A1_tdm47_E4

May 5, 2018

1 EECS 531 - A1

1.0.1 Tristan Maidment (tdm47)

1.0.2 Assignment 1

Exercise 4 To determine the ROC curve for the template matching method in Exercise 3, I first need to calculate the True Positive Rate and False Positive Rate, as defined by the lecture slides.

The first step is to import all of the code needed to do feature detection from the previous exercise. This contains all the methods used, and will be used to calculate the different ROC curves with different values of noise added.

```
In [16]: %matplotlib inline
         import cv2
         import math
         from matplotlib import pyplot as plt
         from matplotlib import patches as patches
         import numpy as np
         def correlation(img, kernel):
             width, height, channels = img.shape
             k_width, k_height = kernel.shape
             k_half_width = math.floor(k_width/2)
             k_half_width_2 = math.ceil(k_width/2)
             k_half_height = math.floor(k_height/2)
             k_half_height_2 = math.ceil(k_height/2)
             ret = np.zeros(img.shape)
             img_pad = np.ones((width + k_width + 1 , height + k_width + 1, channels))
             img_pad[k_half_width : -k_half_width_2 - 1,
                     k_half_height : -k_half_height_2 - 1] = img
             for x in range(width):
                 for y in range(height):
                     for c in range(channels):
                         ret[x, y, c] = np.power(np.subtract(img_pad[x: x + k_width,
```

```
y : y + k_height,
                                                             c], kernel), 2).sum()
    return ret/ret.max()
def create_kernel(img):
    patch = cv2.imread(img, 0)
    patch = patch/patch.max()
    width, height = patch.shape
    k_size = np.amax(patch.shape)
    ret = np.ones((k_size, k_size))
    x_diff = math.floor((k_size - width)/2)
    y_diff = math.floor((k_size - height)/2)
    ret[x_diff:k_size-x_diff,y_diff:k_size-y_diff] = patch
    return ret
def threshold(supp, t2):
    ret = np.zeros_like(supp)
    width, height, channels = supp.shape
    k_width, k_height = kernel.shape
    k_half_width = math.ceil(k_width/2)
    k_half_height = math.ceil(k_height/2)
    for x in range(k_half_width,width-k_half_width):
        for y in range(k_half_height,height-k_half_height):
                theta = np.sum(supp[x, y, :])
                if theta < t2:</pre>
                    ret[x, y, :] = 1.
                else:
                    ret[x, y, :] = 0.
    return ret
# draw rectangles
def plot_squares(img, heatmap):
    p_width, p_height, channels = heatmap.shape
    x_offset = math.floor(kernel.shape[0]/2)
    y_offset = math.floor(kernel.shape[1]/2)
    fig,ax = plt.subplots(1)
    ax.imshow(img)
    for x in range(p_width):
        for y in range(p_height):
            if heatmap[x, y, 0] == 1.:
                rect = patches.Rectangle((y - y_offset,x - x_offset),kernel.shape[0],
                ax.add_patch(rect)
    plt.show()
    #fig.savefig("./output.png", dpi=240) # save the image to the working directory
```

The entirety of Example 3 is run here to ensure that everything works after moving the code

over. As expected, all 38 detections are present, with no false positives.

```
In [2]: img = cv2.imread('./characters.png', 1) #load all three channels, due to rendering pro
          if img is not None:
               b,g,r = cv2.split(img)
               img = cv2.merge((r,g,b))
          img = img/img.max()
          kernel = create_kernel('./h.png')
          probability = correlation(img, kernel)
          heatmap = threshold(probability, 0.075 * 3)
          plot_squares(img, heatmap)
                            does seem to work well in the face of these chillenges is:
                            time imman visual system. It makes eminent sense,
                            (herefore, to attempt to understand the strategies (his bio-
                      100
                            logical system employs, as a first step towards eventually
                            translating from into machine-based algorithms. With this
                            objective in mind, we review here 19 important results
                      200
                            regarding face recognition by formans. While these
                            observations do not constitute a comment throny of face
                            recognition in firman vision (we simply do not have all the
                      300 -
                            pieces yet to construct such a favory), they do provide
                            useful fints and constraints for one. We believe that for
                            this reason, they are likely to be useful to computer vision.
                      400
                            researchers in guiding their ongoing efforts. Of course, the
                            success of madiline vision systems is not dependent on a
```

The function roc_data is used the FPR and TPR, as discussed earlier. These will be used to plot the ROC curves.

300

slavis<mark>ta</mark> imitation of their biological counterparts. Insights

into the functioning of the latter serve primarily as potentially fruitful starting points for computational

400

500

600

To determine the false positives (fp), false negatives (fn), true positives (tp), and true negatives (tn), the predicted and real values using the detection matrix from Exerise 3 are used as the baseline. If both values are the same, it is classified as a true positive or true negative respectively. If the values differ, it is classified as a false positive or false negative.

```
In [17]: def roc_data(heatmap, test):
    h_width, h_height, h_channels = heatmap.shape
    fp = 0
    fn = 0
    tp = 0
    tn = 0
    for w in range(h_width):
```

investigations.
100

200

500

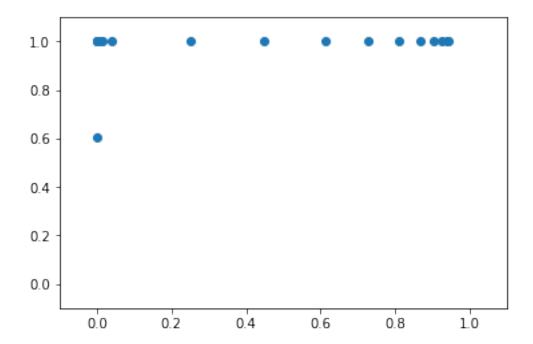
```
for h in range(h_height):
        real = heatmap[w, h, 0]
        pred = test[w, h, 0]
        if real == pred:
            if pred > 0:
                tp += 1
                tn += 1
        else:
            if pred > 0:
                fp += 1
            else:
                fn += 1
#38 0 0 394762
#print(fp, fn, tp, tn)
fpr = fp/(fp + tn)
tpr = tp/(tp + fn)
return fpr, tpr
```

Using the roc_data function, the function create_ROC loops through different detection thresholds and plot the FPR and TPR respectively. The TPR is represented on the Y axis and FPR on the X axis.

```
In [4]: def create_ROC(heatmap, probability_test):
    x_list = []
    y_list = []
    for theta in np.arange(0.05, 0.9, 0.05):
        heatmap_test = threshold(probability_test, theta * 3)
        fpr, tpr = roc_data(heatmap, heatmap_test)
        x_list.append(fpr)
        y_list.append(tpr)
    plt.scatter(x_list, y_list)
    plt.xlim( -0.1, 1.1 )
    plt.ylim( -0.1, 1.1 )
    plt.show()
```

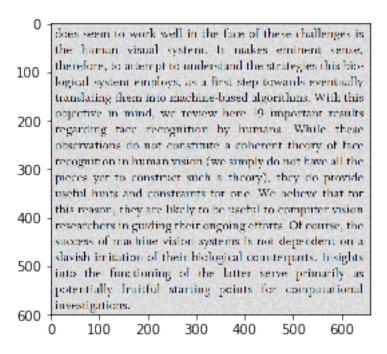
As we can see, the "perfect" threshold that was found in Exercise 3 is following the trend of "ideal behavior", according to the slides. There are no false positives or false negatives.

```
In [5]: create_ROC(heatmap, probability)
```



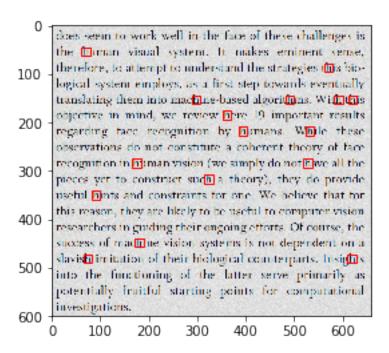
The next step is to introduce noise into the situation to see how it alters the ROC curve. The goal is to increase FP and FN rates. To aid in runtime and simplicity, gray Gaussian noise is added to the image. RGB noise requires an extra three iterations per pixel to check, and would make the runtime very long.

```
In [18]: gray_noise = np.random.normal(0, 0.03, (img.shape[0], img.shape[1])) #mean = 0, std =
    rgb_noise = np.asarray(np.dstack((gray_noise, gray_noise, gray_noise)))
    img_noise = img + rgb_noise
    img_noise = np.absolute(img_noise/img_noise.max())
    plt.imshow(img_noise)
    plt.show()
```



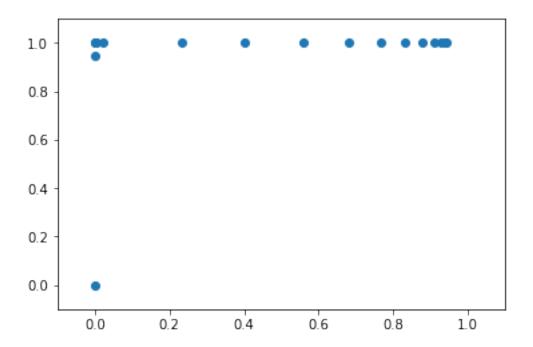
As we can see the number of detections has been lowered to 16. There are no false negatives with the given threshold, but we can determine if some exist given the ROC curve.

```
In [7]: probability_noise = correlation(img_noise, kernel)
    heatmap_noise = threshold(probability_noise, 0.075 * 3)
    plot_squares(img_noise, heatmap_noise)
```



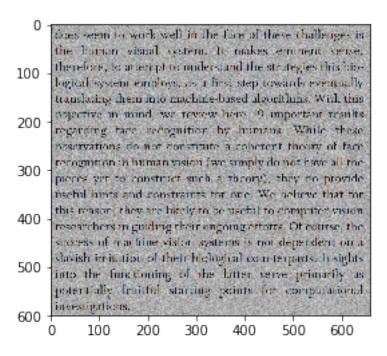
As we can see, there is just one case where false positives occur. False negatives are more common, however.

In [8]: create_ROC(heatmap, probability_noise)

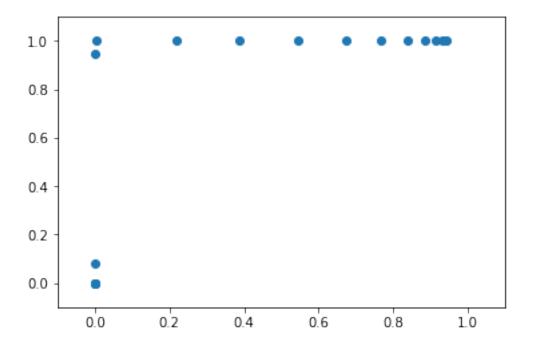


We repeat the test with a higher level of noise.

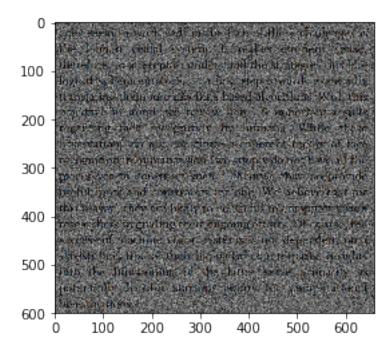
```
In [9]: gray_noise = np.random.normal(0, 0.1, (img.shape[0], img.shape[1]))
    rgb_noise = np.asarray(np.dstack((gray_noise, gray_noise, gray_noise)))
    img_noise = img + rgb_noise
    img_noise = np.absolute(img_noise/img_noise.max())
    plt.imshow(img_noise)
    plt.show()
```



This time, we can visibly see that the curve is no longer at the "ideal", which indicates lower detector performance. It appears to be directly related to amount of noise, as expected.



Let us try an example with a lot of noise. We should expect to see greatly reduced detector performance.



As expected, detector performance has been greatly diminished, and the effects are very noticable on the ROC curve.

