Learning to Rank for Blind Image Quality Assessment

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Abstract

Blind image quality assessment (BIQA) aims to predict perceptual image quality scores without access to reference images. State-of-the-art BIQA methods typically require subjects to score a large number of images to train a robust model. However, the acquisition of image quality scores has several limitations: 1) scores are not precise, because subjects are usually uncertain about which score most precisely represents the perceptual quality of a given image; 2) subjective judgments of quality may be biased by image content; 3) the quality scales between different distortion categories are inconsistent, because images corrupted by different types of distortion are evaluated independently in subjective experiments; and 4) it is challenging to obtain a large scale database, or to extend existing databases, because of the inconvenience of collecting sufficient images associated with different kinds of distortion that have diverse levels of degradation, training the subjects, conducting subjective experiments, and realigning human quality evaluations.

To combat these limitations, this paper explores and exploits preference image pairs (PIPs) such as “the quality of image $I_a$ is better than that of image $I_b$” for training a robust BIQA model. The preference label, representing the relative quality of two images, is generally precise and consistent, and is not sensitive to image content, distortion type, or subject identity; such PIPs can be generated at very low cost. The proposed BIQA method is one of learning to rank. We first formulate the problem of learning the mapping from the image features to the preference label as one of classification. In particular, we investigate the utilization of a multiple kernel learning algorithm based on group lasso (MKLGL) to provide a solution. A simple but effective strategy to estimate perceptual image quality scores is then presented. Experiments show that the proposed BIQA method is highly effective and achieves comparable performance to state-of-the-art BIQA algorithms. Moreover, the proposed method can be easily extended to new distortion categories.
I. INTRODUCTION

Blind image quality assessment (BIQA) aims to predict perceptual image quality scores without access to reference images. Because reference images are usually unavailable in most practical applications, BIQA is of great significance and has consequently received tremendous attention over the past decades. To date, a number of universal BIQA methods that work well for various distortion types have been deployed [1]-[10]. These methods typically require subjects to score a large number of images to learn a robust model, but the acquisition of image quality scores in this way has several limitations.

First, scores are not precise. In standard subjective studies [11]-[16], the assessor assigns each image a number within a range, e.g. 0 to 100, that reflects its perceived quality. There is usually uncertainty about which score most precisely represents the perceptual quality of a given image [17], however, and an arbitrary score from the selected range [18]. Consequently, this score may not capture subtle differences in the perceived quality of images [19]. Fig. 1 illustrates two images from the Laboratory for Image and Video Engineering (LIVE) database [11]. For each image, the difference mean opinion score (DMOS), $s$, and the realigned DMOS, $s_r$, is listed. DMOS is the difference between the mean opinion score (MOS) and perfect quality [11]. The realigned DMOS is obtained by realigning the DMOSs between different subjects and between different distortion types. A larger $s$ or $s_r$ value indicates poorer quality. It is notable that the left image is subjectively better than the right one, but the DMOSs do not accurately reflect their relative quality. The success of $s_r$ implies the necessity for realignment.

Fig. 1: Illustrations of images contained in the LIVE database. Left: JPEG2000 compressed image, $s = 60.77$, $s_r = 70.84$; right: JPEG compressed image, $s = 58.63$, $s_r = 88.92$.

Fig. 2: Two undistorted images with different content. Both are of perfect quality.
Second, subjective judgments of quality may be biased by image content. Images whose impairment scales are similar but whose content is different may be assigned different scores by observers as a result of personal image content preferences. Fig. 2 illustrates two undistorted images, both of which are of perfect quality. However, individuals may prefer the left image and evaluate its quality with a higher score because they find it aesthetically more pleasing than the right image. The content-dependent problem further decreases the reliability of subjective scores.

Third, the quality scales between different sessions are inconsistent. In subjective experiments, the evaluation of all the test images is divided into several independent sessions with respect to the distortion type [11][15] or image content [13][16] to minimize the effects of observer fatigue [20]. Thus, images which have similar quality scores but are evaluated in different sessions may not be perceptually similar to each other. Fig. 3 shows two images included in the LIVE database. Fig. 3a is an image distorted in a fast fading Rayleigh (wireless) channel (FF), with $s = 43.69$ and $s_r = 35.42$. Fig. 3b is an image corrupted by the adaptive Gaussian white noise (WN), with $s = 32.03$ and $s_r = 35.32$. Although their realigned DMOSs are almost the same, the impairment in Fig. 3b is much more annoying than it is in Fig. 3a. Again, the raw DMOS fails to reflect the relative quality.

Finally, it is challenging to obtain a large scale database or to extend existing databases, mainly for the following inconvenient reasons: 1) The test organizer has to collect sufficient images associated with each kind of distortion at diverse levels of degradation to yield a meaningful evaluation [20]; 2) One observer has to continuously evaluate many images in a single session to minimize the influence of contextual effects that the score to an image is highly biased by recently scored images [21][22]; 3)
Subjective experiments should be conducted in critical viewing conditions and in critical procedures, because judgments of perceptual quality scores are sensitive to the viewing environment [19][20]; 4) It is necessary to recruit many observers and train them before the experiment, to minimize quality scale mismatch errors [11][19][20]; and 5) The process of realigning raw human responses is complicated [11].

These limitations constrain the reliability and extension of existing BIQA methods which utilize subjective quality scores as the benchmark for learning a model. Although there have been several BIQA methods that do not require learning from subjectively evaluated images [23]-[25], their performance is still inferior to state-of-the-art algorithms.

To combat the aforementioned limitations, this paper explores and exploits preference image pairs (PIPs) such as “the quality of image \( I_a \) is better than that of image \( I_b \)” for training a robust BIQA model. The preference label, representing the relative quality of two images, is generally precise and consistent, and is not sensitive to image content, distortion type, viewing conditions, or subject identity [19][26], and such PIPs can be generated at very low cost [26]. Consider the image pairs derived from the images shown in Figs. 3 and 4. They are easy to discriminate in terms of perceived quality. Moreover, different subjects may give consistent preference labels [27], even though the images in a certain pair may vary in the types of distortion afflicting them (e.g. Figs. 4a and 4b), or in image content (e.g. Figs. 3a and 4b), or both (e.g Figs. 3b and 4b).

The proposed BIQA method is one of learning to rank [28]. We first formulate the problem of learning the mapping from the image features to the preference label as one of classification. In particular, we investigate using a multiple kernel learning algorithm based on group lasso (MKLGL) [29] to solve it. A simple but effective strategy is then presented to estimate perceptual image quality scores. Thorough experiments conducted on the largest four standard databases show that the proposed BIQA method is highly effective and achieves comparable performance to state-of-the-art BIQA algorithms. Moreover, the proposed method can be easily extended to new distortion categories by simply adding the corresponding PIPs into the training set.

The rest of the paper is organized as follows. In Section II we briefly introduce standard IQA databases and learning to rank. Section III discusses how to generate PIPs with valid preference labels. The framework of the proposed BIQA method is detailed in Section IV. Our experiments are described and analyzed in Section V and an extensive subjective study of PIPs is presented in Section VI. Section VII concludes this paper.
II. RELATED WORK

A. Existing IQA Databases

Subjective IQA studies are of fundamental importance for the development of IQA. Over the years, many researchers have contributed significant research in this area through the construction of various IQA databases. In this section, we introduce the composition of the four largest datasets, i.e. the LIVE dataset [12], Tampere Image Database 2013 (TID2013) [13], Categorical Subjective Image Quality (CSIQ) database [14], and LIVE multiple distorted (LIVEMD) database [15].

The LIVE database includes 808 images, which are generated from 29 original images by corrupting them with five types of distortion, i.e. JPEG2000 compression (JP2k), JPEG compression (JPEG), WN, Gaussian blur (Gblur), and FF. The DMOS and realigned DMOS of each image are available. Because the realigned DMOS is more precise than the DMOS, we adopt the realigned DMOS in our work. The realigned DMOS ranges from -3 to 112. For convenience, we refer to the realigned DMOS as DMOS in the remainder of this paper unless otherwise indicated. In addition, we refer to all the images associated with the same reference image as a “group”.

The TID2013 includes 3,000 images in sum. These images are generated by corrupting 25 original images with 24 types of distortion at 5 different levels. The distortion types include: WN (#1), additive white Gaussian noise which is more intensive in color components than in the luminance component (#2), additive Gaussian spatially correlated noise (#3), masked noise (#4), high frequency noise (#5), impulse noise (#6), quantization noise (#7), Gblur (#8), image denoising (residual noise, #9), JPEG (#10), JP2k (#11), JPEG transmission errors (#12), JPEG2000 transmission errors (#13), non-eccentricity pattern noise (#14), local block-wise distortion of different intensity (#15), mean shift (#16), contrast change (#17), change of color saturation (#18), multiplicative Gaussian noise (#19), comfort noise (#20), lossy compression of noisy images (#21), image color quantization with dither (#22), chromatic aberrations (#23), and sparse sampling and reconstruction (#24). The MOS of each image is available and ranges from 0.2 to 7.3.

The CSIQ database consists of 866 images which are derived from 30 original images. Six types of distortion are considered in CSIQ, i.e. WN, JPEG, JP2k, additive Gaussian pink noise (PN), Gblur, and global contrast decrements (GCD). The DMOS of each image is available and ranges from 0 to 1.

The LIVEMD database includes images distorted by multiple types of distortion. There are two subsets, one of which is associated with the images corrupted by Gblur followed by JPEG (GblurJPEG), and one which is associated with images corrupted by Gblur followed by WN (GblurWN). Each subset includes
225 images. The DMOS of each image is released and ranges from 0 to 85.

B. Learning to Rank

Learning to rank [28] involves learning a function that can predict the ranking list of a given set of stimuli. It is the central issue in web page ranking, document retrieval, image searching and other applications. Of the existing methods, the pairwise approach has been well deployed and successfully applied to information retrieval [30].

Without loss of generality, we take document retrieval as an example to introduce the pairwise approach. In pairwise approaches, document pairs are adopted as instances for learning a ranking function. What is needed is to construct a training set by first collecting document pairs from the ranking lists, and then calculating the corresponding difference feature vector for each pair of documents and assigning a preference label that represents their relative relevance [31]. In this way, the problem of learning-to-rank is reduced to a classification problem, also called preference learning problem [32], and existing classification models, such as support vector machines (SVMs) and Neural Network, can be directly applied to solve it [33]-[35].

Intuitively, the problem addressed in the pairwise approach is similar to the problem we aim to work out in this paper, and is our inspiration for thinking we may succeed in learning a robust BIQA model from PIPs. The proposed BIQA model is detailed in Section VI.

III. GENERATION OF PREFERENCE IMAGE PAIRS

Generating PIPs in an easy and efficient way is of fundamental significance in this research. In this section, we investigate how to produce image pairs with valid preference labels through paired comparisons and from existing IQA databases, based on the quality scores.

A. Paired Comparisons

In paired comparisons, observers are asked to choose which image in a pair has better perceived quality [26]. Consider a pair of images. If the difference in perceived quality between them is large enough, observers can easily discriminate between them, regardless of any type of corruption through distortion, image content, or even viewing conditions. Moreover, different observers may offer a unanimous judgment [27]. In this case, we can obtain valid preference labels by arranging only one observer to assign each image pair. This is consistent with the samples illustrated in Figs. 3 and 4.
However, when the distortion strengths of the two images are similar, it becomes much more difficult for observers to judge their relative quality, especially when the image content of the two images is diverse (e.g. Figs. 5b and 5c), or because of the distortion type (e.g. Figs. 5c and 5f), or both (e.g. Figs. 5b and 5f), or because there is subtle difference between them (e.g. Figs. 5a and 5b), numbers are incorrect. In this case, it is unreasonable neither to force observers to choose a “better” image [13][15], nor to label the relative quality as “the same” [20]. Otherwise, discrepancies are caused and responses from many observers are needed [19] so as to produce valid preference labels. The huge number of comparisons coupled with the need to recruit many observers [13][15] lead to the same limitations as introduced previously.

![Fig. 5: Distorted images: (a) JPEG compressed image; (b) JPEG compressed image; (c) image corrupted by FF; (d) image corrupted by WN.](image)

Thus we propose to allow observers to only judge the image pairs that are easy to distinguish in terms of perceived quality in paired comparisons. Since we aim to learn a BIQA model from image pairs, we do not require a complete comparison, only to label a sample of all possible pairs. We conclude the procedures of generating PIPs through paired comparisons as follows:

First, collect $n$ images, $I = \{I_1, \ldots, I_n\}$, $n > 2$, are diverse in the types of corrupting distortion, image content, and distortion strength.

Second, randomly construct $N$, $1 \leq N \leq n(n-1)/2$, pairs from the collected images. The image pair set $\mathcal{P}$ is given by

$$\mathcal{P} \subseteq \{(I_i, I_j) \mid i, j = 1, 2, \ldots, n\}.$$  \hspace{1cm} (1)

Note that if $(I_i, I_j) \in \mathcal{P}$, then $(I_j, I_i) \notin \mathcal{P}$, to reduce the number of comparisons.

Finally, recruit $S$, $S \geq 1$, subjects to assign preference labels for the image pairs. Given an pair $(I_i, I_j)$, if subject $s$, $s = 1, \ldots, S$, considers $I_i$ to be better than $I_j$ in quality, then the corresponding preference label $l_{k,s}$, $k = 1, 2, \ldots, N$, is set as $+1$; if $I_j$ is subjectively better than $I_i$, $l_{k,s} = -1$; and if subject $s$
does not label the pair \((I_i, I_j)\) or is uncertain about the relative quality of the corresponding images, let \(l_{k,s} = 0\). For simplicity, the final preference label for the pair \((I_i, I_j)\) is calculated as

\[
y_k = \text{sign} \left( \sum_{s=1}^{S} l_{k,s} \right).
\]

If \(y_k = 0\), it means \(I_i\) and \(I_j\) are difficult to discriminate in terms of perceived quality, and \((I_i, I_j)\) will not be adopted for training.

The procedure of the paired comparison experiment is flexible \[26\]. Observers do not have to evaluate many pairs in a single session, because there is practically no contextual effect or scale mismatch problem, and PIPs evaluated in different sessions can be directly aggregated into a single dataset for future applications without complex processing. In this paper, we present an extensive subjective paired-comparison study which we detail in Section [VI]

### B. Collecting PIPs from Existing IQA Databases

Since many efforts have contributed to the construction of various IQA databases \[12\]-\[15\], it is meaningful to find a method to generate reliable PIPs from existing databases. Previous discussions imply that a small difference between two quality scores may not accurately reflect the corresponding relative quality; thus, we recommend selecting the image pairs whose corresponding difference quality scores reach a threshold. The generation of PIPs from IQA databases is therefore formulated as below.

Assume that an IQA database comprises: 1) \(n\) labeled images: \(I = \{I_1, \ldots, I_n\}\); 2) the subjective quality scores of the images \(Q = \{q_1, \ldots, q_n\}\), \(q_i\) is the quality score of \(I_i\). Then we can construct \(N\) image pairs from these labeled images. Let \(P\) be the pair set, we have

\[
P \subseteq \{(I_i, I_j) \mid |q_i - q_j| > T, \ i, j = 1, \ldots, n\},
\]

where, \(T\) is the threshold of the difference quality score, and \(T \geq 0\); and \(|q_i - q_j|\) denotes the absolute value of \((q_i - q_j)\). Because a greater MOS value indicates better quality but a greater DMOS value is associated with poorer quality, a preference label \(y_k, k = 1, \ldots, N\) for each image pair \((I_i, I_j) \in P\) is assigned as:

\[
y_k = \begin{cases} 
\text{sign} (q_i - q_j), & \text{if the quality score is MOS} \\
-\text{sign} (q_i - q_j), & \text{if the quality score is DMOS}
\end{cases}
\]

Thus, \(y_k = 1\) indicates that \(I_i\) is better than \(I_j\) in terms of quality, but \(y_k = -1\) indicates that \(I_j\) is better. Let \(Y\) be the set of preference labels, such that
It is notable that the PIPs separately extracted from different IQA databases can be combined with no realignment. Moreover, we can aggregate the PIPs produced through paired comparisons and those collected from IQA databases into one single dataset for training a BIQA model. Based on all the previous discussions, we conclude that the generation of PIPs is easy and convenient, avoiding the limitations in the acquisition of human quality scores.

IV. THE PROPOSED BIQA METHOD

In this section, we present a BIQA framework that learns to predict perceptual quality scores from PIPs. Inspired by the pairwise learning-to-rank approaches, we first formulate the problem of learning the mapping from difference feature vectors to preference labels as one of classification. We utilize natural scene statistics (NSS) \([36]\)-based features to represent an image and investigate using the MKLGL approach \([29]\) to solve the classification problem. A simple but effective strategy is subsequently presented to estimate perceptual image quality scores. Fig. 6 shows the diagram of the proposed BIQA method, and details are introduced in the following subsection.

A. Integrated NSS Features

NSS-based image features have been widely explored for IQA and have shown promising performance \([1-7]\). However, the features utilized in these methods are generally representative of some distortion...
categories but have relatively weaker correlation with others. In this paper, therefore, we adopt a fusion of the features that have been utilized in several state-of-the-art BIQA methods, i.e. BLIINDS-II [1], BRISQUE-L [6], and SRNSS [7], to represent an image. For the integrity of this paper, we briefly introduce these features in this subsection. For detail, we refer to [1], [6], and [7].

Both the BLIINDS-II features and BRISQUE-L features are extracted over several scales, where the feature extraction is repeated after downsampling it by a factor of 2. BLIINDS-II extracts features over 3 scales and computes 8 features in the discrete cosine transform (DCT) domain at each scale. The BRISQUE-L features are exacted over 2 scales, and 18 spatial-domain features are extracted at each scale. The BLIINDS-II and BRISQUE-L features extracted at each scale are concentrated into a vector and denoted as \( f_{n}^{BLD} \), \( n = 1, 2, 3 \), and \( f_{m}^{BRL} \), \( m = 1, 2 \), respectively. In addition, let \( f^{BLD} \) and \( f^{BRL} \) be all the BLIINDS-II and BRISQUE-L features, respectively.

In SRNSS, He et al. [7] employed the mean, variance, and entropy of wavelet coefficients in each sub-band to encode the generalized spectral behavior, the energy fluctuations, and the generalized information, respectively. In the implementation, an image is decomposed into 4 scales. The coefficients at LH (low, high) and HL (high, low) sub-bands at a particular scale are combined to calculate a group of features due to their similarity in statistics. Let \( m \), \( v \), and \( e \) respectively denote all the means, variances, and entropies extracted from an image, and \( f^{SRN} \) be all the SRNSS features extracted from an image.

Finally we denote all the features extracted from an image as \( f^{all} \), such that

\[
\begin{align*}
    f^{all} = [f^{BLD}, f^{BRL}, f^{SRN}] = [f^{BLD}_{1}, f^{BLD}_{2}, f^{BLD}_{3}, f^{BRL}_{1}, f^{BRL}_{2}, m, v, e].
\end{align*}
\]  

Thus \( f^{all} \) is of the dimension 84 (8\( \times \)3 of \( f^{BLD} \) + 18\( \times \)2 of \( f^{BRL} \) + 8\( \times \)3 of \( f^{SRN} \)).

We calculated Pearson’s linear correlation coefficient (PLCC) between each of these features and the quality score, and plotted the maximum PLCC values of \( f^{BLD} \), \( f^{BRL} \), and \( f^{SRN} \), respectively, across all the images contained in each type of distortion and the whole image set (ALL) in the LIVE, CSIQ, TID2013, and LIVEMD databases in Fig. 7. As is clear, \( f^{BLD} \), \( f^{BRL} \), and \( f^{SRN} \) do not correlate highly with all the distortion categories. It is therefore reasonable to combine these features as the representation of the image [37]. In Section [V] we compare the performance of the BIQA methods that utilize \( f^{BLD} \), \( f^{BRL} \), \( f^{SRN} \), and \( f^{all} \), respectively, to verify the effectiveness of this feature fusion strategy.

B. Preparation of Training Data

In line with the discussions in Section [III] we first collect a number of PIPs with valid preference labels. We then calculate the features of each image and the difference feature vector of each PIP. Thus
Fig. 7: Maximum PLCCs of $f^{BLD}$, $f^{BRL}$, and $f^{SRN}$ across all the images contained in each type of distortion and the whole image set in the LIVE, LIVEMD, CSIQ, and TID2013 databases.

The training data comprises:

1) $n$ images: $\mathcal{I} = \{I_1, \ldots, I_n\}$;
2) for each image an integrated feature vector $f^\text{all}_i \in \mathbb{R}^d$, $i=1,\ldots,n$, $d$ is the dimension of the feature vector;
3) $N$ PIPs generated from the $n$ images. The PIP set is denoted as $\mathcal{P}$: $\mathcal{P} \subseteq \{(I_i, I_j) \mid i, j = 1, \ldots, n\}$; and
4) for each PIP $(I_i, I_j) \in \mathcal{P}$, a difference feature vector $x_k$ and a preference label $y_k$, $k = 1, \ldots, N$:

$$x_k = f^\text{all}_i - f^\text{all}_j,$$

and

$$y_k \in \{-1, +1\}.$$
where $x_{N+k} = -x_k$, $y_{N+k} = -y_k$. The mapping from difference feature vectors to preference labels can then be realized by a binary classifier trained on $\{X, Y\}$.

### C. Preference Learning via MKLGL

Previous discussion in Part A implies that the difference feature vector includes 8 portions in accordance with $f_{BLD}^1$, $f_{BLD}^2$, $f_{BLD}^3$, $f_{BRL}^1$, $f_{BRL}^2$, $m$, $v$, and $e$, each of which represents the image at a particular scale or captures one particular property. We therefore introduce multiple kernel learning (MKL) to measure the similarity of different portions of features using different kernels [38]. Of the various MKL algorithms, the MKLGL approach [29] has shown its efficiency and effectiveness across a wide range of applications [39]. Thus we adopt MKLGL to solve the preference learning problem in the proposed BIQA framework.

The prediction function that maps the difference feature vector to the preference label can then be represented by

$$f(x) = \sum_{k=1}^{2N} \alpha_k y_k \sum_{m=1}^{M} \theta_m K_m(x, x_k),$$

where the optimal parameters $\alpha^* = \{\alpha_k^*\}_{k=1}^{2N}$, $\theta = \{\theta_m^*\}_{m=1}^{M}$, and $b^*$ are learned from the training data $\{X, Y\}$; $K_m$, $m = 1, \ldots, M$, is a base kernel and defines a feature mapping from the original input space to a reproducing kernel Hilbert space (RKHS) $H_m$.

Base kernels may be diverse in the type or parameters of the kernel function, or in the portion of features on which the kernel function operates [38]. In our research, we construct Gaussian kernels with 5 different bandwidths ($\{2^{-2}, 2^{-1}, \ldots, 2^2\}$) on each portion of the difference feature vector. In addition, we exploit Gaussian kernels with 5 different bandwidths ($\{2^{-2}, 2^{-1}, \ldots, 2^2\}$) using the entire difference feature vector to encode the potential correlations between different portions of features. In sum, we have 45 kernels (8 portions of features $\times$ 5 bandwidths Gaussian kernels + 5 bandwidths Gaussian kernels). We construct a kernel in this way according to [29][39] to avoid using a large memory.

The learned MKLGL model can estimate the preference label of a given pair of images based on the corresponding difference feature vector $x_{test}$. In addition, if $x_{test} = 0$, we consider the two images in the test pair to be of the same perceived quality, and assign the predicted preference label as 0.

### D. Quality Prediction

Consider a full round of comparisons of all the training images. We term the sum of the ideal preference labels $\{y_{i,j}\}_{j=1, j\neq i}^{n} \in \{1, -1\}^{n-1}$ associated with $\{(I_i, I_j)\}_{j=1, j\neq i}^{n}$ as the "gain" of $I_i$ and denote it as


g_i, i = 1, \ldots, n, \text{ such that }

\begin{equation}
g_i = \sum_{j=1, j\neq i}^{n} y_{i,j}, \text{ and } (n - 1) \leq g_i \leq (n - 1), \forall i. \tag{12}
\end{equation}

The gain of a training image is proportional to its perceived quality, because $y_{i,j}$ essentially reflects the quality of $I_i$ relative to $I_j$, and $y_{i,j} = 1$ indicates $I_i$ is better than $I_j$ in terms of quality. For simplicity, we assume a linear mapping between the gain value and the ideal perceived quality scores of the training images: $q_i = ag_i + b$, where $q_i$ is the quality score of $I_i$. Because neither all the ideal preference labels nor the ideal quality scores are available in the scenario of the proposed BIQA framework, we investigate to estimate the parameters $a$ and $b$ based on the following two assumptions:

1) The training images cover the full range of possible quality, from the poorest (extremely annoying) to the best (undistorted), with small steps; and

2) The gain value $(n - 1)$ corresponds to the greatest quality score, and $-(n - 1)$ corresponds to the lowest score.

These assumptions are reasonable because it is easy to collect/generate sufficient images that diverse greatly in perceived quality, in which each pair includes an undistorted image and a heavily distorted one. Without loss of generalization, we choose the continuous quality scale of $[0, 100]$. $a$ and $b$ can then be estimated by solving the following linear equation:

\begin{equation}
\begin{bmatrix}
  n - 1 & 1 \\
  -(n - 1) & 1
\end{bmatrix}
\begin{bmatrix}
a \\
b
\end{bmatrix} =
\begin{bmatrix}
100 \\
0
\end{bmatrix}.
\tag{13}
\end{equation}

We have $a = 50/(n - 1)$ and $b = 50$.

