Deep Learning and Convolutional Neural Network (42028)

Week 2 Lecture:

Machine Learning Basics

Introduction to Image Processing

Type of Machine Learning Systems

Type of Machine Learning Systems

Supervised Learning
Unsupervised Learning
Semi-supervised Learning
Reinforcement Learning

Depending on whether the system is trained with human supervision

Whether System can learn on the fly

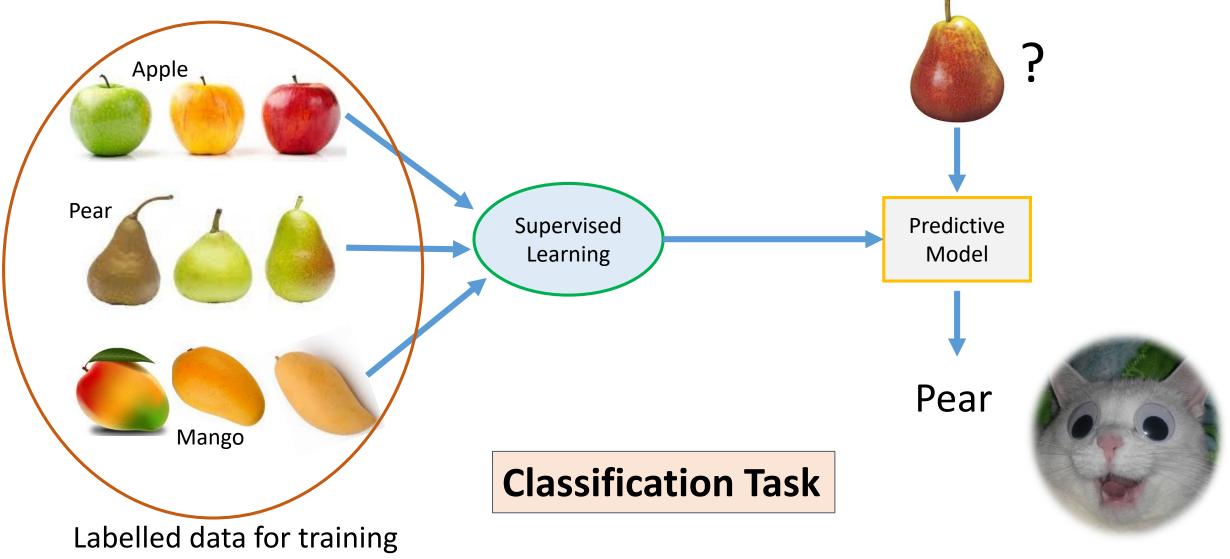
Batch and Online Learning

Instance-based and Model-based Learning

Comparing data points or detect patterns in training data to build a predictive model

Supervised Learning

(Object + Desired Output Label)



Supervised Learning

House Price prediction

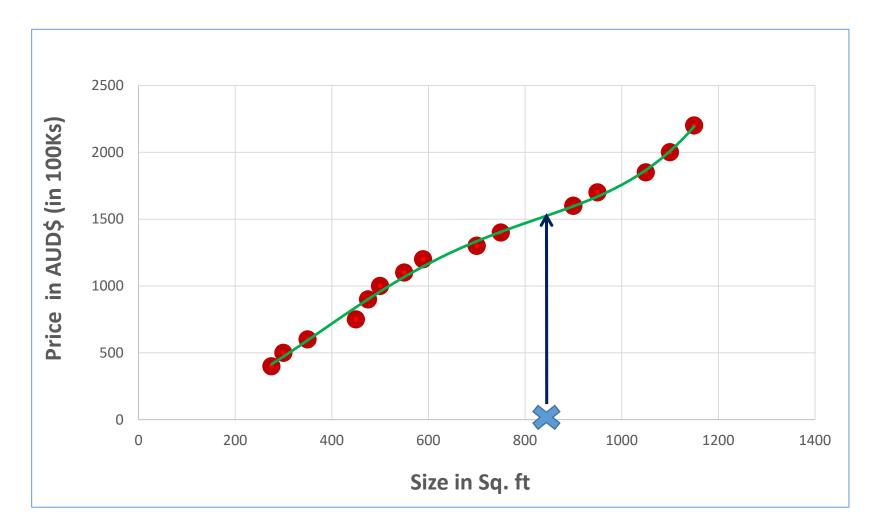
Feature:

Size of the house

To Predict:

- Price of the house

Regression Task

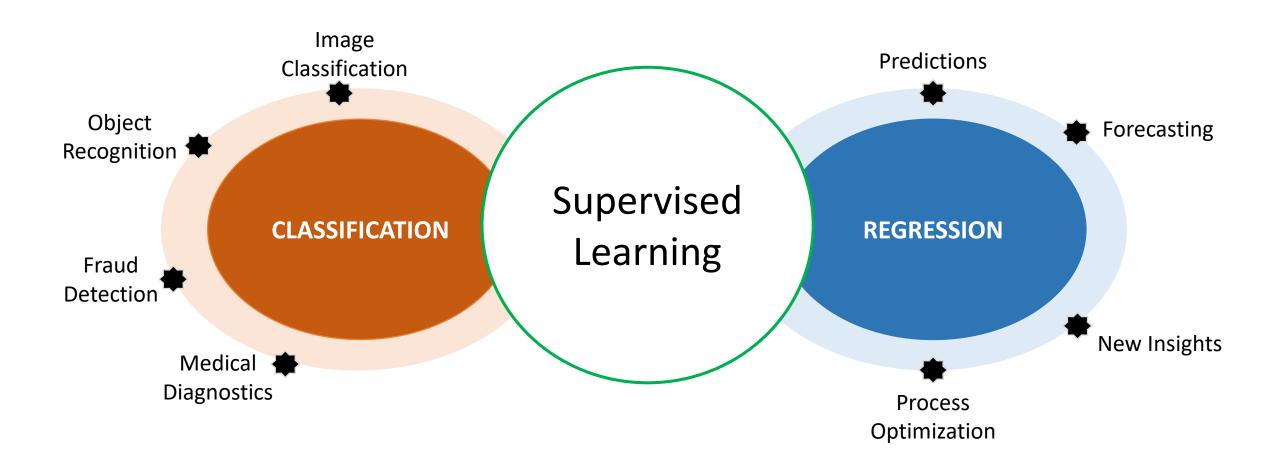


Supervised Learning

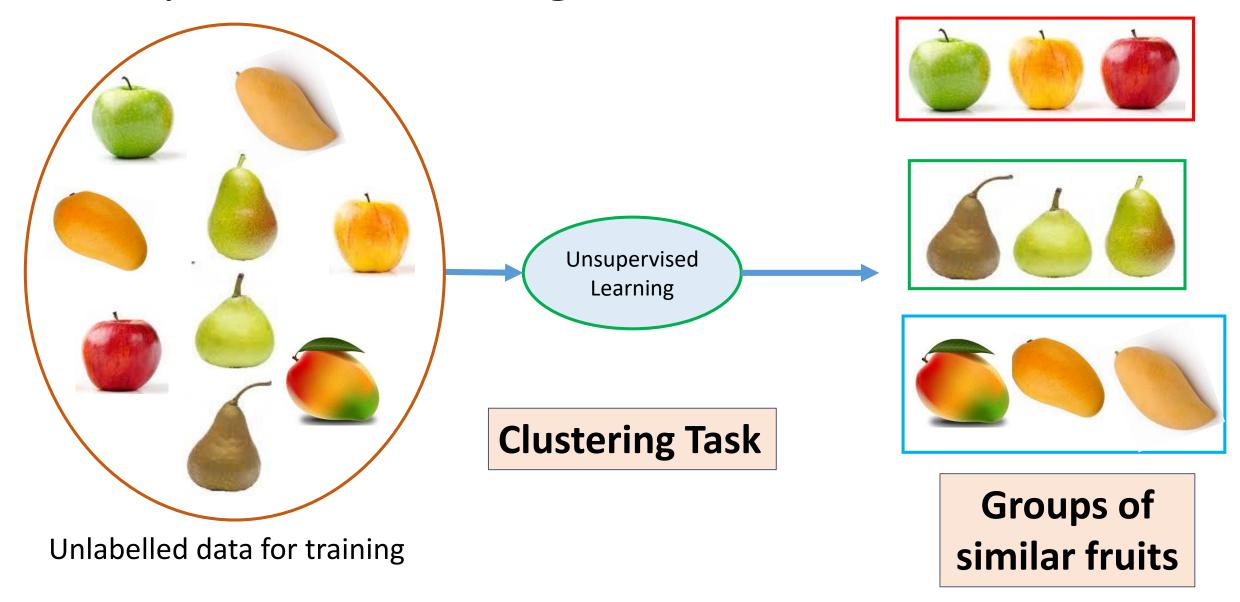
Important Algorithms:

- K-Nearest Neighbours
- Logistic regression
- Support Vector Machines (SVMs)
- Neural Networks (*some of them can be unsupervised)

Supervised Learning Examples



Unsupervised Learning



Unsupervised Learning

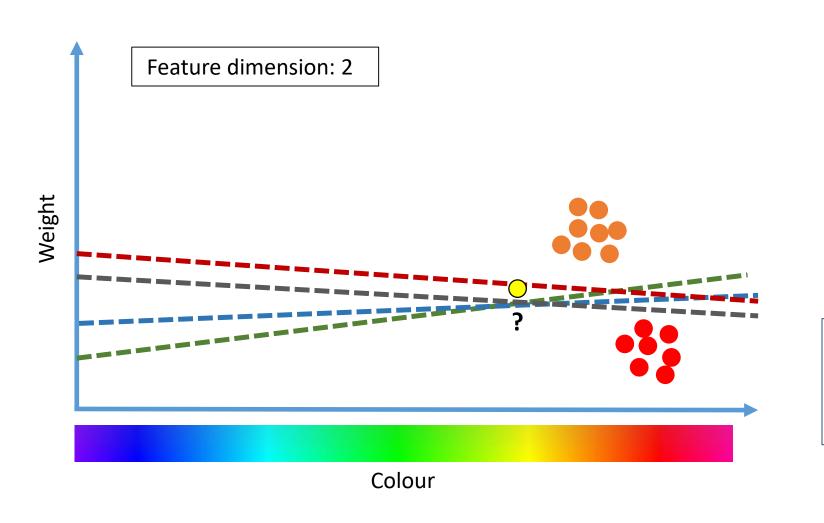
Important Algorithms:

- k-means
- Expectation Maximization

- A Support Vector Machine is a very powerful and versatile Machine Learning model, capable of performing linear or non-linear classification, regression, and also outlier detection.
- Defined by a separating hyperplane
- Suitable for small or medium sized datasets

Reference and Pre-Reading:

Theory: https://medium.com/machine-learning-101/chapter-2-svm-support-vector-machine-coding-edd8f1cf8f2d
Implementation: https://medium.com/machine-learning-101/chapter-2-svm-support-vector-machine-coding-edd8f1cf8f2d



Orange

Apple

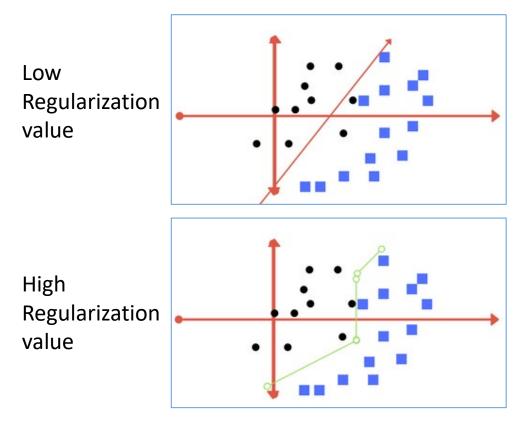
SVM finds the best line or hyper-plane which will fairly separates the classes

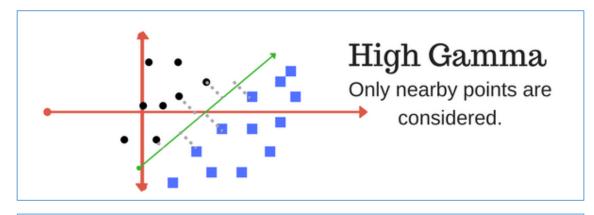
Example: Using sklearn for SVM classification (Partial code snippet)

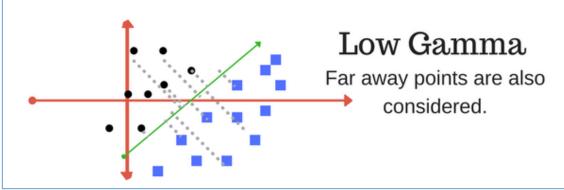
```
import numpy as np
import matplotlib.pyplot as plt
from sklearn import svm, datasets
# import some data to play with
iris = datasets.load iris()
# Take the first two features. We could avoid this by using a two-dim dataset
X = iris.data[:, :2]
v = iris.target
# we create an instance of SVM and fit out data. We do not scale our
# data since we want to plot the support vectors
C = 1.0 # SVM regularization parameter
models = (svm.SVC(kernel='linear', C=C),
          svm.LinearSVC(C=C),
          svm.SVC(kernel='rbf', gamma=0.7, C=C),
          svm.SVC(kernel='poly', degree=3, C=C))
models = (clf.fit(X, y) for clf in models)
```

Reference: https://en.wikipedia.org/wiki/Iris flower data set

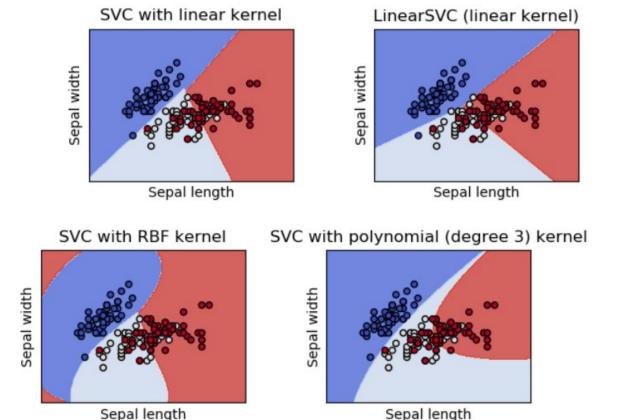
SVM Parameters: Kernel, Gamma, Regularization (C)







Example: Using sklearn for SVM classification



Iris flower data set



Reference: https://scikit-learn.org/stable/auto examples/svm/plot iris svc.html https://en.wikipedia.org/wiki/Iris flower data set

K-Nearest Neighbour (KNN)

• A simple supervised learning algorithm.

Can be used for both classification and regression

 Non-parametric: doesn't make any assumption on the data distribution

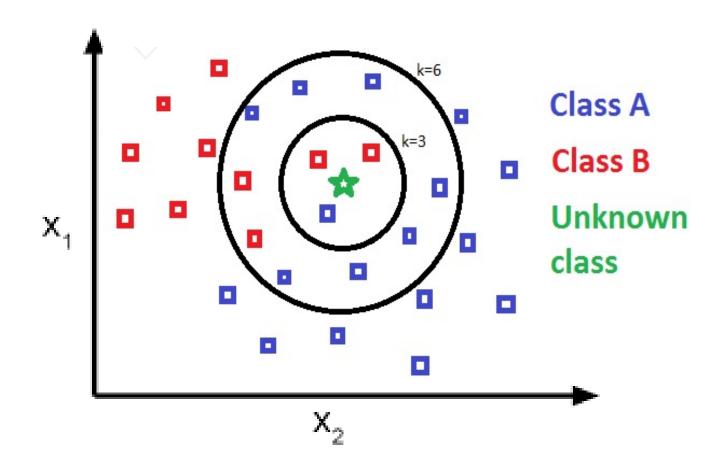
Training data is retained to make future predictions

K-Nearest Neighbour (KNN)

How does it work?

- Computes distance between the new sample and all training samples
- Distance measure: Euclidean, Manhattan etc.
- Picks 'k' entries in the training set which are closest to the new sample
- Majority voting decides the class of the new sample

K-Nearest Neighbour (KNN)



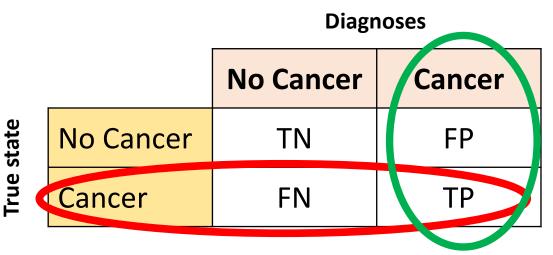
Evaluation Metrics

Precision & Recall

What are the "correct" cells?

- *TN*: (Number of True Negatives), i.e., patients who did *not* have cancer whom we correctly diagnosed as *not* having cancer.
- TP: (Number of True Positives), i.e., patients who did have cancer whom we correctly diagnosed as having cancer

Precision: TP/Cancer Diagnoses



Recall: TP/Cancer True States

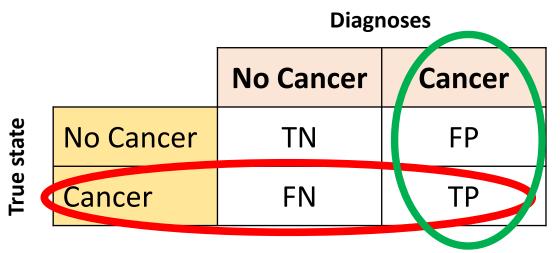
Evaluation Metrics

Precision & Recall

what are the "error" cells are:

- FN: (Number of False Negatives), i.e., patients who did have cancer whom we incorrectly diagnosed as *not* having cancer
- FP: (Number of False Positives), i.e., patients who did not have cancer whom we incorrectly diagnosed as having cancer

Precision: TP/Cancer Diagnoses



Recall: TP/Cancer True States

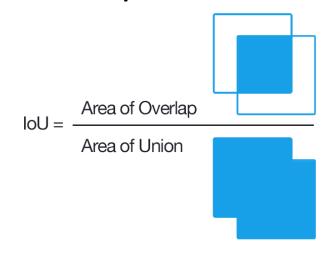
Precision=
$$(TP)/(TP+FP)$$

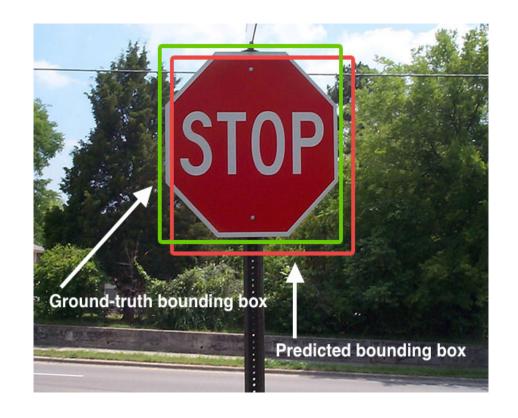
Recall =
$$(TP)/(TP+FN)$$

Evaluation Metrics

Intersection over Union (IoU):

Intersection over Union is a metric used for the evaluation of an object detector, i.e. how good is the predicted bounding box for an object detected closely matches





What is a digital image?

- Digital images are made of picture elements called Pixels.
- It is an array, or a matrix of *Pixels* arranges in *columns* and *rows*.
- Each *Pixel* has its own *intensity* value, or *brightness*

- Intensity values in digital images are defined by bits
- For a standard 8 bits image, a pixel can have $2^8 = 256 (0 255)$ values.
- Black & White images have a single 8-bits intensity range.

Image dimension = 5×5 f(2, 3) = 170 (Pixel/intensity value)

Hence, an image may be defined as a 2D function f(x, y), where, x and y are spatial co-ordinates, and the amplitude of f at (x, y) is the intensity or Gray level of the image at that point/pixel.

170	170	55	170	170
170	55	170	55	170
170	55	170	55	170
55	140	140	140	55
55	170	170	170	55

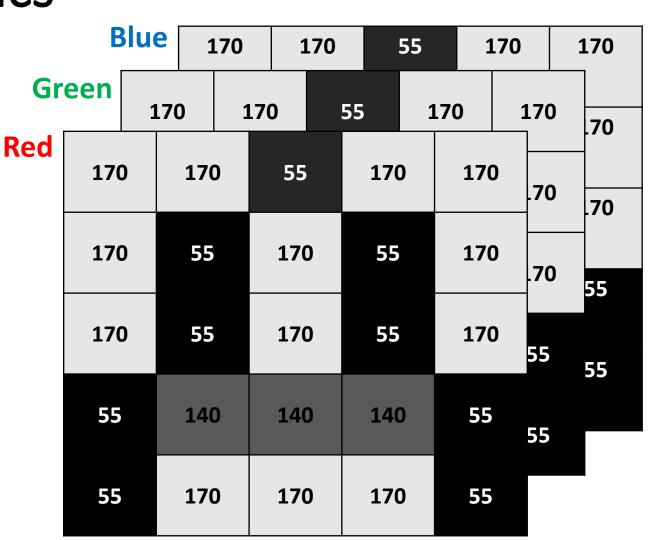
5 X 5 Gray scale image (8 bit)

Image dimension = 5 X 5 X 3 No. of Channels = 3

Since, RGB image contains 3 X 8-bits of intensities, they are referred to as 24-bit colour images.

So, 24-bit colour depth

- = 8 X 8 X 8 bits
- = 256 X 256 X 256 colours
- = ~16 million colours



5 X 5 X 3 colour image (24 bit)

Image Thresholding

- Easiest method for image segmentation!
- Converts gray-scale image into a binary image If f(x,y) > Threshold, then f(x,y) = 0 else f(x,y) = 255

Binary Image (8-bit) has only two possible values of pixel intensity (0 and 1, or B & W)

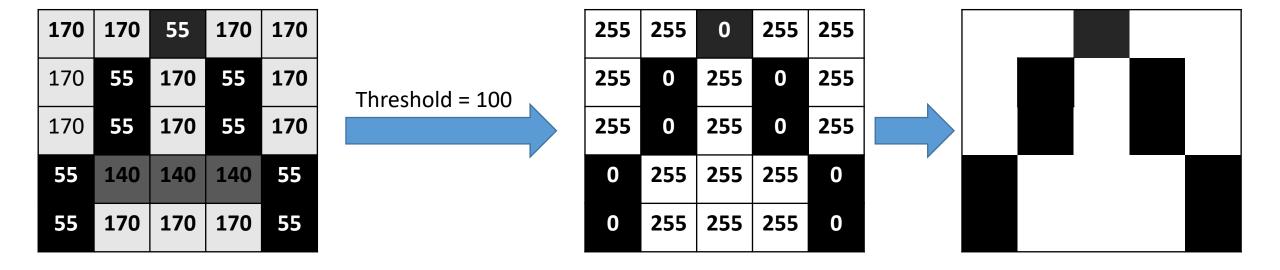


Image Thresholding



Thresholding

Original Image

Binary Image

Image Thresholding methods

- Histogram shape:

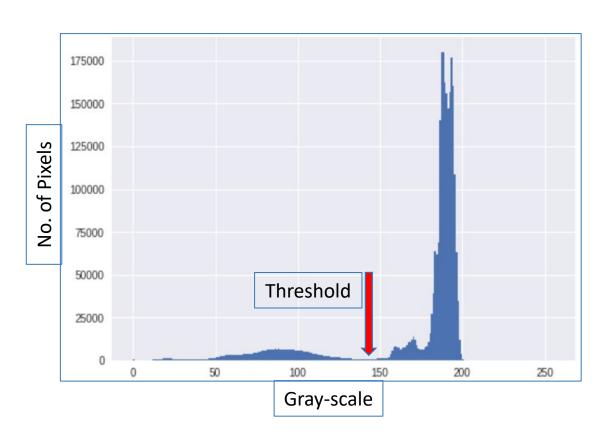
Peaks, valleys and curvature of the histogram are analysed.

- Clustering based:

The ¹Otsu method, very good for bimodal distribution

- Adaptive thresholding:

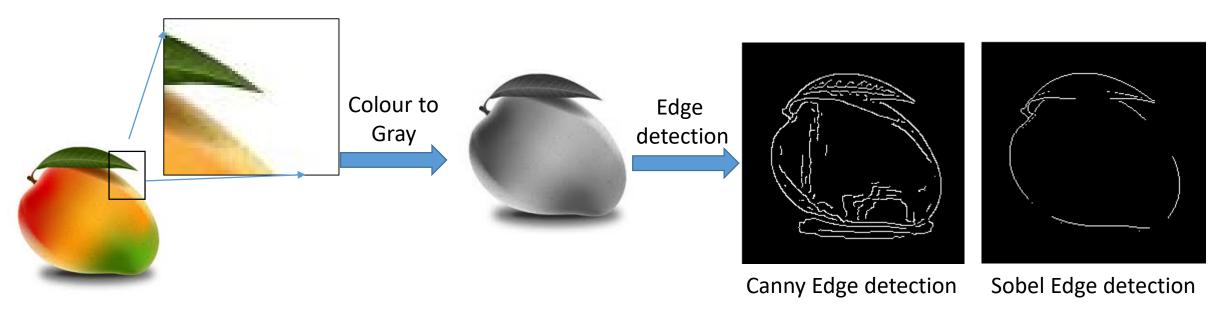
Instead of a single threshold, have thresholds for different regions in the image



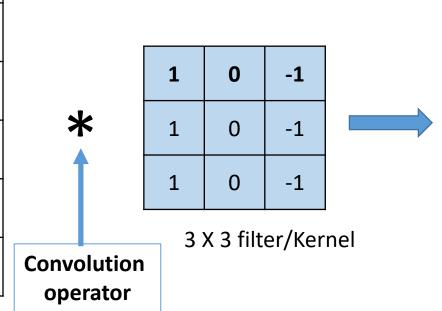
Edge Detection

What is an edge?

- The points/pixels in an image where brightness/intensities changes sharply
- A simple and fundamental tools in image processing and computer vision, useful in feature detection/extraction

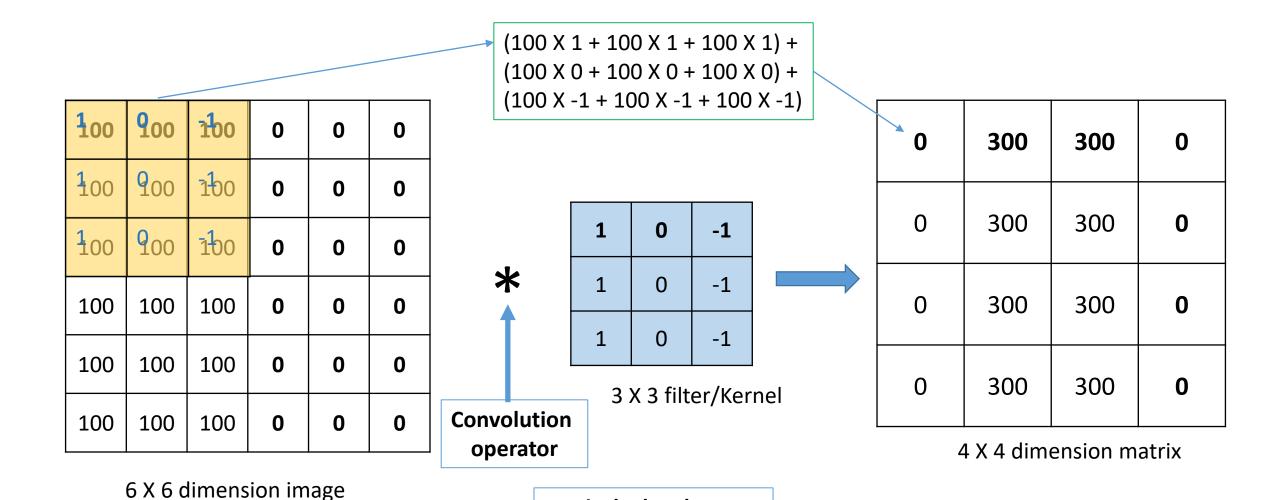


100	100	100	0	0	0
100	100	100	0	0	0
100	100	100	0	0	0
100	100	100	0	0	0
100	100	100	0	0	0
100	100	100	0	0	0

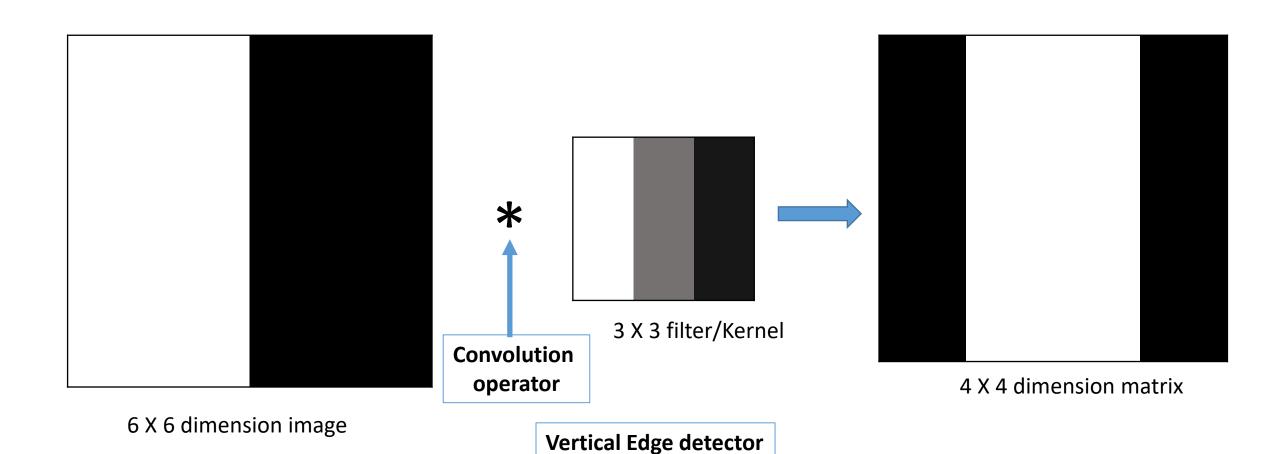


	0	300	300	0
	0	300	300	0
•	0	300	300	0
	0	300	300	0

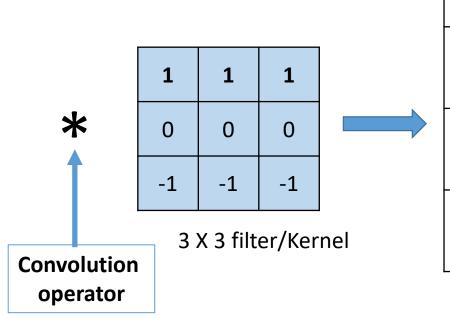
4 X 4 dimension matrix



Vertical Edge detector



100	100	100	100	100	100
100	100	100	100	100	100
100	100	100	100	100	100
0	0	0	0	0	0
0	0	0	0	0	0
0	0	0	0	0	0

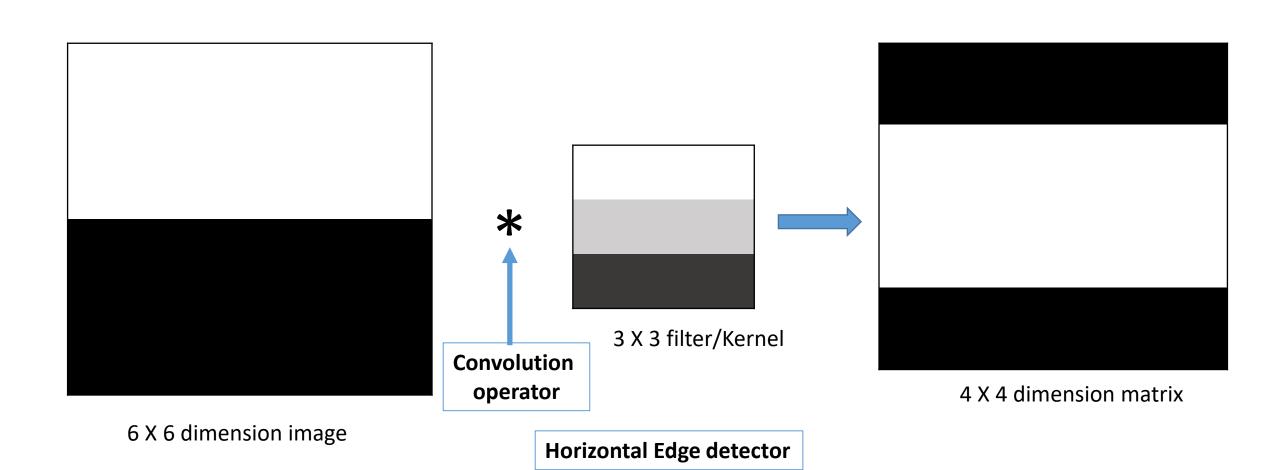


0	0	0	0
300	300	300	300
300	300	300	300
0	0	0	0

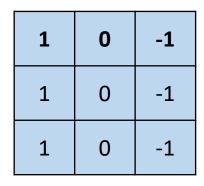
4 X 4 dimension matrix

6 X 6 dimension image

Horizontal Edge detector



Sobel edge detection



3 X 3 filter/Kernel For Vertical edges

1	1	1
0	0	0
-1	-1	-1

3 X 3 filter/Kernel For Horizontal edges

Prewitt Filters

1	2	1
0	0	0
-1	-2	-1

3 X 3 filter/Kernel For Horizontal edges

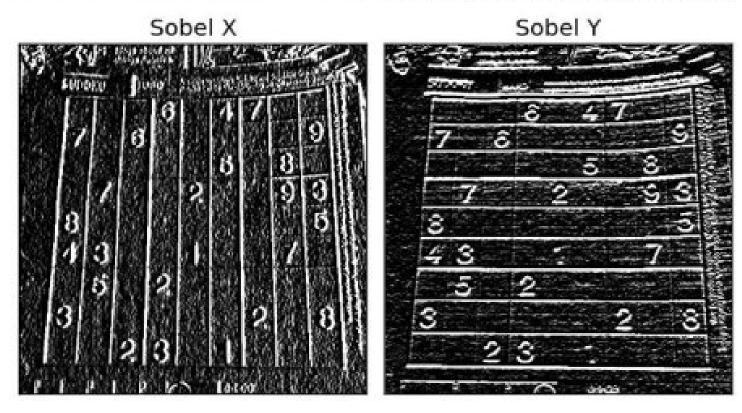
1	0	-1
2	0	-2
1	0	-1

3 X 3 filter/Kernel For Vertical edges

Sobel Filters

Sobel edge detection





Convolutions in CNN

 Convolutions are very important operation in a Convolutional Neural Networks (CNN)

 Filters weights are not fixed, but learned during the training operations of a CNN for a specific task!

Multiple filters are used in CNNs