

BRAIN COMMUNICATIONS

Identifying the neural network for neuromodulation in epilepsy through connectomics and graphs

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Deep brain stimulation is a treatment option for patients with drug-resistant epilepsy. The precise mechanism of neuromodulation in epilepsy is unknown, and biomarkers are needed for optimizing treatment. The aim of this study was to describe the neural network associated with deep brain stimulation targets for epilepsy and to explore its potential application as a novel biomarker for neuromodulation. Using seed-to-voxel functional connectivity maps, weighted by seizure outcomes, brain areas associated with stimulation were identified in normative resting state functional scans of 1000 individuals. To pinpoint specific regions in the normative epilepsy deep brain stimulation network, we examined overlapping areas of functional connectivity between the anterior thalamic nucleus, centromedian thalamic nucleus, hippocampus and less studied epilepsy deep brain stimulation targets. Graph network analysis was used to describe the relationship between regions in the identified network. Furthermore, we examined the associations of the epilepsy deep brain stimulation network with disease pathophysiology, canonical resting state networks and findings from a systematic review of resting state functional MRI studies in epilepsy deep brain stimulation patients. Cortical nodes identified in the normative epilepsy deep brain stimulation network were in the anterior and posterior cingulate, medial frontal and sensorimotor cortices, frontal operculum and bilateral insulae. Subcortical nodes of the network were in the basal ganglia, mesencephalon, basal forebrain and cerebellum. Anterior thalamic nucleus was identified as a central hub in the network with the highest betweenness and closeness values, while centromedian thalamic nucleus and hippocampus showed average centrality values. The caudate nucleus and mammillothalamic tract also displayed high centrality values. The anterior cingulate cortex was identified as an important cortical hub associated with the effect of deep brain stimulation in epilepsy. The neural network of deep brain stimulation targets shared hubs with known epileptic networks and brain regions involved in seizure propagation and generalization. Two cortical clusters identified in the epilepsy deep brain stimulation network included regions corresponding to resting state networks, mainly the default mode and salience networks. Our results were concordant with findings from a systematic review of resting state functional MRI studies in patients with deep brain stimulation for epilepsy. Our findings suggest that the various epilepsy deep brain stimulation targets share a common cortico-subcortical network, which might in part underpin the antiseizure effects of stimulation. Interindividual differences in this network functional connectivity could potentially be used as biomarkers in selection of patients, stimulation parameters and neuromodulation targets.

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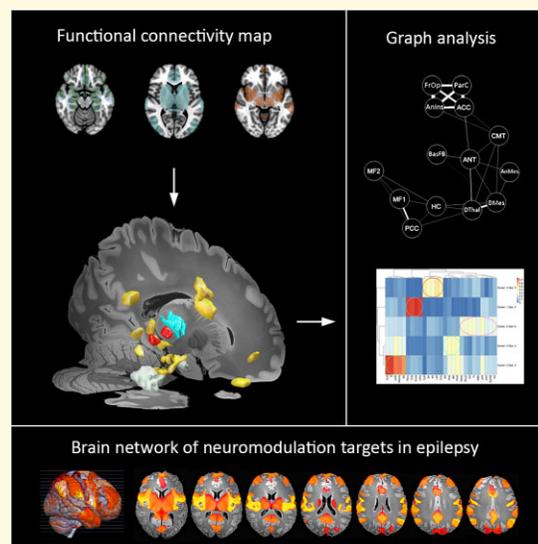
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Abbreviations: 3D=three-dimensional; ACC=anterior cingulate cortex; ANT=anterior nucleus of the thalamus; ARAS=ascending reticular activating system; BOLD=blood oxygenation level dependent imaging; cZI=caudal zona incerta; CMT=centromedian nucleus of the thalamus; DBS=deep brain stimulation; DMN=default mode network; DN=dentate nucleus; CB=cerebellum; Fx=fornix; fMRI=functional magnetic resonance imaging (fMRI); HCN=head of caudate nucleus; MEG=magnetoencephalography; MB=mammillary body; MMT=mammillothalamic tract; MS=medial septum (medial septum); NA=nucleus accumbens; NBM=nucleus basalis of Meynert; PPN=pedunculopontine nucleus; PCC=posterior cingulate cortex; PH=posterior hypothalamus; RR=response rate; SF=seizure free; SR=seizure reduction; SPECT=single photon emission computed tomography; SANTE=Stimulation of the Anterior Nucleus of the Thalamus for Epilepsy study; SNr=substantia nigra pars reticulata; STN=subthalamic nucleus; SUDEP=sudden unexpected death in epilepsy; VNS=vagal nerve stimulation; VTA=volume of tissue activated

Graphical Abstract



Introduction

Deep brain stimulation (DBS) is an alternative or adjuvant treatment for patients with drug-resistant epilepsy (DRE) who are not candidates for resective surgery. Approximately 30% of patients with epilepsy continue to have debilitating seizures despite best medical management.¹ DBS has the potential to stop the propagation of epileptic seizures or increase the threshold for generalization.² Possible mechanisms of DBS include activation of axons, local inhibition, effects on astrocytes, and disturbance of network oscillations.²

Various brain targets have been explored for DBS in epilepsy. Clinically used targets include the anterior nucleus of the thalamus (ANT), centromedian nucleus of the thalamus (CMT) and the hippocampus (HC), all of which have been investigated in randomized controlled trials.³ ANT-DBS received Food and Drug Administration approval in adult focal DRE. A randomized trial (SANTE) reported a median seizure reduction (SR) of 40% in patients who received active stimulation for 3 months, compared to 15% in the control group.⁴ In SANTE's long-term follow-up, a median SR of 75% was reported at 7 years.⁵ CMT-DBS, in contrast, has been commonly used in patients with generalized

epilepsies, particularly in the pediatric population.^{6,7} Hippocampal DBS has been performed in patients with seizures originating from the temporal lobe.^{8,9} Smaller, uncontrolled studies have probed the therapeutic potential of numerous other targets involving the circuit of Papez, limbic or cortico-subcortical circuits, including the subthalamic nucleus (STN), substantia nigra pars reticulata (SNr), caudal zona incerta (cZI), posterior hypothalamus (PH), fornix (Fx), nucleus accumbens (NAs), head of caudate nucleus (HCN), dentate nucleus (DN) and cerebellum (CB) (Fig. 1).³ Additional hypothetical targets for DBS in epilepsy have been studied in animal research.³

Disruption or modulation of epileptogenic networks most likely plays an important role in the therapeutic effect of DBS.^{10–12} Clinical efficacy has been associated with multiple DBS targets that belong to different anatomical circuits (motor, limbic, memory), suggesting the possibility of a common functional neural network responsible for the antiseizure effect associated with stimulation in epilepsy.¹³ Altered functional connectivity in patients with epilepsy, including deactivation of the default mode network (DMN), was described in simultaneous electroencephalography/functional MRI (EEG/fMRI) and MEG studies.^{14–17} Brain activity

measured with blood oxygen level-dependent (BOLD) signal during focal epileptic discharges is significantly altered in wide cortical and subcortical regions that extend beyond the epileptogenic zone.¹⁵ Alterations in resting state functional connectivity, such as activation of DMN and limbic networks, are described in patients receiving DBS for epilepsy.^{18–23} Other treatment options for DRE such as vagal nerve stimulation (VNS), responsive neural stimulation (RNS) and resective surgery also lead to reorganization of neural networks.^{24–26} Data gathered from EEG/fMRI and MEG studies is used in the evaluation of epileptic patients and network analysis techniques can be applied to surgical planning.^{25,27} The importance of functional and structural connectivity has gained recognition in neurologic and psychiatric disorders such as Parkinson's disease, Alzheimer's disease, obsessive-compulsive disorder and epilepsy, and the term circuitopathies has been introduced to describe the effects of neural network alterations in disease and treatment.^{28,29} Epilepsy is considered to be a network disease with different cortico-subcortical hubs responsible for seizure evolution and maintenance.¹⁴

The aim of this study was to describe the functional brain networks associated with clinically important DBS targets

