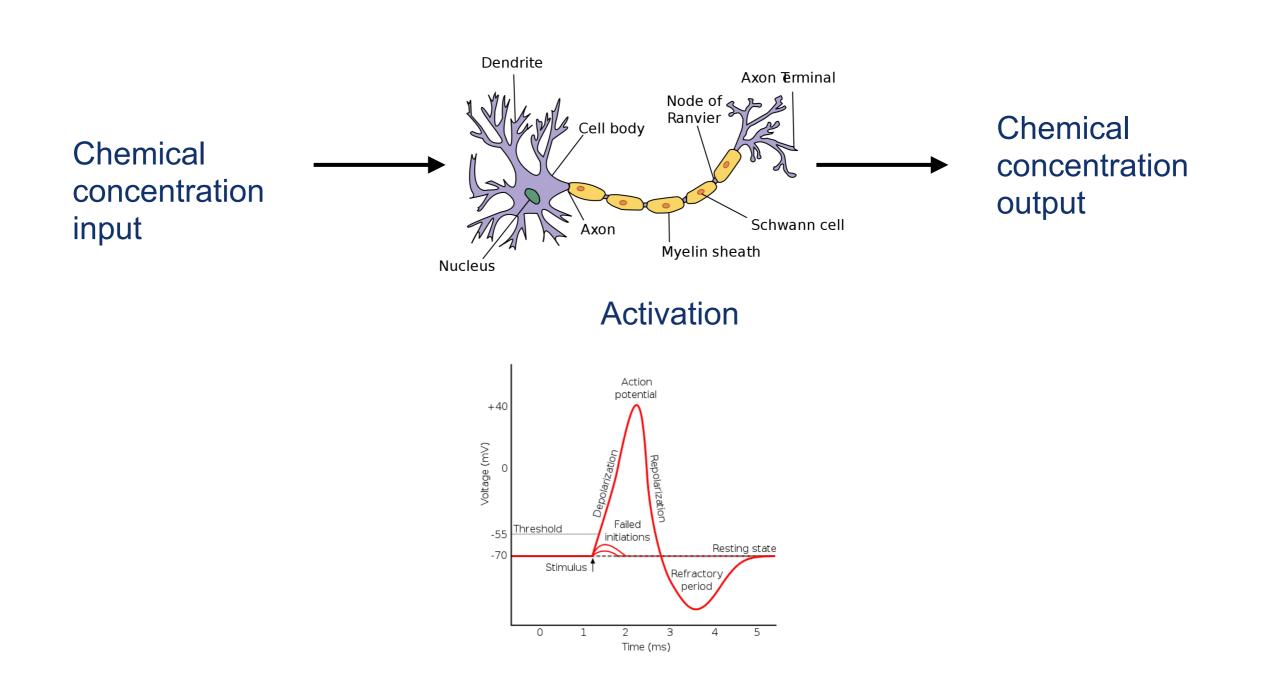
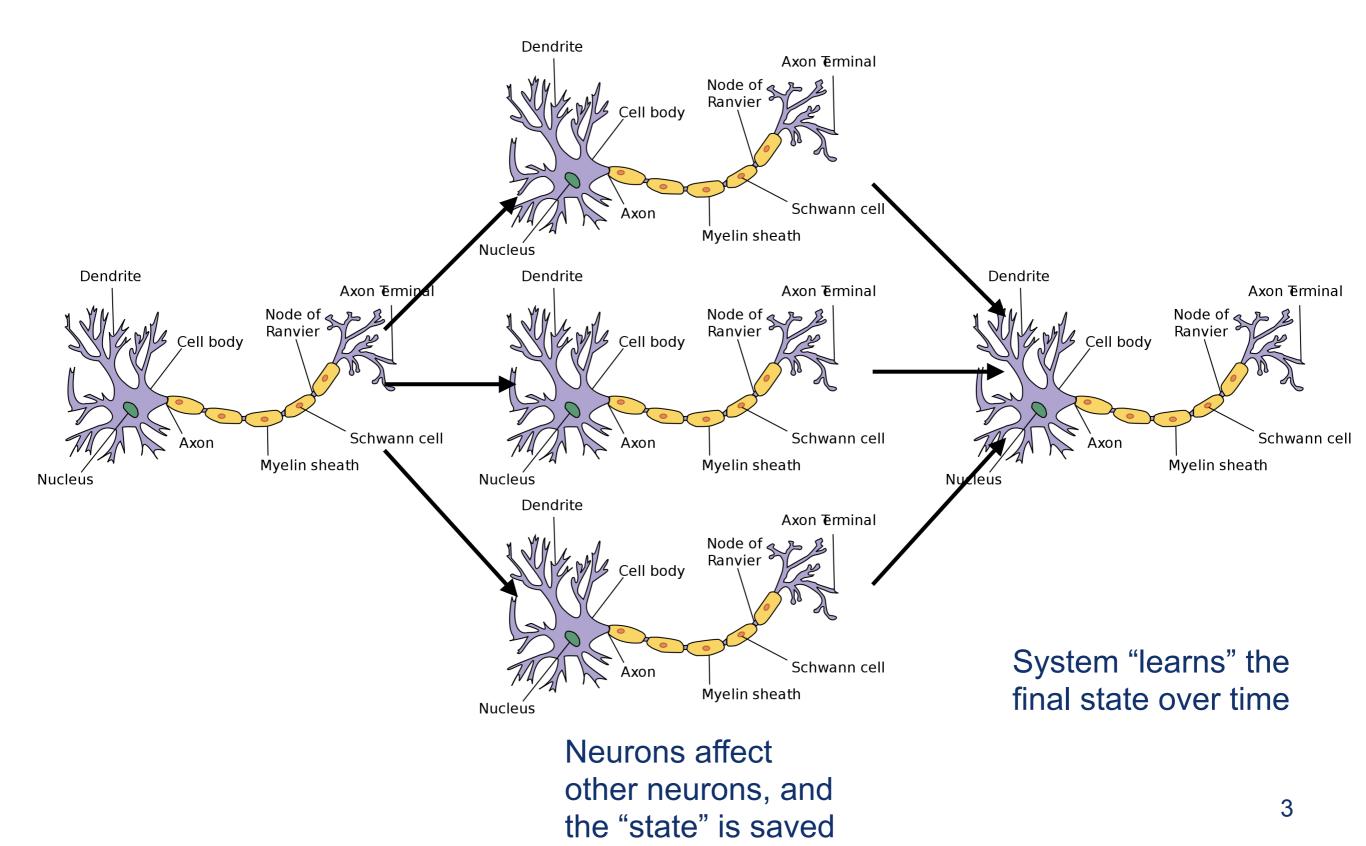
# PY410 / 505 Computational Physics 1

**Salvatore Rappoccio** 

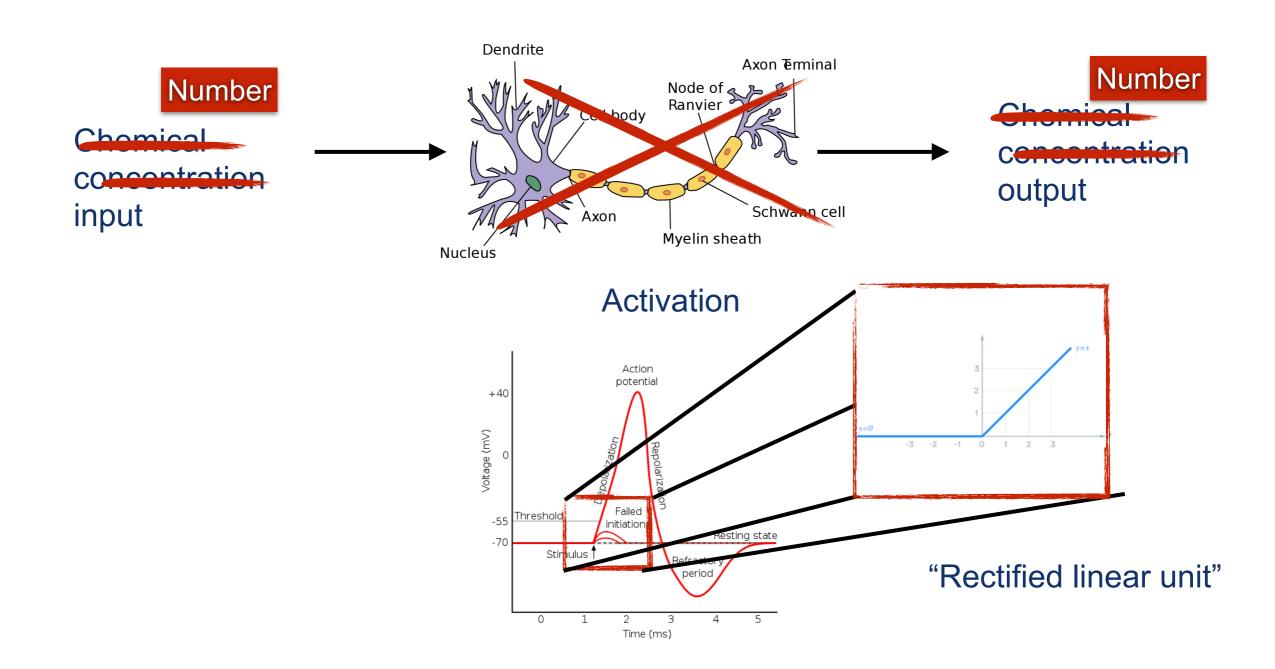
## Real Neural Networks



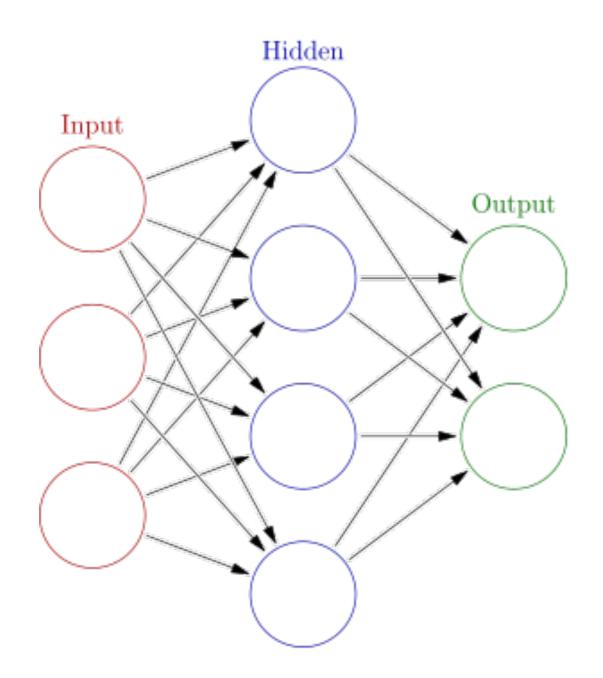
# Real Neural Networks

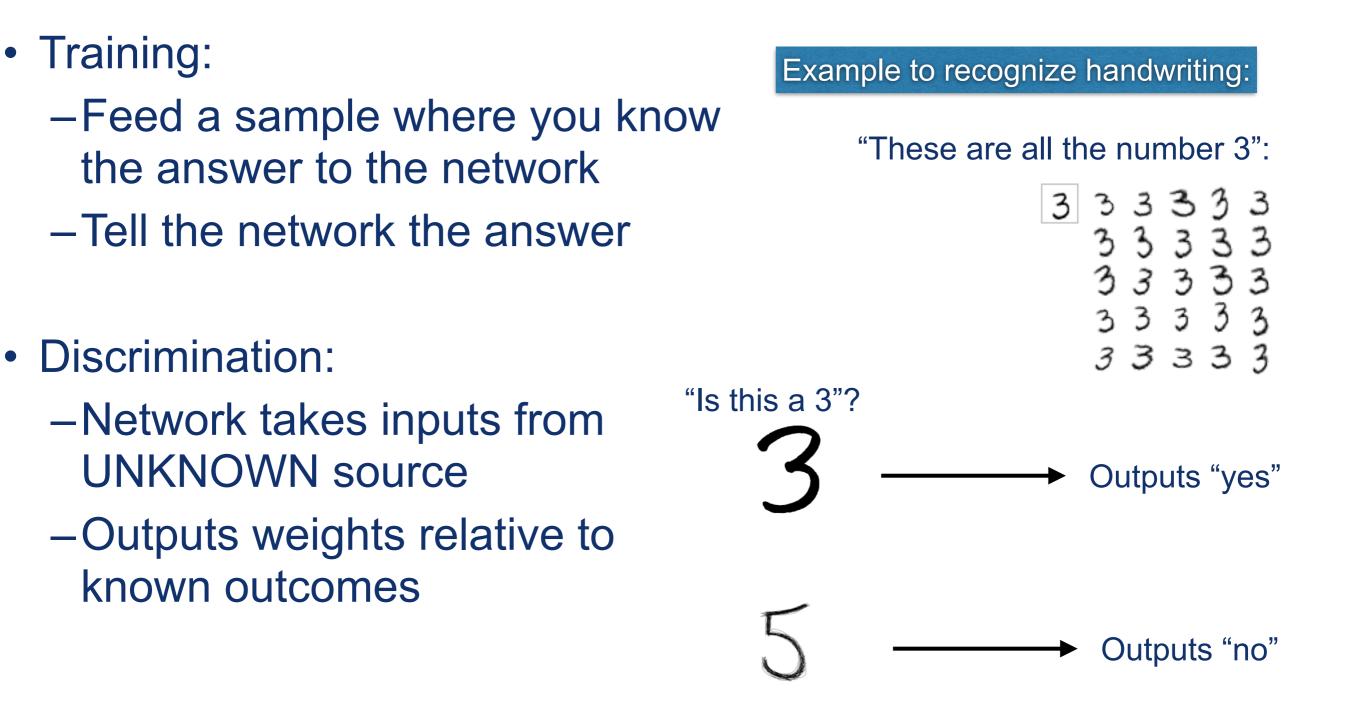


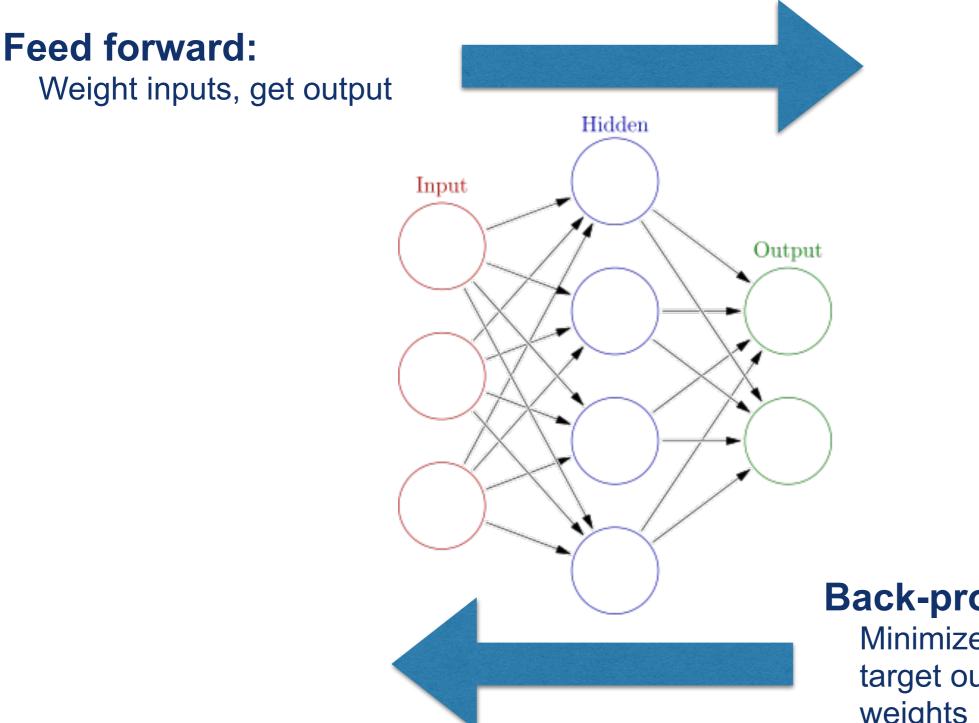
# Artificial Neural Networks



• The neural network is a group of neurons working together in such a persistent state:







#### **Back-propagate:**

Minimize difference between target output and current weights

Weights 1

Weights 2

Output Layer

2

4

8

-2

-4

 Consider a 2-layer network • A bit more formally: -Input vector  $\vec{x}$ -Output vector  $\vec{y}$ –Weights and biases between layers W and b-Activation function  $\sigma$ Input Layer Hidden Layer 10 Usually rectified linear unit (ReLU), but we will use sigmoid because it is differentiable. x 0.5 0.0

Adapted from <u>https://towardsdatascience.com/how-to-build-your-own-neural-network-from-scratch-in-python-68998a08e4f6</u>

- For each training step:
  - -Feed forward  $\vec{x} \rightarrow \hat{y}$ :
  - $\hat{y} = \sigma(W_2\sigma(W_1\vec{x} + b_1) + b_2)$
  - -Compute loss function (least squares):

$$L(y, \hat{y}) = \sum_{i} (y - \hat{y})^2$$

- -Compute gradient of loss function wrt weights  $\frac{\partial L(y, \hat{y})}{\partial W}$
- -Back propagate :
  - -Optimize using gradient descent (like BFGS!)

Caveats about optimization in multiple dimensions hold here!

9

Adapted from

https://towardsdatascience.com/how-to-build-your-own-neural-network-from-scratch-inpython-68998a08e4f6

# **Deep Learning and Neural Networks**

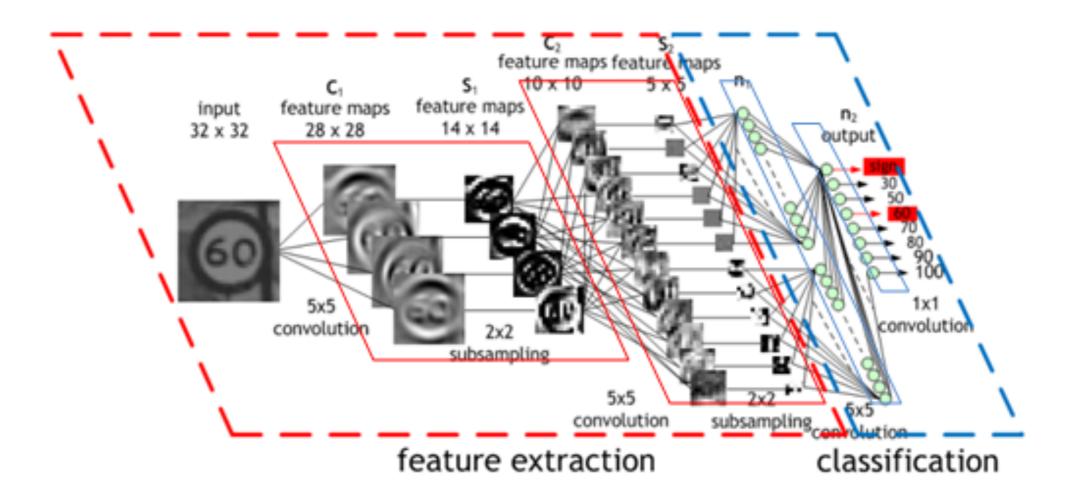
- Next step: instead of teaching, let the neural network learn on its own
  - -Example 1: Convolutional Neural Network
    - Extract features using convolution, pass features to neural network
    - Fixed-size inputs: Suitable for images
  - -Example 2: Recurrent Neural Network
    - Instead of "feed forward" like last time, has possible recurrent links
    - Any size inputs: Suitable for text / speech / handwriting recognition
    - Can also be Recursive (don't get confused!)

https://en.wikipedia.org/wiki/Deep\_learning https://en.wikipedia.org/wiki/Recurrent\_neural\_network https://en.wikipedia.org/wiki/Recursive\_neural\_network https://en.wikipedia.org/wiki/Convolutional\_neural\_network https://devblogs.nvidia.com/parallelforall/deep-learning-nutshell-core-concepts/

# **Deep Learning and Neural Networks**

- Options to train:
- Supervised
  - -Give inputs, tell it what the output is
- Unsupervised
  - -Give inputs, tell it to optimize a function

- Preprocessing to perform feature extraction
  Convolution or otherwise
- Classification
  - -Standard neural network



Does "preprocessing" of the image by convoluting with kernels:

#### 2-d Fourier transform!

Input image



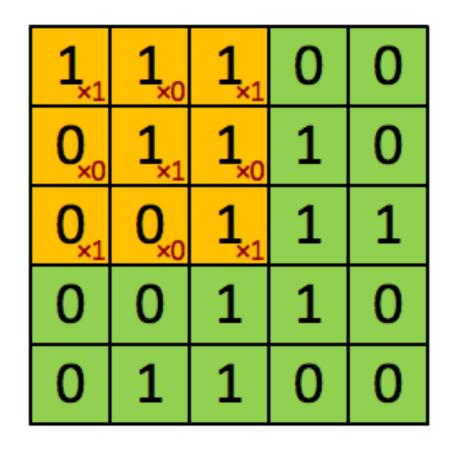
Convolution Kernel

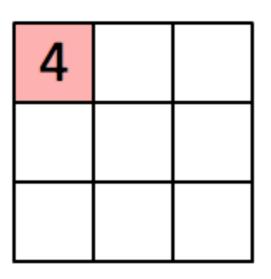
$$\begin{bmatrix} -1 & -1 & -1 \\ -1 & 8 & -1 \\ -1 & -1 & -1 \end{bmatrix}$$

#### Feature map



- Animation of convolution:
- Kernel: 1 0 1 0 0 1 0 1 0 1 0

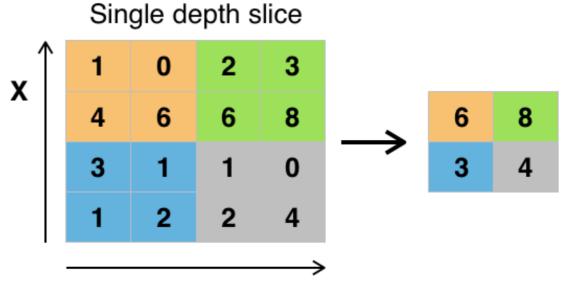




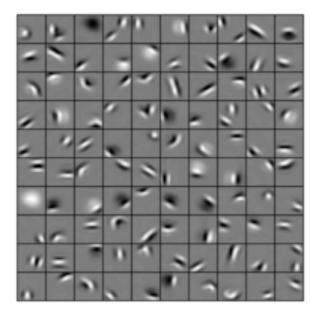
Image

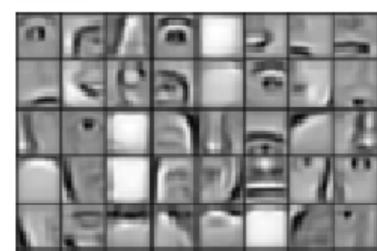
Convolved Feature

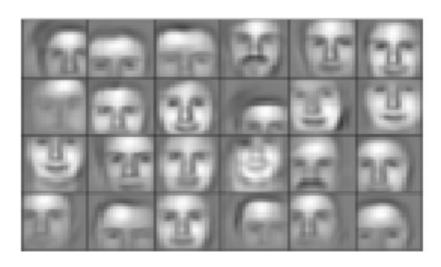
- Pooling layer : used to obtain very big features
  - -Downsampled to extract gross features
  - -ls it "pointy"?
  - -ls it "round"?
  - -Does it have a face?



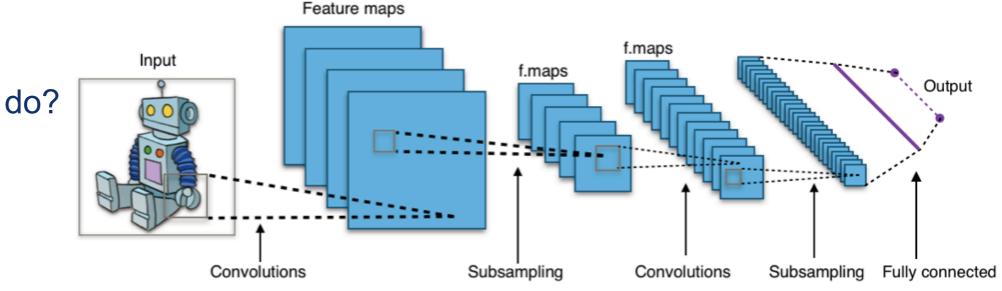






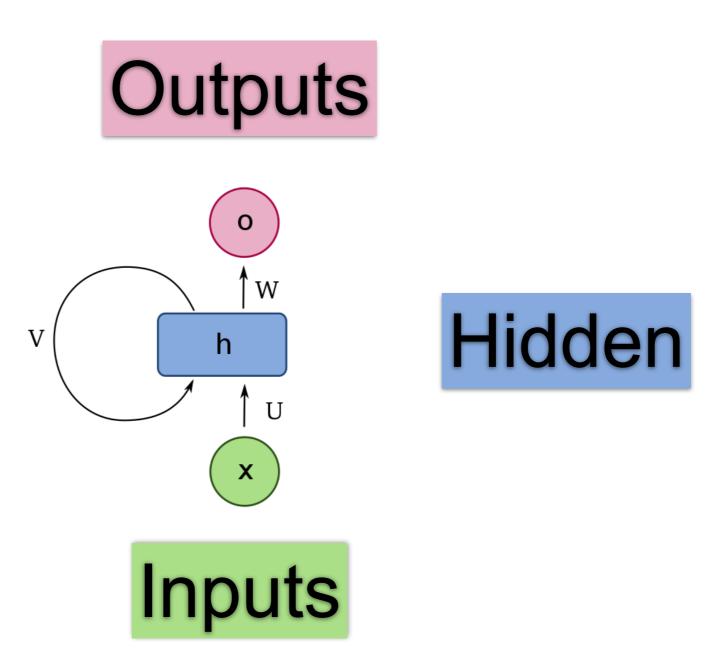


- Putting it together:
  - Convolutional layer
    - Discern features
  - Pooling layer
    - Extract global features
  - Activation function ("Rectified Linear Units")
    - Standard NN
  - Fully connected layer
    - Standard NN
  - Loss layer
    - How well did it do?



#### https://en.wikipedia.org/wiki/Convolutional\_neural\_network

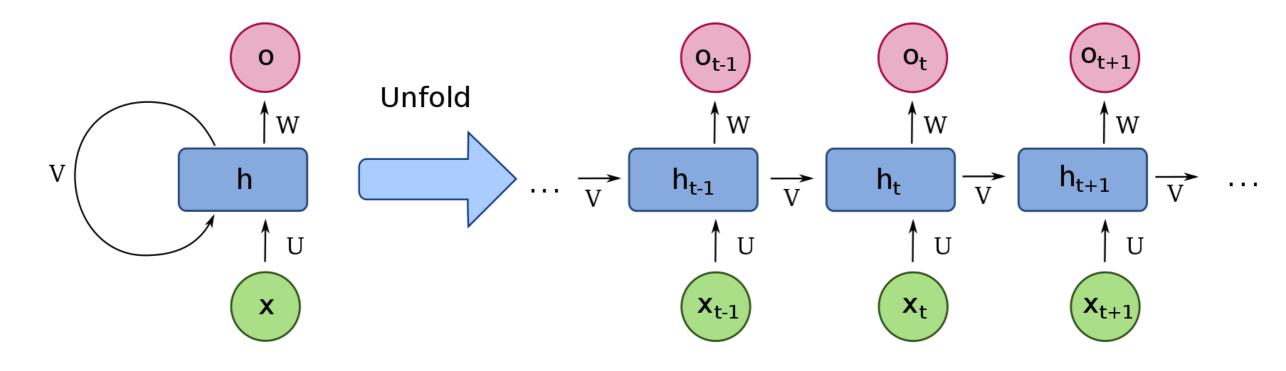
Dynamic directed graph to process sequences of inputs



https://en.wikipedia.org/wiki/Recurrent\_neural\_network

Dynamic directed graph to process sequences of inputs

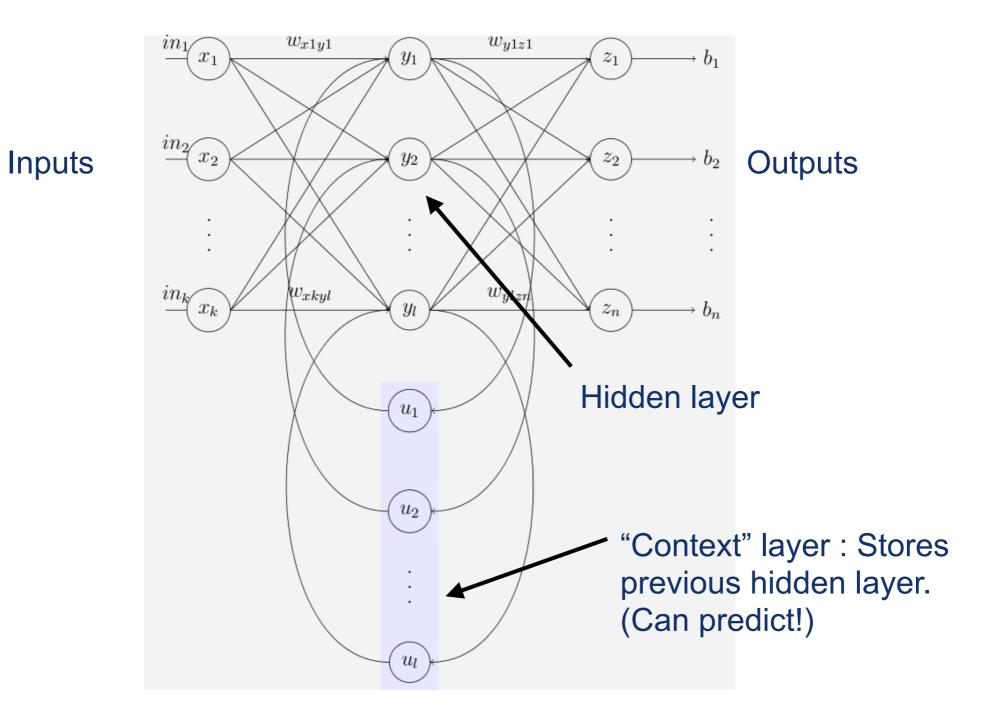
"Dynamic" in the sense that it changes over time



https://en.wikipedia.org/wiki/Recurrent\_neural\_network

- "Recurrent": Can "save state" of the inputs –(CNNs do not)
- Has use in predicting the next value in a sequence (speech, text recognition)
- Can also see how similar two sequences are

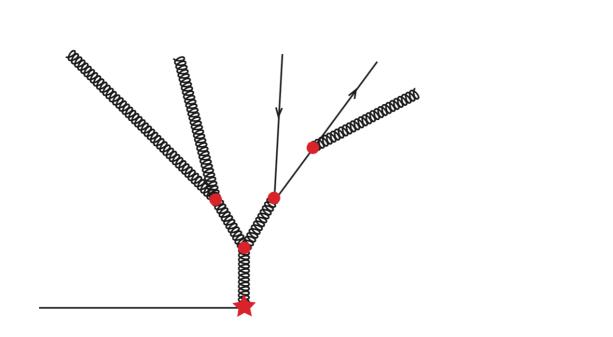
• Example: Elman network

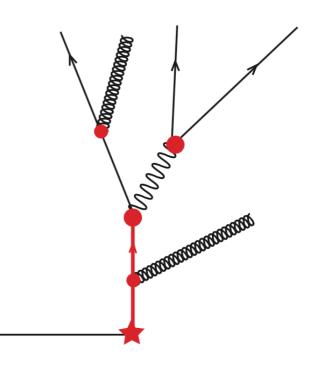


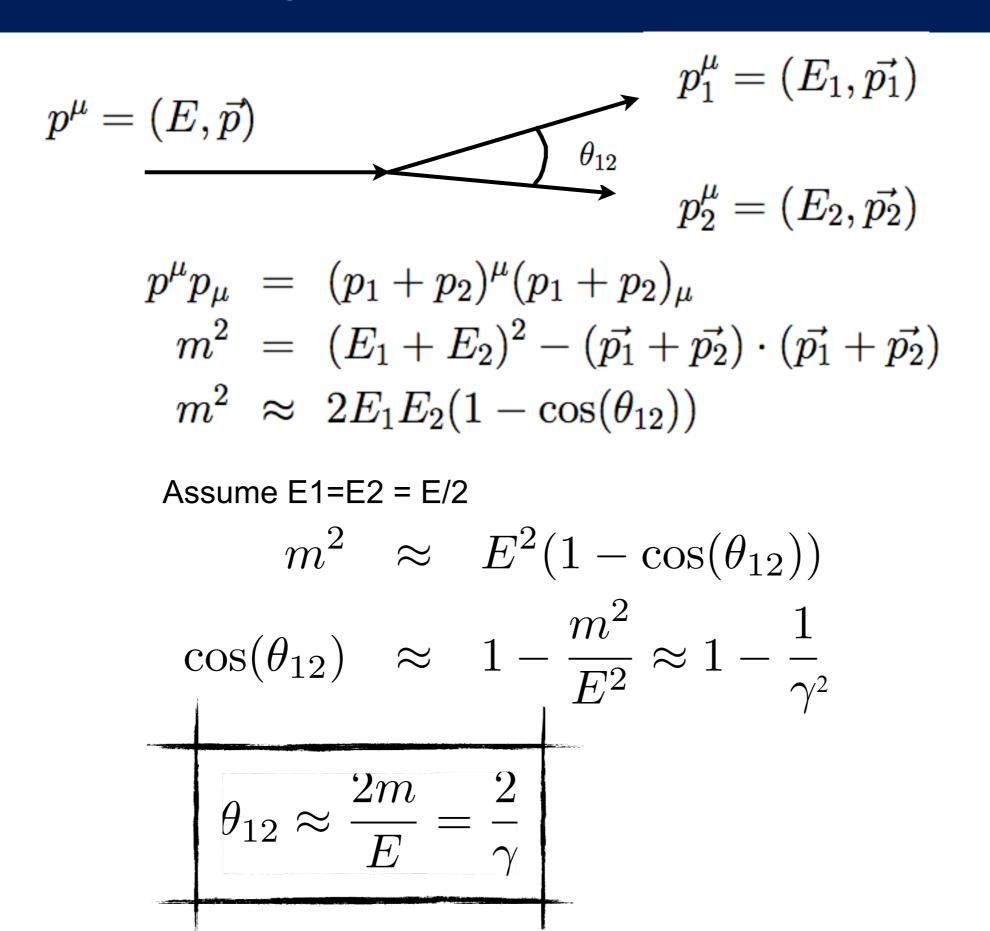
• Example from physics:

-"Learning" the origin of a jet at the LHC

- W bosons can decay to quarks
  - If they obtain a high energy, their decay products merge together
  - -How to distinguish between this and standard QCD jets?



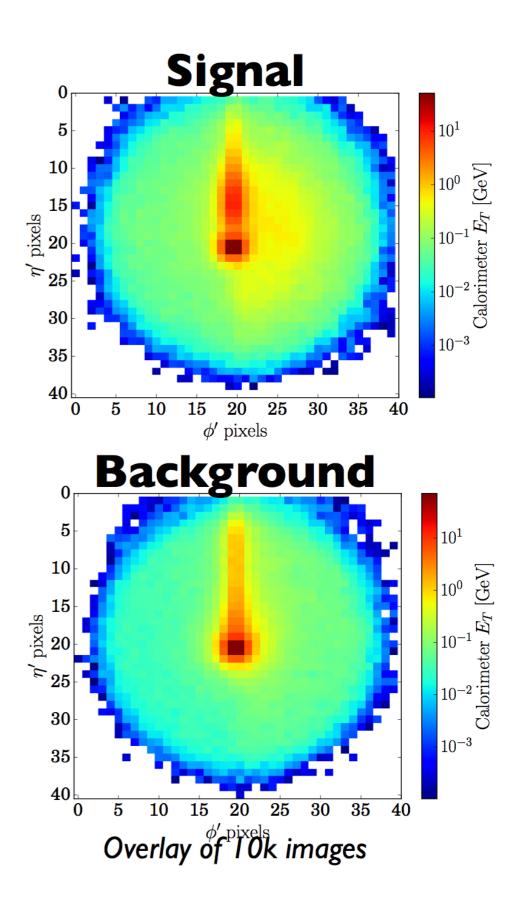


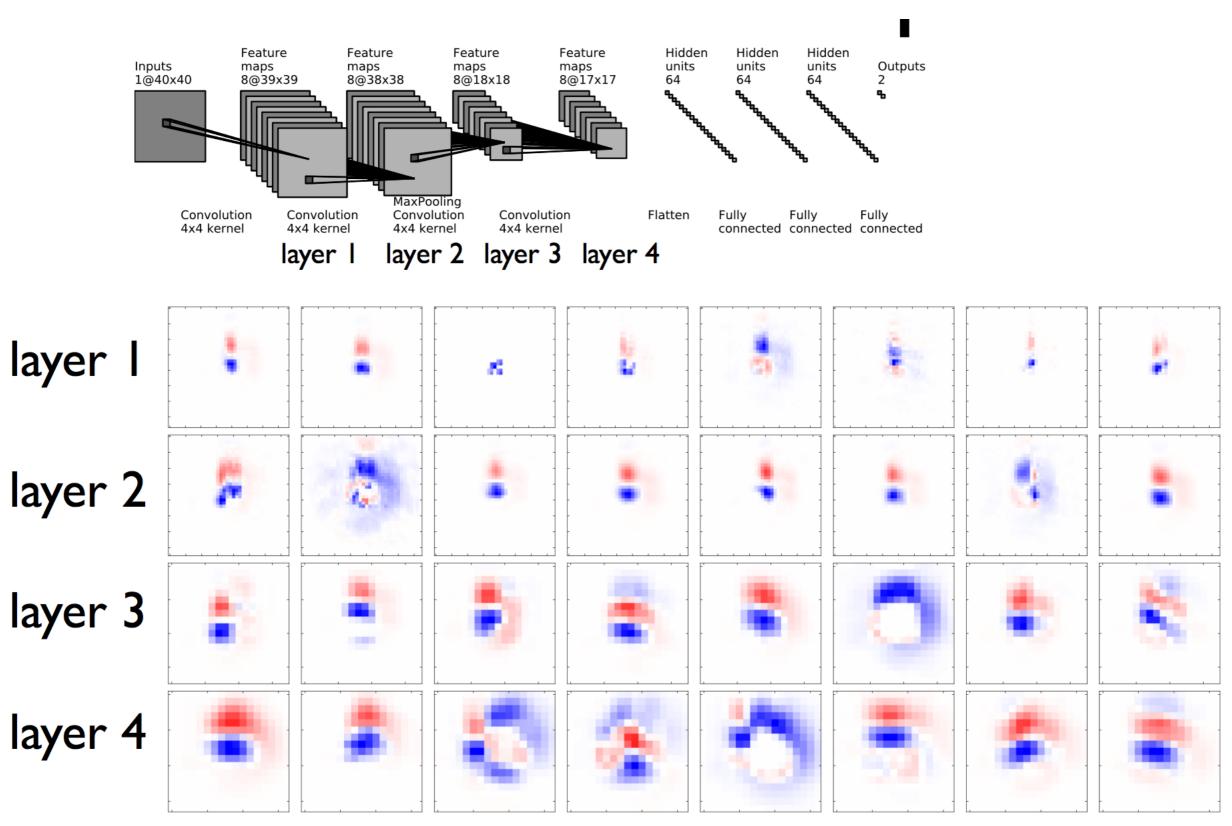


22

- Plot energy of particles as a 2d projection
- Take average, put at center
- Take "next blob", put at top center
- Look at sum of many images
- Use image processing techniques!

arXiv:1407.5675 JHEP 1607 069 JHEP 05 (2017) 006





# Software

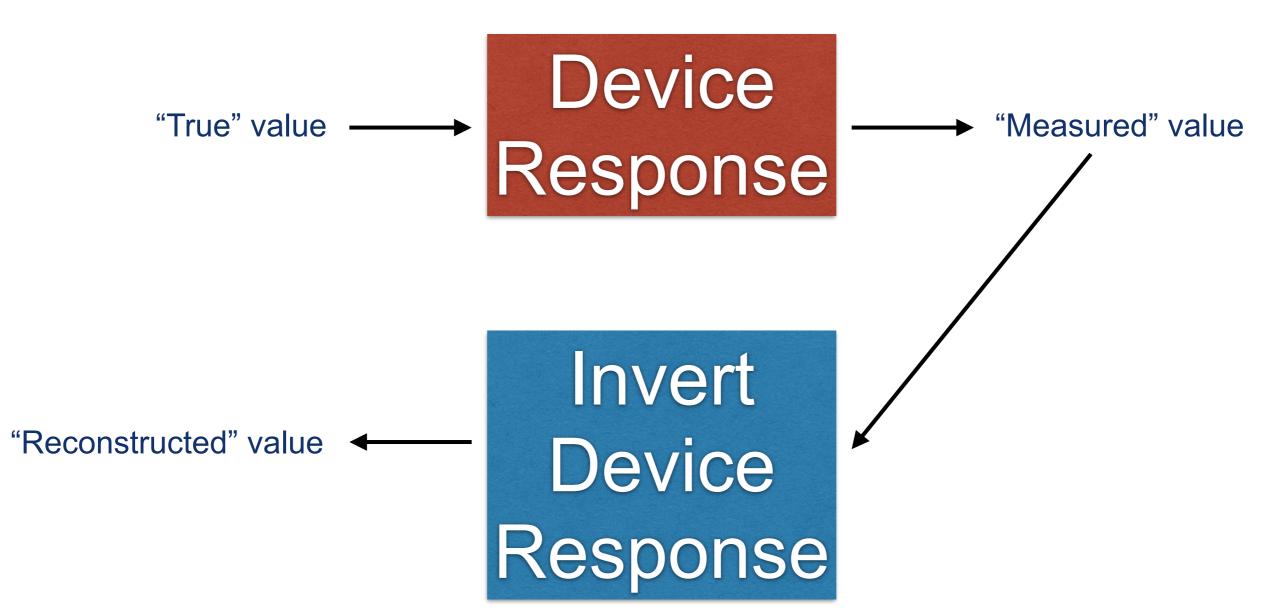
- Lots of Deep Network software out there
- Popular ones:
  - -Theano
  - -TensorFlow
  - -Scikit-learn
  - -etc
- Theano and tensorflow are both installed in vidia, so we can work through an example there:

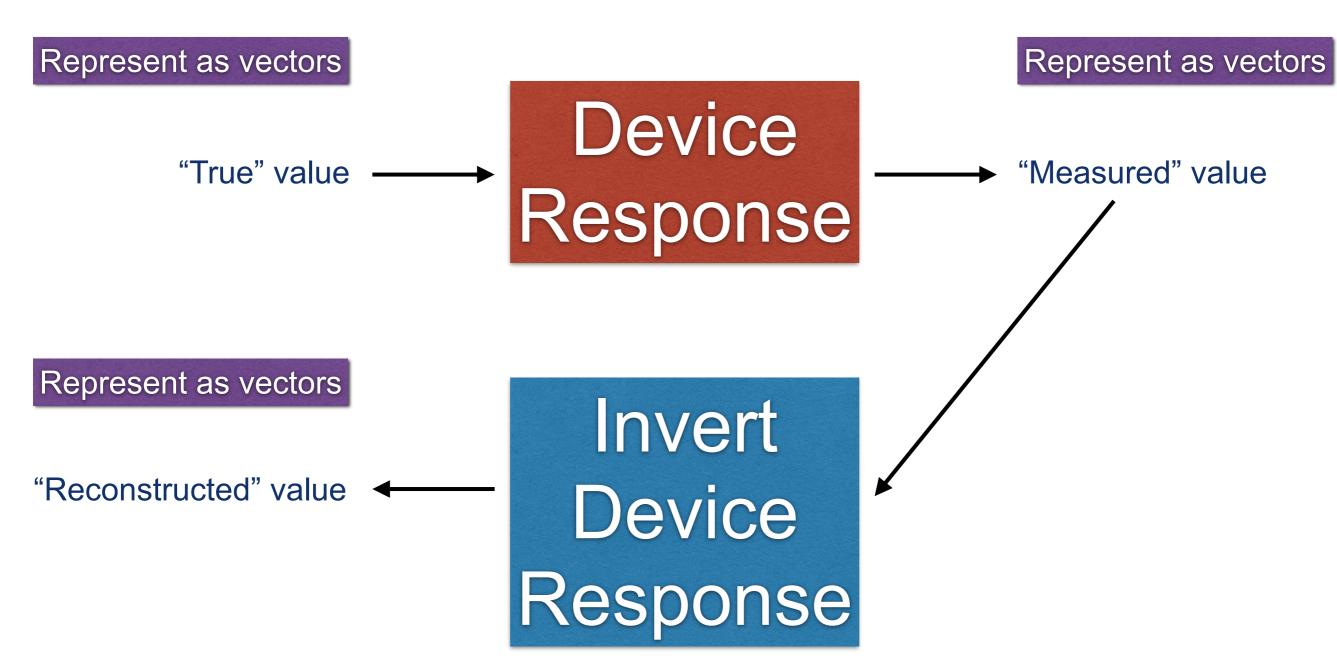
-<u>https://www.tensorflow.org/tutorials/</u>

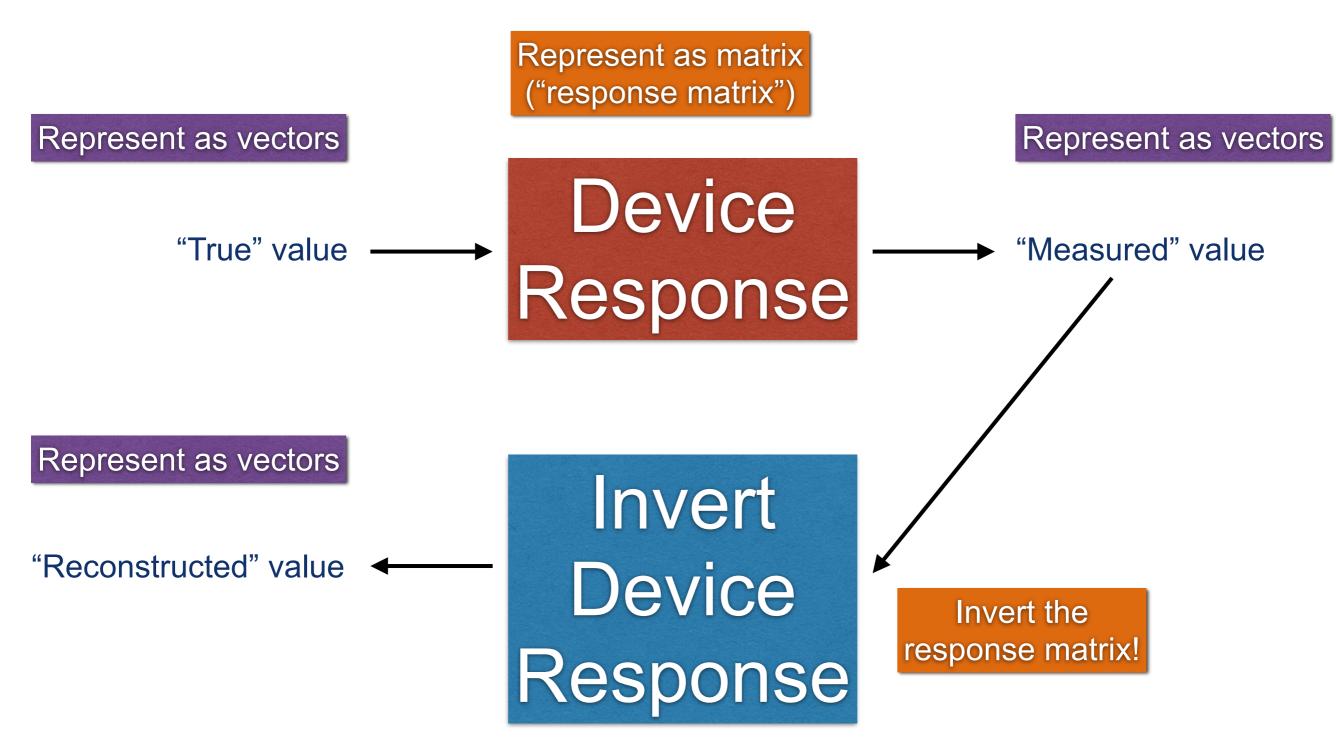
- The inverse problem is, to first order, "simply" inverting functions
- Linear functions: trivial: f(x) = mx + b-But! If m = 0, ????  $f^{-1}(x) = \frac{1}{m}x - \frac{b}{m}$
- Everything else: not even guaranteed to exist!
  –Can be many-to-one mapping
  - -Can be non-invertible

• Becomes even trickier in multiple dimensions!

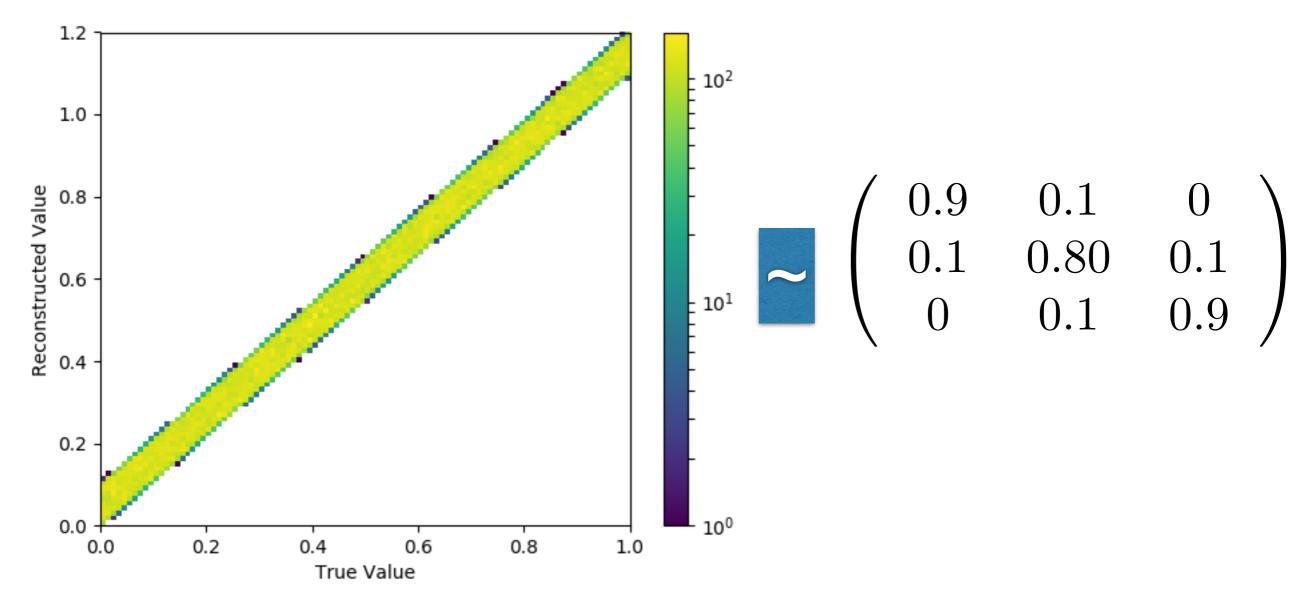




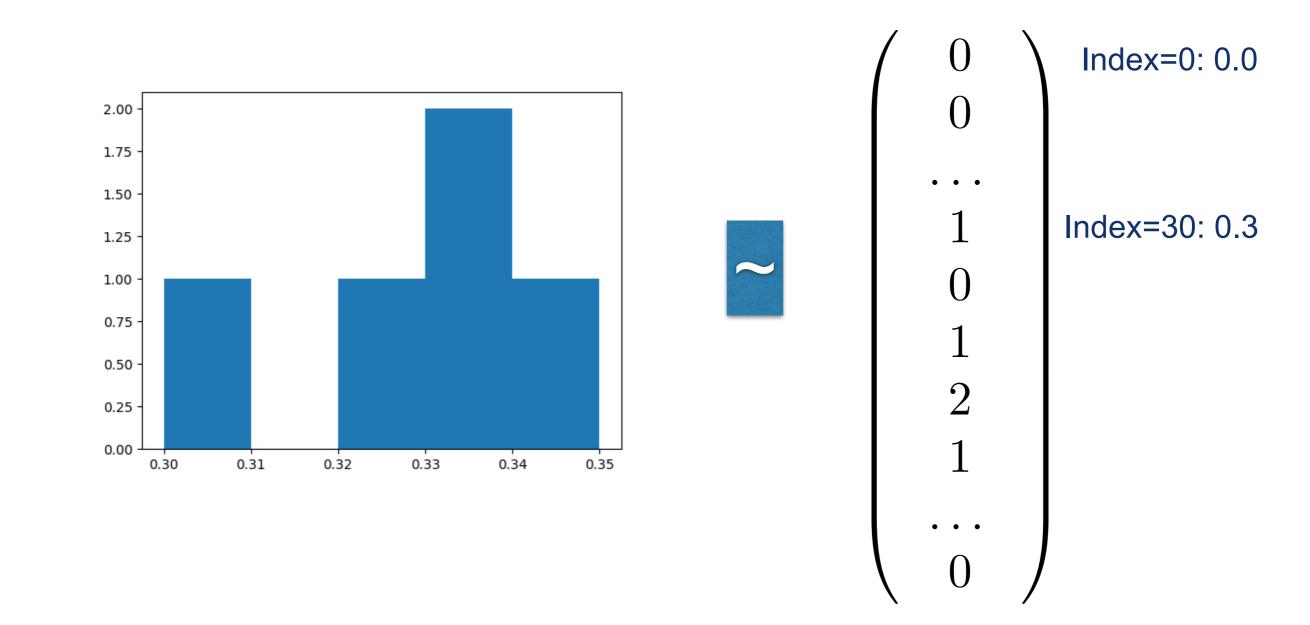


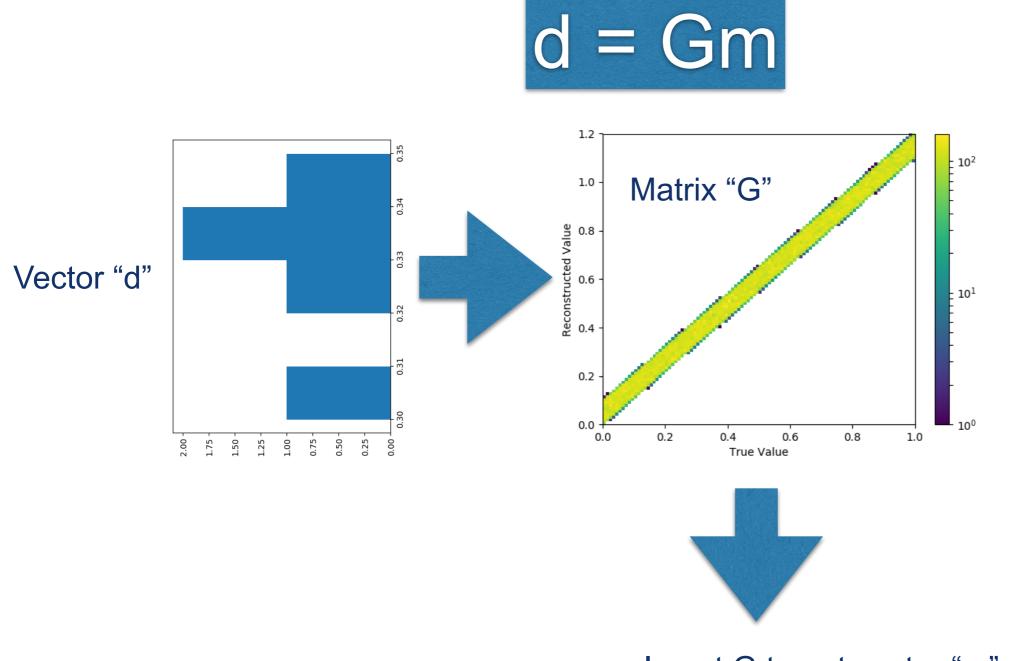


- But! Need one of these for EVERY possible input (0, 0.1, 0.1... 1.0)
- Represent as a matrix:



• Represent our histogram as a vector





Invert G to get vector "m"

- You're aware of the first case: Linear inverses (i.e. inverting matrices)
- There are generalizations
- Reminder:
  - -To solve  $\vec{y} = A\vec{x} + \vec{b}$ :
  - -Invert the matrix:

$$\vec{x} = A^{-1} \left( \vec{y} - \vec{b} \right)$$

-Gauss-Jordan elimination, other techniques we did last semester

- What if the matrix is not invertible?
  - -Can still get information, but not perfectly determined
  - -Often sufficient to have partial information
- Examine the over-constrained case:
- Suppose we have a matrix G (the "observation matrix") with data "d" and true value "m".
- Want to minimize the difference between the prediction (Gm) and the data:  $\label{eq:phi} \phi = |d-Gm|^2$

Where 
$$|a|^2 = (a^T a)$$

• Want to find the place where the difference is minimized!

Difference minimized when

 $\nabla_m \phi = 0$ 

- That is, the gradient wrt the "m" components is zero
- Using chain rules for matrix functions:

$$\nabla_m \phi = 2 \left( G^T G m - G^T d \right) = 0$$

 Needs to be satisfied for all of the components, so require each to vanish:

$$G^T G m = G^T d$$

• So we solve for m:

$$m = \left( G^T G \right)^{-1} G^T a$$

Least squares distance from last semester!

https://en.wikipedia.org/wiki/Inverse\_problem https://atmos.washington.edu/~dennis/MatrixCalculus.pdf

- So the inverse problem is similar to least squares
- Don't get the "full inverse" but do get the closest to it
- For intuition, remember our "design matrix" from last semester:  $V_i(x_i)$

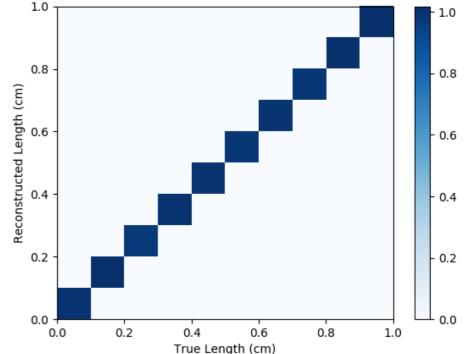
$$A_{ij} = \frac{Y_j(x_i)}{\sigma_i}$$

$$\mathbf{A} = \begin{bmatrix} Y_1(x_1)/\sigma_1 & Y_2(x_1)/\sigma_1 & \dots \\ Y_1(x_2)/\sigma_2 & Y_2(x_2)/\sigma_2 & \dots \\ \dots & \dots & \dots & \dots \end{bmatrix}$$

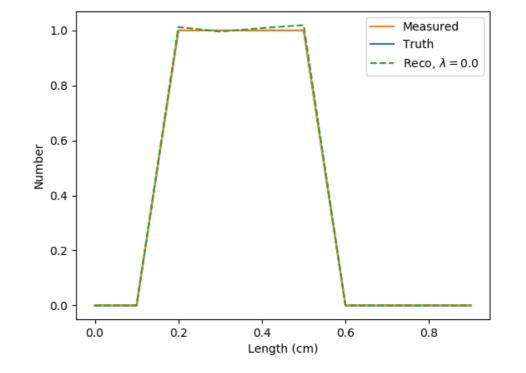
 $\vec{a} = (\mathbf{A}^T \mathbf{A})^{-1} \mathbf{A}^T \vec{b}$ 

• Then for polynomial fits  $A_{ij} = a_j x_i^j / \sigma_i$ 

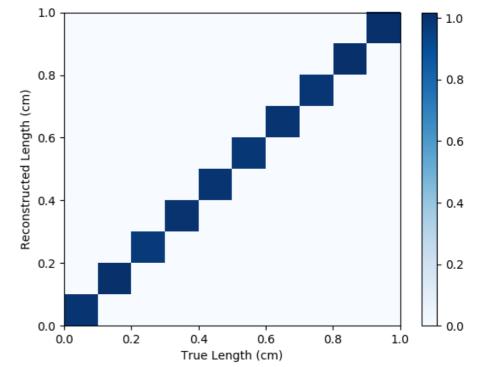
- Try to invert the response of some experimental device or detector to get "true" values
  - -Example 1:
    - Device to measure length is perfect.
      - Response created from 100000 "pseudo-experiments":



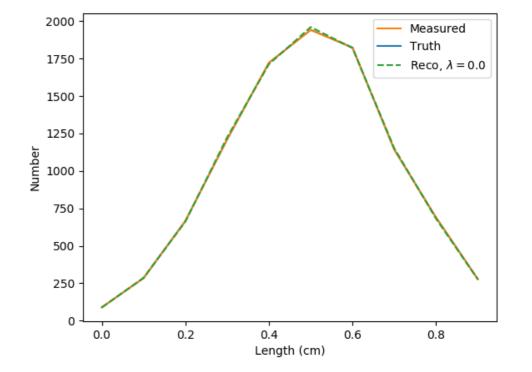
- You measure 4 objects, and obtain:
  - -0.2 cm, 0.3 cm, 0.4 cm, 0.5 cm
  - Measured vs true histogram:



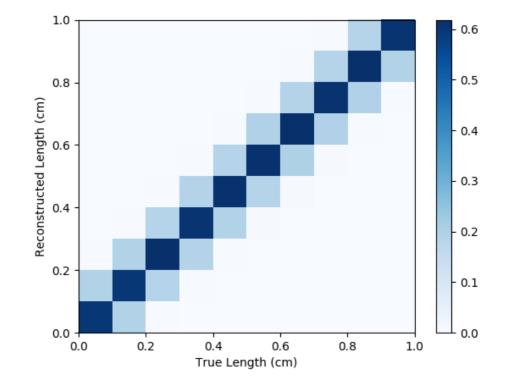
- Try to invert the response of some experimental device or detector to get "true" values
  - -Example 2:
    - Device to measure length is perfect.
      - Response created from 100000 "pseudo-experiments":



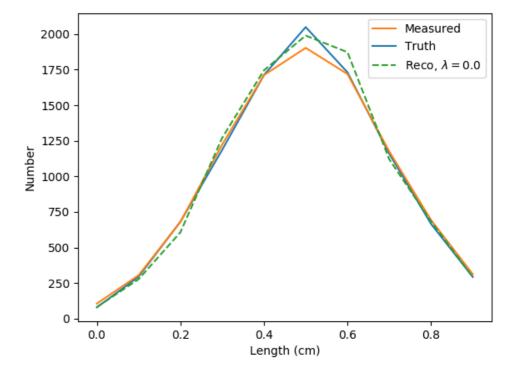
- You measure 10000 objects, and obtain:
  - Gaussian with width = 0.2, mean = 0.5
  - Measured histogram:



- Try to invert the response of some experimental device or detector to get "true" values
  - -Example 3:
    - Device to measure length has resolution of 0.05 cm
      - Response created from 100000 "pseudo-experiments":



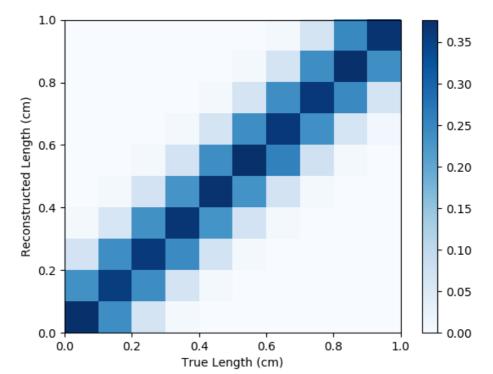
- You measure 10000 objects, and obtain:
  - Gaussian with width = 0.2, mean = 0.5
  - Measured histogram:

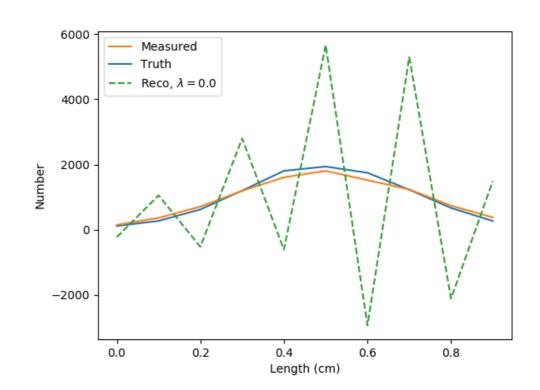


- Try to invert the response of some experimental device or detector to get "true" values
  - -Example 4:
    - Device to measure length has resolution of 0.1 cm
      - Response created from
        100000 "pseudo-experiments":



- Gaussian with width = 0.2, mean = 0.5
- Measured histogram:





- What is going on???
- Statistical uncertainties bin-to-bin are greatly amplified by inversion
  - -Need to damp those out... "regularization" (from Tikhonov)

- How to formally do this?
- Ordinary formula for the inversion is:

$$m = \left( G^T G \right)^{-1} G^T d$$

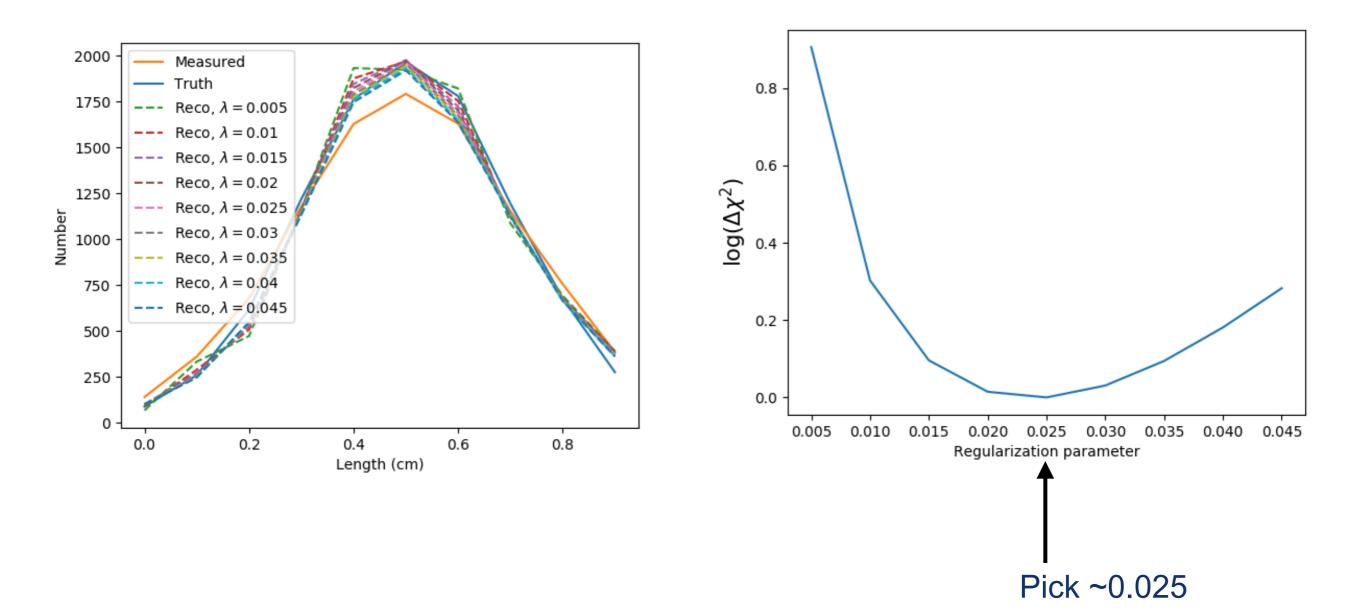
 Introduce a penalty term (here, a Lagrange multiplier) to solve:

$$m = \left( G^T G + \lambda I \right)^{-1} G^T d$$

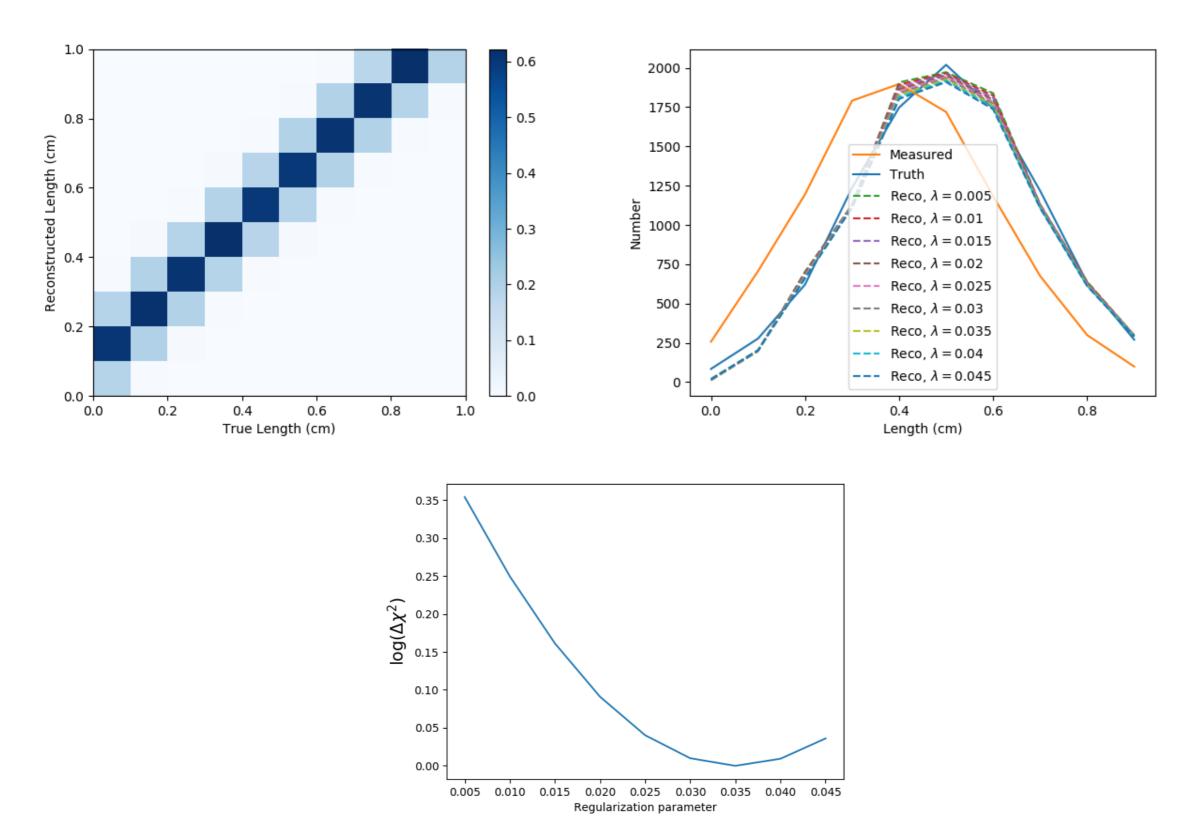
Can then vary as a function of lambda to pick when the regularization is complete

https://www.researchgate.net/publication/ 274138835\_NumPy\_SciPy\_Recipes\_for\_Dat a\_Science\_Regularized\_Least\_Squares\_Opti mization

• Example 4: vary regularization parameter for our example:

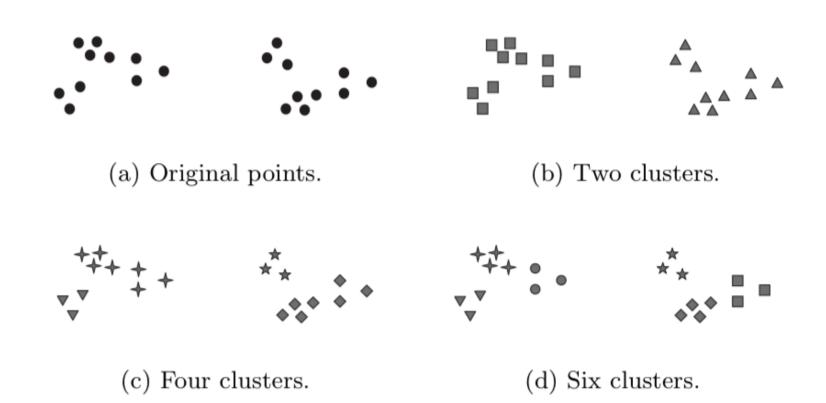


• Example 5: Can also have biases



- Sometimes people decide to put the biases in a separate matrix so it is a "diagonal matrix"
- In practice it doesn't matter.

- Goal: Given N points in space, associate to k partitions
- Many applications:
  - -Data classification
  - -Galaxy clustering
  - -Jet clustering
  - -Sociology
  - -Social networks



https://www-users.cs.umn.edu/~kumar001/dmbook/ch8.pdf https://en.wikipedia.org/wiki/Cluster\_analysis

- Goal: Given N points in space, associate to k partitions
- Many applications:
  - -Data classification
  - -Galaxy clustering
  - -Jet clustering
  - -Sociology
  - -Social networks



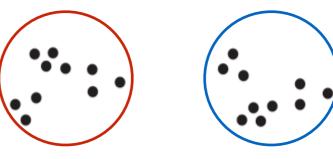
Basket case

- Goal: Given N points in space, associate to k partitions
- Many applications:
  - -Data classification
  - -Galaxy clustering
  - -Jet clustering
  - -Sociology
  - -Social networks



- Uses:
  - Summarize / compress "spatial" information
  - Find nearest neighbors
- Types of clustering algorithms:
  - Exclusive:
    - Partitional: divide into k exclusive categories
    - Hierarchical: can have sub-clusters
  - Non-exclusive:
    - Add same element to more than
      one cluster
  - Fuzzy:
    - Weight elements according to cluster
- Each can be either complete or partial

### Partitional

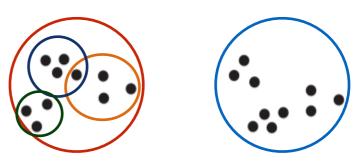


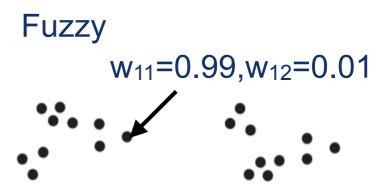
### Hierarchical





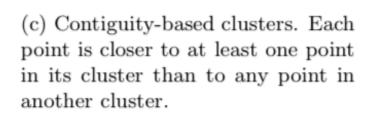
### Non-exclusive

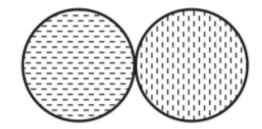




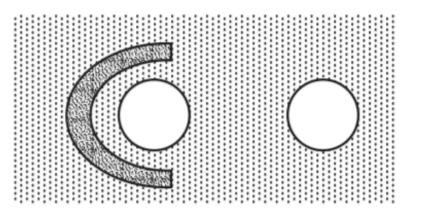
• Types of clusters:

(a) Well-separated clusters. Each point is closer to all of the points in its cluster than to any point in another cluster.

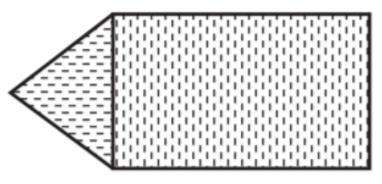


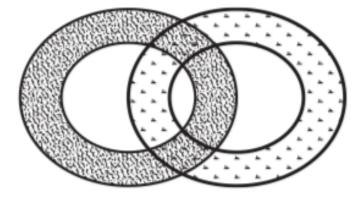


(b) Center-based clusters. Each point is closer to the center of its cluster than to the center of any other cluster.



(d) Density-based clusters. Clusters are regions of high density separated by regions of low density.





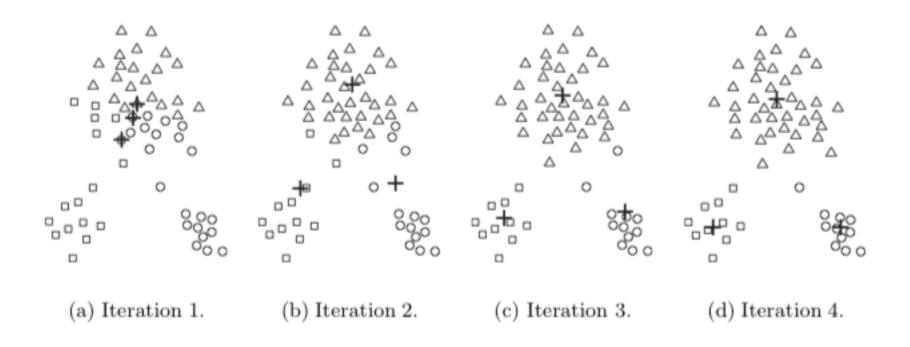
(e) Conceptual clusters. Points in a cluster share some general property that derives from the entire set of points. (Points in the intersection of the circles belong to both.)

- Popular categories :
  - -K-means
    - Partition into k clusters using mean central values (usually exclusively)
  - -Agglomerative hierarchical clustering
    - Pair individual elements into clusters given some distance metric
  - -Density based scan
    - Considers low-density regions to be noise, not exclusive clustering

### • K-means algorithm:

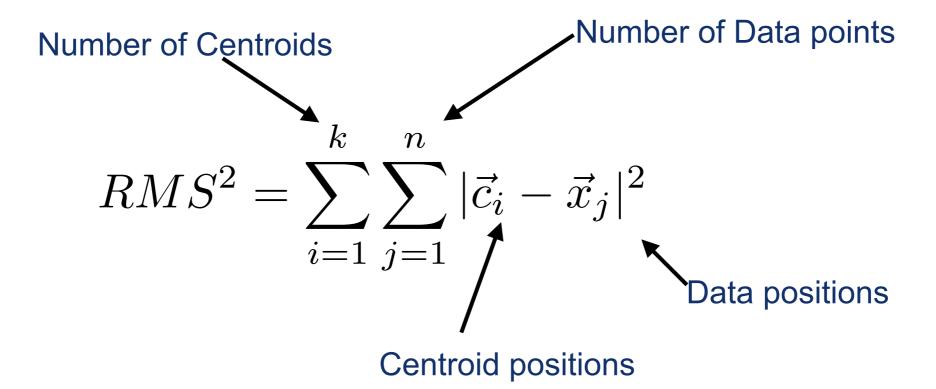
Algorithm 8.1 Basic K-means algorithm.

- 1: Select K points as initial centroids.
- 2: repeat
- 3: Form K clusters by assigning each point to its closest centroid.
- 4: Recompute the centroid of each cluster.
- 5: **until** Centroids do not change.



• K-means algorithm:

-Formally, computes RMS:



- -Minimize the RMS by adjusting the centroids
- -Note: Other distance metrics can be used, but the principle is the same (minimize the metric)

- Computational complexity is ~linear in product of:
  - -Number of points
  - -Number of "dimensions" (or attributes)
  - -Number of clusters
  - -Number of iterations to converge
- Shortcomings of k-means:
  - -As in all minimization routines, danger of local minima
    - Need heuristic methods to avoid them
  - -Can result in empty clusters if initialized poorly
  - -Outliers have disproportionate impact
- Can try to split and merge centroids to mitigate these!

• Alternative to k-means: Bisecting k-means:

Algorithm 8.2 Bisecting K-means algorithm.

1: Initialize the list of clusters to contain the cluster consisting of all points.

### 2: repeat

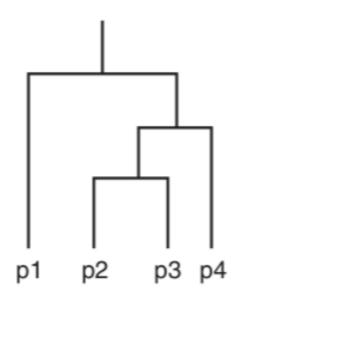
- 3: Remove a cluster from the list of clusters.
- 4: {Perform several "trial" bisections of the chosen cluster.}
- 5: for i = 1 to number of trials do
- 6: Bisect the selected cluster using basic K-means.
- 7: end for
- 8: Select the two clusters from the bisection with the lowest total SSE.
- 9: Add these two clusters to the list of clusters.
- 10: **until** Until the list of clusters contains K clusters.

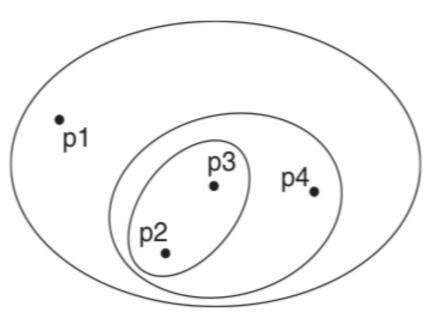
- Result is a Voronoi diagram
- Each point is closer to all points in its cell than other cells
- Also referred to as "cachement areas"
  - -From river basins, water tables, etc... where does the water pool when it rains?



https://en.wikipedia.org/wiki/Voronoi\_diagram

- Hierarchical clustering:
  - -Agglomerative ("bottom up"):
    - Start with individual constituents, merge until criteria met
  - -Divisive ("top down"):
    - Start with conglomerate, split until criteria met (or you get to individual constituents)
  - –Represent by a tree ("dendrogram") or Venn diagram ("nested cluster diagram"):





(a) Dendrogram. (b) Nested cluster diagram.

Basically the same, but in reverse

Algorithm 8.3 Basic agglomerative hierarchical clustering algorithm.

- 1: Compute the proximity matrix, if necessary.  $O(n^2)$ , done once
- 2: repeat
- 3: Merge the closest two clusters. O(1), done n times
- 4: Update the proximity matrix to reflect the proximity between the new cluster and the original clusters.  $O(n^2)$ , done n times
- 5: **until** Only one cluster remains.



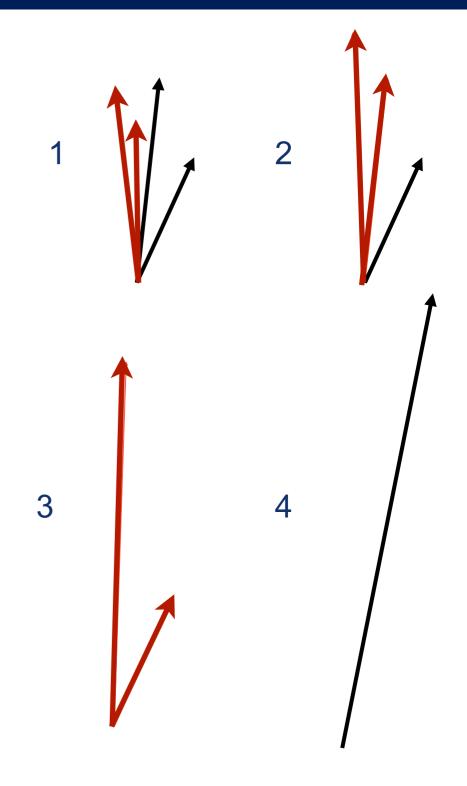
- Faster clustering algorithms (i.e. fastjet):
  - Can precompute nearest neighbors in the metric, and then only look at those instead of all (nearest is nearest is nearest!)

Calculate nearest neighbors while inputs are left:

Compute distances to neighbors, and self

If distance to neighbor is smallest, merge + iterate

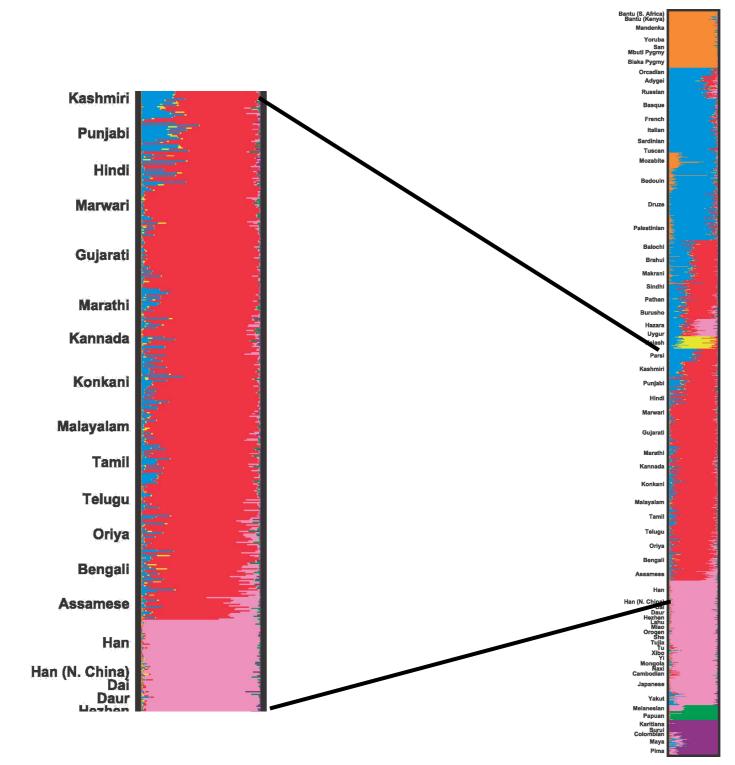
Else if distance to self is smallest, stop



https://arxiv.org/pdf/hep-ph/0512210.pdf

- This reduces to O(n<sup>2</sup>) complexity
- Can reduce further!
  - -To find nearest neighbors, can use Voronoi diagrams (like we had before)
  - -This reduces complexity to O(n ln(n))!

- Example: Genetic clustering
  - -Including those "who are your ancestors" DNA kits!



- Example: Finding galaxy clusters using Voronoi tessellations
  - -https://www.aanda.org/articles/aa/pdf/2001/12/ aa10522.pdf

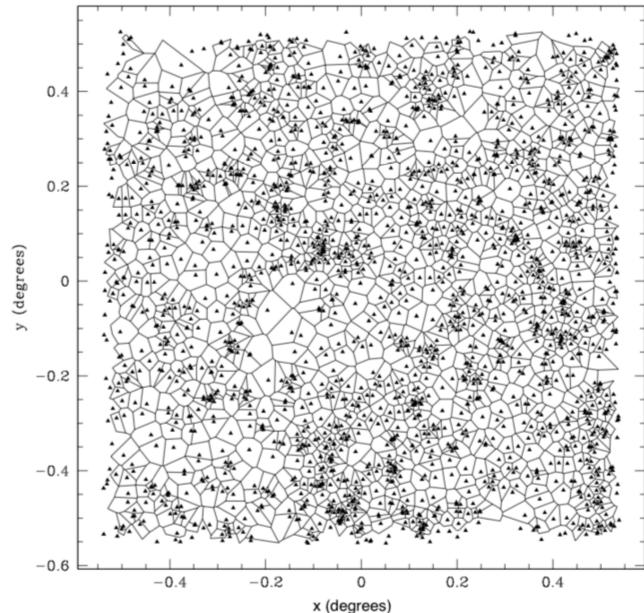


Fig. 1. Voronoi tessellation of a galaxy field

- Example: Finding hadronic jets (The Anti-kT Jet Clustering Algorithm)
- <u>http://inspirehep.net/record/779080</u>
- Hierarchical clustering with various metrics

