

ROC Curves

Miguel Angel
Lozano

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Plotting a
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Analyzing a
curve

An example:
Kernelized
Graph
Matching

ROC Curves

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ROC Curves

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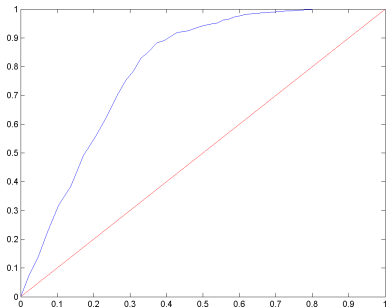
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An example:
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- Receiver Operating Characteristics curves
- Useful for evaluating the quality of a test
- E.g. a test for matching two points
- Is part of a field called *Signal Detection Theory*



History

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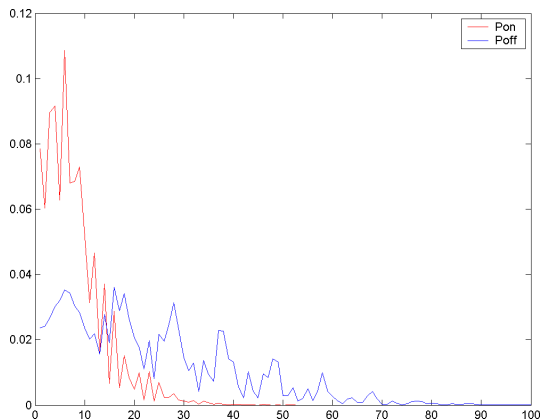
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- Developed during World War II for the analysis of radar images.
- Radar operators had to decide whether a blip on the screen represented an enemy target, a friendly ship, or just noise.
- Signal detection theory measures the ability of radar receiver operators to decide whether a blip on the screen represents an enemy, a friend, or noise. Their ability to do so was called the Receiver Operating Characteristics.
- It was not until the 1970's that signal detection theory was recognized as useful for interpreting medical test results.

Probability density functions

- Given a continuous value X returned by our test, we can define the following PDFs:
 - $P(X|on)$
 - $P(X|off)$



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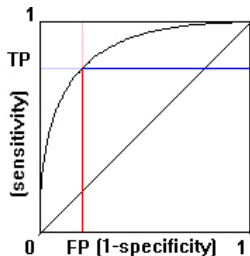
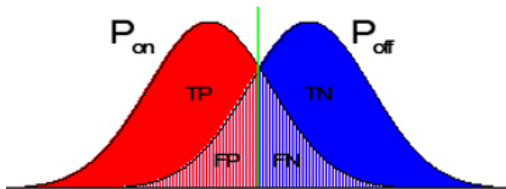
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Sensitivity and Specificity

- **TP** - True positive fraction (*sensitivity*)
- **FP** - False positive fraction
- **TN** - True negative fraction (*specificity*)
- **FN** - False negative fraction

$$TP = 1 - FN \quad (1)$$

$$TN = 1 - FP \quad (2)$$



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Curve points

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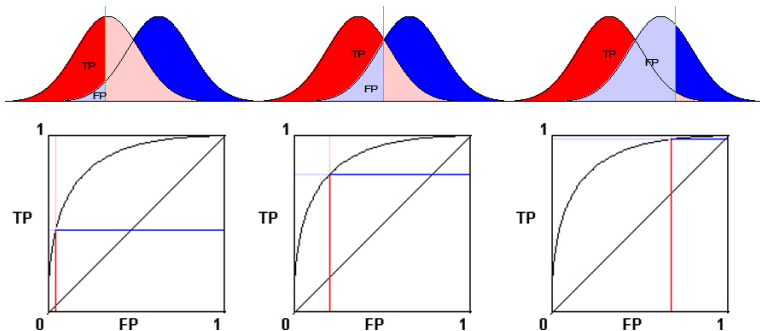
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- Each point from the curve represents a threshold



Plotting a curve

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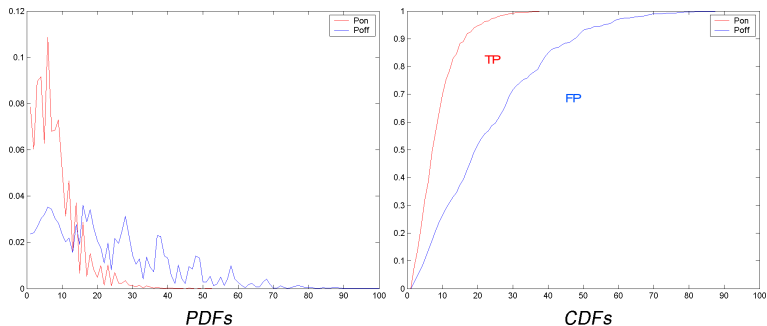
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- Obtain the CDFs corresponding to PDFs $P(X|on)$ and $P(X|off)$
 - The CDF from $P(X|on)$ corresponds to **TP**
 - The CDF from $P(X|off)$ corresponds to **FP**



Plotting a curve

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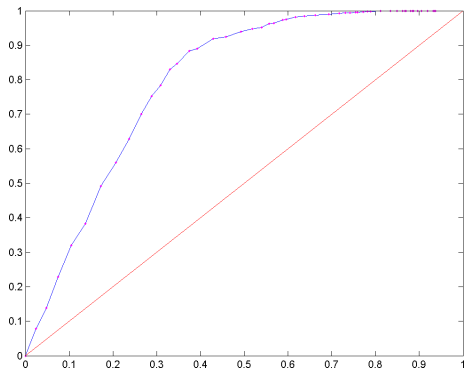
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- x coordinates are taken from the $P(X|off)$ CDF (**FP**)
- y coordinates are taken from the $P(X|on)$ CDF (**TP**)



ROC curves are useful for ...

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- Measuring test accuracy.
 - Area under the curve (**AUC**).
 - AUC is non-parametrical.
 - Probability of classifying successfully both a positive and a negative example.
- Comparing two tests.
 - Compare the shapes of their corresponding curves.
- Choosing a suitable threshold.
 - A point near to the upper left corner.

Analyzing results

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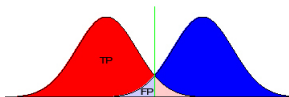
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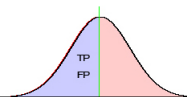
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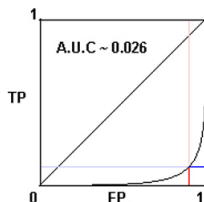
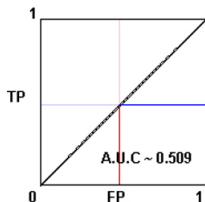
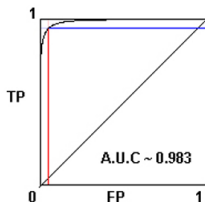
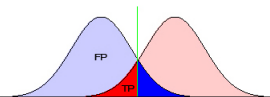
A good test



Flip a coin!



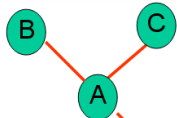
Wrong polarity



Adjacency matrices

- Two undirected and unweighted graphs G_X and G_Y are defined by their adjacency matrices.

$$G_X = (V_X, E_X)$$



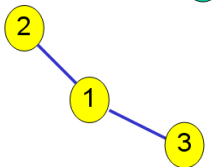
$$V_X = \{A, B, C, D\}$$

$$E_X = \{(A, B), (A, C), (A, D)\}$$

$$X_{ab} = \begin{cases} 1 & \text{if } (a, b) \in E_X \\ 0 & \text{otherwise} \end{cases}$$

$$\mathbf{X} = \begin{bmatrix} 0 & 1 & 1 & 1 \\ 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 \end{bmatrix}$$

$$G_Y = (V_Y, E_Y)$$



$$V_Y = \{1, 2, 3\}$$

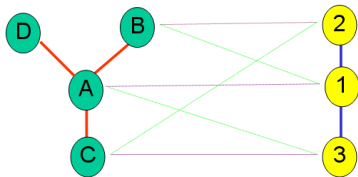
$$E_Y = \{(1, 2), (1, 3)\}$$

$$Y_{ij} = \begin{cases} 1 & \text{if } (i, j) \in E_Y \\ 0 & \text{otherwise} \end{cases}$$

$$\mathbf{Y} = \begin{bmatrix} 0 & 1 & 1 \\ 1 & 0 & 0 \\ 1 & 0 & 0 \end{bmatrix}$$

