

Model-based comparison of connectomes: applications in a whole insect brain

Benjamin D. Pedigo

(he/him) -  bpedigo@jhu.edu

[NeuroData lab](#)

Johns Hopkins University - Biomedical Engineering

Acknowledgements



Mike
Powell



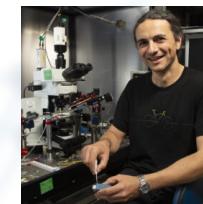
Eric
Bridgeford



Michael
Winding



Marta
Zlatic



Albert
Cardona



Carey
Priebe



Joshua
Vogelstein

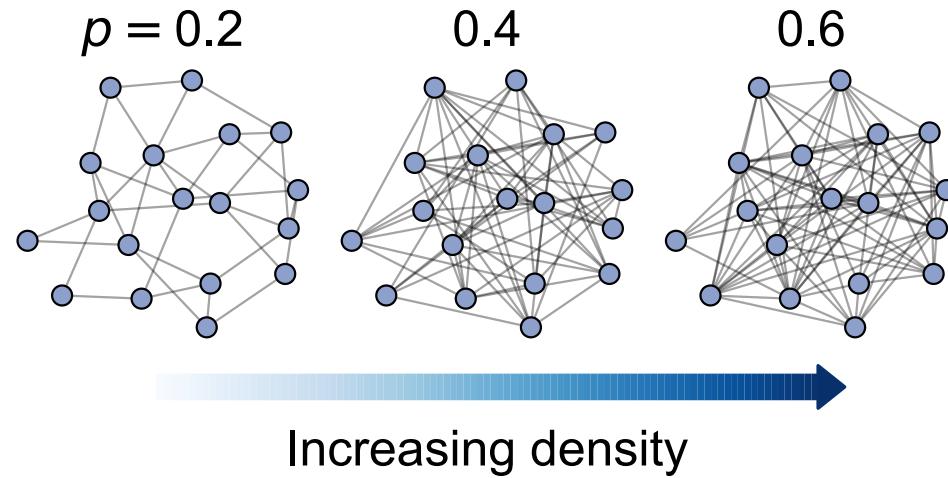
Larval *Drosophila* brain connectome



Are the **left** and **right** sides of this connectome
different?

We reject even the simplest notion of symmetry

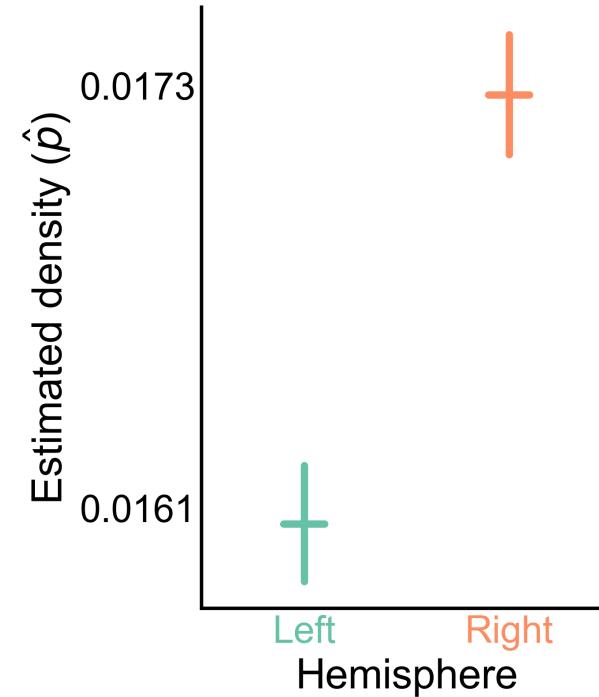
- Fit Erdos-Renyi models to the left and the right brain networks



- Compare densities:

$$H_0 : p^{(L)} = p^{(R)}$$

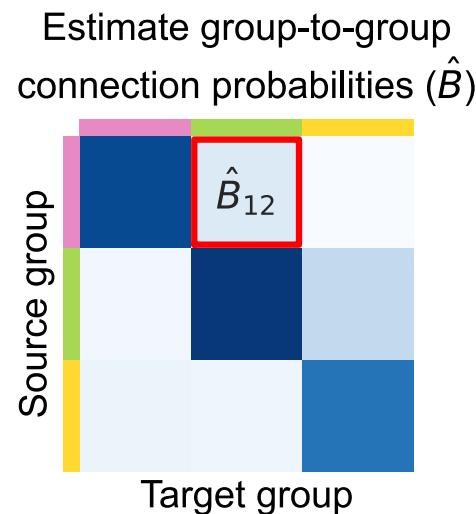
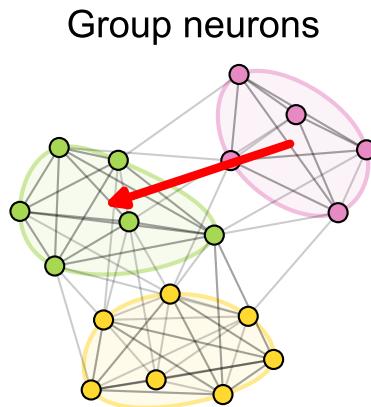
$$H_A : p^{(L)} \neq p^{(R)}$$



p-value: $< 10^{-23}$

Localizing differences to cell type connections

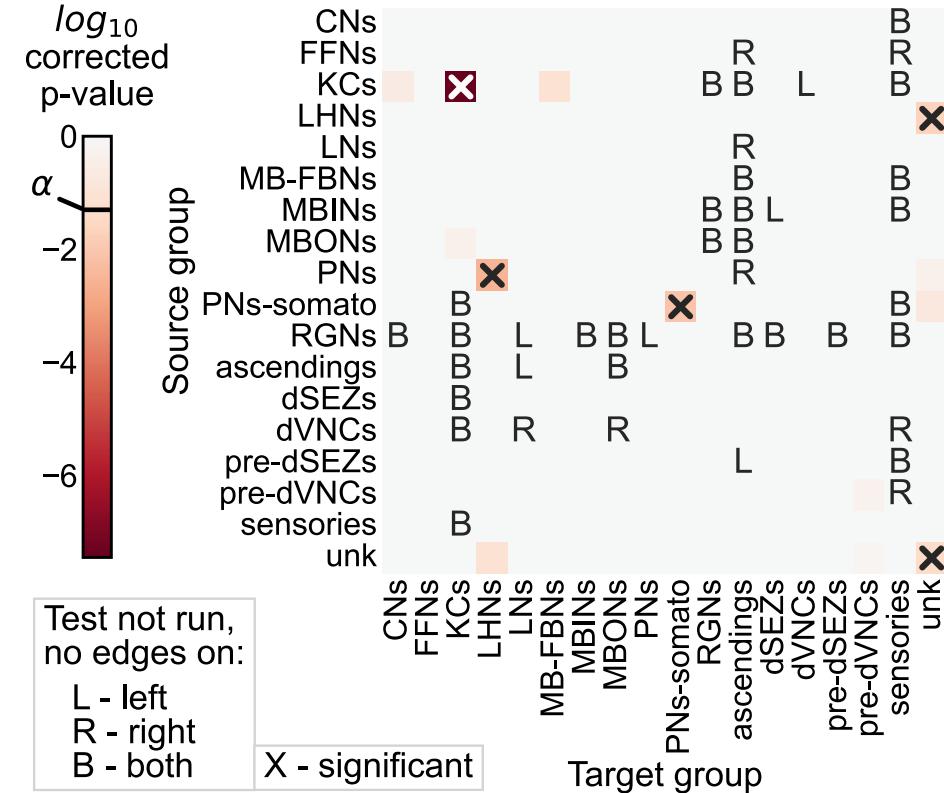
- Fit block models to both hemispheres



- Compare connection probabilities:

$$H_0 : B^{(L)} = B^{(R)}$$

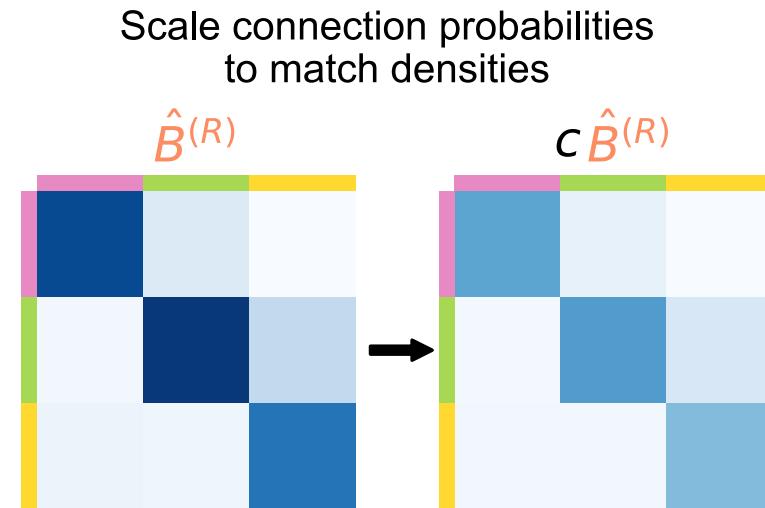
$$H_A : B^{(L)} \neq B^{(R)}$$



p-value: $<10^{-7}$

Modified definitions of symmetry which ARE exhibited

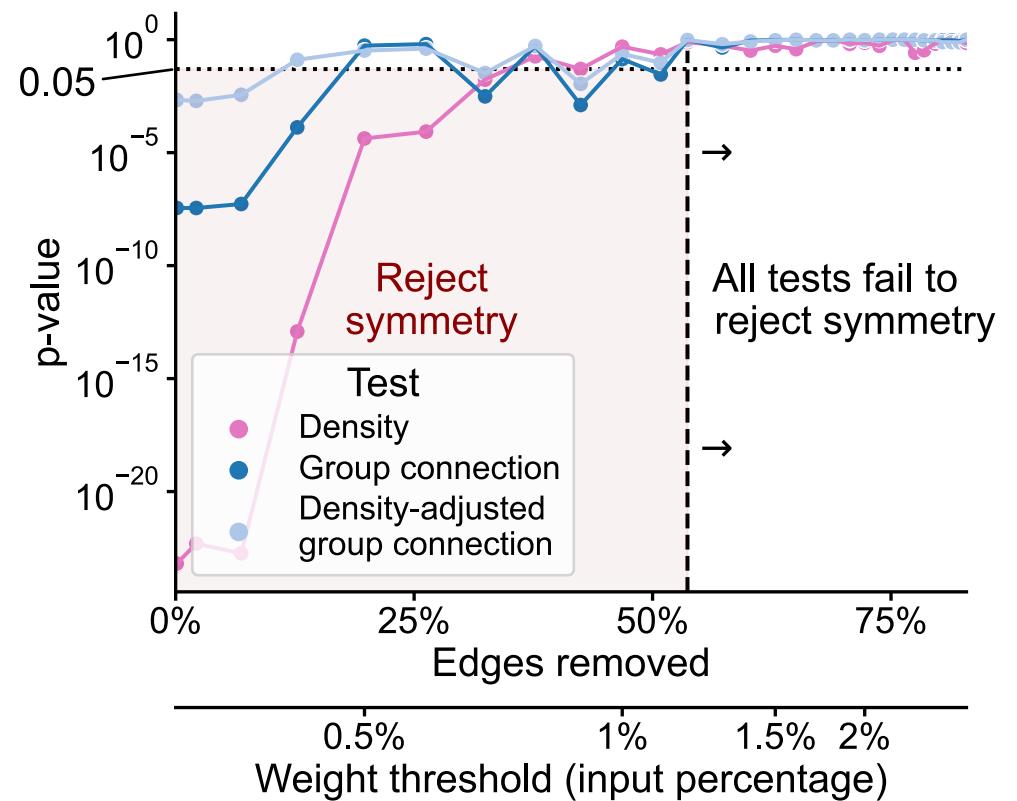
Rescale connection probabilities AND
remove Kenyon cells



$$\begin{aligned} H_0: & B^{(L)} = cB^{(R)} \\ H_A: & B^{(L)} \neq cB^{(R)} \end{aligned}$$

p-value: ~0.51

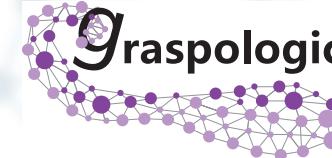
Threshold by edge weight (input
proportion to postsynaptic neuron)



Conclusions

- Testing hypotheses in connectomics requires techniques for networks
 - We presented procedures for comparing connectomes
- Used to evaluate bilateral symmetry, finding how this brain is/is not bilaterally symmetric
- Poised to apply these tools to answer...
 - {Your question here}
 - Get in touch:
 -  bpedigo@jhu.edu
 -  jovo@jhu.edu

More info

-  [Graspologic](#) downloads 121k
- This work:  [jupyter](#) [book](#)
github.com/neurodata/bilateral-connectome
- Chung et al. *Statistical connectomics* (2021)
- Data: Winding, Pedigo et al. *In preparation* (2022)