

# Skin Detection Technique Based on HSV Color Model and SLIC Segmentation Method\*

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**Abstract.** The paper is devoted to new skin detection technique based on the HSV color model and SLIC segmentation method. The algorithm of skin detection is described. Experiments results are presented. The influence of training images on the skin detection is shown. New skin detection algorithm implemented in Python language using OpenCV library is described.

**Keywords:** Skin detection · HSV · SLIC Superpixels · Computer vision · Pattern recognition

## 1 Introduction

Skin detection is the process of finding skin-colored pixels and regions in an image or a video. Skin detection applications are used for personality recognition, body-parts tracking, gesture analysis and adult content filtering and etc [1].

When the standard *RGB* color space is used, the skin detection can be very difficult under conditions of variable lighting and contrast. Therefore, the input image must be converted to another color space [2–4] that is invariant or at least insensitive to lighting changes, such as *HSV* [5].

The implemented skin detector converts the image into required color space and then uses the image histogram to mark each pixel: whether it belongs to skin. Image pixels are grouped in superpixels using the *SLIC* clustering method [6]. Thus, advantages of our skin detector are high processing speed and invariance under rotation and lighting changes.

The main steps of skin detection in the image are:

1. download the input image;
2. convert image to *HSV* color space;
3. generate the image histogram;
4. apply classifier to determine the probability of a given pixel being skin-colored;

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5. divide the image into superpixels;
6. paint out superpixels where sum of probabilities less than the limit.

As a classifier, the *Naive Bayes* algorithm was chosen. The choice is due to the facts that this algorithm is the most accurate [7–9] and a small amount of data is required for it’s learning [10, 11].

*SLIC* algorithm was chosen for image segmentation. It had the fewest errors among those considered in [12], and also it is the fastest one [12, 13].

The novelty of this research is the unique combination of technologies for solving the problem of skin detection in the image.

The paper has the following structure. In Section 2, color model conversion technique is described. In Section 3, algorithm of image histograms generation is given. In Section 4, application of Naive Bayes classifier is shown. In Section 5 superpixel segmentation is conducted. In Section 6, effects of the training example on skin detection are considered and results analysis is given. In Section 7, skin detection algorithm implemented in Python language using OpenCV library is described. In Section 8, the results are summarized and directions for further research are outlined.

## 2 HSV Color Model

The first step of skin detection is to convert the input image to *HSV* color space as the most suitable for our research [14–16]. The *HSV* color model is a cylindrical representation of the standard *RGB* model [17]. *HSV* stands for *Hue*, *Saturation*, and *Value*. The *Hue* is measured in degrees and varies from 0 to 360. It forms the base color. *Saturation* and *Value* (brightness) determine the proximity to white and black respectively. In the basic model they vary from 0 to 100, but in the *OpenCV* library used in the detector, they vary from 0 to 255.

In order to convert the image from the *RGB* to *HSV*, each pixel in the image is subjected to the following transformation [5]:

- Preliminarily, the maximum and minimum of  $R, G, B$  values  $C_{max}$  and  $C_{min}$  should be found and their difference  $M$  is calculated;
- The *Hue* is calculated:

$$H = \begin{cases} 0, & C_{max} = 0, \\ 60 \times \frac{G-B}{M}, & C_{max} = R, \\ 60 \times \frac{B-R}{M} + 120, & C_{max} = G, \\ 60 \times \frac{R-G}{M} + 240, & C_{max} = B. \end{cases} \quad (1)$$

- The *Saturation* is calculated:

$$S = \begin{cases} \frac{M}{C_{max}}, & C_{max} \neq 0, \\ 0, & C_{max} = 0. \end{cases} \quad (2)$$

- The *Value* is calculated:

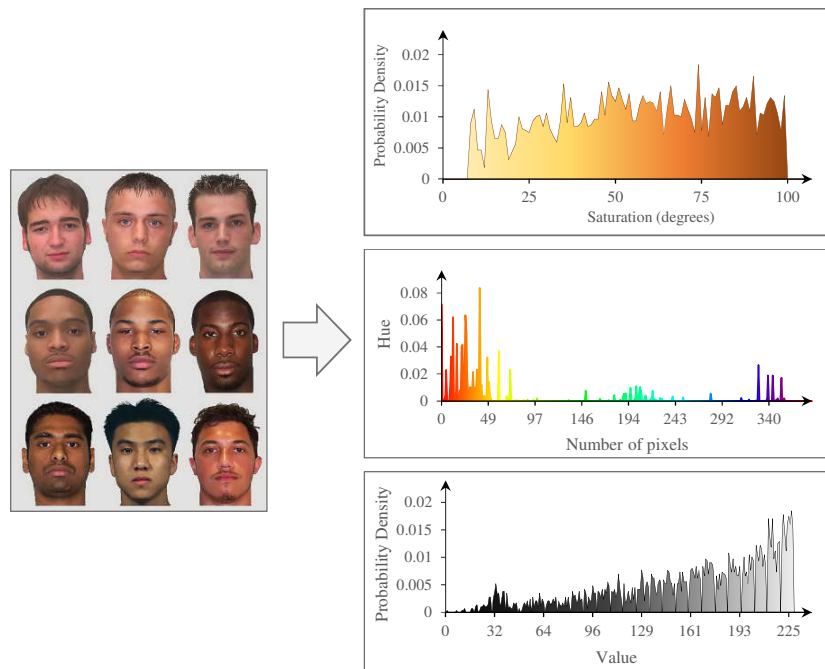
$$V = C_{min}. \quad (3)$$

Calculating the above values for each pixel, we get an image in the *HSV* color space.

### 3 Histogram

The second step is to generate the input image histogram [18]. The histogram [19] is a graph of the distribution of digital image elements with different saturation. The horizontal axis represents pixel property values, and the vertical axis represents the number of pixels. The image representation in the form of a histogram is just another way of displaying the image. It is worth of noting that the color histogram itself is not exhaustive in representing image's features but, nevertheless, it is widely used. In image processing it is used either in combination with others parameters, such as color moments [20], or in its modified version [21], or in its original form with a slightly different approach to data compilation [18].

Let's take an image of people of different nationalities and build a histogram (see Fig. 1). Later, the histogram for the Value will not be taken into account in order to remove the effects of lighting changes.



**Fig. 1.** Sample histogram

## 4 Classifier

The next step is to select a classification function [22] that will determine the probability of a given pixel being skin-colored. Such a function can be Bayesian network [23, 24], *Multilayer Perceptron* [25], *SVM* [26–28], *AdaBoost* [29, 30], *Naive Bayes*, *RBF network* [31] and etc.

In our detector the *Naive Bayes* algorithm is used. This method is based on the Bayes theorem with the assumption that the hypotheses are independent. In other words, the presence of any attribute in the class does not imply the existence of another. For example, if you consider an apple of green color, round in shape, and with a diameter of 5 sm, then all these parameters can be considered as independent, because it is possible to change one or more of them, but the object will still be an apple.

*Naive Bayes* algorithm is easy to implement. It is faster than many other algorithms [32]. Besides, despite very simplified conditions, *Naive Bayes* classifiers often work much better in many difficult situations [6, 33]. Also, the advantage of this algorithm is that it does not require a lot of training examples [34].

By the Bayes theorem, taking into account the variable  $y$  and the dependent characteristic vector  $(x_1, \dots, x_n)$ , we obtain the following relation:

$$p(y|x_1, \dots, x_n) = \frac{p(y)p(x_1, \dots, x_n|y)}{p(x_1, \dots, x_n)} \quad (4)$$

Using the assumption that the parameters are independent

$$p(x_i|x_{i+1}, \dots, y) = p(x_i|y)$$

, the expression can be simplified:

$$p(y|x_1, \dots, x_n) = \frac{p(y) \prod_{i=1}^n p(x_i|y)}{p(x_1, \dots, x_n)} \quad (5)$$

for any  $i$ .

