

# Genuine and Imposter Scores

Example: 2 users, A & B, each has 5 samples ( $A_i$ ,  $B_j$  are feature vectors)

**Genuine Scores:** (Similarity measure,  $S$ )

User A:

$S_A$	$A_1$	$A_2$	$A_3$	$A_4$	$A_5$
$A_1$	$\infty$	$a_{12}$	$a_{13}$	$a_{14}$	$a_{15}$
$A_2$	$a_{21}$	$\infty$	$a_{23}$	$a_{24}$	$a_{25}$
$A_3$	$a_{31}$	$a_{32}$	$\infty$	$a_{34}$	$a_{35}$
$A_4$	$a_{41}$	$a_{42}$	$a_{43}$	$\infty$	$a_{45}$
$A_5$	$a_{51}$	$a_{52}$	$a_{53}$	$a_{54}$	$\infty$

User B:

$S_B$	$B_1$	$B_2$	$B_3$	$B_4$	$B_5$
$B_1$	$\infty$	$b_{12}$	$b_{13}$	$b_{14}$	$b_{15}$
$B_2$	$b_{21}$	$\infty$	$b_{23}$	$b_{24}$	$b_{25}$
$B_3$	$b_{31}$	$b_{32}$	$\infty$	$b_{34}$	$b_{35}$
$B_4$	$b_{41}$	$b_{42}$	$b_{43}$	$\infty$	$b_{45}$
$B_5$	$b_{51}$	$b_{52}$	$b_{53}$	$b_{54}$	$\infty$

**Imposter Scores:**

A & B:

$S_{AB}$	$B_1$	$B_2$	$B_3$	$B_4$	$B_5$
$A_1$	$ab_{11}$	$ab_{12}$	$ab_{13}$	$ab_{14}$	$ab_{15}$
$A_2$	$ab_{21}$	$ab_{22}$	$ab_{23}$	$ab_{24}$	$ab_{25}$
$A_3$	$ab_{31}$	$ab_{32}$	$ab_{33}$	$ab_{34}$	$ab_{35}$
$A_4$	$ab_{41}$	$ab_{42}$	$ab_{43}$	$ab_{44}$	$ab_{45}$
$A_5$	$ab_{51}$	$ab_{52}$	$ab_{53}$	$ab_{54}$	$ab_{55}$

# Genuine and Imposter Scores

Set  $X = \{X_1, \dots, X_M\}$ , is a set of  $M$  match (genuine) scores.