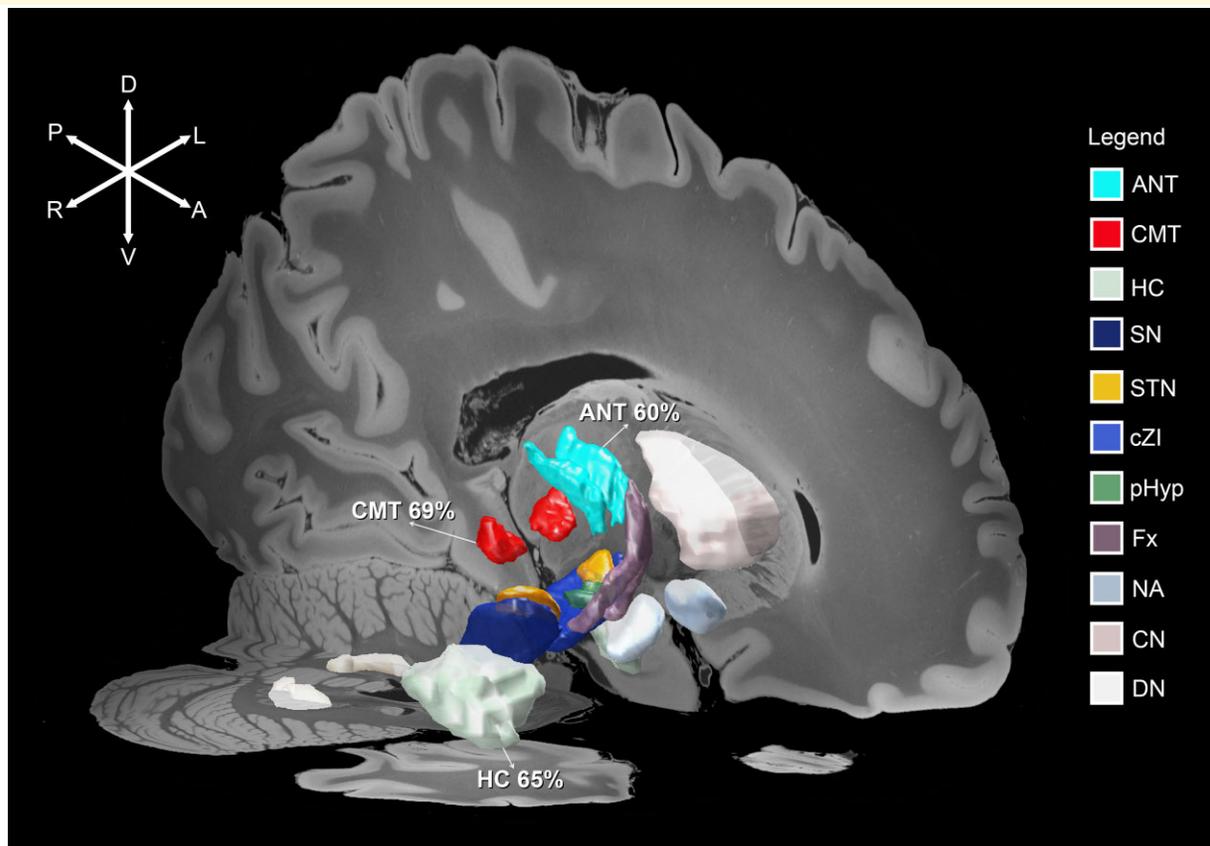


Figure 1 Targets for DBS in epilepsy. 3D representation of anatomical seeds of DBS targets used for the treatment of epilepsy is shown on sagittal and axial 7-Tesla T1 MRI slices of the human brain (100 μ m resolution in MNI152 space). Mean seizure reduction that was calculated from our systematic literature review is shown for ANT – 60%, CMT – 69%, and HC – 65%. Based on our systematic review, higher quality of evidence exists for the use of ANT-DBS in DRE. Other DBS targets that belong to known cortico-subcortical circuits are visualized on the figure.

for epilepsy (ANT, CMT, HC) and to examine how these networks relate: (i) to each other; (ii) to patterns of connectivity associated with less studied epilepsy DBS targets and with epilepsy pathophysiology; and (iii) to canonical resting state networks. We hypothesized that ANT, CMT and HC targets would exhibit connectomic overlap and that their functional networks would overlap with known epileptic circuits and with the DMN.

Materials and methods

PubMed database screening was performed using the terms ‘seizure deep brain stimulation’ and ‘epilepsy deep brain stimulation’. Seizure outcomes for ANT, CMT and HC were gathered from clinical trials that included more than three patients and were published during the last two decades. Seizure outcomes for the less used targets were collected from all identified studies, and hypothetical DBS targets were screened during review. Studies included in calculation of means and identified targets are presented in [Supplementary material 1](#). Mean SR was estimated from medians in studies where means were not reported.^{30,31} Response rate was determined as SR of 50% compared to baseline. DBS targets included in the analysis were classified as: clinically used epilepsy DBS targets (ANT, CMT, HC), less used DBS targets (STN, SNr, cZI, PH, Fx, NA, HCN, DN) and hypothetical targets [mammillothalamic tract (MMT), mammillary body (MB), nucleus basalis of Meynert, pedunculopontine nucleus (PPN) and medial septum]. Seeds of DBS targets were created from anatomical atlases ([Supplementary material 1](#)). A previously published ANT seed was used that covered the anterior nuclei targeted by DBS.²³

Statistical and connectivity analysis

Functional connectivity maps were created for each DBS target region to investigate the brain network associated with SR following DBS. Connectomic analysis has been previously used to describe neural networks associated with neuromodulation and neurological disorders.^{32–34} Functional connectivity maps for each seed (each DBS target) were calculated using a high-quality normative 3 T resting state fMRI data set derived from 1000 healthy subjects (age range: 18–35 years; 57.6% female) of the Brain Genomics Superstruct Project (BGSP, <https://dataverse.harvard.edu/dataverse/GSP>), as previously described.^{35–39} The MRI acquisition and pre-processing parameters for the BGSP normative connectome are described in detail in the original publication.⁴⁰ In short, the data were collected on 3 T Tim Trio scanners (Siemens, Erlangen, Germany) using a 12-channel phased-array head coil. The following EPI parameters were used: repetition time = 3000 ms; echo time = 30 ms; flip angle = 85°; voxel size = 3 × 3 × 3 mm; field of view = 216; slice acquisition = 47 axial slices acquired in interleaved fashion with no gap between slices. Each

participant in the GSP study underwent one or two functional runs (mean of 1.7 runs), each of which lasted 6.2 min (124 time points). The fMRI preprocessing consisted of common methods: the first four volumes of each run were discarded, slice acquisition-dependent time shifts per volume were compensated for, and motion correction was applied. Temporal filtering was also performed, retaining frequencies below 0.08 Hz, and individual scans were normalized to common space. Finally, spatial smoothing of the resting-state data was performed.

Using the DBS target seeds whole-brain connectivity r-maps were generated describing the average correlation of the low-frequency BOLD signal fluctuation between the seed and each voxel. The voxel-wise r-maps show the average voxel to seed correlation across the entire normative dataset. To identify all voxels that meaningfully connected to each seed, the r-maps were converted to t-maps and using the known p-distribution these t-maps were Bonferroni corrected for multiple comparisons at $|t| = 5.1$ (Bonferroni corrected < 0.05 , whole brain). The thresholded t-maps were binarized to capture the brain-wide pattern of regions significantly connected with each seed. Using this conservative approach subsequent results are not driven by subtle variations in local connectivity.³⁸

To identify the brain regions associated with DBS and symptom improvement, a weight of mean SR was assigned to the functional maps, and a voxel efficacy map of percent improvement of ANT, CMT, and HC functional connectivity was created ([Fig. 2](#); in-house MATLAB script, version R2017b; MathWorks, Natick, MA).⁴¹

The overlapping regions between the three binarized maps (ANT, CMT and HC) were used to outline the common network implicated in SR following DBS (the PCC and the frontobasal seed were averaged in size with other seeds for comparison). The identified regions were used to construct a network associated with DBS in epilepsy ([Supplementary Fig. 1](#)). Additional binarized maps of regions of high overlap were calculated for the subsequent network analysis ([Fig. 3](#)). For all seeds and high overlap regions, a Pearson correlation matrix was created between regions of overlapping functional connectivity and targets for DBS in epilepsy using Lead Mapper v2.5.3.⁴² Clustering of the Pearson correlation matrix of functional connectivity between brain regions included in the network was performed with the R package pheatmap (version 1.0.12) based on Euclidian distances where 5 clusters were obvious and chosen for illustration (the simple input correlation matrix is presented as [Supplementary Fig. 2](#)). The weighted network graph was created based on the functional connectivity correlation matrix with qGraph library (version 3.1.1) using the ‘qgraph’ function in R (<https://www.r-project.org/>, version 4.1.1). The regions of anticorrelation were not included in the graph analysis as per Rubinov and Sporns.⁴³ Edge thickness denotes the strength of correlations between nodes (edge thickness cut-off was set at 0.3). Graph centrality measures (betweenness, closeness) were calculated. Higher betweenness shows how often a node is part of the shortest

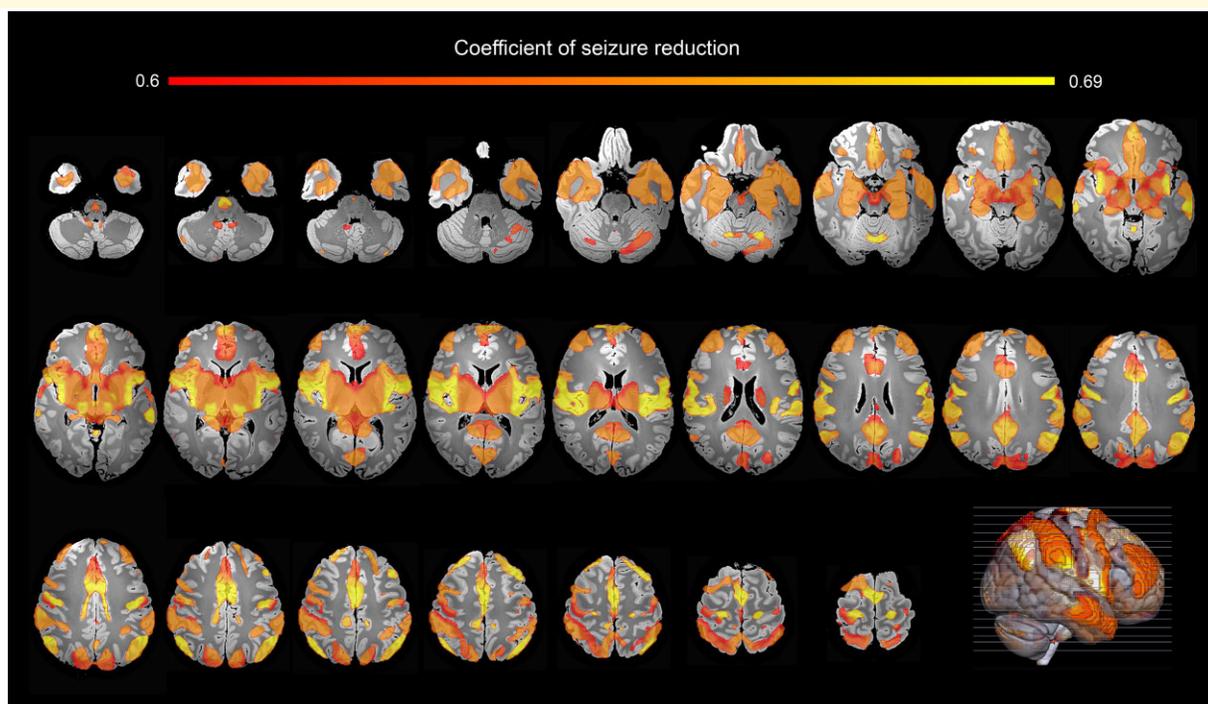


Figure 2 Functional connectivity of clinically used epilepsy DBS targets (ANT, CMT, HC). To identify the brain regions associated with DBS and symptom improvement, a weight equal to targets mean seizure reduction was assigned to the binarized connectivity t-maps calculated from normative resting state fMRI scans of 1000 individuals, and a voxel efficacy map of percent improvement of ANT, CMT, and HC was created. Areas of functional connectivity affected by DBS are shown on the axial 7-Tesla T1 MRI slices. A 3D figure with an overlay of the functional connectivity map represents the level of the axial slices. Cortical regions affected by DBS for epilepsy are the ACC, PCC, insula, medial and lateral frontal region, temporal lobe, superior parietal lobe, angular and supramarginal gyrus, cuneus, sensorimotor and premotor cortex. Subcortical regions affected by stimulation are the basal ganglia, dorsal and ventral mesencephalon, pons and CB.

connection between regions and indicates a stronger influence on the transfer of information within the neural network. Closeness describes the average shortest distance from each node in the network to others and reflects how efficiently a node can spread information within a network.⁴⁴

To compare the normative network model associated with common epilepsy DBS targets with reports from patient studies, we performed a systematic review of resting state fMRI findings in epilepsy DBS (three patient reports and three normative fMRI studies). PubMed search terms ‘epilepsy DBS functional MRI’, ‘epilepsy DBS fMRI’, ‘epilepsy deep brain stimulation fMRI’, were used, and bibliographies of identified articles were searched for additional reports.

Data availability

All data is available upon reasonable request.

Results

Seizure outcomes in DBS for epilepsy

The calculated mean SR after DBS of ANT, CMT and HC in clinical trials involving more than three patients was 60, 69

and 65%, respectively. DBS of ANT is used more often for focal epilepsies of temporal and extratemporal origin, CMT for generalized epilepsies and HC for temporal lobe epilepsy. Higher quality evidence exists for DBS of ANT than for CMT or HC.⁴⁵ Outcomes for less studied epilepsy DBS targets vary and range from 48 to 92% (Table 1). Initial studies of cerebellar DBS showed promising results, but a subsequent randomized trial did not achieve statistical significance.¹³

Areas of functional connectivity associated with DBS in epilepsy

A summed map of ANT, CMT and HC functional connectivity weighted by mean seizure outcome revealed wide cortical and subcortical regions affected by stimulation. Cortical areas with higher time course correlation to DBS targets were in the anterior cingulate cortex (ACC), posterior cingulate cortex (PCC), insula, medial and lateral frontal region, temporal lobe, superior parietal lobe, angular and supramarginal gyrus, cuneus, sensorimotor and premotor cortex. Subcortical areas with higher time course correlation were in the basal ganglia, dorsal and ventral mesencephalon, pons and CB (Fig. 2).

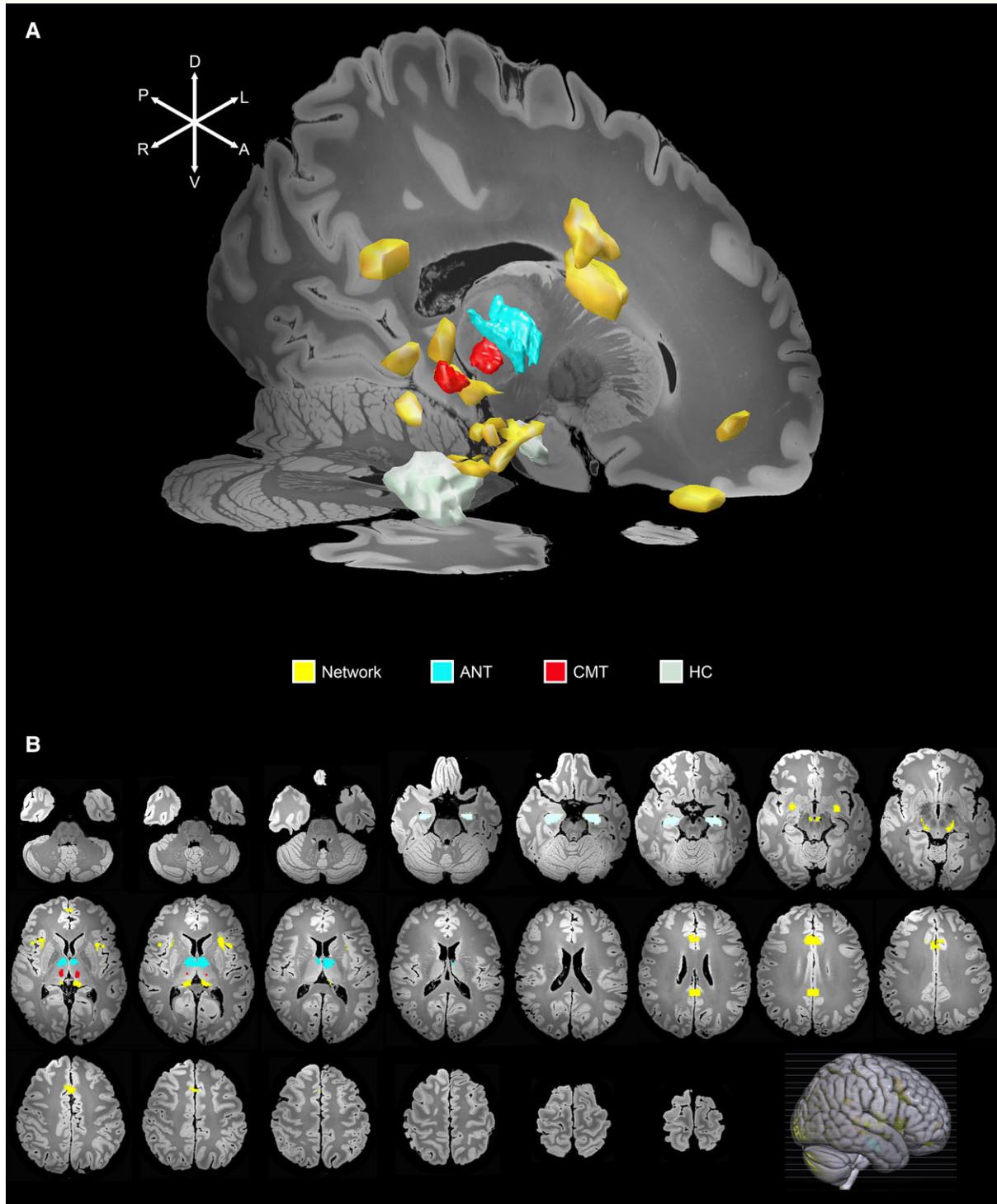


Figure 3 Common brain network of deep brain stimulation in epilepsy. (A) A 3D representation of the anatomical seeds of the clinically used targets for DBS in epilepsy (ANT, CMT, HC) and regions of their functional connectivity overlap calculated from resting state fMRI scans of 1000 individuals are shown on high-resolution sagittal and axial 7-tesla T1 MRI slices of the human brain. The regions that overlapped between the three functional connectivity maps of ANT, CMT, and HC (in yellow) were used to outline the common neural network implicated in seizure reduction following DBS. (B) The regions identified as the normative network model associated with common epilepsy DBS targets are shown on axial 7-Tesla T1 MRI slices. A 3-D figure with an overlay of the network represents the level of the axial slices. Regions of overlap were in the ACC, paracingulate cortex, medial frontal region, PCC, anterior insula, frontal operculum, right sensorimotor cortex, basal forebrain, medial thalamus, dorsal thalamus, dorsal and ventral mesencephalon.

Table 1 Review of outcomes for DBS in epilepsy. Seizure reduction, response rate (SR > 50%), and cohort characteristics of patients with deep brain stimulation for epilepsy were extracted from a systematic literature review

Target	N	Mean age	Follow-up (mo)	Mean seizure reduction	Number of studies
ANT	330	33.5	34.2	59.6	23
CMT	90	23.9	28.7	69.3	8
HC	107	33.8	33.9	64.6	13
STN	22	21.7	26.2	66.5	9
cZI	6	34.7	38.2	87.5 SR in two studies, 3/3 RR in one study	3
SN	1	32	24	100 in GTC, improvement in myoclonic seizures	1
PH	7	24.5	47.3	83.6	2
Fx	7	41	1-9 days	92	1
NA	9	39.5	6	47.5	2
CN	38	None	18	21/38 SF, 14/38 improved	1
DN	95	28.6	36.3	Mixed results	8

Common brain network of DBS in epilepsy

The overlap of the maps of voxels that are significantly connected to the epilepsy DBS targets used for successful treatment overlaps highly and outlines a common brain network underlying treatment (Supplementary Fig. 1). Functional connectivity maps of ANT, CMT and HC seeds had regions of overlap in the ACC, paracingulate cortex, medial frontal region, PCC, anterior insula, frontal operculum, right sensorimotor cortex, basal forebrain, medial thalamus, dorsal thalamus, dorsal and ventral mesencephalon. DBS targets and areas of overlapping functional connectivity between them were identified as a common neural network associated with the action of DBS in epilepsy (Fig. 3). Cortical and subcortical regions of overlapping functional connectivity between clinically used, less used, and hypothetical DBS targets were compared and are presented in Supplementary Table 1.

Graph analysis of the epilepsy DBS network

A correlation matrix of functional connectivity, based on normative resting state fMRI, showed clustering of nodes into five groups (Fig. 4A). Graph analysis revealed a network with two cortical clusters and a subcortical cluster (Fig. 4B). The first cluster largely corresponded to cortical regions of the DMN (medial frontal cortex, PCC, HC). This group of nodes had stronger correlations to dorsal thalamus and dorsal mesencephalon. The second cluster included cortical regions involved in the saliency network (ACC and

paracingulate cortex, frontal operculum, and anterior insula) and had stronger correlations with ANT and CMT. The largest cluster included ANT, CMT, most other DBS targets and subcortical nuclei. A separate cluster of subcortical regions included the anterior mesencephalon structures and the MMT. The fifth cluster contained PPN, PH, MB and DN. The anterior thalamic nucleus had a central position in the network with the highest betweenness and closeness values showing more connections to other nodes in network, while CMT and HC had average centrality values (Fig. 4C). Caudate nucleus and MMT also displayed high centrality values. ACC displayed the highest centrality values from the cortical regions. The expanded graph network with the less used and hypothetical DBS targets is presented in Supplementary Fig. 3.

Review of resting state fMRI studies in patients with DBS for epilepsy

Several recent studies examined the resting state fMRI changes in patients with DBS and showed associations with functional connectivity in the DMN, sensorimotor cortex and other regions (Table 2). A study using resting state fMRI in two ANT DBS patients with DBS on showed activation in regions associated with the limbic and DMNs of the brain.²¹ Functional connectivity associated with regions of DMN was reported from electrode volume of tissue activated (VTA) analyses in three ANT DBS responders.²⁰ The association of ANT stimulation with DMN was further confirmed in patients with long-term ANT DBS.²³ A recent study with low- and high-frequency stimulation during resting state fMRI showed that high frequency DBS produced activation of DMN and limbic networks of the brain, in contrast to low-frequency stimulation.²² Studies of functional connectivity associated with CMT electrode VTAs produced less functional connectivity in regions of DMN, but higher connectivity with sensorimotor, premotor, ACC, ascending reticular activating system (ARAS) and other brain regions (Table 2).^{18, 19}

Discussion

We used the seeds of epilepsy DBS targets and areas of their common functional connectivity, as defined from normative resting state fMRI scans of 1000 individuals, to identify a network associated with neuromodulation in epilepsy. Our results showed that epilepsy DBS targets have common areas of cortical and subcortical resting state functional connectivity in regions associated with propagation and generalization of epileptic seizures. The normative neural network shared by ANT, CMT, HC and other DBS targets overlapped cortical and subcortical regions associated with resting state networks, mainly the DMN and saliency networks. The common functional connectivity network shared by clinically used DBS targets overlapped regions of the epileptic networks identified by previous studies of regions activated

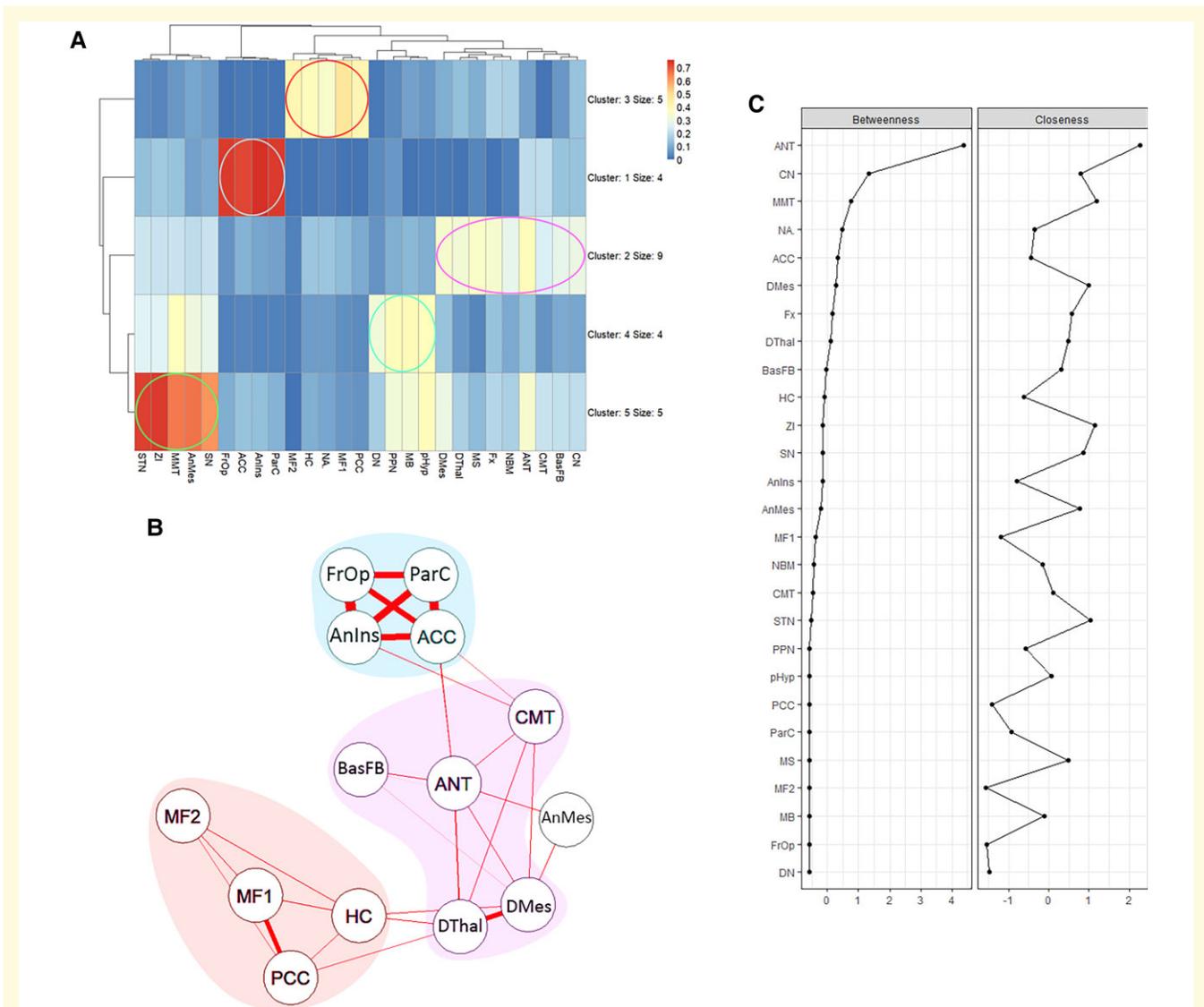


Figure 4 Graph analysis of the normative epilepsy DBS network. (A) A Pearson correlation matrix of functional connectivity between seeds (ANT, CMT, HC, areas of common functional connectivity, less studied and hypothetical DBS targets) was clustered into 5 groups based on the strength of correlations to present the associations between regions involved in DBS for epilepsy. **(B)** A graph network, based on the correlation matrix between the epilepsy DBS targets (ANT, CMT, HC) and regions of their functional connectivity overlap, was created to visualize the relationship between nodes in the epilepsy DBS neural network. Thickness of edges shows the strength of correlations between the nodes (minimum correlations presented >0.2). **(C)** Centrality measures of the epilepsy DBS neural network show that ANT, CN, and MMT act as a central hubs in the network with the highest betweenness and closeness values, while CMT and HC showed average centrality values (standardised z-scores on the x-axis).

during DBS.^{14,15,22,23} This suggests that DBS targets and epilepsy share a common cortico-subcortical network that might be responsible for the antiseizure action of DBS.^{21,46,47}

Identifying the neural network associated with DBS in epilepsy

Cortical nodes identified in both the normative network model and associated with common epilepsy DBS targets were in the ACC and PCC, medial frontal and sensorimotor cortex, frontal operculum and bilateral insulae. Subcortical

nodes of the network were in the basal ganglia, mesencephalon, basal forebrain and CB.

Clustering of the functional connectivity matrix showed two cortical groups and intercorrelated subcortical clusters that included most DBS targets. The first cortical group corresponded to nodes associated with the DMN (medial frontal cortex, PCC and precuneus, HC).⁴⁸ The second cortical group consisted of nodes associated with the saliency network (ACC, anterior insula, frontal operculum).⁴⁹ Overlap between ANT, CMT, and HC functional connectivity in the sensorimotor cortex occurred only on the right side.

Table 2 Review of patient and normative (VTA based) resting state fMRI studies in DBS for epilepsy

Study	Target (N)	fMRI details	Main finding
Middlebrooks et al. 2018 ²⁰	ANT (6)	Electrode VTAs, normative	Responders had connectivity in regions of posterior cingulate cortex, medial prefrontal cortex, inferior parietal lobule and precuneus, which are associated with DMN.
Middlebrooks et al. 2020 ²¹	ANT (2)	ON/OFF in 30 s blocks	Activation in bilateral thalamus, anterior cingulate and posterior cingulate cortex, precuneus, medial prefrontal cortex, amygdala, ventral tegmental area, HC, striatum and right angular gyrus
Middlebrooks et al. 2021 ²²	ANT (5)	ON/OFF in 30 s blocks	High-frequency stimulation led to activation in DMN (PCC, precuneus, angular gyrus, parahippocampal gyrus, medial thalamus, and basal forebrain) and limbic (HC, ACC, amygdala, ANT, medial prefrontal cortex, and ventral tegmental area areas). Low-frequency stimulation produced deactivation in the similar regions.
Sarica et al. 2020 ²³	ANT (2)	ON/OFF in 30 s blocks	Stimulation of ANT contacts produced positive connectivity with bilateral putamen, thalamus, and posterior cingulate cortex, ipsilateral middle cingulate cortex and precuneus, and contralateral medial prefrontal and anterior cingulate cortex
Warren et al. 2020 ¹⁹	CMT (19)	Electrode VTAs, normative	Positive subcortical connectivity in cerebellum, thalamus, brainstem, striatum and subthalamic nuclei. Positive cortical connectivity in auditory cortex, precentral and postcentral gyri, premotor cortex, cingulate cortex, parahippocampal/fusiform cortex and insular cortex.
Diaz et al. 2021 ¹⁸	CMT (10)	Electrode VTAs, normative	Positive subcortical connectivity in thalamus, striatum and STN. Positive cortical connectivity in sensorimotor cortex, SMA, middle frontal cortex, medial temporal cortex, and anterior cingulate. Reponders had DTI fibres passing through VTAs connected to sensorimotor, supplementary motor cortex, middle and superior frontal gyrus, cerebellum and ARAS.

The sensorimotor cortex had a higher time course correlation with CMT and is associated with response to DBS in generalized epilepsy.¹⁸ Areas of resting state functional connectivity activated during successful DBS for epilepsy in patients reviewed from published studies (Table 2 of results) are similar to the network described in our study, suggesting that successful DBS leads to modulation of this network.

We found subcortical areas of common functional connectivity between ANT, CMT and HC in the medial and dorsal thalamus, dorsal and ventral mesencephalon, and CB, regions that have been previously implicated in the pathophysiology of seizures.^{5,50} The region in the dorsal thalamus corresponded to medial pulvinar, and is associated with temporal lobe seizures and status epilepticus.^{51–53} Additionally, the pulvinar nucleus was recently successfully used as a target for responsive neural stimulation (closed loop technique) in patients with posterior quadrant epilepsy.⁵⁴ The identified areas of functional connectivity in the basal forebrain, medial and mediodorsal thalamus are parts of the DMN, and are likely parts of a common subcortical circuit associated with DBS in epilepsy.⁵⁵ The area in the superior dorsal mesencephalon (lateral geniculate body) is associated with epileptic seizures and antiseizure effects.^{56–58} Hyperactivity in the tegmentum is associated with loss of consciousness during seizures and following brain trauma.^{47,59,60} The role of pontine and mesencephalic ascending reticular formation in consciousness supports the cortico-subcortical framework of seizure generalization in epilepsy.² A recent study of CMT DBS showed an association with better outcome when ARAS was covered by the VTAs of electrodes.¹⁸ The long-term follow-up of ANT stimulation in the SANTE trial showed a decrease in sudden unexpected death in epilepsy (SUDEP), which suggests the possibility that this might be related to the effect of stimulation on the brainstem Raphe nuclei.⁵

Graph analysis of the epilepsy DBS network

Graph analysis revealed ANT as a central hub in the normative network model associated with common epilepsy DBS targets with the highest betweenness and closeness centrality values. Mammillothalamic tract shared similar centrality values to ANT. Simultaneous targeting of MMT and ANT is associated with better outcomes after DBS in epilepsy.⁶¹ Adding MMT to the ANT seed increased the area of functional correlation in the brainstem and ACC region.

The anterior thalamic nucleus and CMT had a stronger correlation with ACC, dorsal thalamus, and mesencephalon, which are likely important hubs in the neural network associated with DBS action in epilepsy. CMT receives input from ARAS, which may explain the effectiveness in generalized seizures associated with loss of consciousness. Other DBS targets with high centrality values are the caudate nucleus and NAs. The NAs had a bigger area of functional connectivity in the frontal lobes and had higher correlations with the frontal cortical cluster. The basal forebrain seed had a strong correlation with ANT, dorsal thalamus and dorsal mesencephalon, which potentially makes it a novel target for DBS for epilepsy. The DN had the lowest correlation to the network, which is reflected in its questionable clinical efficacy in DBS for epilepsy.

The ACC was identified as an important hub in the neural network associated with the effect of DBS in epilepsy. Low-frequency DBS (5 Hz) of CMT in 10 patients resulted in a sequential propagation of the EEG source from the ACC to frontal, then to temporal and other cortical regions.⁶² Similar activation in ACC was demonstrated by distributed source modelling of EEG recorded during DBS of ANT in three patients, and during stimulation of different anterior thalamic subnuclei in five patients.^{63,64} In our graph

analysis ACC had a strong correlation with three cortical seeds (frontal operculum, paracingulate cortex, and anterior insula). In the identified normative network model associated with common epilepsy DBS targets ACC acted as a relay for DBS targets to the cortical cluster, similarly to the propagation of electrical signals in neurophysiological studies. Furthermore, enhanced intrinsic functional connectivity between the thalami, ACC and insula predicts response to VNS in children with DRE.⁶⁵

Association of the epilepsy DBS network with resting state fMRI networks

We described an overlap of normative functional connectivity between ANT, CMT and HC seeds in regions corresponding to the DMN. The DMN includes the medial prefrontal cortex and ACC, PCC, precuneus, medial and lateral parietal cortices, and temporal areas, and is involved in introspection and social functions.⁶⁶ A recent study identified additional regions associated with the DMN in the basal forebrain, and anterior and mediodorsal thalamus.⁵⁵ The DMN plays an important role in generalized and focal epilepsy. Reduced functional connectivity between regions of DMN is reported in patients with idiopathic generalized epilepsy.⁶⁷ Decreased fMRI activity in the DMN occurs in patients during bursts of generalized spike-and-wave (GSW) discharges in patients with primary generalized epilepsy.⁶⁸ Deactivation of DMN identified by fMRI and single photon emission computerized tomography (SPECT) is associated with loss of consciousness during generalized and focal impaired awareness seizures (FIAS), which is consistent with the ‘network inhibition hypothesis’.^{68,69} Danielson *et al.*⁷⁰ proposed that increased thalamic activity and inhibition of the reticulate activating system, which otherwise maintains the DMN, could be the common mechanism for loss of consciousness in seizures of different types (FIAS, generalized tonic-clonic, absence). Decrease in DMN connectivity also occurs in other pathological states of decreased consciousness, eg. in minimally conscious patients.^{71,72} Similarly, in a study of focal interictal epileptiform discharges in patients with temporal, frontal and posterior quadrant epilepsy, the common finding was deactivation in the DMN in all three groups, despite the large differences in localization of the epileptiform discharges.⁷³ Patients with focal seizures exhibit fMRI changes not only in the seizure onset zone, but have wider alterations in functional connectivity of the brain during epileptiform discharges, including cortical and subcortical sites in thalamus, basal ganglia, and reticular activating system.⁷⁴

The second cortical cluster in our graph analysis potentially corresponds to the cortical regions of the salience network (ACC, frontoinsula cortex and subcortical structures).⁷⁵ Altered intrinsic connectivity in the salience network was described in patients with childhood absence epilepsy.⁴⁹ Integrity of the salience network is associated with DMN

function.^{76,77} ACC, medial frontal cortex, bilateral insulae, sensorimotor cortex and caudate nucleus have been implicated in initiation and propagation of ictal discharges together with the mediodorsal nuclei of thalamus in an EEG-fMRI study.⁷⁸ The ACC has been functionally connected to the basal forebrain and DMN in recent human and animal studies.⁵⁵ The ACC is an important part of the salience network. As part of the limbic and Papez circuits, ACC is associated with affection, cognition and seizure propagation.^{79,80} Decreased connectivity between thalamus and ACC, measured with positron emission tomography (PET), is associated with the minimally conscious state, and connectivity between them increases with clinical improvement.⁸¹ Children with epilepsy have better neurocognitive outcomes when resting state networks are not perturbed by interictal epileptiform discharges. Consequently, normalization of intrinsic network connectivity through DBS could lead to better cognitive and neuropsychiatric outcomes in patients with epilepsy.⁸²

Role of the anterior and centromedian thalamic nucleus, and HC in DBS

The anterior thalamic nucleus is the most commonly studied target for DBS in epilepsy patients with focal and secondarily generalized seizures.⁵ The anterior thalamic nucleus is part of the Papez circuit, a loop that consists of HC, Fx, mammillary nuclei, ANT and cingulum. The Papez circuit is involved in memory and cognition.⁸³ Remote regions, such as CB, STN and CMT project to nodes within the circuit of Papez.⁸⁴ The anterior thalamic nucleus has wide reciprocal connections to cingulum and the hippocampal area.⁸⁵ The anterior thalamic nucleus is reported to be involved in maintenance and propagation of epileptic seizures.⁸⁶ Multiple studies have reported an association between ANT stimulation and DMN alterations.^{1,20,23}

The centromedian thalamic nucleus is involved in seizure propagation and is probably implicated in loss of consciousness during generalized seizures due to its anatomical connections to the ARAS and wide projections to the cortex, mainly premotor and motor regions.^{86–88} Higher reduction in generalized seizures after CMT-DBS was associated with stimulation affecting discriminative structural connectivity fibres overlapping ARAS, sensorimotor and supplementary motor cortices and CB/brainstem.¹⁸ The centromedian thalamic nucleus is activated earlier during generalized seizures than the ANT, suggesting that early seizure propagation might occur in posterior regions of the thalamus and subsequent maintenance of ictal activity in the anterior regions.⁸⁹ The centromedian thalamic nucleus is likely also involved in focal seizures and their generalization, but studies reported a lower decrease in focal seizures after CMT stimulation.^{6,90} The centromedian thalamic nucleus stimulation was more effective for generalized than frontal lobe epilepsy in a study of 11 patients (84% mean SR in generalized seizures versus

