Initialize libraries and import the dataset

In [1]:

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

In [2]:

```
df = pd.read_csv("penguins_size.csv")
```

In [35]:

```
df.head()
```

Out[35]:

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g
0	Adelie	Torgersen	39.1	18.7	181.0	3750.0
1	Adelie	Torgersen	39.5	17.4	186.0	3800.0
2	Adelie	Torgersen	40.3	18.0	195.0	3250.0
4	Adelie	Torgersen	36.7	19.3	193.0	3450.0
5	Adelie	Torgersen	39.3	20.6	190.0	3650.0

In [4]:

df.shape

Out[4]:

(344, 7)

In [5]:

```
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
                        344 non-null
     island
                                        object
    culmen_length_mm
 2
                        342 non-null
                                        float64
 3
     culmen_depth_mm
                        342 non-null
                                        float64
     flipper_length_mm 342 non-null
                                        float64
 4
                                        float64
 5
                        342 non-null
     body_mass_g
 6
                        334 non-null
                                        object
     sex
dtypes: float64(4), object(3)
memory usage: 18.9+ KB
```

In [6]:

```
df.isna().sum()
```

Out[6]:

dtype: int64

10 rows is to be dropped as it contains null value

In [7]:

```
percent_row_drop = (10/df.shape[0])*100
print("percentage of rows dropped: ", percent_row_drop)
```

percentage of rows dropped: 2.9069767441860463

```
In [8]:
df = df.dropna()
df.info()
<class 'pandas.core.frame.DataFrame'>
Int64Index: 334 entries, 0 to 343
Data columns (total 7 columns):
    Column
                      Non-Null Count Dtype
    ----
                       -----
0
                       334 non-null
    species
                                      object
                      334 non-null
 1
    island
                                      object
 2
    culmen_length_mm 334 non-null
                                      float64
                                      float64
 3
    culmen_depth_mm
                      334 non-null
    flipper_length_mm 334 non-null
                                      float64
 4
 5
    body_mass_g
                       334 non-null
                                      float64
                                      object
    sex
                       334 non-null
dtypes: float64(4), object(3)
memory usage: 20.9+ KB
```

Checking for number of unique values for some columns

```
In [9]:

df['sex'].unique()

Out[9]:
    array(['MALE', 'FEMALE', '.'], dtype=object)

In [10]:

df['island'].unique()

Out[10]:
    array(['Torgersen', 'Biscoe', 'Dream'], dtype=object)
```

Removing data where sex = "."

```
In [11]:

df = df[df['sex'] != '.']
```

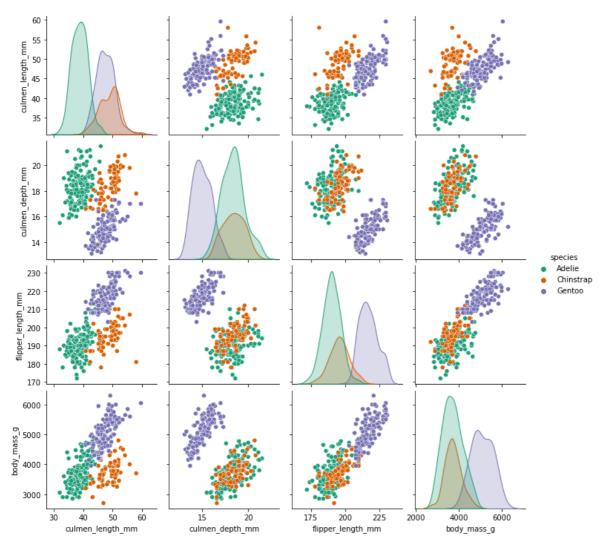
Now Lets Visalize the data

In [16]:

```
sns.pairplot(df,hue='species',palette='Dark2')
```

Out[16]:

<seaborn.axisgrid.PairGrid at 0x1b7089518c8>

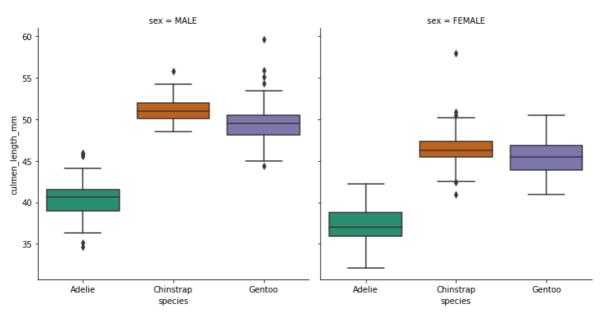


In [17]:

```
sns.catplot(x='species',y='culmen_length_mm',data=df,kind='box',col='sex',palette='Dark2')
```

