Bottom-up sensory processing can decrease

² activity and functional connectivity in the

3 default mode like network in rats.

- 4 Rukun Hinz^{*1}, Lore M. B. Peeters¹, Disha Shah¹, Stephan Missault¹, Michaël Belloy¹, Verdi
- 5 Vanreusel¹, Meriam Malekzadeh¹, Marleen Verhoye¹, Annemie Van der Linden¹, Georgios A.
- 6 *Keliris**¹
- 7 ¹Bio-Imaging Lab, University of Antwerp, Belgium
- 8
- 9 Keywords: Resting state functional MRI, Functional connectivity, Visual stimulation, Default
- 10 mode network, Rats
- 11
- 12 Journal aim: Neuroimage
- 13
- 14 *Corresponding authors
- 15 Rukun Hinz
- 16 Bio-imaging lab, University of Antwerp
- 17 Campus Drie Eiken– Building Uc 1.17
- 18 Universiteitsplein 1 -2610 Wilrijk Belgium
- 19 Tel. +3232652786
- 20 Email: Rukun.Hinz@uantwerpen.be
- 21
- 22

or

- 23
- 24 Prof. Georgios Keliris
- 25 Bio-imaging lab, University of Antwerp
- 26 Campus Drie Eiken– Building Uc 1.07
- 27 Universiteitsplein 1 -2610 Wilrijk Belgium
- 28 Tel. +32322652772
- 29 Email: Georgios.Keliris@uantwerpen.be
- 30

31 Abbreviations

32	AC	Auditory cortex				
33	BOLD	Blood oxygen level dependent				
34	BVS	Blocked visual stimulation				
35	Cg	Cingulate cortex				
36	CVS	Continuous visual stimulation				
37	DMLN	Default mode like-network				
38	DMN	Default mode network				
39	FC	Functional connectivity				
40	fMRI	Functional magnetic resonance imaging				
41	FOV	Field of view				
42	FWE	Family wise error				
43	GLM	General linear model				
44	HRF	Hemodynamic response function				
45	ICA	Independent component analysis				
46	ISO	Isoflurane				
47	LGN	Lateral geniculate nucleus				
48	MD	Matrix dimensions				
49	MED	Medetomidine				
50	PtA	Parietal association cortex				
51	ROI	Regions of interest				
52	RS	Retrosplenial cortex				
53	RSB	Resting state baseline scan				
54	rsfMRI	Resting-state functional magnetic resonance imaging				
55	RSNs	Resting state networks				
56	SC	Superior colliculus				
57	SS	Somatosensory cortex				
58	ST	Slice thickness				
59	TE	Echo time				
60	TR	Repetition time				
61	VC	Visual cortex				

62 Abstract

63 The default mode network is a large-scale brain network that is active during rest and internally 64 focused states and deactivates as well as desynchronizes during externally oriented (top-down) attention demanding cognitive tasks. However, it is not sufficiently understood if unpredicted 65 66 salient stimuli, able to trigger bottom-up attentional processes, could also result in similar reduction 67 of activity and functional connectivity in the DMN. In this study, we investigated whether bottom-68 up sensory processing could influence the default mode like network (DMLN) in rats. DMLN 69 activity was examined using block-design visual functional magnetic resonance imaging (fMRI) 70 while its synchronization was investigated by comparing functional connectivity during a resting 71 versus a continuously stimulated brain state by unpredicted light flashes. We demonstrated that 72 activity in DMLN regions was decreased during visual stimulus blocks and increased during 73 blanks. Furthermore, decreased inter-network functional connectivity between the DMLN and 74 visual networks as well as decreased intra-network functional connectivity within the DMLN was 75 observed during the continuous visual stimulation. These results suggest that triggering of bottom-76 up attention mechanisms in anesthetized rats can lead to a cascade similar to top-down orienting 77 of attention in humans and is able to deactivate and desynchronize the DMLN.

78 **1. Introduction**

79 The brain is a complex network consisting of functionally interconnected regions that dynamically 80 communicate with each other. Part of these interactions can be observed non-invasively using 81 resting-state functional magnetic resonance imaging (rsfMRI) (Damoiseaux et al., 2006; Salvador 82 et al., 2005; van den Heuvel and Hulshoff Pol, 2010). This technique relies on the detection of low 83 frequency fluctuations (0.01-0.2 Hz) in the blood oxygen level dependent (BOLD) signal while the 84 subject is at rest, *i.e.* not performing any task. The coordinated fluctuations in the signals of 85 anatomically separated regions have been shown to reflect intrinsic brain networks and evidence 86 suggests that they correspond to spontaneous neuronal activity (Krishnan et al., 2018; Ma et al., 87 2016; Petridou et al., 2013). The regions that show temporally highly correlated activity are 88 considered to be functionally connected and are referred to as resting state networks (RSNs) 89 (Friston, 2011).

90 Since its discovery, rsfMRI has been widely used in human research to study RSNs in the healthy 91 brain as well as their alterations in neuropathologies (Greicius, 2008; Hull et al., 2017; Zhou et al., 92 2017). More recently, comparable RSNs have also been detected in rodents (Gozzi and Schwarz, 93 2016; Jonckers et al., 2011; C.P. Pawela et al., 2008; Sierakowiak et al., 2015). This finding has 94 been very important as it opened a new window of pre-clinical investigations in (genetic) animal 95 models of disease that can be investigated with different modalities at multiple scales, providing 96 additional information about the underlying mechanisms (Nestler and Hyman, 2010; Trancikova 97 et al., 2011). In addition, pre-clinical rsfMRI shows great potential in identifying early biomarkers 98 for multiple neuropathologies and can be used as an excellent theranostic tool (Bertero et al., 2018; 99 Li et al., 2017; Shah et al., 2016, 2013). However, the translation and interpretation of RSNs 100 between rodents and humans remains challenging, among other reasons, due to the differences in 101 anatomy, physiology and the required use of anesthesia in rodents (Pan et al., 2015). It is therefore of utmost importance to investigate and improve our understanding of specific rodent RSNs thathave been suggested to be homologous to RSNs in humans.

104 A RSN that has raised a lot of interest in humans is the default mode network (DMN), which has 105 been shown to be most active during rest and internally focused tasks and less active during 106 externally oriented attention demanding cognitive tasks (Greicius et al., 2003). Thus, it has been 107 classified as a "task-negative network" and has been shown to alternate its activity with "task-108 positive networks" (Fox et al., 2005). The DMN is thought to play a fundamental role in self-109 referential thought, mind-wandering, internally-oriented cognition, and autobiographical memory 110 (Lin et al., 2017). In humans, this network consists of regions in the anterior pre-frontal cortex, 111 posterior cingulate cortex/retrosplenial cortex (precuneus), hippocampal formation, medial and 112 lateral parietal regions (Buckner et al., 2008; Laird et al., 2009; Liska et al., 2015). A default mode 113 like network (DMLN) suggested to be homologous to the human DMN has also been identified in 114 rodents (Lu et al., 2012; Stafford et al., 2014). This DMLN comprises comparable regions, *i.e.* 115 orbital cortex, prelimbic cortex, cingulate cortex, temporal association cortex, auditory cortex, 116 posterior parietal cortex and parietal association cortex, retrosplenial cortex and hippocampus. 117 Furthermore, besides anatomical similarities, few studies could indicate functional similarities such 118 as the higher activity of the DMLN during rest vs task and the anti-correlation relationship of the 119 DMLN with the task positive network (Rohleder et al., 2016; Schwarz et al., 2013).

