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# Assignment_3
# Name: Keshav Goyal
# Roll No: 21BEC2297
import pandas as pd
\stackrel{\cdot}{\text{import matplotlib.pyplot as plt}}
```

from matplotlib import rcParams

import seaborn as sns import numpy as np

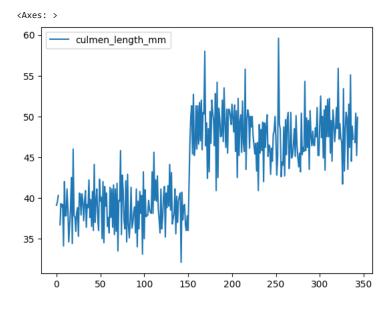
df = pd.read_csv('/content/penguins_size.csv') # Importing the dataset

df

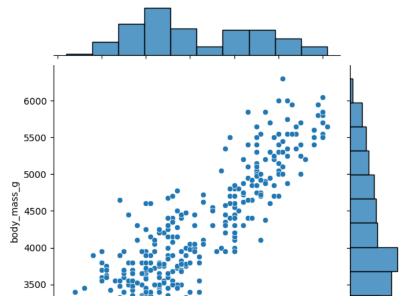
₽		species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_
	0	Adelie	Torgersen	39.1	18.7	181.0	3750
	1	Adelie	Torgersen	39.5	17.4	186.0	3800
	2	Adelie	Torgersen	40.3	18.0	195.0	3250
	3	Adelie	Torgersen	NaN	NaN	NaN	Na
	4	Adelie	Torgersen	36.7	19.3	193.0	3450
	339	Gentoo	Biscoe	NaN	NaN	NaN	Na
	340	Gentoo	Biscoe	46.8	14.3	215.0	4850
	341	Gentoo	Biscoe	50.4	15.7	222.0	5750
	342	Gentoo	Biscoe	45.2	14.8	212.0	5200
	343	Gentoo	Biscoe	49.9	16.1	213.0	5400
	344 ro	v 7 ool	Imno				>

Univariate Analysis

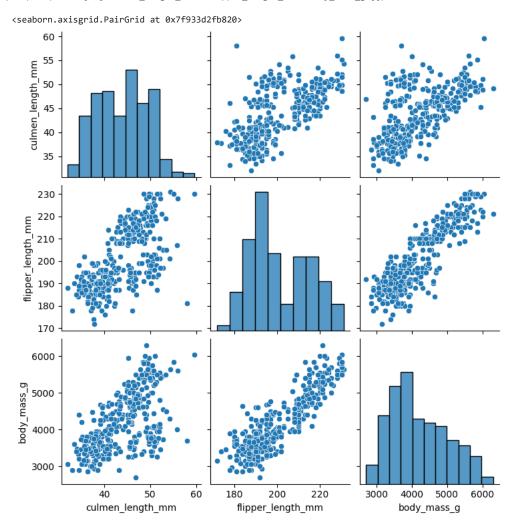
df.loc[:, ['culmen_length_mm']].plot()



Bi- Variate Analysis $\verb|sns.jointplot(x='flipper_length_mm', y='body_mass_g', data=df)|\\$ <seaborn.axisgrid.JointGrid at 0x7f933f4bee90>



Multi-Variate Analysis
sns.pairplot(df.loc[:,['culmen_length_mm','flipper_length_mm','body_mass_g']])



df.describe() # Descriptive statistics

count			flipper_length_mm	body_mass_g
count	242 000000			
	342.000000	342.000000	342.000000	342.000000
mean	43.921930	17.151170	200.915205	4201.754386
std	5.459584	1.974793	14.061714	801.954536
min	32.100000	13.100000	172.000000	2700.000000
25%	39.225000	15.600000	190.000000	3550.000000
species island culmen_ler culmen_der flipper_le body_mass_ sex	Fals Fals gth_mm Tru th_mm Tru ngth_mm Tru	se se ue ue	values in our data	set

Deleting rows with Null values
df=df.dropna()
df

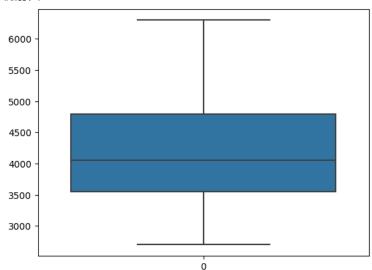
	species	island	culmen_length_mm	${\tt culmen_depth_mm}$	flipper_length_mm	body_mass_g	sex	
0	Adelie	Torgersen	39.1	18.7	181.0	3750.0	MALE	ıl.
1	Adelie	Torgersen	39.5	17.4	186.0	3800.0	FEMALE	
2	Adelie	Torgersen	40.3	18.0	195.0	3250.0	FEMALE	
4	Adelie	Torgersen	36.7	19.3	193.0	3450.0	FEMALE	
5	Adelie	Torgersen	39.3	20.6	190.0	3650.0	MALE	
338	Gentoo	Biscoe	47.2	13.7	214.0	4925.0	FEMALE	
340	Gentoo	Biscoe	46.8	14.3	215.0	4850.0	FEMALE	
341	Gentoo	Biscoe	50.4	15.7	222.0	5750.0	MALE	
342	Gentoo	Biscoe	45.2	14.8	212.0	5200.0	FEMALE	
343	Gentoo	Biscoe	49.9	16.1	213.0	5400.0	MALE	

334 rows × 7 columns

```
# Outlier detection and removal
q1 = df.body_mass_g.quantile(0.25) #Q1
q3 = df.body_mass_g.quantile(0.75) #Q3
IQR = q3-q1
upper_limit = q3+1.5*IQR
lower_limit =q1-1.5*IQR
df.median()
df['body_mass_g'] = np.where(df['body_mass_g']>upper_limit,4050,df['body_mass_g'])
df['body_mass_g'] = np.where(df['body_mass_g']<lower_limit,4050,df['body_mass_g'])
sns.boxplot(df.body_mass_g)</pre>
```

<ipython-input-34-8d2012ab2219>:7: FutureWarning: The default value of numeric_only in DataFrame.median is deprecated. In a
 df.median()

<Axes: >



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