Facial Age Estimation by Learning from Label Distributions

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Abstract

One of the main difficulties in facial age estimation is the lack of sufficient training data for many ages. Fortunately, the faces at close ages look similar since aging is a slow and smooth process. Inspired by this observation, in this paper, instead of considering each face image as an example with one label (age), we regard each face image as an example associated with a label distribution. The label distribution covers a number of class labels, representing the degree that each label describes the example. Through this way, in addition to the real age, one face image can also contribute to the learning of its adjacent ages. We propose an algorithm named IIS-LLD for learning from the label distributions, which is an iterative optimization process based on the maximum entropy model. Experimental results show the advantages of IIS-LLD over the traditional learning methods based on single-labeled data.

Introduction

With the progress of aging, the appearance of human faces exhibits remarkable changes, as the typical example shown in Figure 1. The facial appearance is a very important trait when estimating the age of a human. However, the human estimation of age is usually not as accurate as other facial information such as identity, expression and gender. Hence developing automatic facial age estimation methods that are comparable or even superior to the human ability in age estimation becomes an attractive yet challenging topic emerging in recent years.

An early work on *exact* age estimation was reported by Lanitis et al. (2002; 2004), where the aging pattern was represented by a quadratic function called *aging function*, and the WAS (Weighted Appearance Specific) method (Lanitis, Taylor, and Cootes 2002) and AAS (Appearance and Age Specific) method (Lanitis, Draganova, and Christodoulou 2004) were proposed. Geng et al. (2006; 2007) proposed the AGES algorithm based on the subspace trained on a data

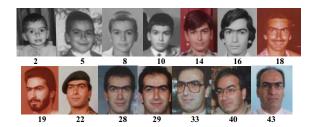


Figure 1: The aging faces of one subject in the <u>FG-NET</u> <u>Database</u>. The real ages are given at the bottom.

structure called aging pattern vector. Later, various methods have been developed for facial age estimation. For example, Fu et al. (2007; 2008) proposed an age estimation method based on multiple linear regression on the discriminative aging manifold of face images. Guo et al. (Guo et al. 2008) used the SVR (Support Vector Regression) method to design a locally adjusted robust regressor for prediction of human ages. Yan et al. (2007b) regarded age estimation as a regression problem with nonnegative label intervals and solved the problem through SDP (semidefinite programmingP). They also proposed an EM (Expectation-Maximization) algorithm to solve the regression problem and speed up the optimization process (Yan et al. 2007a). By using SFP (Spatially Flexible Patch) as the feature descriptor, the age regression was further improved with patchbased GMM (Gaussian Mixture Model) (Yan et al. 2008) and patch-based HMM (Hidden Markov Model) (Zhuang et al. 2008).

One of the main challenges of facial age estimation is the lack of sufficient training data (Geng, Zhou, and Smith-Miles 2007). A suitable training set should contain multiple images from the same person covering a wide age range. Since the aging progress cannot be artificially controlled, the collection of such a dataset usually requires great efforts in searching for the images taken years ago, and can do nothing to acquire future images. Consequently, the available datasets typically just contain a limited number of aging images for each person, and the images at the higher ages are especially rare.

Without sufficient training data, additional knowledge about the aging faces can be introduced to reinforce the learning process. By another close look at Figure 1, one

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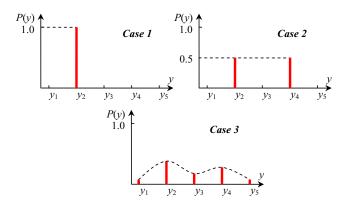


Figure 2: Three cases of label distribution.

may find that the faces at the close ages look quite similar, which results from the fact that aging is a slow and gradual progress. Inspired by this observation, the basic idea behind this paper is to utilize the images at the neighboring ages while learning a particular age.

The utilization of adjacent ages is achieved by introducing a new labeling paradigm, i.e., assigning a label distribution to each image rather than a single label of the real age. A suitable label distribution will make a face image contribute to not only the learning of its real age, but also the learning of its neighboring ages. Accordingly, a learning algorithm different from traditional learning schemes is proposed for learning from the label distributions.

The rest of the paper is organized as follows. First, the concept of label distribution is introduced. Then, an algorithm named IIS-LLD is proposed for learning from label distributions. After that, the new learning scheme is evaluated in the experiments. Finally, conclusions are drawn.

Label Distribution

In a label distribution, a real number $P(y) \in [0,1]$ is assigned to each label y, representing the degree that the corresponding label describes the instance. The numbers for all the labels sum to 1, meaning full description of the instance. Under this definition, the traditional ways to label an instance with a single label or multiple labels can be viewed as special cases of label distribution. Some examples of typical label distributions for five class labels are shown in Figure 2. For $Case\ I$, a single label is assigned to the instance, so $P(y_2)=1$ means the class label y_2 fully describes the instance. For $Case\ 2$, two labels $(y_2\ and\ y_4)$ are assigned to the instance, so each of them only describes 50% of the instance, i.e., $P(y_2)=P(y_4)=0.5$. $Case\ 3$ represents a general case of label distribution with the only constraint $\sum_y P(y)=1$.

Special attention should be paid to the meaning of P(y), which is *not* the *probability* that class y correctly labels the instance, but the *degree* that y describes the instance, or in other words, the *proportion* of y in a full class description of the instance. Thus all the labels with a non-zero P(y) in the distribution are 'correct' labels for the instance but just with different importance reflected by P(y). Recognizing this, one can distinguish label distribution from the previous

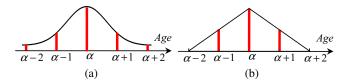


Figure 3: Typical label distributions for age α : (a) Gaussian distribution, (b) Triangle distribution. Both of them are used in the proposed algorithm and tested in the experiments.

studies on *possibilistic labels* (Denoeux and Zouhal 2001) or *uncertain labels* (Quost and Denoeux 2009), where the basic assumption is that there is only one 'correct' label for each instance. Although not a probability by definition, the operations on P(y) are similar to those for probability since $P(y) \in [0,1]$ and $\sum_y P(y) = 1$. Thus, without confusion, the terminology for probability is also used for label distribution in the rest of the paper.

It is also worthwhile to distinguish the 'label distribution' from the 'category membership' used in *fuzzy classification* (Duda, Hart, and Stork 2001). The 'category membership' function is introduced into fuzzy classification to covert an objective measure (e.g., 'length = 1cm') of the instance into a subjective category (e.g., 'short length'), which is a conceptual characteristic of the instance. Thus 'category membership' embodies the ambiguity in the features of the instance, while the final class label of the instance is unambiguous. Since the features of the instance are conceptual, pure fuzzy methods are usually based on the knowledge of the designer rather than learning from examples (Duda, Hart, and Stork 2001). On the other hand, 'label distribution' represents the ambiguity in the class label of the instance, while the features of the instance are unambiguous. Thus the learning from examples style is feasible for the 'label distribution' problem.

As to the problem of facial age estimation, for a particular face image, not only its real age, but also the adjacent ages are used to describe the class of the image. The label distribution assigned to a face image with the real age α should satisfy the following two properties: 1. The probability of α in the distribution is the highest, which ensures the leading position of the real age in the class description; 2. The probability of other ages decreases with the distance away from α , which makes the age closer to the real age contribute more to the class description. While there are many possibilities for this, Figure 3 shows two typical label distributions for the images with the real age α , i.e., the Gaussian distribution and the Triangle distribution. The parameters of the distributions are determined by the training data (further explored in the later experiments).

Learning from Label Distributions

Let $\mathcal{X} = \mathbb{R}^d$ denote the input space and $\mathcal{Y} = \{y_1, y_2, \cdots, y_c\}$ denotes the finite set of possible class labels. The problem of learning from label distributions (LLD) can be formally described as: Given a training set $S = \{(\boldsymbol{x}_1, P_1(y)), (\boldsymbol{x}_2, P_2(y)), \cdots, (\boldsymbol{x}_n, P_n(y))\}$, where $\boldsymbol{x}_i \in \mathcal{X}$ is an in-

stance, $P_i(y)$ is the distribution of the random variable $y \in \mathcal{Y}$ associated with x_i , and the goal is to learn a conditional p.d.f. p(y|x), where $x \in \mathcal{X}$ and $y \in \mathcal{Y}$.