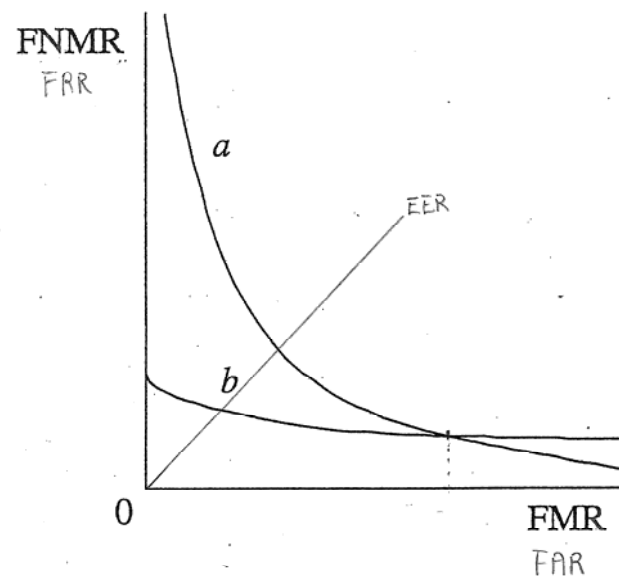
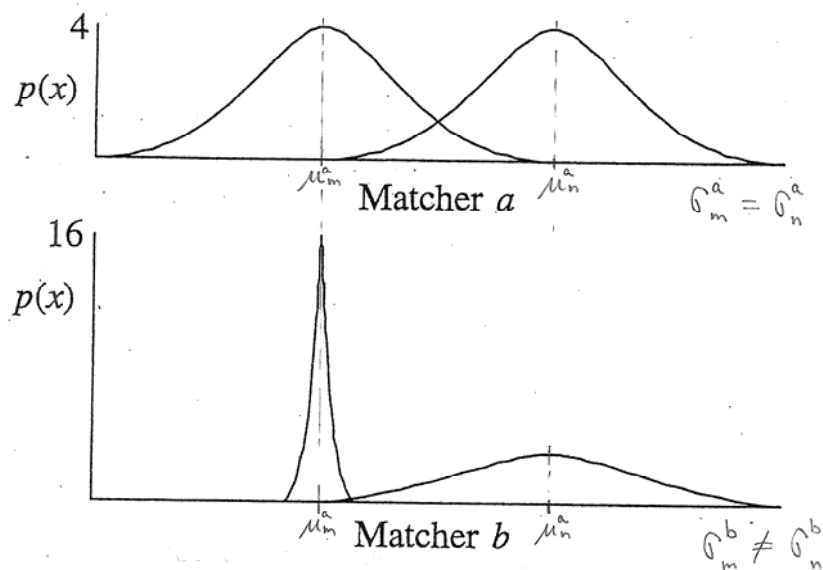
Example:  $\{a_{12}, a_{13}, \dots, a_{45}, b_{12}, b_{13}, \dots, b_{45}\}$ ,  $M = 20$

Set  $Y = \{Y_1, \dots, Y_N\}$ , is a set of  $N$  non match (imposter) scores.

