

SVC EEG

September 9, 2022

1 Support Vector Machine: EEG of pediatric subjects with epilepsy

Support Vector Machine or SVM is a type of algorithm that establish continuous functions that create a surface in the space of the input data and optimize the classification of labeled data. To do this several hyperparameters are needed. We must define what type of function will be optimized(e.g. polynomial), the starting coefficients, and the weights of each class.

The advantages of support vector machines are:

- Effective in high dimensional spaces.
- Still effective in cases where the number of dimensions is greater than the number of samples.
- Uses a subset of training points in the decision function (called support vectors), so it is also memory efficient.
- Versatile: different Kernel functions can be specified for the decision function. Common kernels are provided, but it is also possible to specify custom kernels.

The disadvantages of support vector machines include: * If the number of features is much greater than the number of samples, avoid over-fitting in choosing Kernel functions and the regularization term is crucial. * SVMs do not directly provide probability estimates, these are calculated using an expensive five-fold cross-validation (see [Scores and probabilities](#)).

1.0.1 Importing the packagery:

```
[1]: from seaborn import load_dataset, pairplot, heatmap
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from matplotlib.colors import ListedColormap
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.datasets import make_moons, make_circles, make_classification
from sklearn.neural_network import MLPClassifier
from sklearn.neighbors import KNeighborsClassifier
from sklearn.svm import SVC
from sklearn.gaussian_process import GaussianProcessClassifier
from sklearn.gaussian_process.kernels import RBF
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier, AdaBoostClassifier
```

```

from sklearn.naive_bayes import GaussianNB
from sklearn.discriminant_analysis import QuadraticDiscriminantAnalysis
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score

```

1.0.2 Imprting the data:

```
[2]: data=pd.read_csv("Data_medians_4.csv")
```

```
[3]: del data['Unnamed: 0']
del data['38']
```

```
[4]: for i in range(0,len(data)):
    if data.iloc[i,-1]!='Awake':
        if data.iloc[i,-1]!='N1':
            if data.iloc[i,-1]!='N2':
                if data.iloc[i,-1]!='N3':
                    if data.iloc[i,-1]!='REM':
                        data.iloc[i,-1]=np.nan
```

```
[5]: data_2=data.dropna()
```

```
[6]: data_2
```

```
[6]:
```

	0	1	2	3	4	5 \
0	19.911429	21.826667	19.901111	19.076667	21.918889	14.850000
1	40.660000	35.816667	38.405556	42.591111	40.958889	37.191667
2	28.203333	24.951111	25.603333	33.302222	28.286667	24.416667
3	18.286190	16.575556	17.173333	22.212222	18.586667	15.928333
4	11.339524	10.327778	11.073333	9.653333	12.070000	10.046667
..
160	23.572857	21.060000	22.835556	24.652222	19.970000	18.556667
161	18.736190	16.664444	17.992222	19.476667	16.011111	14.800000
162	14.926190	13.685556	14.278889	15.434444	12.957778	11.726667
163	10.816190	9.702222	10.213333	10.985556	9.072222	8.145000
164	4.886190	4.464444	4.621111	4.972222	4.191111	3.780000

	6	7	8	9 ...	29 \
0	58.638571	60.644444	61.956667	57.580000	... 210.934721
1	148.707143	114.740000	116.255556	180.642222	... 601.949326
2	104.157619	78.117778	77.450000	141.416667	... 395.701329
3	72.619048	55.015556	55.386667	98.113333	... 258.530174
4	42.745238	33.640000	34.700000	37.300000	... 156.242749
..
160	62.023810	55.311111	62.484444	65.027778	... 233.876713
161	47.885238	42.373333	47.960000	49.993333	... 179.285611
162	37.418571	33.524444	37.350000	38.766667	... 137.755238

163	27.838095	24.458889	27.936667	28.534444	...	99.599288
164	12.803333	12.212222	13.241111	14.298889	...	41.261791

	30	31	32	33	34	\
0	627.235043	648.175431	626.022091	643.377231	540.470154	
1	2122.510839	1710.063114	1684.448166	3062.885569	2100.222584	
2	1433.984345	1147.643210	1131.199674	2268.983257	1375.903119	
3	951.861362	774.702891	776.389821	1493.072896	943.001124	
4	587.817203	465.337699	467.104798	546.078070	612.918988	
..	
160	956.606279	901.840306	1079.626406	1000.445620	627.724242	
161	731.916047	694.028468	835.901439	759.237127	474.306587	
162	558.178258	505.701924	648.375509	566.989604	363.330987	
163	449.385031	411.347601	560.246779	459.417619	270.892424	
164	165.142991	188.748088	225.965814	206.167366	122.058242	

