et al., 2006). Significant claustrum activation was only observed at the switch from
430 the stimulus-response-based tapping task to the difficult condition of MSIT. As subjects
431 transitioned to this new rule set, average reaction time decreased significantly compared with
432 later trials in the task block. This transition also was met with an emergence of the task
433 positive/extrinsic mode network. A possible interpretation of these data could be that the
434 claustrum is involved in action inhibition as cognitive demand soars. However, neither
435 optogenetic inhibition nor activation of axon terminals of a major excitatory input source to the
436 claustrum, the ACC, affects motor activity in mice (White et al., 2018). Our results cannot be
437 explained through alterations in sensory binding or motor processes, which are both proposed
438 roles for the claustrum, as we observed claustrum activation when controlling for sensory input
439 and motor responses. The data also do not suggest a role for claustrum in resolving cognitive
440 conflict, as claustrum did not show sustained activation during the block of the difficult condition,
441 whereas the EMN did (Fig 3A).

442 In summary, we showed that even at 1.5mm spatial resolution attained with 7T fMRI, the
443 claustrum BOLD signal bears unsettling similarity with the insula and putamen, and an
444 additional processing step of SRCC allowed us to isolate a claustrum signal independent of the
445 surrounding regions. Using this method we find that in in the human – like in other species – the
446 claustrum has wide-ranging cortical connectivity, including default and extrinsic mode networks,
447 to sensory regions. Additionally, we show that claustrum activity peaks when switching to a
448 cognitive conflict task and that this activity could not be explained by changes in consciousness
449 or sensorimotor processing. These data broadly support a role of the claustrum in cognitive
450 control and are consistent with recent studies in mice (White et al., 2017).

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