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Nodal statistics-based structural pattern detection for graph collections characterization

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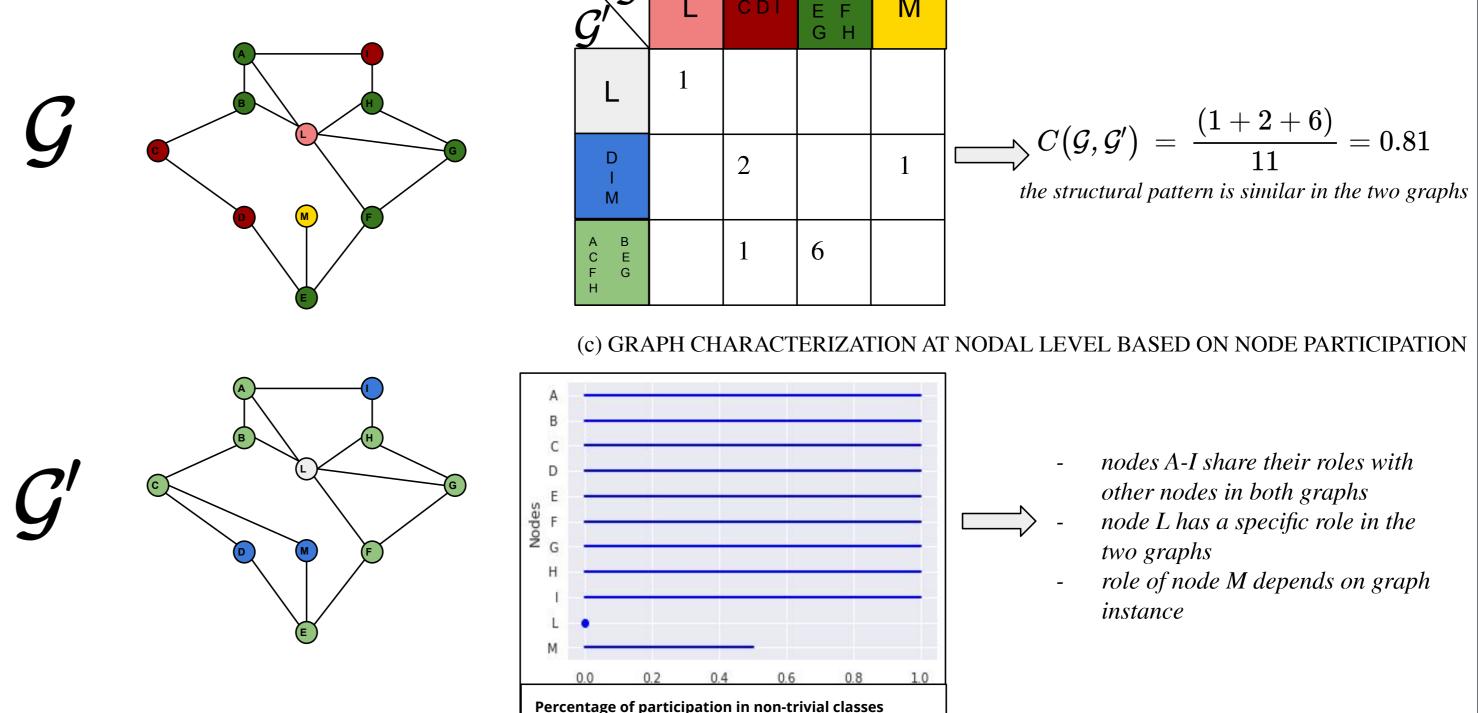


