Brain structures and networks responsible for stimulation-induced memory flashbacks during fornical deep brain stimulation for Alzheimer’s disease

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Abstract

Introduction: Fornix deep brain stimulation (fx-DBS) is under investigation for treatment of Alzheimer’s disease (AD). We investigated the anatomic correlates of flashback phenomena that were reported previously during acute diencephalic stimulation.

Methods: Thirty-nine patients with mild AD who took part in a prior fx-DBS trial (NCT01608061) were studied. After localizing patients’ implanted electrodes and modeling the volume of tissue activated (VTA) by DBS during systematic stimulation testing, we performed (1) voxel-wise VTA mapping to identify flashback-associated zones; (2) machine learning–based prediction of flashback occurrence given VTA overlap with specific structures; (3) normative functional connectomics to define flashback-associated brain-wide networks.

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**Results:** A distinct diencephalic region was associated with greater flashback likelihood. Fornix, bed nucleus of stria terminalis, and anterior commissure involvement predicted memory events with 72% accuracy. Flashback-inducing stimulation exhibited greater functional connectivity to a network of memory-evoking and autobiographical memory-related sites.

**Discussion:** These results clarify the neuroanatomical substrates of stimulation-evoked flashbacks.

**KEYWORDS**
Alzheimer’s disease, brain connectivity, deep brain stimulation, fornix, magnetic resonance imaging, memory

1 **INTRODUCTION**

Deep brain stimulation targeting the fornix region (fx-DBS) is currently under investigation for the treatment of Alzheimer’s disease (AD).1–4 A recent paper reported on acute flashback-like phenomena—the involuntary recall of autobiographical memories described by earlier authors as reminiscences5,6—that were experienced by a subset of AD patients during initial postoperative fx-DBS programming; these were associated with specific stimulation settings and appeared to be unrelated to intrinsic whole-brain or hippocampal volume.7 Pioneering 19th and 20th century experiments involving intraoperative direct electrical stimulation of the exposed cortex8–12 provided tremendous insight into the causal relationships between cortical areas and reminiscences, and these insights still stand.5,6 However, the relationship between these memory phenomena and deep brain structures has not been systematically investigated and it remains unknown which specific structures give rise to the flashbacks reported in the aforementioned fx-DBS population. Although the fornix is the designated target for therapeutic neuromodulation and is well-known to be a critical component of the brain’s memory circuits,13–15 the stimulated region encompasses a number of other white matter tracts and nuclei that could also conceivably play a role. These include the anterior commissure,14,16 septal nuclei,17 and bed nucleus of stria terminalis,18 each of which has been implicated in memory function and—like the fornix—is intimately connected to medial temporal lobe structures like the hippocampus and entorhinal/perirhinal cortex.19–21

To address this question and further elucidate the architecture of human memory experience, we investigated the neuroanatomical substrate of stimulation-induced flashbacks in patients with mild AD who were undergoing bilateral fx-DBS. We hypothesized that the fornix itself would be a key contributor to this phenomenon. However, given that the fornix is presumably engaged in most cases of fx-DBS and yet not all patients experience flashbacks, we expected that nearby structures might also be necessary substrates. To facilitate this investigation, we performed (1) volume of tissue activated modeling and voxel-wise linear modeling of stimulation resulting in memory events as compared with stimulation without events; (2) machine learning-based prediction of flashback occurrence given involvement of specific brain structures; (3) normative resting state functional magnetic resonance imaging (rsfMRI) connectomics involving the aforementioned stimulation volumes; and (4) validation of the normative connectivity results by comparison with both brain areas previously shown to elicit flashbacks when electrically stimulated and also with regions and networks heavily implicated in autobiographical memory and memory retrieval.

2 **MATERIALS AND METHODS**

The analysis involved behavioral observations and pre- and postoperative structural MRI data from 39 patients with mild AD (Table 1) who were treated with bilateral fx-DBS as part of a previously described, 42-patient, multicenter clinical trial (ClinicalTrials.gov)
number: NCT01608061). Each patient who was enrolled in this trial, which was approved by independent research ethics boards at every participating site, provided written informed consent. As specified previously, patients were diagnosed by standardized criteria and expert examination, with the criteria for mild probable AD being scores of 0.5 or 1 on the Clinical Dementia Rating scale and scores of 12-24 on the Alzheimer’s Disease Assessment Scale-11. Additional inclusion and exclusion criteria for enrollment are outlined in Table S1. All patients were implanted with bilateral quadripolar (four contacts each) DBS electrodes (model 3387, Medtronic, Minneapolis, MN) and connected to an implantable pulse generator (Table S2). Of the 42 patients enrolled in the clinical trial, 3 were excluded due to inadequate image quality, which precluded precise electrode localization.

During initial postoperative programming of the device, each electrode contact (four per lead) was tested with high-frequency (130 Hz, 90 μs) stimulation beginning at a low voltage (~1 volt) and increasing in 1 volt increments up to the maximal voltage tolerated (max = 10 volts). For each contact (eight per patient), if any setting induced a flashback, the lowest flashback-inducing voltage setting was sampled, along with the voltage setting(s) immediately below and—if present—above that did not induce flashbacks. For all contacts without induced memory events, the largest voltage setting tested was utilized. This conservative selection method was designed to avoid false-positive results. Classification of memory events was determined using the TEMPau scale.

Although the quality of reminiscences elicited by electrical stimulation varies, this paper’s primary aim was to elucidate the neural correlates of flashback phenomena in general rather than those underlying subtle variation in memory quality; as such, only presence (TEMPau score 1-4; “memory-yes”) or absence of memory events (TEMPau score 0; “memory-no”) were considered for analysis.

### 2.1 Volume of tissue activated modeling

First, we used patient-specific anatomic MRI, stimulation settings, and volume of tissue activated (VTA) modeling techniques to estimate the extent of tissue directly modulated by DBS during each observation. Following non-uniformity correction of all MR images, VTA modeling was conducted using a well-described pipeline (lead-DBS v2.0; https://www.lead-dbs.org/). This involved localization of electrode contacts on postoperative MRI acquisitions by two experienced users (AH and GJBE), non-linear normalization to MNI152 standard space (using “low variance” ANTs (http://stnava.github.io/ANTs/) with an additional subcortical affine transformation when necessary) via coregistered preoperative images, and estimation of the shape/extent of the electrical field using finite element method modeling with 0.2 V/mm gradient thresholding (FieldTrip-SimBio pipeline; http://fieldtriptoolbox.org). A VTA was estimated for each of the sampled “memory-yes” and “memory-no” observations using the corresponding stimulation setting (contact and voltage) and peri-electrode conductivity estimates (gray matter: 0.33 S/m; white matter: 0.14 S/m; cerebrospinal fluid: 1.79 S/m; electrode contact: 10⁹ S/m; insulated electrode components: 10⁻¹⁴ S/m) derived from standard space tissue priors. Left-sided VTAs were flipped in the sagittal plane to facilitate group-level analysis. Figure 1 provides a visual summary of the major neuroimaging processing steps used in this paper.

### 2.2 Whole-brain voxel-wise analysis of flashback-inducing VTAs

Next, “memory-yes” and “memory-no” VTAs were stratified by contact and stimulation voltage in order to examine the possible effects of these factors independent of VTA location. Simple linear models were estimated to assess the relationship between contact and voltage, and between voltage and memory events. Subsequently, whole-brain voxel-wise logistic regression comparing “memory-yes” and “memory-no” VTAs was conducted to identify brain areas associated with flashbacks while controlling for stimulation voltage.

### 2.3 Support-vector machine classification

Support-vector machine (SVM) learning was then used to further interrogate the brain structures driving flashbacks and determine the extent to which their involvement could predict memory events. Specifically, the presence and extent (in mm³) of overlap between VTAs and structures (as defined using a manually segmented high-fidelity diencephalic atlas) within memory-associated regions were calculated and used to classify VTAs as “memory-yes” or “memory-no”. Modeling was performed with balanced data sets of 343 observations for...
FIGURE 1 Visual summary of neuroimaging methods. The major methodological steps (colored arrows) and their corresponding descriptions are shown alongside exemplar brain images. First, patient-acquired pre- and postoperative anatomic MRI scans were coregistered together, and each patient’s leads were precisely localized in patient space based on the post-operative acquisition (red arrows). Next, the coregistered patient scans were normalized to a standard MNI152 template, and the resultant transforms were used to warp the lead models to MNI space (green arrows). VTAs were then modeled for each “memory-yes/memory-no” observation in MNI space using the corresponding stimulation settings and conductivity estimates from standard space tissue priors (turquoise arrow). Finally, the created VTAs ($n = 386$) were employed as (1) inputs for local voxel-wise mapping analysis (yellow arrow) and (2) seeds for rsfMRI functional connectivity mapping (magenta arrow). MNI = Montreal Neurological Institute; rsfMRI = resting state functional magnetic resonance imaging; VTA = volume of tissue activated both “memory-yes” and “memory-no” groups; additional observations for the “memory-yes” cohort were created by random sampling with replacement. The most parsimonious model that best classified these observations was identified and validated using a 10-fold (random split in 10 balanced (“memory-yes” vs “memory-no”) groups, 3 with 35 members per group and 7 with 34) cross-validation approach. In addition, an alternative model classifying memory events on the basis of voltage and electrode contact alone was created for comparative purposes.
2.4 | Connectomic mapping of flashback-inducing VTAs

To investigate wider brain networks associated with flashback-inducing stimulation, whole-brain connectivity maps were derived for each VTA using a high-quality normative 3 Tesla rsfMRI data set (http://neuroinformatics.harvard.edu/gsp/) as described previously.25,28–31 Normative data were used for the primary analysis instead of patient-derived rsfMRI images because the latter were not acquired in the majority of patients and were of low fidelity (eg, 1.5 Tesla) when present. Per this connectomic method, correlations with the seed region (ie, the VTA) were obtained for each voxel in the brain based on the time course of low-frequency blood oxygen level–dependent (BOLD) signal fluctuations across 1000 healthy subjects (age range: 18–35 years; 57% female) (in-house MATLAB script, The MathWorks, Inc., Version R2018a. Natick, MA). Whole-brain voxel-wise logistic regression was then conducted to identify brain areas whose connectedness was associated with incidence of memory events. Finally, to validate these normative results, a supplemental connectivity analysis was performed using a disease-specific connectome assembled from 12 AD-DBS patients with available preoperative rsfMRI imaging.

2.5 | Connectomic overlap with canonical memory networks

To explore how these findings related to the relevant human literature, we analyzed the spatial overlap between our DBS-induced flashback functional connectivity network and (1) brain structures identified through meta-analysis as evoking memory events when stimulated;32 and (2) brain regions whose BOLD response is associated with memory as per Neurosynth (http://neurosynth.org) meta-analyses of published task-based fMRI studies.32 For the former, we selected bilateral probabilistic regions-of-interest (ROIs) using a standardized atlas (Harvard-Oxford cortical-subcortical atlas) in MNI space (Figure 4A).33 For the latter, we used meta-analytic association maps of voxels linked to autobiographical memory and memory retrieval across 84 and 228 pre-existing fMRI studies, respectively. To assess whether overlap with these entities was non-random, we permuted the voxels in the DBS-induced flashback connectivity network 1000 times and determined the extent of each permutation’s overlap with the aforementioned ROIs and meta-analytic association maps. As an additional validation, we used the Neurosynth “decoder” to identify behavioral functional networks—derived from all available fMRI meta-analyses—with the greatest spatial similarity to the flashback network.32,34

2.6 | Statistics

All statistical analyses were performed using R (v3.4.4; https://www.r-project.org) and RMINC (https://github.com/Mouse-Imaging-Centre/RMINC). The pROC (version 1.16.2) package was used to calculate the receiver-operating characteristic (ROC) curve and the e1071 (version 1.7-3) package was used for the support vector machine (SVM) analysis. Whole-brain correction for multiple comparisons was performed using the false discovery rate (FDR; voxel-wise threshold of $P_{FDR} < 0.05$). To strengthen any voxel-wise VTA mapping results and address the presence of non-independent observations in our data, we also conducted a non-parametric permutation analysis at the cluster level. Following a previously described approach, the clinical score associated with each VTA was randomly shuffled across all lead-contact combinations, creating 10,000 new permuted data sets. Summary Q statistics were obtained for each data set and the summary statistic magnitudes of the actual voxel-wise map and the permuted maps were compared to discern the validity of the observed results.35–37

3 | RESULTS

Of the 39 patients included for analysis, 18 (46%) patients experienced flashback phenomena at least once, whereas 21 patients (54%) never experienced flashbacks. Baseline demographic characteristics were similar between these two groups of patients (Table 1). In total, 43 “memory-yes” and 343 “memory-no” observations were sampled, and a separate VTA was created for each observation. Stratification of “memory-yes” and “memory-no” VTAs by contact and stimulation voltage revealed that stimulation delivered from the more dorsal three contacts (contacts “1-3”) on occasion produced acute memory events, while stimulation at the ventral-most contact (contact “0”) never did (Figure 2A). There was a significant effect of contact on voltage, with mean voltage increasing incrementally as stimulation moved dorsally ($P < 0.001$, voltage at contacts 0-3 [ventral to dorsal, mean±standard deviation]: contact 0: 5.16±1.22 volts, contact 1: 5.66±1.61 volts; contact 2: 6.36±2.01 volts; contact 3: 7.57±2.26 volts). This likely reflected a greater tendency for stimulation to evoke unpleasant autonomic side-effects at ventral contacts (thus limiting the voltage tolerated), which were in proximity to hypothalamic nuclei.7 Voltage was significantly lower for “memory-yes” compared to “memory-no” VTAs both overall (mean±standard deviation, “memory-yes”: 5.67±2.01 volts; “memory-no”: 6.36±2.05 volts, $P < 0.01$) and individually for contacts 2 (mean±standard deviation, “memory-yes”: 5.63±1.86 volts; “memory-no”: 6.54±2.00 volts, $P < 0.05$) and 3 (mean±standard deviation, “memory-yes”: 6.14±2.29 volts; “memory-no”: 7.81±2.18 volts, $P < 0.01$) (Figure 2B).

3.1 | Whole-brain voxel-wise VTA analysis

Using whole-brain voxel-wise logistic regression to investigate the association of VTA location and memory events, we identified two significant clusters (each voxel passed FDR correction at $P_{FDR} < 0.05$): a dorsal cluster in the anterior diencephalon, impinging on the column of the fornix, septal region, bed nucleus of stria terminalis (BNST), and anterior commissure, associated with greater likelihood of memory events; and a ventral cluster in the hypothalamus associated with a lower likelihood of memory events (Figure 2C; Table S3). To confirm that these results were not driven by patient-specific
FIGURE 2  Results of voxel-wise VTA mapping of flashback-inducing stimulation. (A) Bar graph showing the number of stimulations (count) at each contact (ventral to dorsal, “0” to “3”) that did (red) or did not (blue) produce acute memory events. Note that no memory events were reported during stimulation of the ventral-most contact. (B) Boxplot showing the minimal voltage (memory event, red) and maximal voltage (no memory event, blue) for the stimulations at each contact. (C) Results of voxel-wise VTA mapping show voxels significantly ($P_{FDR} < 0.05$) positively (warm colors) and negatively (cool colors) associated with memory flashbacks. Only the dorsal cluster of significant voxels lay within a region (outlined in green) established as non-random by non-parametric permutation testing ($P_{permutation} < 0.05, n = 10,000$ permutations). The fornix (red), BNST (yellow), and anterior commissure (green) are overlaid in faded colors on sagittal (left), coronal (middle), and axial (right) T1-weighted MNI152 brain slices. BNST = bed nucleus of stria terminalis; FDR = false discovery rate; MNI = Montreal Neurological Institute; VTA = volume of tissue activated; *$P < 0.05$; **$P < 0.01$

