NEWS BROADCAST ANALYSIS

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1 Shot detection

A score is calculated for each frame using either the sum of absolute differences method or the histogram differences method. Given the mean μ and standard deviation σ of all the scores, a threshold is chosen as

threshold =
$$\mu + k \times \sigma$$

where k is an integer. A shot is a series of frames taken by continuously by one camera. A shot change is declared when the value of the score becomes greater than the threshold and then becomes smaller than the threshold (similar to a peak above the threshold line).

Sum of absolute differences

A score is assigned to each frame by calculating the sum of the absolute differences between consecutive frames for every pixel. This value is then normalized by the size of the frame. When the score of a frame is greater than the threshold, a shot change is declared. This method works quite well with simple videos but it is not robust against movements and changes in lighting.

Histogram differences

Each frame is converted into a gray-scale image. A histogram with 256 bins, representing all the possible values of a pixel, is created for each frame. Then, a score is assigned by calculating the sum of the absolute differences between histograms of consecutive frames. This method is more robust against small changes in a frame.

Performance

Let C be the number of correctly identified cuts, M be the number of cuts that are not identified, and F be the number of falsely identified cuts. To evaluate how well the algorithm is detecting the shots, the recall (V), precision (P) and F_1 scores for each clip are calculated. They are defined as below.

$$V = \frac{C}{C + M}$$

$$P = \frac{C}{C + F}$$

$$F_1 = \frac{2 \times P \times V}{P + V}$$
(1)

Figure 1: Sum of absolute differences scores of clip 1

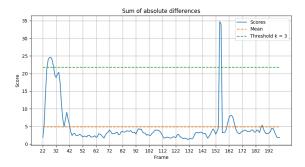


Figure 2: Histogram differences scores of clip 1

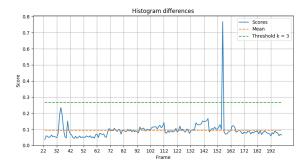


Figure 3: Sum of absolute differences scores of clip 2

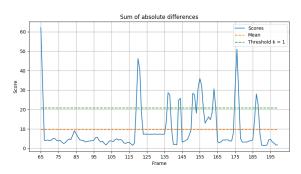


Figure 5: Sum of absolute differences scores of clip 3

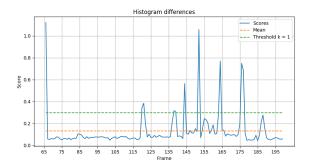
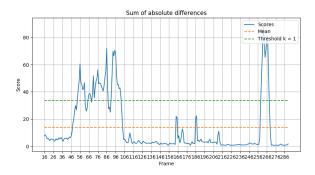


Figure 4: Histogram differences scores of clip 2

Figure 6: Histogram differences scores of clip 3



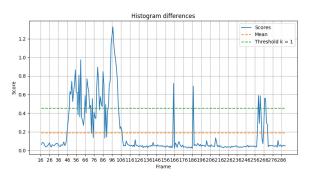


Table 1 and Table 2 shows the respective scores for both methods. On average, histogram differences achieve a higher score across all three metrics comparing to sum of absolute differences method. For example, the transition from frame 164 (Figure 7) to frame 165 (Figure 8) is not detected by the latter but it is by the former. It performs especially better in the montage section in clip 3, where there are frequent shot changes.

Table 1: Shot detection performance with sum of absolute differences

Clip	Correct (C)	Missed (M)	Falsely detected (F)	Recall (V)	Precision (P)	F_1
1	1	0	1	1	0.5	0.6667
2	7	0	2	1	0.7778	0.8750
3	5	16	0	0.2381	1	0.3846

Table 2: Shot detection performance with histogram differences

Clip	Correct (C)	Missed (M)	Falsely detected (F)	Recall (V)	Precision (P)	F_1
1	1	0	0	1	1	1
2	6	1	1	0.8571	0.8571	0.8571
3	15	6	1	0.7142	0.9375	0.8118

Figure 7: Frame 164 of clip 3



Figure 8: Frame 165 of clip 3



The relevant code is in shot.py. To get the graphs of shot detection, run the command below.

```
python3 run.py shot_detection -t <type> -i <path to frames>
```

To add shot numbers into frames, run the command below. The shot number will appear on the bottom left corner of each frame.

```
python3 run.py add_shot_numbe -i <input directory> -o <output directory>
-k <k for thresholding>
```

2 Logo detection

Template matching is an object detection algorithm which is translation invariant but not scale or rotation invariant. As we are detecting the news company's logo, we can assume that the target of detection will be in a known orientation. Due to the possibility that there may be multiple occurrences of the logo in a frame, we cannot simply match SIFT features between the logo template and a frame. Template matching is run on templates of different sizes because the size of the logo in a frame is unknown. Then, a score is calculated for all the matches by normalized cross-correlation. The normalized version is chosen because brighter patches will not have a higher score. Also, the score obtained will be in the range [0, 1] and this makes choosing a threshold more intuitive. As the logo in the template and the one in the frame might be slightly different, possibly due to the differences in resolutions or styles, a looser threshold is first used to filter out the irrelevant matches.

For each match, if its score is greater than the loose threshold, it is kept. Then, the algorithm checks if that particular match is a slight translation of a match we have already decided to keep. If it is, the match is discarded. This prevents having multiple boxes around one logo. Afterwards, SIFT descriptors are calculated for the remaining matches and a score is calculated using feature matching and Lowe's ratio test with the logo template. If a match's score is above a different threshold, it is declared as a match of the template and a box is put around the match.

Initially, I only did one pass with either normalized cross-correlation or SIFT feature matching. However, this led to fairly poor results, with the algorithm often unable to detect multiple logos and including many irrelevant matches. Therefore, the two passes approach, first with a looser threshold using normalized cross-correlation and then with another threshold with SIFT feature matching, is used. As can be seen in Figure 10, the algorithm can detect multiple logos. In hindsight, it would have been more efficient to scale down the images to multiple scales instead of scaling

down the templates. This is because sliding the template across the image takes much longer than resizing every frame to different scales.

Figure 9: Logo detection on frame 104 of clip 1



Figure 10: Logo detection on frame 52 of clip 1



The relevant code is in logo.py. To run logo detection, run the command below.

```
python3 run.py logo_detection -i <input directory> -o <output directory>
-d <logo path> -t <min threshold for NCC>
```

3 Face detection and tracking

There are 260 images in the female and male classes respectively. Each image is accompanied by a .mat file specifying the coordinates of the left eye, right eye, nose and mouth. The following rules are used to crop the images in order to obtain the faces.

$$start_{x} = left \ eye_{x} - 0.5 \times (right \ eye_{x} - left \ eye_{x})$$

$$end_{x} = right \ eye_{x} + 0.5 \times (right \ eye_{x} - left \ eye_{x})$$

$$start_{y} = eyes_{y} - (mouth_{y} - eyes_{y})$$

$$end_{y} = mouth_{y} + (mouth_{y} - eyes_{y})$$

$$(2)$$

The relevant code for face cropping is in crop_images() in face.py.

The initial attempt to detect faces is to use skin detection - trying to filter out skin in images. Figure 11 and Figure 12 show the color distributions of faces in RGB and HSV color spaces. The HSV color space has narrower distributions, especially with hue. Although this method works sometimes as seen in Figure 13, it is not successful in general. It fails to detect a large area of the face of the man on the right in Figure 14 and includs a lot of the background. This model is especially poor when other things in the frame are very similar to human skin tone.

Figure 11: RGB distribution of fe male training images

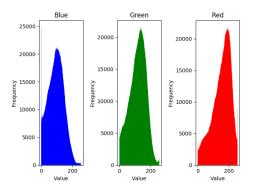


Figure 12: HSV distribution of female training images

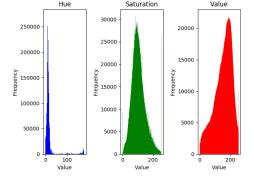


Figure 13: HSV colour detection on frame 160 of clip 1

Figure 14: HSV colour detection on frame 50 of clip 1





The relevant code for HSV face detection is in visualize_distributions() and face_detection_hsv() in face.py.

I ended up using cv2.CascadeClassifier for detecting faces. The full name of this classifier is Haar feature-based cascade classifier for object detection. It is a OpenCV pre-trained classifier for face stored in an XML file. It works fairly well. It has no trouble detecting multiple people in a frame or people of colour, like in Figure 15. However, it is usually unable to detect faces turned to the side. The detector was unable to detect the person on the left in any of the frames he appeared in that particular position in Figure 16.

After obtaining the faces in a frame, for each face in the current frame, the SIFT descriptors are found. Then, they are matched to the SIFT descriptors of all the faces in the previous frame using feature matching and Lowe's ratio test. Then, a score is obtained from the number of matches. If the score is above a set threshold, the algorithm declares that the face we are looking at is found in the previous frame and we will display the index assigned to that particular face in the previous frame. If the face is not found, we assign a new index to the face. The index of each face is labelled on top of the box.

Figure 15: Face detection on frame 69 of clip 2



Figure 16: Face detection on frame 110 of clip 1



4 Gender classification

90% of the images (234 images from each class) are used for training, whereas the other 10% (26 images from each class) are used for testing the accuracy of the model. The accuracy is defined as the percent of correctly identified faces, not SIFT descriptors, in the cases of support-vector machine and neural network.

SVM

The SIFT descriptors of the training images are passed into the SVM model for training. A radial basis function kernel is used. Then, for each detected face in a frame, the SIFT descriptors are extracted and fed to the trained SVM model. A prediction for each descriptor is obtained. If more descriptors are predicted as female than male, the image is classified as female. If more descriptors are predicted as male than female, the image is classified as male. If there are equal number of descriptors being predicted as both female and male, the image is then classified as unknown.

Neural network

The same as SVM, except with a neural network model instead. The model uses a binary crossentropy loss, adam for optimization, and accuracy as the metric. The neural network used is shown in Figure 17.

CNN

All the training and testing images are padded with black borders to obtain a square shape and then resized to be 72 pixels by 72 pixels. The training images are then passed to the CNN model shown in Figure 18 for training. The model uses a binary crossentropy loss, stochastic gradient descent for optimization, and accuracy as the metric. Each detected face is padded to obtain a square shape. Then, the image is resized to be 72 pixels by 72 pixels. The resized image is then passed to the trained CNN model and a category prediction is obtained.

Figure 18: CNN model

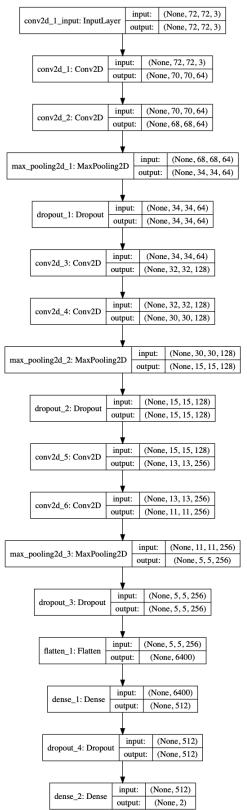
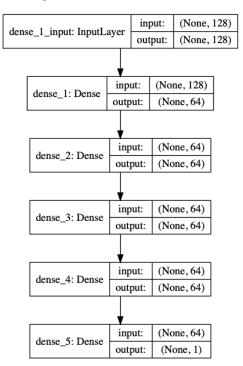


Figure 17: Neural network model



The relevant code for training any of the three models is in train_model() in face.py. To train a gender classification model, run the command below.

```
python3 run.py train -m <model path after training>
-c <classification model (SVM, NN_SIFT, or CNN)>
```

The relevant code for face detection (including gender classification and face tracking) is in face_detection() in face.py. Run the command below.

```
python3 run.py face_detection -i <input directory> -o <output directory>
-c <classification model (SVM, NN_SIFT, or CNN)> -m <trained model path>
-s <min size for face detection>
```

Performance

Table 3 shows the test accuracies of the three models. As expected, CNN performed poorly, achieving an accuracy that is equivalent to random guesses, due to the very small training data size of 468. It consistently classify all the faces as either female and male.

Table 3: Gender classification performance

Model	Description	Accuracy on test set
SVM	Using SIFT descriptors of faces	100.00%
Neural network	Using SIFT descriptors of faces	92.30%
CNN	Using cropped and resized faces	50.00%

5 Make video

The relevant code for combining all the frames to a video is in make_video() in utils.py. To do logo detection, face detection, face tracking and gender classification and make a video, run the command below.

```
python3 run.py run_all -i <input directory> -o <output directory>
-d <logo path> -t <min threshold for NCC> -m <trained model path>
-c <classification model (SVM, NN_SIFT, or CNN)> -v <name of output video>
-f <frame per second> -s <min size for face detection>
```

6 References

Video shot boundary detection based on color histogram

Wikipedia - Shot transition detection

Shot detection using pixel wise difference with adaptive threshold and color histogram method in compressed and uncompressed video

7 Code

```
Listing 1: shot.py
 1 import cv2
 2 import matplotlib.pyplot as plt
 3 import numpy as np
 4 import os
 6
    def add_shot_number(input_dir, output_dir, shots):
 7
 8
        Add shot number using the shot changes specified in variable shots.
9
10
        print("add_shot_number")
        if not os.path.isdir(output_dir):
11
           os.mkdir(output_dir)
12
13
14
        exts = [".jpg", ".png"]
15
        img_names = [img for img in os.listdir(input_dir) if img.endswith(tuple(exts))]
        # Sort images by name in ascending order
16
        img_names.sort(key=lambda img: int("".join(filter(str.isdigit, img))))
17
18
19
        i = 0
        shot = 0
20
21
        for filename in img_names:
           image_num = int("".join(filter(str.isdigit, filename)))
22
           img = cv2.imread(os.path.join(input_dir, filename))
23
24
           if i < len(shots) and image_num == shots[i]:</pre>
25
               shot += 1
26
               i += 1
27
           # Add shot number to frame
28
           text = str(shot)
29
           # Get width and height of the text box
           text_width, text_height = cv2.getTextSize(text, cv2.FONT_HERSHEY_PLAIN, fontScale
30
                \hookrightarrow =1.5, thickness=2)[0]
31
           # Set the text start position
32
           text_x = 10
33
           text_y = img.shape[0] - 10
34
           box_coords = ((text_x, text_y + 2), (text_x + text_width - 2, text_y -
                \hookrightarrow text_height - 4))
35
           cv2.rectangle(img, box_coords[0], box_coords[1], (255, 255, 255), cv2.FILLED)
           cv2.putText(img, text, (text_x, text_y), cv2.FONT_HERSHEY_PLAIN, fontScale=1.5,
36
                \hookrightarrow color=(0, 0, 0), thickness=2)
37
           # Save frame
38
           cv2.imwrite(os.path.join(output_dir, filename), img)
           # cv2.imshow("A box!", img)
39
40
           \# cv2.waitKey(0)
41
    def shot_detection(input_dir, method, k):
42
43
44
        Get graphs of scores of shot changes using the method specify in method.
45
        Method should be either SAD2 or HD.
46
47
        scores = []
        exts = [".ipq", ".pnq"]
48
        img_names = [img for img in os.listdir(input_dir) if img.endswith(tuple(exts))]
49
        # Sort images by name in ascending order
50
        img_names.sort(key=lambda img: int("".join(filter(str.isdigit, img))))
51
        start_idx = int(img_names[0][:img_names[0].find(".jpg")])
52
        end_idx = int(img_names[-1][:img_names[-1].find(".jpg")])
53
```

```
54
 55
        # Sum of absolute differences
 56
        if method == "SAD2":
57
            prev_img = None
58
            for i in range(len(img_names)):
59
               # Default type is numpy.uint64
               curr_img = cv2.imread(os.path.join(input_dir, img_names[i])).astype(np.int64)
60
               if i == 0:
61
62
                   r, c, d = curr_img.shape
                   next_img = cv2.imread(os.path.join(input_dir, img_names[i + 1])).astype(np.
63
                       \hookrightarrow int64)
64
                   score = np.sum(np.abs(curr_img - next_img))
65
               elif i == len(img_names) - 1:
                   score = np.sum(np.abs(curr_img - prev_img))
66
67
               else:
                   next_img = cv2.imread(os.path.join(input_dir, img_names[i + 1])).astype(np.
68
                       \rightarrow int64)
69
                   score = 0.5 * np.sum(np.abs(curr_img - prev_img)) + 0.5 * np.sum(np.abs(
                       70
               scores.append(score)
               prev_img = curr_img
71
 72
            x = np.arange(start_idx, end_idx + 1)
            scores = np.array(scores) / (r * c * d)
 73
 74
            title = "Sum of absolute differences"
 75
            new_filename = "output/" + input_dir.name + "_score_sad2.png"
76
        elif method == "SAD":
77
78
            prev_img = None
 79
            for i in range(len(img_names)):
80
               # Default type is numpy.uint64
               curr_img = cv2.imread(os.path.join(input_dir, img_names[i])).astype(np.int64)
81
82
               if i == 0:
83
                   r, c, d = curr_img.shape
84
                   prev_img = curr_img
85
                   continue
86
               score = np.sum(np.abs(curr_img - prev_img))
87
               scores.append(score)
88
               prev_img = curr_img
89
            x = np.arange(start_idx + 1, end_idx + 1)
90
            scores = np.array(scores) / (r * c * d)
91
            title = "Sum of absolute differences"
            new_filename = "output/" + input_dir.name + "_score_sad2.png"
92
93
94
        # Histogram differences
        elif method == "HD":
95
            prev_histogram = None
96
97
            for i in range(len(img_names)):
98
               # Default type is numpy.uint8
               curr_img = cv2.imread(os.path.join(input_dir, img_names[i]))
99
100
               # Default type is numpy.uint8
               curr_img_g = cv2.cvtColor(curr_img, cv2.COLOR_BGR2GRAY).astype(np.int16)
101
               histogram = np.histogram(np.ravel(curr_img_g), bins=np.arange(-1, 256))
102
               if i == 0:
103
104
                   r, c = curr_img_g.shape
105
                   prev_histogram = histogram[0]
106
107
               score = np.sum(np.abs(histogram[0] - prev_histogram))
108
               scores.append(score)
109
               prev_histogram = histogram[0]
```

```
110
111
            x = np.arange(start_idx + 1, end_idx + 1)
            scores = np.array(scores) / (r * c)
112
            title = "Histogram differences"
113
            new_filename = "output/" + input_dir.name + "_score_hd.png"
114
115
         else:
116
            raise ValueError("Illegal method value")
117
118
119
        y_mean = [np.mean(scores)] * len(x)
         threshold = [np.mean(scores) + k * np.std(scores)] * len(x)
120
121
         f = plt.figure(figsize=(10, 5))
122
         ax = f.gca()
123
        ax.set_xticks(np.arange(start_idx, end_idx, 10))
124
        plt.title(title)
        plt.xlabel("Frame")
plt.ylabel("Score")
125
126
        plt.plot(x, scores, label="Scores")
127
        # Plot the average line
128
        plt.plot(x, y_mean, label="Mean", linestyle="--")
129
        plt.plot(x, threshold, label="Threshold k = " + str(k), linestyle="--")
130
        plt.legend(loc="upper right")
131
132
        plt.grid()
133
         f.savefig(new_filename)
134
        print("Output saved to " + new_filename)
     Listing 2: logo.py
  1 import cv2
     import imutils
     import numpy as np
  4
     import os
  5
  6
     def get_score(img1, img2):
  7
  8
         Get similarity score between img1 and img2 using SIFT features matching and
 9
         Lowe's ratio testing.
 10
 11
        sift = cv2.xfeatures2d.SIFT_create()
 12
        kp1, des1 = sift.detectAndCompute(img1, None)
 13
        kp2, des2 = sift.detectAndCompute(img2, None)
 14
         if len(kp1) < 2 or len(kp2) < 2:
 15
            return 0
 16
         index_params = dict(algorithm=0, trees=5)
 17
        flann = cv2.FlannBasedMatcher(index_params, None)
         matches = flann.knnMatch(des1, des2, k=2)
 18
        good_matches = []
 19
 20
         for m, n in matches:
 21
            if m.distance < 0.6*n.distance:</pre>
 22
                good_matches.append(m)
 2.3
        num_kps = 0
 24
        if len(kp1) <= len(kp2):</pre>
            num_kps = len(kp1)
 25
 26
         else:
 27
            num_kps = len(kp2)
        score = len(good_matches) / num_kps * 100
 28
 29
        return score
 30
31
     def logo_detection(input_dir, output_dir, logo_path, min_threshold):
        print("detect_logo")
 32
```

```
33
34
        if not os.path.isdir(output dir):
35
           os.mkdir(output_dir)
36
37
        template = cv2.imread(str(logo_path))
38
        template_g = cv2.cvtColor(template, cv2.COLOR_BGR2GRAY)
39
40
        w, h = template_g.shape[::-1]
41
42
        # Compute template of different sizes
        if template.shape[0] < 50:</pre>
43
           \# scales = np.linspace(0.8, 1.0, 6)[::-1]
44
45
           scales = np.linspace(0.9, 1.0, 2)[::-1]
46
        else:
47
           \# scales = np.linspace(0.1, 1.0, 25)[::-1]
           scales = np.linspace(0.5, 1.0, 10)[::-1]
48
49
        templates = []
50
        ratios = []
        for scale in scales:
51
52
           resized = imutils.resize(template_g, width=int(template_g.shape[1] * scale))
53
           templates.append(resized)
54
           ratios.append(resized.shape[1] / float(template_g.shape[1]))
55
56
        for img_name in os.listdir(input_dir):
57
           if not img_name.endswith(".jpg"):
58
               continue
59
           img = cv2.imread(os.path.join(input_dir, img_name))
60
           img_g = cv2.cvtColor(img, cv2.COLOR_BGR2GRAY)
           img_canny = cv2.Canny(img_g, 100, 200)
61
62
63
           p = -1
64
           q = -1
65
           matches = []
66
           # Loop through the templates
67
           for i in range(len(templates) -1, -1, -1):
68
               # Stop when template is bigger than image
69
               if img_g.shape[0] < templates[i].shape[0] or img_g.shape[1] < templates[i].</pre>
                   \hookrightarrow shape[1]:
70
                   break
71
               # First pass — normalized cross correlation
72
               match = cv2.matchTemplate(img_g, templates[i], cv2.TM_CCORR_NORMED) # img_g.shape
73
                   \hookrightarrow - template_g.shape + 1
               if p == -1 and q == -1:
74
                   p, q = match.shape
75
76
               m, n = match.shape
77
               matches.append(np.pad(match, ((0, p - m), (0, q - n)), mode="constant",

    constant_values=0))
78
79
           boxes = []
80
           matches = np.array(matches)
81
           r, max_y, max_x = np.unravel_index(np.argmax(matches), matches.shape)
82
           r = ratios[len(ratios) - 1 - r]
83
84
           max_val = np.max(matches)
           max_thresh = max(max_val * 0.95, min_threshold)
85
86
           # If the match with the highest score is smaller than the min threshold,
87
           # there is no match in this image and just saves the input image.
           if max_val < min_threshold:</pre>
88
```

```
89
                cv2.imwrite(os.path.join(output_dir, img_name), img)
 90
                continue
            start_x, start_y = max_x, max_y
 91
            end_x, end_y = int((max_x + w * r)), int((max_y + h * r))
 92
 93
            match_score = get_score(template, img[start_y:end_y, start_x:end_x])
 94
            if match_score < 5:</pre>
 95
                cv2.imwrite(os.path.join(output_dir, img_name), img)
                continue
 96
 97
            boxes.append((r, max_y, max_x, 1))
 98
            # Matches obtained from first pass
 99
100
            match_locations = np.where(matches >= max_thresh)
101
            for i in range(len(match_locations[0])):
102
                r1, y1, x1 = ratios[len(ratios) - 1 - match_locations[0][i]], match_locations
                    \hookrightarrow [1][i], match_locations[2][i]
                found = False
103
104
                for j in range(len(boxes)):
                    r2, y2, x2, count = boxes[j]
105
106
                    # Check if two boxes of the same size overlap or
107
                    # if a smaller one is contained in the bigger one
                    if (r1 == r2 \text{ and } np.abs(x1 - x2) < w * r1 \text{ and } np.abs(y1 - y2) < h * r1) or
108
                        ((r1 < r2)) and (x1 <= (x2 + w * r2) <= (x1 + w * r1)) and (y1 <= (y2 + h))
109
                            \rightarrow * r2) <= (y1 + h * r1))) or
110
                        (np.abs(x1 - x2) < 0.5 * w * r1 and np.abs((x1 + w * r1) - (x2 + w * r2))
                            \hookrightarrow )) < 0.5 * w * r1 and
111
                        np.abs(y1 - y2) < 0.5 * h * r1 and np.abs((y1 + h * r1) - (y2 + h * r2))
                            \hookrightarrow ) < 0.5 * h * r1):
                        boxes[j] = (r2, y2, x2, count + 1)
112
113
                        found = True
                        break
114
115
                if not found:
116
                    start_x, start_y = x1, y1
117
                    end_x, end_y = int((x1 + w * r1)), int((y1 + h * r1))
118
                    # Second pass — SIFT features matching
119
                    match_score = get_score(template, img[start_y:end_y, start_x:end_x])
120
                    if match_score > 5:
121
                       boxes.append((r1, y1, x1, 1))
122
123
            for r, y, x, count in boxes:
124
                start_x, start_y = x, y
125
                end_x, end_y = int((x + w * r)), int((y + h * r))
126
                cv2.rectangle(img, (start_x, start_y), (end_x, end_y), (0, 255, 0), thickness
                    \hookrightarrow =2)
127
            cv2.imwrite(os.path.join(output_dir, img_name), img)
128
129
            # cv2.imshow("img", img)
130
            # cv2.waitKey(0)
     Listing 3: face.py
  1 import copy
  2 import cv2
     import matplotlib.pyplot as plt
  4 import numpy as np
  5 import os
  6 import pickle
  7 import scipy.io, scipy.misc
  8 from skimage import io
  9 from skimage import transform as tf
```

```
from skimage.color import rgb2hsv
   from sklearn import svm
12
   from utils import shuffle
13
14
   def visualize_distributions(imgs_dir):
15
       Visualize the distributions of values in HSV and BGR of pictures in imgs_dir.
16
17
18
       H, S, V = None, None, None
19
       B, G, R = None, None, None
       for filename in os.listdir(imgs_dir):
20
21
              if not filename.endswith(".png"):
22
                  continue
23
              img = cv2.imread(imgs_dir + filename)
24
              hsv_img = cv2.cvtColor(img, cv2.COLOR_BGR2HSV)
25
              if H is None:
                  H = hsv_img[..., 0].flatten()
26
                  S = hsv_img[..., 1].flatten()
27
                  V = hsv_img[..., 2].flatten()
28
                  B = img[..., 0].flatten()
29
                  G = img[..., 1].flatten()
30
                  R = img[..., 2].flatten()
31
32
              else:
33
                  H = np.concatenate([H, hsv_img[..., 0].flatten()])
34
                  S = np.concatenate([S, hsv_img[..., 1].flatten()])
                  V = np.concatenate([V, hsv_img[..., 2].flatten()])
35
                  B = np.concatenate([B, img[..., 0].flatten()])
36
37
                  G = np.concatenate([G, img[..., 1].flatten()])
                  R = np.concatenate([R, img[..., 2].flatten()])
38
39
       # Plot
40
41
       f = plt.figure()
       ax1 = f.add\_subplot(1, 3, 1)
42
43
       ax1.hist(H, bins=180,
44
           range=(0.0, 180.0), histtype="stepfilled", color="b", label="Hue")
45
       plt.title("Hue")
46
       plt.xlabel("Value")
       plt.ylabel("Frequency")
47
48
       ax2 = f.add\_subplot(1, 3, 2)
49
       ax2.hist(S, bins=256,
           range=(0.0, 255.0), histtype="stepfilled", color="g", label="Saturation")
50
51
       plt.title("Saturation")
       plt.xlabel("Value")
52
       plt.ylabel("Frequency")
53
       ax3 = f.add_subplot(1, 3, 3)
54
       ax3.hist(V, bins=256,
55
           range=(0.0, 255.0), histtype="stepfilled", color="r", label="Value")
56
57
       plt.title("Value")
       plt.xlabel("Value")
58
59
       plt.ylabel("Frequency")
60
       f.tight_layout()
       f.savefig(imgs_dir + "_hsv_distributions.png")
61
       plt.show()
62
63
64
       f = plt.figure()
       ax1 = f.add\_subplot(1, 3, 1)
65
66
       ax1.hist(B, bins=256,
67
           range=(0.0, 255.0), histtype="stepfilled", color="b", label="Blue")
68
       plt.title("Blue")
```

```
plt.xlabel("Value")
 69
 70
         plt.ylabel("Frequency")
 71
         ax2 = f.add_subplot(1, 3, 2)
 72
        ax2.hist(G, bins=256,
 73
            range=(0.0, 255.0), histtype="stepfilled", color="g", label="Saturation")
        plt.title("Green")
 74
        plt.xlabel("Value")
 75
        plt.ylabel("Frequency")
 76
 77
         ax3 = f.add\_subplot(1, 3, 3)
 78
         ax3.hist(R, bins=256,
 79
            range=(0.0, 255.0), histtype="stepfilled", color="r", label="Red")
 80
        plt.title("Red")
        plt.xlabel("Value")
 81
        plt.ylabel("Frequency")
 82
 83
         f.tight_layout()
         f.savefig(imgs_dir + "_rgb_distributions.png")
 84
 85
         plt.show()
 86
 87
     def crop_images(old_dir, new_dir):
 88
 89
         Crop .jpg images in old_dir given coordinates of left eye, right eye,
 90
         nose and mouth in .mat files. Save cropped images in new_dir.
 91
 92
        print("crop_images")
 93
        os.mkdir(new_dir)
 94
 95
        H, S, V = None, None, None
 96
         for filename in os.listdir(old_dir):
            if not filename.endswith(".jpg"):
 97
 98
                continue
 99
100
            index = filename.find(".jpg")
101
            name = filename[:index]
102
103
            # Approximate coordinates of face
104
            coords = scipy.io.loadmat(old_dir + name + ".mat")
            start_x = int(coords["x"][0][0] - 0.5*(coords["x"][1][0] - coords["x"][0][0]))
105
106
            end_x = int(coords["x"][1][0] + 0.5*(coords["x"][1][0] - coords["x"][0][0]))
            start_y = int(coords["y"][0][0] - (coords["y"][3][0] - coords["y"][0][0])
107
            end_y = int(coords["y"][3][0] + (coords["y"][3][0] - coords["y"][2][0]))
108
            img = io.imread(old_dir + filename)
109
110
            face = img[start_y:end_y, start_x:end_x]
            # Save cropped image
111
            scipy.misc.imsave(new_dir + name + ".png", face)
112
113
     def resize_images(input_dir, output_dir, size=72):
114
115
116
         Resize images in input dir to size x size.
117
118
        print("resize_images")
         if not os.path.isdir(output_dir):
119
            os.mkdir(output_dir)
120
121
         for filename in os.listdir(input_dir):
122
123
            if not filename.endswith(".png"):
124
                continue
125
            img = cv2.imread(os.path.join(input_dir, filename))
126
            x, y, d = img.shape
127
            # Pad image to square
```

```
128
            if x > y:
129
               padded_img = np.pad(img, ((0, 0), (0, x - y), (0, 0)),
130
                   mode="constant", constant_values=0)
131
            else:
132
               padded_img = np.pad(img, ((0, y - x), (0, 0), (0, 0)),
                   mode="constant", constant_values=0)
133
            resized_img = cv2.resize(padded_img, (size, size))
134
135
            scipy.misc.imsave(output_dir + filename, cv2.cvtColor(resized_img, cv2.
                136
     def get_features(input_dir):
137
138
139
        Get keypoints and descriptors of the images in input_dir with SIFT.
140
141
        kps, des = None, None
        sift = cv2.xfeatures2d.SIFT_create()
142
143
        for filename in os.listdir(input_dir):
144
            if not filename.endswith(".png"):
145
146
                continue
            img = io.imread(input_dir + filename)
147
            kp, d = sift.detectAndCompute(img, None)
148
149
150
            if des is None:
151
               kps = kp
               des = d
152
153
            else:
154
               kps = np.concatenate([kps, kp], axis=0)
155
               des = np.concatenate([des, d], axis=0)
156
        return kps, des
157
158
     def get_data(f_dir, m_dir):
159
160
        Get the images of the F and M classes from their respective directories and shuffle.
161
162
        import keras
163
        x = []
164
        y = [0] * len(os.listdir(f_dir)) + [1] * len(os.listdir(m_dir))
165
        imgs = os.listdir(f_dir) + os.listdir(m_dir)
166
        for filename in os.listdir(f_dir):
            if not filename.endswith(".png"):
167
168
                continue
169
            x.append(cv2.imread(os.path.join(f_dir, filename)))
        for filename in os.listdir(m_dir):
170
            if not filename.endswith(".png"):
171
172
                continue
173
            x.append(cv2.imread(os.path.join(m_dir, filename)))
174
175
        x, y = shuffle(x, y)
176
        y = keras.utils.to_categorical(y, num_classes=2)
177
        return x, y
178
179
     def train_model(model_path, classification):
180
181
        Train either a SVM or neural network model using SIFT features or
182
        a CNN using face images.
183
184
        OLD_F_DIR = "original_data/female/"
        OLD_M_DIR = "original_data/male/"
185
```

```
F_TRAIN_DIR = "data/female_train/"
186
        M TRAIN DIR = "data/male train/"
187
        F_TEST_DIR = "data/female_test/"
188
        M_TEST_DIR = "data/male_test/"
189
        F_TRAIN_CNN_DIR = "data/female_cnn_train/"
190
        M_TRAIN_CNN_DIR = "data/male_cnn_train/"
191
        F_TEST_CNN_DIR = "data/female_cnn_test/"
192
        M_TEST_CNN_DIR = "data/male_cnn_test/"
193
194
195
        if not os.path.isdir("data/"):
            os.mkdir("data/")
196
197
            crop_images(OLD_F_DIR, F_TRAIN_DIR)
198
            crop_images(OLD_M_DIR, M_TRAIN_DIR)
199
         if classification == "SVM":
200
            f_train_kps, f_train_des = get_features(F_TRAIN_DIR)
201
202
            m_train_kps, m_train_des = get_features(M_TRAIN_DIR)
            x_train = np.concatenate([f_train_des, m_train_des], axis=0)
203
            y_{train} = [-1] * len(f_{train_des}) + [1] * len(m_{train_des})
204
            model = svm.SVC(kernel="rbf", gamma="scale", C=10.0)
205
206
            model.fit(x_train, y_train)
207
208
            # Save model
209
            pickle.dump(model, open(model_path, "wb"))
210
            print("Model saved as " + str(model_path))
211
        elif classification == "NN_SIFT":
212
            import keras
213
            from keras.layers import Conv2D, Dense, Dropout, Flatten, MaxPooling2D
214
            from keras.models import Sequential
215
            from keras.optimizers import Adam, SGD
216
217
            from keras.utils import plot_model
218
219
            f_train_kps, f_train_des = get_features(F_TRAIN_DIR)
220
            m_train_kps, m_train_des = get_features(M_TRAIN_DIR)
            x_train = np.concatenate([f_train_des, m_train_des], axis=0)
221
222
            y_train = [0] * len(f_train_des) + [1] * len(m_train_des)
223
            x_train, y_train = shuffle(x_train, y_train)
224
            f_test_kps, f_test_des = get_features(F_TEST_DIR)
225
            m_test_kps, m_test_des = get_features(M_TEST_DIR)
226
227
            x_test = np.concatenate([f_test_des, m_test_des], axis=0)
228
            y_{test} = [0] * len(f_{test_des}) + [1] * len(m_{test_des})
            x_test, y_test = shuffle(x_test, y_test)
229
230
231
            model = Sequential()
232
            model.add(Dense(64, activation='relu', kernel_initializer='random_normal', input_dim
                 \rightarrow =128))
            model.add(Dense(64, activation='relu', kernel_initializer='random_normal'))
model.add(Dense(64, activation='relu', kernel_initializer='random_normal'))
233
234
            model.add(Dense(64, activation='relu', kernel_initializer='random_normal'))
235
            model.add(Dense(1, activation='sigmoid', kernel_initializer='random_normal'))
236
            model.compile(optimizer ='adam',loss='binary_crossentropy', metrics =['accuracy'])
237
238
239
            model.fit(x_train, y_train, batch_size=10, epochs=20)
            score = model.evaluate(x_test, y_test)
240
            print("score:", score)
241
242
243
            model.save(model_path)
```

```
print("Model saved as " + str(model_path))
244
245
            plot model(model, to file="output/nn model.png", show shapes=True)
246
        elif classification == "CNN":
247
248
            import keras
            from keras.layers import Conv2D, Dense, Dropout, Flatten, MaxPooling2D
249
            from keras.models import Sequential
250
            from keras.optimizers import Adam, SGD
251
            from keras.utils import plot_model
252
253
254
            np.random.seed(123)
255
            size = 72
256
257
            # Pad and resize images to size by size
            # resize_images(F_TRAIN_DIR, F_TRAIN_CNN_DIR, size)
258
            # resize_images(M_TRAIN_DIR, M_TRAIN_CNN_DIR, size)
259
            # resize images(F_TEST_DIR, F_TEST_CNN_DIR, size)
260
            # resize_images(M_TEST_DIR, M_TEST_CNN_DIR, size)
261
262
263
            x_train, y_train = get_data(F_TRAIN_CNN_DIR, M_TRAIN_CNN_DIR)
            x_test, y_test = get_data(F_TEST_CNN_DIR, M_TEST_CNN_DIR)
264
265
266
            model = Sequential()
267
            model.add(Conv2D(32, (3, 3), activation="relu", input_shape=(size, size, 3)))
268
            model.add(Conv2D(32, (3, 3), activation="relu"))
            model.add(MaxPooling2D(pool_size=(2, 2)))
269
270
            model.add(Dropout(0.25))
271
272
            model.add(Conv2D(64, (3, 3), activation="relu"))
           model.add(Conv2D(64, (3, 3), activation="relu"))
273
            model.add(MaxPooling2D(pool_size=(2, 2)))
274
275
           model.add(Dropout(0.25))
276
277
            model.add(Conv2D(128, (3, 3), activation="relu"))
278
            model.add(Conv2D(128, (3, 3), activation="relu"))
279
            model.add(MaxPooling2D(pool_size=(2, 2)))
280
            model.add(Dropout(0.25))
281
282
            model.add(Flatten())
283
            model.add(Dense(256, activation="relu"))
            model.add(Dropout(0.5))
284
            model.add(Dense(2, activation="softmax"))
285
286
287
            adam = Adam(1r=0.0005)
            model.compile(loss="binary_crossentropy", optimizer=adam, metrics=["accuracy"])
288
289
290
            model.fit(x_train, y_train, epochs=10)
291
            score = model.evaluate(x_test, y_test)
            print("score:", score)
292
293
            model.save(model_path)
294
            print("Model saved as " + str(model_path))
295
           plot_model(model, to_file="output/cnn_model.png", show_shapes=True)
296
297
298
        else:
299
            raise ValueError("Illegal classification value")
300
301
     def predict_model(model_path):
302
```

```
303
        Get accuacy of test set for either the SVM or neural network model.
304
305
        print("predict_model")
        F_TEST_DIR = "data/female_test/"
306
        M_TEST_DIR = "data/male_test/"
307
308
        ext = os.path.splitext(model_path)[1]
309
        if ext == ".sav":
310
            model = pickle.load(open(model_path, "rb"))
311
        elif ext == ".h5":
312
            from keras.models import load_model
313
314
            model = load_model(model_path)
315
        else:
            raise ValueError("Not valid model extension")
316
317
        sift = cv2.xfeatures2d.SIFT_create()
318
319
        correct = 0
320
        for filename in os.listdir(F_TEST_DIR):
321
322
            if not filename.endswith(".png"):
323
                continue
            img = cv2.imread(os.path.join(F_TEST_DIR, filename))
324
            kps, des = sift.detectAndCompute(img, None)
325
326
            result = model.predict(des)
327
            if ext == ".sav":
                f_{\text{count}} = np.count_{\text{nonzero}}(result == -1)
328
329
               m_count = np.count_nonzero(result == 1)
            elif ext == ".h5":
330
                f_count = np.sum(result < 0.5)</pre>
331
332
               m_{count} = np.sum(result > 0.5)
            if f_count > m_count:
333
334
                correct += 1
335
336
        for filename in os.listdir(M_TEST_DIR):
337
            if not filename.endswith(".png"):
338
                continue
339
            img = cv2.imread(os.path.join(M_TEST_DIR, filename))
            kps, des = sift.detectAndCompute(img, None)
340
341
            result = model.predict(des)
            if ext == ".sav":
342
                f_{count} = np.count_{nonzero}(result == -1)
343
               m_count = np.count_nonzero(result == 1)
344
            elif ext == ".h5":
345
                f_{count} = np.sum(result < 0.5)
346
347
               m_{count} = np.sum(result > 0.5)
348
            if m_count > f_count:
349
               correct += 1
        score = correct / (len(os.listdir(F_TEST_DIR)) + len(os.listdir(M_TEST_DIR))) * 100
350
351
        print(str(score) + "% of images are categorized correctly")
352
353
     def face_detection_hsv(input_dir, output_dir):
354
        Face detection with HSV color detection (does not work well).
355
356
        print("face_detection_hsv")
357
358
359
        if not os.path.isdir(output_dir):
360
            os.mkdir(output_dir)
        for filename in os.listdir(input_dir):
361
```

```
if not filename.endswith(".jpg"):
362
363
                continue
364
            img = cv2.imread(os.path.join(input_dir, filename))
365
366
            # HSV color detection
            hsv_img = cv2.cvtColor(img, cv2.COLOR_BGR2HSV)
367
368
            lower1 = np.array([0, 48, 80])
369
            upper1 = np.array([20, 255, 255])
370
371
            mask1 = cv2.inRange(hsv_img, lower1, upper1) # img hsv.shape
            lower2 = np.array([170, 0, 0])
372
373
            upper2 = np.array([180, 255, 255])
374
            mask2 = cv2.inRange(hsv_img, lower2, upper2) # img hsv.shape
375
            mask = cv2.bitwise_or(mask1, mask2)
376
            kernel = cv2.getStructuringElement(cv2.MORPH_ELLIPSE, (10, 10))
377
378
            new_img = cv2.bitwise_and(img, img, mask=mask)
379
            cv2.imwrite(os.path.join(output_dir, filename), new_img)
380
381
     def face_detection_cascade(input_dir, output_dir, model_path, classification, min_size):
382
         Face detection with classifier, face tracking, and gender classification
383
384
         using the method specified in classification variable using model_path.
385
386
        print("face_detection_cascade")
        XML_FILENAME = "models/haarcascade_frontalface_default.xml"
387
388
         F_TEXT = "Female"
389
        M_TEXT = "Male"
        O_TEXT = "Not sure"
390
391
        F_{COLOR} = (0, 0, 255)
        M_{COLOR} = (255, 0, 0)
392
393
         O_{COLOR} = (0, 255, 0)
394
395
         if classification == "SVM":
396
            model = pickle.load(open(model_path, "rb"))
397
        elif classification == "NN_SIFT":
398
            from keras.models import load_model
399
            model = load_model(model_path)
