# **Step 1: Import the Required Libraries**

#### In [1]:

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
import warnings
warnings.filterwarnings("ignore")
```

# Step 2: Load the Dataset's

#### In [2]:

```
penguins_iter_df = pd.read_csv('penguins_lter.csv')
penguins_size_df = pd.read_csv('penguins_size.csv')
```

# **Step 3: Explore the Datasets**

#### In [3]:

```
# Display the first few rows of the DataFrames
penguins_iter_df.head()
```

### Out[3]:

	studyName	Sample Number	Species	Region	Island	Stage	Individual ID	Clutch Completion	Dat Eg
0	PAL0708	1	Adelie Penguin (Pygoscelis adeliae)	Anvers	Torgersen	Adult, 1 Egg Stage	N1A1	Yes	11/11/0
1	PAL0708	2	Adelie Penguin (Pygoscelis adeliae)	Anvers	Torgersen	Adult, 1 Egg Stage	N1A2	Yes	11/11/0
2	PAL0708	3	Adelie Penguin (Pygoscelis adeliae)	Anvers	Torgersen	Adult, 1 Egg Stage	N2A1	Yes	11/16/0
3	PAL0708	4	Adelie Penguin (Pygoscelis adeliae)	Anvers	Torgersen	Adult, 1 Egg Stage	N2A2	Yes	11/16/0
4	PAL0708	5	Adelie Penguin (Pygoscelis adeliae)	Anvers	Torgersen	Adult, 1 Egg Stage	N3A1	Yes	11/16/0
4									<b>&gt;</b>

## In [4]:

```
penguins_size_df.head()
```

## Out[4]:

	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
4						<b>&gt;</b>

### In [5]:

penguins\_iter\_df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 344 entries, 0 to 343
Data columns (total 17 columns):

#	Column	Non-Null Count	Dtype
0	studyName	344 non-null	object
1	Sample Number	344 non-null	int64
2	Species	344 non-null	object
3	Region	344 non-null	object
4	Island	344 non-null	object
5	Stage	344 non-null	object
6	Individual ID	344 non-null	object
7	Clutch Completion	344 non-null	object
8	Date Egg	344 non-null	object
9	Culmen Length (mm)	342 non-null	float64
10	Culmen Depth (mm)	342 non-null	float64
11	Flipper Length (mm)	342 non-null	float64
12	Body Mass (g)	342 non-null	float64
13	Sex	334 non-null	object
14	Delta 15 N (o/oo)	330 non-null	float64
15	Delta 13 C (o/oo)	331 non-null	float64
16	Comments	26 non-null	object

dtypes: float64(6), int64(1), object(10)

memory usage: 45.8+ KB

## In [6]:

```
penguins_size_df.info()
```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 344 entries, 0 to 343
Data columns (total 7 columns):

#	Column	Non-Null Count	Dtype
0	species	344 non-null	object
1	island	344 non-null	object
2	culmen_length_mm	342 non-null	float64
3	culmen_depth_mm	342 non-null	float64
4	flipper_length_mm	342 non-null	float64
5	body_mass_g	342 non-null	float64
6	sex	334 non-null	object

dtypes: float64(4), object(3)

memory usage: 18.9+ KB

## In [7]:

penguins\_iter\_df.describe()

## Out[7]:

	Sample Number	Culmen Length (mm)	Culmen Depth (mm)	Flipper Length (mm)	Body Mass (g)	Delta 15 N (o/oo)	Delta 13 C (o/oo)
count	344.000000	342.000000	342.000000	342.000000	342.000000	330.000000	331.000000
mean	63.151163	43.921930	17.151170	200.915205	4201.754386	8.733382	-25.686292
std	40.430199	5.459584	1.974793	14.061714	801.954536	0.551770	0.793961
min	1.000000	32.100000	13.100000	172.000000	2700.000000	7.632200	-27.018540
25%	29.000000	39.225000	15.600000	190.000000	3550.000000	8.299890	-26.320305
50%	58.000000	44.450000	17.300000	197.000000	4050.000000	8.652405	-25.833520
75%	95.250000	48.500000	18.700000	213.000000	4750.000000	9.172123	-25.062050
max	152.000000	59.600000	21.500000	231.000000	6300.000000	10.025440	-23.787670

#### In [8]:

