

```
import pandas as pd
import io
import numpy as np
import matplotlib.pyplot as plt
%matplotlib inline
from sklearn.model_selection import train_test_split
from sklearn.impute import KNNImputer
from sklearn import preprocessing
from sklearn.preprocessing import StandardScaler
```

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

```
uploaded = files.upload()
```

No file chosen

rerun this cell to enable.

Upload widget is only available when the cell has been executed in the current browser session. Please

Saving chronic kidney disease.csv to chronic kidney disease (1).csv

```
df = pd.read_csv(io.BytesIO(uploaded['chronic_kidney_disease.csv']))
```

```
X=df.iloc[:,0:24]
```

```
Y=df["class"]
```

```
df.head()
```

	age	bp	sg	al	su	rbc	pc	pcc	ba	bgr	...	pcv	wbcc	rbcc	htn	dm	cad	appet	pe	ane	class
0	40	80	100	1	0	0	0	1	1	121		44	7000	52	0	0	1	1	1	1	old

```

knn_missing_values_imputer = KNNImputer(n_neighbors=5)
df = df.replace('?', np.nan)
X = pd.DataFrame(knn_missing_values_imputer.fit_transform(X), columns = X.columns)
standard_feature_scaler = StandardScaler()
X = standard_feature_scaler.fit_transform(X)
X = pd.DataFrame(X, columns=['age', 'bp', 'sg', 'al', 'su', 'rbc', 'pc', 'pcc', 'ba', 'bgr', 'bu', 'sc', 'sod', 'pot',
                             'hemo', 'pcv', 'wbcc', 'rbcc', 'htn', 'dm',
                             'cad', 'appet', 'pe', 'ane'])

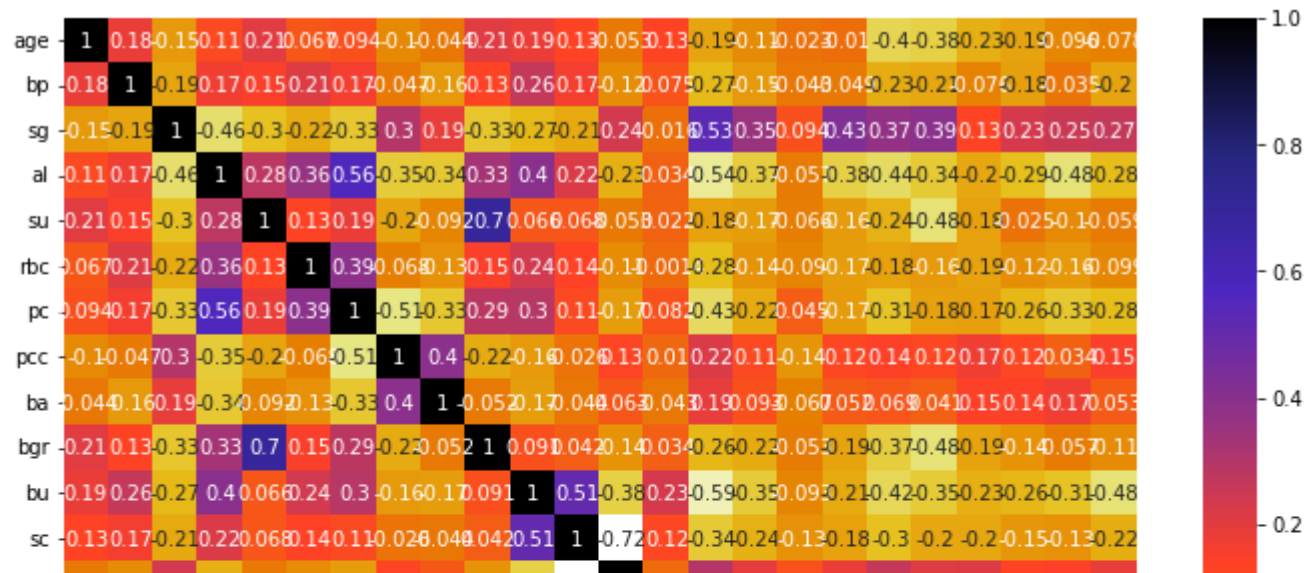
X_train, X_test, Y_train, Y_test = train_test_split(X, Y, train_size = 0.7, test_size = 0.3)

```

```

import seaborn as sns
plt.figure(figsize=(12,10))
cor=X_train.corr()
sns.heatmap(cor,annot=True,cmap=plt.cm.CMRmap_r)
plt.show()

```



```
def correlate(dataset,threshold):
    col_corr=set()
    corr_matrix=dataset.corr()
    for i in range(len(corr_matrix.columns)):
        for j in range(i):
            if abs(corr_matrix.iloc[i,j])>threshold:
                colname=corr_matrix.columns[i]
                col_corr.add(colname)
    return col_corr
```

```
corr_features=correlate(X_train,0.6)
print(len(set(corr_features)))
print(corr_features)
```

```
4
{'rbcc', 'pcv', 'sod', 'bgr'}
```

```
X_train.drop(corr_features,axis=1)
X_test.drop(corr_features,axis=1)
```

	age	bp	sg	al	su	rbc	pc	pcc	ba	bu	sc	
291	-0.257518	0.264037	1.413392	-0.795017	-0.431146	-0.364890	-0.484322	0.360477	0.263664	-0.268332	-0.367591	0.
250	-0.668232	0.264037	1.413392	-0.795017	-0.431146	-0.364890	-0.484322	0.360477	0.263664	-0.955146	-0.332030	0.
107	0.211870	1.745312	-0.409164	-0.023156	3.388529	-0.364890	-0.484322	0.360477	0.263664	-0.086528	-0.047541	-0.
55	-0.961600	0.264037	-2.231720	1.520567	-0.431146	2.740554	-0.484322	0.360477	0.263664	0.123556	-0.043985	-0.
38	1.033299	0.264037	0.502114	1.520567	-0.431146	2.740554	-0.484322	0.360477	0.263664	0.923492	0.183606	0.
...
54	0.681258	0.264037	-1.320442	0.748706	1.478692	-0.364890	-0.484322	0.360477	0.263664	-0.240052	0.059143	-0.
352	-0.844253	-1.217238	0.502114	-0.795017	-0.431146	-0.364890	-0.484322	0.360477	0.263664	-0.207731	-0.349810	0.
325	0.387890	0.264037	0.502114	-0.795017	-0.431146	-0.364890	-0.484322	0.360477	0.263664	-0.147130	-0.332030	-0.
12	0.974625	-0.476600	-0.409164	1.520567	0.523773	-0.364890	-0.484322	-2.774104	0.263664	0.297279	-0.172005	0.
0	-0.198844	0.264037	0.502114	-0.023156	-0.431146	-0.364890	-0.484322	0.360477	0.263664	-0.429935	-0.332030	-0.

120 rows × 20 columns

```

from sklearn.svm import SVC
from sklearn.metrics import classification_report

linear_svm=SVC(C=1.0,kernel='linear',random_state=0)
linear_svm=linear_svm.fit(X_train,Y_train)

print(linear_svm.score(X_train,Y_train))
print(linear_svm.score(X_test,Y_test))

1.0
0.975

predicted =linear_svm.predict(X_test)
report = classification_report(Y_test, predicted)

```

```
print(report)
```

	precision	recall	f1-score	support
ckd	1.00	0.96	0.98	81
notckd	0.93	1.00	0.96	39
accuracy			0.97	120
macro avg	0.96	0.98	0.97	120
weighted avg	0.98	0.97	0.98	120