Given a test image $I_t$, we then pair it with each training image, and calculate the corresponding difference feature vector:

\begin{equation}
x_{t,i} = f_{t}^{all} - f_{i}^{all} \tag{14}
\end{equation}

where $f_{t}^{all}$ is the integrated feature vector of the test image, $f_{i}^{all}$ is the feature vector of the $i$th training image $I_i$, $i = 1, 2, \ldots, n$. The preference labels associated with the test pairs are estimated by the trained MKLGL. Let $\hat{Y}_t = \{\hat{y}_{t,1}, \ldots, \hat{y}_{t,n}\} \in \{+1, 0, -1\}^n$ denote the predicted preference labels. Finally, the gain of the test image is calculated by $\hat{g}_t = \sum_{i=1}^{n} \hat{y}_{t,i}$, and the quality score is then predicted by:

\begin{equation}
\hat{q}_t = 50 \left( \frac{\hat{g}_t}{n - 1} + 1 \right). \tag{15}
\end{equation}
It is notable that $\hat{g}_t$ ranges from $-n$ to $n$, thus $\hat{q}_t$ may become less than 0, which is a situation that occurs when the test image is perceptually worse than the poorest training image. Conversely, when the test image is better than all the training images, the predicted quality score becomes greater than 100, which is taken to indicate quality improvement. This is an interesting feature, albeit possibly controversial, since it is not clear whether this is due to the absence of the potentially most/least annoying image in the training set or a true improvement of the test image in terms of quality.

Even though this quality prediction approach seems rather ad-hoc at first sight, it correlates highly with human perceptions of quality, as verified by the thorough experiments presented in the following two sections.

V. IQA Experiments

We conduct a series of experiments on the four largest datasets, i.e. the LIVE dataset, TID2013, CSIQ, and LIVEMD databases, with four objectives.

1) The first objective is to show how the performance of the proposed BIQA method varies with the parameters included in it. This is illustrated through the experiment on the LIVE database, as presented in Part A.

2) The second objective is to evaluate the effectiveness of our BIQA model. This is verified by the experiments where we train and test the BIQA models on two distinct subsets of each database. Details are described in Part B.

3) The third objective is to show that the proposed BIQA approach is database independent. In this case, we train the model on the whole LIVE database and then test it on the other databases, as introduced in Part C.

4) The fourth objective is to show that it is easy to extend the proposed BIQA framework to emerging distortion categories. This is demonstrated by the experiment in which we aggregate the PIPs separately extracted from the LIVE, CSIQ, and TID2013 databases into a single training set for learning a robust model. Details are presented in Part D.

We compare the performance of the proposed BIQA method with state-of-the-art BIQA algorithms which have been reported as having the best performance \[6,8\], i.e., BLIINDS-II \[1\], BRISQUE \[4\], BRISQUE-L \[6\], and CORNIA \[8\]. In particular, we adopt the version of BLIINDS-II that utilizes support vector regression (SVR) with a radial basis function (RBF) kernel to model the relationship between $f^{BLD}$ and quality scores. All of these algorithms are implemented based on the codes provided by the authors and the corresponding literature.
To verify the efficacy of adopting a combination of NSS features, we implement two BIQA algorithms that use $f^{SRN}$ and $f^{all}$ to represent an image, respectively, and utilize SVR with a RBF kernel to learn a quality prediction function. For convenience, we refer to them as $f^{SRN}$SVR and $f^{all}$SVR, respectively. It is worth declaring that the performance of $f^{SRN}$ SVR is somewhat better than SRNSS [7], thus we do not report the results of SRNSS in this paper owing to space constraints. In addition, to verify the dependency between $f^{all}$ and the MKLGL approach, we test the performance of the proposed framework using a support vector machine (SVM) with a RBF kernel instead of MKLGL.

In our experiments, the LIBrary for Support Vector Machines (LIBSVM) package [40] is used to implement the SVR and SVM algorithms. The optimal parameters of SVR and SVM algorithms are learned by 5-fold cross validation on the training set. MKLGL is implemented by the MKL toolbox [39] with default settings.

To evaluate the performances of the proposed method, three indexes are adopted as the criteria between the predicted quality scores by the BIQA algorithm and DMOS/MOS: Kendall’s Rank Correlation Coefficient (KRCC), Pearson’s Linear Correlation Coefficient (PLCC), and Spearman’s Rank Correlation Coefficient (SRCC). Greater KRCC, PLCC and SRCC values indicate better consistency with human opinions of quality. We perform a nonlinear mapping using a logistic function as described in [41] before computing these indexes.

In all the experiments, we conducted the train-test procedures 100 times to verify the robustness of the proposed method. In Parts A, B, and D, the database in each train-test procedure was randomly split into distinct training and test subsets and $N$ PIPs were randomly generated from the training images for training the proposed BIQA model. In Part C, we randomly produced $N$ PIPs from the entire LIVE database and used all the images contained in the TID2013, CSIQ, and LIVEMD databases as the test set in each train-test procedure. All the other BIQA methods adopt the training images as the instances in learning, thus we only needed to run the train-test procedure once for them in Part C. The median performance indexes across the 100 trials were reported for comparison.

A. Variation with Algorithm Parameters

In this subsection, we focus on the impact of choosing different algorithm parameters: 1) the threshold $T$ for generating PIPs from existing IQA databases, 2) the number of training images, $n$, and 3) the number of PIPs, $N$. We randomly choose $N_g$, $1 \leq N_g \leq 28$, groups of images contained in the LIVE database for training, and use the rest for testing. Thus $n \approx [808 \times N_g/29]$. In the experiments, we considered the parameters of: $N_g \in \{5, 10, 15, 20\}$, $T = 0, 10, \ldots, 70$, and $N \in \{500, 1000, 1500, 2000\}$.
TABLE I: Maximum Number of Pairs That Can Be Generated ($\times 10^3$).

<table>
<thead>
<tr>
<th>$N_g$</th>
<th>$T$</th>
<th>0</th>
<th>10</th>
<th>20</th>
<th>30</th>
<th>40</th>
<th>50</th>
<th>60</th>
<th>70</th>
</tr>
</thead>
<tbody>
<tr>
<td>5</td>
<td></td>
<td>9.3</td>
<td>7.6</td>
<td>5.9</td>
<td>4.3</td>
<td>3.0</td>
<td>2.0</td>
<td>1.2</td>
<td>0.6</td>
</tr>
<tr>
<td>10</td>
<td></td>
<td>38.2</td>
<td>31.1</td>
<td>24.4</td>
<td>18.3</td>
<td>13.1</td>
<td>8.8</td>
<td>5.5</td>
<td>3.2</td>
</tr>
<tr>
<td>15</td>
<td></td>
<td>86.7</td>
<td>70.4</td>
<td>55.2</td>
<td>41.4</td>
<td>29.7</td>
<td>19.9</td>
<td>12.4</td>
<td>7.1</td>
</tr>
<tr>
<td>20</td>
<td></td>
<td>155.4</td>
<td>125.9</td>
<td>98.9</td>
<td>74.4</td>
<td>53.5</td>
<td>36.1</td>
<td>22.5</td>
<td>12.8</td>
</tr>
<tr>
<td>29</td>
<td></td>
<td>326.0</td>
<td>264.5</td>
<td>207.6</td>
<td>156.2</td>
<td>111.9</td>
<td>75.7</td>
<td>47.1</td>
<td>27.0</td>
</tr>
</tbody>
</table>

Fig. 8: Effect of the threshold $T$ used to generate PIPs from the LIVE database.

It is worth noting that the number of PIPs adopted for training is $N_p = \min (N, M)$, where $M$ is the maximum number of pairs we can generate in accordance with certain $T$ and $N_g$. We report the proximate $M$ in the LIVE databases in Table I. It is clear that in most cases, we only utilized a very small portion of all the possible pairs for training the proposed BIQA model.

Fig. 8 shows how the performance on the entire test set varies with the settings of $T$ when $N_g = 20$ and $N = 2000$. As is clear, the values of all the criteria first slightly increase and then gently descend as the thresholds become greater. This is mainly because we can reduce the noisy data contained in the training set by merely choosing the pairs whose quality scores differ sufficiently. However, when the thresholds become too great, the loss of information caused by abandoning image pairs plays a more significant role. As a result, the performance of the proposed BIQA method decreases. It is inspiring that the performance of the proposed BIQA method is acceptable even when $T = 70$. Similar phenomena were observed under other settings of $N_g$ and $N$ as well as across various IQA databases.

To verify the effect of $n$ ($N_g$) and $N$, we fixed $T = 10$. As is shown in Fig. 9, the performance of the proposed BIQA method monotonously increases as the number of images or PIPs included in the training set rises. Moreover, the number of images has more significant impact on the performance than the number of PIPs. For example, the performance when $N_g = 20$ and $N = 500$ is much better than
Fig. 9: Effect of the number of images and the number of PIPs contained in the training set (tested on the LIVE database).

when $N_g = 5$ and $N = 2000$. This is mainly because more comparisons are conducted for a test image when there are more training images, leading to a more precise estimation of its quality score. This result suggests that when researchers try to construct a PIP dataset, they can collect many training images but only randomly label a very small portion of all the possible image pairs.

B. Consistency with Human Opinions

We utilized all four databases to test the performance of the proposed BIQA method and compare it with BLIINDS-II, BRISQUE-L, CORNIA, $f^{SRN}$SVR, and $f^{all}$SVR.

On the LIVE database, we chose $N_g \in \{5, 10, 15, 20\}$ groups of images for training and the remaining images for the test set. The threshold and number of PIPs were fixed as $T = 10$ and $N = 2000$. The performance indexes of these BIQA methods on the entire test dataset are tabulated in Table III. In addition, the best BIQA method for each index is highlighted in boldface.

It is notable that the performance of the proposed method is much better than BRISQUE-L, BLIINDS-II, and $f^{SRN}$SVR when $N_g = 5$. When the number of training images increases, it still performs better than BLIINDS-II, and $f^{SRN}$SVR and is highly comparable to BRISQUE-L, CORNIA, and $f^{all}$SVR. Given that the proposed method utilizes much less information than the other methods which use DMOS in learning (as little as 1.3% of all the possible image pairs when $T = 10$ and $N_g = 20$), this is a solid achievement.

The fact that $f^{all}$SVR performs consistently better than BRISQUE-L, BLIINDS-II, and $f^{SRN}$SVR verifies the effectiveness of combining existing NSS features to represent an image. Further, the algorithm using the proposed framework and MKLGL (Proposed) consistently outperforms that using SVM (Proposed$^SVM$), demonstrating the important correlation between MKLGL and the integrated features.
### TABLE II: Overall Performance of BIQA Methods on the LIVE Database

<table>
<thead>
<tr>
<th>Method</th>
<th>SRCC</th>
<th>KRCC</th>
<th>PLCC</th>
<th>SRCC</th>
<th>KRCC</th>
<th>PLCC</th>
<th>SRCC</th>
<th>KRCC</th>
<th>PLCC</th>
<th>SRCC</th>
<th>KRCC</th>
<th>PLCC</th>
</tr>
</thead>
<tbody>
<tr>
<td>BRISQUE-L</td>
<td>0.872</td>
<td>0.689</td>
<td>0.883</td>
<td>0.919</td>
<td>0.755</td>
<td>0.923</td>
<td>0.934</td>
<td>0.780</td>
<td>0.938</td>
<td>0.942</td>
<td>0.795</td>
<td>0.945</td>
</tr>
<tr>
<td>BLINDS-II</td>
<td>0.849</td>
<td>0.663</td>
<td>0.858</td>
<td>0.885</td>
<td>0.706</td>
<td>0.892</td>
<td>0.913</td>
<td>0.750</td>
<td>0.920</td>
<td>0.927</td>
<td>0.771</td>
<td>0.930</td>
</tr>
<tr>
<td>f^{SRN}SVR</td>
<td>0.890</td>
<td>0.714</td>
<td>0.894</td>
<td>0.915</td>
<td>0.749</td>
<td>0.918</td>
<td>0.927</td>
<td>0.772</td>
<td>0.929</td>
<td>0.938</td>
<td>0.787</td>
<td>0.940</td>
</tr>
<tr>
<td>f^{all}SVR</td>
<td>0.907</td>
<td>0.741</td>
<td>0.913</td>
<td>0.927</td>
<td>0.774</td>
<td>0.932</td>
<td>0.938</td>
<td>0.790</td>
<td>0.940</td>
<td>0.946</td>
<td>0.803</td>
<td>0.949</td>
</tr>
<tr>
<td>CORNIA</td>
<td>0.908</td>
<td>0.733</td>
<td>0.909</td>
<td>0.931</td>
<td>0.767</td>
<td>0.930</td>
<td>0.936</td>
<td>0.777</td>
<td>0.936</td>
<td>0.942</td>
<td>0.787</td>
<td>0.942</td>
</tr>
<tr>
<td>Proposed</td>
<td>0.909</td>
<td>0.743</td>
<td>0.911</td>
<td>0.925</td>
<td>0.767</td>
<td>0.927</td>
<td>0.933</td>
<td>0.778</td>
<td>0.933</td>
<td>0.938</td>
<td>0.790</td>
<td>0.940</td>
</tr>
<tr>
<td>Proposed-SVM</td>
<td>0.883</td>
<td>0.706</td>
<td>0.888</td>
<td>0.914</td>
<td>0.747</td>
<td>0.917</td>
<td>0.924</td>
<td>0.763</td>
<td>0.929</td>
<td>0.930</td>
<td>0.774</td>
<td>0.932</td>
</tr>
</tbody>
</table>

### Table III: Settings for Constructing Training Data In Each Database.

<table>
<thead>
<tr>
<th>Database</th>
<th>Ng</th>
<th>N</th>
<th>MOS/DMOS Scale</th>
<th>T</th>
</tr>
</thead>
<tbody>
<tr>
<td>LIVE</td>
<td>20</td>
<td>2000</td>
<td>(-3, 112)</td>
<td>10</td>
</tr>
<tr>
<td>CSIQ</td>
<td>20</td>
<td>2000</td>
<td>[0, 1]</td>
<td>0.1</td>
</tr>
<tr>
<td>TID2013</td>
<td>17</td>
<td>2000</td>
<td>(0.2, 7.3)</td>
<td>1</td>
</tr>
<tr>
<td>LIVEMD</td>
<td>10</td>
<td>2000</td>
<td>(0, 85)</td>
<td>10</td>
</tr>
</tbody>
</table>

In addition, the performance of Proposed-SVM is also better or highly comparable with state-of-the-art algorithms, corroborating the validity of the proposed BIQA framework.

For each of the CSIQ, TID2013, and LIVEMD databases, we randomly select 2/3 of all the dataset content for training and the remaining 1/3 for test in each trial. There are 6 and 24 distortion types in the CSIQ and TID2013 databases, respectively. However, the features adopted in the proposed BIQA method are not representative of all of them, as shown in Fig. 7. Thus we test all the BIQA methods only on the distortion subsets with which f^{all} correlates highly, i.e. WN, JPEG, JP2k, and Blur in the CSIQ database, and ## 1-11, 19, 21-24 in TID2013. The parameters for constructing the training data were chosen according to the experiments on the LIVE database as well as the quality scale in the corresponding database. Details are listed in Table [III](#).

Table [II](#) shows that the SRCC, KRCC and PLCC produce unanimous comparison results, thus we only report the SRCC values in this subsection for clarity. The SRCCs of the BIQA methods for each distortion category and the entire dataset (ALL or ALLsub, where ALLsub refers to “all the subsets”) are listed in Table [IV](#). The performance on the LIVE database corresponds to Ng = 20. In addition, we compute the average SRCC over the four databases for each method and report it in the last column.
### TABLE IV: Performance of BIQA Methods Across Various Distortion Categories of The TID2013, LIVE, CSIQ, LIVEMD Databases.

<table>
<thead>
<tr>
<th></th>
<th>TID2013</th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>#1</td>
<td>#2</td>
<td>#3</td>
<td>#4</td>
<td>#5</td>
<td>#6</td>
<td>#7</td>
<td>#8</td>
<td>#9</td>
<td>#10</td>
<td>#11</td>
<td>#12</td>
<td>#13</td>
<td>#14</td>
<td>#15</td>
<td>#16</td>
<td>ALLsub</td>
</tr>
<tr>
<td>BRISQUE-L</td>
<td>0.808</td>
<td>0.682</td>
<td>0.815</td>
<td>0.489</td>
<td>0.871</td>
<td>0.847</td>
<td>0.744</td>
<td>0.853</td>
<td>0.652</td>
<td>0.831</td>
<td>0.818</td>
<td>0.787</td>
<td>0.652</td>
<td>0.860</td>
<td>0.757</td>
<td>0.864</td>
<td>0.759</td>
</tr>
<tr>
<td>BLIINDS-II</td>
<td>0.543</td>
<td>0.444</td>
<td>0.602</td>
<td>0.335</td>
<td>0.700</td>
<td>0.632</td>
<td>0.641</td>
<td>0.825</td>
<td>0.744</td>
<td>0.806</td>
<td>0.854</td>
<td>0.515</td>
<td>0.174</td>
<td>0.733</td>
<td>0.614</td>
<td>0.830</td>
<td>0.615</td>
</tr>
<tr>
<td>f^SRN^SVR</td>
<td>0.686</td>
<td>0.534</td>
<td>0.637</td>
<td>0.459</td>
<td>0.846</td>
<td>0.820</td>
<td>0.790</td>
<td>0.922</td>
<td>0.796</td>
<td>0.870</td>
<td>0.895</td>
<td>0.685</td>
<td>0.472</td>
<td>0.853</td>
<td>0.786</td>
<td>0.920</td>
<td>0.750</td>
</tr>
<tr>
<td>f^all^SVR</td>
<td>0.744</td>
<td>0.573</td>
<td>0.765</td>
<td>0.392</td>
<td>0.851</td>
<td>0.787</td>
<td>0.753</td>
<td>0.878</td>
<td>0.767</td>
<td>0.841</td>
<td>0.887</td>
<td>0.703</td>
<td>0.677</td>
<td>0.859</td>
<td>0.726</td>
<td>0.898</td>
<td>0.771</td>
</tr>
<tr>
<td>CORNIA</td>
<td>0.666</td>
<td>0.511</td>
<td>0.772</td>
<td>0.424</td>
<td>0.789</td>
<td>0.624</td>
<td>0.747</td>
<td>0.876</td>
<td>0.808</td>
<td>0.841</td>
<td>0.824</td>
<td>0.617</td>
<td>0.775</td>
<td>0.766</td>
<td>0.719</td>
<td>0.892</td>
<td>0.779</td>
</tr>
<tr>
<td>Proposed</td>
<td>0.701</td>
<td>0.564</td>
<td>0.692</td>
<td>0.409</td>
<td>0.837</td>
<td>0.776</td>
<td>0.665</td>
<td>0.898</td>
<td>0.776</td>
<td>0.832</td>
<td>0.901</td>
<td>0.620</td>
<td>0.615</td>
<td>0.815</td>
<td>0.754</td>
<td>0.892</td>
<td>0.779</td>
</tr>
</tbody>
</table>

of the table. The average SRCC is calculated by weighting the SRCC values according to the number of images contained in each database according to [42]. The best BIQA algorithm for each dataset is highlighted in boldface.

It is rather inspiring that our approach statistically yields the best result across all databases. The proposed method coupled with CORNIA shows the best performance on the TID2013 and LIVEMD databases. Moreover, across all the distortion categories of all the databases, the performance of the proposed method is highly comparable with state-of-the-art BIQA algorithms. It is notable that BRISQUE-L performs poorly on the CSIQ database. This is mainly due to the weak correlation between f^BRL^ and the quality scores of the CSIQ database, as illustrated in Fig. 7. In contrast, the proposed method combats this drawback, again demonstrating the efficacy of employing the fused NSS features.

Scatter plots (for each of the entire LIVE, LIVEMD, CSIQ, and TID2013 databases) of the predicted quality scores versus DMOS/MOS on the test sets are shown in Fig. [10]. The tight clustering property and the monotonic relationship compared to DMOS/MOS demonstrate the impressive consistency between the predicted quality scores by the proposed BIQA approach and the human opinions of quality.

We report the standard deviations of SRCC values on the entire dataset over the 100 trials and the
Fig. 10: Scatter plots of the predicted results $Q$ vs. $DMOS/MOS$ for each of the entire LIVE, LIVEMD, CSIQ, and TID2013 databases (from left to right).

TABLE V: Standard Deviations of The SRCC Values for The Entire Database Over 100 Trials.

<table>
<thead>
<tr>
<th>Method</th>
<th>LIVE</th>
<th>CSIQ</th>
<th>LIVEMD</th>
<th>TID2013</th>
<th>Avg.</th>
</tr>
</thead>
<tbody>
<tr>
<td>BRISQUE-L</td>
<td>0.012</td>
<td>0.076</td>
<td>0.041</td>
<td>0.053</td>
<td>0.046</td>
</tr>
<tr>
<td>BLIINDS-II</td>
<td>0.022</td>
<td>0.030</td>
<td>0.060</td>
<td>0.084</td>
<td>0.060</td>
</tr>
<tr>
<td>$f^{SRN}$SVR</td>
<td>0.019</td>
<td>0.031</td>
<td>0.064</td>
<td>0.049</td>
<td>0.041</td>
</tr>
<tr>
<td>$f^{all}$SVR</td>
<td>0.014</td>
<td>0.041</td>
<td>0.043</td>
<td>0.064</td>
<td>0.048</td>
</tr>
<tr>
<td>CORNIA</td>
<td>0.012</td>
<td>0.026</td>
<td>0.027</td>
<td>0.063</td>
<td>0.042</td>
</tr>
<tr>
<td>Proposed</td>
<td>0.029</td>
<td>0.045</td>
<td>0.046</td>
<td>0.072</td>
<td>0.056</td>
</tr>
</tbody>
</table>

weighted averages across the four databases in Table V. The randomness of the train-test split coupled with that of constructing PIPs leads to higher fluctuations in the performance of the proposed approach compared to the best BIQA methods. However, the standard deviation of the proposed method is still comparative with those of BLIINDS-II and $f^{all}$SVR. From Table V, Fig. 10, and Table V, we conclude that our method correlates highly with human perception of image quality and is insensitive to datasets.

C. Database Independence

Because the proposed BIQA method is learning-based, it is necessary to verify whether its performance is bound to the database on which it is trained. To show this, we trained the BIQA method on the entire LIVE database and then tested it on the TID2013, CSIQ, and LIVEMD databases. In Table VI, we report the performance of the BIQA methods across all the distortion categories and the entire database (ALL). In addition, the overall performance on the distortion subsets of which $f^{all}$ is representative was evaluated (ALLsub). Similarly, we computed the weighted average SROCCs across the three databases, and these are listed in the last two columns. In particular, the calculation of Avg. ALLsub considered all 1-11, 19, 21-24 in the TID2013, WN, JPEG, JP2k, and Blur in the CSIQ database, and GblurJPEG in...
TABLE VI: Performance of BIQA Methods When Trained on The LIVE database But Test on The TID2013, CSIQ, and LIVEMD Databases.

| Method          | TID2013 #1 | TID2013 #2 | TID2013 #3 | TID2013 #4 | TID2013 #5 | TID2013 #6 | TID2013 #7 | TID2013 #8 | TID2013 #9 | TID2013 #10 | TID2013 #11 | TID2013 #12 | TID2013 #13 | TID2013 #14 | TID2013 #15 | TID2013 #16 | TID2013 #17 | TID2013 #18 | TID2013 #19 | TID2013 #20 |
|-----------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| BRISQUE-L       | 0.889      | 0.784      | 0.380      | 0.674      | 0.888      | 0.686      | 0.757      | 0.769      | 0.559      | 0.842      | 0.855      | 0.045      | 0.390      | 0.183      | 0.215      | 0.097      | 0.189      | 0.183      | 0.781      | 0.194      |
| BLINDS-II       | 0.788      | 0.562      | 0.369      | 0.668      | 0.802      | 0.632      | 0.518      | 0.828      | 0.703      | 0.645      | 0.721      | 0.112      | 0.305      | 0.119      | 0.251      | 0.083      | 0.052      | 0.296      | 0.731      | 0.095      |
| \( f^SRN \) SVR | 0.468      | 0.303      | 0.244      | 0.546      | 0.694      | 0.465      | 0.401      | 0.613      | 0.560      | 0.383      | 0.848      | 0.169      | 0.190      | 0.131      | 0.054      | 0.131      | 0.393      | 0.162      | 0.406      | 0.158      |
| \( f^{all} \) SVR | 0.511      | 0.293      | 0.257      | 0.517      | 0.694      | 0.281      | 0.496      | 0.675      | 0.602      | 0.705      | 0.753      | 0.030      | 0.027      | 0.050      | 0.170      | 0.397      | 0.427      | 0.271      | 0.344      | 0.225      |
| CORNIA          | 0.761      | 0.679      | 0.615      | 0.686      | 0.828      | 0.741      | 0.399      | 0.915      | 0.834      | 0.885      | 0.899      | 0.622      | 0.655      | 0.371      | 0.168      | 0.123      | 0.173      | 0.071      | 0.659      | 0.483      |
| Proposed        | 0.764      | 0.727      | 0.505      | 0.664      | 0.736      | 0.732      | 0.768      | 0.818      | 0.742      | 0.873      | 0.908      | 0.105      | 0.408      | 0.082      | 0.358      | 0.208      | 0.099      | 0.332      | 0.657      | 0.096      |

It is notable that the proposed BIQA method obtains the best performance on the TID2013, both on the subsets and the entire database. On the CSIQ database, its performance is acceptable and is comparable to CORNIA and \( f^{all} \) SVR. On the LIVEMD database, all the BIQA methods that employ NSS features fail to precisely predict the quality scores for the images corrupted by GblurWN. This is possibly because Gblur and WN have contrary effects on the NSS features, causing the numerical relation between the features and the quality score of GblurWN images to differ significantly from those of the singly distorted images \[15\]. In contrast, Gblur and JPEG cause similar deviation trends on the features; thus the BIQA model learned from singly distorted images works well for GblurJPEG, and the proposed method outperforms most of the existing BIQA algorithms for GblurJPEG.

Statistically, the proposed method obtains the best SRCC values over all databases. The high consistency between the predicted quality scores and DMOS across various distortion categories and databases corroborate the expectation that the parameters are not over-fitted and the learned model is insensitive to different databases.
TABLE VII: Performance of The Proposed BIQA Method When PIPs Were Separately Generated from Each Distortion Subset in The LIVE Database.