```
penguins_size_df.describe()
```

### Out[8]:

	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g
count	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
50%	44.450000	17.300000	197.000000	4050.000000
75%	48.500000	18.700000	213.000000	4750.000000
max	59.600000	21.500000	231.000000	6300.000000

# **Step 4: Handle Missing Data**

```
In [9]:
```

```
penguins_size_df.isnull().sum()
```

## Out[9]:

```
species 0
island 0
culmen_length_mm 2
culmen_depth_mm 2
flipper_length_mm 2
body_mass_g 2
sex 10
```

dtype: int64

#### In [10]:

```
penguins_size_df = penguins_size_df.dropna()
```

#### In [11]:

```
penguins_size_df.isnull().sum()
```

#### Out[11]:

```
species 0
island 0
culmen_length_mm 0
culmen_depth_mm 0
flipper_length_mm 0
body_mass_g 0
sex 0
dtype: int64
```

#### In [12]:

```
# Check for missing values in Penguins iter dataset
print(penguins_iter_df.isnull().sum())
studyName
Sample Number
                          0
Species
                          0
Region
                          0
Island
                          0
                          0
Stage
Individual ID
                          0
Clutch Completion
                          0
                          0
Date Egg
Culmen Length (mm)
                          2
Culmen Depth (mm)
                          2
Flipper Length (mm)
                          2
Body Mass (g)
                          2
                         10
Sex
Delta 15 N (o/oo)
                         14
                         13
Delta 13 C (o/oo)
Comments
                        318
dtype: int64
In [13]:
penguins_iter_df = penguins_iter_df.dropna()
```

#### In [14]:

```
print(penguins_iter_df.isnull().sum())
studyName
                        0
Sample Number
                        0
                        0
Species
Region
                        0
                        0
Island
                        0
Stage
Individual ID
                        0
Clutch Completion
                        0
                        0
Date Egg
Culmen Length (mm)
                        0
Culmen Depth (mm)
                        0
                        0
Flipper Length (mm)
Body Mass (g)
                        0
                        0
Sex
Delta 15 N (o/oo)
                        0
                        0
Delta 13 C (o/oo)
Comments
dtype: int64
```

## **Step 5: Data Cleaning**

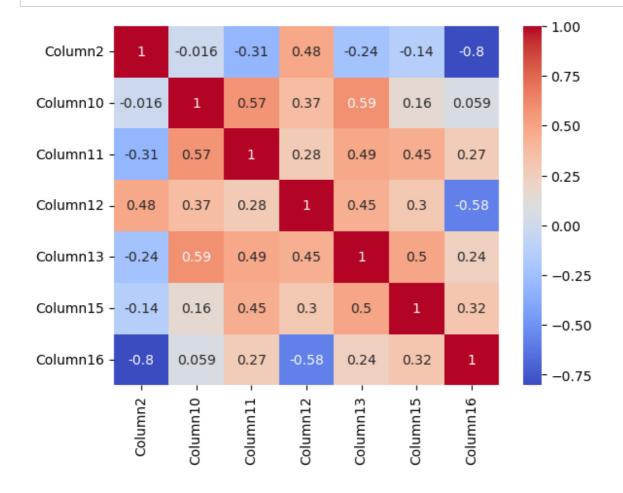
#### In [15]:

```
# Assign unique column names to Penguins Iter dataset
new_column_names = ['Column1', 'Column2', 'Column3', 'Column4', 'Column5', 'Column6', 'Column5_iter_df.columns = new_column_names
```

# **Step 6: Exploratory Data Analysis**

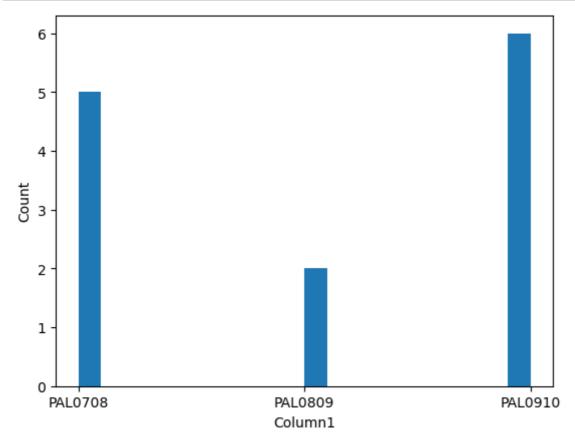
## In [16]:

```
correlation = penguins_iter_df.corr()
sns.heatmap(correlation, annot=True, cmap='coolwarm')
plt.show()
```



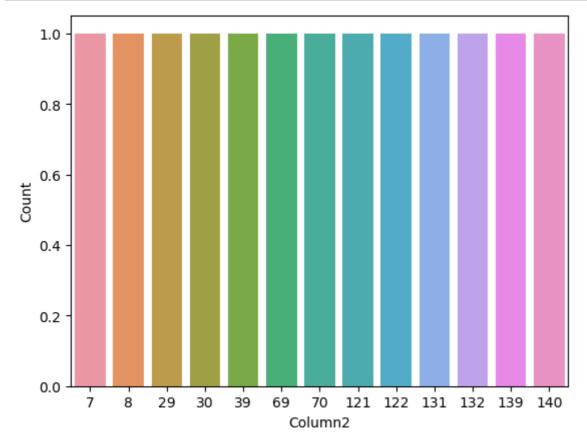
## In [17]:

```
# Plot Different Features
plt.hist(penguins_iter_df['Column1'], bins=20)
plt.xlabel('Column1')
plt.ylabel('Count')
plt.show()
```



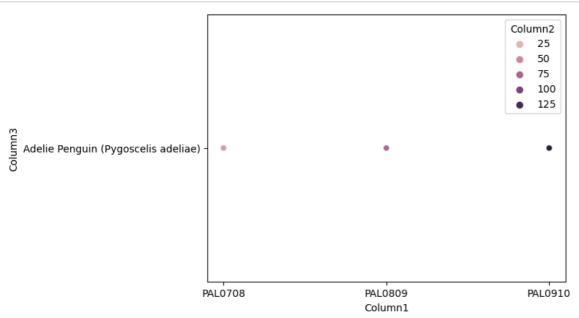
#### In [18]:

```
sns.countplot(x='Column2', data=penguins_iter_df)
plt.xlabel('Column2')
plt.ylabel('Count')
plt.show()
```



## In [19]:

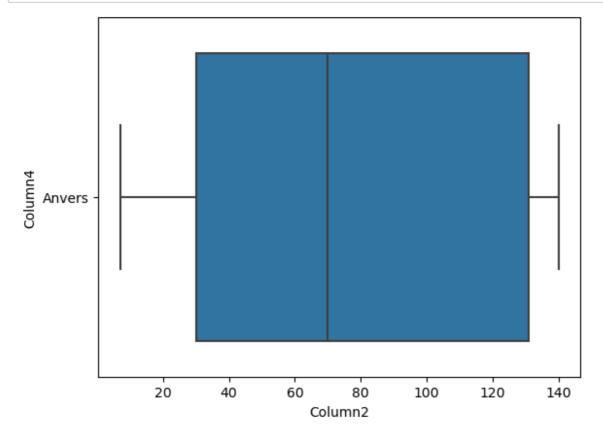
```
sns.scatterplot(x='Column1', y='Column3', hue='Column2', data=penguins_iter_df)
plt.xlabel('Column1')
plt.ylabel('Column3')
plt.show()
```



# **Detect Outliers / Missing Values**

## In [20]:

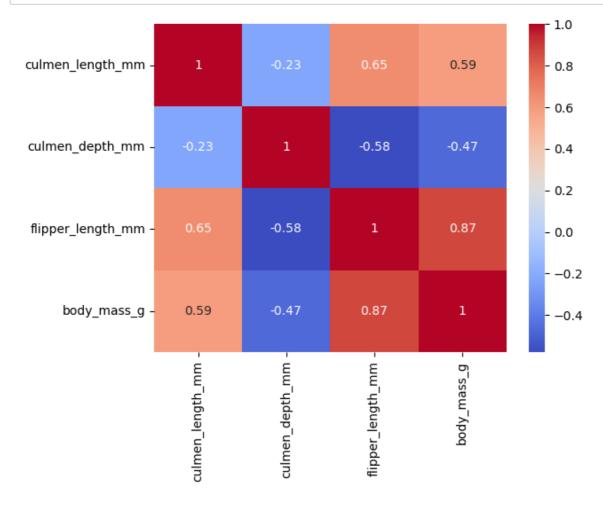
```
# Detect Outliers / Missing Values
sns.boxplot(x='Column2', y='Column4', data=penguins_iter_df)
plt.xlabel('Column2')
plt.ylabel('Column4')
plt.show()
```



## **Find Correlation Between Variables**

## In [21]:

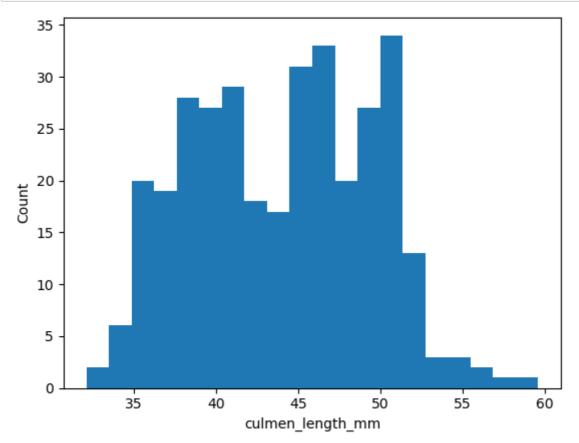
```
correlation = penguins_size_df.corr()
sns.heatmap(correlation, annot=True, cmap='coolwarm')
plt.show()
```



## **EDA ON PENGUINS SIZE**

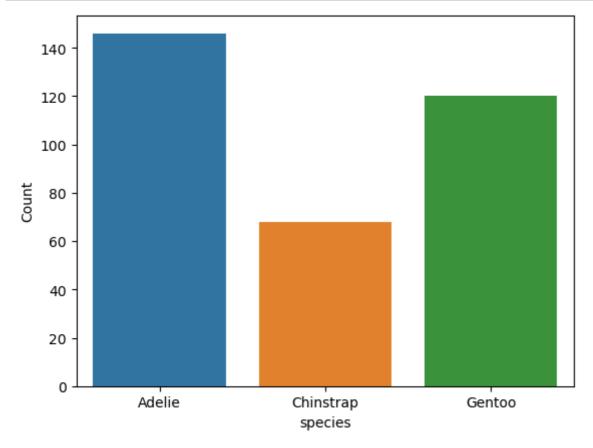
## In [22]:

```
# Histograms
plt.hist(penguins_size_df['culmen_length_mm'], bins=20)
plt.xlabel('culmen_length_mm')
plt.ylabel('Count')
plt.show()
```



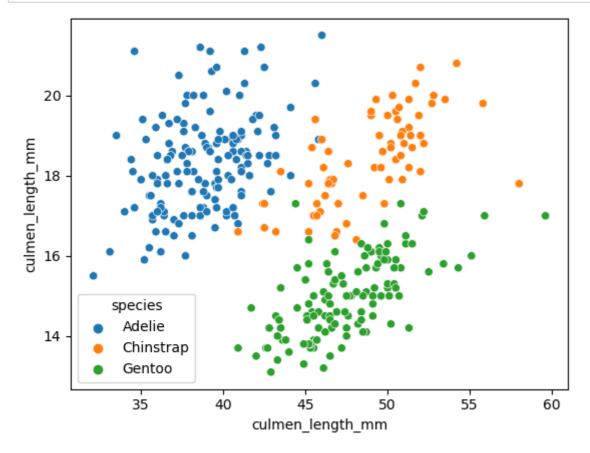
## In [23]:

```
# Bar plots
sns.countplot(x='species', data=penguins_size_df)
plt.xlabel('species')
plt.ylabel('Count')
plt.show()
```



## In [24]:

```
# Scatter plots
sns.scatterplot(x='culmen_length_mm', y='culmen_depth_mm', hue='species', data=penguins_s
plt.xlabel('culmen_length_mm')
plt.ylabel('culmen_length_mm')
plt.show()
```



# **Detect Outliers / Missing Values**

## In [25]:

```
# Detect outliers using box plots
sns.boxplot(x='species', y='body_mass_g', data=penguins_size_df)
plt.xlabel('species')
plt.ylabel('body_mass_g')
plt.show()
```

