Project

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EE475 Group Project: Machine Learning Based Face Mask Recognition

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As the COVID-19 has brought great disaster to human beings, personal protection has become particularly important. For the purpose of controlling the spread of the epidemic, every one of us has the obligation and responsibility to wear masks. The objective of our project is to proposed a system that can monitor people's mask wearing status (Correct, Incorrect and No mask). In our project, we reduce the dimension of input data space by using pre-processing methods: convert into hog image, Histogram of Oriented Gradients algorithm and Canny edge detector algorithm. We firstly implement linear model from scratch, then implement SVM, Decision Tree and Random Forest using scikit learn library.

```
[]: from lib_fun import *
     import numpy as np
     import matplotlib.pyplot as plt
     import sklearn
     import sklearn.svm as svm
     import sklearn.tree as tree
     import sklearn.ensemble as ensemble
```

```
[]: # Load the original face mask data with pixel of 20*20, 50*50 and 100*100
    # We visulaize these image of Correct, Incorrect and No mask with each
     \rightarrow resolution.
    data20, labels20, data50, labels50, data100, labels100 = loads('.../Data/Pixel20/
```

X shape: (4559, 20, 20, 3), Y shape: (4559,) X shape: (4559, 50, 50, 3), Y shape: (4559,) X shape: (4559, 100, 100, 3), Y shape: (4559,)



















[]: # In order to reduce the input dimension and keep critical information, well →transfer the RGB image to gray image data20_gray, data50_gray, data100_gray = RGBtoGrays(data20, data50, data100)



















[]: # Using Histogram of Oriented Gradients to extract features from RGB image # Compute 8 direction in each 2*2 pixel





















[]: # Using Canny edge detector to extract features from gray image data20_edge, data50_edge, data100_edge = GraytoCanny(data20_gray, data50_gray, \rightarrow data100_gray, sigma=[1,3,5])



















Firstly We implement one vs rest SVM model. SVM maps training examples to points in space so as to maximise the width of the gap between the two categories. New examples are then mapped into that same space and predicted to belong to a category based on which side of the gap they fall. As we are using a soft margin SVM, we set the penalty term C as 1. A low C makes the decision surface smooth, while a high C aims at classifying all training examples correctly. And we are using a kernel function to map the input features to a higher dimension space. In this experiment, we compare the linear, polynomial and rbf kernel. We choose gray and hog data as out training data in this experiment.

[]: # SVM with linear kernel, using gray images as data. cf1, cf2, cf3 = train models(svm.SVC(C=1, kernel='linear', cache_size=4000), __ →data20 gray, data50 gray, data100 gray, labels20, labels50, labels100)

accuracy: 0.7905701754385965 accuracy: 0.7324561403508771 accuracy: 0.7214912280701754 It is not difficult to find that with the increase of image size, the accuracy of our model even decreases, which is very counterintuitive.

```
[]: # SVM with linear kernel, using hog images as data.

cf1, cf2, cf3 = train_models(svm.SVC(C=1, kernel='linear', cache_size=4000),

data20_hog, data50_hog, data100_hog, labels20, labels50, labels100)
```

accuracy: 0.8125

accuracy: 0.7521929824561403 accuracy: 0.8004385964912281

Although the classification accuracy was slightly improved after the HoG feature was used, the overall performance of the model was still unsatisfactory, which indicates that Linear kernel has reached its performance limit, and we must find an alternative methodology to replace it.

```
[]: # SVM with 3 degree polynomial kernel, using gray images as data.

cf1, cf2, cf3 = train_models(svm.SVC(C=1, kernel='poly', degree=3,

cache_size=4000), data20_gray, data50_gray, data100_gray, labels20,

labels50, labels100)
```

accuracy: 0.868421052631579

accuracy: 0.875

accuracy: 0.8739035087719298

Here we introduced the 3 degree class polynomial kernel. As can be seen from the figure below, the classification accuracy of all categories has been significantly improved.

accuracy: 0.8530701754385965 accuracy: 0.8464912280701754 accuracy: 0.8037280701754386

Different from the previous control groups, when we replaced gray graphics with Hog features, the classification accuracy did not significantly improve

```
[]: # SVM with rbf kernel, using gray images as data. Beta is set to 1 / n_features. cf1, cf2, cf3 = train_models(svm.SVC(C=1, kernel='rbf', gamma='auto', u → cache_size=4000), data20_gray, data50_gray, data100_gray, labels20, u → labels50, labels100)
```

accuracy: 0.8859649122807017 accuracy: 0.8914473684210527 accuracy: 0.8903508771929824

