

## Introduction to Neural Networks

Dr. Manuel Gehl



#### **Overview**

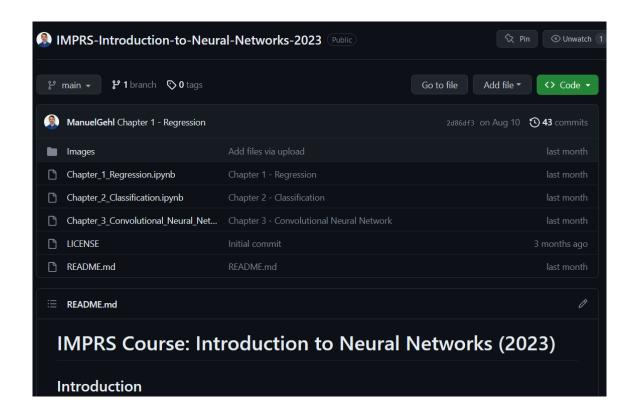
#### Lecture

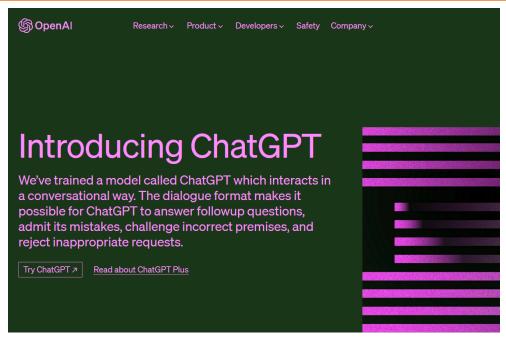
- What are Artificial Neural Networks?
- What is an Artificial Neuron?
- How Do Artificial Neurons Learn?
- What next?

#### **Notebooks**

- Chapter 1 Regression
- Chapter 2 Classification
- Chapter 3 –
   Convolutional Neural
   Networks

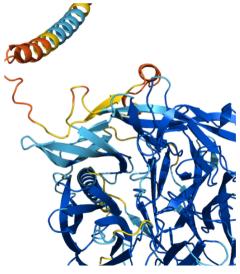
#### **Course Website**





AlphaFold is an AI system developed by DeepMind that predicts a protein's 3D structure from its amino acid sequence. It regularly achieves accuracy competitive with experiment.

DeepMind and EMBL's European Bioinformatics
Institute (EMBL-EBI) have partnered to create
AlphaFold DB to make these predictions freely
available to the scientific community. The latest
database release contains over 200 million entries,
providing broad coverage of UniProt (the standard
repository of protein sequences and annotations). We
provide individual downloads for the human proteome
and for the proteomes of 47 other key organisms
important in research and global health. We also
provide a download for the manually curated subset
of UniProt (Swiss-Prot).



Q8I3H7: May protect the malaria parasite against attack by the immune system. Mean pLDDT 85.57.



#### Deep Learning for Protein-Protein Interaction Site Prediction

Arian R. Jamasb, Ben Day, Cătălina Cangea, Pietro Liò & Tom L. Blundell

Protocol Open Access First Online: 09 July 2021

**5070** Accesses **7** <u>Citations</u> **10** <u>Altmetric</u>

Part of the Methods in Molecular Biology book series (MIMB, volume 2361)

### Classification and mutation prediction from non-small cell lung cancer histopathology images using deep learning

Nicolas Coudray, Paolo Santiago Ocampo, Theodore Sakellaropoulos, Navneet Narula, Matija Snuderl, David Fenyö, Andre L. Moreira, Narges Razavian ≅ & Aristotelis Tsirigos ≅

Nature Medicine 24, 1559–1567 (2018) Cite this article

SCIENCE • 21 Jul 2022 • Vol 377, Issue 6604 • pp. 387-394 • DOI: 10.1126/science.abn2100

#### Scaffolding protein functional sites using deep learning



#### De novo design of luciferases using deep learning

Andy Hsien-Wei Yeh , Christoffer Norn, Yakov Kipnis, Doug Tischer, Samuel J. Pellock, Declan Evans, Pengchen Ma, Gyu Rie Lee, Jason Z. Zhang, Ivan Anishchenko, Brian Coventry, Longxing Cao, Justas Dauparas, Samer Halabiya, Michelle DeWitt, Lauren Carter, K. N. Houk & David Baker

*Nature* **614**, 774–780 (2023) Cite this article

#### Geometric deep learning of RNA structure

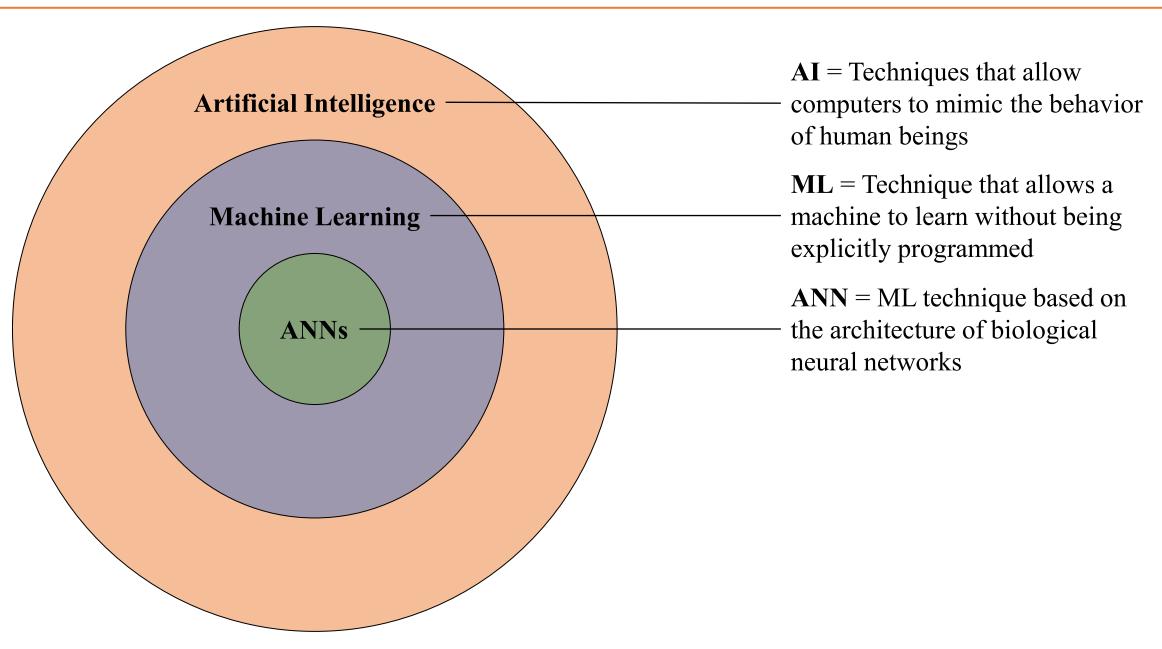


STRUCTURAL BIOLOGY

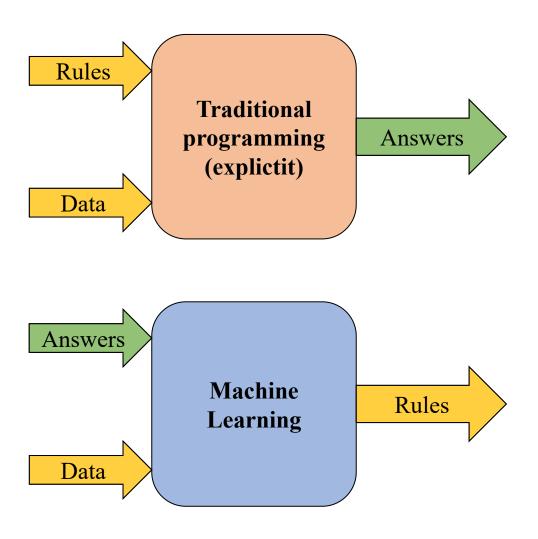
#### Protein sequence design by deep learning

Jue Wang <sup>™</sup>

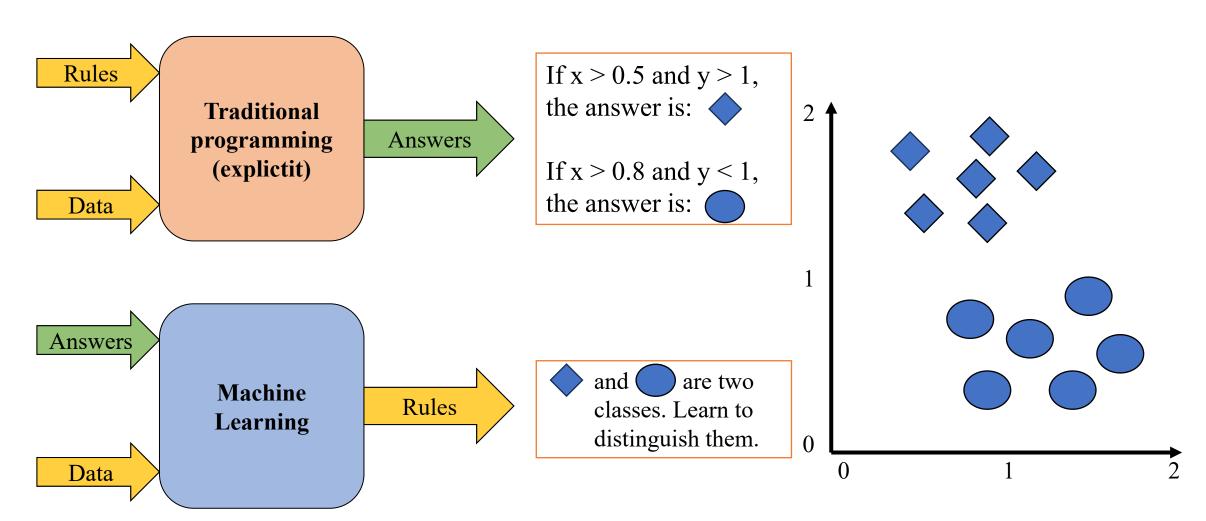
Nature Computational Science 2, 416–417 (2022) | Cite this article

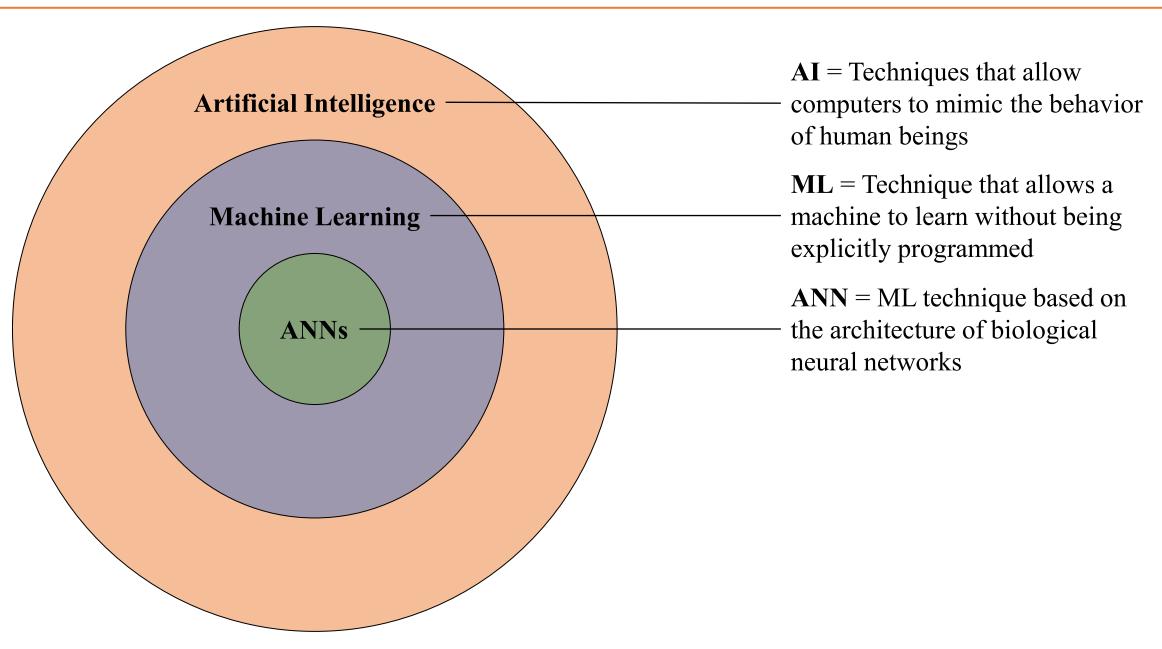


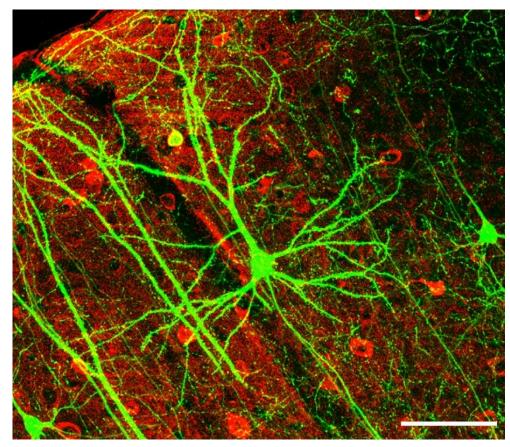
ML = Technique that allows a machine to learn without being explicitly programmed



ML = Technique that allows a machine to learn without being explicitly programmed







Wei-Chung Allen Lee, Hayden Huang, Guoping Feng, Joshua R. Sanes, Emery N. Brown, Peter T. So, Elly Nedivi, PLoSBiol4.e126.Fig6fNeuron, CC BY 2.5

- 1. Neurons (units): Basic processing unit.
- 2. Connectivity: Neurons are connected to each other.
- 3. Activation: Neurons "fire" after exceeding threshold.
- **4.** Layers: Neurons are organized in layers.
- 5. Feedforward information flow: From input to output.
- **6.** Feedback information flow: From output to input.
- 7. Learning and Adaptation: Adapting connections to stimuli.
- **8.** Parallel processing: Processing information simultaneously.

#### 1. Data

# WVIDIA XGPU S G46608.1 0337A3 S TAIWAN

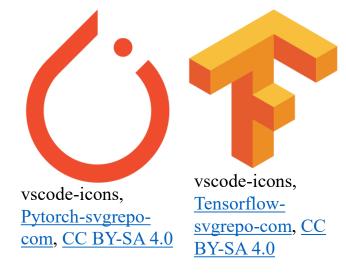
A7N8X, Xbox GPU, CC BY-SA 4.0

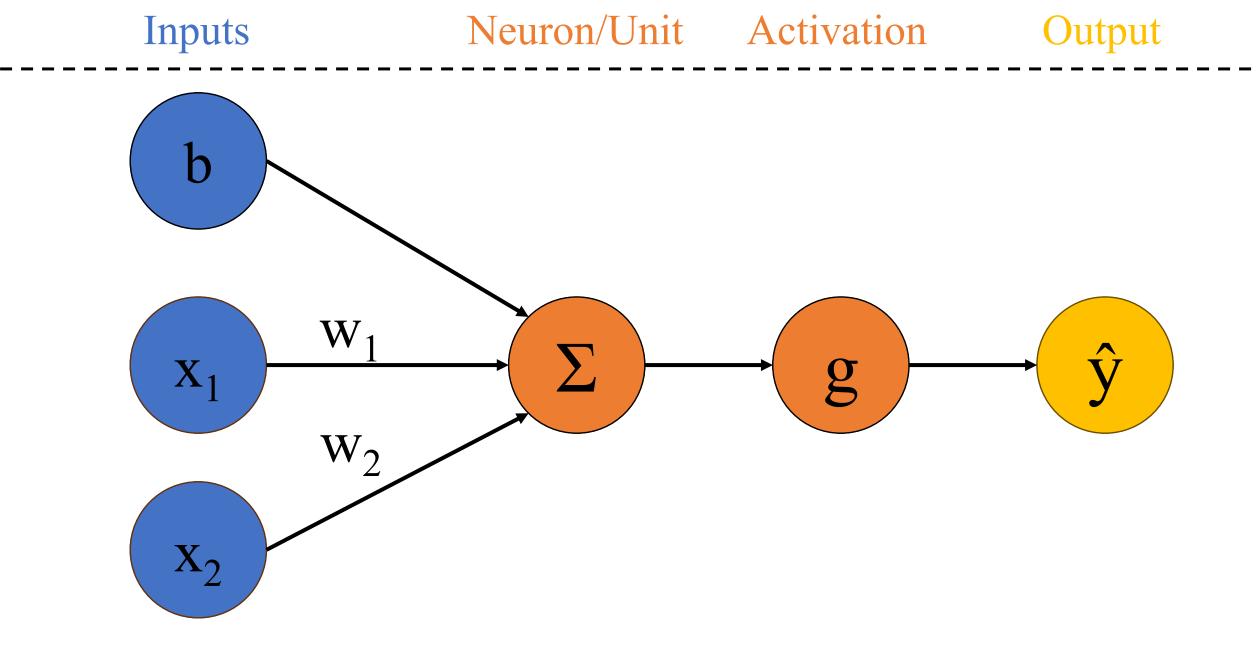
#### 2. Hardware

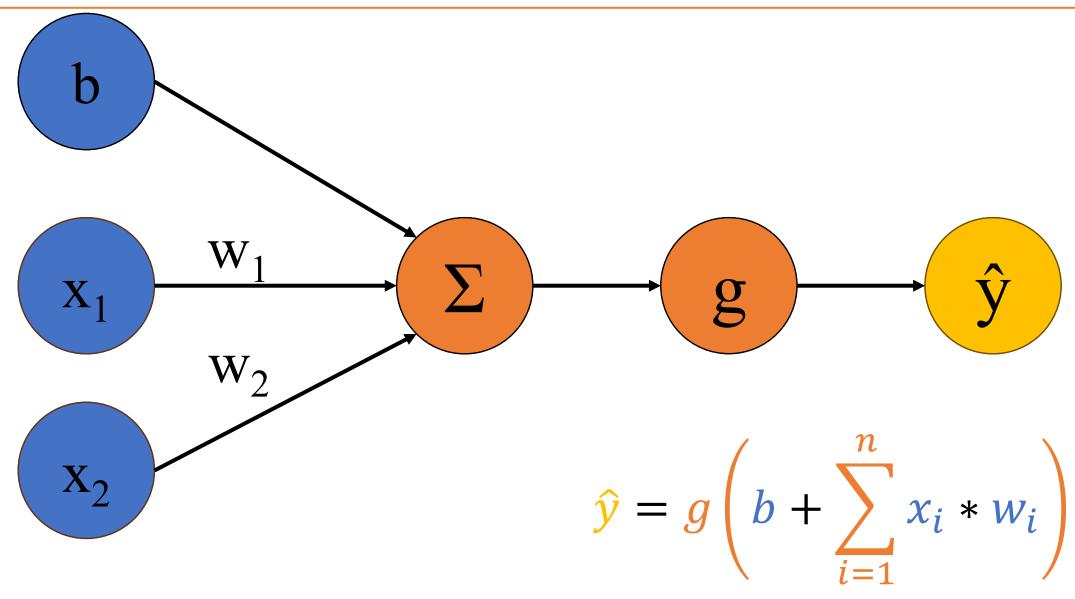


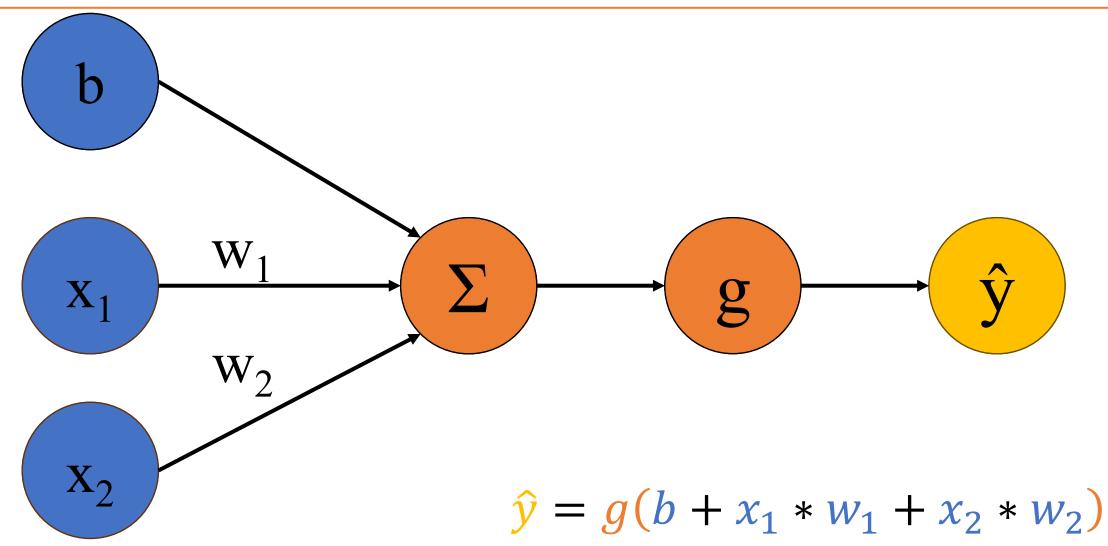


Zxb, Storage size comparison, CC BY-SA 2.0 AT









#### **Example: Oktoberfest**



<u>Heribert Pohl --- Thanks for half a million clicks!</u> from Germering bei München, Bayern, <u>O'zapft is! Münchens 5 Jahreszeit hat begonnen - O'zapft is! Munich 5 season, the Oktoberfest has begun (9855483374), CC BY-SA 2.0</u>

## **Prediction:** Are you going to the Oktoberfest this Saturday?

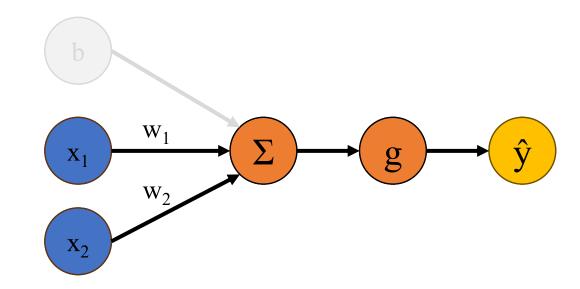
#### Criteria (weights):

