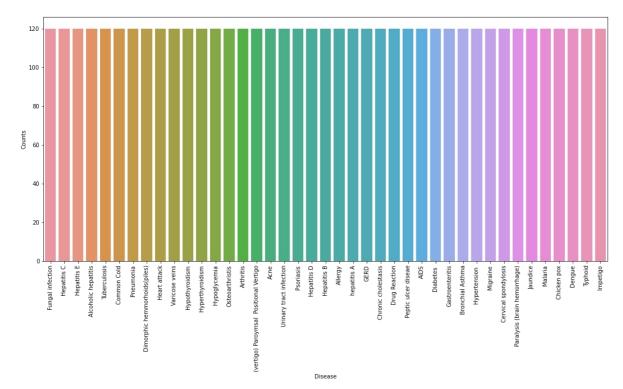
```
In [1]:
# Importing
libraries import
numpy as np
import pandas
as pd
from scipy.stats
import mode import
matplotlib.pyplot as
plt import seaborn as
sns
from sklearn.preprocessing import LabelEncoder
from sklearn.model_selection import
train_test_split, cross_val_score from
sklearn.svm import SVC
from sklearn.naive_bayes import GaussianNB
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score, confusion_matrix
% matplotlib inline
In [3]:
# Reading the train.csv by
removing the # last column
since it's an empty column
DATA_PATH =
"C:/Users/ambee/OneDrive/Desktop/Trai
ning.csv" data =
pd.read_csv(DATA_PATH).dropna(axis
=1)
```

```
# Checking whether the dataset is
balanced or not disease_counts =
data["prognosis"].value_counts()
temp_df = pd.DataFrame({
    "Disease":
    disease_counts.index,
    "Counts":
    disease_counts.values
})
plt.figure(figsize = (18,8))
sns.barplot(x = "Disease", y = "Counts", data = temp_df) plt.xticks(rotation=90)
plt.show()
```



In [4]:

xxxxxxxxx

```
# Encoding the target value
```

into numerical # value using

LabelEncoder

encoder = LabelEncoder()

data["prognosis"] = encoder.fit_transform(data["prognosis"])

In [5]:

XXXXXXXXXX

```
X = data.iloc[:,:-1]
```

y = data.iloc[:, -1]

X_train, X_test, y_train, y_test

=train_test_split(X, y, test_size = 0.2,

 $random_state = 24$)

print(f"Train: {X_train.shape},

{y_train.shape}") print(f"Test:

{X_test.shape}, {y_test.shape}")

Train: (3936, 132), (3936,) Test: (984, 132), (984,)

```
In [6]:
XXXXXXXXX
# Defining scoring metric for k-fold
cross validation def
cv_scoring(estimator, X, y):
   return accuracy_score(y, estimator.predict(X))
# Initializing
Models models =
{
   "SVC":SVC(),
   "Gaussian NB":GaussianNB(),
   "Random Forest":RandomForestClassifier(random_state=18)
}
# Producing cross validation score
for the models for model_name in
models:
   model = models[model_name]
   scores = cross\_val\_score(model, X, y, cv = 10,
                    n_jobs = -1,
                    scoring = cv_scoring)
   print("=="*30)
   print(model_name)
   print(f"Scores:
   {scores}")
```

```
print(f"Mean Score: {np.mean(scores)}")
Scores: [1. 1. 1. 1. 1. 1. 1. 1. 1. 1.]
Mean Score: 1.0
Gaussian NB
Scores: [1. 1. 1. 1. 1. 1. 1. 1. 1. 1.]
Mean Score: 1.0
Random Forest
Scores: [1. 1. 1. 1. 1. 1. 1. 1. 1. 1.]
Mean Score: 1.0
In [7]:
XXXXXXXXX
# Training and testing
SVM Classifier svm_model
= SVC()
svm_model.fit(X_train,
y_train) preds =
svm_model.predict(X_test)
print(f"Accuracy on train data by SVM Classifier\
: {accuracy_score(y_train, svm_model.predict(X_train))*100}")
print(f"Accuracy on test data by SVM Classifier\
: {accuracy_score(y_test,
preds)*100}") cf_matrix =
confusion_matrix(y_test, preds)
plt.figure(figsize=(12,8))
sns.heatmap(cf_matrix,
annot=True)
plt.title("Confusion Matrix for SVM Classifier on
Test Data") plt.show()
# Training and testing Naive
```

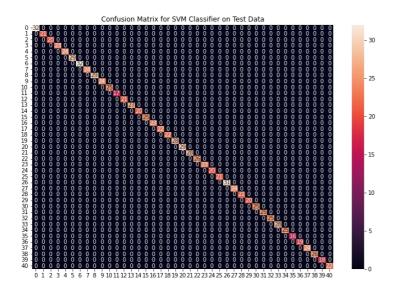
Bayes Classifier nb_model =

```
nb_model.fit(X_train, y_train)
preds = nb_model.predict(X_test)
print(f"Accuracy on train data by Naive Bayes Classifier\
: {accuracy_score(y_train, nb_model.predict(X_train))*100}")
print(f"Accuracy on test data by Naive Bayes Classifier\
: {accuracy_score(y_test,
preds)*100}") cf_matrix =
confusion_matrix(y_test, preds)
plt.figure(figsize=(12,8))
sns.heatmap(cf_matrix,
annot=True)
plt.title("Confusion Matrix for Naive Bayes
Classifier on Test Data") plt.show()
# Training and testing Random Forest
Classifier rf model =
RandomForestClassifier (random\_state
=18) rf_model.fit(X_train, y_train)
preds = rf_model.predict(X_test)
print(f"Accuracy on train data by Random Forest Classifier\
: {accuracy_score(y_train, rf_model.predict(X_train))*100}")
print(f"Accuracy on test data by Random Forest Classifier\
: {accuracy_score(y_test, preds)*100}")
cf_matrix =
confusion_matrix(y_test, preds)
plt.figure(figsize=(12,8))
sns.heatmap(cf_matrix,
annot=True)
plt.title("Confusion Matrix for Random Forest Classifier on
```

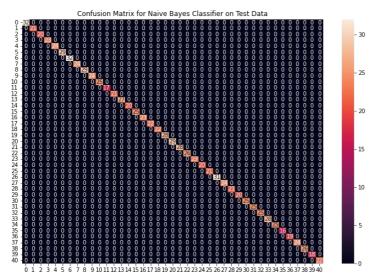
GaussianNB()

Test Data") plt.show()

Accuracy on train data by SVM Classifier: 100.0 Accuracy on test data by SVM Classifier: 100.0



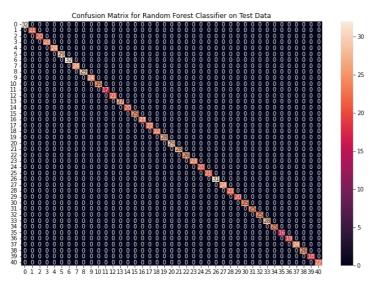
Accuracy on train data by Naive Bayes



Classifier: 100.0 Accuracy on test data by

Naive Bayes Classifier: 100.0

Accuracy on train data by Random Forest



Classifier: 100.0 Accuracy on test data by

Random Forest Classifier: 100.0

In [9]:

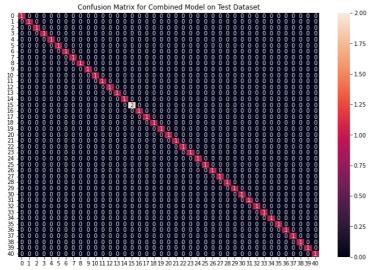
xxxxxxxxx

```
whole data
final_svm_model = SVC()
final_nb_model =
GaussianNB()
final_rf_model =
RandomForestClassifier(random_state=18)
final_svm_model.fit(X, y)
final_nb_model.fit(X, y)
```

Reading the test data

```
test_data = pd.read_csv("C:/Users/ambee/OneDrive/Desktop/Testing.csv").dropna(axis=1)
test_X = test_data.iloc[:, :-1]
test_Y = encoder.transform(test_data.iloc[:, -1])
# Making prediction by take mode
of predictions # made by all the
classifiers
svm_preds =
final_svm_model.predict(test_X)
nb\_preds =
final_nb_model.predict(test_X)
rf_preds =
final_rf_model.predict(test_X)
final\_preds = [mode([i,j,k])[0][0] for i,j,
        k in zip(svm_preds, nb_preds, rf_preds)]
print(f"Accuracy on Test dataset by the combined model\
: {accuracy_score(test_Y, final_preds)*100}")
cf_matrix = confusion_matrix(test_Y,
final_preds) plt.figure(figsize=(12,8))
sns.heatmap(cf_matrix, annot = True)
plt.title("Confusion Matrix for Combined Model on
```

Test Dataset") plt.show()



Accuracy on Test dataset by the combined model: 100.0 In [10]:

XXXXXXXXX

```
symptoms = X.columns.values

# Creating a symptom index
dictionary to encode the # input
symptoms into numerical form
symptom_index = {}

for index, value in enumerate(symptoms):
    symptom = " ".join([i.capitalize() for i in
        value.split("_")]) symptom_index[symptom] =
        index

data_dict = {
        "symptom_index":symptom_index,
        "predictions_classes":encoder.classes_
}
```

Defining the Function

Input: string containing symptoms

separated by commas # Output:

Generated predictions by models

def

predictDisease(sympto

```
ms): symptoms =
symptoms.split(",")
```

creating input data for the models

```
input_data = [0] *
    len(data_dict["symptom_index"]) for
    symptom in symptoms:
      index =
      data_dict["symptom_index"][sympt
      om] input_data[index] = 1
    # reshaping the input data and
    converting it # into suitable
    format for model predictions
    input_data = np.array(input_data).reshape(1,-1)
    # generating individual outputs
    rf_prediction =
    data_dict["predictions_classes"][final_rf_model.predict(input_data)[0]]
    nb_prediction =
    data_dict["predictions_classes"][final_nb_model.predict(input_data)[0]]
    svm_prediction =
    data_dict["predictions_classes"][final_svm_model.predict(input_data)[0]]
    # making final prediction by taking mode of all predictions
    final_prediction = mode([rf_prediction, nb_prediction,
    svm_prediction])[0][0] predictions = {
       "rf_model_prediction":
      rf_prediction,
       "naive_bayes_prediction":
      nb_prediction,
       "svm_model_prediction":
      svm_prediction,
       "final_prediction":final_prediction
    }
return predictions
```

Testing the function

print(predictDisease("Itching,Skin Rash,Nodal Skin Eruptions"))

{'rf_model_prediction': 'Fungal infection', 'naive_bayes_prediction': 'Fungal infection', 'svm_model_prediction': 'Fungal infection', 'final_prediction': 'Fungal infection'}

In []:

XXXXXXXXXX