Assignment-3

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```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns

df=pd.read_csv('/penguins_size.csv')
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
244 =	v 7 aali	imno				>

```
from matplotlib import rcParams
rcParams['figure.figsize']=8,8
fig,axes=plt.subplots(2,2)
sns.histplot(data=df['body_mass_g'],ax=axes[0,0])
sns.distplot(df['culmen_depth_mm'],ax=axes[1,1])
sns.barplot(x=df['culmen_length_mm'],y=df['culmen_length_mm'],ax=axes[0,1])
sns.boxplot(data=df['flipper_length_mm'],ax=axes[1,0])
```

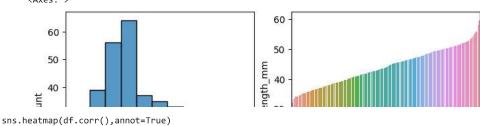
<ipython-input-3-5906f08a3de5>:5: UserWarning:

`distplot` is a deprecated function and will be removed in seaborn v0.14.0.

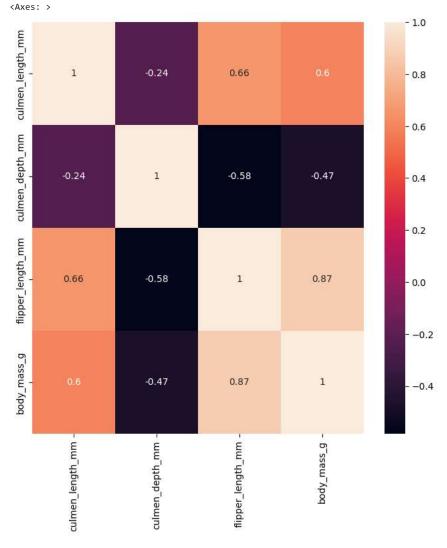
Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

For a guide to updating your code to use the new functions, please see $\underline{\text{https://gist.github.com/mwaskom/de44147ed2974457ad6372750bbe5751}}$

sns.distplot(df['culmen_depth_mm'],ax=axes[1,1])
<Axes: >



<ipython-input-4-8df7bcac526d>:1: FutureWarning: The default value of numeric_only in DataFrame.corr is deprecated. In a fu
sns.heatmap(df.corr(),annot=True)



df.describe()

		culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	===		
	count	342.000000	342.000000	342.000000	342.000000	11.		
	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			
	0=0/	00 005000	15 000000	100 000000	0550 000000			
<pre>df.isnull().any() df['culmen_length_mm'].fillna(df['culmen_length_mm'].median(),inplace=True) df['flipper_length_mm'].fillna(df['flipper_length_mm'].median(),inplace=True) df['culmen_depth_mm'].fillna(df['culmen_depth_mm'].median(),inplace=True) df['body_mass_g'].fillna(df['body_mass_g'].median(),inplace=True) most_frequent_category = df['sex'].mode()[0] df['sex'].fillna(most_frequent_category, inplace=True) df</pre>								

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	sex	\blacksquare	
0	Adelie	Torgersen	39.10	18.7	181.0	3750.0	MALE	11.	
1	Adelie	Torgersen	39.50	17.4	186.0	3800.0	FEMALE		
2	Adelie	Torgersen	40.30	18.0	195.0	3250.0	FEMALE		
3	Adelie	Torgersen	44.45	17.3	197.0	4050.0	MALE		
4	Adelie	Torgersen	36.70	19.3	193.0	3450.0	FEMALE		
339	Gentoo	Biscoe	44.45	17.3	197.0	4050.0	MALE		
340	Gentoo	Biscoe	46.80	14.3	215.0	4850.0	FEMALE		
341	Gentoo	Biscoe	50.40	15.7	222.0	5750.0	MALE		
342	Gentoo	Biscoe	45.20	14.8	212.0	5200.0	FEMALE		
343	Gentoo	Biscoe	49.90	16.1	213.0	5400.0	MALE		
344 rows × 7 columns									

sns.boxplot(df.body_mass_g)

```
<Axes: >
q1=df.body_mass_g.quantile(0.25)
\verb|q3=df.body_mass_g.quantile(0.75)|\\
IQR=q3-q1
upper_limit = q3+1.5*IQR
lower_limit =q1-1.5*IQR
df.median()
     <ipython-input-8-ef88e168ffba>:6: FutureWarning: The default value of numeric_only in DataFrame.median is deprecated. In a future versic
       df.median()
     culmen_length_mm
                              44.45
     culmen_depth_mm
flipper_length_mm
                             17.30
                             197.00
     body_mass_g
                            4050.00
     dtype: float64
\label{eq:df_body_mass_g'} $$ df['body_mass_g']>upper_limit, 30, df['body_mass_g']) $$
df
```

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	sex	
0	Adelie	Torgersen	39.10	18.7	181.0	3750.0	MALE	ıl.
1	Adelie	Torgersen	39.50	17.4	186.0	3800.0	FEMALE	
2	Adelie	Torgersen	40.30	18.0	195.0	3250.0	FEMALE	
3	Adelie	Torgersen	44.45	17.3	197.0	4050.0	MALE	
4	Adelie	Torgersen	36.70	19.3	193.0	3450.0	FEMALE	
339	Gentoo	Biscoe	44.45	17.3	197.0	4050.0	MALE	
340	Gentoo	Biscoe	46.80	14.3	215.0	4850.0	FEMALE	
341	Gentoo	Biscoe	50.40	15.7	222.0	5750.0	MALE	
342	Gentoo	Biscoe	45.20	14.8	212.0	5200.0	FEMALE	

16.1

213.0

5400.0

MALE

49.90

344 rows × 7 columns

Biscoe

343 Gentoo

sns.boxplot(df.body_mass_g)

```
<Axes: >
      6000
      5500
from sklearn.preprocessing import LabelEncoder
le=LabelEncoder()
df.sex=le.fit_transform(df.sex)
df.species=le.fit_transform(df.species)
df.island=le.fit_transform(df.island)
correlation_matrix = df.corr()
correlation_with_target = correlation_matrix['species']
print(correlation_with_target)
     species
                         1.000000
                         -0.635659
     island
     culmen_length_mm
                         0.728706
     culmen_depth_mm
                         -0.741282
                         0.850819
     flipper_length_mm
     body_mass_g
                          0.747547
                         -0.003823
     Name: species, dtype: float64
X=df.drop('body_mass_g',axis=1)
y=df['body_mass_g']
from sklearn.preprocessing import StandardScaler
sc=StandardScaler()
X_scaled= pd.DataFrame(sc.fit_transform(X),columns =X.columns)
X_scaled.head()
from sklearn.model_selection import train_test_split
X_train,X_test,y_train,y_test=train_test_split(X_scaled,y,test_size=0.2,random_state=42)
X_{train.shape}
     (275, 6)
X_test.shape
     (69, 6)
```

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