Exploring Residual and Spatial Consistency for Object Detection

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Keywords: Visual Object Detection, Wide Baseline Matching, Local Image Feature, Spatial Consistency.

Abstract: Local image features show a high degree of repeatability, while their local appearance usually does not bring enough discriminative pattern to obtain a reliable matching. In this paper, we present a new object matching algorithm based on a novel robust estimation of residual consensus and flexible spatial consistency filter. We evaluate the similarity between different homography model via two-parameter integrated Weibull distribution and inlier probabilities estimates, which can select uncontaminated model to help eliminating outliers. Spatial consistency test was encoded by the geometric relationships of domain knowledge in two directions, which is invariant to scale, rotation, and translation especially robust to the flipped image. Experiment results on nature images with clutter background demonstrate our method effectiveness and robustness.

1 INTRODUCTION

With the increasing popularity of image applications on internet such as Google Image and Flickr, considerable attention has been directed to image detection and classification in multimedia research communities. However, accurate object recognition remains a challenging problem, because the target object may be small in size with cluttered background, or it is significantly different from the query image in color, scale, and orientation.

Among existing methods, the bag-of-features model (Sivic and Zisserman, 2003; Lazebnik et al., 2006) has been a popular technique because of its simplicity and effectiveness. This method quantizes local image descriptors into distinct visual words for scalable image indexing and searching. With the inverted index of visual words, one not only avoids storing and comparing high dimensional local descriptors, but also reduces the number of candidate images because only images sharing common words with the query image are considered. To improve the precision of the matching, a visual word may be augmented with compact information from its original local descriptor, including a Hamming code (Jegou et al., 2008), descriptor scale and angle, and the distance (in descriptor space) to its neighboring visual words.

Geometric verification such as RANSAC is a crucial step after retrieval from the inverted index. Many extensions of RANSAC have been proposed in order to improve the accuracy of the matching. MultiGS (Chin et al., 2010) accelerates hypothesis sampling by guiding it with information derived from residual sorting. The LO-RANSAC (Chum et al., 2003) method introduces an inner RANSAC loop into the main RANSAC algorithm such that hypotheses may be generated from the set of inliers. Guided-MLESAC (Tordoff and Murray, 2005) and PROSAC (Chum and Matas, 2005) focus on sampling more confident keypoint matches. The above two methods are essentially guided only by the prior inlier probabilities and do not conduct conditional sampling to further improve efficiency. GroupSAC (Ni et al., 2009) focuses on sampling groups of data obtained using image segmentation. SCRAMSAC (Sattler et al., 2009) introduces a spatial filtering step such that matches with similar local geometry are considered. ARRSAC (Raguram et al., 2008) performs a partially breadth-first verification such that the number of hypotheses may be modified according to the inlier ratio estimation while still bounding the runtime.

In recent years, spatial context have been proven to be useful for enhancing the discriminative power of individual local features. Features that are close to each other are grouped to form a visual phrase. Visual phrase is formed by two different methods. The first type of methods mainly relies on object segmentation or region detection, which finds spatial context of object locating area by detecting contour (Russell et al., 2006; Wu et al., 2009). An example of such meth-
ods is bundled-feature which groups features in local MSER regions described by SIFT into a local group to increase the discriminative power of local features (Zhang et al., 2011). However, object need to be segmented accurately. Otherwise the visual phrase will be far away from the correct area. The second type of methods selects the visual phrase at a fixed area, such as geometry-preserving visual phrases that capture long-range spatial layouts of the words (Sivic and Zisserman, 2009). They were generating a higher-level lexicon, i.e. visual phrase lexicon, where a visual phrase is a meaningful spatially co-occurrent pattern of visual words (Jiang et al., 2011; Li et al., 2011). In (Lowe, 2004), local spatial consistency from some spatial nearest neighbors is used to filter false visual-word matches.

In this work, we propose a novel robust estimation of residual consensus and flexible spatial consistency filter for object matching. Specifically, we evaluate the similar structure from different homography models in two aspects. Firstly, We extend RANSAC to model residuals distribution using two-parameter integrated Weibull which is between a power-law and a Gaussian distribution. Secondly, we calculate the normalized overlap of sorted residuals obtained from hypothesis generation. For spatial consistency filter, previous literatures usually describe the relative spatial positions between each feature pair along the horizontal (X-axis) and vertical (Y-axis) directions or measuring angles only in one direction. However, we measure spatial consistency using geometric relationships of globally point distribution in both clockwise and counterclockwise, which is invariant to scale, rotation, and translation especially robust to flipped image.

2 RESIDUAL AND SPATIAL CONSISTENCY

The SIFT feature is one of the most robust and distinctive features (Kalantidis et al., 2011). SIFT feature descriptor is invariant to uniform scaling orientation, and partially invariant to affine distortion and illumination changes. Local interest points are extracted by DOG detector and described by 128-dimensional SIFT descriptor. A bag of visual words (BOW) is obtained by quantizing high-dimensional local image descriptors through clustering. An inverted index of domain knowledge in both clockwise and counterclockwise, which is measured by relative orientation and distance order. The use of spatial configuration of local features aims at reducing the number of mismatches in the correspondence set.

2.1 Residual Consensus

Residual ordering distribution is useful for distinguishing between contaminated models and uncontaminated models. Naturally, residual errors obtained from uncontaminated models are approximate similar with each other. Thus, potential patterns can be discovered by evaluating the similarity between these structures to help finding the optimal models.

Formally, given a set of N input data. Under the hypothesize-and-verify framework, a model is fitted by a minimal sample. Let \( R = \{r_1, r_2, ..., r_N\} \) be absolute residuals corresponding to the input data. We then find the permutation \( S = \{s_1, s_2, ..., s_N\} \) such that the residuals are sorted in increasing order. We define the intersection set \( \Theta_m^j \) as:

\[
\Theta_m^j = S_{1:n} \cap S_{1:n}^j,
\]

where \( S_{1:n} \) denote the subset containing the first \( n \) points of \( S \) and \( S_{1:n}^j \) finds the number of overlap elements from \( S_{1:n} \) and \( S_{1:n}^j \). We also give the normalized overlap \( \Theta_n^j \):

\[
\Theta_n^j = \frac{1}{n} |\Theta_m^j|,
\]

Intuitively, Similar models should have the most overlap of small value residuals. To verify this method for distinguish uncontaminated and contaminated models, an example is shown to illustrates this intuition explained. In details, we generate 400 homography hypotheses on the real images and select
the uncontaminated models to compare with the random models. Fig.1(a) plots the normalized overlap $\Theta_u^{i,j}$ of sorted residuals from uncontaminated models and random models, respectively. Fig.1(b) shows a block diagonal pattern of $\Theta_u^{i,j}(1 \leq i, j \leq 400)$, in which, the points are scattered according to their structure membership. The results show that, the values for an inlier concentrate mostly on other inliers from the same distribution, while for an outlier the value are generally low and appear to be randomly distributed.

We use a statistical model as another means to characterize the qualitative differences between models. By observing the data, the distribution of residuals is usually between a power-law and a Gaussian distribution, can be well modeled by a two-parameter integrated Weibull distribution,

$$p(x) = \frac{\gamma}{2\beta^2 \Gamma\left(\frac{1}{\beta}\right)} \exp\left(-\frac{1}{2\beta^2} x^\gamma\right), \quad (3)$$

where $x$ is the residuals of geometric model. $\gamma > 0$ denotes the peakness of the distribution. $\beta > 0$ represents the scale parameter of the distribution. $\Gamma(x)$ is the complete Gamma function.

$$\Gamma(x) = \int_0^\infty t^{x-1} \exp(-t)dt, \quad (4)$$

Given the observed data $X = x_1, x_2, \ldots, x_n$. The best fit is obtained when model parameters maximize the log-likelihood function, in which case their respective derivatives should equal zero.

$$\frac{\partial}{\partial \beta} \ln L_w(\beta, \gamma|X) = -\frac{1}{\beta} + \frac{1}{\beta} \sum_{i=1}^n \frac{|x_i|^\gamma}{\beta} = 0, \quad (5)$$

$$\frac{\partial}{\partial \gamma} \ln L_w(\beta, \gamma|X) = \frac{1}{\beta^2} (\gamma - 1 + \Psi\left(\frac{1}{\gamma}\right)) + \ln(\gamma) + \sum_{i=1}^n \frac{|x_i|^\gamma}{\beta}, \quad (6)$$

where $\ln L_w(\beta, \gamma|X)$ is the residuals of geometric model. $\Psi(\gamma) = \frac{d}{d\gamma} \ln \Gamma(\gamma) = \frac{d}{d\gamma} \frac{\Gamma(\gamma)}{\Gamma(\gamma)}$, \quad (7)\]

The parameter is obtained by eliminating from Eq.(6):

$$f(\gamma, X) = -\frac{1}{\gamma} \sum_{i=1}^n \frac{|x_i|^\gamma}{\Gamma\left(\frac{1}{\gamma}\right)} \ln \left(\frac{|x_i|^\gamma}{\Gamma\left(\frac{1}{\gamma}\right)}\right) + 1 + \frac{1}{\gamma} \Psi(\frac{1}{\gamma}) = 0. \quad (8)$$

Eq.(6) is solved using standard iterative procedures, for example, the Newton-Raphson. Fig.2(a) displays the fitting result using an integrated Weibull distribution. The residual is subtracted by median because we want the peak of probability density is located on where residual equal zero. As in Fig.2(b), we can see that the two parameters of models build up a 2-D map and the most uncontaminated models come from the intensive area. Therefore, we can use this property to capture uncontaminated models. Given discrete data points from 2-D map, the center of the intensive area is located using mean-shift method (Comaniciu and Meer, 2002). Here, the mean-shift path toward the mode follows a smooth trajectory, and the angle between two consecutive vectors being always less than 90 degrees. The search procedure can be interpreted as kernel density estimation for the position of the model distribution points. Moreover, we define an elliptical area lying on the center of the 2-D map, such that, the major axis $l_1$ parallel to scale axis and minor axis $l_2$ parallel to shape axis. In this case, the models within the elliptical area are kept for spatial consistency test and unpromising hypotheses can be quickly filtered out.

$$\text{(a)}\quad \text{(b)}$$

**Figure 1:** (a) Normalized overlap vs. subset size $n$ for uncontaminated models and random models. (b) The corresponding matrix of size 400x400 with subset is set to 80.

**Figure 2:** (a) Residual distribution is fitted by an integrated Weibull model. (b) The parameter map of uncontaminated models and contaminated models.

### 2.2 Spatial Consistency Analysis

Given an image $I$, we extract a set of visual words $U(I) = \{u_i\}$ with center coordinates $(x_i,y_i)$ and $U(I) = \{v_j\}$ is another visual word. In an image pair
(I_1, I_2), all the point correspondences are established by visual words. Thus, we obtain a correspondence set C:

\[ C = \{(u_i, v_j) | u_i \in U \land v_j \in V\}. \]  

(9)

We define a matching score \( M(I_1, I_2) \), which consists of a membership term \( M_m(I_1, I_2) \) and a geometric \( M_g(I_1, I_2) \):

\[ M(U, V) = M_m(U, V) + \lambda M_g(U, V), \]  

(10)

where \( \lambda \) is a weighting parameter. For membership term, we compute inlier scores to define the membership term \( M_m(U, V) \).

Our geometric term performs a weak geometric verification between training image \( I_1 \) and query image \( I_2 \) using relative ordering, and we incorporate orientation configuration into distance configuration. In detail, the minimal value of two kinds of configuration is the final value,

\[ M_g(U, V) = \min(M_g^1(U, V), M_g^2(U, V)), \]  

(11)

where, \( M_g^1(U, V) \) is orientation configuration term and \( M_g^2(U, V) \) is distance configuration term. As in Fig.3, the centroid of all the features is located as a reference point. Starting from zero degree, the whole visual words position is scanned clockwise. As the angle increases, each visual word is assigned a number with ascending order. Note that the visual word from testing image has the same number as training one if they are the matching pair. The inconsistency is accumulated by the difference of neighbor number between training image and testing image,

\[ M_g^1(U, V) = - \sum_i \delta(O_i[u_i] > O_i[u_i+1]), \]  

(12)

where \( O \) is a pre-defined geometric sorting order, and \( \delta(O_i[u_i] > O_i[u_i+1]) \) is an indicator function that measures the consistency between the order \( i < i + 1 \) (before matching) and the order \( O_i[u_i] > O_i[u_i+1] \) (after matching). In other words, we penalize geometric inconsistency of the visual word matching between two images and put a weight to inconsistency avoid for weak discrimination by small difference. In order to match flipped image, the minimal inconsistency is set as final scores from clockwise and counterclockwise image scan,

\[ M_g(U, V) = \min(M_g^e(U, V), M_g^c(U, V)), \]  

(13)

where \( M_g^c(U, V) \) is computed by a geometric verification via clockwise, and \( M_g^e(U, V) \) by a counterclockwise geometric verification. For spatial consistency filter with distance, the only difference between orientation is that we compare the distance from reference point to the position of visual words.

Figure 3: The ordering constraint for local features. (a) Correct matches preserve the relative orientation orders; (b) Wrong matches results in inconsistent relative orientation orders; (c) Wrong matches results in inconsistent relative distance orders.

2.3 Algorithm

Consider the residual consensus aforementioned, a statistical rest that evaluates the similarity between distributions can give the optimal homography model. Specifically, we first measure the fraction of models by Weibull distribution intensive area which is located using mean-shift method. Then we restrict all further processing to correspondences where this fraction surpasses a threshold \( T \) by inlier probabilities estimates, resulting in a reduced set of models. Our approach combines the robust estimation of residual consensus and spatial consistency filter in a unified framework. The details of algorithm are shown in Algorithm 1.

Generally, image retrieval system uses criteria whether the absolute matching number exceeds a certain threshold to decide the final matching results. However, taking into account the different complexity of the images and different number of features, which lead to more correspondence generating in images with large number features. In that case, even if the query image does not have the object, the features of the object still have high probabilities to match the other parts of the image. In our approach, we use the total number of features as penalty factor to eliminate this problem. Note that the importance of inliers is different because of residuals of geometric model have diverse values. The residual with small values is more important for correspondence selection, because they have high probabilities to become true inliers. We follow MLESAC(Maximum Likelihood SAC) (Torr and Zisserman, 2000) to define the
loss function. It models inliers error as unbiased Gaussian distribution and outlier error as uniform distribution.

3 EXPERIMENTAL RESULTS

In this section, experiments were conducted to demonstrate the effectiveness of the proposed residual and spatial consistency method on real image datasets. Yannis et al. [16] built the FlickrLogos dataset by downloading real-world images from Flickr containing 27 logos covering various aspects of life. The 27 logo classes is Adidas, Apple, BMW, Citroen, Coca-Cola, DHL, Fedex, Ferrari, Ford, Google, Heineken, HP, Intel, McDonalds, Mini, Nbc, Nike, Pepsi, Porsche, Puma, RedBull, Sprite, Starbucks, Texaco, Unicef, Vodafone, and Yahoo. The dataset selected 40 images per class from Flickr, and every selected image include at least one instance of the brands logo. All 1080 images were annotated with bounding boxes. The annotated logo was partitioned into 2 subsets: The training set has 30 randomly selected images from the 40 images per brand, and the rest were the test set. To further verify the scalability and effectiveness of our approach, we add 500 crawled Flickr images with noting logos and 230 with logos to the test set.

Baseline. We use Trademark Matching (Bagdanov et al., 2007) approach with RANSAC as the baseline approach, which is denoted as “RANSAC”. The method is a compact representation of trademarks and video frame content based on SIFT feature points, which is state-of-the-art logo detection methods, and has been used in commercial application. In the paper, a visual vocabulary of 2M visual words is adopted. In fact, we have experimented with different visual codebook sizes, and have found the 2M vocabularies yield the best overall performance for the baseline.

Comparisons. Another two compared algorithms are improved versions of baseline by adding different geometry consistency. The first one is Hamming Embedding [3] by adding a hamming code to filter out matched features that have the same number of quantized visual words but have a large hamming distance from the query feature. We denote this method as “HE.” The second one is accelerate hypothesis sampling by guiding it with information derived from residual sorting [6], which is denoted as “Multiasc”.

In this paper, we set a fraction of inliers $T$ to 0.95 in Residual consensus evaluation. For pairs of uncontaminated models, we found that when subset $n$ close to the true inlier ratio, $\theta_{ni}^*$ is approaching $T$. For parameter of elliptical area, we set the value of $l_1$ and $l_2$ to 40 and 0.25. We look for $K$ good models to provide robustness structure. In our experiments, setting $K = 3$ was sufficient in practice. We empirically examine the ratio of optimal RANSAC score and geometric inconsistency. The parameter is set in the range of $\lambda = [0, 0.5, ..., 2.5]$. For each varying of $\lambda$, we run the residual and spatial consistency approach. Table 1 shows that $\lambda = 1.0$ is the most effective value. Parameters $\gamma$ and $\sigma$ are set to 0.7 and 0.3 for Loss function.

Table 1: Comparing the performance for varying $\lambda$.

<table>
<thead>
<tr>
<th>$\lambda$</th>
<th>0</th>
<th>0.5</th>
<th>1.0</th>
<th>1.5</th>
<th>2.0</th>
<th>2.5</th>
</tr>
</thead>
<tbody>
<tr>
<td>Accuracy</td>
<td>73.41</td>
<td>75.95</td>
<td>76.42</td>
<td>74.57</td>
<td>72.49</td>
<td>70.52</td>
</tr>
</tbody>
</table>

Performance results when varying the number of training images are presented in Fig.4. To vary the number of training images, we split the training set into 5 random subsets of 5 images per class. Fig.5 shows performance comparison with Recall and Precision curves. We can see that our approach leads to a better detection performance compared with the other algorithms. Fig.6 displays the matching results between the same objects. It can be observed
that most of matched ones locate on the object. As a reminder, these curves are generated by varying the matching threshold and computing the following values:(TP:True Positive, FP:False Positive, FN:False Negative)

\[
\text{Recall}(\%) = \frac{TP}{TP + FN} \times 100, \\
\text{Precision}(\%) = \frac{TP}{TP + FP} \times 100, \tag{14}
\]

4 CONCLUSIONS

In this paper, we present a new object matching algorithm based on robust estimation of residual consensus and flexible spatial consistency filter. For residual consensus, we model residuals distribution using two-parameter integrated Weibull which is between a power-law and a Gaussian distribution. Then we estimate a series of inlier probabilities, which are updated on the fly. For spatial consistency filter, we measure spatial consistency using geometric relationships of global point distribution in both clockwise and counterclockwise, which is invariant to scale, rotation, and translation especially robust to flipped image. Experiments on clutter background images show that the proposed method leads to improvements in the performance of object detection system.

ACKNOWLEDGEMENTS

This research was supported by the High Technology Research and Development Program of China (2012AA011702, 2011AA01A107).

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