	35	36	37	39	
0	495.039453	4.422272e+07	0.001152	Awake	
1	1750.616297	2.508360e+06	0.002201	N2	
2	1170.567469	1.469640e+06	0.001673	REM	
3	787.208156	5.040967e+06	0.002785	N2	
4	465.555425	9.224363e+06	0.001658	N3	
..	
160	663.898963	1.033058e+05	0.000996	REM	
161	509.926345	3.701008e+04	0.000528	N1	
162	390.608436	9.521701e+04	0.000363	N2	
163	296.452757	6.754039e+06	0.000063	N3	
164	109.744217	2.577399e+06	0.000092	N2	

[157 rows x 39 columns]

1.0.3 Splitting the data

```
[7]: X=data_2.iloc[:, :-1]
      Y=data_2.iloc[:, -1]

[8]: X_train, X_test, y_train, y_test = train_test_split(X, Y, train_size=0.75,
      ↪ random_state=100)
```

1.1 Meaning of each hyperparameter:

Warning: DO NOT RUN THE NEXT LINE OF CODE

```
[9]: # The SVC Class from Sklearn
      SVC(C=1.0,                # The regularization parameter
          kernel='rbf',         # The kernel type used
          degree=3,             # Degree of polynomial function)
```

```

gamma='scale',          # The kernel coefficient
coef0=0.0,              # If kernel = 'poly'/'sigmoid'
shrinking=True,         # To use shrinking heuristic
probability=False,      # Enable probability estimates
tol=0.001,              # Stopping criterion
cache_size=200,         # Size of kernel cache
class_weight=None,      # The weight of each class
verbose=False,          # Enable verbose output
max_iter=-1,            # Hard limit on iterations
decision_function_shape='ovr', # One-vs-rest or one-vs-one
break_ties=False,       # How to handle breaking ties
random_state=None       # Random state of the model
)

```

[9]: SVC()

1.2 Defining the model

1.2.1 Weights:

We expect the algorithm to have difficulties distinguishing between N2 and N3 segments due to the epileptic “noise”. To avoid overfitting the data, we will lower the weight of these two classes.

```

[9]: weights = {
    "Awake": 1,
    "N1": 0.5,
    "N2": 0.5,
    "N3": 0.5,
    "REM": 1
}

```

1.2.2 kernel and degree:

We will use a polynomial kernel of 3rd degree

```

[ ]: clf = SVC(kernel='poly', degree=5, gamma='auto', C=1.0, class_weight=weights)
      clf.fit(X_train, y_train)

```

```

[11]: predictions_2 = clf.predict(X_train)

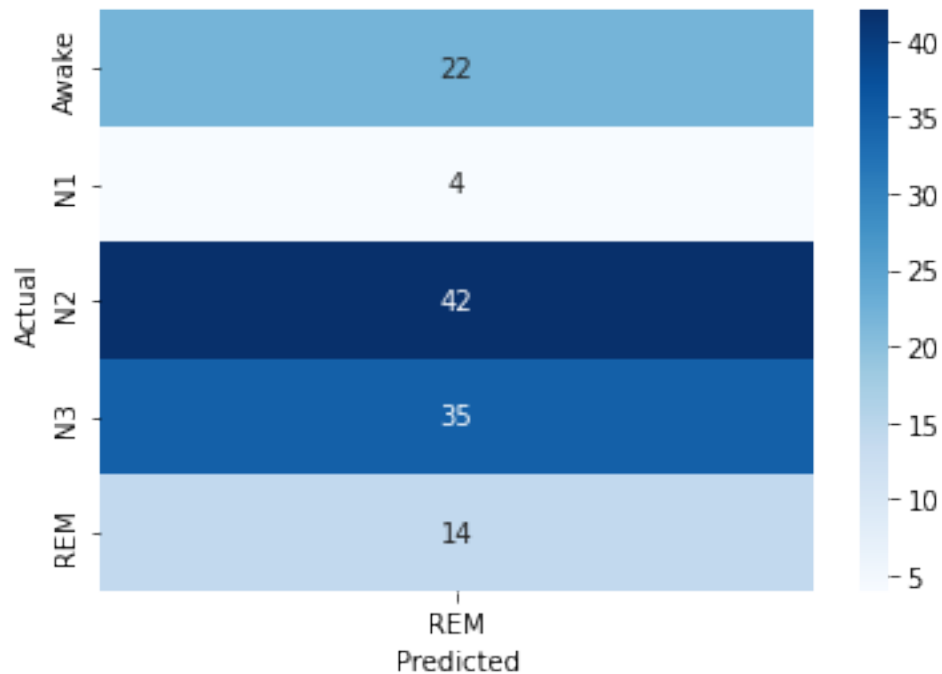
```

```

[12]: confusion_matrix = pd.crosstab(y_train, predictions_2, rownames=['Actual'],
    ↪ colnames=['Predicted'])

      heatmap(confusion_matrix, annot=True, cmap="Blues")
      plt.show()

