

```
In [1]: import pandas as pd
import numpy as np

import matplotlib.pyplot as plt
import seaborn as sns
%matplotlib inline
```

```
In [24]: dataset = pd.read_csv('Heart.csv')
```

```
In [25]: dataset.head(10)
```

Out[25]:

	Unnamed: 0	Age	Sex	ChestPain	RestBP	Chol	Fbs	RestECG	MaxHR	ExAng	Oldpeak	Slope	Ca
0	1	63	1	typical	145	233	1	2	150	0	2.3	3	0.0
1	2	67	1	asymptomatic	160	286	0	2	108	1	1.5	2	3.0
2	3	67	1	asymptomatic	120	229	0	2	129	1	2.6	2	2.0
3	4	37	1	nonanginal	130	250	0	0	187	0	3.5	3	0.0
4	5	41	0	nontypical	130	204	0	2	172	0	1.4	1	0.0
5	6	56	1	nontypical	120	236	0	0	178	0	0.8	1	0.0
6	7	62	0	asymptomatic	140	268	0	2	160	0	3.6	3	2.0
7	8	57	0	asymptomatic	120	354	0	0	163	1	0.6	1	0.0
8	9	63	1	asymptomatic	130	254	0	2	147	0	1.4	2	1.0
9	10	53	1	asymptomatic	140	203	1	2	155	1	3.1	3	0.0

```
In [26]: dataset2 = pd.read_csv('heart.csv')
```

```
In [27]: dataset2.head(10)
```

Out[27]:

	Unnamed: 0	Age	Sex	ChestPain	RestBP	Chol	Fbs	RestECG	MaxHR	ExAng	Oldpeak	Slope	Ca
0	1	63	1	typical	145	233	1	2	150	0	2.3	3	0.0
1	2	67	1	asymptomatic	160	286	0	2	108	1	1.5	2	3.0
2	3	67	1	asymptomatic	120	229	0	2	129	1	2.6	2	2.0
3	4	37	1	nonanginal	130	250	0	0	187	0	3.5	3	0.0
4	5	41	0	nontypical	130	204	0	2	172	0	1.4	1	0.0
5	6	56	1	nontypical	120	236	0	0	178	0	0.8	1	0.0
6	7	62	0	asymptomatic	140	268	0	2	160	0	3.6	3	2.0
7	8	57	0	asymptomatic	120	354	0	0	163	1	0.6	1	0.0
8	9	63	1	asymptomatic	130	254	0	2	147	0	1.4	2	1.0
9	10	53	1	asymptomatic	140	203	1	2	155	1	3.1	3	0.0

```
In [28]: dataset2.isna().sum()
```

```
Out[28]: Unnamed: 0      0
Age          0
Sex          0
ChestPain    0
RestBP       0
Chol         0
Fbs          0
RestECG      0
MaxHR        0
ExAng        0
Oldpeak      0
Slope        0
Ca           4
Thal         2
AHD          0
dtype: int64
```

```
In [29]: dataset2 = dataset2.dropna(axis=0)
```

```
In [30]: dataset2.isnull().sum()
```

```
Out[30]: Unnamed: 0      0
Age          0
Sex          0
ChestPain    0
RestBP       0
Chol         0
Fbs          0
RestECG      0
MaxHR        0
ExAng        0
Oldpeak      0
Slope        0
Ca           0
Thal         0
AHD          0
dtype: int64
```

```
In [31]: dataset2.describe()
```

Out[31]:

	Unnamed: 0	Age	Sex	RestBP	Chol	Fbs	RestECG	MaxHR	
count	297.000000	297.000000	297.000000	297.000000	297.000000	297.000000	297.000000	297.000000	29
mean	150.673401	54.542088	0.676768	131.693603	247.350168	0.144781	0.996633	149.599327	
std	87.323283	9.049736	0.468500	17.762806	51.997583	0.352474	0.994914	22.941562	
min	1.000000	29.000000	0.000000	94.000000	126.000000	0.000000	0.000000	71.000000	
25%	75.000000	48.000000	0.000000	120.000000	211.000000	0.000000	0.000000	133.000000	
50%	150.000000	56.000000	1.000000	130.000000	243.000000	0.000000	1.000000	153.000000	
75%	226.000000	61.000000	1.000000	140.000000	276.000000	0.000000	2.000000	166.000000	
max	302.000000	77.000000	1.000000	200.000000	564.000000	1.000000	2.000000	202.000000	

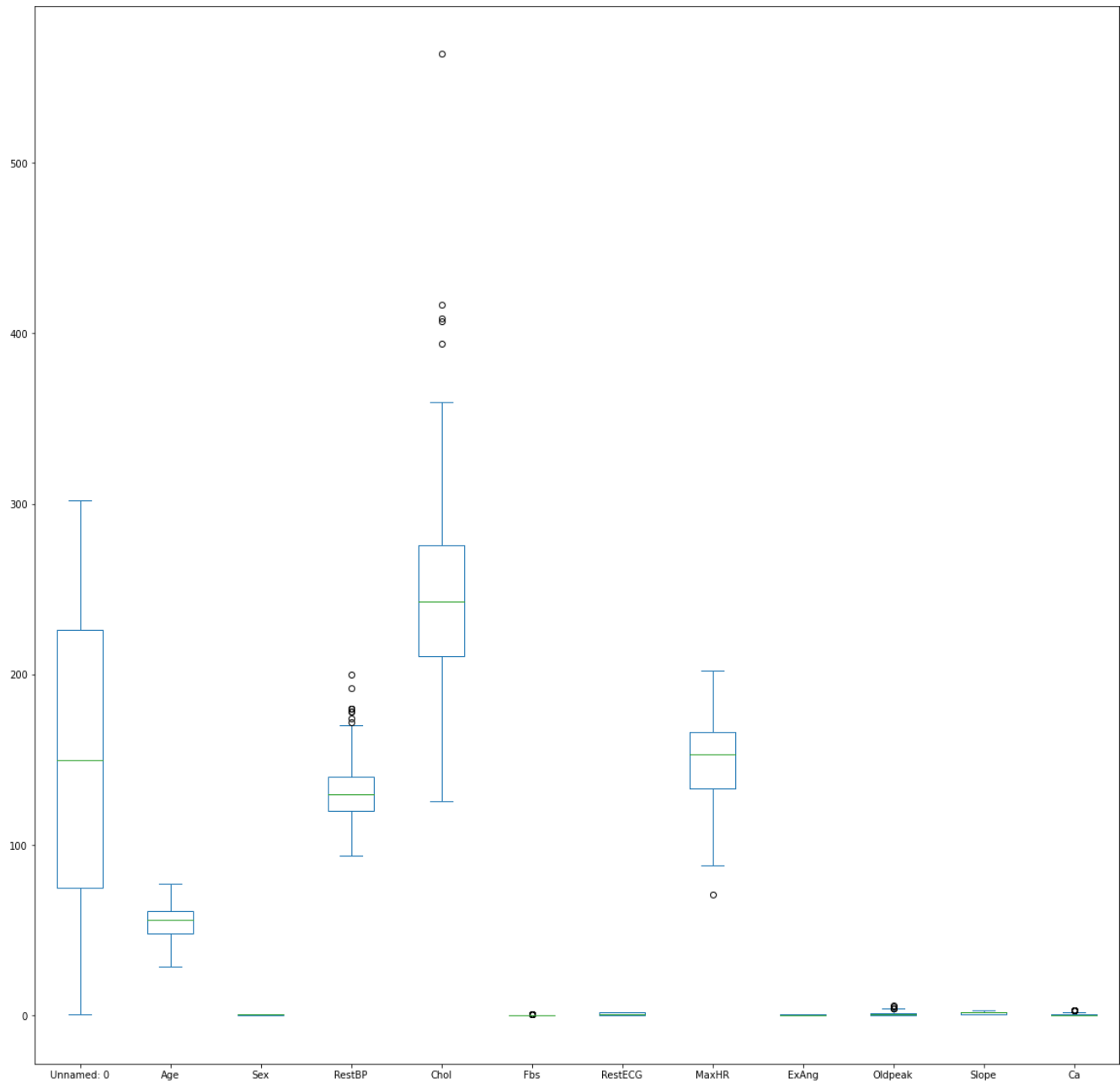
```
In [32]: dataset.count()
```

```
Out[32]: Unnamed: 0      303  
Age          303  
Sex          303  
ChestPain    303  
RestBP       303  
Chol         303  
Fbs          303  
RestECG      303  
MaxHR        303  
ExAng        303  
Oldpeak      303  
Slope        303  
Ca           299  
Thal         301  
AHD          303  
dtype: int64
```

```
In [33]: plt.figure(figsize=(9,3))
dataset2.plot(kind='box',figsize=(20,20))
```

```
Out[33]: <AxesSubplot:>
```

```
<Figure size 648x216 with 0 Axes>
```



```
In [34]: percentile25 = dataset2['Chol'].quantile(0.25)
percentile75 = dataset2['Chol'].quantile(0.75)
```

```
In [35]: iqr = percentile75 - percentile25
```

```
In [36]: upper_limit = percentile75 + 1.5 * iqr
lower_limit = percentile25 - 1.5 * iqr
```

```
In [37]: dataset2[dataset2['Chol'] > upper_limit]
dataset2[dataset2['Chol'] < lower_limit]
```

