Main goal	Hypotheses test	Simulation	EEG Data	References
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First Young Researchers Workshop

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Main Goal

- The construction of a non parametric hypotheses test for samples of graphs;
- Application of this test to analyse brain functional networks constructed from electroencephalographic (EEG) data.

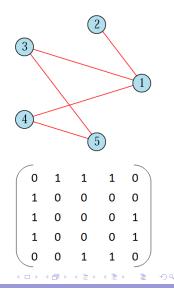
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Graph

- A simple graph is a pair (V, E), where V is a finite set of vertices and E ⊆ V × V is a set of edges;
- The graph can be represented by its adjacency matrix, where

$$g_{ij} = \left\{ egin{array}{ll} 1, & ext{if there is an edge} \\ & ext{between } i ext{ and } j \\ 0, & ext{otherwise} \end{array}
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Hypotheses test. Given two samples of graphs $\mathbf{g} = (g_1, \ldots, g_n)$ and $\mathbf{g}' = (g_1', \ldots, g_m')$, we want to test if they were originated from the same probability distribution, that is

$$\left(\begin{array}{c} \mathbf{H}_{0}: \pi = \pi' \\ \mathbf{H}_{A}: \pi \neq \pi' \end{array}\right)$$

where π is the distribution which originated **g** and π' is the distribution which originated **g**'.

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Definition

Given two samples of graphs $\mathbf{g} = (g_1, \ldots, g_n)$ and $\mathbf{g}' = (g'_1, \ldots, g'_m)$ we define the two-samples test statistic by

$$T(\mathbf{g},\mathbf{g}') = \max_{g \in \mathbb{G}(v)} |\overline{D}_{\mathbf{g}}(g) - \overline{D}_{\mathbf{g}'}(g)| \; ,$$

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where
$$\overline{D}_{\mathbf{g}}(g) = \frac{1}{n} \sum_{k=1}^{n} D(g, g_k)$$
 and $D(g, g_k) = \sum_{i < j} (g_{ij} - g_{ij}^k)^2$.

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The critical region of the test is

$$\mathbf{R} = \{t : t(\mathbf{g}, \mathbf{g}') > q_{(1-\alpha)}\},\$$

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where $q_{(1-\alpha)}$ is the $(1-\alpha)$ -quantile of the distribution of T under the null hypothesis (H₀).

Remark: $t \in R \Rightarrow$ we reject H_0

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It is important to remark that

- We need to know the set $\mathbb{G}(V)$ to compute T;
- The set $\mathbb{G}(V)$ has $2^{\binom{|V|}{2}}$ graphs;
- If |V| = 20, then $|\mathbb{G}(V)| = 2^{190}$ this is extremely LARGE !!!

How do we compute T ?

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Proposition

Given two samples of graphs $\mathbf{g} = (g_1, \ldots, g_n)$ and $\mathbf{g}' = (g_1', \ldots, g_m')$ we have that

$$\mathcal{T}(\mathbf{g},\mathbf{g}') = \sum_{i < j} |\overline{\mathbf{g}}_{ij} - \overline{\mathbf{g}}'_{ij}| \; ,$$

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where

$$\overline{\mathbf{g}}_{ij} = \frac{1}{n} \sum_{k=1}^{n} g_{ij}^{k}.$$

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► To compute the critical region of the statistical test we need to know the distribution of *T*.

What is the distribution of T?

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Proposition

Let two samples of graphs $\mathbf{g} = (g_1, \dots, g_n)$ and $\mathbf{g}' = (g'_1, \dots, g'_m)$. Under the null hypothesis H_0 we have

$$\mathcal{T}(\mathbf{g},\mathbf{g}') = \sum_{i < j} |\mathcal{T}_{ij}|$$

$$\sqrt{\left(\frac{nm}{n+m}\right)(T_{ij})_{ij}} \xrightarrow[m \to \infty]{D} N(0,\Sigma)$$

where
$$\Sigma_{ij,kl} = \pi G_{ij,kl} - (\pi G_{ij})(\pi G_{kl})$$
 and $\pi G_{ij,kl} = \sum_{g \in \mathbb{G}(v)} g_{ij}g_{kl}\pi(g)$.

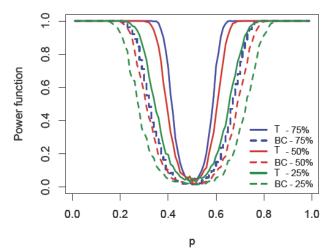
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Simulation

We compared the power function of our test with the power function of the simultaneous testing procedure with Bonferroni correction (BC). The null model is the Erdös-Rényi model with parameter $p_0 =$ 0.5 and the alternative hypothesis is (modified) Erdös-Rényi model with v = 10 nodes and q% of edges with parameter p and the remaining edges with parameter $p_0 = 0.5$. The sample size was n=20.

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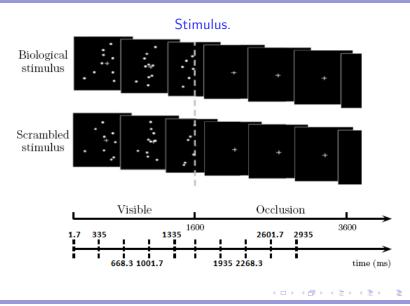
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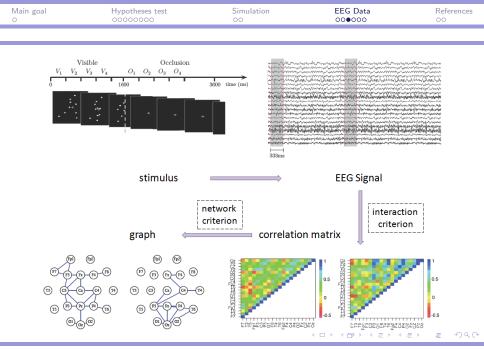
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Discrimination of EEG brain networks.

We want to compare graphs built from EEG data collected during the observation of videos depicting human locomotion.

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Visible x Occlusion

- Biological movement

 - visible: 132 graphs for each window occlusion: 132 graphs for each window
- Non-Biological Movement

 - visible: 132 graphs for each window occlusion: 137 graphs for each window
- p-value of the test

Visible vs Occlusion	Windows				
	V_1 vs O_1	V_2 vs O_2	V_3 vs O_3	V ₄ vs O ₄	
Biological	0.0019	0.4294	0.1984	0.0278	
Non-biological	0.0016	0.8278	0.1249	0.6673	

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Our paper: A test of hypotheses for random graph distributions built from EEG data.

http://arxiv.org/abs/1504.06478

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Acknowledgments







modelagem estocástica e complexidade



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References

E. Bullmore e O. Sporns.

Complex brain networks: graph theoretical analysis of structural and functional systems.

Nat Rev Neurosci, 10:186–198.

J. R. Busch, P. A. Ferrari, A. G. Flesia, R. Fraiman, S. P. Grynberg e F. G. Leonardi.

Testing statistical hypothesis on random trees and applications to the protein classification problem.

The Annals of Applied Statistics, 3(2):542–563.

D. Fraiman, G. Saunier, E. F. Martins, e C. D. Vargas. Biological motion coding in the brain: analysis of visually-driven eeg funcional networks.

Plos One. No prelo 2014.

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References



M. E. J. Newman.

Networks: An Introduction.

Oxford University Press.

 G. Saunier, E. F. Martins, E. C. Dias, J. M. de Oliveira, Thierry Pozzo e Claudia D. Vargas.
 Electrophysiological correlates of biological motion permanence in humans.

Behavioural Brain Research, 236:166–174.