Supplementary Material for BioNet: A Biologically-inspired Network for Face Recognition

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This supplementary material includes:

- 1. The influence of the attribute number (K) when ResNet-101 employs the Visual Cortex Network.
- 2. The Top-K accuracy on IJB-B [13] and IJB-C [10] datasets.
- 3. The performance on LFW [4], CFP-FP [11], CALFW [16], CPLFW [15], SLLFW [1], and YTF [14].
- 4. We list the performance of previous attribute-enhanced FR models [3, 8, 12]. Their theoretical limitations are introduced in the Related Work of the submitted paper.
- 5. The experiments with ResNet-101 also support the conclusion that the vanilla deep FR models and their straightforward improvement with borrowed mechanisms do not capture the human biological characteristics and can not boost FR performance as significantly as our BioNet.
- 6. The detailed illustration of the Straightforward Improvement with borrowed mechanisms in the submitted Tab.3 and our BioNet.

1. Influence of the Attribute Number (K) when ResNet-101 employs the VCN.

We employ the ResNet-101 and conduct the same experiments of the submitted Tab.6. The results in the following Tab.1 support the conclusion in the submitted paper, *i.e.* increasing the number of attributes is an effective method to improve the performance of BioNet.

2. Top-K Accuracy on IJB datasets

The top-K accuracies on IJB-B [13] and IJB-C [10] datasets are presented in Tab.2.

	#attr	IJB-A	IJB-B		IJB-C		MegaFace	
	πatu	@1e-4	@1e-4	@1e-5	@1e-4	@1e-5	rank1	
Baseline: ResNet-101	0	96.83	95.93	85.45	97.18	91.98	98.65	
BioNet: Observed	4	97.51	96.13	92.18	97.32	94.57	99.19	
	4	97.27	96.13	90.66	97.19	93.89	99.03	
BioNet: Latent	10	97.35	96.10	91.36	96.95	93.41	98.83	
	16	97.69	95.97	92.96	97.91	94.58	99.21	

Table 1. Influence of attribute number (K).

		IJB-B		IJB-C			
	rank1	rank5	rank10	rank1	rank5	rank10	
Baseline: CASIA-Net	88.03	92.47	94.21	89.05	93.35	94.51	
BioNet: Latent	89.25	94.06	95.89	90.19	94.02	95.47	
BioNet: Observed	89.32	94.11	95.68	90.28	94.09	95.51	
Baseline: ResNet-101	93.54	95.51	96.93	94.51	95.49	97.09	
BioNet: Latent	94.62	97.35	97.99	95.52	97.46	98.09	
BioNet: Observed	94.86	97.52	98.08	96.77	97.62	98.19	

Table 2. Top-K accuracy on IJB-B and IJB-C.

3. Performance on LFW, CFP-FP, CALFW, CPLFW, SLLFW, and YTF datasets

The performance on the evaluation datasets are presented in the following Tab.3. We observed our BioNet boost the GroupFace [6] with a marginal. We hypothesize it is because these small-scale evaluation datasets are not as challenging as the large-scale ones, which leads our improvement is not as significant as on the IJB-A/B/C and MegaFace.

	LFW	CFP-FP	CALFW	CPLFW	SLLFW	YTF
Baseline: CASIA-Net	99.30	94.00	91.60	80.44	97.28	95.12
BioNet: Latent	99.47	94.59	92.33	81.18	98.03	95.44
BioNet: Observed	99.48	94.71	92.48	81.21	97.87	95.42
Baseline: ResNet-101	99.82	98.14	95.28	92.01	99.33	97.76
BioNet: Latent	99.80	98.41	95.47	93.18	99.35	97.62
BioNet: Observed	99.83	98.60	96.23	93.23	99.38	97.78

Table 3. Performance on LFW, CFP, CALFW, CPLFW, SLLFW, and YTF

4. Performance of previous attribute-enhanced FR models

All the previous attribute-enhanced FR models have not evaluated their FR performance seriously. Besides, their performance is much inferior to the current status. This is the reason why we did not report their performance in the main paper. Their evaluation results are listed in the following:

- 1. Taherkhani *et al.* [12] proposed the PM method and evaluated it only on MegaFace [5]. Its performance is 78.82%.
- Hu *et al.* [3] proposed GTNN and evaluated it on LFW [4], NIR-VIS 2.0 [9], and Multi-PIE [2]. We list its performance on LFW which is 99.65% and on NIR- VIS 2.0 which is 99.94%. For the performance on Multi-PIE, we refer to their paper because the evaluation protocol of Multi-PIE is complex.
- 3. Kumar *et al.*, [8] evaluated their proposals only on LFW [4] which is 85.29%

In the view of attribute-enhanced FR, we carefully evaluate the performance of our proposals on all mainstream evaluation datasets, e.g., IJB-A/B/C [7, 10, 13], MegaFace [5], LFW [4], CFP-FP [11], CALFW [16], CPLFW [15], SLLFW [1], and YTF [14]. What's more, our BioNet significantly boosts the latest state-of-the-arts.

5. Studies about the Vanilla ResNet-101 Model and Its Straightforward Improvement

We employ the ResNet-101 and conduct the same experiments in the submitted Tab.3. The results in the following Tab.4 support the conclusion in the submitted paper.

		IJB-A	IJB-B	IJB-C	MegaFace	Feat	Attr4
		@1e-4	@1e-5	@1e-5	rank1	From	acc
Bacalina	FA	-	-	-	-		95.22
ResNet-101	FR	96.83	85.45	91.98	98.65	feat_id	72.44
						layer4	84.18
Straight- forward	Multitask	94.99	74.04	87.75	98.41	feat id	94.03
	I	///	/	01110	20111	Teur_ru	2
	Multitask	95.12	86.64	92.09	98.60	feat_attrs	95.55
	<u> </u>						
•	Self-	96.69	87.76	92.16	98.76	feat_id	64.56
Improve- ment	Attention					layer4	69.69
	Supervised-	96.88	87.99	92.06	98.63	feat attrs	95 53
	Attention				,		
ment	Avg-	97 10	86 64	91.83	98.83	feat attrs	95 46
	Ensemble	27.10	00.01	/1.00	20105	reation	20.10
ment	Adaptive-	97 35	89.78	93.14	98.89	feat attre	95.48
	Ensemble	77.55	07.70	22.14	20.02	icui-attrs	22.40
Ours: BioNet		97.51	92.18	94.57	99.19	feat_attrs	95.56

Table 4. Studies about the vanilla FR model and its straightforward improvement.

6. Detailed Illustration of the *Straightforwad Improvement* of Previous CNNs

The submitted Tab.3 provides six straightforward improvement settings of the vanilla CNNs. They borrowed lessons from the attention mechanism, multi-task learning mechanism, and ensemble mechanism. The detailed illustrations of the settings are shown in the following Fig.1(A-F)



Figure 1. Straightforward improvement with borrowed mechanisms(A-F). (G) is the legend.

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