

Project Development Phase - Sprint Delivery Plan

Sprint 3 – Model Building

Date	18 November 2022
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Project Name	Statistical Machine Learning Approaches to Liver Disease Prediction

Machine learning Algorithms for Model Building:

Train and Test the model using classification algorithm:

- Import the classification algorithm:

```
In [17]: from sklearn.svm import SVC
from sklearn.ensemble import RandomForestClassifier
from sklearn.neighbors import KNeighborsClassifier
from sklearn.preprocessing import LabelEncoder
le = LabelEncoder()
le.fit(xtrain['Gender'].astype(str))
xtrain['Gender'] = le.transform(xtrain['Gender'].astype(str))
xtest['Gender'] = le.transform(xtest['Gender'].astype(str))
```

- Initialize the model

```
In [18]: svm=SVC()
RFmodel=RandomForestClassifier()
KNNmodel=KNeighborsClassifier()
```

- Training model with our data
 - SVC Model
 - Random Forest Model
 - K-Nearest Neighbors

```
In [19]: from sklearn.svm import SVC
svm=SVC(gamma='auto')
svm.fit(xtrain,ytrain)
```

```
Out[19]: SVC(gamma='auto')
```

```
In [20]: from sklearn.ensemble import RandomForestClassifier
RFmodel=RandomForestClassifier()
RFmodel.fit(xtrain,ytrain)
```

```
Out[20]: RandomForestClassifier()
```

```
In [21]: from sklearn.neighbors import KNeighborsClassifier
KNN=KNeighborsClassifier()
KNN.fit(xtrain,ytrain)
```

```
Out[21]: KNeighborsClassifier()
```

- Evaluation Metrics:

```
In [22]: import warnings
warnings.filterwarnings("ignore")
#accuracies
from sklearn.metrics import accuracy_score
SVMpred=svm.predict(xtest)
SVMaccuracy=accuracy_score(SVMpred,ytest)
print("SVM accuracy:",SVMaccuracy*100)

RFpred=RFmodel.predict(xtest)
RFaccuracy=accuracy_score(ytest,RFpred)
print("RFC accuracy:",RFaccuracy*100)

KNNpred=KNN.predict(xtest)
KNNaccuracy=accuracy_score(KNNpred,ytest)
print("KNN accuracy:",KNNaccuracy*100)

SVM accuracy: 73.50427350427351
RFC accuracy: 75.21367521367522
KNN accuracy: 75.21367521367522
```

```
In [23]: #confusion matrix
from sklearn.metrics import confusion_matrix
cm1=confusion_matrix(ytest,SVMpred)
cm2=confusion_matrix(ytest,RFpred)
cm3=confusion_matrix(ytest,KNNpred)
print("SVM Confusion matrix",cm1)
print("RFC Confusion matrix",cm2)
print("KNN Confusion matrix",cm3)

SVM Confusion matrix [[86  0]
 [31  0]]
RFC Confusion matrix [[73 13]
 [16 15]]
KNN Confusion matrix [[69 17]
 [12 19]]
```

- Save the model:

```
In [24]: import pickle
pickle.dump(svm,open('liver_analysis.pkl','wb'))
```