Matching matrices

- Encode the correspondence between vertices in the first graph (rows) and vertices in the second one (columns).
 - The solutions to matching.
- Constraints:
 - A given vertex in the first graph can match either a unique vertex in the second or none of them, and viceversa.



$$\mathbf{M}_1 = \begin{matrix} & \begin{matrix} 1 & 2 & 3 \end{matrix} \\ \begin{matrix} A \\ B \\ C \\ D \end{matrix} & \begin{bmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \\ 0 & 0 & 0 \end{bmatrix} \end{matrix} \quad \mathbf{M}_2 = \begin{bmatrix} 0 & 0 & 1 \\ 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 0 \end{bmatrix}$$

$$\forall a \sum_{i=1}^I M_{ai} \leq 1, \quad \forall i \sum_{a=1}^A M_{ai} \leq 1, \\ \forall ai M_{ai} \in \{0, 1\}$$

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Gold & Rangarajan

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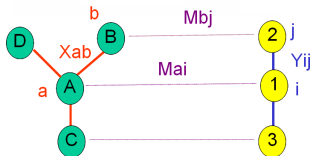
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- Gold & Rangarajan proposed a cost function for quantifying the rectangle rule.

$$F(M) = \sum_{a=1}^A \sum_{i=1}^I \sum_{b=1}^A \sum_{j=1}^I M_{ai} M_{bj} \underbrace{X_{ab} Y_{ij}}_{C_{abij}} \quad (3)$$



- G&R proposed a continuation method for maximizing $F(M)$ which updates all assignments iteratively.

Diffusion Kernels

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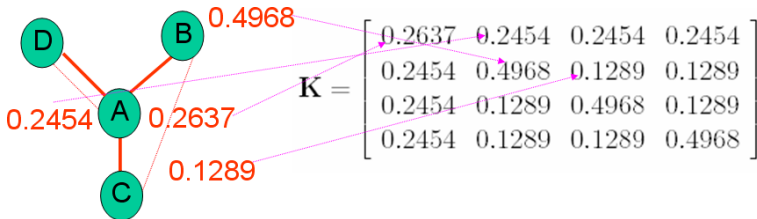
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- We use the values from the graph kernel as attributes for non-attributed graphs.



- Probabilistic interpretation:
 - K_{ij} is the probability that a lazy random walk starting at vertex i reaches vertex j . Such a probability decays with the distance between i and j .

Kernelized cost function

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- The G&R cost function is weighted by information coming from distributional attributes:

$$F(M) = \sum_{a=1}^A \sum_{i=1}^I \sum_{b=1}^A \sum_{j=1}^I M_{ai} M_{bj} C_{abij} \quad (4)$$

$$C_{abij} = X_{ab} Y_{ij} \exp -[(H_a^{K_x} - H_i^{K_y})^2 + (H_b^{K_x} - H_j^{K_y})^2] \quad (5)$$

Attribute comparative

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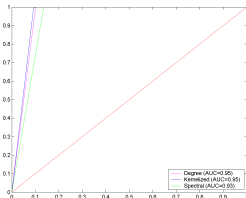
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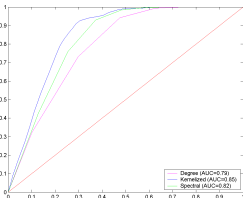
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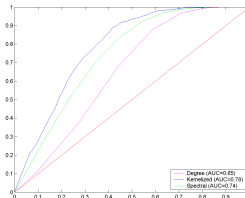
An example: Kernelized Graph Matching



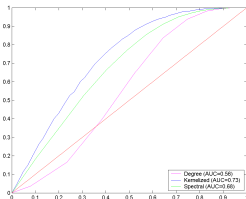
Node noise: 0%



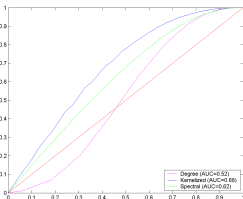
Node noise: 10%



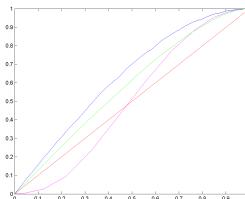
Node noise: 20%



Node noise: 30%



Node noise: 40%



Node noise: 50%