

Image Process and Feature Matching

Dataset in use: <https://susanqq.github.io/UTKFace/> (<https://susanqq.github.io/UTKFace/>)

In-the-wild Faces is used and part-2 is selected for train, part-3 is selected for test set.

```
In [1]: import os
import cv2
import glob
import logging
import numpy as np
import pandas as pd
from tqdm import tqdm
from sklearn import metrics
from skimage import feature
import matplotlib.pyplot as plt
```

```
In [2]: import warnings
warnings.filterwarnings("ignore")
```

```
In [3]: def mkdir(path):
    try:
        if not os.path.exists(path):
            os.makedirs(path)
            logging.debug(f"Folder '{path}' created successfully.")
        else:
            raise RuntimeError(f"Folder '{path}' already exists.")
    except Exception as e:
        logging.error(f"Error creating folder '{path}': {e} You are going to modify an existing file")
        raise
```

```
In [4]: '''
Raporda belirtildiği gibi, girdi olarak net çekilen 1 adet yüz görüntüsü verilir ve çıktı olarak da
SIFT, SURF, distance vb. için tabi ki hazır fonksiyonları kullanacaksınız ama göz bulma, yüz bulma v
hazır kütüphane kullanamazsınız.

ibaresi proje ile alakalı değildir. Projemiz yüz tespiti değil, girdi olarak verilen yüz görüntüsünü
Girdi hatası olması durumunda kodun yine de çalışması amacıyla hazır kütüphane kullanılmaktadır.
'''

def face_extractor(source_dir, dest_dir, ext='jpg'):
    mkdir(dest_dir)
    haar_cascade = cv2.CascadeClassifier(cv2.data.harcascades + 'haarcascade_frontalface_default.xml')

    pimgs = glob.glob(f"{source_dir}/*.{ext}")

    for pimg in tqdm(pimgs, desc="Cropping faces from wild images"):
        img=cv2.imread(pimg)
        gray_img = cv2.cvtColor(img, cv2.COLOR_BGR2GRAY)
        faces_rect = haar_cascade.detectMultiScale(gray_img, 1.2, 5)

        if faces_rect is ():
            logging.debug('No face detected')
            continue

        for (x, y, w, h) in faces_rect:
            cv2.rectangle(img, (x, y), (x+w, y+h), (0, 255, 0), 2)
            cropped_face = gray_img[y:y+h, x:x+w]
            img_name = os.path.splitext(os.path.basename(pimg))[0]
            cv2.imwrite(f'{dest_dir}/{img_name}_cropped.jpg', cropped_face)
```

```
In [5]: def metric_report(actual, predicted):
        acc = metrics.accuracy_score(actual, predicted)
        precision = metrics.precision_score(actual, predicted)
        recall = metrics.recall_score(actual, predicted)
        f1 = metrics.f1_score(actual, predicted)
        return (acc, precision, recall, f1)
```

```
In [6]: source_dir='./utk_train'
        dest_dir='./utk_train_cropped'
        ext='jpg'
```

```
In [7]: test_source_dir='./utk_test'
        test_dest_dir='./utk_test_cropped'
```

```
In [8]: test_dir = './test_images'
```

