

# Introduction of Deep Learning

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PLPTH813

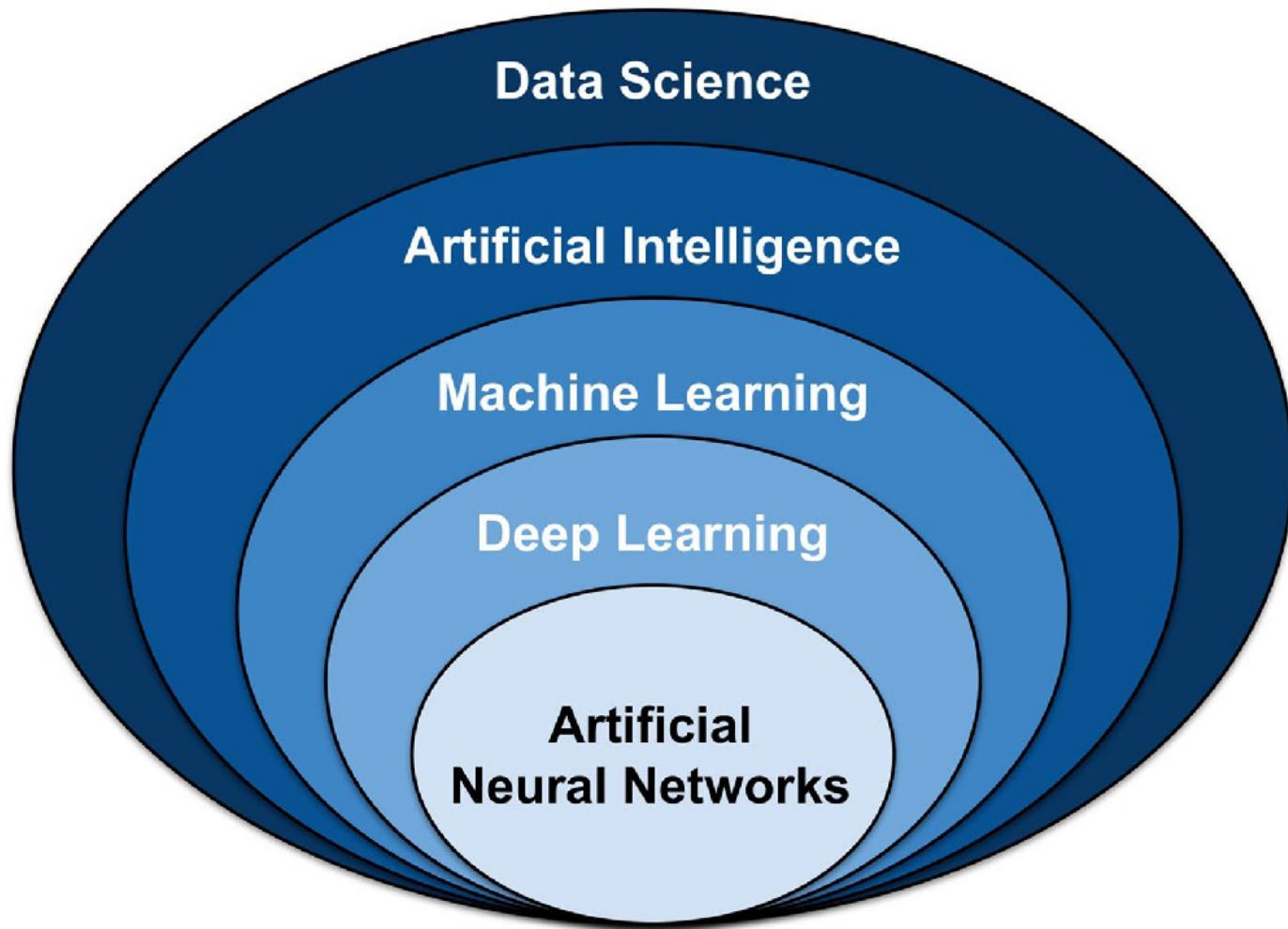
4/29/2021

# Goals

To learn:

- What is machine learning or deep learning?
- How does neural network work?
- What do we need to know for applying machine learning in our studies?
- How to run a simple machine learning using R? (lab)

# Machine learning



# Supervised and unsupervised (major methods)

- Unsupervised (to learn inherent patterns within data)  
e.g., Clustering, principal component analysis
- Supervised: to predict response (regression) or classification of each data point by using a provided set of labeled training examples

# Supervised: regression and classification

A **regression** model predicts continuous values.

- What is the value of a house in California?
- What is the probability that a user will click on this ad?

A **classification** model predicts discrete values.

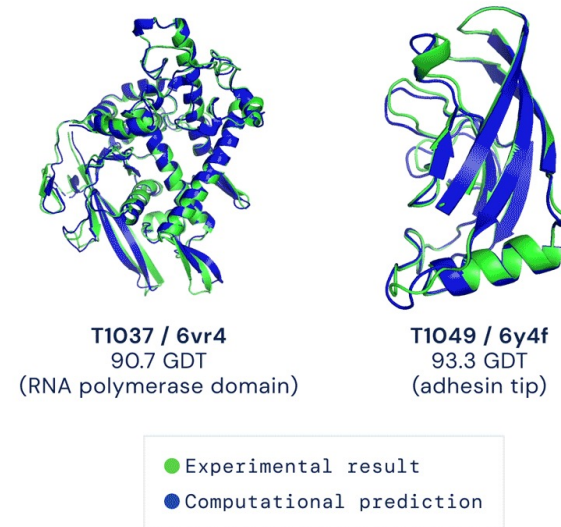
- Is a given email message spam or not spam?
- Is this an image of a dog, a cat, or a hamster?

# Why machine learning

- Advance of algorithms and theories
- Increasing computational power
- Large data

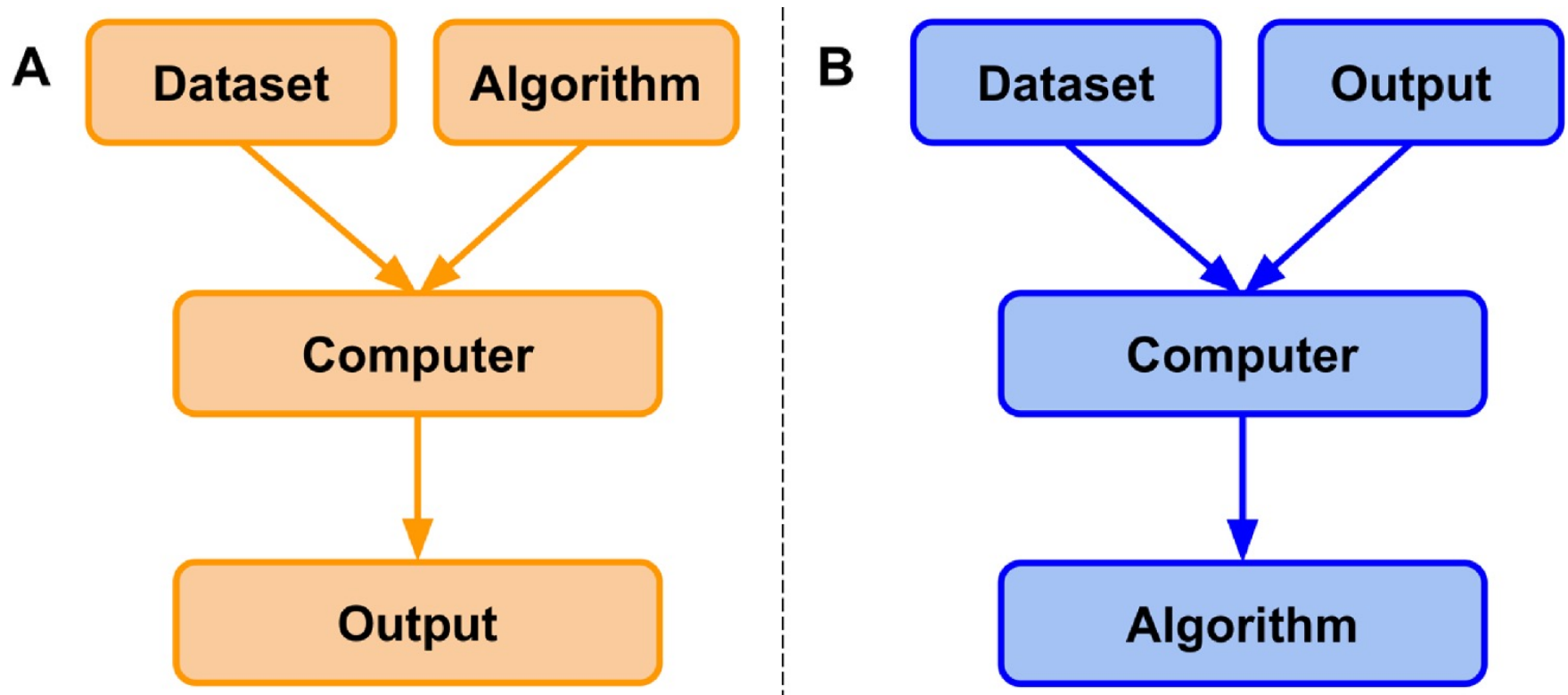


autopilot

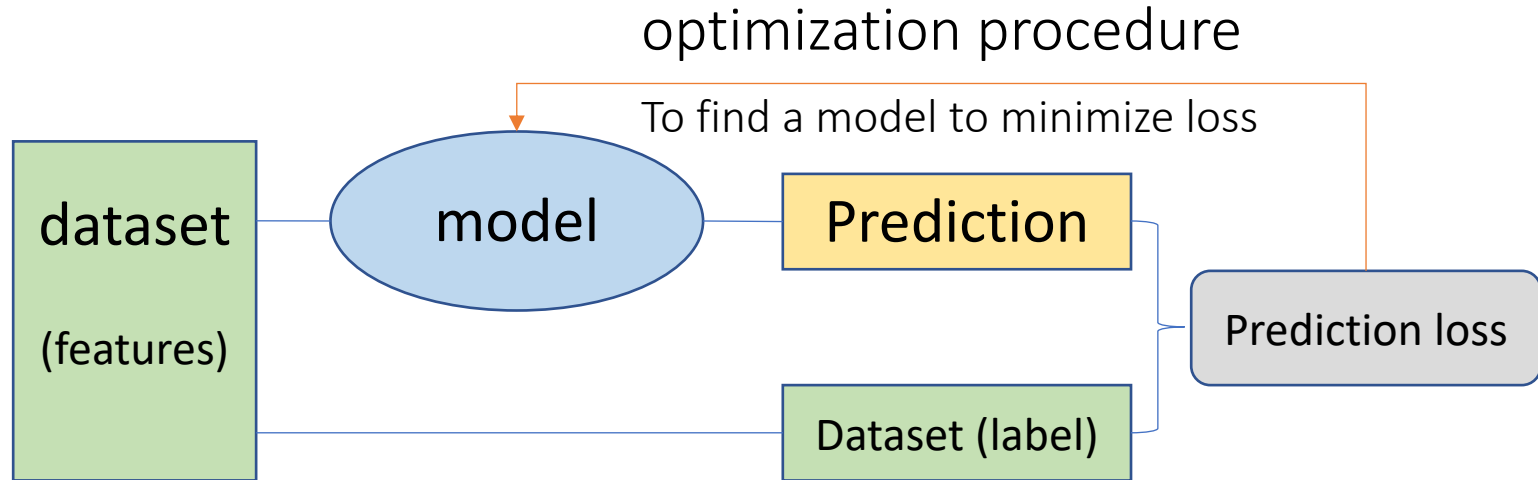


AlphaFold

# Classical programming vs. machine learning paradigm



# General procedure



location, room, area, ...

house price

edu, org, spam\_words, ...

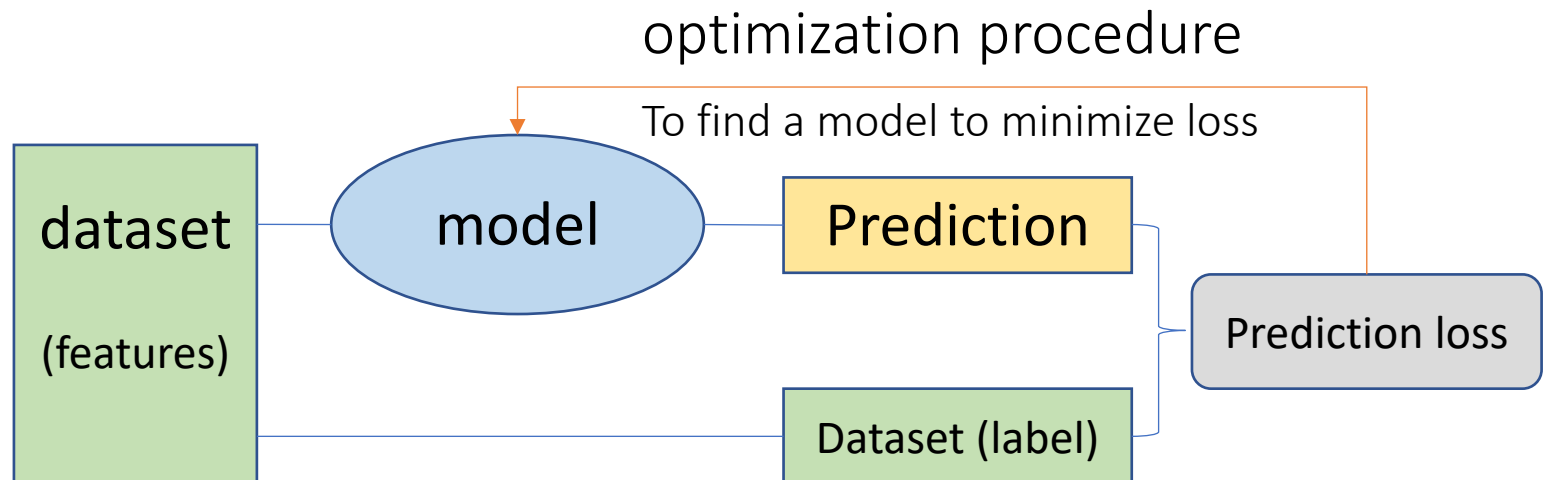
spam (0 or 1)



