

Notebook:

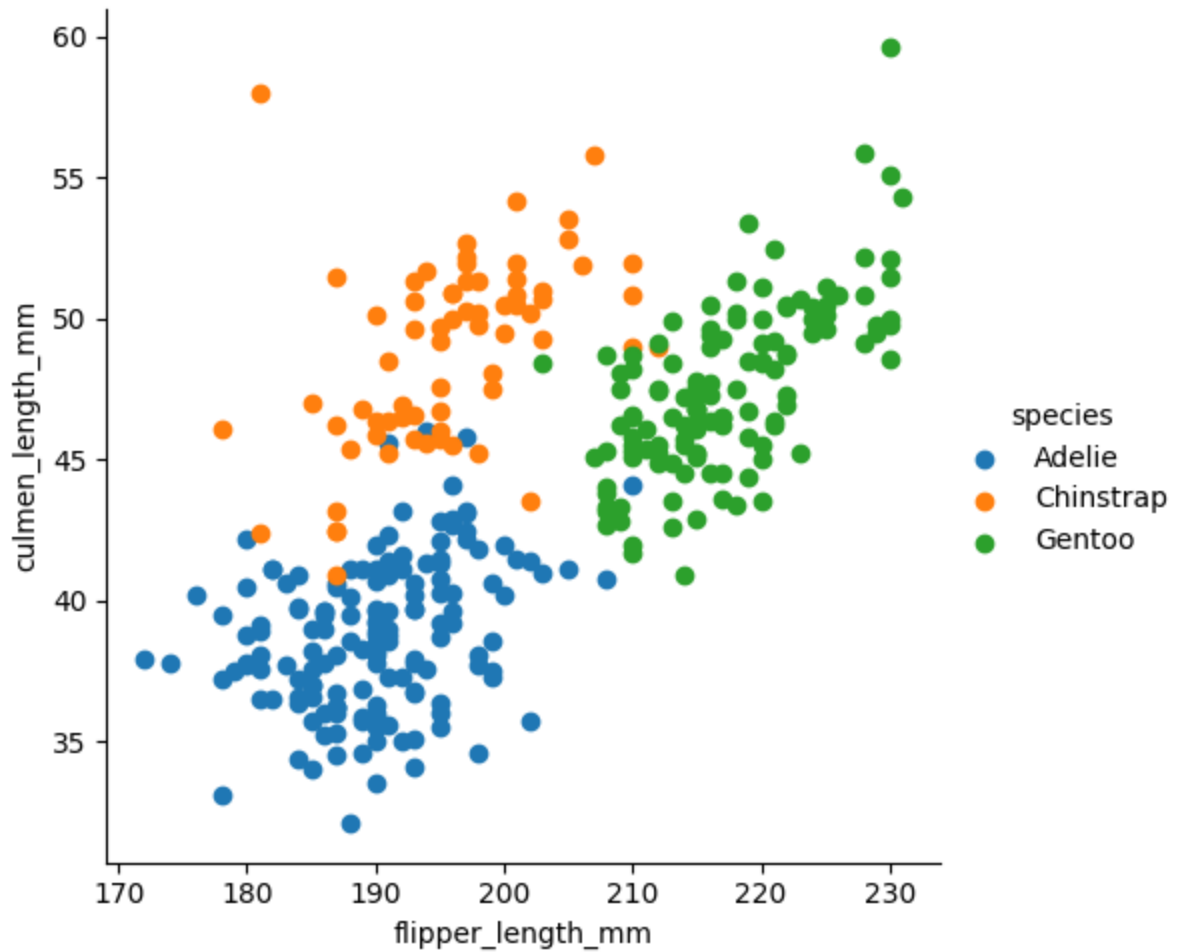
```
In [ ]: import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
%matplotlib inline
```

```
In [ ]: df = pd.read_csv('./penguins_size.csv')
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 [ ]: sns.FacetGrid(df, hue="species", height=5) \
.map(plt.scatter, "flipper_length_mm", "culmen_length_mm") \
.add_legend()
```

```
Out[ ]: <seaborn.axisgrid.FacetGrid at 0x7f849c6bd4d0>
```