Notice that the variable  $p(x_1, \dots, x_n)$  can be eliminated from the original equation (4), because it is a constant:

$$p(y|x_1, \dots, x_n) \propto p(y) \prod_{i=1}^n p(x_i|y) \quad (6)$$

Using the transformations described above, the classification rule can be obtained [35]:

$$Y = \operatorname{argmax}_y p(y) \prod_{i=1}^n p(x_i|y) \quad (7)$$

This rule will be applied to each pixel. Thus, we obtain the *probability map (PMap)* [36] of the image (see Fig. 2).



Fig. 2. Probability map of the image

## 5 Superpixel Segmentation

Superpixel segmentation algorithms combine the pixels into the atomic regions, which will later be checked on being skin-colored. There are various algorithms for segmentation [13, 37]. One of them is *SLIC Superpixels* algorithm. It is an adaptation of the *k-means* method for superpixels [6]. The difference is that the algorithm is more efficient because the detection area is reduced to the size of a superpixel.

The algorithm is simple for understanding and implementation [12]. Initially, only the number of regions  $k$  is specified. The image is divided into  $k$  segments of size  $S$ , and center is located at the point with the lowest gradient. Then, for each pixel, a  $2S \times 2S$ -sized area is considered and it is attached to the segment with the smallest difference in the distance  $D_s$ , calculated from the equation (8) [38]. After that, in each cluster, the center moves to the point with the lowest gradient and again all the pixels are rearranged [12, 4]. This occurs until a certain threshold of the minimum distance is reached.

$$D_s = d_{lab} + \frac{m}{s} d_{xy}, \quad (8)$$

$$d_{lab} = \sqrt{(l_k - l_i)^2 + (a_k - a_i)^2 + (b_k - b_i)^2}, \quad (9)$$

$$d_{xy} = \sqrt{(x_k - x_i)^2 + (y_k - y_i)^2}, \quad (10)$$

where  $d_{lab}$  is the distance in *CIELAB* color space between  $k$ - and  $i$ -pixels (color distance);  $d_{xy}$  is the distance between coordinates  $k$ - and  $i$ -pixels (spatial distance);  $m$  is the parameter that affects the size of a superpixel. The more  $m$ , the more the spatial proximity affects the overall distance.

After all transformations of the image used in the previous section we get Figure 3 (for  $m = 10$ ).

The dependence of the segmentation on the coefficient  $m$  is shown on Figure 4. It is clear, that the greater the  $m$ , the smaller the segments. The optimal is  $m = 10$ .

During the segmentation, unnecessary regions was removed and possible image noises were excluded. Thus, skin will be selected entirely, and the other pixels, which falsely have a high probability, will not be colored out. For comparison, in Figure 5 there are two methods results. It is noticeable that the left image is less informative. Segments were selected by threshold function, i.e. if the sum of the probabilities of the whole segment was greater than a certain



**Fig. 3.** SLIC superpixels segmentation



**Fig. 4.** Dependence of image partitioning on the coefficient  $m$  (right image  $m = 5$ , left image  $m = 20$ )

value, then the pixel was not colored out and recognized as the skin. The right image shows that not only the skin was colored out. This is due to the people of the training example are with hair, so the hair in Figure 5 was also recognized, as well as parts of the clothing, because its color is similar to the dark skin color (see Fig. 5).



**Fig. 5.** Methods comparison

## 6 The Effect of the Training Example on the Result

In this section, let us consider the dependence of the detection result on a variety of skin colors on the training example. Let us take the original image with people of different races (see Fig. 6) and test different training examples on it.



Fig. 6. Input image

Comparative figures are presented below (see Fig. 7, Fig. 8, Fig. 9, Fig. 10).



Fig. 7. 1st experiment

Experiments show that the image on which the histogram is generated significantly affects the skin detection. For the training images of people of different races should be chosen. The first and second tests are the examples of incorrectly selected training images. The most accurate result is obtained in the fourth test.

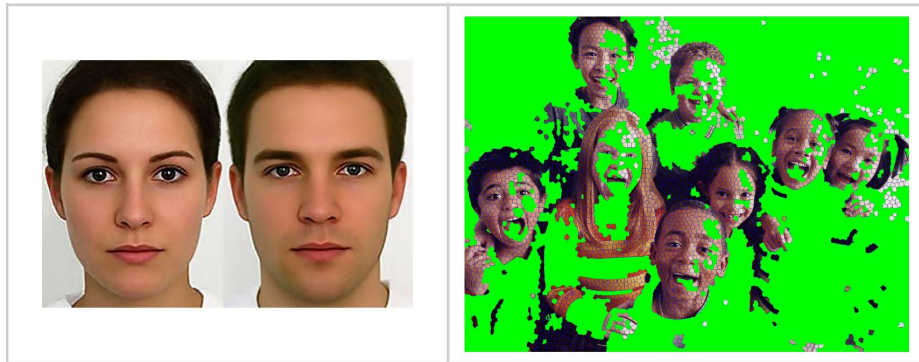


Fig. 8. 2nd experiment

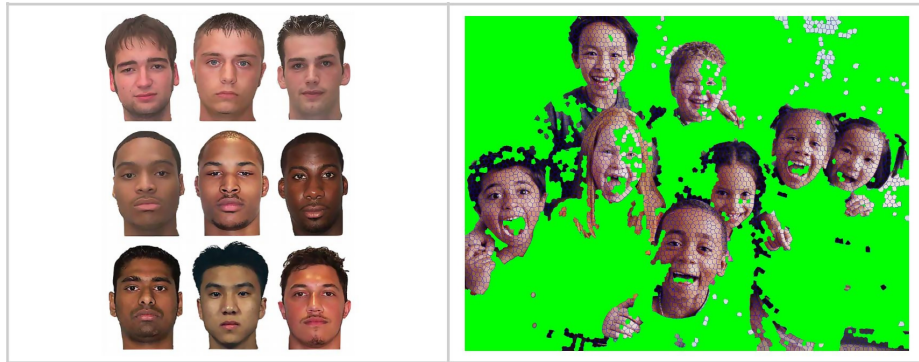


Fig. 9. 3rd experiment

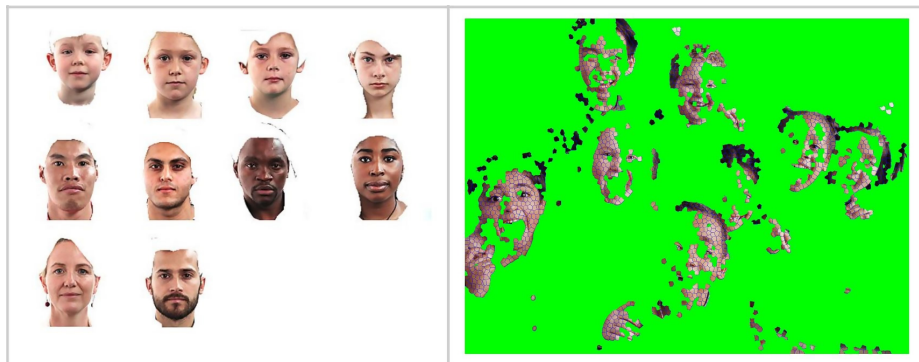


Fig. 10. 4th experiment



However, there are also false positives, for example, then the clothes color is similar to the skin color.

## 7 Skin Detector Implementation

The program is implemented in Python (v3.6.1) programming language. This language is chosen because it is commonly used for solving the pattern recognition problem, a large number of libraries are freely available - it greatly simplifies the development. The following libraries are used in the program:

- OpenCV (v3.2.0) is applied for the basic calculations;
- Numpy (v1.12.1) accelerates and simplifies operations with arrays [39], which OpenCV operates with [40];
- Matplotlib (v2.0.2) is used for plotting Numpy arrays as images [41].

