

# ISO/IEC 30136

## Performance Testing of Template Protection Schemes

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- Since 2023 (before @UAM, h\_da, HSAN)
- Research on (but not limited to!):
  - ❖ Different biometric modalities, and multi-biometrics
  - ❖ Biometric Template Protection and Attack Detection
  - ❖ Synthetic data
  - ❖ Explainability

➤ Chair BIOSIG

➤ Involved in EAB and ISO/IEC SC 37

➤ More details on:

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## BioML Lab

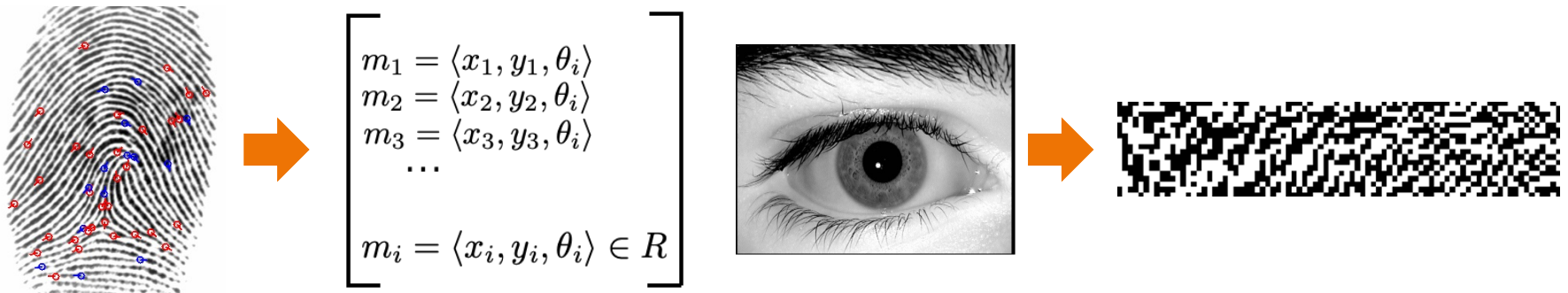
# Introduction



- Biometric data is classified as **sensitive personal data** by the European Data Protection Regulation (GDPR)
- Different legislations across the globe, adding more protection:
  - ❖ Japan Act on the Protection of Personal Information (APPI) also includes extra-territorial protection since 2022
  - ❖ Australian Privacy Act is undergoing reforms
  - ❖ The California Consumer Privacy Act (CCPA), based on the GDPR, aims to empower consumers with new rights in order to protect their privacy
  - ❖ ...



- It was a common belief that the stored templates revealed no information about the biometric characteristics:



- However, biometric samples can be recovered from the stored unprotected templates

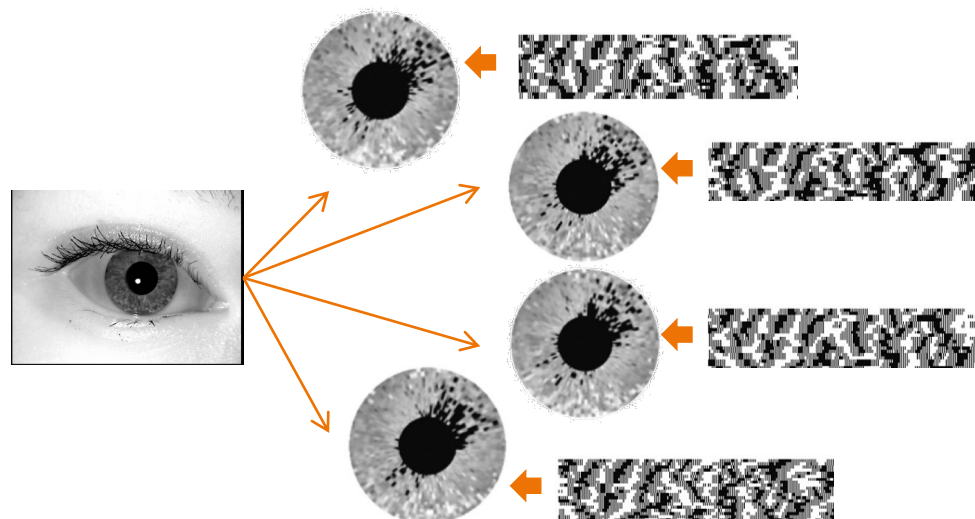
- Based on the HC algorithms, we can reconstruct biometric samples:



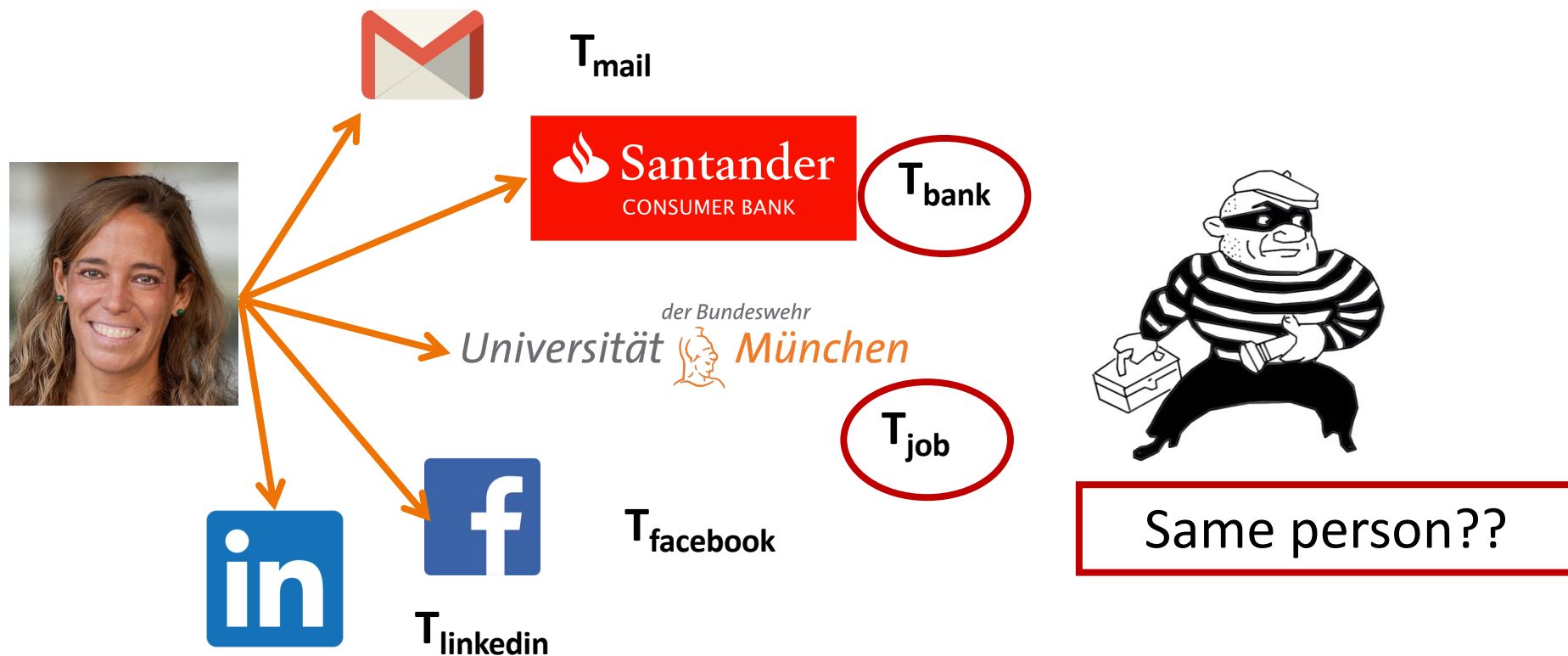
[M. Gomez-Barrero *et al.*, *Int. Conf. on Biometrics*, 2012]