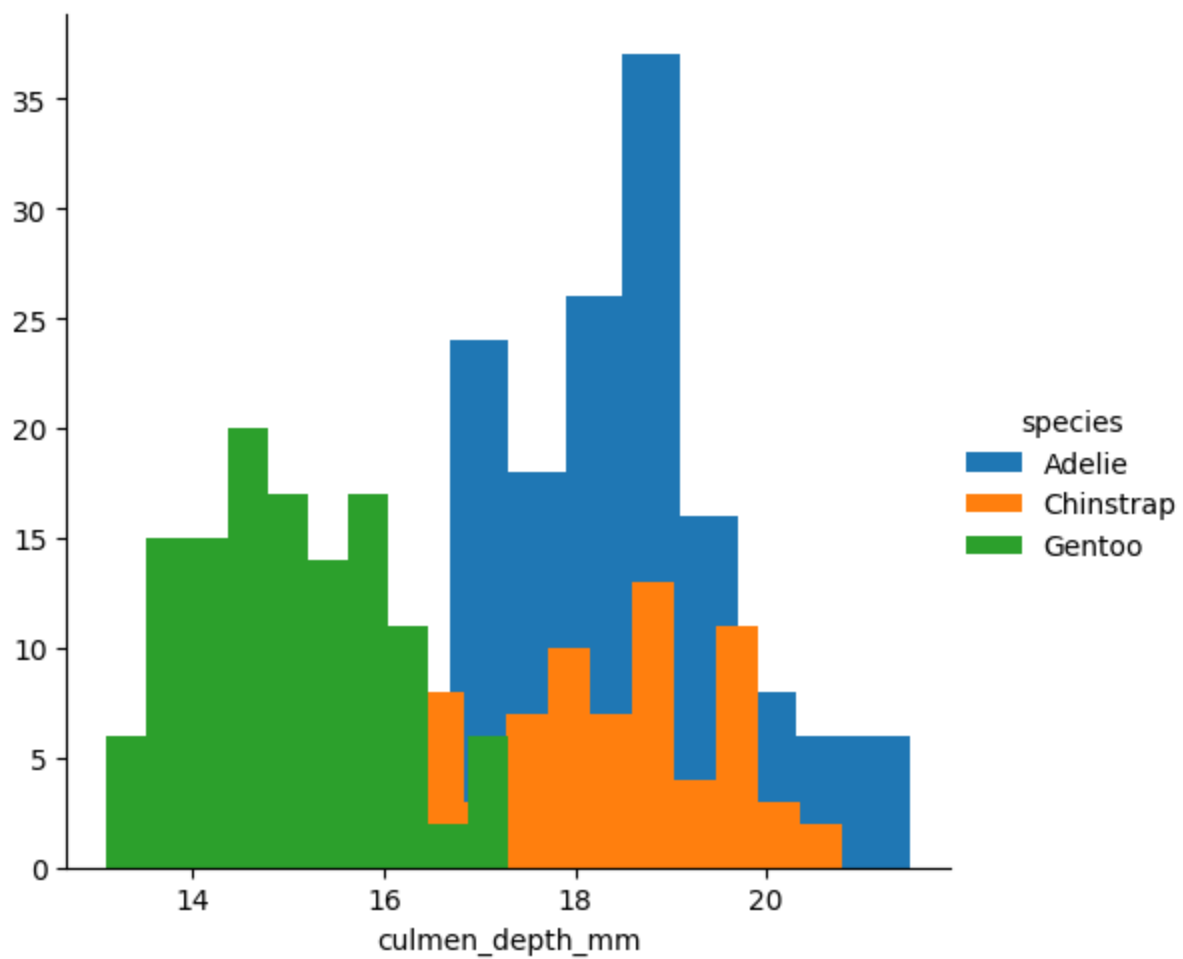


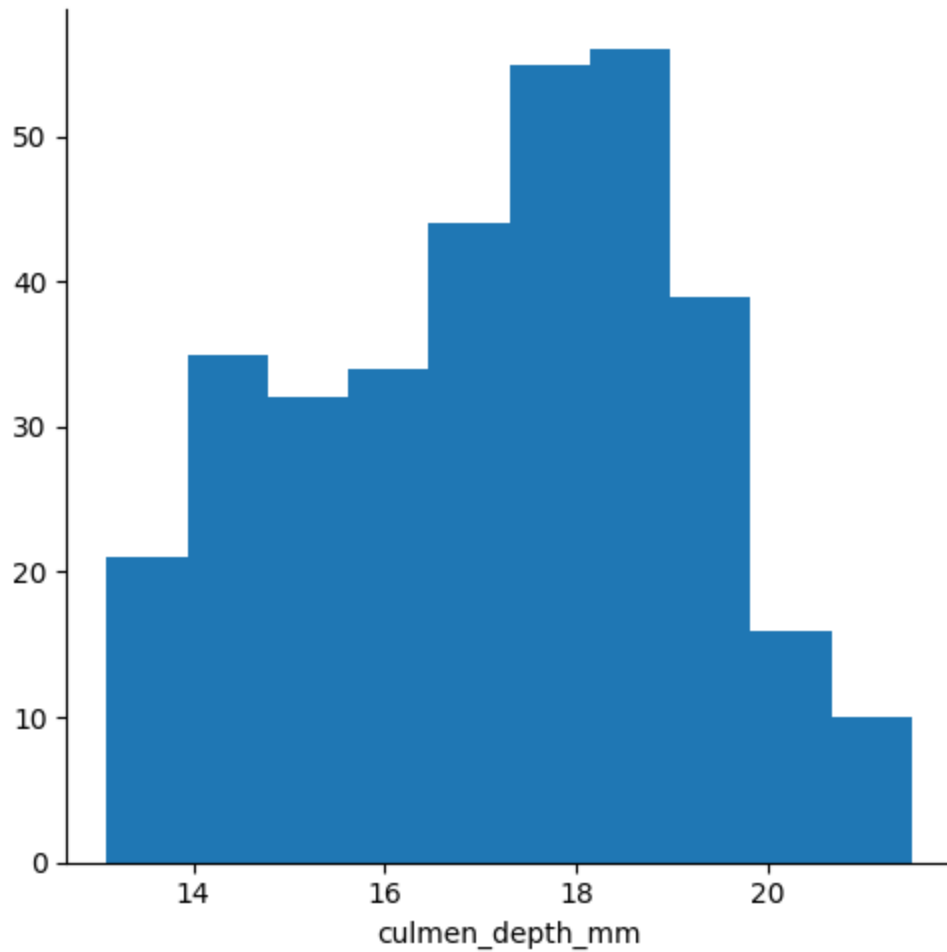
From this graph we can infer the following :

1. The ratio between flipper length and bill length is relatively similar
2. The ratio between flipper length and bill length is approximately 5.14
3. The data is similar to that of a regression problem

```
In [ ]: sns.FacetGrid(df, hue="species", height=5) \
        .map(plt.hist, "culmen_depth_mm") \
        .add_legend()
sns.FacetGrid(df, height=5) \
        .map(plt.hist, "culmen_depth_mm")
```

```
Out[ ]: <seaborn.axisgrid.FacetGrid at 0x7f8500237dd0>
```



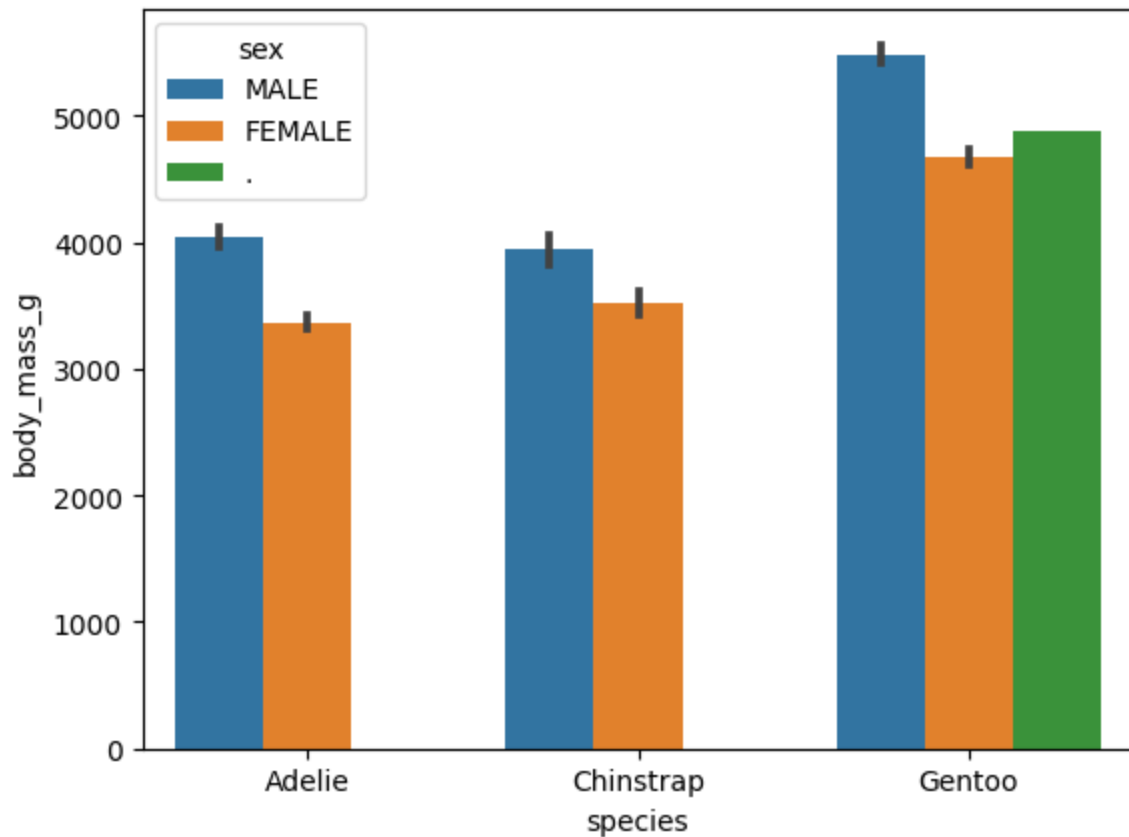


From this graph we can infer the following :

1. The bill depth has a substantial variance
2. The Adelie species has the largest bill depth amongst other species
3. The Chinstrap species has the smallest average bill depth

```
In [ ]: sns.barplot(df, x="species", y="body_mass_g", hue="sex")
```

```
Out[ ]: <Axes: xlabel='species', ylabel='body_mass_g'>
```

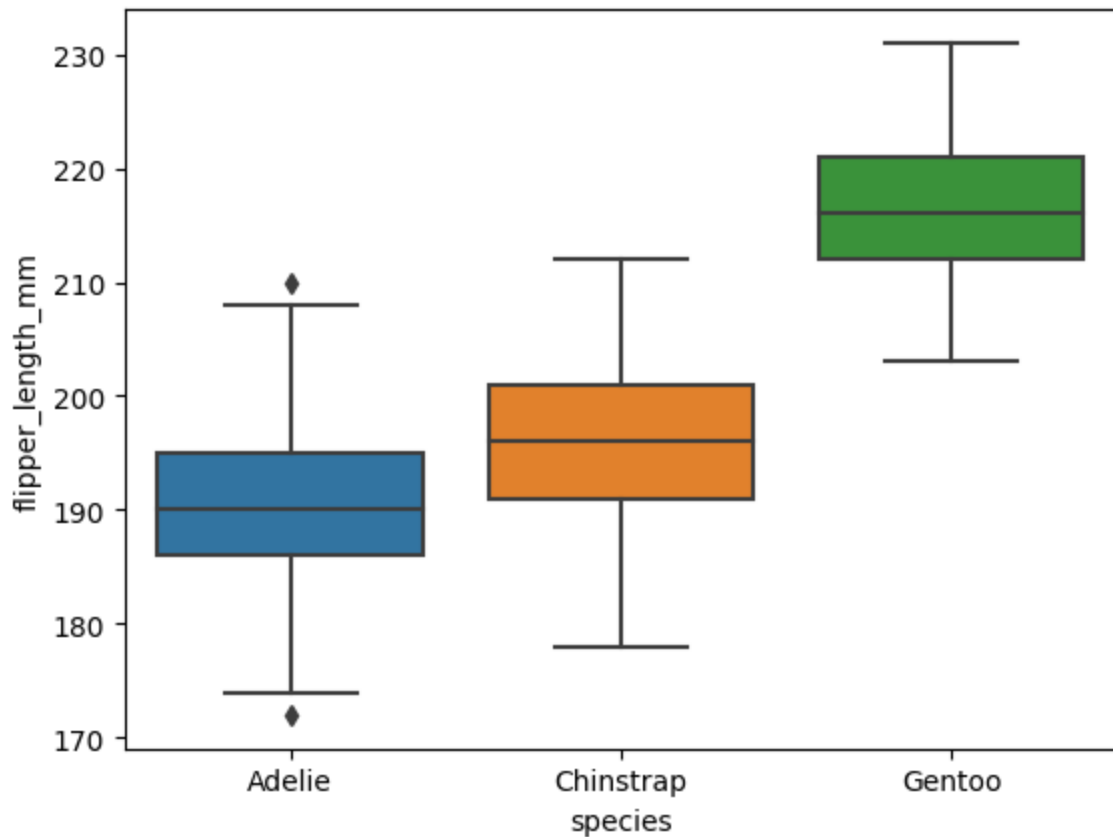


From this graph we can infer the following :

1. The males of all species have larger average body masses
2. The gentoo species have the largest body mass
3. The chinstrap has the smallest body mass on average
4. The gentoo species has 3 genders

```
In [ ]: sns.boxplot(df, x="species", y="flipper_length_mm")
```

```
Out[ ]: <Axes: xlabel='species', ylabel='flipper_length_mm'>
```

