

Motivation

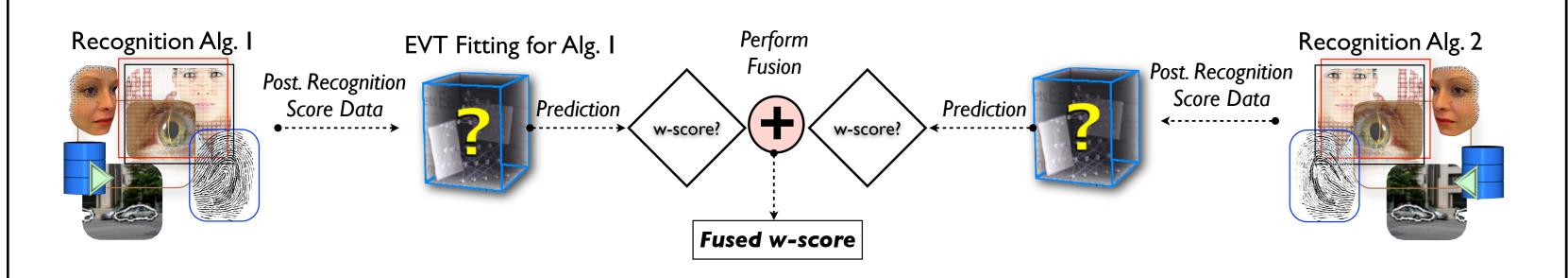
We introduce the w-score, a new technique for robust score normalization that does not assume a match or non-match distribution, but instead relies on the observation that the scores of a recognition system's non-match scores follow the statistical Extreme Value Theory.

Our goal is to achieve **robust fusion**: a fusion process that is insensitive to errors in its distributional assumptions on the data, has simple parameter estimation, and a high input failure tolerance.

Robustness in score level fusion is impacted by normalization in two ways:

- 1. The varying nature of the underlying distribution of scores across different recognition algorithms often leads to inconsistent normalization results.
- 2. Complications arise when one or more sensors or recognition algorithms considered for fusion fail or are deceived.

The w-score is designed to address both of these problems



An overview of the w-score process. Our robust fusion is inspired by postrecognition score analysis^{1,2}

Recognition Systems

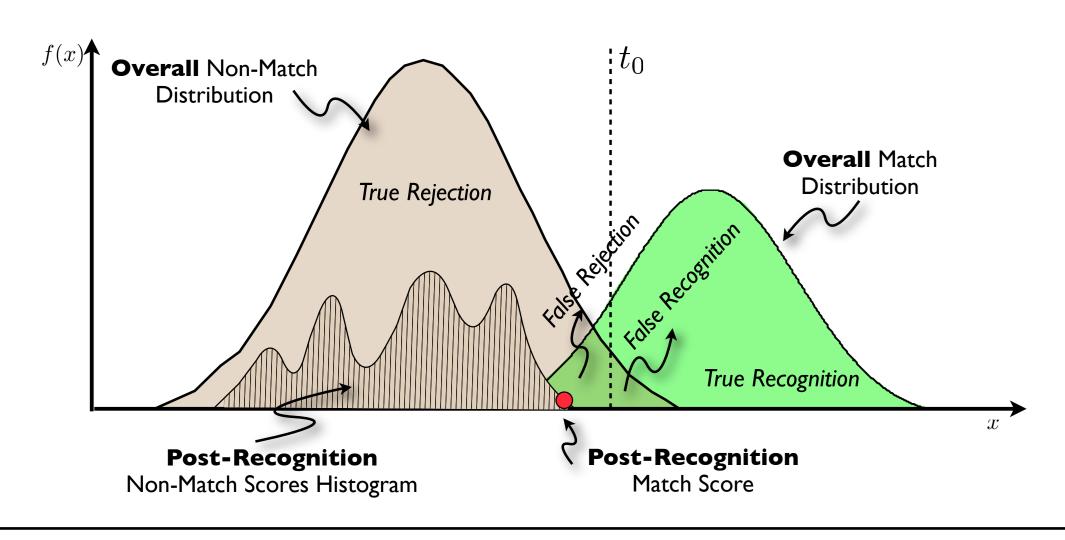
The task of a recognition system³ is to find the class label c^* , where p_k is an underlying probability rule and p_0 is the input distribution, satisfying

$$c^* = \operatorname*{argmax}_{class c} Pr(p_0 = p_c)$$

subject to $Pr(p_0 = p^*) \ge 1 - \delta$ for a given confidence threshold δ , or to conclude the lack of such a class.

probe is defined as the input image distribution p_0 submitted to the recognition system in order to find its corresponding class label c^* .

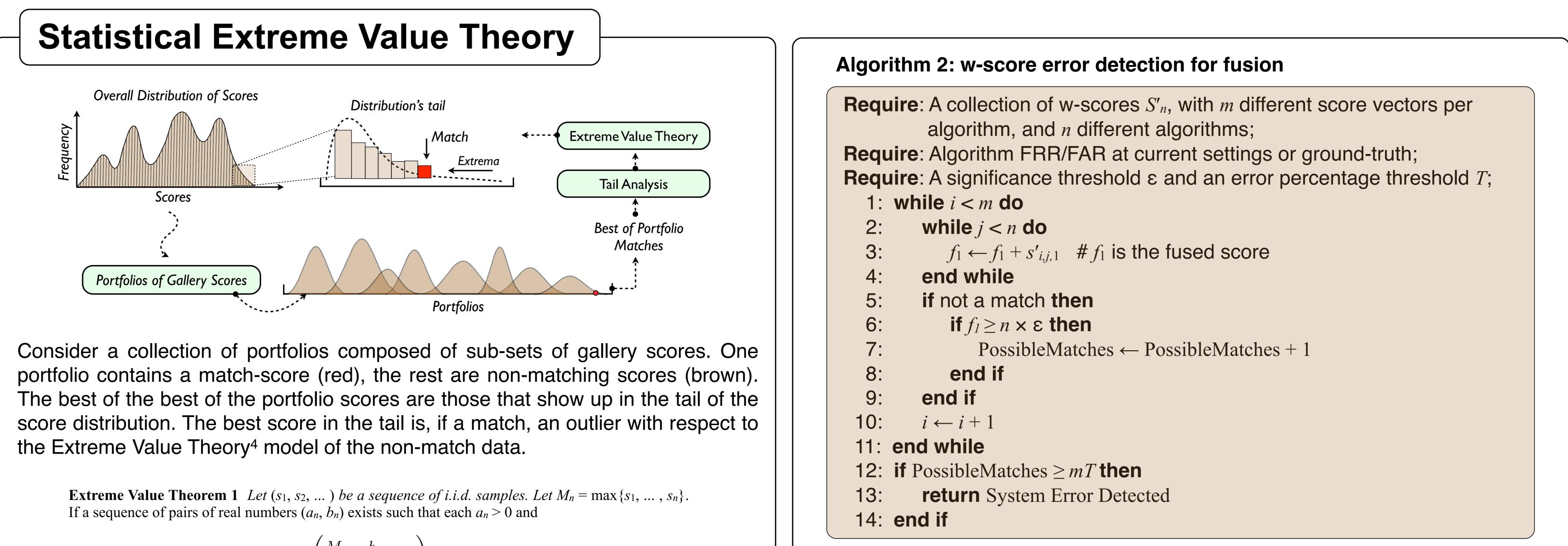
gallery is defined to be all the classes c^* known to the recognition system.



Where the tails of the match and non-match distributions overlap is where we find False Rejection and False Recognition.

Robust Fusion: Extreme Value Theory for Recognition Score Normalization

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$$\lim_{x \to \infty} P\left(\frac{M_n - b_n}{a_n} \le x\right) = F(x)$$

then if *F* is a non-degenerate distribution function, it belongs to one of three extreme value distributions.

Theorem 1 tells us that a large set of individual maximums M_n from the portfolios must converge to an extreme value distribution. These maxima are most accurately modeled by the Weibull Distribution (Extreme Value Type III Distribution).

W-Score Normalization

Algorithm 1: w-score normalization

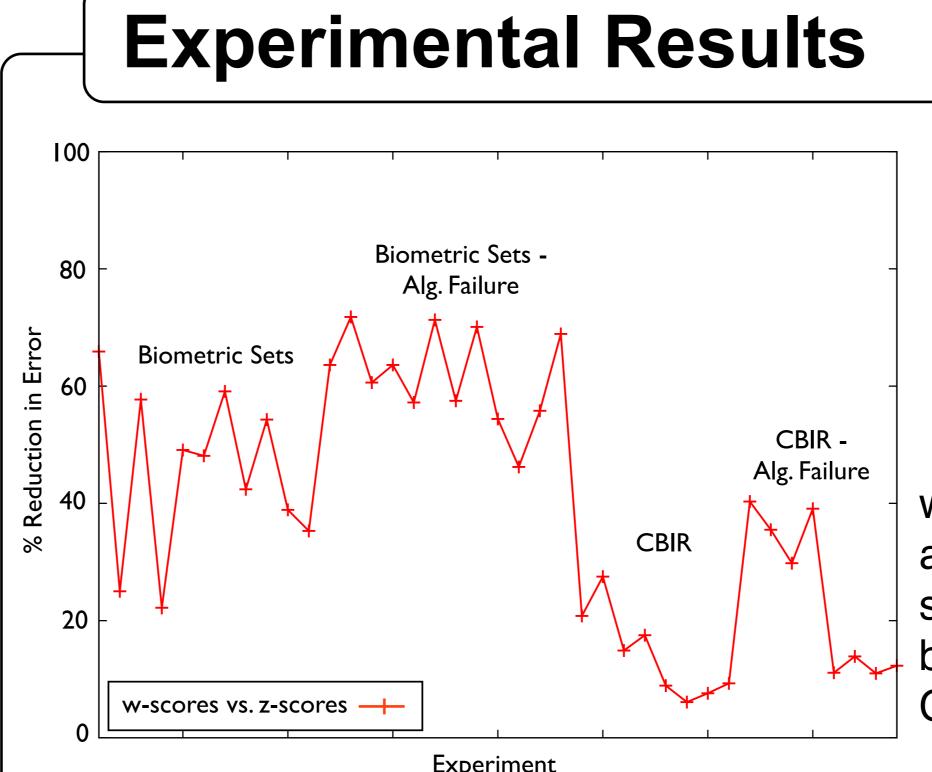
Require: A collection of scores S, of vector length m, from a single recognition algorithm *j*;

- : Sort and retain the *n* largest scores, $s_1, \ldots, s_n \in S$;
- 2: Fit a GEV or Weibull distribution W_S to s_2, \ldots, s_n , skipping the hypothesized outlier;
- 3: **while** *k* < *m* **do**
- 4: $s'_k = \text{CDF}(s_k, W_S)$
- 5: $k \leftarrow k+1$
- 6: end while

The w-score re-normalizes the data based on its formal probability of being an outlier in the extreme value "non-match" model, and hence its chance of being a successful recognition.

We use the CDF defined by the parameters of the fitting to produce the normalized probability score. The tail size used for fitting is the only parameter that must be estimated empirically.