Out[37]:

Unnamed: 0	Age	Sex	ChestPain	RestBP	Chol	Fbs	RestECG	MaxHR	ExAng	Oldpeak	Slope	Ca	Th
------------	-----	-----	-----------	--------	------	-----	---------	-------	-------	---------	-------	----	----

```
In [38]: dataset2['Chol'] = dataset2[dataset2['Chol'] < upper_limit]
```

```
-----
ValueError                                Traceback (most recent call last)
~\AppData\Local\Temp\ipykernel_6648\3294842726.py in <module>
----> 1 dataset2['Chol'] = dataset2[dataset2['Chol'] < upper_limit]

~\AppData\Local\Programs\Python\Python310\lib\site-packages\pandas\core\frame.py in __
setitem__(self, key, value)
    3600         self._setitem_array(key, value)
    3601     elif isinstance(value, DataFrame):
-> 3602         self._set_item_frame_value(key, value)
    3603     elif (
    3604         is_list_like(value)

~\AppData\Local\Programs\Python\Python310\lib\site-packages\pandas\core\frame.py in _s
et_item_frame_value(self, key, value)
    3727         len_cols = 1 if is_scalar(cols) else len(cols)
    3728         if len_cols != len(value.columns):
-> 3729             raise ValueError("Columns must be same length as key")
    3730
    3731         # align right-hand-side columns if self.columns
```

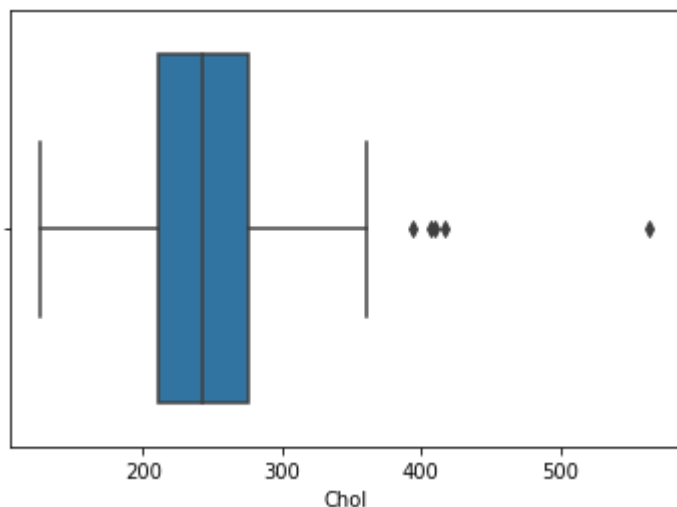
ValueError: Columns must be same length as key

```
In [39]: sns.boxplot(dataset2['Chol'])
```

C:\Users\rapen\AppData\Local\Programs\Python\Python310\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

warnings.warn(

Out[39]: <AxesSubplot:xlabel='Chol'>



```
In [40]: percentile25 = dataset2['RestBP'].quantile(0.25)
percentile75 = dataset2['RestBP'].quantile(0.75)
```

```
In [41]: iqr = percentile75 - percentile25
```

```
In [42]: upper_limit = percentile75 + 1.5 * iqr
lower_limit = percentile25 - 1.5 * iqr
```

```
In [43]: dataset2[dataset2['RestBP'] > upper_limit]
dataset2[dataset2['RestBP'] < lower_limit]
```

Out[43]:

Unnamed: 0	Age	Sex	ChestPain	RestBP	Chol	Fbs	RestECG	MaxHR	ExAng	Oldpeak	Slope	Ca	Th
------------	-----	-----	-----------	--------	------	-----	---------	-------	-------	---------	-------	----	----



```
In [44]: dataset2['RestBP'] = dataset2[dataset2['RestBP'] < upper_limit]
```

```
-----
ValueError                                Traceback (most recent call last)
~\AppData\Local\Temp\ipykernel_6648\2386516796.py in <module>
----> 1 dataset2['RestBP'] = dataset2[dataset2['RestBP'] < upper_limit]

~\AppData\Local\Programs\Python\Python310\lib\site-packages\pandas\core\frame.py in __
setitem__(self, key, value)
    3600         self._setitem_array(key, value)
    3601     elif isinstance(value, DataFrame):
-> 3602         self._set_item_frame_value(key, value)
    3603     elif (
    3604         is_list_like(value)

~\AppData\Local\Programs\Python\Python310\lib\site-packages\pandas\core\frame.py in _s
et_item_frame_value(self, key, value)
    3727         len_cols = 1 if is_scalar(cols) else len(cols)
    3728         if len_cols != len(value.columns):
-> 3729             raise ValueError("Columns must be same length as key")
    3730
    3731         # align right-hand-side columns if self.columns
```

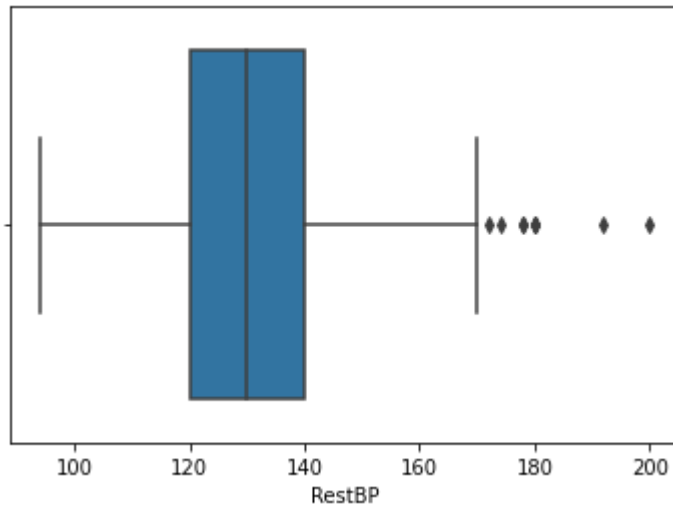
ValueError: Columns must be same length as key

```
In [45]: sns.boxplot(dataset2['RestBP'])
```

