```
# Importing Libraries:
import pandas as pd
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt

dataset=pd.read_csv(r"/content/kidney_disease.csv")

# Top 5 records:
dataset.head()
```

	id	age	bp	sg	al	su	rbc	рс	рсс	ba	bgr	bu	sc	sod	pot	hemo	pcv	WC	rc	htn
0	0	48.0	80.0	1.020	1.0	0.0	NaN	normal	notpresent	notpresent	121.0	36.0	1.2	NaN	NaN	15.4	44	7800	5.2	yes
1	1	7.0	50.0	1.020	4.0	0.0	NaN	normal	notpresent	notpresent	NaN	18.0	8.0	NaN	NaN	11.3	38	6000	NaN	no
2	2	62.0	80.0	1.010	2.0	3.0	normal	normal	notpresent	notpresent	423.0	53.0	1.8	NaN	NaN	9.6	31	7500	NaN	no
3	3	48.0	70.0	1.005	4.0	0.0	normal	abnormal	present	notpresent	117.0	56.0	3.8	111.0	2.5	11.2	32	6700	3.9	yes
4	4	51.0	80.0	1.010	2.0	0.0	normal	normal	notpresent	notpresent	106.0	26.0	1.4	NaN	NaN	11.6	35	7300	4.6	no

Saved successfully!

```
# Dropping unneccsary feature :
dataset = dataset.drop('id', axis=1)
```

```
# Shape of dataset:
dataset.shape
```

(400, 25)

```
# Cheaking Missing (NaN) Values:
dataset.isnull().sum()
                          9
     age
     bp
                         12
                         47
     sg
     al
                         46
                         49
     su
     rbc
                        152
                         65
     рс
     рсс
                          4
     ba
                          4
     bgr
                         19
     bu
                         17
     SC
                         87
     sod
     pot
                         88
                         52
     hemo
                         70
     pcv
                        105
     WC
                        130
     rc
     htn
                          2
     dm
                          2
     cad
                          1
     appet
                          1
     pe
     ane
                          1
     classification
 Saved successfully!
```

# Description:
dataset.describe()

	age	bp	sg	al	su	bgr	bu	sc	sod	pot	
count	391.000000	388.000000	353.000000	354.000000	351.000000	356.000000	381.000000	383.000000	313.000000	312.000000	348.00
mean	51.483376	76.469072	1.017408	1.016949	0.450142	148.036517	57.425722	3.072454	137.528754	4.627244	12.52
std	17.169714	13.683637	0.005717	1.352679	1.099191	79.281714	50.503006	5.741126	10.408752	3.193904	2.9′
min	2.000000	50.000000	1.005000	0.000000	0.000000	22.000000	1.500000	0.400000	4.500000	2.500000	3.1(
25%	42.000000	70.000000	1.010000	0.000000	0.000000	99.000000	27.000000	0.900000	135.000000	3.800000	10.30

# Datatypes:
dataset.dtypes

age bp	float64 float64
sg	float64
al	float64
su	float64
rbc	object
рс	object
рсс	object
ba	object
bgr	float64
bu	float64
SC	float64
sod	float64

Saved successfully! object WC rc object object htn object dm object cad appet object object pe object ane classification object

dtype: object

## dataset.head()

	age	bp	sg	al	su	rbc	рс	рсс	ba	bgr	• • •	pcv	WC	rc	htn	dm	cad	appet	pe	ane
0	48.0	80.0	1.020	1.0	0.0	NaN	normal	notpresent	notpresent	121.0		44	7800	5.2	yes	yes	no	good	no	no
1	7.0	50.0	1.020	4.0	0.0	NaN	normal	notpresent	notpresent	NaN		38	6000	NaN	no	no	no	good	no	no
2	62.0	80.0	1.010	2.0	3.0	normal	normal	notpresent	notpresent	423.0		31	7500	NaN	no	yes	no	poor	no	yes
3	48.0	70.0	1.005	4.0	0.0	normal	abnormal	present	notpresent	117.0		32	6700	3.9	yes	no	no	poor	yes	yes
4	51.0	80.0	1.010	2.0	0.0	normal	normal	notpresent	notpresent	106.0		35	7300	4.6	no	no	no	good	no	no
5 rows × 25 columns																				

```
dataset['rbc'].value counts()
     normal
                 201
     abnormal
                  47
     Name: rbc, dtype: int64
dataset['rbc'] = dataset['rbc'].replace(to replace = {'normal' : 0, 'abnormal' : 1})
 Saved successfully!
     HUI IIIa±
     abnormal
                  76
     Name: pc, dtype: int64
dataset['pc'] = dataset['pc'].replace(to replace = {'normal' : 0, 'abnormal' : 1})
dataset['pcc'].value_counts()
     notpresent
                   354
```

```
42
     present
     Name: pcc, dtype: int64
dataset['pcc'] = dataset['pcc'].replace(to replace = {'notpresent':0,'present':1})
dataset['ba'].value counts()
     notpresent
                   374
     present
                    22
     Name: ba, dtype: int64
dataset['ba'] = dataset['ba'].replace(to replace = {'notpresent':0,'present':1})
dataset['htn'].value counts()
     no
            251
            147
     yes
     Name: htn, dtype: int64
dataset['htn'] = dataset['htn'].replace(to_replace = {'yes' : 1, 'no' : 0})
dataset['dm'].value counts()
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     \tno
                3
     \tyes
                2
                1
      yes
     Name: dm, dtype: int64
dataset['dm'] = dataset['dm'].replace(to_replace = {'\tyes':'yes', ' yes':'yes', '\tno':'no'})
dataset['dm'] = dataset['dm'].replace(to_replace = {'yes' : 1, 'no' : 0})
```

```
dataset['cad'].value_counts()
             362
     no
              34
     ves
     \tno
               2
     Name: cad, dtype: int64
dataset['cad'] = dataset['cad'].replace(to replace = {'\tno':'no'})
dataset['cad'] = dataset['cad'].replace(to replace = {'yes' : 1, 'no' : 0})
dataset['appet'].unique()
     array(['good', 'poor', nan], dtype=object)
dataset['appet'] = dataset['appet'].replace(to replace={'good':1,'poor':0,'no':np.nan})
dataset['pe'].value_counts()
            323
     no
     yes
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dataset['pe'] = dataset['pe'].replace(to_replace = {'yes' : 1, 'no' : 0})
dataset['ane'].value_counts()
            339
     no
     yes
             60
     Name: ane, dtype: int64
```

```
dataset['ane'] = dataset['ane'].replace(to_replace = {'yes' : 1, 'no' : 0})
dataset['classification'].value_counts()
     ckd
               248
     notckd
               150
     ckd\t
     Name: classification, dtype: int64
dataset['classification'] = dataset['classification'].replace(to replace={'ckd\t':'ckd'})
dataset["classification"] = [1 if i == "ckd" else 0 for i in dataset["classification"]]
dataset.head()
```

	age	bp	sg	al	su	rbc	рс	рсс	ba	bgr	• • •	pcv	WC	rc	htn	dm	cad	appet	pe	ane	classification
0	48.0	80.0	1.020	1.0	0.0	NaN	0.0	0.0	0.0	121.0		44	7800	5.2	1.0	1.0	0.0	1.0	0.0	0.0	1
1	7.0	50.0	1.020	4.0	0.0	NaN	0.0	0.0	0.0	NaN		38	6000	NaN	0.0	0.0	0.0	1.0	0.0	0.0	1
2	62.0	80.0	1.010	2.0	3.0	0.0	0.0	0.0	0.0	423.0		31	7500	NaN	0.0	1.0	0.0	0.0	0.0	1.0	1
3	48.0	70.0	1.005	4.0	0.0	0.0	1.0	1.0	0.0	117.0		32	6700	3.9	1.0	0.0	0.0	0.0	1.0	1.0	1
4	51 N	80 N	1 010	2 0	0 0	0.0	0.0	0.0	0.0	106.0		35	7300	4.6	0.0	0.0	0.0	1.0	0.0	0.0	1
vod :	CLICCACC	fullyl			×																

Saved successfully!

# Datatypes: dataset.dtypes

age	float64
bp	float64
sg	float64
al	float64
su	float64

```
float64
rbc
                 float64
рс
                 float64
pcc
                 float64
ba
                 float64
bgr
                 float64
bu
                 float64
sc
                 float64
sod
                 float64
pot
                 float64
hemo
                  object
pcv
                  object
WC
                  object
rc
                 float64
htn
                 float64
dm
                 float64
cad
appet
                 float64
                 float64
pe
                 float64
ane
classification
                   int64
dtype: object
```

atype. object

```
dataset['pcv'] = pd.to_numeric(dataset['pcv'], errors='coerce')
dataset['wc'] = pd.to_numeric(dataset['wc'], errors='coerce')
dataset['rc'] = pd.to_numeric(dataset['rc'], errors='coerce')
```

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age	float64
bp	float64
sg	float64
al	float64
su	float64
rbc	float64
рс	float64
рсс	float64
ba	float64
bgr	float64

float64 bu float64 sc float64 sod float64 pot hemo float64 float64 pcv float64 WC float64 rc htn float64 float64 dm cad float64 float64 appet float64 pe float64 ane classification int64

dtype: object

# Description:
dataset.describe()

Saved successfully!

рс

pcc

ba

al bp rbc age sg su # Cheaking Missing (NaN) Values: dataset.isnull().sum().sort\_values(ascending=False) rbc 152 rc 131 WC 106 88 pot sod 87 71 pcv 65 рс 52 hemo 49 su 47 sg al 46 bgr 44 bu 19 17 SC bp 12 9 age ba рсс htn 2 dm 2 cad Saved successfully! classification 0 dtype: int64 dataset.columns Index(['age', 'bp', 'sg', 'al', 'su', 'rbc', 'pc', 'pcc', 'ba', 'bgr', 'bu', 'sc', 'sod', 'pot', 'hemo', 'pcv', 'wc', 'rc', 'htn', 'dm', 'cad', 'appet', 'pe', 'ane', 'classification'], dtype='object')

bgr ...

Saved successfully!

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