
A deep learning approach to infer connectivity and neuronal dynamics from spike trains

Machine Learning and Optimization Seminar
02/23/2023

Rodrigo FO Pena

e-mail: pena@njit.edu

BioDatanamics Lab (H. G. Rotstein), Federated Department of Biological Sciences

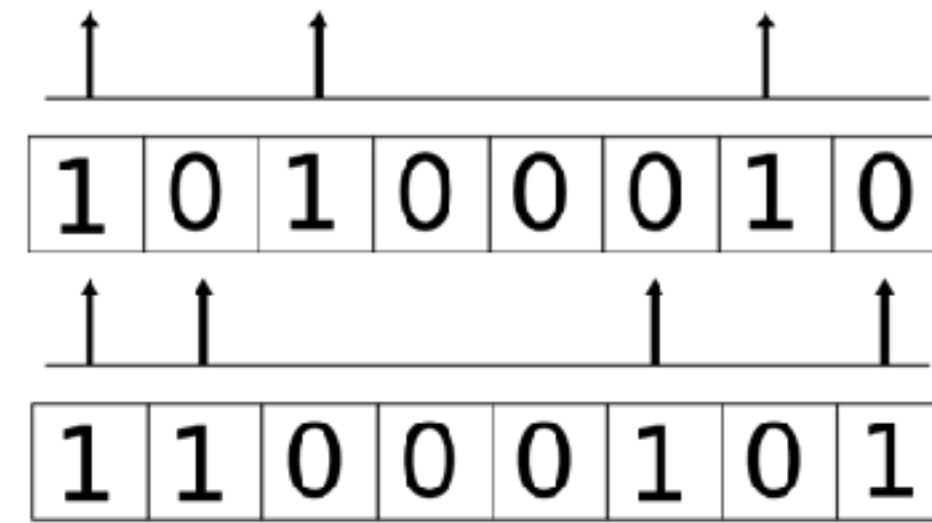
New Jersey Institute of Technology & Rutgers University



Spike-train cross-correlation function (CCF)

$$x(t) = \sum_i \delta(t - t_i^f)$$

$$y(t) = \sum_j \delta(t - t_j^f)$$



$$\rightarrow c_{xy}(\tau) = \langle x(t)y(t + \tau) \rangle - \langle x(t) \rangle \langle y(t + \tau) \rangle,$$

where $\langle . \rangle$ is an ensemble average.

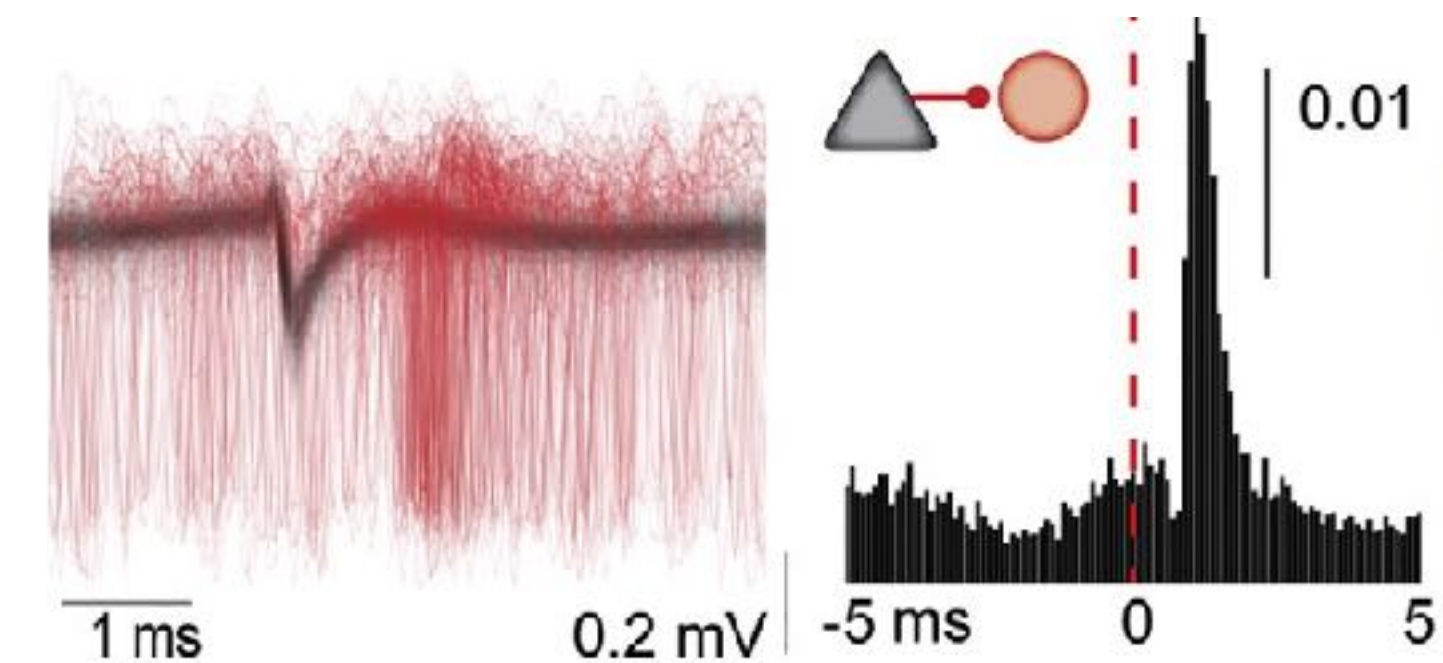
- How spike-trains are temporarily related;
- Provide information to infer connectivity;

Spike-train cross-correlation functions (CCF) are very informative

- ▶ A sharp peak within a few milliseconds in the CCF indicates the presence of a connection.

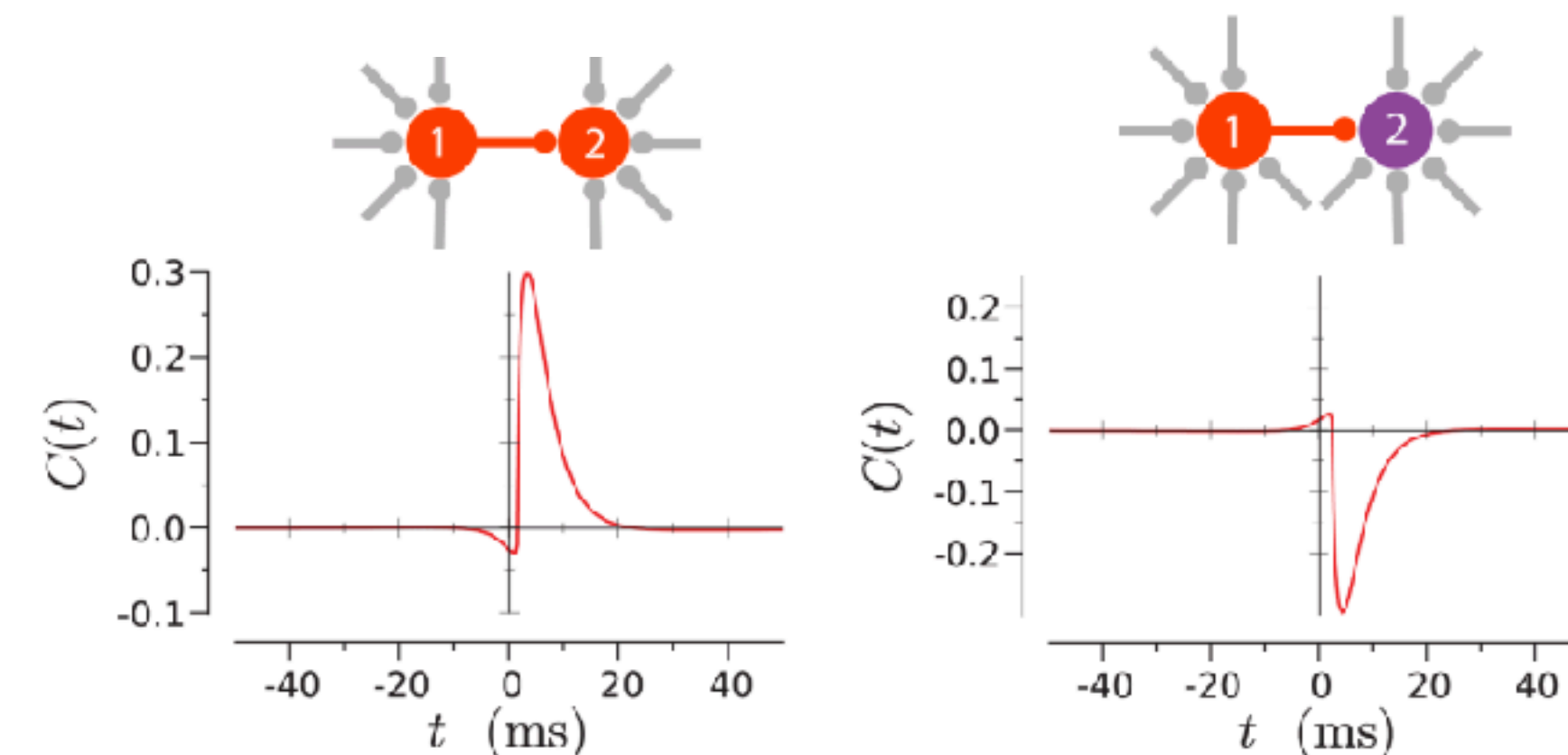
in vivo multi-electrode extracellular recordings dorsal CA1 pyramidal layer

Cross-correlation



English et al., *Neuron* 96: 505–520 (2017)

- ▶ Synaptic properties can be observed in the CCF.

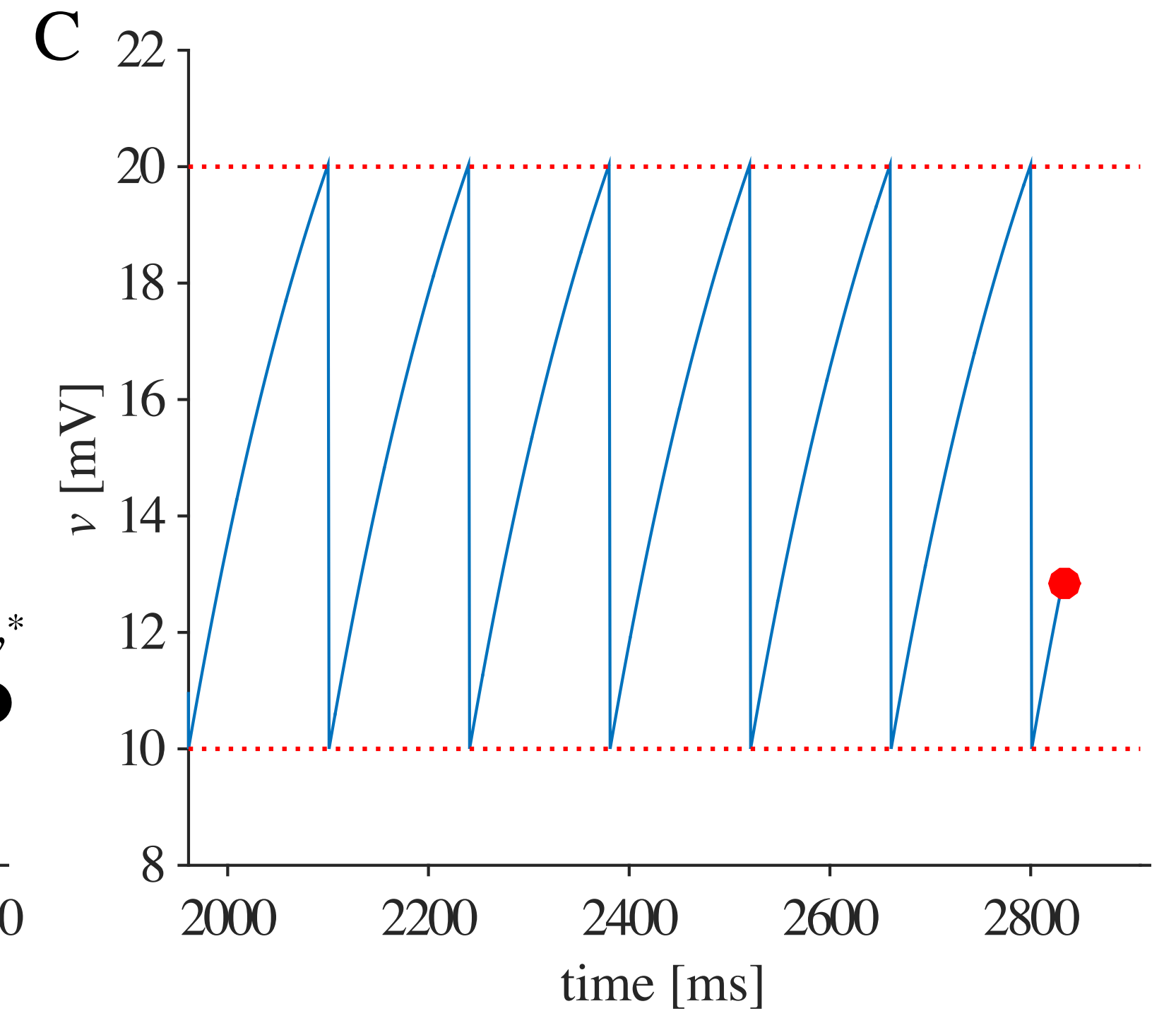
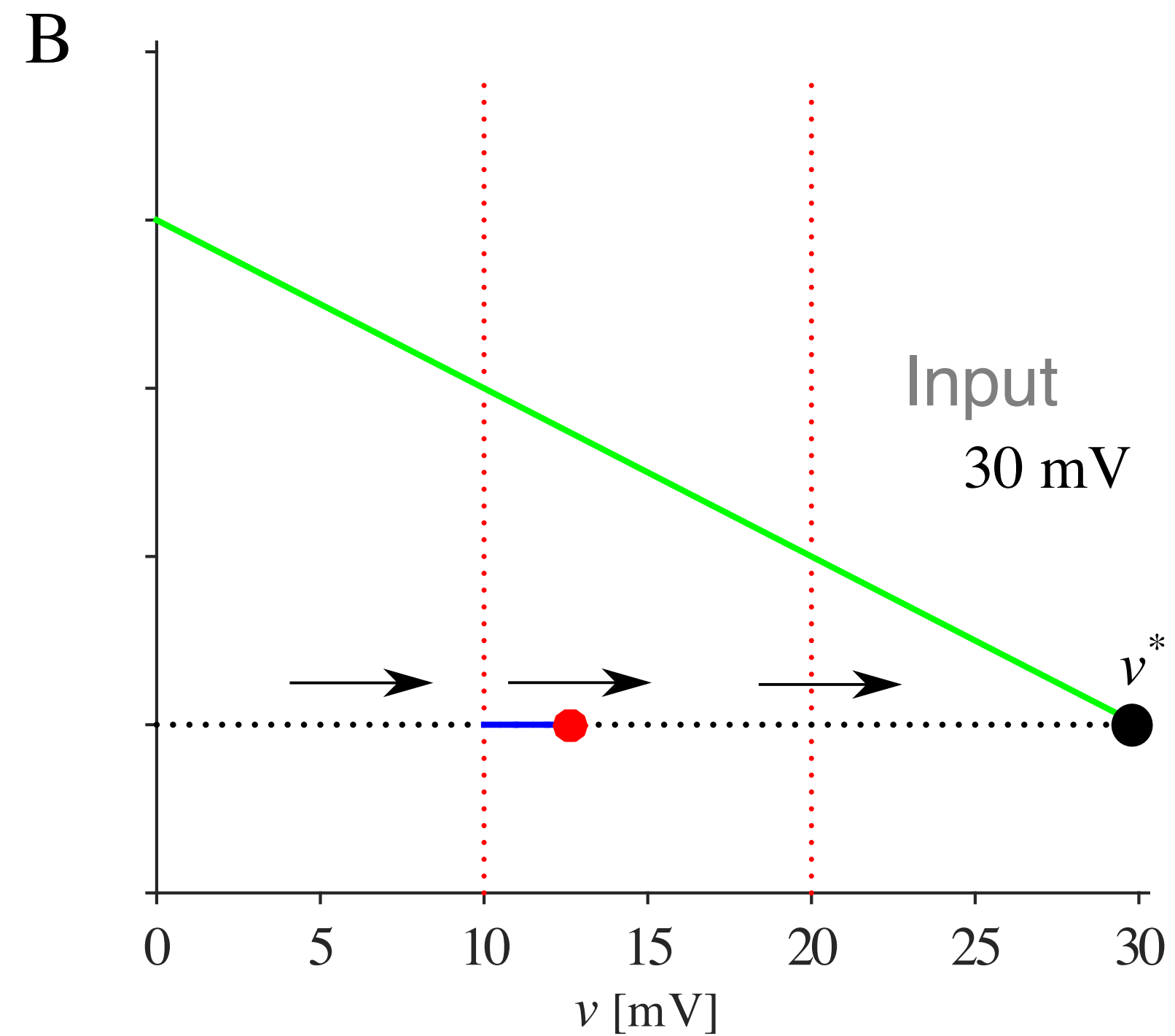
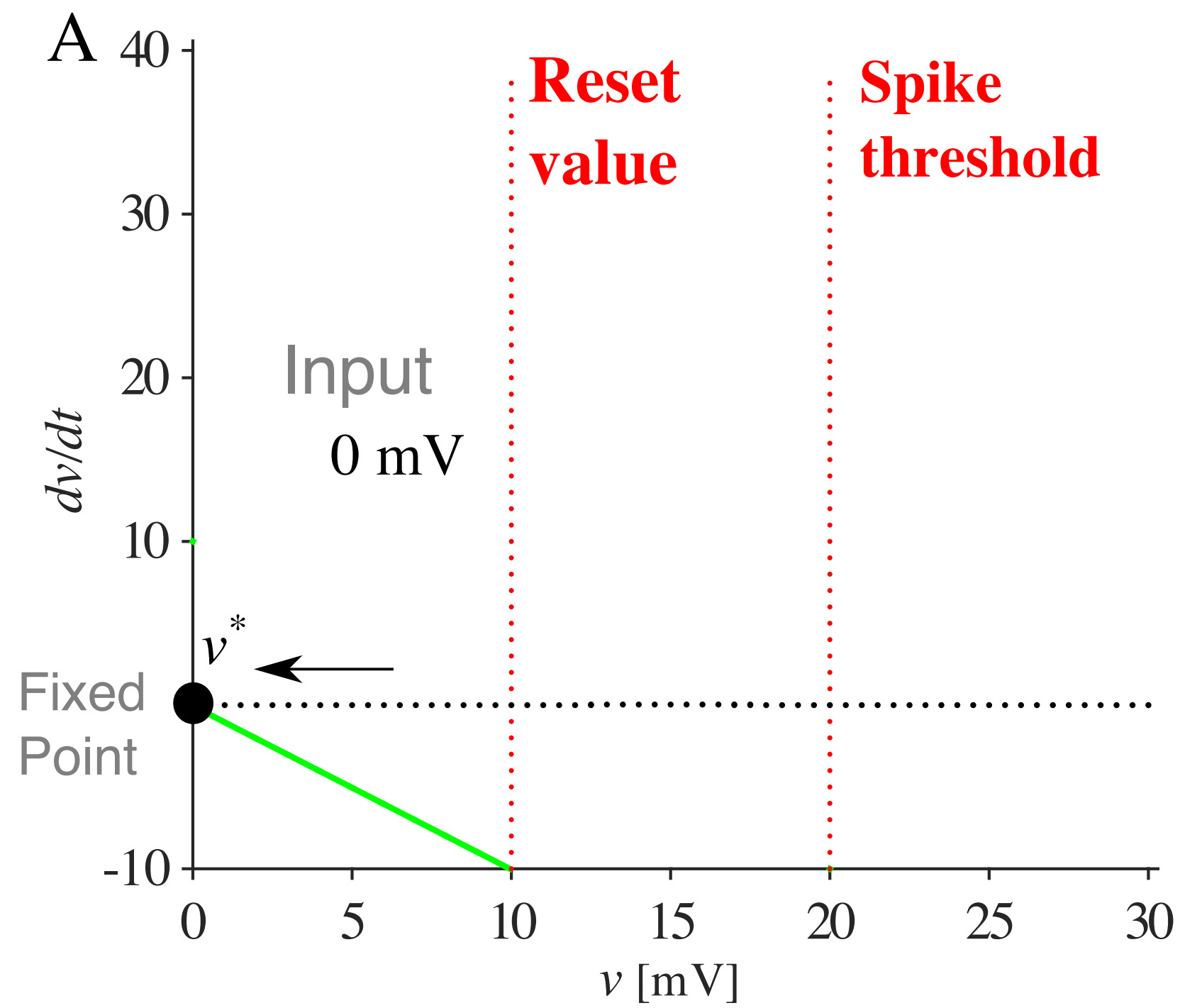


Ostojic et al., *J. Neurosci.* 29: 10234–10253 (2009)

Neurons have a very rich individual neuronal dynamics.
Can CCF say something about **intrinsic dynamics**?

Dynamical systems

$$\tau_m \dot{v} = f(v) + \text{input} + \text{update rule}$$



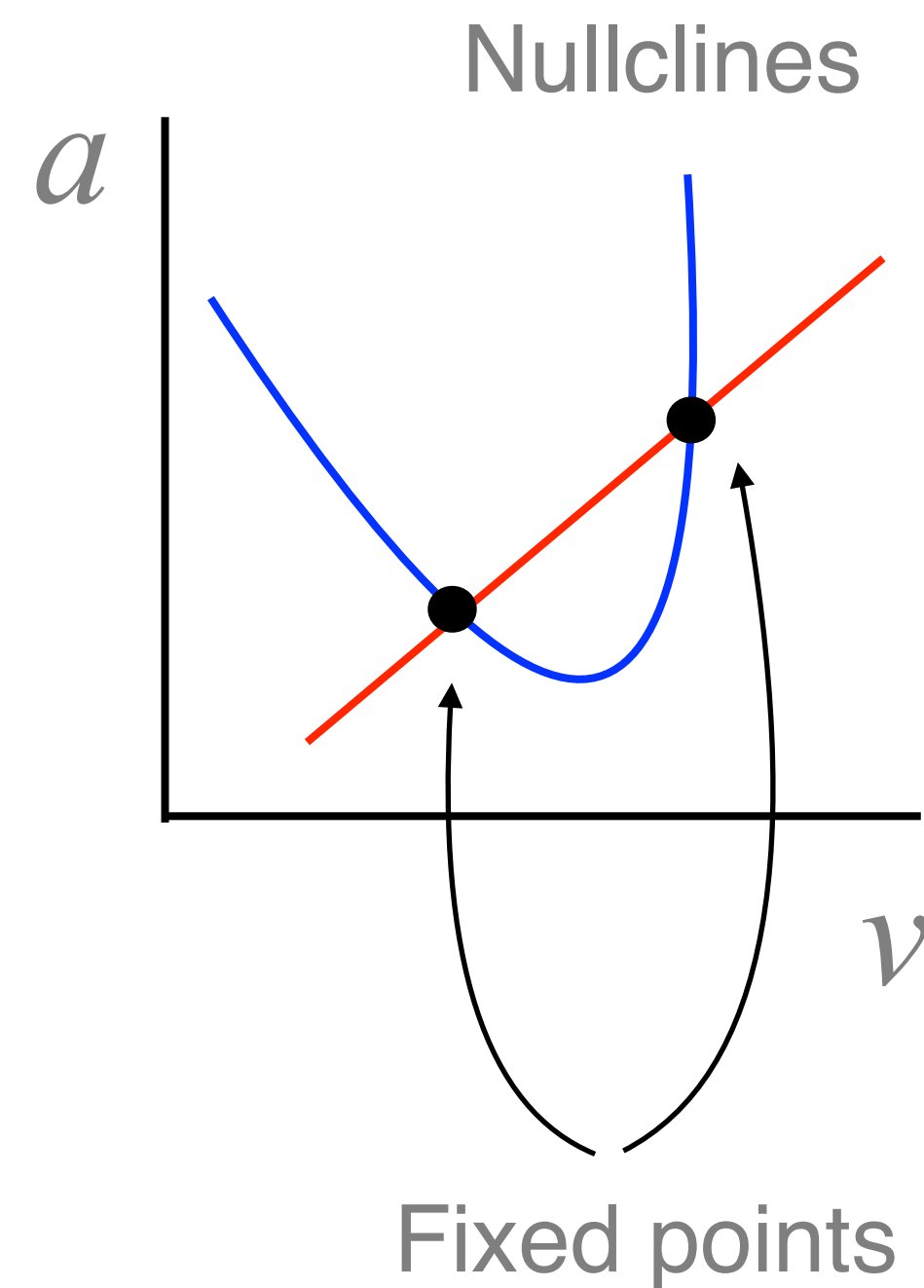
Nullcline!

$$\tau_m \dot{v} = -v + \text{input} + \text{update rule} = 0$$

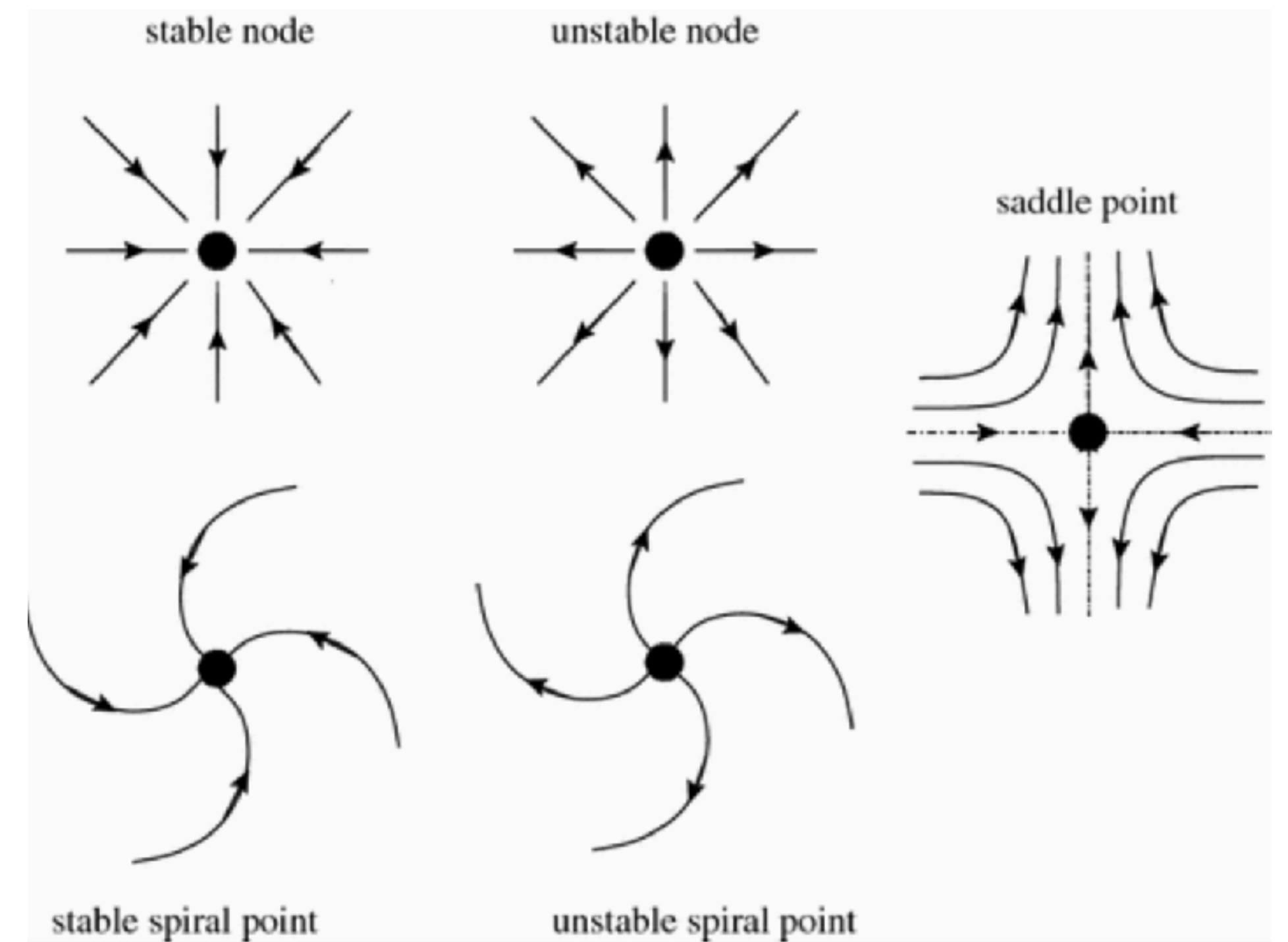
Dynamical systems

$$\tau_v \dot{v} = v^2 - a + RI = 0$$

$$\tau_a \dot{a} = a + v = 0$$



Fixed points can attract or repel trajectories!