[M. Gomez-Barrero *et al.*, *Information Sciences*, 2014]

[J. Galbally, *et al.*, *Computer Vision & Image Understanding*, 2013]

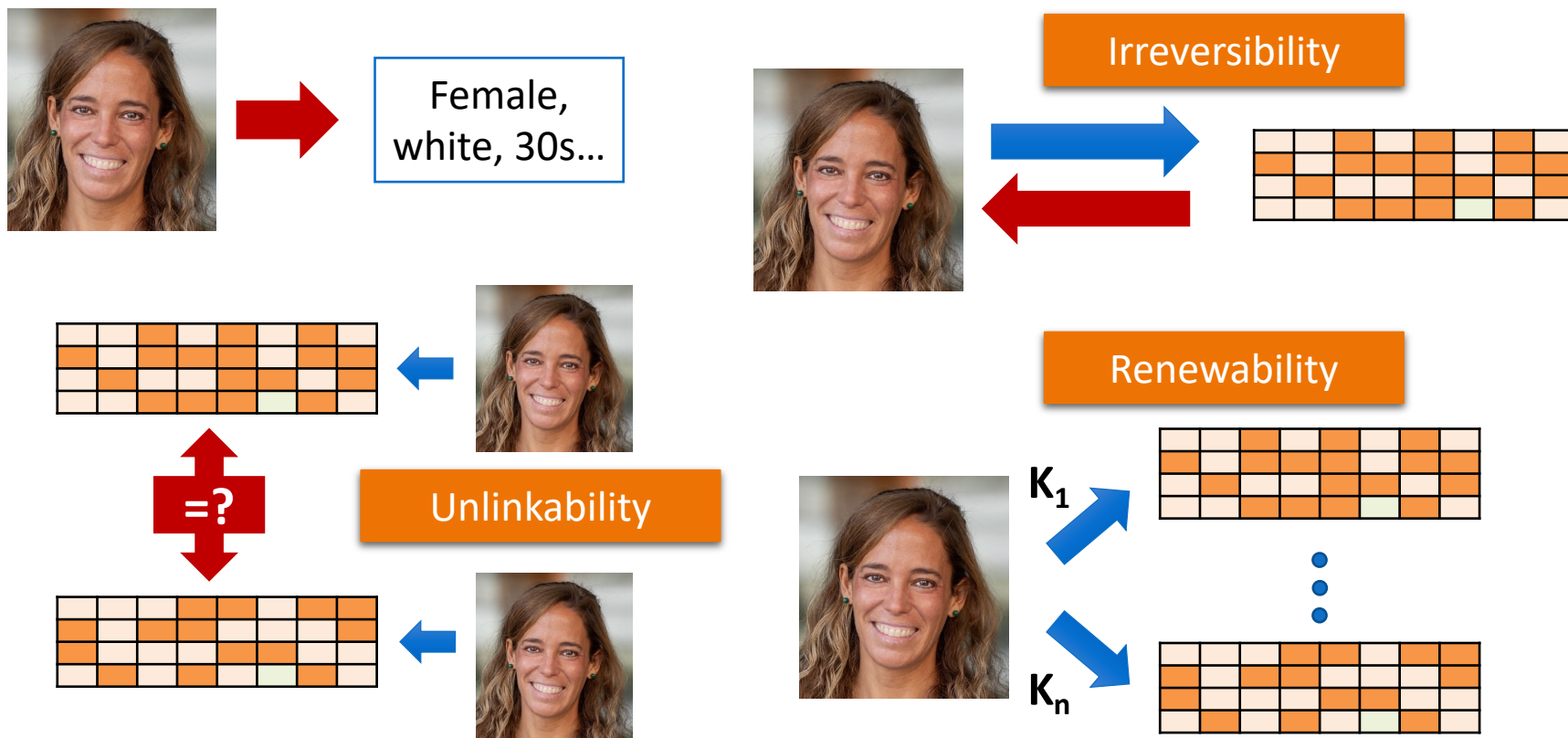


- We can enroll with a single instance in different applications



Templates need to be protected, so that no one can find out on which applications we are enrolled

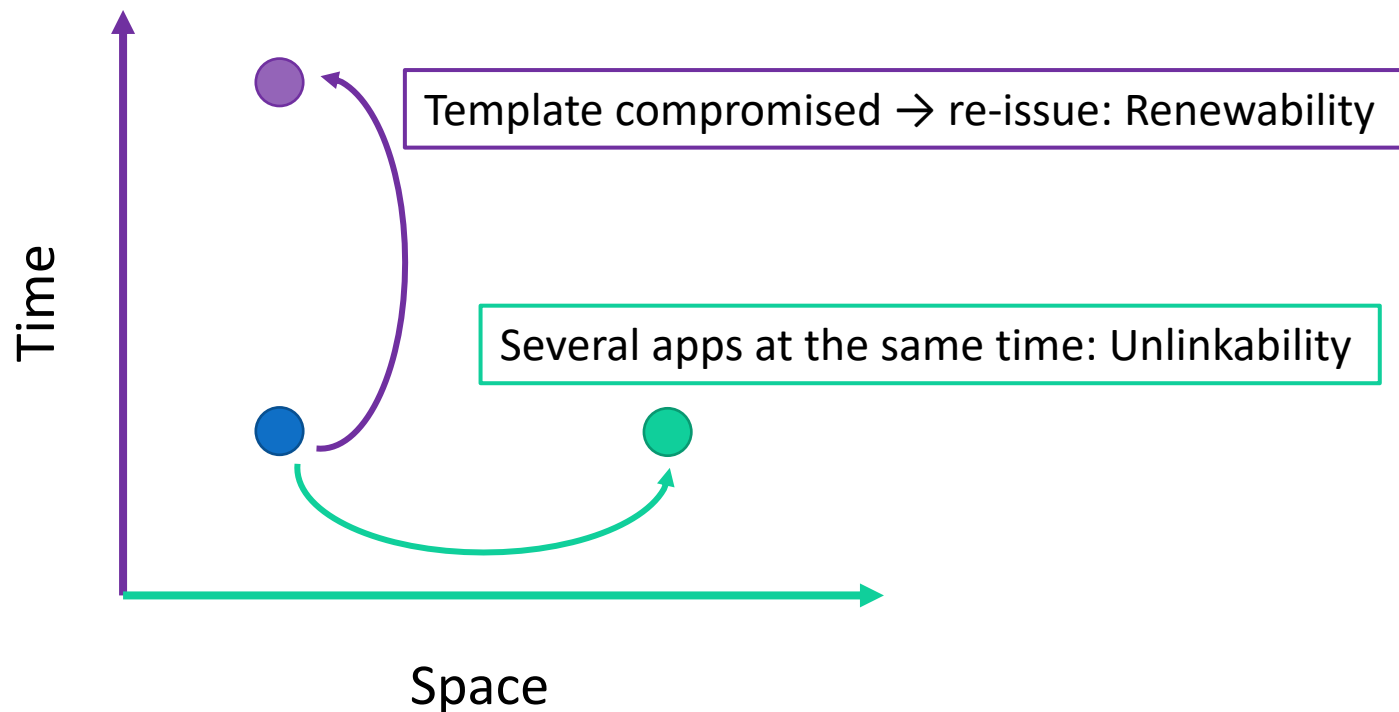
[ISO/IEC IS 24745 on Biometric Information Protection]

