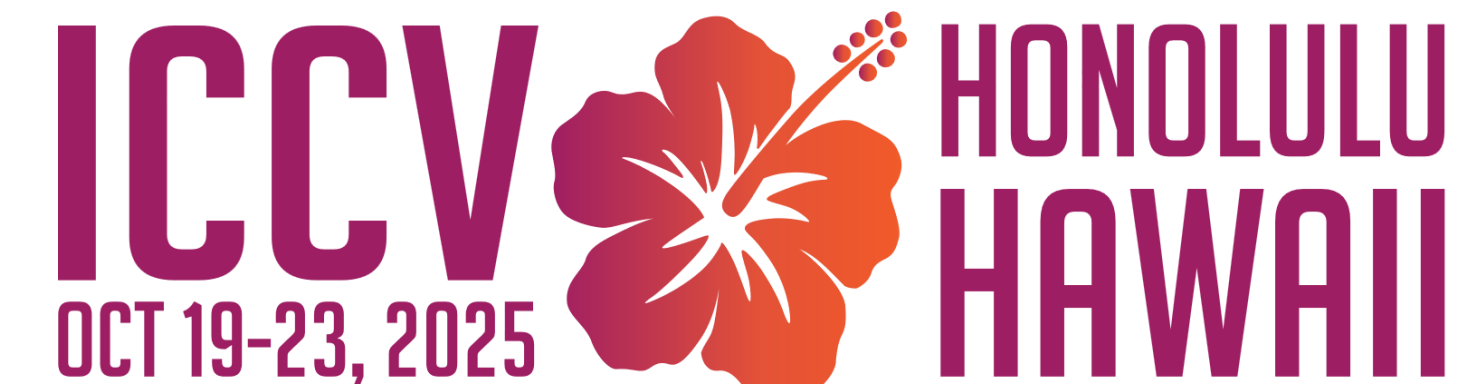




A Quality-Guided Mixture of Score-Fusion Experts Framework for Human Recognition

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Code



Motivation

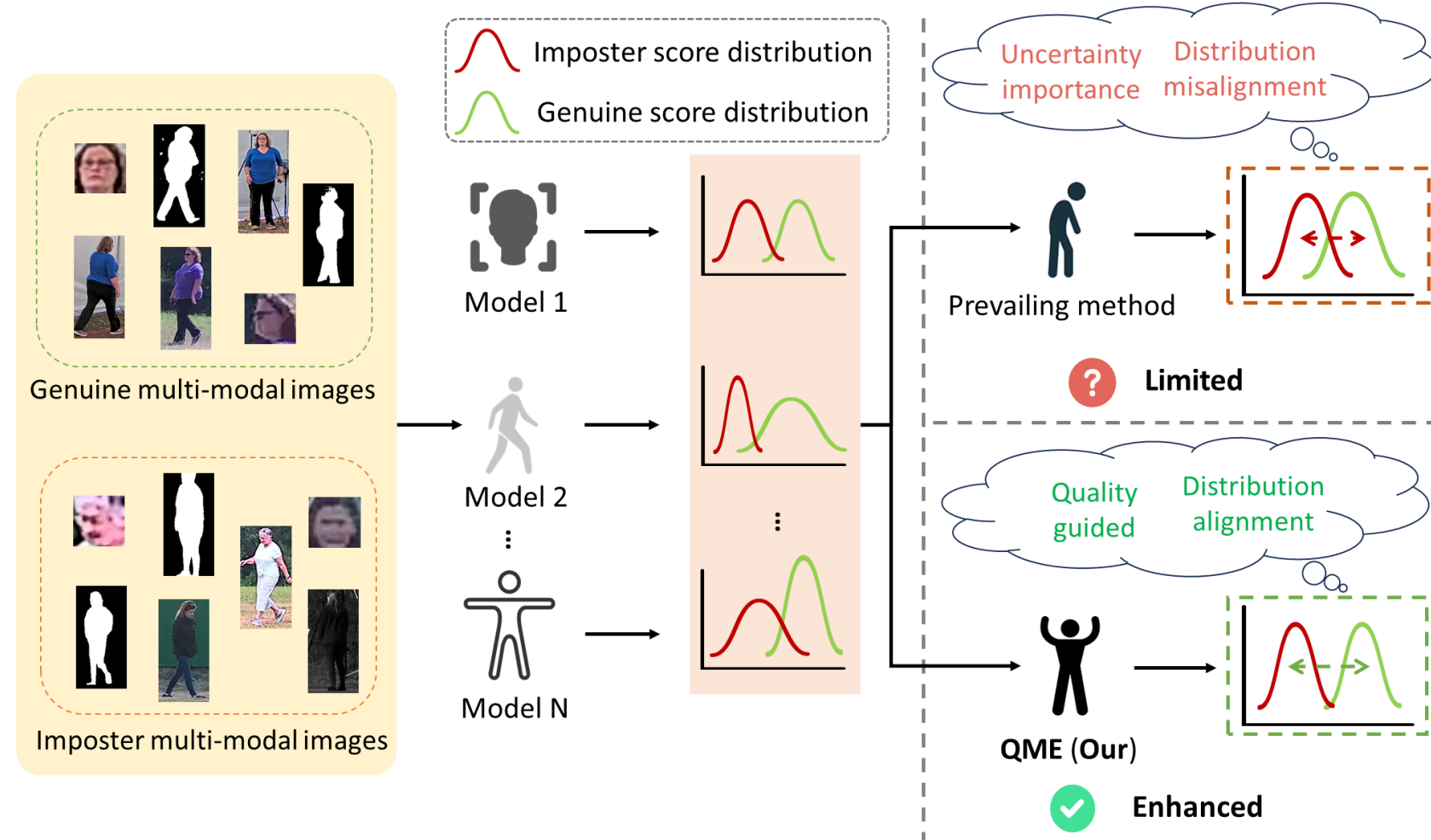
1. Low quality vs high quality modality:



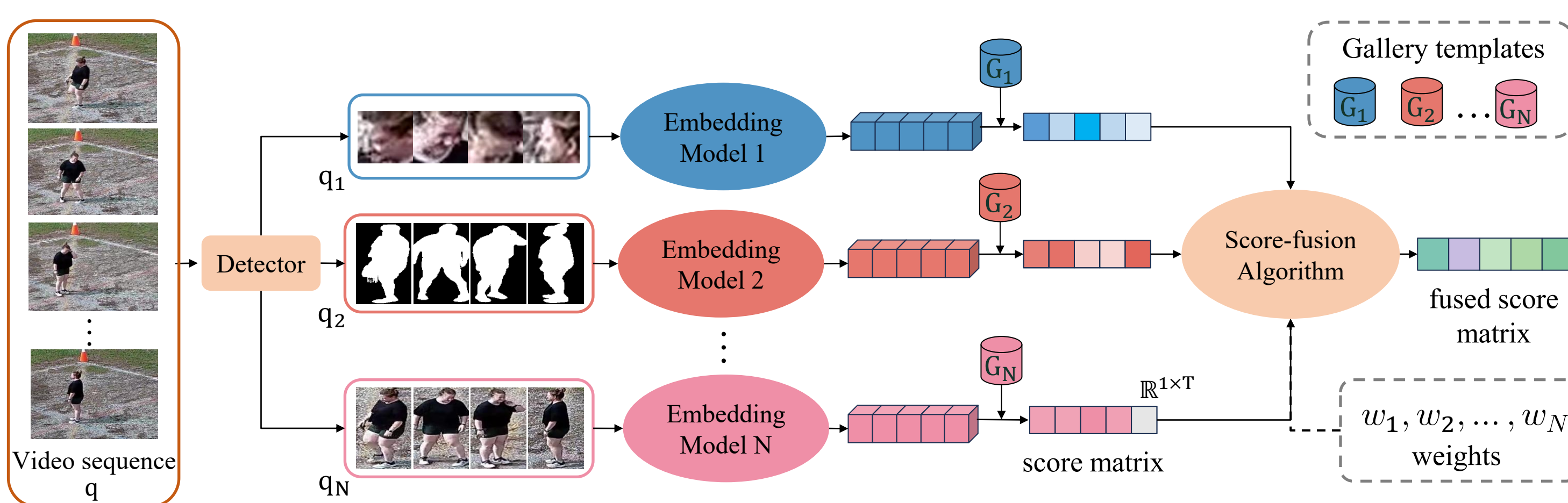
For a probe with **low-quality**, we should give a **lower weight**.

For a probe with **high-quality**, we should give a **higher weight**.

2. Distribution misalignment & uncertainty importance:

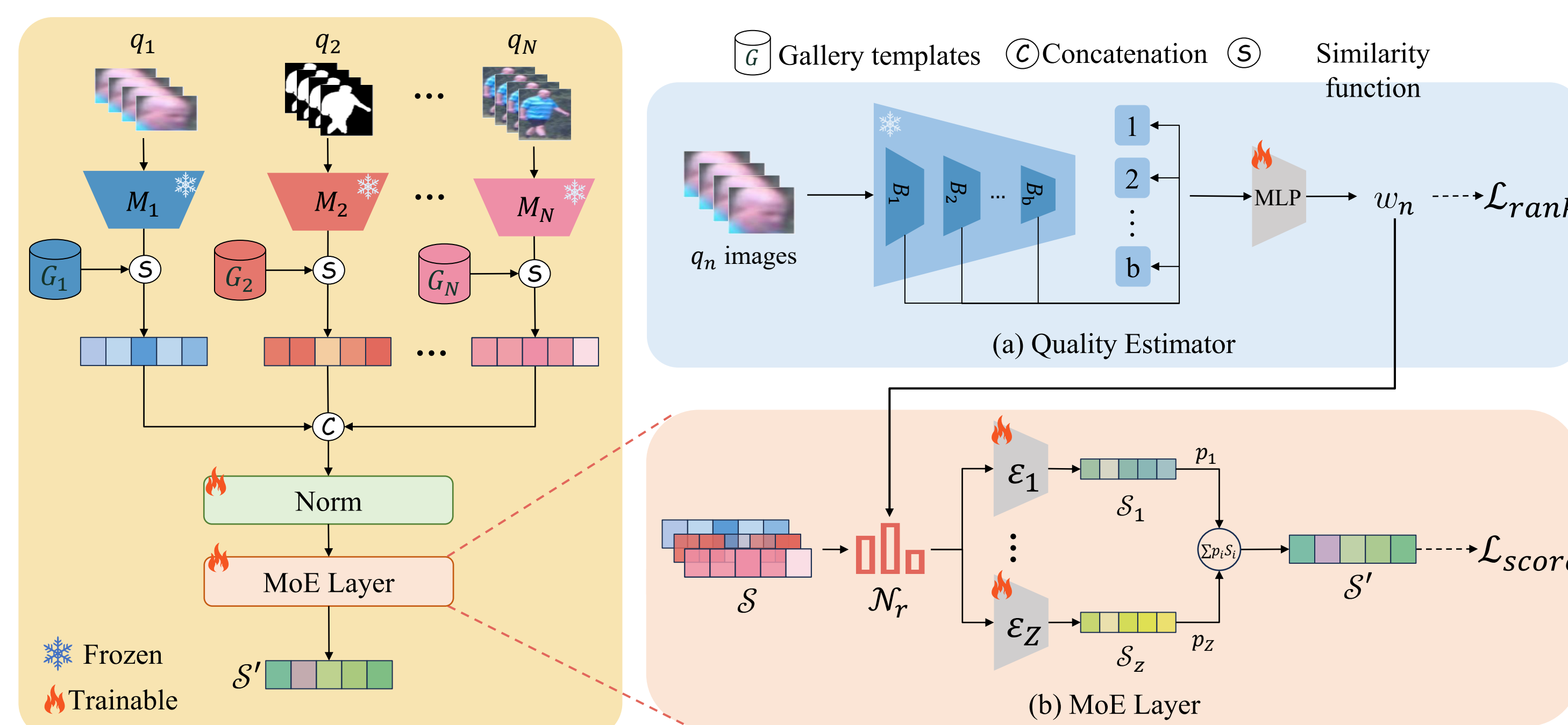


Systematic Whole-body Biometric Recognition



QME Framework

We propose a **Quality-Guided** Mixture of Score-Fusion Experts (QME) framework to fused the score matrices.



QME dynamically assigns weights to each expert based on **input quality**, encouraging **diverse specializations**.

Loss Functions

Pseudo Quality Loss

Provide pseudo quality labels based on the **ranking result** of the sample:

$$\mathcal{L}_{rank} = \sum_{i \in L} \text{MSELoss} \left(w_i, \text{ReLU} \left(\frac{\delta - r_i}{\delta - 1} \right) \right)$$

ranking result of the query feature
 ranking threshold

Examples:

$r_i=1, \delta=10$, Pseudo quality label: 1

$r_i=5, \delta=10$, Pseudo quality label: 0.56

$r_i=20, \delta=10$, Pseudo quality label: 0

Score Triplet Loss

Traditional triplet loss optimizes relative distances between samples:

$$\mathcal{L}_{tri} = \text{ReLU}(d(a, p) - d(a, n) + m),$$

but it does not constrain the **values of negative samples**, which are crucial for calculating the **threshold** in verification and open-set search metrics.

To optimize these metrics, we introduce the score triplet loss:

$$\mathcal{L}_{score} = \text{ReLU}(\mathcal{S}'_{nm}) + \text{ReLU}(m - \mathcal{S}'_{mat})$$

Experiments

Our method consistently outperforms existing approaches across all four benchmarks, achieving SoTA performance in challenging metrics.

MEVID

Method	Comb.	Rank1↑	mAP↑	TAR↑	FNIR↓
AdaFace* [22]	♦	25.0	8.1	5.4	98.8 ± 1.2
CAL [15]	♦	52.5	27.1	34.7	67.8 ± 7.3
AGRL [53]	♦	51.9	25.5	30.7	69.4 ± 8.9
Z-score [47]		54.1	27.4	30.7	66.5 ± 7.0
Min-max [47]		52.8	24.7	25.0	71.3 ± 6.1
RHE [17]		52.8	24.8	25.3	71.2 ± 6.2
Weighted-sum [35]	♦	54.1	27.3	30.3	66.3 ± 7.0
Asym-AOI [18]		52.5	22.9	23.6	71.7 ± 5.8
BSSF [49]		53.5	27.4	30.5	65.9 ± 7.2
Farsight [32]		53.8	25.4	26.6	69.8 ± 6.4
Ours (AdaFace-QE)		55.7	28.2	32.9	64.6 ± 8.2
Ours (CAL-QE)		55.4	27.9	32.5	64.3 ± 8.7

CCVID

Method	Comb.	Rank1↑	mAP↑	TAR↑	FNIR↓
AdaFace* [22]	♦	94.0	87.9	75.7	13.0 ± 3.5
CAL [15]	♦	81.4	74.7	66.3	52.8 ± 13.3
BigGait* [57]	♦	76.7	61.0	49.7	71.1 ± 6.1
Z-score [47]		92.2	90.6	73.9	15.1 ± 1.5
Min-max [47]		91.8	90.9	73.9	15.4 ± 2.5
RHE [17]		91.7	90.2	73.1	16.6 ± 2.5
Weighted-sum [35]	♦	91.7	90.6	73.6	15.4 ± 1.8
Asym-AOI [18]		92.3	90.0	74.0	15.9 ± 1.7
BSSF [49]		91.8	91.1	73.9	14.1 ± 1.3
Farsight [31]		92.0	91.2	73.9	13.9 ± 1.1
Ours (AdaFace-QE)		92.6	91.6	75.0	13.3 ± 1.2
Ours (CAL-QE)		94.1	90.8	76.2	12.3 ± 1.4

BRIAR

Method	Comb.	Face Incl. Trt.			Face Restr. Trt.		
		TAR↑	R20↑	FNIR↓	TAR↑	R20↑	FNIR↓
KPRPE [24]	♦	66.5	80.5	54.8	31.5	44.5	81.3
BigGait [57]	♦	66.3	93.1	72.7	61.0	90.4	76.3
CLIP3DReID [32]	♦	55.8	83.5	80.1	47.9	79.3	83.4
Farsight [31]		82.4	95.8	46.1	65.7	91.0	68.2
Ours		84.5	96.0	41.2	67.9	90.6	64.1

LTCC

Method	Comb.	Rank1↑	mAP↑	TAR↑	FNIR↓
AdaFace* [22]	♦	18.5	5.9	2.4	99.8 ± 0.2
CAL [15]	♦	74.4	40.6	36.7	59.7 ± 7.3
AIM [56]	♦	74.8	40.9	37.0	66.2 ± 9.2
Asym-AOI [18]		71.2	32.9	19.1	76.3 ± 8.9
BSSF [49]		73.5	39.1	34.2	68.9 ± 8.5
Farsight [31]		73.2	37.8	31.3	72.4 ± 8.6
Ours		73.8	39.6	35.0	64.3 ± 8.0

Ablation Studies

(1) Effects of proposed component.