47% mean SR in frontal seizures), which could be related to higher anatomical connectivity of CMT to premotor, motor, and primary somatosensory areas.^{7,88} Patients with genetic (primary) generalized epilepsy studied with EEG-fMRI during GSW discharges had a substantial BOLD activation in the striatum and motor/premotor cortex.⁹¹ In a recent study, functional connectivity associated with CMT-DBS electrode regions of interest showed an overlap with the functional network described for generalized epilepsy.¹⁹

In cases of temporal lobe epilepsy, HC stimulation could potentially stop seizure propagation by interrupting the early spread of ictal activity from the seizure onset zone to other regions associated with seizure maintenance (i.e. ANT).⁹²

The association between basal ganglia, ARAS and resting state networks (DMN, salience network) in epileptic seizures provides a common physiological substrate for the antiseizure effect of neuromodulation in epilepsy.^{18–21} The nodes in the normative network model associated with common epilepsy DBS targets can have different functions, i.e. some nodes can be directly associated with seizure emergence and propagation, while others and white matter tracts can be relays to functional hubs in the neural circuit. The effect of DBS might be produced by local changes in key hubs leading to neural network modulation and plasticity. In the future, surgical targets and stimulation parameters in epilepsy DBS could be individually selected according to the connectivity of the hubs to patient specific epileptic networks. Alternatively, hubs in the identified normative network model associated with common epilepsy DBS targets could be used as targets for DBS, non-invasive neuromodulation, or for sensing electrodes in closed loop stimulation or even for the prediction of SUDEP.⁹³

Approach to epilepsy as a network disorder

Neural network alterations that occur in patients with epilepsy are described both in the ictal and interictal state.^{94,95} Ponten *et al.*⁹⁶ have demonstrated that epileptic seizures are characterized by a loss of normal balance between local and global connectivity compared to the interictal state based on the correlations between EEG signals from different brain regions. A meta-analysis of functional and structural connectivity in epilepsy summarized that a pattern of increased local connectivity and decreased global connectivity is present in epilepsy.⁹⁷ Hyper-connected and hypersynchronized brain regions, that are associated with the epileptogenic zone, are likely important hubs in the epilepsy networks, and resection of these regions is connected to seizure freedom.^{98–100} In addition, an altered level of activity and connectivity is present in physiological hubs, most commonly in the DMN.⁹⁴ Both surgical and medical treatment modalities can lead to reorganization of neural connectivity in epilepsy.^{24,26,101} The current understanding of the mechanism of neuromodulation, such as DBS, supports the notion that functional treatments exert their effects through modulation of neural networks.²⁸ The technique of neural network interogation

using connectivity of neuromodulation targets through normative datasets gives valuable insights into the neuroscientific basis of disease and treatment mechanisms.¹⁰² Previous research in Parkinson's disease showed that normative functional connectivity studies and patient specific fMRI connectivity have a good overlap.¹⁰³ Furthermore, previous work has demonstrated that while age or disease processes are associated with detectable differences in connectivity between hub regions within networks, these do not change the general network structure.^{104–106} Furthermore, no general change in functional network structure is associated with treatment resistant epilepsy, according to a recent review.⁹⁵ The general physiological hubs and structure of the networks are consistent between individuals, but topological changes are possible in the epileptic networks.⁹⁹ As the core question of the present study was the identification of the brain network associated with epilepsy treatment, the use of normative data should not be detrimental to achieving this goal. The significance of the network hubs identified in the epilepsy DBS network can be explained by their role in epilepsy networks, and in seizure spread and maintenance, but it has to be further confirmed by examining the connectivity alterations in patients with epilepsy. The demonstration of the role functional and structural networks in neuromodulation for epilepsy requires extensive research in long-term prospective patient cohorts to establish its clinical significance.

Limitations

The use of normative fMRI data has some limitations. First normative data does not include any information about the individual patients included in the specific study. Furthermore, since the normative dataset was acquired using healthy control subjects, any disease specific structural and functional abnormalities are also not reflected in the data. The notion of network reorganization in epilepsy (i.e. due to disease related pathological changes or antiseizure medication) cannot be accounted for in the current study. However, this approach allows for a solution to the complex issue of acquiring fMRI scans in a small DBS population with epilepsy and the typically encountered questionable test-retest reliability and poor signal to noise ratio in fMRI patient scans. While normative datasets cannot provide the same level of specificity as patients studies, previous research has demonstrated that analysis using either normative, patient-specific data or a disease specific connectome lead to the identification of the same brain networks.^{39,103} We described the epilepsy DBS network in normative scans, but our results are supported by fMRI-EEG studies in epilepsy patients, and by fMRI studies in patients with epilepsy and DBS. Using anatomical labels might lead to different results compared to active stimulation or VTAs of electrodes but allows for a standardized approach. Anatomical labels were selected to reduce the heterogenous results created by varying electrode positions. Combining targets that belong to different anatomical circuits (motor, limbic, Papez) does not allow identification of direct anatomical connections, but

this approach allows for a study of shared functional connectivity that is present in the central nervous system based on the hodological framework.¹⁰⁷ Our approach allowed us to describe the common functional connectivity network between DBS targets and epilepsy, with the limitation that we cannot pinpoint specific function of the nodes and their role in the therapeutic effect. Retrospective review of clinical studies can lead to bias (heterogeneous groups of patients, differences in follow-up and programming, and stimulation parameters) and overestimation of clinical outcomes for less studied targets, an unavoidable limitation. Finally, the significance of our findings in different types of epilepsy, and for different subtypes of seizures requires further research in patient populations.

Conclusion

We described a novel brain network associated with targets used for neuromodulation in epilepsy. The epilepsy DBS brain network shared hubs with known epileptic networks and regions involved in seizure propagation and generalization. The normative network model associated with common epilepsy DBS targets demonstrated a partial overlap with regions of the DMN and saliency network. Our results are supported by previous findings from fMRI studies in patients with DBS, RNS and VNS for epilepsy. Functional connectivity may be used as a biomarker in selection of targets or adjustment of DBS programming parameters. In the future, DBS treatment could be tailored to individual patients and disease-specific networks in epilepsy or other pathologies.

Ethical publication statement

We confirm that we have read the Journal's position on issues involved in ethical publication and affirm that this report is consistent with those guidelines.

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Competing interests

C.S. has been receiving fellowship grants from Michael and Amira Dan Foundation and Turkish Neurosurgical Society. T.V. is an investor in the company Neurescence, who has no involvement in this study. T.V. provides consulting to the company Panaxium, who has no involvement in this study. S.K.K. has received consulting and speakers' fees from Medtronic. A.M.L. is Scientific Director for

Functional Neuromodulation Ltd. and a consultant to Medtronic, Abbott, Boston Scientific, Insightec, and Focused Ultrasound Foundation. The remaining authors report no competing interests.

Supplementary material

Supplementary material is available at *Brain Communications online*.

References

1. Kalilani L, Sun X, Pelgrims B, Noack-Rink M, Villanueva V. The epidemiology of drug-resistant epilepsy: a systematic review and meta-analysis. *Epilepsia* 2018;59(12):2179–2193.
2. Gummadaelli A, Kundishora AJ, Willie JT, et al. Neurostimulation to improve level of consciousness in patients with epilepsy. *Neurosurg Focus*. 2015;38(6):E10.
3. Zangiabadi N, Ladino LD, Sina F, Orozco-Hernández JP, Carter A, Téllez-Zenteno JF. Deep brain stimulation and drug-resistant epilepsy: a review of the literature. *Front Neurol*. 2019;10:601.
4. Fisher R, Salanova V, Witt T, et al. Electrical stimulation of the anterior nucleus of thalamus for treatment of refractory epilepsy. *Epilepsia* 2010;51(5):899–908.
5. Salanova V, Sperling MR, Gross RE, et al. The SANTÉ study at 10 years of follow-up: effectiveness, safety, and sudden unexpected death in epilepsy. *Epilepsia* 2021;62(6):1306–1317.
6. Velasco F, Velasco M, Jiménez F, et al. Predictors in the treatment of difficult-to-control seizures by electrical stimulation of the centromedian thalamic nucleus. *Neurosurgery* 2000;47(2):295–305.
7. Valentín A, García Navarrete E, Chelvarajah R, et al. Deep brain stimulation of the centromedian thalamic nucleus for the treatment of generalized and frontal epilepsies. *Epilepsia* 2013;54(10):1823–1833.
8. Cukiert A, Cukiert CM, Burattini JA, et al. Seizure outcome after hippocampal deep brain stimulation in patients with refractory temporal lobe epilepsy: a prospective, controlled, randomized, double-blind study. *Epilepsia* 2017;58(10):1728–1733.
9. Téllez-Zenteno JF, Wiebe S. Hippocampal stimulation in the treatment of epilepsy. *Neurosurg Clin N Am*. 2011;22(4):465–475.
10. Li DH, Yang XF. Remote modulation of network excitability during deep brain stimulation for epilepsy. *Seizure* 2017;47:42–50.
11. Zhu J, Xu C, Zhang X, et al. The thalamus-precentral gyrus functional connectivity changes in epilepsy patients following vagal nerve stimulation. *Neurosci Lett*. 2021;751(45):135815.
12. Zhu J, Wang J, Xu C, et al. The functional connectivity study on the brainstem-cortical/subcortical structures in responders following cervical vagus nerve stimulation. *Int J Dev Neurosci*. 2020;80(8):679–686.
13. Laxpati NG, Kasoff WS, Gross RE. Deep brain stimulation for the treatment of epilepsy: circuits, targets, and trials. *Neurotherapeutics*. 2014;11(3):508–526.
14. Kay B, Szaflarski JP. EEG/fMRI contributions to our understanding of genetic generalized epilepsies. *Epilepsy Behav*. 2014;34:129–135.
15. Sadjadi SM, Ebrahimzadeh E, Shams M, Seraji M, Soltanian-Zadeh H. Localization of epileptic foci based on simultaneous EEG–fMRI data. *Front Neurol*. 2021;12:645594.
16. Goodman AM, Szaflarski JP. Recent advances in neuroimaging of epilepsy. *Neurotherapeutics* 2021;18(2):811–826.