Example:  $\{ab_{11}, ab_{12}, \dots, ab_{55}\}$ ,  $N = 25$

# d-prime Value

Ref. [2] p. 76

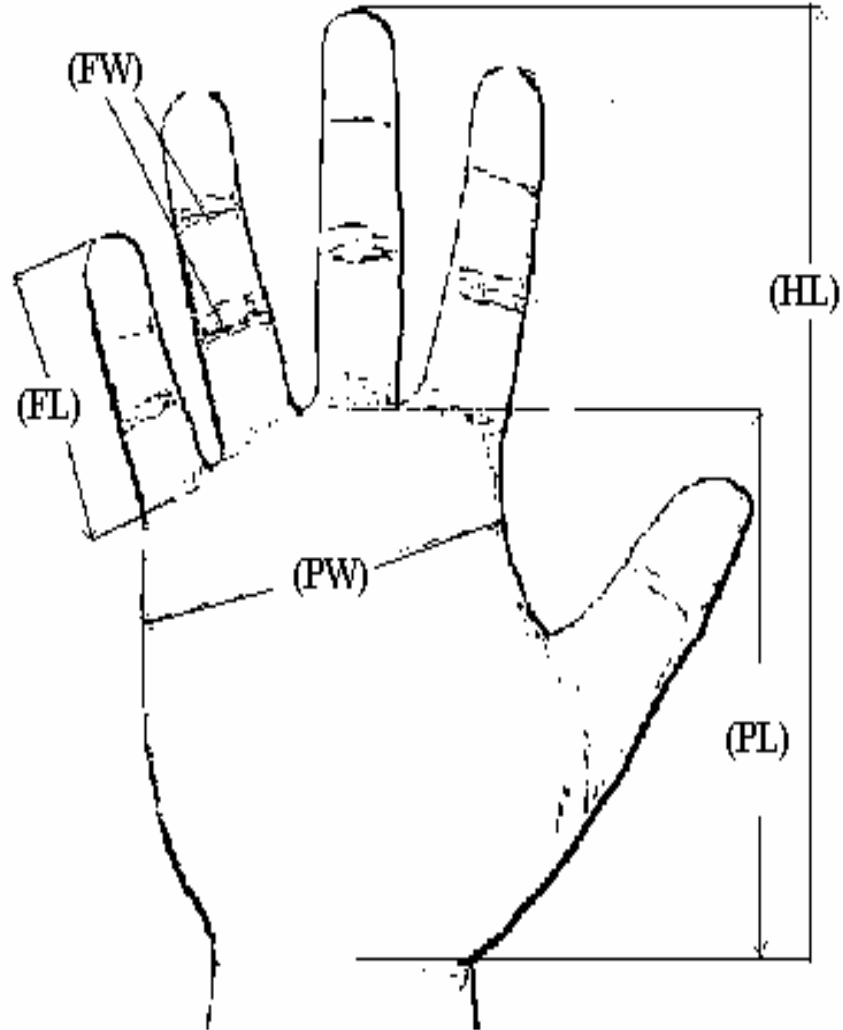


These two systems (matchers) have the same d-prime values  
But their ROC are quite different. Which is better?

# Representation: Hand Geometry features

## Vector of 17 features

- Finger length (FL) – 4
- Finger width (FW) – 8
- Palm width (PW)
- Palm Length (PL)
- Hand Length (HL)
- Hand Contour (HC)
- Hand Area (HA)



