

# High-precision Immune Computation for Secure Face Recognition

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## Abstract

*The accuracy of face recognition is very important for its security in many applications, because wrong face recognition may cause such security problems as authorization. To increase the recognition rate in such face database as ORL, a face recognition algorithm should be good at minimizing the disturbances of facial pose, illumination and expression (PIE) to this recognition. In this paper, the improved clonal selection algorithm and diverse samples are designed. The improved clonal selection algorithm searches the most similar sample for unknown face image, according to the affinity between the unknown one and the most similar sample. The affinity is newly designed to improve the adaptive matching between the object and the samples. Compared with some state-of-the-art algorithms on the ORL face database, the proposed approach outperforms the other algorithms in the recognition rate, based on the experimental results.*

**Keywords:** Face recognition; PIE; high-precision immune computation; security; clonal selection.

## 1. Introduction

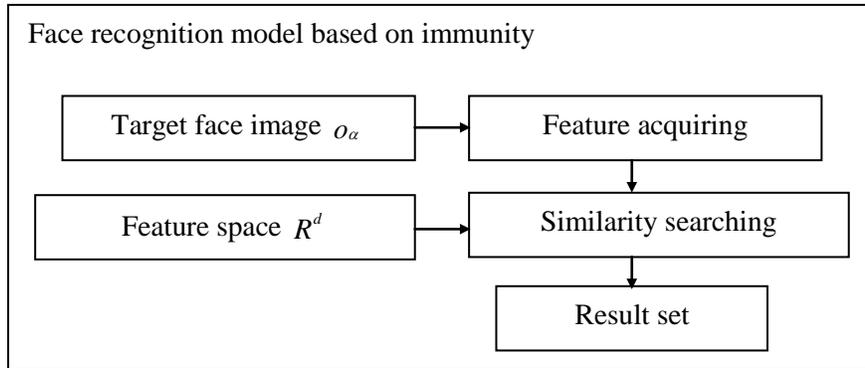
The high-precision face recognition is of high recognition rate of faces, and this is important for many applications of security, management and service. To increase the recognition rate, face recognition algorithms have to minimize the disturbances of facial pose, illumination, and expression (PIE) [1]. He et al. proposed a Laplacianface method of locality preserving projection (LPP) to preserve the local structure of training samples [2]. This approach was improved by Cai et al. with an orthogonal locality preserving projection (OLPP) to use more local information [3]. Jiang et al. proposed a method to enhance maximum likelihood face recognition [4]. In this paper, a novel high-precision face recognition approach is proposed with an improved clonal selection algorithm of immune computation. It filters some disturbances of PIE and hence boosts the high accuracy of the face recognition approach.

## 2. Face Recognition Model based on High-precision Immune Computation

The face recognition model based on immunity is proposed and comprised of feature acquiring of the target face images, similarity searching in the feature space  $R^d$  of known face images, and classification of the target face images, as shown in Figure 1. Consider  $N$   $d$ -dimensional samples  $x_1, x_2, \dots, x_N$ , which constitute  $N_c$  classes of faces. Let  $x_j^i$  denote a sample in the feature space  $R^d$  representing the  $j$ th sample in the  $i$ th

class of size  $N_i$ . The known samples  $x_1, x_2, \dots, x_N$  are used to train the face recognition model based on immunity, and this model is built on the strategy for searching the most similar sample of the unknown objects  $o_\alpha, \alpha = 1, 2, \dots, N_o$  in the feature space  $R^d$  of the samples. The antigens  $Ag_\alpha, \alpha = 1, 2, \dots, N_o$  and the antibodies  $Ab_m, m = 1, 2, \dots, N$  are the two repertoires of face image matrixes in the improved clonal selection algorithm.

$$N = \sum_{i=1}^{N_c} N_i. \quad (1)$$



**Figure 1. Face Recognition Model based on Immunity**

In the pattern recognition cases of traditional clonal selection algorithms, the affinity measure was calculated with the Hamming distance between the antigen  $Ag_\alpha$  and an antibody  $Ab_m$  [5]. Because the binary coding of the face images causes the loss of some image pixel data, the affinity measure is improved and calculated below.

$$M_m = \sum_{k=1}^{Ltp} (\delta_{mk} + \rho_{mk}),$$

where  $Ltp = Mtp \times Ntp, \delta_{mk} = \begin{cases} u, & \text{if } Ab_{mk} \pm \varepsilon \in Ag_\alpha \\ 0, & \text{otherwise} \end{cases},$