```

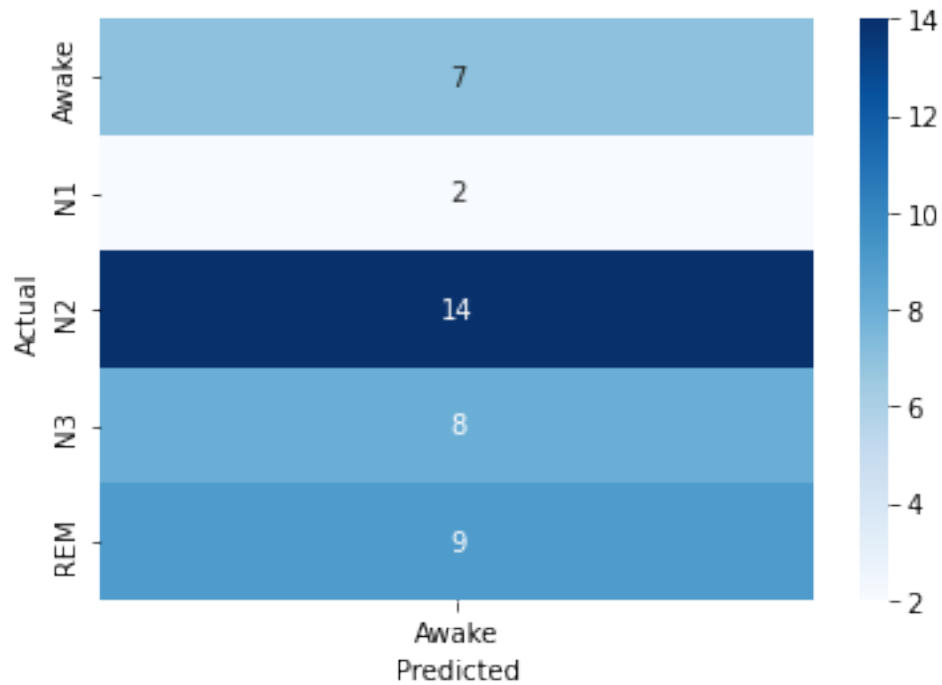


```
[111]: predictions = clf.predict(X_test)
print(predictions)
```

```
['Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake'
'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake'
'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake'
'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake' 'Awake'
'Awake' 'Awake' 'Awake' 'Awake']
```

```
[112]: confusion_matrix = pd.crosstab(y_test, predictions, rownames=['Actual'],
    ↪ colnames=['Predicted'])

heatmap(confusion_matrix, annot=True, cmap="Blues")
plt.show()
```



```
[65]: from sklearn.gaussian_process import GaussianProcessClassifier
      from sklearn.gaussian_process.kernels import RBF
```

```
[66]: gauss=GaussianProcessClassifier(1.0 * RBF(1.0),random_state=0)
```

