SVM_Models_VGG19

March 28, 2019

1 Import Library

In [1]:
```
import os
import pandas as pd
import numpy as np
import pickle
import time
# Machine Learning Algorithms
from sklearn.linear_model import LogisticRegression
from sklearn.svm import LinearSVC
from sklearn.metrics import confusion_matrix
from sklearn.pipeline import make_pipeline
from sklearn.model_selection import GridSearchCV
from sklearn.metrics import confusion_matrix
from sklearn.metrics import precision_score,recall_score,f1_score
from sklearn.metrics import precision_recall_fscore_support
from sklearn.model_selection import validation_curve,learning_curve
import matplotlib.pyplot as plt
from sklearn.metrics import precision_recall_fscore_support
from sklearn.svm import SVC
from joblib import dump,load
```

2 Loading Paths

In [2]:
```
train_path="A:\\Projects\\Major Project\\Extracted CNN Features\\VGG19\\train"
test_path="A:\\Projects\\Major Project\\Extracted CNN Features\\VGG19\\test"
```

In [3]:
```
param_range = [0.001, 0.01, 0.1, 1.0, 10.0,]

param_grid = [{'svc__C': param_range,
               'svc__kernel': ['linear']},
              {'svc__C': param_range,
               'svc__gamma': ['auto'],
               'svc__kernel': ['rbf']}]
pipe_svc = make_pipeline(SVC(random_state=1))
```

In [4]:
```
# Training Paths
X_train=np.load(train_path+"\\data_cnn_VGG19_train.npy")
```

```
Y_train=np.load(train_path+"\\data_mag_VGG19_train.npy")
# Cancer class
cancerclass_train=np.load(train_path+"\\data_cancerclass_VGG19_train.npy")
# Cancer type
cancertype_train=np.load(train_path+"\\data_cancertype_VGG19_train.npy")
# Testing Paths
X_test=np.load(test_path+"\\data_cnn_VGG19_test.npy")
Y_test=np.load(test_path+"\\data_mag_VGG19_test.npy")
# Cancer class
cancerclass_test=np.load(test_path+"\\data_cancerclass_VGG19_test.npy")
# Cancer type
cancertype_test=np.load(test_path+"\\data_cancertype_VGG19_test.npy")
```

2.1 Magnification classification

```
In [5]: start_time = time.clock()


        param_grid = [{'svc__C': param_range,
                       'svc__kernel': ['linear']},
                      {'svc__C': param_range,
                       'svc__gamma': ['auto'],
                       'svc__kernel': ['rbf']}]

        gs = GridSearchCV(estimator=pipe_svc,
                          param_grid=param_grid,
                          scoring='accuracy',
                          cv=10,
                          n_jobs=-1)
        gs = gs.fit(X_train, Y_train)
        print(gs.best_score_)
        print(gs.best_params_)
        print(time.clock() - start_time, "seconds")



        clf = gs.best_estimator_
        clf.fit(X_train, Y_train)
        print('Test accuracy: %.3f' % clf.score(X_test, Y_test))
```

```
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche
  """Entry point for launching an IPython kernel.


0.8354430379746836
{'svc__C': 0.001, 'svc__kernel': 'linear'}
20291.82806508 seconds
```

```
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche


Test accuracy: 0.895


In [6]: clf = gs.best_estimator_
        clf.fit(X_train, Y_train)
        print('Test accuracy: %.3f' % clf.score(X_test, Y_test))
        print(time.clock() - start_time, "seconds")

Test accuracy: 0.895
20456.541849765 seconds


c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche
  after removing the cwd from sys.path.


In [7]: dump(clf,'models/SVM/SVM_Models_VGG19_Magnification.joblib')

Out[7]: ['models/SVM/SVM_Models_VGG19_Magnification.joblib']

In [8]: pred=clf.predict(X_test)

In [9]: con=confusion_matrix(Y_test,pred)

In [10]: print(con)

[[378  19   0   0]
 [ 19 336  36   1]
 [  2  28 337  27]
 [  2   2  30 363]]


In [11]: precision_recall_fscore_support(Y_test,pred)

Out[11]: (array([0.94264339, 0.87272727, 0.83622829, 0.92838875]),
          array([0.95214106, 0.85714286, 0.85532995, 0.91435768]),
          array([0.94736842, 0.86486486, 0.84567127, 0.9213198 ]),
          array([397, 392, 394, 397], dtype=int64))
```