http://arxiv.org/abs/2210.01053

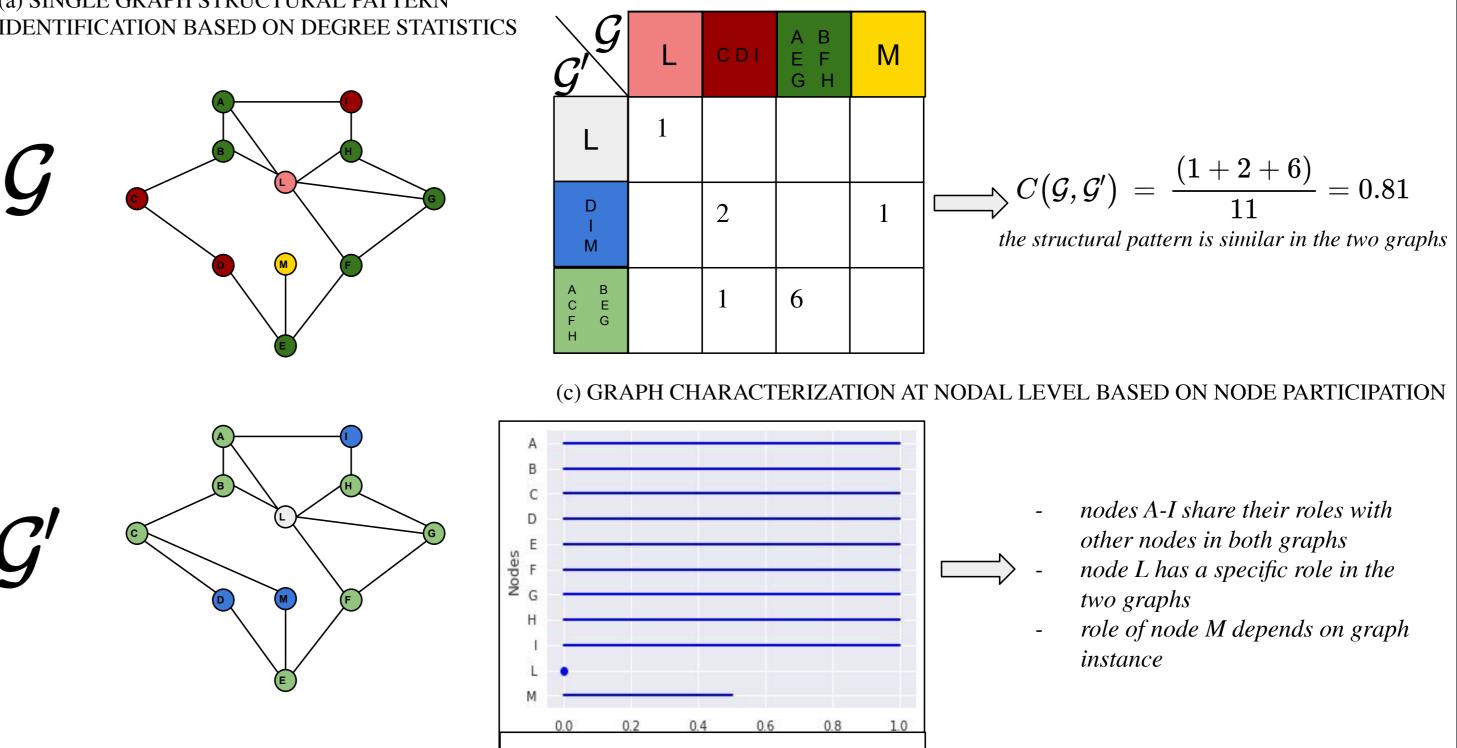
Motivations

o Human consciousness states can be differentiated by nodal organization in brain functional connectivity networks [1]

IDENTIFICATION BASED ON DEGREE STATISTICS



(b) COMPARISON AT GLOBAL LEVEL BASED ON GRAPH STRUCTURAL PATTERNS



- structural pattern definition
- ▶ extend node role discovery to graph collections [2, 3] graph collections comparison based on structural pattern

Nodal statistics-based structural pattern on single graph

Def. Nodal statistics based equivalence relation. We consider undirected unweigthed graph $\mathcal{G} = (\mathcal{V}, \mathcal{E})$ and refer to a nodal statistics $s: \mathcal{V} \to s(\mathcal{V})$ any function of the adjacency matrix. The equivalence relation \sim_s associated with a nodal statistics s, on the nodes set \mathcal{V} of a graph is:

 $v \sim_s u \iff s(u) = s(v).$

The equivalence relation associated with any collection of statistics $S = \{s_i\}_{i=1,..,n}$, is defined as:

$$a \sim_{\mathcal{S}} b \iff a \sim_{s_1} b, \ a \sim_{s_2} b, \ \ldots, \ a \sim_{s_n} b.$$

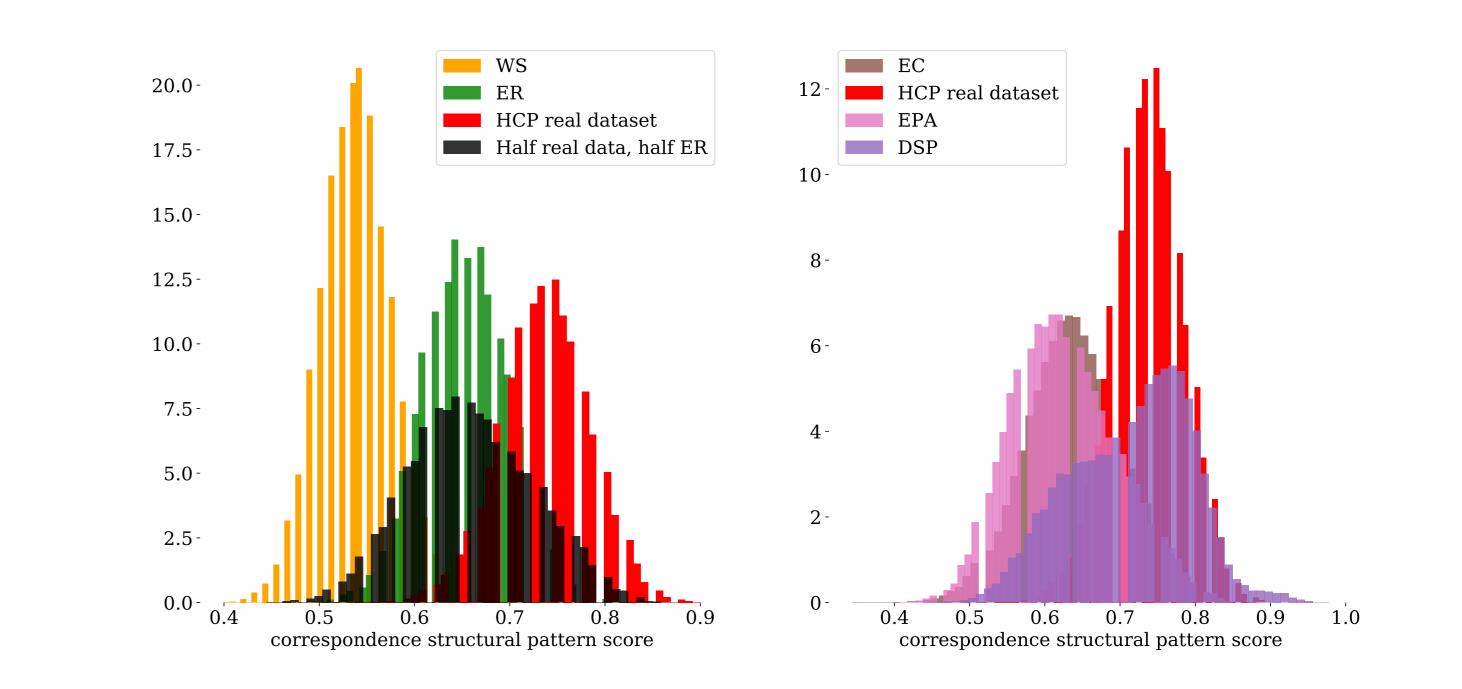
Def. Structural Pattern. Its induced partition P on \mathcal{V} ,

$$P_{\mathcal{S}} = rac{\mathcal{V}}{\sim_{\mathcal{S}}} = \{[a], \quad \forall a \in \mathcal{V}\}$$
 (2)

(1)

defines the **structural pattern** of \mathcal{G} associated with the statistics

Correspondence structural pattern score differentiates models



collection \mathcal{S} .

Def. Node role. The class of equivalence $[a] = \{b \in \mathcal{V} | a \sim_s b \iff s(a) = s(b)\}$ corresponds to **node role**.

