

# Lecture 3: Model Evaluation

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## Binary Classification

- **Binary** classification problem in Machine Learning (ML)
  - identifying if a certain patient has *some disease* using his *health record*
  - Credit versus No Credit (*using Decision Tree lecture 2*)
  - Class a versus Class b
- ML trained on a training set  $D_t$  tested on a test set  $D_{test}$  with  

$$\emptyset = D_t \cap D_{test}$$

$$accuracy = \frac{\text{Correctly Classified}}{\text{All}} \quad \text{error rate} = 1 - accuracy$$

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## Evaluating classification models

- Some types of mistakes that are worse than others
- We are choosing between two models A and B that diagnose a given infectious disease
  - positive if the disease present, negative if the disease not
  - present both models have the **same accuracy**, which model is better?
- model A's mistakes are all **false positives**
  - cases where the patient is not sick but the model *predicted disease*
- model B where all mistakes are **false negatives**
  - *contagious people are told they are healthy*

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## Confusion Matrix

|             |   | true/actual/target   |                      |         |
|-------------|---|----------------------|----------------------|---------|
|             |   | P                    | N                    |         |
| predicted { | P | True Positives (TP)  | False Positives (FP) | TP+FP   |
|             | N | False Negatives (FN) | True Negatives (TN)  | FN+TN   |
|             |   | P=TP+FN              | N=FP+TN              | All=P+N |

### Recall/sensitivity

% of positive observations predicted as positive

$$Recall = \frac{TP}{P} = \frac{TP}{TP+FN}$$

### Precision

% of positive observations among the observations predicted as positive

$$Precision = \frac{TP}{TP+FP}$$

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## Precision and Recall

- A high recall value without a high precision does not give us any confidence about the quality of the binary classifier
  - High recall value by classifying all patterns as positive (the recall value will be one); however, the precision value will be very low

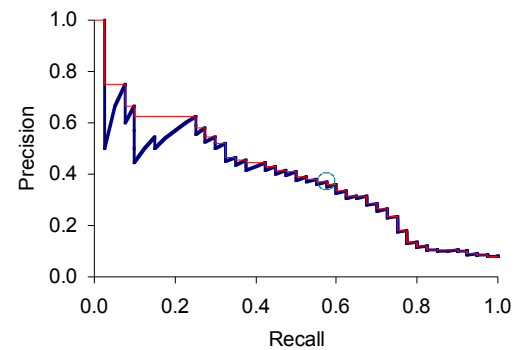
$$Recall = \frac{TP}{P} = \frac{TP}{TP+FN} = \frac{All}{All} = 1$$

- By classifying only one pattern correctly as positive, we obtain the maximal precision value of one but a low recall value.

$$Precision = \frac{TP}{TP+FP} = \frac{1}{1+0} = 1$$

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## A precision-recall curve



Both values have to be simultaneously interpreted

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## Balanced Measure

- Precision and Recall have to be simultaneously interpreted.
- We can combine both values with the *harmonic mean*

$$F = 2 \cdot \frac{\text{Precision} \cdot \text{Recall}}{\text{Precision} + \text{Recall}}$$

- Both values are evenly weighted.
- This measure is also called the balanced measure.

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## A combined measure: $F$

- Combined measure that assesses this tradeoff is  $F$  measure (weighted harmonic mean):

$$F = \frac{1}{\alpha \frac{1}{P} + (1-\alpha) \frac{1}{R}} = \frac{(\beta^2 + 1)PR}{\beta^2 P + R}$$

- However, usually use **balanced  $F_1$  measure**
  - i.e., with  $\beta = 1$  or  $\alpha = \frac{1}{2}$
  - $P = \text{Precision}$ ,  $R = \text{Recall}$

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## ROC curve

- For binary classifier indicates the probability of two classes:

$C_1$  and **not**  $C_1$       positive class :=  $C_1$   
 $p(C_1)$  and **not**  $C_1 = 1 - p(C_1)$       negative class := not  $C_1$

If  $p(C_1) \geq \text{threshold}$  then class  $C_1$   
 If  $p(C_1) < \text{threshold}$  then class **not**  $C_1$

Usually the threshold is 0.5

- Niave Bayes, Perceptron, Logistic Regression
  - introduced later in the course

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## ROC Curve

### Receiver Operating Characteristic

#### Recall/sensitivity

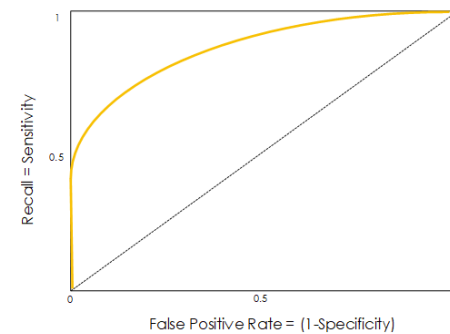
- % of positive observations predicted as positive

$$\text{Recall} = \frac{TP}{P} = \frac{TP}{TP + FN}$$

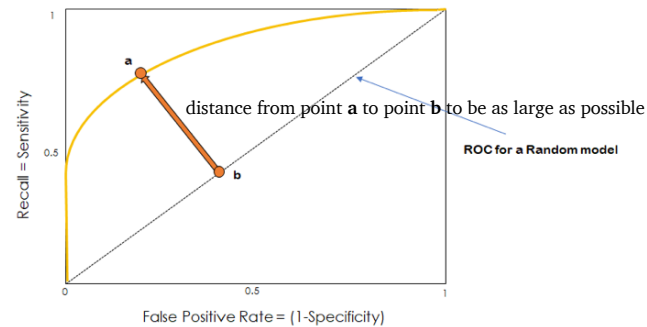
#### Fallout/specificity

- % of negative observations predicted as negative

$$\text{False Positive Rate} = \text{Specificity} = \frac{TN}{N} = \frac{TN}{TN + FP}$$



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- To plot the ROC curve, we must first calculate the *Recall* and the *Specificity* for **various thresholds**, and then plot them against each other
- The further away we are to the curve of the random model, the better

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## Various thresholds for ROC curve

- For binary classifier indicates the probability of two classes:

$C_1$  and **not**  $C_1$                       positive class :=  $C_1$   
 $p(C_1)$  and **not**  $C_1 = 1 - p(C_1)$                       negative class := not  $C_1$

If  $p(C_1) \geq \text{threshold}$  then class  $C_1$   
 If  $p(C_1) < \text{threshold}$  then class **not**  $C_1$

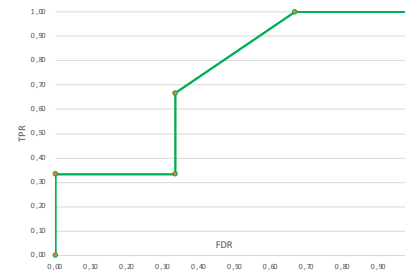
Usually the threshold is 0.5

- To compute the ROC curve we chose **various thresholds**  $\in [0,1]$
- We chose threshold=0, then threshold=0.1,..., threshold=0.9, threshold=1

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$z$  is the true value and  $\hat{z}$  the classifier prediction

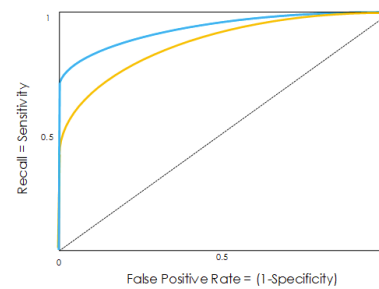
| $z$      | $\hat{z}$ | 0    | >0.3 | >0.4 | >0.45 | >0.6 | >0.8 |
|----------|-----------|------|------|------|-------|------|------|
| 1        | 0.5       | TP   | TP   | TP   | FN    | FN   | FN   |
| 1        | 0.8       | TP   | TP   | TP   | TP    | TP   | FN   |
| 1        | 0.45      | TP   | TP   | FN   | FN    | FN   | FN   |
| 0        | 0.4       | FP   | FP   | TN   | TN    | TN   | TN   |
| 0        | 0.3       | FP   | TN   | TN   | TN    | TN   | TN   |
| 0        | 0.6       | FP   | FP   | FP   | FP    | TN   | TN   |
| FPR=FP/N |           | 1.00 | 0.67 | 0.33 | 0.33  | 0.00 | 0.00 |
| TPR=TP/P |           | 1.00 | 1.00 | 0.67 | 0.33  | 0.33 | 0.00 |



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## AUC metric (Area Under the Curve)

- *ACU* quantifies in a **single metric** how well our model classifies the True and False data points.
- *AUC* goes from values of 0.5 (random classifier) to 1 (perfect classifier)



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## Lift Charts

- Comparing classifiers:
  - 1,000,000 prospective respondents
  - prediction that 0.1% of **all households** (1,000,000) will respond
  - prediction that 0.4% of a **specified** 100,000 homes will respond.
  - lift factor=increase in response rate=4
  - Given a classifier that outputs *probabilities* for the predicted class value for each test instance, what to do?

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## Lift Factor

**sample success proportion=**  
**(number of positive instances in sample) / sample size**

**lift factor=**  
**(sample success proportion) / (total test set success proportion)**

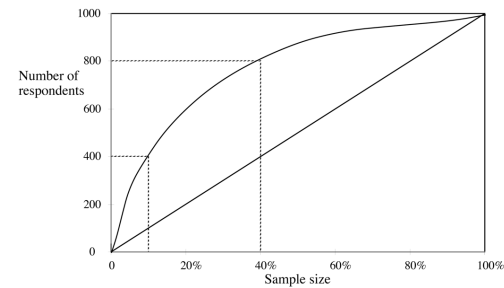
- Plot the number of respondents as a function of the number of mailings
- Why is response rate dropping with increasing number of mailings?

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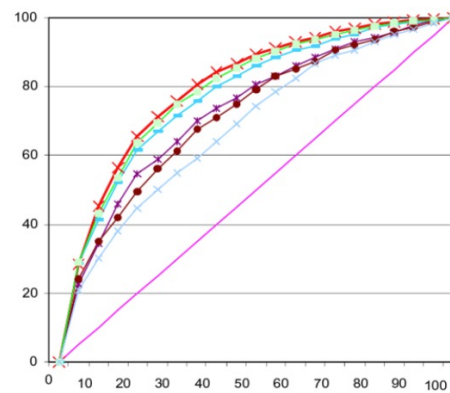
## Evaluation of Lift Chart

- Two extreme points:
  - Lower left: if no solicitations are sent - no respondents
  - Upper right: if all households receive offers - 1000 respondents
- What is the ideal point in the chart?
- Best to be in the upper left-hand corner of the chart: mail only to those 1000 (out of a million!) who would respond.



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## Lift charts of two classifiers: which one is better?



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## Evaluating multiclass classifiers

- Most real-world classification problems have more than two classes
  - e.g. identifying risk groups, categorizing documents, recommending products
- Extend binary confusion matrices

|                  |   | <i>true/actual/target</i> |              |              |
|------------------|---|---------------------------|--------------|--------------|
|                  |   | A                         | B            | C            |
| <i>predicted</i> | P | True A (TA)               | False A (FA) | False A (FA) |
|                  | B | False B (FB)              | True B (TB)  | False B (FB) |
|                  | C | False C (FC)              | False C (FC) | True C (TC)  |

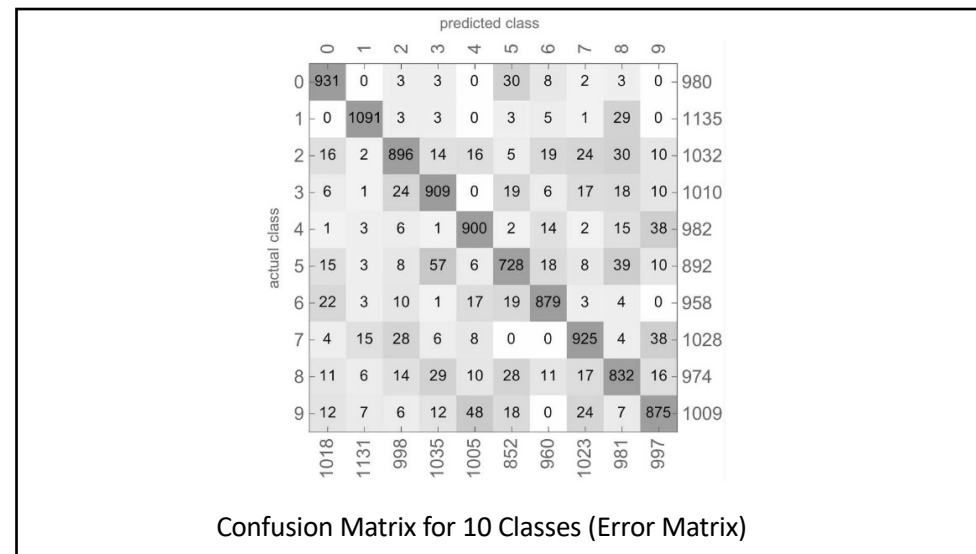
- Accuracy is the % of observations along the diagonal