Suppose p(y|x) is a parametric model $p(y|x;\theta)$, where θ is the vector of the model parameters. Given the training set S, the goal of LLD is to find the θ that can generate a distribution similar to $P_i(y)$ given the instance x_i . Here the Kullback-Leibler divergence is used as the measurement of the similarity between two distributions. Thus the best model parameter vector θ^* is determined by

$$\boldsymbol{\theta}^* = \underset{\boldsymbol{\theta}}{\operatorname{argmin}} \sum_{i} \sum_{y} \left(P_i(y) \log \frac{P_i(y)}{p(y|\boldsymbol{x}_i; \boldsymbol{\theta})} \right)$$
$$= \underset{\boldsymbol{\theta}}{\operatorname{argmax}} \sum_{i} \sum_{y} P_i(y) \log p(y|\boldsymbol{x}_i; \boldsymbol{\theta}). \tag{1}$$

It is interesting to examine the traditional learning paradigms under the optimization criterion shown in Eq. (1). For *supervised learning*, each instance is associated with a single label (see *Case 1* in Figure 2), thus $P_i(y) = \delta(y, y_i)$, where $\delta(\cdot, \cdot)$ is the Kronecker function and y_i is the class label of x_i . Consequently, Eq. (1) can be simplified to the maximum likelihood criterion.

For *multi-label learning* (Zhang and Zhou 2007), each instance is associated with a label set (see *Case 2* in Figure 2). Consequently, Eq. (1) can be changed into

$$\theta^* = \underset{\theta}{\operatorname{argmax}} \sum_{i} \frac{1}{p_i} \sum_{y \in Y_i} \log p(y|x_i; \theta),$$
 (2)

where Y_i is the label set associated with x_i and $p_i = |Y_i|$. Eq. (2) can be viewed as a maximum likelihood criterion weighted by the reciprocal cardinality of the label set associated with each instance. In fact, this is equivalent to first applying the Entropy-based Label Assignment (ELA) (Tsoumakas and Katakis 2007), a well-known technique dealing with multi-label data, to transform the multi-label instances into the weighted single-label instances, and then optimizing the maximum likelihood criterion based on the weighted single-label instances.

Suppose $f_k(x, y)$ is a *feature function* which depends on both the instance x and the label y. Then, the expected value of f_k w.r.t. the empirical joint distribution $\tilde{p}(x, y)$ in the training set is

$$\tilde{f}_k = \sum_{\boldsymbol{x}, y} \tilde{p}(\boldsymbol{x}, y) f_k(\boldsymbol{x}, y). \tag{3}$$

The expected value of f_k w.r.t. the conditional model $p(y|x;\theta)$ and the empirical distribution $\tilde{p}(x)$ in the training set is

$$\hat{f}_k = \sum_{\boldsymbol{x}, y} \tilde{p}(\boldsymbol{x}) p(y|\boldsymbol{x}; \boldsymbol{\theta}) f_k(\boldsymbol{x}, y). \tag{4}$$

One reasonable choice of $p(y|x; \theta)$ is the one that has the maximum *conditional entropy*

$$H = -\sum_{\boldsymbol{x},y} \tilde{p}(\boldsymbol{x}) p(y|\boldsymbol{x};\boldsymbol{\theta}) \log p(y|\boldsymbol{x};\boldsymbol{\theta}), \tag{5}$$

subject to the constraint $\tilde{f}_k = \hat{f}_k$. It can be proved (Berger, Pietra, and Pietra 1996) that such a model (a.k.a. *maximum entropy model*) has the exponential form

$$p(y|\boldsymbol{x};\boldsymbol{\theta}) = \frac{1}{Z} \exp\left(\sum_{k} \theta_{k} f_{k}(\boldsymbol{x}, y)\right),$$
 (6)

where $Z = \sum_{y} \exp\left(\sum_{k} \theta_{k} f_{k}(\boldsymbol{x}, y)\right)$ is the normalization factor and θ_{k} 's are the model parameters. In practice, the features usually depend only on the instance but not on the class label, thus Eq. (6) can be rewritten as

$$p(y|\boldsymbol{x};\boldsymbol{\theta}) = \frac{1}{Z} \exp\left(\sum_{k} \theta_{y,k} g_{k}(\boldsymbol{x})\right), \tag{7}$$

where $g_k(x)$ is a <u>class-independent feature function</u>.

