



Neuroinformatics and Brain Connectivity Lab

HABENULAR ALTERATIONS IN RESTING STATE FUNCTIONAL CONNECTIVITY AMONG AUTISTIC INDIVIDUALS

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To address a critical gap in autism reward-based literature, this study examined habenula resting AIM state functional connectivity (rsFC) using functional magnetic resonance imaging (fMRI) data.

BACKGROUND

neurodevelopmental disability Autism IS a characterized by social communication differences + restrictive interests and/or repetitive behaviors [1]. The **reward-based theoretical framework** suggests that altered reward circuitry contributes to core autism symptoms [2,3], which has been supported by robust evidence of altered reward-based corticostriatal circuitry [4] across development [5].

Recent research has revealed autism-related structural alterations in the habenula [6], a small

Neuroimaging Data

Resting state functional MRI (rs-fMRI) data were accessed for 1,584 ABIDE participants (N=705 autism; M=16.26±8.15 years) [8, 9].

rs-fMRI data for autistic (ASD) and neurotypical (NT) individuals were preprocessed and quality controlled using fMRIPrep [10] and MRIQC [11]. Data Exchange



METHODS

Habenula Region of Interest

Voxelwise time series were extracted from a handdrawn bilateral habenula ROI defined using the MNI 152subject averaged T1 template brain [12, 13].



Analyses

modulates that epithalamic structure the dopaminergic pathways of the reward network and is associated with **motivation and emotion**[7].

Despite these findings, the potential alterations in habenular functional connectivity (FC) remain unexplored.

1. Mapped habenula FC in autistic individuals by conducting a whole-brain rsFC analysis (ASD+NT, ASD>NT).

2. Probed for age-related changes in habenula FC from childhood to early adulthood (5–21 years) between ASD and NTs.

3. Examined relationships between altered habenula FC and behavior with measures social motivation (SM), of social communication (SC), executive functioning (EF), and daily living skills (DLS).

All thresholded rsFC maps were generated using Nilearn [14] Python package and functionally decoded with Neurosynth [15].



were observed, aligning with hypotheses that cerebellar connectivity alterations are a key component of autism [19].

Habenula hyperconnectivity was significantly associated with SM, EF, and DLS, emphasizing the critical role of the habenula in motivated and adaptive behaviors, which are crucial for positive life outcomes and pose challenges for autistic individuals in achieving independence [20]. Taken together, these results contribute to the emerging evidence that the dopaminergic reward system plays a critical role in the pathophysiology of autism.



ACKNOWLEDGMENTS

Primary funding for this project was provided by the FIU Embrace Center for Advancing Inclusive Communities, the FIU Office of Research & Economic Development, and the FIU Undergraduate to Graduate Program. Contributions from co-authors were provided with support from NIH R01-MH09606 (JAP, ARL), U01-DA041156 (ARL), U54-MD012393 (RA), and NIH R35GM153434. This project was also supported by funding provided to each of the contributing ABIDE research institutions, which are listed for each site at https://fcon_1000.projects.nitrc.org/indi/abide.

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