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0 → 0, 3 → 3, 9 → 9, 0 → 0, 2 → 2, 1 → 1, 1 → 1, 3 → 3, 9 → 9  
 4 → 4, 1 → 1, 2 → 2, 2 → 2, 1 → 1, 4 → 4, 8 → 8, 0 → 0, 4 → 4  
 4 → 4, 7 → 7, 7 → 7, 2 → 2, 9 → 9, 6 → 6, 5 → 5, 5 → 5, 4 → 4  
 8 → 8, 2 → 2, 5 → 5, 9 → 9, 5 → 5, 4 → 4, 1 → 1, 3 → 3, 7 → 7  
 8 → 8, 0 → 0, 7 → 7, 4 → 4, 4 → 4, 7 → 7, 4 → 4, 7 → 7, 9 → 9  
 8 → 8, 9 → 9, 9 → 9, 2 → 2, 2 → 2, 0 → 0, 1 → 1, 6 → 6, 5 → 5  
 4 → 4, 4 → 4, 3 → 3, 9 → 9, 9 → 9, 1 → 1, 1 → 1, 5 → 5, 9 → 9  
 2 → 2, 7 → 7, 0 → 0, 3 → 3, 4 → 4, 7 → 7, 5 → 5, 8 → 8, 7 → 7  
 9 → 9, 0 → 0, 2 → 2, 8 → 8, 1 → 1, 2 → 2, 2 → 2, 7 → 7, 8 → 3

Example of MNIST digits represented by gray images of size 28 × 28

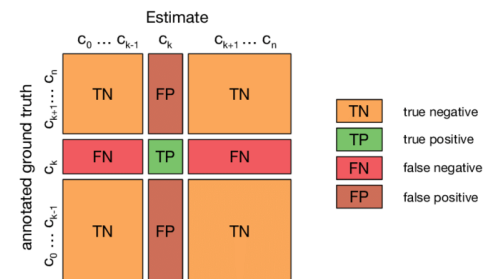
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## Evaluating multiclass classifiers

- Recall/sensitivity, specificity and precision *per class*
  - the target class is seen as positive
  - the negative class is the **union** of the remaining classes



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## Overfitting

- The training data contains information about the regularities in the mapping from input to output, but it also contains noise
- There is sampling error and a flexible architecture can model the sampling error really well
- However, we cannot tell which regularities are real and which are caused by sampling error

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- In general, we try to learn a function  $f : \mathbb{R}^n \rightarrow \mathbb{R}^m$

$$\mathbf{y} = f(\mathbf{x})$$

- that is described by a sample of training data  $D_t$  of the labeled data set

$$D_t = \{(\mathbf{x}_1, \mathbf{y}_1), (\mathbf{x}_2, \mathbf{y}_2), \dots, (\mathbf{x}_N, \mathbf{y}_N)\}$$

- Labels can include multiple things like faces vs. non-faces or man-made objects vs. non-man-made objects

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- After learning, the trained network can be seen as an hypothesis  $h$  that tries to represent the function  $f$  and it can be then used for mapping new examples
- The hypothesis  $h$  should represent the function  $f$  well on the training set. However, ideally, it should generalize from the training data set to unseen **future data points**.
- To try to make sure this is the case, we can validate on an unseen validation (or test set) data set  $D_v$

$$D_v = \{(\mathbf{x}'_1, \mathbf{y}'_1), (\mathbf{x}'_2, \mathbf{y}'_2), \dots, (\mathbf{x}'_M, \mathbf{y}'_M)\} \quad \emptyset = D_t \cap D_v.$$

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## Mean Squared Error (MSE)

- The validation of the model is done by comparing the hypothesis  $h$  outputs

$$\mathbf{o}_k = h(\mathbf{x}'_k)$$

- with the correct values  $\mathbf{y}'_k$  of the validation data set  $D_v$  by the mean squared error

$$MSE_{D_v}(h) = \sum_{k=1}^M \frac{1}{M} \|\mathbf{y}'_k - \mathbf{o}_k\|^2.$$

- The smaller the  $MSE(D_v)$  the better the hypothesis  $h$  describing the function  $f$

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- We can define the mean squared error for the training data set  $D_t$

$$MSE_{D_t}(h) = \frac{1}{N} \cdot \sum_{k=1}^N \|\mathbf{y}_k - \mathbf{o}_k\|^2,$$

- usually

$$MSE_{D_v}(h) > MSE_{D_t}(h).$$

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- If we have two hypothesis  $h_1$  and  $h_2$  with

$$MSE_{D_t}(h_1) < MSE_{D_t}(h_2), \quad MSE_{D_v}(h_1) > MSE_{D_v}(h_2).$$

- then we say that the hypothesis  $h_1$  overfits the training data set  $D_t$ 
  - $h_1$  fits better the training examples than  $h_2$  performs more poorly over examples it didn't learn.
- It seems as if  $h_1$  learned  $D_t$  **by heart** and not the topological structure that describes the function  $f$
- $h_2$  learned the corresponding structure and can **generalize**

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## Cross-Validation

- Estimate the accuracy of a hypothesis induced by a supervised learning algorithm
- Predict the accuracy of a hypothesis over future unseen instances
- Select the optimal hypothesis from a given set of alternative hypotheses
  - Model selection
  - Feature selection
- Combining multiple classifiers (boosting)

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## Holdout Method

- Partition data set  $D = \{(v_1, y_1), \dots, (v_n, y_n)\}$  into *training*  $D_t$  and *validation* set  $D_h = D \setminus D_t$



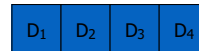
Problems:

- makes insufficient use of data
- training and validation set are correlated

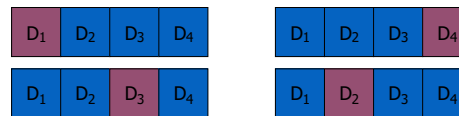
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## Cross-Validation

- $k$ -fold cross-validation splits the data set  $D$  into  $k$  mutually exclusive subsets  $D_1, D_2, \dots, D_k$



- Train and test the learning algorithm  $k$  times, each time it is trained on  $D \setminus D_i$  and tested on  $D_i$



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## Cross-Validation

- Uses all the data for training and testing
- Complete  $k$ -fold cross-validation splits the dataset of size  $m$  in all  $(m \text{ over } m/k)$  possible ways (choosing  $m/k$  instances out of  $m$ )
- Leave  $n$ -out cross-validation sets  $n$  instances aside for testing and uses the remaining ones for training (leave one-out is equivalent to  $n$ -fold cross-validation)
- In stratified cross-validation, the folds are stratified so that they contain approximately the same proportion of labels as the original data set

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- One major drawback of cross-validation is that the number of training runs that must be performed is increased by a factor of  $k$
- How to Evaluate cross-validation for different models ( $h_1, h_2, h_3$ )?
  - We will use t-statistics

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## The logic of hypothesis testing

- Example: toss a coin ten times, observe eight heads. Is the coin fair (i.e., what is its long run behavior?) and what is your residual uncertainty?
- You say, "If the coin were fair, then eight or more heads is pretty unlikely, so I think the coin isn't fair."
- Like proof by contradiction: Assert the opposite (the coin is fair) show that the sample result ( $\geq 8$  heads) has low probability  $p$ , **reject** the assertion, with residual uncertainty related to  $p$ .
- Estimate  $p$  with a *sampling distribution*.

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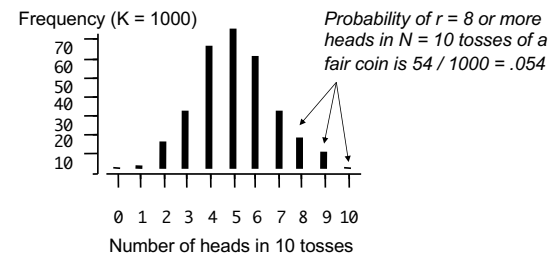
## Probability of a sample result under a null hypothesis

- If the coin were fair ( $p = .5$ , the *null hypothesis*) what is the probability distribution of  $r$ , the number of heads, obtained in  $N$  tosses of a fair coin? Get it analytically or estimate it by simulation (on a computer):

- Loop  $K$  times
  - $r := 0$  //  $r$  is num.heads in  $N$  tosses
  - Loop  $N$  times // simulate the tosses
    - Generate a random  $0 \leq x \leq 1.0$
    - If  $x \geq p$  increment  $r$  //  $p$  is the probability of a head
  - Push  $r$  onto `sampling_distribution`
- Print `sampling_distribution`

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## Sampling distributions



The estimation is constructed by *Monte Carlo sampling*.

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## The t test

- Same logic as the Z test, but appropriate when **population standard deviation** is unknown, samples are small, etc.
- Sampling distribution is t, not normal, but approaches normal as samples size increases
- Test statistic has very similar form but probabilities of the test statistic are obtained by consulting tables of the t distribution, not the normal

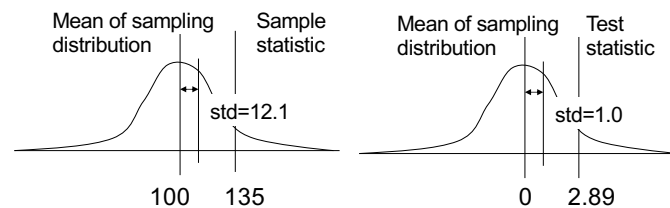
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## The t test

Suppose  $N = 5$  students have mean IQ = 135, std = 27

Estimate the standard deviation of sampling distribution using the sample standard deviation

$$t = \frac{\bar{x} - \mu}{\frac{s}{\sqrt{N}}} = \frac{135 - 100}{\frac{27}{\sqrt{5}}} = \frac{35}{12.1} = 2.89$$



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## $p$ Values

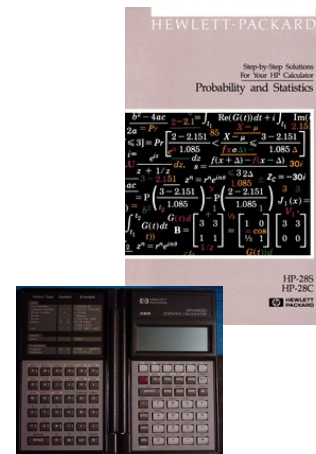
- We find the probabilities by looking them up in tables, or statistics packages provide them
  - The probability of obtaining a particular sample given the null hypothesis is called the  $p$  value
- Commonly we reject the  $H_0$  when the probability of obtaining a *sample statistic* given the null hypothesis is low, say  $p < 0.05$
- The null hypothesis is rejected but might be true

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## Paired Sample t Test

- Given a set of paired observations
  - (from two normal populations)

| A  | B  | $\delta = A - B$ |
|----|----|------------------|
| x1 | y1 | x1-y1            |
| x2 | y2 | x2-y2            |
| x3 | y3 | x3-y3            |
| x4 | y4 | x4-y4            |
| x5 | y5 | x5-y5            |



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- Calculate the mean  $\bar{x}_\delta$  and the standard deviation  $s_\delta$  of the differences  $\delta$
- $H_0: \mu_\delta = 0$  (no difference)
- $H_0: \mu_\delta = k$  (difference is a constant)

$$t_\delta = \frac{\bar{x}_\delta - \mu_\delta}{\hat{\sigma}_\delta} \quad \hat{\sigma}_\delta = \frac{s_\delta}{\sqrt{N_\delta}}$$

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## Paired sample t Test

- We have two rows of data  
94, 86, 12, 90, 66, 40  
10, 20, 22, 26, 6, 18
- Are the two rows significantly different?

$$\delta: 84, 66, -10, 64, 60, 22 \quad \frac{47.6667}{34.8119 / \sqrt{6}} = 3.3540$$

- For five degrees of freedom in t-student table between  $p=0.01$  and  $p=0.02$ , which is less than 0.05, for this reason we have to reject  $H_0$ ! The two rows are significantly different!

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## Paired sample t-test

| Partition Index | Partition Size | Test           | Measure            | Value  |
|-----------------|----------------|----------------|--------------------|--------|
| 1               | 125            | Classification | True Positive      | 79     |
| 2               | 125            | Classification | True Positive      | 79     |
| 3               | 125            | Classification | True Positive      | 72     |
| 4               | 125            | Classification | True Positive      | 80     |
| 5               | 125            | Classification | True Positive      | 75     |
| 6               | 125            | Classification | True Positive      | 81     |
| 7               | 125            | Classification | True Positive      | 64     |
| 8               | 125            | Classification | True Positive      | 72     |
| classifier A    |                |                | Average            | 75.25  |
|                 |                |                | Standard Deviation | 5.3794 |

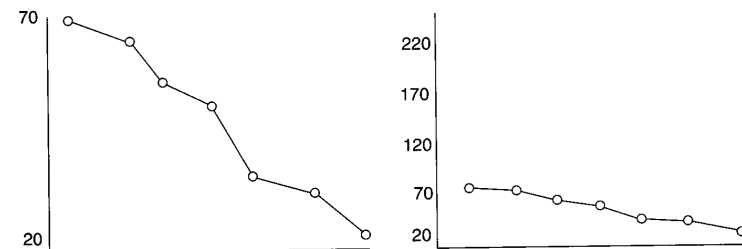
| Partition Index | Partition Size | Test           | Measure            | Value  |
|-----------------|----------------|----------------|--------------------|--------|
| 1               | 125            | Classification | True Positive      | 63     |
| 2               | 125            | Classification | True Positive      | 55     |
| 3               | 125            | Classification | True Positive      | 70     |
| 4               | 125            | Classification | True Positive      | 58     |
| 5               | 125            | Classification | True Positive      | 67     |
| 6               | 125            | Classification | True Positive      | 70     |
| 7               | 125            | Classification | True Positive      | 55     |
| 8               | 125            | Classification | True Positive      | 61     |
| classifier C    |                |                | Average            | 62.375 |
|                 |                |                | Standard Deviation | 5.7866 |

| Partition Index | Partition Size | Test           | Measure            | Value  |
|-----------------|----------------|----------------|--------------------|--------|
| 1               | 125            | Classification | True Positive      | 75     |
| 2               | 125            | Classification | True Positive      | 73     |
| 3               | 125            | Classification | True Positive      | 80     |
| 4               | 125            | Classification | True Positive      | 71     |
| 5               | 125            | Classification | True Positive      | 75     |
| 6               | 125            | Classification | True Positive      | 80     |
| 7               | 125            | Classification | True Positive      | 67     |
| 8               | 125            | Classification | True Positive      | 77     |
| classifier B    |                |                | Average            | 74.75  |
|                 |                |                | Standard Deviation | 4.1458 |

using Cross Validation, determine if the classifier A, B, C are significantly different  
Compare (A,B), (A,C), (B,C)

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## Confidence Intervals



- Just looking at a figure representing the mean values, we can not see if the differences are significant

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## Confidence Intervals ( $\sigma$ known)

- Standard error from the standard deviation

$$\sigma_{\bar{x}} = \frac{\sigma_{Population}}{\sqrt{N}}$$

- 95 Percent confidence interval for normal distribution is about the mean

$$\bar{x} \pm 1.96 \cdot \sigma_{\bar{x}}$$

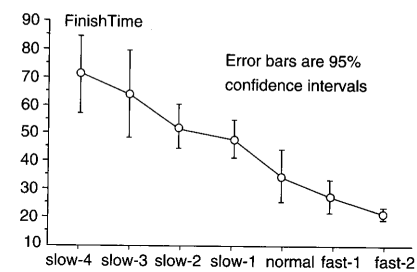
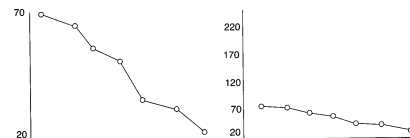
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## Confidence interval when ( $\sigma$ unknown)

- Standard error from the sample standard deviation
- 95 Percent confidence interval for t distribution ( $t_{0.025}$  from a table) is

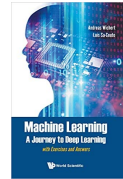
$$\bar{x} \pm t_{0.025} \cdot \hat{\sigma}_{\bar{x}}$$

$$\hat{\sigma}_{\bar{x}} = \frac{s}{\sqrt{N}}$$



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## Literature



- Machine Learning - A Journey to Deep Learning, A. Wichert, Luis Sa-Couto, World Scientific, 2021
  - Chapter 8