Out[17]:

<seaborn.axisgrid.FacetGrid at 0x1b708867708>



So, now that we have the data, lets split it into X and y and also convert the categories to seperate columns containing 0 and 1

```
In [18]:
```

```
X = pd.get_dummies(df.drop('species',axis=1),drop_first=True)
y = df['species']
```

Split train and test data

In [19]:

```
from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=101)
```

Now train the model with Decision Tree Classifier

In [20]:

```
from sklearn.tree import DecisionTreeClassifier

Model = DecisionTreeClassifier()
Model.fit(X_train,y_train)
```

Out[20]:

DecisionTreeClassifier()

In [21]:

```
prediction = Model.predict(X_test)
```

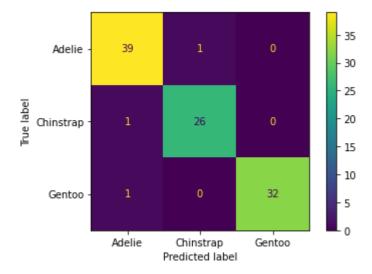
Now evaluate the predictions

In [25]:

```
from sklearn.metrics import confusion_matrix,classification_report,plot_confusion_matrix
confusion_matrix(y_test,prediction)
plot_confusion_matrix(Model,X_test,y_test)
```

Out[25]:

<sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x1b70b4ce
248>



We can see that almost all the predictions are correct except some are wrongly predicted

Lets get our classification report

In [27]:

print(classification_report(y_test,prediction))

	precision	recall	f1-score	support
Adelie	0.95	0.97	0.96	40
Chinstrap	0.96	0.96	0.96	27
Gentoo	1.00	0.97	0.98	33
accuracy			0.97	100
macro avg	0.97	0.97	0.97	100
weighted avg	0.97	0.97	0.97	100

Lets see how important each feature is in predicting the species

In [29]:

pd.DataFrame(index=X.columns,data=Model.feature_importances_,columns=['Feature Importance']

Out[29]:

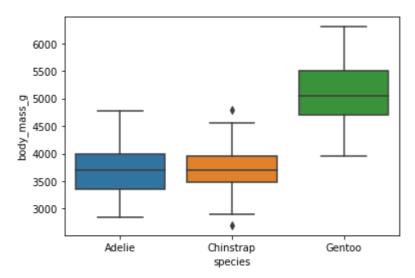
	Feature Importance
culmen_length_mm	0.333501
culmen_depth_mm	0.026964
flipper_length_mm	0.575758
body_mass_g	0.000000
island_Dream	0.038061
island_Torgersen	0.000000
sex_MALE	0.025717

In [30]:

```
sns.boxplot(x='species',y='body_mass_g',data=df)
```

Out[30]:

<AxesSubplot:xlabel='species', ylabel='body_mass_g'>



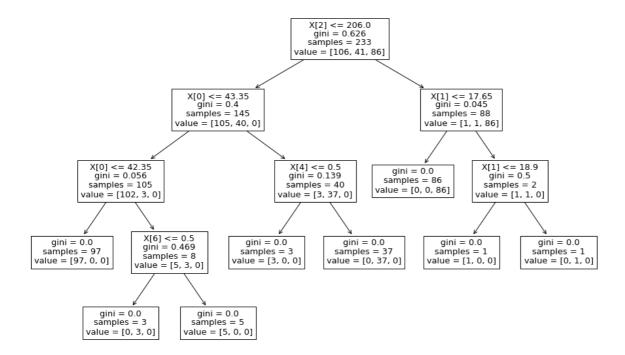
Above graph indates that the body mass has no importance in differenciating first 2 species hence it has 0 importance in table above the graph.

Finally, lets visualize the decision tree

In [34]:

```
from sklearn.tree import plot_tree

plt.figure(figsize=(16,10))
plot_tree(Model);
```



In []: