

# Tutorial for Computer Vision Technical Project

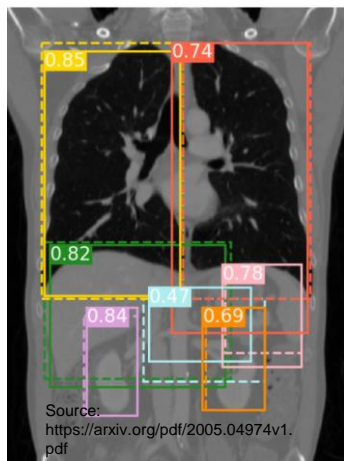
03/02/2025

Chengliang Dai

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# Key Tasks in Medical Image Analysis

Localisation



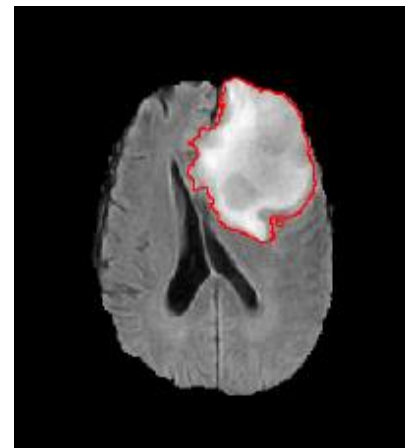
Output: bounding boxes

Classification



Output: tumour/no tumour

Semantic Segmentation



Output: segmentation map

## Other Tasks in Computer Vision

Image Classification



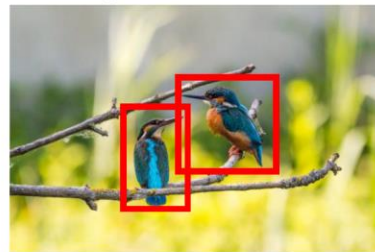
Output: Category (e.g., "bird")

Object Detection



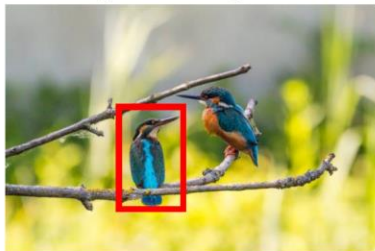
Output: Coordinates (e.g., centroid)

Object Localisation



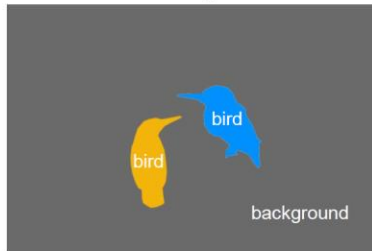
Output: Coordinates (e.g., bounding box)

Object Recognition



Output: Category (e.g., "kingfisher")

Semantic Segmentation

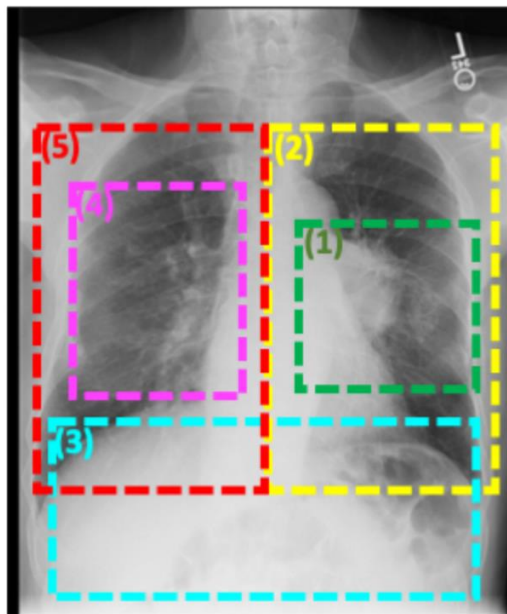


Output: Labelmap

Image Captioning



Output: Text  
(e.g., "two birds sitting on a branch")



- (1) A mass is present in the superior segment of the left lower lobe and therefore malignancy must be considered.
- (2) Elsewhere, the left lung appears clear.
- (3) There is no pleural effusion. (4) Calcified pleural plaque is present in the right mid zone. (5) The right lung appears clear.

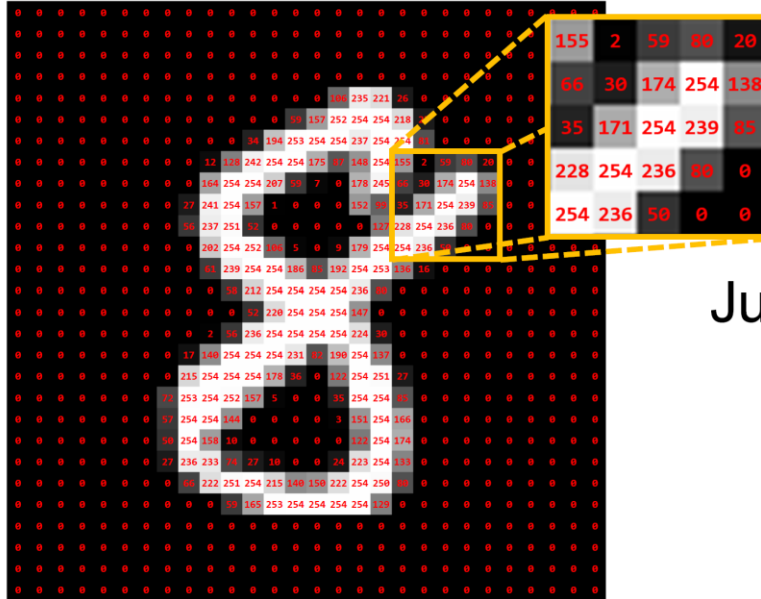
## Example: Image Classification Algorithms



Image  
Classification  
Algorithms

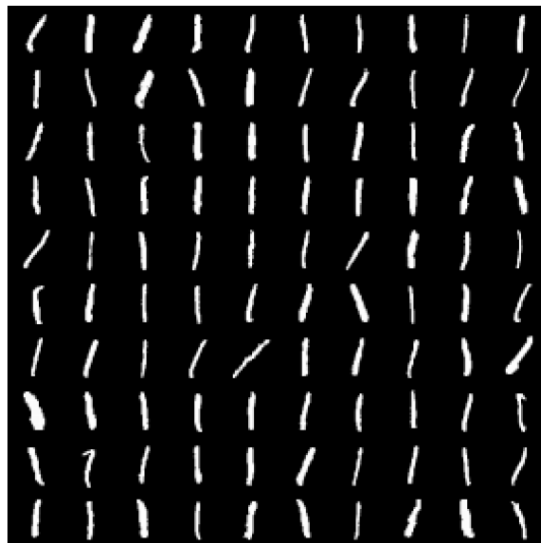


With/without  
tumour



Just an array of numbers

## Classification Example : Digit Recognition

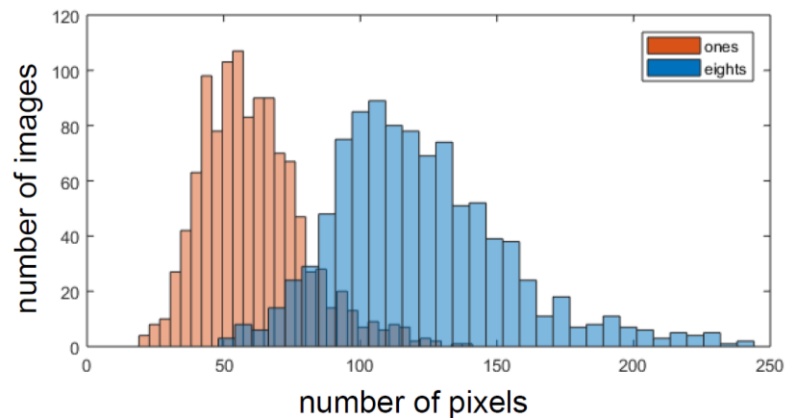


ones



eights

## Classification Example : Digit Recognition



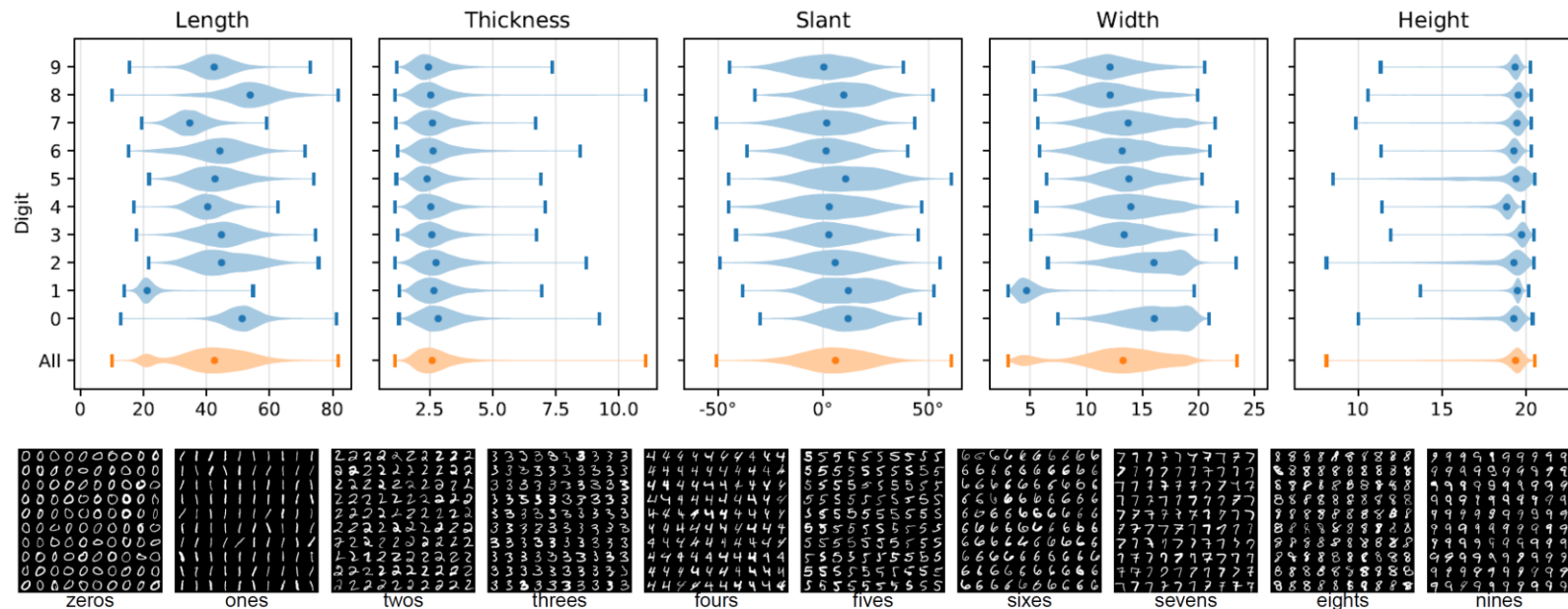
ones



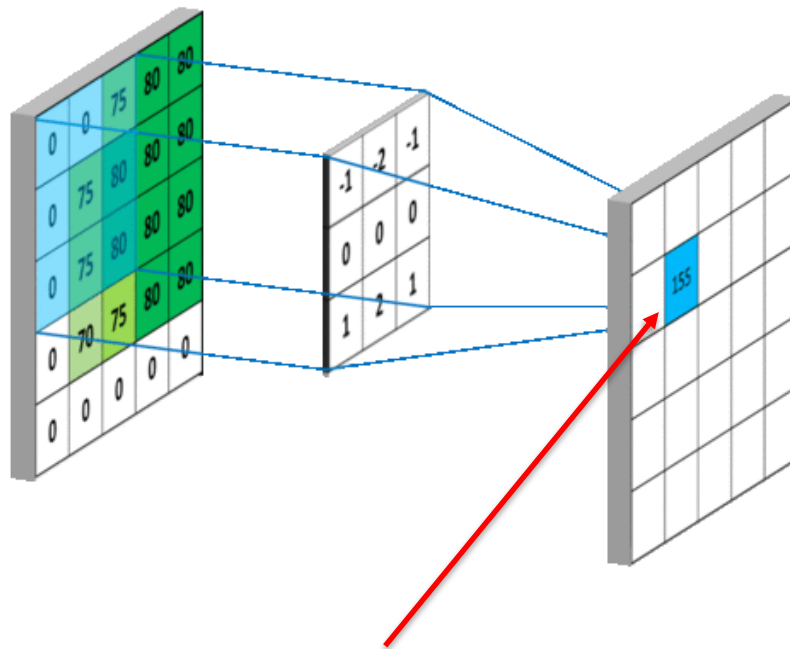
eights



# Classification Example : Digit Recognition



## Convolutional Operations in Image Processing



$$-1 * 0 + (-2) * 0 + (-1) * 75 + 0 * 0 + 0 * 75 + 0 * 80 + 1 * 0 + 2 * 75 + 1 * 80 = 155$$

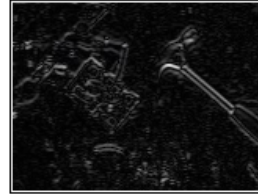
## Convolutional Operations with a Sobel Filter

$$\begin{Bmatrix} -1 & -2 & -1 \\ 0 & 0 & 0 \\ 1 & 2 & 1 \end{Bmatrix} \times$$

*Horizontal Sobel*



=



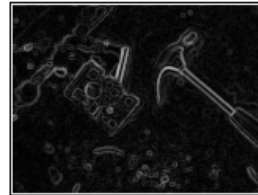
+

$$\begin{Bmatrix} -1 & 0 & 1 \\ -2 & 0 & 2 \\ -1 & 0 & 1 \end{Bmatrix} \times$$

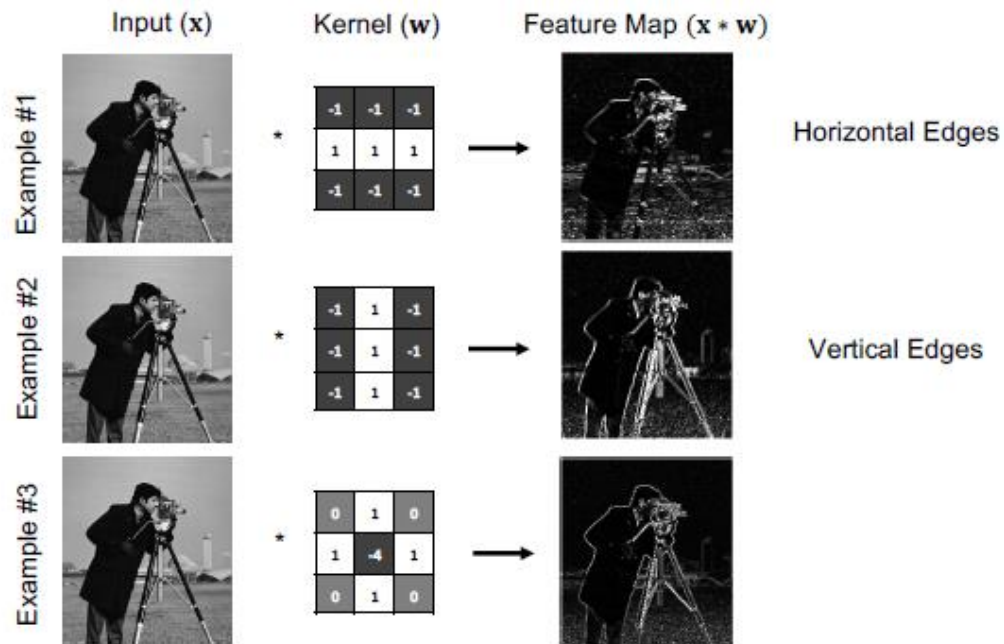
*Vertical Sobel*



=

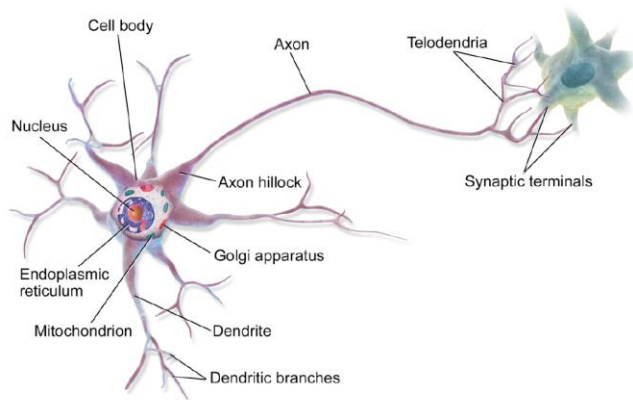


# Applying Convolutional Operations

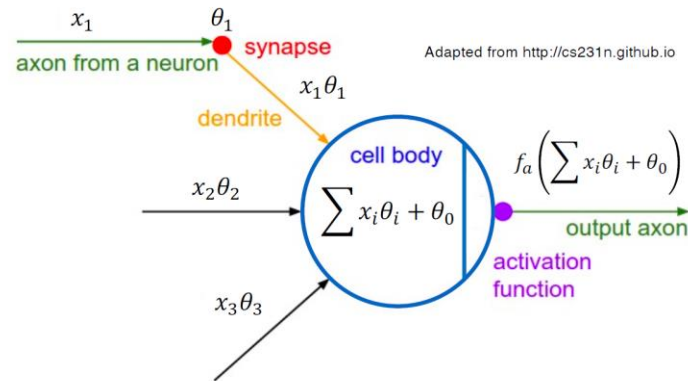


# Neural Network

- First machine learning methods were inspired by how the brain works:



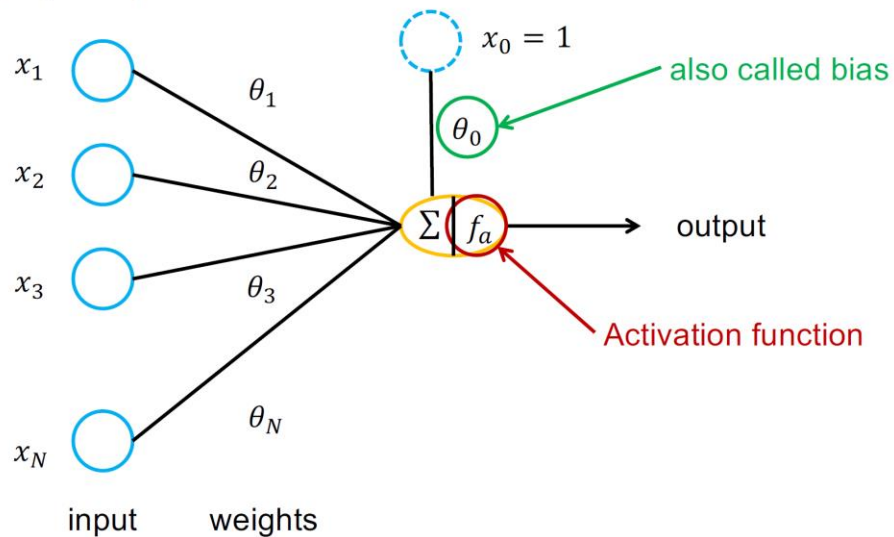
Biological neuron



Artificial neuron

# Single-layer Perceptron

- A single-layer perceptron looks as follows:

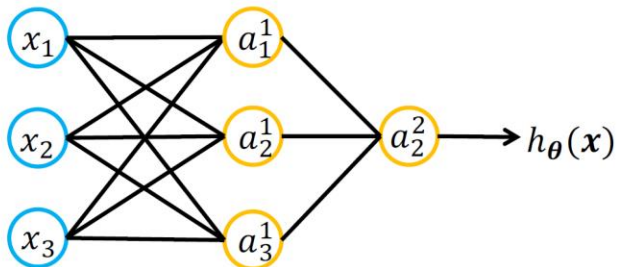


## Single-layer Perceptron

- Training a perceptron means to iteratively updating the weights associated with each of its inputs
- This allows to progressively approximate the underlying relationship in the given training dataset
- Once properly trained, it can be used to classify entirely new samples



# Neural Network



$a_j^i$  activation of neuron  $j$  in layer  $i$

$\theta^i$  matrix of weights controlling mapping from layer  $i - 1$  to  $i$

$$a_1^1 = f_a(\theta_{0,0}^1 + \theta_{1,0}^1 x_1 + \theta_{2,0}^1 x_2 + \theta_{3,0}^1 x_3)$$

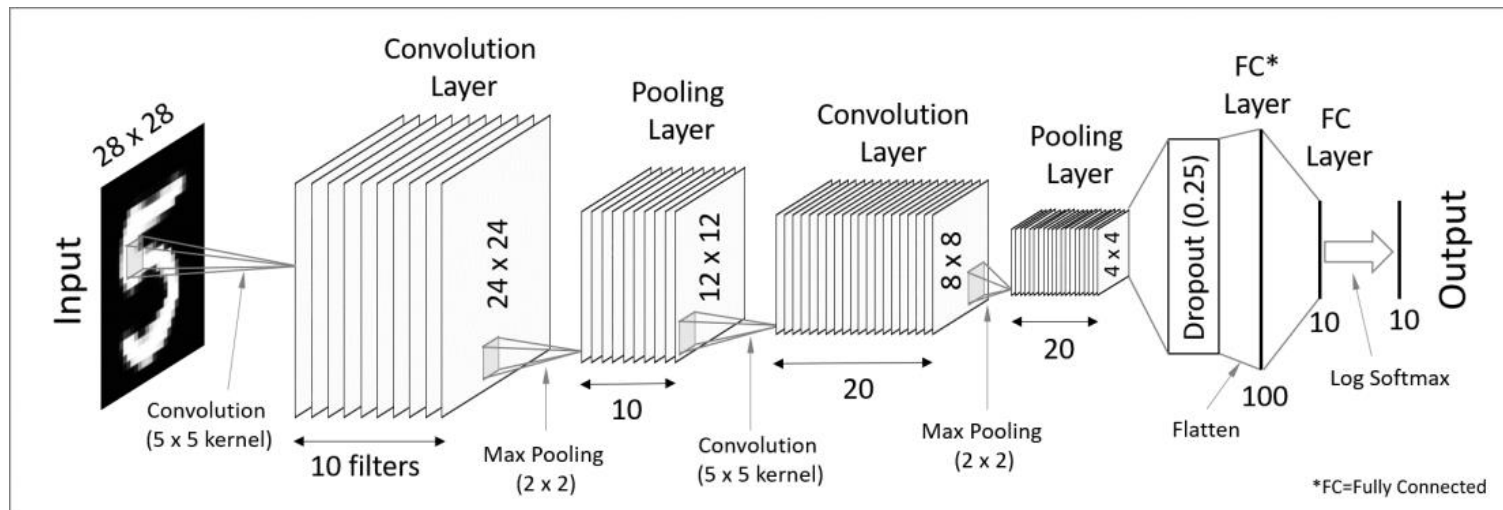
$$a_2^1 = f_a(\theta_{0,1}^1 + \theta_{1,1}^1 x_1 + \theta_{2,1}^1 x_2 + \theta_{3,1}^1 x_3)$$

$$a_3^1 = f_a(\theta_{0,2}^1 + \theta_{1,2}^1 x_1 + \theta_{2,2}^1 x_2 + \theta_{3,2}^1 x_3)$$

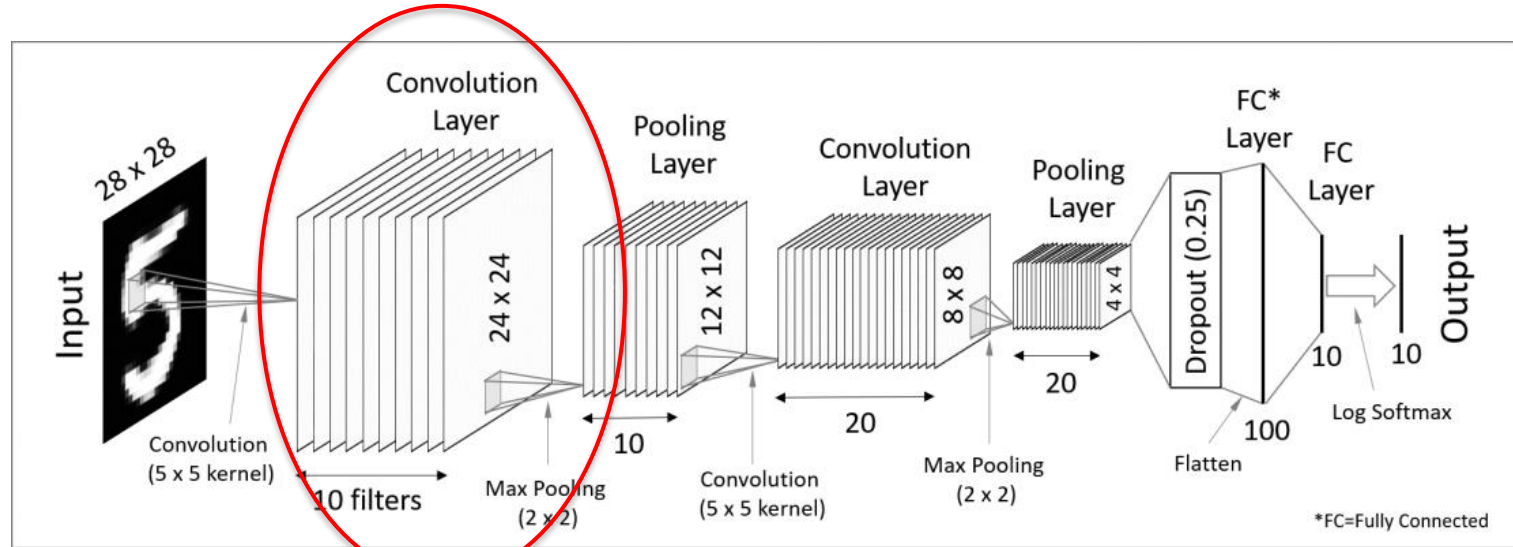
$$h_{\theta} = a_1^2 = f_a(\theta_{0,0}^2 + \theta_{1,0}^2 a_1 + \theta_{2,0}^2 a_2 + \theta_{3,0}^2 a_3)$$



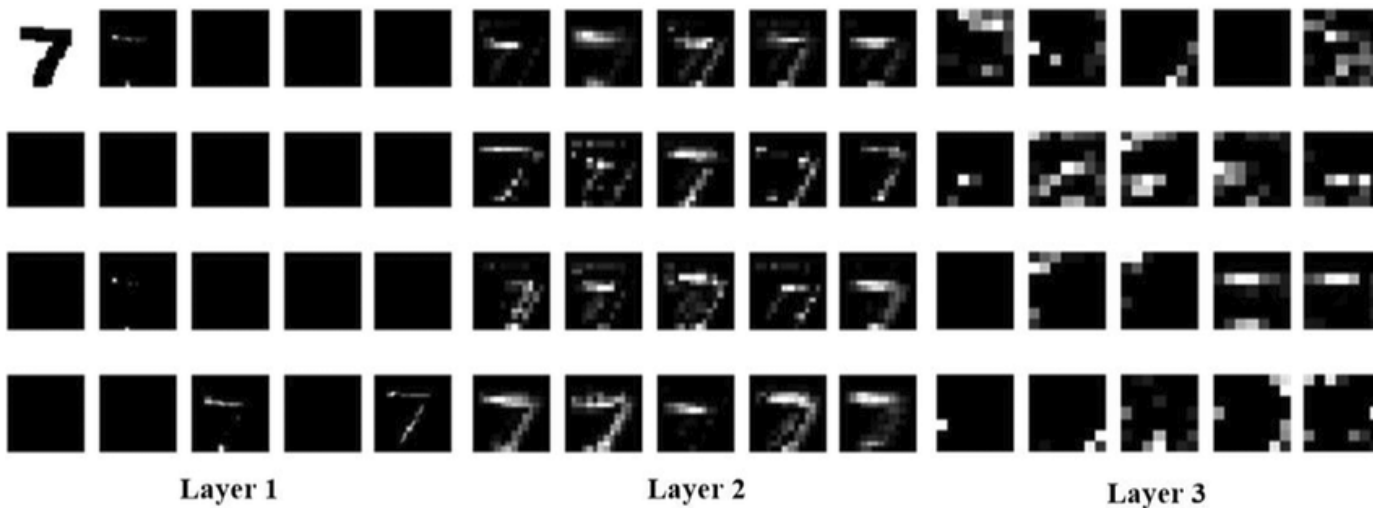
# Convolutional Neural Network (CNN) Model



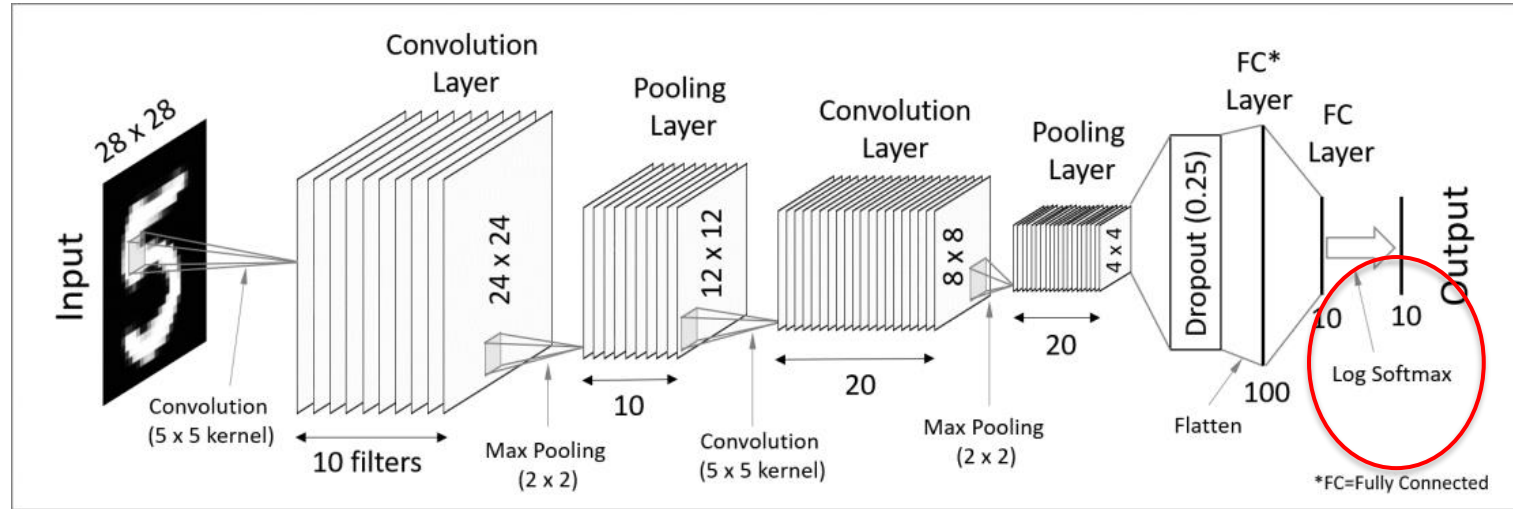
# Convolution Layer



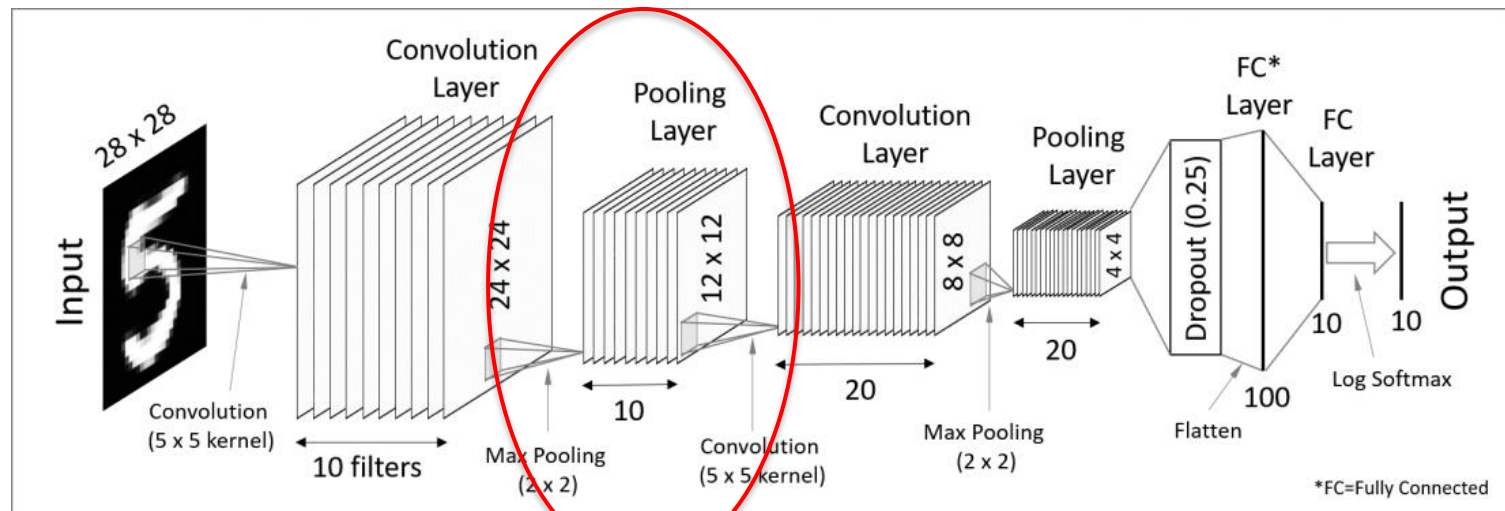
## Feature maps



# Activation Functions

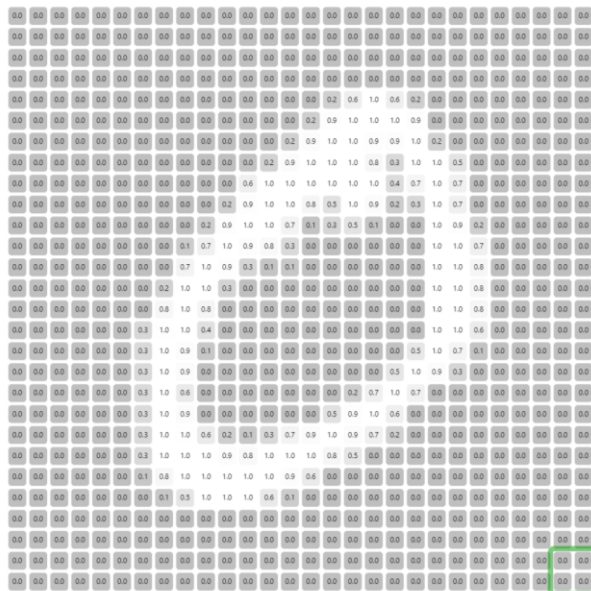


# Pooling Layer

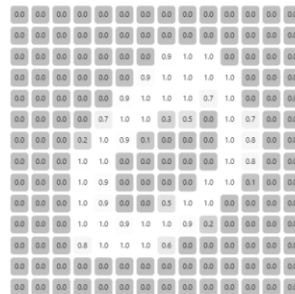


# Max Pooling

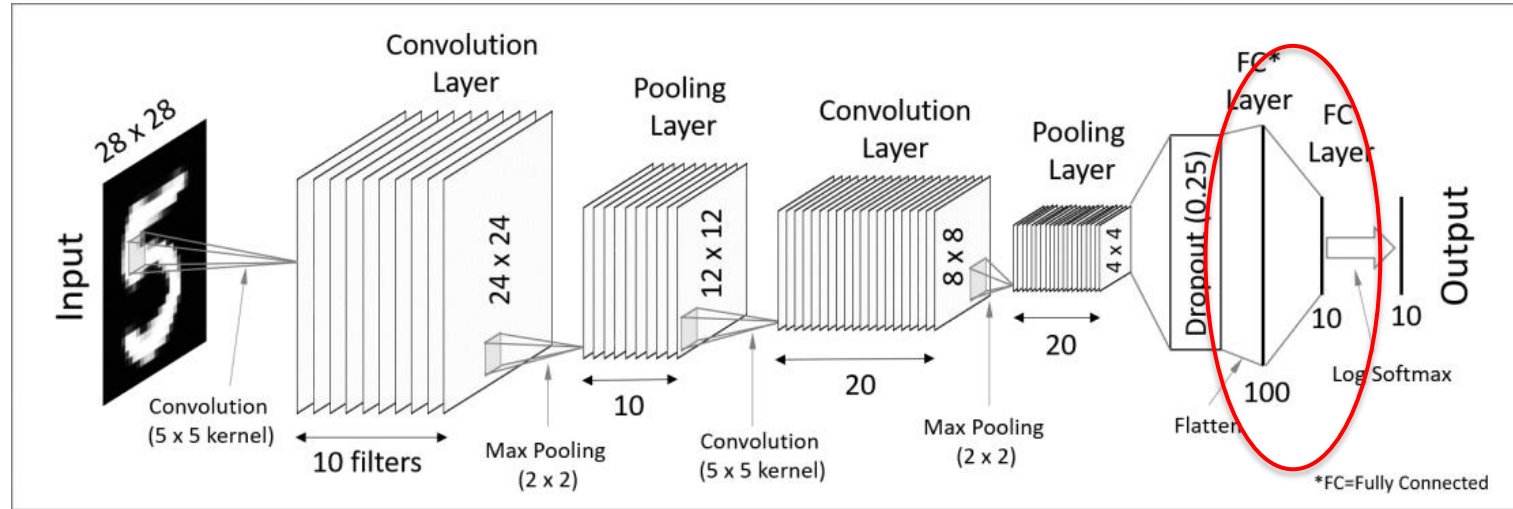
Input



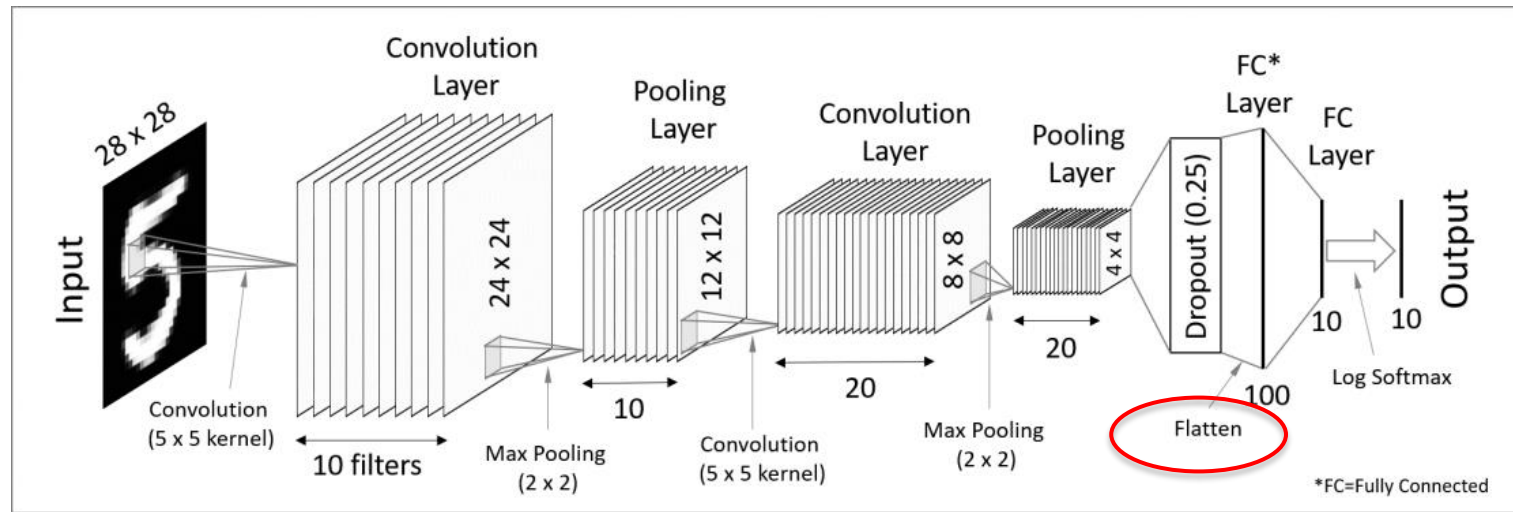
Output



# Fully Connected Layer

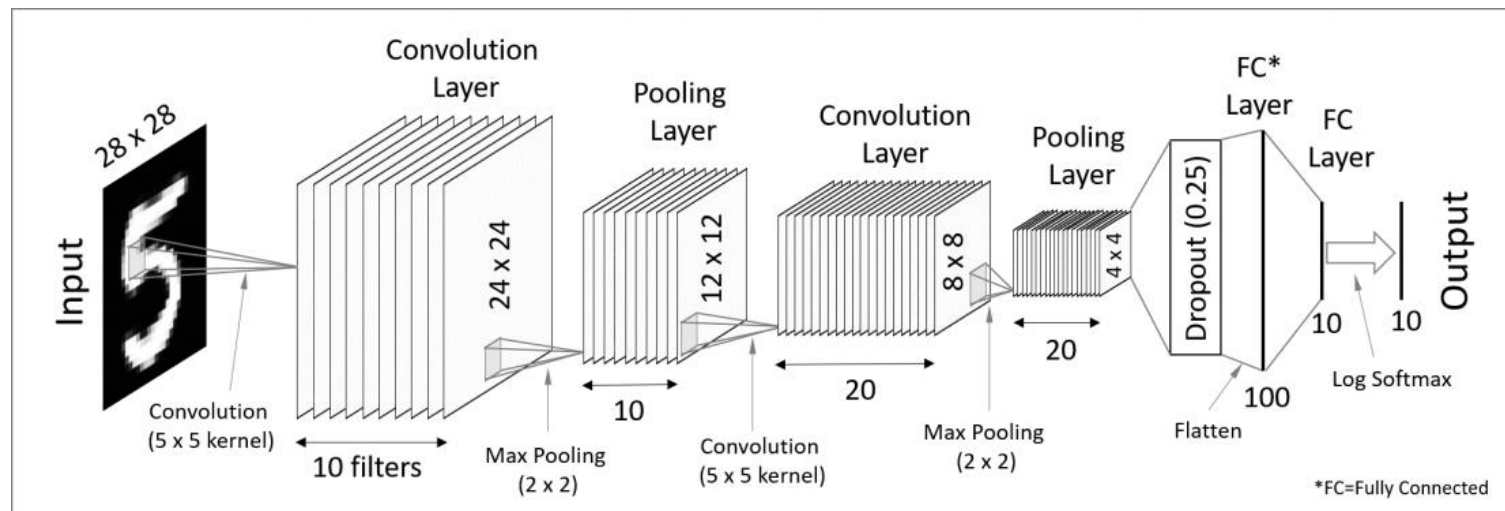


# Flattening

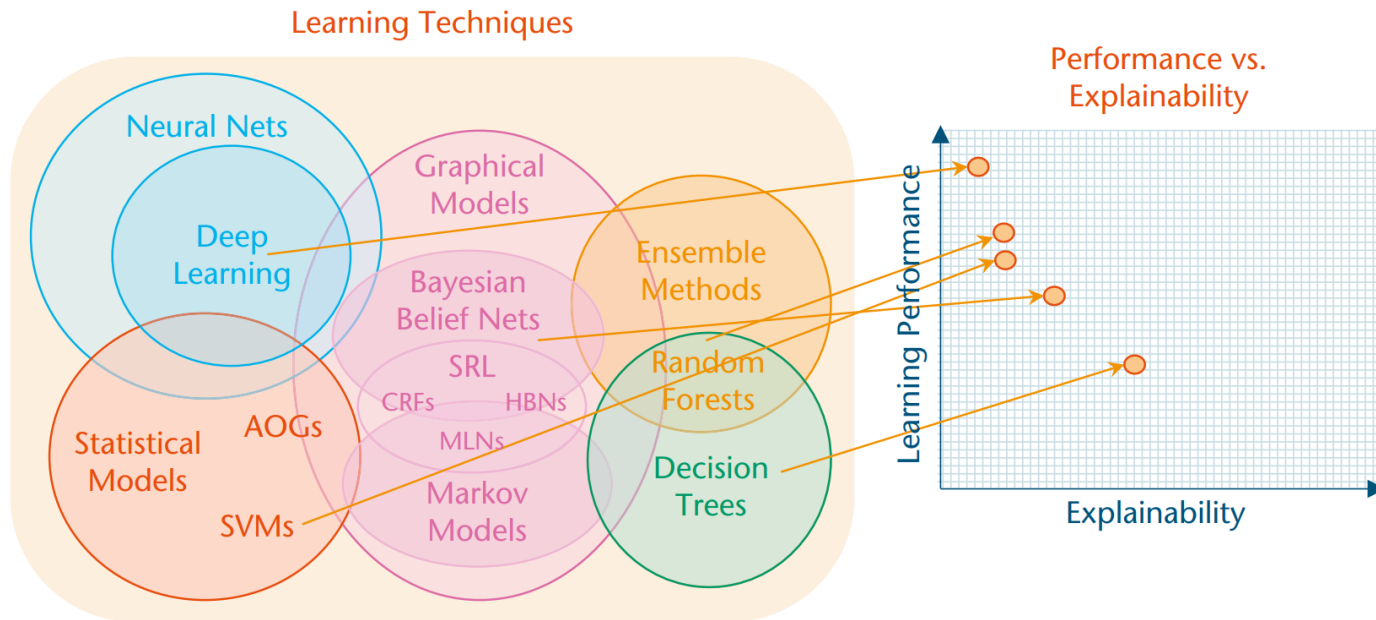




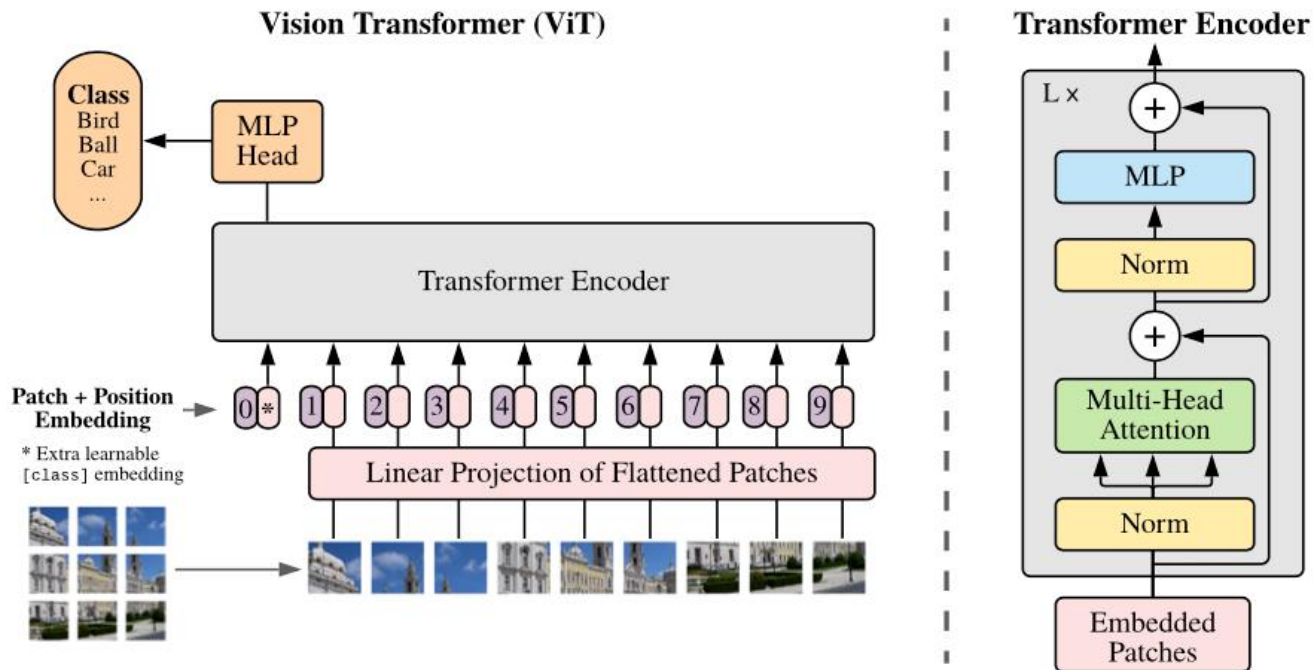
# Dropout Layer



# Performance vs Explainability



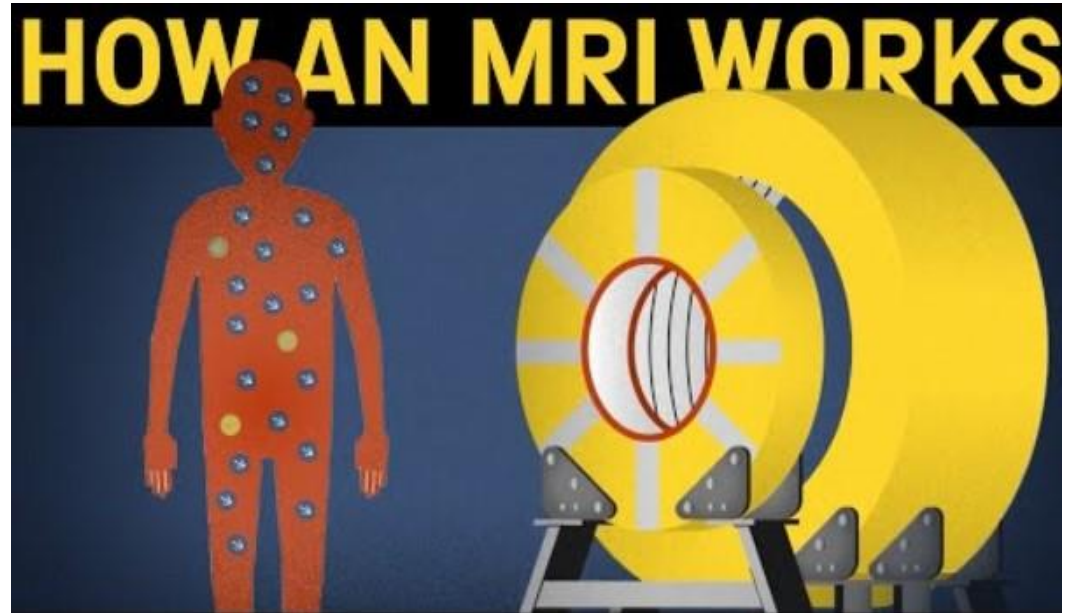
# Vision Transformer (ViT)



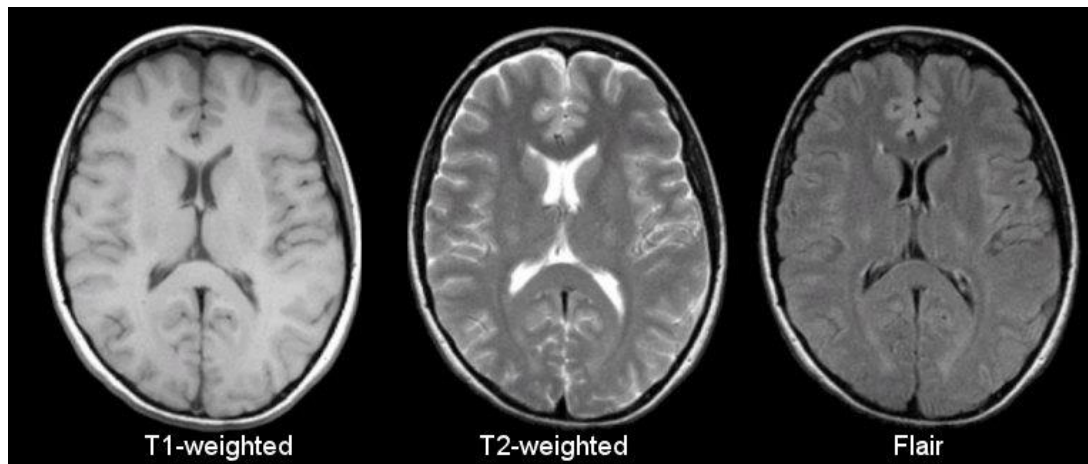
## Project Workflow

- Understanding and Preprocessing Your Dataset
  - Building and Training the Model
  - Optimising Hyperparameters
  - Evaluating Performance
  - Testing the Model
-

- Non-invasive imaging technology
- Providing three dimensional detailed anatomical images
- Used for disease detection, diagnosis, and treatment monitoring

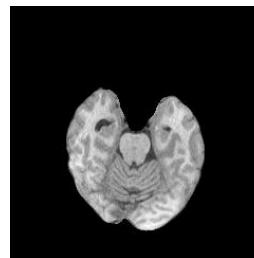


## Different Sequences of Brain MR Images

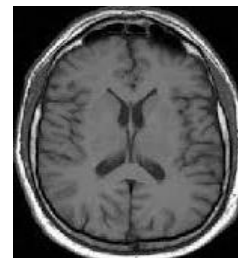


Different MRI sequences

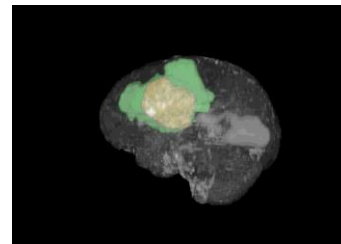
- 210 MRI volumes will be given to you for training and evaluating the segmentation model.



Brain image with  
tumour



Brain image w/o  
tumour



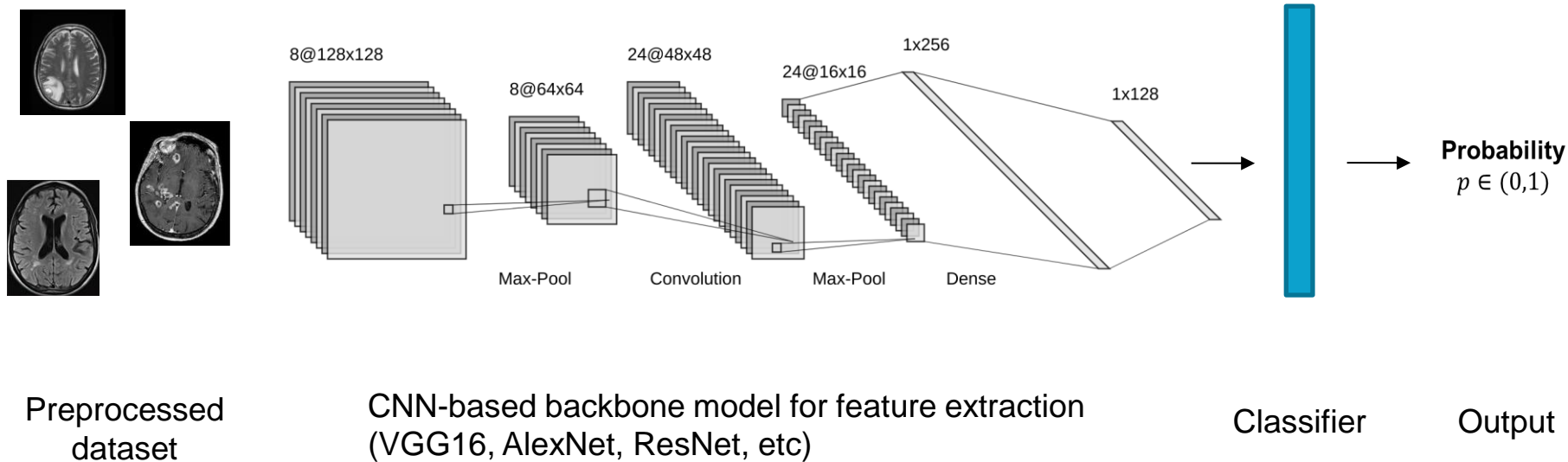
Brain tumour in 3D

## Data preprocessing

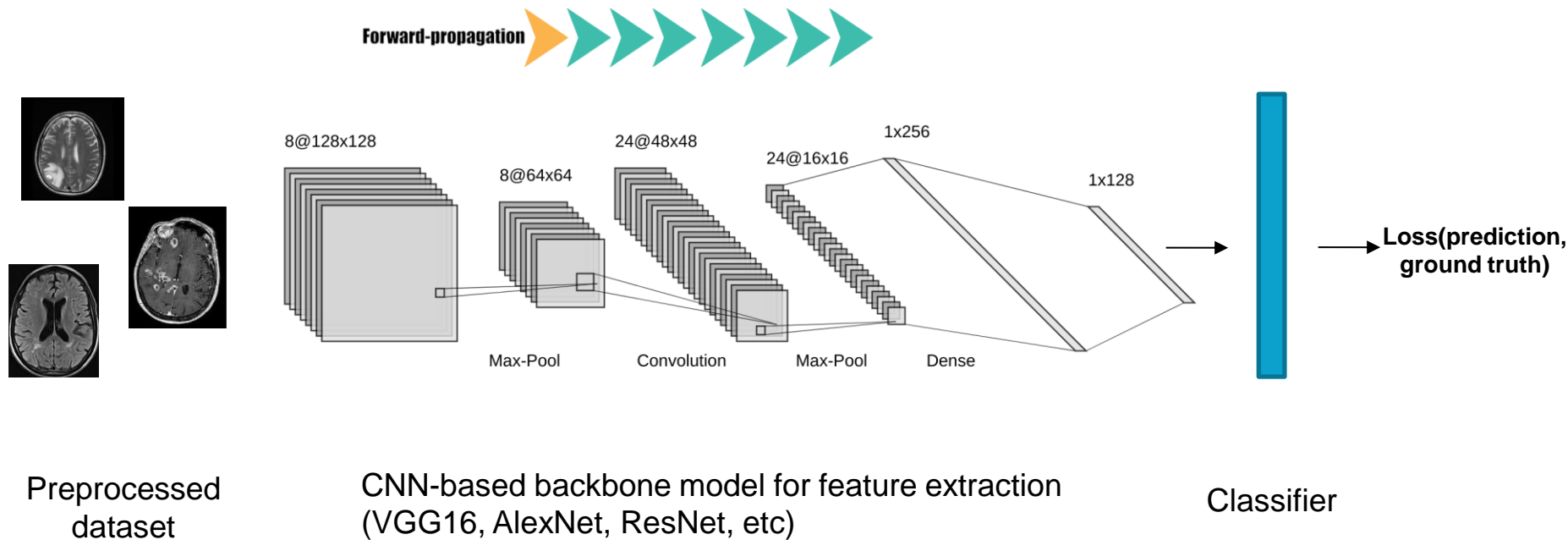
- Denoising
  - Image Standardisation (Resizing , Normalisation, etc.)
  - Data Augmentation (Rotation, Shearing, Scaling, etc.)
-



# Building the Classification Model



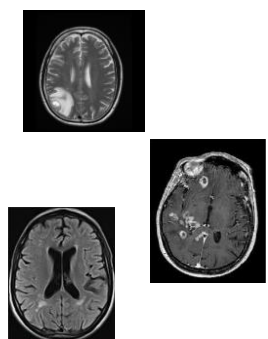
# Training the Classification Model



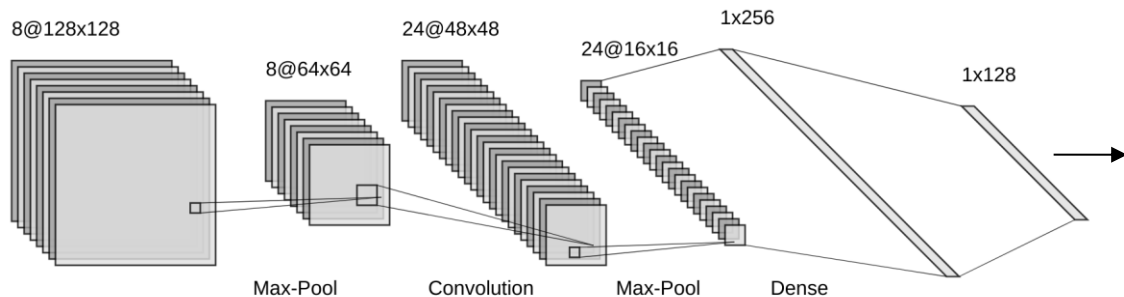
## Improving Prediction Accuracy

- Choose the right data augmentation methods;
- Find the optimal backbone model;
- Choose to finetune the model or train it from the scratch;
- Set the number of epochs to train the model and early stopping criteria;
- Tune other hyperparameters to achieve the best performance.

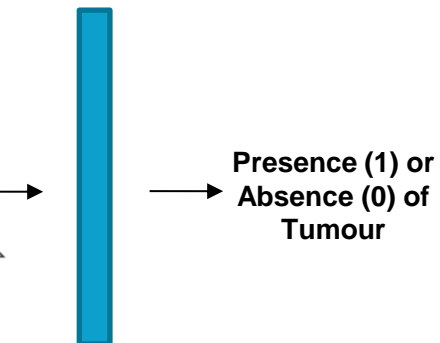
# Test/Evaluate the Classification Model



Preprocessed  
dataset



CNN-based backbone model for feature extraction  
(VGG16, AlexNet, ResNet, etc)



Classifier

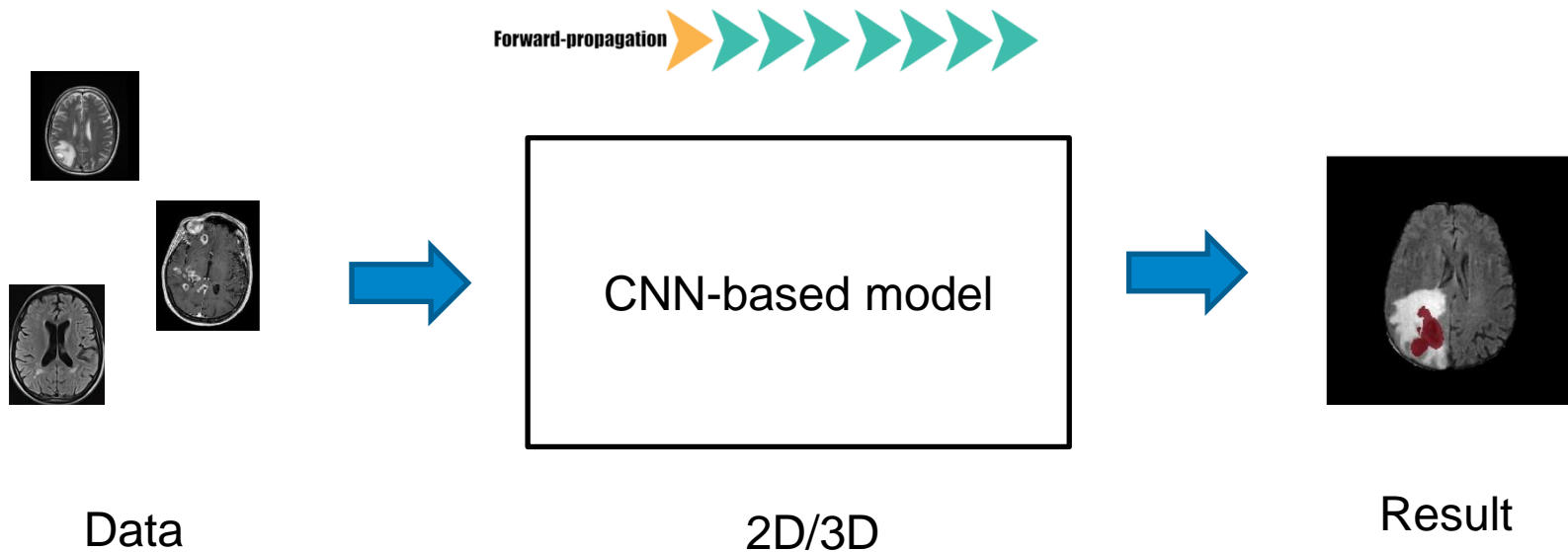
Result

## Result Evaluation

Accuracy is used to evaluate the performance of your classification model, which is defined as

$$\text{Accuracy} = \frac{\text{Number of correctly classified images}}{\text{Number of total images}} \times 100\%$$

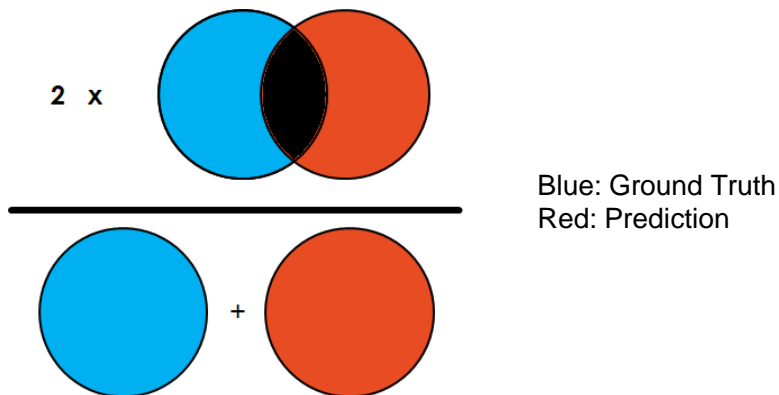
## Building/Training Segmentation Model



# Result evaluation

Dice score and 95% Hausdorff Distance are used to evaluate the performance of your segmentation model. Dice score is defined as

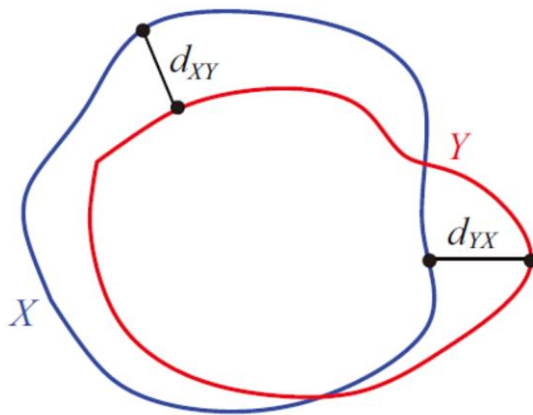
$$\text{Dice score} = (2 * \text{Area of Overlap}) / (\text{total pixels combined})$$



## Result evaluation

The maximum Hausdorff Distance is defined as:

$$d_H(X, Y) = \max\{d_{XY}, d_{YX}\} = \max\left\{\max_{x \in X} \min_{y \in Y} d(x, y), \max_{y \in Y} \min_{x \in X} d(x, y)\right\}$$



95% Hausdorff Distance it is based on the calculation of the 95th percentile of the distances between boundary points in X and Y.



# Choice Considerations

- Model architecture: depth, width, scales, residuals,...
- Loss function: (weighted) cross-entropy, IoU, Dice,...
- Sampling strategy: equally per class, fore/background, uniform,...
- Optimization: optimizer, learning rate, momentum, regularization,...
- Data normalization/standardisation: z-score, bias field correction, histogram matching,...
- Post-processing: conditional random fields (CRF), smoothing,...

# Setting Up Python Development Environment

Anaconda and Tensorflow Installation (2020.02)

<https://www.h2kinfosys.com/blog/how-to-download-and-setup-tensorflow-with-anaconda/>

<https://repo.anaconda.com/archive/>

For using Nvidia GPU

<https://saturncloud.io/blog/how-to-install-tensorflow-with-anaconda-on-windows/>

Key Packages:

numpy>=1.19.2

tensorflow>=2.4.0

matplotlib>=3.3.2

scikit-learn>=0.24.0

tqdm>=4.50.2