<table>
<thead>
<tr>
<th></th>
<th>JP2k</th>
<th>JPEG</th>
<th>WN</th>
<th>Gblur</th>
<th>FF</th>
<th>ALL</th>
</tr>
</thead>
<tbody>
<tr>
<td>Proposed</td>
<td>0.945</td>
<td>0.936</td>
<td>0.957</td>
<td>0.949</td>
<td>0.921</td>
<td>0.875</td>
</tr>
</tbody>
</table>

D. Easy Extension of the Proposed Method

Having evaluated the effectiveness and database-independent property of our method, we now demonstrate that the proposed BIQA framework can be easily extended to emerging distortion categories by simply adding a small number of corresponding PIPs into the training set. To show this, we conducted the following two experiments.

a) On the LIVE database: We separately generated 400 PIPs from each distortion subset in the LIVE database with \( N_g = 20 \) and \( T = 10 \), and then aggregated them into a single training set in learning. Subsequently, we applied the learned BIQA model to the test images. Similar to previous experiments, the train-test procedure was repeated 100 times and the median SRCC values for each distortion category and the entire database are calculated and reported in Table VII. Scatter plots of the predicted quality scores and DMOS for each distortion category are shown in Fig. 12.

From Table VII and Fig. 12 we conclude that our method has an impressive consistency with human perception across all the distortion subsets. The acceptable performance on the whole demonstrates that we can extend the proposed framework to various distortion types by simply adding the corresponding PIPs into the training set. It is encouraging that only hundreds of PIPs for each distortion category can lead to such a satisfactory performance. This will facilitate the application of BIQA methods since the generation of PIPs is very easy and efficient.

The overall performance is inferior to the performance when we used 2000 PIPs generated from the entire LIVE database without distinguishing the distortion categories, as shown in Table VII. This is mainly because the information about the relative quality between images afflicted by different types of distortion in the training set of this experiment is not considered, leading to quality scale mismatch errors between different distortion subsets, i.e. the predicted scores for WN diverge from the other distortion categories, as shown in Fig. 11. This result suggests that, to extend the proposed BIQA method to new distortion types, it is necessary to add both intra-distortion and inter-distortion PIPs.

b) On Hybrid Databases: We also conducted experiments on the LIVE, TID2013, and CSIQ databases, which differ greatly in image content, distortion categories and quality scales. In each train-
test procedure, we constructed 1000 PIPs from each database and then aggregated them into a single training set with no post-processing. The parameters $N_g$ and $T$ used to generate PIPs from each database are the same as shown in Table III. Similar to previous experiments, we only considered the distortion categories in which the features were representative.

For comparison purposes, we linearly mapped the TID2013 MOS scores and the CSIQ database DMOS scores to the range of LIVE database DMOS scores. State-of-the-art BIQA models were then learned on the training images, based on the realigned quality scores, and applied to the test images. We randomly extracted 3000 PIPs from all the training images based on the realigned quality scores with $T = 10$, and trained the proposed approach on them. Similarly, the train-test procedure was repeated 100 times and the median SRCC values are reported in Table VIII. The proposed approach trained in the first case is referred to as $Proposed_1$ in Table VIII and the approach in the second case is referred to as $Proposed_2$.

For comparison purposes, we linearly mapped the TID2013 MOS scores and the CSIQ database DMOS scores to the range of the LIVE database DMOS scores, respectively. State-of-the-art BIQA models were then learned on the training images based on the realigned quality scores and then applied to the test images. In addition, we randomly extracted 3000 PIPs from all the training images based on the realigned quality scores with $T = 10$, and trained the proposed approach on them. Similarly, the train-test procedure was repeated 100 times and the median SRCC values were reported in Table VIII. The proposed approach trained in the first case is referred to as $Proposed_1$, and that in the second case is referred to as $Proposed_2$ in Table VIII.
TABLE VIII: Performance of BIQA Methods When The Training Data Were Collected From Hybrid Databases

<table>
<thead>
<tr>
<th></th>
<th>#1</th>
<th>#2</th>
<th>#3</th>
<th>#4</th>
<th>#5</th>
<th>#6</th>
<th>#7</th>
<th>#8</th>
<th>#9</th>
<th>#10</th>
<th>#11</th>
<th>#19</th>
<th>#21</th>
<th>#22</th>
<th>#23</th>
</tr>
</thead>
<tbody>
<tr>
<td>BRISQUE-L</td>
<td>0.737</td>
<td>0.604</td>
<td>0.804</td>
<td>0.429</td>
<td>0.770</td>
<td>0.719</td>
<td>0.742</td>
<td>0.815</td>
<td>0.702</td>
<td>0.845</td>
<td>0.870</td>
<td>0.720</td>
<td>0.822</td>
<td>0.797</td>
<td>0.734</td>
</tr>
<tr>
<td>BLIINDS-II</td>
<td>0.544</td>
<td>0.320</td>
<td>0.607</td>
<td>0.425</td>
<td>0.655</td>
<td>0.590</td>
<td>0.528</td>
<td>0.780</td>
<td>0.666</td>
<td>0.779</td>
<td>0.804</td>
<td>0.536</td>
<td>0.594</td>
<td>0.701</td>
<td>0.539</td>
</tr>
<tr>
<td>f^{SRN}SVR</td>
<td>0.558</td>
<td>0.423</td>
<td>0.678</td>
<td>0.309</td>
<td>0.763</td>
<td>0.686</td>
<td>0.708</td>
<td>0.875</td>
<td>0.786</td>
<td>0.776</td>
<td>0.847</td>
<td>0.570</td>
<td>0.448</td>
<td>0.777</td>
<td>0.770</td>
</tr>
<tr>
<td>f^{all}SVR</td>
<td>0.612</td>
<td>0.449</td>
<td>0.718</td>
<td>0.326</td>
<td>0.750</td>
<td>0.697</td>
<td>0.689</td>
<td>0.865</td>
<td>0.739</td>
<td>0.818</td>
<td>0.871</td>
<td>0.577</td>
<td>0.730</td>
<td>0.806</td>
<td>0.721</td>
</tr>
<tr>
<td>CORNIA</td>
<td>0.726</td>
<td>0.582</td>
<td>0.720</td>
<td>0.553</td>
<td>0.803</td>
<td>0.644</td>
<td>0.745</td>
<td>0.877</td>
<td>0.812</td>
<td>0.859</td>
<td>0.856</td>
<td>0.641</td>
<td>0.789</td>
<td>0.730</td>
<td>0.706</td>
</tr>
<tr>
<td>Proposed_{1}</td>
<td>0.743</td>
<td>0.639</td>
<td>0.688</td>
<td>0.388</td>
<td>0.837</td>
<td>0.790</td>
<td>0.740</td>
<td>0.896</td>
<td>0.786</td>
<td>0.882</td>
<td>0.926</td>
<td>0.639</td>
<td>0.570</td>
<td>0.845</td>
<td>0.718</td>
</tr>
<tr>
<td>Proposed_{2}</td>
<td>0.734</td>
<td>0.609</td>
<td>0.688</td>
<td>0.388</td>
<td>0.837</td>
<td>0.790</td>
<td>0.740</td>
<td>0.896</td>
<td>0.786</td>
<td>0.882</td>
<td>0.926</td>
<td>0.639</td>
<td>0.570</td>
<td>0.845</td>
<td>0.718</td>
</tr>
</tbody>
</table>

As can be seen, both Proposed_{1} and Proposed_{2} are highly comparable to state-of-the-art BIQA methods across all three databases. In particular, they outperform the other algorithms for most distortion categories in the TID2013, i.e. #1, 2, 5, 6, 8-10, 22, and 24. On the CSIQ and LIVE database, they perform better than BRISQUE-L and BLIINDS but are inferior to f^{SRN}SVR, f^{all}SVR, and CORNIA, indicating that there is still scope to improve performance. It is important to note that better performance can be obtained by adding more PIPs to the training data.

Proposed_{2} yields better performance than Proposed_{1}, because the information about the relative quality between images from different databases is not included in the training set of Proposed_{1}. This is similar to the previous experiment on the LIVE database; thus, to apply the proposed BIQA framework to emerging distortion categories or to extend existing PIP databases with new image pairs, it would better to add PIPs composed of both new and existing images besides those composed purely of new images.

Although f^{all}SVR and CORNIA obtain better performance by using the realigned quality scores, it should be noted that the realignment is not precise and relies heavily on the hypothesis that the images included in each database cover the full range of quality with small intervals. Thus for the extension of...
these BIQA approaches, it is necessary to collect sufficient images associated with emerging kinds of distortion at diverse levels of degradation and to recruit many observers to evaluate their quality scores. As has been discussed, the collection of such images and the acquisition of the quality scores is difficult and inconvenient, leading to a huge cost to extend these algorithms.