Local Binary Patterns

```
In [9]: class LocalBinaryPatterns:
        def __init__(self, numPoints, radius):
            self.numPoints = numPoints
            self.radius = radius

        def describe(self, image, eps=1e-7):
            lbp = feature.local_binary_pattern(image, self.numPoints, self.radius, method="uniform")
            (hist, _) = np.histogram(lbp.ravel(),
                                    bins=np.arange(0, self.numPoints + 3),
                                    range=(0, self.numPoints + 2))

            hist = hist.astype("float")
            hist /= (hist.sum() + eps)
            return hist
```

```
In [10]: def single_image_pipeline(img, desc, male_hist_path, female_hist_path):
        fimg = cv2.imread(img, cv2.COLOR_BGR2GRAY)

        lbp_hist = desc.describe(fimg)
        lbp_hist = lbp_hist.astype(np.float32)

        male_lbp_hist = np.load(male_hist_path).astype(np.float32)
        female_lbp_hist = np.load(female_hist_path).astype(np.float32)

        male_distance = cv2.compareHist(lbp_hist, male_lbp_hist, cv2.HISTCMP_INTERSECT)
        female_distance = cv2.compareHist(lbp_hist, female_lbp_hist, cv2.HISTCMP_INTERSECT)
        # HISTCMP_INTERSECT, HISTCMP_CORREL, HISTCMP_BHATTACHARYYA, HISTCMP_HELLINGER, HISTCMP_CHISQR

        return 0 if male_distance >= female_distance else 1
```

```
In [11]: '''
        UTK Train setindeki görüntülerden yüzler elde edilir ve dest_dir üzerine kaydedilir.
        '''
        #face_extractor(source_dir, dest_dir)
```

```
Out[11]: '\nUTK Train setindeki görüntülerden yüzler elde edilir ve dest_dir üzerine kaydedilir.\n'
```

```
In [12]: '''
        Elde edilen yüz görüntüleri cinsiyete göre ayrıştırılır.
        '''
        male_images=[]
        female_images=[]
        for pimg in tqdm(glob.glob(f"{dest_dir}/*.{ext}"), desc="Creating Male and Female Arrays"):
            img = cv2.imread(pimg, cv2.COLOR_BGR2GRAY)
            female_images.append(img) if int(os.path.basename(pimg).split('_')[1]) else male_images.append(
            ...
```

```
Out[12]: '\nmale_images=[]\nfemale_images=[]\nfor pimg in tqdm(glob.glob(f"{dest_dir}/*.{ext}"), desc="Crea
        ting Male and Female Arrays"):\n    img = cv2.imread(pimg, cv2.COLOR_BGR2GRAY)\n    female_images.
        append(img) if int(os.path.basename(pimg).split(\'_\'[1]) else male_images.append(img)\n'
```

```
In [13]: #print(f'Number of male images in trainset: {len(male_images)}\nNumber of female images in trainset
```

```
In [14]: desc=LocalBinaryPatterns(24, 8) # define descriptor instance
```

```
In [15]: '''
Male-Female LBP histogramları elde edilir ve kaydedilir.
'''
'''
male_histograms = [desc.describe(img) for img in tqdm(male_images, desc="Computing LBP histogram - Male")]
female_histograms = [desc.describe(img) for img in tqdm(female_images, desc="Computing LBP histogram - Female")]

np.save('male_lbp_histograms.npy', male_histograms)
np.save('female_lbp_histograms.npy', female_histograms)

male_lbp_hist = np.mean(male_histograms, axis=0)
female_lbp_hist = np.mean(female_histograms, axis=0)

np.save('male_lbp_hist.npy', male_lbp_hist)
np.save('female_lbp_hist.npy', female_lbp_hist)
'''
```

```
Out[15]: '\nmale_histograms = [desc.describe(img) for img in tqdm(male_images, desc="Computing LBP histogram - Male")]\nfemale_histograms = [desc.describe(img) for img in tqdm(female_images, desc="Computing LBP histogram - Female")]\n\nnp.save(\'male_lbp_histograms.npy\', male_histograms)\nnp.save(\'female_lbp_histograms.npy\', female_histograms)\n\nmale_lbp_hist = np.mean(male_histograms, axis=0)\nfemale_lbp_hist = np.mean(female_histograms, axis=0)\n\nnp.save(\'male_lbp_hist.npy\', male_lbp_hist)\nnp.save(\'female_lbp_hist.npy\', female_lbp_hist)\n'
```

```
In [16]: #face_extractor(source_dir, dest_dir)
```

```
In [17]: '''
Test veri setinden rastgele görüntü seçilerek cinsiyet tahmini yapılır.
'''

random_image_pick = np.random.choice(glob.glob(test_dest_dir + f'/*.{ext}'))
pred = single_image_pipeline(random_image_pick, desc, 'male_lbp_hist.npy', 'female_lbp_hist.npy')
```

```
In [18]: '''
Tüm test verisi üzerinde tahminler gerçekleştirilir.
'''

timgs = glob.glob(f"{test_dest_dir}/*.{ext}") #test images

actual, predicted = [], []
for timg in tqdm(timgs, desc="Inference in progress"):
    gender = single_image_pipeline(timg, desc, 'male_lbp_hist.npy', 'female_lbp_hist.npy')
    label = int(os.path.basename(timg).split('_')[1])

    actual.append(label)
    predicted.append(gender)

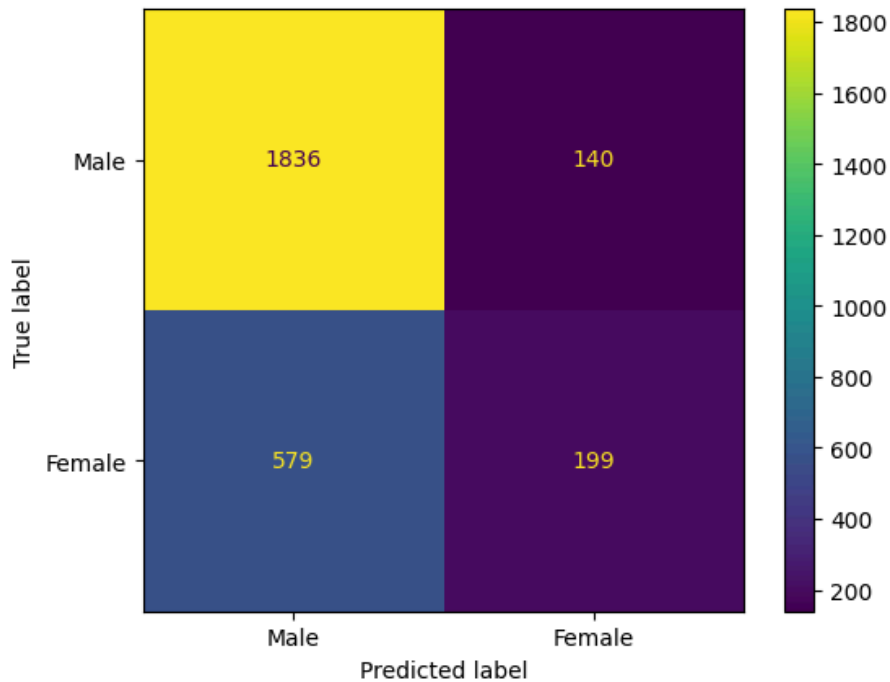
Inference in progress: 100%|████████████████████████████████████████| 2754/2754 [00:55<00:00, 49.79it/s]
```

```
In [19]: metric_report(actual, predicted)
```

```
Out[19]: (0.7389251997095134,
0.5870206489675516,
0.25578406169665807,
0.35631154879140553)
```

```
In [20]: confusion_matrix = metrics.confusion_matrix(actual, predicted)
```

```
In [21]: cm_display = metrics.ConfusionMatrixDisplay(confusion_matrix = confusion_matrix, display_labels = [
cm_display.plot()
plt.show())
```



```
In [22]: female_data = np.sum(actual); male_data = len(actual) - female_data
female_data, male_data
```

```
Out[22]: (778, 1976)
```

```
In [23]: # Asagidaki img_path degiskenine istenen görüntülerin pathi verilir ve lbp ile gender detect edilir
```

```
In [24]: img_path = 'test_images/22_0_0_20170119151204830_cropped.jpg'
actual = int(os.path.basename(img_path).split('_')[1])
```

```
In [25]: pred = single_image_pipeline(img_path, desc, 'male_lbp_hist.npy', 'female_lbp_hist.npy')
```

```
In [26]: if actual == pred:
print('Gender Detected!')
```

