

# Lecture 3

## Spike Stimulus Analysis

# About me

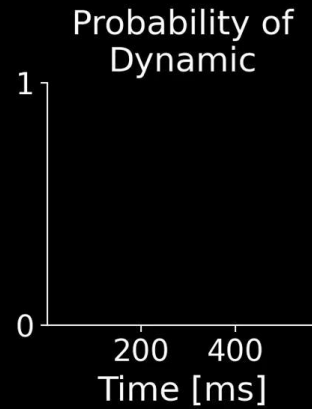
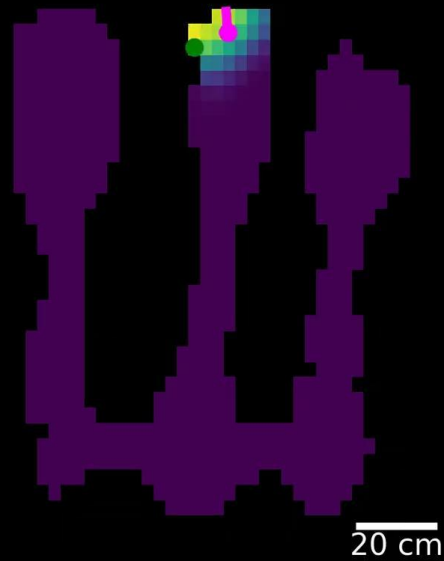
Computational Research Scientist in the lab of Dr. Loren Frank.

Postdoc with Dr. Uri Eden in the Department of Mathematics and Statistics at Boston University

Ph.D. in Computational Neuroscience from Boston University working with Drs. Daniel H. Bullock and Earl K. Miller.

Undergraduate degrees in Mathematics and Philosophy

# My Research



Animal's  
Position

Decoded  
Position

# Jupyter Notebooks and Google Colab

<https://github.com/edeno/ncbs-neural-circuits-navigation>







Click on the **Open in Colab** button

## Neural Circuits for Navigation

Course materials for **Neural Circuits for Navigation: Anatomy, Physiology, and Computational Methods** at NCBS.

**Instructors:** Eric Denovellis and Abhilasha Joshi

### Course Structure

Week	Topic	Notebook	Colab
1	Loading and Exploring NWB Data	<a href="#">01-loading-nwb-data.ipynb</a>	 <a href="#">Open in Colab</a>
2a	Spike-Stimulus Analysis	<a href="#">02a-spike-stimulus-analysis.ipynb</a>	 <a href="#">Open in Colab</a>
2b	Poisson Regression / GLMs	<a href="#">02b-poisson-regression.ipynb</a>	 <a href="#">Open in Colab</a>
3	Spectral properties of LFP	<a href="#">03-spectral-lfp.ipynb</a>	 <a href="#">Open in Colab</a>
4	Decoding + Open data use and visualization	<a href="#">04-decoding.ipynb</a>	 <a href="#">Open in Colab</a>
5	Clusterless Decoding Approaches	<a href="#">05-clusterless-decoding.ipynb</a>	 <a href="#">Open in Colab</a>
6	Project + Presentation	—	—



# Overall: How do we understand and characterize brain data and behavior with statistics?

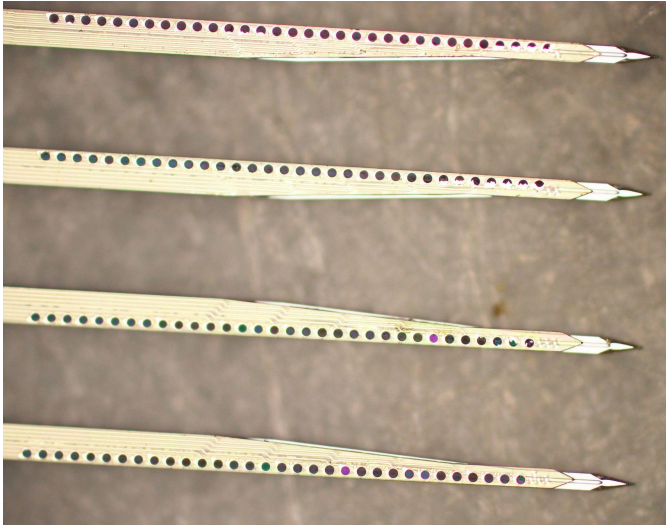
Next four weeks:

1. Relating spikes to stimuli (Encoding)
2. Basic signal processing
3. Relating Local Field Potentials (LFPs) to stimuli and analyzing oscillations
4. Decoding brain states from spikes

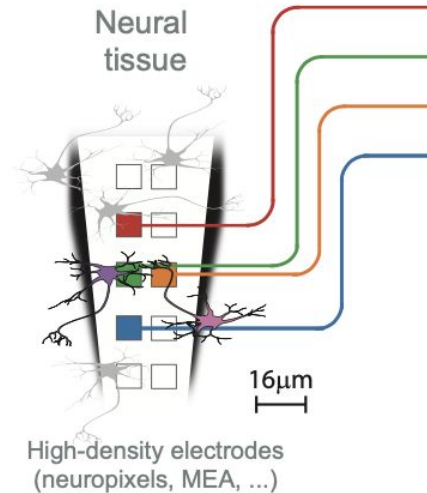
Focus on extracellular recordings (but of course there are many types of data including calcium imaging, functional neuroimaging (fMRI), EEG, etc)

# Overview: what do we record when we record extracellularly in the brain?

Stick electrode in brain



4 shank, 32 channel polymer probes

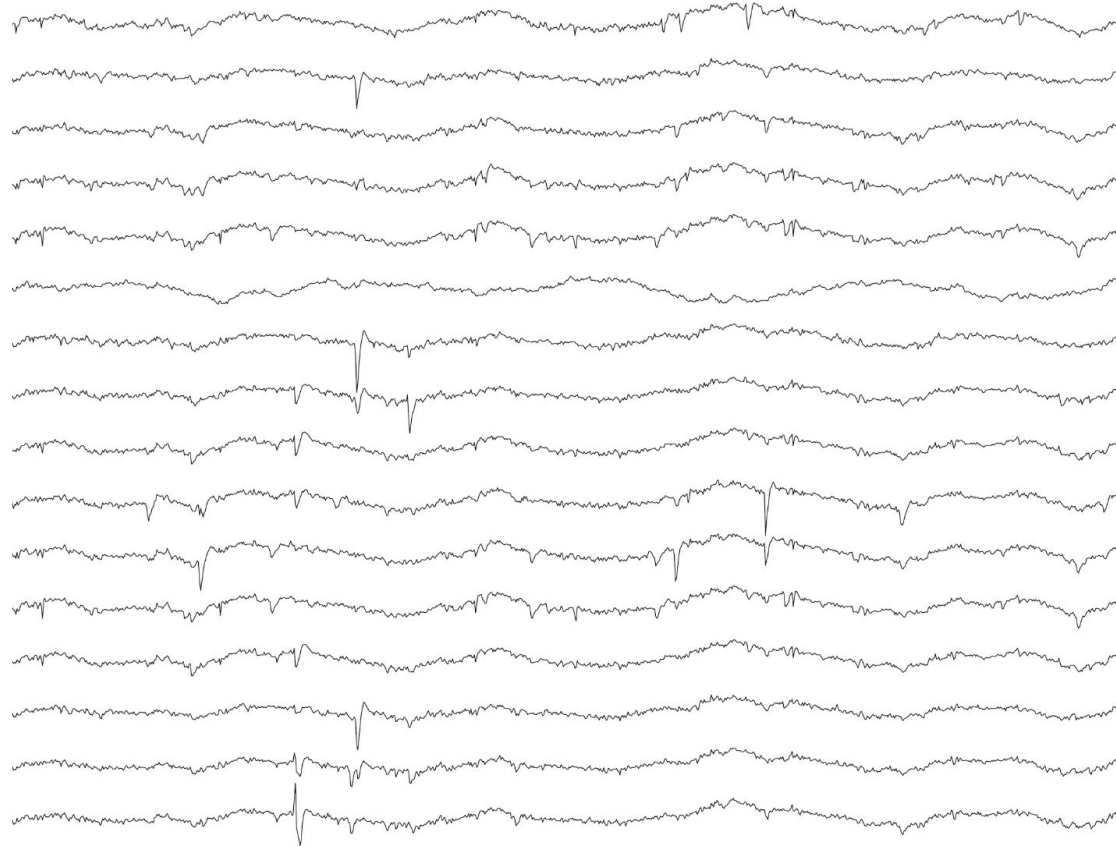


100s of neurons

Einevoll et al. 2012;  
Chung et al. 2019

# Extracellular signals

16 channel  
trace from  
polymer probes

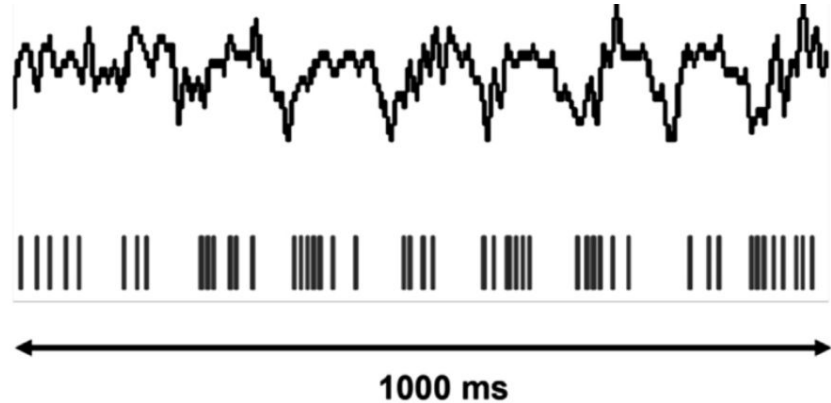


# Two Types of Data: Spikes and Local Field Potentials

Two different types of data:

- Spikes (list of times, point process)
- Local Field Potentials (continuous)

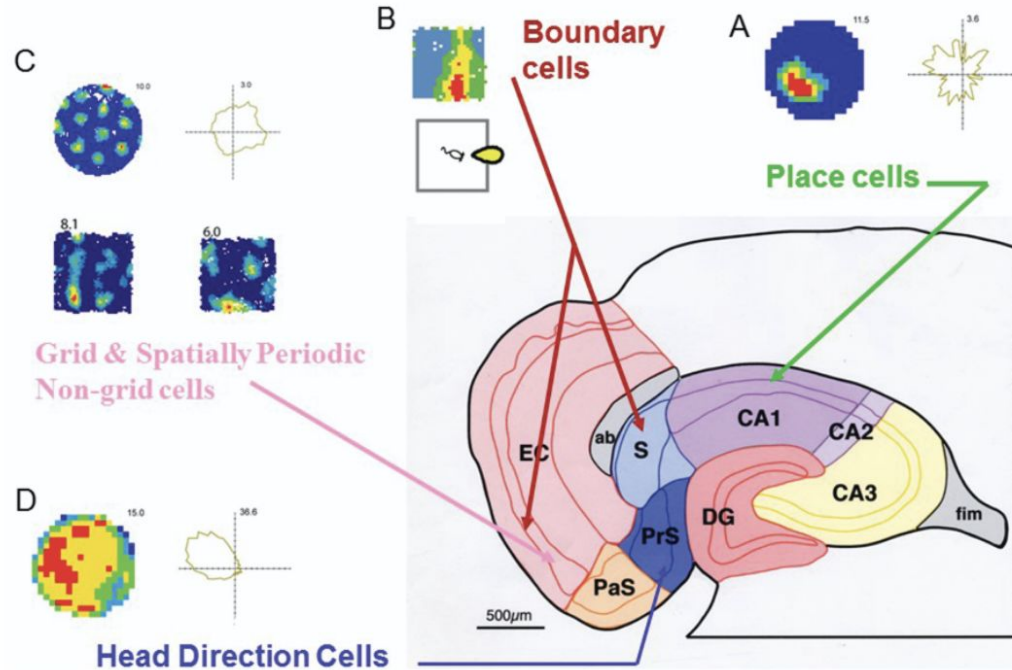
We are going to first focus on spikes (LFPs next week)



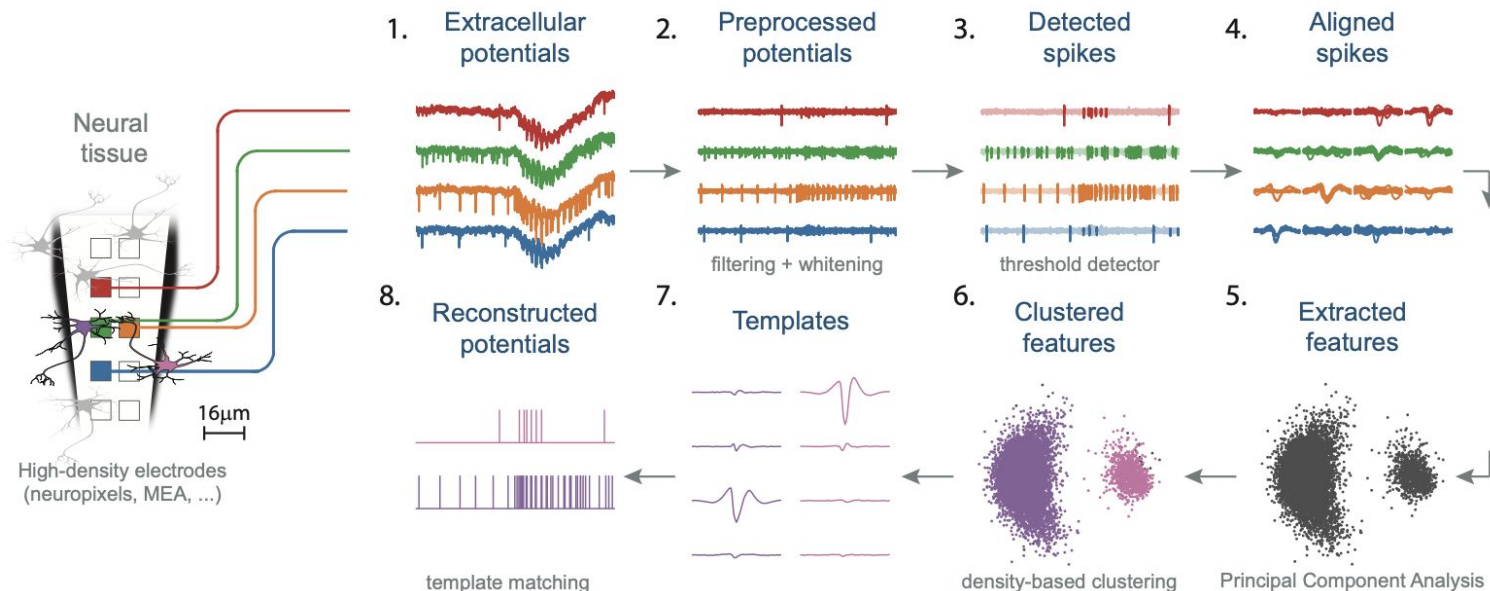


# Motivation: We want to relate this to some representation of the external world

Spatial cells in the hippocampal formation

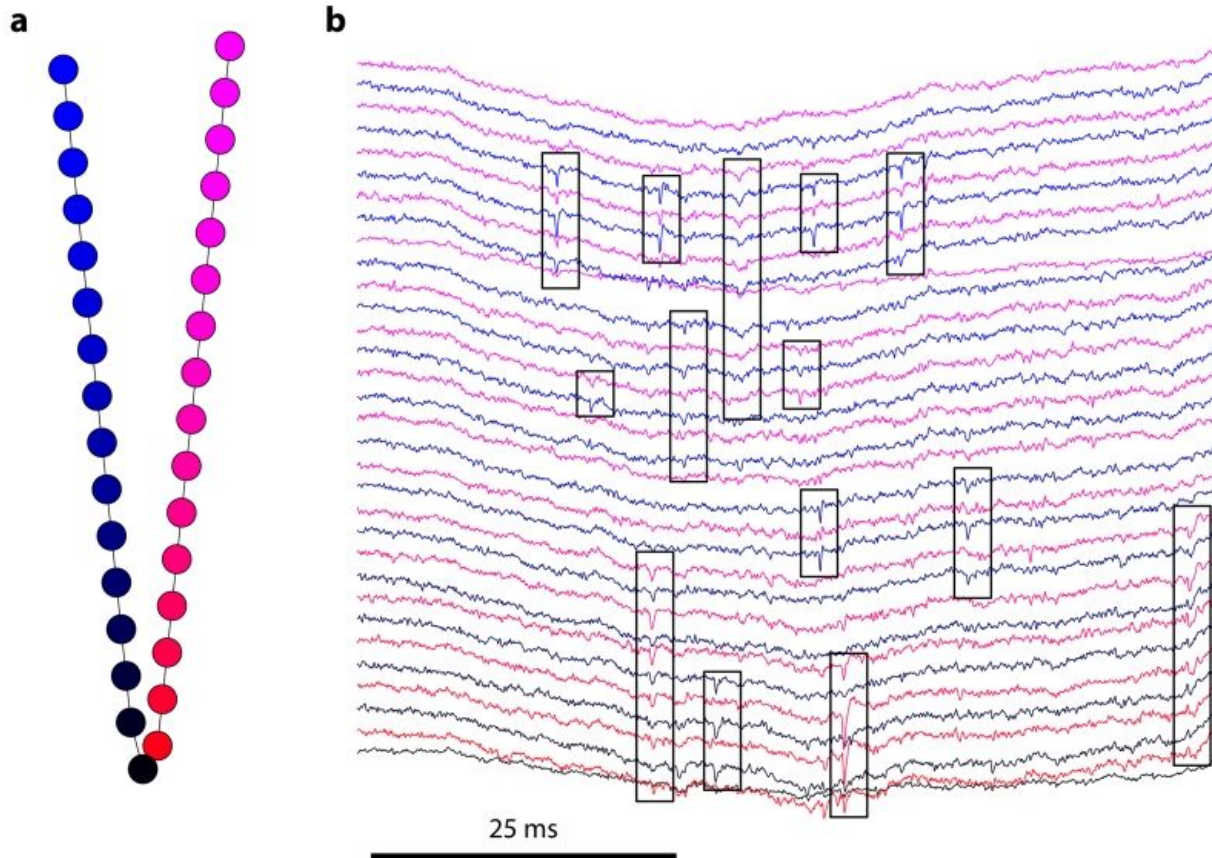


# Preprocessing: to get spikes, we must spike sort



*The spike sorting problem: 1) identify the spikes in the multi-channel voltage recording and 2) assign those spikes to individual neurons*

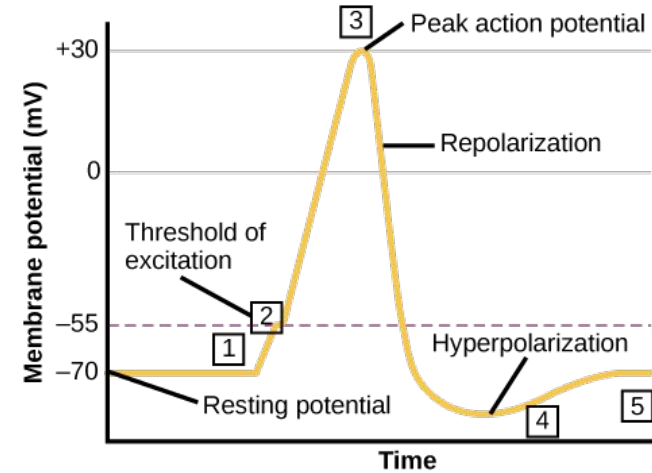
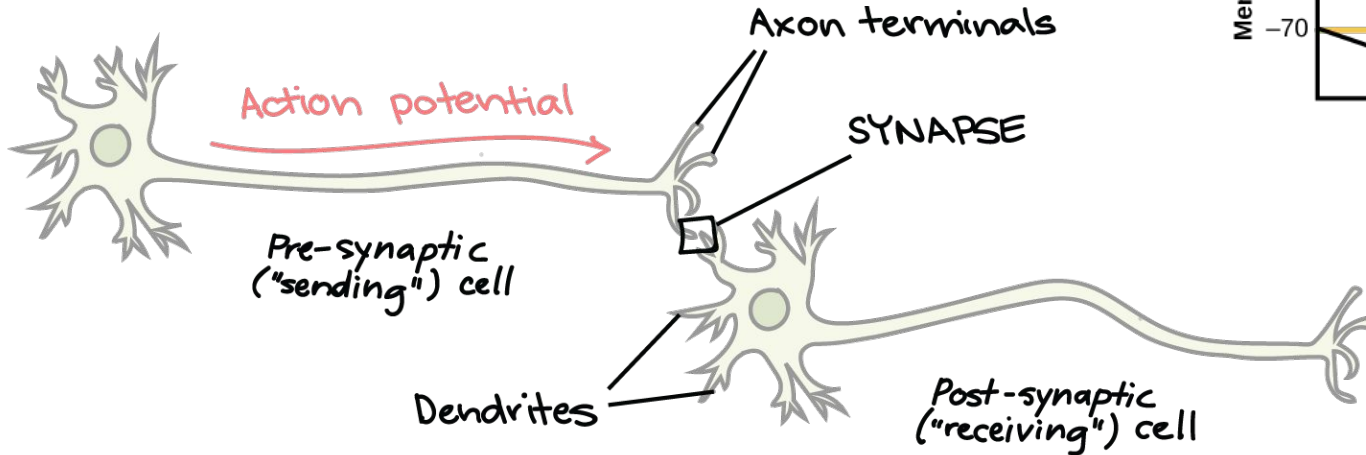
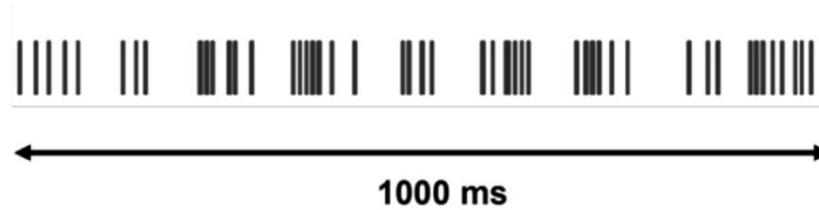
# Spike sorting is non-trivial



- Temporally overlapping spikes on contacts
- Simultaneous spiking from multiple neurons
- Accounting for spatio-temporal correlation given probe geometry.
- Autoregressive process (bursts)
- Electrode drift

# What does a spike represent?

Spikes have a shape, we are ignoring that and only looking at the spike time



# Summary: a spike is not a spike is not a spike

Spike times and neuron identities derived from extracellular data are created through **non-trivial preprocessing**.

We make necessary and useful simplifications but with that comes uncertainty.

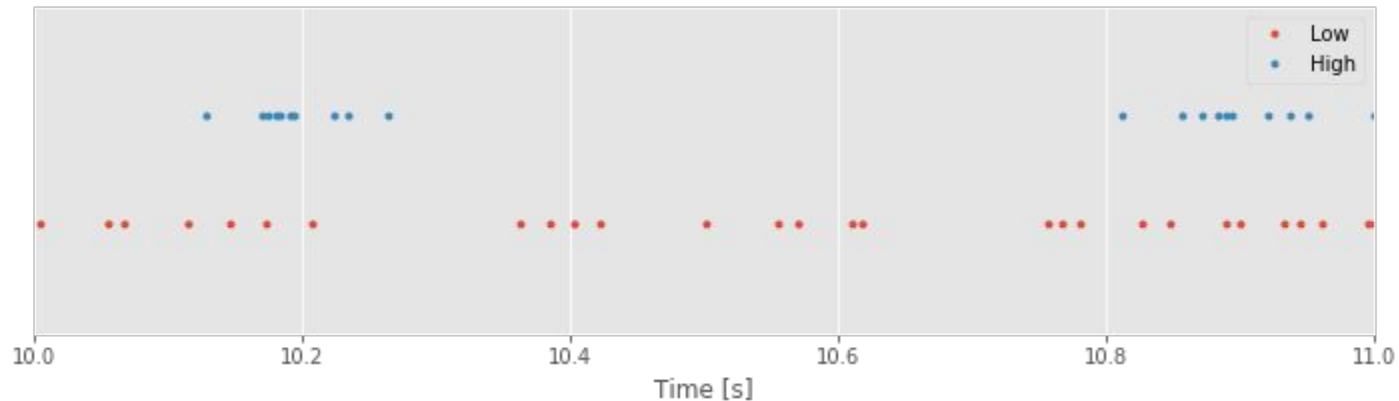
This should not stop us from analyzing spikes, but **it is important to understand that there is uncertainty and be careful with what we are analyzing.**

# What makes a neuron(s) spike?

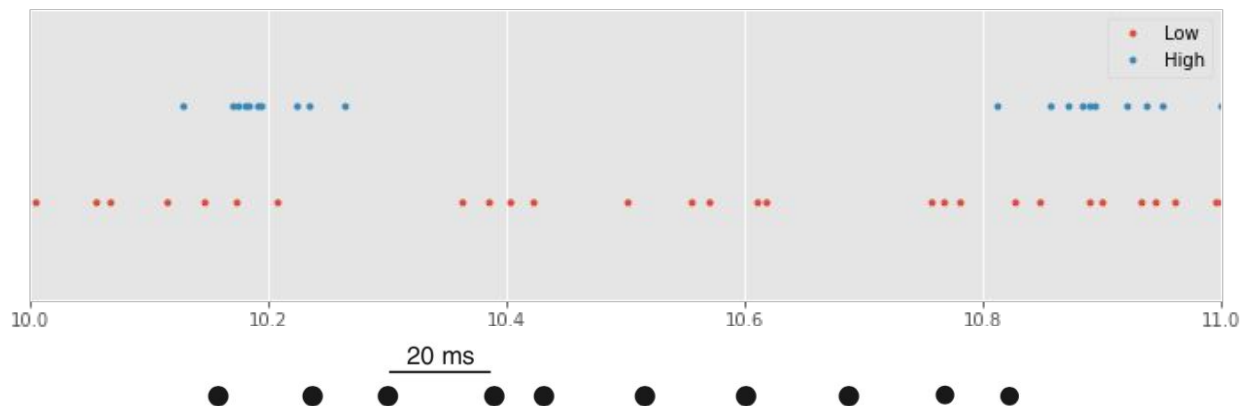
List of spike times: [0.1, 0.8, 0.9]