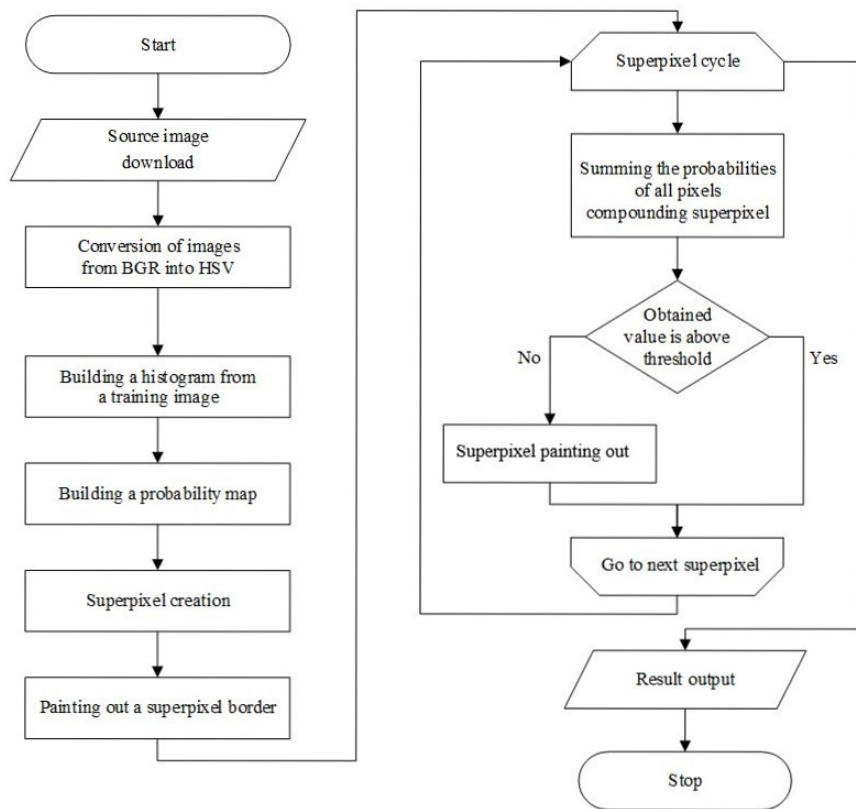
To download input image the function `cv2.imread` used. It is worth noting that the downloaded image is in *BGR* color space [42] by default, so it should be converted from the *BGR* color space to *HSV* (see Fig. 11). To do this, we used the function `cv2.cvtColor`. Next, a mask of the training image is generated using the function `cv2.inRange`, which sets the lower and upper bounds of colors involved in generating the histogram. This step is needed to remove the background from the image. Then, the training image histogram is generated in two channels: *Hue* and *Saturation*, via the function `cv2.calcHist`. The next step is the probability map calculation. This operation is performed by the function `cv2.calcBackProject`, to which the input image and the histogram are given as a parameters. It is indicated, through which channels of the color space the back projection should be made. Back projection is the only way to determine how much are the pixels correspond to the histogram [43]. At the end of the preparatory stage, the function `cv2.ximgproc.createSuperpixelSLIC` initialize the object `SSLIC`, and the `SSLIC.iterate` method refine the boundaries.

After the above steps, we got the *PMap* and segmented image. Using them, we check the sum in each superpixel and color it out, if it does not exceed the threshold, which depends on image sizes and the number of superpixels.

The source code of the program is freely available on GitHub at <https://github.com/AntonSinkov/skin-recognition>.

## 8 Conclusion

During the research, the skin detection algorithm was developed. The influence of training images on the result is shown. The more people of different races (in approximately equal proportions), the more accurate the results of skin detection. The usage of segmentation allowed selecting skin segments, without dividing into individual pixels, which improves perception.



**Fig. 11.** Activity diagram of the developed algorithm

However, this algorithm is not absolutely accurate. Segments with a color similar to the skin color will be recognized as the skin. Nevertheless, this algorithm is suitable for preprocessing, because it is fast and accurate enough if the correct training example is provided.

In the future, it is planned to use artificial neural networks in the algorithm to speed up the information processing.

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