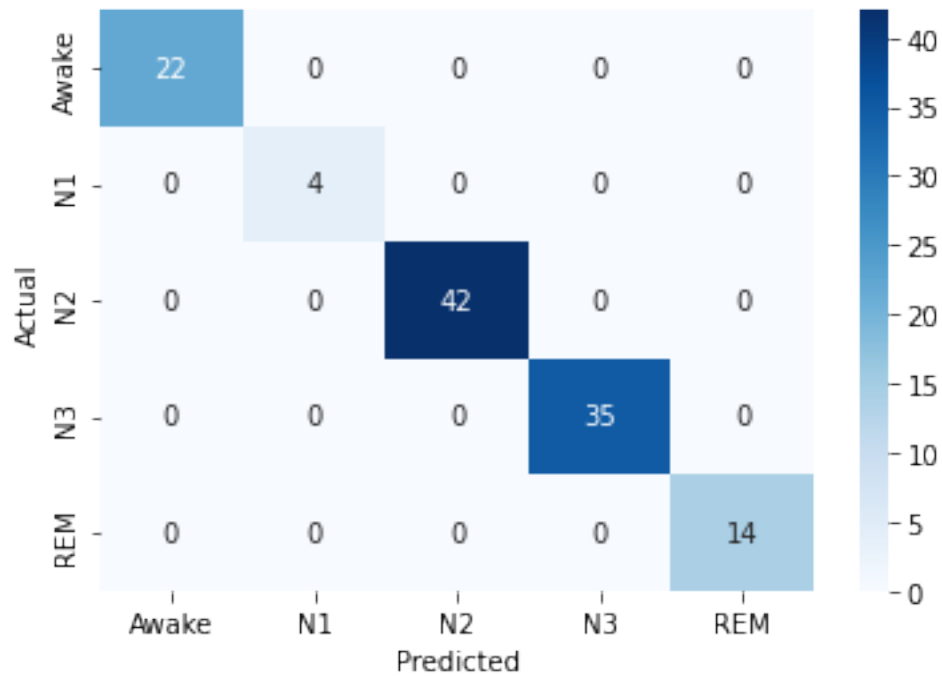
```
[67]: gauss.fit(X_train, y_train)
```

```
[67]: GaussianProcessClassifier(kernel=1**2 * RBF(length_scale=1), random_state=0)
```

```
[70]: predictions_4 = gauss.predict(X_train)
```

```
[74]: confusion_matrix = pd.crosstab(y_train, predictions_4, rownames=['Actual'],
      ↪ colnames=['Predicted'])

      heatmap(confusion_matrix, annot=True, cmap="Blues")
      plt.show()
```

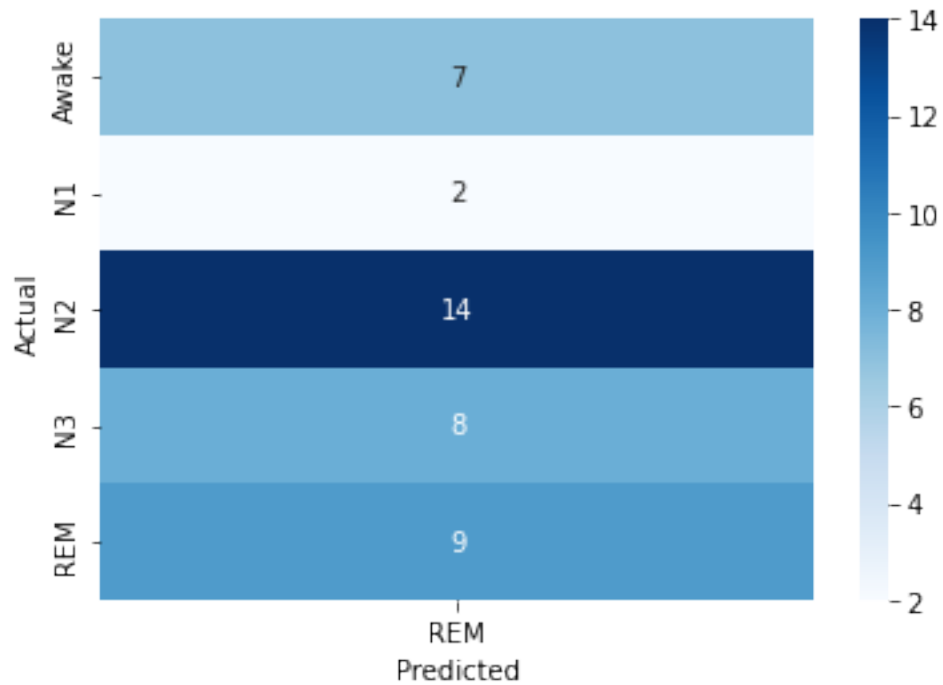


```
[75]: predictions_3 = gauss.predict(X_test)
      print(predictions_3)
```

```
['REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM'
 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM'
 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM' 'REM'
 'REM' 'REM' 'REM' 'REM']
```

```
[76]: confusion_matrix = pd.crosstab(y_test, predictions_3, rownames=['Actual'],
    ↪ colnames=['Predicted'])

heatmap(confusion_matrix, annot=True, cmap="Blues")
plt.show()
```



Observations:

- Note that the training set has a 100% accuracy, however the test set lets us know that all the data is being classified as REM. This means that the input data is biased and the algorithm tends to overfit the training set.
- To avoid this we must use a simpler algorithm

[]: