Optimal Stimulation Sites and Networks for Deep Brain Stimulation of the Fornix in Alzheimer's Disease

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METHODS

Retrospective analysis of a multi-center cohort of 50 patients (25 female, mean age: 67.5 ± 7.9 years) who underwent fornix-DBS to treat mild AD. Pre- and post-operative MRI volumes of the participants were processed using the lead-DBS pipeline (lead-dbs.org)⁷, normalization and electrode localization were manually refined (WarpDrive tool) and clinical outcomes were measured by changes in the Alzheimer's Disease Assessment Scale-cognitive subscale 11 (ADAS-cog score).

At the fibertract level, the subjects were randomly assigned to a Training cohort (n=30) or Test cohort (n=20). Streamlines correlated to clinical improvement were identified in the Training cohort (Leave-one-out and K-fold approaches) and then used to predict clinical improvement of the Test cohort (cross-prediction). For the sweet spot and network mapping analyses, optimal sites and networks were investigated using the whole cohort and results were cross-validated with a Leave-one-out and multiple K-fold approaches.

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INTRODUCTION

Alzheimer's Disease (AD) is the most common neurodegenerative disease burdening healthcare attention without an effective treatment to date¹. Deep brain stimulation (DBS) to the fornix is under investigation for mild AD with completed phase I and II trials^{2,3}, showing cognitive improvement in some patients and deterioration in others³. As observed in other conditions treated with DBS⁴⁻⁶, an explanation of these outcomes could lie in variance in electrode placement engaging distinct neural circuits.

OBJECTIVES

To identify optimal electrode location by investigating effects of stimulation on three levels: 1) Fibertract: white matter tracts traversing Electric-fields informed by a high resolution normative connectome⁸.

2) Sweetspot: local-level voxel-wise analysis to identify an optimal stimulation site. 3) Network Mapping: whole-brain network effect informed by resting state fMRI data of 1000 healthy subjects⁹.

DISCRIMINATIVE **NORMATIVE CONNECTOME: TRACT-WISE COLOR EACH FIBER TRACT TRACT-WISE SPLIT OF E-FIELDS SPEARMAN CORRELATION** FIBER TRACT PROFILE **BY R-VALUE EXAMPLE FIBERTRACT EXAMPLE FIBERTRACT PREDOMINANTLY MODULATED** MAGNITUDE 0.8 V/M %CLINICAL IPROVEMEN **BY TOP RESPONDERS** E-FIELD SWEET SOUR 0.4 V/M **E-FIELD MAGNITUDE** TRACTS TRACTS **ELECTRODE RECONSTRUCTION** B **E-FIELDS IN STANDARD SPACE VOXEL-WISE SWEETSPOTS VOXEL-WISE GRID SPEARMAN CORRELATION & E-FIELD ESTIMATION** MAGNITUDE 0.8 V/M %CLINICAL APROVEMEN SWEET SOUR 0.4 V/M **E-FIELD MAGNITUDE** E-FIELD MAGNITUDE **SPOTS SPOTS CORRELATE TIME-SERIES FUNCTIONAL CONNECTIVITY CORRELATE INDIVIDUAL "FINGERPRINTS" "FINGERPRINT"** WITH %-IMPROVEMENT **R-MAP** CONNEC ΤΙVITY IMPROVE TO VTA MENT "OPTIMAL"

RESULTS

These analyses demonstrated that:

- 1) Modulation of the Papez' circuit and stria terminalis associated with cognitive improvement (R = 0.45 at p = 0.026).
- 2) Optimal stimulation site resided at the interface between fornix and bed nucleus of the stria terminalis (R = 0.29 at p 0.016).

SWEETSPOT LEVEL

× 1000 FC-CONNECTOMES



NETWORK MAPPING LEVEL

CONNECTIVITY PROFILE FOR

MAXIMAL CLINICAL IMPROVEMENT



3) Modulating specific distributed brain networks accounted for optimal outcomes (R = 0.30 at p = 0.015).





0.1 -0.1 -0.2

SPATIAL SPATIAL

-200

CONCLUSIONS

A potential optimal stimulation target for Alzheimer's Disease treatment with fx-DBS is proposed.

- Stimulation of Papez' circuit and bed nucleus of the stria terminalis associated with cognitive improvement. 1)
- Optimal stimulation site: intersection between fornix and bed nucleus of the stria terminalis. 2)

Modulating specific whole-brain networks seems crucial for DBS-induced positive effects on cognition. 3)

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SWEETSPOT 0.3 0.2 R = 0.41LOOCV: p = 0.001R = 0.29, p = 0.01610-fold CV: R = 0.33, p = 0.013 \geq

RESULTS SUMMARY



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Results summary including DBS fiber filtering, sweetspot mapping and network-mapping models. Three levels of analysis explain similar amount of variance of clinical outcomes when analyzed in circular nature (scatterplots ~16-19%) and led to significant cross-validation of clinical outcomes across leave-one-patient-out and k-fold designs. For visualization, patients were distributed into three groups based on their ADAS-cog score results a year after stimulation (poor responders: blue, middle responders: yellow and top responders: red). Gray shaded areas represent 95% confidence intervals. Analysis results were superimposed on slices of Big Brain atlas in MNI 152 space.



