

CS 260: Foundations of Data Science

Prof. Sara Mathieson

Fall 2023



HAVERFORD
COLLEGE

Admin

- Exam will be handed back on Tuesday
- Last candidate talk at **4:15pm TODAY**
 - Tea at 4pm
 - Student lunch Friday 12:30-1:30pm

Outline for November 30

- Gaussian Mixture Models (GMMs)
- Kernel Density Estimation (KDE)
- Missing data
- Begin: neural networks

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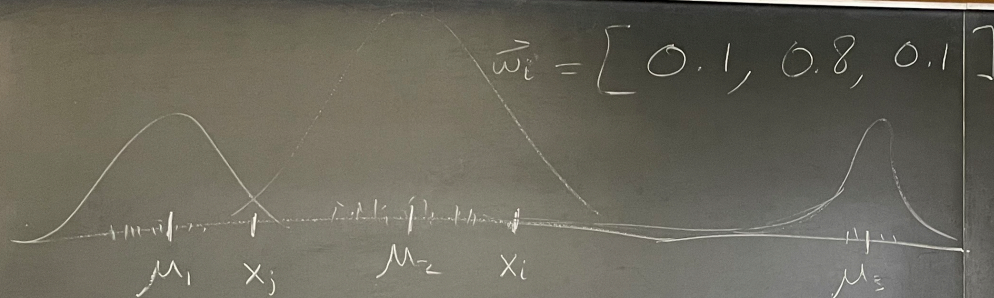
Gaussian Mixture Models | E-step "soft" assignment

- π_k = prob of class k
= $\frac{1}{K}$ for all k
 - $\vec{\mu}_k$ = mean of cluster k
= random points
 - σ_k^2 = variance of cluster k
= sample variance of pts closest to each mean.
- initialization

w_{ik} = prob that \vec{x}_i came from cluster k

$$= p(k | \vec{x}_i) = \frac{p(k) p(\vec{x}_i | k)}{p(\vec{x}_i)}$$

$$= \frac{\pi_k N(\vec{x}_i; \vec{\mu}_k, \sigma_k^2)}{\sum_{j=1}^K \pi_j N(\vec{x}_i; \vec{\mu}_j, \sigma_j^2)}$$



$$\vec{w}_j = [0.49, 0.49, 0.02]$$

$$W = \begin{bmatrix} \mu_1 & \mu_2 & \mu_3 \\ 0.1 & 0.8 & 0.1 \\ 0.49 & 0.49 & 0.02 \end{bmatrix} \begin{matrix} \rightarrow 1 \\ \rightarrow 1 \end{matrix}$$

$n \times K$

M-step param update

$$M_k = \sum_{i=1}^n w_{ik} \quad \left. \vphantom{\sum_{i=1}^n} \right\} \begin{matrix} \# \text{ pts} \\ \text{assigned to} \\ \text{cluster } k \end{matrix}$$

$$\textcircled{1} \pi_k = \frac{M_k}{n}$$

$$\textcircled{2} \vec{\mu}_k = \frac{1}{M_k} \sum_{i=1}^n w_{ik} \vec{x}_i$$

$$\textcircled{3} \sigma_k^2 = \text{weighted sample variance.}$$

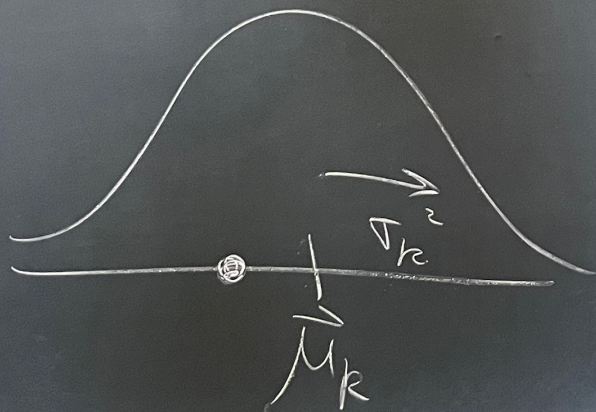
K-means

$$\frac{1}{|C_k|} \sum_{\vec{x}_i \in C_k} \vec{x}_i$$

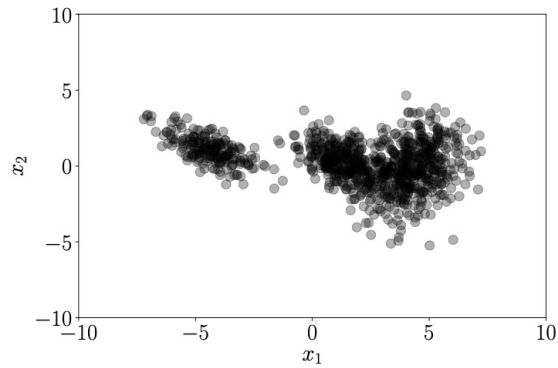
Generative process

① choose a cluster k
using $[\pi_1, \pi_2, \dots, \pi_K]$

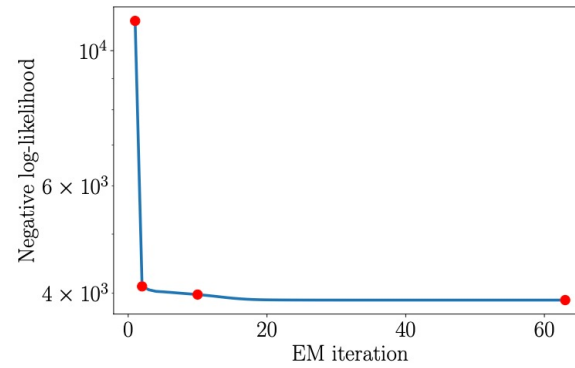
② use $\vec{\mu}_k + \sigma_k^2$ to sample
from $N(\vec{\mu}_k, \sigma_k^2)$



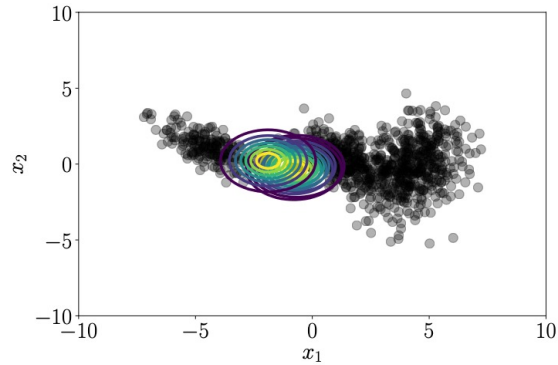
$$\frac{x_i}{\sqrt{\sum_{j=1}^n x_j^2}}$$



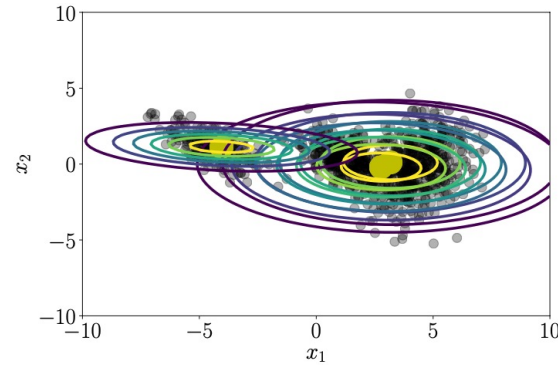
(a) Dataset.



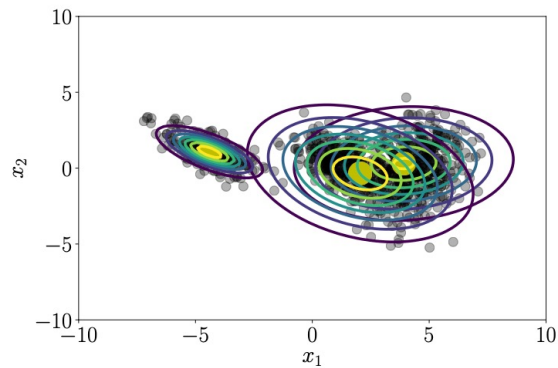
(b) Negative log-likelihood.



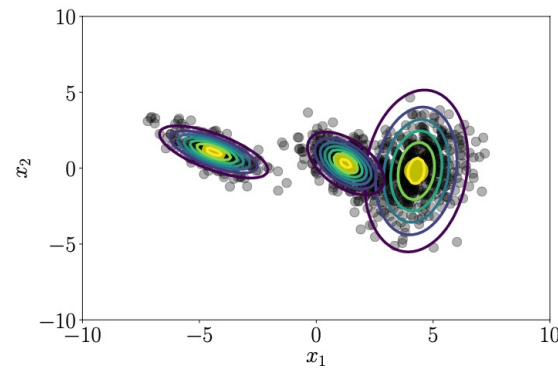
(c) EM initialization.



(d) EM after one iteration.



(e) EM after 10 iterations.

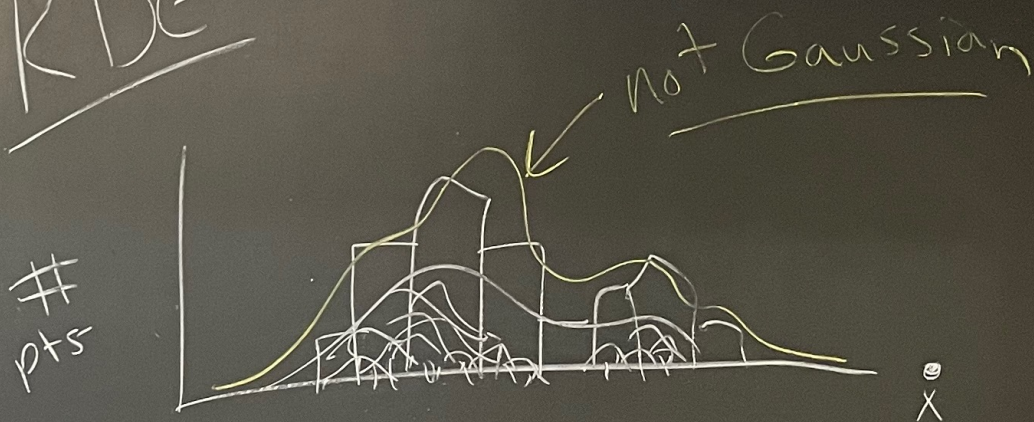


(f) EM after 62 iterations.

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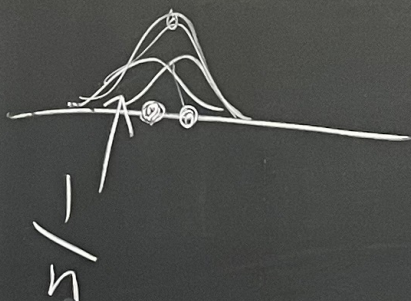
KDE



$$p(x) = \frac{1}{nh} \sum_{i=1}^n k\left(\frac{x - x_i}{h}\right)$$

kernel

width



KDE (Kernel Density Estimation)

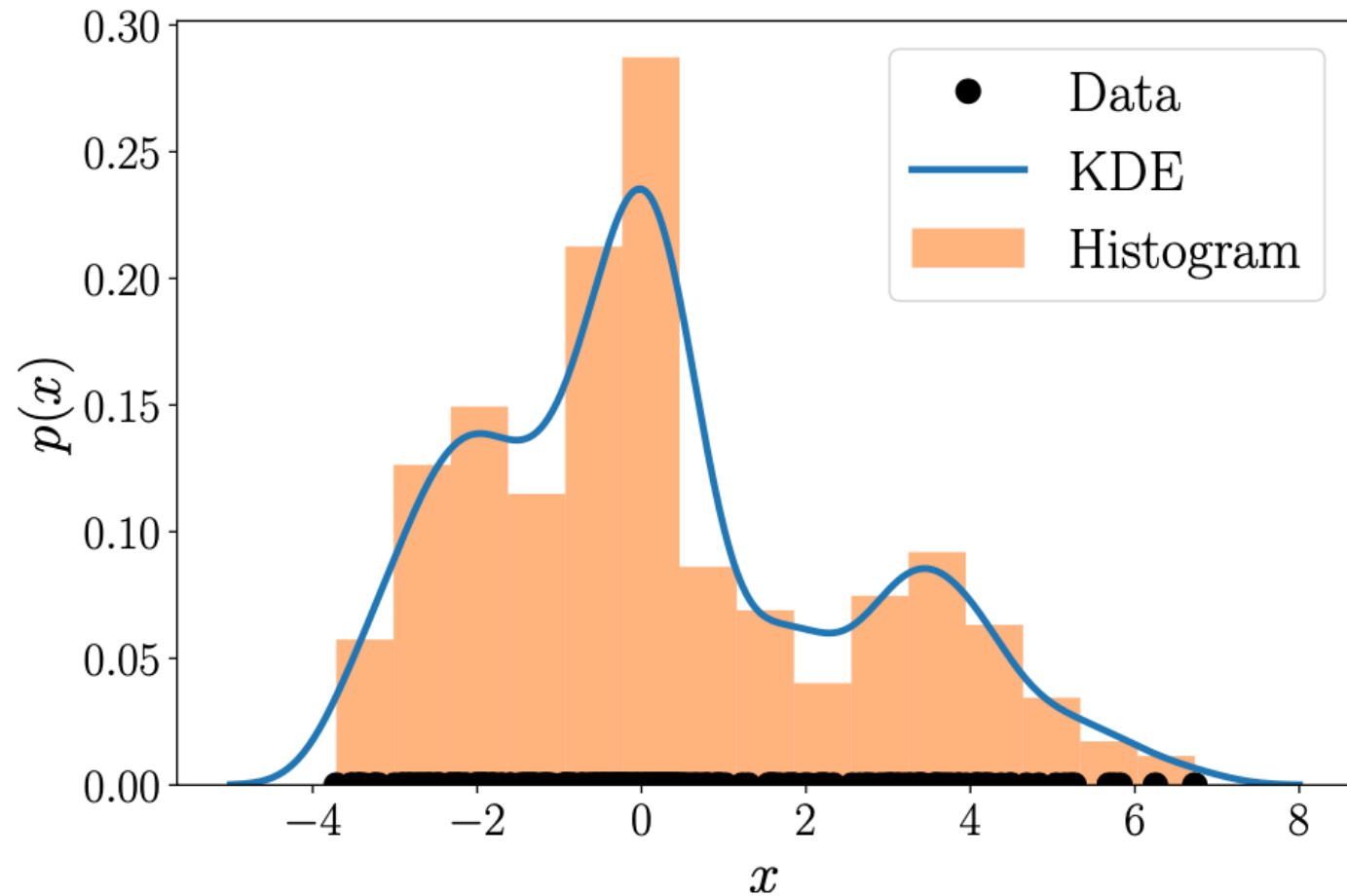


Figure 11.9 from MML textbook

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- **Missing data**
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Types of missing data

- MCAR: Missing Completely At Random. Not related to:
 - Specific values
 - Observed responses
- MAR: Missing At Random. Not related to:
 - Specific values
- MNAR: Missing Not At Random

Techniques for handling missing data

- Try to prevent the problem in the first place
 - Careful study design, follow-up with participants, etc
- Omit rows with missing data (reduces n)
- Omit only when value is needed
 - i.e. Naïve Bayes, per-feature estimates
- Mean substitution (per feature)

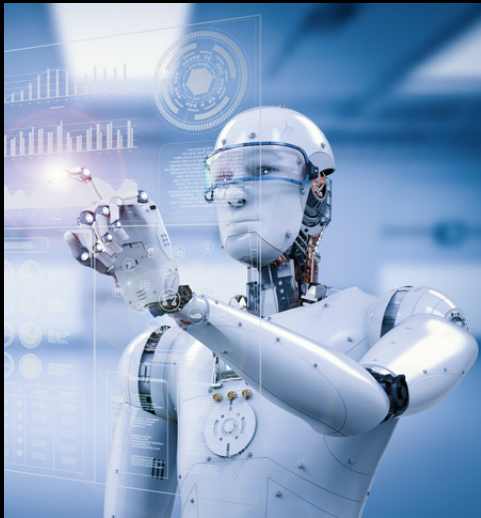
Techniques for handling missing data

- Imputation
 - Use similar examples to guess the missing values
 - Can be done locally or globally
- Last observation carried forward
 - Useful for time-series data

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MACHINE LEARNING



What society thinks I do

$$\nabla_w \mathcal{L}(w, b, \alpha) = w - \sum_{i=1}^m \alpha_i y^{(i)} x^{(i)} = 0$$

This implies that

$$w = \sum_{i=1}^m \alpha_i y^{(i)} x^{(i)}.$$

As for the derivative with respect to b , we obtain

$$\frac{\partial}{\partial b} \mathcal{L}(w, b, \alpha) = \sum_{i=1}^m \alpha_i y^{(i)} = 0.$$

If we take the definition of w in Equation (9) and plug that back into the Lagrangian (Equation 8), and simplify, we get

$$\mathcal{L}(w, b, \alpha) = \sum_{i=1}^m \alpha_i - \frac{1}{2} \sum_{i,j=1}^m y^{(i)} y^{(j)} \alpha_i \alpha_j (x^{(i)})^T x^{(j)} - b \sum_{i=1}^m \alpha_i y^{(i)}.$$

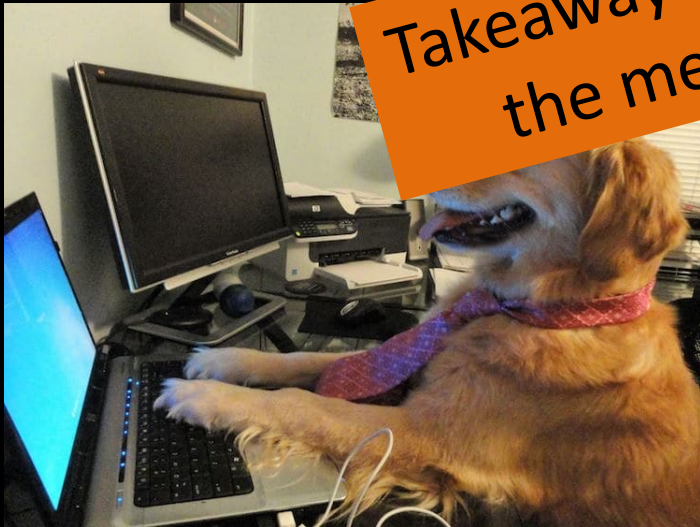
But from Equation (10), the last term must be zero, so we obtain

$$\mathcal{L}(w, b, \alpha) = \sum_{i=1}^m \alpha_i - \frac{1}{2} \sum_{i,j=1}^m y^{(i)} y^{(j)} \alpha_i \alpha_j (x^{(i)})^T x^{(j)}.$$

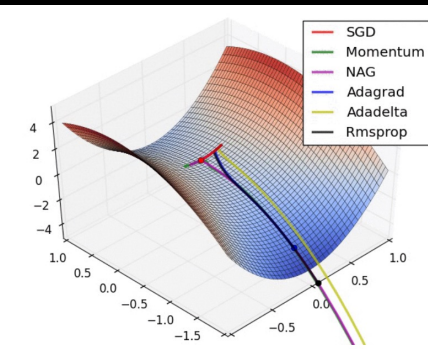
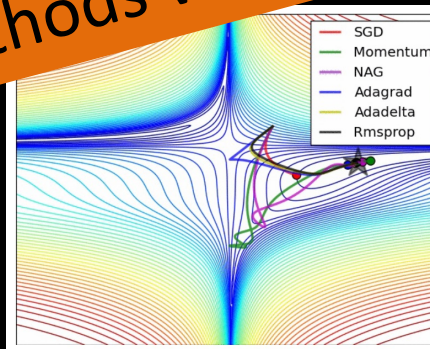


What other computer scientists think I do

Takeaway: we should understand the methods we are using!



What mathematicians think I do

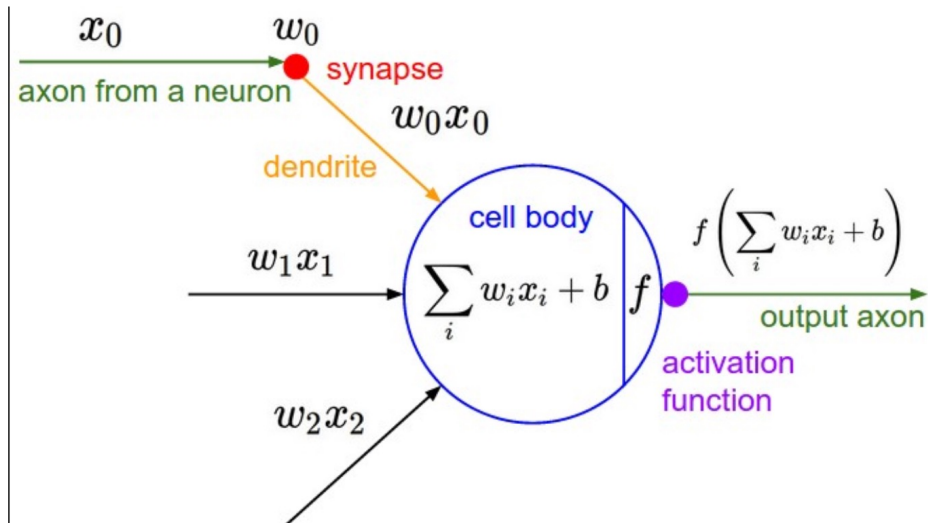
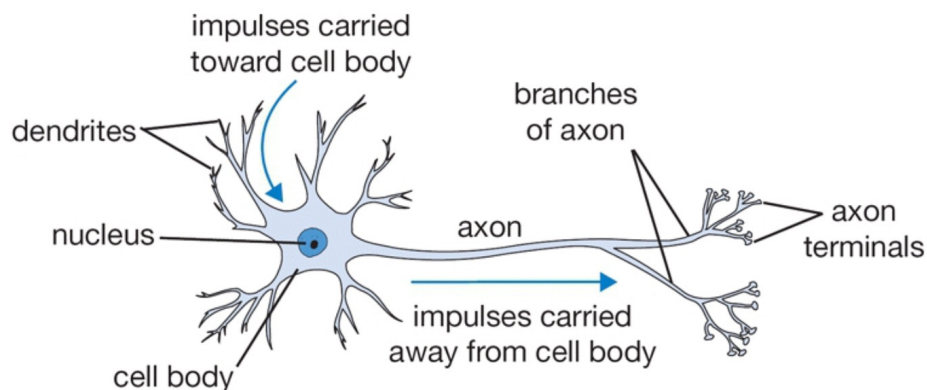


What I think I do

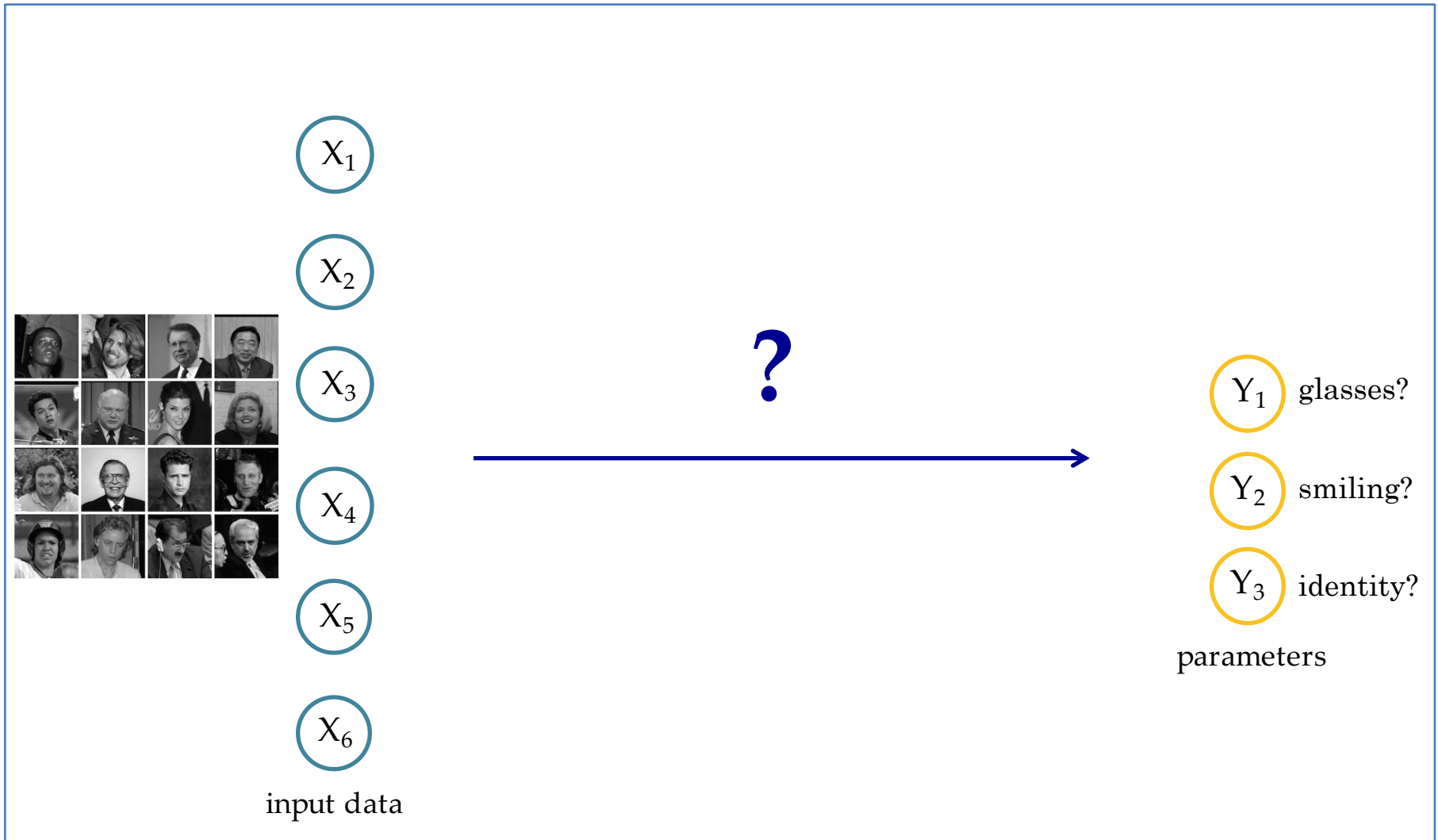
```
>>> from sklearn import svm
>>> import tensorflow as tf
```

What I really do

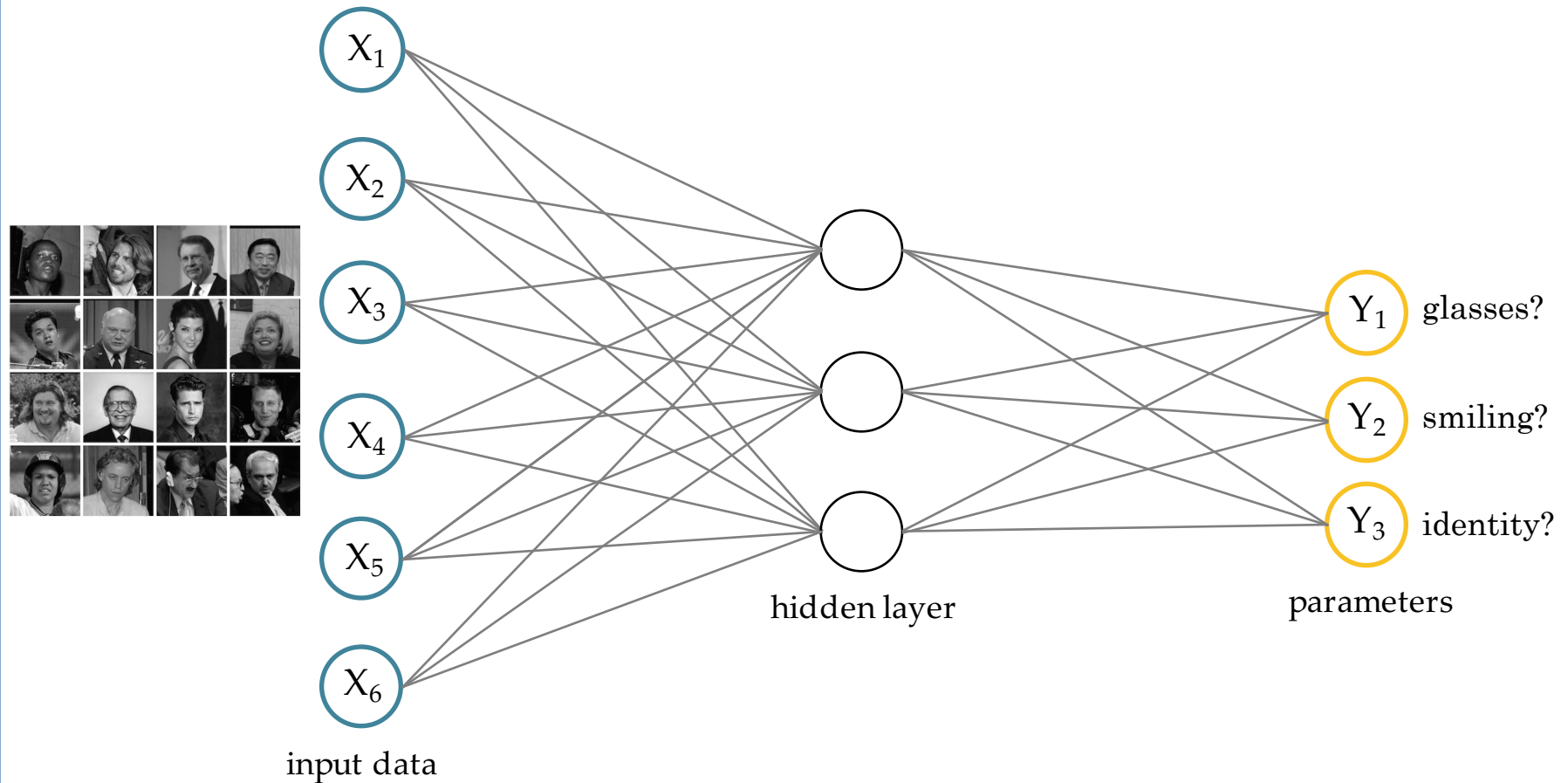
Biological Inspiration



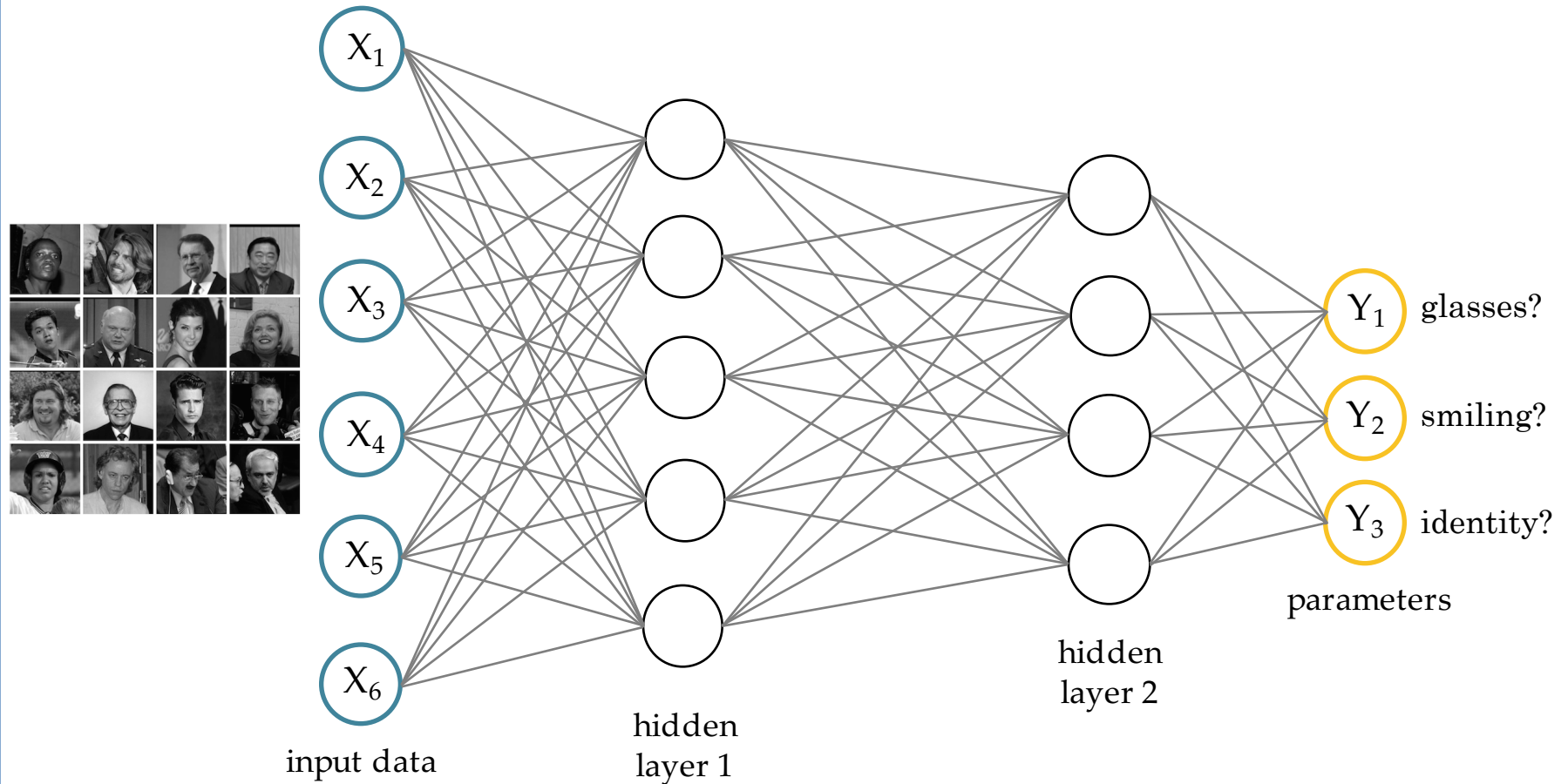
Goal: learn from complicated inputs



Idea: transform data into lower dimension



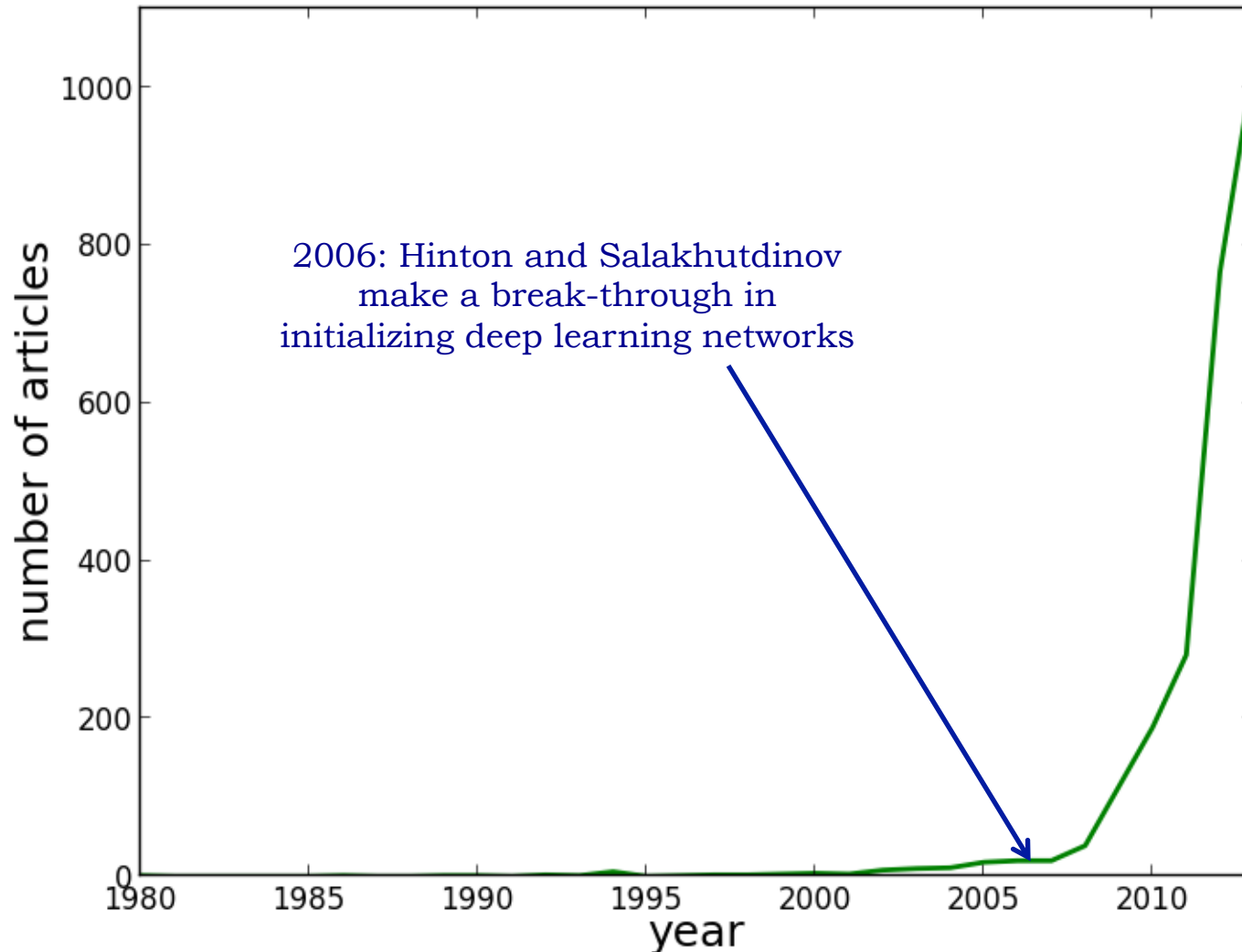
Multi-layer networks = “deep learning”



History of Neural Networks

- Perceptron can be interpreted as a simple neural network
- Misconceptions about the weaknesses of perceptrons contributed to declining funding for NN research
- Difficulty of training multi-layer NNs contributed to second setback
- Mid 2000's: breakthroughs in NN training contribute to rise of “deep learning”

Number of papers that mention “deep learning” over time



Big picture for today

- Neural networks can approximate any function!

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- For our purposes in ML, we want to use them to approximate a function from our inputs to our outputs

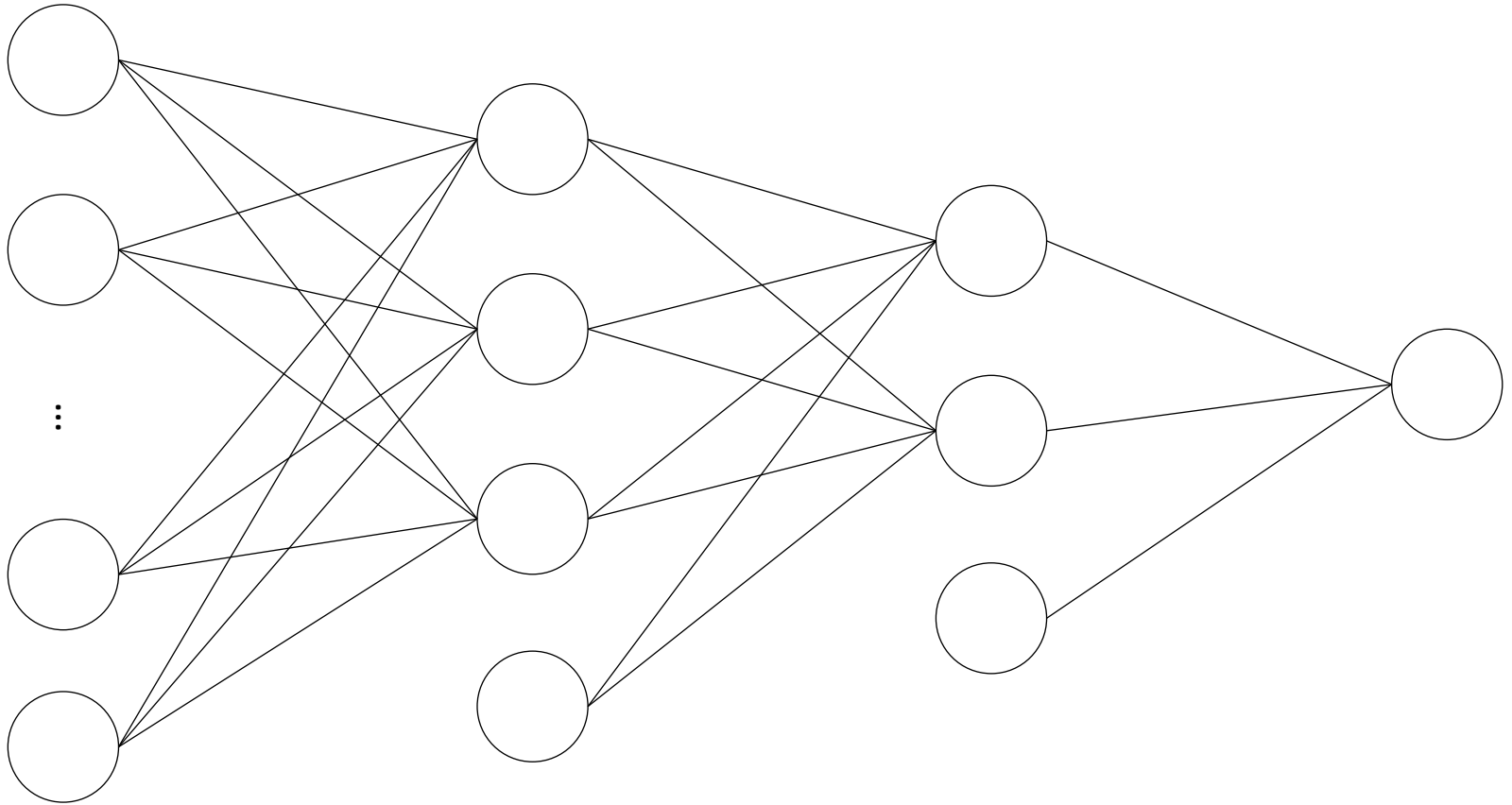
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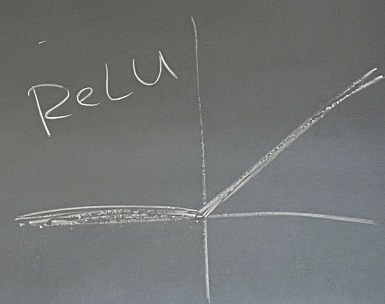
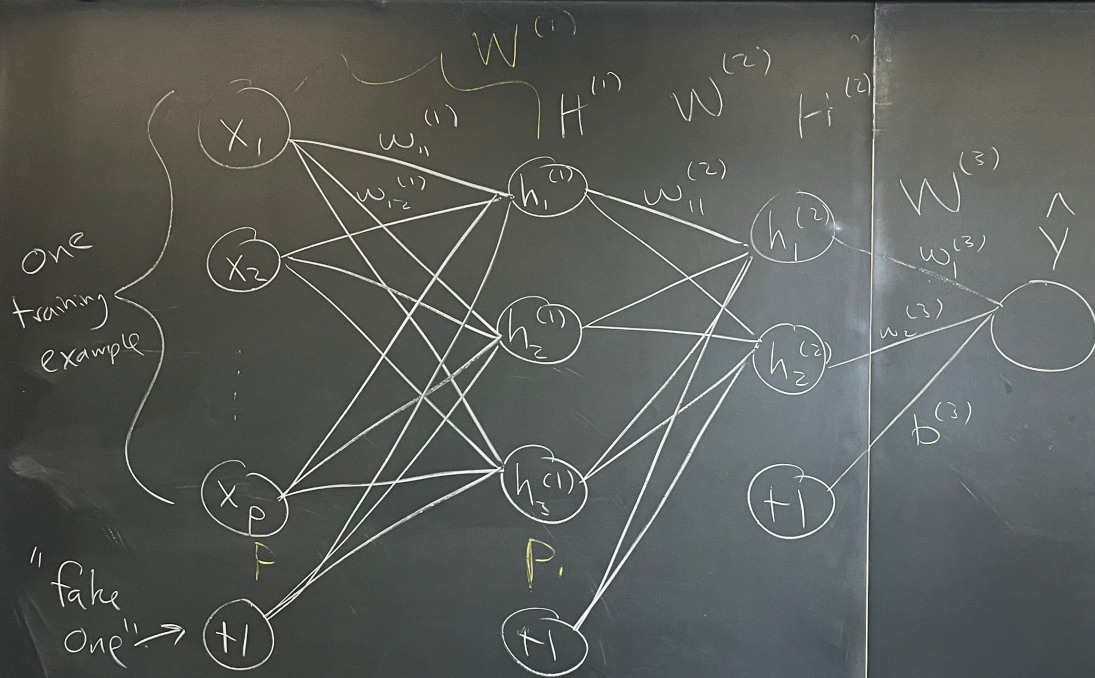
- Neural networks can approximate any function!
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- We will train our network by asking it to minimize the loss between its output and the true output

Big picture for today

- Neural networks can approximate any function!
- For our purposes in ML, we want to use them to approximate a function from our inputs to our outputs
- We will train our network by asking it to minimize the loss between its output and the true output
- We will use SGD-like approaches to minimize loss

Fully Connected Neural Network Architecture





Matrix a :

$$a = \begin{bmatrix} -1 & 2 & 5 \\ 0 & -2 & 1 \\ -3 & -4 & 8 \end{bmatrix}$$

Matrix a (repeated):

$$\begin{bmatrix} 0 & 0 & 2 & 5 \\ 0 & 0 & -2 & 1 \\ 0 & 0 & -3 & 8 \end{bmatrix}$$

$$H^{(1)} = a \left(W^{(1)} X + \vec{b}^{(1)} \right)$$

activation function

 $\underbrace{p \times p \quad p \times n}_{p \times n}$

 $\underbrace{\quad}_{p \times 1}$

$$\begin{bmatrix} | & | & | \\ x_1 & x_2 & \dots \\ | & | & | \end{bmatrix}_{p \times n}$$

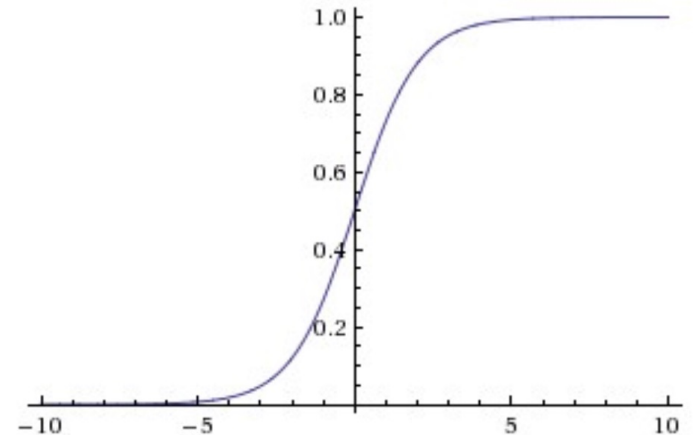
$$H^{(2)} = a \left(W^{(2)} H^{(1)} + \vec{b}^{(2)} \right)$$

$$\hat{Y} = a \left(W^{(3)} H^{(2)} + \vec{b}^{(3)} \right)$$

Option 1: sigmoid function

- Input: all real numbers, output: $[0, 1]$

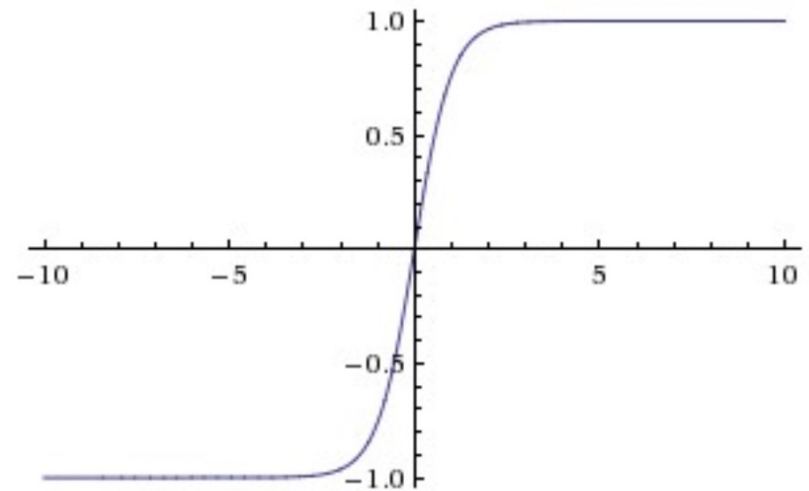
$$\sigma(x) = \frac{1}{1 + e^{-x}}$$



Option 2: hyperbolic tangent

- Input: all real numbers, output: $[-1, 1]$

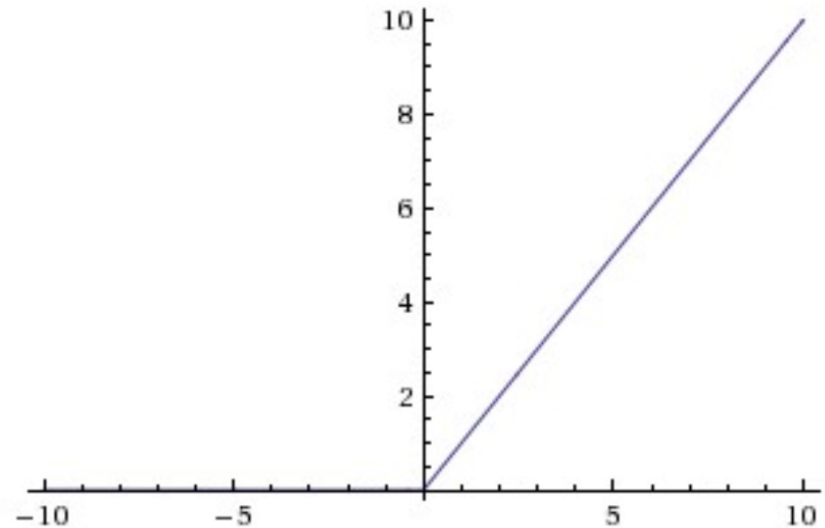
$$\tanh(x) = \frac{e^x - e^{-x}}{e^x + e^{-x}}$$



Option 3: Rectified Linear Unit (ReLU)

- Return x if x is positive (i.e. threshold at 0)

$$f(x) = \max(0, x)$$



Pros and Cons of Activation Functions

1) Sigmoid

- (-) When input becomes very positive or very negative, gradient approaches 0 (saturates and stops gradient descent)
- (-) Not zero-centered, so gradient on weights can end up all positive or all negative (zig-zag in gradient descent)
- (+) Derivative is easy to compute given function value!

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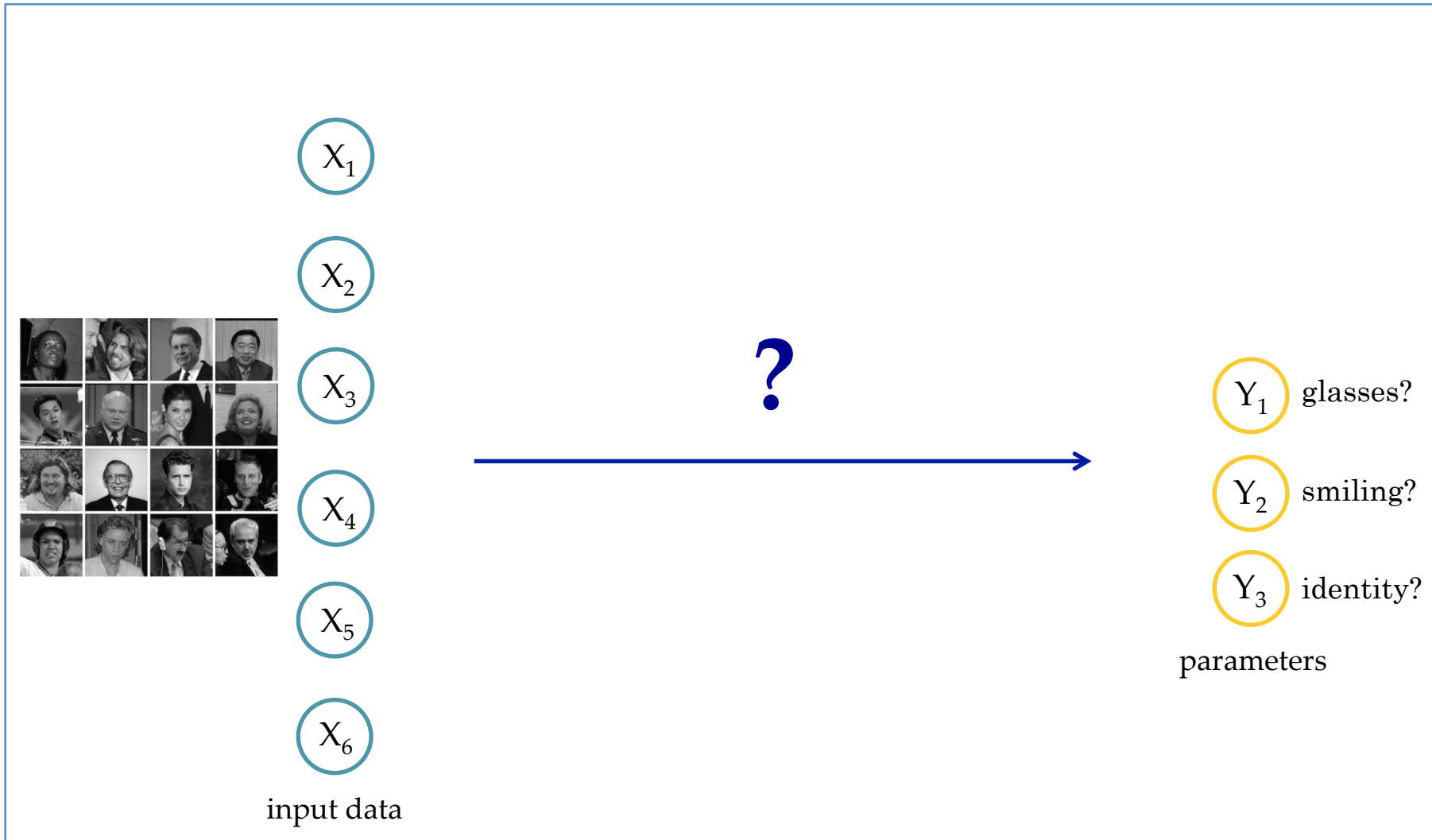
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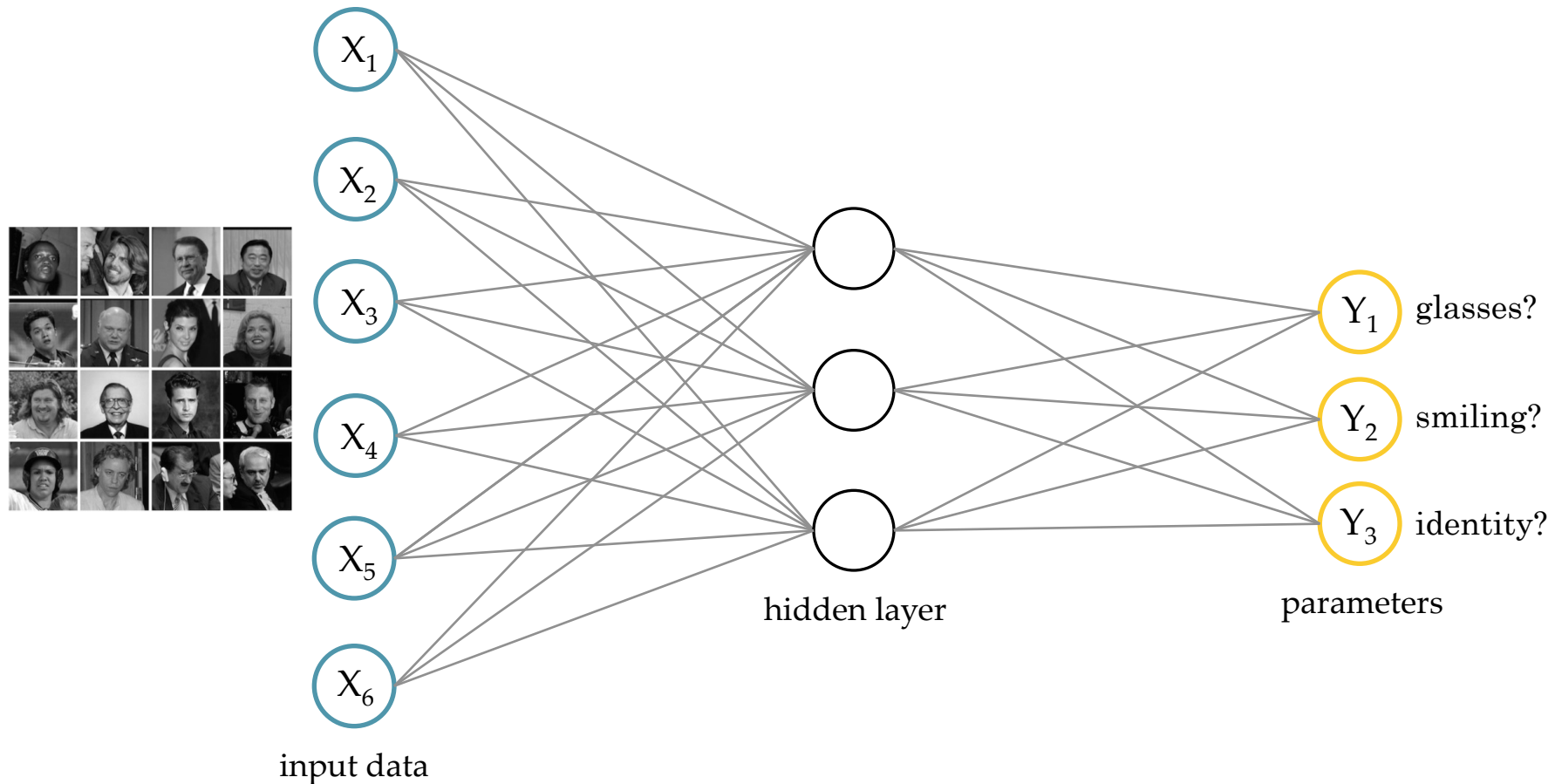
3) ReLU

- (+) Works well in practice (accelerates convergence)
- (+) Function value very easy to compute! (no exponentials)
- (-) Units can have no signal if input becomes too negative throughout gradient descent

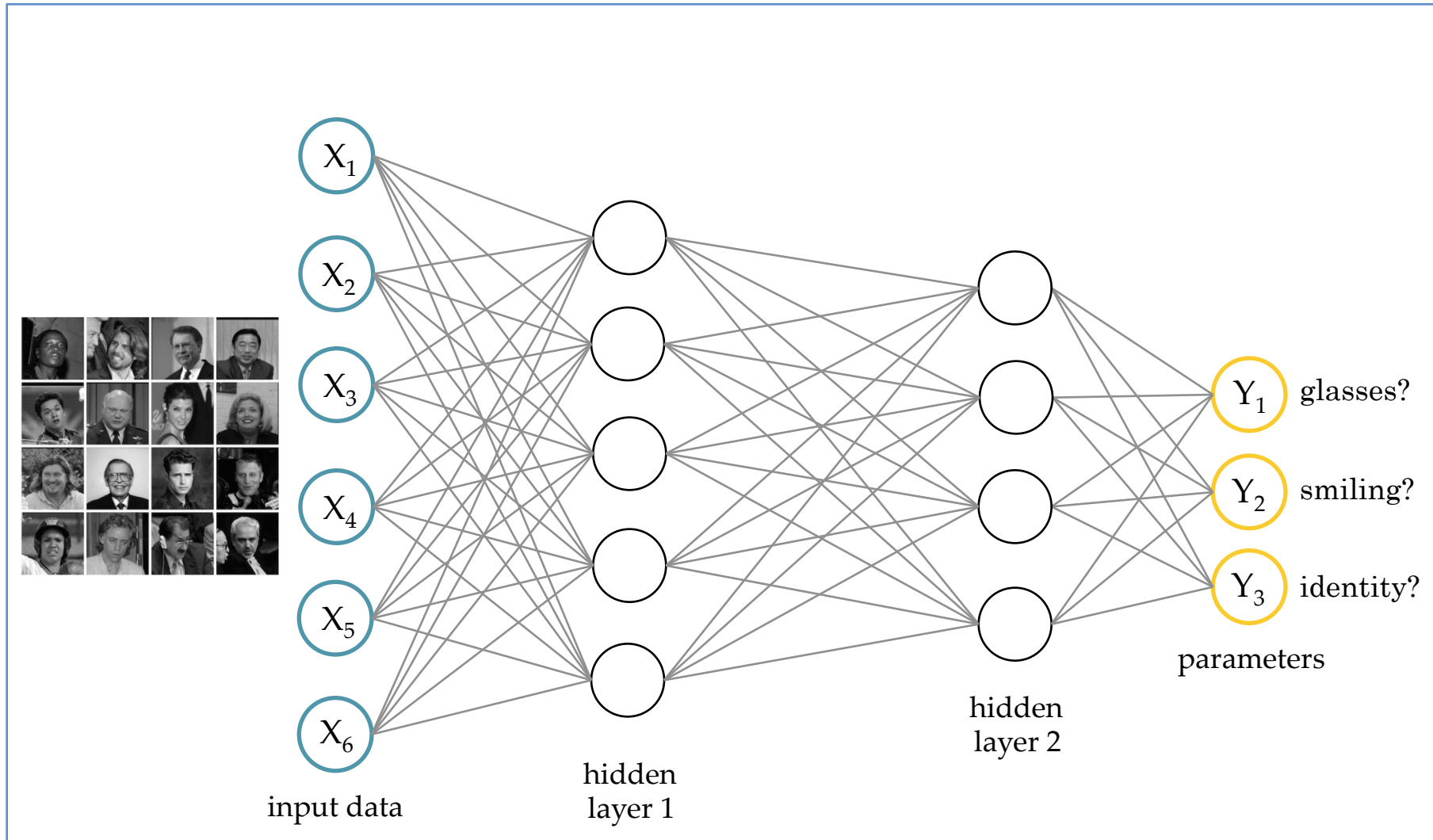
Goal: find a function between input and output



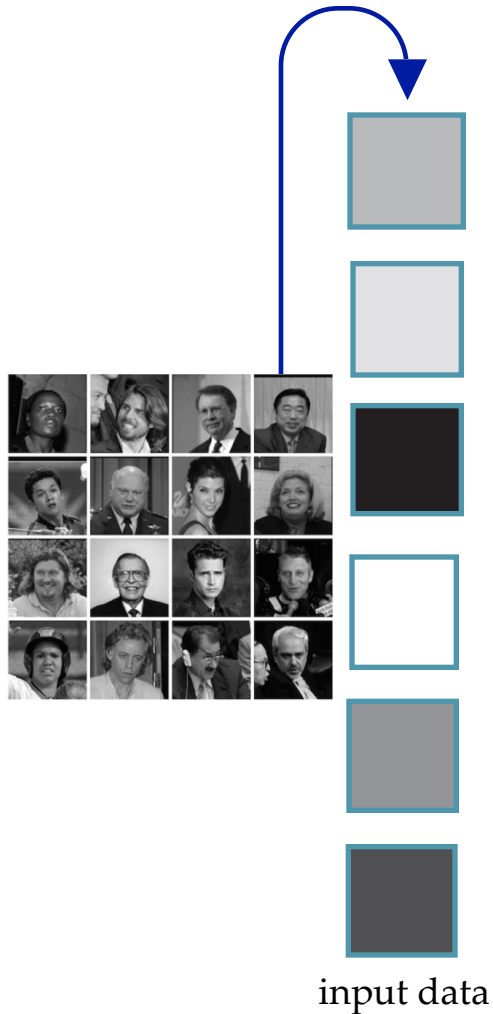
First idea: one hidden layer



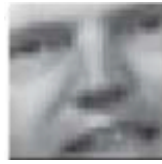
Second idea: more hidden layers (“deep” learning)



Flatten pixels of image into a single vector



Detour to autoencoders



x_1

x_2

x_3

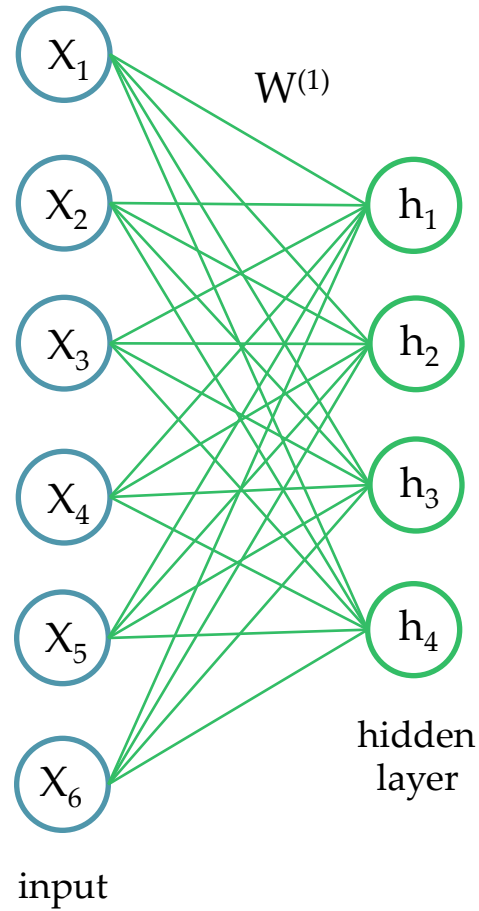
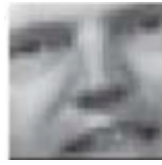
x_4

x_5

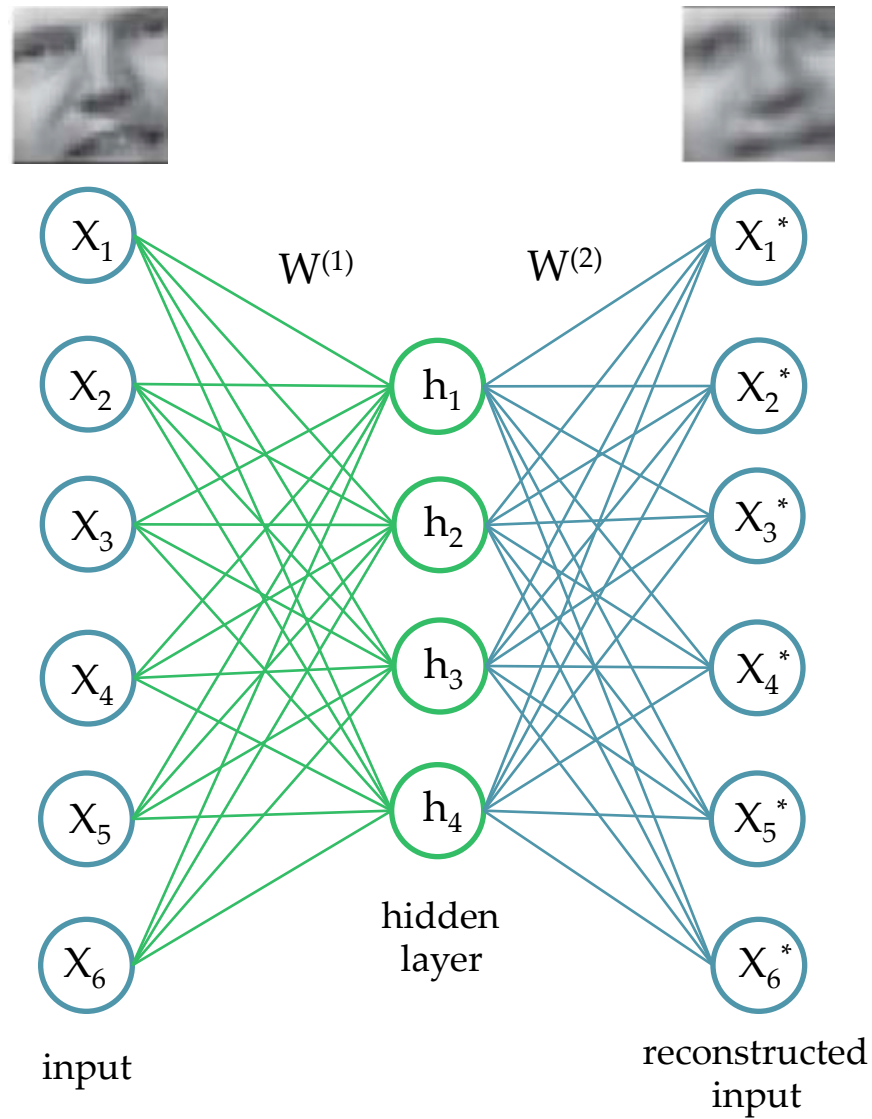
x_6

input

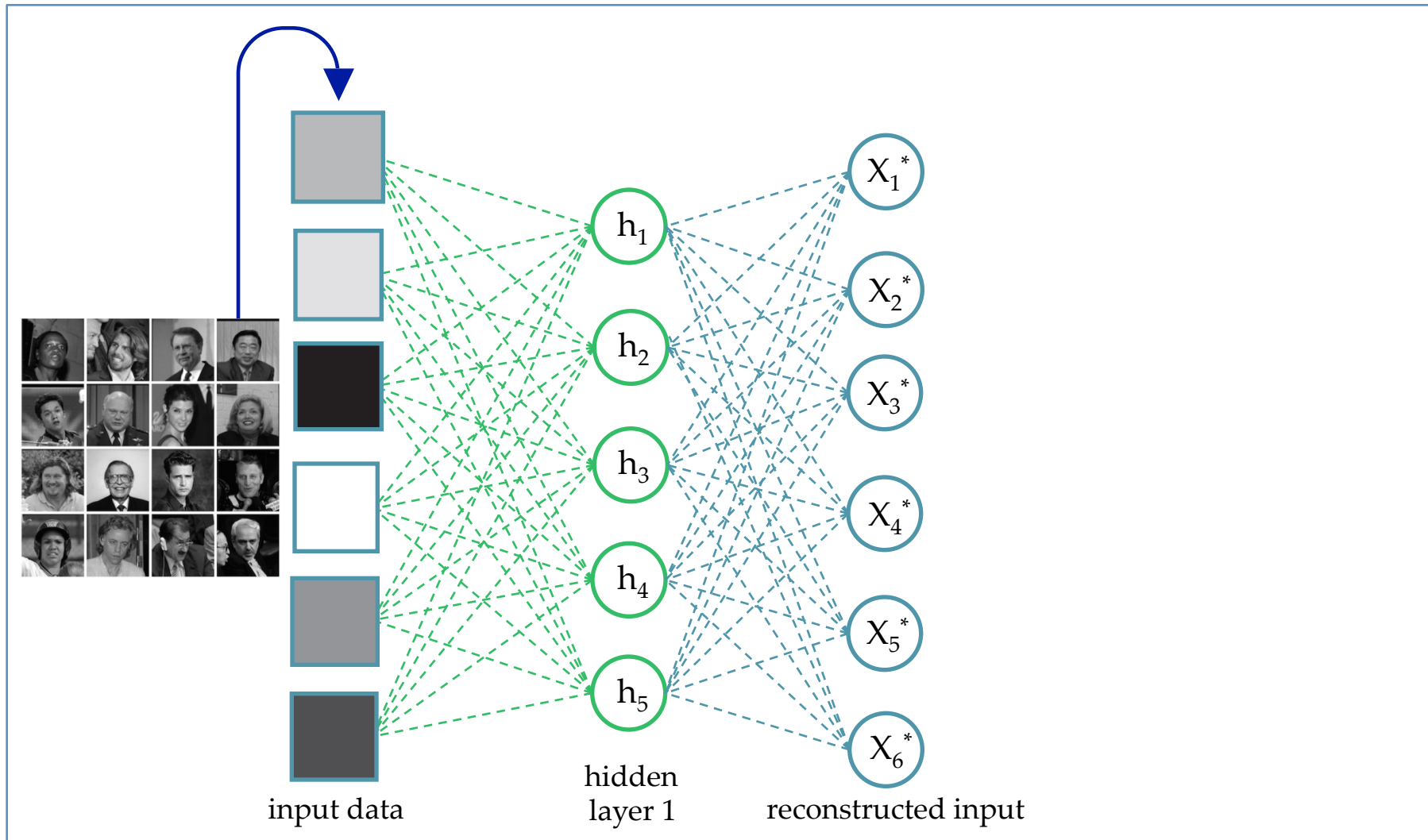
Detour to autoencoders



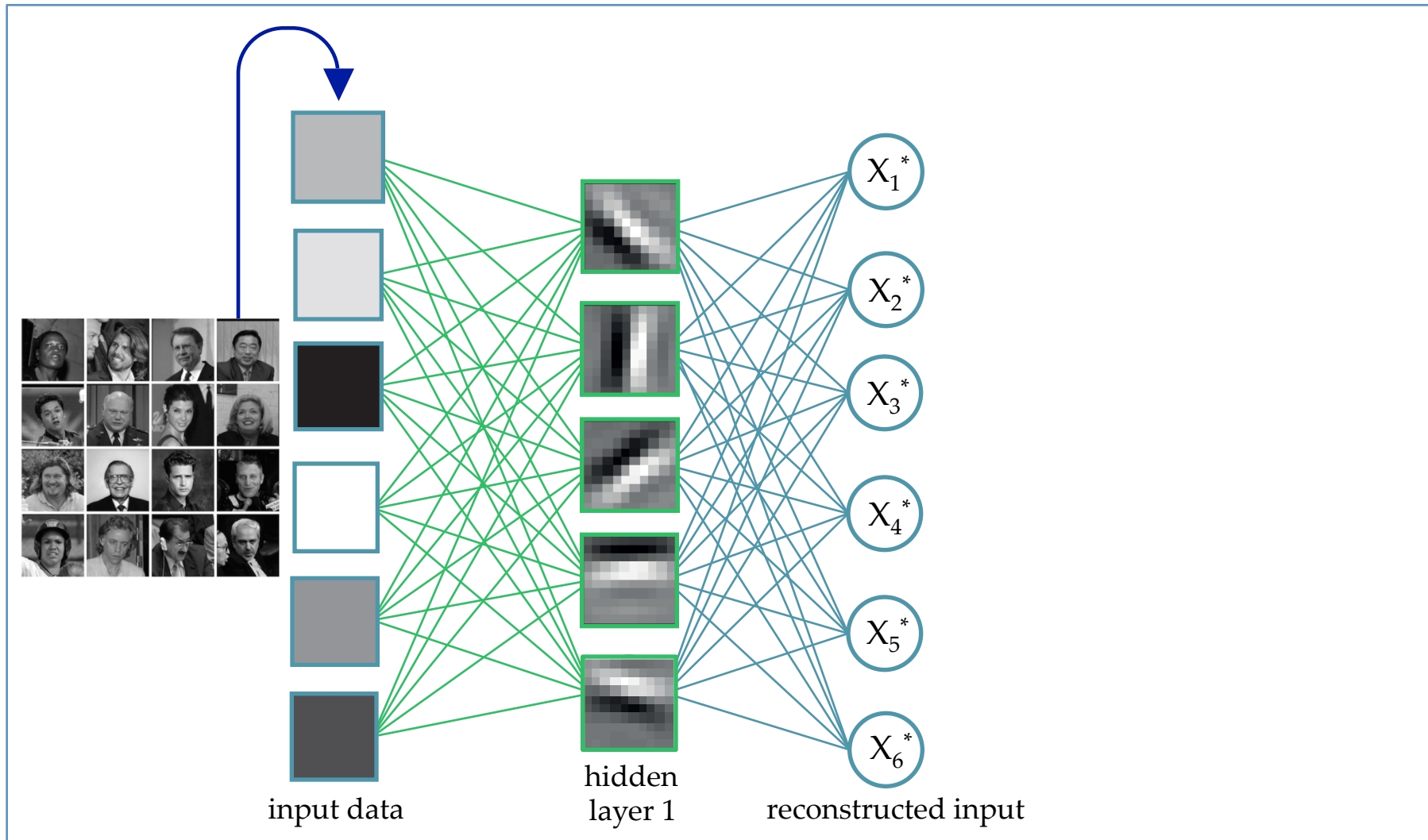
Detour to autoencoders



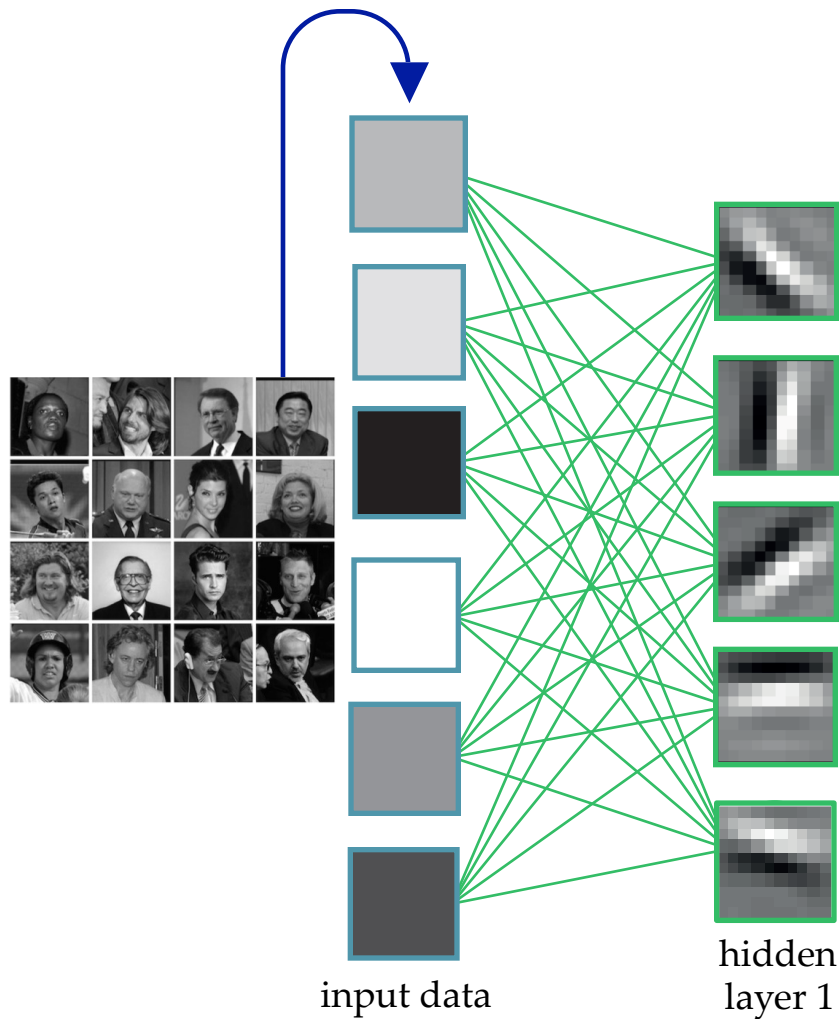
Use unsupervised pre-training to find a function from the input to itself



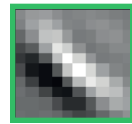
Hidden units can be interpreted as edges



Now: throw away reconstruction and input

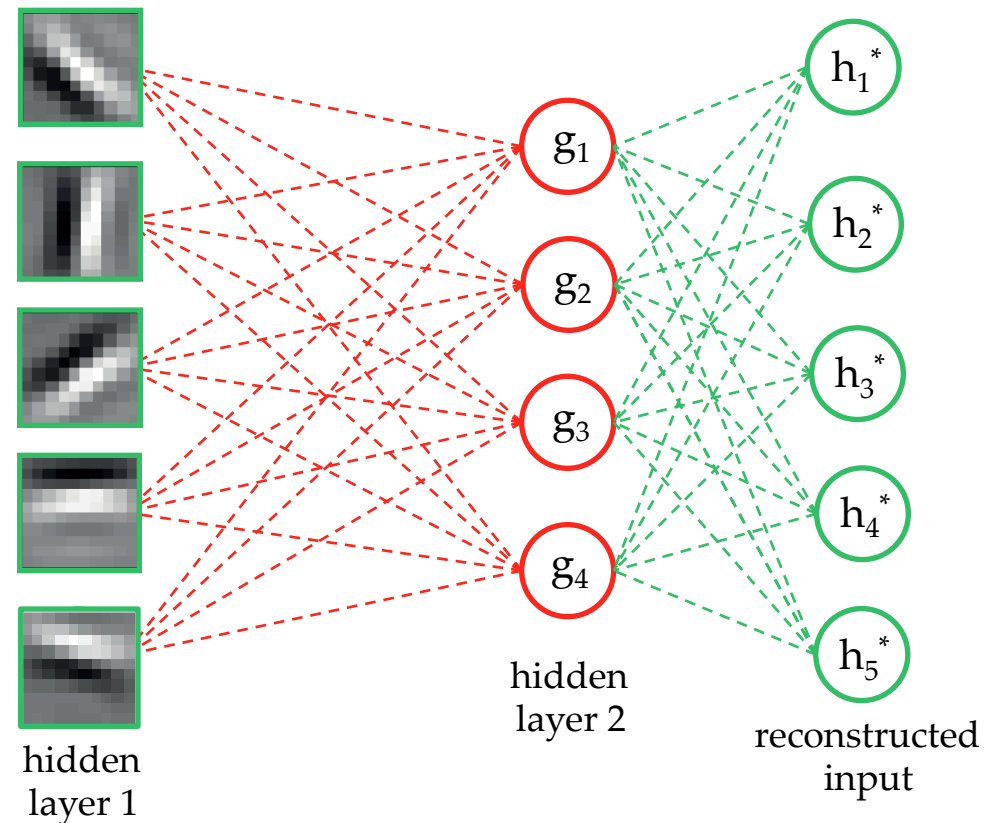


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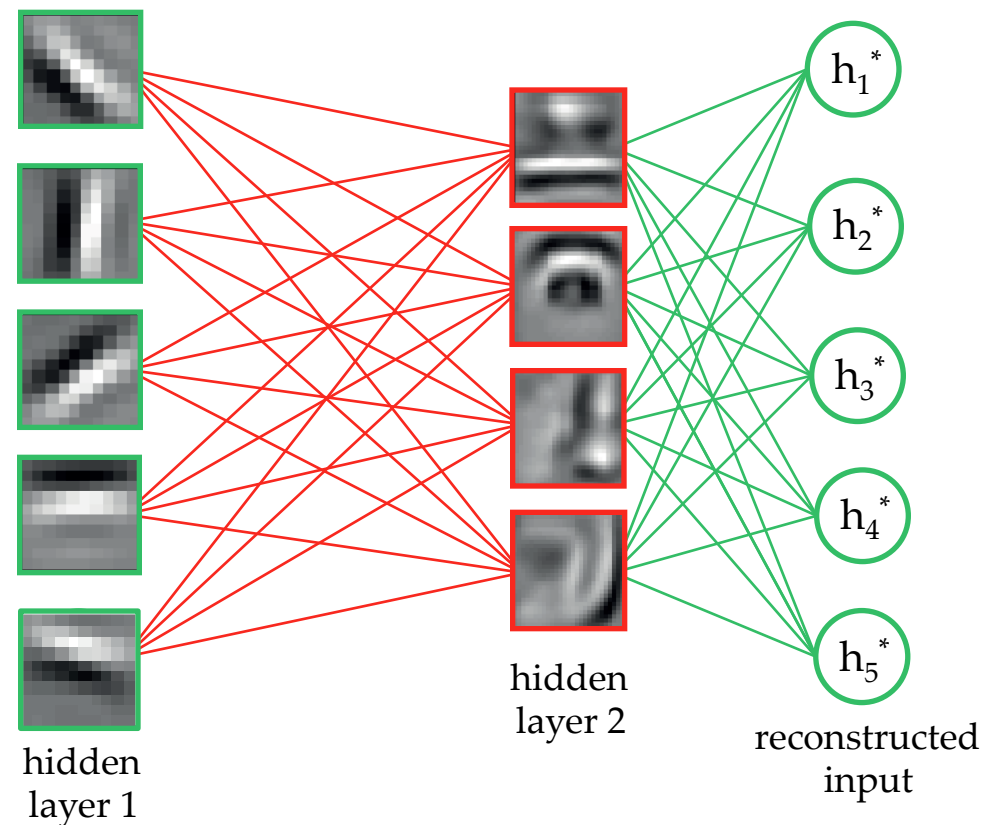


hidden
layer 1

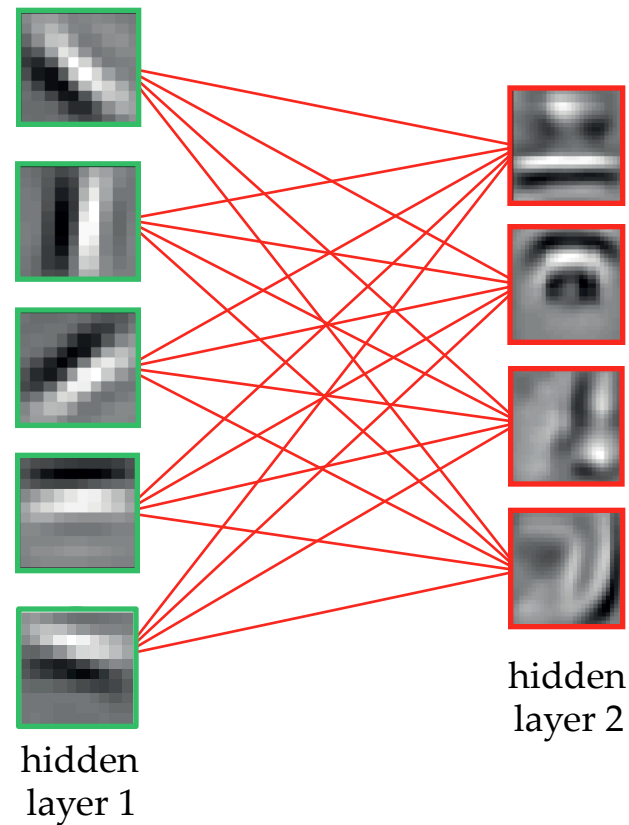
Then repeat the entire process for each layer



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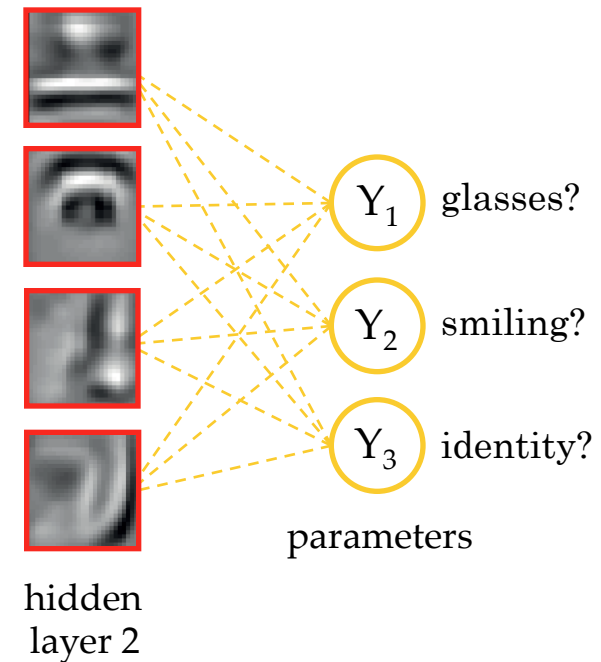


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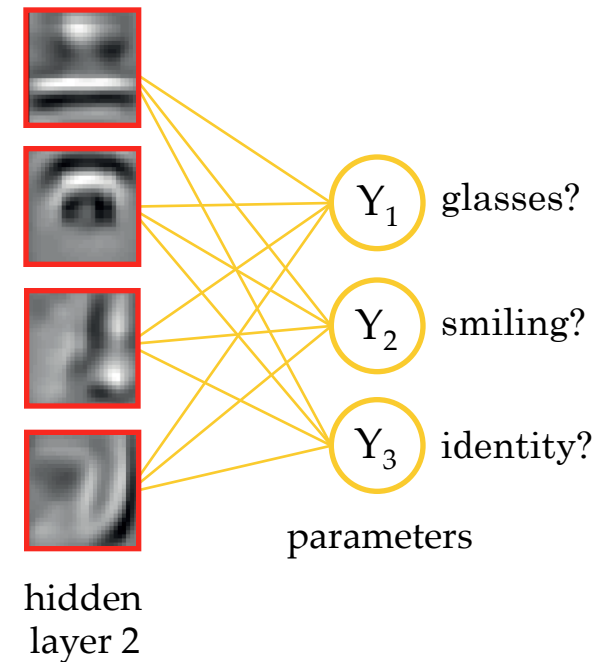


hidden
layer 2

In the last layer, use the outputs (supervised)



In the last layer, use the outputs (supervised)



Finally, “fine-tune” the entire network!