In contrast, we can generate PIPs with an arbitrary number of images afflicted by the new distortion category at very low cost, and then simply add them to the existing PIP dataset to learn a robust BIQA model. Moreover, only a few subjects are needed to label the relative quality of each PIP; thus it is easy and efficient to extend the proposed framework, and our method can predict perceptual quality scores in high consistency with human perception, as verified in both Tables VII and VIII.

VI. SUBJECTIVE STUDY PIPS

In this section, we present an extensive subjective study of PIPs through a paired-comparison experiment. We adopted the 808 images (29 reference images and 779 distorted ones) contained in the LIVE database to construct image pairs. In total, there are 326,028 (= 808 × 807/2) possible pairs, from which we randomly selected approximately 240,000 pairs for evaluation in accordance with [20]. We conducted such a large number of comparisons with the objective of statistically investigating the relationship between subjective quality scores and preference labels, and validating the efficiency and efficacy of our recommended PIP-generation methods in Section III.

We randomly divided all the pairs into 10 portions and appointed one subject for each portion. The subjects for the study are students between 20 and 30 years old at Xidian University. Most image pairs were evaluated by only one subject each. In addition, 2000 image pairs were evaluated by all the subjects. The average preference labels for the 2000 pairs labeled by all the subjects is regarded as the benchmark for evaluating the validity of the raw labels assigned by each subject.

In the experiment, subjects were shown the prepared image pairs in random order and could divide image pairs into an arbitrary number of sessions. The maximum time of each session was limited to 30 minutes to minimize the effect of observer fatigue [20]. The experimental setup we used was a double-stimulus methodology in which both of the images in a certain pair were simultaneously shown on a monitor displaying at resolution of 1680 × 1080 pixels. Each image was displayed at its original resolution. The test framework was developed in MATLAB. The procedures of the experiments and the processing of the preference labels are the same as presented in Section III.A.

The evaluation of an image pair in our experiment cost 3 seconds on average, demonstrating the efficiency of generating PIPs. Analysis of the obtained data is detailed in the following.
Fig. 12: The relation between the difference DMOS and the preference label.

### TABLE IX: Accuracies of The Responses from Each Subject and Preference Labels Derived from DMOS. (%)

<table>
<thead>
<tr>
<th>intervals of difference DMOS values</th>
<th>Subject</th>
<th>DMOS</th>
</tr>
</thead>
<tbody>
<tr>
<td>(0,10)</td>
<td>91.4</td>
<td>90.5</td>
</tr>
<tr>
<td>(10,20)</td>
<td>95.4</td>
<td>97.2</td>
</tr>
<tr>
<td>(20,30)</td>
<td>99.3</td>
<td>100</td>
</tr>
<tr>
<td>(30,40)</td>
<td>99.9</td>
<td>100</td>
</tr>
<tr>
<td>(40,50)</td>
<td>100</td>
<td>100</td>
</tr>
<tr>
<td>(50,120)</td>
<td>100</td>
<td>100</td>
</tr>
</tbody>
</table>

### A. Relationship between Subjective Preference Labels and Quality Scores

To simplify our representation, we refer to the difference between the DMOS values associated with an image pair as *difference DMOS* in the remainder of this paper. To visualize the relationship between the subjective preference labels and the quality scores, we plot the distributions of the preference labels versus the difference DMOS values for the 2000 pairs labeled by all the subjects (all) and for the other pairs labeled by one subject (one), respectively, in Fig. 12. The x-coordinate shows the interval of the difference DMOS values, and the y-coordinate is the percentage of corresponding pairs assigned as each kind of preference label.

In Table IX we report the median accuracy of the responses by each subject and the accuracy of the preference labels derived from DMOS for the 2,000 pairs. In Table IX we tabulate the accuracies without considering the pairs associated with an average preference label of $y = 0$. As is clear, when the difference DMOS is small, it cannot precisely reflect the relative quality of the corresponding images, but if the difference DMOS is large enough, e.g. greater than 10, it is generally consistent with the human judgment of relative quality. This agrees with the impact of the threshold $T$ on the performance of the proposed BIQA method, as discussed in Section V.A.

It is remarkable that the judgments made by a single subject yield a highly comparable level of accuracy, given that the DMOS are obtained at much higher cost. In addition, the distributions of the singly assigned...
TABLE X: The Accuracies Without Considering the Pairs Associated With an Average Preference Label of $y = 0$. (%)

<table>
<thead>
<tr>
<th>intervals of difference DMOS values</th>
<th>[0,10)</th>
<th>[10,20)</th>
<th>[20,30)</th>
<th>[30,40)</th>
<th>[40,50)</th>
<th>[50,120)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Subject</td>
<td>92.2</td>
<td>95.8</td>
<td>99.3</td>
<td>99.9</td>
<td>100</td>
<td>100</td>
</tr>
<tr>
<td>DMOS</td>
<td>91.5</td>
<td>97.7</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
</tr>
</tbody>
</table>

labels approach those of the average responses of many observers. These results demonstrate that we can obtain valid preference labels by making only one subjective judgment for each image pair. Compared with the acquisition of DMOS, this is much easier and more efficient; we can learn an efficient BIQA model from the labeled image pairs, as shown in Section IV and the following subsection.

B. IQA Performance

In order to further test the validity of the obtained subjective preference labels, we conducted IQA experiments on the LIVE database similarly as introduced in Section V.B. Specially, we randomly selected 20 groups of images from the LIVE database and 2000 PIPs from the corresponding subjectively labeled image pairs for training. Afterward, we applied the learned model to the test images to predict the corresponding quality scores. For the purpose of comparison, we also constructed training set based on the DMOS. The median KRCC, SRCC, and PLCC values on the entire LIVE database were reported in Table XI.

In addition, we calculated the difference feature vectors of the subjectively labeled image pairs which are purely composed of the test images and then fed them into the learned classifier to estimate the corresponding preference labels. We computed four accuracies to evaluate the prediction precision. The first one is the accuracy on all the corresponding test image pairs and is denoted as $Acc^{All}$. The second one is operated only on the pairs of which the difference DMOS is consistent with the subjective judgment, and is denoted as $Acc^{cnst}$. Moreover, we estimated the corresponding preference labels by comparing the predicted quality scores, and denote the corresponding preference accuracies $Acc^{Q}_{Q}$ and $Acc^{Q}_{cnst}$.

It is obvious that the proposed BIQA method obtains almost the same performance by adopting the subjective preference labels or the labels derived from DMOS, demonstrating the validity of the preference labels obtained through paired comparisons. It is also notable that the performance here is similar to that reported in Section V.B. This corroborates the expectation that by choosing a proper threshold, we can
TABLE XI: Performance of the Proposed BIQA Method on the Entire LIVE Database.

<table>
<thead>
<tr>
<th></th>
<th>KRCC</th>
<th>SRCC</th>
<th>PLCC</th>
<th>Acc\textsuperscript{All}</th>
<th>Acc\textsuperscript{cnst}</th>
<th>Acc\textsuperscript{All}</th>
<th>Acc\textsuperscript{cnst}</th>
</tr>
</thead>
<tbody>
<tr>
<td>DMOS</td>
<td>0.783</td>
<td>0.935</td>
<td>0.938</td>
<td>0.949</td>
<td>0.957</td>
<td>0.956</td>
<td>0.963</td>
</tr>
<tr>
<td>Subject</td>
<td>0.783</td>
<td>0.935</td>
<td>0.937</td>
<td>0.948</td>
<td>0.956</td>
<td>0.956</td>
<td>0.963</td>
</tr>
</tbody>
</table>

generate valid PIPs from existing IQA databases for learning a robust BIQA model. In addition, $Acc\textsuperscript{All}_Q$ and $Acc\textsuperscript{cnst}_Q$ are slightly greater than $Acc\textsuperscript{All}$ and $Acc\textsuperscript{cnst}$. The reason is that more comparisons are performed on each test image in the quality prediction procedure, indicating the efficacy of the proposed quality prediction approach.

VII. CONCLUSION

In this paper, we presented a BIQA framework that learns to predict perceptual image quality scores from preference image pairs. Thorough experiments on the four largest standard databases demonstrate that our method correlates highly with human perceptions of quality and that the framework can be easily extended to emerging distortions. In addition, an extensive subjective study corroborates the efficiency and validity of generating PIPs through paired comparisons or from existing IQA databases. In our continued search for convenient but effective PIP-generation methods and BIQA metrics, we have deployed the simplest operations in both the subjective study and the proposed BIQA approach in this paper. There is still room for the optimization of the subjective study and the improvement of IQA performance. In addition, the low cost of the generation of PIPs, coupled with the efficacy of the proposed BIQA approach, make the deployment of this framework to solve other quality assessment (QA) problems, e.g. other types of images, videos, etc., a promising option. Further research is needed to explore quality-relevant features and generate corresponding stimulus pairs with valid preference labels.

REFERENCES