2.2 CancerClass Magnification Classification

```
In [12]: Y_train_40=[]
         X_train_40=[]

         Y_train_100=[]
         X_train_100=[]
```

```
Y_train_200=[]
X_train_200=[]

Y_train_400=[]
X_train_400=[]

for i in range(0,len(Y_train)):
    if(Y_train[i]==40):
        Y_train_40.append(cancerclass_train[i])
        X_train_40.append(X_train[i])
    if(Y_train[i]==100):
        Y_train_100.append(cancerclass_train[i])
        X_train_100.append(X_train[i])
    if(Y_train[i]==200):
        Y_train_200.append(cancerclass_train[i])
        X_train_200.append(X_train[i])
    if(Y_train[i]==400):
        Y_train_400.append(cancerclass_train[i])
        X_train_400.append(X_train[i])

X_train_40=np.array(X_train_40)
X_train_100=np.array(X_train_100)
X_train_200=np.array(X_train_200)
X_train_400=np.array(X_train_400)
Y_train_40=np.array(Y_train_40)
Y_train_100=np.array(Y_train_100)
Y_train_200=np.array(Y_train_200)
Y_train_400=np.array(Y_train_400)
print(Y_train_40.size)

Y_test_40=[]
X_test_40=[]

Y_test_100=[]
X_test_100=[]

Y_test_200=[]
X_test_200=[]

Y_test_400=[]
X_test_400=[]

for i in range(0,len(Y_test)):
    if(Y_test[i]==40):
        Y_test_40.append(cancerclass_test[i])
        X_test_40.append(X_test[i])
    if(Y_test[i]==100):
        Y_test_100.append(cancerclass_test[i])
```

```
            X_test_100.append(X_test[i])
        if(Y_test[i]==200):
            Y_test_200.append(cancerclass_test[i])
            X_test_200.append(X_test[i])
        if(Y_test[i]==400):
            Y_test_400.append(cancerclass_test[i])
            X_test_400.append(X_test[i])

X_test_40=np.array(X_test_40)
X_test_100=np.array(X_test_100)
X_test_200=np.array(X_test_200)
X_test_400=np.array(X_test_400)
Y_test_40=np.array(Y_test_40)
Y_test_100=np.array(Y_test_100)
Y_test_200=np.array(Y_test_200)
Y_test_400=np.array(Y_test_400)
```

```
1596
```

2.3 CancerClass Magnification Classification-40

```
In [13]: start_time = time.clock()
         pipe_svc = make_pipeline(SVC(random_state=1))

         param_range = [0.001, 0.01, 0.1, 1.0, 10.0, 100.0]

         param_grid = [{'svc__C': param_range,
                        'svc__kernel': ['linear']},
                       {'svc__C': param_range,
                        'svc__gamma': ['auto'],
                        'svc__kernel': ['rbf']}]

         gs = GridSearchCV(estimator=pipe_svc,
                           param_grid=param_grid,
                           scoring='accuracy',
                           cv=10,
                           n_jobs=-1)
         gs = gs.fit(X_train_40, Y_train_40)
         print(gs.best_score_)
         print(gs.best_params_)
         print(time.clock() - start_time, "seconds")


         clf = gs.best_estimator_
         clf.fit(X_train_40, Y_train_40)
         print('Test accuracy: %.3f' % clf.score(X_test_40, Y_test_40))
```

```
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche
```

```
  """Entry point for launching an IPython kernel.


0.8402255639097744
{'svc__C': 0.001, 'svc__kernel': 'linear'}
553.6712695530005 seconds


c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche


Test accuracy: 0.773


In [14]: dump(clf,'models/SVM/SVM_Models_VGG19_Magnification_40.joblib')

Out[14]: ['models/SVM/SVM_Models_VGG19_Magnification_40.joblib']
```

2.4 CancerClass Magnification Classification-100

```
In [15]: start_time = time.clock()
         pipe_svc = make_pipeline(SVC(random_state=1))

         param_range = [0.001, 0.01, 0.1, 1.0, 10.0, 100.0]

         param_grid = [{'svc__C': param_range,
                        'svc__kernel': ['linear']},
                       {'svc__C': param_range,
                        'svc__gamma': ['auto'],
                        'svc__kernel': ['rbf']}]

         gs = GridSearchCV(estimator=pipe_svc,
                           param_grid=param_grid,
                           scoring='accuracy',
                           cv=10,
                           n_jobs=-1)
         gs = gs.fit(X_train_100, Y_train_100)
         print(gs.best_score_)
         print(gs.best_params_)
         print(time.clock() - start_time, "seconds")


         clf = gs.best_estimator_
         clf.fit(X_train_100, Y_train_100)
         print('Test accuracy: %.3f' % clf.score(X_test_100, Y_test_100))

c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche
  """Entry point for launching an IPython kernel.
```

```
0.8298755186721992
{'svc__C': 0.001, 'svc__kernel': 'linear'}
666.2640964229977 seconds


c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche


Test accuracy: 0.791


In [16]: print(gs.best_estimator_)

Pipeline(memory=None,
     steps=[('svc', SVC(C=0.001, cache_size=200, class_weight=None, coef0=0.0,
  decision_function_shape='ovr', degree=3, gamma='auto_deprecated',
  kernel='linear', max_iter=-1, probability=False, random_state=1,
  shrinking=True, tol=0.001, verbose=False))])


In [17]: dump(clf,'models/SVM/SVM_Models_VGG19_Magnification_100.joblib')

Out[17]: ['models/SVM/SVM_Models_VGG19_Magnification_100.joblib']
```

2.5 CancerClass Magnification Classification-200

```
In [18]: tart_time = time.clock()
         pipe_svc = make_pipeline(SVC(random_state=1))

         param_range = [0.001, 0.01, 0.1, 1.0, 10.0, 100.0]

         param_grid = [{'svc__C': param_range,
                        'svc__kernel': ['linear']},
                       {'svc__C': param_range,
                        'svc__gamma': ['auto'],
                        'svc__kernel': ['rbf']}]

         gs = GridSearchCV(estimator=pipe_svc,
                           param_grid=param_grid,
                           scoring='accuracy',
                           cv=10,
                           n_jobs=-1)
         gs = gs.fit(X_train_200, Y_train_200)
         print(gs.best_score_)
         print(gs.best_params_)
         print(time.clock() - start_time, "seconds")


         clf = gs.best_estimator_
         clf.fit(X_train_200, Y_train_200)
         print('Test accuracy: %.3f' % clf.score(X_test_200, Y_test_200))
```

```
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche
  """Entry point for launching an IPython kernel.


0.8695114409400123
{'svc__C': 0.01, 'svc__kernel': 'linear'}
1274.3814800069995 seconds


c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche


Test accuracy: 0.830


In [19]: dump(clf,'models/SVM/SVM_Models_VGG19_Magnification_200.joblib')

Out[19]: ['models/SVM/SVM_Models_VGG19_Magnification_200.joblib']
```

2.6 CancerClass Magnification Classification-400

```
In [20]: tart_time = time.clock()
         pipe_svc = make_pipeline(SVC(random_state=1))



         gs = GridSearchCV(estimator=pipe_svc,
                           param_grid=param_grid,
                           scoring='accuracy',
                           cv=10,
                           n_jobs=-1)
         gs = gs.fit(X_train_200, Y_train_200)
         print(gs.best_score_)
         print(gs.best_params_)
         print(time.clock() - start_time, "seconds")


         clf = gs.best_estimator_
         clf.fit(X_train_200, Y_train_200)
         print('Test accuracy: %.3f' % clf.score(X_test_400, Y_test_400))

c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche
  """Entry point for launching an IPython kernel.


0.8695114409400123
{'svc__C': 0.01, 'svc__kernel': 'linear'}
1880.8733263979993 seconds
```

```
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche
```

```
Test accuracy: 0.788
```

```
In [21]: dump(clf,'models/SVM/SVM_Models_VGG19_Magnification_400.joblib')
```

```
Out[21]: ['models/SVM/SVM_Models_VGG19_Magnification_400.joblib']
```

2.7 Benign Sub-Classification Using Cancer Classification

```
In [22]: Y_train_1=[]
         X_train_1=[]

         for i in range(0,len(Y_train)):
             if(cancerclass_train[i]==1):
                 Y_train_1.append(cancertype_train[i])
                 X_train_1.append(X_train[i])

         X_train_1=np.array(X_train_1)
         Y_train_1=np.array(Y_train_1)
         print(Y_train_1.size)

         Y_test_1=[]
         X_test_1=[]

         for i in range(0,len(Y_test)):
             if(cancerclass_test[i]==1):
                 Y_test_1.append(cancertype_test[i])
                 X_test_1.append(X_test[i])

         X_test_1=np.array(X_test_1)
         Y_test_1=np.array(Y_test_1)
```

```
1683
```

```
In [ ]:
```

```
In [23]: classes=[11,12,13,14]
```

```
In [ ]:
```

```
In [24]: from sklearn.utils.class_weight import compute_class_weight
```

```
In [25]: class_weight=compute_class_weight(class_weight='balanced', classes=classes,y=Y_train_
```

```
In [26]: print(class_weight)
```

```
[1.66964286 0.51752768 1.67629482 1.14645777]
```

```
In [27]: print(np.unique(Y_train_1))
```

```
[11 12 13 14]
```

```
In [28]: print(len(X_train_1))
```

```
1683
```

```
In [29]: print(len(Y_test_1))
```

```
792
```

```
In [30]: d = dict(enumerate(class_weight, 1))
```

```
In [31]: print(d)
```

```
{1: 1.6696428571428572, 2: 0.5175276752767528, 3: 1.6762948207171315, 4: 1.146457765667575}
```

```
In [32]: d1={1:11,2:12,3:13,4:14}
```

```
In [33]: d=dict((d1[key], value) for (key, value) in d.items())
```

```
In [34]: d
```

```
Out[34]: {11: 1.6696428571428572,
          12: 0.5175276752767528,
          13: 1.6762948207171315,
          14: 1.146457765667575}
```

```
In [ ]:
```

```
In [ ]:
```

```
In [35]: pipe_svc = make_pipeline(SVC(random_state=1,class_weight=d))
         gs3=GridSearchCV(estimator=pipe_svc,
                          param_grid=param_grid,
                          scoring='accuracy',
                          cv=10,
                          n_jobs=-1)
         start_time = time.clock()
         #Training of Model
         gs3.fit(X_train_1,Y_train_1)
         print(time.clock() - start_time, "seconds")

         print(gs3.best_score_)
         print(gs3.best_params_)
```

```
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche
  import sys
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\sklearn\model_sel
  DeprecationWarning)


1052.7684747269996 seconds
0.6868686868686869
{'svc__C': 0.01, 'svc__kernel': 'linear'}


c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche
  # Remove the CWD from sys.path while we load stuff.


In [36]: clf4=gs3.best_estimator_
         clf4.fit(X_train_1,Y_train_1)
         print(clf4.score(X_test_1,Y_test_1))

0.43434343434343436


In [37]: dump(clf4,'models/SVM/SVM_Models_VGG19_CancerType_Benign.joblib')

Out[37]: ['models/SVM/SVM_Models_VGG19_CancerType_Benign.joblib']

In [ ]:
```

2.8 Malignant Sub-Classification Using Cancer Classification

```
In [38]: Y_train_2=[]
         X_train_2=[]

         for i in range(0,len(Y_train)):
             if(cancerclass_train[i]==2):
                 Y_train_2.append(cancertype_train[i])
                 X_train_2.append(X_train[i])

         X_train_2=np.array(X_train_2)
         Y_train_2=np.array(Y_train_2)
         print(Y_train_2.size)

         Y_test_2=[]
         X_test_2=[]

         for i in range(0,len(Y_test)):
             if(cancerclass_test[i]==2):
                 Y_test_2.append(cancertype_test[i])
                 X_test_2.append(X_test[i])
```

```
        X_test_2=np.array(X_test_2)
        Y_test_2=np.array(Y_test_2)
```

```
4637
```

```
In [39]: classes=[21,22,23,24]
```

```
In [40]: from sklearn.utils.class_weight import compute_class_weight
```

```
In [41]: class_weight=compute_class_weight(class_weight='balanced', classes=classes,y=Y_train_2
```

```
In [42]: print(class_weight)
```

```
[0.35669231 2.72764706 1.96150592 3.12466307]
```

```
In [43]: print(np.unique(Y_train_2))
```

```
[21 22 23 24]
```

```
In [44]: print(len(X_train_2))
```

```
4637
```

```
In [45]: print(len(Y_test_2))
```

```
788
```

```
In [46]: d = dict(enumerate(class_weight, 1))
```

```
In [47]: print(d)
```

```
{1: 0.3566923076923077, 2: 2.7276470588235293, 3: 1.9615059221658206, 4: 3.1246630727762805}
```

```
In [48]: d1={1:21,2:22,3:23,4:24}
```

```
In [49]: d=dict((d1[key], value) for (key, value) in d.items())
```

```
In [50]: d
```

```
Out[50]: {21: 0.3566923076923077,
          22: 2.7276470588235293,
          23: 1.9615059221658206,
          24: 3.1246630727762805}
```

```
In [51]: pipe_svc = make_pipeline(SVC(random_state=1,class_weight=d))
         gs3=GridSearchCV(estimator=pipe_svc,
                          param_grid=param_grid,
                          scoring='accuracy',
                          cv=10,
                          n_jobs=-1)
         start_time = time.clock()
         #Training of Model
         gs3.fit(X_train_2,Y_train_2)
         print(time.clock() - start_time, "seconds")

         print(gs3.best_score_)
         print(gs3.best_params_)
```

```
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche
  import sys
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\sklearn\model_se
  DeprecationWarning)


9503.054898564998 seconds
0.7170584429588096
{'svc__C': 10.0, 'svc__gamma': 'auto', 'svc__kernel': 'rbf'}


c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launche
  # Remove the CWD from sys.path while we load stuff.
```

```
In [52]: clf4=gs3.best_estimator_
         clf4.fit(X_train_2,Y_train_2)
         print(clf4.score(X_test_2,Y_test_2))
```

```
0.2817258883248731
```

```
In [53]: pred=clf4.predict(X_test_2)
```

```
In [54]: precision_recall_fscore_support(Y_test_2,pred)
```

```
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\sklearn\metrics\c
  'precision', 'predicted', average, warn_for)


Out[54]: (array([0.25955204, 0.94117647, 0.75      , 0.        ]),
          array([0.985, 0.08 , 0.045, 0.   ]),
          array([0.41084463, 0.14746544, 0.08490566, 0.        ]),
          array([200, 200, 200, 188], dtype=int64))
```

```
In [55]: confusion_matrix(Y_test_2,pred)
```

```
Out[55]: array([[197,   1,   2,   0],
                [183,  16,   1,   0],
                [191,   0,   9,   0],
                [188,   0,   0,   0]], dtype=int64)

In [56]: dump(clf4,'models/SVM/SVM_models_VGG19_CancerType_Malignant.joblib')

Out[56]: ['models/SVM/SVM_models_VGG19_CancerType_Malignant.joblib']

In [ ]:
```