# Examples of Feature Vector – Hand Geometry

## User 1:

### **Sample1:**

320,365,354,292,113,89,111,89,112,95,103,91,914,470,584,4275,413574

### **Sample2:**

320,378,361,298,113,89,109,90,113,93,105,90,926,469,590,4478,426774

## User 2:

### **Sample1:**

346,387,349,278,92,80,95,87,86,78,78,70,868,428,538,4279,365147

### **Sample2:**

335,382,349,277,93,84,93,85,87,78,78,69,871,432,542,4241,367027

## User 3:

### **Sample1:**

315,402,380,294,97,79,94,89,88,82,80,75,911,451,544,4519,396858

### **Sample2:**

334,400,382,298,92,80,95,90,97,87,80,75,884,444,519,4421,383321

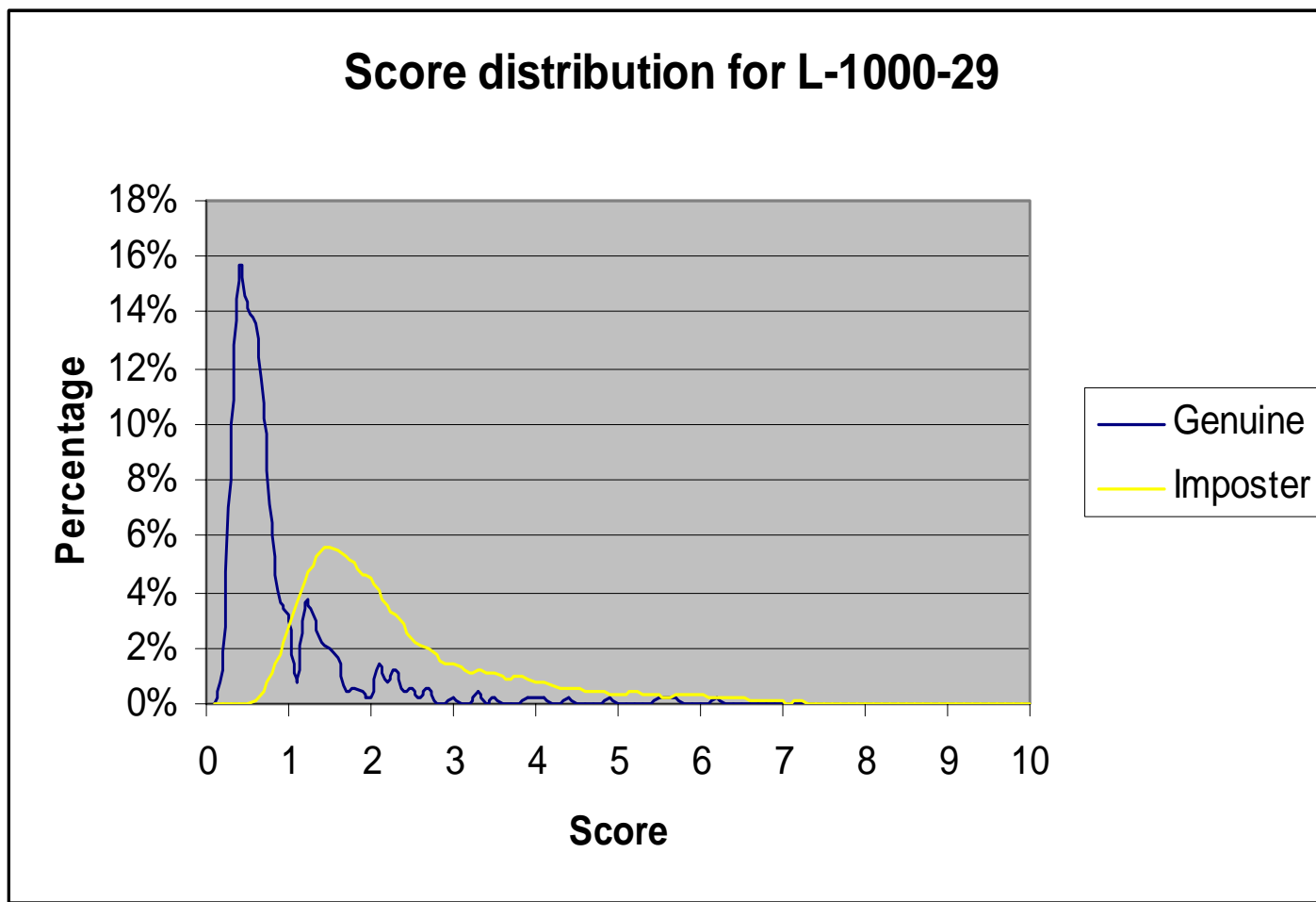
## User 4:

314,371,359,294,92,77,99,82,107,88,87,73,875,445,544,4175,365795

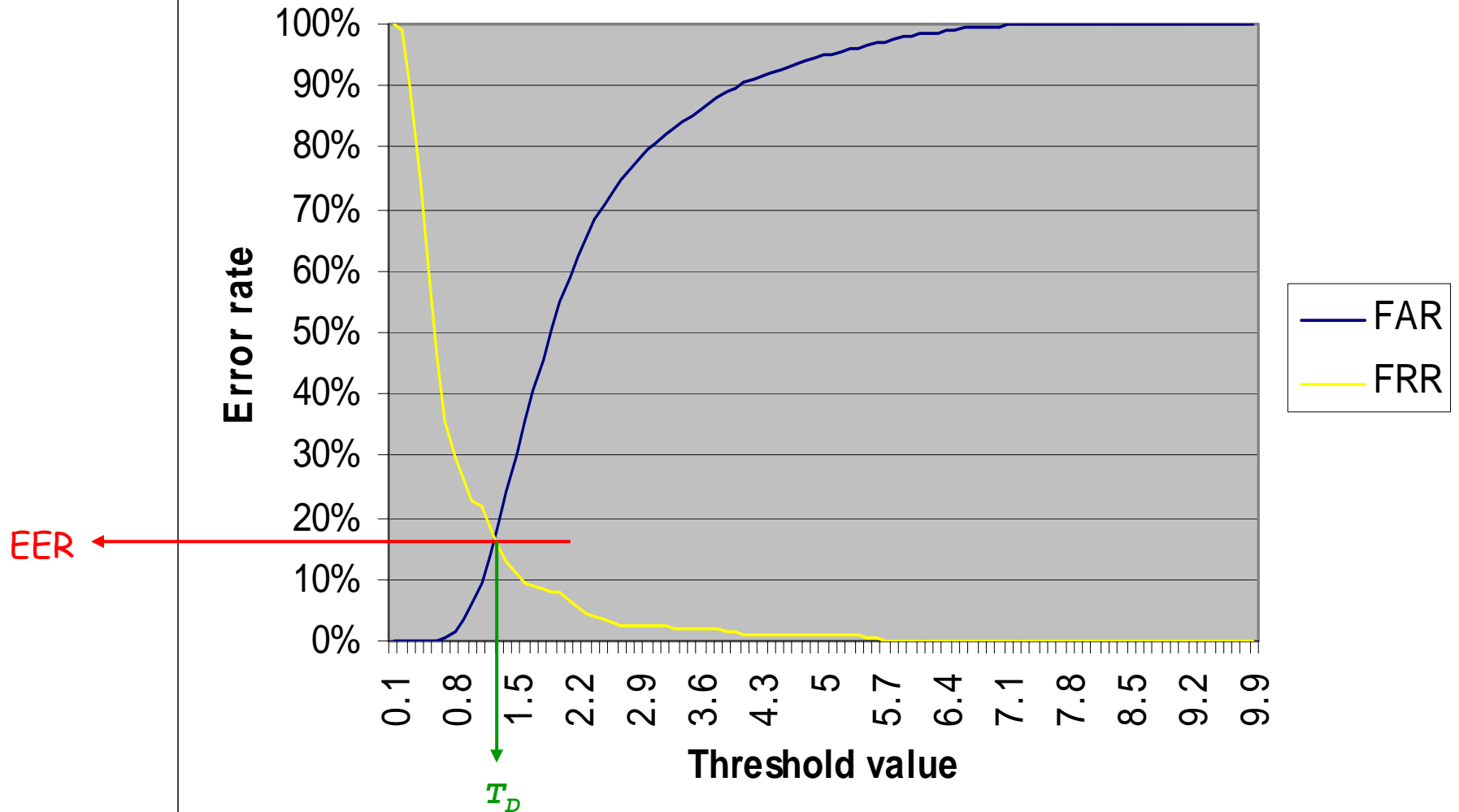
313,375,361,294,94,79,102,84,104,88,87,72,879,447,545,4187,368345

# Matching Score Distribution

Distance Measure - 100 users, 10 samples each,  
29 features



## FAR\_FRR for L-1000-29



# Estimation of the no. of scores from a given set of data

For a given set of feature vector, we have assumed

- set  $X = \{X_1, \dots, X_M\}$ , is a set of  $M$  match (genuine) scores.
- set  $Y = \{Y_1, \dots, Y_N\}$ , is a set of  $N$  non match (imposter) scores.

Q: What are the values of  $M$  and  $N$ , say for a database of 100 users and 10 samples each?



# Estimation of the no. of scores from a given set of data

For a given set of feature vector, we have assumed

- set  $X = \{X_1, \dots, X_M\}$ , is a set of  $M$  match (genuine) scores.
- set  $Y = \{Y_1, \dots, Y_N\}$ , is a set of  $N$  non match (imposter) scores.

For 100 users, 10  
samples each,  $M$  :

$$100 \times C_{10}^2 = 100 \times \frac{(10)!}{(10-2)!(2)!} = 4,500$$

For  $Q$  users,  $P$   
samples each,  $M$  :

$$Q \times C_P^2 = Q \times \frac{(P)!}{(P-2)!(2)!}$$

For 100 users, 10  
samples each,  $N$ :

$$C_{100}^2 \times 10 \times 10 = \frac{(100)!}{(100-2)!(2)!} \times 100 = 495,000$$

For  $Q$  users,  $P$   
samples each,  $N$ :

$$C_Q^2 \times P \times P = \frac{(Q)!}{(Q-2)!(2)!} \times P^2$$

For  $Q$  users,  $P$   
samples each,  $M$ :

$$C_{Q \times P}^2 - M$$