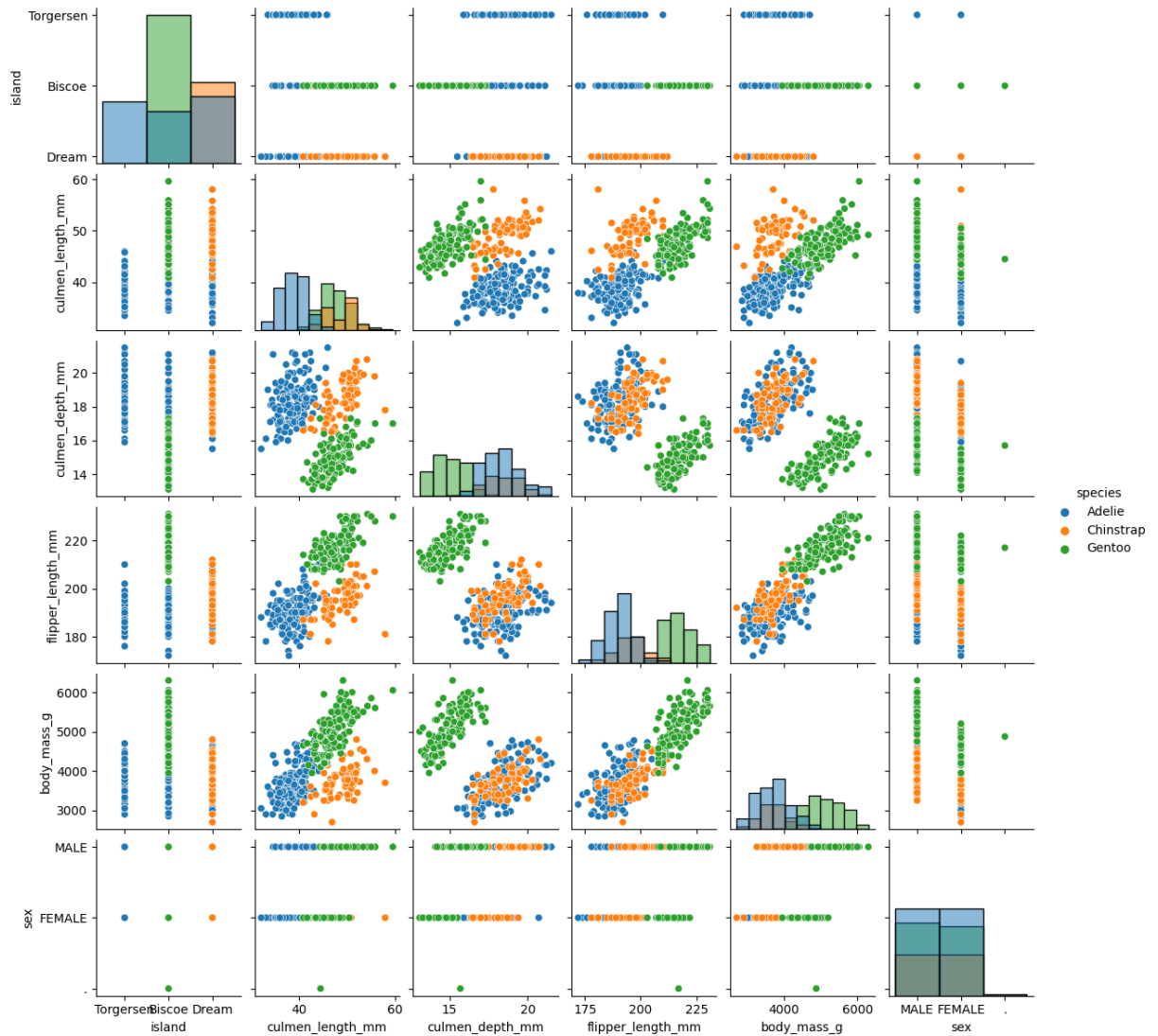


From this graph we can infer the following :

1. The gentoo species have the largest flipper lengths on average
2. The Adelie species has the smallest flipper length on average
3. The chinstrap species has an average flipper length, more on the short side

```
In [ ]: x=["island","culmen_length_mm","culmen_depth_mm","flipper_length_mm","body_m
y=["island","culmen_length_mm","culmen_depth_mm","flipper_length_mm","body_m
sns.pairplot(df,x_vars=x,y_vars=y,hue="species", height=2, diag_kind="hist")
```

```
Out[ ]: <seaborn.axisgrid.PairGrid at 0x7f8498e98d10>
```



From this graph we can infer the following :

1. The gentoo species are the largest among the 3
2. The Adelle species are the smallest on average
3. The males of all species are generally bigger