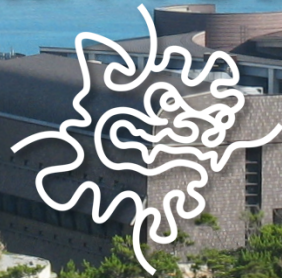


**NIPS 2015 Workshop: Modelling and Inference for
Dynamics on Complex Interaction Networks**

**Inference of Neural Circuit Connectivity
from
High-dimensional Activity Recording
Data: A Survey**



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Brain/MINDS Program (2014~2023)

- Structural/Functional maps of marmoset brain
 - RIKEN + 22 labs in Japan
 - 3 billion yen (~\$30M)/year
 - OIST: data-driven model building

brainminds.jp

Brain Mapping by Integrated Neurotechnologies for Disease Studies

[Japanese](#) [Contact](#)




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[Overview](#)

[Central Institutes](#)

[Clinical Research Group](#)

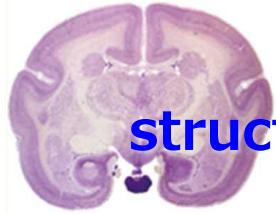
[Technology Development Group](#)



Studying the neural networks controlling higher brain functions in the marmoset, to gain new insights into information processing and diseases of the human brain

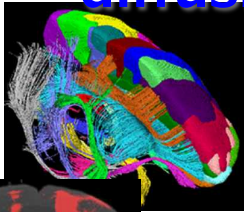
Multi-scale Neural Modeling

Structural Map

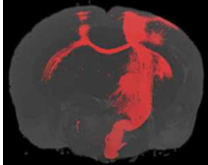


structural MRI

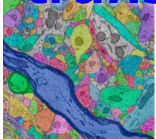
diffusion MRI



tracers



transparent brain



serial section EM

Neural Models

Macroscopic

whole brain connectome
population rate models
behaviors and cognition

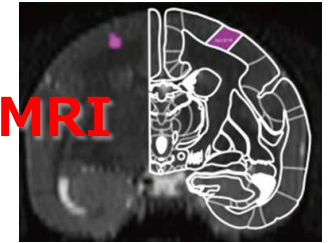
Mesoscale

cell types, synaptic dynamics
spiking neuron models
computation in each area

Microscopic

dendrites, ion channels
conductance/pathway models
cellular/molecular mechanisms

Functional Map



task fMRI

resting fMRI



ECoG

multi-electrodes



Ca²⁺ imaging

molecular imaging



Introduction

Why Neural Circuit Inference?

- To uncover dynamics/algorithms behind neural representation
 - whole brain: functional MRI, MEG, EEG,...
 - local circuit: multi-electrode, optical recording
- Anatomical identification of all connections is costly/impractical/imprecise

This Survey: Local Circuit Inference

- Data processing pipeline
- Biophysical and technical challenges



Data Processing Pipeline



Calcium imaging

- 1000s of neurons
- Good time resolution
- Complex pre-processing

Multiple electrodes

- Very good time resolution
- Few 100s
- Mixed signals
- Highly invasive

Calcium imaging

- Neuron image segmentation
- Fluorescence trace extraction
- Spike train inference

Multiple electrodes

- Sorting required

Model-free:

- Correlation
- Transfer entropy
- Deep learning
- ...

Model-based:

- Generalized linear models
- Max. entropy
- Hawkes processes
- ...

- Confounded effects
- Matrixes combination
- Edge orientation

Simulated vs real data



Model-free Approaches

■ Descriptive statistics

- correlation

$$\rho_{X,Y} = \text{corr}(X, Y) = \frac{\text{cov}(X, Y)}{\sigma_x \sigma_y} = \frac{E[(X - \mu_x)(Y - \mu_y)]}{\sigma_x \sigma_y}$$

- cross correlation

$$\rho_{Y \rightarrow X}(\tau) = \frac{E[(Y_t - \mu_y)(X_{t+\tau} - \mu_x)]}{\sigma_x \sigma_y}$$

- coincidence index

$$CI = \frac{\sum_{d=0}^r \rho_{X,Y}(d)}{\sum_{d=0}^T \rho_{X,Y}(d)}$$

- partial correlation

$$PC_{i,j} = \frac{\sum_{ij}^{-1}}{\sqrt{\sum_{ii}^{-1} \sum_{jj}^{-1}}}$$

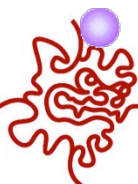
■ Information theoretic methods

- transfer entropy

$$TE_{Y \rightarrow X} = \sum_{x_{t+1}, x_t^k, y_t^l} P(x_{t+1}, x_t^k, y_t^l) \log_2 \frac{P(x_{t+1} | x_t^k, y_t^l)}{P(x_{t+1} | x_t^k)}$$

- information gain

$$IG_{Y \rightarrow X} = G(x_{t+1}) - G(x_{t+1} | y_t)$$

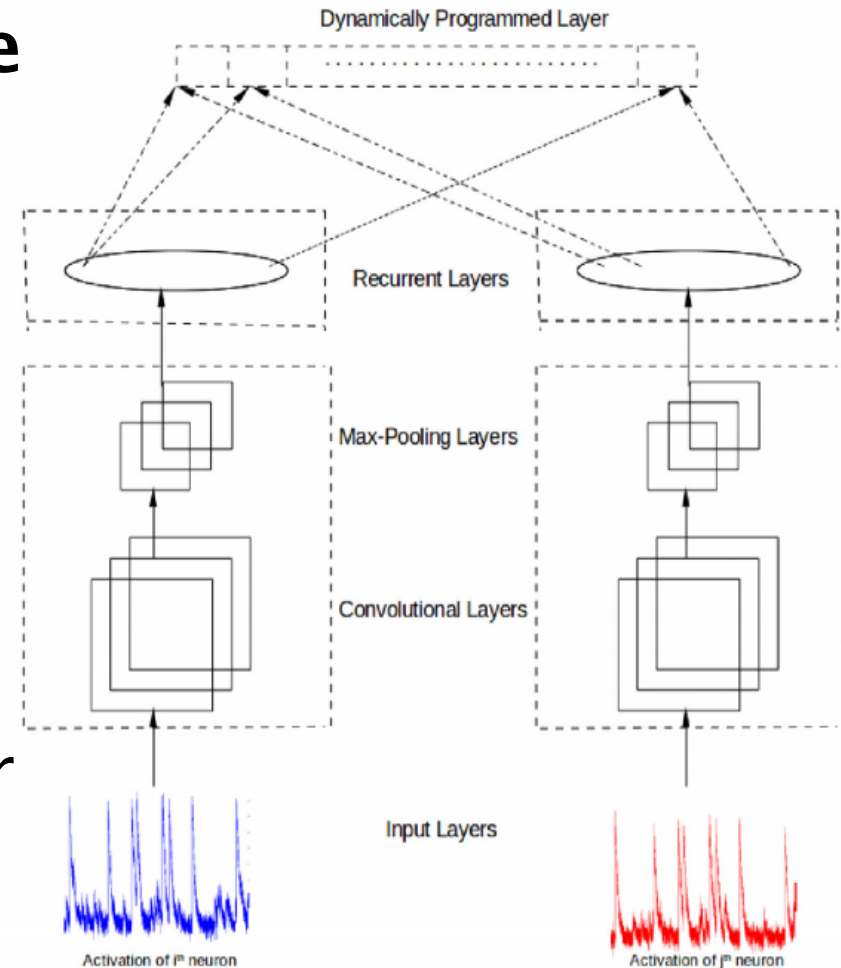


Deep Learning Approach

(Veeriah et al. 2015)

Integrated architecture for spike and connection inference

- Convolutional neural network
 - identify relevant events in fluorescence time series
 - with temporal tolerance
- Recurrent Neural Network
 - model temporal sequences of relevant events
- Dynamically programmed layer
 - compute connection probability



Model-based Approaches

Generative models

- auto-regressive (AR)
$$x_i(t) = A_{i0} + \sum_{j=1}^P \sum_{k=1}^K A_{ij}(k)x_j(t-k) + \epsilon_i(t),$$

- generalized linear (GLM)
$$x_i(t) \sim \text{Ber}(\cdot | \rho_i(t))$$

$$\rho_i(t) \equiv \phi \left(A_{i0} + \sum_{j=1}^P \sum_{k=1}^K A_{ij}(k)x_j(t-k) \right)$$

e.g. stochastic integrate & fire

Inference

- maximum likelihood
$$\theta^* = \underset{\theta}{\operatorname{argmin}} J(\theta|D)$$

- Bayesian
$$\theta^* = \underset{\theta}{\operatorname{argmin}} \{J(\theta|D) + \lambda \mathcal{R}(\theta)\}$$

regularizers

$$\mathcal{R}(\theta) = \sum_r \theta_r^2 \quad \mathcal{R}(\theta) = \sum_r |\theta_r|$$



Neural and Ca²⁺ Dynamics Model

(Mishchenko et al. 2011; Fletcher et al 2014)

Generative model

- stochastic IF

$$\tilde{v}_i^{k+1} = (1 - \alpha_{IF})v_i^k + \sum_{j=1}^N W_{ij}s_j^{k-\delta} + d_{v_i}^k$$

- calcium dynamics

$$(v_i^{k+1}, s_i^{k+1}) = \begin{cases} (\tilde{v}_i^k, 0) & \text{if } v_i^k < \mu \\ (0, 1) & \text{if } \tilde{v}_i^k \geq \mu \end{cases}$$

- fluorescence

$$z_i^{k+1} = (1 - \alpha_{CA,i})z_i^k + s_i^k$$
$$y_i^k = a_{CA,i}z_i^k + b_{CA,i} + d_{y_i}^k$$

Inference

$$\hat{\theta} = \arg \max_{\theta} L(\mathbf{y}|\theta) + \lambda \|\mathbf{W}\|_1$$

- parameters $\theta = \{\mathbf{W}, \tau_{IF}, \tau_{CA}, \alpha_{IF}, b_{IF,i}, \alpha_{CA}, a_{CA,i}, b_{CA,i}, i = 1, \dots, N\}$

- hidden variables $\mathbf{x} = \{\mathbf{v}, \mathbf{z}, \mathbf{q}, \mathbf{s}\}$

- E-step:

$$P(\mathbf{x}|\mathbf{y}, \hat{\theta}^\ell)$$

- M-step:

$$\hat{\theta}^{\ell+1} = \arg \min_{\theta} \mathbb{E} \left[L(\mathbf{x}, \mathbf{y}|\theta) | \hat{\theta}^\ell \right] + \phi(\theta)$$



Biophysical Challenges

- **Apparent/invisible connection:**
 - common inputs cause apparent connections
 - connection is invisible without neural activity
- **Directionality:** with limited temporal resolution
- **Cellular dynamics:** post-spike adaptation, post-inhibitory rebound, etc.
- **Synaptic dynamics:** short-term adaptation, long-term plasticity, etc.
- **Non-stationarity:** network can shift between multiple states, such as synchronized bursting



Technical Challenges

- **Noise:** electric noise, motion artifacts, etc.
- **Time/space resolution:** calcium image $\sim 100\text{ms}$
- **Hidden neurons:** e.g., off the focal plane
- **External inputs:** e.g. spontaneous action
- **Prior knowledge:** sparseness, topography, etc.
- **Accuracy:** existence, sign, magnitude, time delay
- **Scalability:** thousands of neurons,
millions of potential connections...



Challenges and Solutions

	Data Acquisition	Pre-processing	Connection Inference	Post-inference	Validation
Apparent connection			c1 d12 e12	a3	
Directionality			b123 c1 d12 e12	a2	
Cellular dynamics			c1 d12 e12		
Synaptic dynamics			d12 e12		
Non-stationarity		a123 b2	c1		
Noise		a123 b1 c1	c1 d12 e12		
Time/space resolution			c1 e12		
Hidden neurons					
External inputs					
Prior knowledge			a2 d12 e12		
Accuracy			a23 c1 d12 e12	a3	d1
Scalability			a23		



References

Model-free

- Descriptive statistics
 - a1: Cohen & Kohn, Nat Neurosci, 2011
 - a2: Suter et al., ECML, 2014
 - a3: Magrans & Nowe, ECML, 2014
- Information theoretic
 - b1: Stetter et al., PLoS Comp Bio, 2012
 - b2: Garofalo et al., PLoS One, 2009
 - b3: Ito et al., PLoS One, 2011
- Others
 - c1: Veeriah et al., SIGKDD, 2015

Model-based

- Generalized linear models
 - d1: Pillow et al., Nature, 2008
 - d2: Stevenson et al., IEEE TNSR, 2009
- Spike & calcium dynamics
 - e1: Mishchenko et al., Ann Appl Stat, 2011
 - e2: Fletcher et al., NIPS 2014



Conclusion

■ Model-free methods

- less assumptions, less parameters
- cannot reproduce dynamics

■ Model-based methods

- needs hyper parameter tuning
- reproduce dynamics for validation/analysis

■ Limitations and opportunities

- external inputs and hidden neurons
- non-stationarity
- cellular/synaptic diversity and scalability

