## **Descriptive Analysis**

#### Introduction

The provided dataset contains descriptive statistics for several variables related to penguin species, including species, island, bill length, bill depth, flipper length, body mass, and sex. The primary target variable for analysis is species. This analysis aims to provide a comprehensive overview of the data, highlighting key metrics and their implications.

### **Species Distribution**

The species variable has three unique values: Adelie, Chinstrap, and Gentoo. The mode is Adelie, indicating that the majority of the data points belong to this species. The frequency distribution shows that Adelie has the highest frequency (152), followed by Gentoo (124), and then Chinstrap (68). This suggests that Adelie is the most abundant species in the dataset.

#### **Island Distribution**

The island variable has three unique values: Torgersen, Biscoe, and Dream. The mode is Biscoe, indicating that the majority of the data points are from this island. The frequency distribution shows that Biscoe has the highest frequency (168), followed by Dream (124), and then Torgersen (52). This suggests that Biscoe is the most represented island in the dataset.

#### Bill Length and Depth

The bill length and depth variables have similar distributions. The mean, median, and mode of both variables are relatively close, indicating that the data is skewed towards the middle values. The range is relatively small, suggesting that the data is concentrated around the mean. The variance and standard deviation of both variables are also relatively small, indicating that the data is relatively consistent.

The skewness of both variables is positive, indicating that the data is skewed to the right. This means that there are more values above the mean than below it. The kurtosis of both variables is negative, indicating that the data is more spread out than a normal distribution.

### Flipper Length and Body Mass

The flipper length and body mass variables have different distributions. The mean, median, and mode of flipper length are relatively close, indicating that the data is skewed towards the middle values. The range is relatively large, suggesting that the data is more spread out than bill length and depth.

The variance and standard deviation of flipper length are relatively large, indicating that the data is more variable than bill length and depth. The skewness of flipper length is positive, indicating that the data is skewed to the right. The kurtosis of flipper length is negative, indicating that the data is more spread out than a normal distribution.

The body mass variable has a larger range and variance than flipper length, indicating that the data is more spread out and variable. The skewness of body mass is positive, indicating that the data is skewed to the right. The kurtosis of body mass is negative, indicating that the data is more spread out than a normal distribution.

#### **Sex Distribution**

The sex variable has three unique values: Male, Female, and nan (not available). The mode is Male, indicating that the majority of the data points are male. The frequency distribution shows that Male has the highest frequency (168), followed by Female (165). The nan value is likely missing or unavailable data.

#### Conclusion

The analysis suggests that the dataset is dominated by Adelie penguins, with a majority of the data points coming from this species. The island distribution is also skewed towards Biscoe. The bill length and depth variables have similar distributions, with a small range and variance. The flipper length and body mass variables have different distributions, with a larger range and variance. The sex distribution is skewed towards Male.

Overall, the dataset provides a comprehensive overview of penguin characteristics, with a focus on species, island, bill length, bill depth, flipper length, body mass, and sex. The analysis highlights the key metrics and their implications, providing valuable insights into the penguin population.

# **Diagnostic Analysis**

Based on the provided descriptive statistics, I will perform a diagnostic analysis to identify potential issues, anomalies, or areas that require further investigation.

#### Species:

\* The mode of species is 'Adelie', which suggests that the majority of the dataset consists of Adelie species.

- \* The frequency distribution shows a relatively even distribution across the three species, with 'Adelie' having the highest frequency (152) and 'Chinstrap' having the lowest frequency (68).
- \* There are no obvious outliers or anomalies in the frequency distribution.

#### Island:

- \* The mode of island is 'Biscoe', which suggests that the majority of the dataset consists of samples from Biscoe island.
- \* The frequency distribution shows a relatively even distribution across the three islands, with 'Biscoe' having the highest frequency (168) and 'Torgersen' having the lowest frequency (52).
- \* There are no obvious outliers or anomalies in the frequency distribution.

### Bill Length (mm):

- \* The mean, median, and mode of bill length are relatively close, indicating a relatively normal distribution.
- \* The range (27.5) and variance (29.807) suggest a moderate level of dispersion, but not extreme.
- \* The skewness (0.053) and kurtosis (-0.876) suggest a slightly positively skewed and platykurtic distribution, respectively.
- \* There are no obvious outliers or anomalies in the distribution.

## Bill Depth (mm):

- \* The mean, median, and mode of bill depth are relatively close, indicating a relatively normal distribution.
- \* The range (8.4) and variance (3.899) suggest a moderate level of dispersion, but not extreme.
- \* The skewness (-0.143) and kurtosis (-0.906) suggest a slightly negatively skewed and platykurtic distribution, respectively.
- \* There are no obvious outliers or anomalies in the distribution.

#### Flipper Length (mm):

- \* The mean, median, and mode of flipper length are relatively close, indicating a relatively normal distribution.
- \* The range (59.0) and variance (197.731) suggest a moderate to high level of dispersion.
- \* The skewness (0.345) and kurtosis (-0.984) suggest a slightly positively skewed and platykurtic distribution, respectively.
- \* There are no obvious outliers or anomalies in the distribution.

### Body Mass (g):

- \* The mean, median, and mode of body mass are relatively close, indicating a relatively normal distribution.
- \* The range (3600.0) and variance (643131.077) suggest a high level of dispersion.
- \* The skewness (0.470) and kurtosis (-0.719) suggest a slightly positively skewed and platykurtic distribution, respectively.
- \* There are no obvious outliers or anomalies in the distribution.

#### Sex:

- \* The mode of sex is 'Male', which suggests that the majority of the dataset consists of male individuals.
- \* The frequency distribution shows a relatively even distribution across male and female individuals, with a single missing value (nan).
- \* The presence of a missing value in the sex variable may require further investigation to determine the cause and potential impact on analysis.

## **Target Variable (Species):**

\* The target variable, species, appears to have a relatively even distribution across the three species, with no obvious outliers or anomalies.

Overall, the diagnostic analysis suggests that the dataset appears to be relatively clean and free of obvious issues. However, the presence of a missing value in the sex variable may require further investigation. Additionally, the high level of dispersion in the body mass variable may warrant further exploration to determine the underlying causes and potential impact on analysis.

# **Descriptive Analysis in Tabular Format**

## species

Statistic	Value
mode	Adelie
unique_values	Adelie, Chinstrap, Gentoo
frequency_distribution	Adelie, Gentoo, Chinstrap

# island

Statistic	Value
mode	Biscoe
unique_values	Torgersen, Biscoe, Dream
frequency_distribution	Biscoe, Dream, Torgersen

# bill\_length\_mm

Statistic	Value
mean	43.9219298245614
median	44.45
mode	41.1
min	32.1
max	59.6
range	27.5
variance	29.807054329371816
std_dev	5.4595837139265315
iqr	9.2749999999999
25th_percentile	39.225
50th_percentile	44.45
75th_percentile	48.5
skewness	0.05311806699132413
kurtosis	-0.8760269663060134
outlier_percentage	0.0

# bill\_depth\_mm

Statistic	Value
mean	17.151169590643278
median	17.3
mode	17.0
min	13.1
max	21.5

range	8.4
variance	3.8998080122103893
std_dev	1.9747931568167816
iqr	3.099999999999
25th_percentile	15.6
50th_percentile	17.3
75th_percentile	18.7
skewness	-0.1434646251943698
kurtosis	-0.9068660903732537
outlier_percentage	0.0

# flipper\_length\_mm

Statistic	Value
mean	200.91520467836258
median	197.0
mode	190.0
min	172.0
max	231.0
range	59.0
variance	197.73179160021266
std_dev	14.061713679356888
iqr	23.0
25th_percentile	190.0
50th_percentile	197.0
75th_percentile	213.0
skewness	0.34568183286876963
kurtosis	-0.9842728861838852
outlier_percentage	0.0

# body\_mass\_g

Statistic	Value
mean	4201.754385964912

median	4050.0
mode	3800.0
min	2700.0
max	6300.0
range	3600.0
variance	643131.077326748
std_dev	801.9545356980956
iqr	1200.0
25th_percentile	3550.0
50th_percentile	4050.0
75th_percentile	4750.0
skewness	0.470329330480123
kurtosis	-0.7192218658321541
outlier_percentage	0.0

## sex

Statistic	Value
mode	Male
unique_values	Male, Female, nan
frequency_distribution	Male, Female