Structural patterns for graph collections characterization

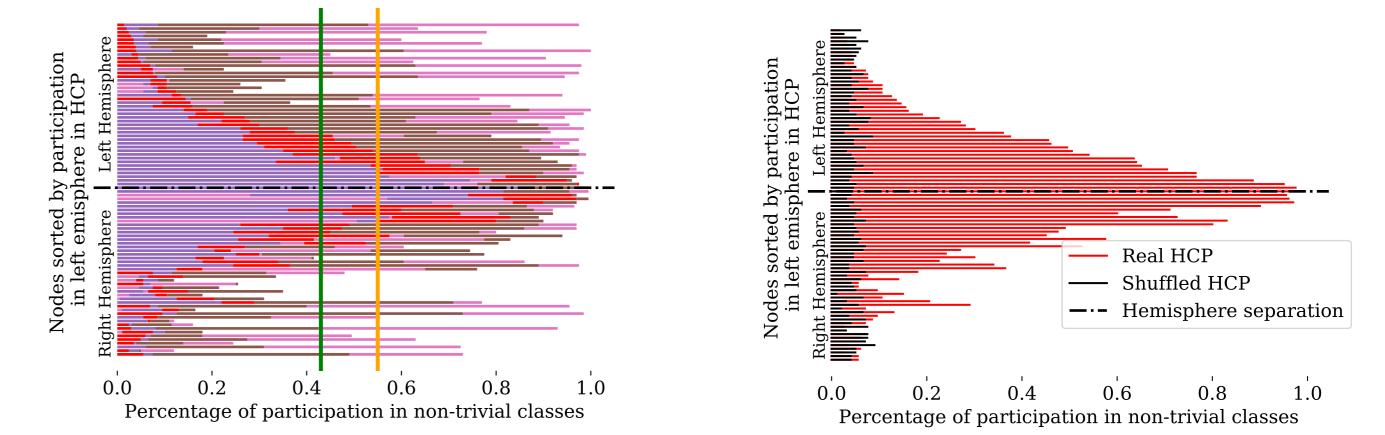
Def. Correspondence structural pattern score. Let $\mathcal{G}, \mathcal{G}'$ be two graphs having same vertices $\mathcal V$ and let $\mathcal S$ be a statistics collection whose associated partitions are $P_{\mathcal{S}}, P'_{\mathcal{S}}$ on $\mathcal{G}, \mathcal{G}'$ respectively. Given bijective mapping from P_S, P'_S to an initial segment of the natural numbers as enumerations, let $c(v_i), c'(v_i)$ be the enumeration of the classes of v_i , the correspondence structural pattern score between $\mathcal{G}, \mathcal{G}'$ is defined as:

$$C(\mathcal{G}, \mathcal{G}') = \max_{\pi \in \Pi} \frac{1}{|\mathcal{V}|} \sum_{i=1}^{|\mathcal{V}|} \mathcal{X}(\pi(c(v_i)) = c'(v_i))$$
 (3)

where Π is the set of all coupling between the elements in P_S and the elements in $P'_{\mathcal{S}}$ and \mathcal{X} is the indicator function. Def. Nodal-percentage of participation. Given a graphs collection $G = \{\mathcal{G}_k = (\mathcal{V}_k, \mathcal{E}_k) \text{ s.t. } \mathcal{V}_k = \mathcal{V}\}$, and a statistics collection \mathcal{S} we count the percentage of participation of each

(Left) WS: Watt-Strogatz model, ER: Erdős-Rényi model, HCP: Human Connectome Project real networks [4]; (Right) Brain inspired models.[5] DSP: Degree sequence preserving model; EPA: Economical preferential attachment model; EC: Economical clustering model.

Nodal-percentage of participation captures brain homotopy



The symmetry in real data and brain models reveals the expected hemisphere similarity in the participation of analogue regions.

node of \mathcal{V} in non-trivial classes:

$$\forall v \in \mathcal{V} \quad \mathsf{PP}_{G}^{\mathcal{S}}(v) = \mathsf{PP}_{G}(v) = \frac{|\{\mathcal{G}_{k} \in G \text{ s.t. } \#[v]_{\sim_{\mathcal{S}}}^{\mathcal{G}_{k}} \neq 1\}|}{|G|}$$

with $[v]_{\sim S}^{\mathcal{G}_k}$ the class of v in \mathcal{G}_k in the partition induced by \mathcal{S} .

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Conclusion

(4)

New mathematical method for exploration and characterization of network collections.

New way to **combine** many graph **nodal** statistics and to associated relative structural patterns.

Correspondence structural pattern score to **compare graphs at global level** based on their structural pattern similarity. Nodal-percentage of participation for nodal level characterization. Validation on generative models and real clinical data.

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