17. Hsiao FJ, Yu HY, Chen WT, *et al.* Increased intrinsic connectivity of the default mode network in temporal lobe epilepsy: evidence from resting-State MEG Recordings. *PLoS One.* 2015;10(6): e0128787.
18. Torres Diaz C V, González-Escamilla G, Ciolac D, *et al.* Network substrates of centromedian nucleus deep brain stimulation in generalized pharmacoresistant epilepsy. *Neurotherapeutics* 2021; 18(3):1665–1677.
19. Warren AEL, Dalic LJ, Thevathasan W, Roten A, Bulluss KJ, Archer J. Targeting the centromedian thalamic nucleus for deep brain stimulation. *J Neurol Neurosurg Psychiatry.* 2020;91(4): 339–349.
20. Middlebrooks EH, Grewal SS, Stead M, Lundstrom BN, Worrell GA, Van Gompel JJ. Differences in functional connectivity profiles as a predictor of response to anterior thalamic nucleus deep brain stimulation for epilepsy: a hypothesis for the mechanism of action and a potential biomarker for outcomes. *Neurosurg Focus.* 2018; 45(2): E7.
21. Middlebrooks EH, Lin C, Okromelidze L, *et al.* Functional activation patterns of deep brain stimulation of the anterior nucleus of the thalamus. *World Neurosurg.* 2020;136:357–363.e2.
22. Middlebrooks EH, Jain A, Okromelidze L, *et al.* Acute brain activation patterns of high- versus low-frequency stimulation of the anterior nucleus of the thalamus during deep brain stimulation for epilepsy. *Neurosurgery.* 2021;89:901–908.
23. Sarica C, Yamamoto K, Loh A, *et al.* Blood oxygen level-dependent (BOLD) response patterns with thalamic deep brain stimulation in patients with medically refractory epilepsy. *Epilepsy Behav.* 2021;122:108153.
24. Wang K, Chai Q, Qiao H, Zhang J, Liu T, Meng F. Vagus nerve stimulation balanced disrupted default-mode network and salience network in a postsurgical epileptic patient. *Neuropsychiatr Dis Treat.* 2016;12:2561–2571.
25. Sinha N, Dauwels J, Kaiser M, *et al.* Predicting neurosurgical outcomes in focal epilepsy patients using computational modelling. *Brain.* 2017;140(2): 319–332.
26. Khambhati AN, Shafi A, Rao VR, Chang EF. Long-term brain network reorganization predicts responsive neurostimulation outcomes for focal epilepsy. *Sci Transl Med.* 2021;13(608): eabf6588.
27. An S, Bartolomei F, Guye M, Jirsa V. Optimization of surgical intervention outside the epileptogenic zone in the virtual epileptic patient (VEP). *PLoS Comput Biol.* 2019;15(6):e1007051.
28. Lozano AM, Lipsman N. Probing and regulating dysfunctional circuits using deep brain stimulation. *Neuron.* 2013;77(3):406–424.
29. Kramer MA, Cash SS. Epilepsy as a disorder of cortical network organization. *Neuroscientist.* 2012;18(4):360–372.
30. Luo D, Wan X, Liu J, Tong T. Optimally estimating the sample mean from the sample size, median, mid-range, and/or mid-quartile range. *Stat Methods Med Res.* 2018;27(6):1785–1805.
31. Wan X, Wang W, Liu J, Tong T. Estimating the sample mean and standard deviation from the sample size, median, range and/or interquartile range. *BMC Med Res Methodol.* 2014;14:135.
32. Germann J, Elias GJB, Neudorfer C, *et al.* Potential optimization of focused ultrasound capsulotomy for obsessive compulsive disorder. *Brain.* 2021;144:3529–3540.
33. Elias GJB, De Vloot P, Germann J, *et al.* Mapping the network underpinnings of central poststroke pain and analgesic neuromodulation. *Pain.* 2020;161(12):2805–2819.
34. Zhu Z, Hubbard E, Guo X, *et al.* A connectomic analysis of deep brain stimulation for treatment-resistant depression. *Brain Stimul.* 2021;14(5):1226–1233.
35. Horn A, Reich M, Vorwerk J, *et al.* Connectivity Predicts deep brain stimulation outcome in Parkinson disease. *Ann Neurol.* 2017;82(1):67–78.
36. Elias GJB, Giacobbe P, Boutet A, *et al.* Probing the circuitry of panic with deep brain stimulation: connectomic analysis and review of the literature. *Brain Stimul.* 2020;13(1):10–14.
37. Boutet A, Jain M, Elias GJB, *et al.* Network basis of seizures induced by deep brain stimulation: literature review and connectivity analysis. *World Neurosurg.* 2019;132:314–320.
38. Mithani K, Boutet A, Germann J, *et al.* Lesion network localization of seizure freedom following MR-guided laser interstitial thermal ablation. *Sci Rep.* 2019;9(1):18598.
39. Germann J, Elias GJB, Boutet A, *et al.* Brain structures and networks responsible for stimulation-induced memory flashbacks during forniceal deep brain stimulation for Alzheimer's disease. *Alzheimers Dement.* 2021;17(5):777–787.
40. Yeo BT T, Krienen FM, Sepulcre J, *et al.* The organization of the human cerebral cortex estimated by intrinsic functional connectivity. *J Neurophysiol.* 2011;106(3):1125–1165.
41. Elias GJB, Boutet A, Joel SE, *et al.* Probabilistic mapping of deep brain stimulation: insights from 15 Years of therapy. *Ann Neurol.* 2021;89(3):426–443.
42. Horn A, Li N, Dembek TA, *et al.* Lead-DBS v2: towards a comprehensive pipeline for deep brain stimulation imaging. *Neuroimage* 2019;184:293–316.
43. Rubinov M, Sporns O. Complex network measures of brain connectivity: uses and interpretations. *Neuroimage* 2010;52(3): 1059–1069.
44. Kwon H, Choi YH, Lee JM. A physarum centrality measure of the human brain network. *Sci Rep.* 2019;9(1):5907.
45. Sprengers M, Vonck K, Carrette E, Marson AG, Boon P. Deep brain and cortical stimulation for epilepsy. *Cochrane Database Syst Rev.* 2017;7(7):CD008497.
46. Výtvarová E, Mareček R, Fousek J, Strýček O, Rektor I. Large-scale cortico-subcortical functional networks in focal epilepsies: the role of the basal ganglia. *NeuroImage Clin.* 2017;14: 28–36.
47. Blumenfeld H, Varghese GI, Purcaro MJ, *et al.* Cortical and subcortical networks in human secondarily generalized tonic-clonic seizures. *Brain.* 2009;132(4):999–1012.
48. Buckner RL, Andrews-Hanna JR, Schacter DL. The brain's default network: anatomy, function, and relevance to disease. *Ann N Y Acad Sci.* 2008;1124:1–38.
49. Luo C, Yang T, Tu S, *et al.* Altered intrinsic functional connectivity of the salience network in childhood absence epilepsy. *J Neurol Sci.* 2014;339(1-2):189–195.
50. Forcelli PA. Serotonin in the dorsal raphe: as i live and breathe. *Epilepsy Curr.* 2018;18(3):191–193.
51. Filipescu C, Lagarde S, Lambert I, *et al.* The effect of medial pulvinar stimulation on temporal lobe seizures. *Epilepsia.* 2019;60(4): e25–e30.
52. Capecchi F, Mothersill I, Imbach LL. The medial pulvinar as a subcortical relay in temporal lobe status epilepticus. *Seizure.* 2020;81: 276–279.
53. Bin G, Wang T, Zeng H, *et al.* Patterns of gray matter abnormalities in idiopathic generalized epilepsy: a Meta-analysis of voxel-based morphology studies. *PLoS One.* 2017;12(1):e0169076.
54. Burdette D, Mirro EA, Lawrence M, Patra SE. Brain-responsive corticothalamic stimulation in the pulvinar nucleus for the treatment of regional neocortical epilepsy: a case series. *Epilepsia Open.* 2021;6(3):611–617.
55. Alves PN, Foulon C, Karolis V, *et al.* An improved neuroanatomical model of the default-mode network reconciles previous neuroimaging and neuropathological findings. *Commun Biol.* 2019; 2:370.
56. Lüders H. Chapter 38 Brain stimulation and epilepsy: novel approaches for seizure control. *Suppl Clin Neurophysiol.* 2004; 57(C):379–382.
57. Redgrave P, Simkins M, Overton P, Dean P. Anticonvulsant role of nigrothalamic projection in the maximal electroshock model of epilepsy-I. Mapping of dorsal midbrain with bicuculline. *Neuroscience.* 1992;46(2):379–390.
58. Soper C, Wicker E, Kulick CV, N'Gouemo P, Forcelli PA. Optogenetic activation of superior colliculus neurons suppresses

