

# Detection & Tests

## IML

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ENIB

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- 1 Classification
  - KNN
- 2 Clustering
  - k-means
  - Hierarchical clustering
  - Distance
- 3 Tests
  - Training vs Testing
  - K-Fold Cross Validation
  - Model performance
- 4 Feature extraction

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## Model : KNN

### Examples

- ▷ predict how I'm going to vote in the next presidential election. If you know nothing else about me, one approach is to look at how my neighbors are planning to vote. Living in Seattle, my neighbors are planning to vote for the Democratic candidate, which suggests that "Democratic candidate" is a good guess for me as well.
- ▷ you know more about me : my age, my income, how many kids I have ... To the extent my behavior is influenced by those things, looking just at my neighbors who are close to me among all those dimensions seems likely to be an even better predictor than looking at all my neighbors. This is the idea behind *nearest neighbors classification*.

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**Examples**

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## Model : KNN

### Requirements

- ▷ Some notion of distance
- ▷ An assumption that points that are close to one another are similar

the prediction for each new point depends only on the handful of points closest to it.

## Model : KNN

- ▷ classify some new data point : find the k nearest labeled points and let them vote on the new output.
- ▷ need a function that counts votes : Reduce k until we find a unique winner.

```
def majority_vote(labels):
    """assumes that labels are ordered from nearest to farthest"""
    vote_counts = Counter(labels)
    winner, winner_count = vote_counts.most_common(1)[0]
    num_winners = len([count for count in vote_counts.values() if count ==
                      winner_count])
    if num_winners == 1:
        return winner # unique winner, so return it
    else:
        return majority_vote(labels[:-1]) # try again without the farthest
```

### Requirements

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  - ▷ An assumption that points that are close to one another are similar
- the prediction for each new point depends only on the handful of points closest to it.

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    vote_counts = Counter(labels)
    winner, winner_count = vote_counts.most_common(1)[0]
    num_winners = len([count for count in vote_counts.values() if count ==
                      winner_count])
    if num_winners == 1:
        return winner # unique winner, so return it
    else:
        return majority_vote(labels[:-1]) # try again without the farthest
```

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## Model : KNN

```
def knn_classify(k, labeled_points, new_point):
    """each labeled point should be a pair (point, label)"""

    # order the labeled points from nearest to farthest
    by_distance = sorted(labeled_points, key=lambda (point, _): distance(point,
        new_point))

    # find the labels for the k closest
    k_nearest_labels = [label for _, label in by_distance[:k]]

    # and let them vote
    return majority_vote(k_nearest_labels)
```

## Example : Favorite Programming Languages

```
# each entry is ([longitude, latitude], favorite_language)
cities = [[[-122.3, 47.53], "Python"], # Seattle
          [[-96.85, 32.85], "Java"], # Austin
          [[-89.33, 43.13], "R"], # Madison
          # ... and so on
          ]
```

```
def knn_classify(k, labeled_points, new_point):
    """each labeled point should be a pair (point, label)"""

    # order the labeled points from nearest to farthest
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          # ... and so on
          ]
```

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# Example : Favorite Programming Languages

## Plotting the data

```
# key is language, value is pair (longitudes, latitudes)
plots = { "Java" : ([], []), "Python" : ([], []), "R" : ([], []) }

# we want each language to have a different marker and color
markers = { "Java" : "o", "Python" : "s", "R" : "^" }
colors = { "Java" : "r", "Python" : "b", "R" : "g" }

for (longitude, latitude), language in cities:
    plots[language][0].append(longitude)
    plots[language][1].append(latitude)

# create a scatter series for each language
for language, (x, y) in plots.iteritems():
    plt.scatter(x, y, color=colors[language], marker=markers[language],
              label=language, zorder=10)

plot_state_borders(plt) # pretend we have a function that does this

plt.legend(loc=0) # let matplotlib choose the location
plt.axis([-130,-60,20,55]) # set the axes

plt.title("Favorite Programming Languages")
plt.show()
```

```
Plotting the data
from mpl_toolkits.basemap import Basemap, projection, contour
import numpy as np
import matplotlib.pyplot as plt

# Create a map of the United States
m = Basemap(projection='merc', llcrnrlon=-130, llcrnrlat=20, urcrnrlon=-60, urcrnrlat=55,
            resolution='l', contour=True)

# Create a scatter plot of the data
plt.figure(figsize=(10, 10))
m.scatter(cities['Java'], cities['Python'], cities['R'], marker='o', color='r', zorder=10)
m.scatter(cities['Python'], cities['Python'], cities['Python'], marker='s', color='b', zorder=10)
m.scatter(cities['R'], cities['R'], cities['R'], marker='^', color='g', zorder=10)

# Add state borders
m.plot_state_borders()

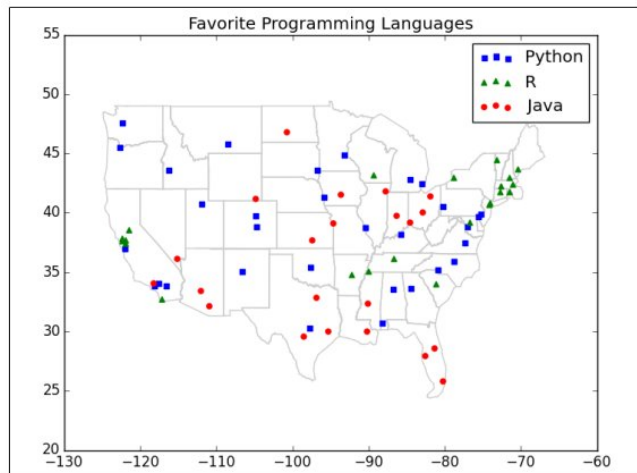
# Add a legend
plt.legend(loc=0)

# Set the axes
plt.axis([-130, -60, 20, 55])

# Title the plot
plt.title("Favorite Programming Languages")

# Show the plot
plt.show()
```

# Result



## Example : Favorite Programming Languages

Try several different values for  $k$

```
for k in [1, 3, 5, 7]:
    num_correct = 0
    for city in cities:
        location, actual_language = city
        other_cities = [other_city
                        for other_city in cities
                        if other_city != city]

        predicted_language = knn_classify(k, other_cities, location)

        if predicted_language == actual_language:
            num_correct += 1

    print k, "neighbor[s]:" , num_correct, "correct_out_of", len(cities)
```

## Example : Favorite Programming Languages

```
plots = { "Java" : ([], []), "Python" : ([], []), "R" : ([], []) }

k = 1 # or 3, or 5, or ...

for longitude in range(-130, -60):
    for latitude in range(20, 55):
        predicted_language = knn_classify(k, cities, [longitude, latitude])
        plots[predicted_language][0].append(longitude)
        plots[predicted_language][1].append(latitude)
```