- 1. Do you like beer?
  - Yes (1) No (0)
- 2. How important is the weather to you?
  - Very (1) Not at all (0)

**Prediction:** Are you going to the Oktoberfest this Saturday?

#### Criteria (weights):

- 1. Do you like beer?
  - Yes (1) No(0)
- 2. How important is the weather to you?
  - Very (1) Not at all (0)

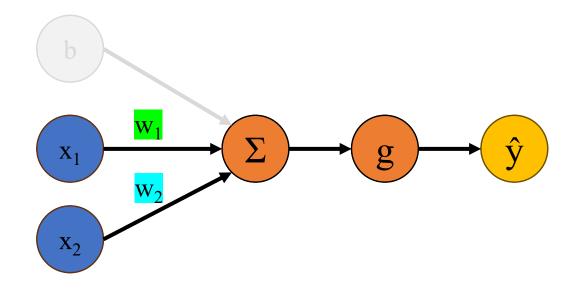


$$\hat{y} = g(x_1 * w_1 + x_2 * w_2)$$

**Prediction:** Are you going to the Oktoberfest this Saturday?

#### **Criteria:**

- 1. Do you like beer?
  - Yes (1) No(0)
- 2. How important is the weather to you?
  - Very (1) Not at all (0)

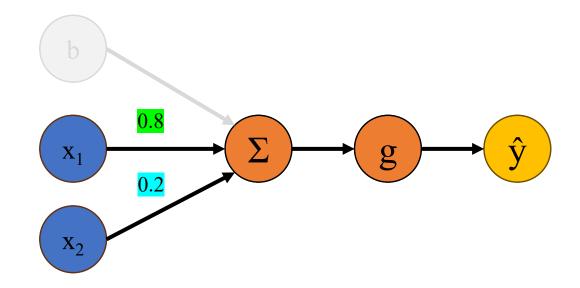


$$\hat{y} = g(x_1 * w_1 + x_2 * w_2)$$

**Prediction:** Are you going to the Oktoberfest this Saturday?

#### Person 1:

- 1. Do you like beer?
  - ,,Pretty much" (0.8)
- 2. How important is the weather to you?
  - "There is no bad weather, only the wrong clothes" (0.2)

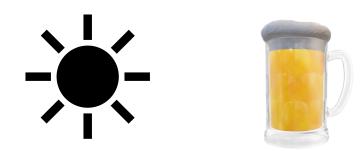


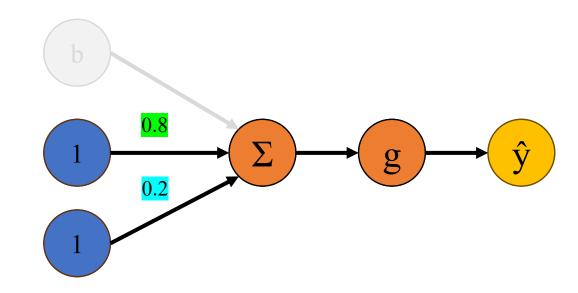
$$\hat{y} = g(x_1 * \mathbf{0.8} + x_2 * \mathbf{0.2})$$

**Prediction:** Are you going to the Oktoberfest this Saturday?

#### Person 1:

- 1. Do you like beer?
  - ,,Pretty much" (0.8)
- 2. How important is the weather to you?
  - "There is no bad weather, only the wrong clothes" (0.2)



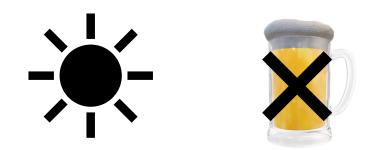


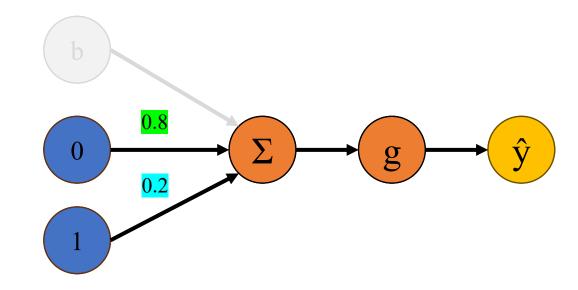
$$\hat{y} = g(1 * 0.8 + 1 * 0.2)$$
  
 $\hat{y} = g(1)$ 

**Prediction:** Are you going to the Oktoberfest this Saturday?

#### **Person 1:**

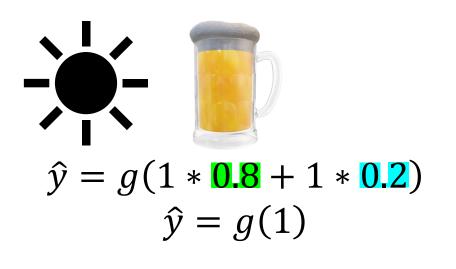
- 1. Do you like beer?
  - ,,Pretty much" (0.8)
- 2. How important is the weather to you?
  - "There is no bad weather, only the wrong clothes" (0.2)

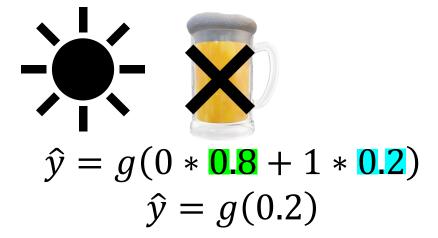


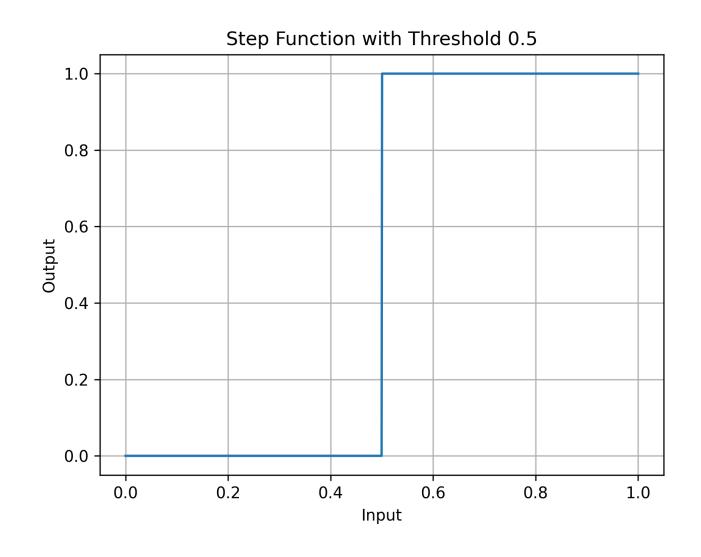


$$\hat{y} = g(0 * 0.8 + 1 * 0.2)$$
  
 $\hat{y} = g(0.2)$ 

**Prediction:** Are you going to the Oktoberfest this Saturday?



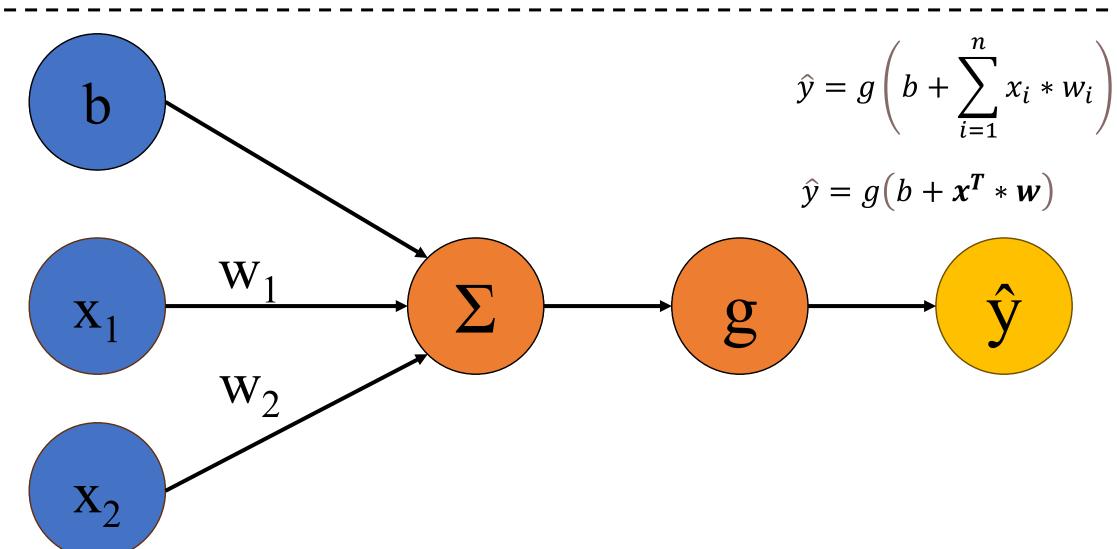


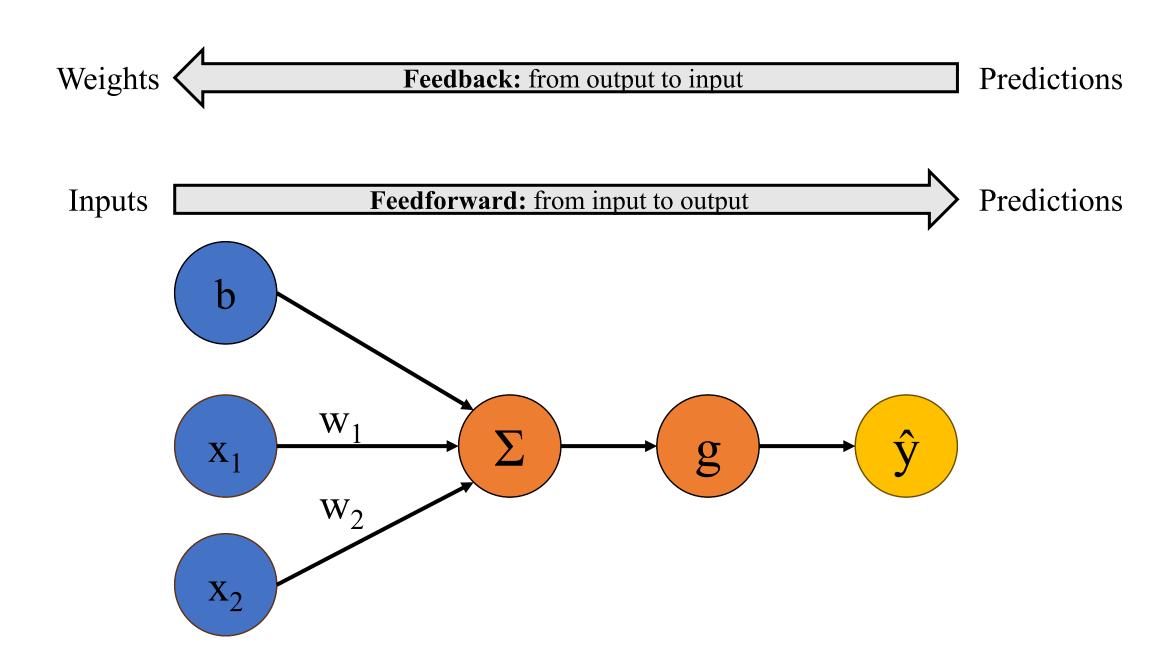




Neuron/Unit Activation

Output





**Loss functions** measure difference between true values (y) and predicted values (ŷ):

Mean Absolute Error (MAE)
$$\sum_{i=1}^{n} \frac{|y_i - y_i|}{n}$$

- Mean Squared Error (MSE)
- Binary Crossentropy
- Categorical Crossentropy

$$\sum_{i=1}^{n} \frac{y_i * \log(p(y_i)) + (1 - y_i) * (\log(1 - p(y_i)))}{n}$$

 $p(y_i)$  = Probability the model assigns the value to be  $\hat{y}$ 

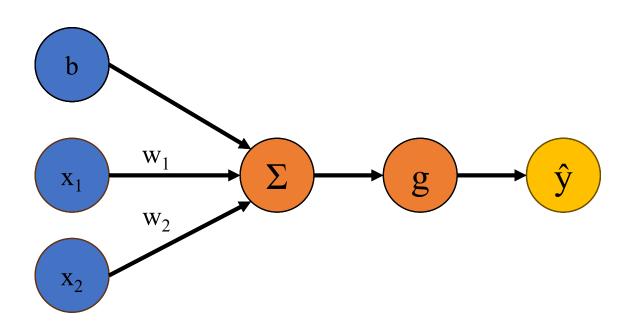
**Loss functions** measure difference between true values (y) and predicted values (ŷ):

- Mean Absolute Error (MAE)
- Mean Squared Error (MSE)
- Binary Crossentropy
- Categorical Crossentropy

•

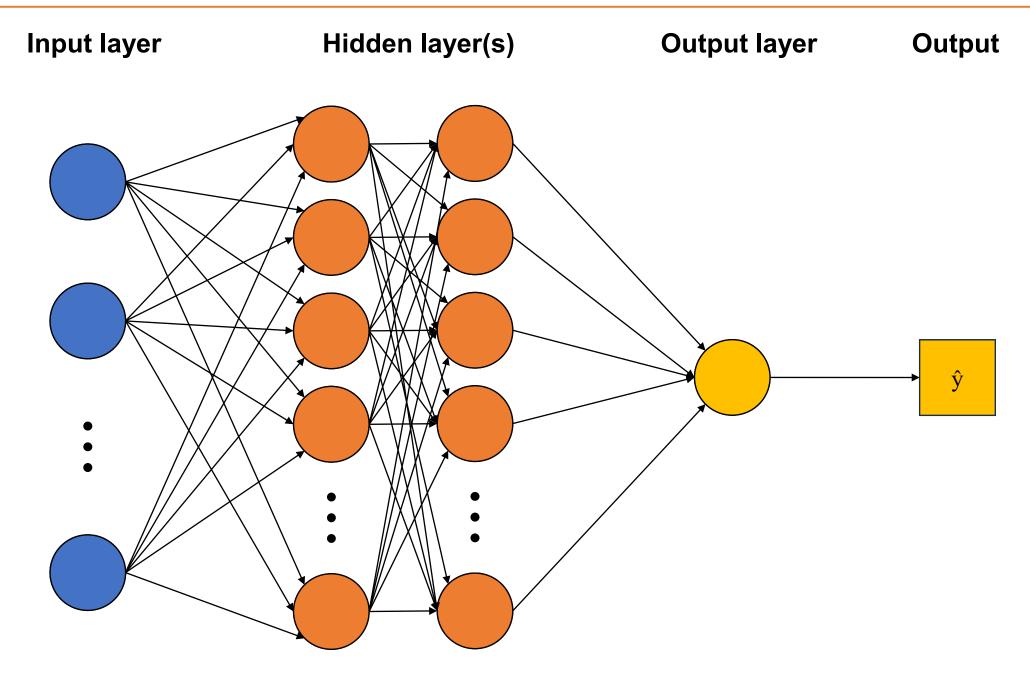
**Training** an artificial neuron (or network; learning) refers to the **minimization of the loss function** 

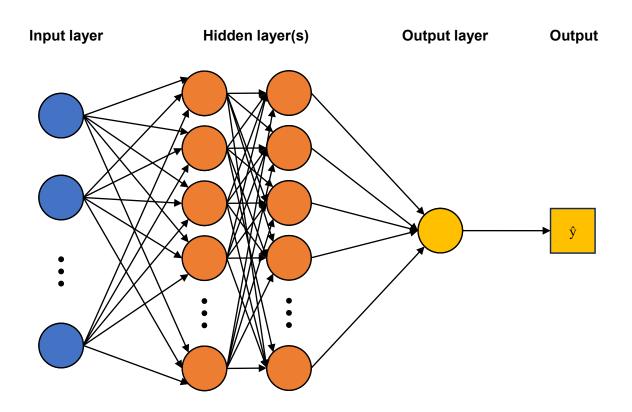
The error is a function of all neurons, which in turn are a function of all weights. The error is **backpropagated** through the network and the weights are adjusted to minimize the loss function.



**Training** an artificial neuron (or network, learning) refers to the **minimization of the loss function** 

The error is a function of all neurons, which in turn are a function of all weights. The error is **backpropagated** through the network and the weights are adjusted to minimize the loss function.

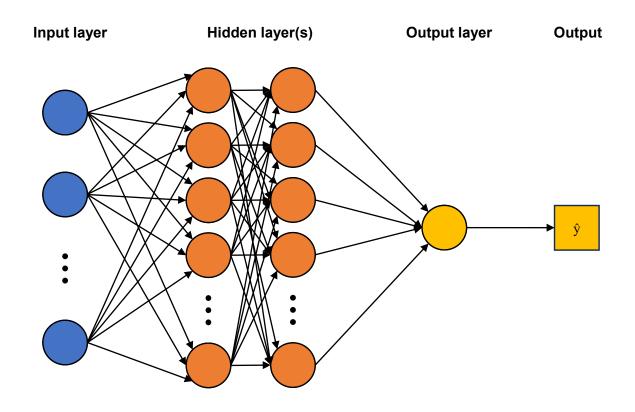




#### Universal approximation theorem:

An ANN with enough neurons in the hidden layer can approximate any continuous function with arbitrary precision.

**Deep Learning:** Multiple hidden layers.

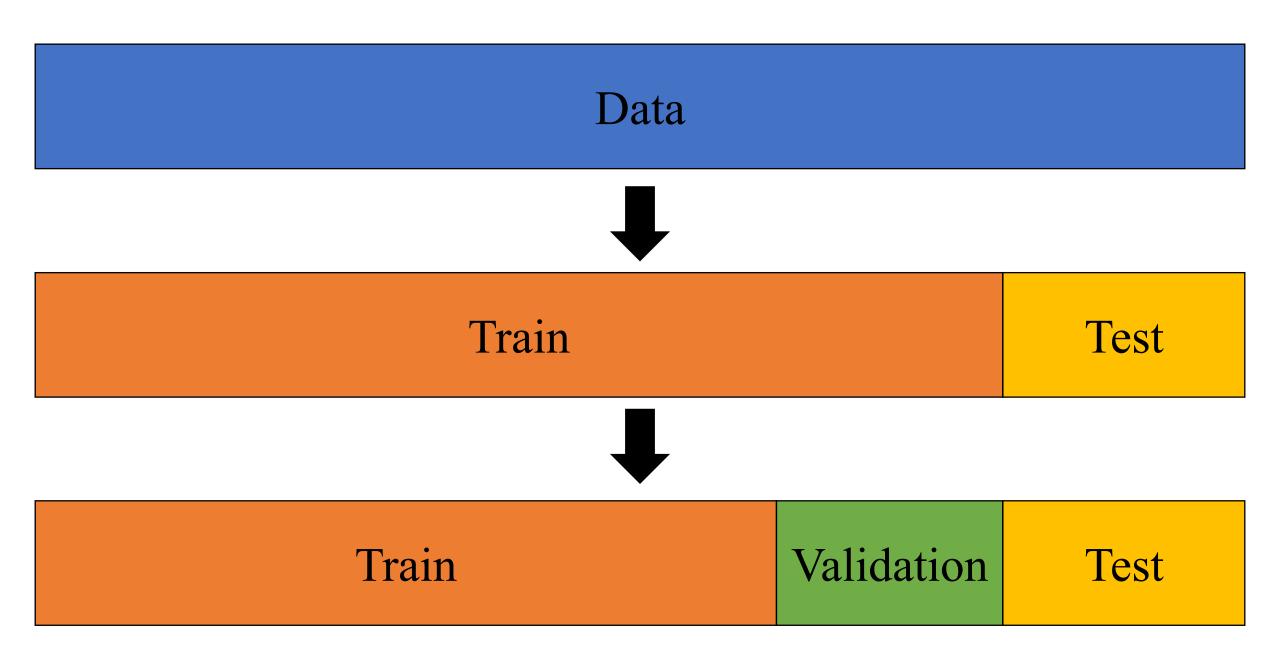


- Pattern Detector
- High-dimensional feature extractor
- Function approximator
- Computational model

#### **Chapter 1 - Regression**

- Regression model for Bradford assay
- Model for growth curve during diauxic growth
- Model for a circadian pattern of enzyme activity
- Human vs. Machine Learning
- The curse of small data sets
- Overfitting, Underfitting, and Holdout Validation

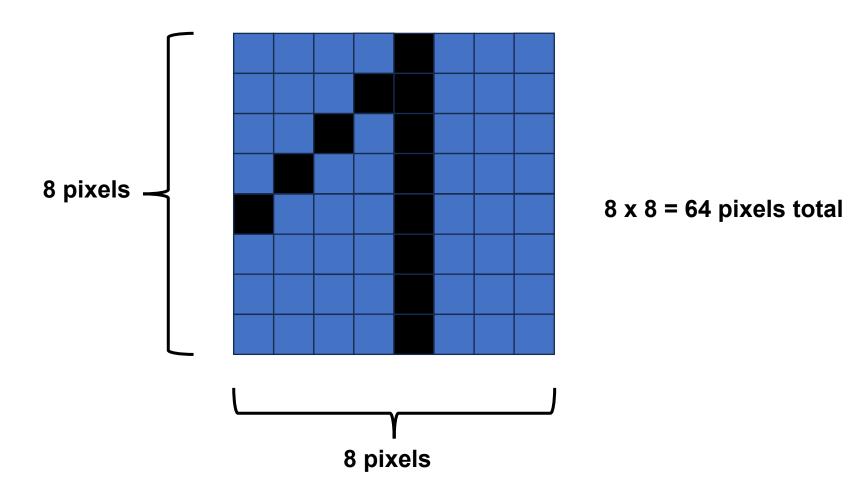
#### **Holdout validation**



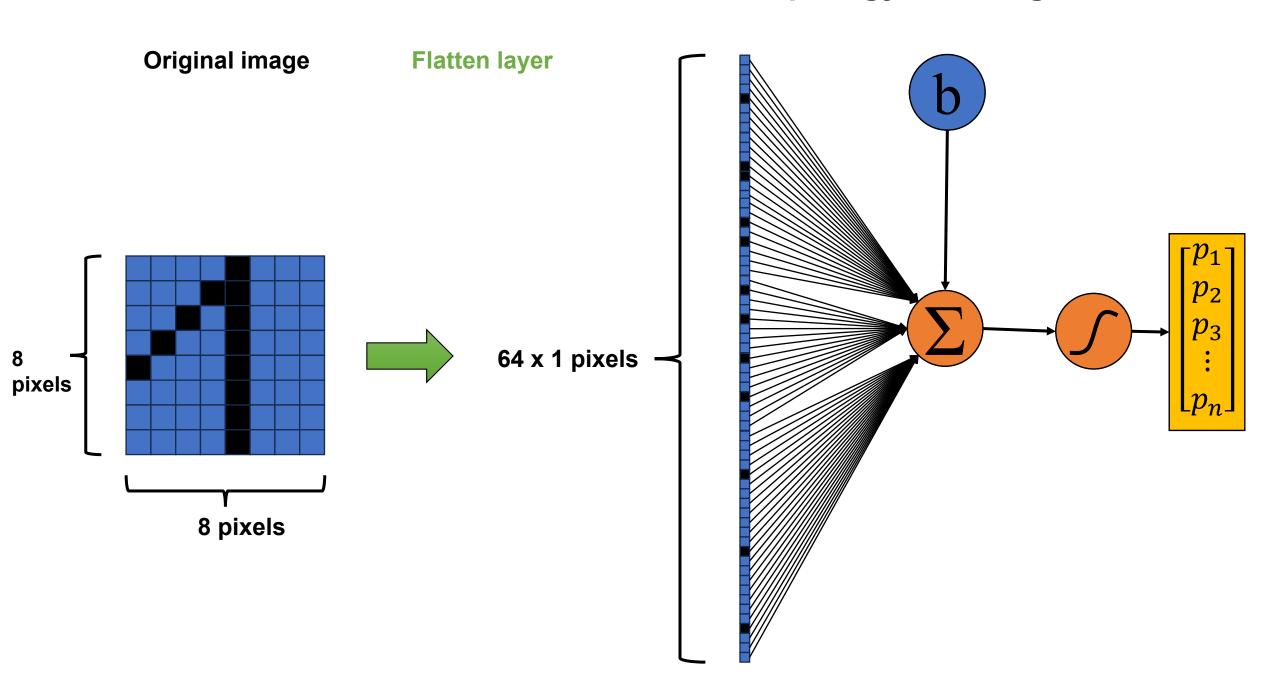
#### **Chapter 2 - Classification**

- Binary classification to distinguish microbial cells
- Multi-class classification to distinguish microbial cells
- Simple image classification using the MNIST dataset
- Precision and Recall, Confusion matrix and decision boundaries
- Black box problem

#### Multi-class classification network topology for images



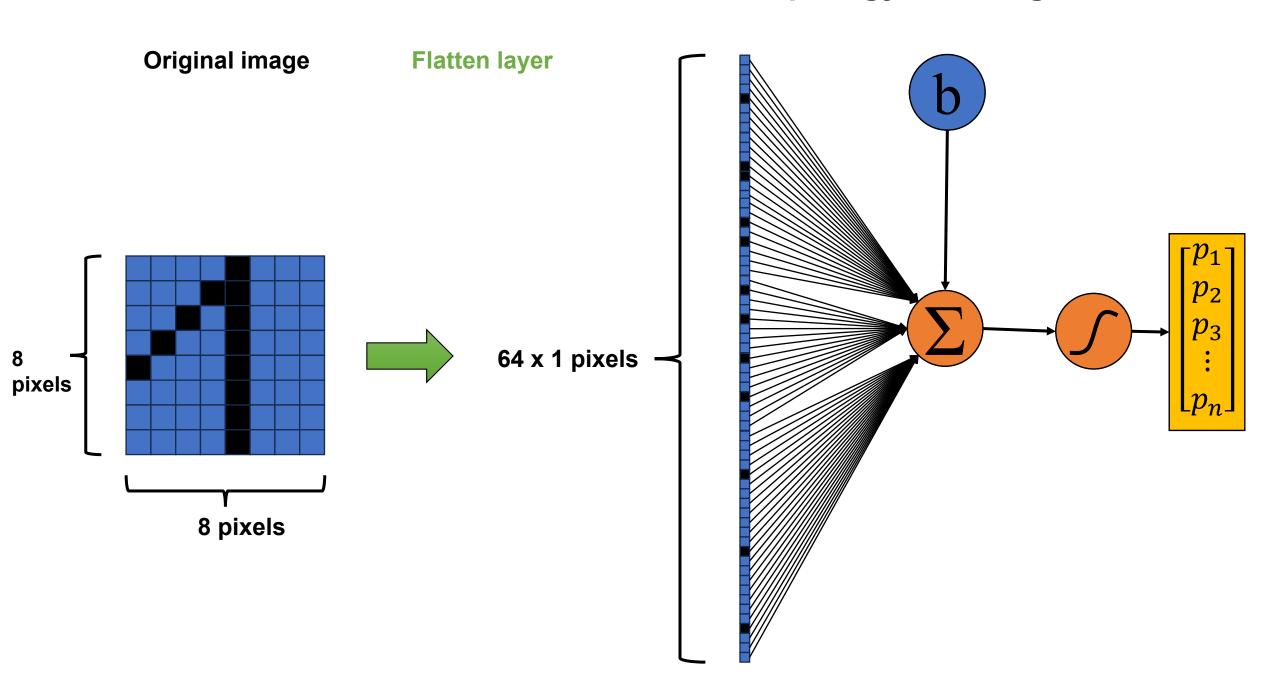
#### Multi-class classification network topology for images

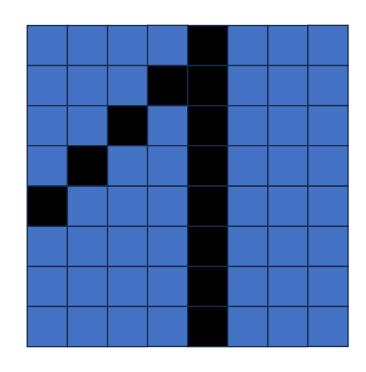


#### **Chapter 3 – Convolutional Neural Networks**

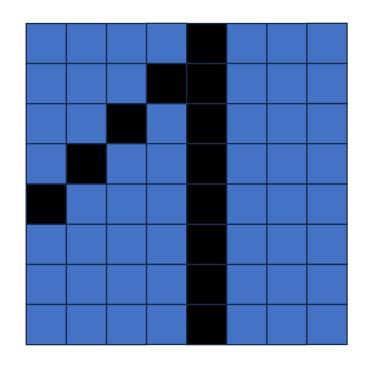
- CNN for healthy and infected cell classification
- Convolution layer and pooling layer
- Transfer learning

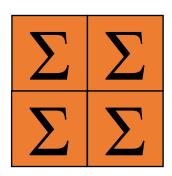
#### Multi-class classification network topology for images

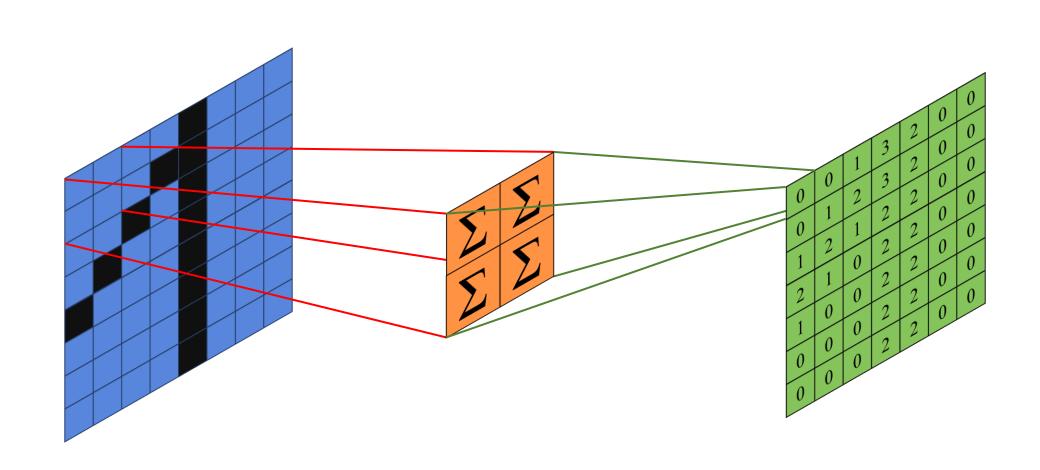


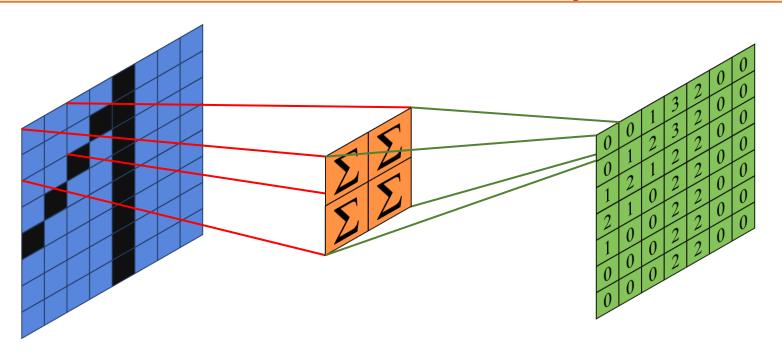










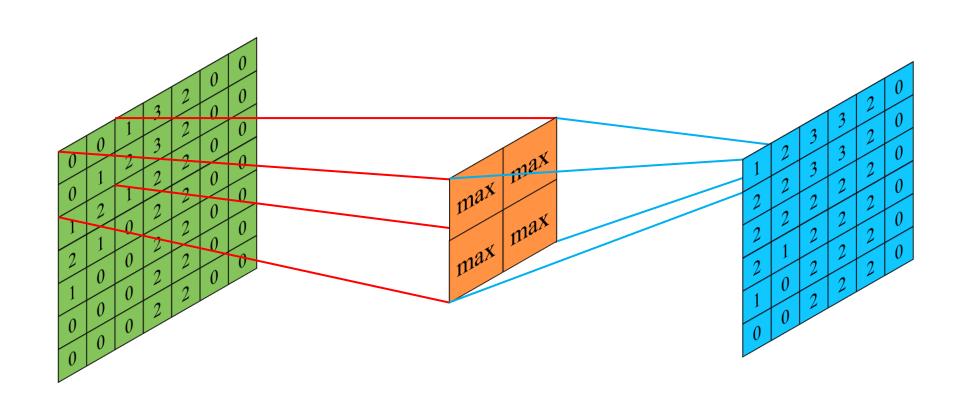


Convolutional layers preserve the intrinsic structure of the data. The output is called a **feature map**. As a result, they perform very well on **structured data**.

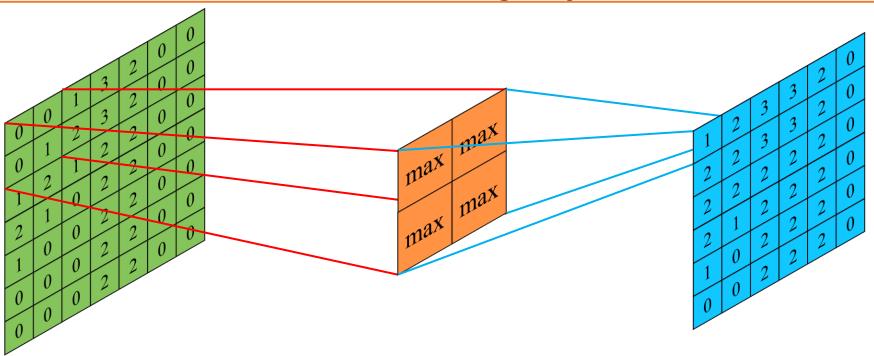
#### Examples include:

- Computer Vision
- Natural Language Processing
- Time series forecasting and classification
- Molecular structure prediction

#### What is a Pooling Layer?

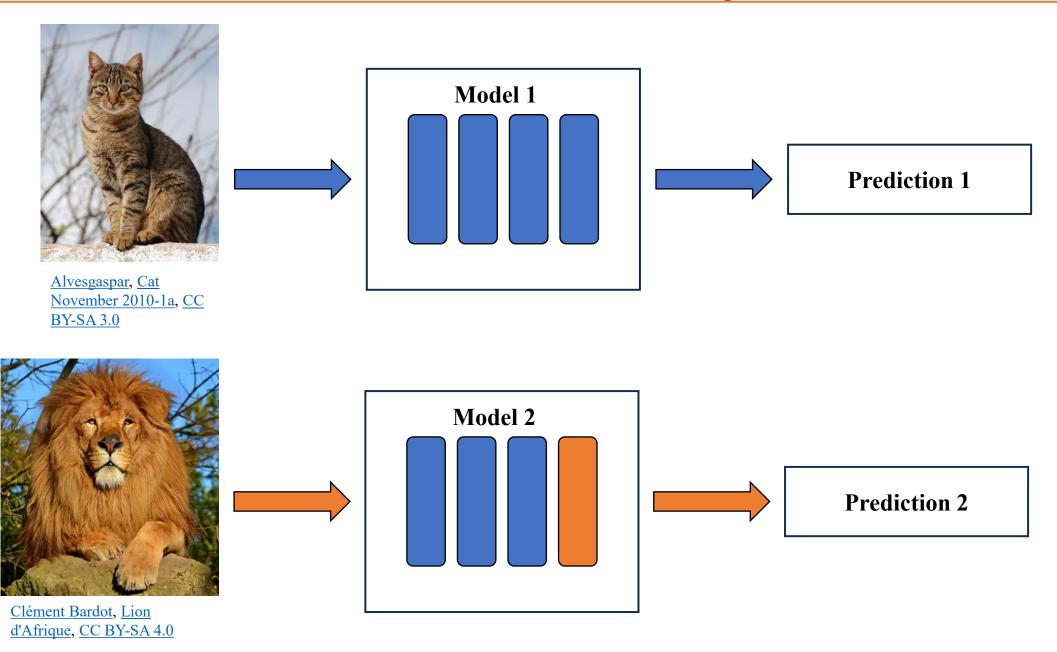


#### What is a Pooling Layer?

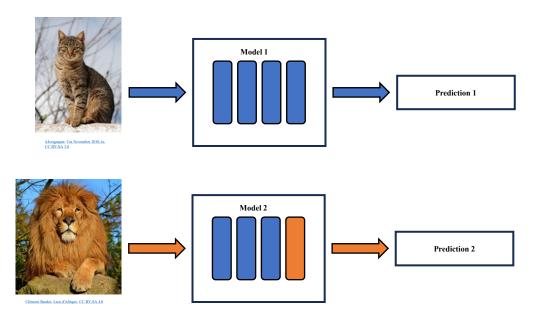


Pooling layers reduce the size of feature maps while preserving **meaningful features**.

#### **What is Transfer Learning?**



#### What is Transfer Learning?



Transfer learning is a technique that uses a **pre-trained model** for a related task.

#### Benefits:

- Faster training
- Improved performance with small amount of data
- Generalization: Pre-trained models have typically learned general features

#### What next?