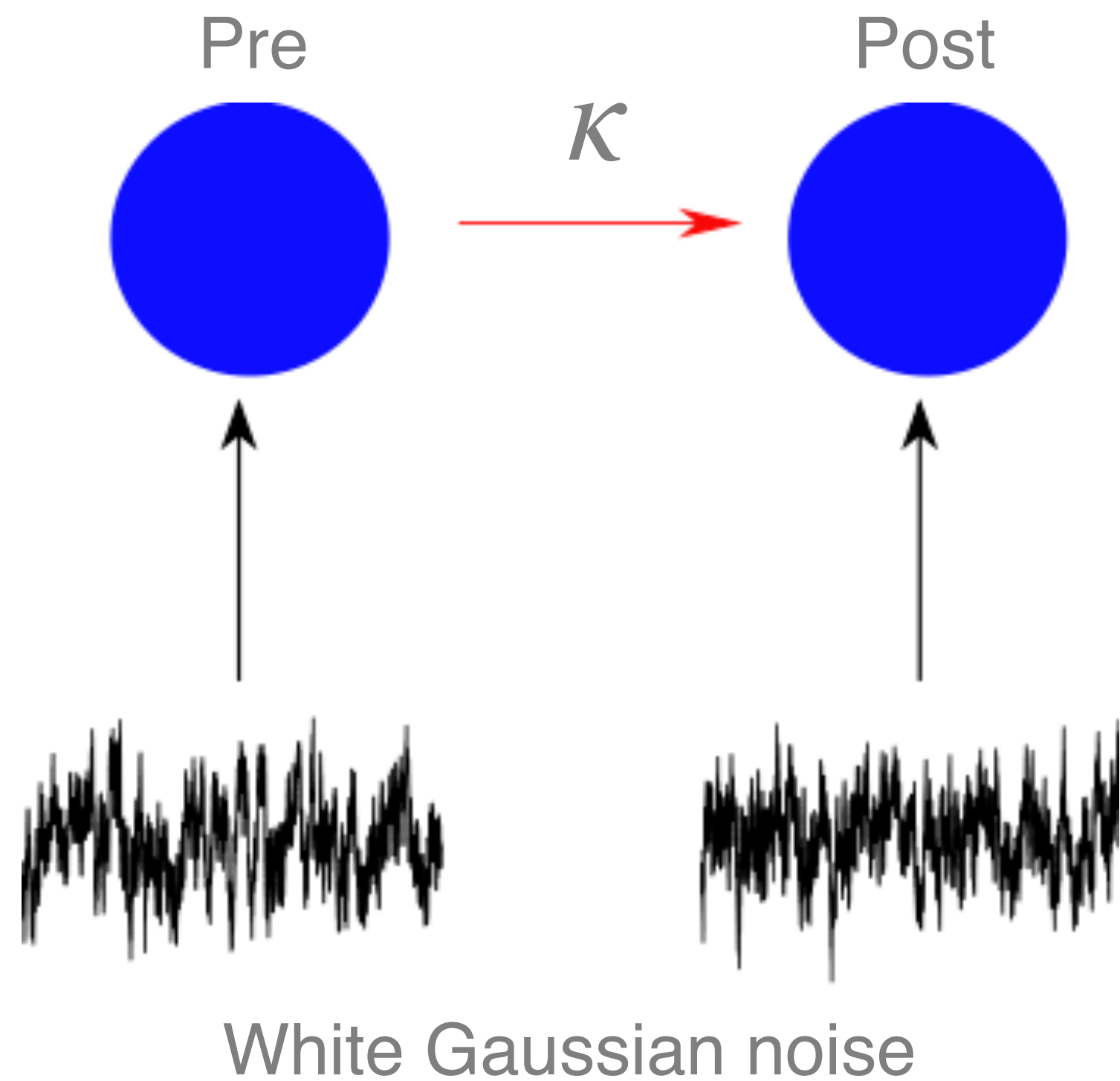


Stauffer D., Stanley H.E., Lesne A. (2017)

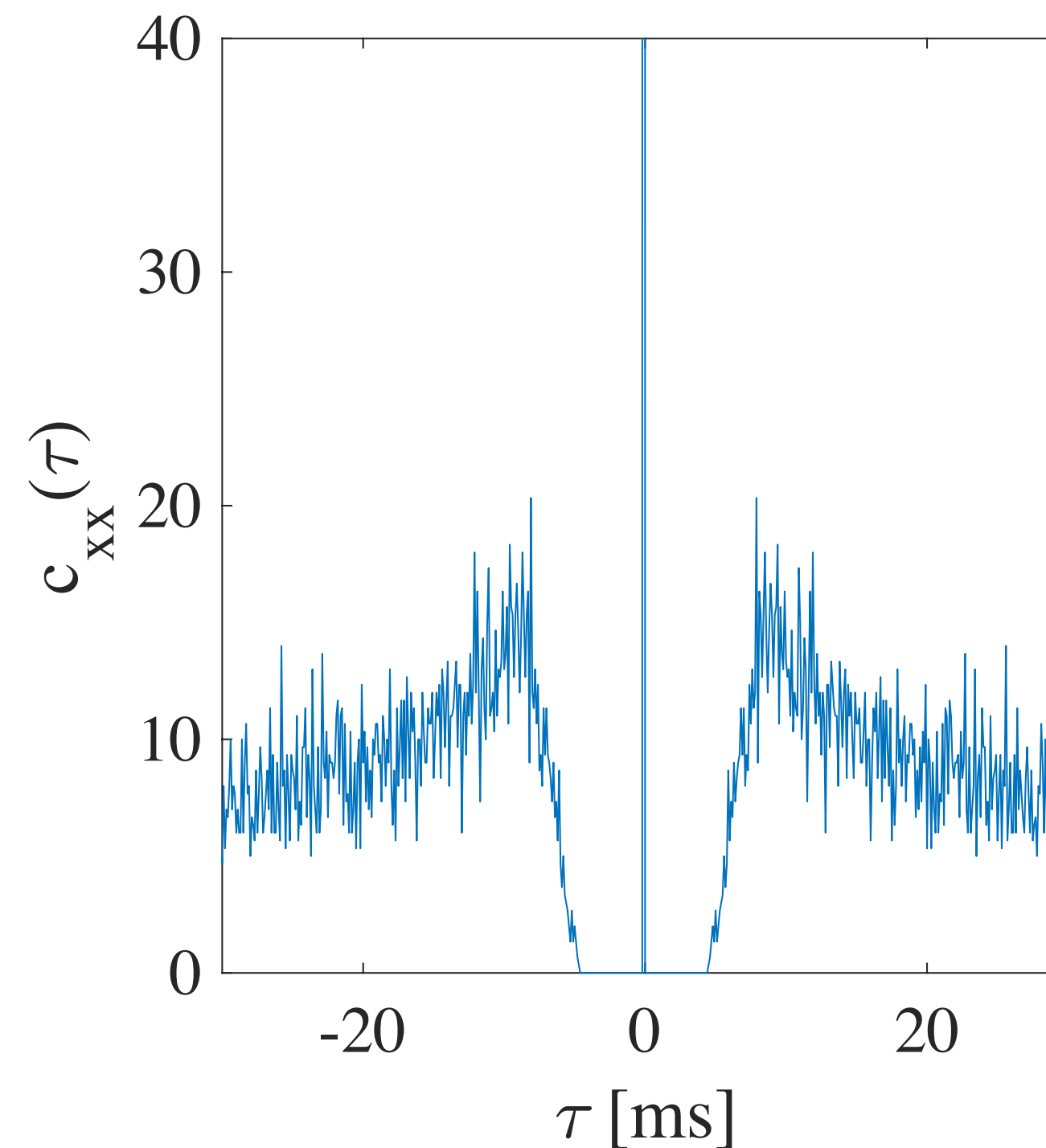
Preliminaries

- ▶ Two neurons are coupled in an ultra-precise monosynaptic model:

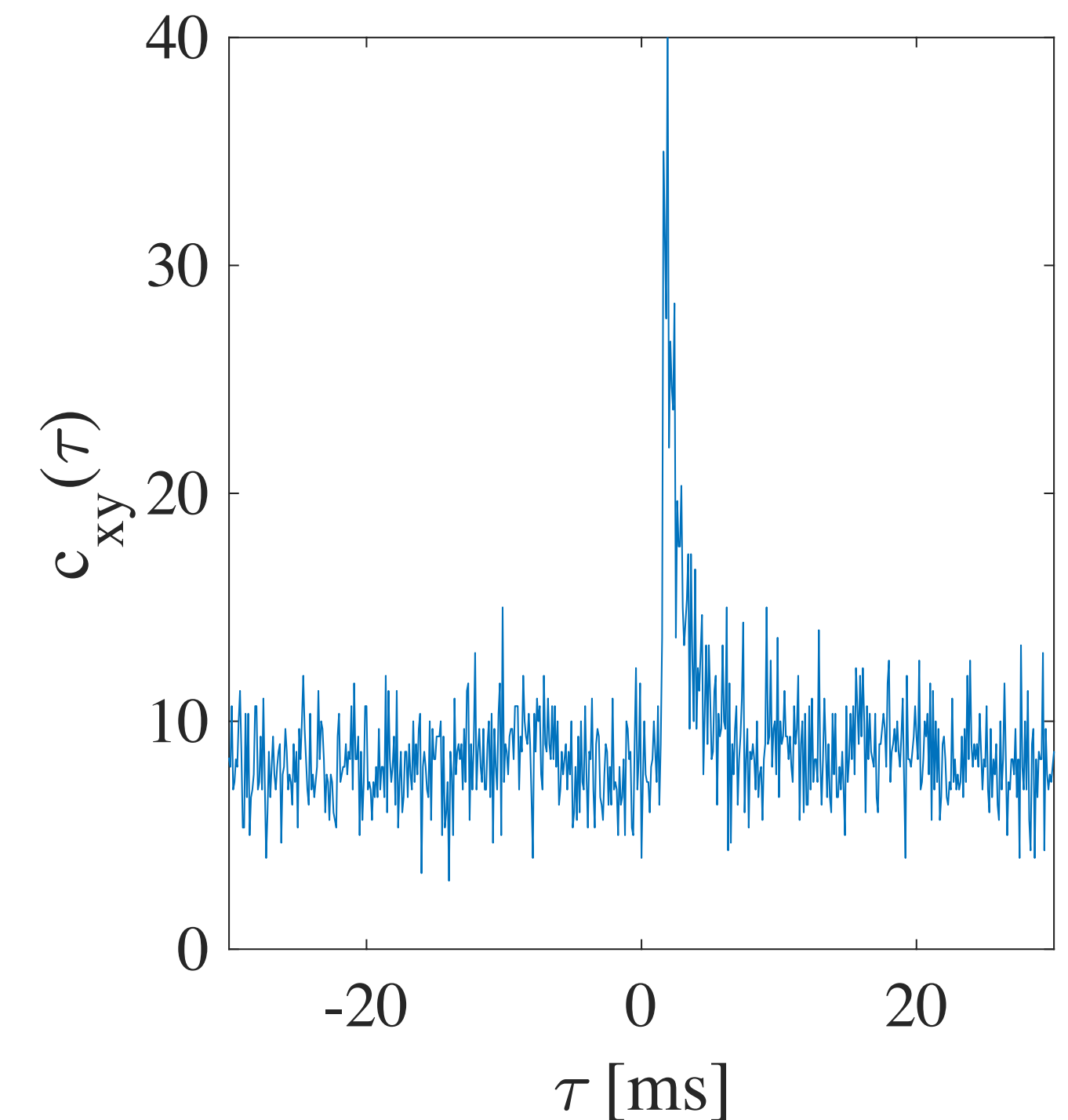
Platkiewicz *et al.* *J Comput Neurosci* **49**, 131–157 (2021)



Autocorrelation



Cross-correlation



Are they model dependent?

- ▶ We will be working with models where nonlinearities $f(v)$ change:

$$C\dot{v} = f(v) + I$$

How these mechanisms connect to biophysics

Depending on parameters, ionic currents can flexibly create multiple nonlinearities which are reflected on c_{xy}

$I_{NaP}+I_h$ model

$$C \frac{dV}{dt} = -I_L - I_h - I_{Nap} + I_{app} + I_{in}(t)$$

$$I_L = G_L (V - E_L)$$

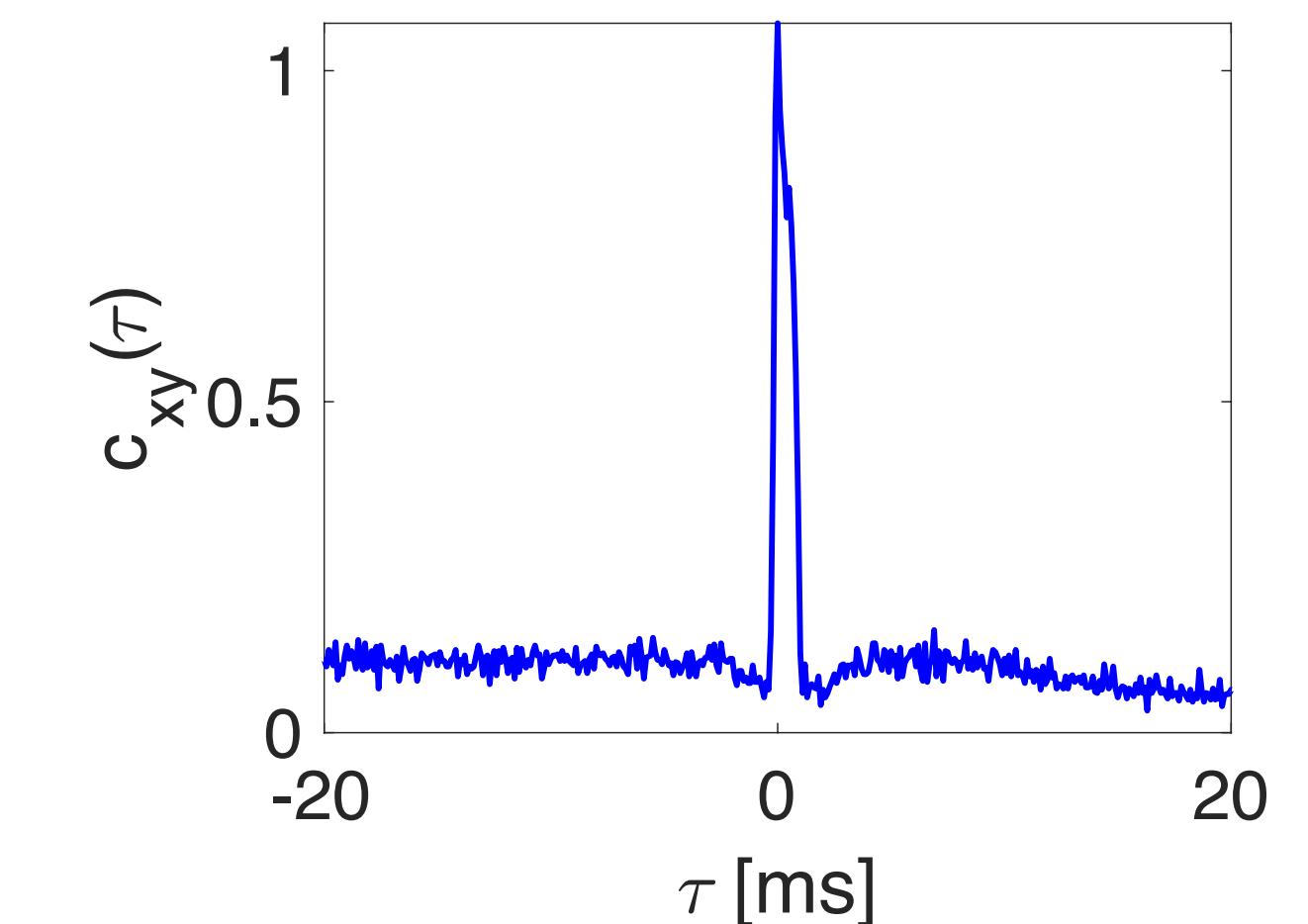
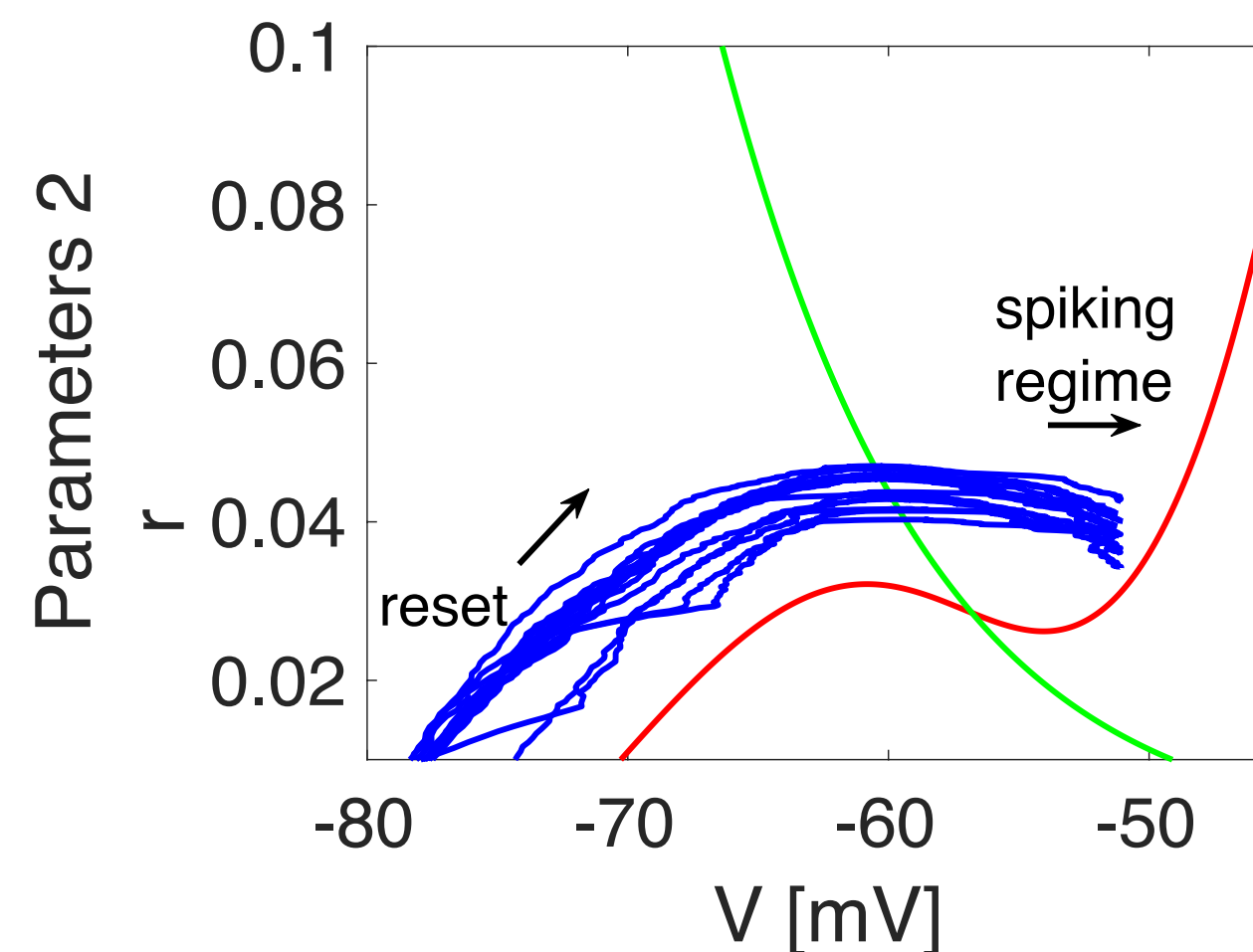
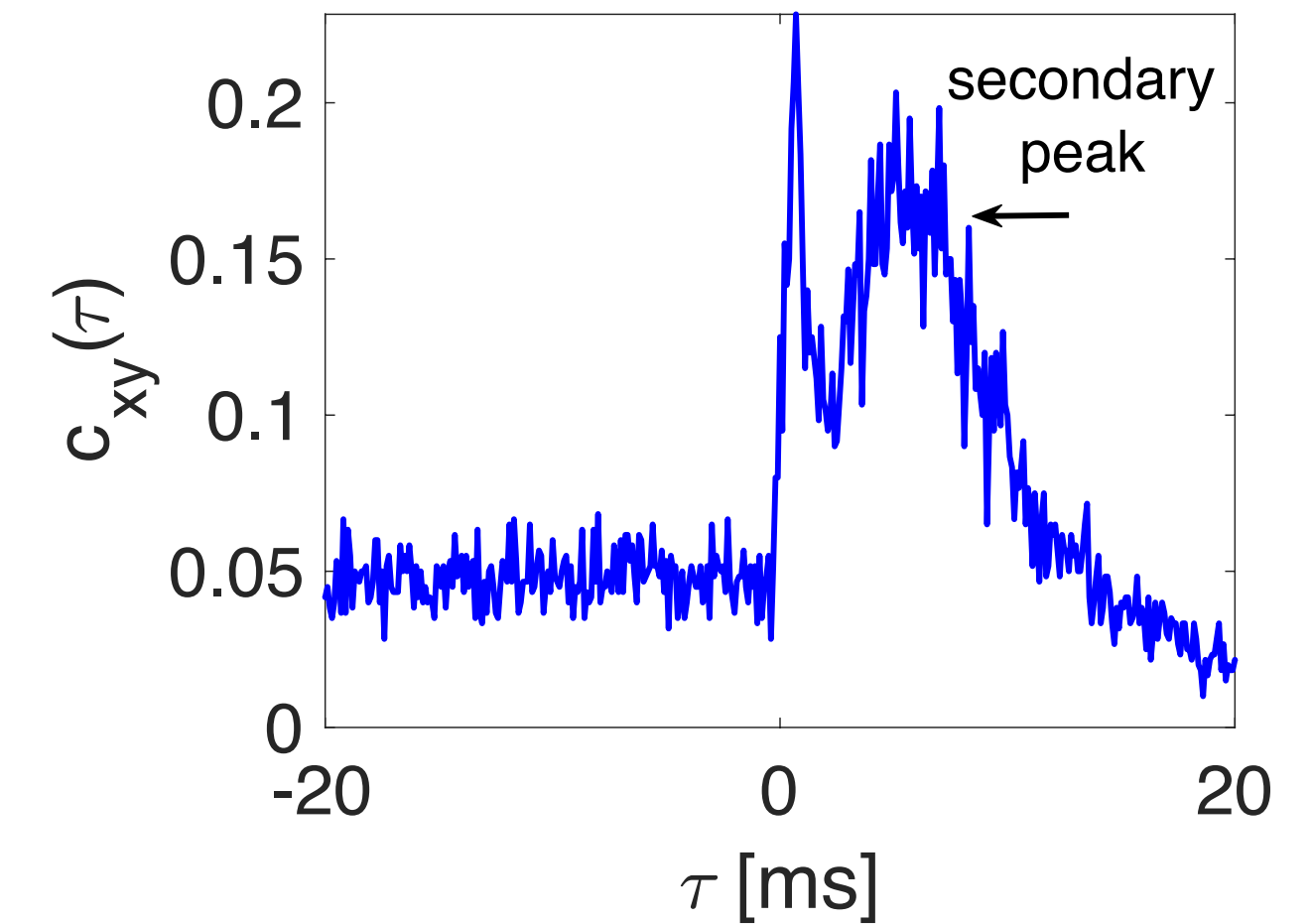
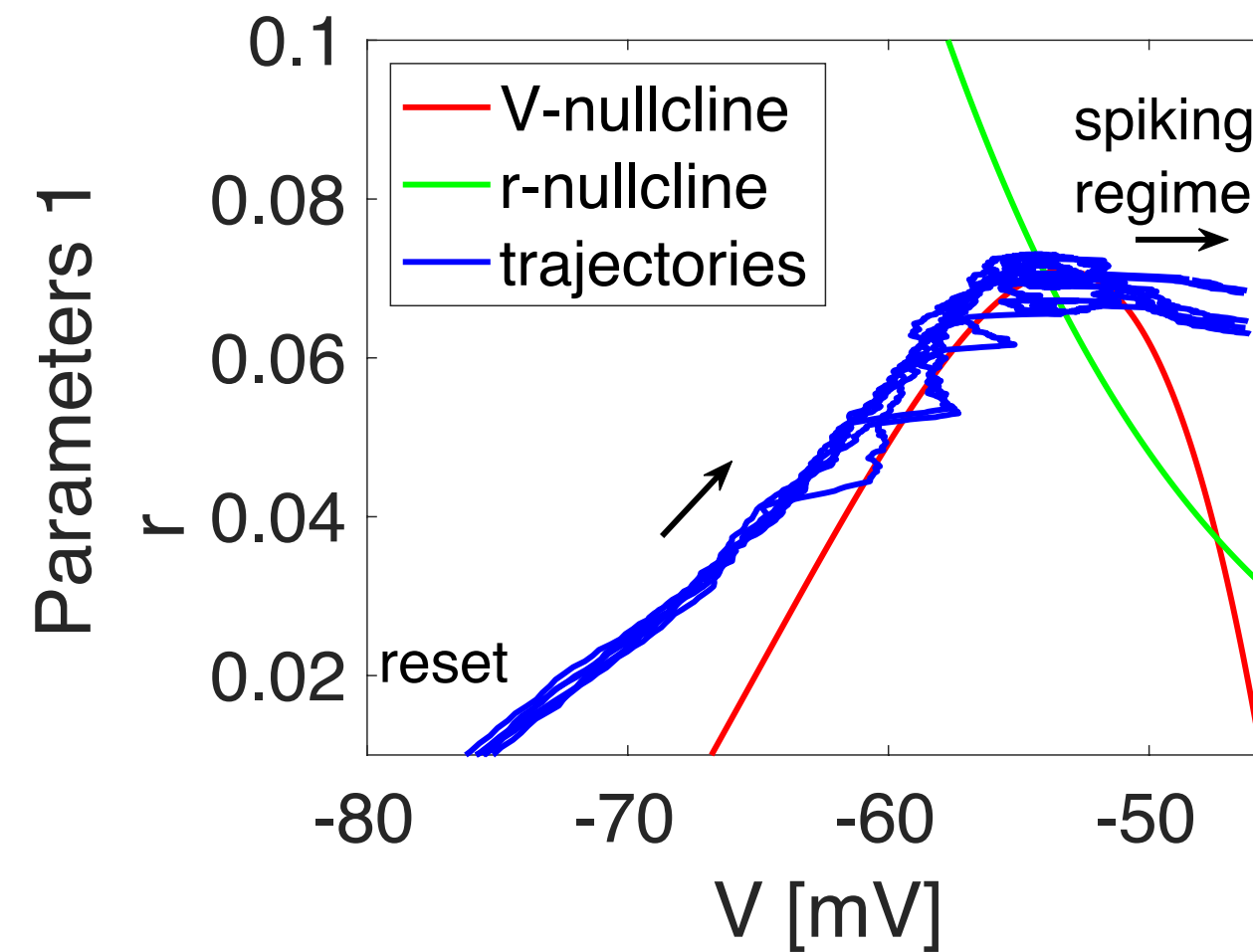
$$I_h = G_h r (V - E_h)$$

$$I_{Nap} = G_p p_\infty(V) (V - E_{Na})$$

$$x (= r, p)$$

$$\frac{dx}{dt} = \frac{x_\infty(V) - x}{\tau_x(V)}$$

Rotstein H.G., *J. Comput. Neurosci.* 43: 243–271 (2017)

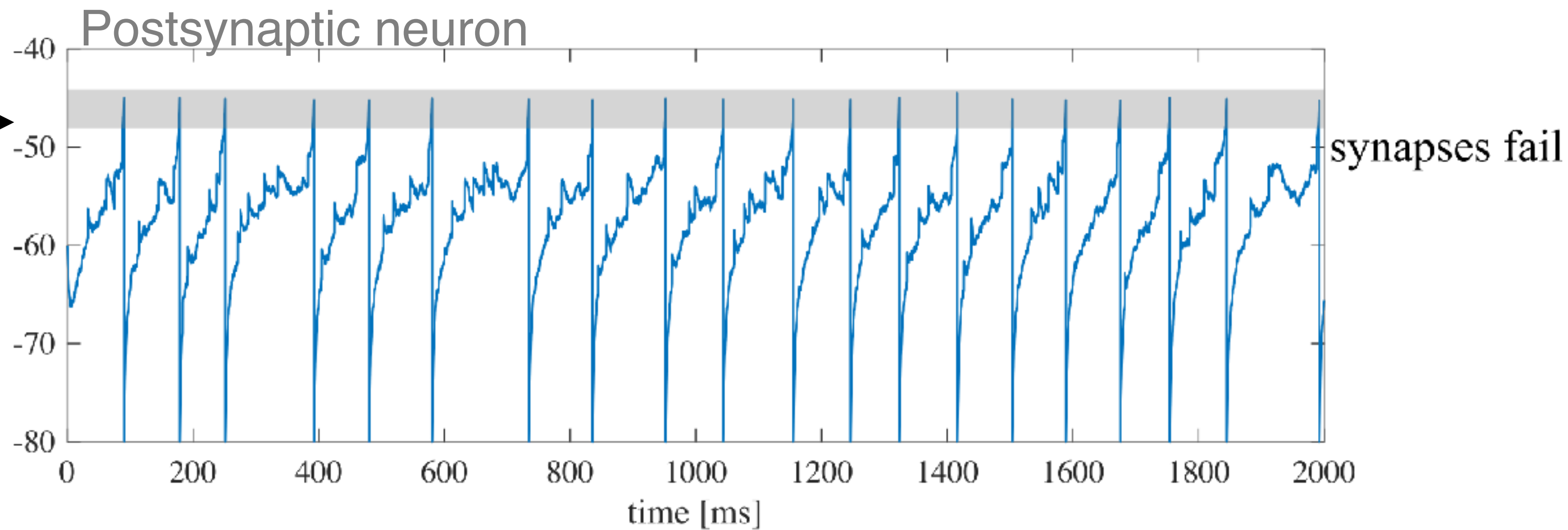


Subthreshold vs. near-action potential voltages

membrane potential at near-action potential voltages

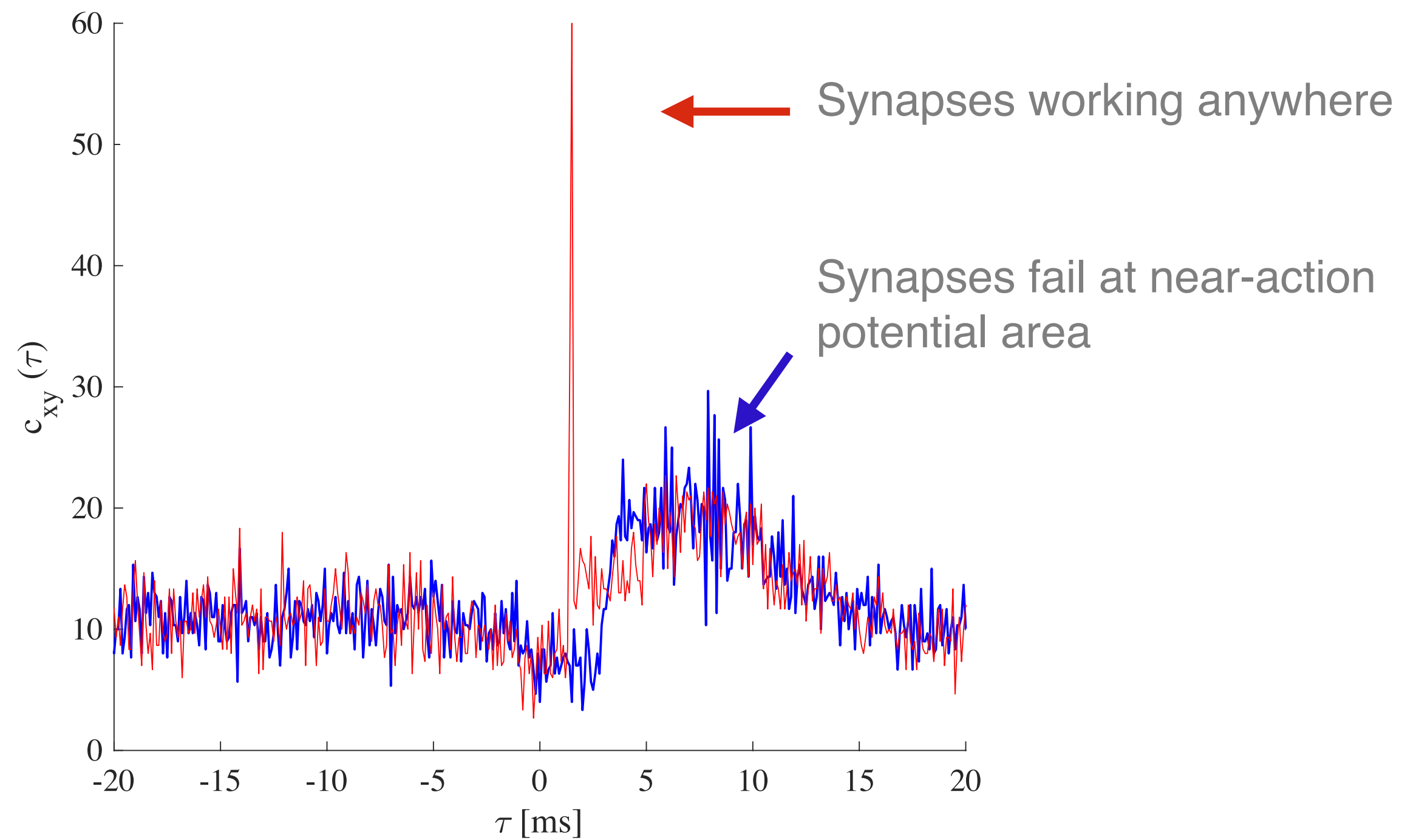


membrane potential at subthreshold voltages

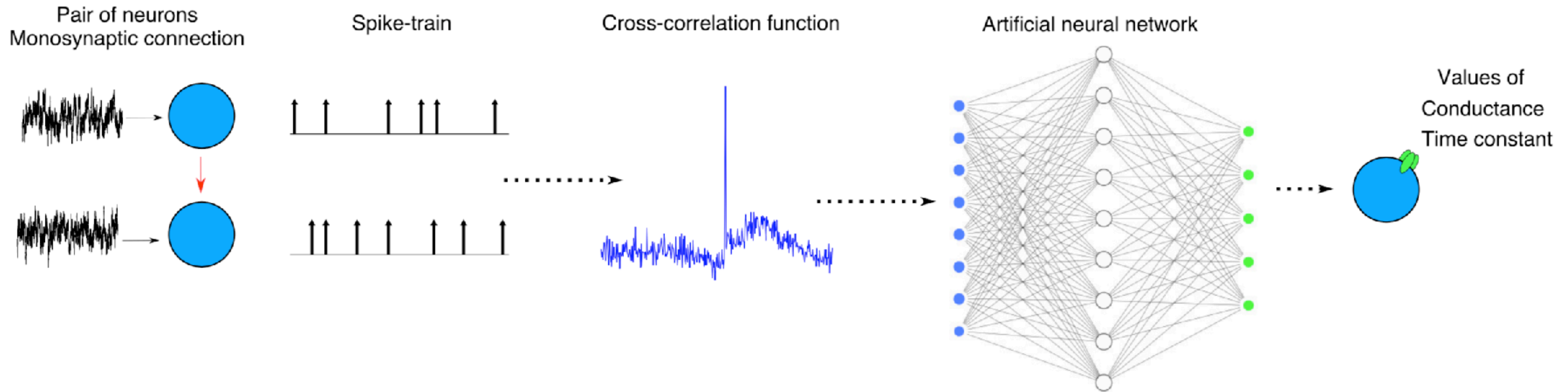


Removing events arriving at near-action potential voltages:

Only second peak shows up!

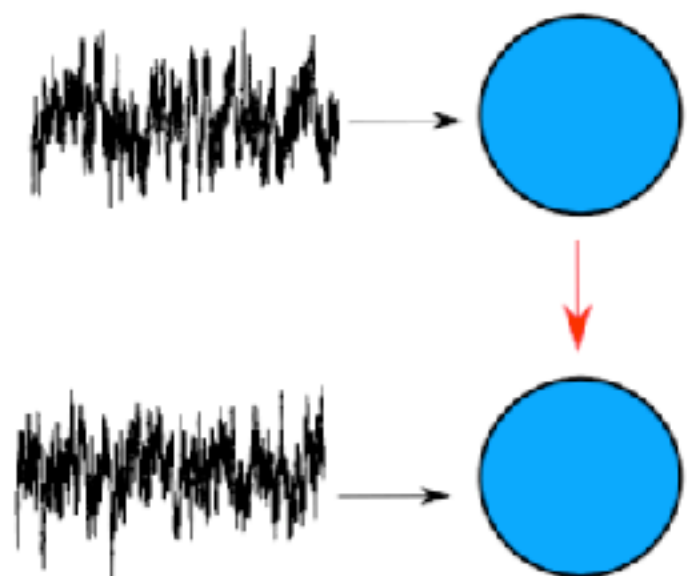


Can we use these signatures?

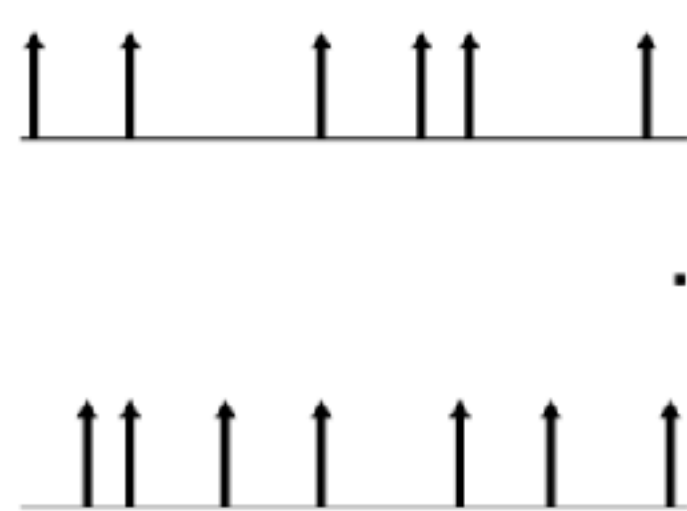


to what extent can **CCFs** be used to capture **intrinsic neuronal dynamics**?
parameters of ionic currents such as **conductance** and **time constant**?

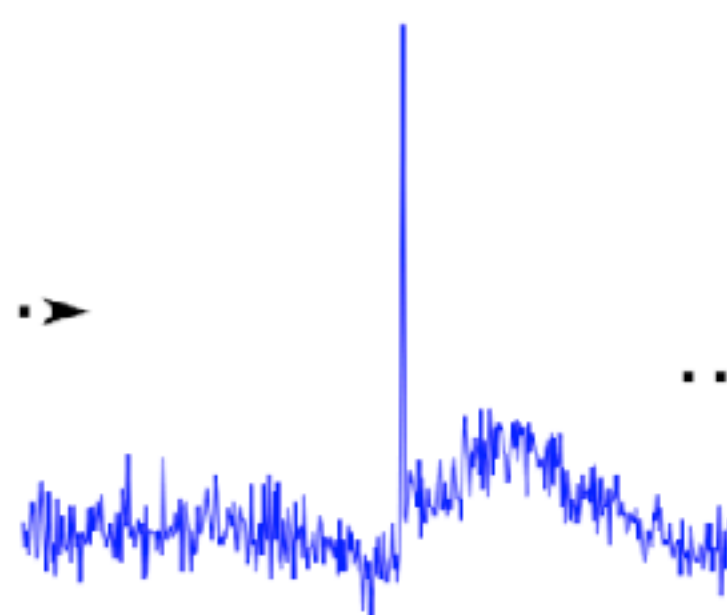
Pair of neurons
Monosynaptic connection



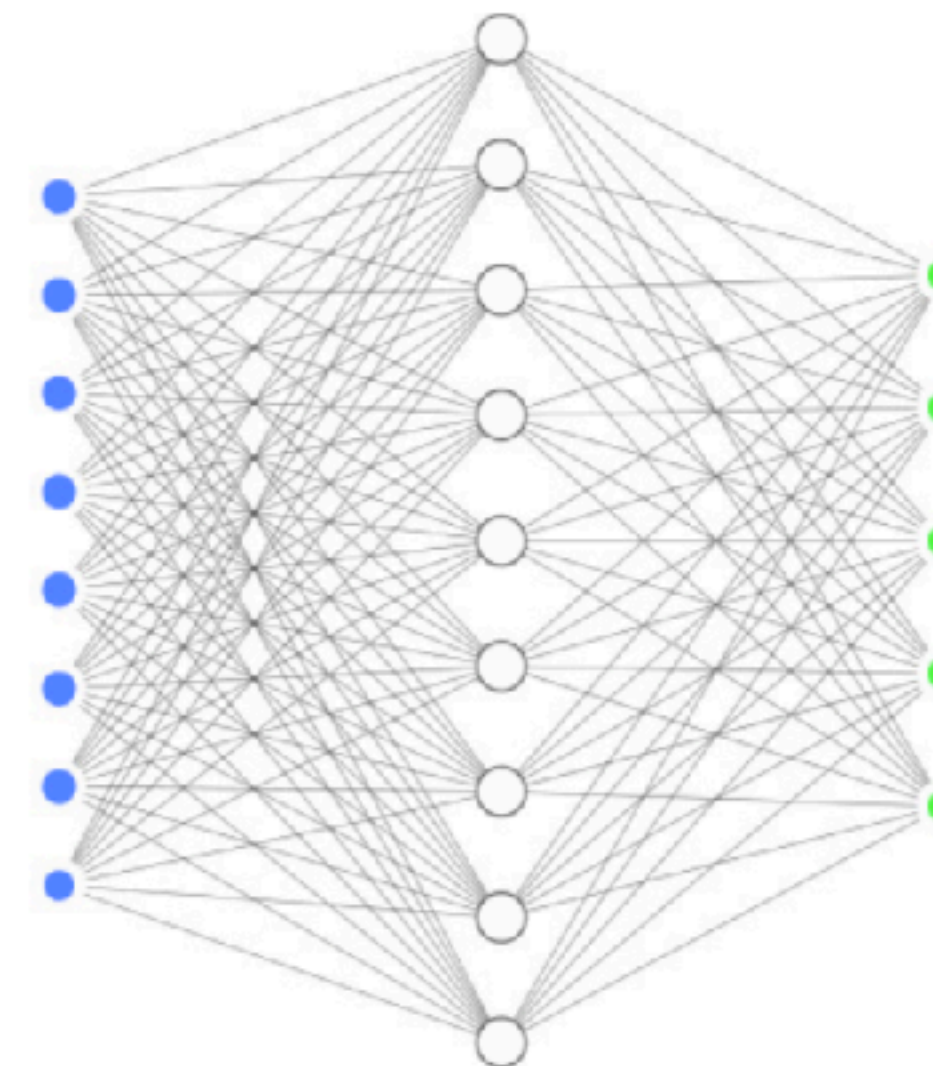
Spike-train



Cross-correlation function



Artificial neural network



Values of
Conductance
Time constant

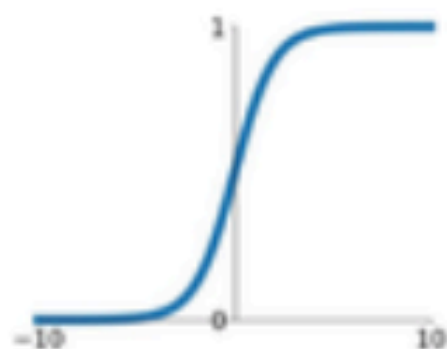


We vary:

- Activation functions

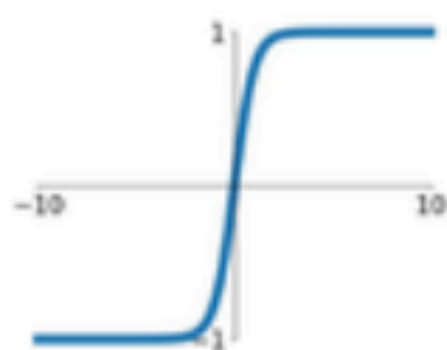
Sigmoid

$$\sigma(x) = \frac{1}{1+e^{-x}}$$



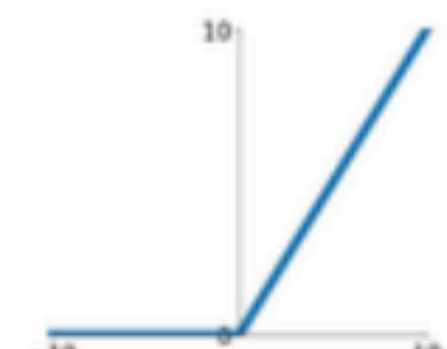
tanh

$$\tanh(x)$$



ReLU

$$\max(0, x)$$



Training procedure:

- Training:validation is 80:20
- Backpropagation algorithm;
- Stochastic gradient descent;
- Cross-entropy loss function;

$$\mathcal{L} = -\frac{1}{m} \sum_{i=1}^m y_i \log(\hat{y}_i),$$

We measure:

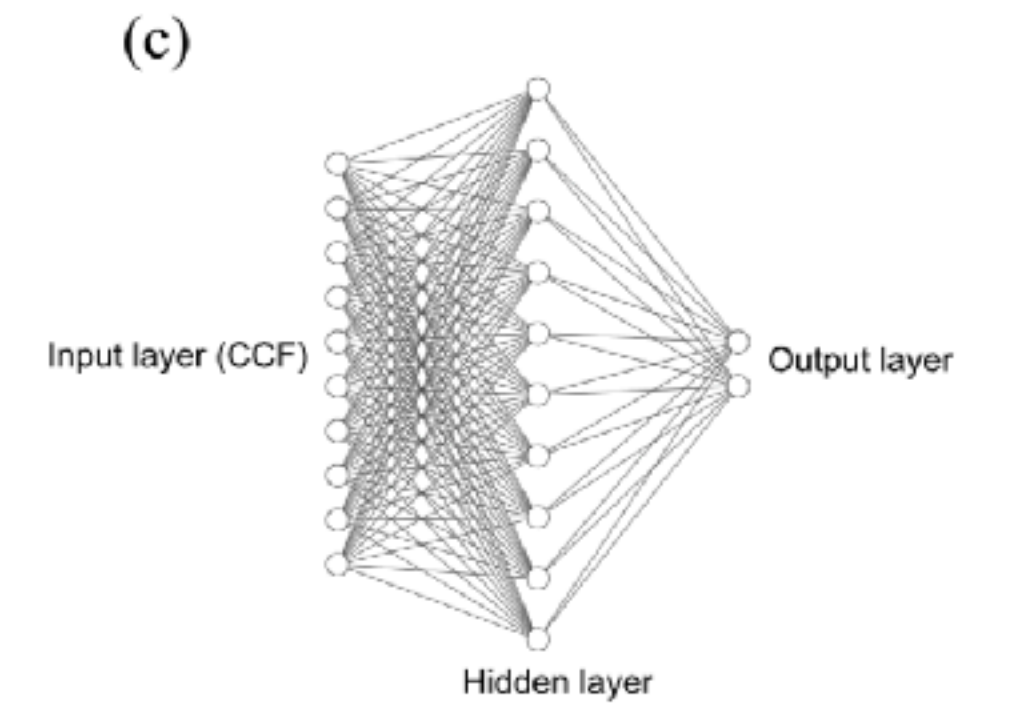
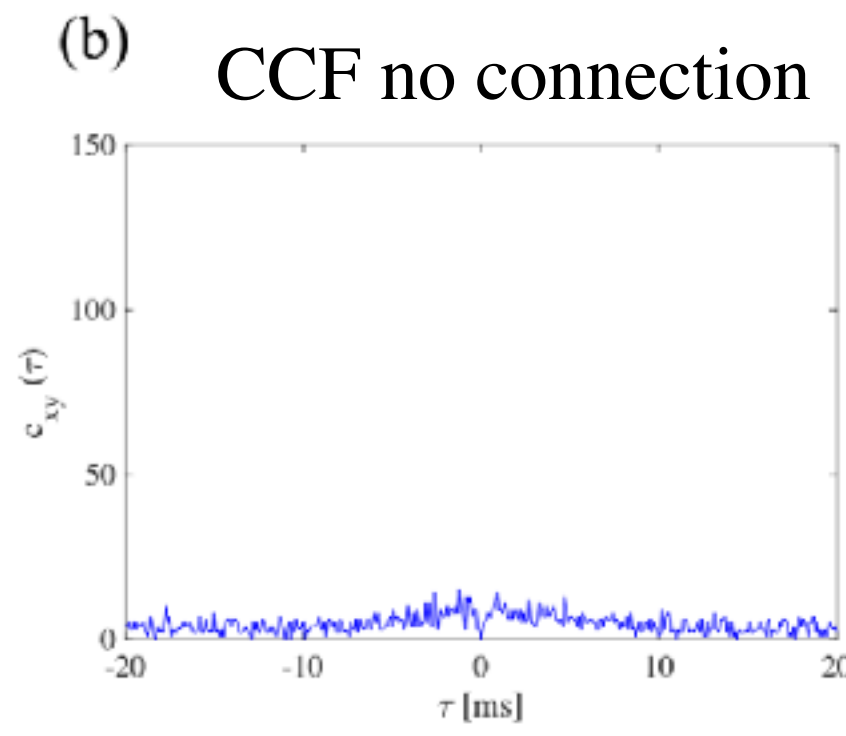
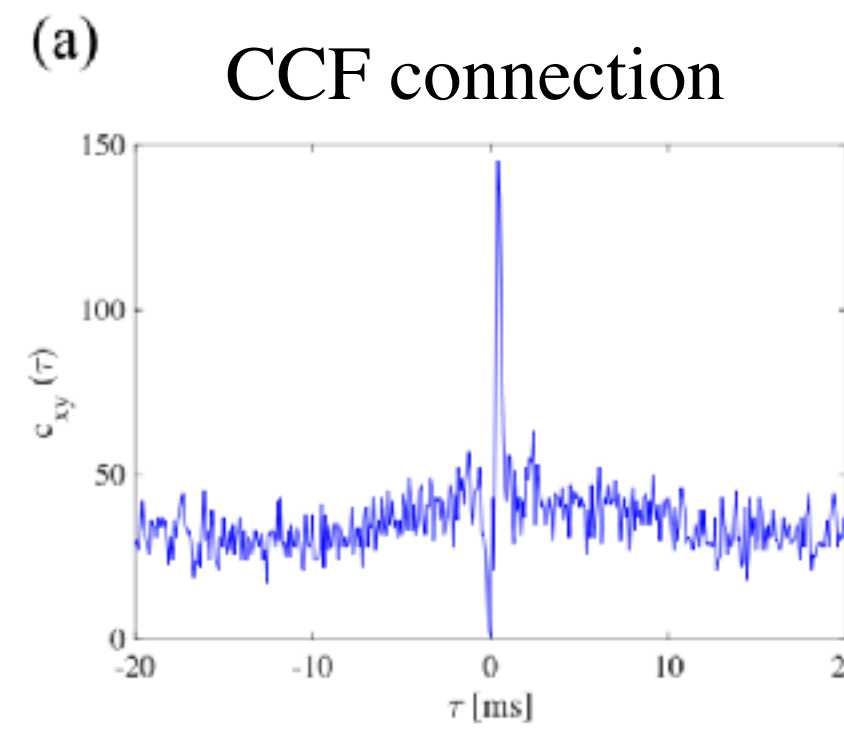
$$\text{Precision} = \frac{TP}{TP + FP}$$

$$\text{Recall} = \frac{TP}{TP + FN}$$

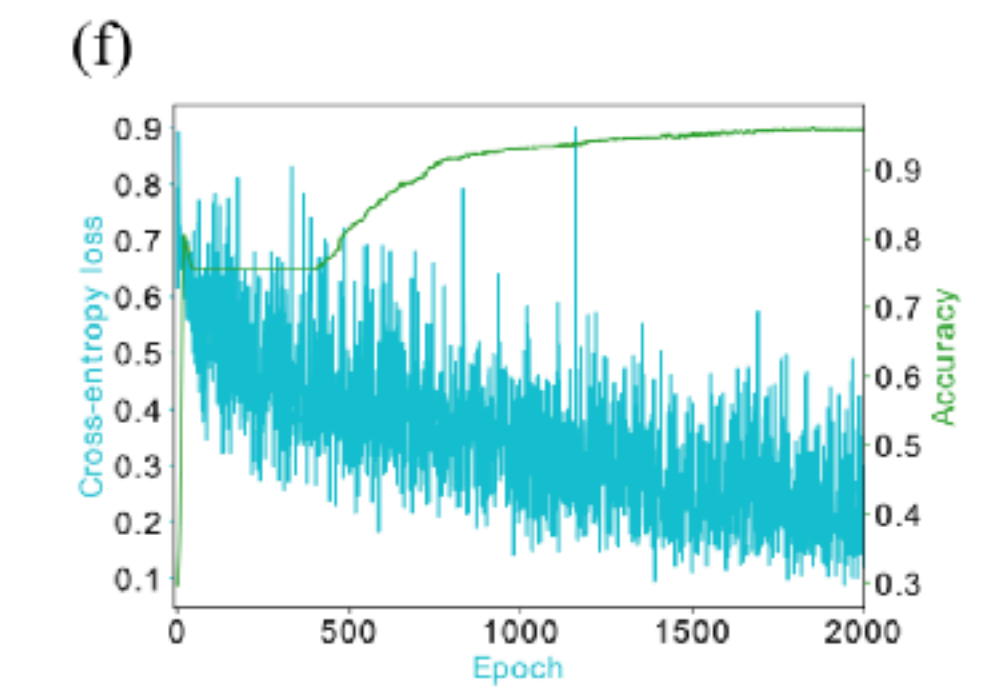
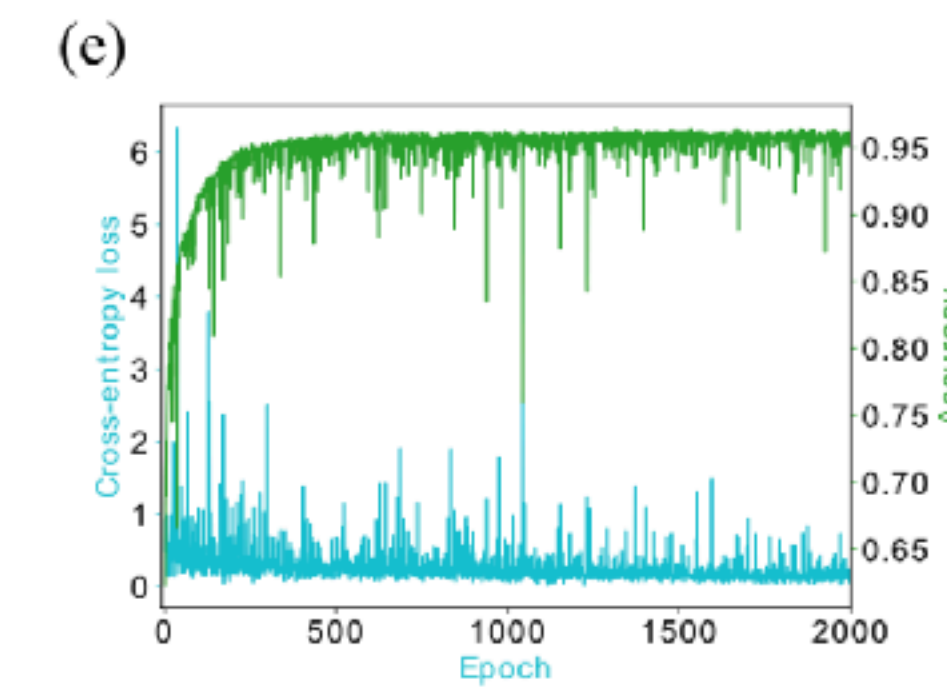
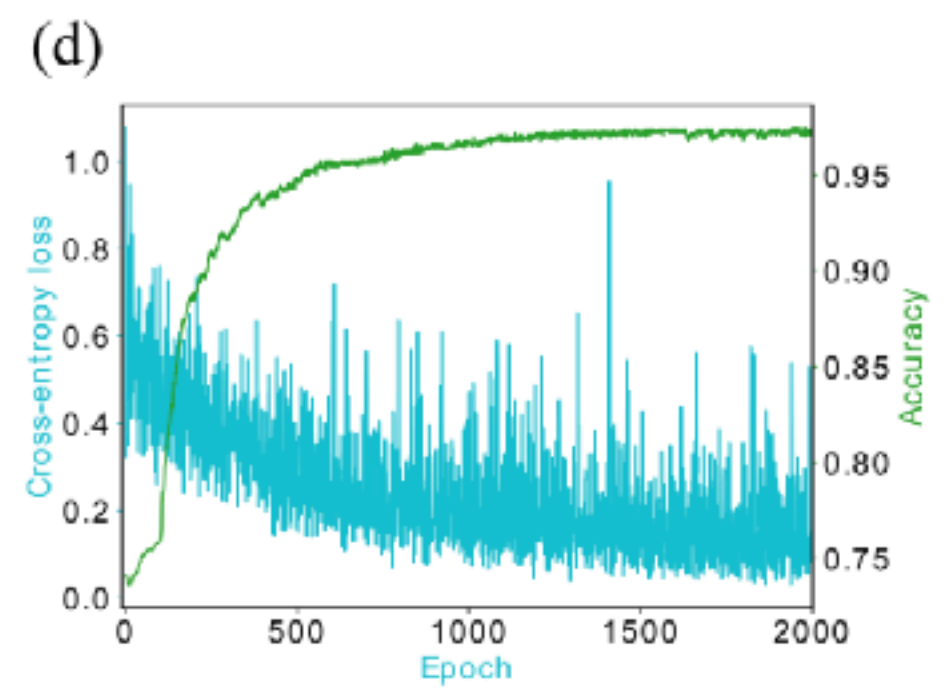
$$\text{F1 - score} = \frac{2 \times (\text{Precision} \times \text{Recall})}{\text{Precision} + \text{Recall}}$$

$$\text{MCC} = \frac{(TP \times TN - FP \times FN)}{\sqrt{(TP + FP)(TP + FN)(TN + FP)(TN + FN)}}$$

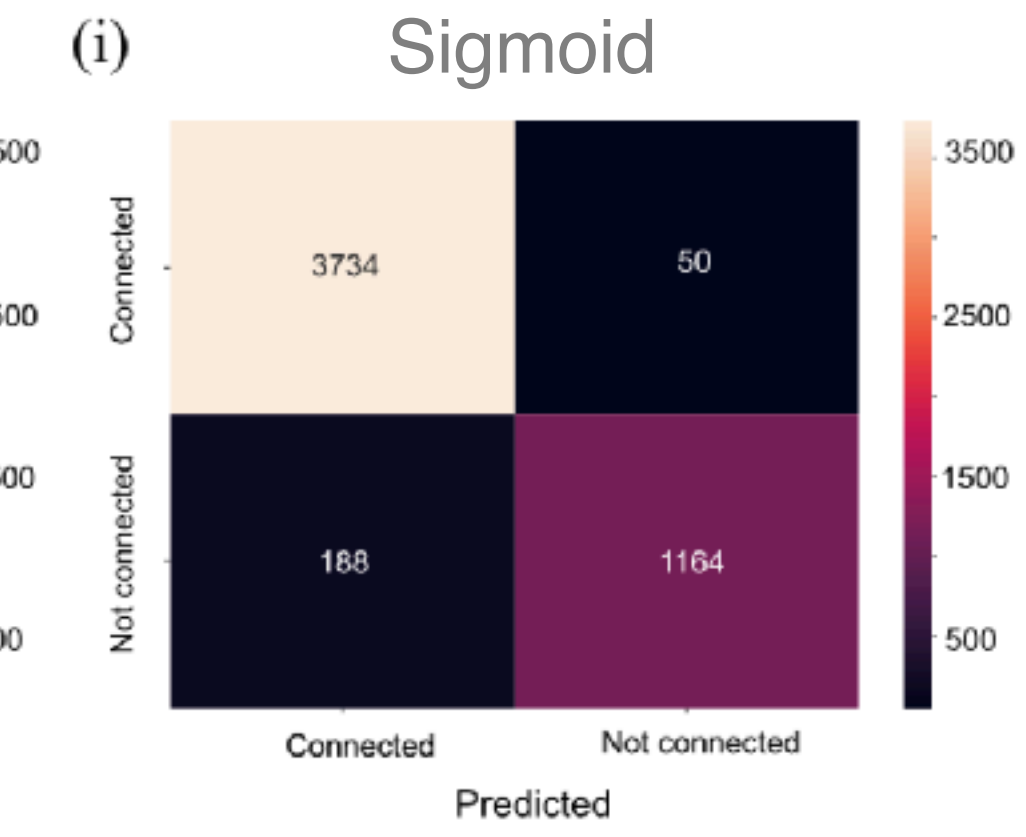
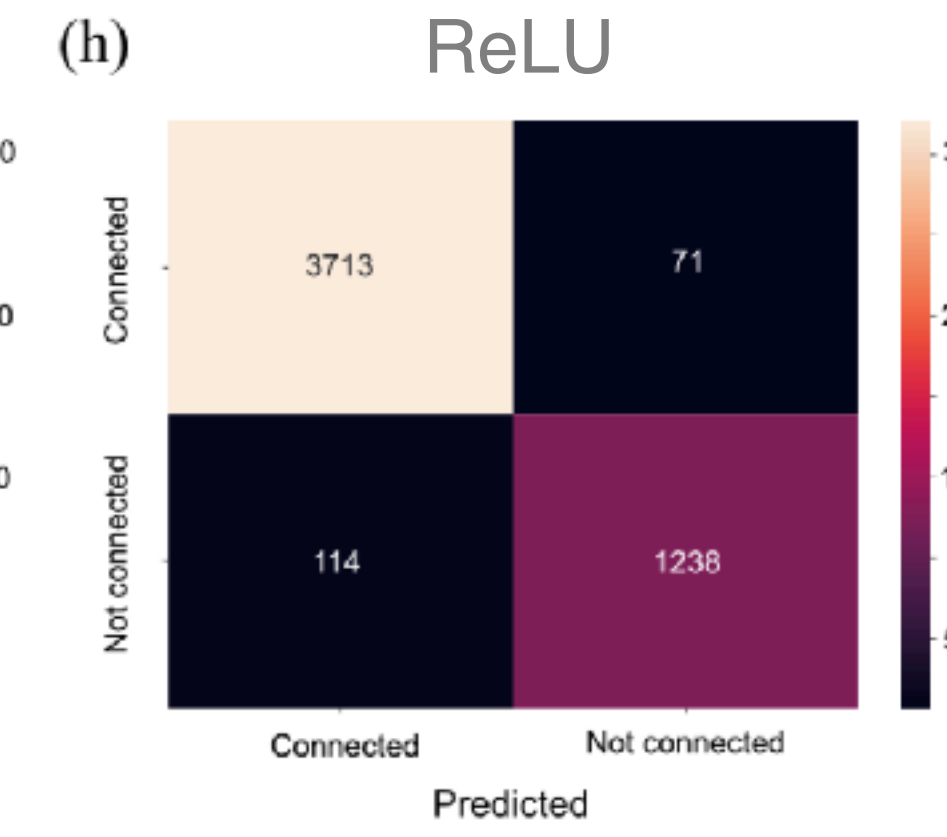
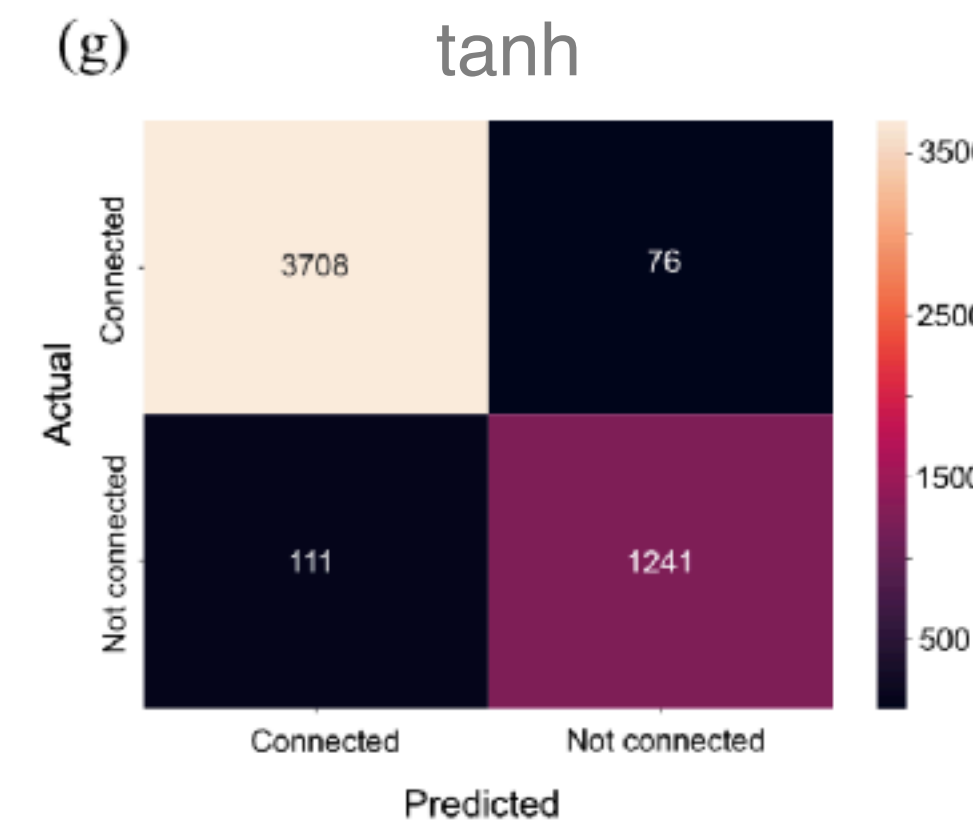
Monosynaptic connections from experimental data
(Sam McKenzie - University of New Mexico)



Training →



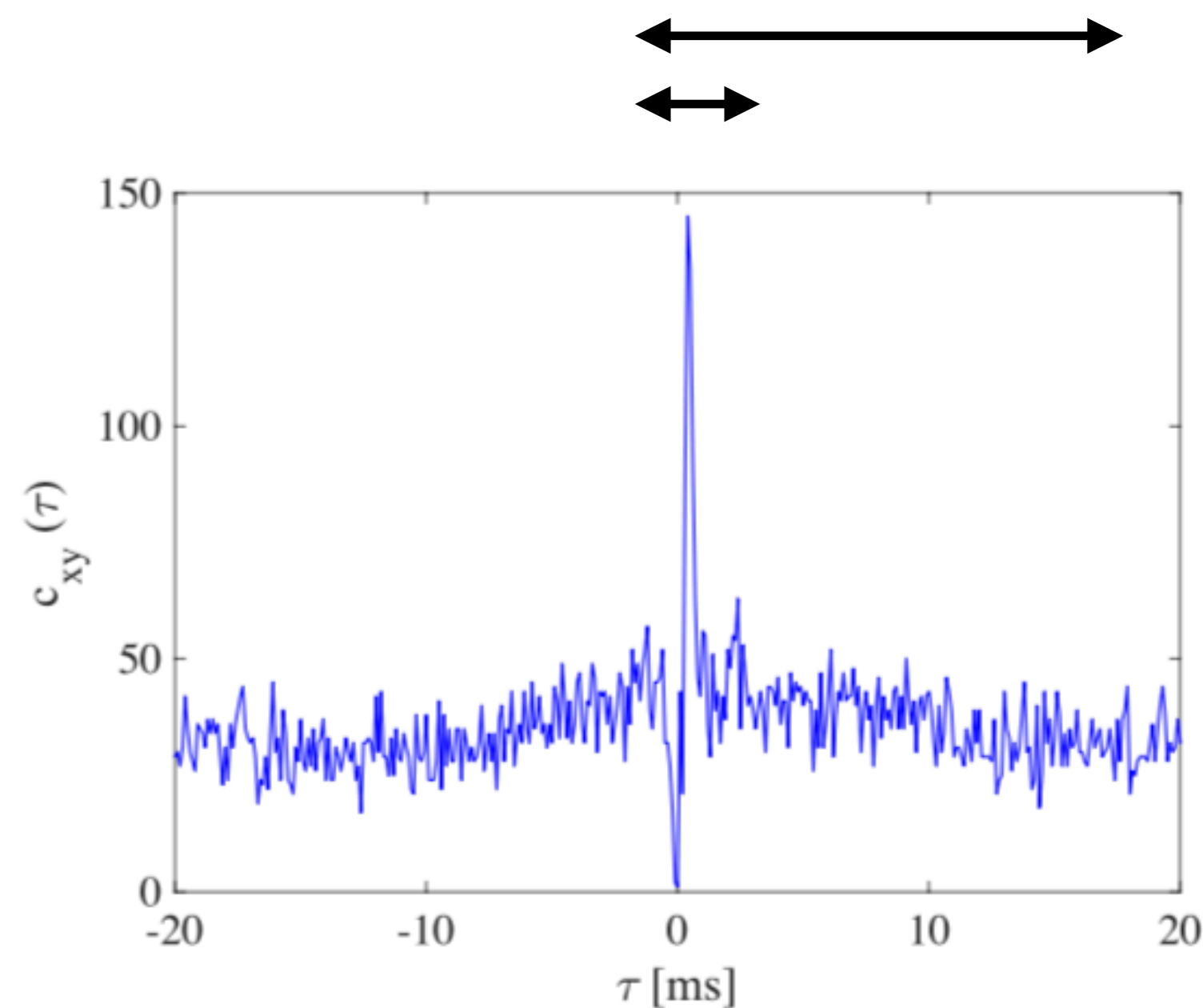
Confusion matrices →



high classification accuracy →

Activation function	Connection	Precision	Recall	F1-score	MCC
tanh	No	0.97	0.98	0.97	0.903
	Yes	0.94	0.92	0.93	
ReLU	No	0.97	0.99	0.98	0.914
	Yes	0.97	0.91	0.94	
Sigmoid	No	0.95	0.99	0.97	0.88
	Yes	0.96	0.86	0.91	

Using information from longer lags change doesn't change results



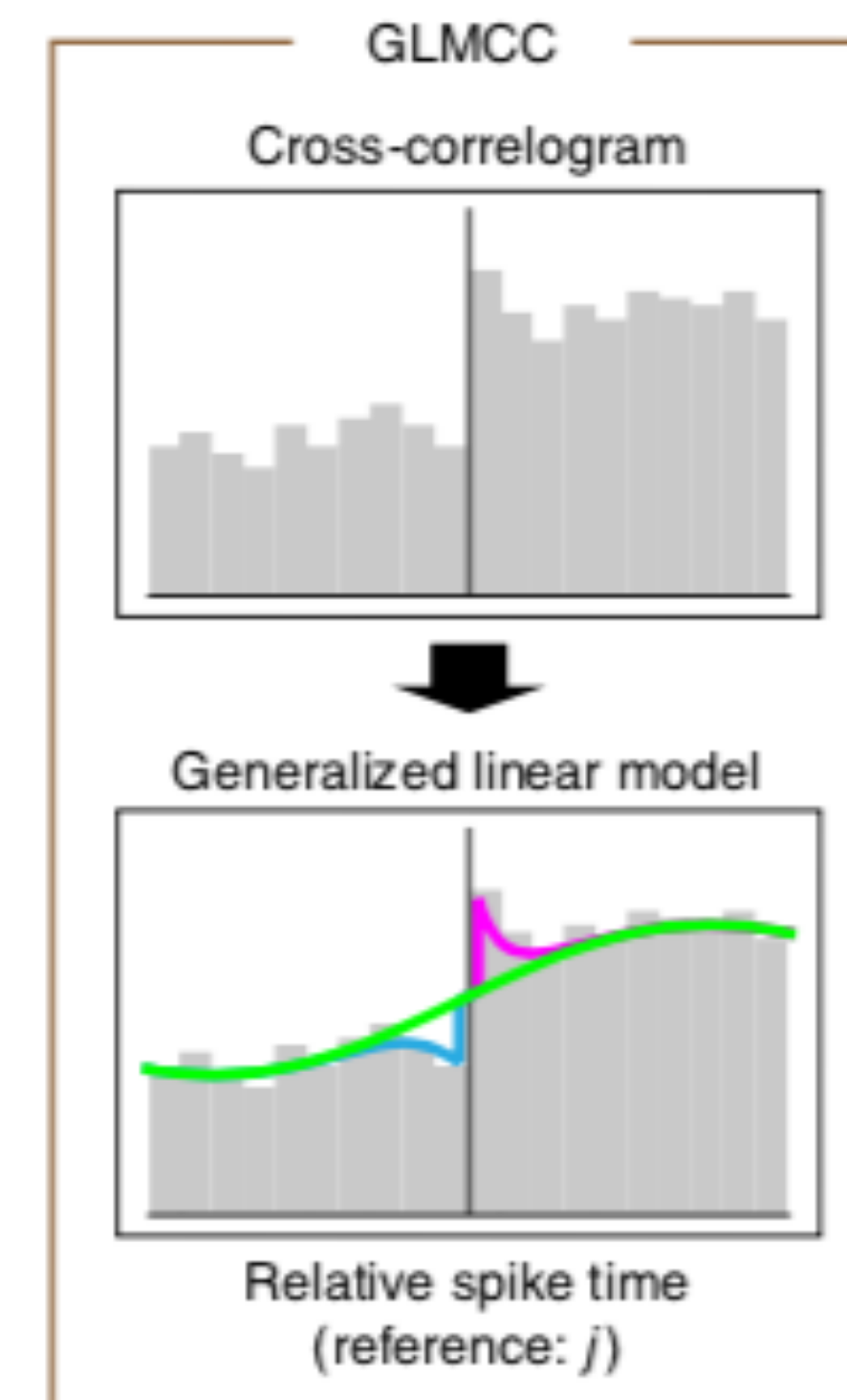
F1-scores and MCCs are slightly lower

Activation function	Connection	Precision	Recall	F1-score	MCC
tanh	No	0.94	0.96	0.95	0.813
	Yes	0.88	0.84	0.86	
ReLU	No	0.99	0.95	0.97	0.881
	Yes	0.87	0.97	0.91	
Sigmoid	No	0.92	0.98	0.95	0.795
	Yes	0.92	0.77	0.84	

Only around 1st peak



This is how it has been done



Kobayashi et al., *Nat Commun.*, 10:4468 (2019)

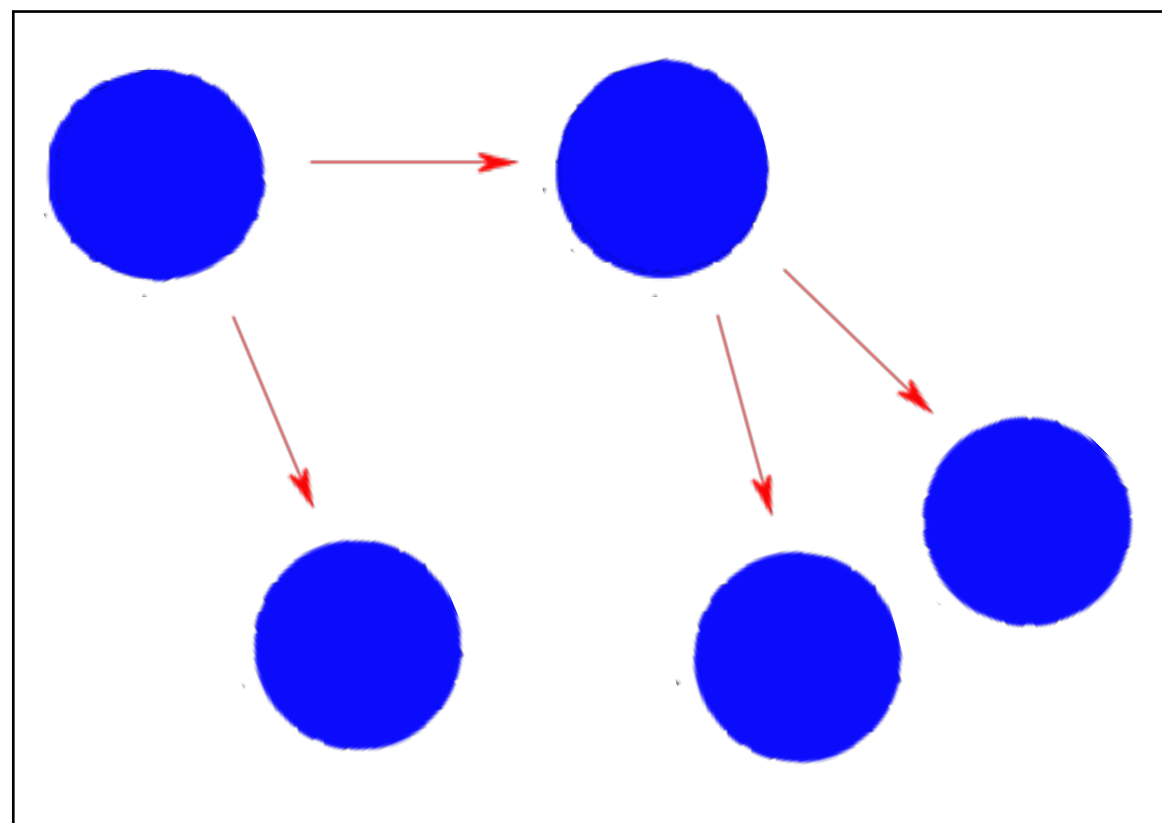
Or others that rely on a single lag...

Artificial neural networks: what about time constants and conductance values?

Generation of synthetic data

- Database with 25,000 CCFs

Random arrangements of a 5x5 network
(always monosynapses)



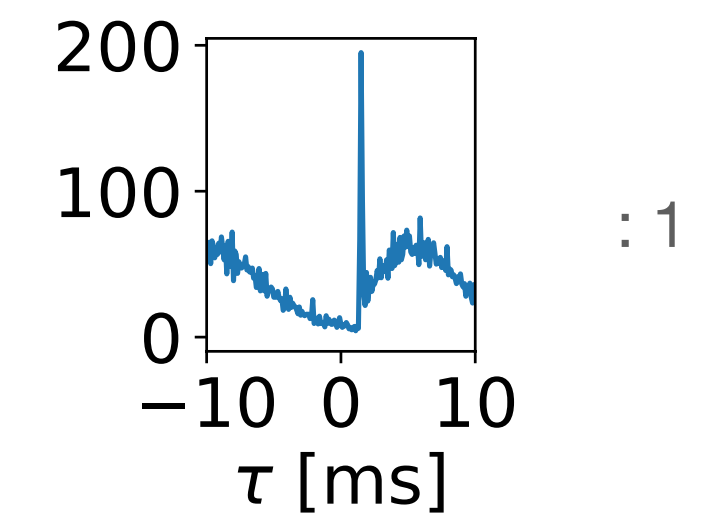
- Each CCF 100 trials per network

G_h in the range [1.5,5.5]

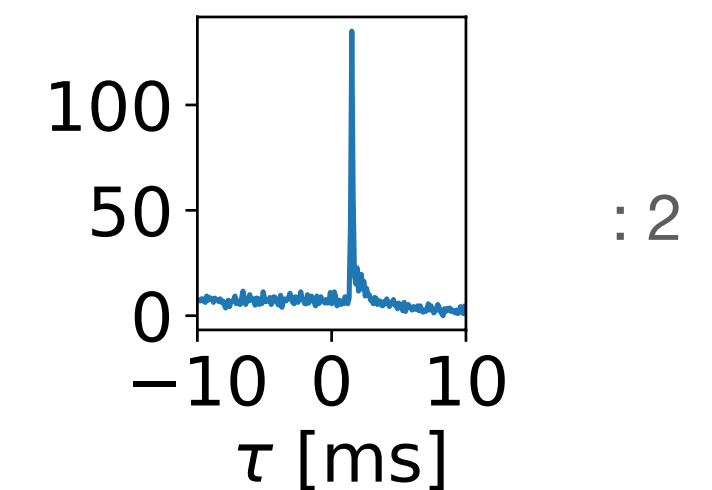
τ_h in the range [40,200]

- Labelling process:

Connection where
 I_h with $G_h = 1.5$



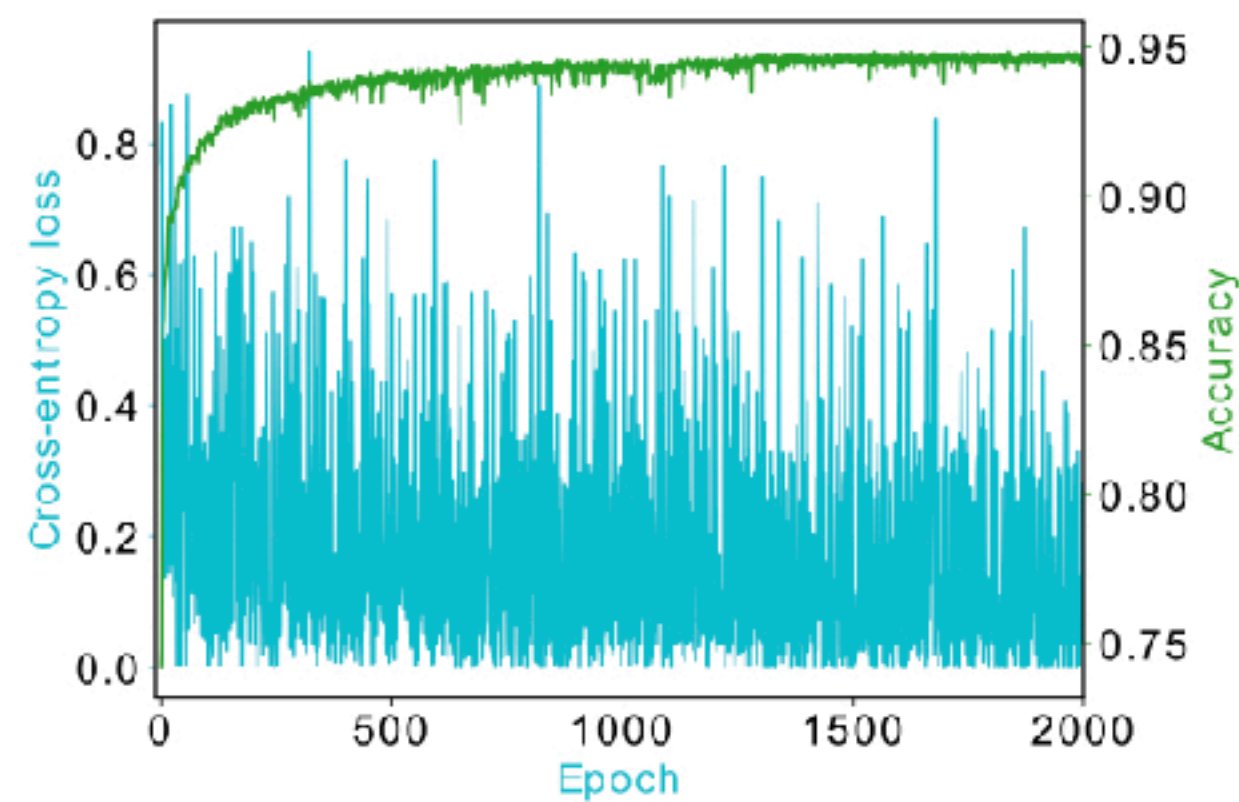
Connection where
 I_h with $G_h = 3.5$



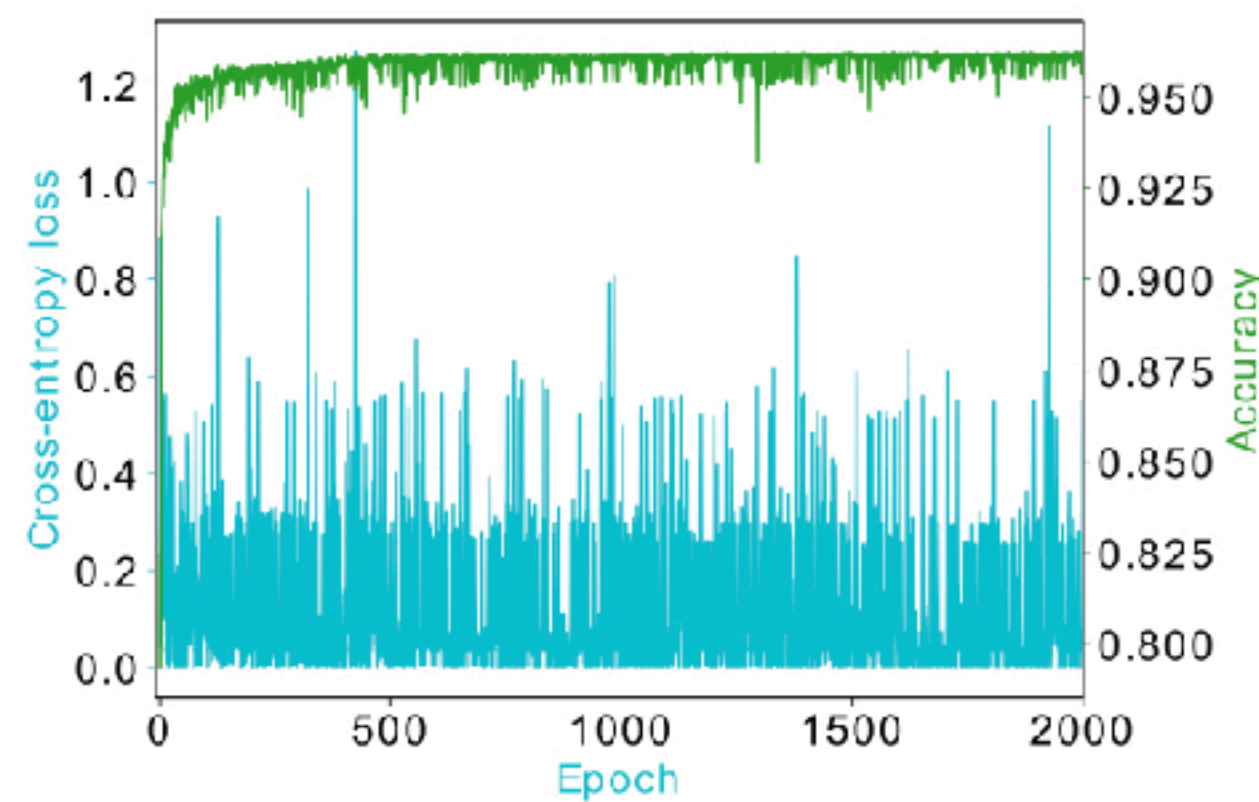
All
other
cases : 0

Conductances and time constants are classified with high accuracy;

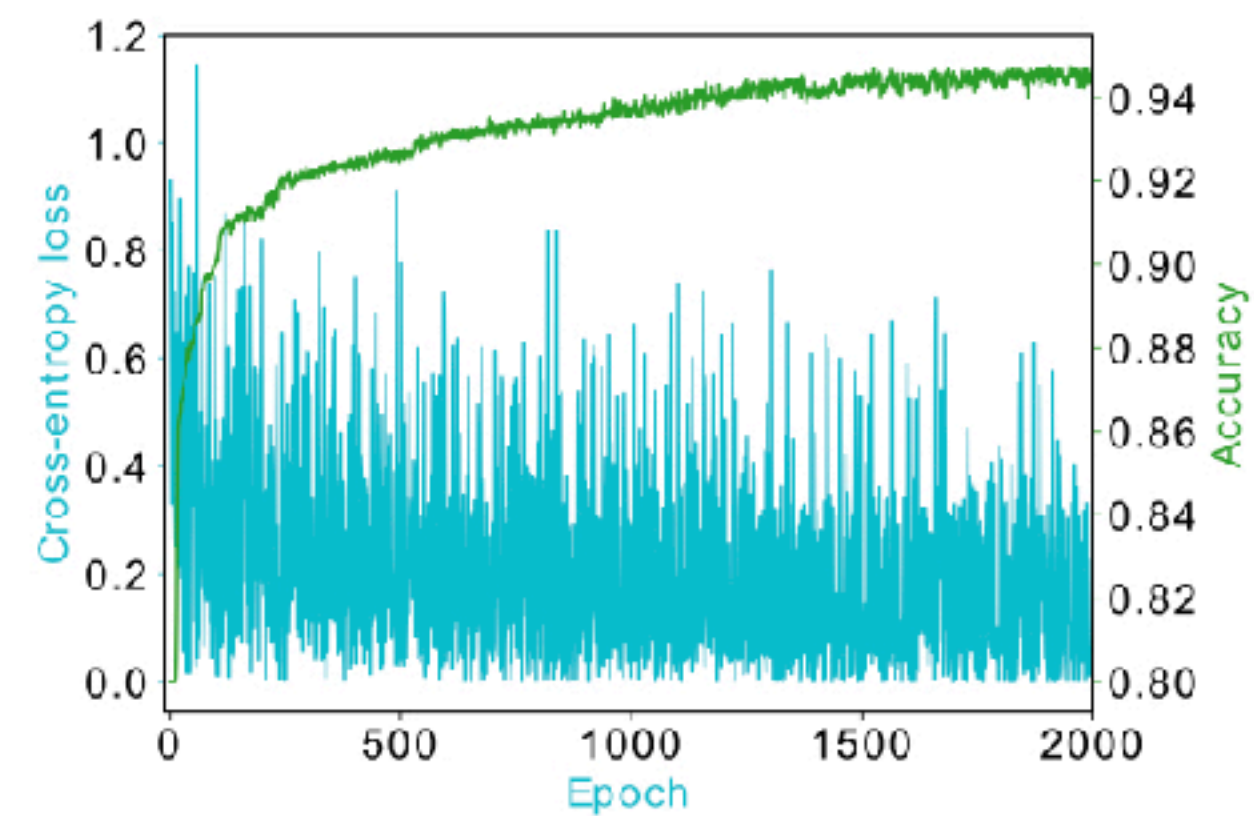
(a) tanh



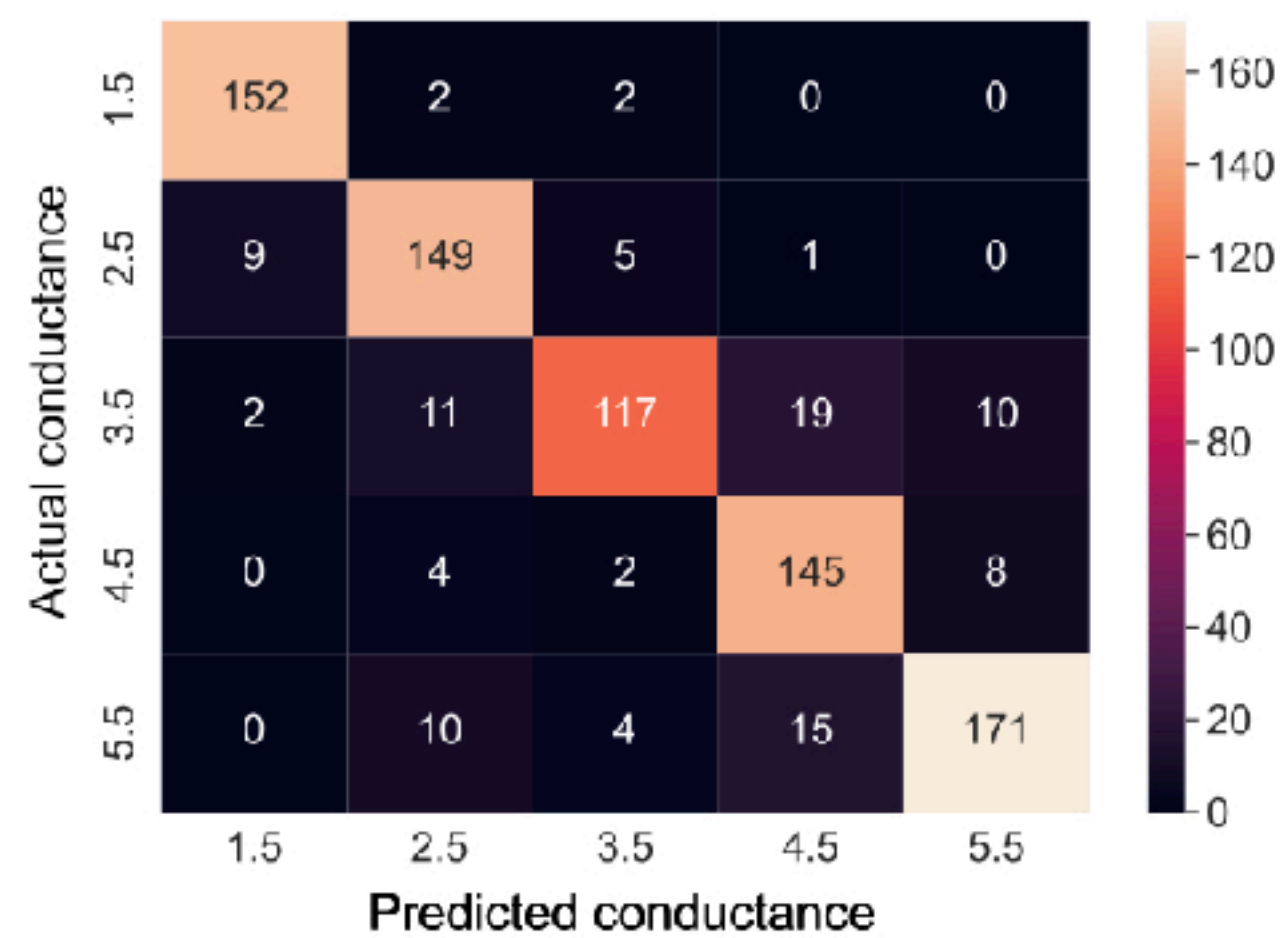
(b) ReLU



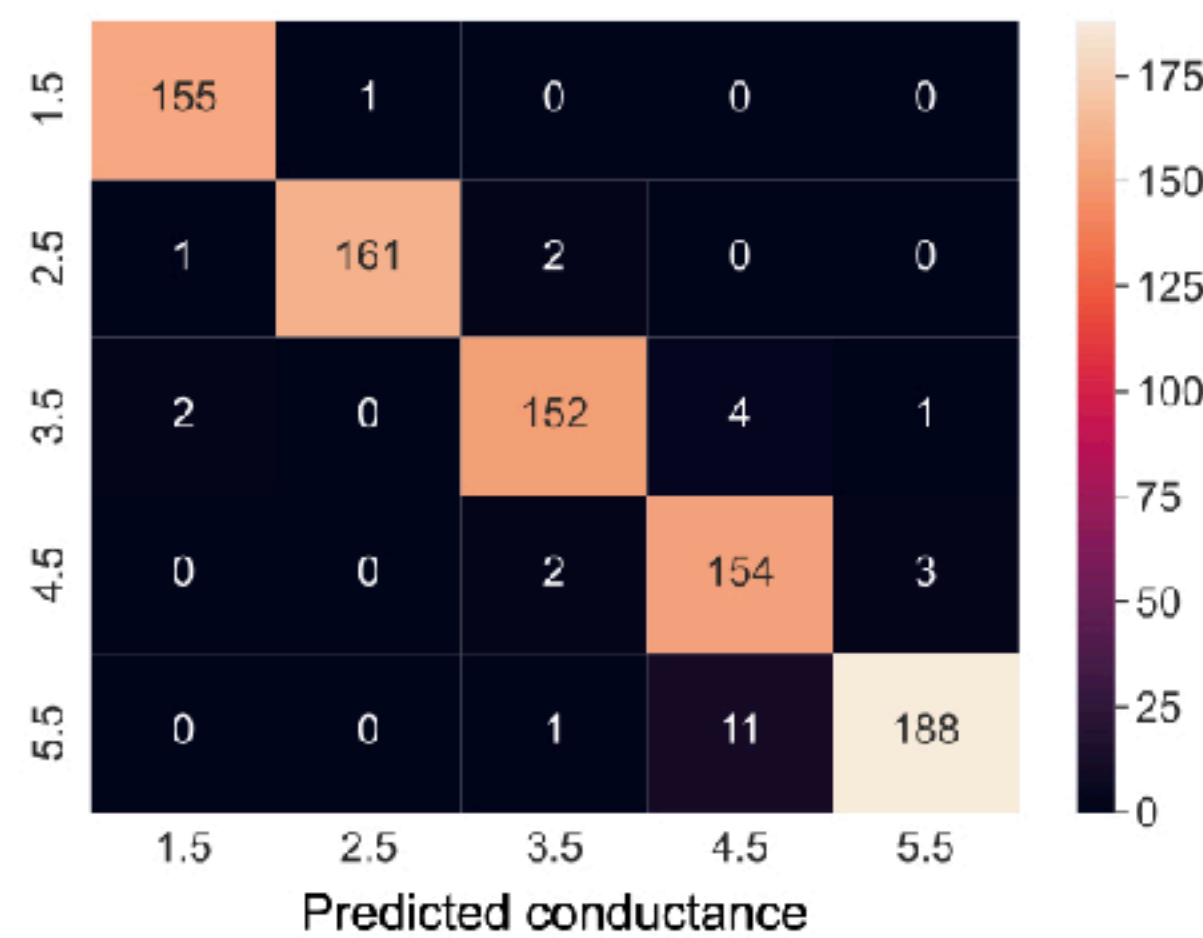
(c) Sigmoid



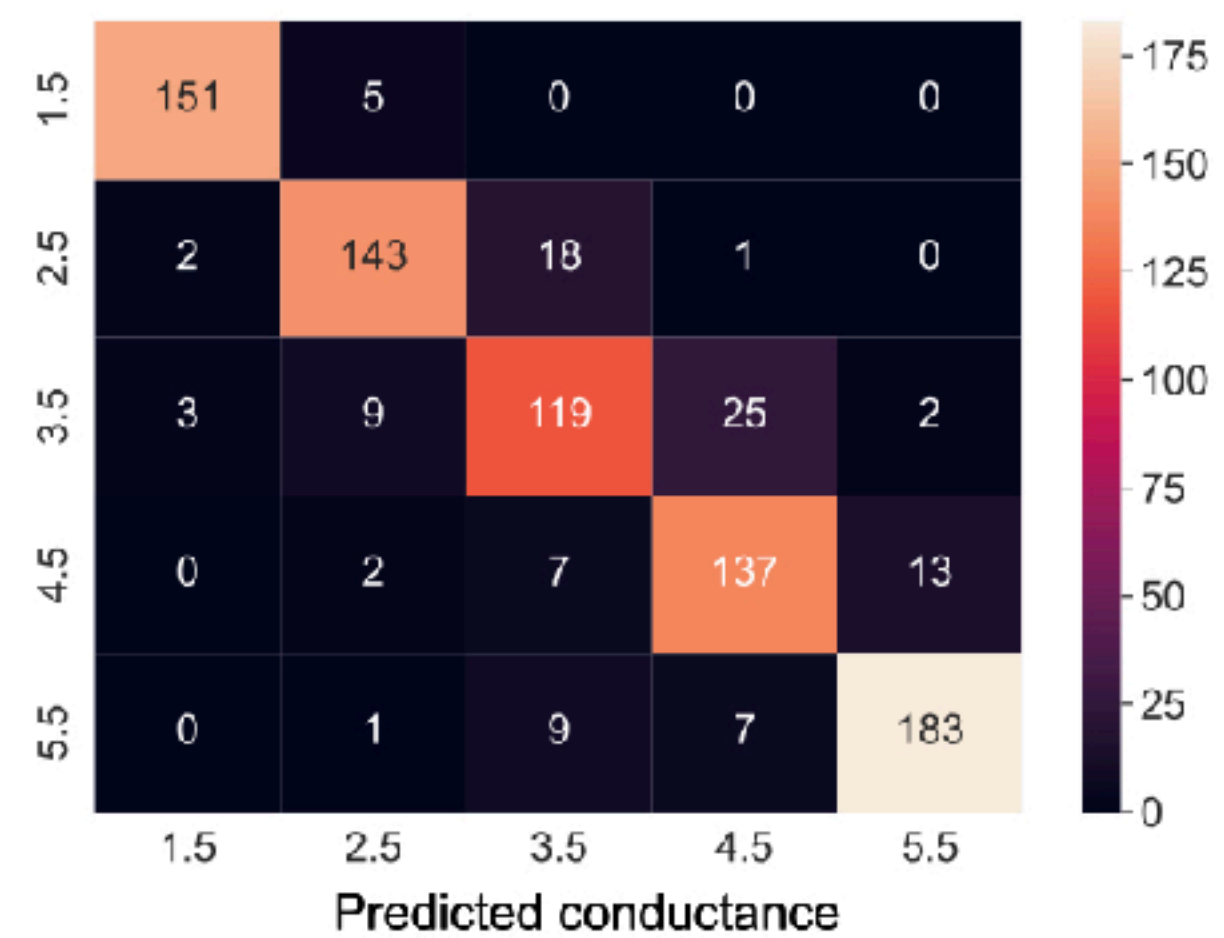
(d)



(e)

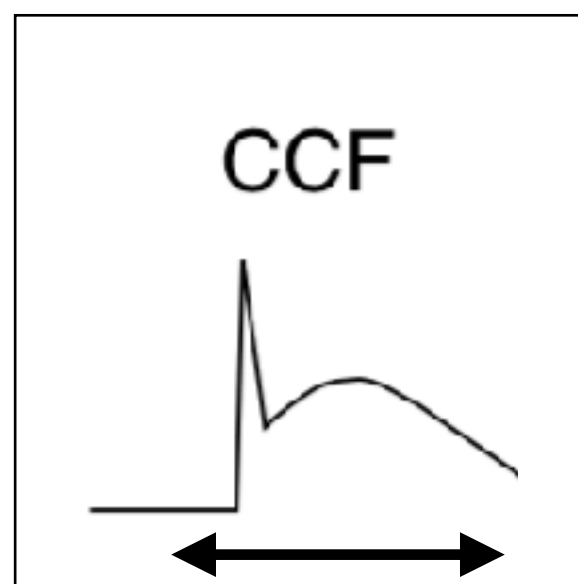


(f)

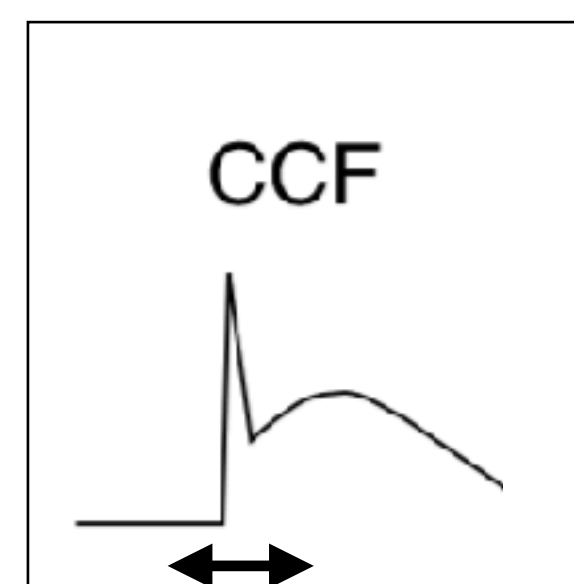


ReLU is more reliable.

... but only if information beyond the first peak is provided



G_h variation					
Activation function	G_h	Precision	Recall	F1-score	MCC
tanh	No connection	0.96	1.0	0.98	0.849
	1.5	0.98	0.78	0.87	
	2.5	0.88	0.75	0.81	
	3.5	0.87	0.68	0.76	
	4.5	0.83	0.70	0.76	
	5.5	0.86	0.86	0.86	
ReLU	No connection	0.95	1.0	0.98	0.876
	1.5	1.0	0.8	0.89	
	2.5	1.0	0.81	0.89	
	3.5	0.99	0.80	0.89	
	4.5	0.95	0.79	0.86	
	5.5	0.99	0.76	0.86	
Sigmoid	No connection	0.96	1.0	0.98	0.856
	1.5	0.94	0.76	0.84	
	2.5	0.90	0.72	0.80	
	3.5	0.89	0.70	0.78	
	4.5	0.88	0.70	0.78	
	5.5	0.88	0.95	0.91	
τ_h variation					
Activation function	τ_h	Precision	Recall	F1-score	MCC
tanh	No connection	0.95	1.0	0.97	0.803
	40	0.86	0.76	0.81	
	80	0.82	0.66	0.73	
	120	0.83	0.67	0.74	
	160	0.93	0.60	0.73	
	200	0.94	0.67	0.78	
ReLU	No connection	0.95	1.0	0.97	0.829
	40	0.97	0.74	0.84	
	80	0.92	0.70	0.79	
	120	0.96	0.66	0.78	
	160	0.96	0.70	0.81	
	200	0.98	0.78	0.81	
Sigmoid	No connection	0.95	1.0	0.97	0.784
	40	0.79	0.76	0.77	
	80	0.81	0.61	0.70	
	120	0.78	0.60	0.68	
	160	0.91	0.60	0.72	
	200	0.92	0.65	0.76	



G_h variation					
Activation function	G_h	Precision	Recall	F1-score	MCC
tanh	No connection	0.95	1.0	0.98	0.704
	1.5	0.82	0.48	0.61	
	2.5	0.56	0.67	0.61	
	3.5	0.57	0.42	0.48	
	4.5	0.77	0.36	0.50	
	5.5	0.62	0.63	0.63	
ReLU	No connection	0.95	1.0	0.98	0.792
	1.5	0.89	0.76	0.82	
	2.5	0.82	0.69	0.75	
	3.5	0.73	0.58	0.65	
	4.5	0.80	0.58	0.68	
	5.5	0.84	0.66	0.74	
Sigmoid	No connection	0.95	1.0	0.98	0.746
	1.5	0.87	0.67	0.75	
	2.5	0.71	0.64	0.68	
	3.5	0.60	0.52	0.56	
	4.5	0.71	0.53	0.61	
	5.5	0.75	0.55	0.64	
τ_h variation					
Activation function	τ_h	Precision	Recall	F1-score	MCC
tanh	No connection	0.95	1.0	0.97	0.634
	40	0.55	0.63	0.59	
	80	0.45	0.20	0.27	
	120	0.44	0.47	0.46	
	160	0.57	0.37	0.45	
	200	0.67	0.38	0.49	
ReLU	No connection	0.94	1.0	0.97	0.659
	40	0.67	0.71	0.69	
	80	0.70	0.32	0.44	
	120	0.44	0.33	0.38	
	160	0.72	0.34	0.46	
	200	0.50	0.54	0.52	
Sigmoid	No connection	0.94	1.0	0.97	0.623
	40	0.68	0.62	0.65	
	80	0.43	0.35	0.38	
	120	0.42	0.36	0.39	
	160	0.42	0.32	0.36	
	200	0.66	0.33	0.44	

Can we use CCFs?

CCFs are essentially linear metrics

Channel dynamics is not linear!

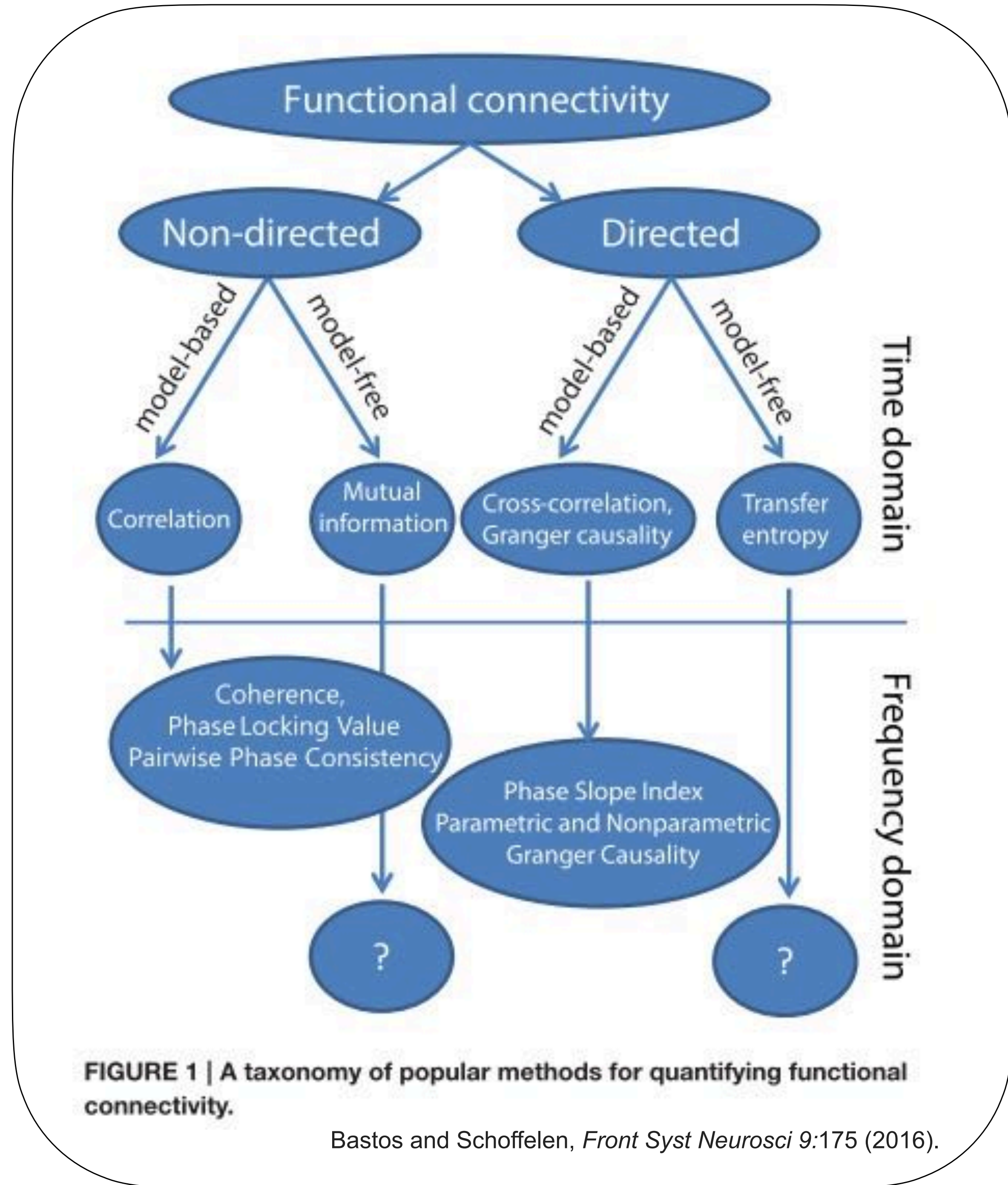
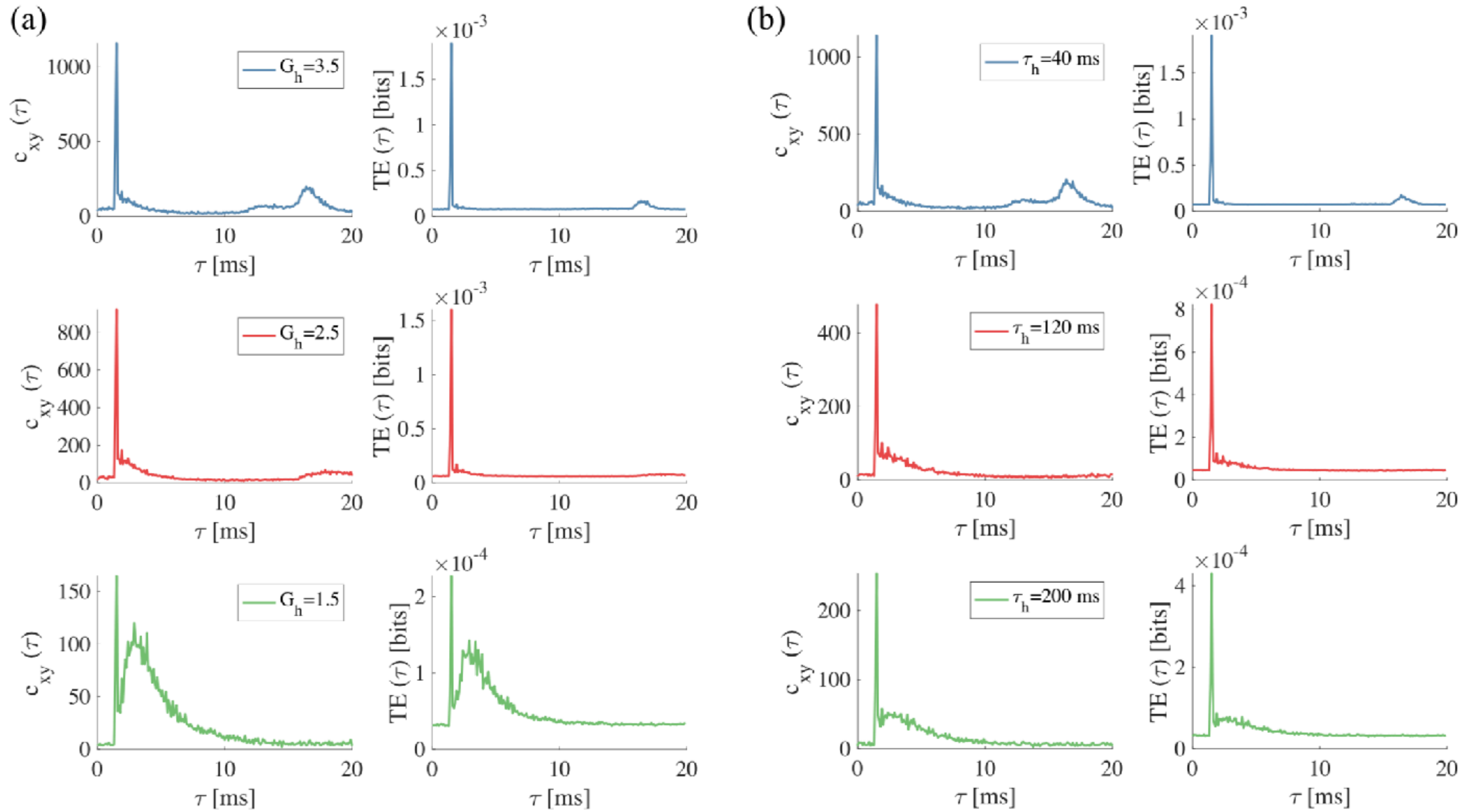


FIGURE 1 | A taxonomy of popular methods for quantifying functional connectivity.

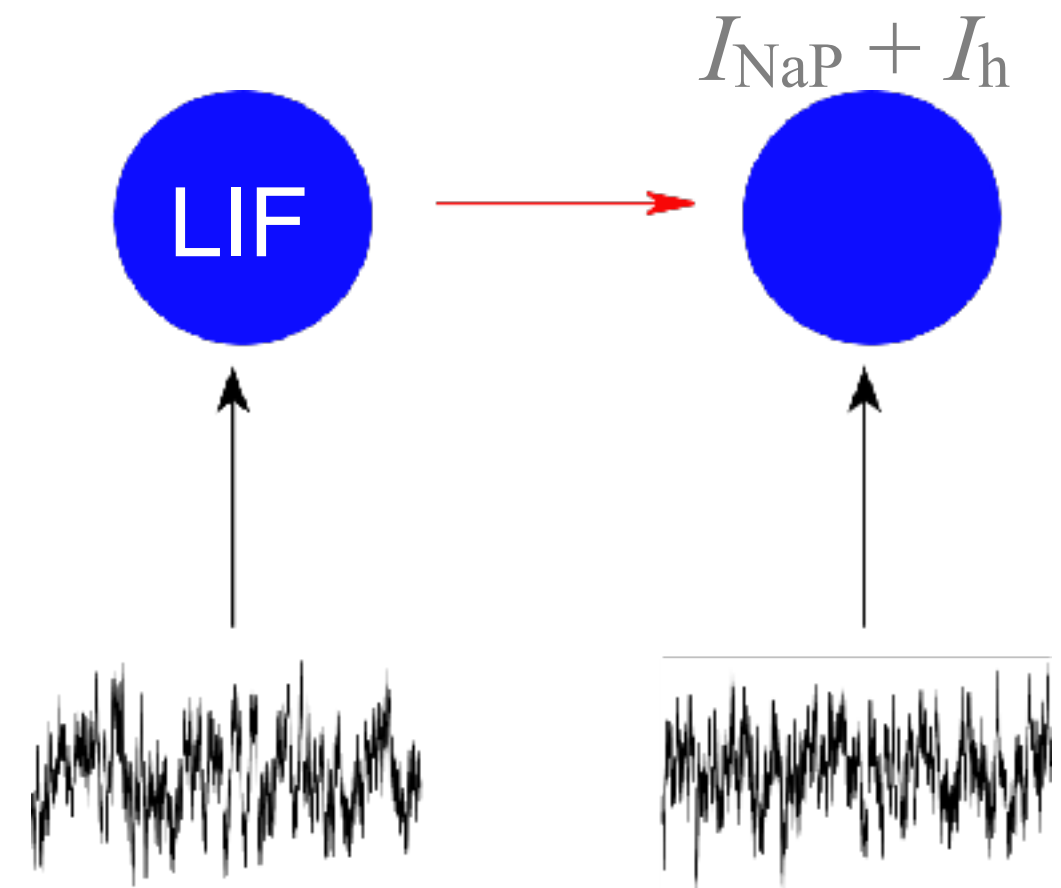
Bastos and Schoffelen, *Front Syst Neurosci* 9:175 (2016).



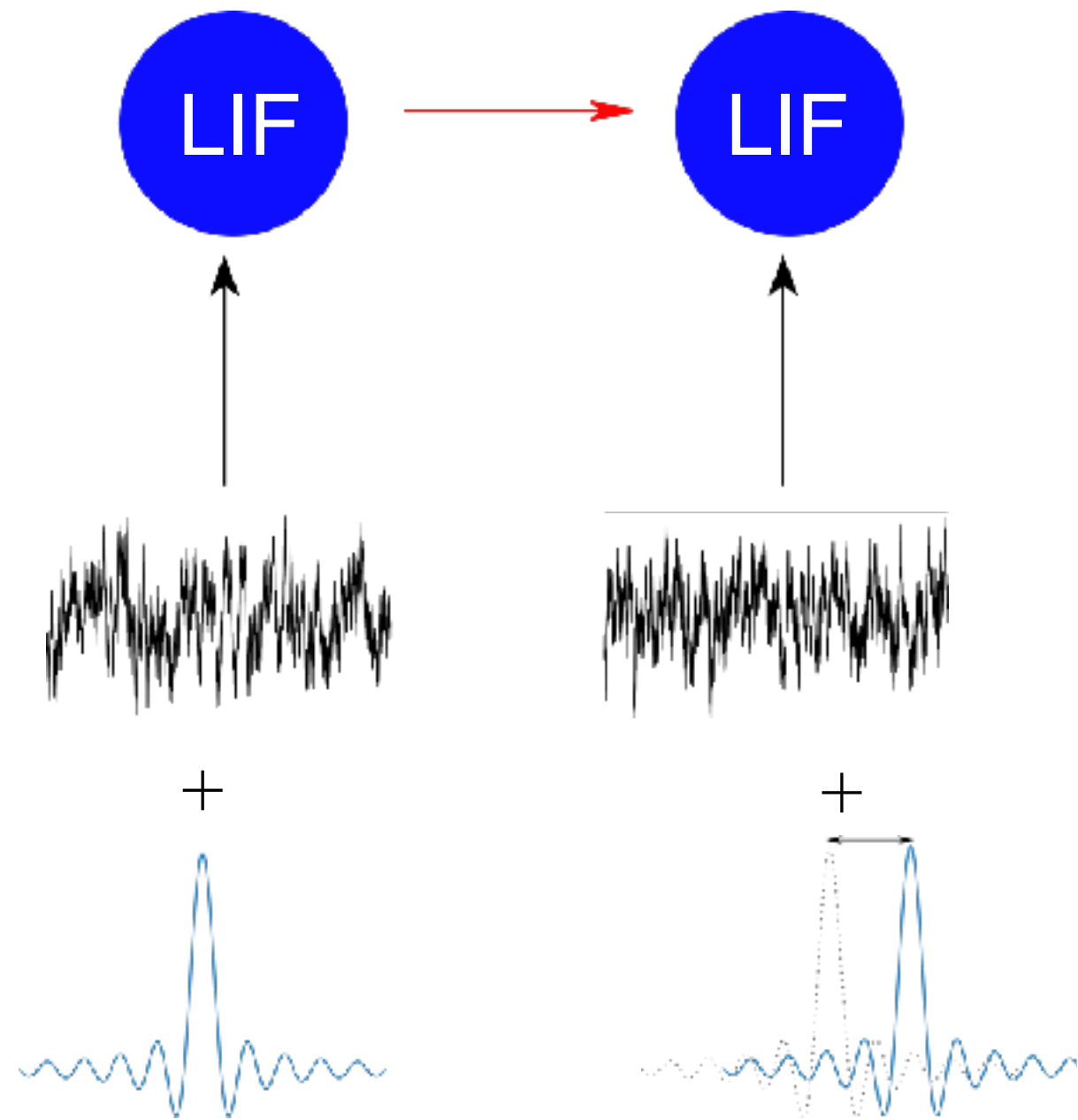
- Similar **variety** of data in TEs is and CCFs,
- Variety is the only input necessary for a **supervised training algorithm**.

Confounding factors

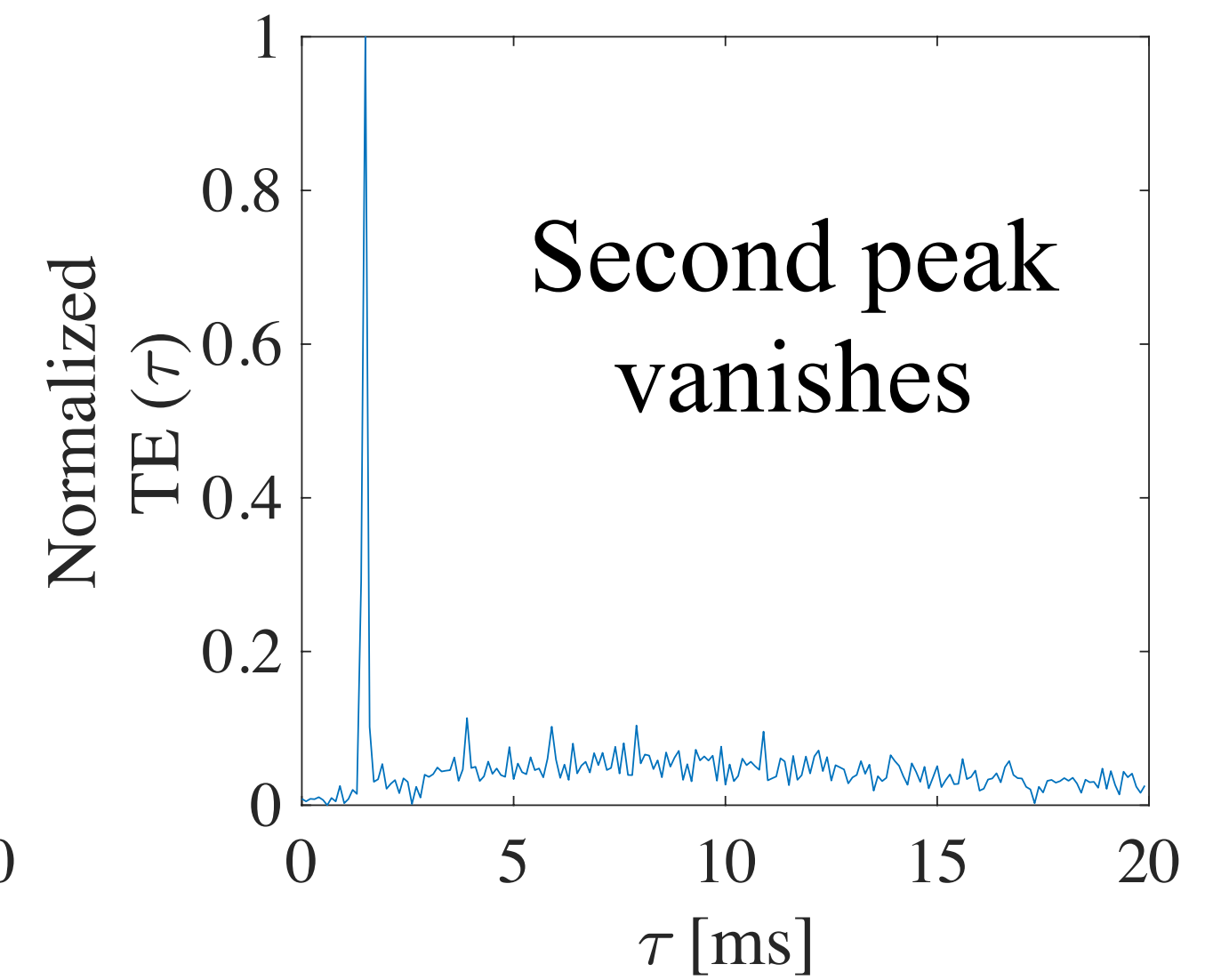
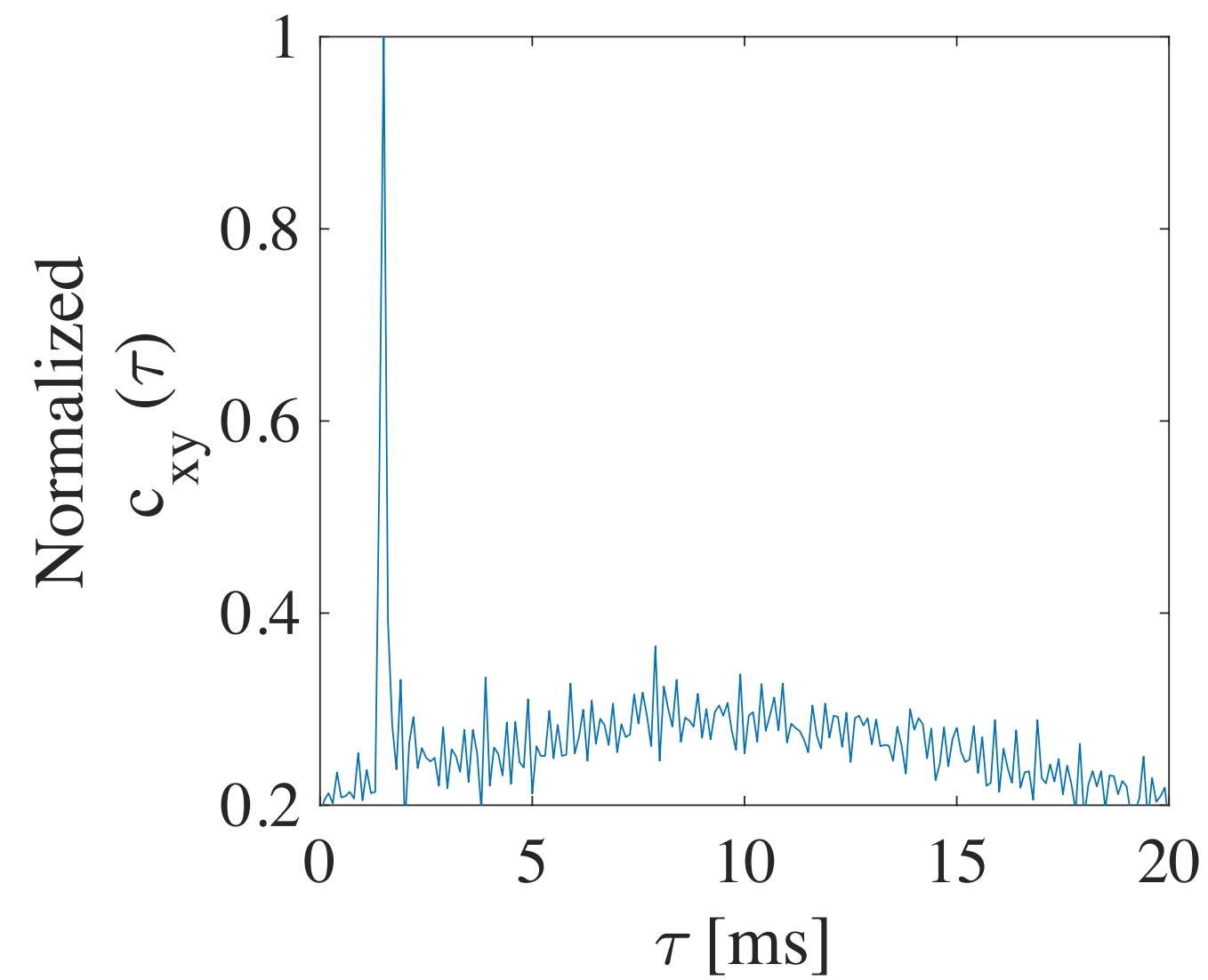
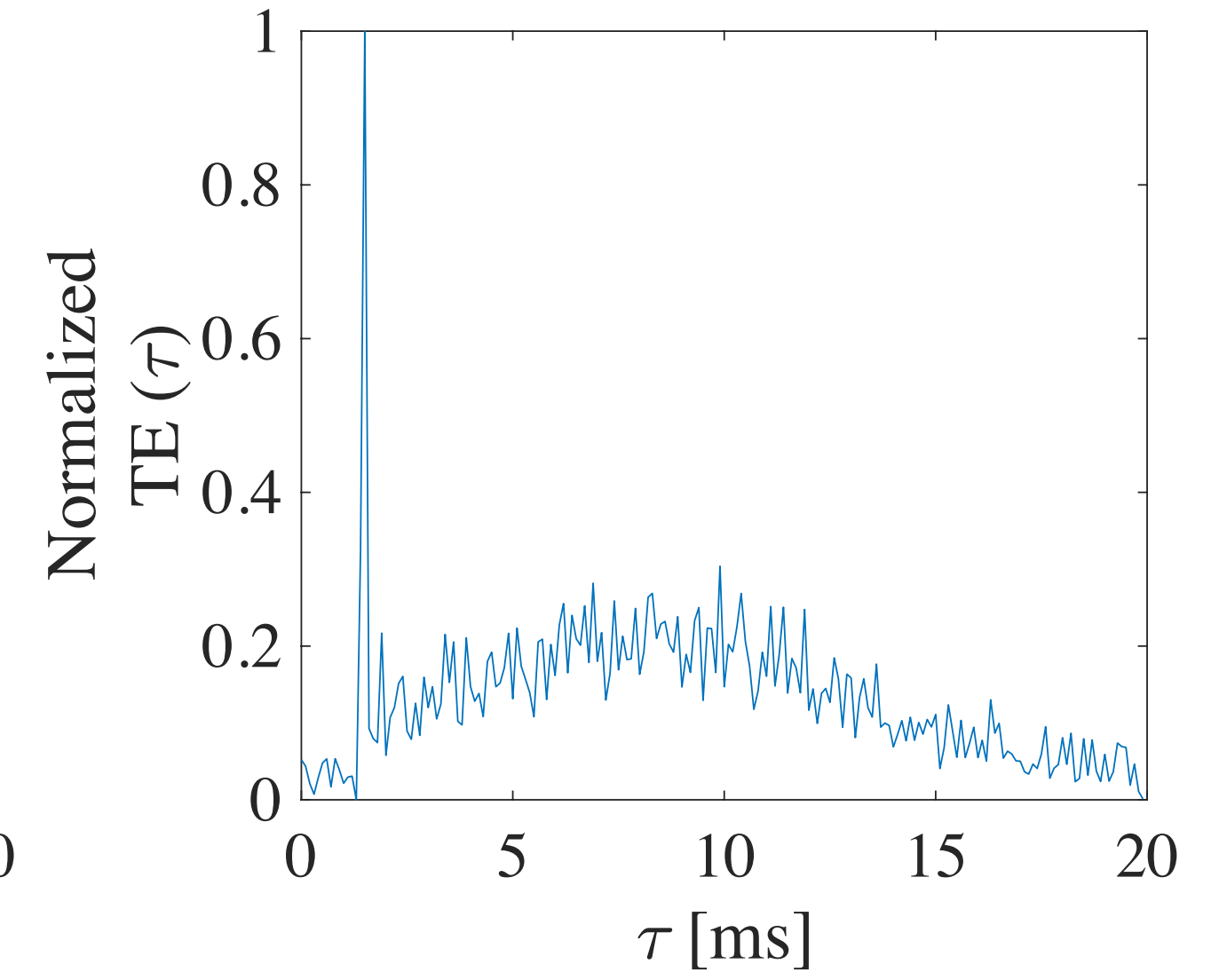
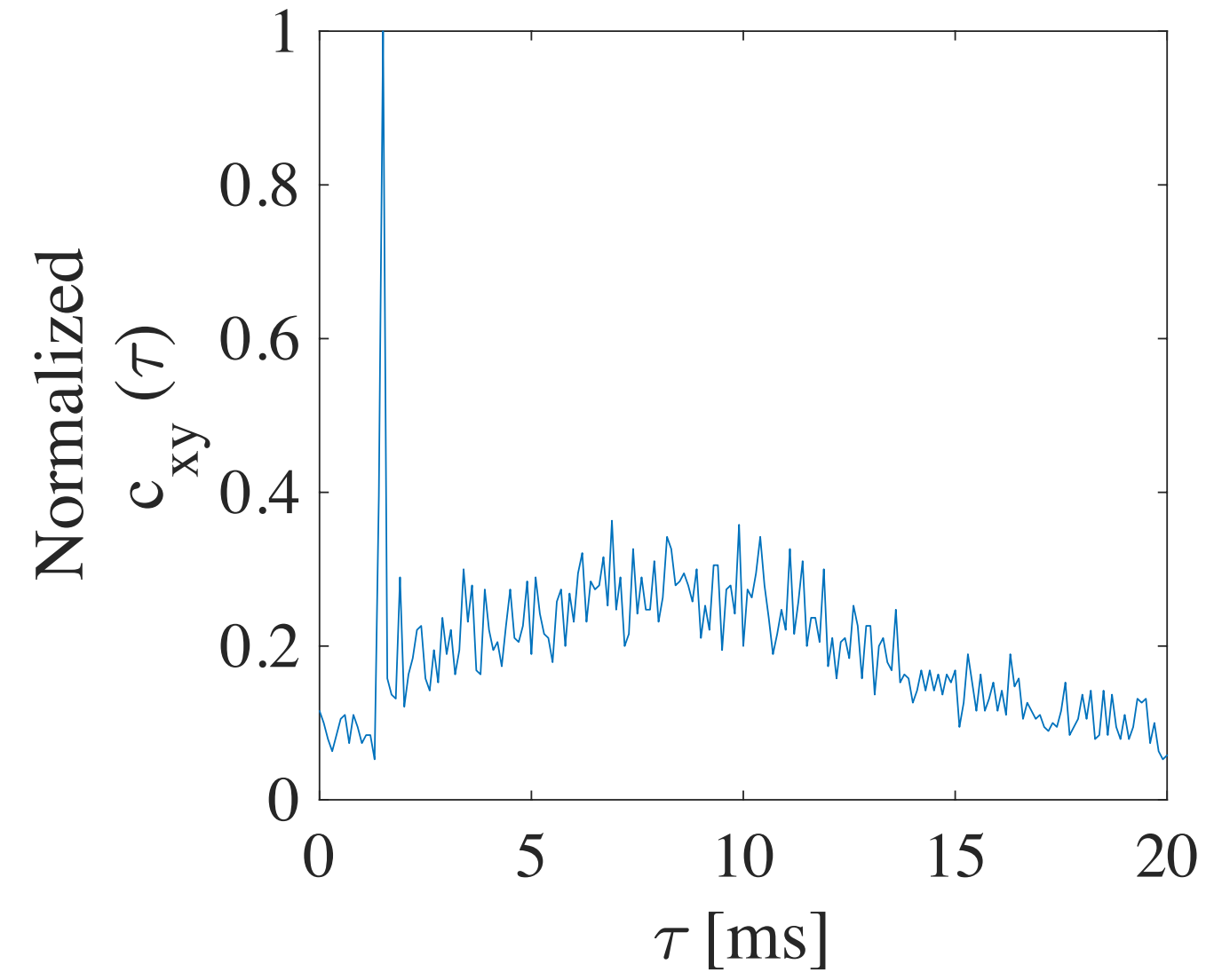
Correlation
and
Causality

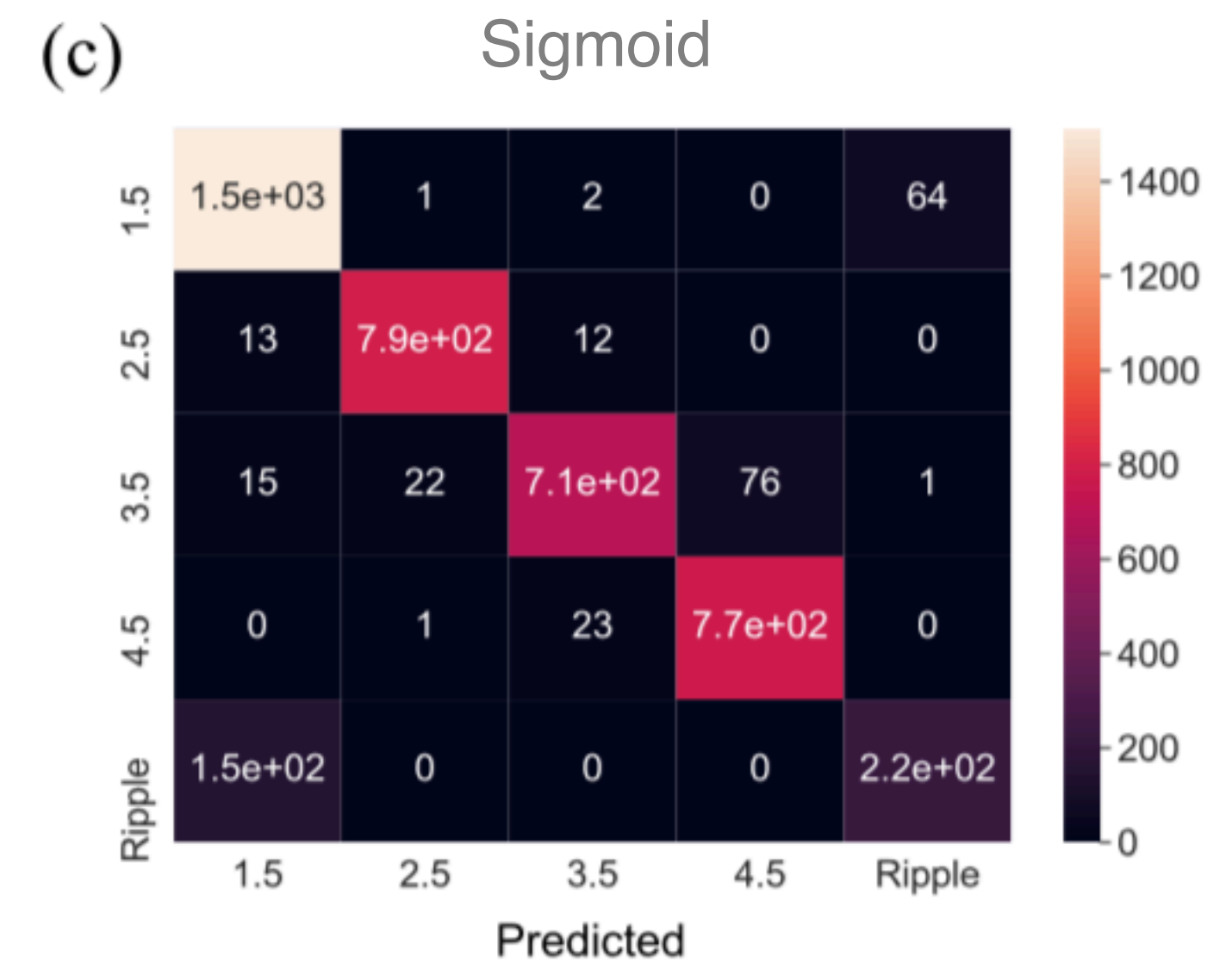
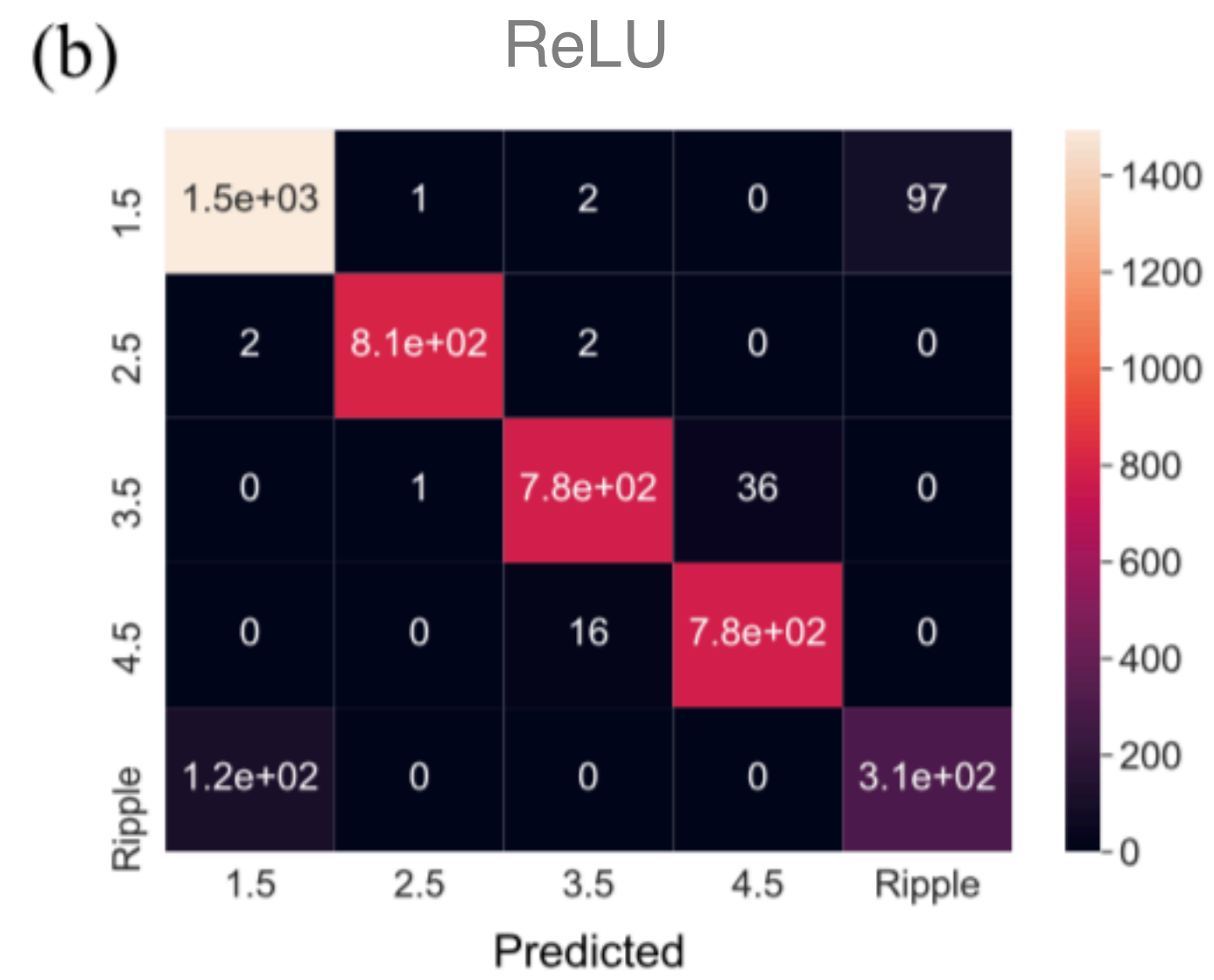
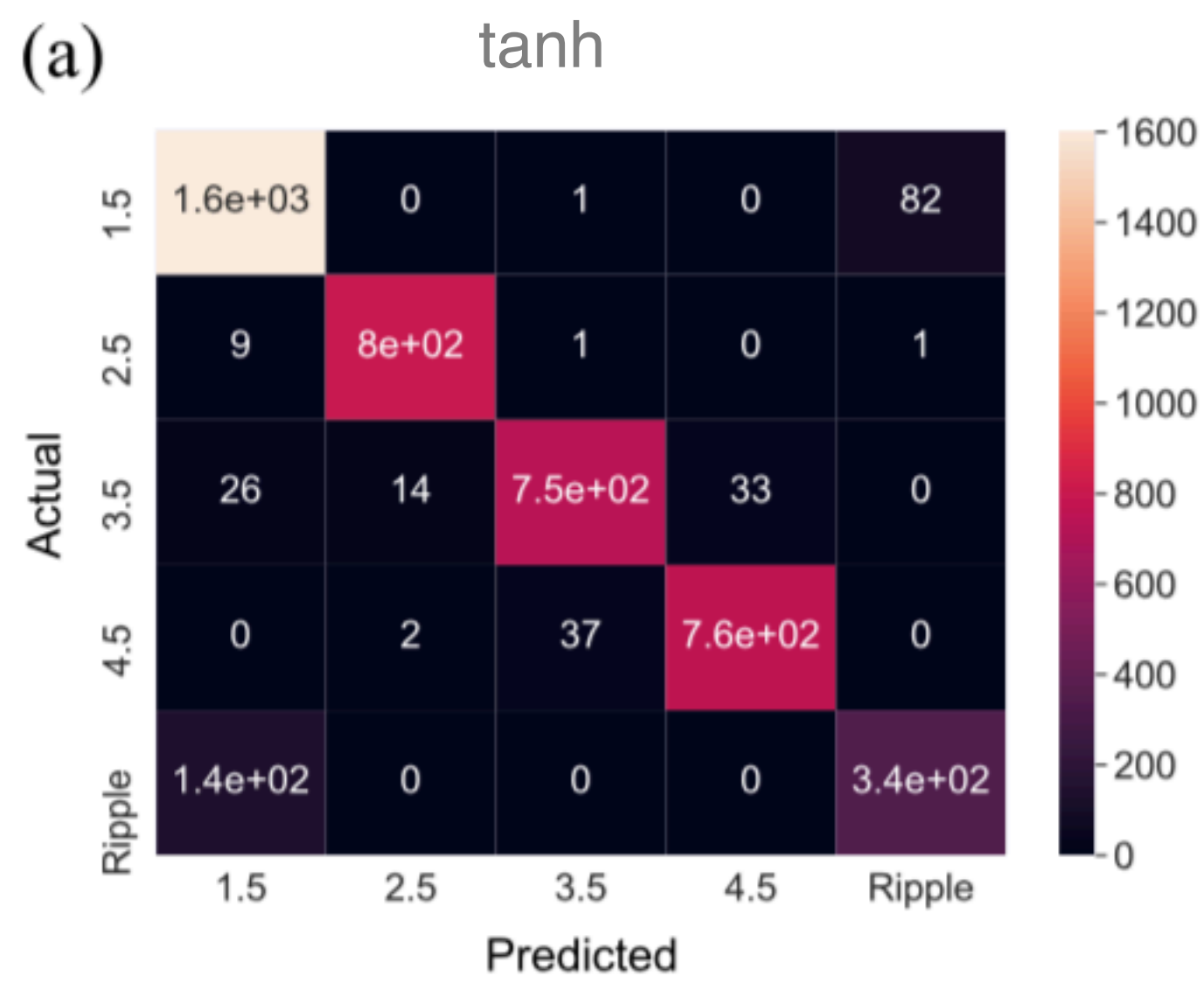


Correlation
but no
Causality



Second peak persists in transfer entropy



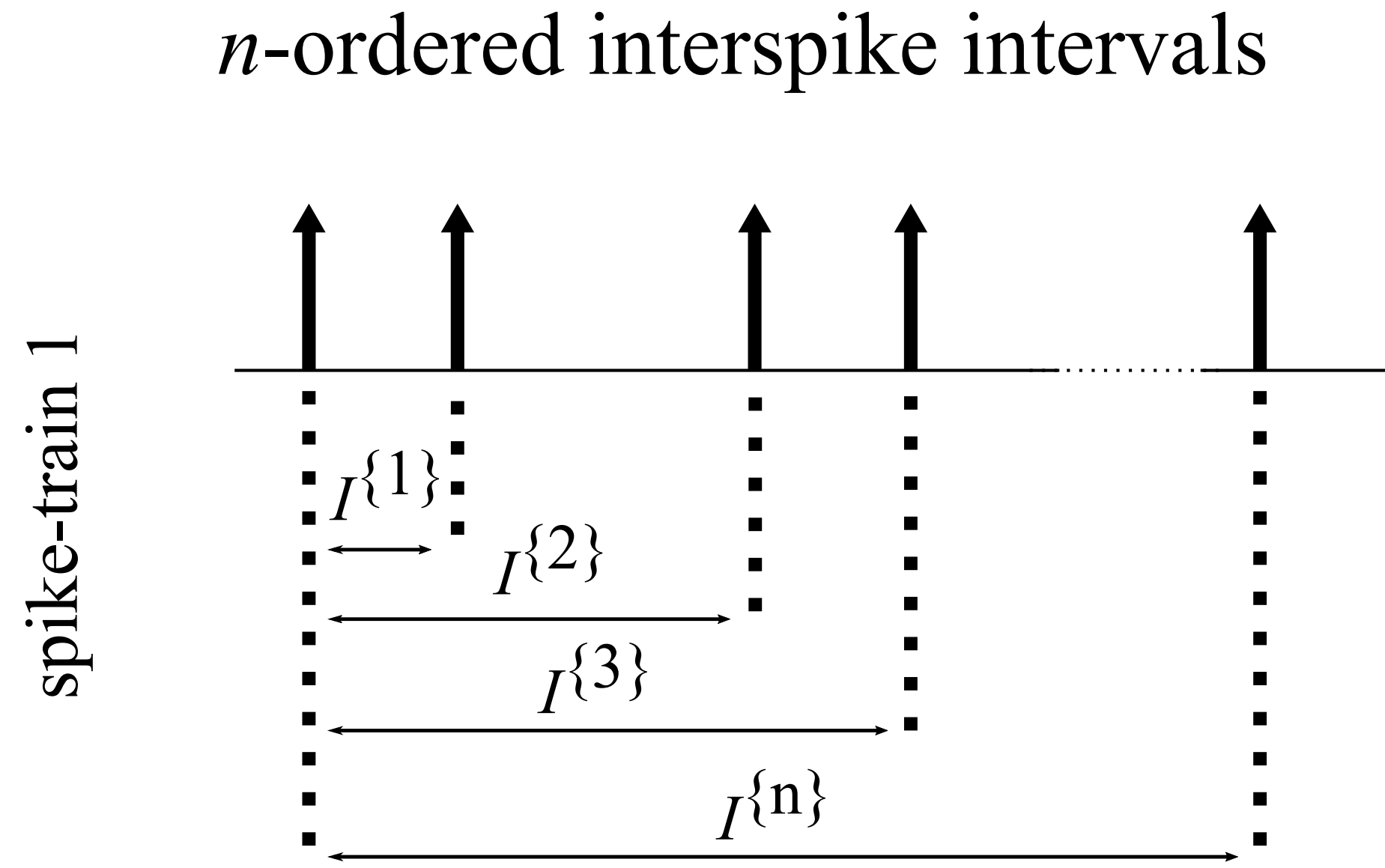


G_h variation					
Activation function	G_h	Precision	Recall	F1-score	MCC
tanh	No connection	0.93	0.99	0.96	0.791
	1.5	0.90	0.80	0.85	
	2.5	0.98	0.80	0.88	
	3.5	0.95	0.75	0.84	
	4.5	0.96	0.76	0.85	
	Ripple	0.48	0.34	0.40	
ReLU	No connection	0.93	0.99	0.96	0.789
	1.5	0.92	0.75	0.83	
	2.5	1.0	0.81	0.89	
	3.5	0.98	0.79	0.87	
	4.5	0.96	0.78	0.86	
	Ripple	0.49	0.31	0.38	
Sigmoid	No connection	0.92	0.99	0.96	0.777
	1.5	0.90	0.76	0.82	
	2.5	0.97	0.79	0.87	
	3.5	0.95	0.71	0.81	
	4.5	0.91	0.77	0.84	
	Ripple	0.50	0.22	0.31	

Confounding factors lower the accuracy
 But how one could improve the classification without TE?

Connection between interval statistics and spike-train statistics

- Perkel , Gerstein, and Moore. *Biophys. J.*, 7(4):419–440, 1967.

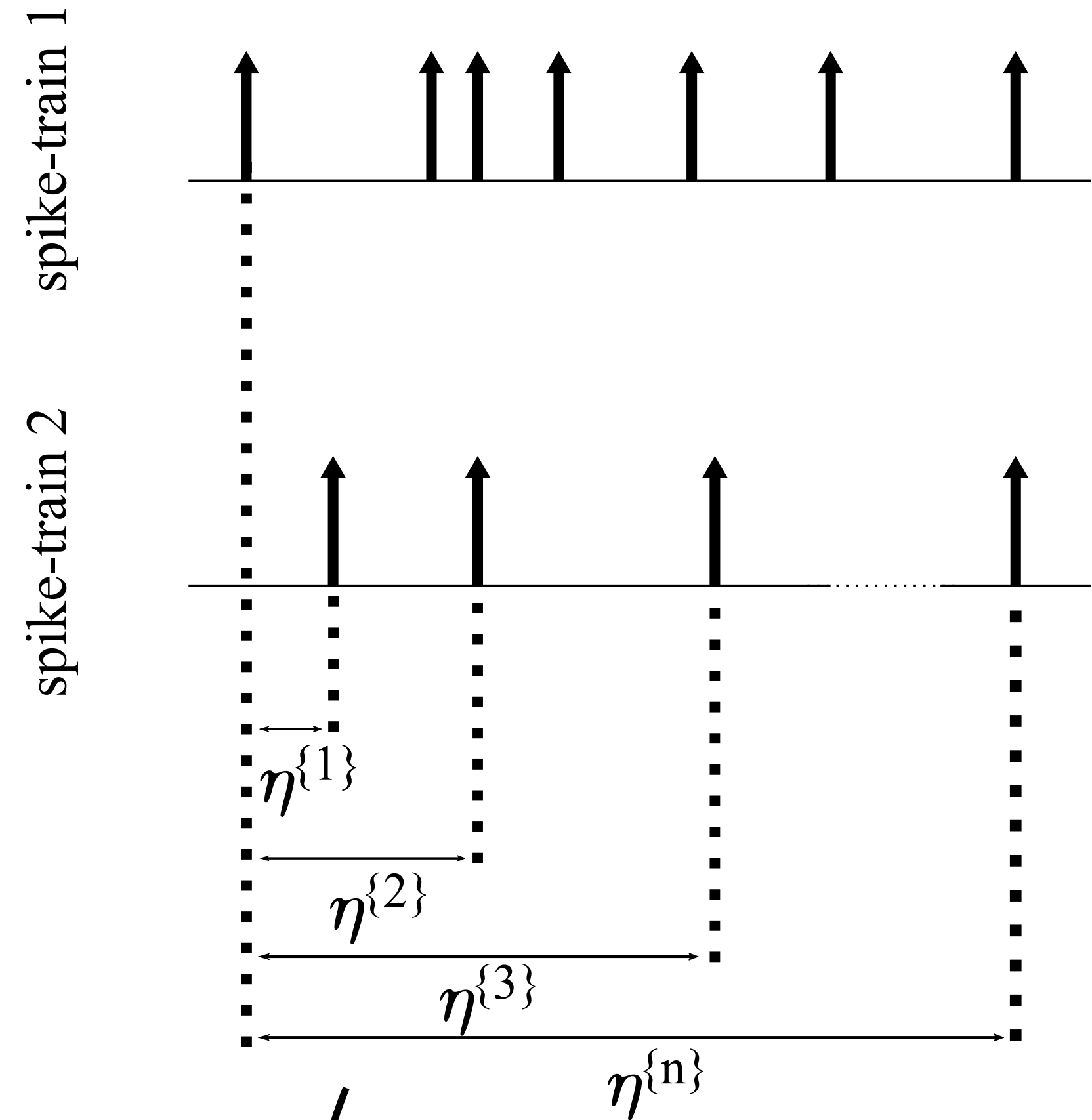


Connected to

Autocorrelation function

$$c_{xx}(\tau) \approx I^1 + I^2 + \dots + I^n$$

n-ordered cross-spike intervals

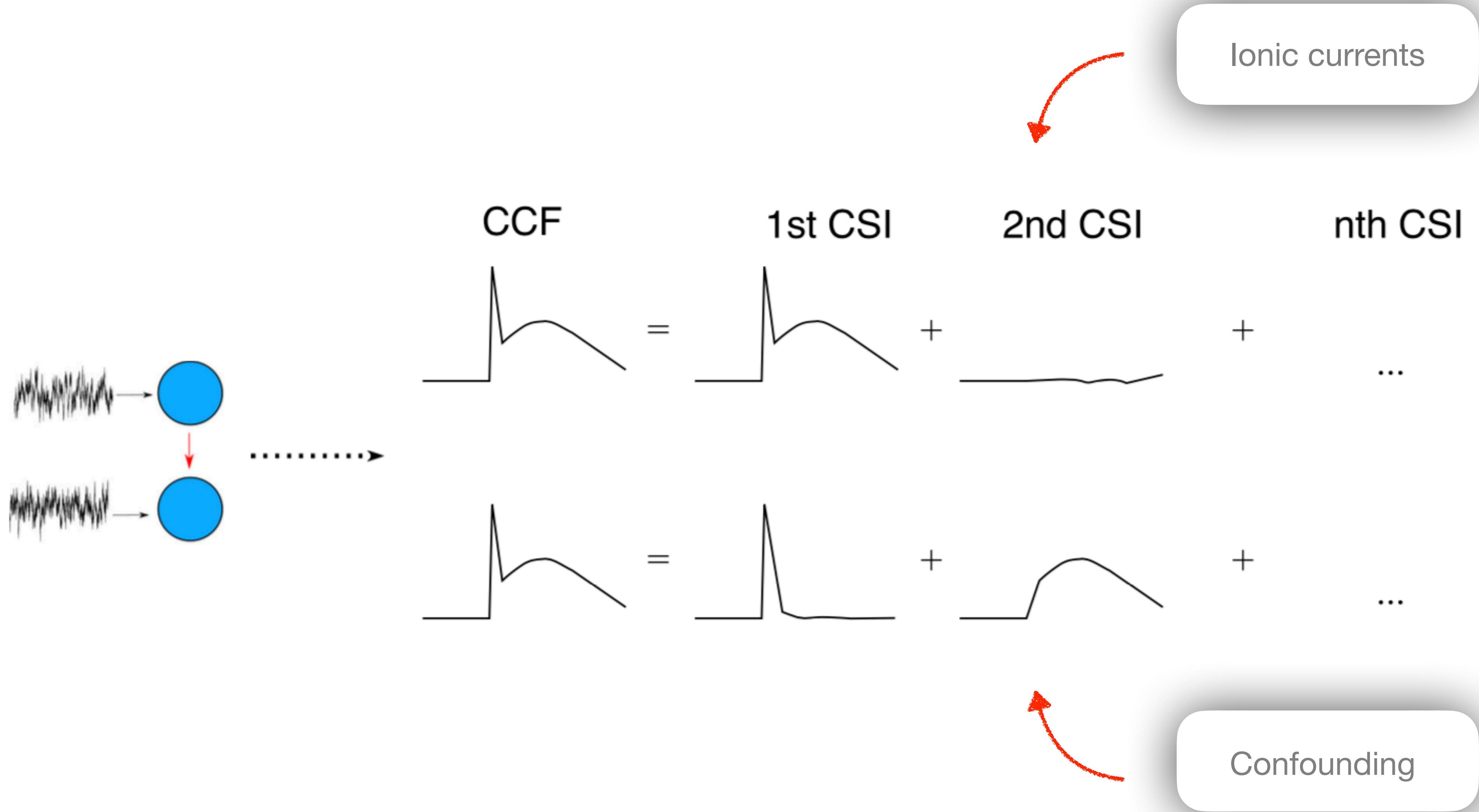


Connected to

Cross-correlation function

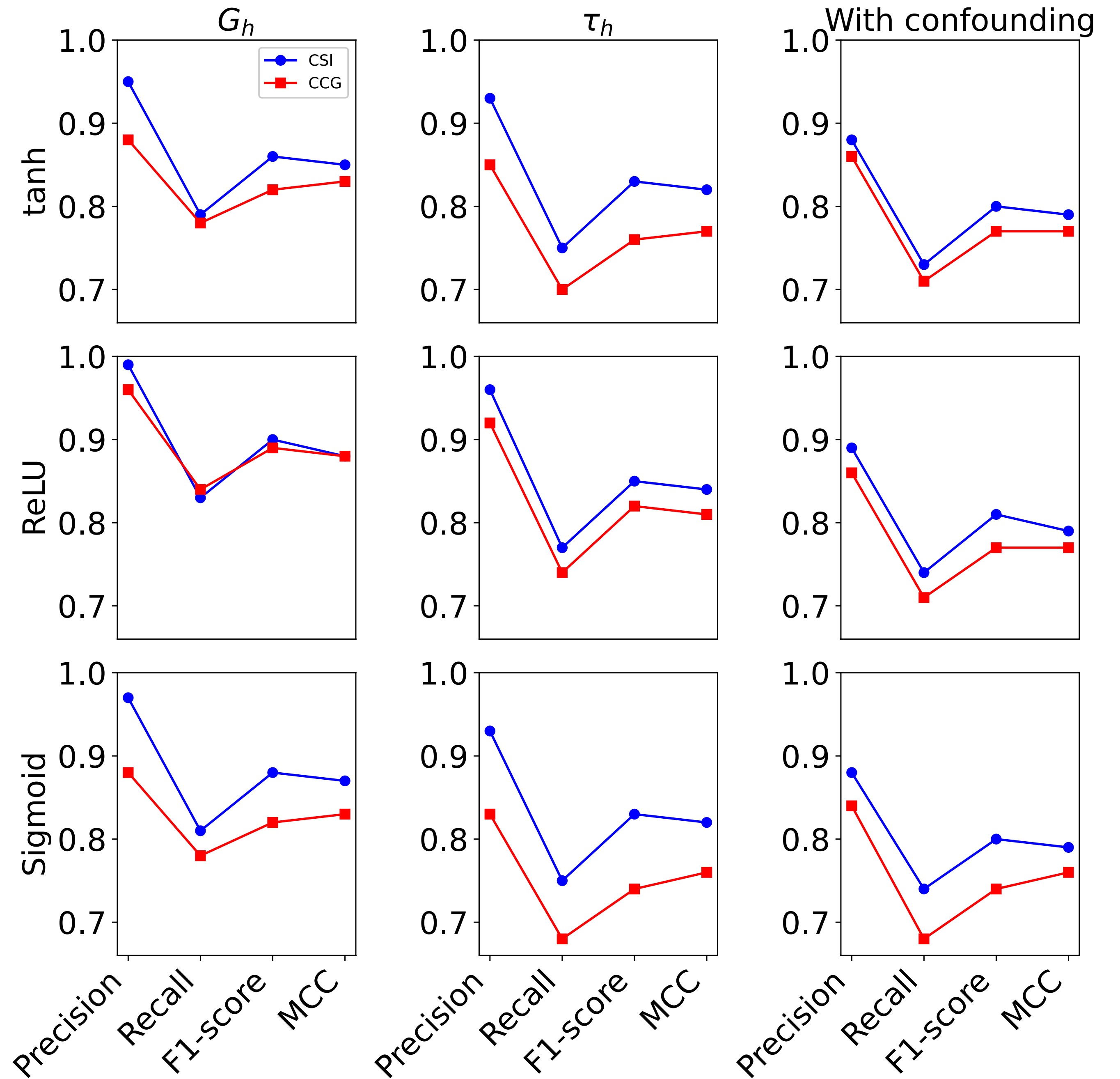
$$c_{xy}(\tau) \approx \eta^1 + \eta^2 + \dots + \eta^n$$

Connection between interval statistics and spike-train statistics

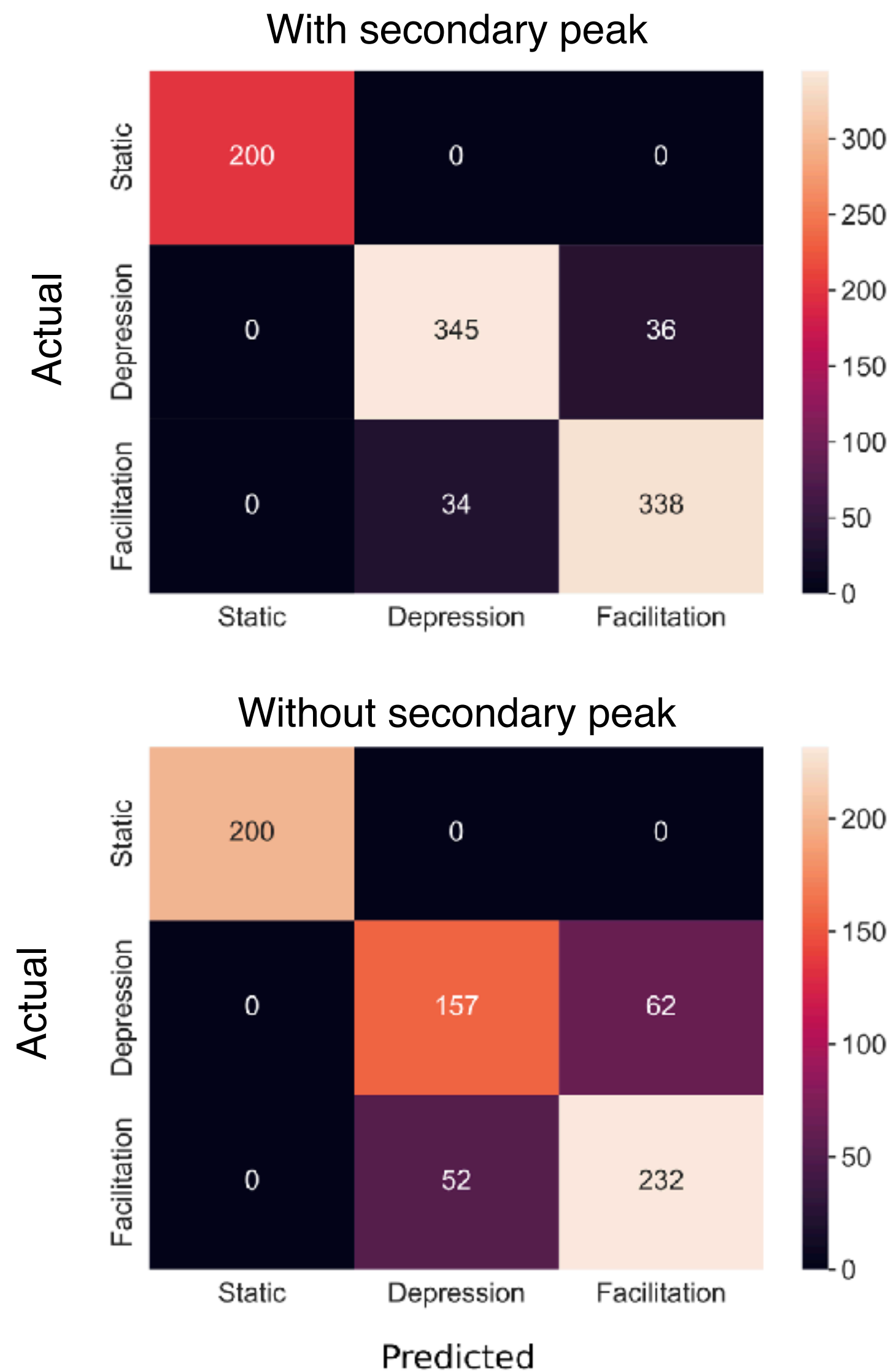
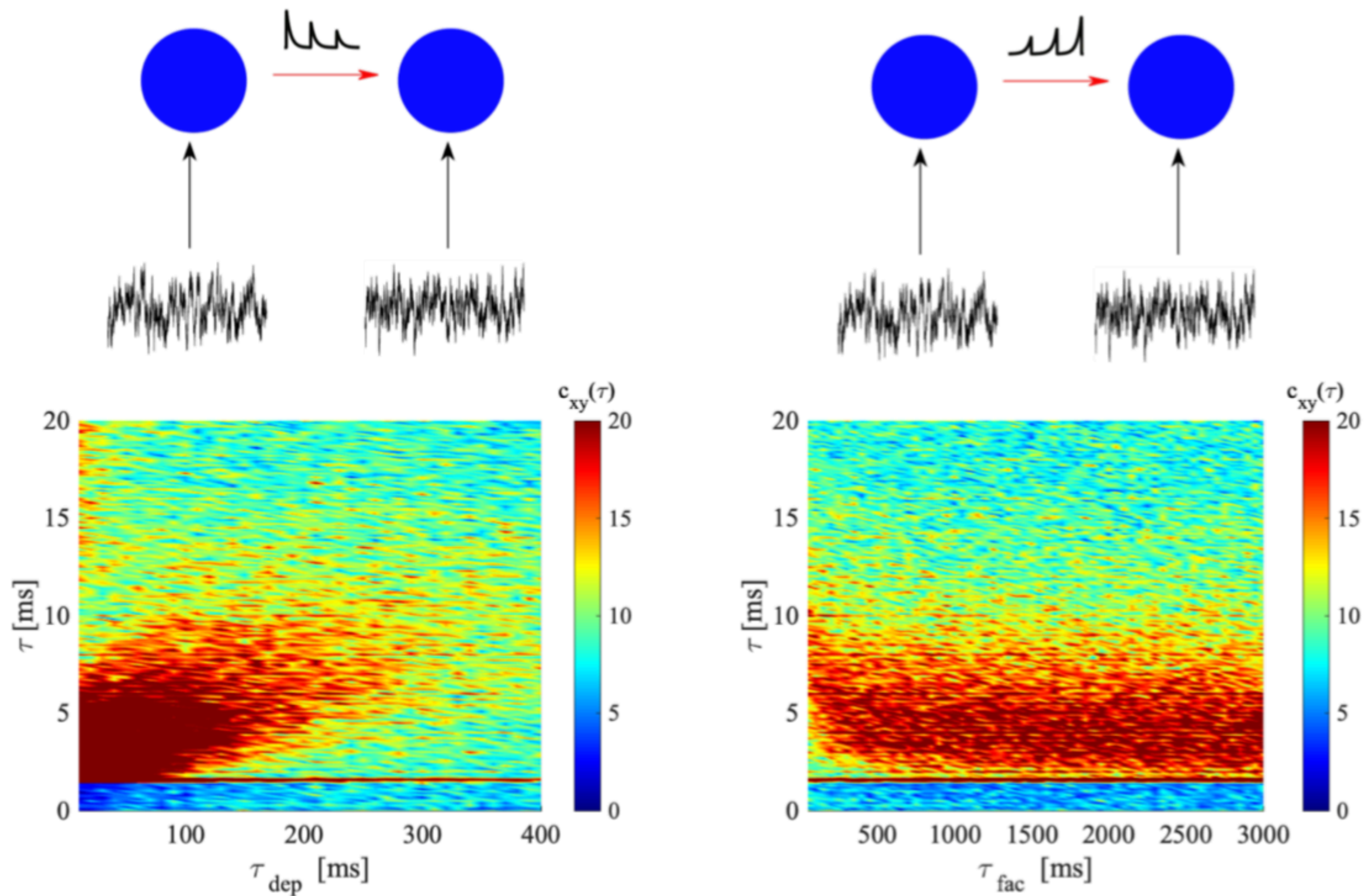


CSIs vs. CCFs

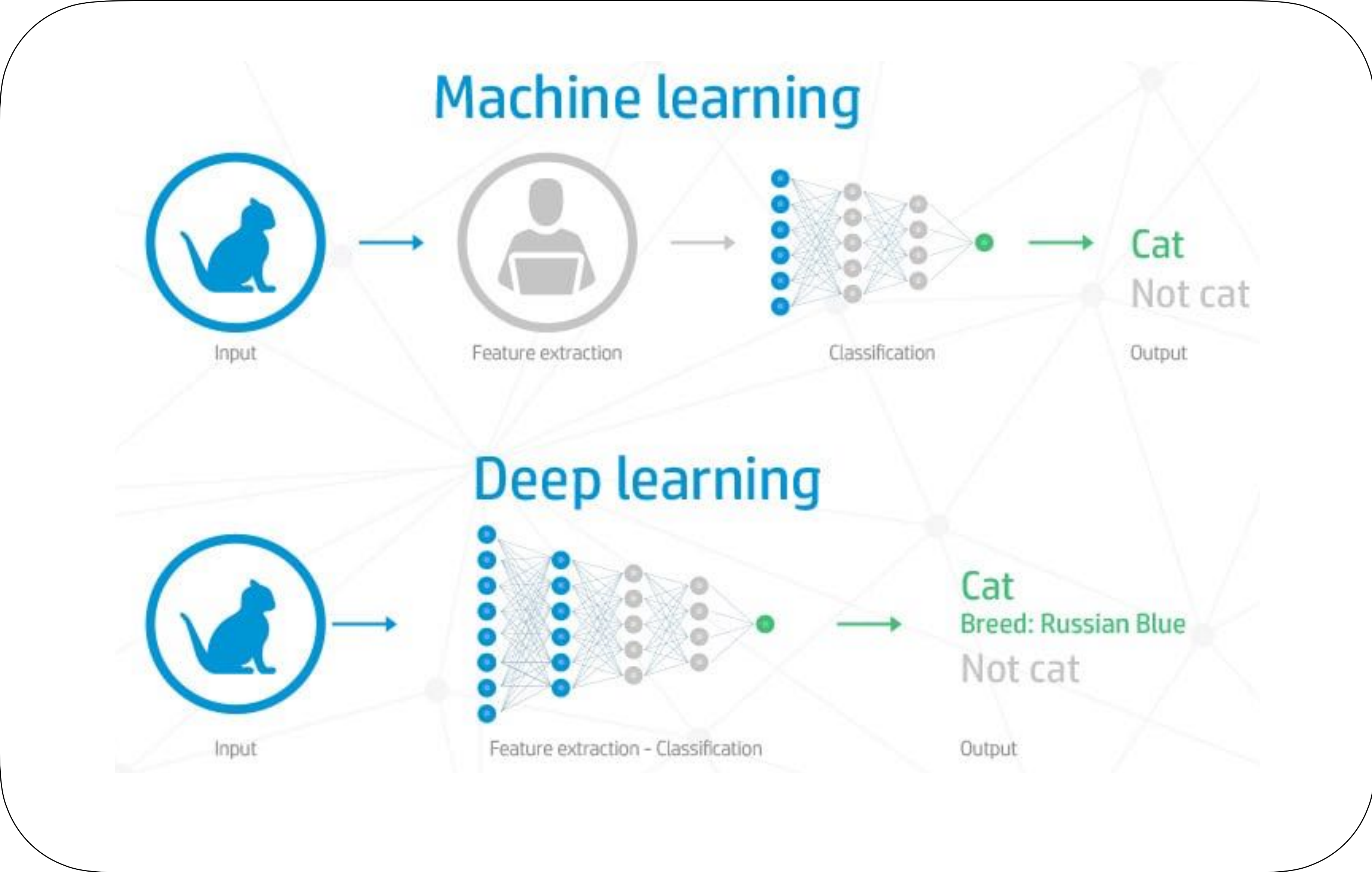
- Classification improves!
- Even with confounding factors



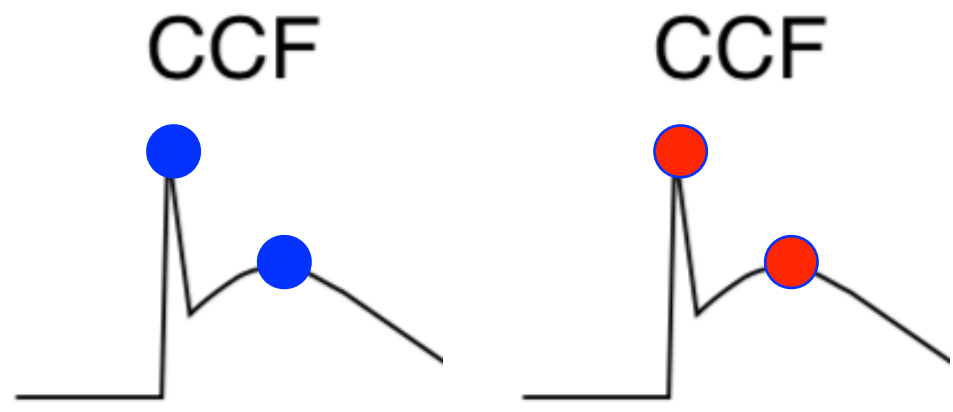
Short-term plasticity can be well classified if CCF includes intrinsic dynamics information



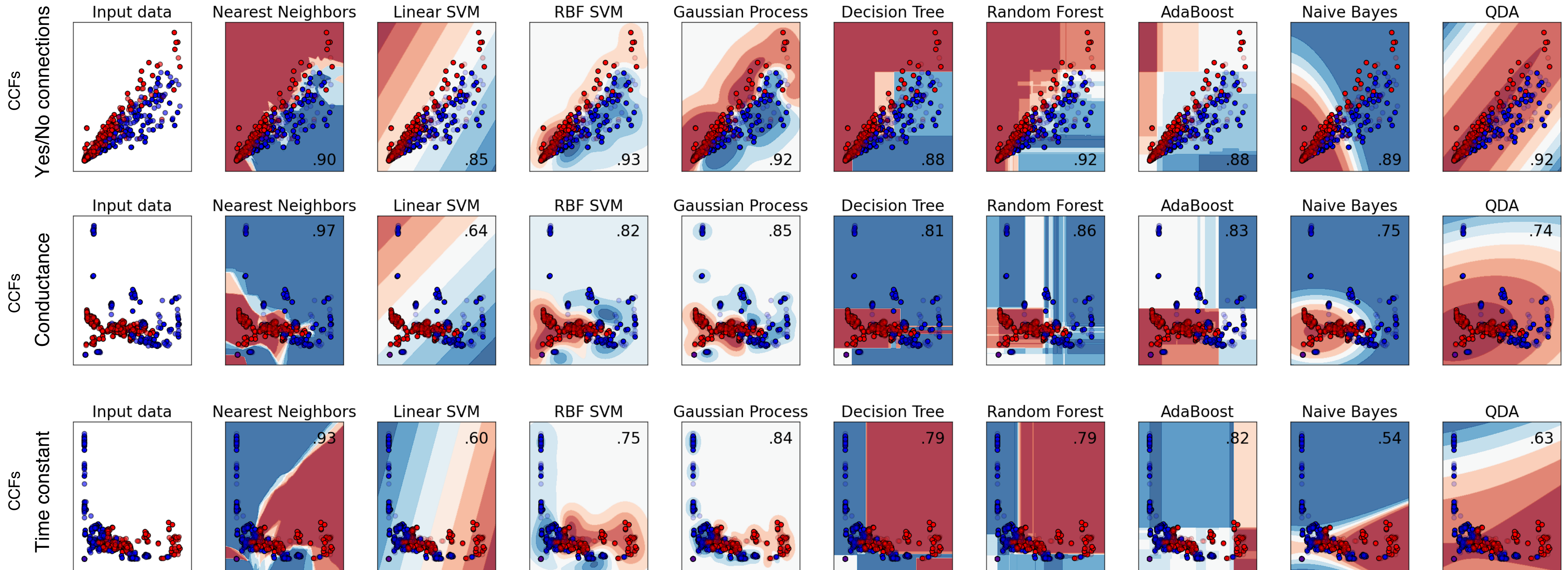
A comparison of several Machine Learning classifiers to distinguish biophysical features



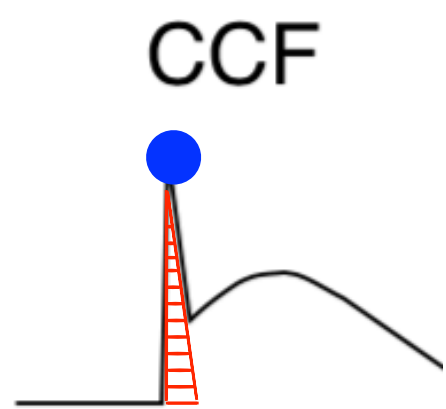
A comparison of several Machine Learning classifiers to distinguish biophysical features



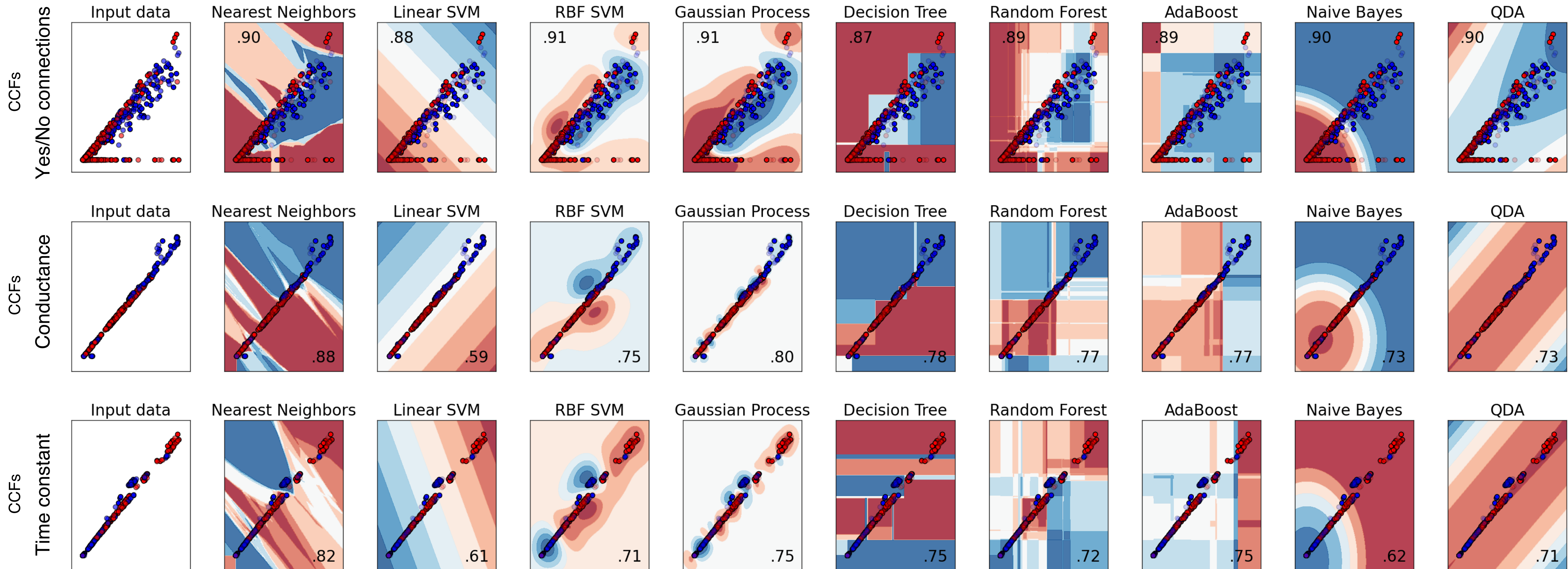
Depending on the classifiers and purpose, two delays can be enough;



A comparison of several Machine Learning classifiers to distinguish biophysical features



Features now are the first peak value and its area

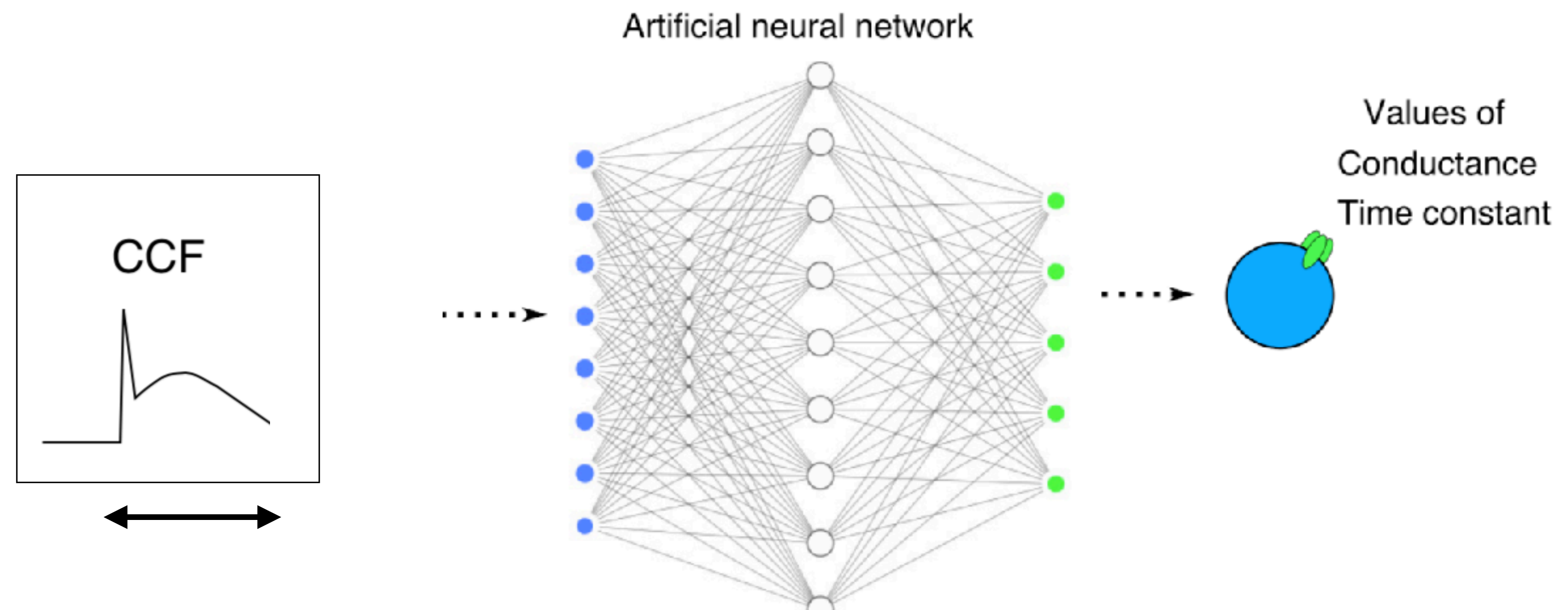


Conclusions

Ionic properties can be extracted from CCFs

... only with information from lags beyond the first peak

Lags with **subthreshold dynamics!!**

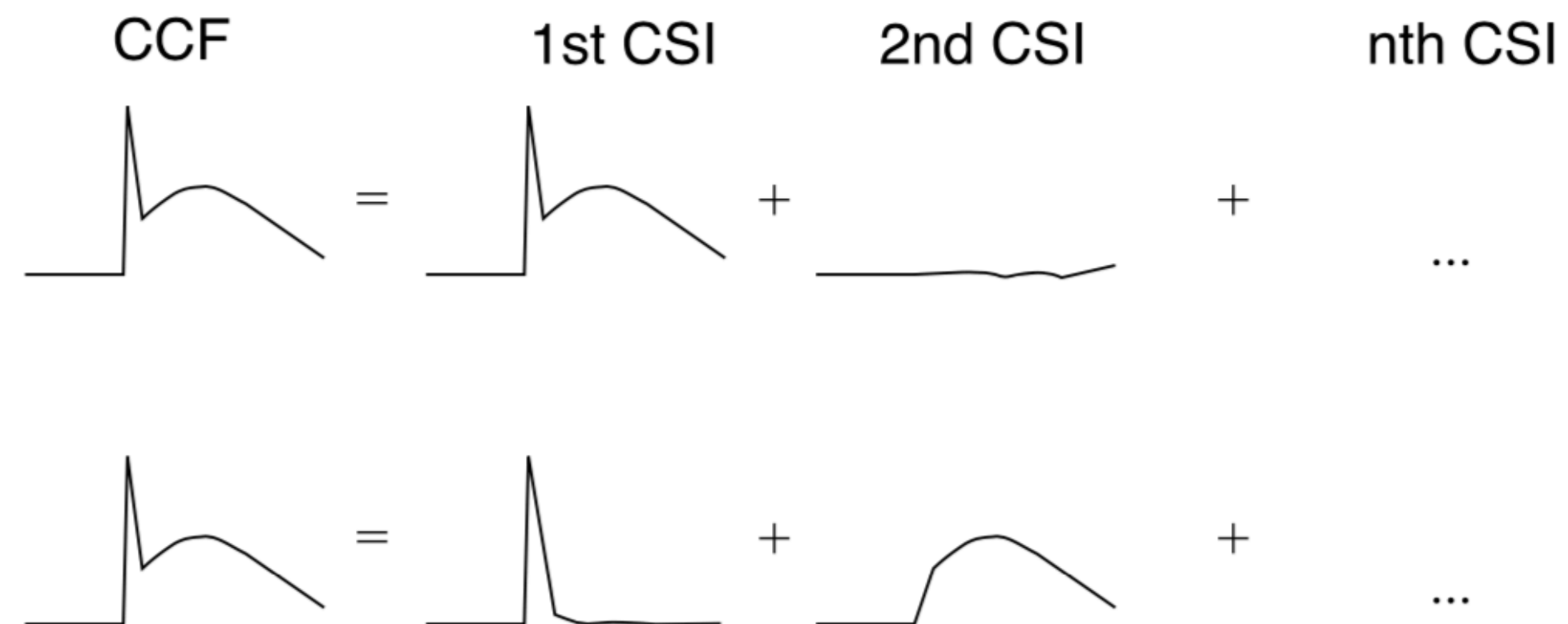


Conclusions

Our results do not require sophisticated information-theoretic metrics such as **transfer entropy**

... but suffer from confounding factors provided by background oscillations.

The cross-spike intervals is an alternative that improves the classification algorithms accuracy.



References

- Toyama, K., Kimura, M., & Tanaka, K., *46(2)*, 191-201, 1981
- English, D. F., McKenzie, S., Evans, T., Kim, K., Yoon, E., & Buzsáki, G., *Neuron*, *96(2)*, 505-520, 2017
- Platkiewicz, J., Saccomano, Z., McKenzie, S., English, D., & Amarasingham, A., *J Comput Neurosci*, *49(2)*, 131-157, 2021
- Rotstein, H. G., *J Comput Neurosci*, *43(3)*, 243-271, 2017
- Perkel, D. H., Gerstein, G. L., & Moore, G. P. *Biophys J*, *7(4)*, 391-418, 1967
- Tsodyks, M., Pawelzik, K., & Markram, H., *Neural Computat*, *10(4)*, 821-835, 1998
- McKenzie, S., Huszár, R., English, D. F., Kim, K., Christensen, F., Yoon, E., & Buzsáki, G., *Neuron*, *109(6)*, 1040-1054, 2021

Many thanks to



Prof Horacio Rotstein

NJIT & Rutgers

- Martin V Ibarra, Universidad de la Patagonia, Argentina
- Sam MacKenzie, University of New Mexico, USA
- Zach Saccomano, CUNY Graduate Center, USA



STG Lab

NJIT & Rutgers



(HGR) DMS-1608077

Thank you very much!
