Neuroscience Flythrough

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Content

Raw Observations

What does the brain look like?

Modeling

How can we model/replicate the behavior?

Testing

Does our method make sense?

Content

Raw Observations

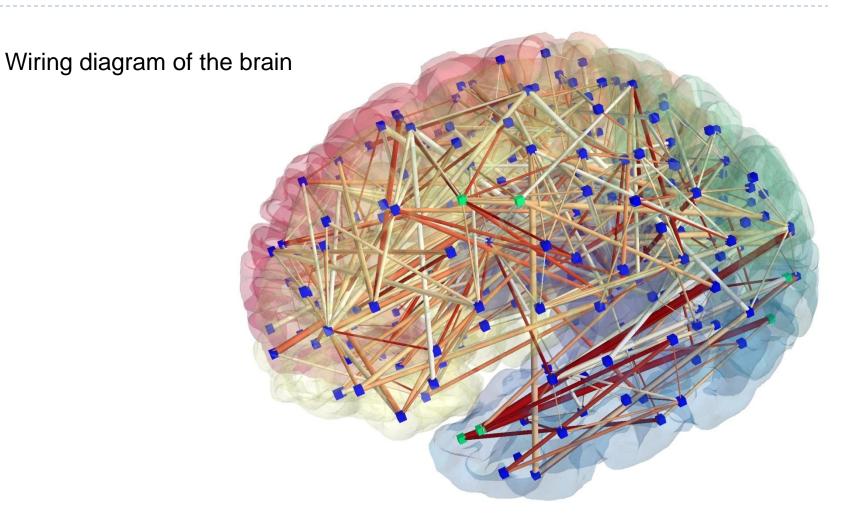
What does the brain look like?

Modeling

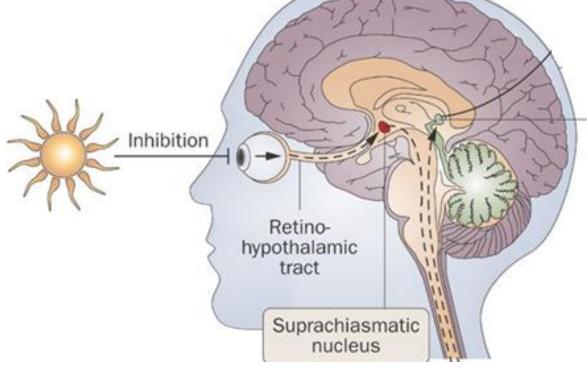
How can we model/replicate the behavior?

Testing

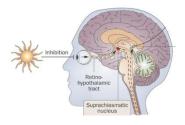
Does our method make sense?



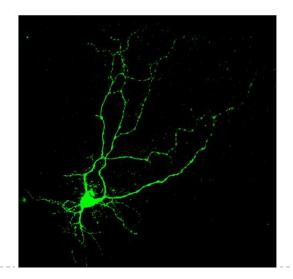
- Macro
 - Node is gray matter region (e.g. retina and suprachiasmic nucleus)

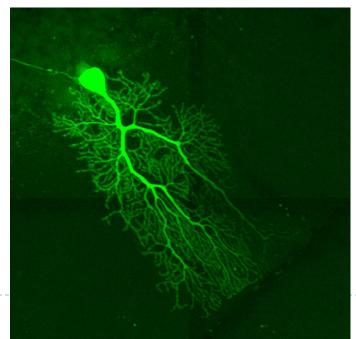


- Macro
 - Node is gray matter region (e.g. retina and suprachiasmic nucleus)



- Meso
 - Node is neuron **type** (e.g. granule cells, Purkinje cells)



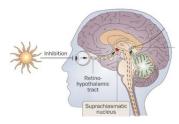


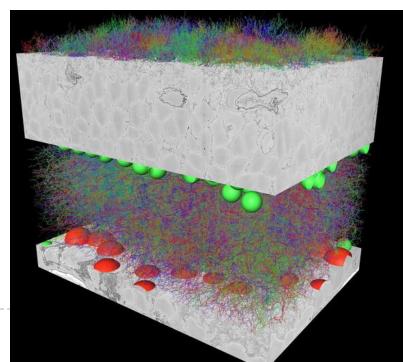
Connectomes

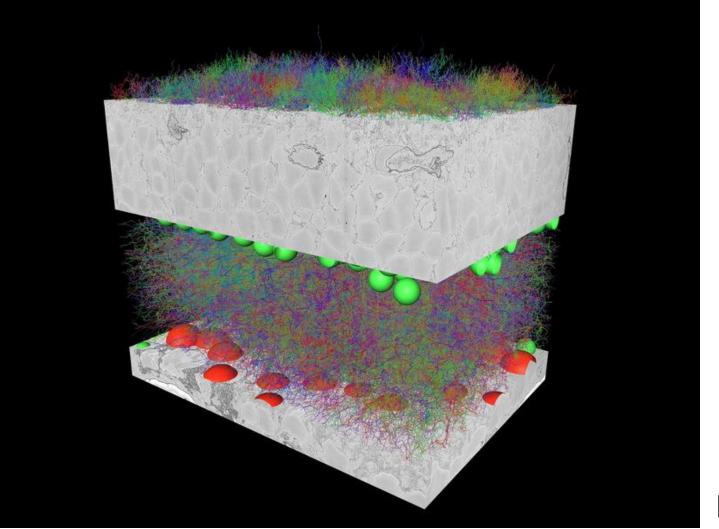
- Macro
 - Node is gray matter region (e.g. retina and suprachiasmic nucleus)

[4]

- Meso
 - Node is neuron type
- Micro
 - Nodes are neurons

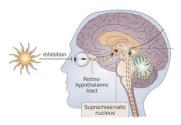


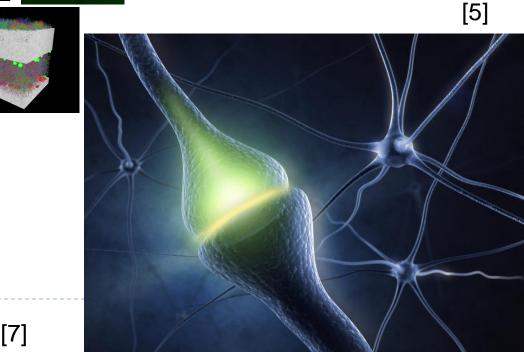




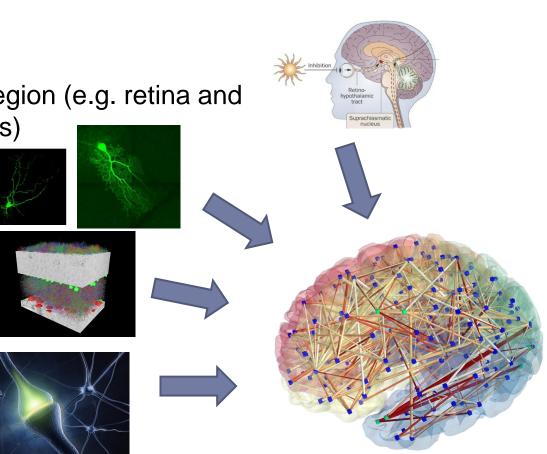
[4]

- Macro
 - Node is gray matter region (e.g. retina and suprachiasmic nucleus)
- Meso
 - Node is neuron type
- Micro
 - Nodes are neurons
- Nano
 - Nodes are synapses

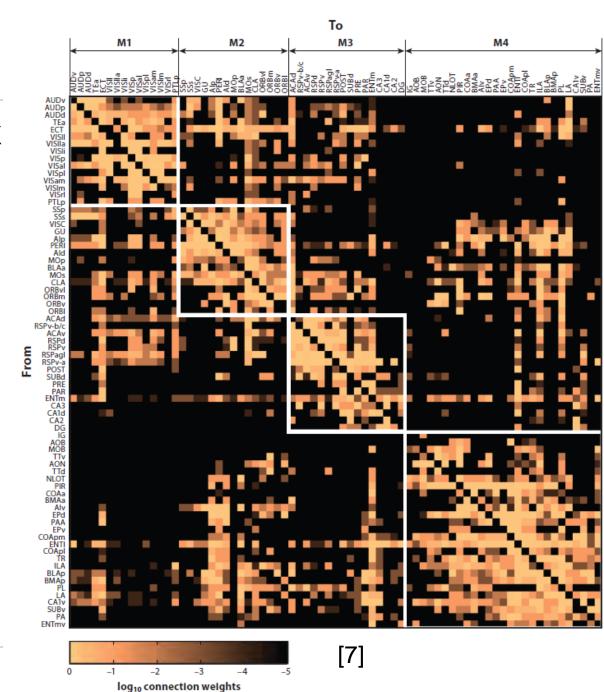




- Macro
 - Node is gray matter region (e.g. retina and suprachiasmic nucleus)
- Meso
 - Node is neuron type
- Micro
 - Nodes are neurons
- Nano
 - Nodes are synapses



- Connectivity Matrix
 + connectivy
 - Distance
 - # of synapses
 - Size of synapse
 - Type of synapse



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Computational Cognitive Neuroscience

- Model Brain activity
 - Single neuron
 - Leaky integrate and fire model
 - Hodkin Huxley
 - Compartments
 - Learning
 - Reinforcment Learning (Homer Bild)
 - Hebbian Rules
 - Long Term Potentation
 - Dopamine as Reward
 - □ Stratium vs Frontal Cortex
 - Long Term Depression

Computational Cognitive Neuroscience

Goal:

- Modeling the neurodynamics of cognition
- Concretely?
 - Single/ multiple neuron/ regions behavior
 - Learning

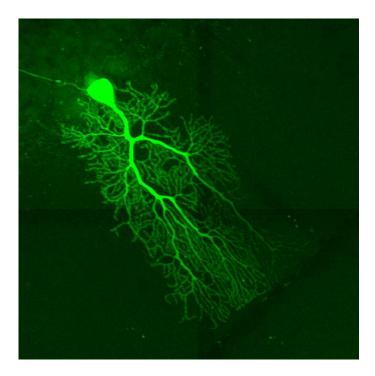
Computational Cognitive Neuroscience

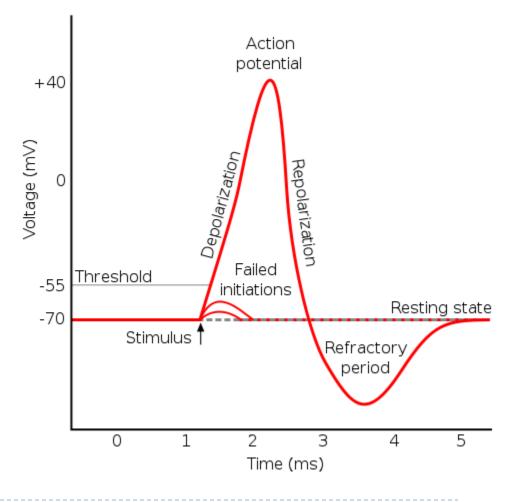
Rules

- Brain area Connections must exist
- Must fit excitatory and inhibitory discoveries
- Must obey single neuron behavior (single neuron meas.)
- Match region activity from fMRI data
- Make testable assumptions and predictions (TMS)

Singe Neuron Model Spiking

- Hodgking Huxley
- Leaky integrate and fire



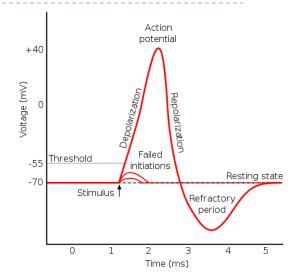


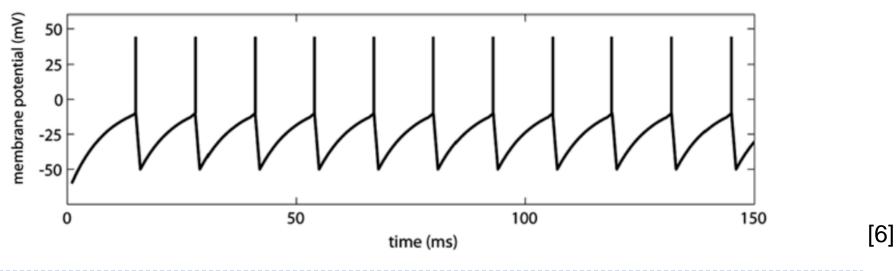
Singe Neuron Model Spiking

- Hodgking Huxley
- Leaky integrate and fire

$$\frac{dV_B(t)}{dt} = \alpha f[V_A(t)] + \beta + \gamma [V_B(t) - V_r][V_B(t) - V_t]$$

$$V_{\text{peak}} = -10 \text{ and } V_{\text{reset}} = -50.$$





Hebbian Learning

What?

- Hebb: Father of synaptic plasticity
- What fires together wires together
- Use it or loose it

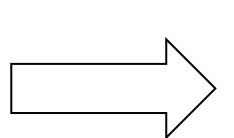
Reinforcement Learning

Why?

Obtained Reward – Predicted Reward







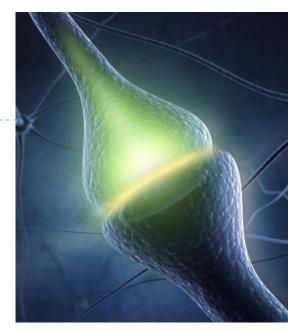




Time Model For Learning

- How?
- LTP and LTD discrete
- Synapses grow with input
- Slow dopamine uptake
- Decay with output, Cortex, Hippocampus

$$w_{A,B}(n+1) = w_{A,B}(n) + \alpha_w \int f[V_A(t)]dt \left[\int [V_B(t)]^+ dt - \theta_{NMDA} \right]^+ [w_{max} - w_{A,B}(n)] - \beta_w \int f[V_A(t)]dt \left\{ \left[\theta_{NMDA} - \int [V_B(t)]^+ dt \right]^+ - \theta_{AMPA} \right\}^+ w_{A,B}(n)$$
[6]



Time Model For Learning

- How?
- LTP and LTD
- Fast dopamine uptake, Stratium
- More like RL
- D for Dopamine

$$w_{A,B}(n+1) = w_{A,B}(n) + \alpha_w \int f[V_A(t)]dt \left[\int [V_B(t)]^+ dt - \theta_{NMDA} \right]^+ [D(n) - D_{base}]^+ [w_{max} - w_{A,B}(n)] - \beta_w \int f[V_A(t)]dt \left[\int [V_B(t)]^+ dt - \theta_{NMDA} \right]^+ [D_{base} - D(n)]^+ w_{A,B}(n) - \gamma_w \int f[V_A(t)]dt \left\{ \left[\theta_{NMDA} - \int [V_B(t)]^+ dt \right]^+ - \theta_{AMPA} \right\}^+ w_{A,B}(n)$$
[6]

Time Model For Learning

Local vs glboal

- All active synapses same update (all models)
- All synapses different update (backpropagation not in CCN, Monte Carlo Policy Gradient)

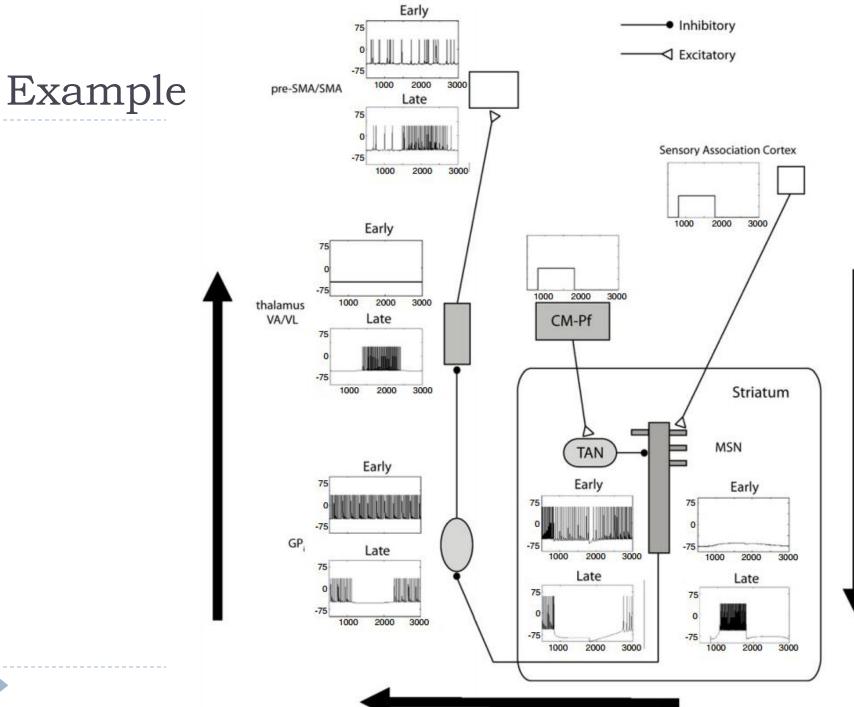
Problems of global Learning

Needs noise (otherwise equal synapses stay equal)

Stuff Left Out

D

Compartment model



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CCN vs ANNs

- No Backprop
- Global
- Spiking

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Mind of a worm

Mind of a worm

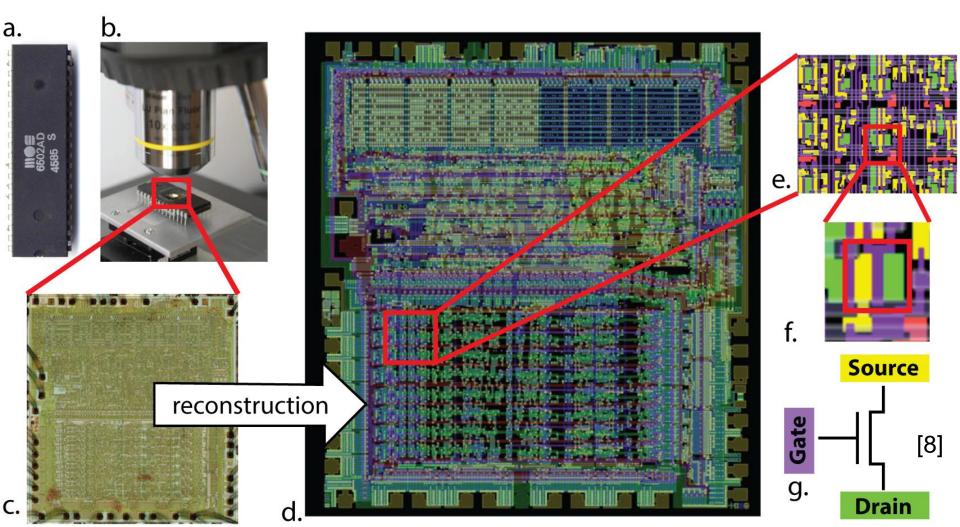
Information in the connectome reproduces behavior

Can we understand a microprocessor with the neuroscientific methods?

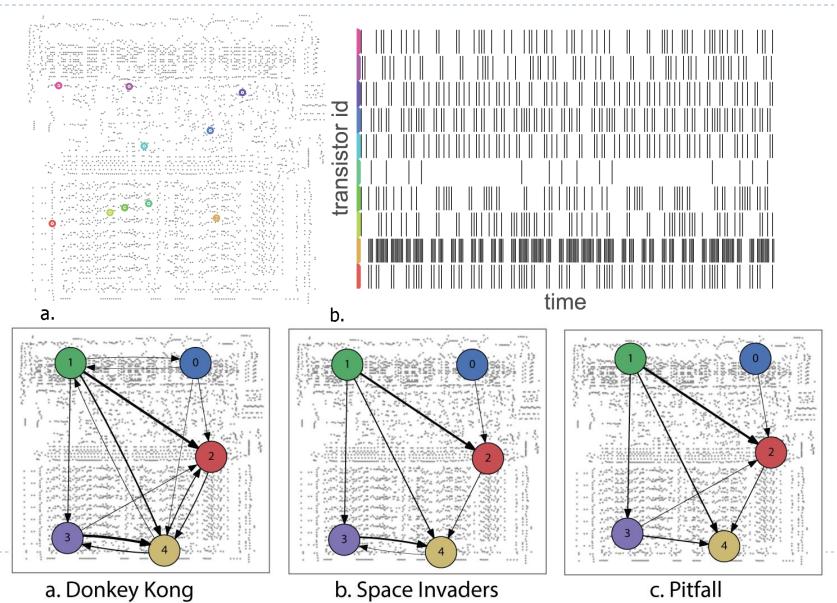
- Can we understand a microprocessor with the neuroscientific methods?
- Brain = Microprocessor?
 - Interconnections of large number of small processing units
 - Specialized modules hierarchically organized
 - Flexibly route information
 - Retain memory over time
- When do we understand something?
 - Labznick

Understanding a Mircroprocessor

MP: MOSA6502 Atari, Apple I

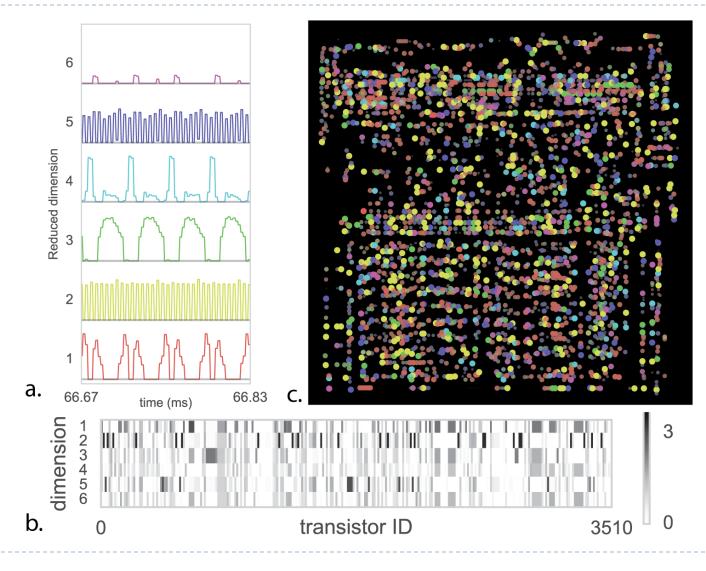


Understanding a Mircroprocessor - Spiking Patterns



[8]

Understanding a Mircroprocessor - NMF



Understanding a Mircroprocessor

We understood many behaviors but fall short understanding Miroprocessers.

Still time

- Tasks proportional to depth
- Short term memory apes

Conclusion

- Brain understanding at different levels
- Modeling of plasticity
- Methodic can be applied but not understood

References

• [1]

https://www.google.de/url?sa=i&rct=j&q=&esrc=s&source=images&cd=&cad=rja&uact =8&ved=0ahUKEwj5str2hbnUAhVGvBoKHVQKD3cQjRwIBw&url=http%3A%2F%2Fw ww.connectomics.org%2Fviewer%2F&psig=AFQjCNESoHcHPQwFPkstwD6fMDr7O4 Xz3A&ust=1497382305361033

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- [3] http://www.cellimagelibrary.org/images/40124
- [4] https://www.mpg.de/7491772/connectome-retina
- [6] Ashby, F. Gregory, and Sebastien Helie. "A tutorial on computational cognitive neuroscience: modeling the neurodynamics of cognition." *Journal of Mathematical Psychology* 55.4 (2011): 273-289.
- [7]Swanson, Larry W., and Jeff W. Lichtman. "From Cajal to connectome and beyond." Annual Review of Neuroscience 39 (2016): 197-216.
- [8] Jonas, Eric, and Konrad Paul Kording. "Could a neuroscientist understand a microprocessor?." PLOS Computational Biology 13.1 (2017): e1005268.
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