400
        elif classification == "CNN":
401
            from keras.models import load_model
            model = load_model(model_path)
402
403
            size = 72
404
         else:
            raise ValueError("Illegal classification value")
405
406
407
         if not os.path.isdir(output_dir):
408
            os.mkdir(output_dir)
409
410
         exts = [".jpg", ".png"]
         img_names = [img for img in os.listdir(input_dir) if img.endswith(tuple(exts))]
411
         # Sort images by name in ascending order
412
         img_names.sort(key=lambda img: int("".join(filter(str.isdigit, img))))
413
        sift = cv2.xfeatures2d.SIFT_create()
414
        index_params = dict(algorithm=0, trees=5)
415
416
         flann = cv2.FlannBasedMatcher(index_params, None)
417
         face count = 0
418
        prev_faces = {}
419
420
         for filename in img_names:
```

```
img = cv2.imread(os.path.join(input_dir, filename))
421
422
           img_g = cv2.cvtColor(img, cv2.COLOR_BGR2GRAY)
423
           424
425
           ### Cascade face detection ###
           426
           face_cascade = cv2.CascadeClassifier(XML_FILENAME)
427
           faces = face_cascade.detectMultiScale(img_g, scaleFactor=1.3,
428
              minNeighbors=5, minSize=(min_size, min_size))
429
430
           curr_faces = {}
431
432
           for x, y, w, h in faces:
433
              434
              ### Face tracking ###
              435
              face = img[y:y+h, x:x+w]
436
437
              kp1, des1 = sift.detectAndCompute(face, None)
              found_face = False
438
              for (x2, y2, w2, h2), index in prev_faces.items():
439
440
                  face2 = img[y2:y2+h, x2:x2+w]
                  kp2, des2 = sift.detectAndCompute(face2, None)
441
                  if len(kp1) < 2 or len(kp2) < 2:
442
                     continue
443
444
                  matches = flann.knnMatch(des1, des2, k=2)
445
                  good_matches = []
                  for m, n in matches:
446
                     if m.distance < 0.6*n.distance:</pre>
447
                        good_matches.append(m)
448
                  num\_kps = 0
449
                  if len(kp1) <= len(kp2):</pre>
450
                     num_kps = len(kp1)
451
452
453
                     num_kps = len(kp2)
454
                  score = len(good_matches) / num_kps * 100
455
                  if score > 45:
456
                     found_face = True
457
                     face_number = index
458
                     curr_faces[(x, y, w, h)] = index
              if not found_face:
459
460
                  face_number = face_count
                  curr_faces[(x, y, w, h)] = face_count
461
462
                  face_count += 1
463
              464
465
              ### Gender classification ###
              466
              if classification == "SVM":
467
468
                  # Get SIFT descriptors
                  kps, des = sift.detectAndCompute(face, None)
469
470
                  # Predict female or male
                  result = model.predict(des)
471
                  f_{count} = np.count_{nonzero}(result == -1)
472
                  m_count = np.count_nonzero(result == 1)
473
                  # Plot color box
474
475
                  if f_count > m_count:
                     text = str(face_number) + " " + F_TEXT + ": " +\
476
477
                        format(f_count/(f_count + m_count)*100, ".2f") + "%"
478
                     color = F_COLOR
                  elif m_count > f_count:
479
```

```
text = str(face_number) + " " + M_TEXT + ": " +\
480
481
                          format(m count/(f count + m count)*100. ".2f") + "%"
482
                       color = M_COLOR
483
                   else:
484
                       text = str(face_number) + " " + O_TEXT
                       color = O_COLOR
485
486
               elif classification == "NN_SIFT":
487
488
                   assert w == h
489
                   kps, des = sift.detectAndCompute(face, None)
490
                   prediction = model.predict(des)
491
                   f_{count} = np.sum(prediction < 0.5)
492
                   m_count = np.sum(prediction > 0.5)
                   if f_count > m_count:
493
                       text = str(face_number) + " " + F_TEXT + ": " +\
494
                          format(f_count/(f_count + m_count)*100, ".2f") + "%"
495
496
                       color = F_COLOR
497
                   elif m_count > f_count:
                       text = str(face_number) + " " + M_TEXT + ": " +\
498
                          format(m_count/(f_count + m_count)*100, ".2f") + "%"
499
500
                       color = M_COLOR
501
                   else:
                       text = str(face_number) + " " + 0_TEXT
502
503
                       color = O_COLOR
504
               elif classification == "CNN":
505
506
                   assert w == h
507
                   face = cv2.resize(face, (size, size))
508
                   prediction = model.predict(np.array([face]), verbose=0)
                   prediction = prediction[0]
509
                   if prediction[0] > 0.5:
510
                       text = str(face_number) + " " + F_TEXT + ": " +\
511
512
                          format(prediction[0]*100, ".2f") + "%"
513
                       color = F_COLOR
514
                   elif prediction[0] < 0.5:</pre>
                       text = str(face_number) + " " + M_TEXT + ": " +\
515
516
                          format(prediction[0]*100, ".2f") + "%"
517
                       color = M_COLOR
518
                   else:
                       text = str(face_number) + " " + O_TEXT
519
520
                       color = 0 COLOR
521
               cv2.rectangle(img, (x, y), (x+w, y+h), color, thickness=2)
522
               cv2.putText(img, text, (x, y-10), color=color,
                   fontFace=cv2.FONT_HERSHEY_PLAIN, fontScale=1)
523
524
525
            prev_faces = curr_faces
526
            cv2.imwrite(os.path.join(output_dir, filename), img)
527
            # cv2.imshow("img", img)
            # cv2.waitKey(0)
528
     Listing 4: utils.py
  1 import cv2
  2 import os
    import numpy as np
    def make_video(imgs_dir, vid_name, fps):
  5
  6
 7
        Make vid_name.mp4 using images in imgs_dir.
  8
```

```
9
       OUTPUT_DIR = "output/"
10
       if not os.path.isdir(OUTPUT DIR):
11
           os.mkdir(OUTPUT_DIR)
12
       exts = [".jpg", ".png"]
13
       imgs = [img for img in os.listdir(imgs_dir) if img.endswith(tuple(exts))]
       # Sort images by name in ascending order
14
       imgs.sort(key=lambda img: int("".join(filter(str.isdigit, img))))
15
       frame = cv2.imread(os.path.join(imgs_dir, imgs[0]))
16
17
       h, w, _ = frame.shape
18
       # Make sure vid_name does not include a file extension
19
20
       index = vid_name.find(".")
21
       if index !=-1:
22
           vid_name = vid_name[:index]
       file_path = OUTPUT_DIR + vid_name + ".mp4"
23
24
       vid = cv2.VideoWriter(file_path, cv2.VideoWriter_fourcc(*"MP4V"), fps, (w, h))
25
       for img in imgs:
           vid.write(cv2.imread(os.path.join(imgs_dir, img)))
26
27
       vid.release()
28
       print("Video is now in ", file_path)
29
30
   def shuffle(x, y):
31
32
       Shuffle data x and their labels y.
33
34
       assert len(x) == len(y)
35
       idx = np.random.permutation(len(x))
       x, y = np.array(x)[idx], np.array(y)[idx]
36
37
       return x, y
    Listing 5: run.py
 1 import click
 2 from pathlib import Path
 3
 4 @click.group()
 5 def main():
 6
       pass
 7
 8 @main.command()
9 #@click.option("--model_path", "-m", type=Path, default="models/cnn_model.h5")
10 @click.option("--model_path", "-m", type=Path, default="models/svm_model.sav")
11 @click.option("--classification", "-c", default="SVM", help="SVM or NN_SIFT or CNN")
12 def train(**kwargs):
       from face import train_model
13
       train_model(**kwargs)
14
15
16 @main.command()
   \# \ @ \ click.option("--model\_path", "-m", \ type=Path, \ default="models/nn\_model.h5")
17
18 @click.option("--model_path", "-m", type=Path, default="models/svm_model.sav")
19
   def predict(**kwargs):
20
       from face import predict_model
21
       predict_model(**kwargs)
22
23 @main.command()
24 @click.option("--model_path", "-m", type=Path, default="models/svm_model.sav")
25 #@click.option("--model_path", "-m", type=Path, default="models/nn_model.h5")
26 @click.option("--classification", "-c", default="SVM", help="SVM or NN_SIFT or CNN")
27 # Clip 1
28 @click.option("--input_dir", "-i", type=Path, default="original_data/clip_1/")
```

```
29  @click.option("--output_dir", "-o", type=Path, default="output/clip_1_face/")
30  @click.option("--min_size", "-s", default=70, type=int, help="Min size for face
          → detection")
31 # Clip 2
32 #@click.option("--input_dir", "-i", type=Path, default="original_data/clip_2/")
33 #@click.option("--output dir", "-o", type=Path, default="output/clip 2/")
34 #@click.option("--min_size", "-s", default=70, type=int, help="Min size for face detection")
    def face_detection(**kwargs):
35
          from face import face_detection_hsv, face_detection_cascade
36
37
          # face detection hsv(**kwargs)
38
          face_detection_cascade(**kwargs)
39
40 @main.command()
41 # Clip 1
## Cup 1

42 #@click.option("--input_dir", "-i", type=Path, default="original_data/clip_1/")

43 #@click.option("--output_dir", "-o", type=Path, default="output/clip_1_logo/")

44 #@click.option("--logo_path", "-d", type=Path, default="data/clip_1_logo2.png")

45 #@click.option("--min_threshold", "-t", type=float, default=0.87)
46 # Clip 2
# @click.option("--input_dir", "-i", type=Path, default="original_data/clip_2/")
# @click.option("--output_dir", "-o", type=Path, default="output/clip_2/")
49 #@click.option("--logo_path", "-d", type=Path, default="data/clip_2_logo.png")
50 #@click.option("--min_threshold", "-t", type=float, default=0.82)
51 # Clip 3
55 @click.option("--min_threshold", "-t", type=float, default=0.82)
     def logo_detection(**kwargs):
56
          from logo import logo_detection
57
58
          logo_detection(**kwargs)
59
60
     @main.command()
61
     @click.option("--input_dir", "-i", type=Path, default="original_data/clip_3/")
     @click.option("--method", "-m", default="SAD2", help="SAD2 or HD")
@click.option("--k", "-k", default=1, help="k used in threshold")
62
63
64
     def shot_detection(**kwargs):
          from shot import shot_detection
65
66
          shot_detection(**kwargs)
67
68 @main.command()
69 @click.option("--imgs_dir", "-i", default="original_data/clip_1/", type=Path)
70 @click.option("--vid_name", "-v", default="clip_1")
71 @click.option("--fps", "-fps", default=6, type=int, help="Frame per second")
72
     def make_video(**kwargs):
          from utils import make_video
73
74
          make_video(**kwargs)
75
76 @main.command()
77 # Clip 1
78 @click.option("--input_dir", "-i", type=Path, default="original_data/clip_1/")
79 @click.option("--output_dir", "-o", type=Path, default="output/clip_1_shots/")
80 @click.option("--shots", "-s", type=list, default=[22, 156])
81 # Clip 2
82 #@click.option("--input_dir", "-i", type=Path, default="original_data/clip_2/")
# @click.option("--output_dir", "-o", type=Path, default="output/clip_2_shots/")
84 #@click.option("--shots", "-s", type=list, default=[65, 120, 138, 144, 152, 164, 177, 188])
85 # Clip 3
86 #@click.option("--input dir", "-i", type=Path, default="original data/clip 3/")
```

```
87 #@click.option("--output_dir", "-o", type=Path, default="output/clip_3_shots/")
      #@click.option("--shots", "-s", type=list, default=[16, 51, 57, 59, 61, 63, 65, 67, 69, 71, 72, 74, 78, 80, 84, 91.
            \hookrightarrow 95, 102, 165, 187, 260, 2671)
      def add_shot_number(**kwargs):
 89
 90
            from shot import add_shot_number
            add_shot_number(**kwargs)
 91
 92
      @main.command()
 93
 94 # Clip 1
 95 #@click.option("--input_dir", "-i", type=Path, default="output/clip_1_shots/")
 96 #@click.option("--output_dir", "-o", type=Path, default="output/clip_1/")
     #@click.option("--logo_path", "-d", type=Path, default="data/clip_1_logo.png")
     #@click.option("--min_threshold", "-t", type=float, default=0.87)
     # @click.option("—min_inreshola", "-i , type=jtotil, default="models/svm_model.sav")

# @click.option("—model_path", "-m", type=Path, default="models/svm_model.sav")

# @click.option("—classification", "-c", default="SVM", help="SVM or NN_SIFT or CNN")

# @click.option("—vid_name", "-v", default="clip_1")

# @click.option("—fps", "-fps", default=6, type=int, help="Frame per second")

# @click.option("—min_size", "-s", default=70, type=int, help="Min size for face detection")
104 # Clip 2
105 @click.option("--input_dir", "-i", type=Path, default="output/clip_2_shots/")
106 @click.option("--output_dir", "-o", type=Path, default="output/clip_2/")
107 @click.option("--logo_path", "-d", type=Path, default="data/clip_2_logo.png")
108 @click.option("--min_threshold", "-t", type=float, default=0.82)
      @click.option("--model_path", "-m", type=Path, default="models/svm_model.sav")
      @click.option("--classification", "-c", default="SVM", help="SVM or NN_SIFT or CNN")
@click.option("--vid_name", "-v", default="clip_2")
111
      @click.option("--fps", "-fps", default=6, type=int, help="Frame per second")
112
      @click.option("--min_size", "-s", default=70, type=int, help="Min size for face
113
            → detection")
114 # Clip 3
      #@click.option("—-input_dir", "-i", type=Path, default="output/clip_3_shots/")
#@click.option("—-output_dir", "-o", type=Path, default="output/clip_3/")
#@click.option("—-logo_path", "-d", type=Path, default="data/clip_3_logo.png")
115
117
118
      # @click.option("--min_threshold", "-t", type=float, default=0.82)
119
      # @click.option("——model_path", "—m", type=Path, default="models/svm_model.sav")
# @click.option("--classification", "-c", default="SVM", help="SVM or NN_SIFT or CNN")
# @click.option("--vid_name", "-v", default="clip_3")
# @click.option("--fps", "-fps", default=8, type=int, help="Frame per second")
123 #@click.option("--min size", "-s", default=40, type=int, help="Min size for face detection")
124
      def run_all(input_dir, output_dir, logo_path, min_threshold, model_path, \
125
            classification, vid_name, fps, min_size):
            from face import face_detection_cascade
126
127
            from logo import logo_detection
            from utils import make_video
128
            logo_detection(input_dir=input_dir, output_dir=output_dir,\
129
                 logo_path=logo_path, min_threshold=min_threshold)
130
131
            face_detection_cascade(input_dir=output_dir, output_dir=output_dir,\
132
            # face detection cascade(input dir=input dir, output dir=output dir,\
                 model_path=model_path, classification=classification, min_size=min_size)
133
134
            make_video(imgs_dir=output_dir, vid_name=vid_name, fps=fps)
135
       if __name__ == "__main__":
136
137
           main()
```