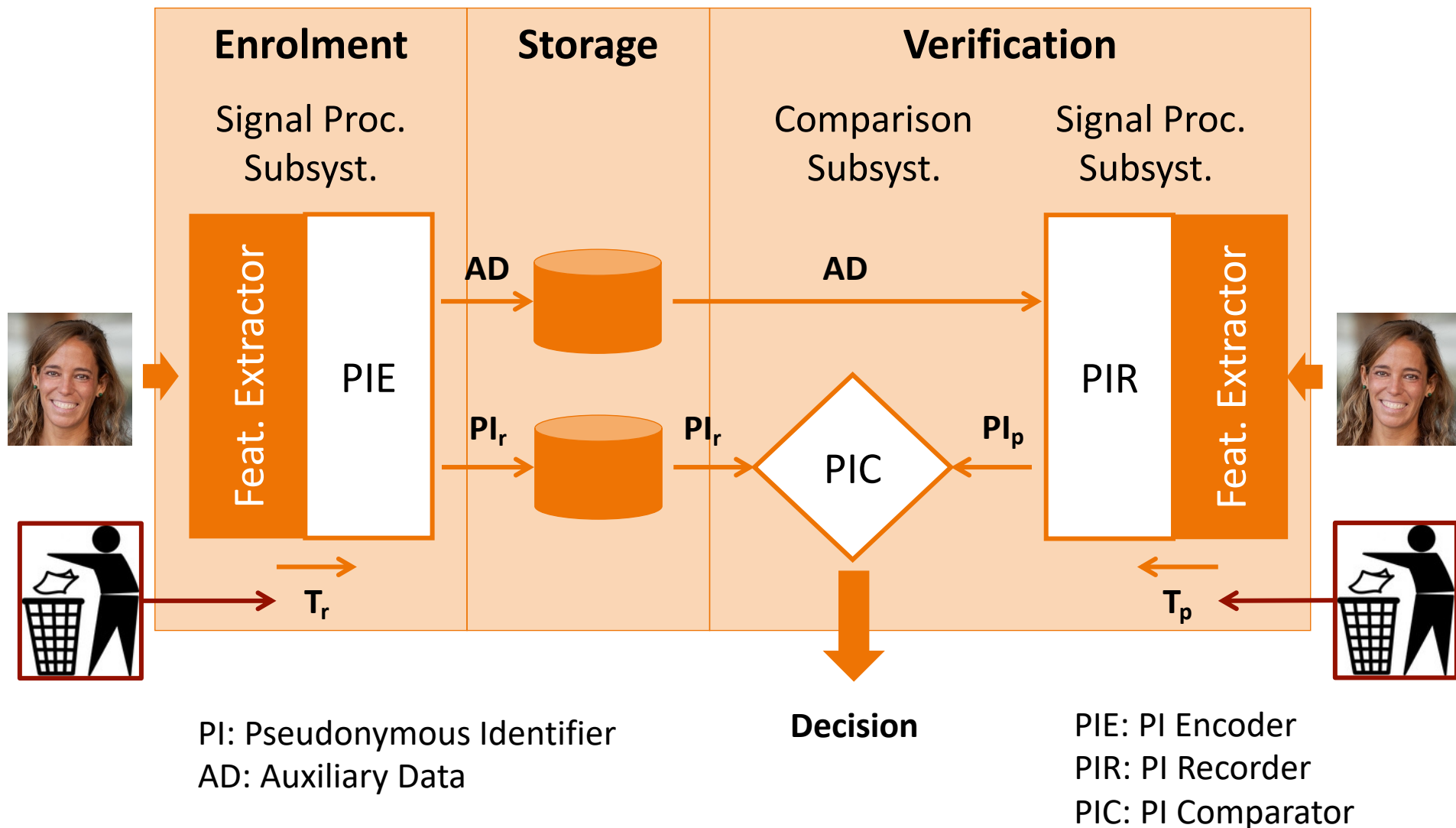


Accuracy, template size and verification speed must be preserved.



➤ Diversification in space and time





# Security and Privacy Evaluation

## Reproducible Evaluation

**BTP System**

**Biometric  
Dataset**

**Threat Model**

**Evaluation  
Protocol**

## ISO Requirements

**Analysis 1:  
Accuracy**

**Analysis 2:  
Irreversibility**

**Analysis 3:  
Unlinkability**

**Analysis 4:  
Computational Load Increase**

- We need to describe the attacker's capabilities and skills:
  - ❖ **Naive model:**
    - No knowledge about the system, no access to large DBs
    - Access to RBRs + black-box system
  - ❖ **Collision model:**
    - Access to large DBs
    - Access to a BTP system that generates similar PIs to the system under attack
  - ❖ **General model:**
    - Full knowledge about the system, statistical properties of the enrolled data, access to protected data
    - **Standard model:** secret key remains secure
    - **Advanced model:** chosen-plaintext and chosen-ciphertext attack
    - **Full disclosure model:** secret key compromised

- Most BTP schemes transform either the sample (e.g. surface folding) or the template (e.g., fuzzy vault)
- That leads to the addition of noise or information loss, which in turn leads to a decrease in accuracy
- We need to assess such performance loss in accordance with the ISO/IEC 19795:
  - ❖ Compute FMR and FNMR for the baseline system AND the BTP scheme
  - ❖ Following a common experimental protocol
  - ❖ Compare in terms of DET plots
    - The Equal Error Rate (EER), where  $FMR = FNMR$ , is not enough!!