[]: # SVM with rbf kernel, using hog images as data. Beta is set to 1 / n_features.

```
cf1, cf2, cf3 = train_models(svm.SVC(C=1, kernel='rbf', gamma='auto', u cache_size=4000), data20_hog, data50_hog, data100_hog, labels20, labels50, u labels100)
```

accuracy: 0.8530701754385965 accuracy: 0.8728070175438597 accuracy: 0.8607456140350878

Finally, we test the effect of RBF kernel. Compared with linear kernel, both RBF kernel and POLY kernel can significantly improve the effect of the model. Meanwhile, when gray Scale image (50×50) was used as the input, we obtained the highest accuracy of the SVM model of 89.1%

In this experiment, we can see that the result of rbf kernel is better than polynomial, which is better than linear kernel. The hog method lead to worse results using each kernel function. The reason is that models need more information other than edge information to make right decision. The most interesting thing we can see is that, the highest resolution of image doesn't produce the best results. The reason may be that it is harder for SVM to split high dimension data space.

The next model we choose is decision tree. Decision Trees (DTs) are a non-parametric supervised learning method used for classification and regression. The goal is to create a model that predicts the value of a target variable by learning simple decision rules inferred from the data features. A tree can be seen as a piecewise constant approximation. In our setting, the maximum depth of the tree is nor specified, nodes are expanded until all leaves are pure or until all leaves contain less than 2 samples. And we compare the results of using gray, hog and canny dataset.

```
[]: # Decision Tree on gray image.
cf1, cf2, cf3 = train_models(tree.DecisionTreeClassifier(), data20_gray,

data50_gray, data100_gray, labels20, labels50, labels100)
```

accuracy: 0.7817982456140351 accuracy: 0.8135964912280702 accuracy: 0.7905701754385965

```
[]: # Decision Tree on hog image.
cf1, cf2, cf3 = train_models(tree.DecisionTreeClassifier(), data20_hog,

data50_hog, data100_hog, labels20, labels50, labels100)
```

accuracy: 0.7719298245614035 accuracy: 0.7631578947368421 accuracy: 0.7182017543859649

```
[]: # Decision Tree on canny image.
cf1, cf2, cf3 = train_models(tree.DecisionTreeClassifier(), data20_edge,
data50_edge, data100_edge, labels20, labels50, labels100)
```

accuracy: 0.5032894736842105 accuracy: 0.5646929824561403 accuracy: 0.48793859649122806 We can see that hog extractor has bad influence on the results, while canny extractor has very bad influence on the results. And the accuarcy of a single tree is worse than SVM with rbf kernel.

Then, we try the random forest algorithm. The Random forest is a special estimator that fits a certain number of single trees on various sub-samples of the dataset and uses averaging to improve the classification accuracy and control the situation of over-fitting. In our setting, The maximum depth of the tree is nor specified, nodes are expanded until all leaves are pure or until all leaves contain less than 2 samples. And we compare the results of 5, 10 and 50 subtrees on only gray dataset.

```
[]: # Random Forest using 5 subtrees on gray image.
cf1, cf2, cf3 = train_models(ensemble.RandomForestClassifier(n_estimators= 5),

data20_gray, data50_gray, data100_gray, labels20, labels50, labels100)
```

accuracy: 0.8267543859649122 accuracy: 0.8388157894736842 accuracy: 0.856359649122807

```
[]: # Random Forest using 10 subtrees on gray image.
cf1, cf2, cf3 = train_models(ensemble.RandomForestClassifier(n_estimators= 10),

data20_gray, data50_gray, data100_gray, labels20, labels50, labels100)
```

accuracy: 0.8793859649122807 accuracy: 0.8706140350877193 accuracy: 0.8760964912280702

```
[]: # Random Forest using 50 subtrees on gray image.
cf1, cf2, cf3 = train_models(ensemble.RandomForestClassifier(n_estimators= 50),

data20_gray, data50_gray, data100_gray, labels20, labels50, labels100)
```

accuracy: 0.8969298245614035 accuracy: 0.9024122807017544 accuracy: 0.9067982456140351

Since the growth of each tree in the random forest has grow to the maximum extent and has a certain degree of randomness, the performance and robustness of the overall model are largely improved when we integrate them together. We can see that random forest has higher accuarcy than a single tree, and the more subtress we use, the higher accuarcy we can gain.

Finally, instead of using scikit learn package, we implement a linear model from scratch. Linear classifier is a very common and efficient classification algorithm in machine learning. In this project we will compare its results as a baseline with the other three classification algorithms. Its result can be illustrated as below: