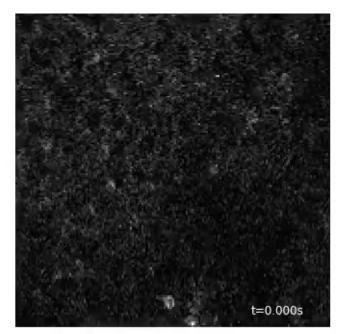
# Machine Learning Methods for Neural Data Analysis

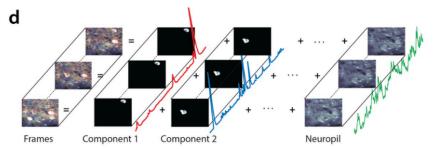
Lecture 5: Deconvolution with a Point Process Prior

## **Announcements**

- Lab 2 due Thursday, 11:59pm.
- My office hours today 1-2pm.

## 2 photon calcium imaging





$$\begin{split} Y &= U^T C + u_0 c_0^\top + \epsilon \\ U &\in \mathbb{R}^{N \times P} \ C \in \mathbb{R}_+^{N \times T} \ u_0 \in \mathbb{R}^P \ c_0 \in \mathbb{R}^T \\ \epsilon_{pt} &\sim \mathcal{N}(0, \sigma^2) \end{split}$$

Sue Ann Koav and David Tank

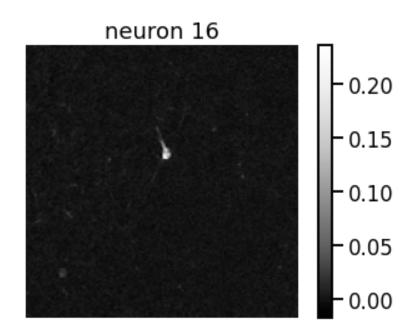
## **Optimizing the footprints**

Given an estimate of the fluorescence trace  $c_n$ , the optimal footprint is,

$$u_n = \frac{R_n c_n}{\|R_n c_n\|}$$

where  $R_n \in \mathbb{R}^{P \times T}$  is the residual.

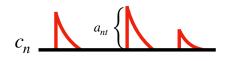
This is assuming a uniform prior on unit-norm footprints.



#### **Prior on calcium traces**

We had an exponential prior on spike amplitudes, i.e. jumps in the fluorescence:

$$p(c_n) = \prod_{t=1}^{T} \text{Exp}(a_{nt})$$
$$= \prod_{t=1}^{T} \text{Exp}(c_{nt} - e^{-1/\tau}c_{n,t-1}; \lambda_n)$$

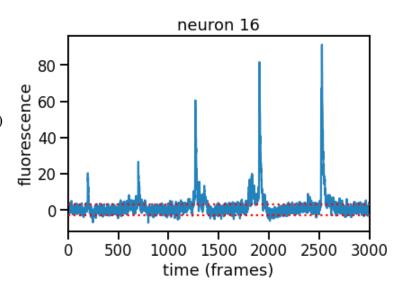


### **Objective for optimizing traces**

Combining the log likelihood and the log prior lead to the following objective:

$$\begin{split} \mathcal{L}(c_n) &= -\frac{1}{2\sigma^2} \|c_n - \mu_n\|_2^2 + \lambda_n \sum_{t=1}^T (c_{nt} - e^{-1/\gamma} c_{n,t-1}) \\ &= -\frac{1}{2\sigma^2} \|c_n - \mu_n\|_2^2 + \lambda_n \|Gc_n\|_1. \end{split}$$

where  $\mu_n = R_n^{\mathsf{T}} u_n$  is the residual projected onto the spatial factor for this neuron and  $Gc_n = a_n$  are the spike amplitudes.



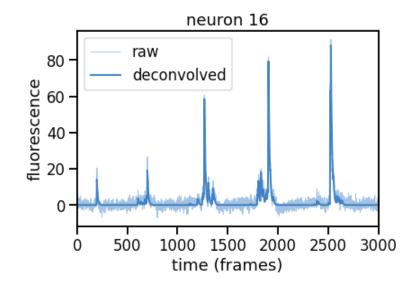
### **Objective for optimizing traces**

• Maximizing  $\mathcal{L}(c_n)$  is equivalent to solving the following convex optimization problem,

$$\begin{split} \hat{c}_n &= \arg\min_{c_n} \|Gc_n\|_1 \\ \text{s.t. } \|c_n - \mu_n\|_2 &\leq \theta \\ c_n &\geq 0, \end{split}$$

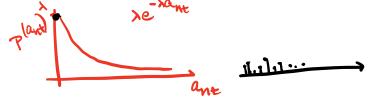
for some threshold  $\theta$ .

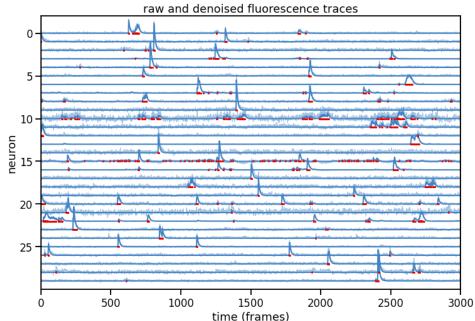
- Idea: set  $\theta = (1 + \epsilon)\sigma\sqrt{T}$ .
- Since G is a sparse banded matrix, we can solve this in O(T) time with CVXpy. (Lab 3)



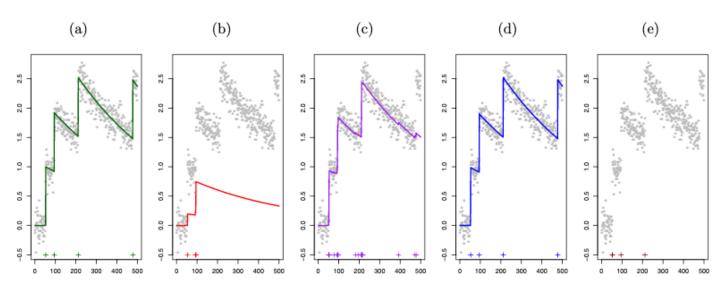
#### Limitations of "L1" deconvolution

- The resulting traces are sparse but still overestimate the number of spikes.
- Likewise, the amplitudes are penalized, shrinking our estimates.
- Many of these issues can be traced back to the exponential prior distribution on amplitudes.





## Limitations of "L1" deconvolution



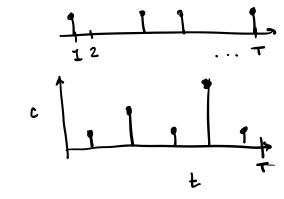
# **Agenda**

- 1. A point process prior
- 2. Efficient optimization via dynamic programming

# Point processes

(More to come next week)

$$\chi \in \{1,...,T\}$$
 eg  $\chi = \{t_k\}_{k=1}^{k}$   
 $1 \le t_1 \le t_2 \le ... \le t_k \le T$ 



# A point process prior for spikes

point process prior for spikes
$$\chi = \left\{ \left( t_{k}, c_{t_{k}} \right) \right\}_{k=1}^{K} \qquad \downarrow$$

$$\gamma(\chi) = \left[ \gamma(t_{i}) \prod_{k=2}^{K} \left[ \gamma(t_{k}) t_{k-1} \right] \times \gamma(t_{k+1}) \right] \times \left[ \prod_{k=1}^{K} \gamma(c_{t_{k}}) \right]$$

Geom(t, |p) Geom(t, -t, -1p) I Geom(t, -t, (p)

Unit (Celi (O, Cmax))

# From points to calcium traces

Goodbye convolutional model!

$$P(X) \propto \left(\frac{f}{(1-\rho)c_{max}}\right)^{K} \qquad \chi \rightarrow \left(c_{1} \cdots c_{T}\right)$$

$$c_{+}(\chi) = \begin{cases} c_{+u} & \text{if } t \in \S_{+u} \end{cases}$$

$$c_{+}(\chi) = \begin{cases} c_{+v} & \text{if } t \in \S_{+v} \end{cases}$$

# **Deriving the prior probability**

# **Rewriting the likelihood**

In terms of spike times and amplitudes

$$\begin{aligned} \log P(M|X) &= -\frac{1}{2\sigma^2} \sum_{t=1}^{T} (M_t - C_t(X))^2 \\ &= -\frac{1}{2\sigma^2} \sum_{k=0}^{K} \sum_{t=k}^{t_{k0}-1} (M_t - C_{t_k} e^{-(t-t_k)/\tau})^2 \end{aligned}$$

# **Defining our objective**

$$\frac{1}{2\sigma^{2}} \sum_{k=0}^{K_{1}} \sum_{t=k}^{t_{k}} \left( \mu_{t} - c_{t_{k}} e^{-(t-t_{k})/t} \right)^{2} + \eta_{K} + const$$

# Optimizing the spikes with dynamic programming

## This looks hard...

- $\mathcal{X}$  is a set of unknown cardinality
- Each entry consists of a time and an amplitude
- Idea: pull the last spike out of  ${\mathcal X}$

# Rewrite the objective in terms of $\mathcal{X}_{[1,t)}$ , t, and $c_t$

$$\mathcal{A}(\chi) = \mathcal{A}(\chi_{[i,t)} \cup \{l^{+}, c_{+}\})$$

$$= \mathcal{A}(\chi_{[i,t)}, +, c_{+})$$

$$= LO(P(M_{l,t}), \chi_{l',t}) - \frac{1}{2\sigma^2} \sum_{t'=t}^{T} (M_{t'} - c_t e^{-(t'-t)/\tau})^2 + \eta$$

# Optimizing wrt the amplitude $c_t$

# Optimizing wrt previous spikes, $\mathcal{X}_{[1,t)}$

It's the same problem on a subset of data!

# Optimizing wrt the time of the last spike, t

# Putting it all together

## **Considerations**

- Complexity?
- Hyperparameters?

## Conclusion

- Point processes offer a truly sparse prior.
- While they seem to pose a hard **combinatorial optimization problem**, in fact we can solve for the optimal spikes in polynomial time.
- This is what the Allen Institute currently uses in their 2P analysis pipelines.

## **Further reading**

- Ch 3 of the course notes (coming later today)
- Jewell, Sean, and Daniela Witten. 2018. "Exact Spike Train Inference via  $\ell_0$  Optimization." The Annals of Applied Statistics 12 (4): 2457–82.