

# Data Structures & Signal Processing

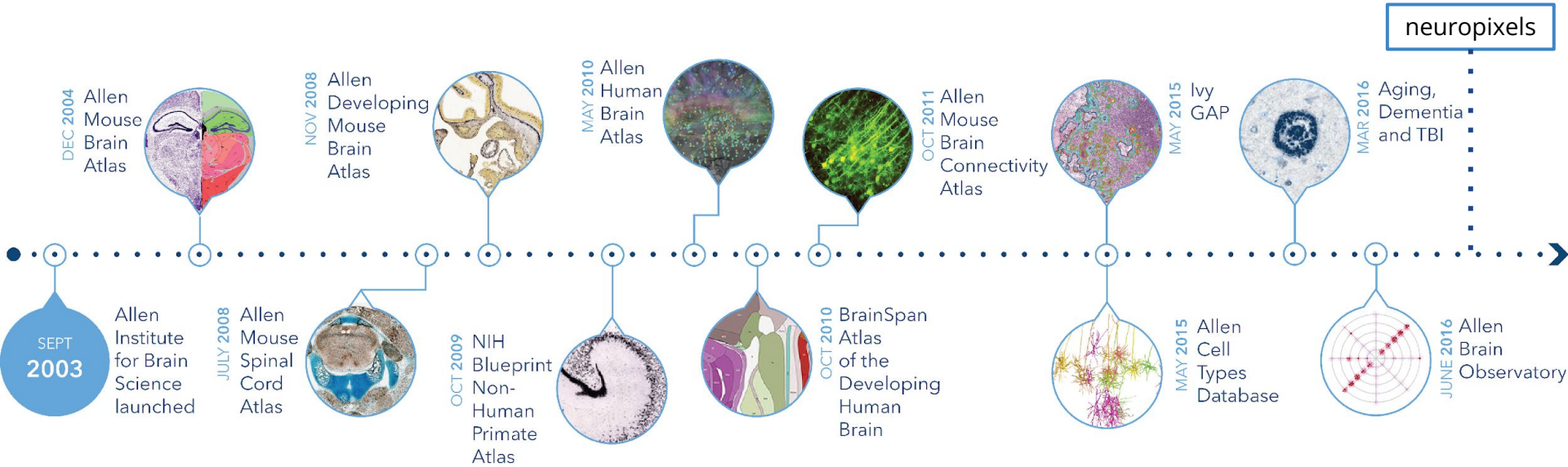
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BIPN 162

# Objectives for today

- Identify the three common features of datasets: **metadata**, **raw data**, and **processed** data
  - Define **metadata** and explore the metadata associated with an NWB file
  - Describe features of **time series** and common ways to analyze them
  - Load & explore an NWB time series
-

# Neuroscience datasets are increasingly being shared online



**NEURODATA**  
WITHOUT BORDERS



**DANDI**  
dandiarchive.org

Neuroscience Datasets ☆ 📁 ☁

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|    | A                         | B   | C  | D   | E                 | F            | G           |
|----|---------------------------|---|--|---|-------------------|--------------|-------------|
|    | Database Name             | URL   | Data Summary   | Student Objective   | Accessing API/SDK | Using Pandas | Text Mining |
| 2  | NeuroSynth                | <a href="https://neurosynth.org/">https://neurosynth.org/</a>   | Summarizes fMRI data from many studies   | Perform meta analyses of fMRI data  |                   |              | X           |
| 3  | LISC                      | <a href="https://lisc-tools.github.io/">https://lisc-tools.github.io/</a>   | Text analysis of papers  | Search publications for terms to identify interesting intersections.  |                   |              | X           |
| 4  | Allen Cell Types Atlas    | <a href="http://celltypes.brain-map.org/">http://celltypes.brain-map.org/</a>   | Whole-cell electrophysiology in genetically-identified cell types in mice and humans | Compare electrophysiology metrics for different cell types in mice and humans.  | X                 | X            |             |
| 5  | Allen Brain Observatory   | <a href="http://observatory.brain-map.org/">http://observatory.brain-map.org/</a>   | In vivo 2p imaging in genetically-identified cell types in mice                      | Compare visual responses of cells recorded via two-photon imaging and analyze correlations.                                       | X                 | X            |             |
| 6  | Allen Neuropixels         | <a href="https://allensdk.readthedocs.io/en/latest/">https://allensdk.readthedocs.io/en/latest/</a>                                       | Extracellular recording & behavior in mice performing a task                         | Compare visual & behavioral responses of cells recorded with high density extracellular recording arrays. Also includes LFP data. | X                 | X            |             |
| 7  | Allen RNAseq              | <a href="https://portal.brain-map.org/datasets/AllenRNAseq">https://portal.brain-map.org/datasets/AllenRNAseq</a>                         | RNAseq in mice & humans  | Compare gene expression in mice and humans in different cells & brain regions.  |                   | X            |             |
| 8  | Allen Connectivity        | <a href="http://connectivity.brain-map.org/">http://connectivity.brain-map.org/</a>   |  |   |                   |              |             |
| 9  | Allen Synaptic Physiology | <a href="https://portal.brain-map.org/datasets/AllenSynapticPhysiology">https://portal.brain-map.org/datasets/AllenSynapticPhysiology</a> |  |   |                   |              |             |
| 10 | Neuropixels Spike Sorting | <a href="http://repository.cshl.edu/handle/123456789/123456789">http://repository.cshl.edu/handle/123456789/123456789</a>                 | Lots of spikes   | PCA analysis to isolate units in extracellular recording  |                   |              |             |

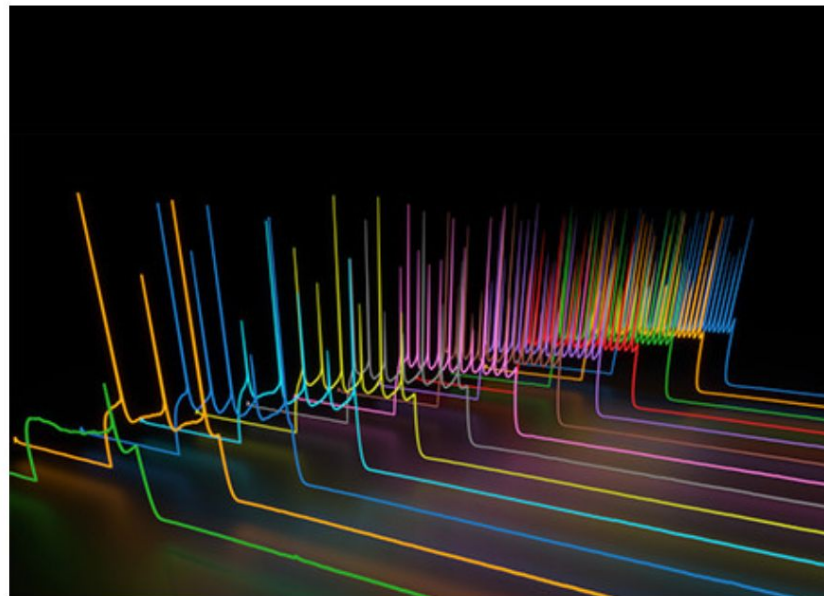
Open Neuroscience Datasets <https://bit.ly/openneurodatasets>



## Neurodata Without Borders

**Neurodata Without Borders** (NWB) is a data standard for neurophysiology, providing neuroscientists with a common standard to share, archive, use, and build analysis tools for neurophysiology data. NWB is designed to store a variety of neurophysiology data, including data from intracellular and extracellular electrophysiology experiments, data from optical physiology experiments, and tracking and stimulus data.