C:\Users\rapen\AppData\Local\Programs\Python\Python310\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

```
warnings.warn(
```

```
Out[45]: <AxesSubplot:xlabel='RestBP'>
```



```
In [ ]:
```

```
In [46]: print(dataset2.ChestPain.unique())  
print(dataset2.Thal.unique())  
print(dataset2.AHD.unique())
```

```
['typical' 'asymptomatic' 'nonanginal' 'nontypical']  
['fixed' 'normal' 'reversible']  
['No' 'Yes']
```

```
In [47]: heart_encoding = pd.get_dummies(dataset2[['ChestPain', 'Thal', 'AHD']])
heart_final = pd.concat([dataset2, heart_encoding],1)
heart_final = heart_final.drop(['ChestPain', 'Thal', 'AHD'], axis = 1)
heart_final.head(10)
```

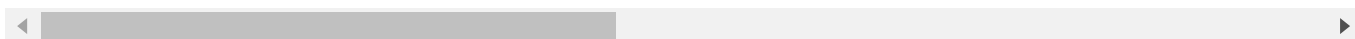
C:\Users\rapen\AppData\Local\Temp\ipykernel_6648\3224836008.py:2: FutureWarning: In a future version of pandas all arguments of concat except for the argument 'objs' will be keyword-only

```
heart_final = pd.concat([dataset2, heart_encoding],1)
```

Out[47]:

	Unnamed: 0	Age	Sex	RestBP	Chol	Fbs	RestECG	MaxHR	ExAng	Oldpeak	...	Ca	ChestPain_asymptomatic
0	1	63	1	145	233	1	2	150	0	2.3	...	0.0	
1	2	67	1	160	286	0	2	108	1	1.5	...	3.0	
2	3	67	1	120	229	0	2	129	1	2.6	...	2.0	
3	4	37	1	130	250	0	0	187	0	3.5	...	0.0	
4	5	41	0	130	204	0	2	172	0	1.4	...	0.0	
5	6	56	1	120	236	0	0	178	0	0.8	...	0.0	
6	7	62	0	140	268	0	2	160	0	3.6	...	2.0	
7	8	57	0	120	354	0	0	163	1	0.6	...	0.0	
8	9	63	1	130	254	0	2	147	0	1.4	...	1.0	
9	10	53	1	140	203	1	2	155	1	3.1	...	0.0	

10 rows × 21 columns



```
In [48]: heart_final.Sex.value_counts()
```

```
Out[48]: 1    201
0     96
Name: Sex, dtype: int64
```

```
In [49]: pd.crosstab(heart_final.AHD_Yes,heart_final.Sex)
```

Out[49]:

	Sex	0	1
AHD_Yes			
0	71	89	
1	25	112	

```
In [50]: heart_final.columns
```

```
Out[50]: Index(['Unnamed: 0', 'Age', 'Sex', 'RestBP', 'Chol', 'Fbs', 'RestECG', 'MaxHR',
               'ExAng', 'Oldpeak', 'Slope', 'Ca', 'ChestPain_asymptomatic',
               'ChestPain_nonanginal', 'ChestPain_nontypical', 'ChestPain_typical',
               'Thal_fixed', 'Thal_normal', 'Thal_reversible', 'AHD_No', 'AHD_Yes'],
              dtype='object')
```



```
In [51]: heart_final.dtypes
```

```
Out[51]: Unnamed: 0          int64
Age          int64
Sex          int64
RestBP       int64
Chol         int64
Fbs          int64
RestECG      int64
MaxHR        int64
ExAng        int64
Oldpeak      float64
Slope        int64
Ca           float64
ChestPain_asymptomatic  uint8
ChestPain_nonanginal    uint8
ChestPain_nontypical    uint8
ChestPain_typical       uint8
Thal_fixed              uint8
Thal_normal             uint8
Thal_reversable         uint8
AHD_No                  uint8
AHD_Yes                 uint8
dtype: object
```

```
In [52]: df = heart_final.drop('AHD_Yes', axis=1)
df_norm = (df-df.min())/(df.max()-df.min())
df_norm = pd.concat((df_norm, heart_final.AHD_Yes), 1)
```

C:\Users\rapen\AppData\Local\Temp\ipykernel_6648\1126837166.py:3: FutureWarning: In a future version of pandas all arguments of concat except for the argument 'objs' will be keyword-only

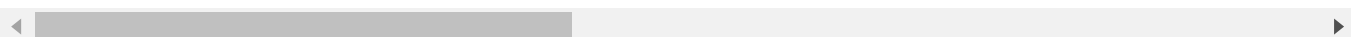
```
df_norm = pd.concat((df_norm, heart_final.AHD_Yes), 1)
```

```
In [53]: df_norm.head(10)
```

```
Out[53]:
```

	Unnamed: 0	Age	Sex	RestBP	Chol	Fbs	RestECG	MaxHR	ExAng	Oldpeak	...	Ca
0	0.000000	0.708333	1.0	0.481132	0.244292	1.0	1.0	0.603053	0.0	0.370968	...	0.000000
1	0.003322	0.791667	1.0	0.622642	0.365297	0.0	1.0	0.282443	1.0	0.241935	...	1.000000
2	0.006645	0.791667	1.0	0.245283	0.235160	0.0	1.0	0.442748	1.0	0.419355	...	0.666667
3	0.009967	0.166667	1.0	0.339623	0.283105	0.0	0.0	0.885496	0.0	0.564516	...	0.000000
4	0.013289	0.250000	0.0	0.339623	0.178082	0.0	1.0	0.770992	0.0	0.225806	...	0.000000
5	0.016611	0.562500	1.0	0.245283	0.251142	0.0	0.0	0.816794	0.0	0.129032	...	0.000000
6	0.019934	0.687500	0.0	0.433962	0.324201	0.0	1.0	0.679389	0.0	0.580645	...	0.666667
7	0.023256	0.583333	0.0	0.245283	0.520548	0.0	0.0	0.702290	1.0	0.096774	...	0.000000
8	0.026578	0.708333	1.0	0.339623	0.292237	0.0	1.0	0.580153	0.0	0.225806	...	0.333333
9	0.029900	0.500000	1.0	0.433962	0.175799	1.0	1.0	0.641221	1.0	0.500000	...	0.000000

10 rows × 21 columns



```
In [54]: df_norm = df_norm.dropna()
```

```
In [55]: X = df_norm.drop(['AHD_Yes', 'Unnamed: 0'], axis=1)
Y = df_norm.AHD_Yes
```

```
In [56]: #data cleaning done
X.isnull().sum()
```

```
Out[56]: Age                0
Sex                0
RestBP            0
Chol              0
Fbs              0
RestECG          0
MaxHR            0
ExAng            0
Oldpeak          0
Slope            0
Ca              0
ChestPain_asymptomatic  0
ChestPain_nonanginal  0
ChestPain_nontypical  0
ChestPain_typical    0
Thal_fixed        0
Thal_normal       0
Thal_reversable    0
AHD_No           0
dtype: int64
```

```
In [ ]:
```

```
In [57]: Y.isnull().sum()
```