First step: Basic descriptive statistics and visualization

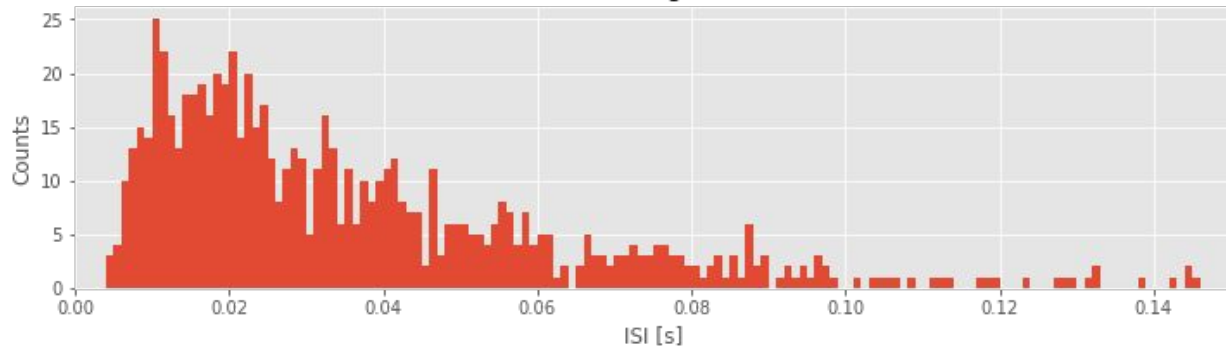
Raster plot



# Interspike intervals (ISIs)

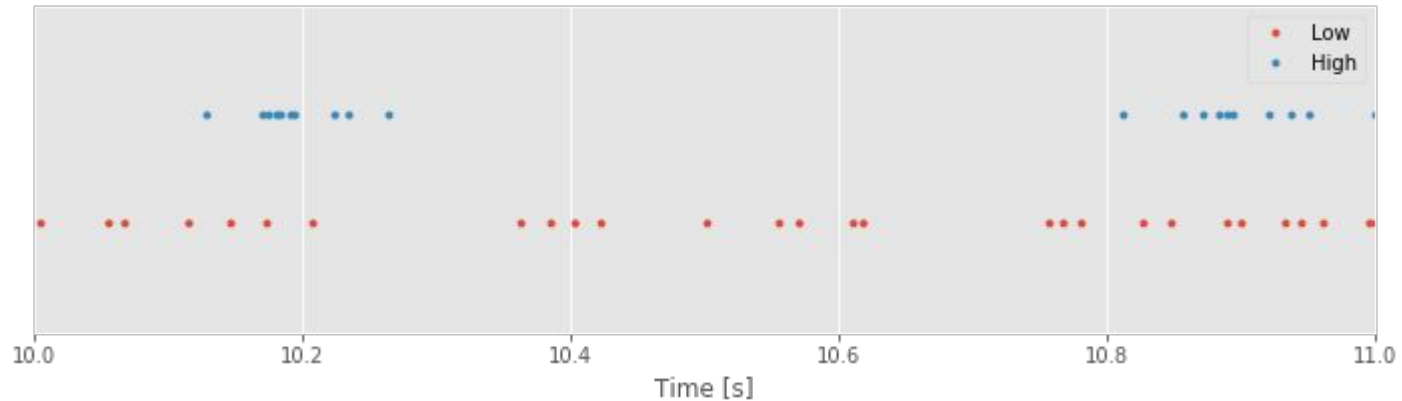


Low-light



$$X_i = S_i - S_{i-1}$$

# Counting spikes



Why is this a good summary of what is happening with a neuron under different conditions?



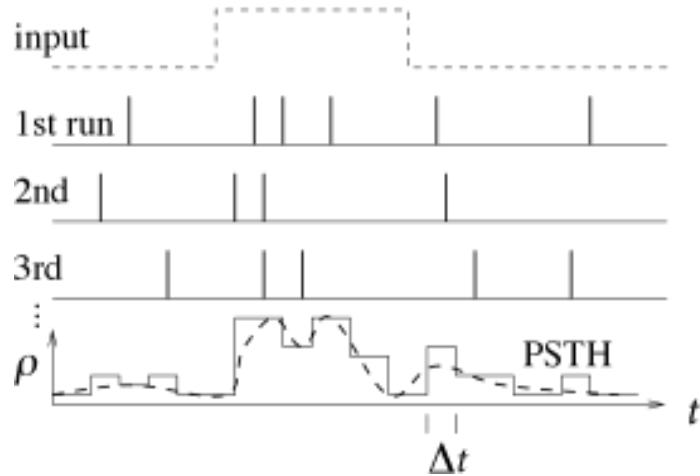
# Spike Rate

Provides a way to normalize different conditions (different length trials, different bin sizes)

$$FR = \frac{\text{number of spikes}}{\Delta t}.$$

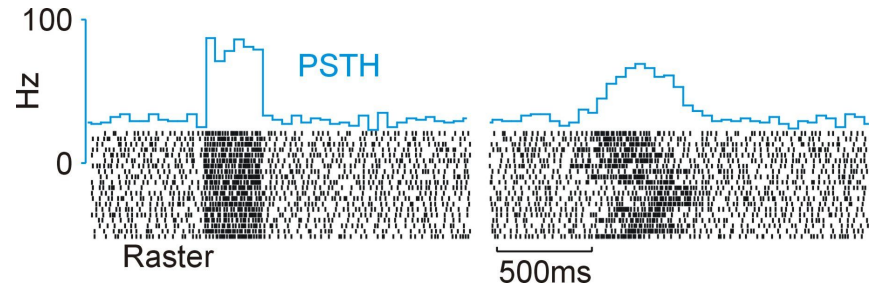
# Peri-event time histograms

rate = average over several runs  
(single neuron, repeated runs)



spike density  
in PSTH

$$\rho = \frac{1}{\Delta t} \frac{1}{K} n_K(t; t + \Delta t)$$

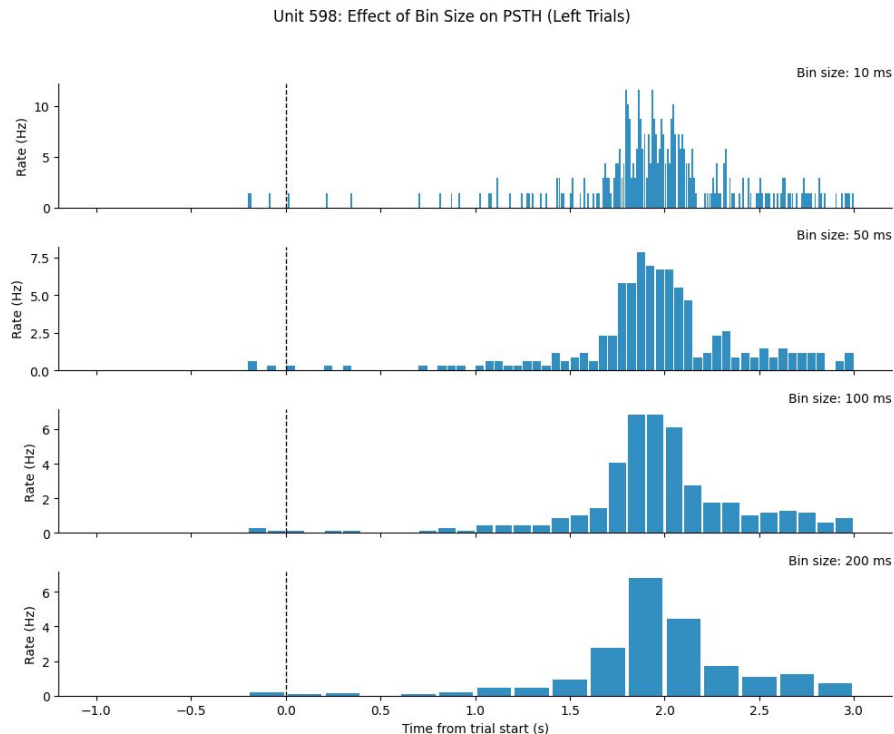


# Bin size matters

It's a choice about how fast you believe the rate is evolving (smaller bin, evolving faster, larger bin, evolving slower).

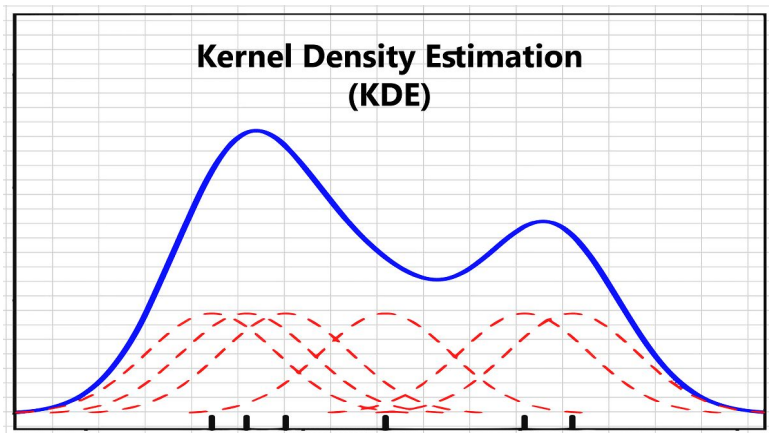
In a histogram, the prediction is a constant rate within the bin.

Bias-variance tradeoff

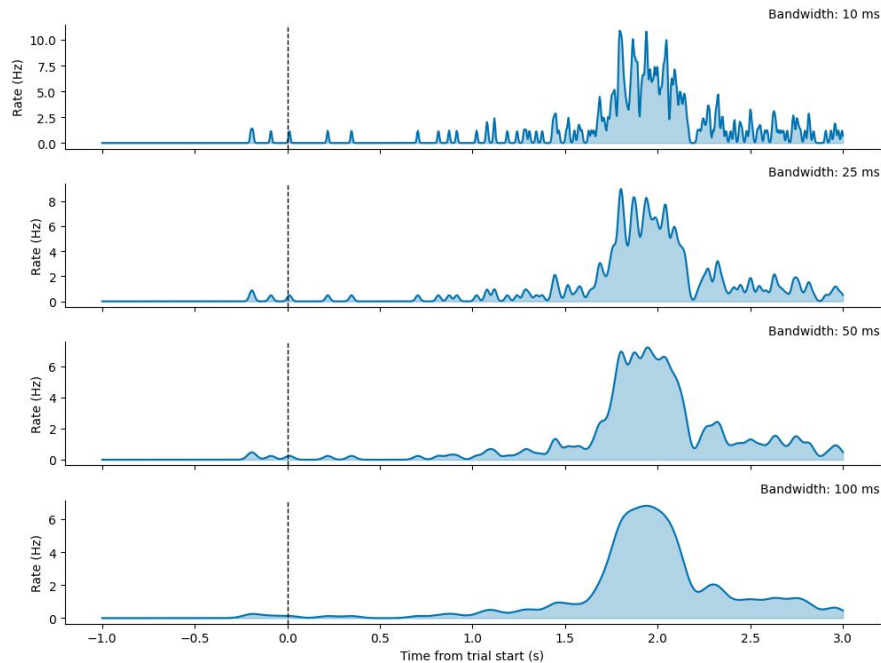


# Kernel Density Estimation

There are other “binless” methods such as kernel density estimation which give you smoother rates



Unit 598: Kernel Density Estimation (Left Trials)

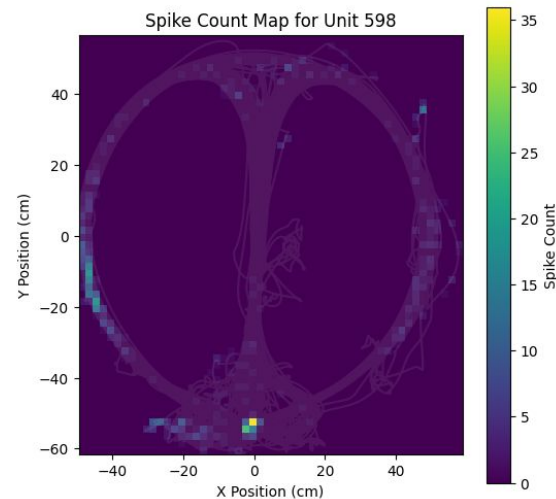
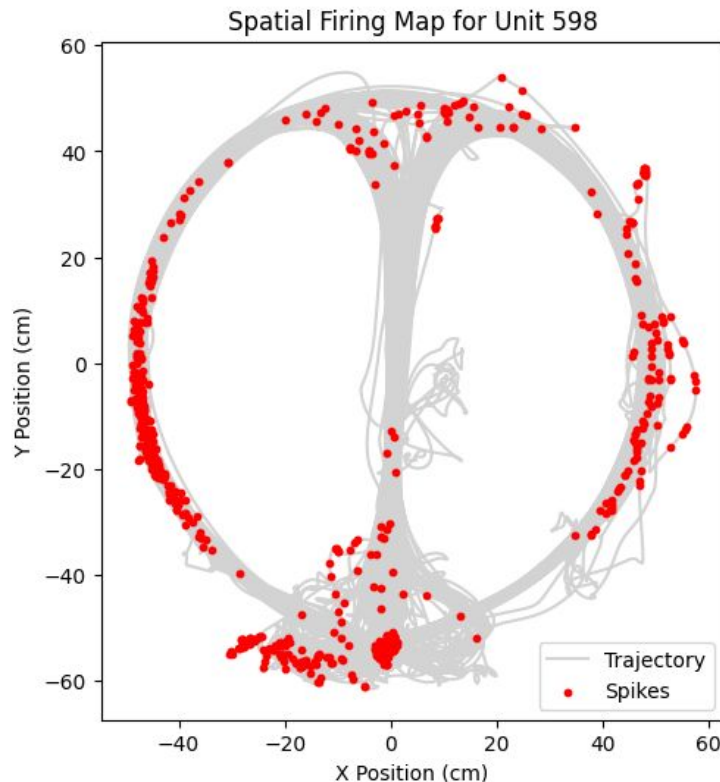


# Rate maps (e.g. place fields)

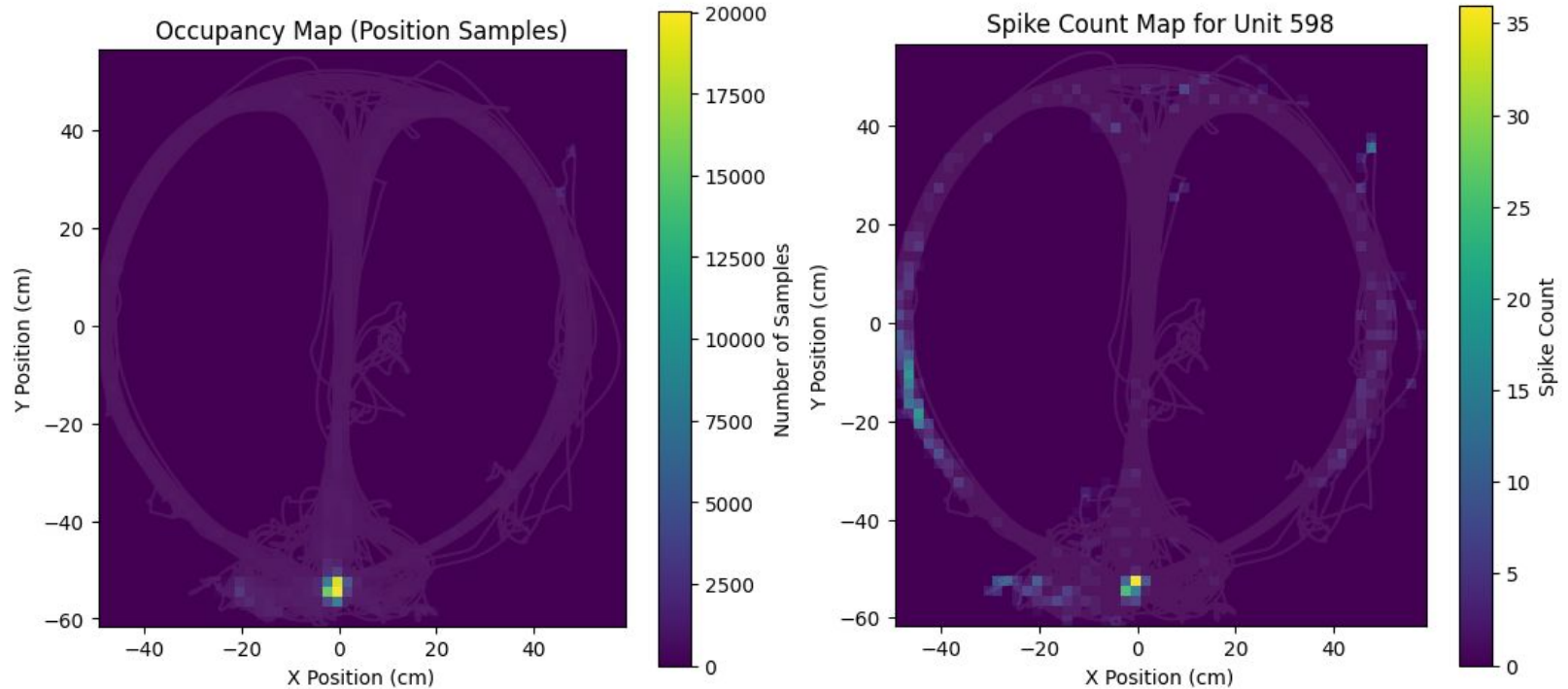
We could take a histogram over position, but this would be just counts of spikes per position. The animal could be in one position longer.

We want spike rate (spikes per second for each position bin)

How do we do this?

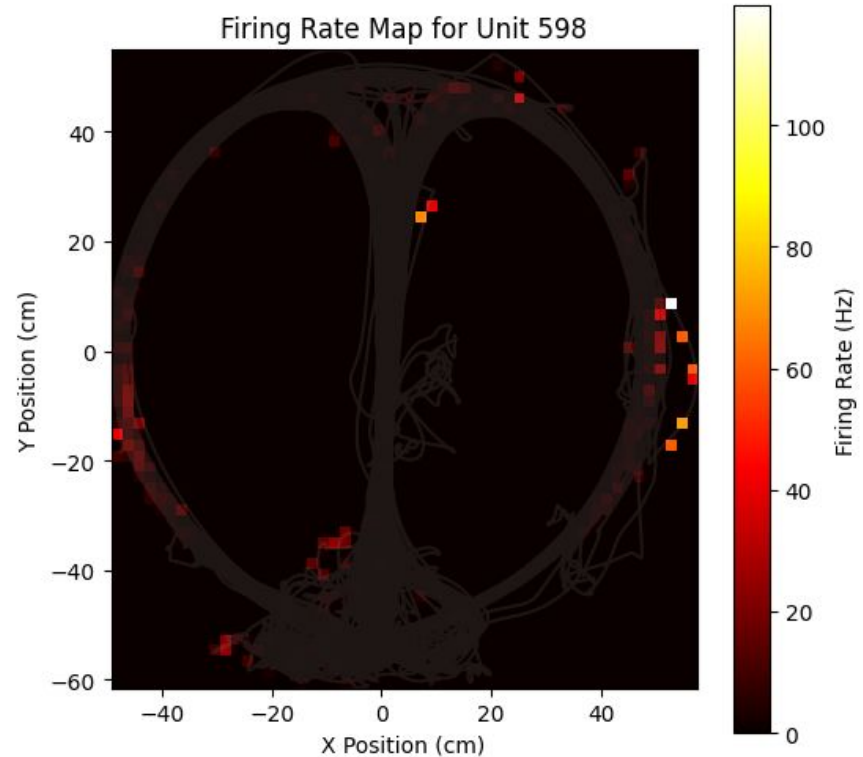


# Occupancy - How often is the animal at a given position



# Rate map - occupancy normalized histogram

Normalize counts per position bin by how often the animal is there.

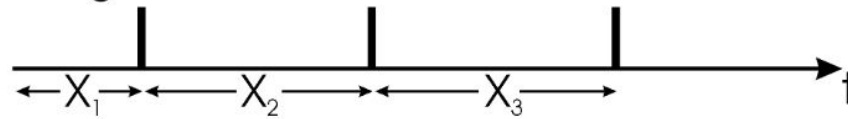


# Summary: four ways to describe a point process

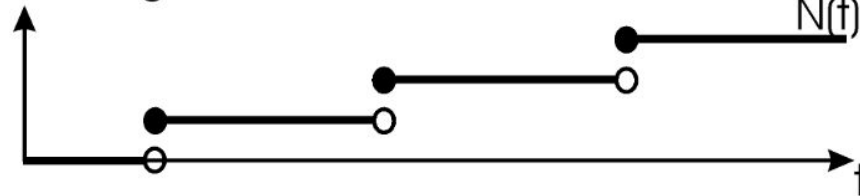
Spike Times:



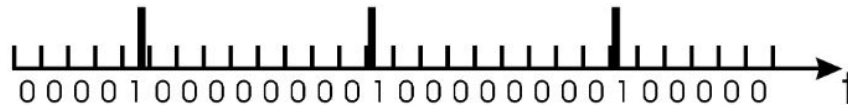
Waiting Times:



Counting Process:

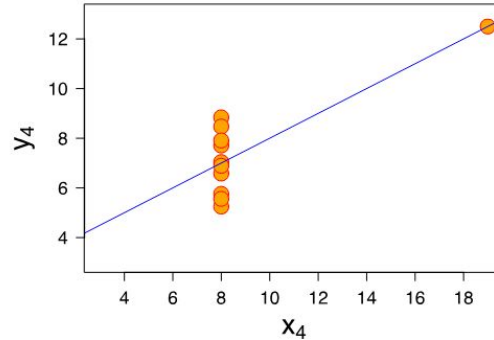
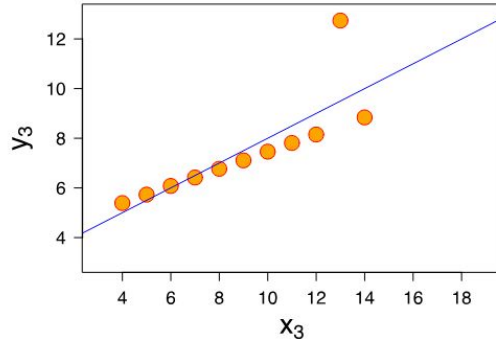
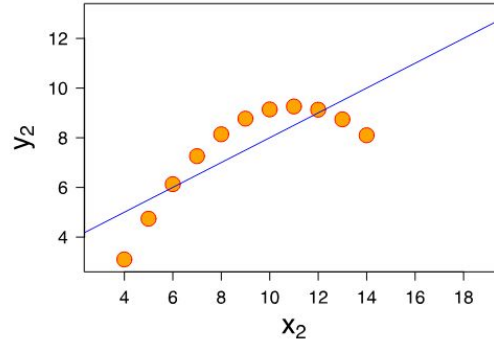
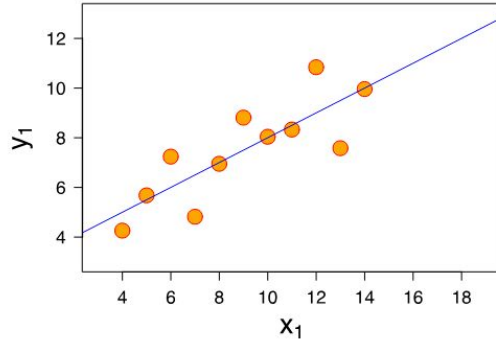


Discrete Increments:





# Importance of basic visualizations - Anscombe's quartet



all four of these have the same mean, variance, correlation, and regression line

So you need to plot and understand your model

# Characterizing variability spike rate

The number of spikes over a period of time (spike rate) exhibits high variability

An individual neuron presented multiple times with the same stimulus will typically produce a different number of spikes following each presentation.

But over many stimulus presentations, the ratio between the sample mean and sample variance of the spike counts over fixed intervals remains constant.

# Fano Factor

The ratio between the sample mean and sample variance of the spike counts

Dimensionless - easy to compare between in different conditions (e.g. different time windows, different stimulus intensities) and different rates (variability can be higher at higher rates)

Overdispersion: Fano Factor  $> 1$

Underdispersion: Fano Factor  $< 1$

Poisson: Fano Factor  $= 1$

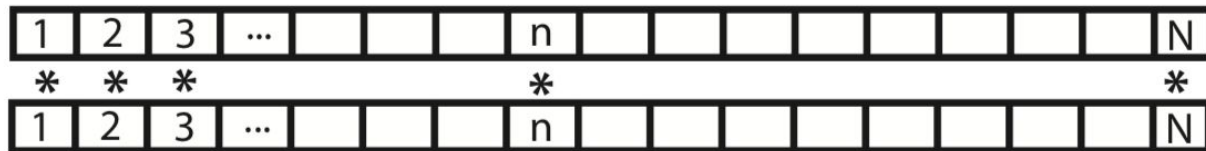
$$F_n = \frac{\mathbb{V}[y_{t,n}]}{\mathbb{E}[y_{t,n}]}$$

Autocorrelation: how is the current spiking related to past spiking

$$\rho_{xx}[L] = \frac{\sum_{i=1}^{N-L} (x_i - \bar{x})(x_{i+L} - \bar{x})}{\sum_{i=1}^N (x_i - \bar{x})^2}$$

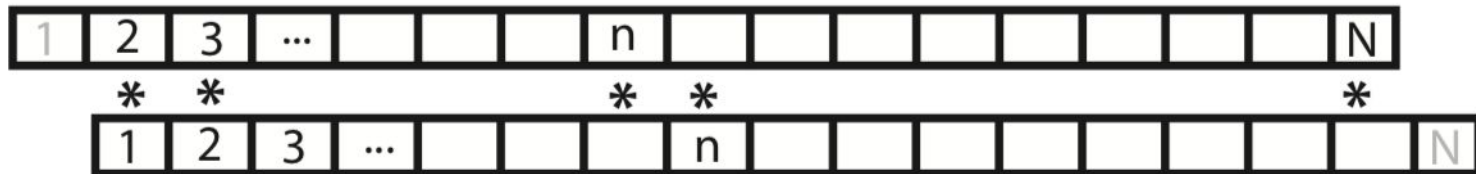
$$\rho_{xx}[L] = \frac{\sum_{i=1}^{N-L} (x_i - \bar{x})(x_{i+L} - \bar{x})}{\sum_{i=1}^N (x_i - \bar{x})^2}$$

L=0



sum  $r_{xx}[0]$

L=1



sum  $r_{xx}[1]$

# slide and inner product

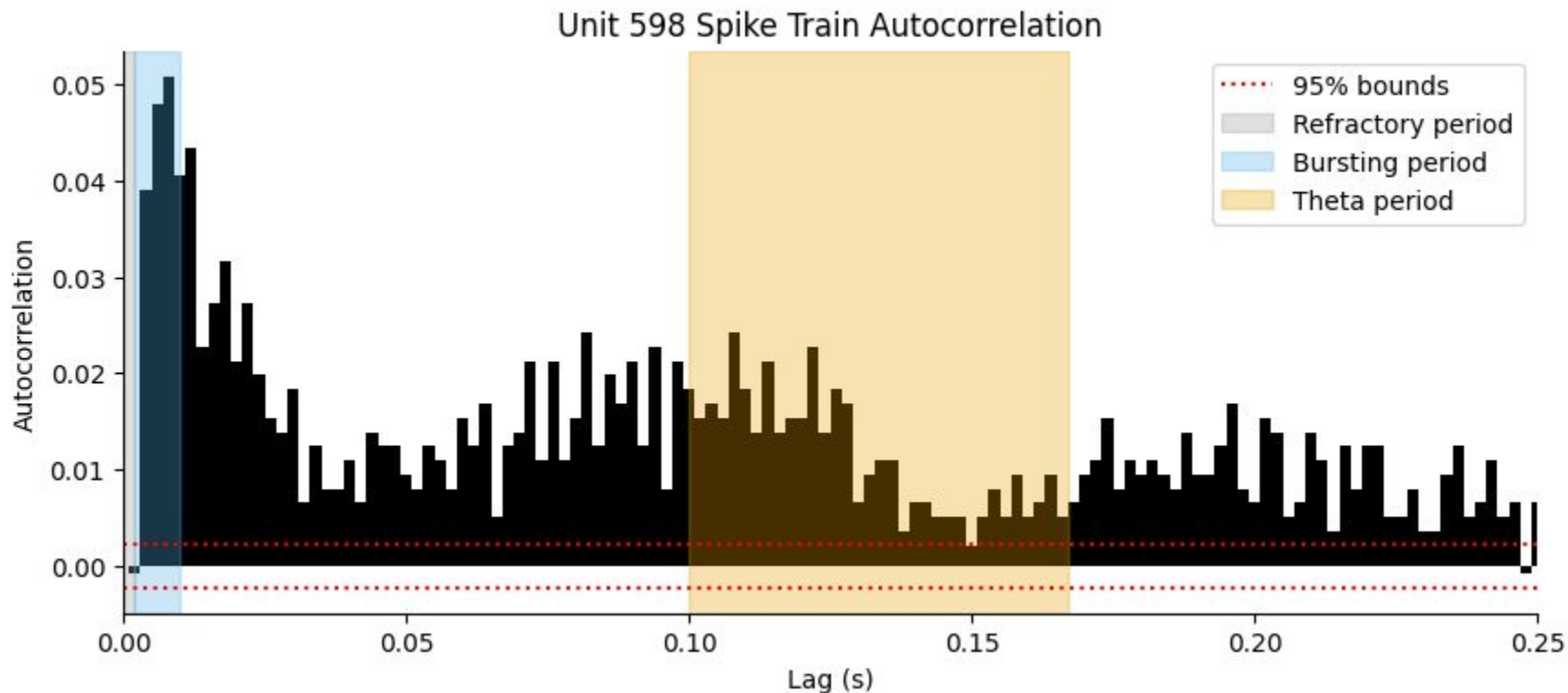
inner product (multiply & sum) measures similarity (how orthogonal vectors are, inner product =0 -> vectors orthogonal, inner product = 1 -> vectors parallel)

We measure similarity at each lag. We look at the original time series versus a lagged copy (the same time timeseries shifted backwards L time steps)

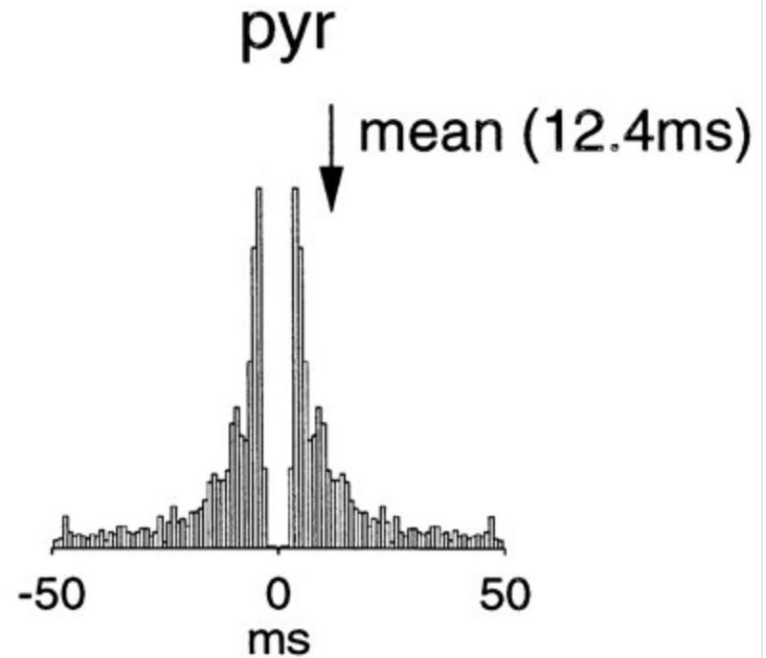
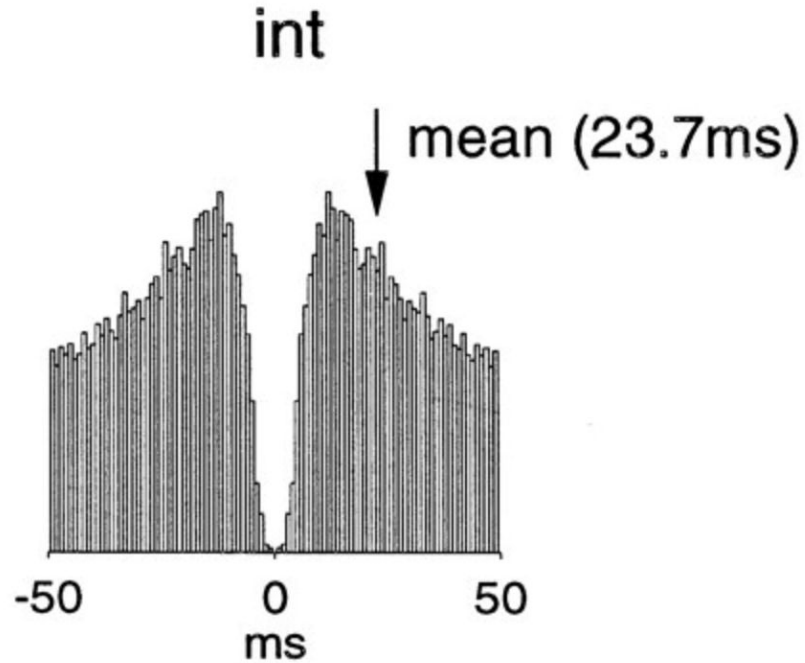
Remember that we have to subtract the mean (we are looking at residuals, similarity of fluctuations)

$$\rho_{xx}[L] = \frac{\sum_{i=1}^{N-L} (x_i - \bar{x})(x_{i+L} - \bar{x})}{\sum_{i=1}^N (x_i - \bar{x})^2}$$