```
Try several different values for k
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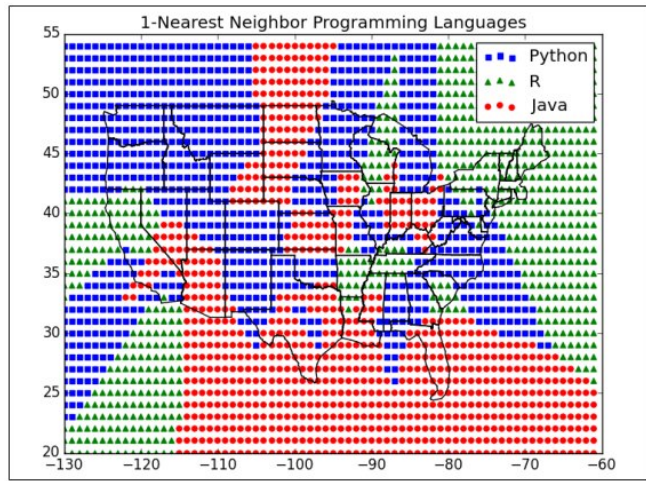
1 neighbor[s] : 40 correct out of 75 3 neighbor[s] : 44 correct out of 75 5 neighbor[s] : 41 correct out of 75  
7 neighbor[s] : 35 correct out of 75

```
plots = { "Java" : ([], []), "Python" : ([], []), "R" : ([], []) }

k = 1 # or 3, or 5, or ...

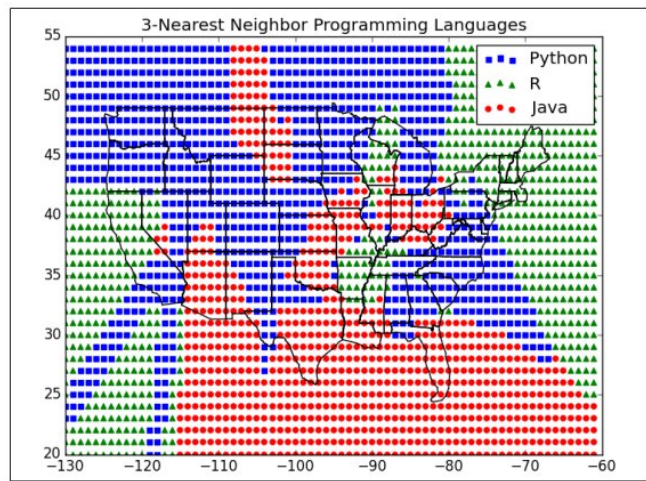
for longitude in range(-130, -60):
    for latitude in range(20, 55):
        predicted_language = knn_classify(k, cities, [longitude, latitude])
        plots[predicted_language][0].append(longitude)
        plots[predicted_language][1].append(latitude)
```

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k = 1

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k = 3

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## Unsupervised learning

### Learning mode

- ▷ supervised learning : set of labeled data for making predictions about new, unlabeled data.
- ▷ unsupervised learning : no label at all
- ▷ Whenever you look at some source of data, the data will somehow form *clusters*.

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Examples

- ▷ A data set showing where millionaires live probably has clusters in places like Beverly Hills and Manhattan.
- ▷ A data set showing how many hours people work each week probably has a cluster around 40.
- ▷ A data set of demographics of registered voters likely forms a variety of clusters (e.g., "soccer moms", "bored retirees" ...)

the clusters won't label themselves. You'll have to do that by looking at the data underlying each one.

- 1 Start with a set of k-means, which are points in d-dimensional space.
- 2 Assign each point to the mean to which it is closest.
- 3 If no point's assignment has changed, stop and keep the clusters.
- 4 If some point's assignment has changed, recompute the means and return to step 2.

Examples

- ▷ A data set showing where millionaires live probably has clusters in places like Beverly Hills and Manhattan.
- ▷ A data set showing how many hours people work each week probably has a cluster around 40.
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Model : k-means

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## Model : k-means

```
def vector_mean(vectors):
    "compute the vector whose ith element is the mean of the ith elements of the
    input vectors"
    n = len(vectors)
    return scalar_multiply(1/n, vector_sum(vectors))

class KMeans:
    """performs k-means clustering"""

    def __init__(self, k):
        self.k = k # number of clusters
        self.means = None # means of clusters

    def classify(self, input):
        """return the index of the cluster closest to the input"""
        return min(range(self.k),
            key=lambda i: squared_distance(input, self.means[i]))
```

## Model : k-means

```
def train(self, inputs):
    # choose k random points as the initial means
    self.means = random.sample(inputs, self.k)
    assignments = None

    while True:
        # Find new assignments
        new_assignments = map(self.classify, inputs)

        # If no assignments have changed, we're done.
        if assignments == new_assignments:
            return

        # Otherwise keep the new assignments
        assignments = new_assignments

        # And compute new means based on the new assignments
        for i in range(self.k):
            # find all the points assigned to cluster i
            i_points = [p for p, a in zip(inputs, assignments) if a == i]

            # make sure i_points is not empty so don't divide by 0
            if i_points:
                self.means[i] = vector_mean(i_points)
```

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## Example : stickers

### Context

- ▷ sticker printer can print at most five colors per sticker.
- ▷ there's some way to take a design and modify it so that it only contains five colors ?

### Data

- ▷ images can be represented as two-dimensional array of pixels, where each pixel is itself a three-dimensional vector (red, green, blue) indicating its color.
- ▷ five-color version of the image
  - 1 Choosing five colors
  - 2 Assigning one of those colors to each pixel

## Example : stickers

```
path_to_png_file = r"C:\images\image.png"
import matplotlib.image as mpimg
img = mpimg.imread(path_to_png_file)

top_row = img[0]
top_left_pixel = top_row[0]
red, green, blue = top_left_pixel

pixels = [pixel for row in img for pixel in row]

clusterer = KMeans(5)
clusterer.train(pixels)

def recolor(pixel):
    cluster = clusterer.classify(pixel)
    return clusterer.means[cluster]

new_img = [[recolor(pixel) for pixel in row]
            for row in img]

plt.imshow(new_img)
plt.axis('off')
plt.show()
```

**Context**

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# Alternative approach

“grow” clusters from the bottom up

- 1 Make each input its own cluster of one.
- 2 As long as there are multiple clusters remaining, find the two closest clusters and merge them.
- 3 At the end, we'll have one giant cluster containing all the inputs. If we keep track of the merge order, we can recreate any number of clusters by unmerging. For example, if we want three clusters, we can just undo the last two merges.

# Example : pizza



Pizza chain

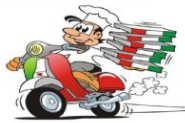
Optimal location ?

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# Example : pizza



Pizza chain

Optimal location ?



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# Example : pizza



How to teach the PC to do that ?



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# Example : pizza



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IML  
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└ Hierarchical clustering  
└ Example : pizza

Example : pizza



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# Example : pizza



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IML  
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└ Hierarchical clustering  
└ Example : pizza

Example : pizza



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# Example : pizza



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Example : pizza



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# Example : pizza



Hierarchical clustering

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└ Example : pizza

Example : pizza



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# Distance

Name	Egg-laying	Scales	Poisonous	Cold-blooded	Legs nb	Reptile
Cobra	True	True	True	True	0	Yes
Rattlesnake	True	True	True	True	0	Yes
Boa	False	False	False	True	0	Yes
Chicken	True	False	False	False	2	No
Alligator	True	False	False	True	4	Yes
Frog	True	True	True	True	4	No
Salmon	True	False	False	True	0	No
Python	True	False	False	True	0	Yes

Features = four binary and one integer

Boa = (0,1,0,1,0)

Frog =(1,0,1,1,4)

Distance to separate ?

# Distance : Euclidean

	rattlesnake	boa	frog
rattlesnake		1.4	4.2
boa	1.4		4.4
frog	4.2	4.4	

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Features : four binary and one integer  
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Distance to separate ?

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	rattlesnake	boa	frog
rattlesnake		1.4	4.2
boa	1.4		4.4
frog	4.2	4.4	

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## Distance : Euclidean

	rattlesnake	boa	frog	Alligator
rattlesnake		1.4	4.2	4.1
boa	1.4		4.4	4.1
frog	4.2	4.4		1.7
Alligator	4.1	4.1	1.7	

Alligator is closer to a frog than a snake

## Distance : Euclidean

	rattlesnake	boa	frog	Alligator
rattlesnake		1.4	1.7	1.4
boa	1.4		2.2	1.4
frog	1.7	2.2		1.7
Alligator	1.4	1.4	1.7	

Using binary Feature : Alligator is closer to a snake than a frog  
 Feature Engineering Matters

	rattlesnake	boa	frog	Alligator
rattlesnake		1.4	4.2	4.1
boa	1.4		4.4	4.1
frog	4.2	4.4		1.7
Alligator	4.1	4.1	1.7	

Alligator is closer to a frog than a snake

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	rattlesnake	boa	frog	Alligator
rattlesnake		1.4	1.7	1.4
boa	1.4		2.2	1.4
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Using binary Feature - Alligator is closer to a snake than a frog  
 Feature Engineering Matters

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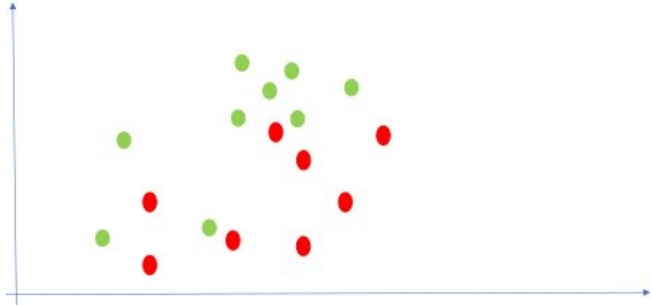
## Testing

- ▷ How well is my model doing ?
- ▷ How do I improve it ?

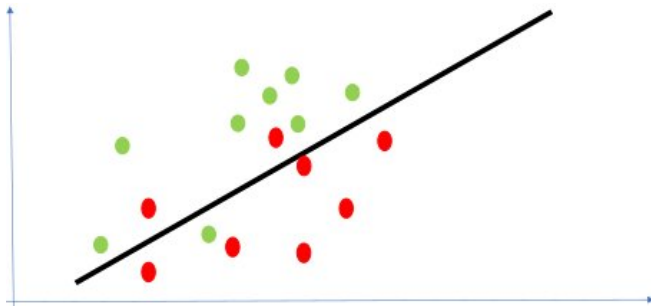
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# Which model is better ?

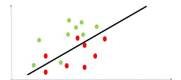


# Which model is better ?



└ Which model is better ?

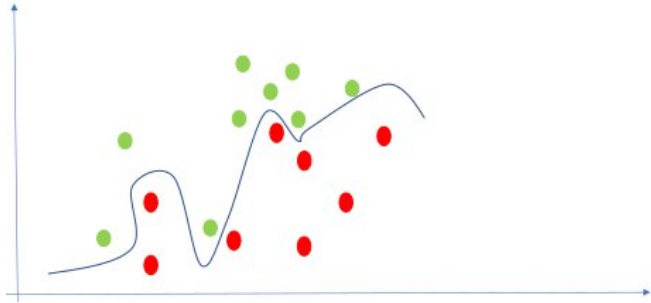
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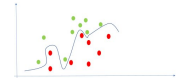
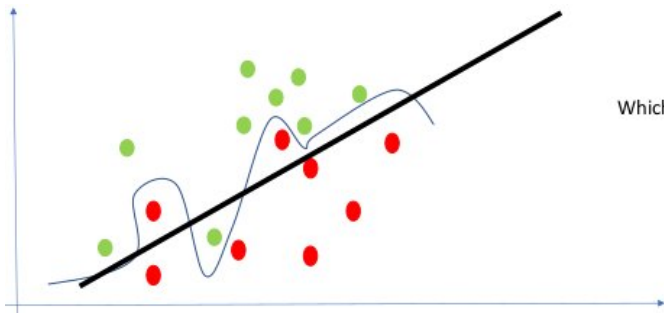
└ Which model is better ?

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# Which model is better ?

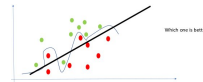


# Which model is better ?



Which model is better ?

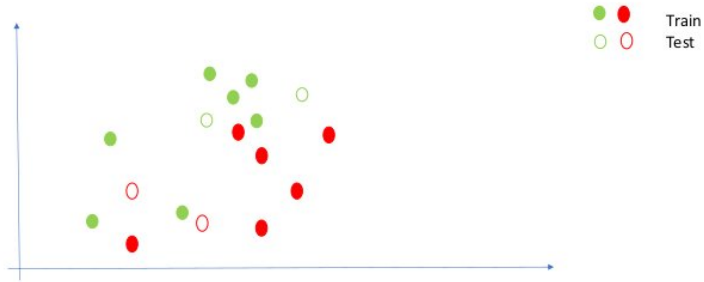
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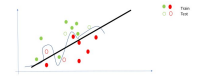
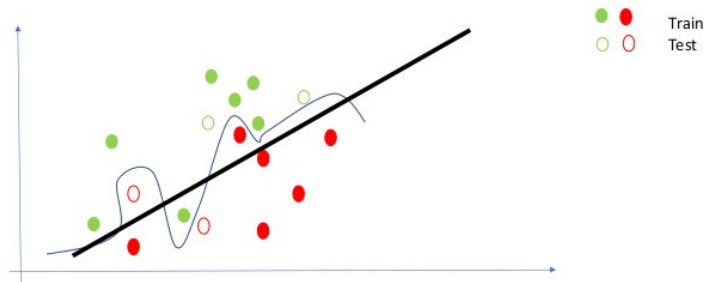
Which model is better ?

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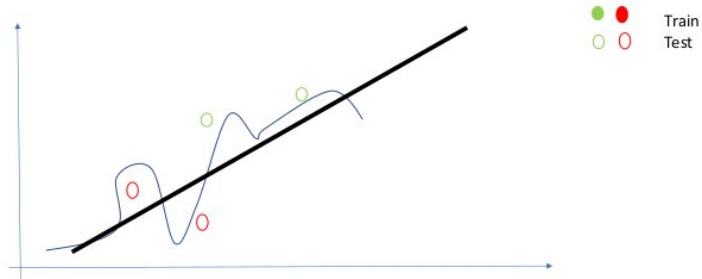
# Training vs Testing



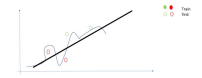
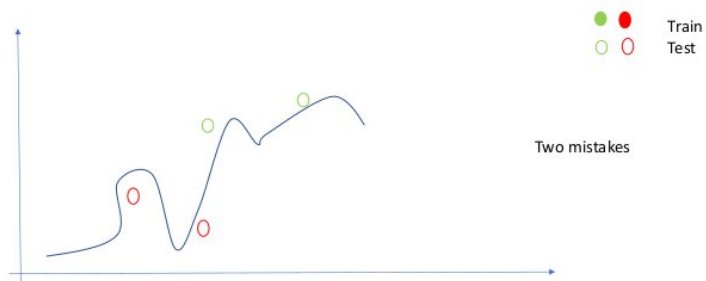
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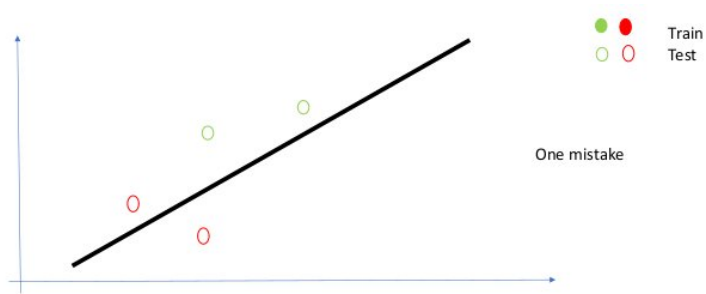
# Training vs Testing



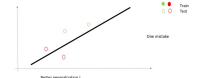
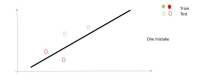
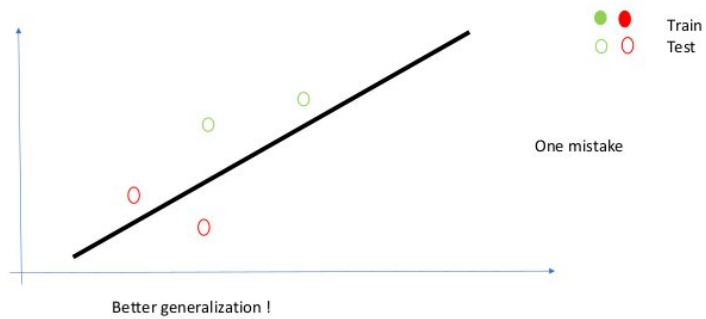
# Training vs Testing



# Training vs Testing



# Training vs Testing



# Learning Rule

▷ NEVER use your testing data for training



# Training vs Testing

```
def split_data(data, prob):
    results = [], []
    for row in data:
        results[0 if random.random() < prob else 1].append(row)
    return results

def train_test_split(x, y, test_pct):
    data = zip(x, y)
    train, test = split_data(data, 1 - test_pct) # pair corresponding values
    x_train, y_train = zip(*train) # split the data set of pairs
    x_test, y_test = zip(*test) # magical un-zip trick
    return x_train, x_test, y_train, y_test

model = SomeKindOfModel()
x_train, x_test, y_train, y_test = train_test_split(xs, ys, 0.33)
model.train(x_train, y_train)
performance = model.test(x_test, y_test)
```

▷ NEVER use your testing data for training

Page 47 :

```
def split_data(data, prob):
    results = [], []
    for row in data:
        results[0 if random.random() < prob else 1].append(row)
    return results

def train_test_split(x, y, test_pct):
    data = zip(x, y)
    train, test = split_data(data, 1 - test_pct) # pair corresponding values
    x_train, y_train = zip(*train) # split the data set of pairs
    x_test, y_test = zip(*test) # magical un-zip trick
    return x_train, x_test, y_train, y_test

model = SomeKindOfModel()
x_train, x_test, y_train, y_test = train_test_split(xs, ys, 0.33)
model.train(x_train, y_train)
performance = model.test(x_test, y_test)
```

Page 48 :



# Learning Rule

▷ NEVER use your testing data for training



How not losing data ?

# K-Fold Cross Validation



# K-Fold Cross Validation

Training

Testing



# K-Fold Cross Validation

Training

Testing



Page 51 :

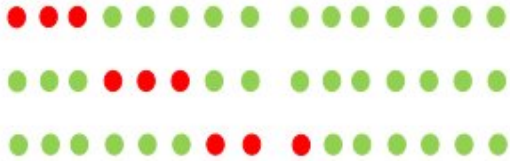


Page 52 :

# K-Fold Cross Validation

Training

Testing



Page 53 :

# K-Fold Cross Validation

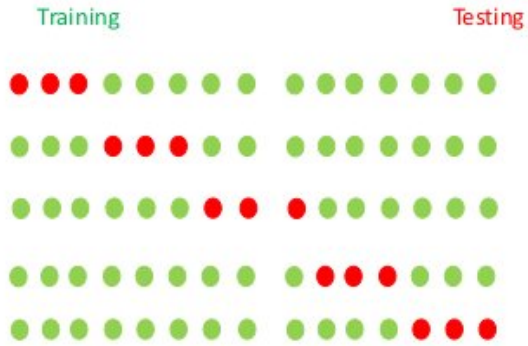
Training

Testing

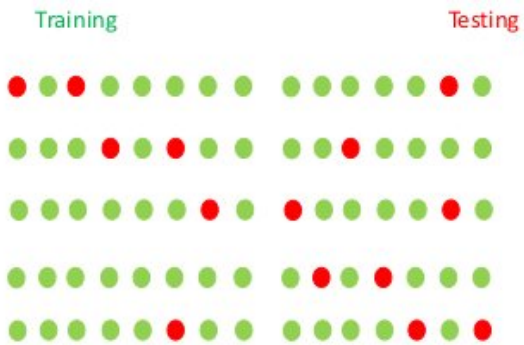


Page 54 :

# K-Fold Cross Validation



# K-Fold Cross Validation



- ▷ How well is my model doing?
  - ◇ Tricky question

## Example : Credit Card Fraud



Page 57 :

Page 58 :

# Example : Credit Card Fraud



284 335



472



Page 59 :

# Example : Credit Card Fraud



284 335



472

Model : All transactions are good



Page 60 :

# Example : Credit Card Fraud



284 335



472

Model : All transactions are good  
Correct :  $284\,335 / (284\,335 + 472) = 99.83\%$



Page 61 :

# Example : Credit Card Fraud



284 335



472

Model : All transactions are good  
Correct :  $284\,335 / (284\,335 + 472) = 99.83\%$   
What about bad transactions ??



Page 62 :

# Example : Credit Card Fraud



284 335



472

Model : All transactions are fraudulent  
Catching all bad transactions  
But ...

# Example : medical model







Page 63 :







Page 64 :










# Confusion Matrix

	Diagnosed SICK	Diagnosed HEALTHY
SICK	True Positive 	False Negative 
HEALTHY	False Positive 	True Positive 

	Diagnosed SICK	Diagnosed HEALTHY
SICK	True Positive 	False Negative 
HEALTHY	False Positive 	True Positive 

# Confusion Matrix

	Diagnosed SICK	Diagnosed HEALTHY
SICK	True Positive 	False Negative 
HEALTHY	False Positive 	True Positive 

	Diagnosed SICK	Diagnosed HEALTHY
SICK	True Positive 	False Negative 
HEALTHY	False Positive 	True Positive 

# Confusion Matrix

	Diagnosed SICK	Diagnosed HEALTHY
SICK	1000	200
HEALTHY	800	8000

# Example : spam model







	Diagnosed SICK	Diagnosed HEALTHY
SICK	1000	200
HEALTHY	800	8000

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Page 68 :

# Confusion Matrix

	Diagnosed SPAM	Diagnosed NON SPAM
SPAM	True Positive 	False Negative 
NON SPAM	False Positive 	True Positive 

# Confusion Matrix

	Diagnosed SPAM	Diagnosed NON SPAM
SPAM	100	170
NON SPAM	30	700

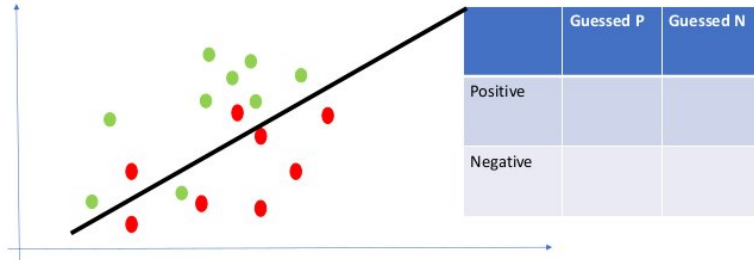
	Diagnosed SPAM	Diagnosed NON SPAM
SPAM	True Positive 	False Negative 
NON SPAM	False Positive 	True Positive 

Page 69 :

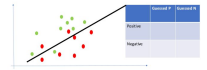
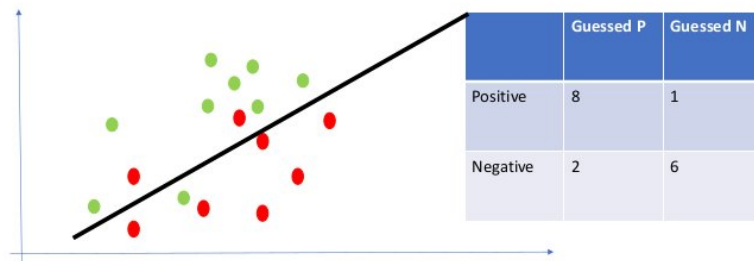
	Diagnosed SPAM	Diagnosed NON SPAM
SPAM	100	170
NON SPAM	30	700

Page 70 :

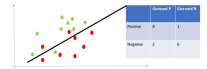
# Confusion Matrix



# Confusion Matrix



Page 71 :



Page 72 :

# Accuracy

How many did we classify correctly ?

	Diagnosed SICK	Diagnosed HEALTHY
SICK	1000	200
HEALTHY	800	8000

# Accuracy

How many did we classify correctly ?

	Diagnosed SICK	Diagnosed HEALTHY
SICK	1000	200
HEALTHY	800	8000

$$\text{Accuracy} = (1000+8000)/10000 = 90\%$$

How many did we classify correctly ?

	Diagnosed SICK	Diagnosed HEALTHY
SICK	1000	200
HEALTHY	800	8000

Page 73 :

How many did we classify correctly ?

	Diagnosed SICK	Diagnosed HEALTHY
SICK	1000	200
HEALTHY	800	8000

Accuracy = (1000+8000)/10000 = 90%

Page 74 :

# Accuracy

How many did we classify correctly?

	Diagnosed SPAM	Diagnosed NON SPAM
SPAM	100	170
NON SPAM	30	700

Accuracy = 80%

Accuracy = Correctly classified / all

# Accuracy

```
def accuracy(tp, fp, fn, tn):
    correct = tp + tn
    total = tp + fp + fn + tn
    return correct / total
```

How many did we classify correctly?

	Diagnosed SPAM	Diagnosed NON SPAM
SPAM	100	170
NON SPAM	30	700

Accuracy = 80%  
Accuracy = Correctly classified / all

Page 75 :



How many did we classify correctly?

	Diagnosed SPAM	Diagnosed NON SPAM
SPAM	100	170
NON SPAM	30	700

Accuracy = 80%  
Accuracy = Correctly classified / all



Page 76 :

# Confusion Matrix

	Diagnosed SICK	Diagnosed HEALTHY
SICK		False Negative 
HEALTHY	False Positive 	

	Diagnosed SICK	Diagnosed HEALTHY
SICK		False Negative 
HEALTHY	False Positive 	

# Confusion Matrix

	Diagnosed SICK	Diagnosed HEALTHY
SICK		False Negative 
HEALTHY	False Positive 	

	Diagnosed SICK	Diagnosed HEALTHY
SICK		False Negative 
HEALTHY	False Positive 	



# Precision

	Diagnosed SPAM	Diagnosed NON SPAM
SPAM		False Negative 
NON SPAM	False Positive 	

# Precision

▷ high PRECISION 

▷ high RECALL 

	Diagnosed SPAM	Diagnosed NON SPAM
SPAM		False Negative 
NON SPAM	False Positive 	

Page 79 :

	Diagnosed SPAM	Diagnosed NON SPAM
SPAM		
NON SPAM		

Page 80 :



# Precision

How many did we classify correctly ?

	Diagnosed SICK	Diagnosed HEALTHY
SICK	1000	200
HEALTHY	800	8000

Precision =  $1000 / (1000 + 800) = 55,7\%$

# Precision

How many did we classify correctly ?

	Diagnosed SPAM	Diagnosed NON SPAM
SPAM	100	170
NON SPAM	30	700

Precision = 76.8%

Precision =  $\text{True Positives} / (\text{True Positives} + \text{False Positives})$

How many did we classify correctly ?

	Diagnosed SICK	Diagnosed HEALTHY
SICK	1000	200
HEALTHY	800	8000

Precision =  $1000 / (1000 + 800) = 55,7\%$

Page 81 :

How many did we classify correctly ?

	Diagnosed SPAM	Diagnosed NON SPAM
SPAM	100	170
NON SPAM	30	700

Precision = 76.8%  
Precision =  $\text{True Positives} / (\text{True Positives} + \text{False Positives})$

Page 82 :

# Precision

```
def precision(tp, fp, fn, tn):
    return tp / (tp + fp)
```

# Recall

How many did we classify correctly ?

	Diagnosed SICK	Diagnosed HEALTHY
SICK	1000	200
HEALTHY	800	8000

$$\text{Recall} = 1000 / (1000 + 200) = 83.3\%$$

Page 83 :

How many did we classify correctly ?

	Diagnosed SICK	Diagnosed HEALTHY
SICK	1000	200
HEALTHY	800	8000

Recall = 1000 / (1000 + 200) = 83.3%

Page 84 :

# Recall

How many did we classify correctly?

	Diagnosed SPAM	Diagnosed NON SPAM
SPAM	100	170
NON SPAM	30	700

Recall = 37%

Recall = True Positives / (True Positives + False Negatives)

# Recall

```
def recall(tp, fp, fn, tn):
    return tp / (tp + fn)
```

2018-08-30

IML  
└─ Tests  
    └─ Model performance  
        └─ Recall

Recall

How many did we classify correctly?

	Diagnosed SPAM	Diagnosed NON SPAM
SPAM	100	170
NON SPAM	30	700

Recall = 37%  
Recall = True Positives / (True Positives + False Negatives)

Page 85 :

2018-08-30

IML  
└─ Tests  
    └─ Model performance  
        └─ Recall

Recall

How many did we classify correctly?

	Diagnosed SPAM	Diagnosed NON SPAM
SPAM	100	170
NON SPAM	30	700

Recall = 37%  
Recall = True Positives / (True Positives + False Negatives)

Page 86 :

# Precision and Recall



- ◇ Precision : 76,9%
- ◇ Recall : 37%



- ◇ Precision : 55,7%
- ◇ Recall : 83,3%

# Average



- ◇ Precision : 76,9%
- ◇ Recall : 37%
- ◇ Average : 56,9%

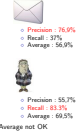


- ◇ Precision : 55,7%
- ◇ Recall : 83,3%
- ◇ Average : 69,5%

Average not OK



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Page 88 :



Page 89 :



284 335

472

Model : All transactions are good  
Precision = 100%  
Recall = 0%  
Average = 50%

Page 90 :



284 335



472

Model : All transactions are fraudulent  
Precision = .016%  
Recall = 100%  
Average = 50%

Page 91 :

$$F1 \text{ Score} = (2 \times \text{Precision} \times \text{Recall}) / (\text{Precision} + \text{Recall})$$



- ◇ Precision : 76,9%
- ◇ Recall : 37%
- ◇ Average : 56,9%
- ◇ F1 Score = 50%



- ◇ Precision : 55,7%
- ◇ Recall : 83,3%
- ◇ Average : 69,5%

$$F1 \text{ Score} = (2 \times 55,7 \times 83,3) / (55,7 + 83,3) = 66\%$$

Page 92 :

# F1 Score

```
def f1_score(tp, fp, fn, tn):  
    p = precision(tp, fp, fn, tn)  
    r = recall(tp, fp, fn, tn)  
    return 2 * p * r / (p + r)
```

# Underfitting/overfitting

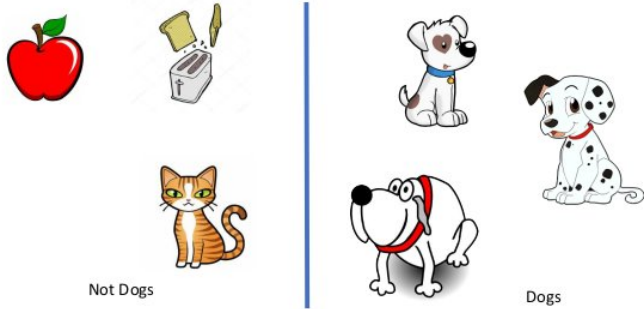


Page 93 :

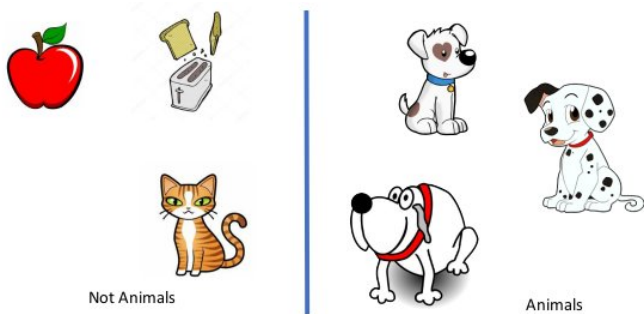


Page 94 :

# Underfitting/overfitting

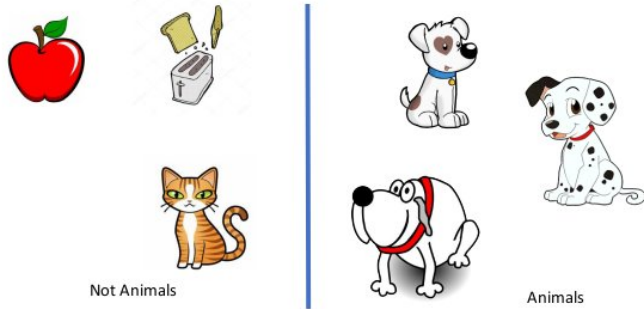


# Underfitting/overfitting

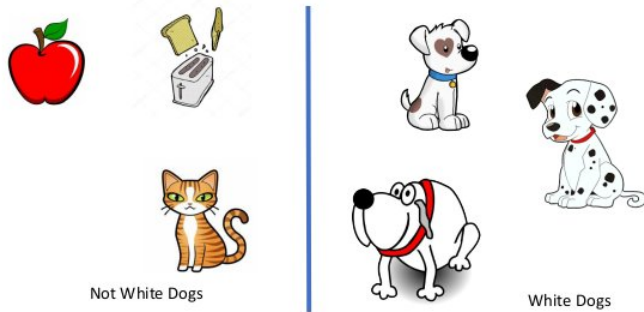




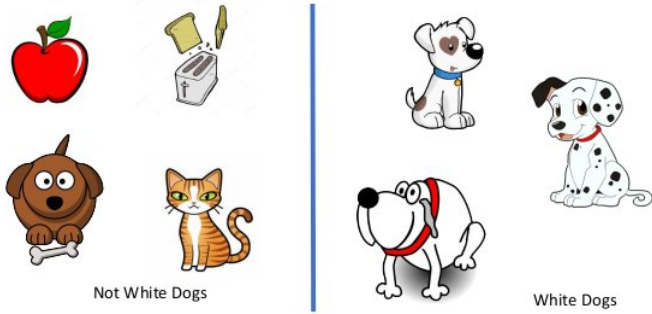
# Underfitting/overfitting



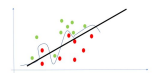
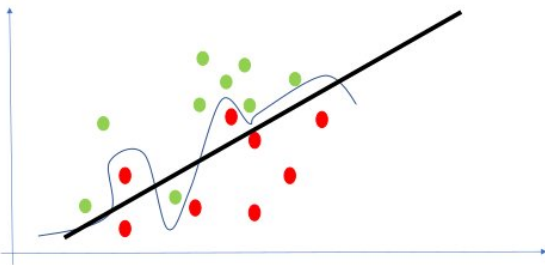
# Underfitting/overfitting



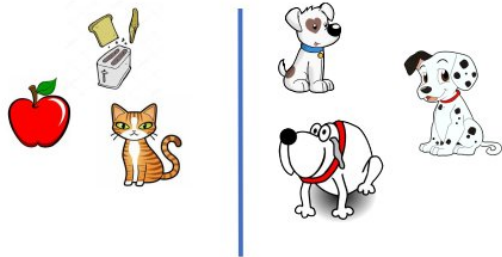
# Underfitting/overfitting



# Underfitting/overfitting



# Underfitting/overfitting



Underfitting : Not Animals  
Overfitting : Not White Dogs  
OK : Not Dogs

Underfitting : Animals  
Overfitting : White Dogs  
OK : Dogs

# Underfitting/overfitting



Training set :  
Bad  
Great  
Good

Underfitting : Not Animals  
Overfitting : Not White Dogs  
OK : Not Dogs

Underfitting : Animals  
Overfitting : White Dogs  
OK : Dogs

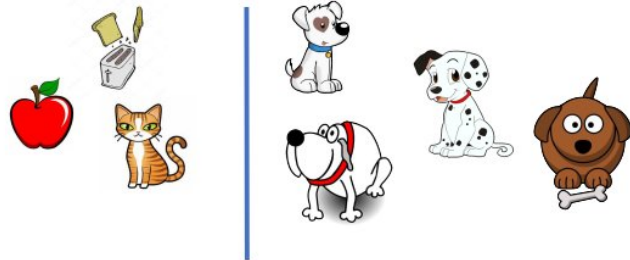


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Page 102 :

# Underfitting/overfitting



Testing set :

Bad Underfitting : Not Animals  
Bad Overfitting : Not White Dogs  
Good OK : Not Dogs

Underfitting : Animals  
Overfitting : White Dogs  
OK : Dogs

the more data you have, the harder it is to over- fit.

- 1 Classification
  - KNN
- 2 Clustering
  - k-means
  - Hierarchical clustering
  - Distance
- 3 Tests
  - Training vs Testing
  - K-Fold Cross Validation
  - Model performance
- 4 Feature extraction



the more data you have, the harder it is to over- fit.

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## Features

As we mentioned, when your data doesn't have enough features, your model is likely to underfit. And when your data has too many features, it's easy to overfit. But what are features and where do they come from ?

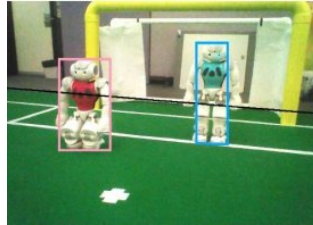
*Features* are whatever inputs we provide to our model.

## Type of features

Type of features we have constrains the type of models we can use :

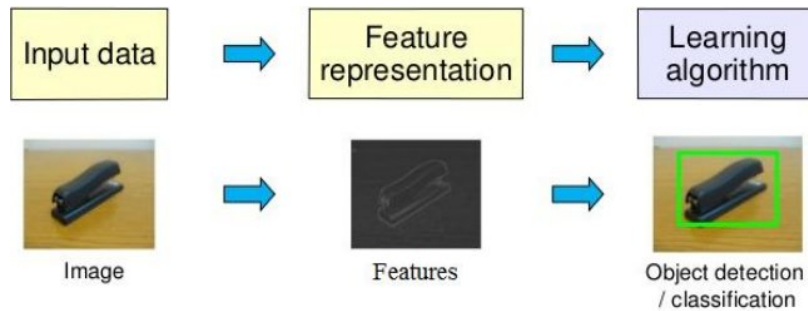
- ▷ The Naive Bayes classifier is suited to yes-or-no features
- ▷ Regression models require numeric features
- ▷ Decision trees can deal with numeric or categorical data.

# Example : robot detection



Can we detect robot using low quality images ?

# Example : robot detection



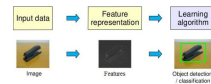
# Example : robot detection



Can we detect robot using low quality images ?

Page 107 :

# Example : robot detection

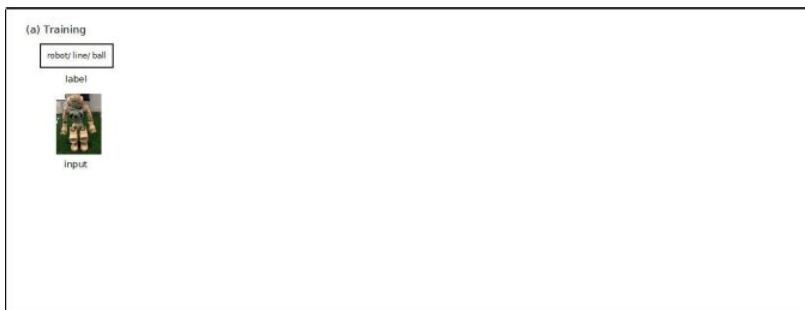


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# Example : robot detection



# Example : robot detection

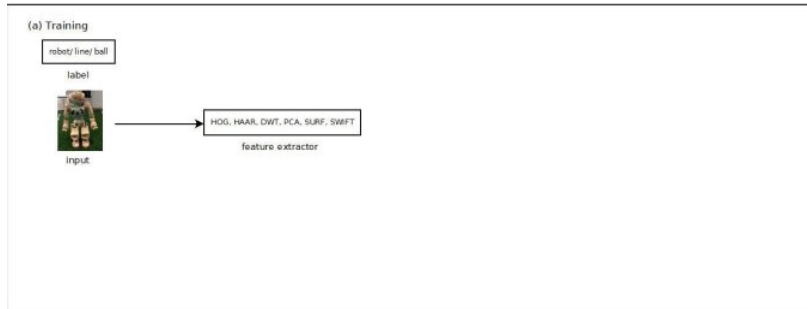


Page 109 :

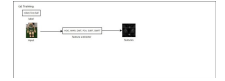
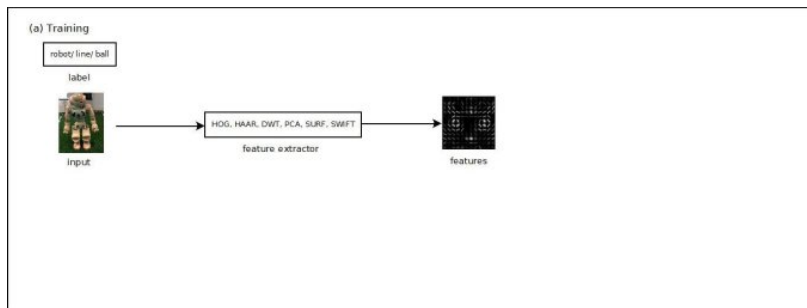


Page 110 :

# Example : robot detection

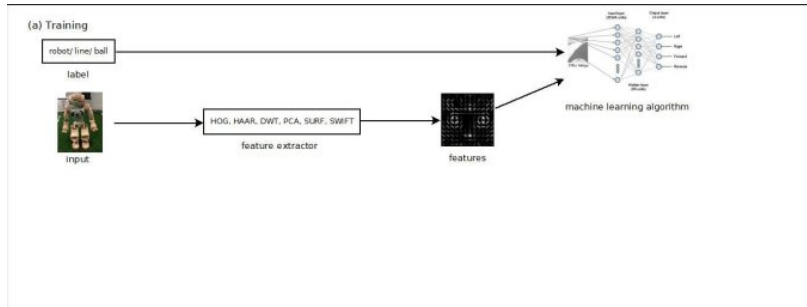


# Example : robot detection

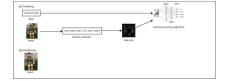
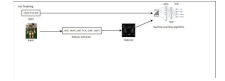
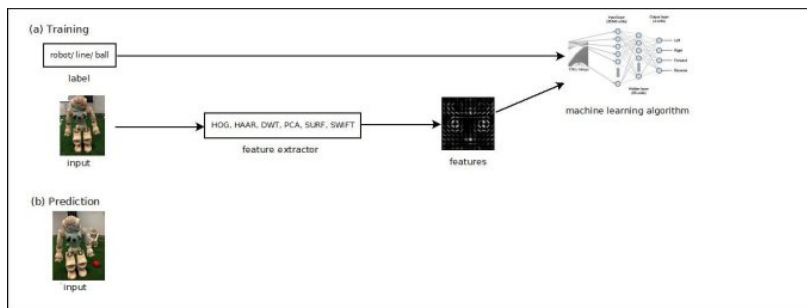




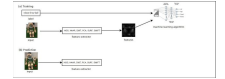
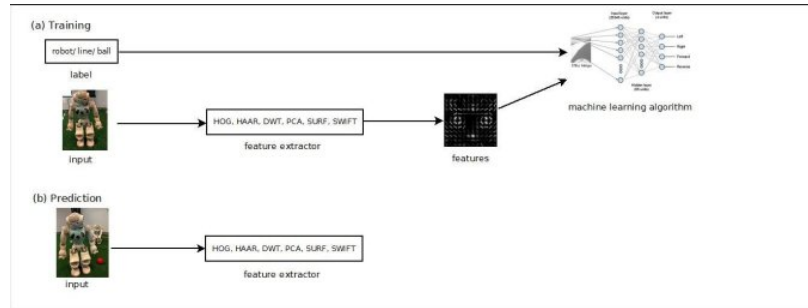
# Example : robot detection



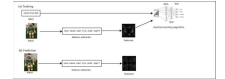
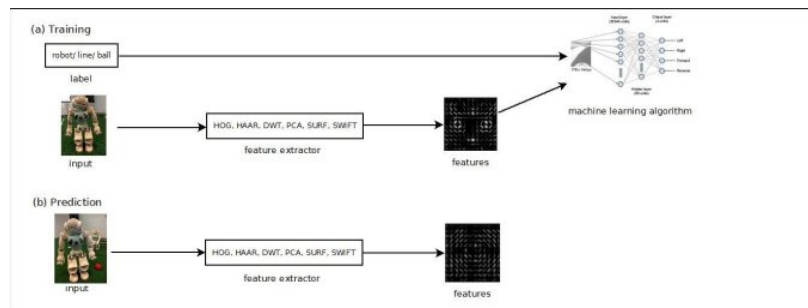
# Example : robot detection



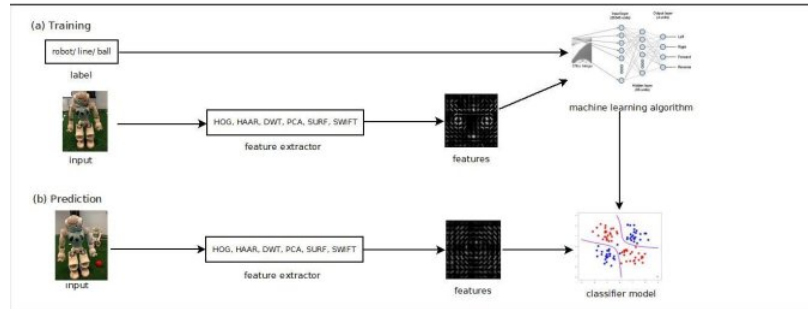
# Example : robot detection



# Example : robot detection

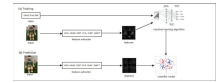


## Example : robot detection



## HOG+SVM

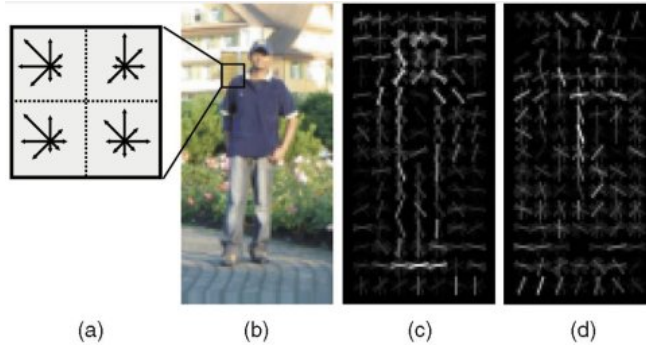
- ▷ Application : Persons detector
- ▷ HOG : Histograms of Oriented Gradients
- ▷ The intent of a feature descriptor is to generalize the object in such a way that the same object (in this case a person) produces as close as possible to the same feature descriptor when viewed under different conditions. This makes the classification task easier.
- ▷ The creators of this approach trained a Support Vector Machine (a type of machine learning algorithm for classification), or "SVM", to recognize HOG descriptors of people.



- ▷ Application : Persons detector
- ▷ HOG : Histograms of Oriented Gradients
- ▷ The intent of a feature descriptor is to generalize the object in such a way that the same object (in this case a person) produces as close as possible to the same feature descriptor when viewed under different conditions. This makes the classification task easier.
- ▷ The creators of this approach trained a Support Vector Machine (a type of machine learning algorithm for classification), or "SVM", to recognize HOG descriptors of people.

# HOG+SVM

HOG : entire person is represented by a single feature vector

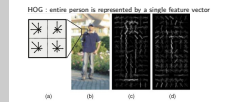


# HOG+SVM

The HOG person detector uses a sliding detection window which is moved around the image.

```

1 from skimage.transform import pyramid_gaussian
2 from PIL import Image
3 import cv2
4
5 im = cv2.imread('data/images/robot.jpg')
6 min_w, min_h = 100, 100
7 step_size = (10, 10)
8 Downscale = 1.5
9
10 def sliding_window(image, window_size, step_size):
11
12     This function returns a patch of the input image image of size equal
13     to window_size. The first image returned top-left co-ordinates (x, y),
14     and the increment in both x and y directions by the step_size supplied
15     in the next subsequent call.
16     * image - Input image
17     * window_size - Size of sliding window
18     * step_size - Incremental size of window
19
20     The function returns a tuple
21     (x, y, window)
22
23     where
24     * x is the top-left x co-ordinate
25     * y is the top-left y co-ordinate
26     * window is the sliding window image
27
28     for y in xrange(0, image.shape[0], step_size):
29         for x in xrange(0, image.shape[1], step_size):
30             yield (x, y, image[y:y+window_size[0], x:x+window_size[1]])
31
32 python sliding-window.py
    
```



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# HOG+SVM

- ▷ At each position of the detector window, a HOG descriptor is computed for the detection window.
- ▷ This descriptor is then shown to the trained SVM, which classifies it as either “person” or “not a person” .
- ▷ To recognize persons at different scales, the image is subsampled to multiple sizes. Each of these subsampled images is searched

# HOG+SVM



data

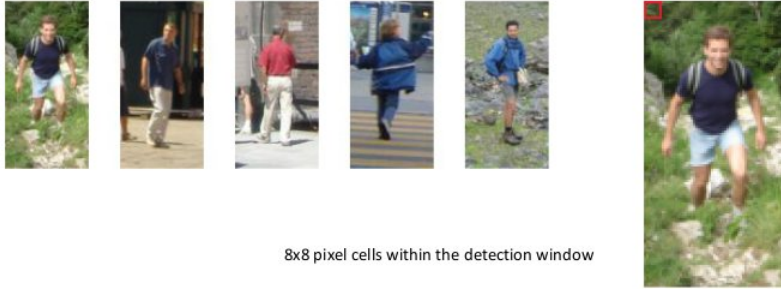
- ▷ At each position of the detector window, a HOG descriptor is computed for the detection window.
- ▷ This descriptor is then shown to the trained SVM, which classifies it as either “person” or “not a person” .
- ▷ To recognize persons at different scales, the image is subsampled to multiple sizes. Each of these subsampled images is searched

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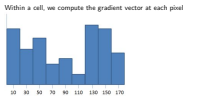
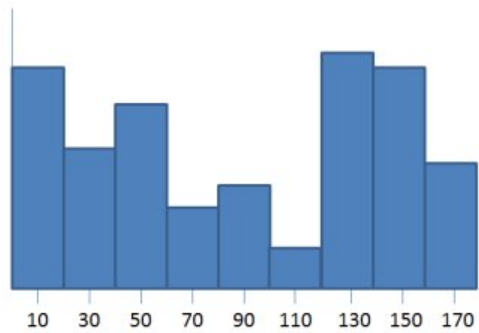
# HOG+SVM



8x8 pixel cells within the detection window

# HOG+SVM

Within a cell, we compute the gradient vector at each pixel



## To sum up

- ▷ Define problem (data)
- ▷ List tools (algorithms)
- ▷ Evaluate tools to find the best one
  - ◇ Accuracy
  - ◇ Precision
  - ◇ Recall
  - ◇ F1

## Detection & Tests *IML*

Cédric Buche

ENIB

30 août 2018

- ▷ Define problem (data)
- ▷ List tools (algorithms)
- ▷ Evaluate tools to find the best one
  - ◇ Accuracy
  - ◇ Precision
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  - ◇ F1

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