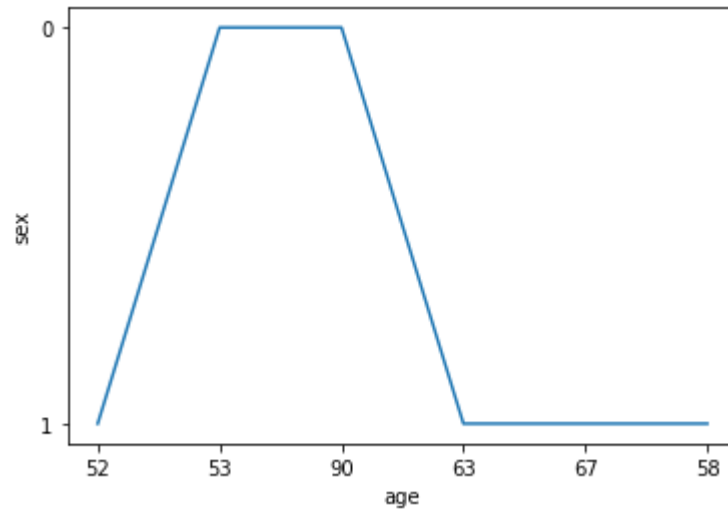
```
Out[57]: 0
```

```
In [58]: df_norm.isnull().sum()
```

```
Out[58]: Unnamed: 0        0
Age                0
Sex                0
RestBP            0
Chol              0
Fbs              0
RestECG          0
MaxHR            0
ExAng            0
Oldpeak          0
Slope            0
Ca              0
ChestPain_asymptomatic  0
ChestPain_nonanginal  0
ChestPain_nontypical  0
ChestPain_typical    0
Thal_fixed        0
Thal_normal       0
Thal_reversable    0
AHD_No           0
AHD_Yes          0
dtype: int64
```

```
In [59]: # d] line charts
age=["52","53","90","63","67","58"]
sex=['1','0','0','1','1','1']
plt.plot(age,sex)
plt.xlabel("age")
plt.ylabel("sex")
x = np.array([52, 53, 90, 63,67]) # X-axis points
y = x*2 # Y-axis points

plt.show()
```



```
In [61]: #b]histogram
plt.hist(dataset['Age'],dataset['Sex'])
```

```
-----
ValueError                                Traceback (most recent call last)
~\AppData\Local\Temp\ipykernel_6648\1216958322.py in <module>
      1 #b]histogram
----> 2 plt.hist(dataset['Age'],dataset['Sex'])

~\AppData\Local\Programs\Python\Python310\lib\site-packages\matplotlib\pyplot.py in hi
st(x, bins, range, density, weights, cumulative, bottom, histtype, align, orientation,
rwidth, log, color, label, stacked, data, **kwargs)
    2588     orientation='vertical', rwidth=None, log=False, color=None,
    2589     label=None, stacked=False, *, data=None, **kwargs):
-> 2590     return gca().hist(
    2591         x, bins=bins, range=range, density=density, weights=weights,
    2592         cumulative=cumulative, bottom=bottom, histtype=histtype,

~\AppData\Local\Programs\Python\Python310\lib\site-packages\matplotlib\__init__.py in
inner(ax, data, *args, **kwargs)
    1410     def inner(ax, *args, data=None, **kwargs):
    1411         if data is None:
-> 1412             return func(ax, *map(sanitize_sequence, args), **kwargs)
    1413
    1414         bound = new_sig.bind(ax, *args, **kwargs)

~\AppData\Local\Programs\Python\Python310\lib\site-packages\matplotlib\axes\_axes.py i
n hist(self, x, bins, range, density, weights, cumulative, bottom, histtype, align, or
ientation, rwidth, log, color, label, stacked, **kwargs)
    6637         # this will automatically overwrite bins,
    6638         # so that each histogram uses the same bins
-> 6639         m, bins = np.histogram(x[i], bins, weights=w[i], **hist_kwargs)
    6640         tops.append(m)
    6641         tops = np.array(tops, float) # causes problems later if it's an int

<__array_function__ internals> in histogram(*args, **kwargs)

~\AppData\Local\Programs\Python\Python310\lib\site-packages\numpy\lib\histograms.py in
histogram(a, bins, range, normed, weights, density)
    791     a, weights = _ravel_and_check_weights(a, weights)
    792
--> 793     bin_edges, uniform_bins = _get_bin_edges(a, bins, range, weights)
    794
    795     # Histogram is an integer or a float array depending on the weights.

~\AppData\Local\Programs\Python\Python310\lib\site-packages\numpy\lib\histograms.py in
_get_bin_edges(a, bins, range, weights)
    429     bin_edges = np.asarray(bins)
    430     if np.any(bin_edges[:-1] > bin_edges[1:]):
--> 431         raise ValueError(
    432             '`bins` must increase monotonically, when an array')
    433

ValueError: `bins` must increase monotonically, when an array
```