## ➤ Theoretical evaluation:

- ❖ “Proving a computational or an information-theoretic property limiting the success probability or the advantage of any computationally-bounded adversary”
- ❖ Example for unlinkability: *indistinguishability game describes the advantage of an attacker with respect to a perfect indistinguishable system*

## ➤ Empirical evaluation:

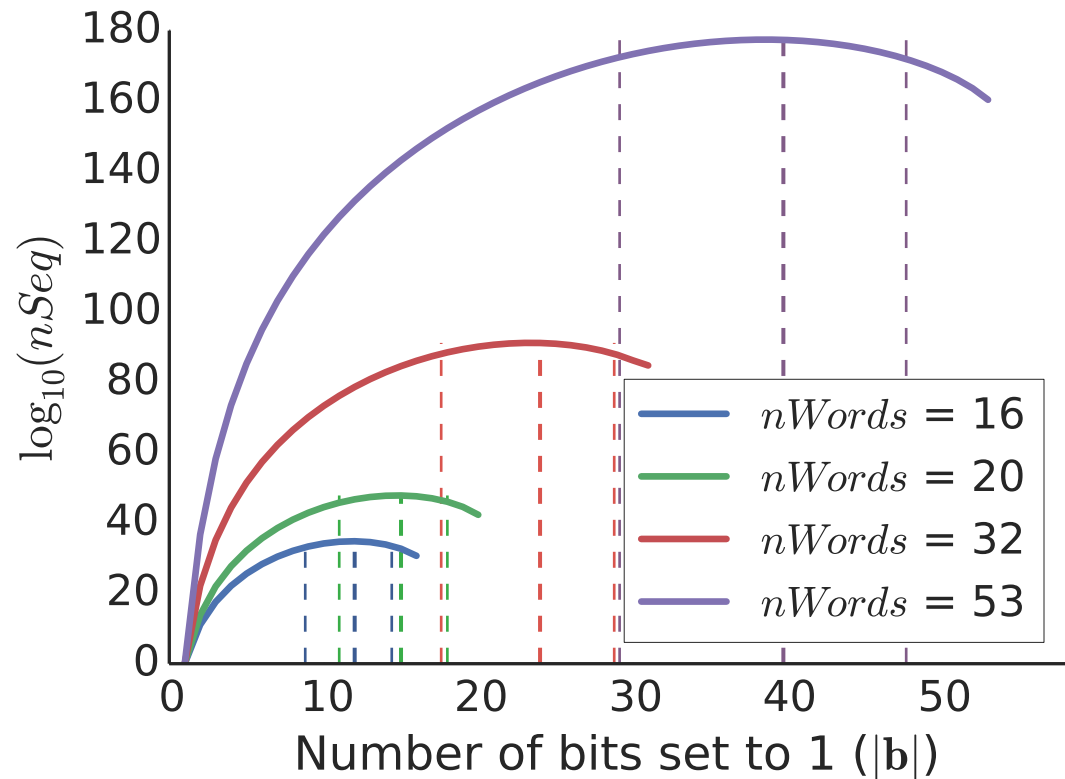
- ❖ “The experimenter shall specify an empirical method of demonstrating irreversibility / unlinkability, for example, empirically estimating the success probability or the advantage over a random guess for some specific adversary in an attack scenario”

[ISO/IEC IS 30136 on Performance Testing of BTP Schemes]

- How can we analyse irreversibility? Following cryptographic paradigms?
- Careful! Some assumptions are not valid:
  - ❖ Uniformity of data – neighbouring bits are correlated!!
  - ❖ In fact, some biometric templates (e.g., finger vein or fingerprint minutiae spectral representation) are compared in terms of their cross correlation!
  - ❖ There are also symmetries
- Therefore, we need to model such correlations and take them into account in the computations
  - ❖ No general method proposed so far

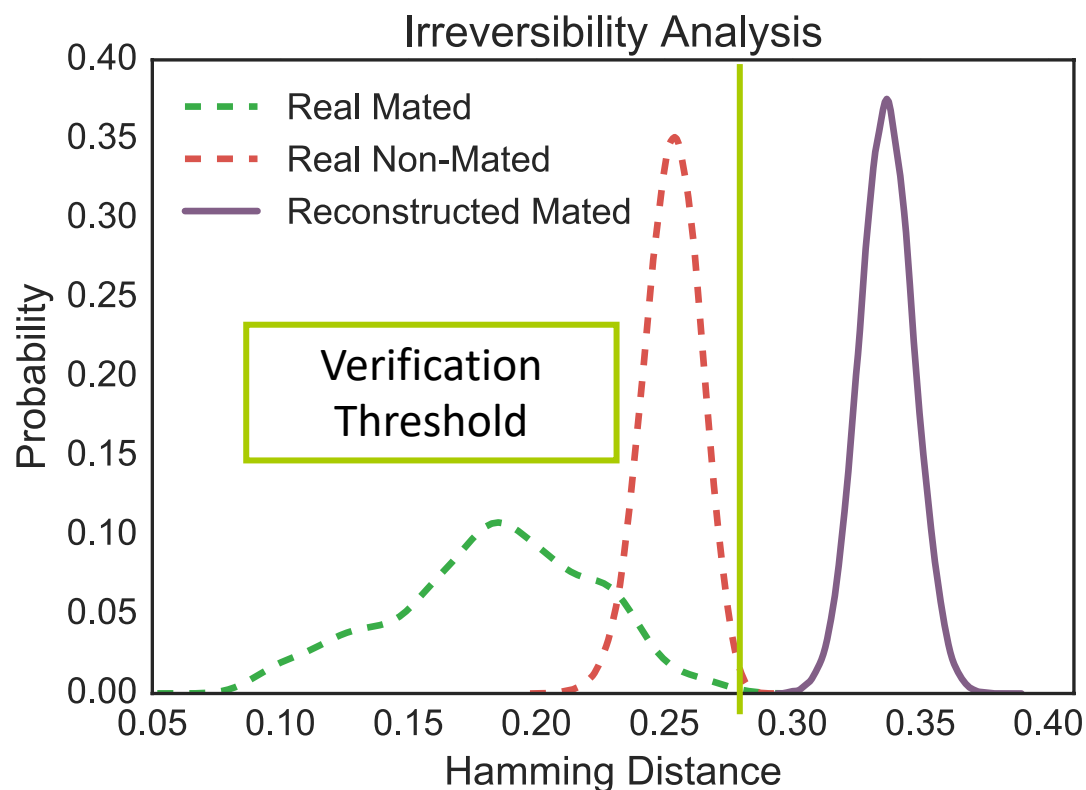


- $nSeq$ : number of original sequences leading to a single template.
- For a full disclosure model, the probability of a reconstruction is  $2^{-40,960}$



[Gomez-Barrero et al., Information Sciences 2016]

- Are the reconstructed unprotected templates similar to the original ones?



**Irreversible:** HD  
bigger than impostor  
comparisons

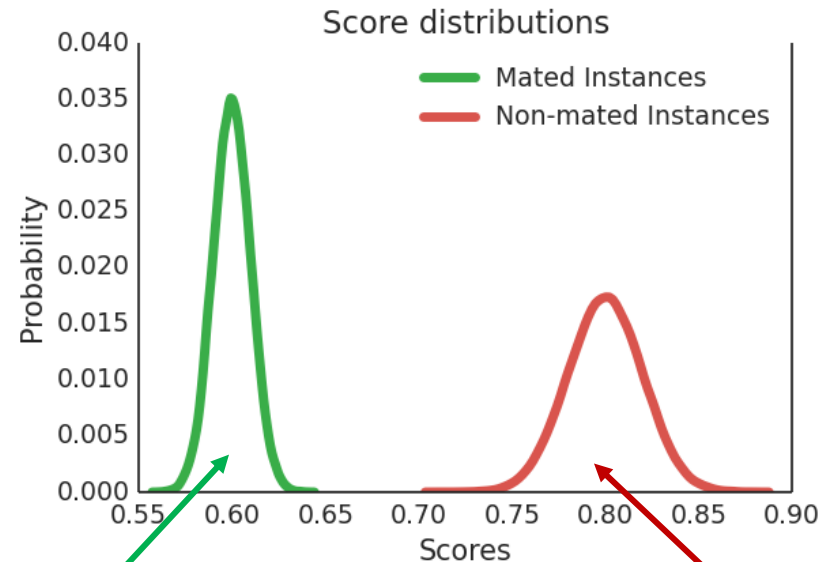
[Bringer *et al.*, ICB 2015]



$T_{\text{job}}$     $T_{\text{bank}}$



$$s = LF(T_{\text{job}}, T_{\text{bank}})$$



$s$  here → success!! 😊

$s$  here → try again!! ☹

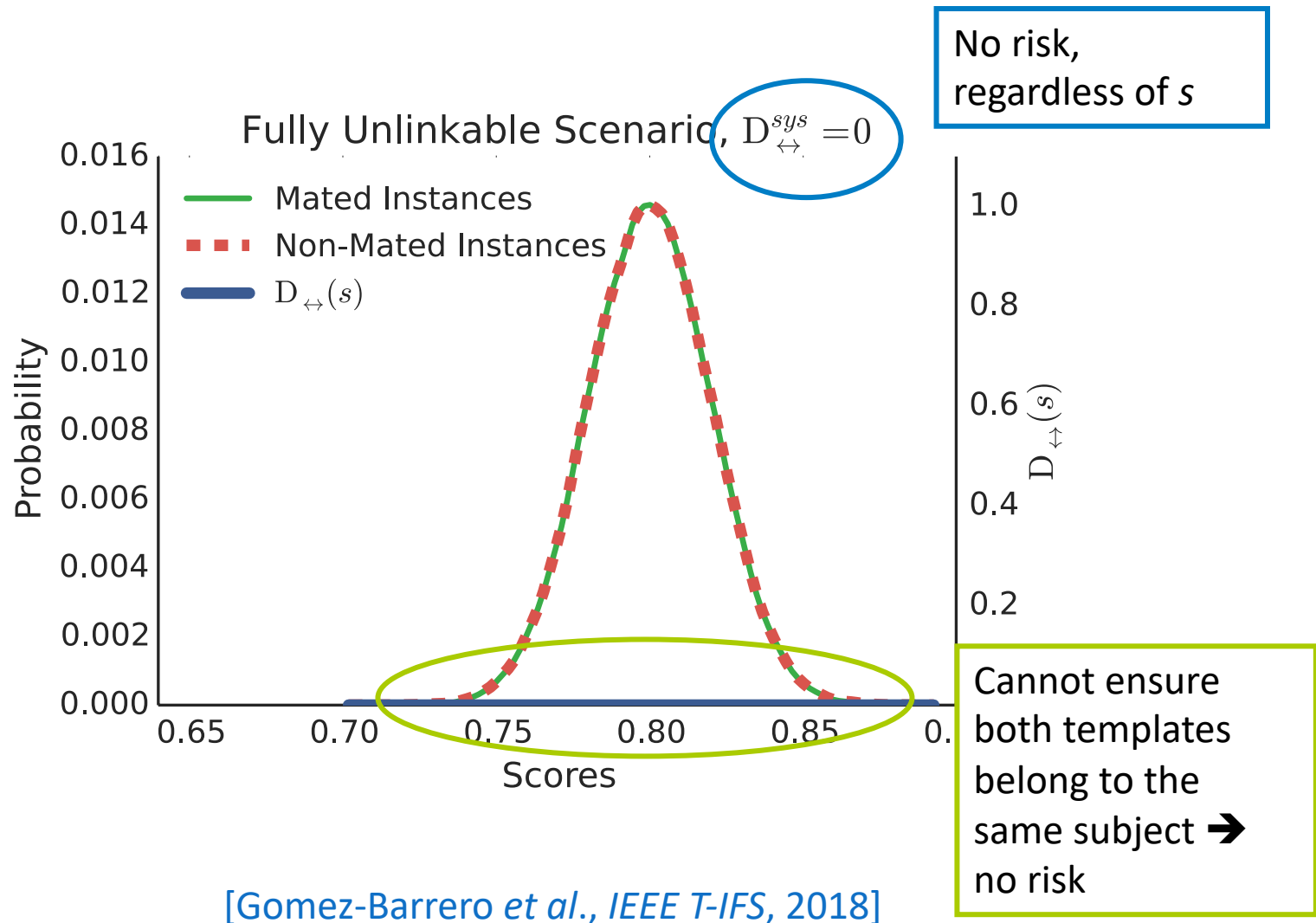
$s$  can be the dissimilarity score of the system or any other dissimilarity score, such as values extracted from partial decoding in fuzzy schemes