The **NWB team** consists of neuroscientists and software developers who recognize that adoption of a unified data format is an important step toward breaking down the barriers to data sharing in neuroscience.



*(Courtesy of the Allen Institute for Brain Science)*

See also Rübel et al. (2022):  
<https://elifesciences.org/articles/78362>

d.

### Experimental Subject

- Species
- Genotype
- Age and weight
- Custom subject-specific fields



**NEURODATA**  
WITHOUT BORDERS

### Behavior

- Position and speed
- Choice and response time
- Video and audio
- Eye tracking
- Experiment-specific measures



### Experimental Design

- Stimuli
- Environment
- Trial structure
- Epochs
- Perturbations



### Data Acquisition

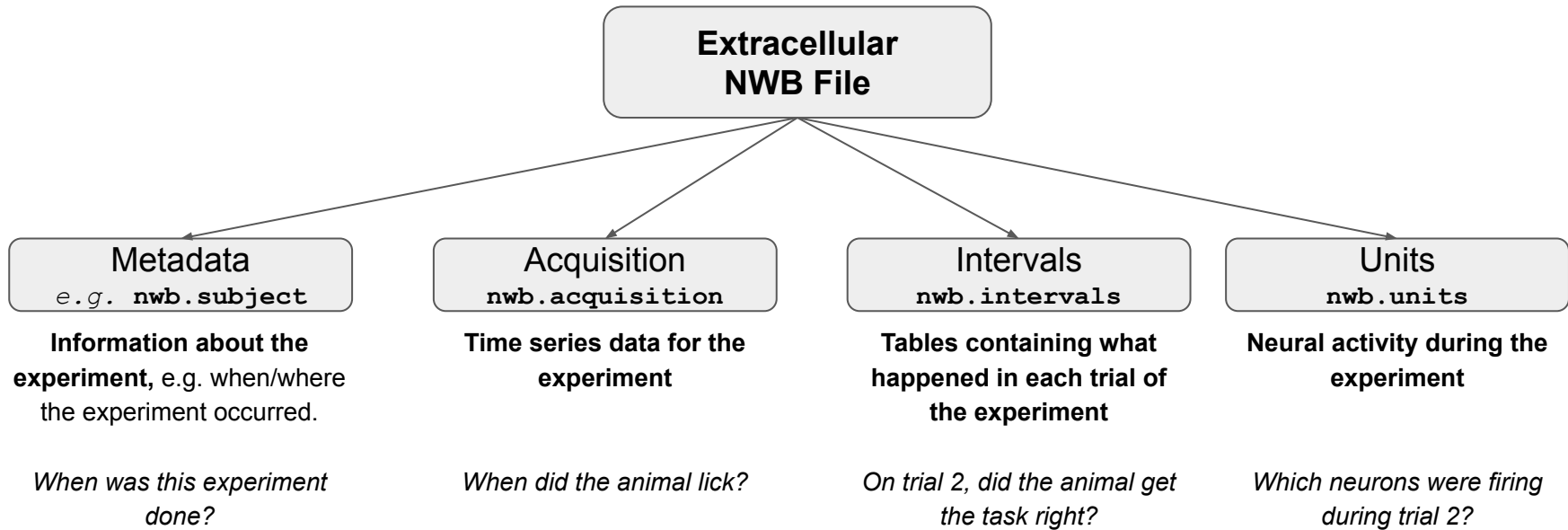
- Device settings
- Filtering parameters
- Sampling rate
- Recording/imaging area
- Emerging technologies



### Neuronal Activity

- Extracellular electrophysiology
- Intracellular electrophysiology
- Optophysiology
- Pre-processed data
- New data modalities







The BRAIN Initiative archive for publishing and sharing neurophysiology data including electrophysiology, optophysiology, and behavioral time-series, and images from immunostaining experiments.



# Signal processing

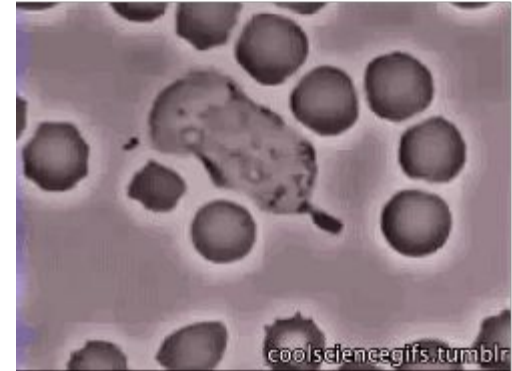
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Anything recorded  
continuously over  
time is a **time series**  
(a set of data points  
generated from successive  
measurements over time)



# Commonly encountered time series data in biology

- Gene expression data over time
- Neurophysiology recordings (e.g. electrophysiology, imaging)
- Circadian rhythm data
- Medical observations over time
- Animal movement
- Physiology data (e.g. heart rate/ECG, pulse rate, respiration, etc.)
- Molecules/proteins/cells moving



White blood cell tracking bacteria

[Image info](#)

# Common signal processing approaches

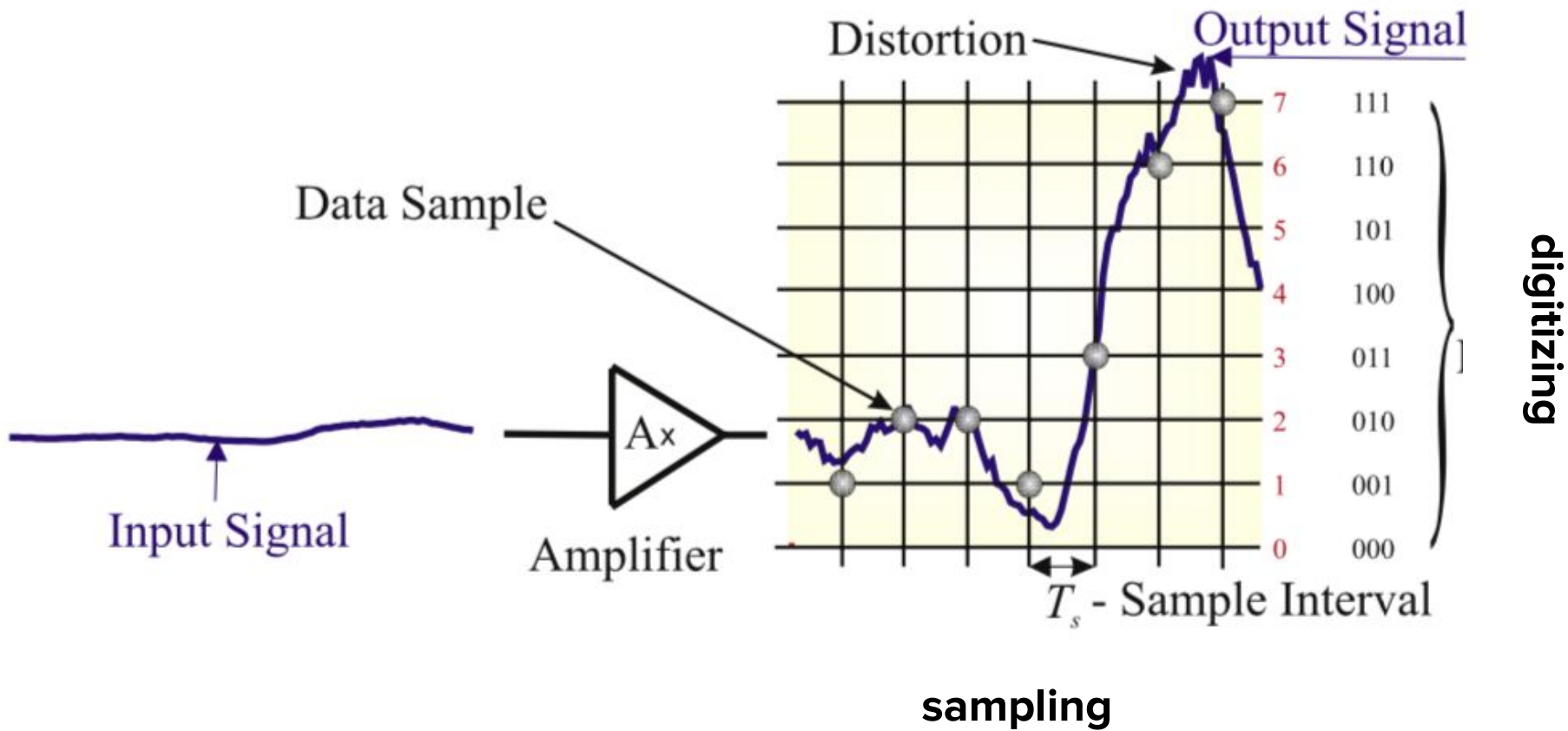
- Preprocessing & data cleaning
  - Removing outliers and/or noise
- Filtering
  - Using convolution
  - Using frequency
- **Looking for correlations in time**
- Clustering & classification
- Dimensionality reduction or segmentation
- Prediction
- Anomaly or peak detection — ***spike sorting***