Detection for three possible errors: (1) inaccurate Weibull fitting, (2) invalid score data produced CDF that returns too many high w-scores (3) data alignment error

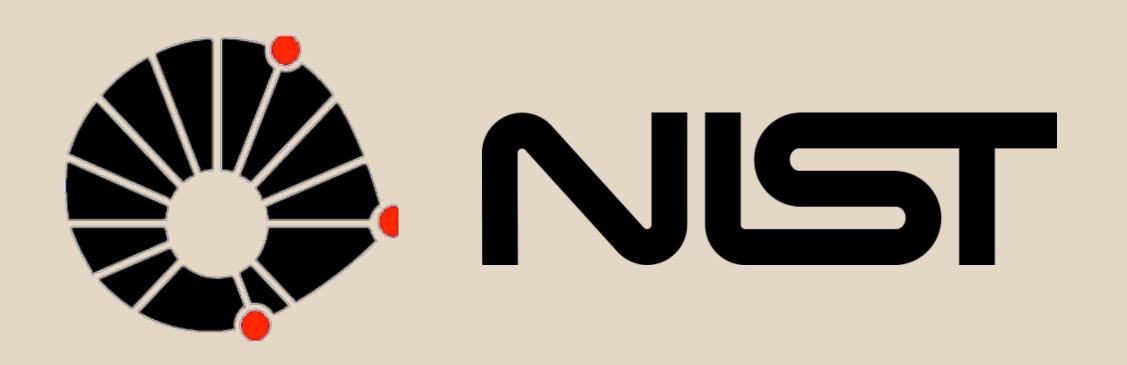


w-scores reduce the margin of error after fusion, when compared to zscores (baseline), for a variety of biometric recognition algorithms and CBIR descriptors.

Biometric Recognition: NIST BSSR1 biometric score set; 2 face recognition algorithms (labeled C & G) and 1 fingerprint recognition algorithm applied to two different fingers (labeled LI & RI); 517 scores across common subjects in true multibiometric set, 3000 scores across common subjects in our Chimera set.

Content Based Image Retrieval: Corel "Relevants" set, containing 50 unique classes, and the INRIA "Holidays" set, containing 500 unique classes; using a variety of descriptors, we generated 1624 score sets for Corel Relevants and 1491 score sets for INRIA Holidays.

All sum rule results are presented as a percentage of error reduction (improvement) compared to z-scores⁵, the most popular type of adaptive score normalization, calculated as: % reduction = ($\% e_z - \% e_w$) / $\% e_z$



w-scores are most useful when some of the data considered for fusion is "failing".

Algorithms	Improvement	%c*
!C & LI	63.6%	2.0%
!C & RI	71.8%	2.0%
!G & LI	60.6%	2.0%
!C & RI	63.6%	2.0%
Chimera !C & LI	57.2%	0.3%
Chimera !C & RI	71.3%	0.3%

Algorithms	Improvement	%c*
Chimera !G & LI	57.5%	0.3%
Chimera !G & RI	70.1%	0.3%
Chimera LI & !RI	54.4%	0.3%
Chimera RI & !LI	46.2%	0.3%
Chimera !C & !G & LI	55.8%	0.3%
Chimera !C & !G & RI	68.9%	0.3%

Rank-1 fusion results, compared to z-scores, for the BSSR1 multibiometric and the BSSR1 "Chimera" data sets, fusing with failing algorithms (marked with !).

CBIR Descriptors	Improvement	%c*
Relevants !csd & gch	40.3%	6.0%
Relevants csd & !jac	35.5%	6.0%
Relevants <i>cwhsv</i> & ! <i>cwluv</i>	29.8%	6.0%
Relevants ! <i>cwhsv</i> & <i>cwluv</i>	39.1%	6.0%

CBIR Descriptors	Improvement	%c*
Relevants !csd & gch	11.1%	0.6%
Relevants csd & !jac	13.9%	0.6%
Relevants <i>cwhsv</i> & ! <i>cwluv</i>	11.0%	0.6%
Relevants ! <i>cwhsv</i> & <i>cwluv</i>	12.3%	0.6%

Rank-1 CBIR fusion results, compared to z-scores, for the Corel Relevants and INRIA Holidays data sets, fusing with failing algorithms.

%c* represents the tail size equal to a percentage of the total number of classes

References

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