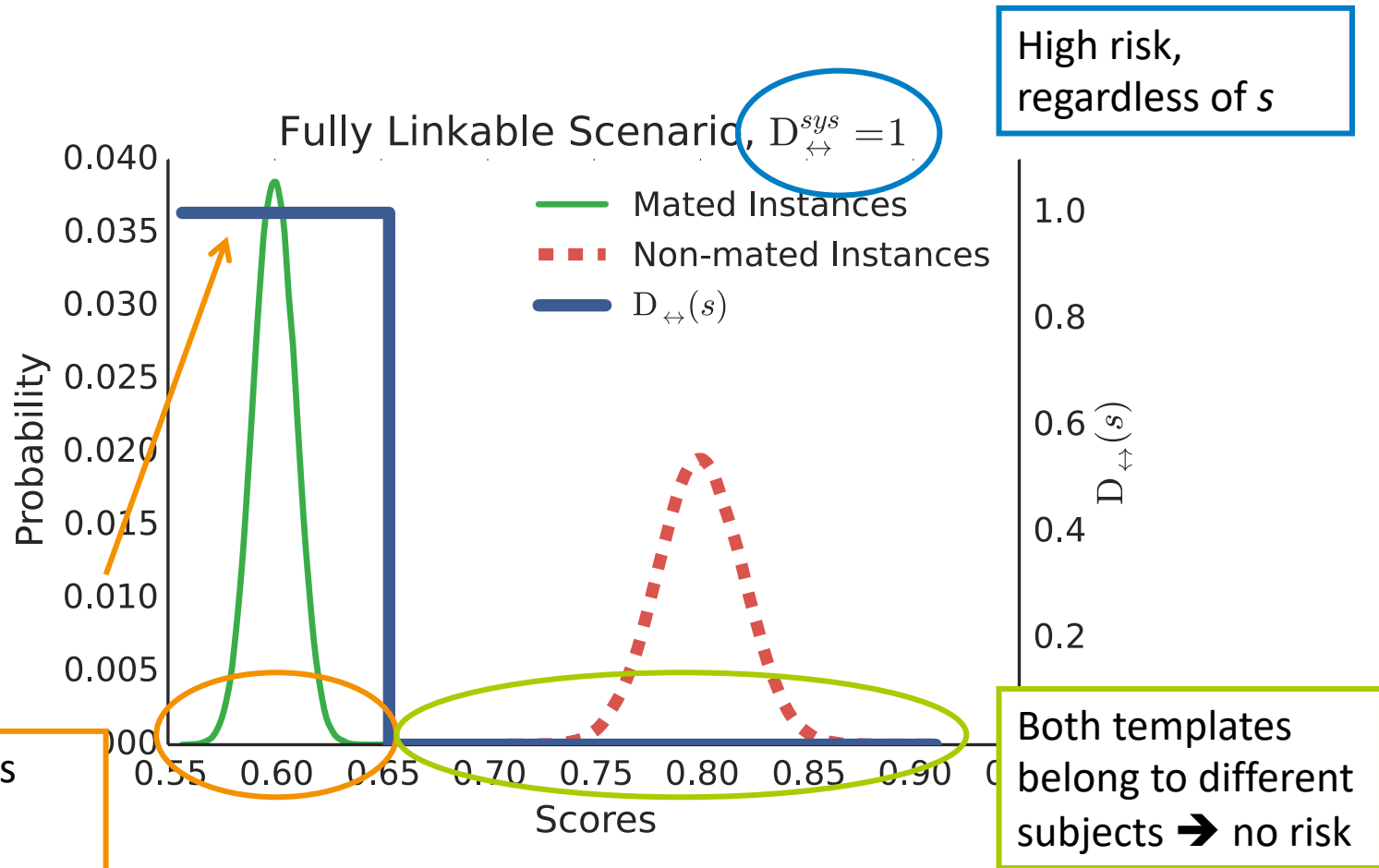
- Plot *Mated* and *Non-mated instances* distributions, for templates protected with different keys, and analyse them
- Two measures:
  - ❖ Local measure  $D_{\leftrightarrow}(s) \rightarrow$  for which scores is the system vulnerable?
  - ❖ Global measure  $D_{\leftrightarrow}^{sys} \rightarrow$  how can we compare two systems globally?
- Both bounded in  $[0,1]$ , and defined for all dissimilarity scores
- General measures, valid for all BTP schemes
- Available at: <https://github.com/dasec/unlinkability-metric>
- Included in the ongoing revision of ISO/IEC 30136

[Gomez-Barrero *et al.*, *IEEE T-IFS*, 2018]

- We need to simulate the enrolment of the subjects in different applications, and then apply the linkage function to those protected templates
- We need access to:
  - ❖ Biometric database (**DB**)
  - ❖ BTP scheme to be evaluated (**BTP**)
- Define a Linkage Function (**LF**)
  - ❖ Robustness to one **LF** does not imply robustness to all possible **LF**s!
- Compute score distributions of **LF** using **DB** and **BTP**

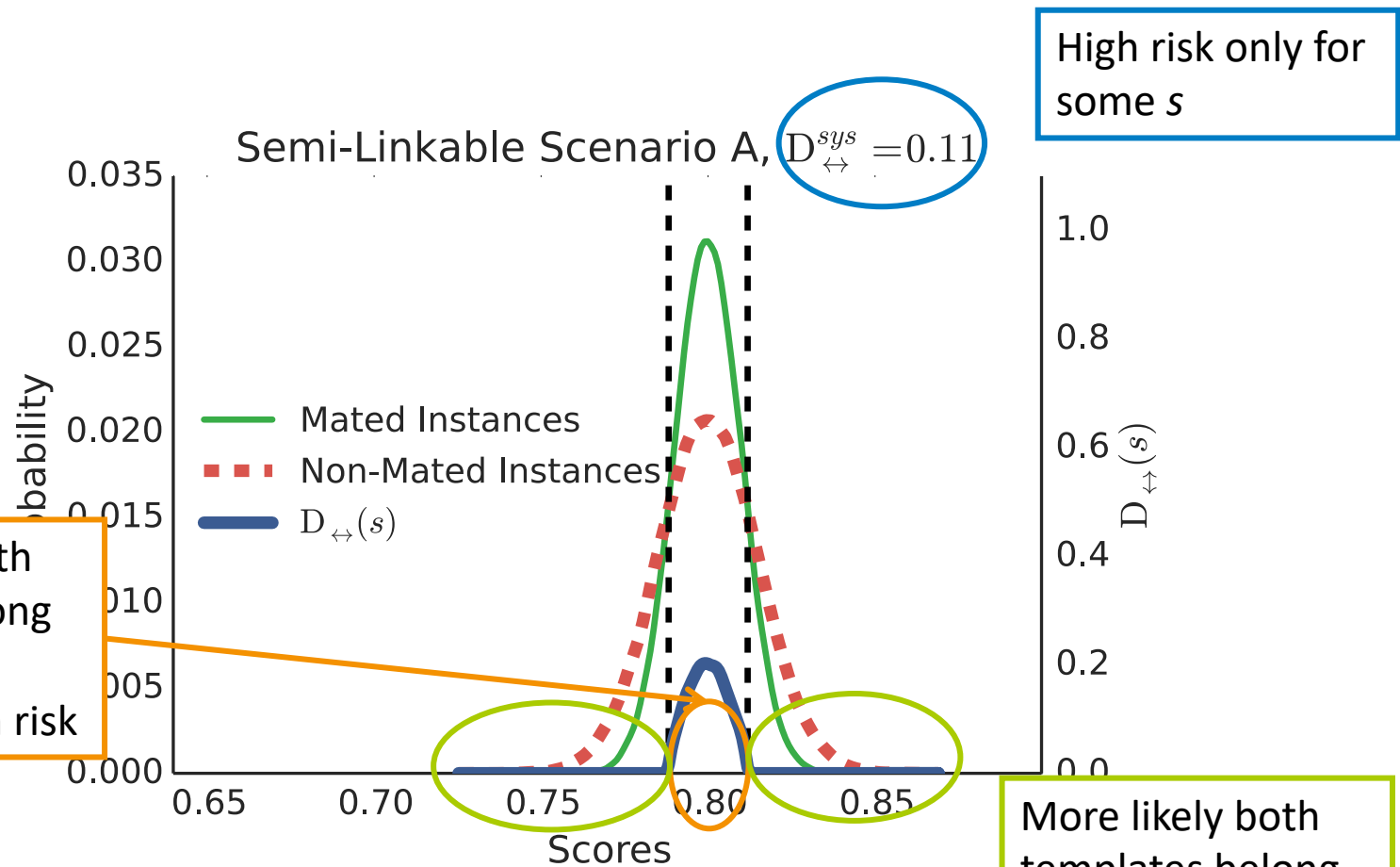
- Simulate  $K$  different applications
  - ❖ Choose a reasonably big  $K$  - e.g.,  $K = 10$
- Enrol the data subjects into all  $K$  applications: for sample  $j$  of subject  $i$ , we have  $K$  templates:  $T_1^{i,j}, T_2^{i,j}, \dots, T_k^{i,j}, \dots, T_K^{i,j}$ 
  - ❖ For subject  $i$ , having  $J$  samples, we have a total of  $J \times K$  templates:  
 $T_1^{i,1}, T_1^{i,2}, \dots, T_1^{i,J}, T_2^{i,1}, \dots, T_2^{i,J}, \dots, T_K^{i,1}, \dots, T_K^{i,J}$
- Design your protocol for mated and non-mated trials
- Compute mated and non-mated linkage scores by comparing:
  - ❖ Mated scores: samples of the **same** subject, enrolled in **different** applications:  $s = LF(T_1^{1,1}, T_2^{1,2})$
  - ❖ Non-Mated scores: samples of the **different** subjects, enrolled in **different** applications:  $s = LF(T_1^{1,1}, T_2^{2,2})$



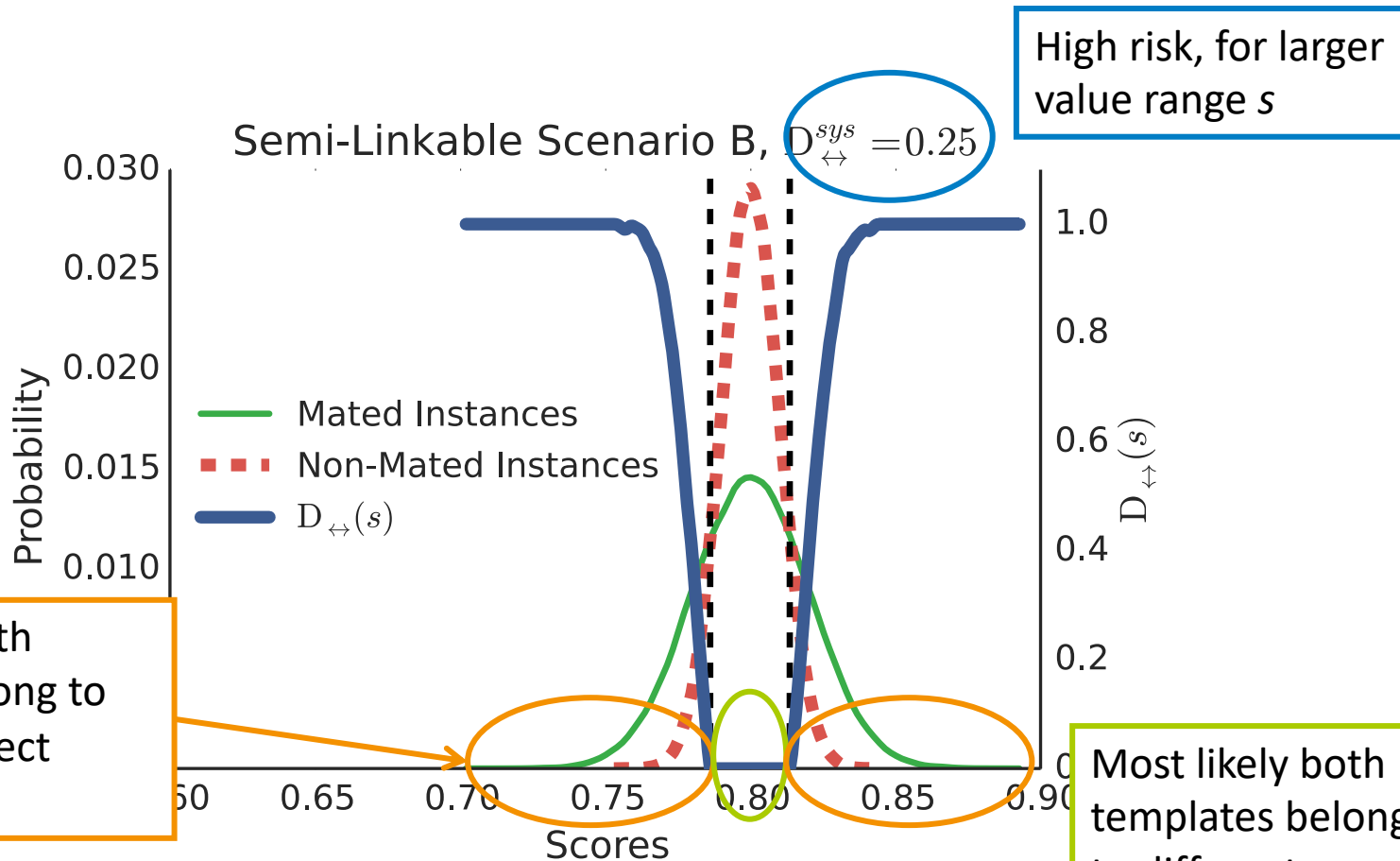


[Gomez-Barrero *et al.*, *IEEE T-IFS*, 2018]





[Gomez-Barrero et al., IEEE T-IFS, 2018]



[Gomez-Barrero et al., IEEE T-IFS, 2018]

- We are interested in evaluating:  $D_{\leftrightarrow}(s) = p(H_m|s) - p(H_{nm}|s)$
- Doing some math tricks, we get:

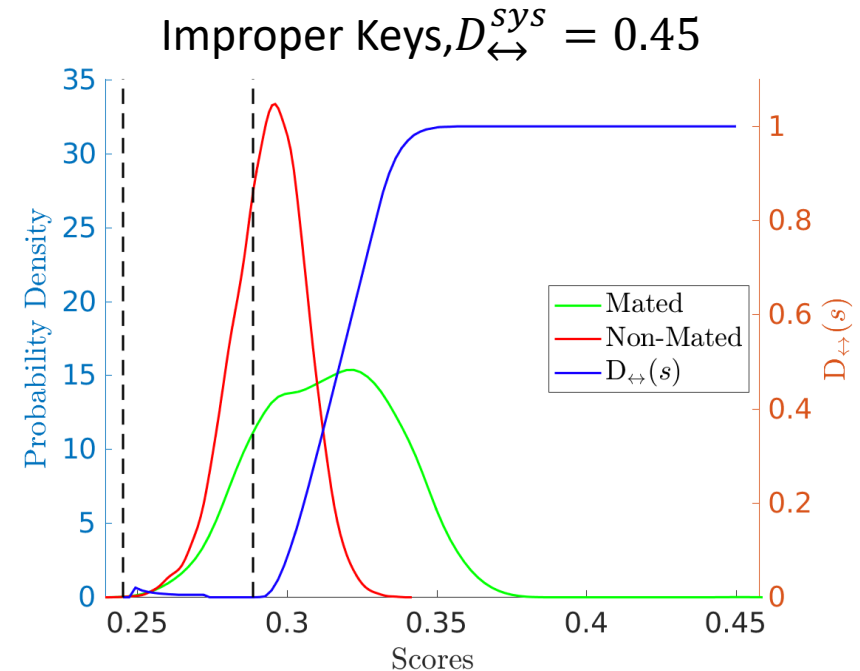
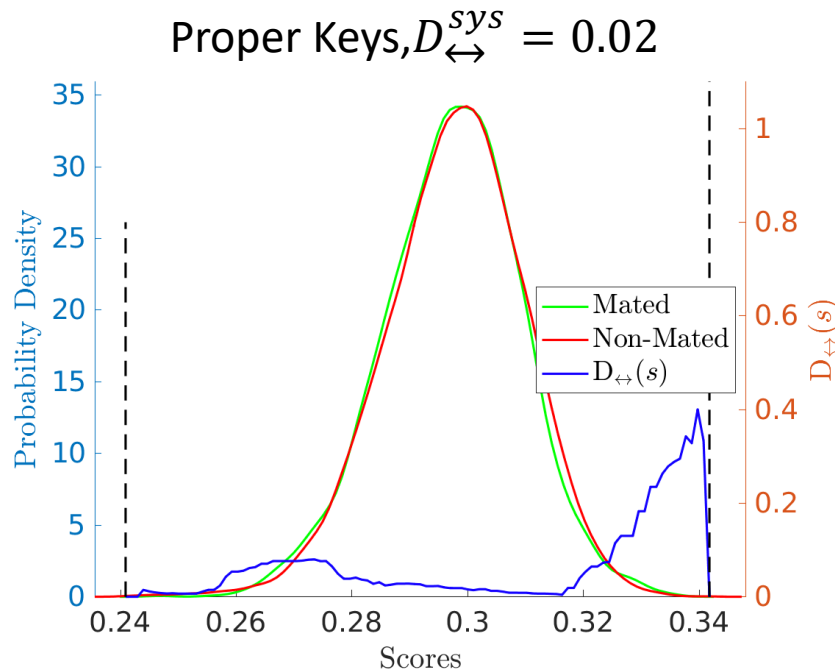
$$D_{\leftrightarrow}(s) = \begin{cases} 0 & \text{if } LR(s) \cdot \omega \leq 1 \\ 2 \frac{LR(s) \cdot \omega}{1 + LR(s) \cdot \omega} - 1 & \text{if } LR(s) \cdot \omega > 1 \end{cases}$$

- Where  $\omega = p(H_m) / p(H_{nm})$
- And we can define a global metric as:

$$D_{\leftrightarrow}^{sys} = \int_{s_{min}}^{s_{max}} p(s|H_m) \cdot D_{\leftrightarrow}(s) ds$$

[Gomez-Barrero *et al.*, *IEEE T-IFS*, 2018]

- Assumption: Key selection is done *properly*
  - ❖ What means properly?
  - ❖ Improper: leading to template collisions – can be spotted using  $D_{\leftrightarrow}^{sys}$



[S. Kirchgasser & A. Uhl, Proc. BIOSIG 2022]

# Summary

➤ Do the stored templates reveal any information about the original biometric samples?

IRREVERSIBILITY

➤ Are my enrolled templates in different recognition systems somehow related to each other?

UNLINKABILITY

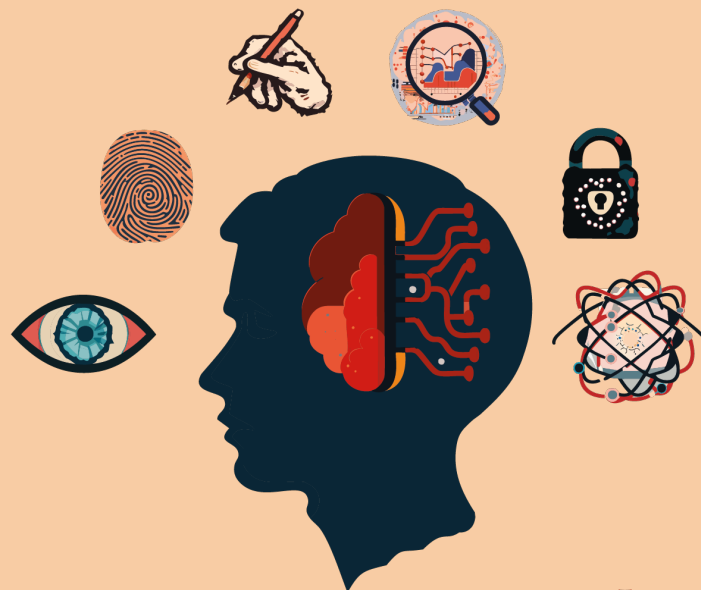
➤ What if someone steals a template extracted from my face? Has it been permanently compromised?

RENEWABILITY

[ISO/IEC IS 24745 on Biometric Information Protection]

[ISO/IEC IS 30136 on Performance Testing of BTP Schemes]

- M. Gomez-Barrero, J. Galbally. “Reversing the irreversible: A survey on inverse biometrics” *Computers & Security*, vol. 19, pp. 101700, 2020
- M. Gomez-Barrero, J. Galbally, C. Rathgeb, C. Busch, “General Framework to Evaluate Unlinkability in Biometric Template Protection Systems”, *IEEE Trans. on Information Forensics and Security*, vol. 3, no. 6, pp. 1406-1420, 2018
- M. Gomez-Barrero, C. Rathgeb, J. Galbally, C. Busch, J. Fierrez, “Unlinkable and irreversible biometric template protection based on Bloom filters”, *Information Sciences*, vol. 370-371, pp. 18-32, 2016
- M. Gomez-Barrero, C. Rathgeb, G. Li, R. Raghavendra, J. Galbally, C. Busch, “Multi-Biometric Template Protection Based on Bloom Filters”, *Information Fusion*, vol. 42, pp. 37-50, 2018
- S. Kirchgasser, A. Uhl, “Template Protection: On the need to adapt the current Unlinkability Evaluation Protocol”, *Proc. BIOSIG*, 2022
- ISO/IEC 24745 on Biometric information protection
- ISO/IEC 30136 on Performance testing of biometric template protection schemes - **IN REVISION!!!**
- S. Rane, “Standardization of biometric template protection”, *IEEE MultiMedia*, vol. 21, no. 4, pp. 94-99, 2014
- **Handbook of Biometric Template Protection, Springer, eds. V. Krivokuca Hahn, M. Gomez-Barrero, A. Ross, S. Marcel, 2025 (soon)**
  - ❖ **EAB Workshop on BTP on October 29 and 30, 2025**



# BioML Lab



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