The trend of disruption in the functional brain network topology of Alzheimer's disease

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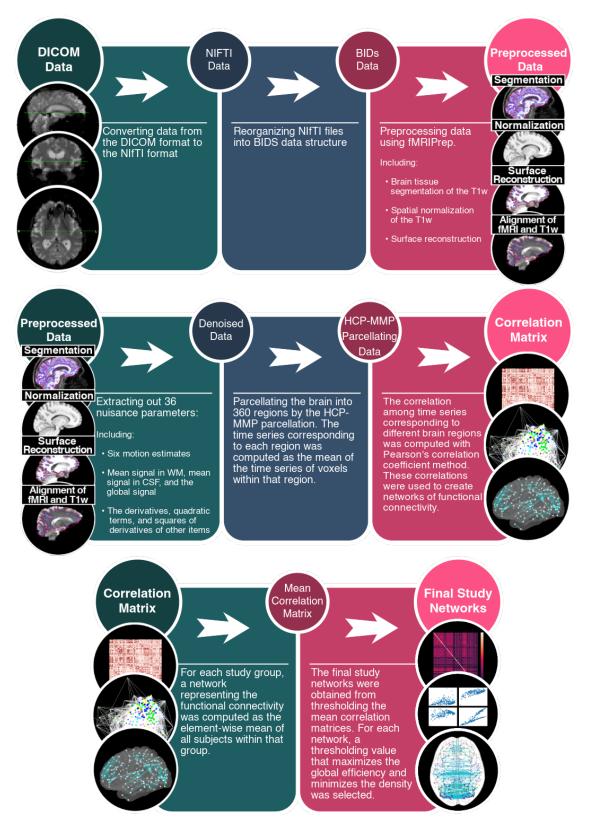


Figure S1. The overall processing pipeline of the present study. *Nilearn*, *fMRIPrep*, and *draw.io* were used to create this figure.

a					b				
CN	1.000	0.882	0.824	0.683	CN	0.869±0.017	0.858±0.012	0.809±0.028	0.704±0.031
EMCI	0.882	1.000	0.840	0.710	EMCI	0.861±0.011	0.861±0.011	0.821±0.026	0.725±0.03
LMCI	0.824	0.840	1.000	0.799	LMCI	0.807±0.03	0.819±0.027	0.813±0.016	0.775±0.024
AD	0.683	0.710	0.799	1.000	AD	0.707±0.034	0.728±0.033	0.778±0.026	0.786±0.018
	CN	EMCI	LMCI	AD		CN	EMCI	LMCI	AD

Figure S2. (a) The PCC between the four main study networks. (b) For each study group, the 23 subjects of each group were randomly divided into two subgroups of 11 and 12 networks. Then, the mean networks of these 12 and 11 subjects were constructed, and the PCC between these two mean networks was computed. This process was repeated 100 times resulting in 100 PCC matrices. This panel shows the element-wise mean and SD of these matrices.

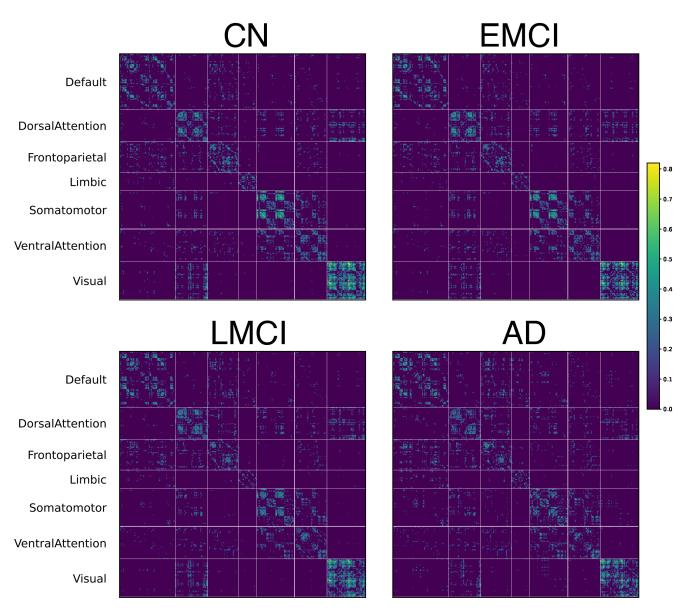


Figure S3. The correlation matrix of each study group

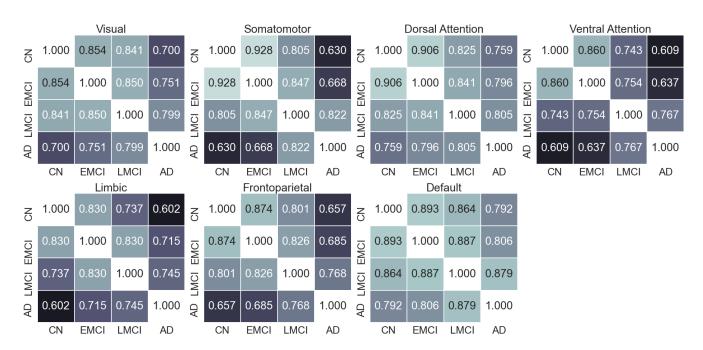


Figure S4. For each study group, the modular sub-networks containing the vertices belong to each module were computed. Then, for each module, the PCC was computed among the adjacency matrices of the modular sub-network of each study network.

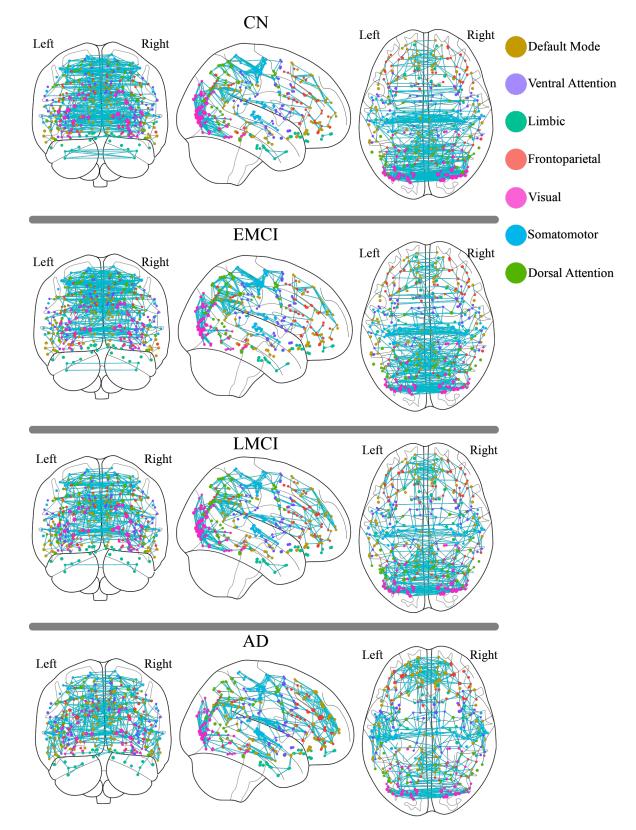


Figure S5. The top 1 percent highly-weighted links of each group network. Vertex color represents the module the vertex belongs to, and the vertex size is proportional to the vertex strength. It shows that there is a trend towards the increase in the number of default mode links as well as the decrease in the inter-hemisphere links in dorsal attention.

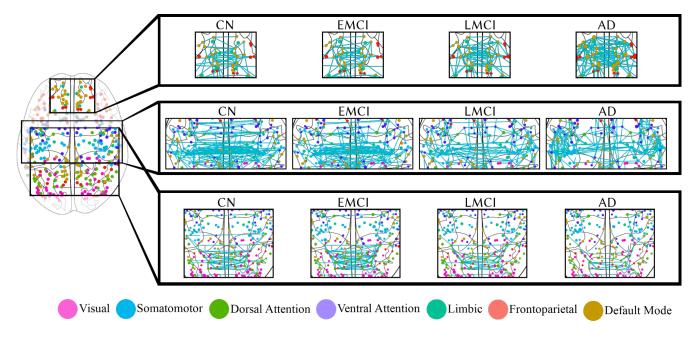


Figure S6. The axial view of the overall gradually changing patterns of top 1 percent highly-weighted links.