# SVM\_Models\_VGG16

March 29, 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

```
Y_train=np.load(train_path+"\\data_mag_VGG16_train.npy")
# Cancer class
cancerclass_train=np.load(train_path+"\\data_cancerclass_VGG16_train.npy")
# Cancer type
cancertype_train=np.load(train_path+"\\data_cancertype_VGG16_train.npy")
# Testing Paths
X_test=np.load(test_path+"\\data_cnn_VGG16_test.npy")
Y_test=np.load(test_path+"\\data_mag_VGG16_test.npy")
# Cancer class
cancerclass_test=np.load(test_path+"\\data_cancerclass_VGG16_test.npy")
# Cancer type
cancertype_test=np.load(test_path+"\\data_cancertype_VGG16_test.npy")
```

#### 2.1 Magnification classification

```
In [18]: 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\_launch """Entry point for launching an IPython kernel.

```
0.8313291139240506
{'svc_C': 0.001, 'svc_kernel': 'linear'}
```

```
NameError
                                                  Traceback (most recent call last)
        <ipython-input-18-685885a6b1f9> in <module>
        24 clf = gs.best_estimator_
   ---> 25 clf.fit(X_train, y_train)
         26 print('Test accuracy: %.3f' % clf.score(X_test, y_test))
         27 print(time.clock() - start_time, "seconds")
        NameError: name 'y_train' is not defined
In [19]: 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: 0.902
28811.304737958 seconds
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launch
  after removing the cwd from sys.path.
In [20]: pred=clf.predict(X_test)
In [21]: con=confusion_matrix(Y_test,pred)
In [22]: print(con)
[[375 22
           0
                07
 [ 15 347 30
                07
 [ 5 18 344 27]
 [ 1 0 37 359]]
In [23]: precision_recall_fscore_support(Y_test,pred)
Out[23]: (array([0.9469697, 0.89664083, 0.83698297, 0.93005181]),
          array([0.94458438, 0.88520408, 0.87309645, 0.90428212]),
          array([0.94577554, 0.89088575, 0.85465839, 0.91698595]),
          array([397, 392, 394, 397], dtype=int64))
```

## 2.2 CancerClass Magnification Classification

```
In [6]: 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)
```

#### 2.3 CancerClass Magnification Classification-40

1596

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

0.8596491228070176
{'svc__C': 0.001, 'svc__kernel': 'linear'}
632.631926201 seconds
```

Test accuracy: 0.786

#### 2.4 CancerClass Magnification Classification-100

print(gs.best\_params\_)

```
In [7]: 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_launch
  """Entry point for launching an IPython kernel.
0.8601066982809722
{'svc C': 0.01, 'svc kernel': 'linear'}
807.2974345060001 seconds
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launch
Test accuracy: 0.852
In [8]: print(gs.best_estimator_)
Pipeline (memory=None,
     steps=[('svc', SVC(C=0.01, 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))])
2.5 CancerClass Magnification Classification-200
In [9]: 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\_launch """Entry point for launching an IPython kernel.

```
0.8670377241805813
{'svc__C': 0.001, 'svc__kernel': 'linear'}
1584.021985761 seconds
```

Test accuracy: 0.820

### 2.6 CancerClass Magnification Classification-400

c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel\_launchers' c:\users\karan gupta\appdata\python\python37\lib\site-packages\python\python37\lib\site-packages\ipykernel\_launchers' c:\users\karan gupta\appdata\python37\lib\site-packages\python\python37\lib\site-packages\python\python37\lib\site-packages\python\python37\lib\site-packages\python\python37\lib\site-packages\python\python37\lib\site-packages\python\python37\lib\site-packages\python\python37\lib\site-packages\python\python37\lib\site-packages\python\python37\lib\site-packages\python\python37\lib\site-packages\python\python37\lib\site-packages\python\python37\lib\site-packages\python\python37\lib\site-packages\python37\lib\site-packages\python37\lib\site-packages\python37\lib\site-packages\python37\lib\site-packages\python37\lib\site-packages\python37

```
0.8549295774647887
{'svc__C': 0.001, 'svc__kernel': 'linear'}
536.3881946619999 seconds
```

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

Test accuracy: 0.796

In [ ]: Y\_train\_1=[]

### 2.7 Benign Sub-Classification Using Cancer Classification

```
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)
In []:
In [51]: classes=[11,12,13,14]
In [42]:
[1.]
In [52]: from sklearn.utils.class_weight import compute_class_weight
In [53]: class_weight=compute_class_weight(class_weight='balanced', classes=classes,y=Y_train_
In [54]: print(class_weight)
[1.66964286 0.51752768 1.67629482 1.14645777]
```

```
In [55]: print(np.unique(Y_train_1))
[11 12 13 14]
In [56]: print(len(X_train_1))
1683
In [57]: print(len(Y_test_1))
792
In [58]: d = dict(enumerate(class_weight, 1))
In [59]: print(d)
{1: 1.6696428571428572, 2: 0.5175276752767528, 3: 1.6762948207171315, 4: 1.146457765667575}
In [60]: d1={1:11,2:12,3:13,4:14}
In [61]: d=dict((d1[key], value) for (key, value) in d.items())
In [62]: d
Out[62]: {11: 1.6696428571428572,
          12: 0.5175276752767528,
          13: 1.6762948207171315,
          14: 1.146457765667575}
In []:
In []:
In [20]: 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_launch
import sys
```

c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\sklearn\model\_se
DeprecationWarning)

#### 2.8 Malignant Sub-Classification Using Cancer Classification

Out[64]: ['models/SVM/SVM\_Models\_VGG16\_CancerType\_Benign.joblib']

```
In [5]: 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 [6]: classes=[21,22,23,24]
In [7]: from sklearn.utils.class_weight import compute_class_weight
In [8]: class_weight=compute_class_weight(class_weight='balanced', classes=classes,y=Y_train_2
In [9]: print(class_weight)
[0.35669231 2.72764706 1.96150592 3.12466307]
In [10]: print(np.unique(Y_train_2))
[21 22 23 24]
In [11]: print(len(X_train_2))
4637
In [12]: print(len(Y_test_2))
788
In [13]: d = dict(enumerate(class_weight, 1))
In [14]: print(d)
{1: 0.3566923076923077, 2: 2.7276470588235293, 3: 1.9615059221658206, 4: 3.1246630727762805}
In [15]: d1={1:21,2:22,3:23,4:24}
In [16]: d=dict((d1[key], value) for (key, value) in d.items())
In [17]: d
```

```
Out[17]: {21: 0.3566923076923077,
          22: 2.7276470588235293,
          23: 1.9615059221658206,
          24: 3.1246630727762805}
In [18]: 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_launch
  import sys
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\sklearn\model_se
  DeprecationWarning)
9759.513311783001 seconds
0.7088634893249947
{'svc__C': 10.0, 'svc__gamma': 'auto', 'svc__kernel': 'rbf'}
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\ipykernel_launch
  # Remove the CWD from sys.path while we load stuff.
In [19]: clf4=gs3.best_estimator_
         clf4.fit(X_train_2,Y_train_2)
         print(clf4.score(X_test_2,Y_test_2))
0.2779187817258883
In [20]: pred=clf4.predict(X_test_2)
In [21]: precision_recall_fscore_support(Y_test_2,pred)
c:\users\karan gupta\appdata\local\programs\python\python37\lib\site-packages\sklearn\metrics\
  'precision', 'predicted', average, warn_for)
```

```
Out[21]: (array([0.25945241, 1.
                                 , 0.88888889, 0.
                                                              ]),
         array([0.995, 0.06, 0.04, 0.
         array([0.41158221, 0.11320755, 0.07655502, 0.
                                                              ]),
         array([200, 200, 200, 188], dtype=int64))
In [22]: confusion matrix(Y test 2,pred)
Out[22]: array([[199, 0,
                            1,
                                 0],
                            Ο,
                                 0],
                [188, 12,
                [192, 0, 8,
                                 0],
                [188, 0, 0,
                                 0]], dtype=int64)
In [49]: dump(clf4, 'models/SVM/SVM models_VGG16_CancerType_Malignant.joblib')
Out[49]: ['models/SVM/SVM_models_VGG16_CancerType_Malignant.joblib']
   Dumping Models
3
In [27]: clf=SVC(C=.001,kernel='linear')
         clf.fit(X train, Y train)
         clf.score(X_test,Y_test)
Out [27]: 0.9018987341772152
In [28]: dump(clf,'models/SVM/SVM models_VGG16_Magnification.joblib')
Out[28]: ['models/SVM/SVM_models_VGG16_Magnification.joblib']
In [40]: clf=SVC(C=.001,kernel='linear')
         clf.fit(X_train_40,Y_train_40)
         clf.score(X test 40,Y test 40)
Out [40]: 0.7858942065491183
In [41]: dump(clf, 'models/SVM/SVM models VGG16 Magnification 40.joblib')
Out[41]: ['models/SVM/SVM_models_VGG16_Magnification_40.joblib']
In [42]: clf=SVC(C=.01,kernel='linear')
         clf.fit(X_train_100,Y_train_100)
         clf.score(X_test_100,Y_test_100)
Out[42]: 0.8520408163265306
In [43]: dump(clf, 'models/SVM/SVM models_VGG16 Magnification_100.joblib')
Out[43]: ['models/SVM/SVM_models_VGG16_Magnification_100.joblib']
In [44]: clf=SVC(C=.001,kernel='linear')
         clf.fit(X_train_200,Y_train_200)
         clf.score(X_test_200,Y_test_200)
```

```
Out[44]: 0.8197969543147208
In [45]: dump(clf,'models/SVM/SVM_models_VGG16_Magnification_200.joblib')
Out[45]: ['models/SVM/SVM_models_VGG16_Magnification_200.joblib']
In [46]: clf=SVC(C=.001,kernel='linear')
         clf.fit(X_train_400,Y_train_400)
         clf.score(X_test_400,Y_test_400)
Out [46]: 0.7959697732997482
In [47]: dump(clf, 'models/SVM/SVM_models_VGG16_Magnification_400.joblib')
Out[47]: ['models/SVM/SVM_models_VGG16_Magnification_400.joblib']
In []:
```