

# Assessment of Functional Connectome in End-Stage Organ Disease Patients after Life-Threatening Surgery

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## Purpose

The purpose of this study was to assess the organization of intrinsic functional brain networks (functional connectome) in neurologically asymptomatic patients with end-stage organ disease who had undergone major surgery for life-threatening conditions and compare it to a control group.

## Materials and Methods

Resting-state functional magnetic resonance imaging (rest-fMRI) was performed in 10 adult, post-operative patients with end-stage organ disease. The patients (7 men, 3 women, mean age  $57.9 \pm 7.4$  years) had undergone: n=4 coronary artery bypass graft for heart failure, n=2 orthotopic liver transplantation for non-alcoholic cirrhotic liver failure, n=3 mitral valve repair for heart failure, n=1 pancreaticoduodenectomy for pancreatic papillary tumor. Rest-fMRI was acquired within 48 hours after intensive care unit discharge. Structural brain MR imaging was acquired with T1-weighted, T2-weighted, diffusion weighted imaging, and susceptibility weighted imaging along with 3D isotropic T1-weighted MR images. MR examinations were performed on a 3T MR scanner (Discovery 750w, General Electric Healthcare, Milwaukee, WI, USA). Ten age- and sex-matched healthy controls were studied with the same protocol. Brain functional networks were analyzed by calculating the interregional correlation of low-frequency fluctuations in spontaneous brain activity. Weighted graph-based models were employed to topologically recognize reproducibly determined large-scale functional networks, including default mode, salience,

dorsal attention, sensorimotor, visual and language networks using a seed-based approach with Functional Connectivity Toolbox (CONN) (<https://web.conn-toolbox.org/>) running under MatLab (The MathWorks Inc., Natick, MA, USA).

A  $p$ -uncorrected  $< 0.0025$  was used to threshold connectomes for voxel-wise paired statistical analysis to take in account the lateralization effect. Network comparisons were thresholded using a False Discovery Rate (FDR) cluster-level correction approach in patients and controls using the MatLab function "mafdr" for  $p$ -uncorrected  $< 0.0025$  obtaining the  $p$ -FDR  $< 0.077$ .

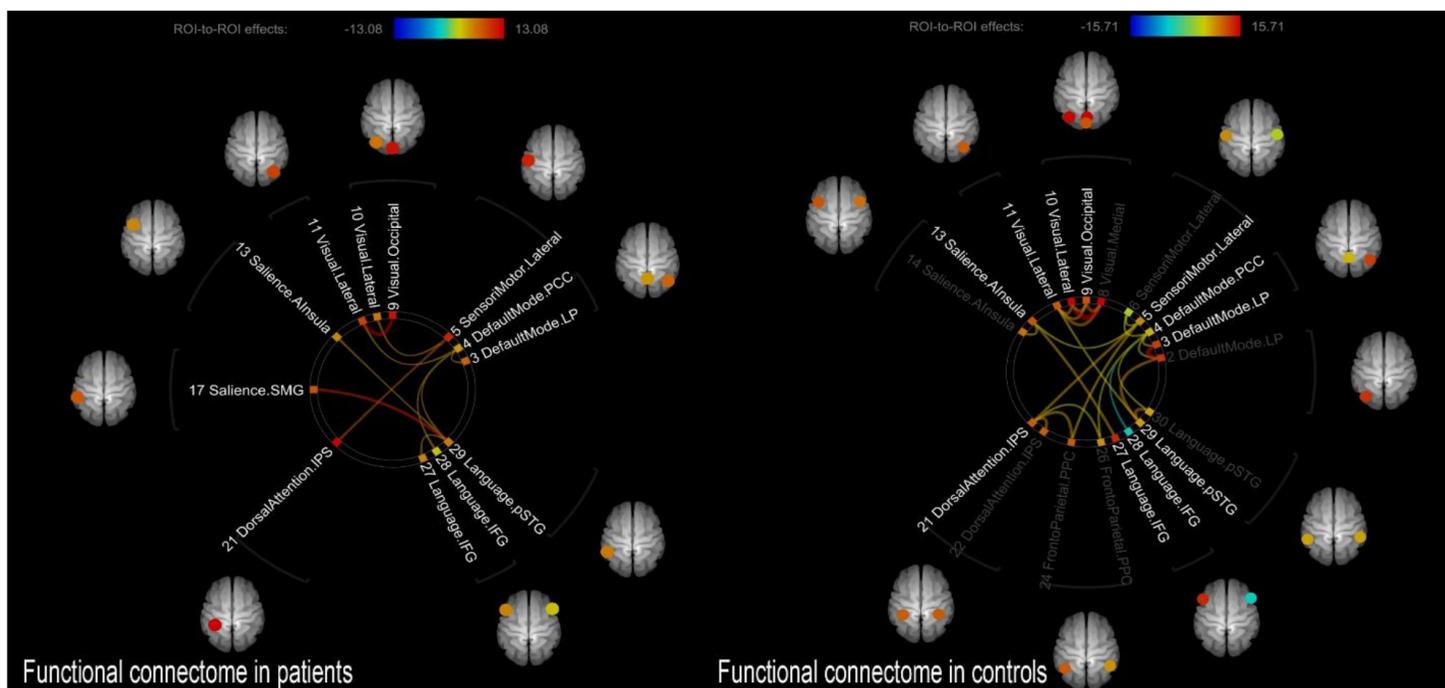
## Results

Functional connectivity in patients and controls was successfully assessed for the default mode, salience, dorsal attention, sensorimotor, visual, and language networks of the brain. There was statistically significant ( $p < 0.05$ ) reduced connectivity between seeds in the default mode, salience, sensorimotor, and language networks in patients compared to controls (Fig. 1).

No acute structural lesions were observed in patients at MR imaging.

## Conclusion

Assessment of functional connectome in end-stage organ disease patients is feasible in routine clinical setting. Rest-fMRI can demonstrate reduced connectivity of several intrinsic functional networks in critically ill, post-operative patients and it may be a useful prognostic indicator of early and long-term clinical outcome.



## References

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**Fig. 1 - Functional Connectome.** There was statistically significant ( $p < 0.05$ ) reduced connectivity between seeds in the default mode, salience, sensorimotor, and language networks in patients compared to the controls