Real world signals (e.g., brain and sound waves) are continuous in time. In other words, they're **analog**.

However, our computers are **digital** — based on **binary** representations of information.

When we work with continuous signals, then, we're always **sampling** and **digitizing** them.





# Important signal processing terms

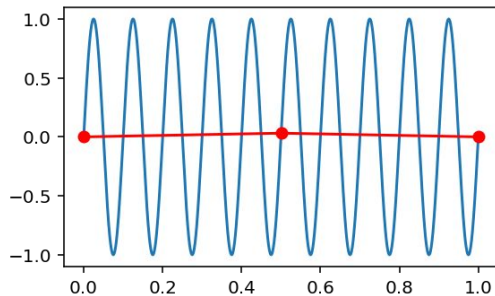
**Sampling frequency** (or rate): how often was a sample collected?

- Lower for EEG (typically around 100 Hz), very high for spiking data — why?
- Trade off: higher sampling rate means bigger data!

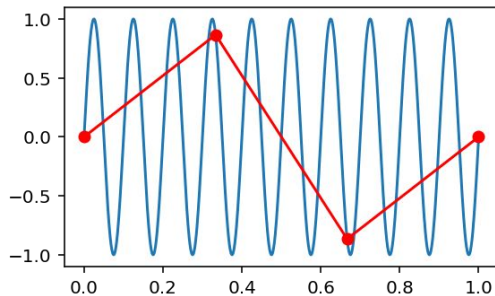
**Nyquist rate:** to accurately represent a periodic signal with a frequency, you have to sample ***at least*** twice as fast, otherwise you will get aliasing

- Relatedly, we can only detect frequencies at half the sampling frequency: the **Nyquist frequency**

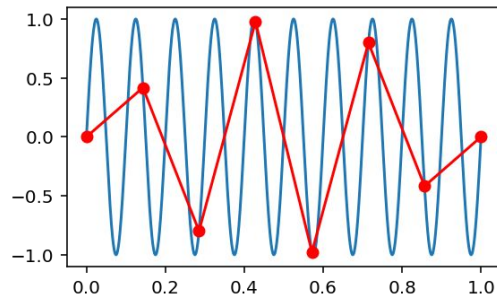
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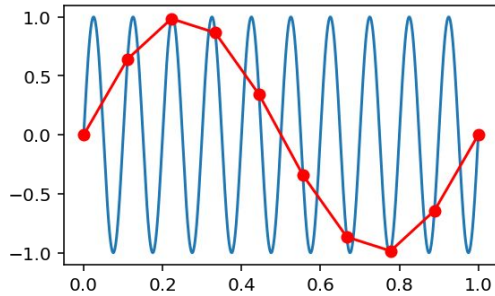
fs = 4



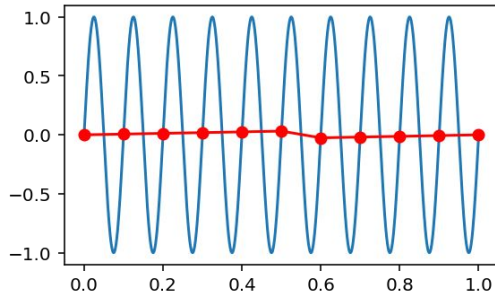
fs = 8



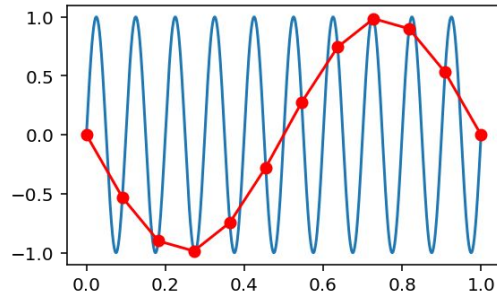
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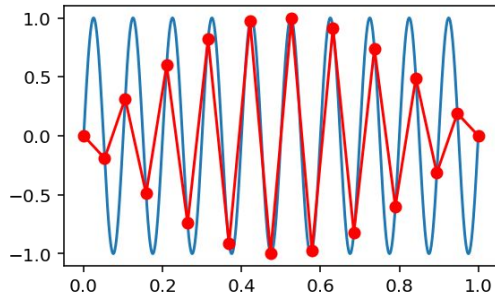
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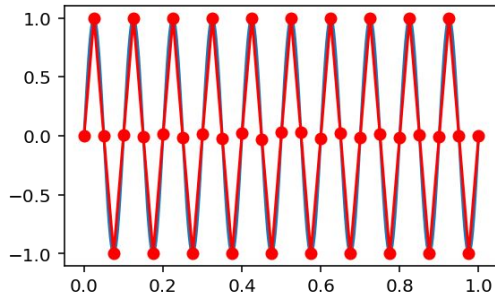
fs = 12



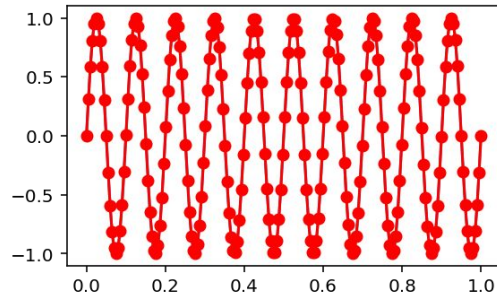
fs = 20



fs = 41



fs = 200



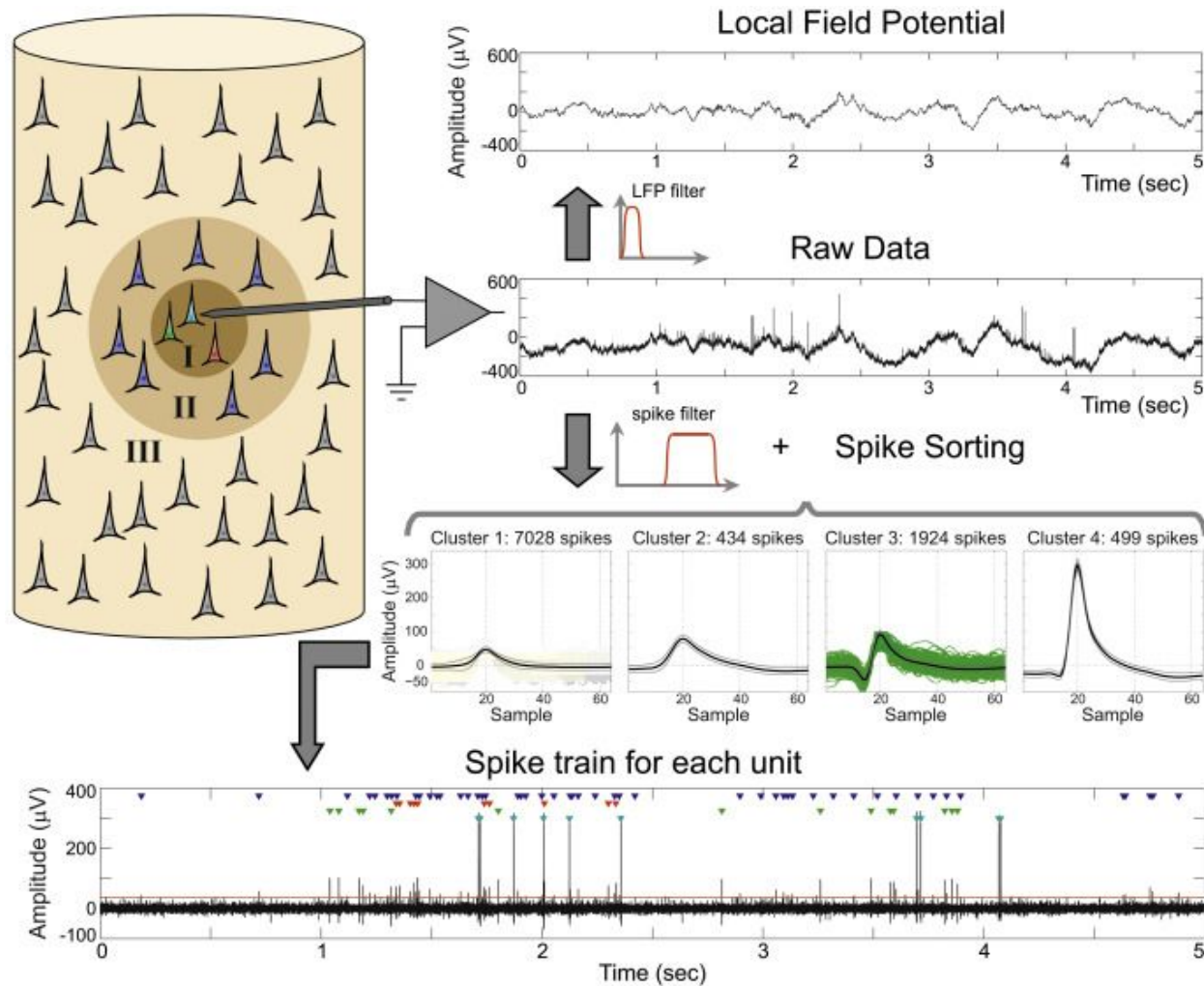
**Blue** = the  
signal we're  
trying to  
measure

**Red** = our  
sampling of  
that signal

**fs** = sampling  
frequency



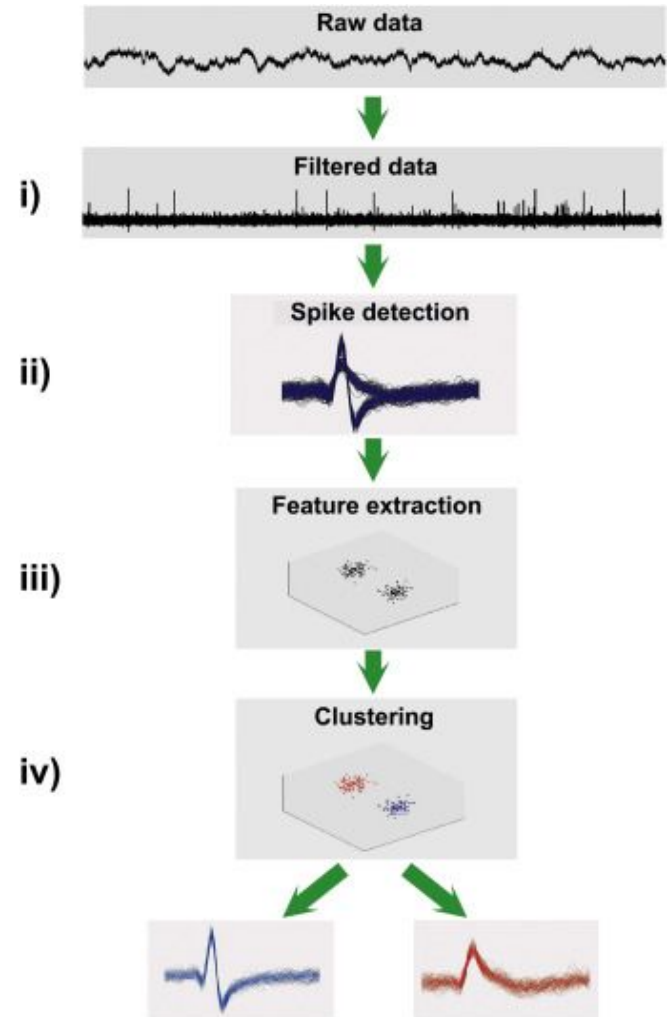
# Extracellular recordings require spike sorting

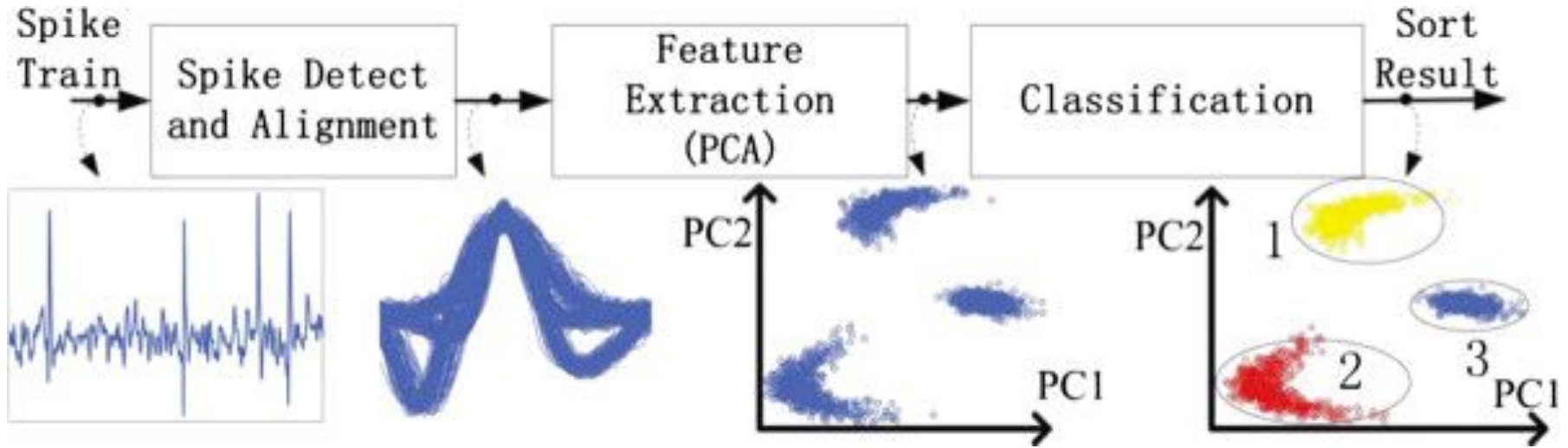


From Rey et al. (2015) "Past, present and future of spike sorting techniques"

# Basic steps for spike sorting

1. a bandpass **filter** is applied, e.g., between 300 Hz and 3000 Hz, to keep the most useful part of the spectrum for spike sorting
2. **spikes** are detected, usually with an amplitude threshold applied to the filtered data
3. relevant **features** of the spike shapes are extracted via a dimensionality reduction technique (PCA)
4. those features are the input to a **clustering algorithm** that performs the classification of the waveforms and associate each cluster to a unit.





PCA is also commonly (traditionally) used for extracellular spike sorting. After projecting onto these dimensions, you can then use a clustering algorithm (or detect them manually...)

# Spike Correlations

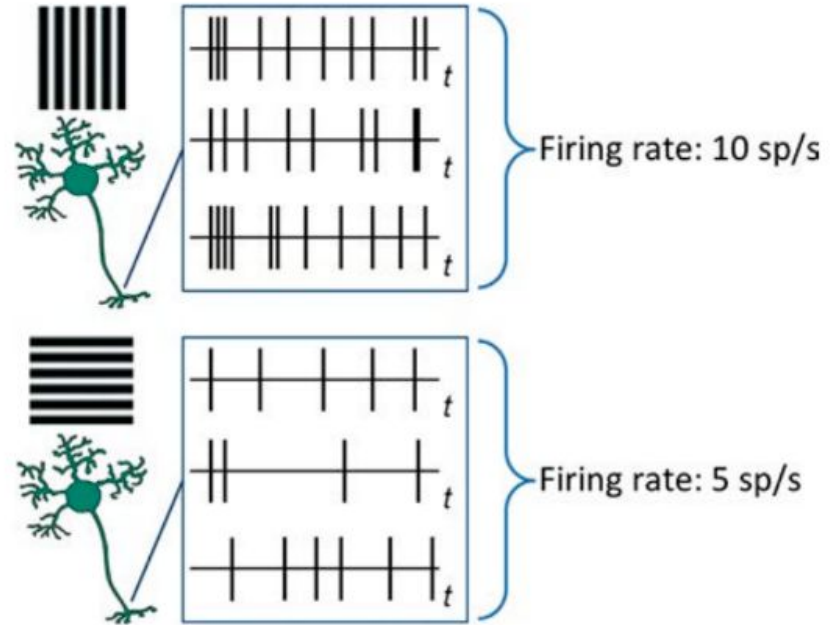
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# Correlations can provide information about the **functional architecture** of networks

For example, **connectivity in the retina**  
(Greschner et al. 2011),

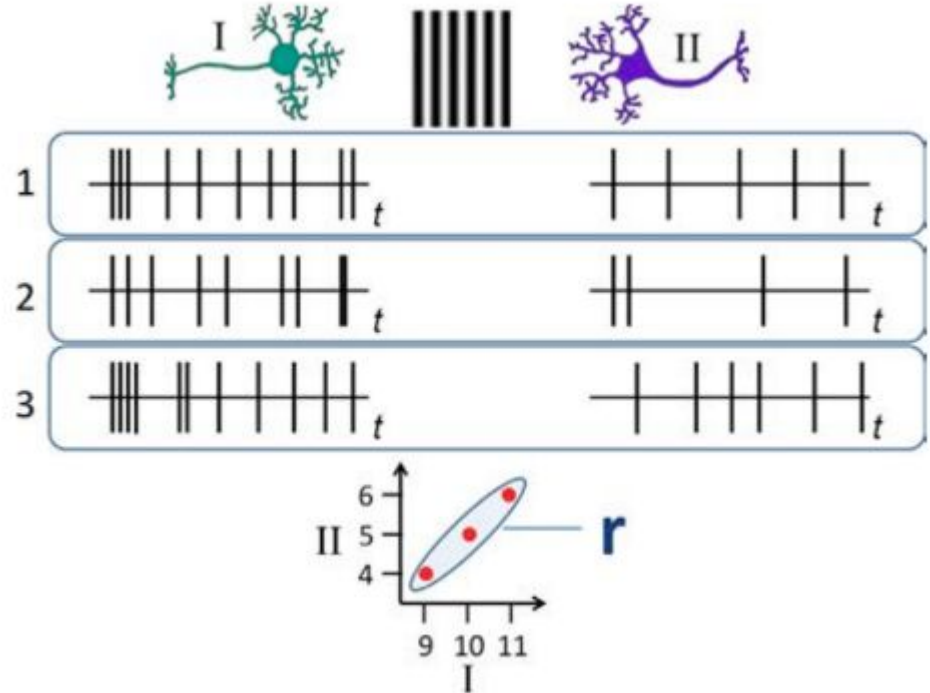
between the **visual thalamus and cortex**  
(Reid & Alonso, 1995),

and **between neurons in cortex** (Aertsen  
et al., 1989 & Alonso & Martinez, 1998)



## Correlations between pairs of neurons can arise over a range of timescales

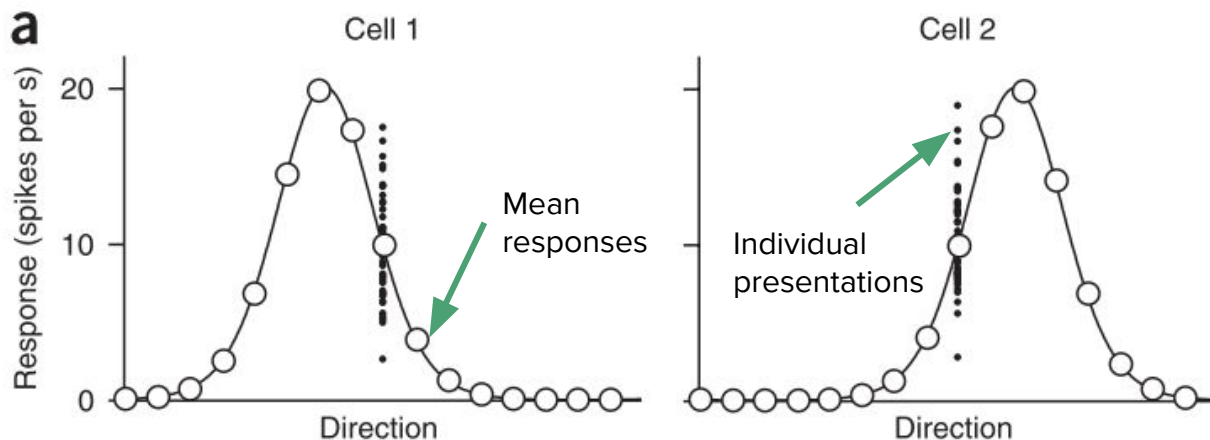
- Spikes may be precisely aligned in time, down to the millisecond
- Excitability may change on the order of seconds or longer



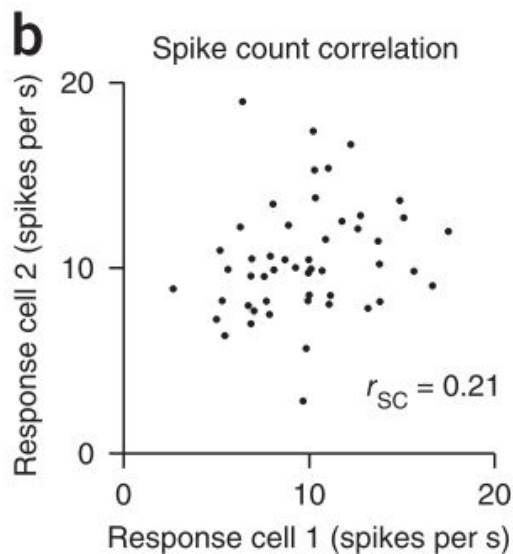
Neuroscientists often look for two different types of correlations: **signal** and **noise** correlations.

Small dots = individual presentations;  
Big circles = mean responses

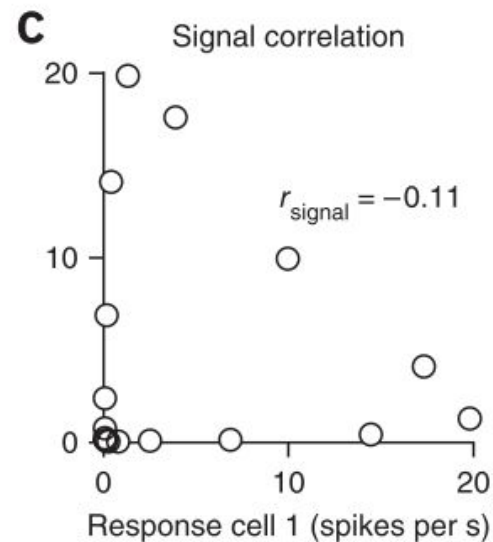
**a.** Tuning curves for two hypothetical cells



**b.** Spike count correlation (**noise** correlation) for those cells



**c.** Signal correlation



From Cohen & Kohn, 2011



## Signal correlations

measures the correlation coefficient between the cells' mean responses to different stimuli.

often used to quantify the extent to which a pair of neurons has similar tuning or other functional properties.

decreases in this type of correlation can lead to sparsening of population responses

## Noise correlations

measures the correlation between responses to repeated presentations of identical stimuli, under the same behavioral conditions

“spike count correlations”

Noise, or **spike count correlations** ( $r_{sc}$ ) tend to be small but positive

and tend to be higher for neurons that are near each other

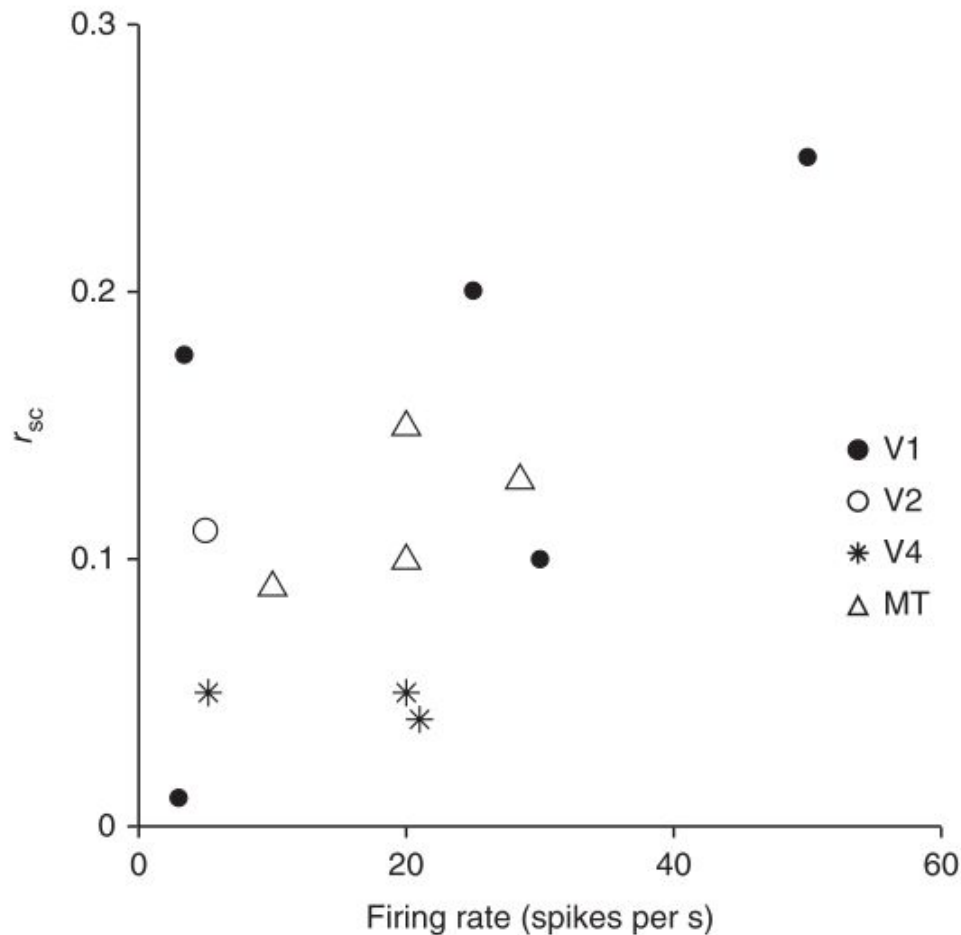
From Cohen & Kohn, 2011

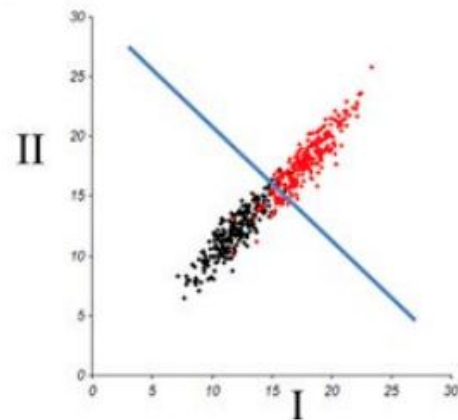
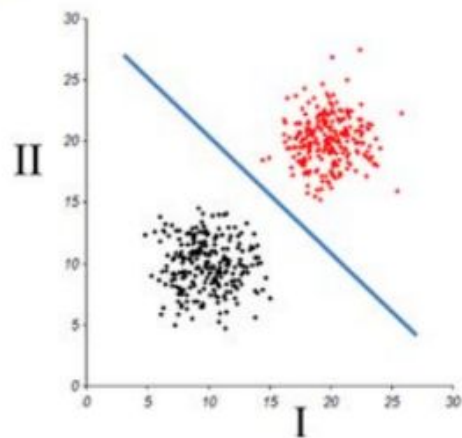
**Table 1 Summary of studies measuring spike count correlations in primates**

| Reference number   | Area                      | Firing rate (spikes per s) | Duration (ms) | State (task, anesthesia, etc.) | $r_{sc}$  |
|--|---------------------------|----------------------------|---------------|--------------------------------|-----------|
| 12*  | V1                        | ~25                        | 2,560         | Anesthetized                   | 0.2       |
| 26*  | V1                        | 3.4                        | 1,280         | Anesthetized                   | 0.18      |
| 23   | V1                        |                            | 1,894         | Anesthetized                   | 0.25      |
| 31   | V1                        |                            |               | Anesthetized                   | 0.26      |
| 13*  | V1                        | ~50                        | 1,860         | Fixation                       | 0.25      |
| 28*  | V1                        | ~3                         | 500           | Fixation                       | 0.01      |
| 82   | V1                        |                            | 400           | Tracing                        | 0.18      |
| 83*  | V1                        | 30                         | 1,000         | Discrimination                 | 0.1       |
| A. Zandvakili and A.K., unpublished data*                                  | V2                        | 5                          | 1,000         | Anesthetized                   | 0.11      |
| M. Smith and M. Sommer (University of Pittsburgh), personal communication* | V4                        | 5.2                        | 1,000         | Fixation                       | 0.05      |
| 7*   | V4                        | 21                         | 200           | Attention/detection task       | 0.04      |
| 8*   | V4                        | >5, ~20                    | 800           | Attention/tracking task        | 0.05      |
| A.B.G. Graf (New York University), personal communication*                 | MT                        | ~10                        | 300           | Anesthetized                   | 0.09      |
| 29*  | MT                        | ~20                        | 500           | Fixation                       | 0.1       |
| 15*  | MT                        | 28.5                       | 500           | Discrimination                 | 0.13      |
| 6/22*  | MT                        | ~20                        | 1,000         | Discrimination                 | 0.15      |
| 84   | Perirhinal                | ~12                        | 200–500       | Fixation/matching task         | 0.02      |
| 85   | Supp motor area           |                            | 66 or 200     | Serial reaching                | 0.013     |
| 27   | Supp motor area           | ~15                        | 200           | Reaching                       | 0.02      |
| 86   | Premotor areas            | ~5                         | 400           | Grasping/imagery task          | 0.02      |
| 87   | M1                        | ~20                        | 600           | Reaching                       | 0.1–0.2   |
| 25   | Motor/parietal; areas 2/5 | ~5                         | 1,000         | Reaching                       | 0.02–0.04 |
| 88   | Substantia nigra          | 58                         | 500           | Cue matching                   | 0.01–0.04 |

Variability in  $r_{sc}$  across studies can largely be explained by differences in firing rate

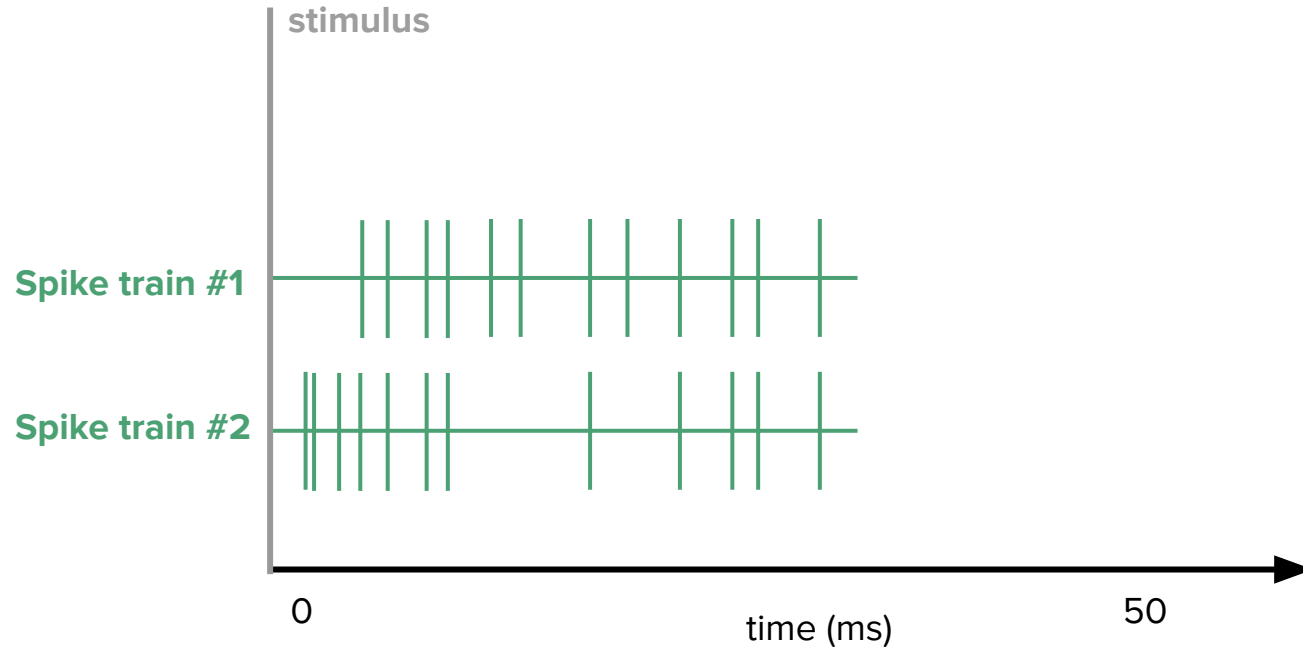
From Cohen & Kohn, 2011





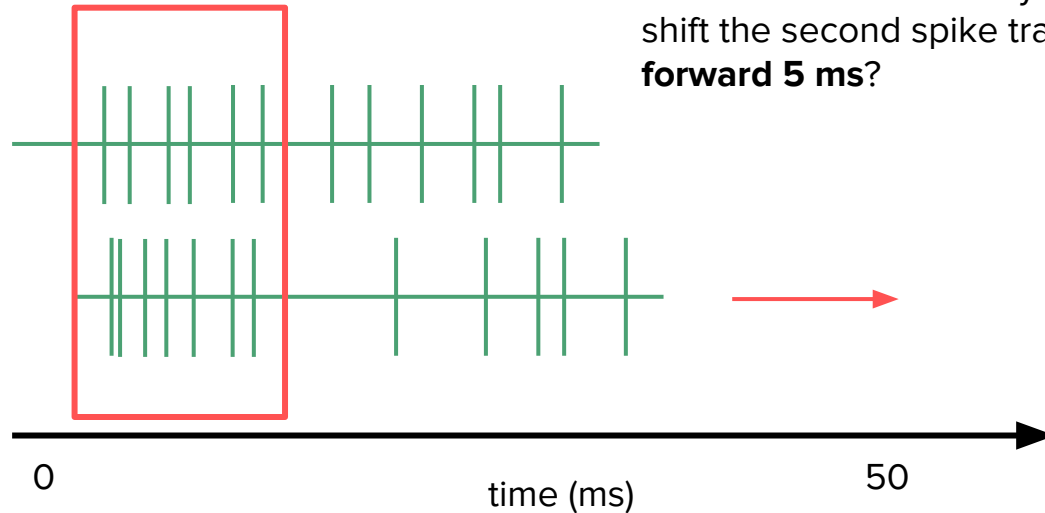
Spike count correlations change how well a population of neurons can decode a stimulus

We can also ask how correlated spike trains are, by shifting them forward and backward in time

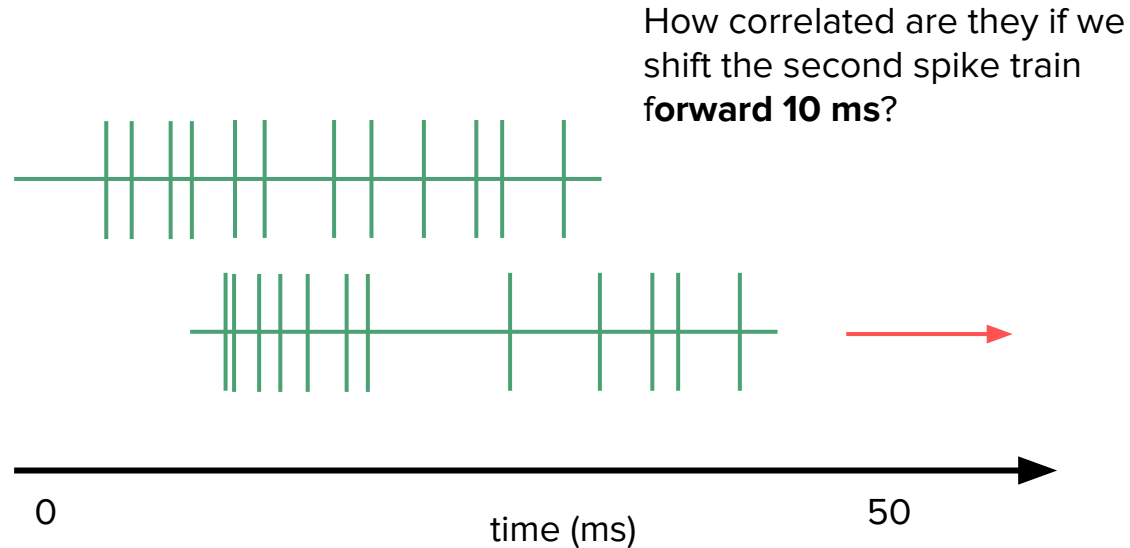


We can also ask how correlated spike trains are, by shifting them forward and backward in time

bin & compute a  
Pearson's correlation

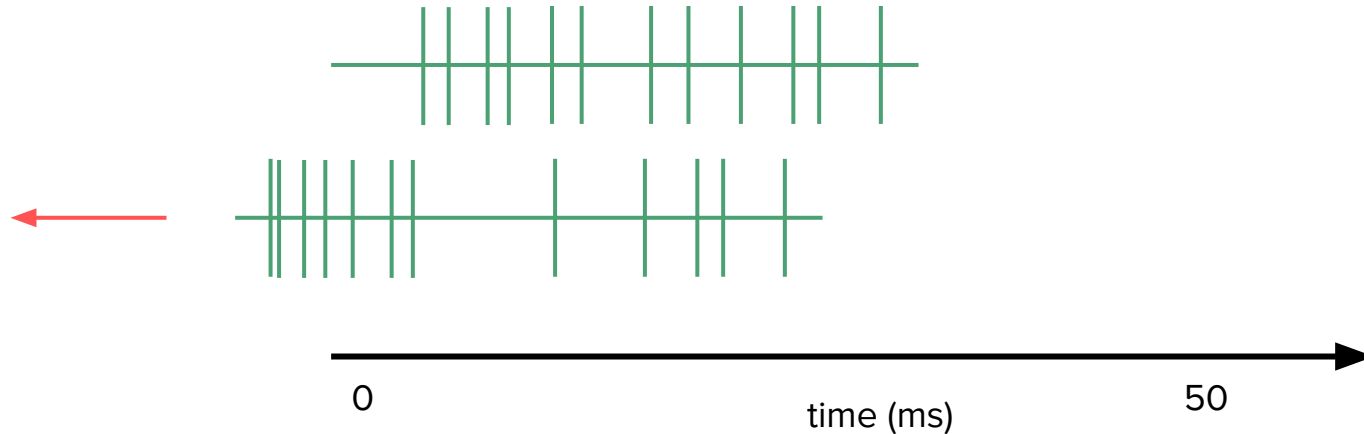


We can also ask how correlated spike trains are, by shifting them forward and backward in time



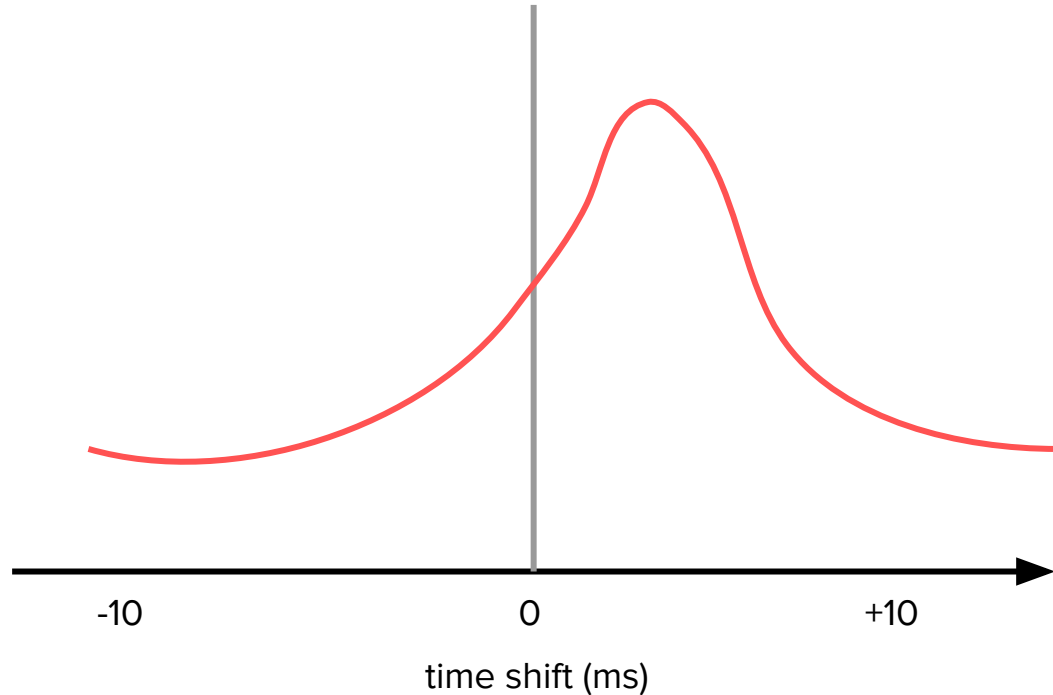
We can also ask how correlated spike trains are, by shifting them forward and backward in time

How correlated are they if we  
shift the second spike train  
**back 10 ms?**





We can also ask how correlated spike trains are, by shifting them forward and backward in time



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Article | [Open access](#) | Published: 28 January 2021

## Mouse entorhinal cortex encodes a diverse repertoire of self-motion signals

[Caitlin S. Mallory](#), [Kiah Hardcastle](#), [Malcolm G. Campbell](#), [Alexander Attinger](#), [Isabel I. C. Low](#), [Jennifer L. Raymond](#) & [Lisa M. Giocomo](#) 

[Nature Communications](#) **12**, Article number: 671 (2021) | [Cite this article](#)

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### Abstract

Neural circuits generate representations of the external world from multiple information streams. The navigation system provides an exceptional lens through which we may gain insights about how such computations are implemented. Neural circuits in the medial



Caitlin Mallory  
(Postdoc @  
UC Berkeley)



Lisa Giocomo  
(Professor @  
Stanford)

Dataset that we'll explore today!

## Additional Resources (Signal processing)

<https://mark-kramer.github.io/Case-Studies-Python/03.html>

<https://voyteklab.com/oscillations/publications/interpreting-spectrum/>

### **Related UCSD classes:**

COGS 118C. Neural Signal Processing

DSC 120. Signal Processing for Data Analysis