# Components

Common components of nearly all machine learning algorithms:

- Dataset (training data)
- cost function
- optimization procedure
- model



# Training data

- **Labels**

A **label** is the thing we're predicting—the  $y$  variable in simple linear regression.

e.g., the price of wheat, the kind of animal shown in a picture

- **Features**

A **feature** is an input variable—the  $x$  variable in simple linear regression. A machine learning project could use millions of features, specified as:







$$x_1, x_2, \dots x_N$$

- **Examples**

An **example** is a particular entry of data  $(\mathbf{x}, \mathbf{y})$

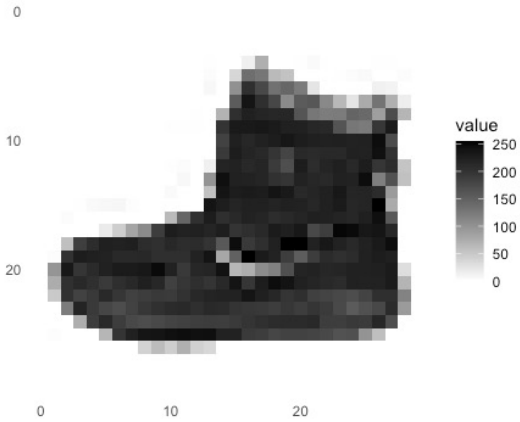
# Training data - example

Table 2: Class names and example images in Fashion-MNIST dataset.

Label	Description	Examples
0	T-Shirt/Top	
1	Trouser	
2	Pullover	
3	Dress	
4	Coat	
5	Sandals	
6	Shirt	
7	Sneaker	
8	Bag	
9	Ankle boots	

## Features of image 1

28 × 28 grayscale

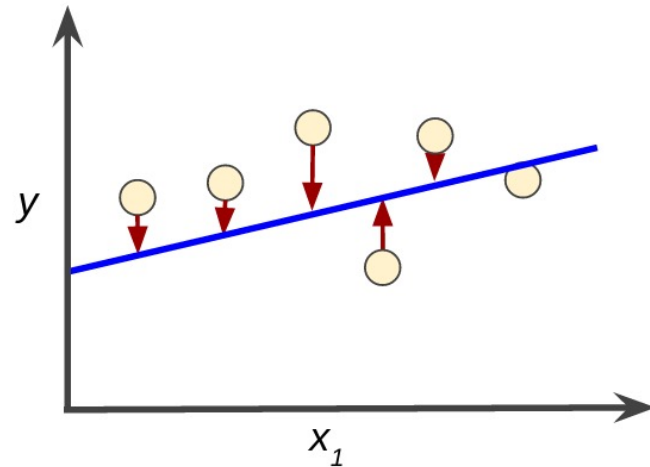
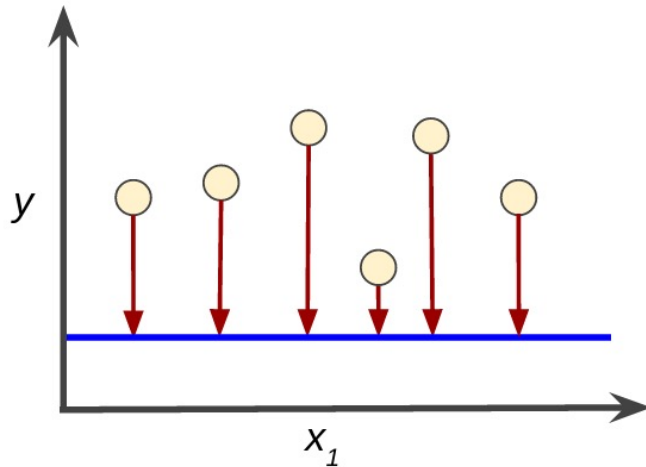


Label = 9

[illegible]

# Cost function for a simple linear regression

Loss is the penalty for a prediction.

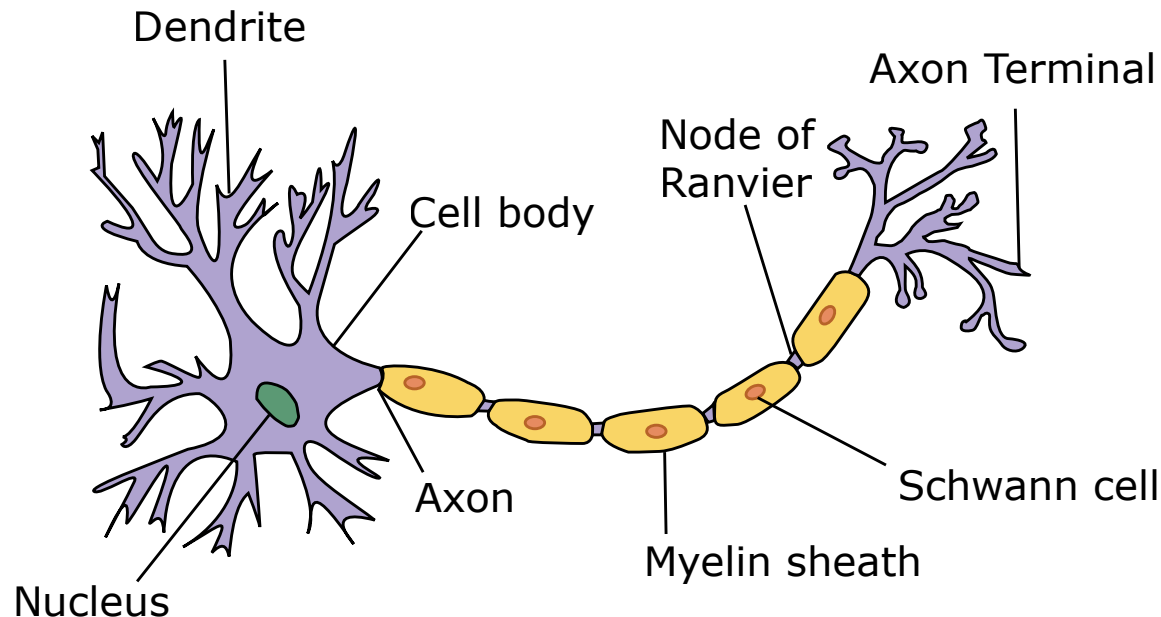


Loss function:  $\frac{1}{N} \sum (y - \text{prediction}(x))^2$

Training a model is to find a set of weights (parameters) that have a *low* loss, on average, across all examples.

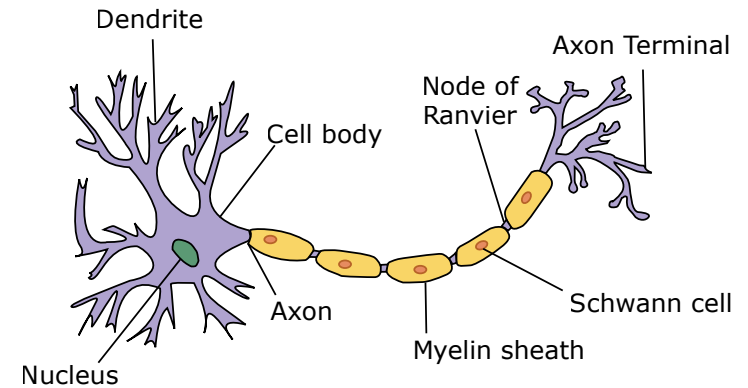
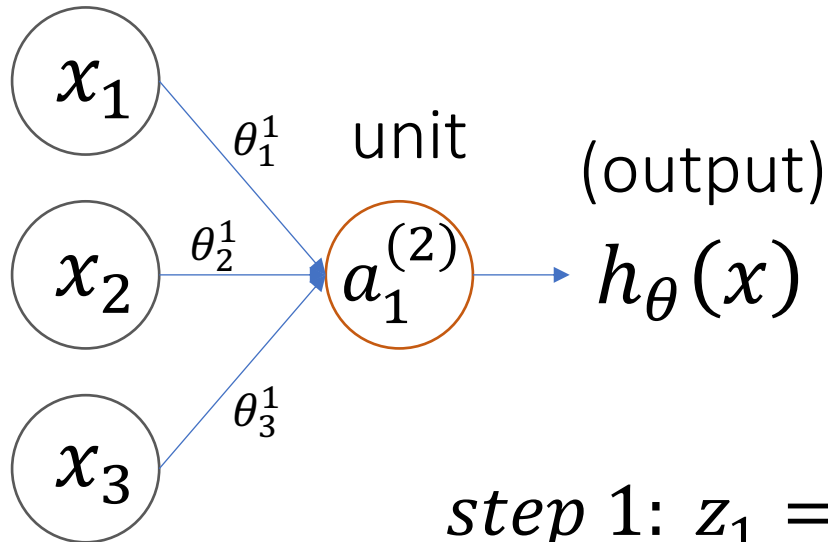
# Neural networks: a main form of deep learning

- Deep learning refers to the recent advances in neural networks and the corresponding training platform
- Algorithms trying to mimic the brain
- Widely used in 80s and early 90s; popularity diminished in late 90s but resurged recently



# Neuron model

Features (input)



$\theta$ : parameters or weights

$$\text{step 1: } z_1 = \theta^T x = \theta_1^1 x_1 + \theta_2^1 x_2 + \theta_3^1 x_3$$

Layer 1

Input layer

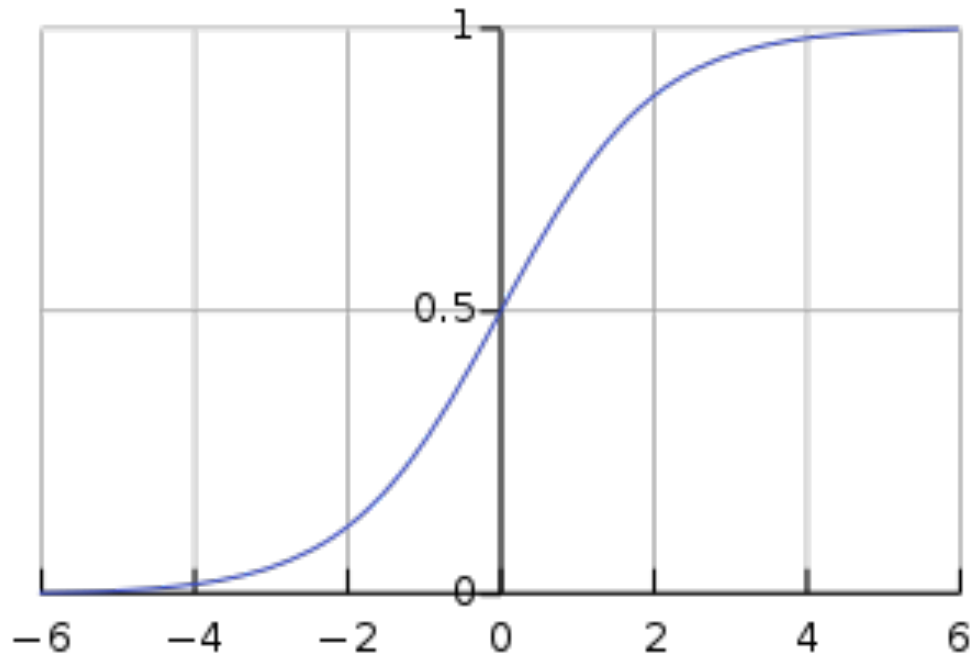
Layer 2

Output layer

Step 2: activation

Activation function:  
Sigmoid (logistic)

$$a_1^{(2)} = h_{\theta}(x) = \frac{1}{1 + e^{-\theta^T x}}$$

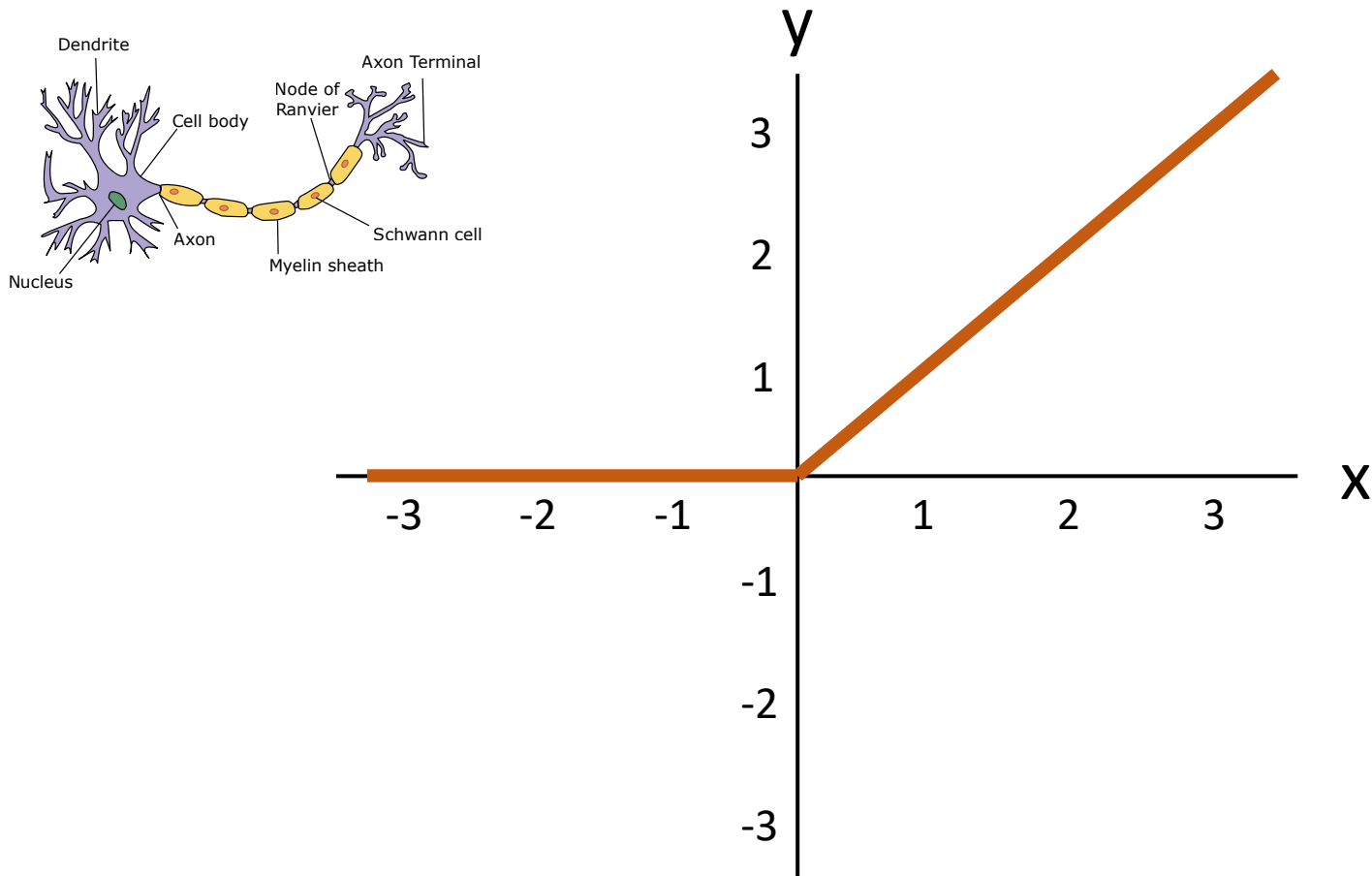


$[0, 1]$



# Activation function:

## ReLU (rectified linear activation function)



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

# Activation function: softmax

$$\sigma(\mathbf{z})_i = \frac{e^{z_i}}{\sum_{j=1}^K e^{z_j}} \quad \text{for } i = 1, \dots, K \text{ and } \mathbf{z} = (z_1, \dots, z_K) \in \mathbb{R}^K.$$

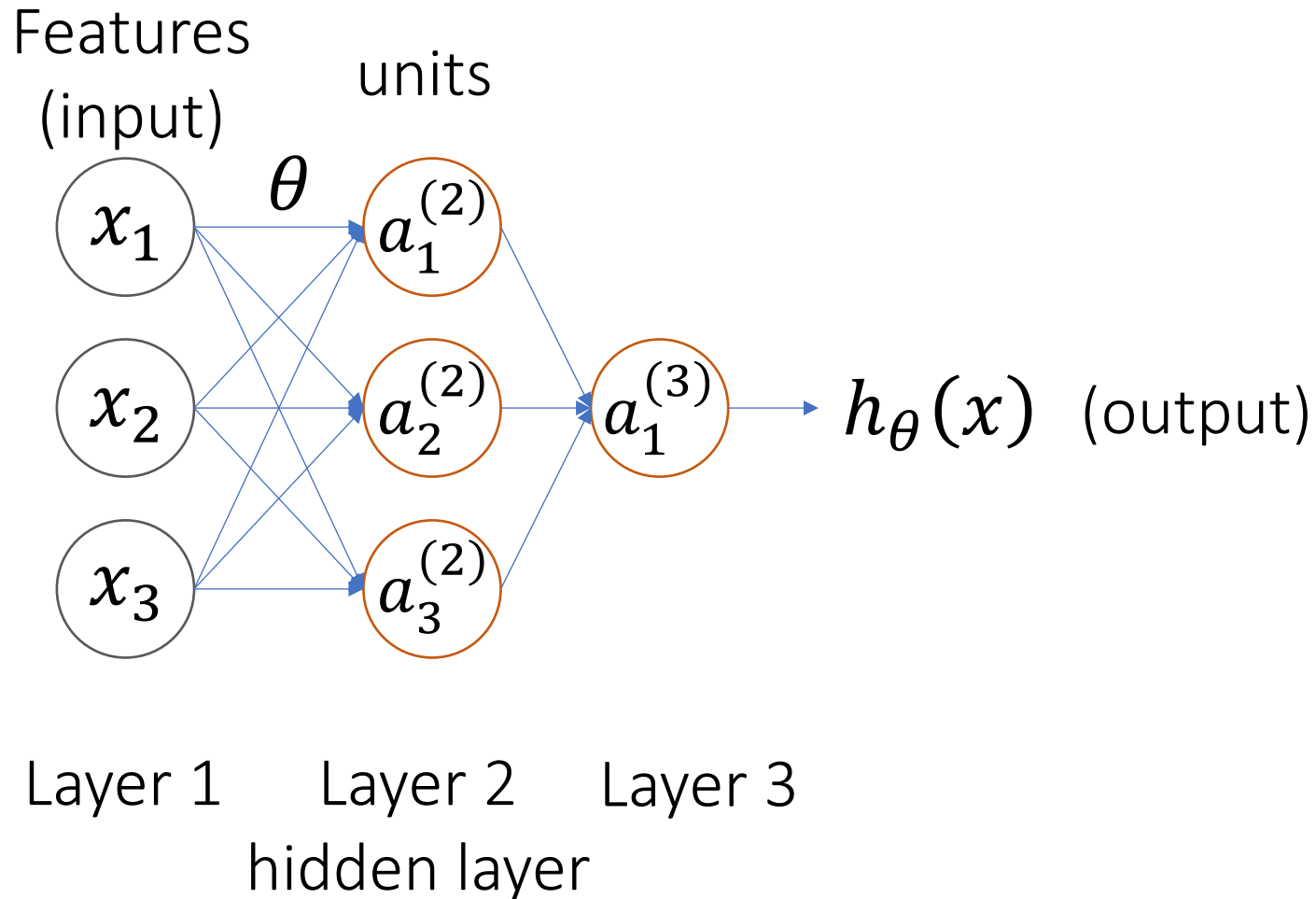
Prior to applying softmax, some vector components could be negative, or greater than one; and might not sum to 1

After applying softmax, each component will be in (0,1), and all values will add up to 1, so that they can be interpreted as probabilities.

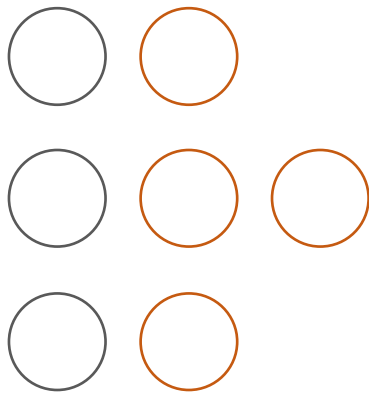
```
softmax <- function(x) { exp(x) / sum(exp(x)) }  
input <- c(0.02, 0.16, 0.10, 0.13, 0.99)  
softmax(input)
```

```
[1] 0.1431069 0.1646121 0.1550259 0.1597471 0.3775080
```

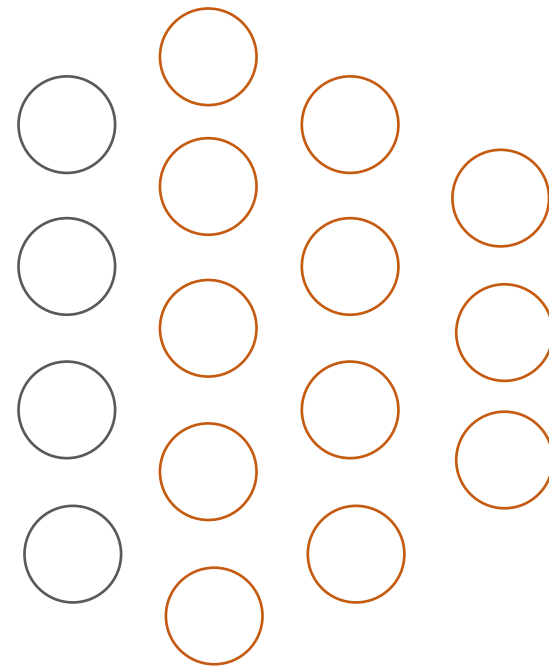
# Neural Network



# Neural network architecture



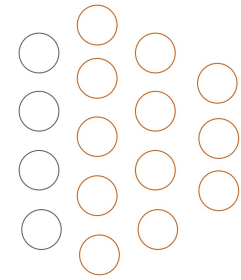
3 layers  
2 units in 2<sup>nd</sup> layer



4 layers  
5 units in 2<sup>nd</sup> layer

...

# Activation function: softmax



$$\sigma(\mathbf{z})_i = \frac{e^{z_i}}{\sum_{j=1}^K e^{z_j}} \quad \text{for } i = 1, \dots, K \text{ and } \mathbf{z} = (z_1, \dots, z_K) \in \mathbb{R}^K.$$

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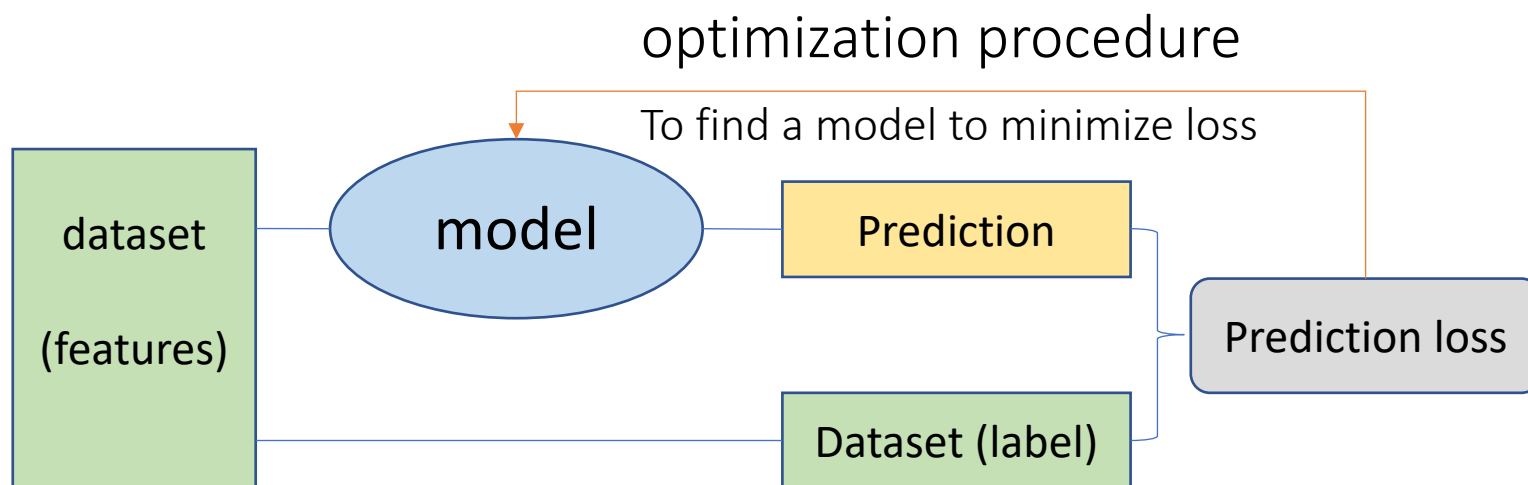
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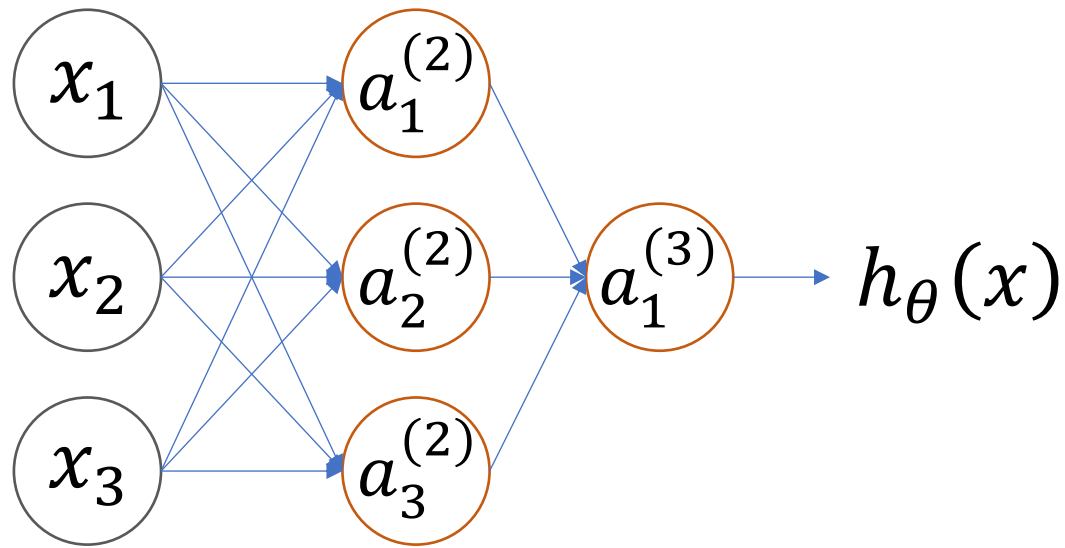
```
[1] 0.1431069 0.1646121 0.1550259 0.1597471 0.3775080
```

# training process

- dataset
- cost function  $\frac{1}{N} \sum (y - \text{prediction}(x, \theta))^2$
- **optimization procedure** To know:  $\frac{\Delta \text{loss}(\theta)}{\Delta \theta}$ , partial derivative
- model (parameters)?



# Forward propagation and backpropagation



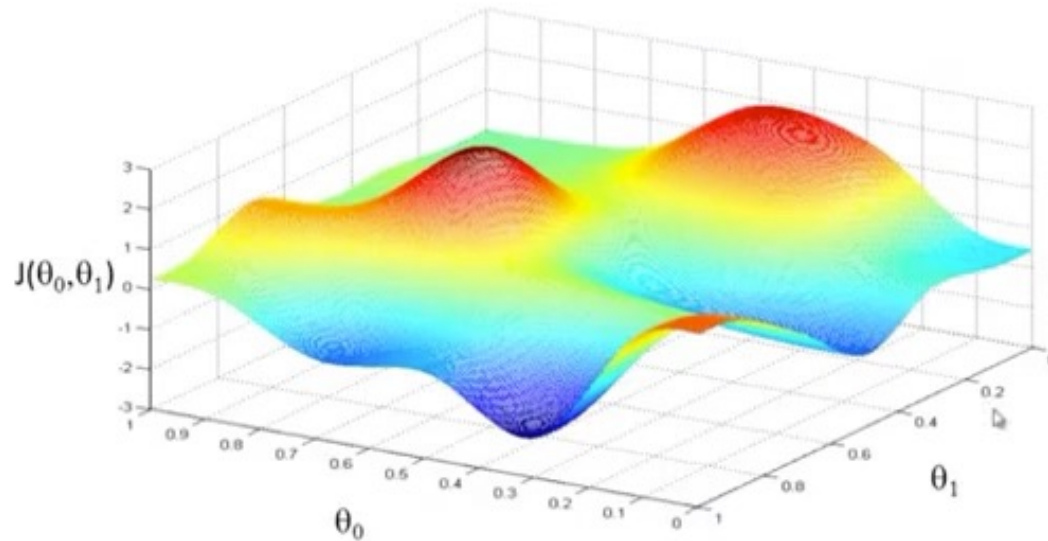
Forward propagation:

To determine  $h_{\theta}(x)$  with the input data  $x$  and parameters  $\theta$  for each example (entry)

Backpropagation:

To compute “error” of each node (e.g.,  $a_1^{(3)}$ ) and then determine partial derivatives of loss function (loss slope for each  $\theta$  or  $\frac{\Delta \text{loss}(\theta)}{\Delta \theta}$ )

# Gradient decent to optimize parameters ( $\theta$ )



Andrew Ng

To find an optimized model is to identify  $\theta$  that can minimize the prediction loss;

Therefore, the key is to understand how the loss changes with the change of each  $\theta$  (partial derivative)



# Procedure of neural network (one typical method)

- Randomly initialization of parameters ( $\theta$ , weights)
- For any input from training data (e.g.,  $(x_i, y_i)$ )
  1. *Forward propagation* to determine  $\hat{y}_i$   
(prediction( $x, \theta$ ))
  2. *Backpropagation* to compute the function of partial derivatives of loss function
- *Gradient descent* (or other optimization methods) to optimize  $\theta$  to minimize the loss

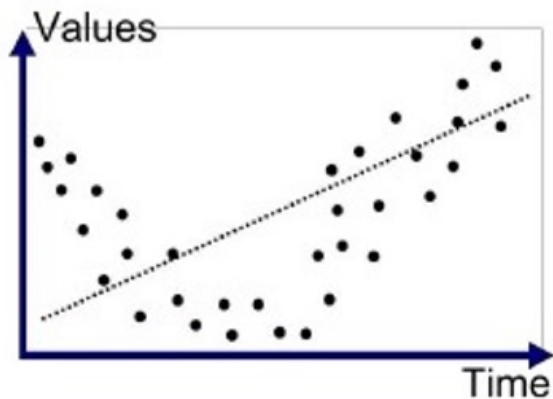
\* my understanding: backpropagation determines how the change of each  $\theta$  changes the loss

# How can we know if a model trained is a good predictor?

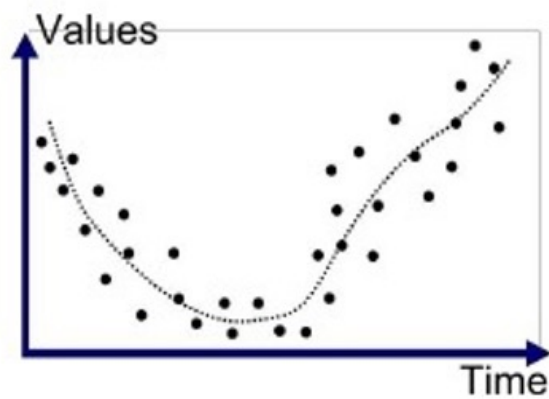
- Training set
  - to learn the model parameters
- Validation set
  - to select the best model
- Test set
  - to estimate the **generalization**

# underfitting and overfitting

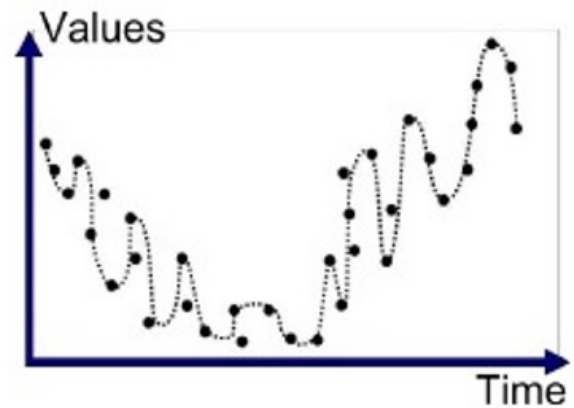
- *Underfitting* occurs when the model obtains an insufficiently low error value on the training set.
- *Overfitting* occurs when the gap between the training error and test/validation error is too large.



Underfitted

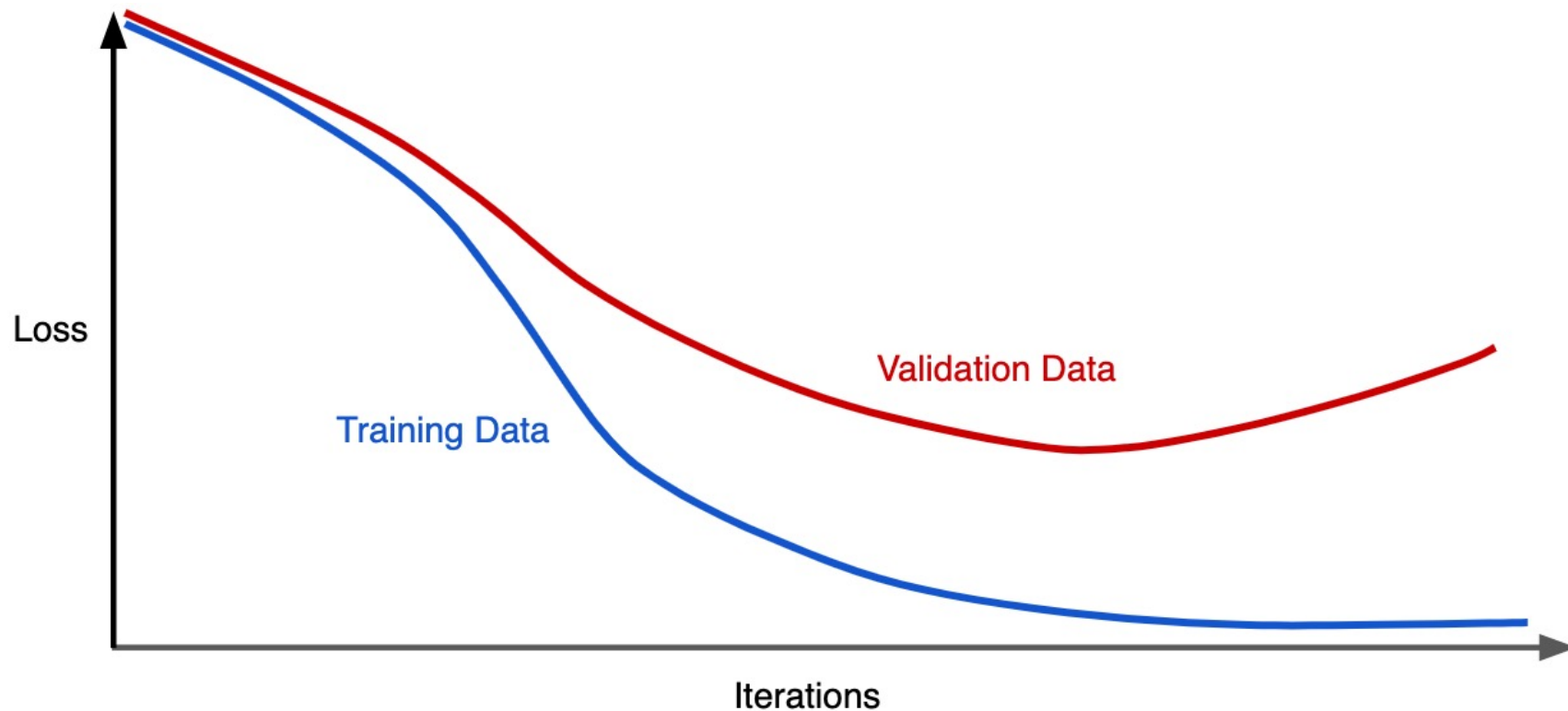


Good Fit/Robust



Overfitted

# underfitting and overfitting



Make the training error small.

Make the gap between training and validation error small.

# regularization

Regularization refers to a set of different techniques that lower the complexity of a neural network model during training, and thus prevent the overfitting.

Large weights ( $\theta$ ) in a neural network are a sign of a complex model that possibly overfits the training data.

Original loss function + regularization term

$$\frac{1}{N} \sum (y - \text{prediction}(x, \theta))^2 + \lambda \|\theta\|^2$$

$$L_2 \text{ regularization term} = \lambda \|\theta\|^2$$

$\lambda$ : weight of regulation  
 $\|\theta\|^2$ : sum of square

# Dropout: a simple and effective regularization method

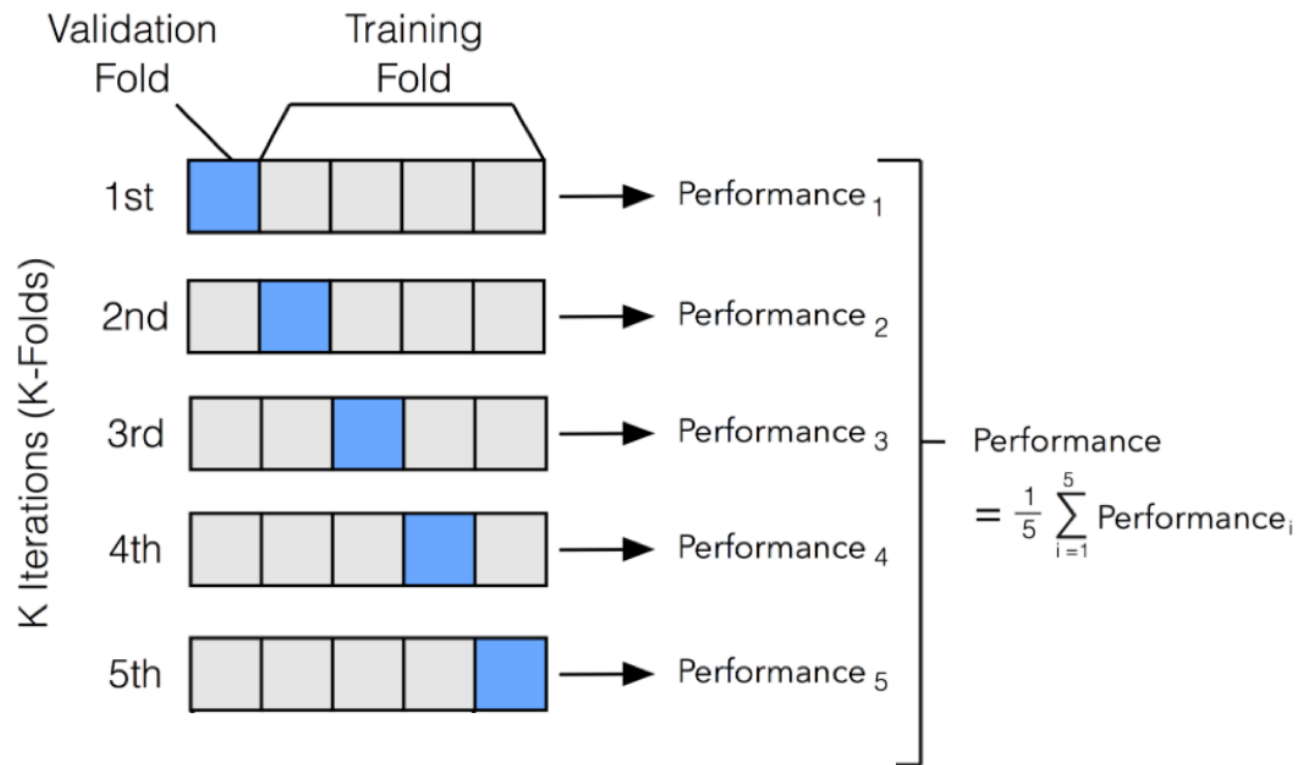
A model is adjusted by dropping out nodes during training. The node is temporarily removed from the network, along with all its incoming and outgoing connections.

Dropout is a computationally cheap and remarkably effective regularization method to reduce overfitting and improve generalization in deep neural networks of all kinds.

# K-fold cross validation

the k-fold cross-validation partitions a dataset into K nonoverlapping subsets. The model performance may then be estimated by taking the average performance across K trials.

5-fold



# software

- Tensorflow: a platform developed by the Google Brain team for machine learning.
- Keras: a deep learning software written in Python, running on top of the machine learning platform TensorFlow.



# Deep learning in genomics

1. Large training datasets (e.g., thousands of examples), curated to remove confounders, are typically required.
2. Most genomic data do not require very deep networks.
3. Researchers must be wary of high accuracy due to data imbalance or bias that makes classification too easy.
4. A good practice is to compare against simpler machine learning models on the same dataset.
5. Deep learning can achieve high accuracy, but the interpretation of results is more challenging than for standard statistical models.

# References

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- Google course: <https://developers.google.com/machine-learning/crash-course/ml-intro>