characteristics of individuals who reported memory flashbacks, a linear mixed-effect model analysis with subject as random variable (repeated measure design) was performed, looking only at patients who had at least one memory event. In these patients, we compared each setting that elicited flashbacks with a matched setting at the same contact, just below in voltage, that did not elicit flashbacks. This supplementary analysis, which used threshold-free cluster enhancement $^{38}$ (TFCE; voxel threshold of $P_{Bonferroni} < 0.0001$) for multiple comparisons correction, reaffirmed the results of the whole sample analysis, identifying a nearly identical cluster of voxels to be significantly associated with memory flashbacks (Figure S1). Only the dorsal cluster lay within a region that was shown by non-parametric permutation testing to be non-random ($P_{permutation} < 0.05, n = 10,000$ permutations).

3.2 Support-vector machine classification

SVM modeling reinforced the role of the fornix, BNST, and anterior commissure in eliciting memory flashbacks upon electrical stimulation.
A model using stimulation voltage, volume overlap (continuous) with BNST, and impingement (binary) on fornix and anterior commissure was found to be most successful at classifying VTAs (Figure 3). This model achieved 72% accuracy (true-negative rate: 0.68, false-negative rate: 0.24, false-positive rate: 0.32, true-positve rate: 0.76) and 77% area under the receiver-operating characteristic (ROC) curve (AUC) compared to chance performance (50%). The addition of other components like the septal region, other diencephalic structures, baseline Alzheimer’s Disease Assessment Scale-Cognitive Subscale scores, or demographic features (eg, age or sex) did not improve performance. The alternative model, which disregarded anatomic structure involvement and used only voltage and electrode contact, performed more poorly (67% accuracy; true-negative rate: 0.46, false-negative rate: 0.11, false-positive rate: 0.54, true-positve rate: 0.89; 67% AUC). Ten-fold cross-validation of the best model classified VTAs with 71% accuracy (true-negative rate: 0.64, false-negative rate: 0.36, true-positve rate: 0.78) (Figure S2).

3.3 | Connectomic mapping of flashback phenomena

Whole-brain voxel-wise logistic regression of the VTA-specific connectivity maps identified a number of brain areas whose
3.4 | Connectomic overlap with canonical memory networks

The extent of overlap between this DBS-induced flashback connectivity profile and previously reported memory-eliciting ROIs and memory-related meta-analytic association maps was then calculated. As verified by permutation testing (n = 1000 permutations), considerable non-random overlap was observed between the flashback...
connectome and several ROIs (amygdala, hippocampus, middle temporal gyrus, parahippocampal gyrus, and insular cortex) as well as both the "memory retrieval" and " autobiographical memory" association maps (Figure 4B-D). Using the Neurosynth " decoder" we identified the top five most similar behavioral networks to be " autobiographical" \(r = 0.24\), " episodic" \(r = 0.20\), " retrieval" \(r = 0.17\), " autobiographical memory" \(r = 0.17\), and " episodic memory" \(r = 0.17\).

4 | DISCUSSION

Using VTA modeling and normative functional connectomics, we uncovered brain areas and networks likely involved in previously reported stimulation-induced flashback phenomena.\(^7\) To our knowledge, this is the first systematic experimental analysis of deep subcortical stimulation causing acute reminiscences in humans. This marks a meaningful step beyond modern causal evidence, which has by necessity derived primarily from animal work or case series of human brain lesions. In doing so, this study marks a rare addition to the research legacy of classical 19th and 20th experiments\(^8–12\) that meticulously elucidated the relationship between various brain areas and memory responses through direct intraoperative stimulation of the exposed cortex.

The region of the antero-dorsal diencephalon emerged as important for inducing memory flashbacks; insights derived from machine learning moreover suggested that BNST,\(^39,40\) fornix,\(^15,41,42\) and anterior commissure,\(^43–45\) in particular, contributed to these events. The fact that a model incorporating overlap with these structures performed better than an alternative model that relied solely on stimulation voltage and contact—specifically avoiding false-positive identification of flashbacks—emphasizes that the occurrence of memory events cannot be explained fully by stimulation intensity or relative depth, instead being more accurately predicted by " hitting" specific neuroanatomical substrates. All three of these structures have been implicated extensively in memory function.\(^14,15,19,39–45\) Of interest, the volume of VTA overlap appeared to make a difference with respect to flashback induction for the gray matter structure (BNST) but not for the two white matter structures. This may reflect the continuous nature of white matter axons, and the notion that impingement on a circumscribed cross-section of a given bundle will propagate along its extent.

Through normative rsfMRI mapping, we found that flashback-associated VTAs were preferentially connected to a wider brain network that primarily comprised the medial and lateral temporal lobes, prefrontal regions, insular cortex, and cingulate areas. These same regions are implicated in autobiographical memory recall by prior brain stimulation work\(^5\) and functional neuroimaging studies,\(^22\) as well as in a recent normative mapping analysis of brain lesions causing amnesia.\(^19\) Indeed, the BNST, fornix, and anterior commissure are key local substrates of flashbacks evoked during fornix region DBS, and that flashback-inducing stimulation interacts with a distributed brain network previously implicated in autobiographical memory retrieval. These findings might provide the basis for future work investigating therapies to stabilize or improve memory in patients with dementia.

AUTHOR CONTRIBUTIONS

Jürgen Germann, Gavin J.B. Elias, and Alexandre Boutet conceived the experiment. Wissam Deeb, Bryan Salvato, Leonardo Almeida, Kelly D. Foote, Paul B. Rosenberg, David F. Tang-Wai, David A. Wolk, Anna D. Burke, Stephen Salloway, Marwan N. Sabbagh, M. Mallar Chakravarty, Gwenn S. Smith, Constantine G. Lyketsos, Michael S. Okun, and Andres M. Lozano collected the data. Andreas Horn performed MR image registration and electrode localization. Alexandre Boutet and Keshav Narang constructed the VTAs. Aaron Loh constructed the patient-specific connectome. Jürgen Germann and Gavin J.B. Elias performed the data analysis. Jürgen Germann, Gavin J.B. Elias, and Clemens Neudorfer wrote the manuscript. All authors edited the manuscript. Andres M. Lozano supervised the project.
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COMPETING INTERESTS
AML is scientific director of Functional Neurumodulation and reports consultant fees from Medtronic, Abbott, and Boston Scientific. CGL reports consultant fees from Biogen, Lilly, Eli Lilly, and the NFL benefits office. MSO reports consultant fees from Biogen, Lilly, Eisai, Genentech, Roche, Novartis, and Avid. DAW reports consultant fees from Lilly, Merck, Jannsen, GE Healthcare, and Neurionx. MNS reports consultant fees from Allergan, Biogen, Roche, Genentech, Cortexyme, Bracket, and Sanofi. The other authors have nothing to disclose.

DATA AVAILABILITY STATEMENT
The data supporting these analyses are available from the corresponding author upon reasonable request.

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SUPPORTING INFORMATION
Additional supporting information may be found online in the Supporting Information section at the end of the article.

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