Substituting Eq. (7) into the optimization criterion used in Eq. (1) and recognizing $\sum_y P_i(y) = 1$ yields the target function of θ

$$T(\boldsymbol{\theta}) = \sum_{i,y} P_i(y) \log p(y|\boldsymbol{x}_i; \boldsymbol{\theta})$$

$$= \sum_{i,y} P_i(y) \sum_k \theta_{y,k} g_k(\boldsymbol{x}_i) - \sum_i \log \sum_y \exp \left(\sum_k \theta_{y,k} g_k(\boldsymbol{x}_i)\right). \quad (8)$$

Directly setting the gradient of Eq. (8) w.r.t. θ to zero does not yield a closed form solution. Thus the optimization of Eq. (8) uses a strategy similar to IIS (Improved Iterative Scaling) (Pietra, Pietra, and Lafferty 1997), a well-known algorithm for maximizing the likelihood of the maximum entropy model. IIS starts with an arbitrary set of parameters; then for each step, it updates the current estimate of the parameters θ with $\theta + \Delta$, where Δ maximizes a lower-bound of the likelihood changes between the adjacent steps. This iterative process, nevertheless, needs to be migrated to the new target function $T(\theta)$. Furthermore, the constraint needed for IIS on the feature functions $f_k(x,y) \geq 0$ (hence $g_k(x) \geq 0$) should be removed to ensure the freedom in choosing any feature extractors suitable for the data.

In detail, the change of $T(\theta)$ between the adjacent steps is

$$T(\boldsymbol{\theta} + \boldsymbol{\Delta}) - T(\boldsymbol{\theta}) = \sum_{i,y} P_i(y) \sum_k \delta_{y,k} g_k(\boldsymbol{x}_i) - \sum_i \log \sum_y p(y|\boldsymbol{x}_i; \boldsymbol{\theta}) \exp\left(\sum_k \delta_{y,k} g_k(\boldsymbol{x}_i)\right), \quad (9)$$

where $\delta_{y,k}$ is the increment for $\theta_{y,k}$. Applying the inequation $-\log x \geq 1-x$ yields

$$T(\boldsymbol{\theta} + \boldsymbol{\Delta}) - T(\boldsymbol{\theta}) \ge \sum_{i,y} P_i(y) \sum_k \delta_{y,k} g_k(\boldsymbol{x}_i) + n - \sum_{i,y} p(y|\boldsymbol{x}_i; \boldsymbol{\theta}) \exp\left(\sum_k \delta_{y,k} g_k(\boldsymbol{x}_i)\right).$$
(10)

Algorithm 1: IIS-LLD

Input: The training set $S = \{(\boldsymbol{x}_i, P_i(y))\}_{i=1}^n$, the feature functions $g_k(\boldsymbol{x})$ Output: The conditional p.d.f. $p(y|\boldsymbol{x};\boldsymbol{\theta})$ 1 Initialize the model parameter vector $\boldsymbol{\theta}^{(0)}$;
2 $i \leftarrow 0$;
3 repeat
4 $i \leftarrow i+1$;
5 Solve Eq. (14) for $\delta_{y,k}$;
6 $\boldsymbol{\theta}^{(i)} \leftarrow \boldsymbol{\theta}^{(i-1)} + \boldsymbol{\Delta}$;
7 until $T(\boldsymbol{\theta}^{(i)}) - T(\boldsymbol{\theta}^{(i-1)}) < \varepsilon$;
8 $p(y|\boldsymbol{x};\boldsymbol{\theta}) \leftarrow \frac{1}{Z} \exp\left(\sum_k \theta_{y,k}^{(i)} g_k(\boldsymbol{x})\right)$;