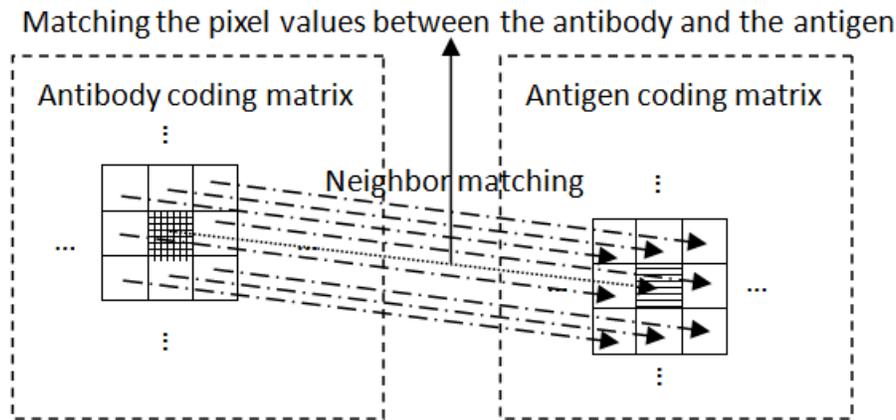
$$\rho_{mk} = \begin{cases} v, & \text{if } Ab_{mk} \pm \varepsilon = Ag_{\alpha\omega}, Ab_{m(k-1)} \pm \varepsilon = Ag_{\alpha(\omega-1)} \\ v, & \text{if } Ab_{mk} \pm \varepsilon = Ag_{\alpha\omega}, Ab_{m(k+1)} \pm \varepsilon = Ag_{\alpha(\omega+1)} \\ v, & \text{if } Ab_{mk} \pm \varepsilon = Ag_{\alpha\omega}, Ab_{m(k-Ntp)} \pm \varepsilon = Ag_{\alpha(\omega-Ntp)} \\ v, & \text{if } Ab_{mk} \pm \varepsilon = Ag_{\alpha\omega}, Ab_{m(k+Ntp)} \pm \varepsilon = Ag_{\alpha(\omega+Ntp)} \\ v, & \text{if } Ab_{mk} \pm \varepsilon = Ag_{\alpha\omega}, Ab_{m(k-Ntp-1)} \pm \varepsilon = Ag_{\alpha(\omega-Ntp-1)}, \\ v, & \text{if } Ab_{mk} \pm \varepsilon = Ag_{\alpha\omega}, Ab_{m(k-Ntp+1)} \pm \varepsilon = Ag_{\alpha(\omega-Ntp+1)} \\ v, & \text{if } Ab_{mk} \pm \varepsilon = Ag_{\alpha\omega}, Ab_{m(k+Ntp-1)} \pm \varepsilon = Ag_{\alpha(\omega+Ntp-1)} \\ v, & \text{if } Ab_{mk} \pm \varepsilon = Ag_{\alpha\omega}, Ab_{m(k+Ntp+1)} \pm \varepsilon = Ag_{\alpha(\omega+Ntp+1)} \\ 0, & \text{otherwise} \end{cases} \quad (2)$$

$$1 \leq \omega \leq Ltp, u + v = 1.$$

Here,  $M_{tp}$  and  $N_{tp}$  represents the pixel lengths of the face images,  $Ab_{mk}$  represents the  $k$ th pixel value of the antibody  $Ab_m$  for matching the face image,  $Ag_{\alpha\omega}$  represents the  $\omega$ th pixel value of the antigen  $Ag_\alpha$  for the face image,  $Ag_{\alpha(\omega-1)}, Ag_{\alpha(\omega+1)}, Ag_{\alpha(\omega-Ntp)}, Ag_{\alpha(\omega+Ntp)},$

$Ag_{\alpha(\omega-Nip-1)}$ ,  $Ag_{\alpha(\omega-Nip+1)}$ ,  $Ag_{\alpha(\omega+Nip-1)}$ , and  $Ag_{\alpha(\omega+Nip+1)}$  respectively represents the values of left-neighbor, right-neighbor, upper-neighbor, bottom-neighbor, left-upper-neighbor, right-upper-neighbor, left-bottom-neighbor, and right-bottom-neighbor for the  $\omega$ th pixel value of the face image antigen  $Ag_{\alpha}$ , and  $Ag_{\alpha\omega}$  represents the  $\omega$ th pixel value of the antigen  $Ag_{\alpha}$  for the face image.

The neighbors of the  $k$ th pixel value of the antibody  $Ab_m$  for matching the face image use the same positioning value in Figure 2 as the  $\omega$ th pixel value of the antigen  $Ag_{\alpha}$  for the face image. Figure 2 shows the similarity matching between the antibodies and the antigens and their small differences are permitted for the similarity.



**Figure 2. Affinity Measure on Matching between the Antibodies and the Antigens for Face Recognition**

### 3. Improved Clonal Selection Algorithm for Face Recognition

Based on the characteristics of the faces with the disturbances of PIE and the improved affinity measure, the clonal selection algorithm is improved for better application in the face recognition below.

Input:  $N$  training samples  $x_1, x_2, \dots, x_N$ , and the target face images  $o_{\alpha}, \alpha = 1, 2, \dots, N_o$ .

Output: The recognition type of the target face images  $\{o_{\alpha}\}$ .

Step 1: Initialize the parameters of the feature space, the feature-acquiring operator and the operator for searching the most similar sample.

Step 2: Build the feature space  $R^d$  with the feature vectors of the samples  $x_1, x_2, \dots, x_N$  and represent the vectors as the antibodies  $Ab_m, m = 1, 2, \dots, N$ .

Step 3: Acquire the feature information of the image  $o_{\alpha}$ , and design the antigen  $Ag_{\alpha}$ .

Step 4: Search the most similar sample of the object  $o_{\alpha}$  with the improved clonal selection algorithm, by matching the antigen  $Ag_{\alpha}$  with any of the antibodies  $Ab_m, m = 1, 2, \dots, N$  according to the improved affinity measure.

Step 5: Output the recognition type of the target face image  $o_{\alpha}$ .

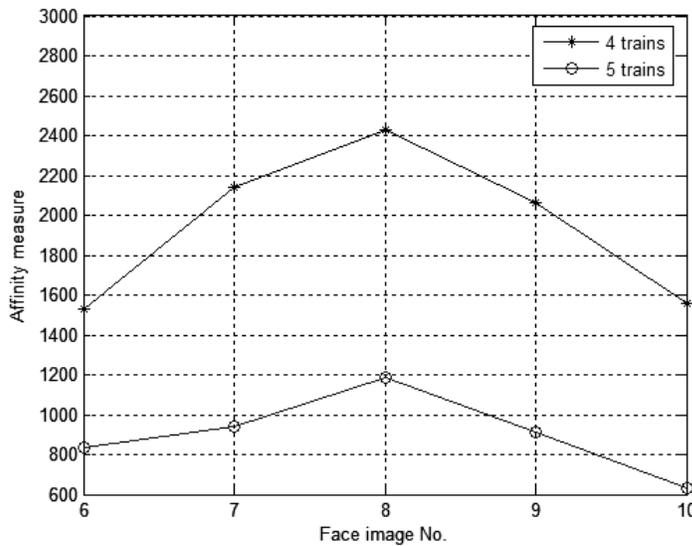
### 4. Experimental Results

Some experiments were conducted to test the effectiveness of the proposed approach, and the benchmark ORL face database was used [6]. The ORL face data set is comprised of 400 images of size 92 by 112 and 40 individuals (10 images per person), with different PIE. All testing samples are matched with the training samples using some matching rules, and the sums of the training samples are respectively 4 and 5. The highest recognition rates of various training samples in the ORL face database for the comparing algorithms are listed in Table 1, and the standard derivations for these data sets are also compared in Table 1 to analyze the robustness of the proposed method.

**Table 1. Recognition Rates on the ORL Databases (per cent)**

Rate type	The highest		Average
Number of trains	4 Trains	5 Trains	4 Trains
PCA+LPPFace	91.25(41)	97.00(51)	$89.95 \pm 1.3$
PCA+OLPPFace	94.17(37)	97.00(22)	$92.37 \pm 1.8$
PCA+ONPDA	91.92(61)	95.25(27)	$89.92 \pm 2.0$
PCA+MFA	89.16(69)	95.00(69)	$87.16 \pm 2.0$
PCA+NFLE $F_1$	94.17(41)	95.00(22)	$91.97 \pm 2.2$
PCA+NFLE $F_2$	94.58(39)	97.00(24)	$92.38 \pm 2.2$
Novel approach	<b>100</b>	<b>100</b>	<b><math>99.17 \pm 0.9</math></b>

Figure 3 shows that the affinity measures of recognizing the face images of No. 6 and No. 10 with failure are almost both the lowest ones in the 4 trains, and the affinity measure of recognizing the face image of No. 10 with failure is also the lowest one in the 5 train. From this table, the recognition rate of the proposed algorithm is maximal and the standard derivation of the proposed algorithm is minimal. Therefore, the performance of the novel face recognition approach is better than those of the other algorithms.



**Figure 3. Affinity measures of recognizing the face images in the 4 trains and the 5 trains of the ORL face database**

## 5. Conclusions

The improved clonal selection frame of immune computation [7-10] and the diverse samples are useful for recognize the face images with high recognition rates. The face recognition model based on the immunity can maximize the effectiveness of the feature search of the most similar samples to the target face images, and the affinity measure should be improved for matching the face feature with small differences due to the disturbances of PIE. The diversity and typicality are important to keep the high recognition rates [11], because the face recognition is sensitive to the unknown features of the face images. The high accuracy of the proposed approach is substantiated by the experiments on the ORL face database.

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