In recent years, multiple human studies have shown that functional connectivity (FC) within the DMN is decreased when subjects are in a higher attentive and cognitive brain state associated with performing an internally guided (top-down) attention-demanding task (Elton and Gao, 2015; Fransson, 2006; Gao et al., 2013; Marrelec and Fransson, 2011). It is thought that this internally guided attention to external sensory input can suppress other internal processes associated with the DMN, resulting in this network's inactivation and relative disconnection (Gao et al., 2013; Mayer 126 et al., 2010). However, it is not vet sufficiently understood if attentional guidance by externally 127 driven factors (bottom-up) could also result in similar reduction of activity and connectivity in the 128 DMN. In humans, some studies suggested that the DMN is not deactivated by simple sensory 129 processing (Greicius et al., 2003). It should be noted, however, that the stimulus design in these 130 studies was simple and predictable and thus not expected to continuously drive bottom-up attention. 131 Interestingly, a study using simple but unpredictable visual stimuli could dynamically activate 132 attention network and DMN indicating their interaction during stimulus-driven processes of 133 attention (Hahn et al., 2007).

134 Neurophysiological experiments in the past few years suggest that top-down and bottom-up 135 processes share overlapping neural systems and in particular the employment of the prefrontal and 136 parietal network (for a review see (Katsuki and Constantinidis, 2014)). We conjectured, that 137 similarly to top-down, bottom-up attention triggering stimuli could also deactivate DMN and 138 reduce its connectivity. To test this hypothesis, we performed fMRI experiments in anesthetized 139 rats driven by randomized (unpredictable) continuous visual stimulation (CVS) and compared 140 DMLN activity and connectivity with a resting state baseline scan (RSB) and a blocked visual 141 stimulation (BVS) design.

142 **2. Material and Methods**

143

2.1. Animals and ethical statement

In this study, we used male Long Evans wild type rats (N=12) of 4 months of age (Long Evans, Janvier). Rats were kept under a normal day/night cycle (12/12) with an average room temperature of 20-24°C and 40% humidity. Furthermore, rats were group housed and had *ad libitum* access to standard rodent chow and water. One animal was excluded from the analysis due to the detection of unilateral ventricular enlargement. All procedures were performed in accordance with the European Directive 2010/63/EU on the protection of animals used for scientific purposes. The

150 protocols were approved by the Committee on Animal Care and Use at the University of Antwerp,

151 Belgium (permit number: 2015-50), and all efforts were made to minimize animal suffering.

152

2 2.2. Animal preparation and anesthesia

153 Rats were first anesthetized using 5% isoflurane for induction and 2% isoflurane for maintenance 154 (IsoFlo, Abbott Illinois, USA) in a mixture of 70% N₂ and 30% O₂. Animals were head fixed in 155 the scanner using bite- and ear-bars and ophthalmic ointment was applied to the eyes. As soon as 156 the animal was fixed in the scanner, medetomidine anesthesia (Domitor, Pfizer, Karlsruhe, 157 Germany) was administered via a subcutaneous bolus of 0.05 mg/kg and isoflurane concentration 158 was decreased to 0% over a time period of 5 minutes. Continuous subcutaneous infusion of 159 medetomidine anesthesia of 0.1 mg/kg.h was started 15 minutes post bolus injection. Functional 160 MRI scans were acquired starting from 30 min post-bolus injection until 1h05 min post bolus 161 injection. The physiological status of the animals was monitored throughout the entire imaging 162 procedure. Respiratory rate was obtained from a pressure sensitive pad (MR-compatible Small 163 Animal Monitoring and Gating system, SA Instruments, Inc.) with a sampling rate of 225 Hz. Body temperature was closely monitored using a rectal thermistor and was maintained between (37.0 \pm 164 165 0.1 °C) using a feed-back controlled warm air heat system (MR-compatible Small Animal Heating 166 System, SA Instruments, Inc.). Furthermore, blood oxygenation was recorded using a pulse 167 oxygenation meter (MR-compatible Small Animal Monitoring and Gating system, SA Instruments, 168 Inc.) with a sampling rate of 450 Hz. After imaging procedures, animals received a subcutaneous 169 bolus injection of 0.1 mg/kg atipamezole (Antisedan, Pfizer, Karlsruhe, Germany) to counteract 170 the effects of medetomidine and were placed in a recovery box with infrared heating for post-scan 171 monitoring until the animal was fully awake.

172 **2.3. MRI**

173 All imaging procedures were performed with Paravision 6.0 using a 9.4 T BioSpec MR system 174 (Bruker, Germany) with an active decoupled rat quadrature surface coil (Rapid biomedical, 175 Germany) and a 98 mm diameter quadrature volume resonator for transmission (Bruker, Germany). 176 First, three orthogonal anatomical multi-slice Turbo RARE T2-weighted images (field of view 177 (FOV): (30x30) mm², matrix dimensions (MD): [256x256], 12 slices, Slice thickness (ST): 0.9 178 mm, echo time (TE)/ repetition time (TR): 33/2500 ms, RARE factor: 8) were acquired to allow 179 reproducible flat skull positioning of coronal slices. Then, a coronal anatomical reference scan was 180 acquired using a multi-slice Turbo RARE T2-weighted sequence (FOV: (30x30) mm², MD: 181 [256x256], 12 slices, ST: 0.9 mm, TE/TR: 33/2500 ms, RARE factor: 8), covering the brain from 182 3.3 mm anterior to bregma to 7.5 mm posterior to bregma (suppl. figure 1). Next, a B₀ field map 183 was acquired to assess field homogeneity, followed by local shimming in a rectangular volume of 184 interest in the brain to correct for the measured inhomogeneities. The RSB scan had an identical 185 geometry to the reference scan and was acquired using a T2*-weighted single shot echo planar 186 imaging (EPI) sequence (FOV: (30x30) mm², MD: [128x98], 12 slices, ST: 0.9 mm, TE/TR: 187 18/2000 ms) resulting in a voxel dimension of (0.234 x 0.313 x 0.9) mm³ and a total scan duration 188 of 10 min (300 volumes). Subsequently, random continuous light flickering (see section 2.4) was 189 turned on and after 1 minute the CVS scan was acquired using the same parameters as the RSB 190 (300 volumes). Last, a BVS data set was acquired using the same parameters and sequence (380 191 volumes; Figure 1A).

192

2.4. Visual stimulation

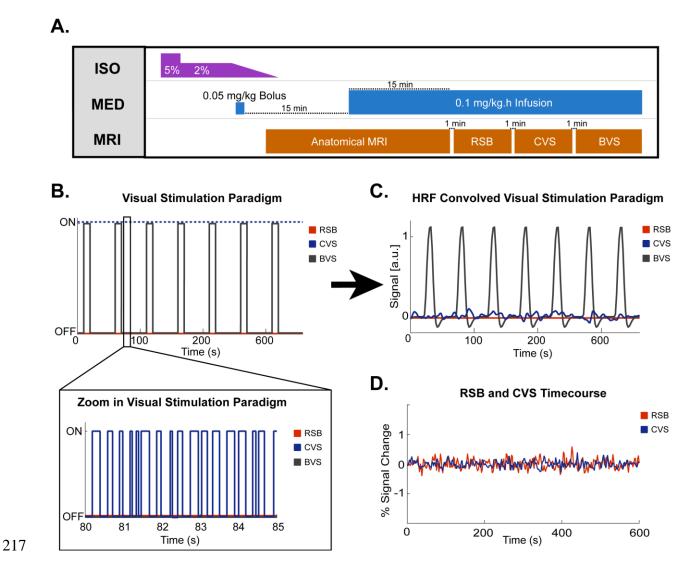
193 Visual stimulation with flickering light was presented using a white LED coupled to a fiber-optic 194 cable, which was centrally placed in front of the animal's head. The LED light was controlled by 195 a voltage-gated device to control the triggering of the LED light (ON-OFF) driven by a RZ2 196 BioAmp Processor (Tucker-Davis, Alachua). Stimulus timing and alignment to the MR imaging 197 was achieved by TTL pulses sent by the scanner at the beginning of every volume of the fMRI198 scan.

199 2.4.1. Continuous visual stimulation

200 The CVS was used to induce a continuous visual sensory drive with randomization of light pulses 201 to avoid sensory adaptation effects and to constantly trigger bottom-up attention mechanisms. This 202 condition can be assumed to create a steady brain state similar to rest and thus could be analyzed 203 the same way as the RSB scan (Figure 1B-D). The CVS was initiated 1 min before the start of the 204 acquisition to avoid the detection of the initial transient increase of brain activity, which occurs 205 when the visual stimulation is turned on. The stimulation paradigm was controlled via Matlab code 206 (MATLAB R2014a, The MathWorks Inc. Natick, MA, USA) using an USB to serial port 207 connection (IC-232A, Rotronic) and consisted of continuous short pulses of light and inter-light 208 intervals (both with a random duration between 50-250 ms) (Figure 1B). Convolution of the CVS 209 stimulus paradigm with the canonical hemodynamic response function (HRF) in SPM12 was 210 performed in order to demonstrate that the expected signal fluctuations from CVS are negligible in 211 comparison to those expected by the BVS paradigm (Figure 1C).

212 2.4.2. Block design visual stimulation

To evoke BOLD responses from the visual system, fMRI scan was acquired during a block design paradigm with a visual stimulation frequency of 4 Hz, duty cycle 50% (125 ms ON/OFF), an initial OFF block of 10 seconds followed by 15 ON/OFF blocks of 10 s and 40 s respectively (Figure 1B-C).



218 Figure 1. Scanning protocol and visual stimulation paradigms. A. Scanning protocol. For 219 handling procedures, animals were first anesthetized using 5% isoflurane (ISO) for induction 220 followed by 2% ISO for maintenance. Once the animal is fixated in the scanner bed, a bolus of 221 0.05mg/kg of medetomidine (MED) was administered and ISO anesthesia was gradually 222 decreased to 0% ISO. After 15 minutes post bolus injection, a continuous infusion of 0.1 223 mg/kg.h MED was administered to the animal. For imaging procedures, first a set of 224 anatomical Turbo RARE T2 scans were acquired and shimming procedures were performed. 225 Next, 30 min post bolus injection a resting state baseline (RSB) scan was acquired. 226 Subsequently, continuous visual stimulation (CVS) paradigm was turned on and after one

227 minute the CVS scan was acquired. Lastly, after a recovery time of 1 minute a block design 228 visual stimulation (BVS) fMRI scan was acquired. B. Visual stimulation paradigm of RSB, CVS 229 and BVS scan indicating when visual stimuli was turned on or off. C. Haemodynamic Response 230 Function (HRF) convolved visual stimulation paradigm of RSB, CVS and BVS scan showing 231 predicted BOLD signal response in arbitrary units (a.u.) from each condition. D. Example of 232 acquired RSB and CVS normalized BOLD signal time course in the visual network.

233

2.5. Breathing rate processing

Breathing rate pressure signals were analyzed to investigate the potential influence of visual stimulation on the animals' physiology. First, breathing rates were calculated for each volume by calculating the median period of the breathing cycle between the initial and the following volume and inverting this value to breaths per minute. For resting state data, averaged breathing rate from the complete scans were compared between the RSB and CVS condition using a paired t-test (p<0.05). For the BVS scans, averaged breathing rate over all visual stimulation blocks was compared ten seconds before with ten seconds during visual stimuli using a paired t-test.

241 2.6. MRI Processing

All data processing was performed using SPM 12 software (Statistical Parametric Mapping, <u>http://www.fil.ion.ucl.ac.uk</u>), REST toolbox (REST1.7, <u>http://resting-fmri.sourceforge.net</u>) and GIFT toolbox (Group ICA of fMRI toolbox version 3.0a: http://icatb.sourceforge.net/).

Pre-processing consisted of realignment of the data towards the first image of each scan using a 6parameter (rigid body) spatial transformation, normalization towards a study specific EPI template using a global 12-parameter affine transformation, followed by a non-linear transformation. Finally, data were smoothed in-plane using a Gaussian kernel with a full width at half maximum of twice the voxel size (0.458 x 0.626 mm). rsfMRI data were then further band pass filtered between 0.01-0.2 Hz using REST toolbox.

251

- 2.6.1. Functional connectivity
- 252 2.6.1.1. ICA components

253 To obtain the RSNs from the rsfMRI data, an independent component analysis (ICA) was 254 performed on RSB data. First, movement was regressed out of the data based on the estimators 255 from the realignment procedure using the REST toolbox. Next, ICA was performed in GIFT using 256 the Infomax algorithm with predefined number of 15 components as used previously (Jonckers et 257 al., 2011). Components representing functional networks were identified by comparison to 258 previous observed RSNs and discarding a minority of artefactual components (e.g. only edge of 259 the brain) by careful visual inspection (Gozzi and Schwarz, 2016; Jonckers et al., 2011; Lu et al., 260 2012; C.P. Pawela et al., 2008; Sierakowiak et al., 2015). Afterwards, the selected mean z-scored 261 RSNs group components were thresholded at z-score > 1 to obtain a mask of each network.

262

2.6.1.2. ICA-based inter-network connectivity

263 To evaluate inter-network connectivity, network masks were used as regions of interest (ROI) in 264 correlation-based analysis per subject. Pairwise Pearson correlation coefficients between the 265 average BOLD time series of each network ROI were calculated and Fisher z-transformed using 266 an in-house Matlab program. Mean Fisher z-transformed inter-network correlation matrices were 267 calculated for RSB and CVS conditions. Statistical analysis between conditions was performed 268 using a repeated measures 2-way ANOVA (p<0.05, Sidak correction for multiple comparisons). 269 To calculate the FC between the DMLN and visual network, z-transformed correlation values of 270 the cingulate-retrosplenial, hippocampal and temporal-prefrontal with visual network were 271 averaged. Differences in correlation between the two conditions were then investigated using 272 pairwise t-test (p<0.05).

273

2.6.1.3. ROI-based intra- and inter-network connectivity

274 To investigate within and between network connectivity differences, specific anatomically defined 275 bilateral ROIs were selected in the DMLN (based upon the RSB ICA networks: cingulate cortex 276 (Cg), retrosplenial cortex (RS), parietal association cortex (PtA), temporal association cortex 277 (TeA)) as well as in the visual system (based upon the visual stimulation correlated ICA component 278 of the BVS: lateral geniculate nucleus (LGN), superior colliculus (SC), visual cortex (VC)) (suppl. 279 figure 2). Furthermore, a ROI in the primary somatosensory cortex (SS) was added as a control 280 region. Pairwise Pearson correlation coefficients between the average BOLD time series of each 281 ROI were calculated and Fisher z-transformed. Mean z-transformed ROI connectivity matrices 282 were calculated for RSB and CVS conditions. To compare RSB and CVS condition, statistical 283 analysis between conditions was performed using a repeated measures 2-way ANOVA (p<0.05, 284 Sidak correction for multiple comparisons).

285

2.6.1.4. Seed-based analysis

286 To investigate how the full-brain FC of the Cg is influenced by CVS, seed-based analysis was 287 performed. To this end, the mean time course of Cg was used as a predictor in a General linear 288 model (GLM) and motion parameters were included as covariates. Each subject's FC between Cg 289 and DMLN or visual network was extracted by using the respective binarized masks derived from 290 the ICA and calculating the mean T-values within this mask for each individual subject. The 291 DMLN network mask was constructed by the union of the cingulate-retrosplenial network mask, 292 hippocampal network mask and temporal-prefrontal network mask. Statistical analysis to compare 293 differences between the conditions was performed by means of a paired t-test (p < 0.05).

294 2.6.2. Block design fMRI

BVS data were processed using a group ICA approach using the GIFT toolbox with an Infomax algorithm and 15 predefined components. Group ICA analysis was chosen instead of the standard GLM to increase sensitivity towards detecting responding brain regions by not relying on 298 predetermined BOLD HRF functions (Calhoun et al., 2009). The components temporal correlation 299 with the visual stimulation paradigm was then calculated using R-square statistic in the GIFT 300 toolbox. The component with highest temporal correlation with the visual stimulation paradigm 301 were regarded as responding regions. Event related BOLD responses were extracted from a binary 302 mask of responding regions, which are either correlated (z-score > 1) or anti-correlated (z-score < 303 -1) with the ICA time course. To confirm our results, GLM analysis was performed for each subject 304 within the ICA component mask (z-score > 1 and < -1) of the responding regions using a canonical 305 HRF function in SPM 12. The model was set to either detect a positive BOLD response during the 306 stimulation period, to locate visual responding regions, or during the rest period, to detect regions 307 which are more active during rest vs stimulation period i.e. DMLN activity. One sample t-test 308 (p<0.001, uncorrected for multiple comparisons) was performed to evaluate BSV group response.

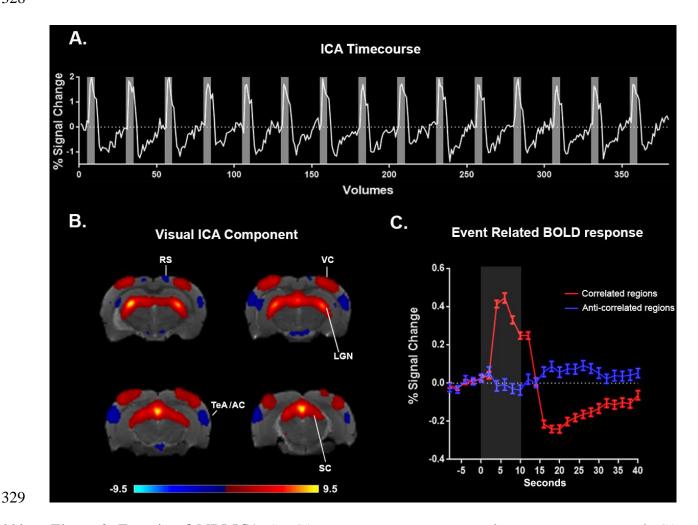
309 **3. Results**

310

3.1. Block-design visual stimulation

311 To investigate whether activity in the DMLN of anesthetized rats is changing between visual 312 sensory stimulation and rest periods, we analyzed the block-design fMRI experiment. To this end, 313 we performed group-ICA analysis and sorted the components based on their temporal correlation 314 with the visual stimulation paradigm. The component with the highest correlation ($R^2=0.64$) 315 showed clear BOLD signal increases during the ON-blocks indicating a strong visual activation of 316 multiple responding regions (Figure 3A). The mean group statistical ICA map (z-score > 1 and < -317 1) of this component revealed that the activated regions were, as expected, areas involved in visual 318 processing, *i.e.* the LGN, SC, VC (Figure 3B). Event related analysis of all regions which are 319 correlated with the ICA time course demonstrated that visual stimulation evoked a positive BOLD 320 response as expected (Figure 3C). Interestingly, we also observed regions such as the temporal 321 association cortex/auditory cortex (TeA/AC) and RS, being anti-correlated to the ICA component's time course, indicating higher BOLD signal during rest vs stimulation (Figure 3B). Event related analysis of anti-correlated regions with the ICA time course showed a subtle BOLD signal decrease during visual stimulation which rebounded and increased during blanks (Figure 3C). To confirm our results, GLM modeling was applied and detected increased BOLD response during visual stimulation in LGN, SC and VC as well as increased BOLD response during blanks in TeA/AC (suppl. figure 3).





330 *Figure 3. Functional MRI ICA. A. ICA Time course. Time course from mean group visual ICA* 331 *component with highest temporal correlation* ($R^2=0.64$) *with the visual stimulation paradigm (grey*

332 blocks). B. Visual ICA Component. Mean statistical group ICA map (z-score -1 < and > 1)

333 demonstrating areas which are correlated to the mean ICA component's time course involved in 334 visual processing, i.e. visual cortex (VC), lateral geniculate nucleus (LGN) and superior colliculus 335 (SC). Regions demonstrating a BOLD time course that is anti-correlated with the mean ICA 336 component's time course, meaning higher BOLD signal during rest than during visual stimulation, 337 were observed in DMLN regions i.e. temporal association cortex/auditory cortex (TeA/AC) and 338 retrosplenial cortex (RS). C. Event related response of the regions correlating (Red) and anti-339 correlating (Blue) to the ICA component's time course. Grey block indicates visual stimulation 340 period

341

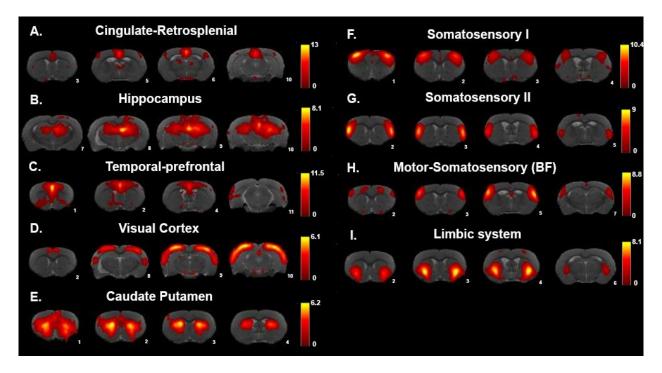
3.2. Resting state vs Continuous Visual Stimulation

342 **3.2.1.** Decreased intra- and inter-network connectivity during CVS

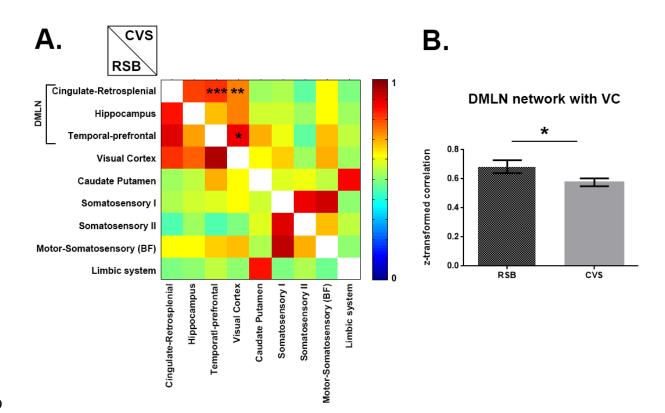
343 To identify potential changes in FC induced by our randomized continuous stimulation paradigm, 344 we first performed group-ICA for the RSB condition and identified commonly observed RSNs. 345 Functionally relevant components included three subnetworks of the DMLN: a cingulate-346 retrosplenial network (with cingulate cortex, retrosplenial cortex and parietal association cortex), a hippocampal network (with subiculum, dentate gyrus, CA1 and CA3), and a temporal-prefrontal 347 348 network (with orbito-frontal cortex, prelimbic cortex and temporal association cortex). 349 Furthermore, ICA analysis detected a visual network, a caudate putamen network, a primary and 350 secondary somatosensory network, a barrel field network and a limbic network (including the 351 amygdala and ventral striatum). Next, to evaluate intra- and inter-network connectivity, we 352 threshold each ICA-component map (z-values > 1), extracted the time-courses and performed 353 pairwise correlations for both the RSB and CVS conditions (Figure 5). Statistical analysis 354 comparing RSB and CVS condition using a repeated measures 2-way ANOVA (multiple 355 comparison correction Sidak p<0.05) detected a significantly decreased intra-DMLN network 356 correlation in the CVS condition (*i.e.* subnetworks cingulate-retrosplenial and temporal-prefrontal

- 357 (p<0.001)) as well as decreased inter-network correlations between DMLN subnetworks and the
- 358 visual network (*i.e.* cingulate-retrosplenial network with visual network (p=0.002) and temporal-
- 359 prefrontal network with visual network (p=0.027).

360



361 Figure 4. ICA of Resting State Baseline (RSB) condition. Mean group statistical ICA maps (z-362 score>1) revealed nine functionally relevant components. A.-C. The DMLN in rats split up in three 363 subcomponents i.e. A. Cingulate-Retrosplenial Network (with cingulate, retrosplenial cortex and 364 parietal association cortex), B. Hippocampal network (with subiculum, dentate gyrus, CA1 and 365 CA3) and C. Temporal-prefrontal network (with orbito-frontal cortex, prelimbic cortex and 366 temporal association cortex). D. Visual network (with visual and somatosensory cortex). E. Caudate Putamen network. F.-G. Primary and secondary somatosensory network. H. Barrel field 367 368 network. I. Limbic network (with amvgdala and ventral striatum).



370 Figure 5. Inter-network functional connectivity. A. Pairwise z-transformed Pearson correlation 371 matrix of network components' time courses of the resting state baseline (RSB) (Bottom) and of the 372 continuous visual stimulation (CVS) condition (Top). Stars indicate significant differences found 373 between the two conditions with repeated measures 2-way ANOVA (p<0.05, with Sidak multiple 374 comparison correction). For CVS, significant decreased inter-network connectivity was found 375 between subnetworks of the DMLN (i.e cingulate-retrosplenial network and temporal-prefrontal 376 network) and between DMLN subnetworks and visual network. Color bar represents z-values. 377 *p<0.05, **p<0.005, ***p<0.001 B. Inter-network connectivity between DMLN and Visual 378 network (average of connectivity between cingulate-retrosplenial network and visual network, 379 hippocampal network and visual network, and temporal-prefrontal network and visual network). 380 Statistical analysis using a paired t-test detected a significant decreased inter-network connectivity 381 in the CVS condition as compared to the RSB condition (p < 0.05).

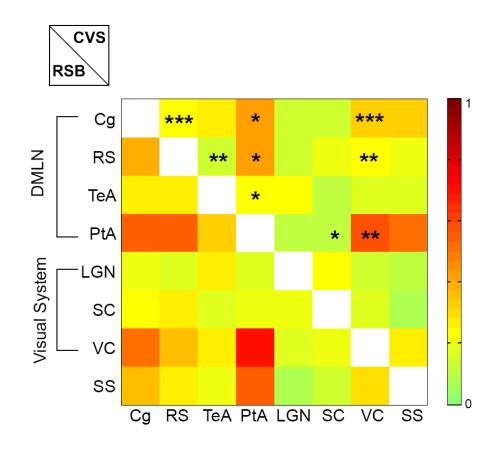
382

383 Since CVS induced decreased inter-network correlation between the DMLN subnetworks and the 384 visual network, we further assessed the pairwise correlation between specific ROIs of these 385 networks to zoom-in and better understand the sources of these decreases. ROIs in DMLN (i.e. Cg, 386 RS, TeA and PtA) and visual system (*i.e.* LGN, SC and VC) were selected. In addition, we included 387 the SS as a control area. Pairwise correlation between each ROI's averaged BOLD time course was 388 performed and compared between RSB and CVS condition (Figure 6). Statistical analysis using a 389 repeated measures 2-way ANOVA (Sidak multiple comparisons correction p<0.05) showed a 390 decreased correlation between the DMLN ROIs *i.e.* Cg-RS (p<0.001), Cg-PtA (p=0.005), RS-TeA 391 (p=0.002) and RS-PtA (p=0.006) as well as between DMLN and visual ROIs *i.e.* Cg-VC (p<0.001), 392 RS-VC (p=0.005), PtA-SC (p=0.036), and PtA-VC (p=0.002). None of these areas showed a 393 significant change in correlation with the SS.

394

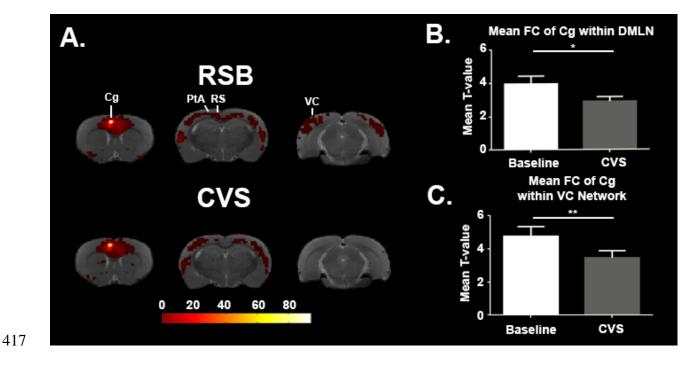
3.2.2. Voxel-based functional connectivity of Cingulate cortex

395 The cingulate cortex, a major node of both DMLN in rodents as well as DMN in humans has been 396 shown to change its activity during unpredictable stimuli (Hahn et al., 2007). Since our stimuli in the CVS condition were similarly designed to be unpredictable we selected this area for seed-based 397 398 analysis. As demonstrated in the statistical FC maps (one sample t-test, p<0.001, family wise error 399 (FWE) corrected for multiple comparisons) presented in Figure 7A, the Cg demonstrated wider 400 brain connectivity during the RSB in comparison to the CVS condition. To further quantify this 401 effect, we performed a paired t-test for the T-values within the DMLN or visual network. We found 402 that CVS condition induced a decreased correlation between Cg and DMLN as well as between Cg 403 and the visual network (Figure 7B-C).



404

405 Figure 6. ROI based analysis. A. Pairwise z-scored Pearson functional connectivity (FC) matrix 406 between time courses of ROIs of the DMLN (i.e. cingulate cortex (Cg), retrosplenial cortex (RS), 407 temporal association cortex (TeA) and parietal association cortex (PtA)), the visual system (i.e. 408 lateral geniculate nucleus (LGN), superior colliculus (SC) and visual cortex (VC)) and the 409 somatosensory cortex (SS) as a control region. Top half of the matrix represents FC of continuous 410 visual stimulation (CVS) condition. Bottom half of the matrix represents FC of resting state 411 baseline (RSB) condition. Color bar represents z-values. Stars indicate significant differences 412 found between the two conditions (diagonally symmetric positions) with repeated measures 2-way 413 ANOVA (p<0.05, with Sidak multiple comparison correction). Decreased FC was detected 414 between DMLN ROIs i.e. Cg-RS, Cg-PtA, RS-TeA and RS-PtA as well as between DMLN and visual system ROIs i.e. Cg-VC, RS-VC, Pta-SC and PtA-VC. *p<0.05, **p<0.005, ***p<0.001 415



418 Figure 7. Seed based analysis of functional connectivity with the cingulate cortex as seed region. 419 A. Statistical maps of functional connectivity (p < 0.05 with family wise error correction (FWE) for 420 multiple comparison correction) of the cingulate cortex (Cg) in the resting state baseline (RSB)421 condition (Top) and continuous visual stimulation (CVS) condition (Bottom). Decreased FC in the 422 parietal association cortex (PtA), retrosplenial cortex (RS- and visual cortex (VC) was observed 423 in the CVS condition. Color bar represents t-values. High t-values indicate high functional 424 connectivity with the seed B. Bar graph of the mean T-value with \pm SEM within the default mode 425 like network. For each subject, mean T-values were extracted within the default mode like network 426 ICA mask. Results show that CVS significantly decreased connectivity of the Cg (paired t-test, 427 p < 0.05) with the default mode like network. C. Bar graph of the Mean T-value with \pm SEM of all 428 subjects within the Visual network (VC). For each subject, mean T-values were extracted within 429 the VC network ICA mask. Results show that continuous visual stimulation (CVS) significantly 430 decreased connectivity of the Cg (paired t-test, p<0.05) with the VC network.

431 **3.3. Unaltered breathing rate during visual stimulation**

The pressure signal for breathing rate of the animals was recorded throughout the whole experiment and analyzed to assess potential changes on the general physiological state due to the visual stimulation. A paired t-test showed that there were no significant changes in breathing rate due to the visual stimulation (p=0.79) (suppl. figure 4A). Breathing rate was further compared between RSB and CVS conditions using a paired t-test (p<0.05). Likewise, CVS did not significantly alter the breathing rate (p=0.1) (suppl. figure 4B).

438 **4. Discussion**

In the current study, we investigated the impact of visual stimulation on the DMLN activity and its
FC in rats. We found that visual stimulation could deactivate nodes of the DMLN and could
decrease FC within DMLN as well as across DMLN and visual networks.

442

4.1. Block design visual stimulation induces deactivation in DMLN regions

Block design visual stimulation was performed a) to identify the visually responsive areas and b)to investigate the influence of visual stimulation on the DMLN.

445 Similar to previous visual fMRI studies in rats, we detected activation of visual processing areas 446 including LGN, SC, and VC (Van Camp et al., 2006), (Chrisopher P. Pawela et al., 2008). In 447 addition to these activating regions, we detected deactivating regions that demonstrated reduced 448 activity during visual stimulation and displayed increased activity during rest i.e. TeA/AC and RC. 449 Interestingly, these areas have been shown to be nodes of the DMLN in rats (Lu et al., 2012). This 450 finding seems in contrast to earlier human studies that did not detect significant reductions in DMN 451 activity during passive sensory processing states that have low cognitive demand e.g. flashing 452 checkerboard pattern presentation (Greicius et al., 2003). However, a number of important 453 differences in our study could explain this discrepancy. Firstly, it should be noted that the effect 454 we found in rats was very subtle. Thus, it is possible that similar effects could be present in humans 455 but not sufficiently strong to be detected during awake conditions that are potentially compromised 456 by additional processes. Previous studies directly demonstrated that the magnitude and extent of 457 the suppression depends on the difficulty of the cognitive task (Leech et al., 2011; Mayer et al., 2010). As our study was performed in anesthetized rats, although not optimal for top-down 458 459 cognitive processing, could be beneficial for identifying subtle bottom-up effects that could be 460 otherwise hindered by additional variability induced by awake conditions. An alternative 461 explanation is that in rodents passive sensory stimulation is more cognitively demanding in 462 comparison to humans. Visual stimulation in rodents is thought to increase behavior mechanisms 463 such as fear, which could be responsible for modulating the DMLN (McClearn, 1960). As previous 464 studies in human have shown, fear can readily deactivate the DMN (Marstaller et al., 2017). Our 465 observation could therefore suggest stronger cognitive involvement in rats during visual 466 stimulation (Anticevic et al., 2013).

467 **4.2. Continuous visual stimulation decreases inter- and intra-network FC of the DMLN**

We explored the reorganization of functional networks during a visually stimulated brain state induced by CVS. The CVS paradigm was specifically developed to get the animal in a visually attentive stimulated steady state and stochasticity was included to avoid habituation towards the stimulus. Firstly, ICA analysis was performed on the RSB data in order to identify the RSNs. These networks showed strong bilateral connectivity and were similar to previously described networks in rats (Hutchison et al., 2010; Jonckers et al., 2011). We then investigated how the activity and connectivity within and across these networks changed in the CVS condition.

By comparing FC between networks in both conditions, we demonstrated that CVS decreased internetwork FC between DMLN subcomponents (cingulate-retrosplenial and temporal-prefrontal networks (Lu et al., 2012)) and with the visual network. The observed decrease in inter-network connectivity during the CVS condition suggests an alteration in communication between the

DMLN and the visual network. This finding could subserve the enhancement of local, input
specific visual processing during CVS versus a higher inter-network communication during rest
conditions.

482 Further, we performed ROI based analysis focusing on areas within the DMLN. This analysis 483 demonstrated that intra-DMLN FC was also decreased. This included connections between 484 multiple major nodes of the DMLN *i.e* Cg-RS, Cg-PtA, RS-PtA and RS-TeA. Similar to activity 485 level decreases, connectivity decreases within human DMN were previously observed only with 486 tasks involving higher cognitive load (Elton and Gao, 2015; Fransson, 2006; Gao et al., 2013; 487 Marrelec and Fransson, 2011), while simple visual stimulation with a constantly flickering 488 checkerboard pattern were not able to induce such deactivation (Di et al., 2015). Interestingly, the 489 results observed in our study are more consistent with human data from subjects in a higher 490 attentive and cognitive brain state.

491

492 **5.** Conclusion

493 In summary, we demonstrated that simple yet stochastic sensory stimulation in anesthetized rats 494 could a) deactivate certain nodes of the DMLN, and b) reduce intra- and inter-DMLN network 495 connectivity simulating similar results in humans performing task involving high cognitive and 496 top-down attentional demand. We conjecture that the stochasticity of our stimulus, may play an 497 important role in consistently and continuously driving bottom-up attention triggering mechanisms. 498 Given that the bottom-up and top-down attentional systems share specific network components 499 (Katsuki and Constantinidis, 2014), we suggest that both attention mechanisms are able to 500 deactivate and reduce functional connectivity of the DMN. These results are very significant and 501 could prove immensely useful not only for our better understanding of the DMN using animal 502 models but are also very promising for being used in human patients that are anesthetized or non-

503	responsive as a result of trauma and or injury. However, to more explicitly demonstrate the link							
504	between attention systems and DMN activity and connectivity, more studies are required both in							
505	humans using stimuli specifically designed to continuously drive bottom-up attention, as well as in							
506	awake and behaving animals including more complicated cognitive tasks.							
507	Acknowledgements							
508	This research was supported by the fund of scientific research Flanders (FWO G048917N),							
509	Flagship ERA-NET (FLAG-ERA) FUSIMICE (grant agreement G.0D7651N) and University							
510	Research Fund of University of Antwerp (BOF DOCPRO FFB150340).							
511 512	Reference list							
513	Anticevic, A., Cole, M., Murray, J., Corlett, P., Wang, XJ., Krystal, J., 2013. The Role of Default							
514	Network Deactiviation in Cognition and Disease. Trends Cogn. Sci. 16, 584-592.							
515	https://doi.org/10.1016/j.tics.2012.10.008.The							
516	Bertero, A., Liska, A., Pagani, M., Parolisi, R., Masferrer, M.E., Gritti, M., Pedrazzoli, M.,							
517	Galbusera, A., Sarica, A., Cerasa, A., Buffelli, A., Tonini, R., Buffo, A., Gross, C., Pasqualetti,							
518	M., Gozzi, A., 2018. Autism-associated 16p11.2 microdeletion impairs prefrontal functional							
519	connectivity in mouse and human. Brain 141, 2055–2065.							
520	https://doi.org/10.1093/brain/awy111							
521	Buckner, R.L., Andrews-Hanna, J.R., Schacter, D.L., 2008. The brain's default network: Anatomy,							
522	function, and relevance to disease. Ann. N. Y. Acad. Sci. 1124, 1-38.							
523	https://doi.org/10.1196/annals.1440.011							
524	Calhoun, V.D., Lui, J., Adali, T., 2009. A review of group ICA for fMRI data and ICA for joint							
525	inference of imaging, genetic, and ERP data. Neuroimage 45, S163-S172.							
526	https://doi.org/10.1016/j.neuroimage.2008.10.057.A							

- 527 Damoiseaux, J.S., Rombouts, S.A.R.B., Barkhof, F., Scheltens, P., Stam, C.J., Smith, S.M.,
- 528 Beckmann, C.F., 2006. Consistent resting-state networks across healthy subjects. Proc. Natl.
- 529 Acad. Sci. 103, 13848–13853. https://doi.org/10.1073/pnas.0601417103
- 530 Di, X., Fu, Z., Chan, S.C., Hung, Y.S., Biswal, B.B., Zhang, Z., 2015. Task-related functional
- 531 connectivity dynamics in a block-designed visual experiment. Front. Hum. Neurosci. 9, 1–11.
- 532 https://doi.org/10.3389/fnhum.2015.00543
- 533 Elton, A., Gao, W., 2015. Task-positive Functional Connectivity of the Default Mode Network
- 534 Transcends Task Domain. J. Cogn. Neurosci. 27, 2369–2381.
 535 https://doi.org/10.1162/jocn_a_00859
- 536 Fox, M.D., Snyder, A.Z., Vincent, J.L., Corbetta, M., Van Essen, D.C., Raichle, M.E., 2005. The
- 537 human brain is intrinsically organized into dynamic, anticorrelated functional networks. Proc.
- 538 Natl. Acad. Sci. 102, 9673–9678. https://doi.org/10.1073/pnas.0504136102
- 539 Fransson, P., 2006. How default is the default mode of brain function?. Further evidence from
- 540 intrinsic BOLD signal fluctuations. Neuropsychologia 44, 2836–2845.
 541 https://doi.org/10.1016/j.neuropsychologia.2006.06.017
- 542 Friston, K.J., 2011. Functional and Effective Connectivity: A Review. Brain Connect. 1, 13–36.
 543 https://doi.org/10.1089/brain.2011.0008
- Gao, W., Gilmore, J.H., Alcauter, S., Lin, W., 2013. The dynamic reorganization of the defaultmode network during a visual classification task. Front. Syst. Neurosci. 7, 1–13.
 https://doi.org/10.3389/fnsys.2013.00034
- 547 Gozzi, A., Schwarz, A.J., 2016. Large-scale functional connectivity networks in the rodent brain.
- 548 Neuroimage 127, 496–509. https://doi.org/10.1016/j.neuroimage.2015.12.017
- 549 Greicius, M., 2008. Resting-state functional connectivity in neuropsychiatric disorders. Curr. Opin.
- 550 Neurol. 24, 424–430. https://doi.org/10.1097/WCO.0b013e328306f2c5

- 551 Greicius, M.D., Krasnow, B., Reiss, A.L., Menon, V., 2003. Functional connectivity in the resting
- brain: a network analysis of the default mode hypothesis. Proc. Natl. Acad. Sci. U. S. A. 100,
- 553 253–8. https://doi.org/10.1073/pnas.0135058100
- Hahn, B., Ross, T.J., Stein, E.A., 2007. Cingulate activation increases dynamically with response
- speed under stimulus unpredictability. Cereb. Cortex 17, 1664–1671.
 https://doi.org/10.1093/cercor/bhl075.Cingulate
- 557 Hull, J. V., Jacokes, Z.J., Torgerson, C.M., Irimia, A., Van Horn, J.D., Aylward, E., Bernier, R.,
- 558 Bookheimer, S., Dapretto, M., Gaab, N., Geschwind, D., Jack, A., Nelson, C., Pelphrey, K.,
- 559 State, M., Ventola, P., Webb, S.J., 2017. Resting-state functional connectivity in autism
- 560 spectrum disorders: A review. Front. Psychiatry 7. https://doi.org/10.3389/fpsyt.2016.00205
- 561 Hutchison, R.M., Mirsattari, S.M., Jones, C.K., Gati, J.S., Leung, L.S., 2010. Functional Networks
- in the Anesthetized Rat Brain Revealed by Independent Component Analysis of Resting-State
 fMRI 3398–3406. https://doi.org/10.1152/jn.00141.2010.
- 564 Jonckers, E., Van Auderkerke, J., De Visscher, G., Van der Linden, A., Verhoye, M., 2011. 565 Functional connectivity fMRI of the rodent brain: comparison of functional connectivity 566 networks **PLoS** 6, e18876. in One and mouse. rat 567 https://doi.org/10.1371/journal.pone.0018876
- Katsuki, F., Constantinidis, C., 2014. Bottom-up and top-down attention: Different processes and
 overlapping neural systems. Neuroscientist 20, 509–521.
 https://doi.org/10.1177/1073858413514136
- 571 Krishnan, G.P., González, O.C., Bazhenov, M., 2018. Origin of slow spontaneous resting-state
 572 neuronal fluctuations in brain networks. Proc. Natl. Acad. Sci. 201715841.
 573 https://doi.org/10.1073/pnas.1715841115
- 574 Laird, A.R., Eickhoff, S.B., Li, K., Robin, D.A., Glahn, D.C., Fox, P.T., 2009. Investigating the

575	Functional	Heterogeneity	of	the	Default	Mode	Network	Using	Coordinate-Based	Meta-

 576
 Analytic
 Modeling.
 J.
 Neurosci.
 29,
 14496–14505.

 577
 https://doi.org/10.1523/JNEUROSCI.4004-09.2009
 14496–14505.
 14496–14505.

578 Leech, R., Kamourieh, S., Beckmann, C.F., Sharp, D.J., 2011. Fractionating the Default Mode

- 579 Network: Distinct Contributions of the Ventral and Dorsal Posterior Cingulate Cortex to
- 580 Cognitive Control. J. Neurosci. 31, 3217–3224. https://doi.org/10.1523/JNEUROSCI.5626581 10.2011
- Li, Q., Li, G., Wu, D., Lu, H., Hou, Z., Ross, C.A., Yang, Y., Zhang, J., Duan, W., 2017. Restingstate functional MRI reveals altered brain connectivity and its correlation with motor
 dysfunction in a mouse model of Huntington's disease. Sci. Rep. 7, 1–9.
 https://doi.org/10.1038/s41598-017-17026-5
- Lin, P., Yang, Y., Gao, J., De Pisapia, N., Ge, S., Wang, X., Zuo, C.S., Jonathan Levitt, J., Niu, C.,
- 587 2017. Dynamic Default Mode Network across Different Brain States. Sci. Rep. 7, 1–13.
 588 https://doi.org/10.1038/srep46088
- Liska, A., Galbusera, A., Schwarz, A.J., Gozzi, A., 2015. Functional connectivity hubs of the
 mouse brain. Neuroimage 115, 281–291. https://doi.org/10.1016/j.neuroimage.2015.04.033
- Lu, H., Zou, Q., Gu, H., Raichle, M.E., Stein, E.A., Yang, Y., 2012. Rat brains also have a default
 mode network. Proc. Natl. Acad. Sci. 109, 3979–3984.
 https://doi.org/10.1073/pnas.1200506109
- 594 Ma, Y., Shaik, M.A., Kozberg, M.G., Kim, S.H., Portes, J.P., Timerman, D., Hillman, E.M.C.,
- 595 2016. Resting-state hemodynamics are spatiotemporally coupled to synchronized and
- 596 symmetric neural activity in excitatory neurons. Proc. Natl. Acad. Sci. 113, E8463–E8471.
- 597 https://doi.org/10.1073/pnas.1525369113
- 598 Marrelec, G., Fransson, P., 2011. Assessing the influence of different ROI Selection Strategies on

- 599 functional connectivity analyses of fMRI Data acquired during steady-state conditions. PLoS
- 600 One 6, 1–14. https://doi.org/10.1371/journal.pone.0014788
- Marstaller, L., Burianová, H., Reutens, D.C., 2017. Adaptive contextualization: A new role for the
- 602 default mode network in affective learning. Hum. Brain Mapp. 38, 1082–1091.
- 603 https://doi.org/10.1002/hbm.23442
- Mayer, J.S., Roebroeck, A., Maurer, K., Linden, D.E.J., 2010. Specialization in the default mode:
- Task-induced brain deactivations dissociate between visual working memory and attention.
- 606 Hum. Brain Mapp. 31, 126–139. https://doi.org/10.1002/hbm.20850
- 607 McClearn, G.E., 1960. Strain differences in activity of mice: Influence of illumination. J. Comp.
- 608 Physiol. Psychol. 53, 142–143. https://doi.org/10.1037/h0042766
- Nestler, E.J., Hyman, S.E., 2010. Animal models of neuropsychiatric disorders. Nat. Neurosci. 13,
 1161–1169. https://doi.org/10.1038/nn.2647
- 611 Pan, W.-J., Billings, J.C.W., Grooms, J.K., Shakil, S., Keilholz, S.D., 2015. Considerations for
- 612 resting state functional MRI and functional connectivity studies in rodents. Front. Neurosci.
- 613 9, 20130152. https://doi.org/10.3389/fnins.2015.00269
- 614 Pawela, C.P., Biswal, B.B., Cho, Y.R., Kao, D.S., Li, R., Jones, S.R., Schulte, M.L., Matloub, H.S.,
- 615 Hudetz, A.G., Hyde, J.S., 2008. Resting-state functional connectivity of the rat brain. Magn.
- 616 Reson. Med. 59, 1021–1029. https://doi.org/10.1002/mrm.21524.Resting-State
- 617 Pawela, C.P., Hudetz, A.G., Ward, D.B., Schulte, M.L., Li, R., Kao, D.S., Mauck, M.C., Cho, Y.R.,
- 618 Neitz, J., James, H.S., 2008. Modeling of region-specific fMRI BOLD neurovascular response
- 619 functions in rat brain reveals residual differences that correlate with the differences in regional
- 620 evoked potentials 41, 525–534. https://doi.org/10.1016/j.neuroimage.2008.02.022.Modeling
- 621 Petridou, N., Gaudes, C.C., Dryden, I.L., Francis, S.T., Gowland, P.A., 2013. Periods of rest in
- 622 fMRI contain individual spontaneous events which are related to slowly fluctuating

623	spontaneous activit	v. Hum. Bra	ain Mapp. 34.	1319–1329. ht	tps://doi.org/10	.1002/hbm.21513

- 624 Rohleder, C., Wiedermann, D., Neumaier, B., Drzezga, A., Timmermann, L., Graf, R., Leweke,
- 625 F.M., Endepols, H., 2016. The Functional Networks of Prepulse Inhibition: Neuronal
- 626 Connectivity Analysis Based on FDG-PET in Awake and Unrestrained Rats. Front. Behav.
- 627 Neurosci. 10, 1–10. https://doi.org/10.3389/fnbeh.2016.00148
- Salvador, R., Suckling, J., Coleman, M.R., Pickard, J.D., Menon, D., Bullmore, E., 2005.
 Neurophysiological architecture of functional magnetic resonance images of human brain.
 Cereb. Cortex 15, 1332–2342. https://doi.org/10.1093/cercor/bhi016
- 631 Schwarz, A.J., Gass, N., Sartorius, A., Risterucci, C., Spedding, M., Schenker, E., Meyer-
- Lindenberg, A., Weber-Fahr, W., 2013. Anti-Correlated Cortical Networks of Intrinsic
 Connectivity in the Rat Brain. Brain Connect. 3, 503–511.
 https://doi.org/10.1089/brain.2013.0168
- 635 Shah, D., Jonckers, E., Praet, J., Vanhoutte, G., Delgado Y Palacios, R., Bigot, C., D'Souza, D. V. 636 Verhoye, M., Van der Linden, A., 2013. Resting state FMRI reveals diminished functional 637 amyloidosis. connectivity in а mouse model of PLoS One 8. e84241. 638 https://doi.org/10.1371/journal.pone.0084241
- 639 Shah, D., Praet, J., Latif Hernandez, A., Höfling, C., Anckaerts, C., Bard, F., Morawski, M., Detrez, 640 J.R., Prinsen, E., Villa, A., De Vos, W.H., Maggi, A., D'Hooge, R., Balschun, D., Rossner, 641 S., Verhoye, M., Van der Linden, A., 2016. Early pathologic amyloid induces hypersynchrony 642 of BOLD resting-state networks in transgenic mice and provides an early therapeutic window 643 before amyloid plaque deposition. Alzheimer's Dement. 12, 964–976. 644 https://doi.org/10.1016/j.jalz.2016.03.010
- 645 Sierakowiak, A., Monnot, C., Aski, S.N., Uppman, M., Li, T.Q., Damberg, P., Brené, S., 2015.
- 646 Default mode network, motor network, dorsal and ventral basal ganglia networks in the rat

- brain: Comparison to human networks using resting state-fMRI. PLoS One 10, 1–20.
 https://doi.org/10.1371/journal.pone.0120345
- 649 Stafford, J.M., Jarrett, B.R., Miranda-Dominguez, O., Mills, B.D., Cain, N., Mihalas, S., Lahvis,
- 650 G.P., Lattal, K.M., Mitchell, S.H., David, S. V, Fryer, J.D., Nigg, J.T., Fair, D.A., 2014. Large-
- scale topology and the default mode network in the mouse connectome. Proc. Natl. Acad. Sci.
- 652 111, 18745–18750. https://doi.org/10.1073/pnas.1404346111
- Trancikova, A., Ramonet, D., Moore, D.J., 2011. Genetic mouse models of neurodegenerative
 diseases, 1st ed, Progress in Molecular Biology and Translational Science. Elsevier Inc.
- 655 https://doi.org/10.1016/B978-0-12-384878-9.00012-1
- Van Camp, N., Verhoye, M., De Zeeuw, C.L., Van der Linden, A., 2006. Light Stimulus Frequency
- 657 Dependence of Activity in the Rat Visual System as Studied With High-Resolution BOLD
 658 fMRI. J. Neurophysiol. 95, 3164–3170. https://doi.org/10.1152/jn.00400.2005
- van den Heuvel, M.P., Hulshoff Pol, H.E., 2010. Exploring the brain network: A review on resting-
- state fMRI functional connectivity. Eur. Neuropsychopharmacol. 20, 519–534.
 https://doi.org/10.1016/j.euroneuro.2010.03.008
- 662 Zhou, J., Liu, S., Ng, K.K., Wang, J., 2017. Applications of Resting-State Functional Connectivity
- to Neurodegenerative Disease. Neuroimaging Clin. N. Am. 27, 663–683.
 https://doi.org/10.1016/j.nic.2017.06.007
- 665

666