Differentiating the right side of Eq. (10) w.r.t. $\delta_{y,k}$ yields coupled equations of $\delta_{y,k}$ which are hard to be solved. To decouple the interaction between $\delta_{y,k}$, Jansen's inequality is applied here, i.e., for a p.d.f. p(x),

$$\exp\left(\sum_{x} p(x)q(x)\right) \le \sum_{x} p(x) \exp q(x). \tag{11}$$

The last term of Eq. (10) can then be written as

$$\sum_{i,y} p(y|\boldsymbol{x}_i;\boldsymbol{\theta}) \exp\left(\sum_k \delta_{y,k} s(g_k(\boldsymbol{x}_i)) g^{\#}(\boldsymbol{x}_i) \frac{|g_k(\boldsymbol{x}_i)|}{g^{\#}(\boldsymbol{x}_i)}\right), (12)$$

where $g^{\#}(\boldsymbol{x}_i) = \sum_k |g_k(\boldsymbol{x}_i)|$ and $s(g_k(\boldsymbol{x}_i))$ is the sign of $g_k(\boldsymbol{x}_i)$. Since $|g_k(\boldsymbol{x}_i)|/g^{\#}(\boldsymbol{x}_i)$ can be viewed as a p.d.f., Eq. (10) can be rewritten as

$$T(\boldsymbol{\theta} + \boldsymbol{\Delta}) - T(\boldsymbol{\theta}) \ge \sum_{i,y} P_i(y) \sum_k \delta_{y,k} g_k(\boldsymbol{x}_i) + n - \sum_{i,y} p(y|\boldsymbol{x}_i; \boldsymbol{\theta}) \sum_k \frac{|g_k(\boldsymbol{x}_i)|}{g^{\#}(\boldsymbol{x}_i)} \exp(\delta_{y,k} s(g_k(\boldsymbol{x}_i)) g^{\#}(\boldsymbol{x}_i)).$$
(13)

Denote the right side of Eq. (13) as $\mathcal{A}(\Delta|\theta)$, which is a lower-bound of $T(\theta + \Delta) - T(\theta)$. Setting the derivative of $\mathcal{A}(\Delta|\theta)$ w.r.t. $\delta_{y,k}$ to zero gives

$$\frac{\partial \mathcal{A}(\boldsymbol{\Delta}|\boldsymbol{\theta})}{\partial \delta_{y,k}} = \sum_{i} P_{i}(y)g_{k}(\boldsymbol{x}_{i}) -$$

$$\sum_{i} p(y|\boldsymbol{x}_{i};\boldsymbol{\theta}) g_{k}(\boldsymbol{x}_{i}) \exp(\delta_{y,k} s(g_{k}(\boldsymbol{x}_{i})) g^{\#}(\boldsymbol{x}_{i})) = 0.$$
 (14)

What is nice about Eq. (14) is that $\delta_{y,k}$ appears alone, and therefore can be solved one by one through nonlinear equation solvers, such as the Gauss-Newton method. This algorithm, called IIS-LLD, is summarized in Algorithm 1.

After p(y|x) is learned from the training set, given a new instance x', its label distribution p(y|x') can be first calculated. The availability of the explicit label distribution for x' provides many possibilities in classifier design. To name just a few, if the expected class label for x' is single, then the predicted label could be $y^* = \operatorname{argmax}_{y} p(y|x')$,

together with an confidence measure $p(y^*|x')$. If multiple labels are allowed, then the predicted label set could be $L = \{y|p(y|x') > \xi\}$, where ξ is a predefined threshold. Moreover, all the labels in L can be ranked according to their probabilities.

Experiments

Methodology

The dataset used in the experiments is the FG-NET Aging Database (Lanitis, Taylor, and Cootes 2002). There are 1,002 face images from 82 subjects in this database. Each subject has 6-18 face images at different ages. Each image is labeled by its real age. The ages are distributed in a wide range from 0 to 69. Besides age variation, most of the age-progressive image sequences display other types of facial variations, such as significant changes in pose, illumination, expression, *etc.* A typical aging face sequence in this database is shown in Figure 1.

According to the real age associated to each image in the FG-NET Database, a label distribution is generated using the Gaussian or Triangle distribution shown in Figure 3. Then the IIS-LLD algorithm is applied to the image set with generated label distributions. The predicted age for a test image x' is determined by $y^* = \operatorname{argmax}_{u} p(y|x')$. To study the usefulness of the adjacent ages, IIS-LLD is also applied to the special label distribution of the single-labeled data shown as Case 1 in Figure 2. The three label distributions are denoted by 'Gaussian', 'Triangle', and 'Single', respectively. Several existing algorithms specially designed for the problem of facial age estimation are compared as baseline methods, which include AGES (Geng, Zhou, and Smith-Miles 2007) and two aging function regression based methods, i.e., WAS (Lanitis, Taylor, and Cootes 2002) and AAS (Lanitis, Draganova, and Christodoulou 2004). Some conventional general-purpose classification methods for singlelabel data are also compared, including KNN (K-Nearest Neighbors), BP (Back Propagation neural network), C4.5 (C4.5 decision tree) and SVM (Support Vector Machine).

As an important baseline, the human ability in age perception is also tested. About 5% samples of the database (51 face images) are randomly selected and presented to 29 human observers. There are two stages in the experiments. In each stage, the images are randomly presented to the observers, and the observers are asked to choose an age from 0 to 69 for each image. The difference of the two stages is that in the first stage (HumanA), only the grayscale face regions (i.e., the color images are converted to grayscale and the background is removed) are shown, while in the second stage (HumanB), the whole color images are shown. HumanA intends to test the age estimation ability purely based on the face image intensity, which is also the input to the algorithms, while HumanB intends to test the human estimation ability based on multiple traits including face, hair, skin color, clothes, and background.

The feature extractor (i.e., $g_k(x)$) used here is the Appearance Model (Edwards, Lanitis, and Cootes 1998). The main advantage of this model is that the extracted features combine the shape and the intensity of the face images, both

Table 1: Mean Absolute Error (in Years) of Age Estimation on the FG-NET Aging Database

Method	IIS-LLD			ACEC	WA C	A A C	1.NINI	DD.		CVM	Human Observers ¹	
	Gaussian	Triangle	Single	AGES	WAS	AAS	kNN	BP	C4.5	SVM	HumanA	HumanB
MAE	<u>5.77</u>	<u>5.90</u>	6.27	6.77	8.06	14.83	8.24	11.85	9.34	7.25	8.13	6.23

 $^{^1}$ The human observers are tested on 5% samples of the database.

are important in the aging progress. The extracted features require 200 model parameters to retain about 95% of the variability in the training data, i.e., $k=1,\cdots,200$.

For IIS-LLD, when generating the label distributions, if not explicitly mentioned, the standard deviation of the Gaussian distribution is 1, and the bottom length of the Triangle distribution is 6. For all other algorithms, several parameter configurations are tested and the best result is reported. For AGES, the aging pattern subspace dimensionality is set to 20. In AAS, the error threshold in the appearance cluster training step is set to 3, and the age ranges for the age specific classification are set as 0-9, 10-19, 20-39 and 40-69. The *K* in *K*NN is set to 30. The BP neural network has a hidden layer of 100 neurons. The parameters of C4.5 are set to the default values of the J4.8 implementation. SVM uses the RBF kernel with the inverse width of 1.

The algorithms are tested through the LOPO (Leave-One-Person-Out) mode (Geng et al. 2006), i.e., in each fold, the images of one person are used as the test set and those of the others are used as the training set. After 82 folds, each subject has been used as test set once, and the final results are calculated from all the estimates.

Results

The performance of the age estimation is evaluated by MAE (Mean Absolute Error), i.e., the average absolute difference between the estimated age and the real age. The results are tabulated in Table 1. The algorithms performing better than HumanA are highlighted by boldface and those better than HumanB are underlined. As can be seen, the overall performance of IIS-LLD is significantly better than those of the single-label based algorithms, either specially designed for age estimation or for general-purpose classification. Except for the 'Single' distribution case, being slightly worse than HumanB, IIS-LLD performs even better than the human observers. Further looking into the three distributions that IIS-LLD works on, the MAE can be ranked as: Gaussian < Triangle < Single. The Gaussian distribution utilizes all ages, the Triangle distribution utilizes those ages within the triangle, and the Single distribution only utilizes the real age. This supports the idea to use suitable label distributions to cover as many as possible correlated class labels.

In addition to the coverage of the distribution, the performance of LLD can also be affected by how the related labels are covered, i.e., the parameters of the label distributions. Figure 4 shows the MAE of IIS-LLD on the Gaussian distribution with different standard deviations $\sigma=0,\cdots,4$, and the Triangle distribution with different bottom length l=2,4,6,8,10. Note that both $\sigma=0$ and l=2 correspond to the 'Single' distribution. Figure 4 reveals that too

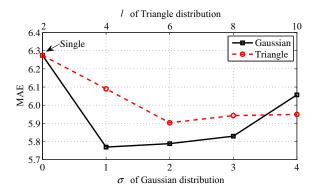


Figure 4: The MAE of IIS-LLD on different Gaussian distributions (with different σ) and different Triangle distributions (with different l).

Table 2: MAE in Different Age Ranges

D	ДС1		ACEC			
Range	#Samples	Gaussian	Triangle	Single	AGES	
0-9	371	2.83	2.83	3.06	2.30	
10-19	339	5.21	5.17	4.99	3.83	
20-29	144	6.60	6.39	6.72	8.01	
30-39	79	11.62	11.66	12.10	17.91	
40-49	46	12.57	15.78	18.89	25.26	
50-59	15	21.73	22.27	27.40	36.40	
60-69	8	24.00	26.25	32.13	45.63	

concentrative ($\sigma=0, l=2,4$) or too dispersive distribution ($\sigma=2,3,4, l=8,10$) would lead to performance deterioration. This is consistent with our intuition that the related classes are helpful but should not threaten the priority of the original class. A balanced choice of the scale of the distribution is crucial to achieve a good performance.

Due to different aging rate in different aging stages (e.g., the facial appearance changes faster during childhood than middle age), it is reasonable to assume different distribution width for different ages. A further experiment on IIS-LLD with the Gaussian distribution uses different σ for different ages based on the similarity among adjacent ages. In detail, the similarity of an image to an adjacent age is calculated as the reciprocal distance to the mean image of that age. Then, σ is obtained by fitting a Gaussian distribution to the normalized similarities. The new algorithm achieves a better performance MAE=5.70.

The MAE of IIS-LLD and the best single-label based method AGES are also compared in different age ranges in Table 2. It is interesting to see that in the age ranges 0-9 and 10-19, all the three cases of IIS-LLD perform worse

than AGES. This is not difficult to understand since IIS-LLD is based on the general-purpose maximum entropy model while AGES builds on a problem-specific data structure called aging pattern vector whose effect becomes more apparent when there are sufficient training data. This can be evidenced by the the 'Single' case of IIS-LLD, which is equivalent to learning a maximum entropy model using the maximum likelihood criterion but performs worse than AGES. A comparison between 'Gaussian'/'Triangle' and 'Single' reveals that the introduction of label distribution does not cause remarkable performance deterioration at the ranges with sufficient training samples. The main advantage of LLD comes from the classes with insufficient training samples. For example, in the range 60-69 with minimum training samples, the MAE of 'Gaussian' is 25% lower than that of 'Single', and 47% lower than that of AGES. This validates that LLD is an effective way to relieve the 'lack of training samples' problem.

Conclusions

This paper proposes a novel method for facial age estimation based on learning from label distributions (LLD). By extending the single label of an instance to a label distribution, one instance can contribute to the learning of multiple classes. It is particularly useful when dealing with the problems where: 1) the classes are correlated to each other, and 2) the training data for some classes are insufficient. An algorithm named IIS-LLD is proposed to learn from the label distributions. Experimental results on facial age estimation validate the advantages of utilizing the related classes via label distributions.

While achieving good performance on facial age estimation, LLD might also be useful for other learning problems. Generally speaking, there are at least three scenarios where LLD could be helpful:

- 1. The instances are initially labeled with class distributions. The class distributions might come from the knowledge of experts or statistics.
- 2. Some classes are highly correlated with other classes. According to the correlation among the classes, a label distribution can be naturally generated.
- The labeling from different sources are controversial. Rather than deciding a single wining label, it might be better to generate a label distribution which incorporates the information from all sources.

Further research on LLD in these cases will be an attractive future work.

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