Gender Detected!

Scale Invariant Feature Transform (SIFT)

```
In [27]: sift = cv2.SIFT_create()
```

```
In [28]: def flann_match(query_descriptors, train_descriptors):
FLANN_INDEX_KDTREE = 1
index_params = dict(algorithm=FLANN_INDEX_KDTREE, trees=5)
search_params = dict(checks=50)

flann = cv2.FlannBasedMatcher(index_params, search_params)
matches = flann.knnMatch(query_descriptors, np.vstack(train_descriptors), k=2)

# Ratio test as per Lowe's paper
good_matches = []
for m, n in matches:
    if m.distance < 0.7 * n.distance:
        good_matches.append(m)
return len(good_matches)
```

```
In [29]: def extract_descriptors(image, extractor):
        gray = cv2.imread(image, 0)
        keypoints, descriptors = extractor.detectAndCompute(gray, None)
        return descriptors
```

```
In [30]: def single_image_pipeline_sift(img, male_desc_path, female_desc_path):
        descriptor = extract_descriptors(img, sift)

        male_desc = np.load(male_desc_path).astype(np.float32)
        female_desc = np.load(female_desc_path).astype(np.float32)

        male_distance = flann_match(descriptor, male_desc)
        female_distance = flann_match(descriptor, female_desc)

        return 0 if male_distance >= female_distance else 1
```

```
In [31]: '''
        Male-Female SIFT descriptor elde edilir.
        '''

        '''
        male_descriptors = []
        female_descriptors = []

        for img in tqdm(male_images, desc= 'Extracting descriptors - Male'):
            desc = extract_descriptors(img)
            if desc is not None:
                male_descriptors.extend(desc)

        for img in tqdm(female_images, desc='Extracting descriptors - Female'):
            desc = extract_descriptors(img)
            if desc is not None:
                female_descriptors.extend(desc)
        '''
```

```
Out[31]: "\n\nmale_descriptors = []\n\nfemale_descriptors = []\n\n\nfor img in tqdm(male_images, desc= 'Extractin
g descriptors - Male'):\n    desc = extract_descriptors(img)\n    if desc is not None:\n        ma
le_descriptors.extend(desc)\n\n\nfor img in tqdm(female_images, desc='Extracting descriptors - Femal
e'):\n    desc = extract_descriptors(img)\n    if desc is not None:\n        female_descriptors.ex
tend(desc)\n"
```

```
In [32]: '''
        Cikartilan descriptorlar kaydedilir.
        '''

        '''
        male_descriptors = np.array(male_descriptors, dtype=np.float32)
        female_descriptors = np.array(female_descriptors, dtype=np.float32)

        np.save('male_descriptors.npy', male_descriptors)
        np.save('female_descriptors.npy', female_descriptors)
        '''
```

```
Out[32]: "\n\nmale_descriptors = np.array(male_descriptors, dtype=np.float32)\n\nfemale_descriptors = np.array
(female_descriptors, dtype=np.float32)\n\n\nnp.save('male_descriptors.npy', male_descriptors)\n\nnp.sa
ve('female_descriptors.npy', female_descriptors)\n"
```

```
In [33]: male_descriptors = np.load('male_descriptors.npy')
        female_descriptors = np.load('female_descriptors.npy')
```

```
In [34]: './utk_test_cropped/'
```

```
Out[34]: './utk_test_cropped/'
```

