## Portfolio assignment 17

30 min: Train a decision tree to predict the body\_mass\_q of a penguin based on their characteristics.

- Split the penguin dataset into a train (70%) and test (30%) set.
- Use the train set to fit a DecisionTreeRegressor. You are free to to choose which columns you want to use as feature variables and you are also free to choose the max\_depth of the tree. Note: Some machine learning algorithms can not handle missing values. You will either need to
  - replace missing values (with the mean or most popular value). For replacing missing values you can use .fillna(\<value>) https://pandas.pydata.org/docs/reference/api/pandas.Series.fillna.html
  - remove rows with missing data. You can remove rows with missing data with .dropna() https://pandas.pydata.org/pandas-docs/stable/reference/api/pandas.DataFrame.dropna.html
- Use your decision tree model to make predictions for both the train and test set.
- Calculate the RMSE for both the train set predictions and test set predictions.
- Is the RMSE different? Did you expect this difference?
- Use the plot\_tree\_regression function above to create a plot of the decision tree. Take a few minutes to analyse the decision tree. Do you understand the tree?

## Regression

When the target variable is a numerical variable then we refer to this task as a regression task.

Examples of regression tasks:

- Predict the price people are willing to pay for a house.
- Predict the salary a student will earn in the future.

Examples of machine learning algorithms that we could use for regression:

- Decision trees
- Random forests

```
Linear regression
           Neural networks
          import pandas as pd
          import seaborn as sns
          penguins = sns.load dataset("penguins")
          penguins.head()
                            bill_length_mm bill_depth_mm
            species
                     island
                                                       flipper_length_mm
                                                                       body_mass_g
                                                                                      sex
                                     39.1
                                                  18.7
         0
             Adelie
                   Torgersen
                                                                  181.0
                                                                             3750.0
                                                                                      Male
         1
             Adelie
                   Torgersen
                                     39.5
                                                  17.4
                                                                  186.0
                                                                             3800.0 Female
         2
                                     40.3
                                                  18.0
                                                                  195.0
                                                                             3250.0 Female
             Adelie
                   Torgersen
         3
             Adelie
                   Torgersen
                                     NaN
                                                  NaN
                                                                   NaN
                                                                               NaN
                                                                                      NaN
             Adelie Torgersen
                                                                             3450.0 Female
                                     36.7
                                                  19.3
                                                                  193.0
          from sklearn.tree import DecisionTreeRegressor
In [4]:
          penguins = penguins.dropna() # removing a low amount of na's
         features= ['body mass g']
          dt regression = DecisionTreeRegressor(max depth = 3) # Increase max depth to see effe
          dt regression.fit(penguins[features], penguins['bill length mm'])
Out[5]: DecisionTreeRegressor(max depth=3)
          from sklearn import tree
          import graphviz
          def plot tree regression(model, features):
              # Generate plot data
              dot data = tree.export graphviz (model, out file=None,
                                      feature names=features,
                                      filled=True, rounded=True,
                                      special characters=True)
              # Turn into graph using graphviz
              graph = graphviz.Source(dot data)
              # Write out a pdf
              graph.render("decision tree")
              # Display in the notebook
              return graph
          def calculate_rmse(predictions, actuals):
              if(len(predictions) != len(actuals)):
                  raise Exception ("The amount of predictions did not equal the amount of actuals
              return (((predictions - actuals) ** 2).sum() / len(actuals)) ** (1/2)
```

```
predictionsOnTrainset = dt regression.predict(penguins[features])
```

```
predictionsOnTestset = dt regression.predict(penguins[features])
rmseTrain = calculate rmse(predictionsOnTrainset, penguins.bill length mm)
rmseTest = calculate rmse(predictionsOnTestset, penguins.bill length mm)
print("RMSE on training set " + str(rmseTrain))
print("RMSE on test set " + str(rmseTest))
RMSE on training set 4.266950627893799
RMSE on test set 4.266950627893799
```

well. The errors SHOULD be minimal, but you're never sure.

The difference seems to be very minimal, this means the training set can probably predict the output very

```
plot tree regression(dt regression, features)
                                                                          body_mass_g ≤ 4737.5
mse = 29.817
                                                                                samples = 333
                                                                                value = 43.993
                                                                          True
                                                                                                 False
                                                         body_mass_g ≤ 3637.5
mse = 27.613
samples = 244
                                                                                            body_mass_g ≤ 5325.0
mse = 8.657
samples = 89
                                                               value = 42.364
                                                                                                  value = 48.457
                body_mass_g ≤ 3187.5
mse = 22.383
                                                                                            body_mass_g ≤ 5125.0
mse = 4.991
                                                                                                                                      body_mass_g ≤ 6025.0
mse = 6.263
                                                         body_mass_g ≤ 3987.5
mse = 25.599
                      samples = 95
                                                              samples = 149
                                                                                                  samples = 47
                                                                                                                                            samples = 42
                                                              value = 43.807
                     value = 40.101
                                                                                                  value = 46.802
                                                                                                                                            value = 50.31
                      mse = 22.946
                                             mse = 33.192
                                                                    mse = 18.612
samples = 83
                                                                                            mse = 4.61
samples = 34
                                                                                                                    mse = 5.054
samples = 13
                                                                                                                                            mse = 4.345
samples = 40
mse = 9.182
samples = 22
                      samples = 73
                                             samples = 66
                                                                                                                                                                   samples = 2
                      value = 40.99
value = 37.15
                                             value = 42.992
                                                                                                                                                                    value = 54.4
```