- seizures originating in diverse brain networks. *Neurobiol Dis.* 2016;87:102–115.
59. Snider SB, Hsu J, Darby RR, et al. Cortical lesions causing loss of consciousness are anticorrelated with the dorsal brainstem. *Hum Brain Mapp.* 2020;41(6): 1520–1531.
 60. Blumenfeld H. Impaired consciousness in epilepsy. *Lancet Neurol.* 2012;11(9):814.
 61. van der Vlis TAM B, Schijns OEMG, Schaper FLWVJ, et al. Deep brain stimulation of the anterior nucleus of the thalamus for drug-resistant epilepsy. *Neurosurg Rev.* 2019;42(2):287–296.
 62. Kim SH, Lim SC, Yang DW, et al. Thalamo-cortical network underlying deep brain stimulation of centromedian thalamic nuclei in intractable epilepsy: a multimodal imaging analysis. *Neuropsychiatr Dis Treat.* 2017;13:2607–2619.
 63. Kim HY, Hur YJ, Kim HD, Park KM, Kim SE, Hwang TG. Modification of electrophysiological activity pattern after anterior thalamic deep brain stimulation for intractable epilepsy: report of 3 cases. *J Neurosurg.* 2017;126(6):2028–2035.
 64. Zumsteg D, Lozano AM, Wieser HG, Wennberg RA. Cortical activation with deep brain stimulation of the anterior thalamus for epilepsy. *Clin Neurophysiol.* 2006;117(1):192–207.
 65. Ibrahim GM, Sharma P, Hyslop A, et al. Presurgical thalamocortical connectivity is associated with response to vagus nerve stimulation in children with intractable epilepsy. *Neuroimage (Amst).* 2017;16:634–642.
 66. Meindl T, Teipel S, Elmouden R, et al. Test-retest reproducibility of the default-mode network in healthy individuals. *Hum Brain Mapp.* 2010;31(2):237–246.
 67. McGill ML, Devinsky O, Kelly C, et al. Default mode network abnormalities in idiopathic generalized epilepsy. *Epilepsy Behav.* 2012;23(3):353–359.
 68. Gotman J, Grova C, Bagshaw A, Kobayashi E, Aghakhani Y, Dubeau F. Generalized epileptic discharges show thalamocortical activation and suspension of the default state of the brain. *Proc Natl Acad Sci U S A.* 2005;102(42):15236–15240.
 69. Englot DJ, Blumenfeld H. Consciousness and epilepsy: why are complex-partial seizures complex? *Prog Brain Res.* 2009;177(C): 147–170.
 70. Danielson NB, Guo JN, Blumenfeld H. The default mode network and altered consciousness in epilepsy. *Behav Neurol.* 2011;24(1): 55–65.
 71. Vanhauzenhuyse A, Noirhomme Q, Tshibanda LJF, et al. Default network connectivity reflects the level of consciousness in non-communicative brain-damaged patients. *Brain.* 2010;133(1):161–171.
 72. Laureys S, Goldman S, Phillips C, et al. Impaired effective cortical connectivity in vegetative state: preliminary investigation using PET. *Neuroimage.* 1999;9(4):377–382.
 73. Fahoum F, Lopes R, Pittau F, Dubeau F, Gotman J. Widespread epileptic networks in focal epilepsies: EEG-fMRI study. *Epilepsia.* 2012;53(9):1618–1627.
 74. Federico P, Archer JS, Abbott DF, Jackson GD. Cortical/subcortical BOLD changes associated with epileptic discharges: an EEG-fMRI study at 3 T. *Neurology.* 2005;64(7):1125–1130.
 75. Snyder W, Uddin LQ, Nomi JS. Dynamic functional connectivity profile of the salience network across the life span. *Hum Brain Mapp.* 2021;42(14):4740–4749.
 76. Bonnelle V, Ham TE, Leech R, et al. Salience network integrity predicts default mode network function after traumatic brain injury. *Proc Natl Acad Sci.* 2012;109(12):4690–4695.
 77. Jilka SR, Scott G, Ham T, et al. Damage to the salience network and interactions with the default mode network. *J Neurosci.* 2014;34(33):10798–10807.
 78. Zhang CH, Sha Z, Mundahl J, et al. Thalamocortical relationship in epileptic patients with generalized spike and wave discharges—A multimodal neuroimaging study. *Neuroimage Clin.* 2015;9: 117–127.
 79. Walker J, Storch G, Quach-Wong B, Sonnenfeld J, Aaron G. Propagation of epileptiform events across the corpus callosum in a cingulate cortical slice preparation. *PLoS One.* 2012;7(2): e31415.
 80. Caruana F, Gerbella M, Avanzini P, et al. Motor and emotional behaviours elicited by electrical stimulation of the human cingulate cortex. *Brain.* 2018;141(10):3035–3051.
 81. Laureys S, Faymonville ME, Luxen A, Lamy M, Franck G, Maquet P. Restoration of thalamocortical connectivity after recovery from persistent vegetative state. *Lancet.* 2000;355(9217):1790–1791.
 82. Ibrahim GM, Cassel D, Morgan BR, et al. Resilience of developing brain networks to interictal epileptiform discharges is associated with cognitive outcome. *Brain.* 2014;137(10):2690–2702.
 83. Shah A, Jhavar SS, Goel A. Analysis of the anatomy of the Papez circuit and adjoining limbic system by fiber dissection techniques. *J Clin Neurosci.* 2012;19(2):289–298.
 84. Lega BC, Halpern CH, Jaggi JL, Baltuch GH. Deep brain stimulation in the treatment of refractory epilepsy: Update on current data and future directions. *Neurobiol Dis.* 2010;38(3):354–360.
 85. Child ND, Benarroch EE. Anterior nucleus of the thalamus: functional organization and clinical implications. *Neurology.* 2013;81(21):1869–1876.
 86. Feng L, Motelow JE, Ma C, et al. Seizures and sleep in the thalamus: focal limbic seizures show divergent activity patterns in different thalamic nuclei. *J Neurosci.* 2017;37(47):11441–11454.
 87. Sadikot AF, Rymar VV. The primate centromedian-parafascicular complex: anatomical organization with a note on neuromodulation. *Brain Res Bull.* 2009;78(2-3):122–130.
 88. Ilyas A, Pizarro D, Romeo AK, Riley KO, Pati S. The centromedian nucleus: anatomy, physiology, and clinical implications. *J Clin Neurosci.* 2019;63:1–7.
 89. Tyvaert L, Chassagnon S, Sadikot A, Levan P, Dubeau F, Gotman J. Thalamic nuclei activity in idiopathic generalized epilepsy: an EEG-fMRI study. *Neurology.* 2009;73(23):2018–2022.
 90. Velasco AL, Velasco F, Jiménez F, et al. Neuromodulation of the centromedian thalamic nuclei in the treatment of generalized seizures and the improvement of the quality of life in patients with Lennox-Gastaut syndrome. *Epilepsia.* 2006;47(7):1203–1212.
 91. Masterton RAJ, Carney PW, Abbott DF, Jackson GD. Absence epilepsy subnetworks revealed by event-related independent components analysis of functional magnetic resonance imaging. *Epilepsia.* 2013;54(5):801–808.
 92. Bubb EJ, Kinnavane L, Aggleton JP. Hippocampal–diencephalic–cingulate networks for memory and emotion: an anatomical guide. *Brain Neurosci Adv.* 2017;1:239821281772344.
 93. Geller EB. Responsive neurostimulation: review of clinical trials and insights into focal epilepsy. *Epilepsy Behav.* 2018;88:11–20.
 94. Middlebrooks EH, Ver Hoef L, Szaflarski JP. Neuroimaging in epilepsy. *Curr Neurol Neurosci Rep.* 2017;17(4):32.
 95. Larivière S, Bernasconi A, Bernasconi N, Bernhardt BC. Connectome biomarkers of drug-resistant epilepsy. *Epilepsia* 2021;62(1):6–24.
 96. Ponten SC, Bartolomei F, Stam CJ. Small-world networks and epilepsy: graph theoretical analysis of intracerebrally recorded mesial temporal lobe seizures. *Clin Neurophysiol.* 2007;118(4):918–927.
 97. Van Diessen E, Zweiphenning WJEM, Jansen FE, Stam CJ, Braun KPJ, Otte WM. Brain network organization in focal epilepsy: a systematic review and meta-analysis. *PLoS One.* 2014;9(12):e114606.
 98. Wilke C, Worrell G, He B. Graph analysis of epileptogenic networks in human partial epilepsy. *Epilepsia* 2011;52(1):84–93.
 99. Yasuda CL, Chen Z, Beltramini GC, et al. Aberrant topological patterns of brain structural network in temporal lobe epilepsy. *Epilepsia* 2015;56(12):1992–2002.
 100. Varotto G, Tassi L, Franceschetti S, Spreafico R, Panzica F. Epileptogenic networks of type II focal cortical dysplasia: a stereo-EEG study. *Neuroimage* 2012;61(3):591–598.
 101. Xiao F, Koepp MJ, Zhou D. PharmacofMRI: a tool to predict the response to antiepileptic drugs in epilepsy. *Front Neurol.* 2019;10:1203.
 102. Fox MD, Buckner RL, Liu H, Mallar Chakravarty M, Lozano AM, Pascual-Leone A. Resting-state networks link invasive and

- noninvasive brain stimulation across diverse psychiatric and neurological diseases. *Proc Natl Acad Sci.* 2014;111(41):E4367–E4375.
103. Wang Q, Akram H, Muthuraman M, *et al.* Normative vs. patient-specific brain connectivity in deep brain stimulation. *Neuroimage* 2021;224:117307.
 104. Cao M, Wang JH, Dai ZJ, *et al.* Topological organization of the human brain functional connectome across the lifespan. *Dev Cogn Neurosci.* 2014;7:76–93.
 105. Yu M, Sporns O, Saykin AJ. The human connectome in Alzheimer disease—relationship to biomarkers and genetics. *Nat Rev Neurol.* 2021;17(9):545–563.
 106. Tinaz S, Lauro PM, Ghosh P, Lungu C, Horovitz SG. Changes in functional organization and white matter integrity in the connectome in Parkinson's disease. *NeuroImage Clin.* 2017;13:395–404.
 107. De Benedictis A, Duffau H. Brain hodotopy: from esoteric concept to practical surgical applications. *Neurosurgery.* 2011;68(6):1703–1723.