

## #Red Wine Quality Dataset

```
from google.colab import files
```

```
uploaded = files.upload()  
import pandas as pd  
wn = pd.read_csv('winequality2.csv')  
wn.head()  
wn.isnull().values.any()  
wn.describe()
```

```
x = wn.drop('quality', axis = 1)  
y = wn['quality']  
from sklearn.model_selection import train_test_split
```

```
from sklearn.pipeline import Pipeline  
from sklearn.feature_selection import VarianceThreshold  
from imblearn.over_sampling import SMOTE  
import numpy as np  
from sklearn.datasets import make_classification  
rng = np.random.RandomState(42)  
x, y = make_classification(random_state=rng)  
oversample=SMOTE()  
wn=oversample.fit_resample(x,y)  
import sklearn.model_selection  
!pip install scikit-optimize  
from skopt import BayesSearchCV  
x_train, x_test, y_train, y_test = sklearn.model_selection.train_test_split(x, y,  
random_state=42, stratify=y)  
from sklearn.ensemble import RandomForestClassifier  
pipe = Pipeline([  
    ('selector', VarianceThreshold()),  
    ('model', RandomForestClassifier())  
)  
param_grid = {  
    'selector__threshold': (0,0.1),  
    'model': [RandomForestClassifier()],  
    'model__max_depth': (1,500),  
    'model__max_leaf_nodes': (10,100),
```

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        'model__max_samples': (0.1,0.8),
        'model__min_samples_leaf': (1,6),
        'model__min_samples_split': (2,8),
        'model__n_estimators': (50,700),
        'model__n_jobs': (1,400)
    }
    opt = BayesSearchCV(
        pipe,
        # (parameter space, # of evaluations)
        [(param_grid, 500)],
        cv=2)
    opt.fit(x_train, y_train)
    print("test score: %s" % opt.score(x_test, y_test))
    clf = RandomForestClassifier(random_state=42)
    clf = clf.fit(x_train, y_train)
    y_hat = clf.predict(x_test)
    print("RF Accuracy", sklearn.metrics.accuracy_score(y_test, y_hat))

```

## # Brain Stroke Dataset

```

from google.colab import files

```

```

uploaded = files.upload()
import pandas as pd
st = pd.read_csv('brain_stroke.csv')
st.head()
print(st['gender'].unique())
st['gender'].value_counts()
print(st['ever_married'].unique())
st['ever_married'].value_counts()
print(st['work_type'].unique())
st['work_type'].value_counts()
print(st['Residence_type'].unique())
st['Residence_type'].value_counts()
print(st['smoking_status'].unique())
st['smoking_status'].value_counts()
ohed = pd.get_dummies(st, columns = ['gender', 'ever_married', 'work_type',
'Residence_type', 'smoking_status'])
ohed.head()

```

```

ohed.describe()
from sklearn.model_selection import train_test_split
x = ohed.drop('stroke', axis = 1)
y = ohed['stroke']
import numpy as np
from sklearn.datasets import make_classification
from imblearn.over_sampling import SMOTE
oversample=SMOTE()
ohed=oversample.fit_resample(x,y)
import sklearn.model_selection
x_train, x_test, y_train, y_test = sklearn.model_selection.train_test_split(x, y,
random_state=42, stratify=y)
from sklearn.ensemble import RandomForestClassifier
!pip install scikit-optimize
from skopt import BayesSearchCV
from sklearn.pipeline import Pipeline
from sklearn.feature_selection import VarianceThreshold
x_train, x_test, y_train, y_test = sklearn.model_selection.train_test_split(x, y,
random_state=42, stratify=y)
pipe = Pipeline([
    ('selector', VarianceThreshold()),
    ('model', RandomForestClassifier())
])
rfc_search = {
    'selector__threshold': (0,0.1),
    'model':([RandomForestClassifier()]),
    'model__max_depth': (1,500),
    'model__max_leaf_nodes': (10,100),
    'model__max_samples': (0.1,0.8),
    'model__min_samples_leaf': (1,6),
    'model__min_samples_split': (2,8),
    'model__n_estimators': (50,700),
    'model__n_jobs': (-1,400)
}

opt = BayesSearchCV(
    pipe,
    # (parameter space, # of evaluations)
    [(rfc_search, 50)],
    cv=3)

```

```

opt.fit(x_train, y_train)
print("test score: %s" % opt.score(x_test, y_test))
clf = RandomForestClassifier(random_state=42)
clf = clf.fit(x_train, y_train)
y_hat = clf.predict(x_test)
print("RF Accuracy", sklearn.metrics.accuracy_score(y_test, y_hat))

```

Table 1: Red Wine Quality Pipeline	
Mean Accuracy Score Before Bayesian Optimization	0.88
Mean Accuracy Score After Bayesian Optimization	0.88

Table 2: Brain Stroke Pipeline	
Mean Accuracy Score Before Bayesian Optimization	0.9478
Mean Accuracy Score After Bayesian Optimization	0.9502