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HELMHOLTZ ZENTRUM DRESDEN ROSSENDORF



TECHNISCHE UNIVERSITÄT CHEMNITZ

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Theoretical research and experiments suggest that the brain operates at or near a **critical state** between sustained activity and an inactive phase, exhibiting optimal computational properties (see: *Beggs & Plenz J. Neurosci. 2003; Chialvo Nat. Phys. 2010; Haimovici et al. PRL 2013*)



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Neurons exhibit oscillatory behavior



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Quasistatic inhomogneity causes dynamical criticality in Griffiths phases



→ Edge of Synchronization and Griffiths phase in brain models ?





















Electrode LFP experiments Since Beggs and Plenz 2003 For humans and animals



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Diverging fluctuations → High sensitivity to stimuli



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Diverging correlation functions → Optimal transmission and storage of information



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#### **Pros:**

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Maximal information processing and computational performance

**Cons:** Tuning to critical point is needed Danger of super-critical (epileptic) behavior Self-organization to criticality (SOC) ?

## **Explanations for tuning to criticality**



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### SOC $\leftrightarrow$ GP do not exclude each other

## **Explanations for tuning to criticality**



# SOC ↔ GP do not exclude each other For SOC we need a responsible feedback mechanism, GP can occur spontaneously in heterogeneous systems







Order parameter : density of active sites (  $\,\rho\,$  )



For m > n: first order phase transition see my book :
#### Discrete, stochastic threshold models on networks



On low dimensional regular, Euclidean lattice: **DP** critical point :  $\lambda_c > 0$  between inactive and active phases (GÓ: PRE 67 (2003) 056114.)



#### Discrete, stochastic threshold models on networks



On low dimensional regular, Euclidean lattice: **DP** critical point :  $\lambda_c > 0$  between inactive and active phases (GÓ: PRE 67 (2003) 056114.)



• Fixed (quenched) disorder/impurity changes the local birth rate  $\Rightarrow \lambda_{c} > \lambda_{c}^{0}$ 









contribute to the density:  $\rho(t) \sim \int dL_R L_R w(L_R) \exp[-t/\tau(L_R)]$ 



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- At  $\lambda_c^0$  the characteristic time scales as:  $\tau (L_R) \sim L_R^Z \Rightarrow \ln \rho(t) \sim t^{d/(d+Z)}$

saddle point analysis: stretched exponential



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saddle point analysis: stretched exponential **Griffiths Phase** 

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• For  $\lambda_c^0 < \lambda < \lambda_c$ :

 ⇒ saddle point analysis: stretched exponential
 L<sub>R</sub>): Griffiths Phase
 continuously changing exponents

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• For  $\lambda_c^0 < \lambda < \lambda_c$ :

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continuously changing exponents

• At  $\lambda_c$ : b may diverge  $\rightarrow \rho(t) \sim \ln(t)^{-\alpha}$  Infinite randomness fixed point scaling

 $\tau$  (  $L_{R}$ ) ~ exp(b  $L_{R}$ ):



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• For  $\lambda_c^0 < \lambda < \lambda_c$ :  $\tau (L_R) \sim \exp(b L_R)$ : Griffiths Phase continuously changing exponents

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- GP: Dynamical (scaling) criticality + susceptibility diverges



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• For  $\lambda_c^0 < \lambda < \lambda_c^c$ :  $\tau (L_R) \sim \exp(b L_R)$ : Griffiths Phase  $\rho(t) \sim t^{-c/b}$  continuously changing exponents

• At  $\lambda_c$ : b may diverge  $\rightarrow \rho(t) \sim \ln(t)^{-\alpha}$  Infinite randomness fixed point scaling

• **GP: Dynamical (scaling) criticality + susceptibility diverges** GP can occur by pure topological disorder in finite dimensional systems

 Mean-field for threshold models with activation :

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λ

0

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providing a threshold within a system of size  $\Lambda_{c} = \frac{8}{5(N-1)(N-2)}$  active phase active phase

and an order parameter for 
$$\Lambda \to \Lambda$$

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 $(\rho - \rho_c) \propto (\lambda - \lambda_c)^{1/2}$ 

#### In heterogenous HMN models HPT + GP + Multistable states !



FIG. 9. Evolution of  $\rho(t)$  for different  $\lambda$  in case of starting from ully active state in the excitatory model with levels: l = 5, 6. From bottom to top symbols:  $\lambda = 0.30, 0.32, 0.321$  (l = 6), 0.322, 0.322 (l = 6), 0.325, 0.33, 0.34 (l = 6), 0.35, 0.4, 0.5, 0.6, 0.7.

#### G.O. B.S: Phys. Rev. Res. 3 (2021) 0131106

The largest precisely explored brain structural networks contains N = 302 neurons (C. Elegans), fruit-fly hemibrain: N = 127,978



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Unknown faithfulness, intensive research to automate image processing



Diffusion and structural MRI images with  $1 \text{ mm}^3$  voxel resolution :  $10^5 - 10^6$  nodes



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fiber  $\rightarrow$  edge (~ 10<sup>10</sup>)

+ noise reduction  $\rightarrow$  graph

undirected, weighted









# SCIENTIFIC **Reports**

### OPEN The topology of large Open Connectome networks for the human brain

Michael T. Gastner<sup>1,2</sup> & Géza Ódor<sup>2</sup>

The structural human connectome (i.e. the network of fiber connections in the brain) can be analyzed at ever finer spatial resolution thanks to advances in neuroimaging. Here we analyze several large data sets for the human brain network made available by the Open Connectome Project. We apply statistical model selection to characterize the degree distributions of graphs containing up to  $\simeq 10^6$  nodes and  $\simeq 10^8$ edges. A three-parameter generalized Weibull (also known as a stretched exponential) distribution is a good fit to most of the observed degree distributions. For almost all networks, simple power laws cannot fit the data, but in some cases there is statistical support for power laws with an exponential cutoff. We also calculate the topological (graph) dimension *D* and the small-world coefficient  $\sigma$  of these networks. While  $\sigma$  suggests a small-world topology, we found that D < 4 showing that long-distance connections provide only a small correction to the topology of the embedding three-dimensional space.

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#### Small world, still finite dimensional, non-scale free, universal modular graphs

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SCIENTIFIC REPORTS 6:27249 DOI: 10.1038/srep27249

KKI-18 graph: *836733 nodes*, 4 *x 10*<sup>7</sup> weighted, undirected edges



FIG. 1. Link weight probability density function of the KKI-18 OCP graph. Dashed line: a PL fit for intermediate  $w_{ij}$ 's. Inset: survival probability in the K = 6 threshold model near the transition point for  $\lambda = 0.003$ ,  $\nu = 0.3, 0.4, 0.45, 0.5, 0.55, 0.6, 0.7$  (top to bottom curves).

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Cluster spreading simulations from randomly selected active nodes



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incoming weights normalized by the sum : to model homogeneous sensitivity of nodes



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Inhibition: randomly selected weights are flipped to negative (quenched)

$$w_{i,j}' = -w_{i,j}'.$$



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FIG. 4. The same as Fig. 2 in the case of the undirected graph. Inset: Local slopes of the curves.



FIG. 2. Avalanche survival distribution of the relative threshold model with K = 0.25, for v = 0.95 and  $\lambda = 0.8, 0.81$ , 0.82, 0.83, 0.835, 0.84, 0.845, 0.85, 0.86, 0.87, 0.9, 0.95, 1 (bottom to top curves). Inset: Local slopes of the same from  $\lambda = 0.835$  to  $\lambda = 1$  (top to bottom curves). Griffiths effect manifests by slopes reaching a constant value as  $1/t \rightarrow 0$ .



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# (a) 20% of links are turned directional, randomly(b) Unidirectional



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# (a) 20% of links are turned directional, randomly(b) Unidirectional

No qualitative difference, but proves robustness (for more long links)



FIG. 3. Avalanche size distribution of the relative threshold model with K = 0.25, for v = 1 and  $\lambda = 1,0.9,0.8$ . Dashed line: PL fit to the  $\lambda = 0.8$  case. Inset: Avalanche shape collapse for T = 25,63,218,404 at  $\lambda = 0.86$  and v = 0.95.











## Scaling near experimental values in the Griffiths Phase ( $\tau \sim 1.5$ )



### **Robustness of Griffiths effects in homeostatic connectome threshold models**

G. Ó. Phys. Rev. E 98 (2018) 042126

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Addition of a third (refractive) state does not destroy GP



FIG. 3: Avalanche size distribution in the relative threshold model with refractory states, for K = 0.2,  $\nu = 1$  and  $\lambda = 0.91, 0.965, 0.985, 0.995$  (bottom to top symbols). Lines: PL fits for  $10^2 < s < 10^5$ , for these curves as shown by the legends.

### **Robustness of Griffiths effects in homeostatic connectome threshold models**

G. Ó. Phys. Rev. E 98 (2018) 042126

Addition of a third (refractive) state does not destroy GP

Time dependent threshold model : GP shrinks, but survives for weak variations





FIG. 3: Avalanche size distribution in the relative threshold model with refractory states, for K = 0.2,  $\nu = 1$  and  $\lambda = 0.91, 0.965, 0.985, 0.995$  (bottom to top symbols). Lines: PL fits for  $10^2 < s < 10^5$ , for these curves as shown by the legends.

FIG. 10: Avalanche size distribution of the time dependent relative threshold model with 30% inhibitory links at K = 0.1,  $\Delta K = 0.01$ ,  $\nu = 0.95$  and  $\lambda = 0.473, 0.478, 0.480, 0.493$  (bottom to top symbols) Dashed lines: PL fits for the tails of the  $\lambda = 0.473$  and  $\lambda = 0.493$  curves (bottom to top).





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Order parameter : average phase:



Non-zero, above critical coupling strength K > K, tends to zero for  $K \le K_c$  as  $R \propto (1/N)^{1/2}$  or exhibits an initial growth:  $R(t, N) = N^{-1/2} t^{\eta} f_{\uparrow}(t/N^{\tilde{z}})$  for incoherent initial state



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The dynamical behavior suffers very strong corrections to scaling and chaoticity

We use this "toy" synchronization model assuming universality,
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Numerical ODE solution of large set of equations via adaptive Bulrisch-Stoer stepper, implemented on HPC GPU-s

# **Kuramoto solution for the** *K***I-18 graph with** *N*= *836 733* nodes and *41 523 931* weighted edges

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The synchronization transition point determined by growth as before



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 $K_c = 1.7$  and growth exponent:  $\eta = 0.6(1)$ 





Measure characteristic times  $t_x$  of first

dip below:  $R_c = (1/N)^{1/2}$ 



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Histogramming of  $t_x$  at the critical point



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Histogramming of  $t_x$  at the critical point



10 Measure characteristic times  $t_x$  of first R(t) 10 dip below:  $R_c = (1/N)^{1/2}$  $10^{-1}$ average over: 10.000 independent  $\omega_i$ realizations  $10^{-2}$ p(t<sub>x</sub>) Histogramming of  $t_x$  at the critical point 10<sup>-3</sup> Critical exponent:  $\tau_t = 1.2$  (D)  $10^{-4}$ obtained by fitting for the PL tails  $10^{-5}$  $10^{1}$  $10^{2}$ 10 t,

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# Inhibitory (negative) links compared to experiments

Inhibitions: 5% of links:  $w_{ij} \rightarrow -w_{ij}$  randomly

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 $K_c = 1.7(1)$  and  $\eta = 0.6(1)$  remains the same. Sub-critically:

**Duration scaling exponent within experimental range:**  $1.5 < \tau_t < 2.4$ *J.M. Palva et al PNAS 110 (2013) 3585* 

$$\dot{\theta}_i(t) = \omega_{i,0} + \frac{K}{k_i} \sum_j W_{ij} \sin[\theta_i(t) - \theta_j(t)]$$

Brain experiments:  $\omega_i > 0$ 

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#### The effect of additive stochastic noise

Gaussian distributed annealed noise is added:

$$\dot{\theta}_i(t) = \omega_{i,0} + \frac{K}{k_i} \sum_j W_{ij} \sin[\theta_j(t) - \theta_i(t)] + s\xi(i)$$

Negligible effect:



## **Comparison with the fruit-fly connecome results**

 $A_{ij}$ 




$A_{ii}$ 



Fruit-fly connectome is the largest exactly known neural network: N = 21.615, L = 3.410.247





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K = 1.60(1) (inflexion curve) Characterized by mean-field growth Exponent  $\eta = 0.7(1)$ 



Fluctuations of R show

extended transition for KKI-18







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extended transition for KKI-18 For FF ~ ER like distro With random inhibitors: wider range The same is true for fluctuations of  $\Omega$ HMN structure of KKI-18 is responsible for the extended critical region and Griffiths Phase of humans As compared to the fly connectome

Fluctuations of R show

Shinomoto-Kuramoto oscillator model synchronization transition:

 $\theta_i$ : angle, *K*: global coupling

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Quenched heterogeneity in self-frequencies and network topology



#### **Force induced synchronization**





FIG. 1: Order parameter dependence on F for the fruit-fly connectome for the noisy (black bullet) and the noiseless (red boxes) cases at K = 1.3. The blue diamonds show the steady-state  $\Omega$  values with noise. Lower inset: Variances of R and  $\Omega$  for the noisy case. Upper inset: Time dependence of the noisy R(t), for F = 0, 0.02, 0.03, 0.04, 0.07, 0.1, 0.2, 0.3, 0.4 (bottom to top curves).

FIG. 3: Fluctuations of R and  $\Omega$  as the function of F for the KKI-18, for the noisy and the noiseless cases at K = 1. Inset: Order parameter R for the noisy and noiseless cases as well as  $\Omega$ , denoted by the same symbols as in the main figure.

## Characteristic time exponent $\tau_t$ results

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The  $p(t_{\star})$  distros exhibit power-law near the synchronization transition point  $F_{\star} \sim 0.1$  for K=1.3characterized by the *exponent:* 2

### Characteristic time exponent $\tau_{t}$ results



FIG. 4: Avalanche duration distributions on the fruit-fly connectome for different forces, shown by the legends and at K = 1.3,  $\epsilon = 0.01$ . Dashed lines are PL fits for  $\Delta t > 100$ . The inset shows the steady state  $\sigma(\Omega)$  as the function of K, for excitation values F = 0.001, 0.0667, 0.1, 0.2, 0.3 (top to bottom).



The  $p(t_{\star})$  distros exhibit power-law near the synchronization transition point  $F_{\star} \sim 0.1$  for K=1.3characterized by the *exponent: 2* 

Similarly as in case of the KKI-18:





FIG. 8: Hurst and beta exponents of all fruit-fly connectome communities. In the forceless case at the critical Hopf transition coupling, the H exponent is the largest for every community. With forces these values drop for each community. This shows a resemblance with the rest and non-rest studies of different brain areas in [63], showing  $\langle H \rangle \approx 1.0$  at resting state and  $\langle H \rangle \approx 0.7$  at task driven states.



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Community dependent synch. Quasi-criticality, like in fMRI experiments: *Ochab et al*, *Sci. Rep. 12, 17866 (2022)*.

### **FMRI experiments**



**Task** ↔ **rest state operation** 

Heterogeneity effects on the scaling can be investigated on **large connectomes** and random small-world graphs for comparison

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- Invariance with respect to frequency distributions
- Insensitivity for the additive Gaussian noise
- Periodic force induces synchronization and higher fluctuations
- Force enhances long-range correlations, i.e. in the task phase operation of brain with respect to resting state

Thank you for your attention ! Recent publications:

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