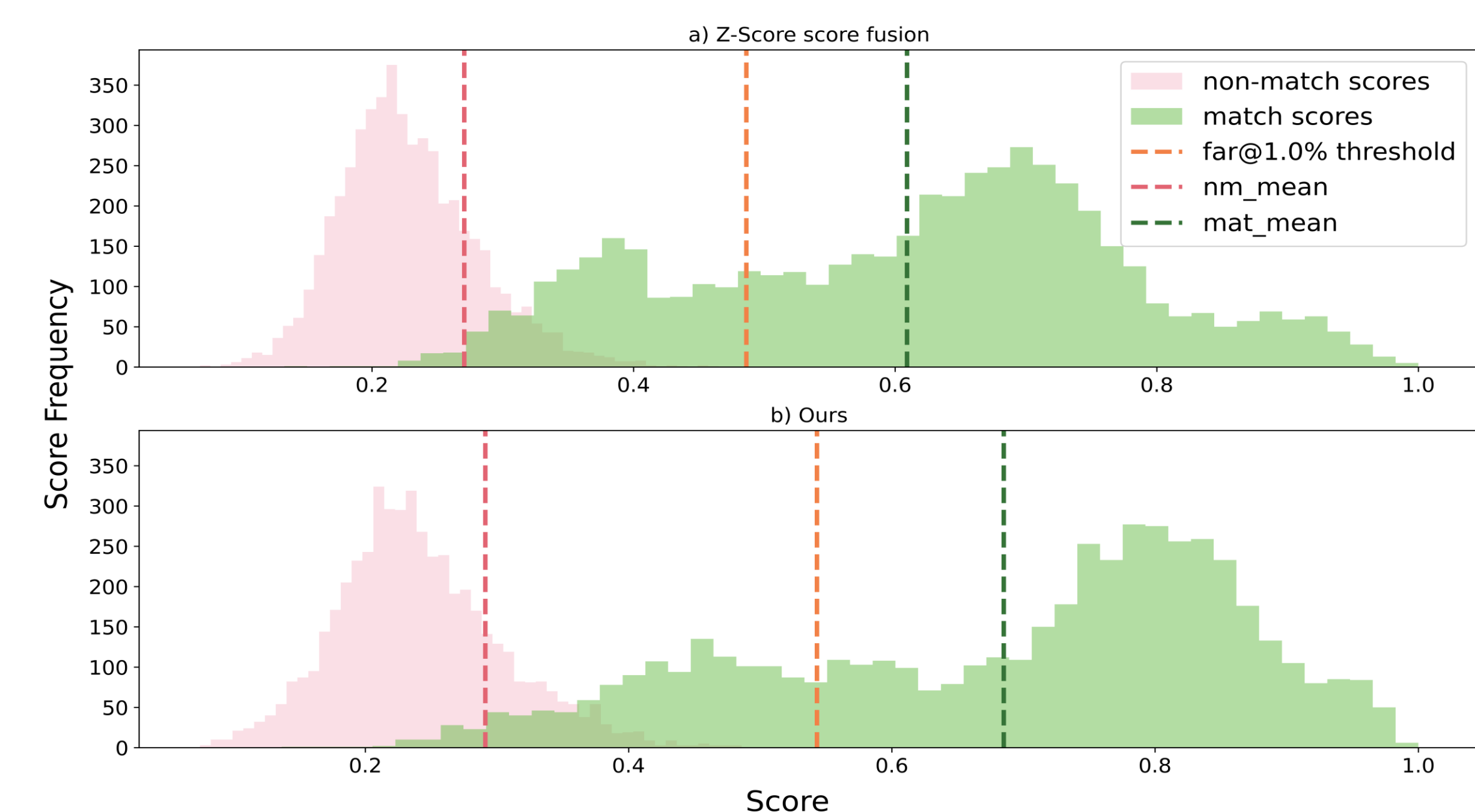
\mathcal{L}_{score}	QE	Z	Rank1↑	mAP↑	TAR↑	FNIR↓
✗	✗	1	49.4	21.6	23.3	84.0
✓	✗	1	53.8	24.5	25.3	70.4
✗	✗	2	54.1	25.5	30.8	65.4
✓	✗	2	55.1	27.0	31.3	66.5
✓	✓	2	55.7	28.2	32.9	64.6

(2) Effects of expert aggregation. Each expert specializes in **specific data scenarios and metrics**, and their aggregation leads to the best overall performance.

Expert	Face Incl. Trt.			Face Restr. Trt.		
	TAR↑	R20↑	FNIR↓	TAR↓	R20↑	FNIR↓
ε_1	<u>83.6</u>	<u>95.5</u>	<u>41.7</u>	62.0	90.6	<u>66.7</u>
ε_2	81.8	<u>95.5</u>	46.6	<u>65.0</u>	90.6	68.4
Ours ($\varepsilon_1 + \varepsilon_2$)	84.5	95.7	41.2	67.9	90.6	64.1

Visualizations

QME has a **clearer boundary** between match scores and non-match scores.



The predicted quality weight can **dynamically reflect** the quality of the input sample:

