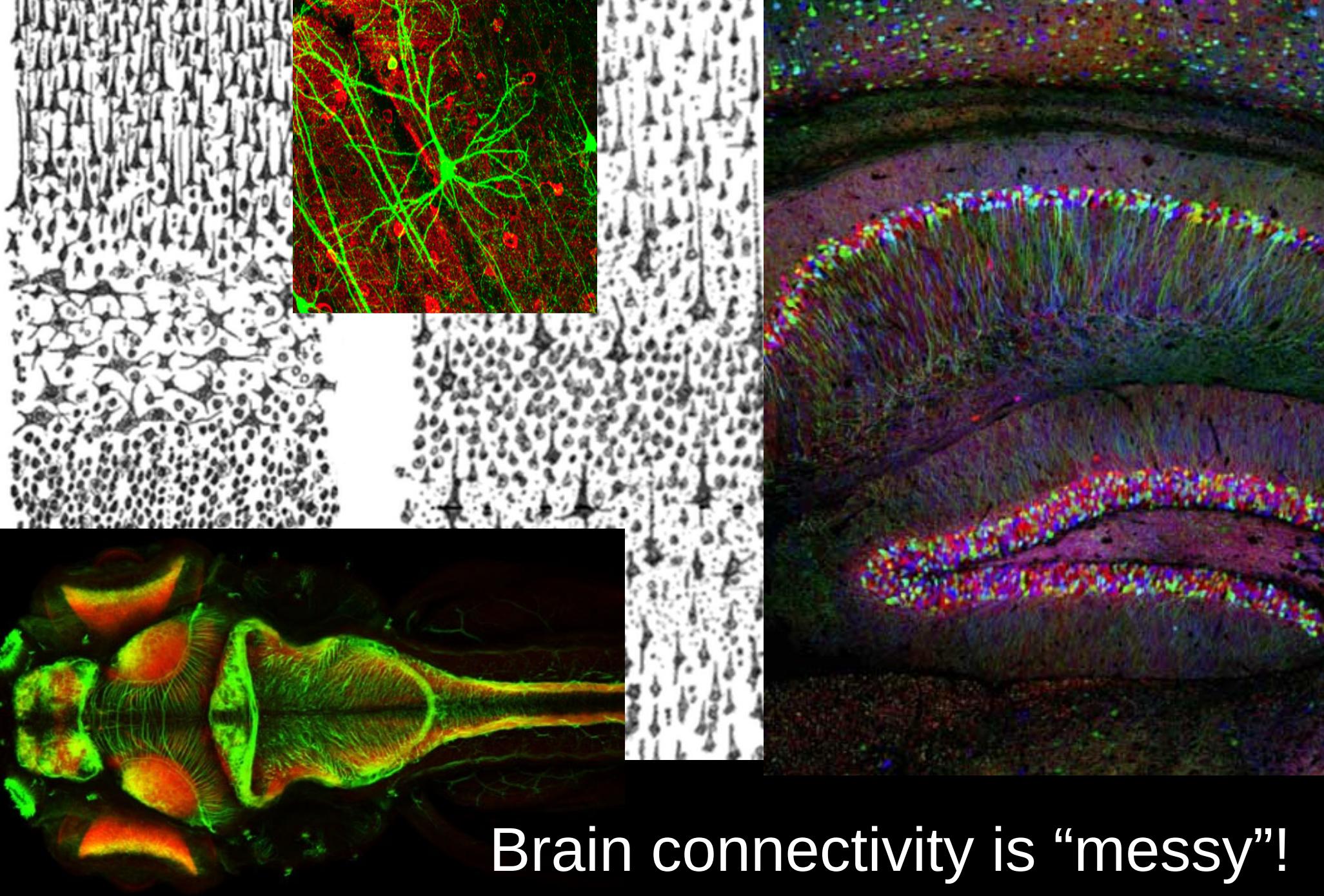


This Brain Is a Mess: Inference, Random Graphs, and Biophysics to Disentangle Neuronal Networks

Kameron Decker Harris
November 30, 2017

Acknowledgements

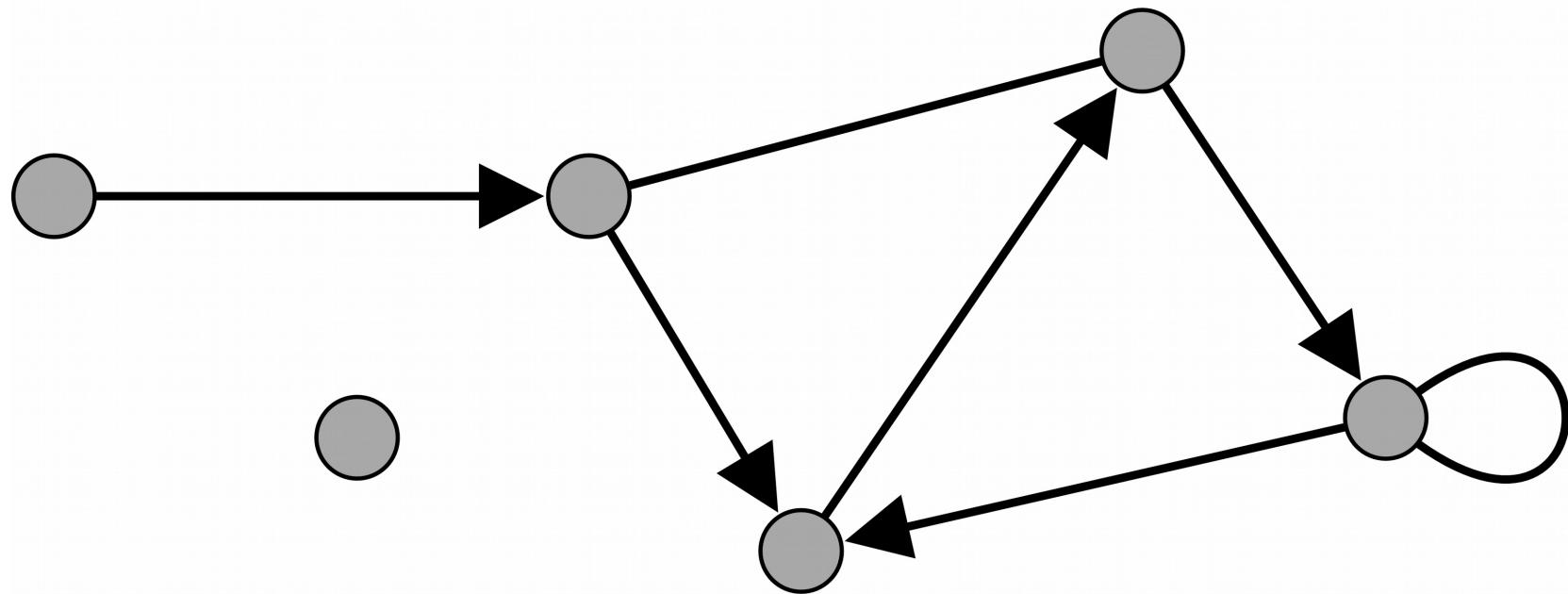
- **Eric Shea-Brown & team:**
 - Josh Mendoza
 - Iris Shi
 - Doris Voina
 - Matt Farrell
 - Ali Weber
 - Kathleen Champion
 - Tim Oleskiw
 - Alex Cayco-Gajic
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 - Hannah Choi
 - Merav Stern
 - Gabrielle Gutierrez
 - Joel Zylberberg
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 - Fred Garcia
 - Aguan Wei
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 - Joe Knox
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 - David Feng
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- **UW AMATH:**
 - Keshanie Dissanayake
 - Lauren Lederer
 - Derek Franz
 - Alan Perry
 - Tony Garcia
 - Everyone else!



Brain connectivity is “messy”!

Credits: Lee et al. (2006) cortical pyramidal cell; Santiago Ramon y Cajal; UCL Zebrafish Group; Tamily Weissman, Jean Livet, Ryan Draft (mouse hippocampus)

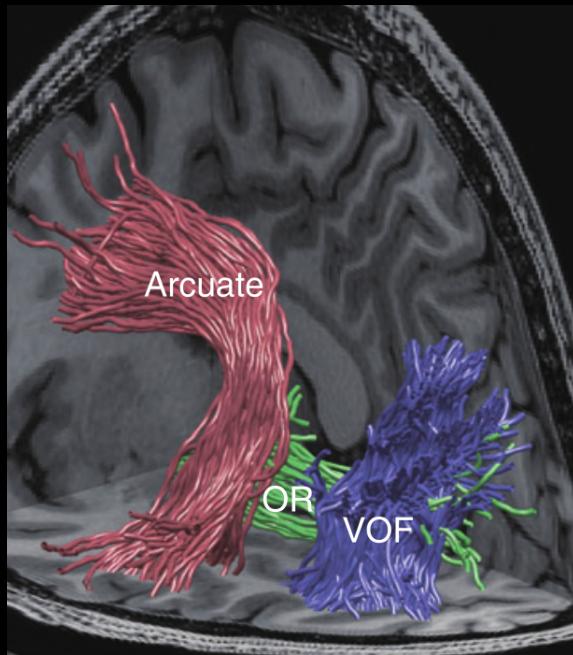
Neuron wiring forms a **network** or **graph**



Structural connectivity data

Structural connectivity data

Diffusion MRI
Macroscopic

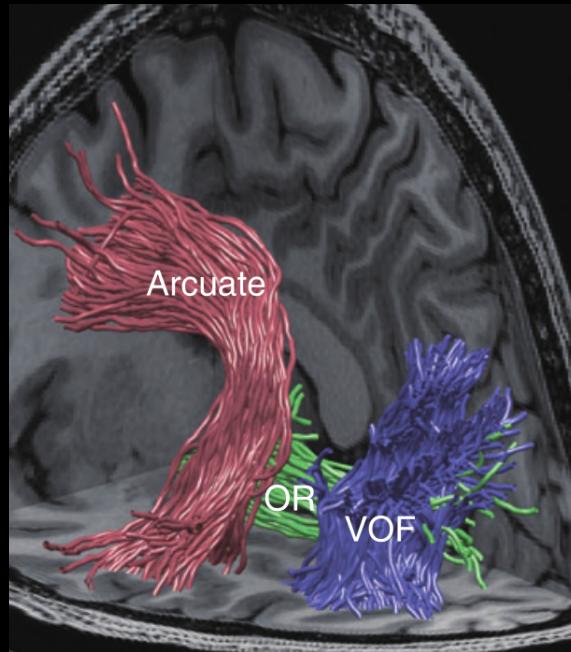


Takemura et al. (2016)

1-3 mm
Undirected
Non-destructive

Structural connectivity data

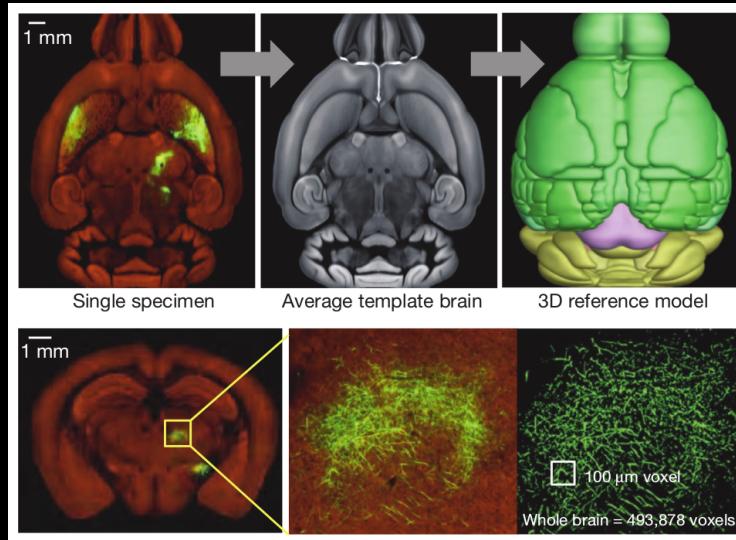
Diffusion MRI
Macroscopic



Takemura et al. (2016)

1-3 mm
Undirected
Non-destructive

Tracing & tomography
Mesoscopic



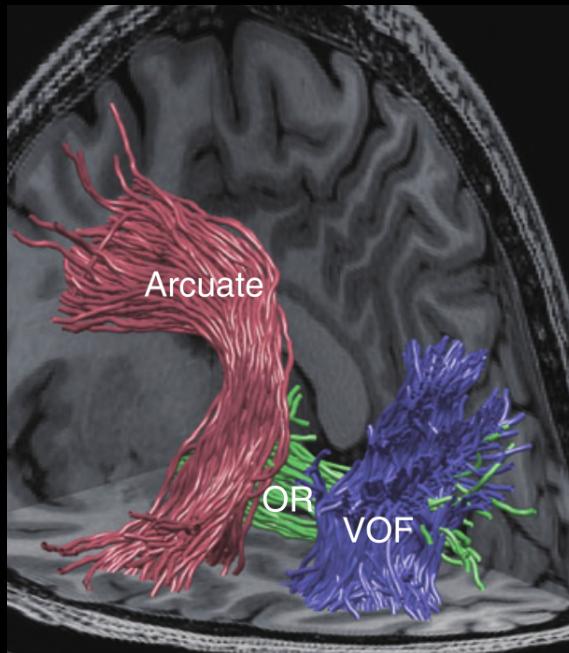
10-100 μm
Directional
100's of neurons

Increasing resolution

All require algorithms!

Structural connectivity data

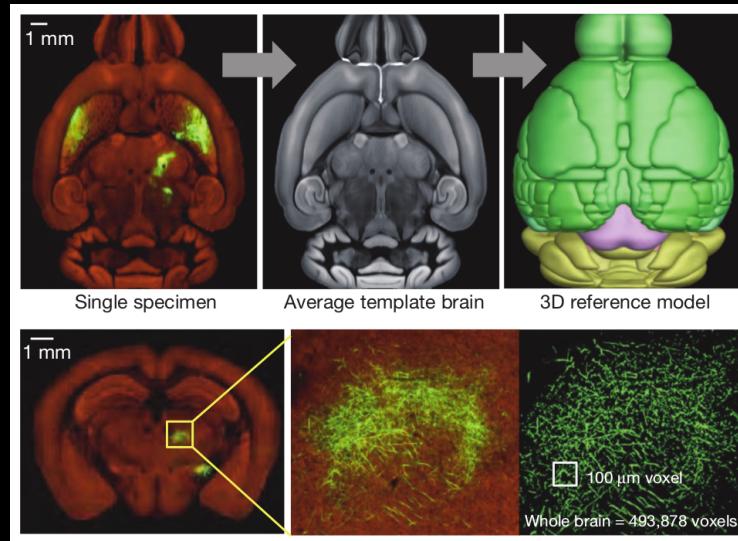
Diffusion MRI
Macroscopic



Takemura et al. (2016)

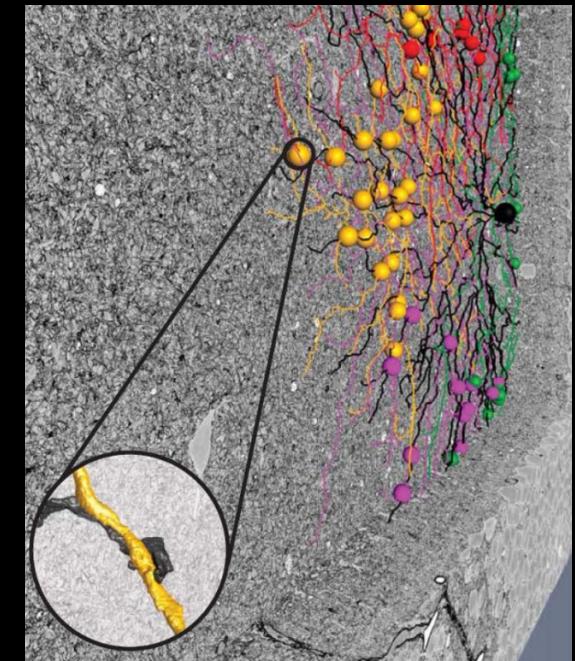
1-3 mm
Undirected
Non-destructive

Tracing & tomography
Mesoscopic



10-100 μm
Directional
100's of neurons

Electron Microscopy
Microscopic



10 nm
Neurons & synapses

Increasing resolution

All require algorithms!

Connectivity questions

Connectivity questions

1) How do we infer mesoscopic connections?

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- 1) How do we infer mesoscopic connections?
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Connectivity questions

- 1) How do we infer mesoscopic connections?
- 2) What are good microscopic models?
- 3) How does connectivity affect brain dynamics?

Machine learning for neuronal network data

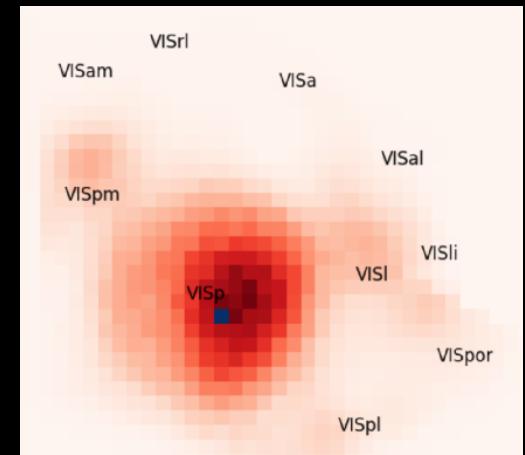
Published as:

Harris, Mihalas, Shea-Brown.

“Nonnegative spline regression of incomplete tracing data reveals high resolution neural connectivity.”

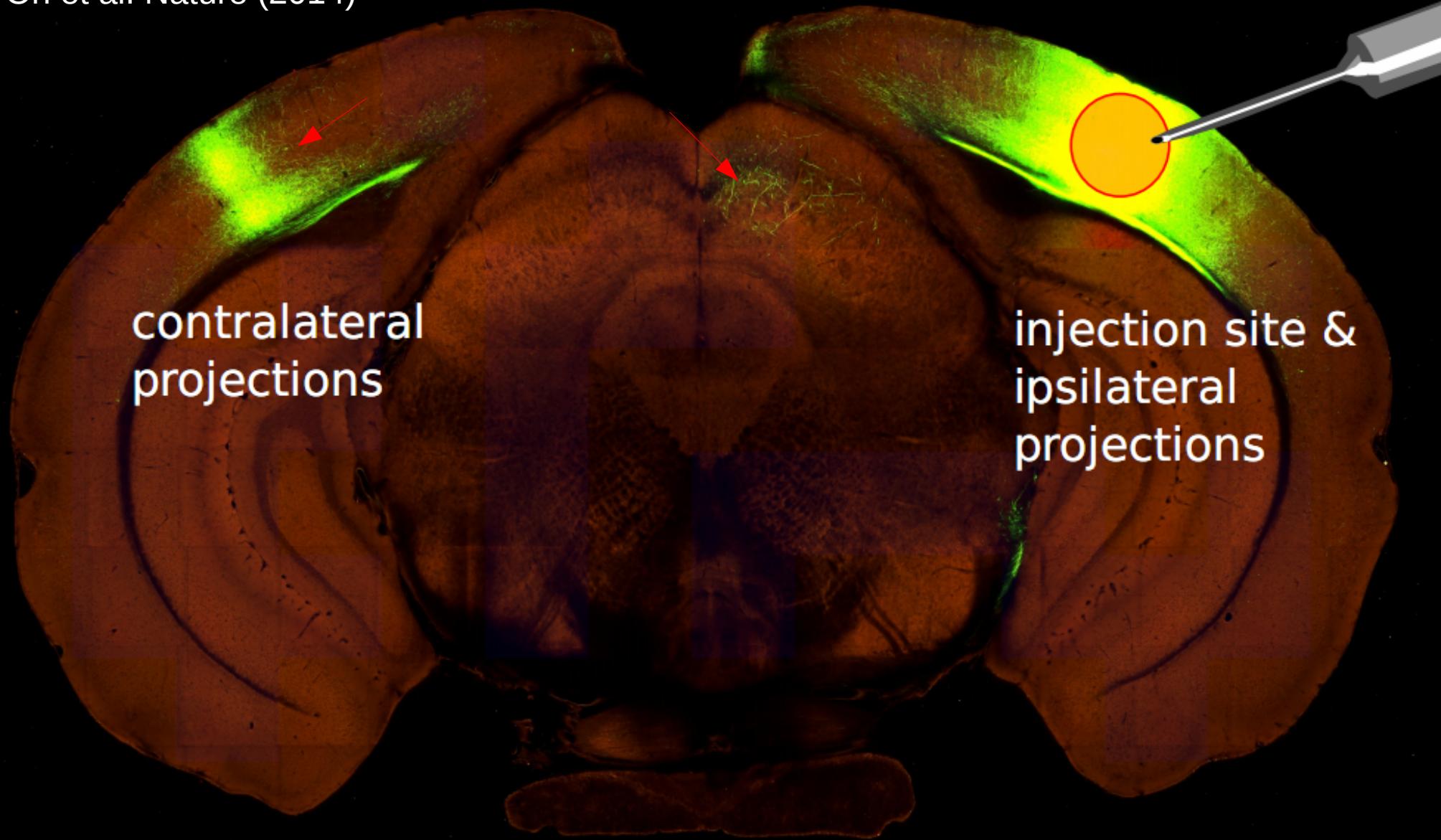
In proceedings of NIPS. 2016.

Ongoing work with Joe Knox and others at Allen Institute

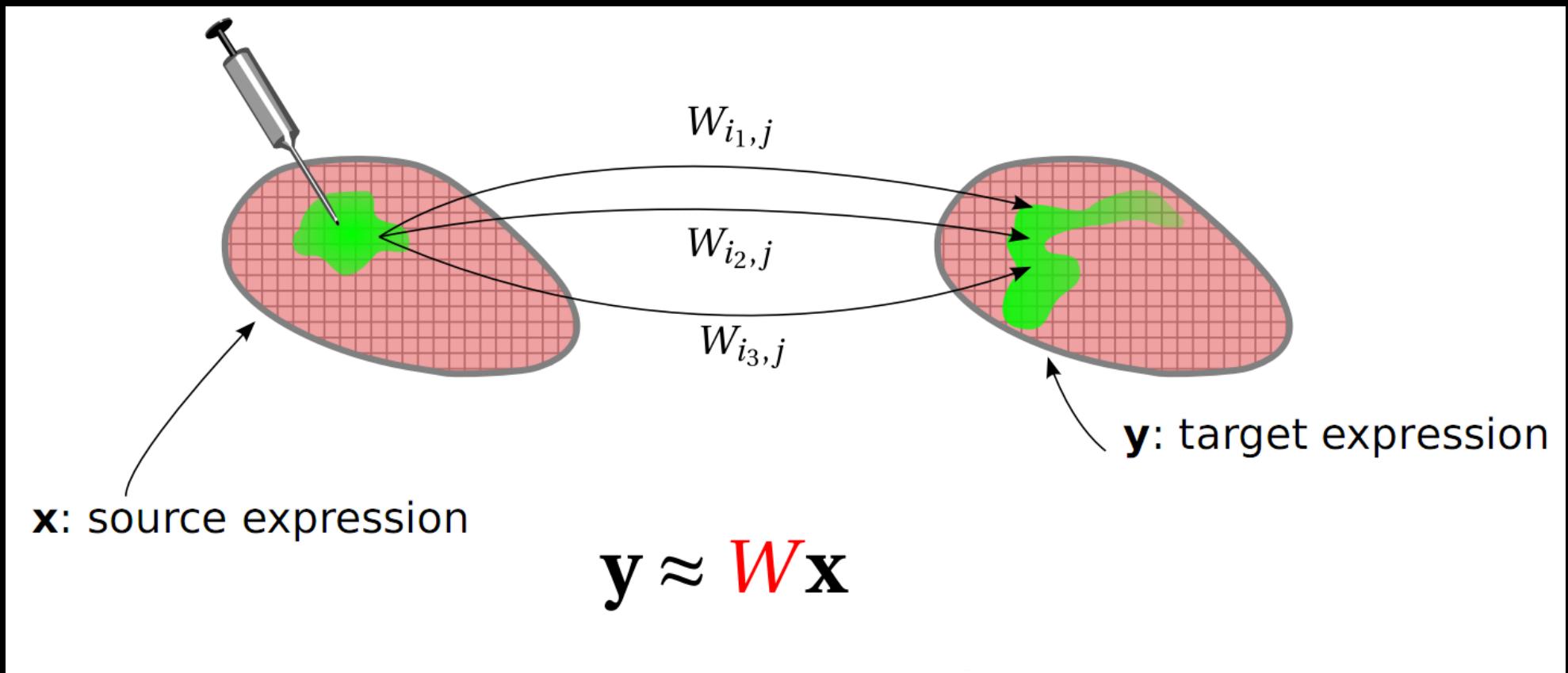


Mesoscale viral tracing experiments Allen Institute for Brain Science (AIBS)

Oh et al. Nature (2014)



Model connectivity as a matrix



Challenges for inference:

Challenges for inference:

- 1) Injection sites do not cover whole brain; model is underdetermined

Challenges for inference:

- 1) Injection sites do not cover whole brain; model is underdetermined
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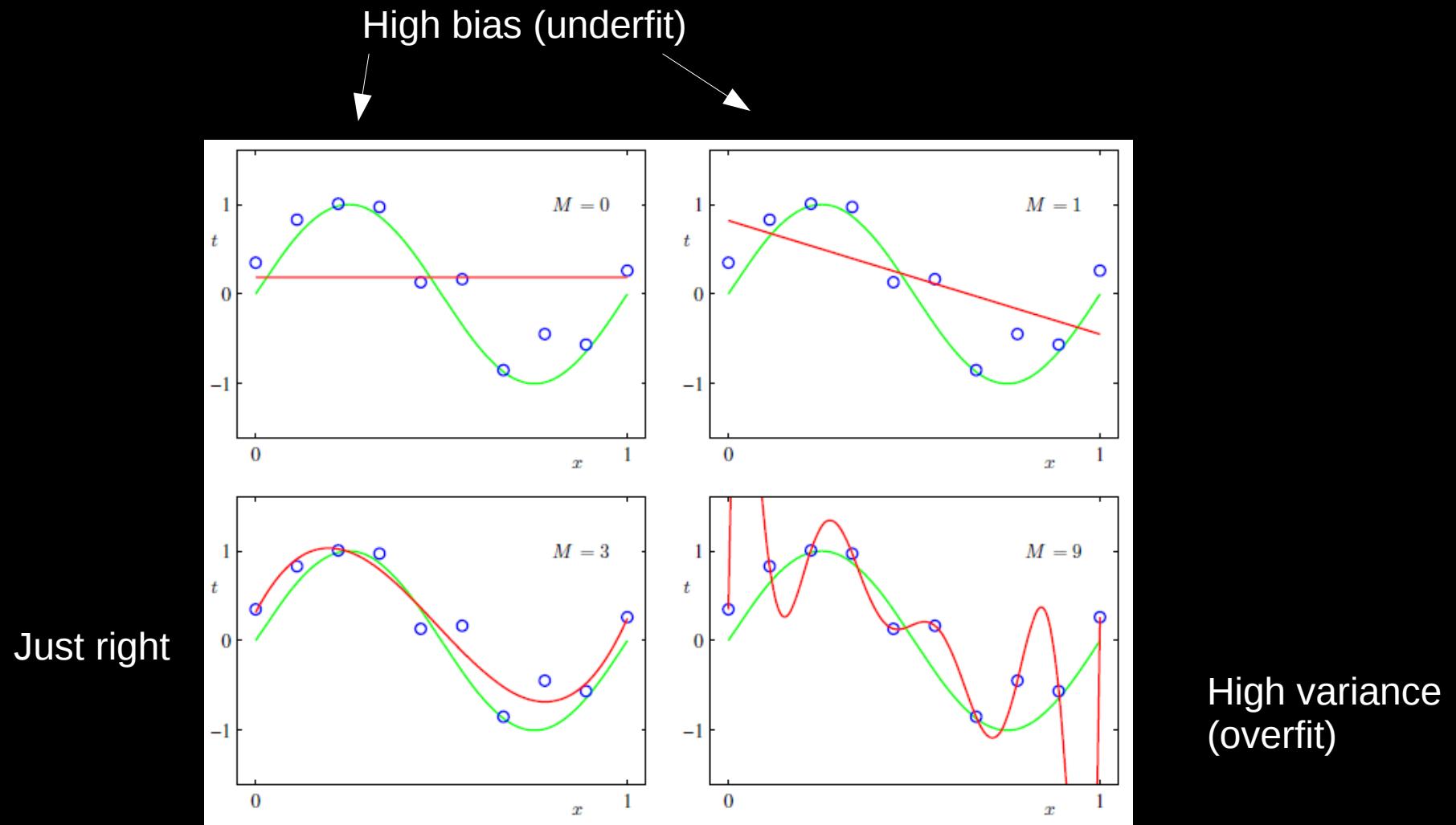
Challenges for inference:

- 1) Injection sites do not cover whole brain; model is underdetermined
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- 3) Dimensionality of unknown W

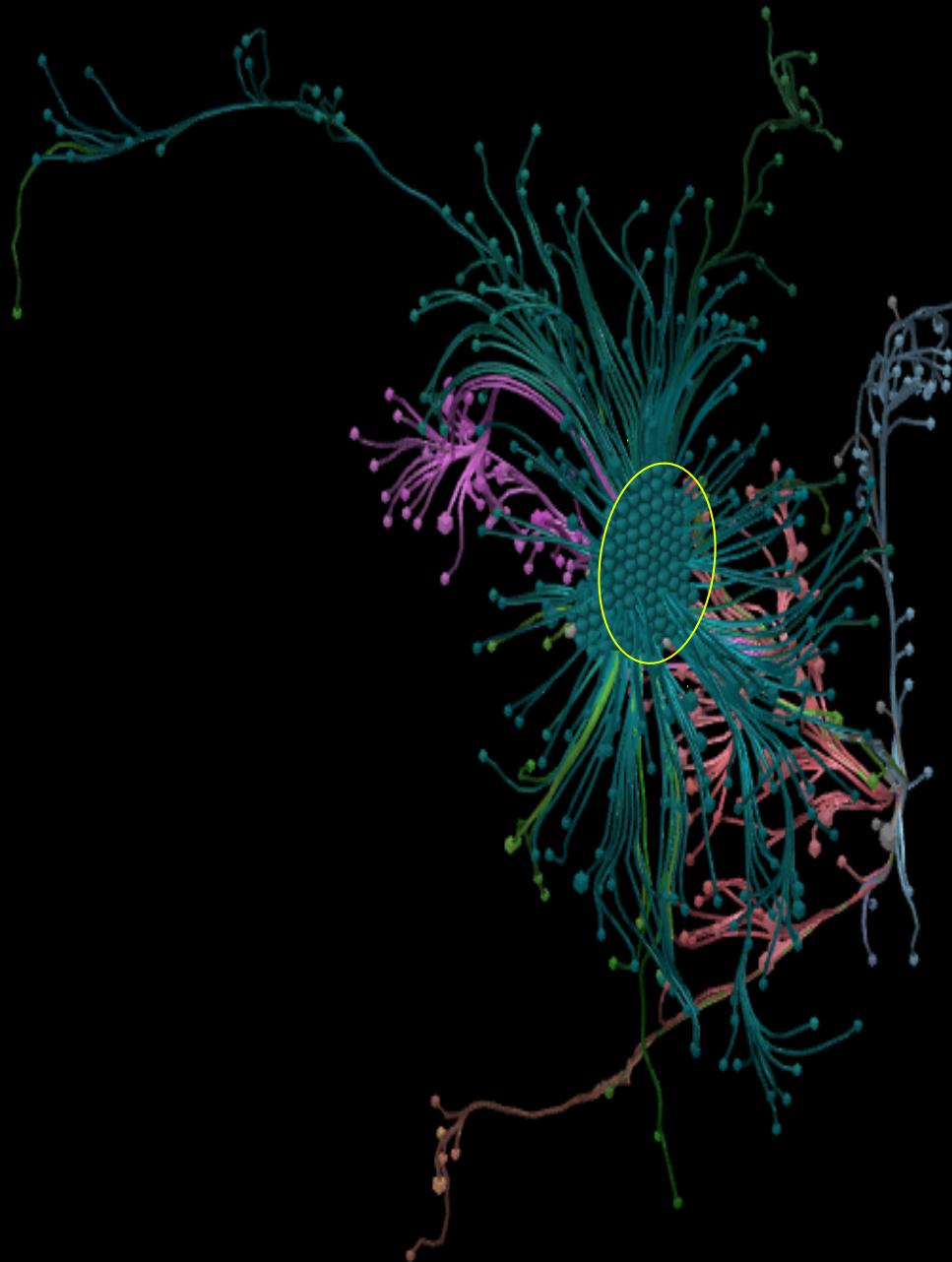
Challenges for inference:

- 1) Injection sites do not cover whole brain; model is underdetermined
 - Fill in gaps with smoothing regularizer
- 2) Projection strength unknown at injection site
- 3) Dimensionality of unknown W

Regularization prevents over-fitting

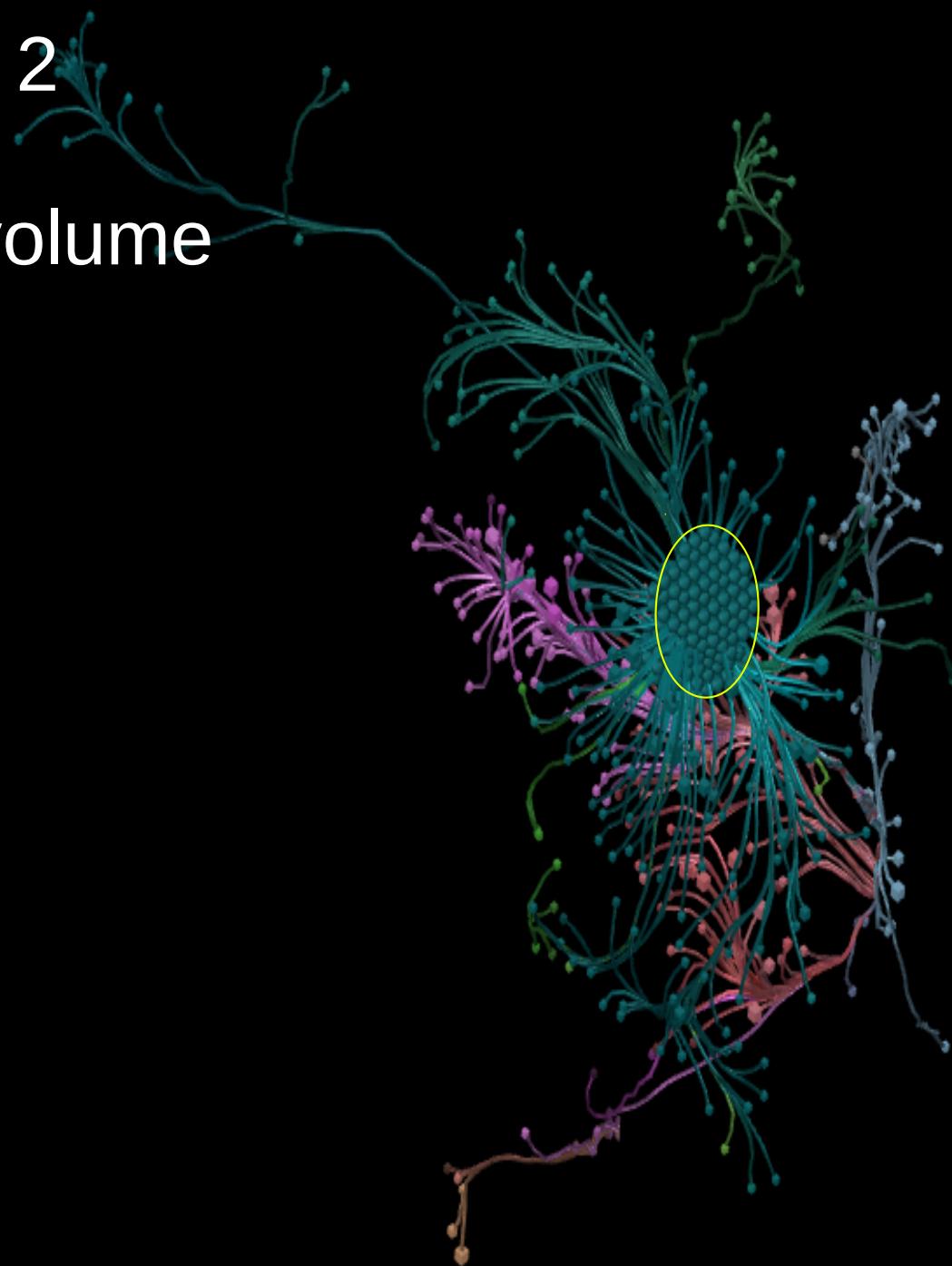


Injection 1 into VISp

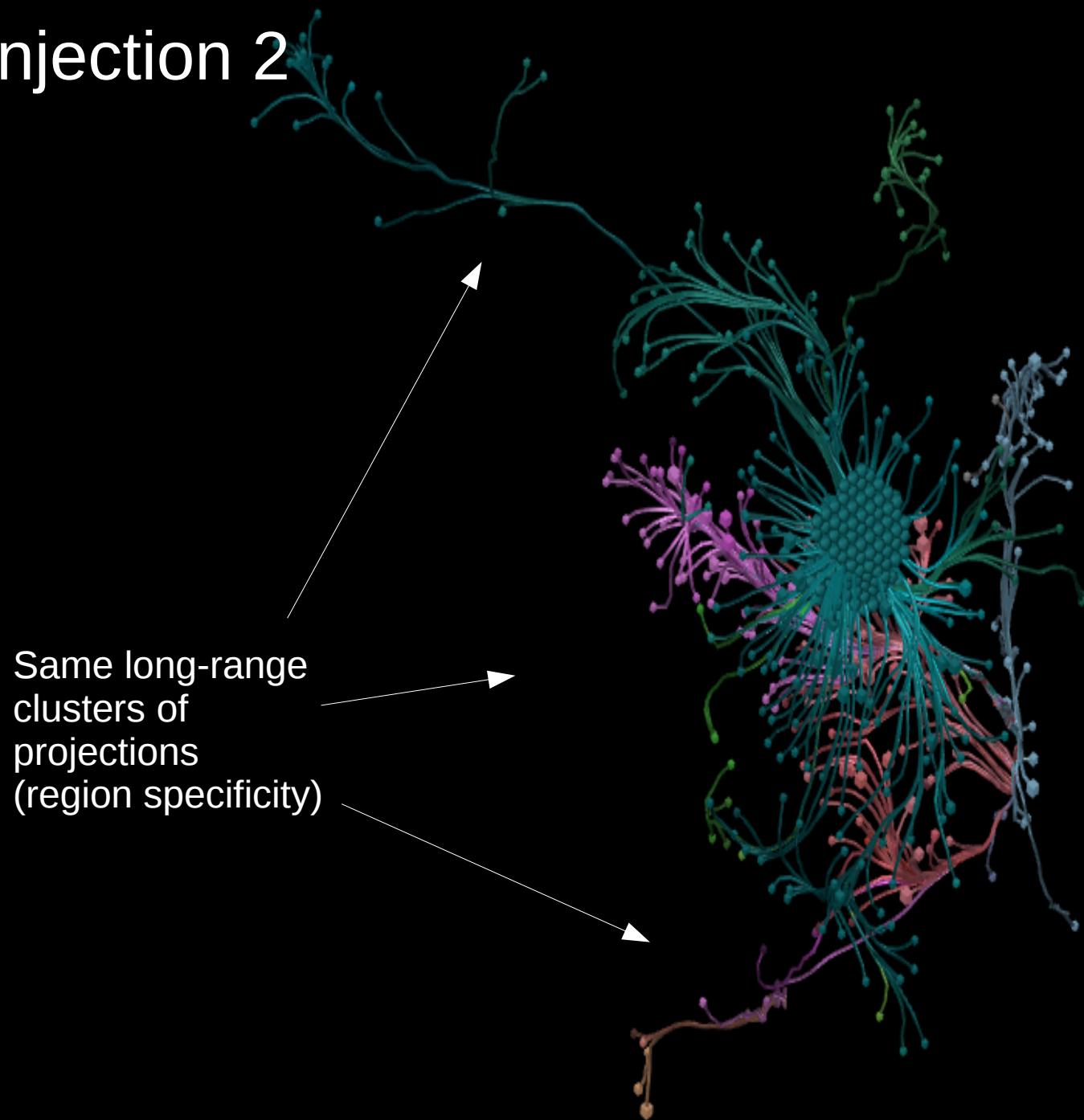


Images made with
AIBS Brain Explorer

Injection 2
Nearby
Similar volume

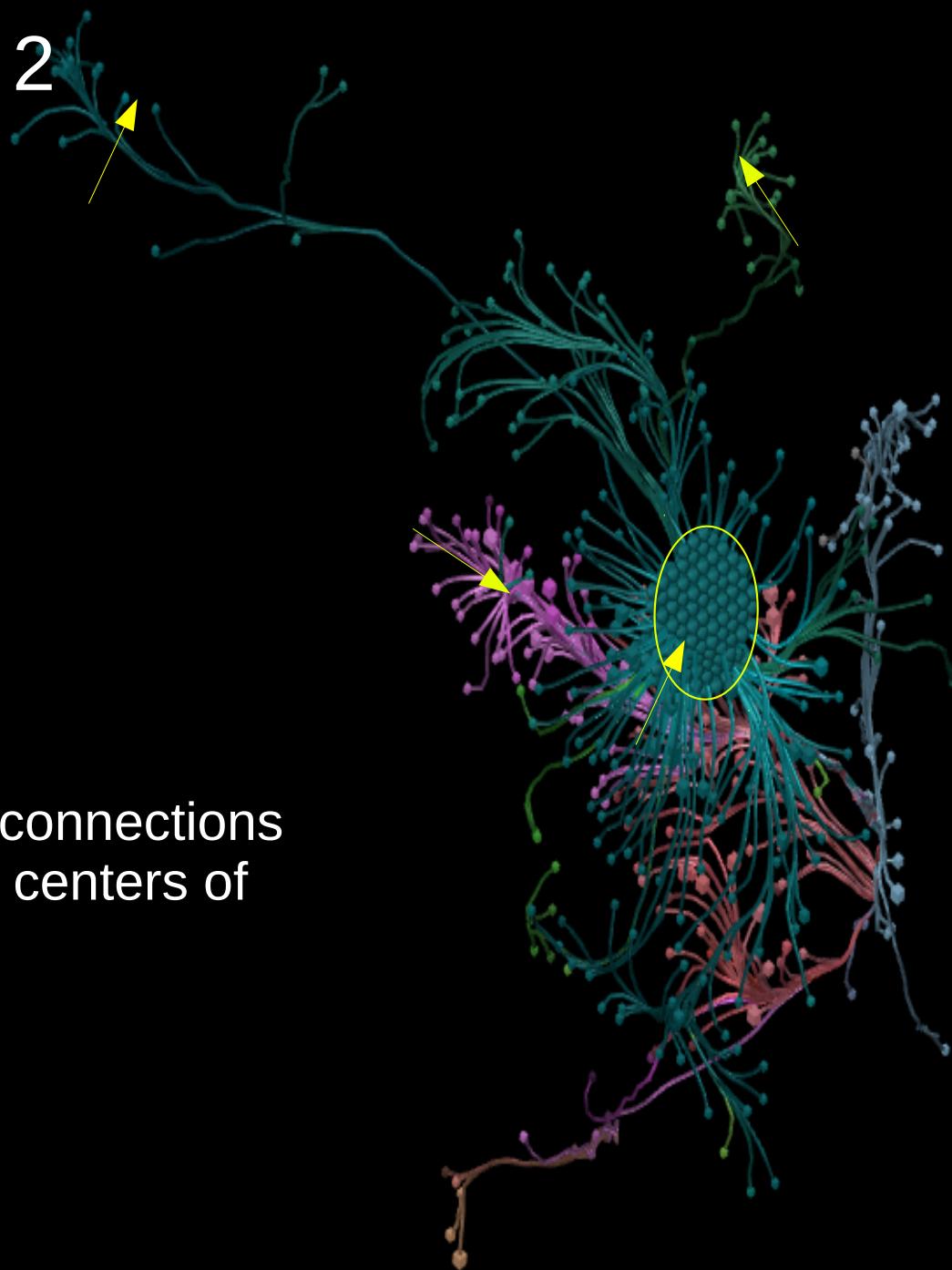


Injection 2

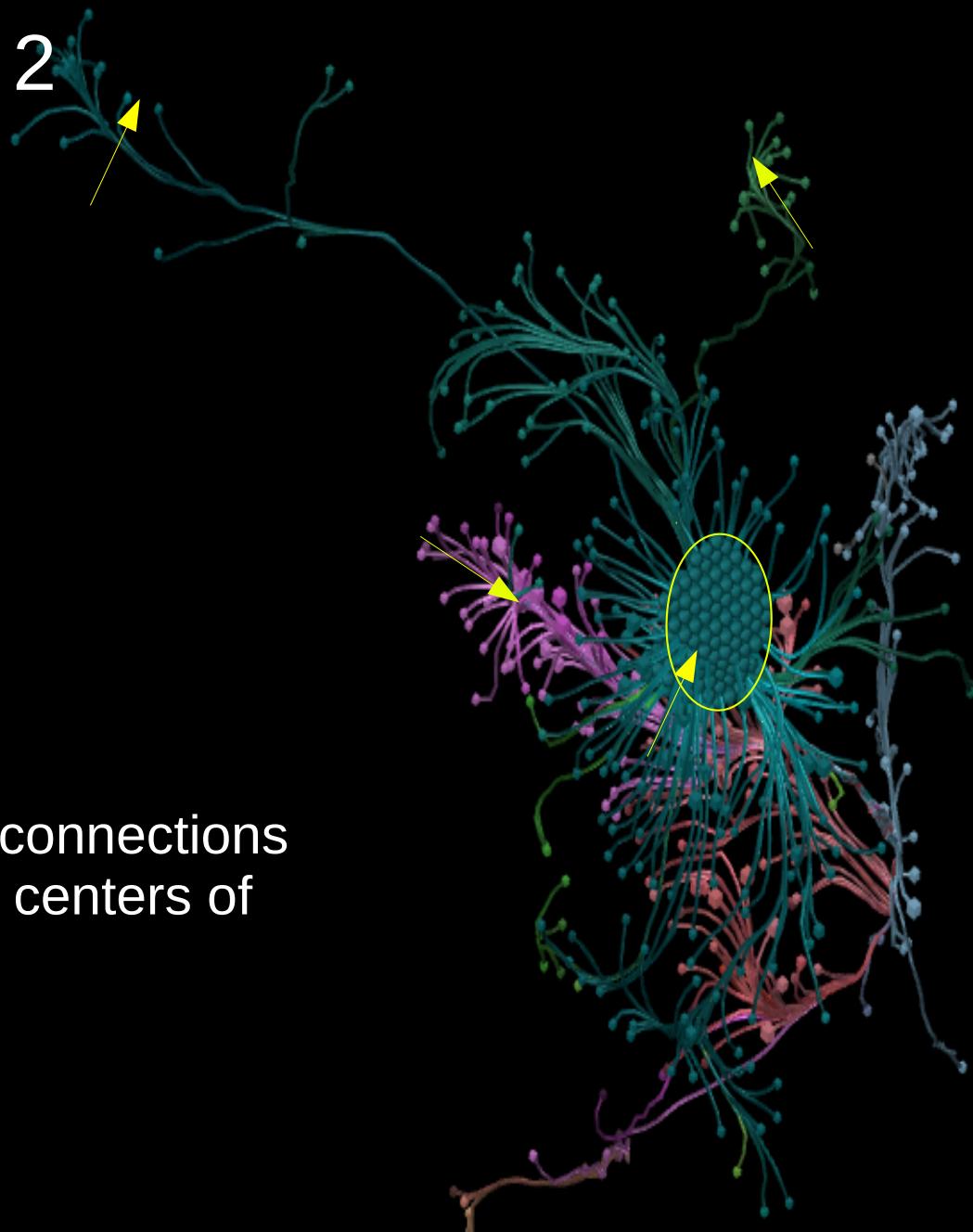


Injection 2

Long-range connections
have shifted centers of
mass



Injection 2



Long-range connections
have shifted centers of
mass

Mesoscale not so messy
Smoothness assumption is reasonable

Challenges for inference:

- 1) Injection sites do not cover whole brain; model is underdetermined
 - Fill in gaps with smoothing regularizer
- 2) Projection strength unknown at injection site
 - Ignore unknown residuals
- 3) Dimensionality of unknown W

Netflix challenge: matrix completion

Ratings of movies →

| Users | | | -1 | | | 1 | 1 | -1 | 1 | -1 |
|-------|---|---|----|---|----|---|---|----|---|----|
| | | | 1 | | | 1 | 1 | -1 | 1 | -1 |
| | 1 | 1 | -1 | 1 | -1 | 1 | 1 | -1 | 1 | -1 |
| | 1 | | | | -1 | 1 | 1 | -1 | 1 | -1 |
| ▼ | | | -1 | | | 1 | 1 | -1 | 1 | -1 |

Lepeisi, <https://commons.wikimedia.org/w/index.php?curid=45676223>

Observation M

Full solution W , rank 1

Netflix challenge: matrix completion

Ratings of movies →

| Users | | | -1 | | | 1 | 1 | -1 | 1 | -1 |
|-------|---|---|----|---|----|---|---|----|---|----|
| | | | | 1 | | 1 | 1 | -1 | 1 | -1 |
| | 1 | 1 | -1 | 1 | -1 | 1 | 1 | -1 | 1 | -1 |
| | 1 | | | | -1 | 1 | 1 | -1 | 1 | -1 |
| ▼ | | | -1 | | | 1 | 1 | -1 | 1 | -1 |

Lepeisi, <https://commons.wikimedia.org/w/index.php?curid=45676223>

Observation M

Full solution W , rank 1

$$\min_W \|W\|_* \text{ where } P(W - M) = 0$$

Complexity of W

Match W to data M where observed

Challenges for inference:

- 1) Injection sites do not cover whole brain; model is underdetermined
 - Fill in gaps with smoothing regularizer
- 2) Projection strength unknown at injection site
 - Ignore unknown residuals
- 3) Dimensionality of unknown W
 - Make $W = UV$ low rank

Both regularization and compression

Spline regression model of connectome

- Find W nonnegative that minimizes the expression

$$\|P(WX - Y)\|_F^2 + \lambda \|L(W)\|_F^2$$

Goodness of fit
(loss)

Roughness penalty
(regularization, prior)

Spline regression model of connectome

- Find W nonnegative that minimizes the expression

$$\|P(WX - Y)\|_F^2 + \lambda \|L(W)\|_F^2$$

Squared Frobenius norms = sum of squares of matrix entries
= Gaussian noise model

Spline regression model of connectome

- Find W nonnegative that minimizes the expression

$$\|P(WX - Y)\|_F^2 + \lambda \|L(W)\|_F^2$$


Unknown weight matrix

Spline regression model of connectome

- Find W nonnegative that minimizes the expression

$$\|P(WX - Y)\|_F^2 + \lambda \|L(W)\|_F^2$$

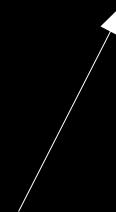


Injection data

Spline regression model of connectome

- Find W nonnegative that minimizes the expression

$$\|P(WX - Y)\|_F^2 + \lambda \|L(W)\|_F^2$$



Projection data

Spline regression model of connectome

- Find W nonnegative that minimizes the expression

$$\|P(WX - Y)\|_F^2 + \lambda \|L(W)\|_F^2$$



Deals with holes in data, just like **matrix completion**

Ignores residuals in injection sites

Spline regression model of connectome

- Find W nonnegative that minimizes the expression

$$\|P(WX - Y)\|_F^2 + \lambda \|L(W)\|_F^2$$

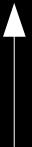


Controls strength of smoothing

Spline regression model of connectome

- Find W nonnegative that minimizes the expression

$$\|P(WX - Y)\|_F^2 + \lambda \|L(W)\|_F^2$$



Laplacian “roughness” of W

Can respect region boundaries

6-dimensional!

Spline regression model of connectome

- Find W nonnegative that minimizes the expression

$$\|P(WX - Y)\|_F^2 + \lambda \|L(W)\|_F^2$$

Problem is convex:

unique global solution and **standard methods to find it**

We found W for mouse visual cortex

- 7497 x 7497 matrix fit with 28 injections
- **Marked improvement** over regional model:

| | | |
|-------------------|--------------|------------|
| - Regional model: | 48% regional | 107% voxel |
| - Voxel model: | 16% regional | 33% voxel |

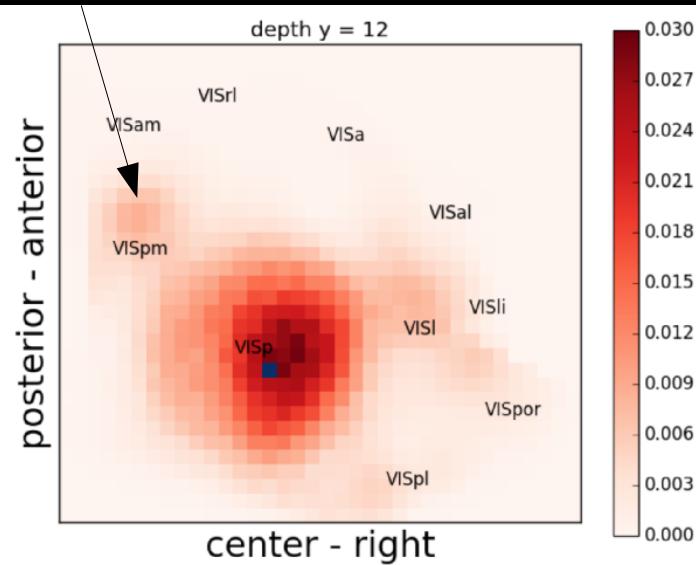
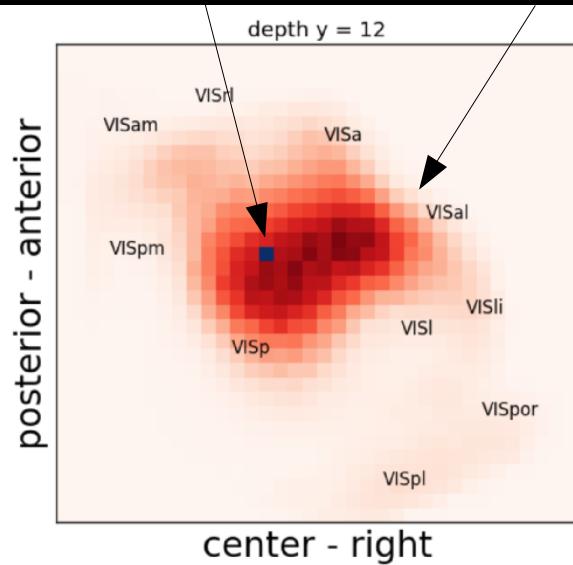
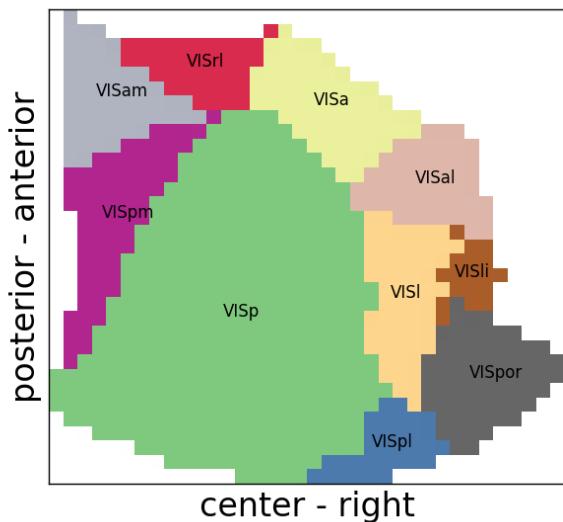
$$\frac{\|Y_{\text{pred}}\|^2}{\frac{1}{2}\|Y_{\text{pred}}\|^2 + \frac{1}{2}\|Y_{\text{true}}\|^2}$$

Applied to visual cortex we see hints of retinotopy

Maps of retinal (visual) space between areas

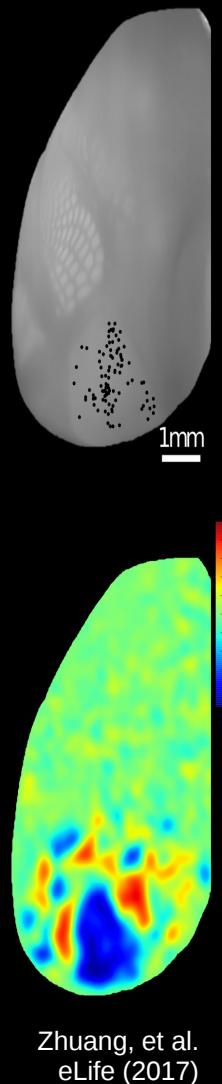
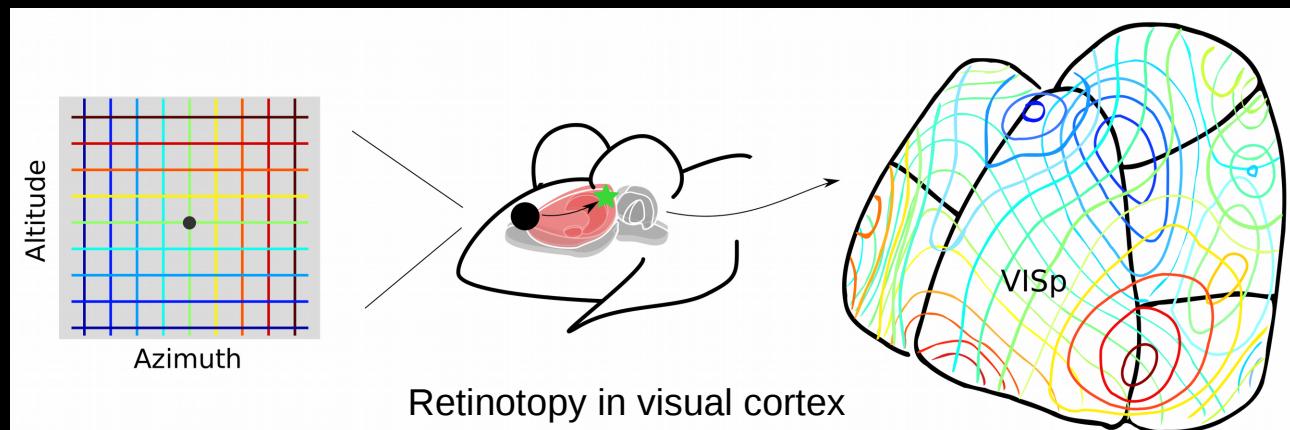
Source voxel

“Blobby” projections move with source



Spatial connectivity opens exciting scientific directions

- Comparable to neuron activity data
 - e.g., Kim et al. *Cell Reports* (2016), “Crystal Skull”
- Topography between regions (retinotopy, tonotopy, etc.)
- Cell-type specific data: E/I, layers, etc.
- Connectivity-defined regions?



Conclusions: mesoscopic inference

Conclusions: mesoscopic inference

- New spatial inference method tailored to data

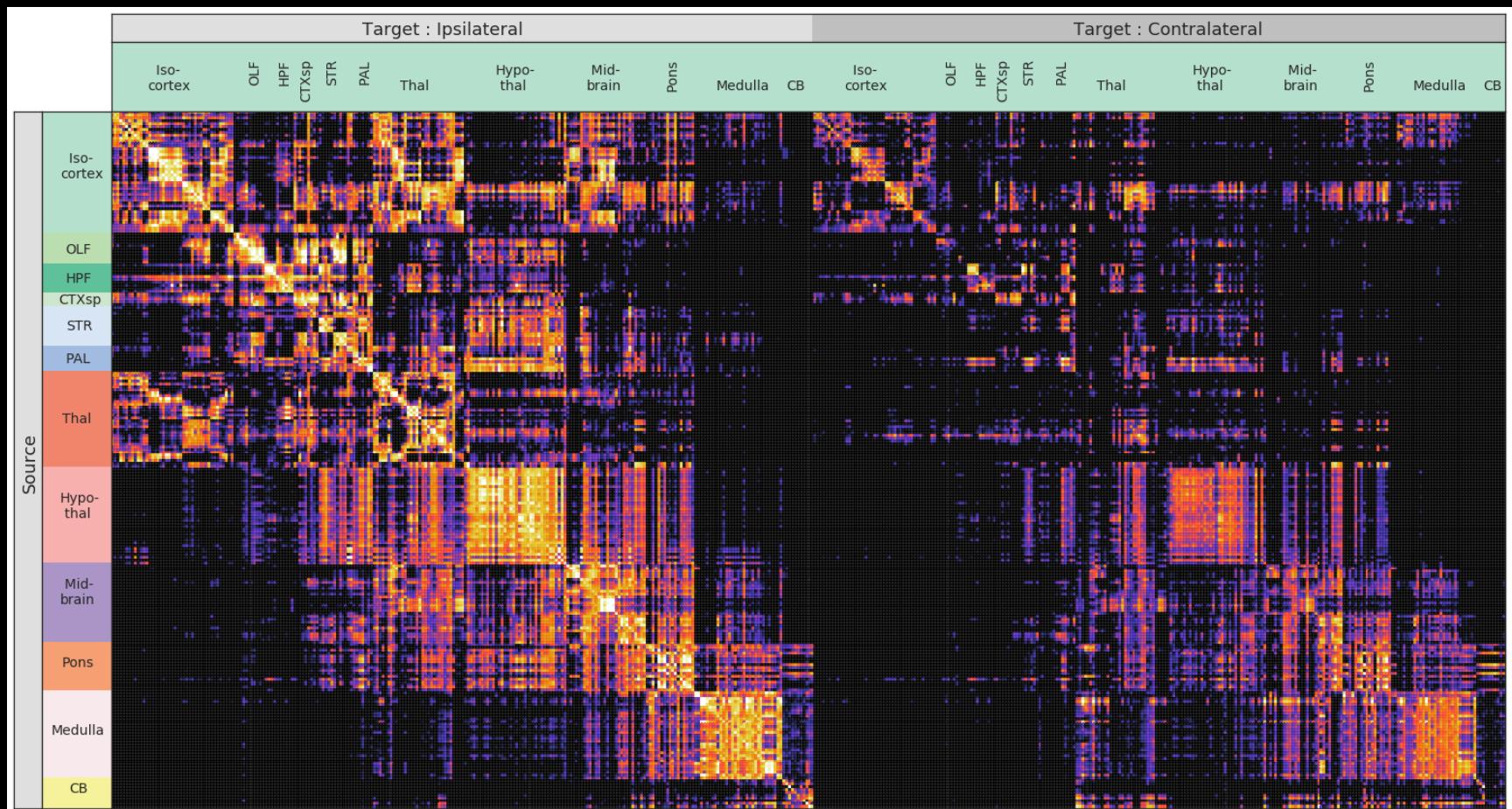
Conclusions: mesoscopic inference

- New spatial inference method tailored to data
 - More predictive than before

Conclusions: mesoscopic inference

- New spatial inference method tailored to data
 - More predictive than before
 - Improves regional models, too!

Joe Knox (AIBS)

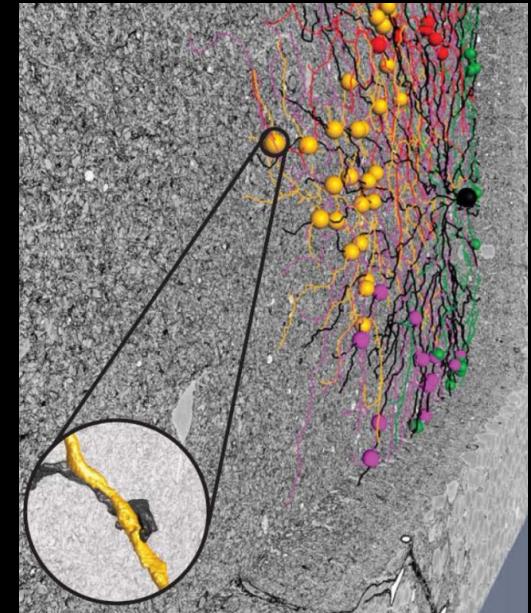


Connectivity questions

- 1) How do we infer mesoscopic connections?
- 2) What are good microscopic models?
- 3) How does connectivity affect brain dynamics?

Zooming in with random graphs

In preparation:
Brito, Dumitriu, Harris.
“Spectral gap in random bipartite biregular graphs and its applications.”

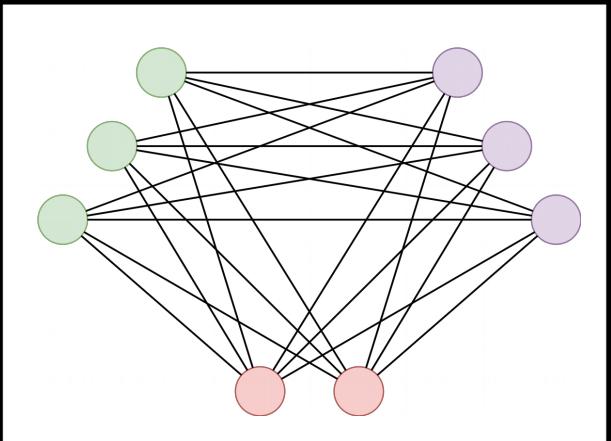
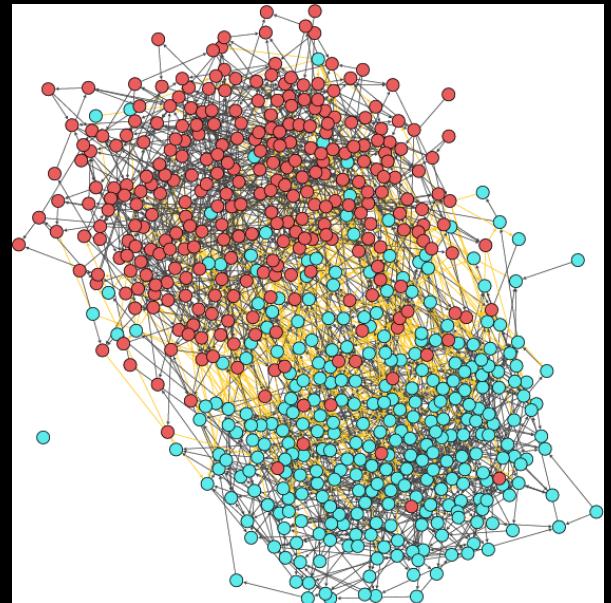


Kleinfeld et al. (2011)

Random graphs can match measureable microscopic features

Joshua Mendoza

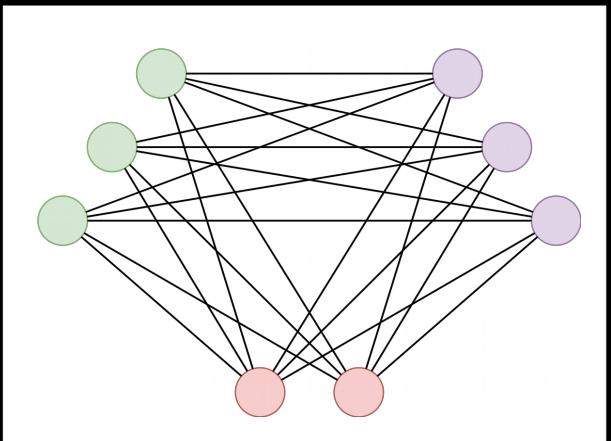
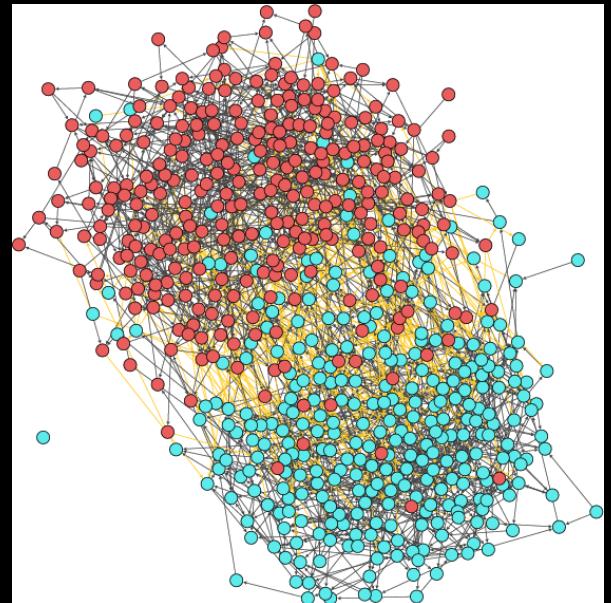
- Example statistics:
 - Average # connections (degree)
 - Number of cycles
 - Other subgraph counts



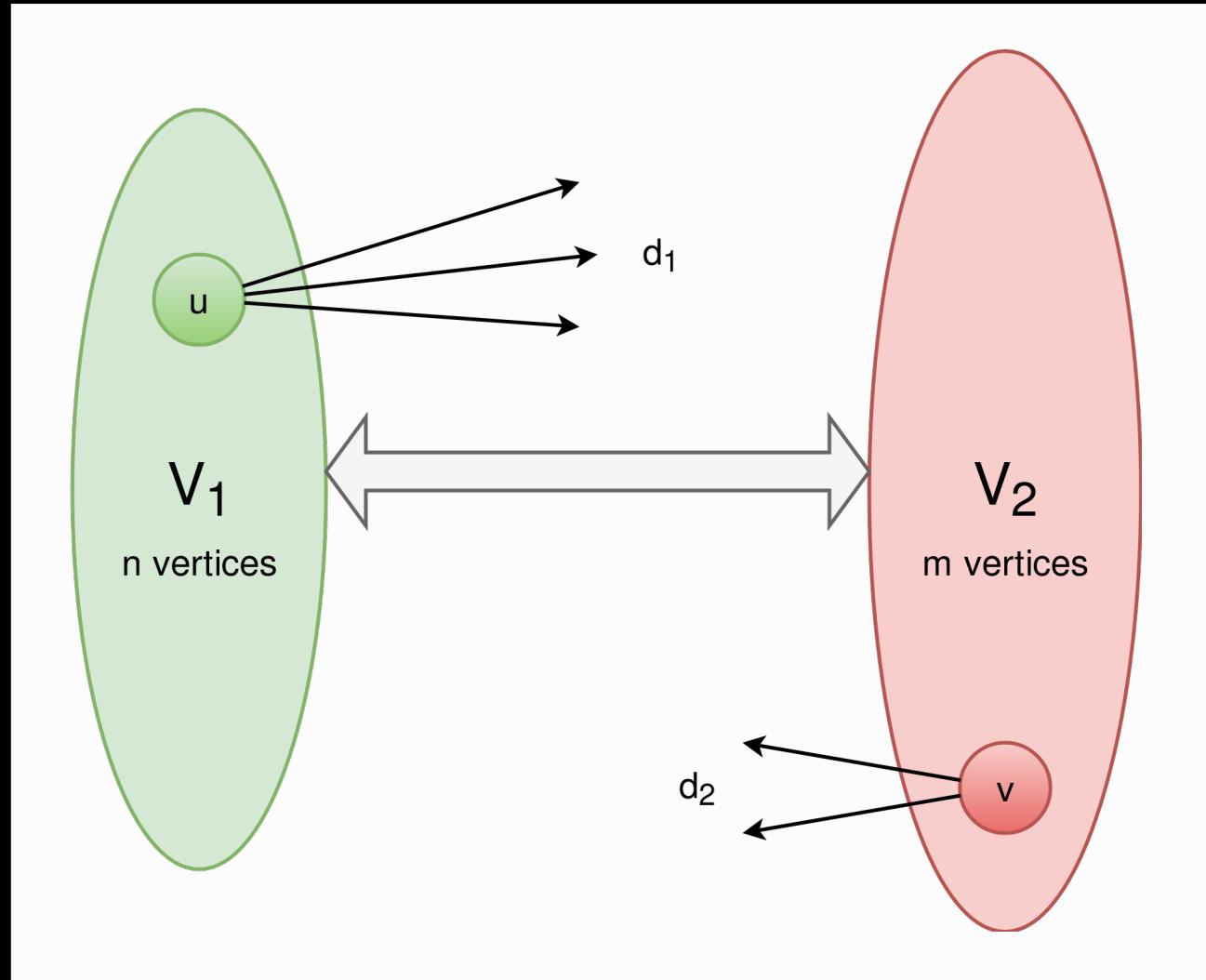
Random graphs can match measureable microscopic features

Joshua Mendoza

- Example statistics:
 - Average # connections (degree)
 - Number of cycles
 - Other subgraph counts
- Community structure
 - Block models
 - Multi-partite graphs



Bipartite, biregular random graphs



Fixed d 's mean these graphs are **very sparse**

We found a spectral “gap!”

- $\lambda_1 \gg \lambda_2 \rightarrow \text{expander}$

Extends past work for d-regular graphs:

Friedman (2003, 2004)

Alon (1986)

Bordenave (2015)

Angel, Friedman, Hoory (2015)

Marcus, Spielman, Srivastava (2013)

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 - Random walks
 - Community detection & spectral clustering
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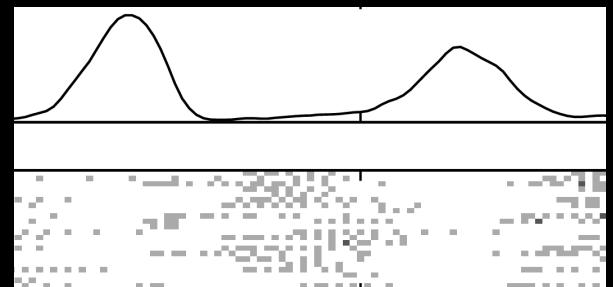
Marcus, Spielman, Srivastava (2013)

Hold that thought!

Connectivity questions

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Effects of sparsity on rhythm generation



Published as:

Harris, Dashevskiy, Mendoza, Garcia III, Ramirez, Shea-Brown.

“Different roles for inhibition in the rhythm-generating respiratory network.”

J Neurophys 118(4), 2070-2088. 2017.

Litwin-Kumar, Harris, Sompolinsky, Abbott.

“Optimal synaptic connectivity.”

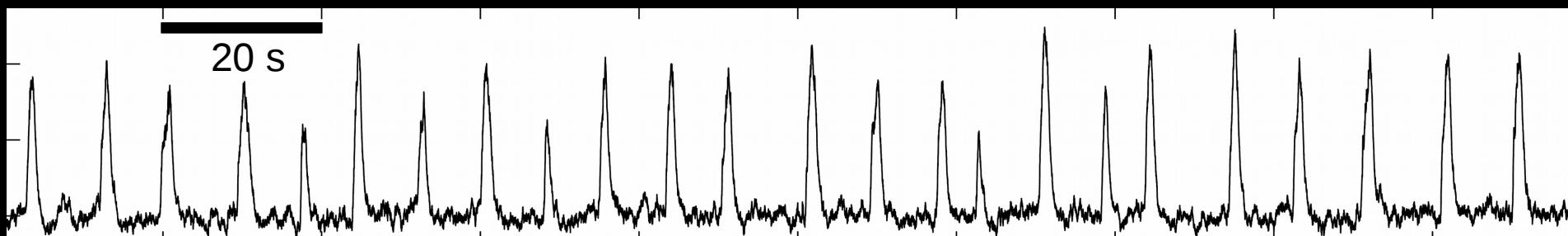
Neuron 93, 1153-1164. 2017.



Sparsity is good for
“classifier” circuits

Breathing arises in the brainstem

- The “pre-Bötzinger Complex”

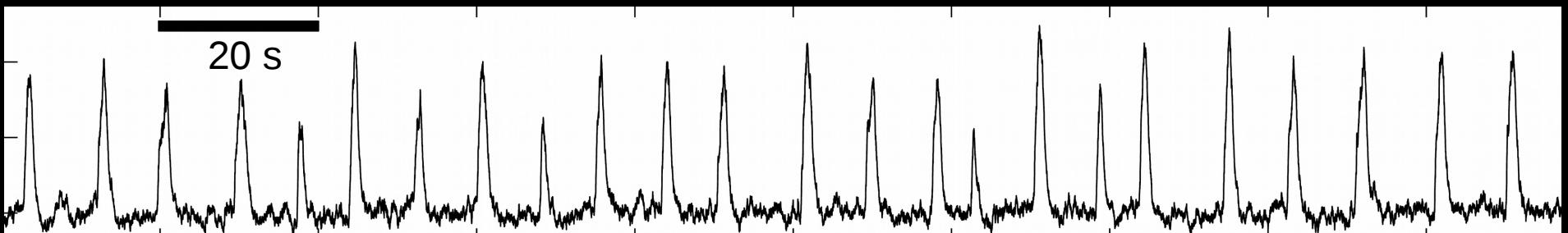


Important related papers:

Smith et al. 1991
Ramirez & Richter 1996
Rekling & Feldman 1998
Feldman et al. 2013

Breathing arises in the brainstem

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 - Bursting, tonic spiking, and quiescent E / I cells

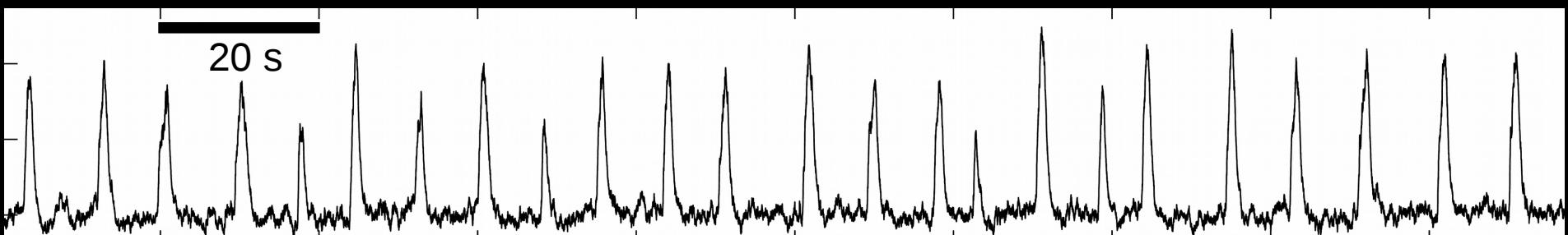


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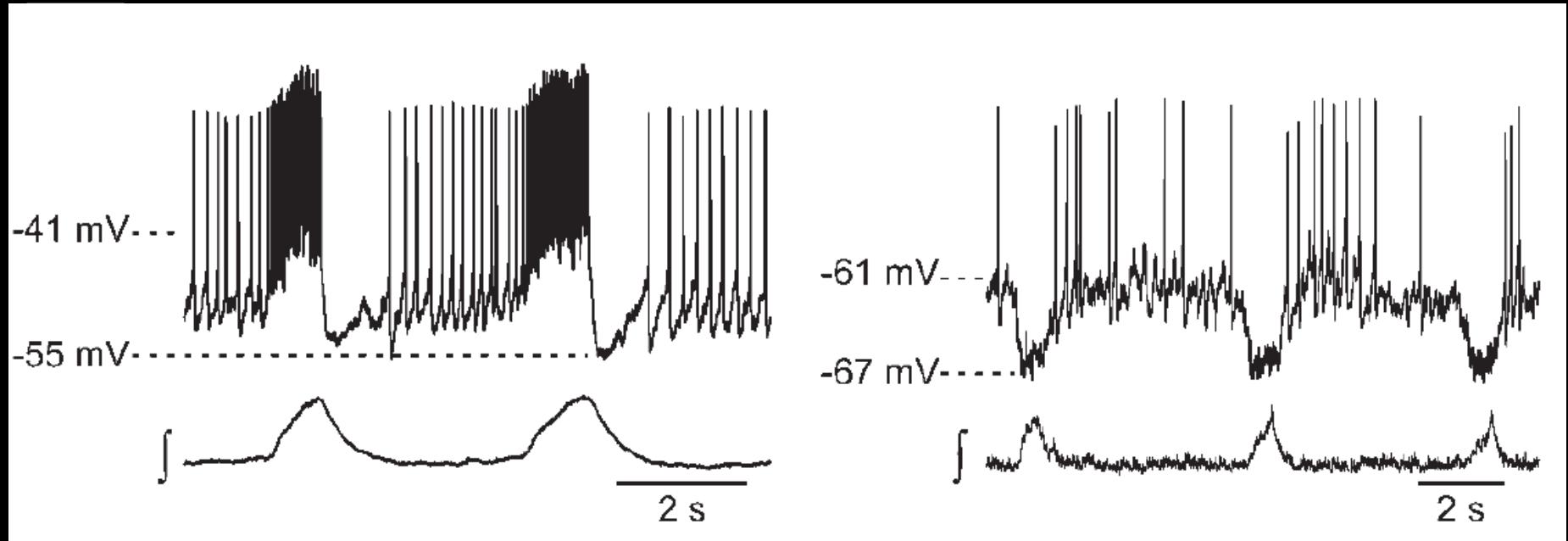
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 - Bursting, tonic spiking, and quiescent E / I cells
 - Synchronized population bursts via excitation



Important related papers:

Smith et al. 1991
Ramirez & Richter 1996
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Feldman et al. 2013

Why do you get expiratory cells?

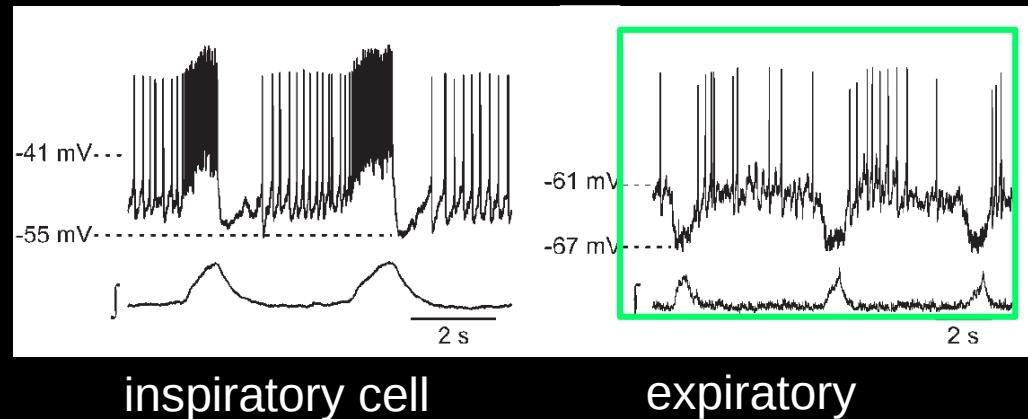


inspiratory

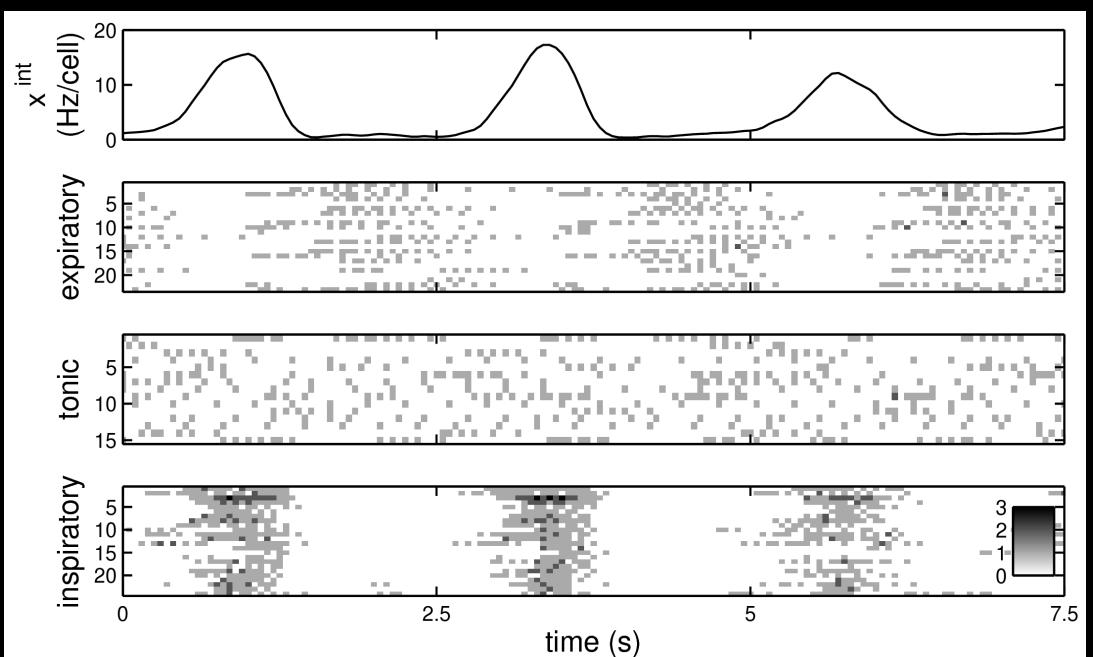
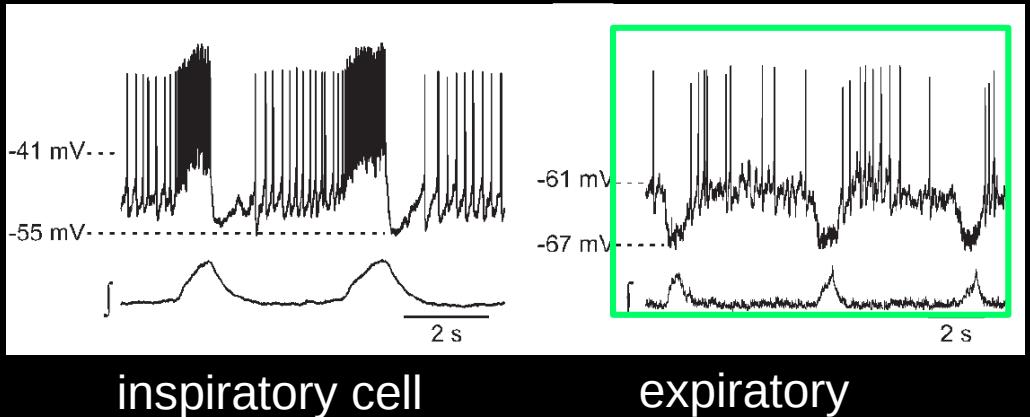
expiratory

Figure from:
Lieske, Thoby-Brisson, Telgkamp, Ramirez (2000)

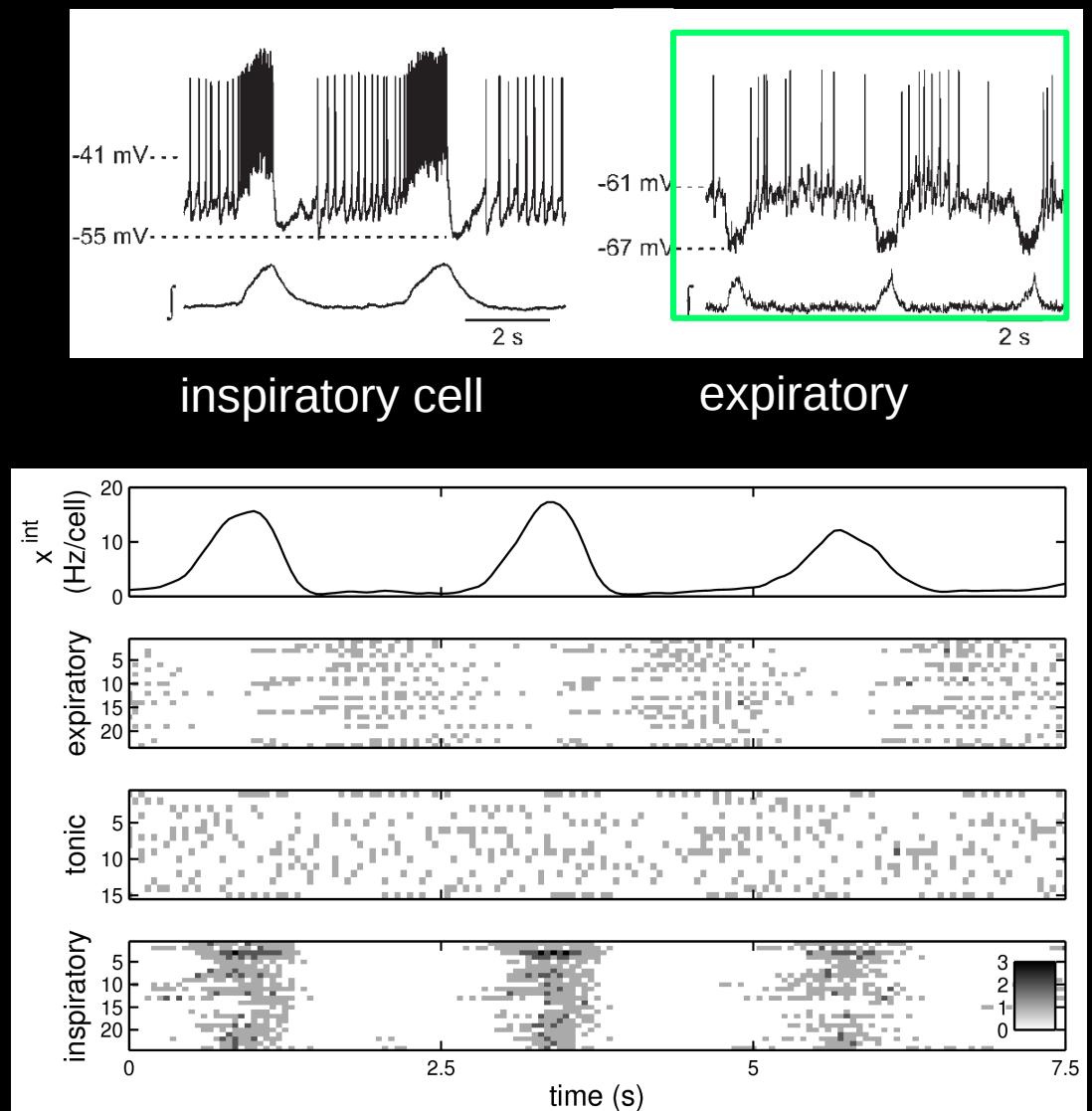
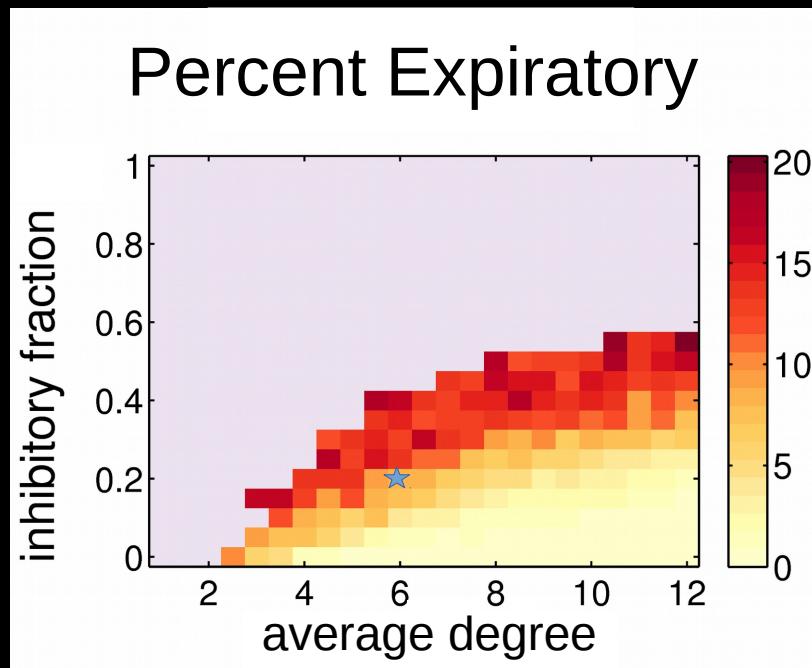
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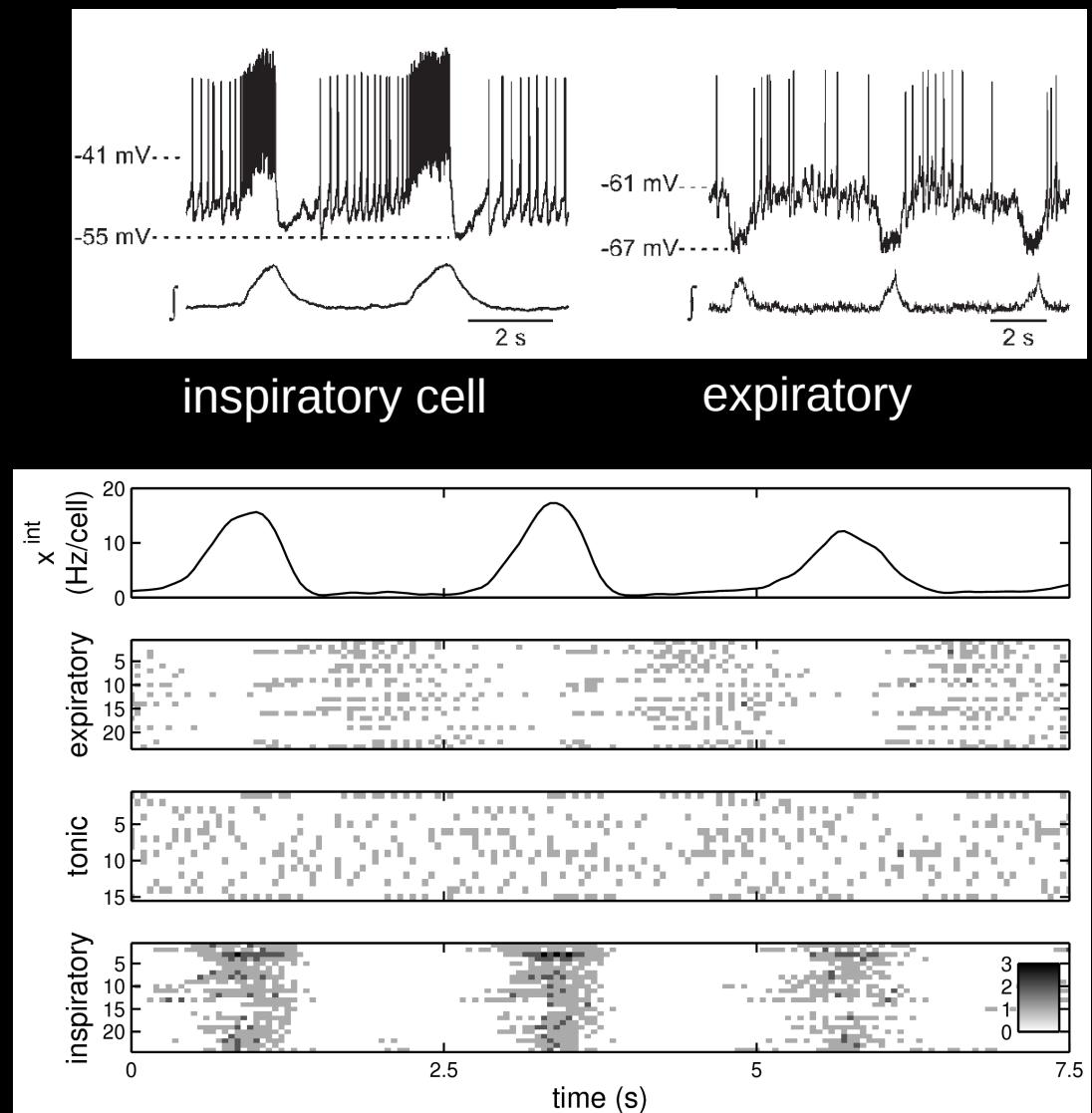
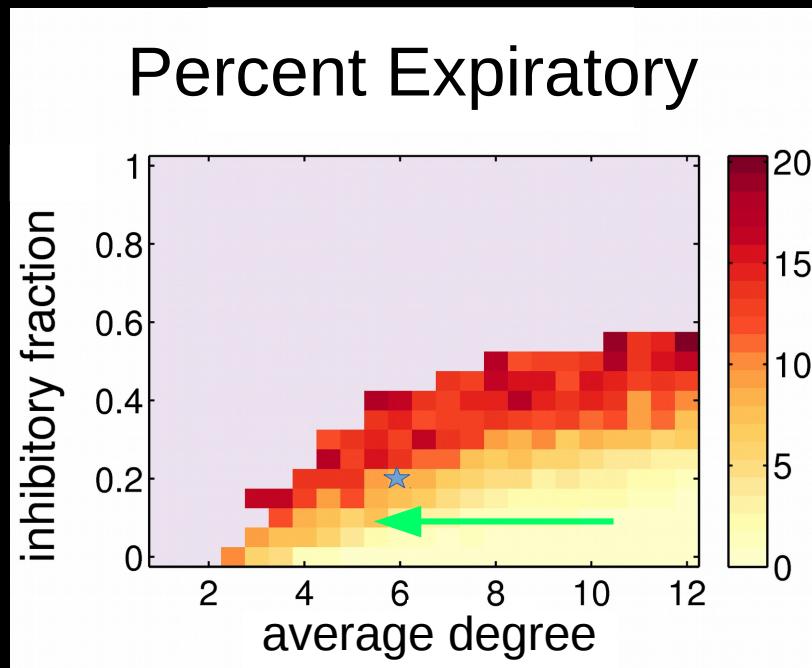
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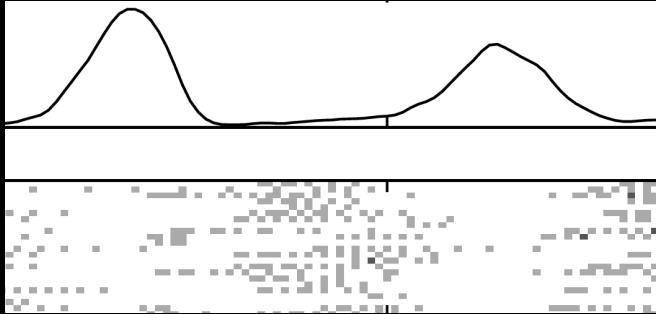
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Why do you get expiratory cells?



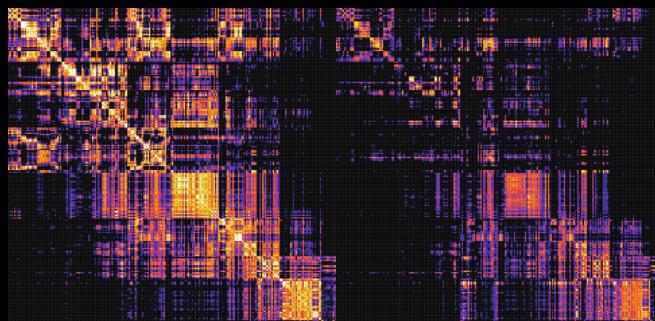
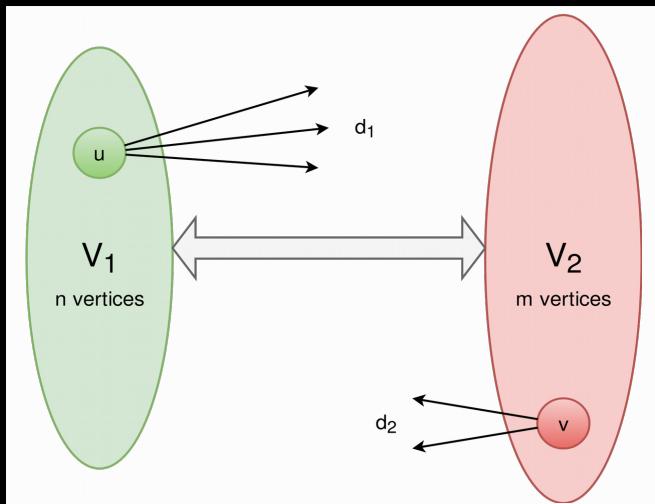
Sparse graph:
some cells randomly
driven into expiratory phase



- Neuronal network dynamics
 - Choose graph class with measurable parameters

- Random graph theory
 - Rich mathematical area
 - Spectra, expansion, & dynamics

- Mesoscopic network inference
 - Methods tailored to new datasets



More acknowledgements



Erik Turner



Cody Lourie





Stuart Ryan



Jason Griffith



Drew Tabke

Thank you for listening!