# Hippocampal example



# Hippocampal example





# Poisson model: Statistical Model of a Neuron

Simple model (not modeling complex biophysics like Hodgkin-Huxley)

Well suited for modeling expected number of counts (number of spikes is discrete non-negative integers  $0, 1, 2, \dots$ )

Neuron firing is stochastic (spike times are variable across conditions)

**The goal is to find a rate parameter that explains the expected number of spikes.**

# Poisson distribution

Probability of k spikes for a given time bin:

$$P(k) = \frac{\lambda^k e^{-\lambda}}{k!}$$

lambda = rate (expected number of events)

P(k) = probability of k spikes

**The goal is to find a rate parameter that explains the expected number of spikes.**

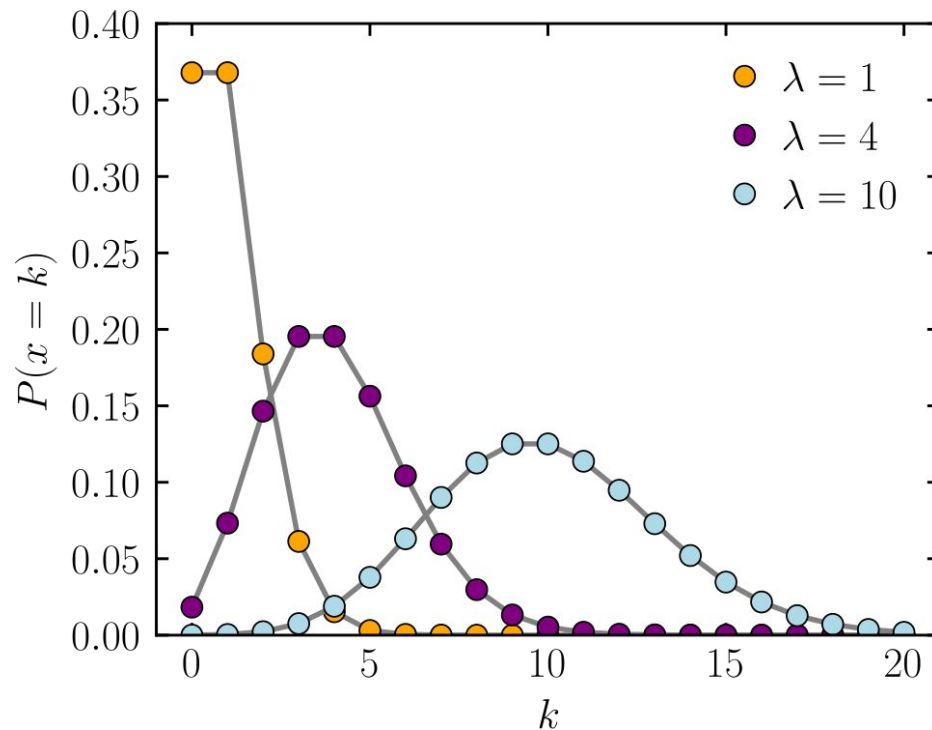
**What is the probability of 0 spikes?**

# Poisson

Low rate ( $\lambda = 1$ ): most probable one spike

Medium rate ( $\lambda = 4$ ):  
most probable 4 spikes but  
long tail

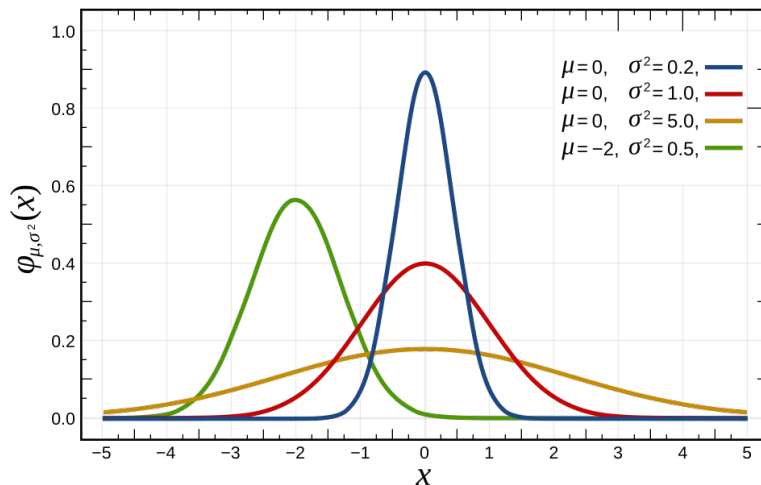
High rate ( $\lambda = 10$ ):  
Gaussian-like



# Why this is better than using a Gaussian

With a Gaussian, you can go below zero but spike rate is non-negative

With a Gaussian, mean and variance are linked: higher rate  $\rightarrow$  higher variability.  
Poisson captures this with one parameter; Gaussian can't.



# Poisson

Number of spikes in any time bin is independent of all previous (and future) spiking (memoryless).

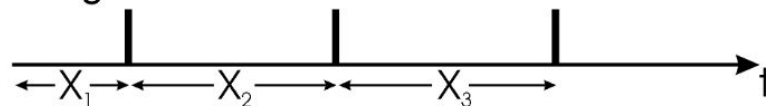
Variance scales with the mean

$$P(k) = \frac{\lambda^k e^{-\lambda}}{k!}$$

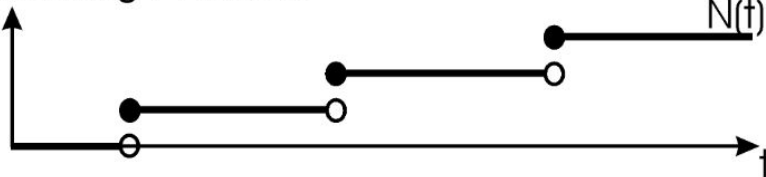
Spike Times:



Waiting Times:



Counting Process:



Discrete Increments:



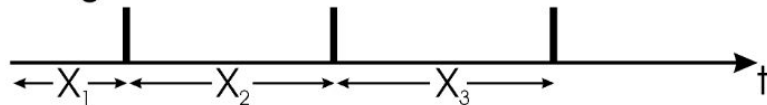
## Exponential waiting times (ISIs)

$$f(x) = \lambda \exp(-\lambda x)$$

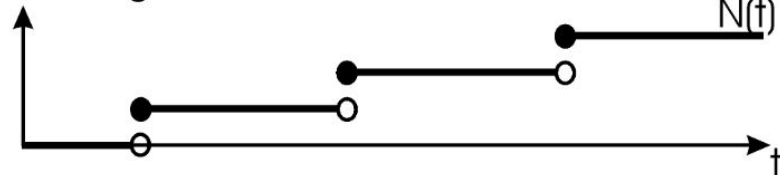
Spike Times:



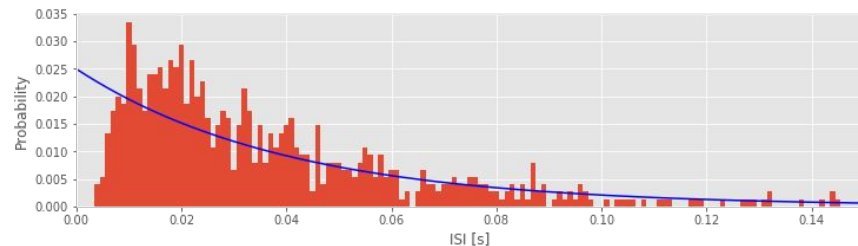
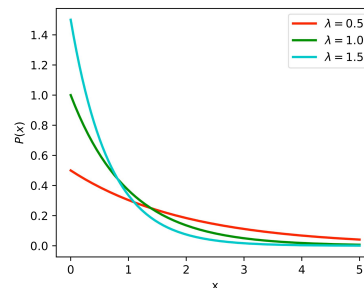
## Waiting Times:



### Counting Process:



### Discrete Increments:



Does our model fit well?

