## Copy\_of\_sklearn\_decisiontree\_baskin\_robbins

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## 1 Classification Tree

- Baskin Robbins nutritional information: http://www.baskinrobbins.ca/nutritional-information/
- ref: https://scikit-learn.org/stable/modules/generated/sklearn.tree.DecisionTreeClassifier.html

```
[]: import pandas as pd
     import seaborn as sns
     import numpy as np
     import matplotlib.pyplot as plt
     %matplotlib inline
     # %config InlineBackend.figure_format = 'retina'
     %config InlineBackend.figure_format = 'svg'
[]: print(f"pandas version: {pd._version_}")
     print(f"numpy version: {np.__version__}")
     print(f"seaborn version: {sns.__version__}}")
[]: url = "https://github.com/sophalITC/datasci/raw/master/baskin_robbins.csv"
     df = pd.read_csv(url)
     df.head()
[]: df.info()
[]: df.columns
[]: cols = [
        "Calories",
         "Total Fat (g)",
         "Trans Fat (g)",
         "Carbohydrates (g)",
         "Sugars (g)",
        "Protein (g)",
[]: fig, ax = plt.subplots(nrows=2, ncols=3, figsize=(20, 9))
     ax = ax.ravel()
     for i, col in enumerate(cols):
```

```
sns.violinplot(x="Category", y=col, data=df, ax=ax[i])
```

## 1.1 Draw Decision Tree

```
[]: from sklearn.model_selection import train_test_split
     from sklearn.tree import DecisionTreeClassifier
     from sklearn.metrics import (
        confusion_matrix,
        ConfusionMatrixDisplay,
        classification_report,
        accuracy_score,
        precision_score,
        recall_score,
        precision_recall_fscore_support,
        f1_score,
[]: df.columns
[]: cols = [
        "Calories",
         "Total Fat (g)",
        "Trans Fat (g)",
         "Carbohydrates (g)",
         "Sugars (g)",
         "Protein (g)",
     ]
[]: X = df[cols] # features
     y = df["Category"] # label
[]: test_size = 0.2
     X_train, X_test, y_train, y_test = train_test_split(
        X, y, test_size=test_size, stratify=y, random_state=7
[]: model = DecisionTreeClassifier(criterion="gini") # qini is a default
     # model=DecisionTreeClassifier(criterion='entropy')
     model.fit(X_train, y_train)
[]: X_train.columns
[]: model.feature_importances_
```

```
[]: fs = pd.Series(model.feature_importances_, index=X_train.columns).sort_values(
         ascending=False
     )
     fs
[]: y_train.value_counts()
    1.1.1 tree diagram (better way)
       • install graphviz on Windows
      1. install graphviz: pip install -U graphviz
      2. download graphviz: https://www2.graphviz.org/Packages/stable/windows/10/msbuild/Release/Win32/
      3. extract and copy to: C:\Program Files (x86)\Graphviz\bin
      4. add C:\Program Files (x86)\Graphviz\bin to PATH environment variable
[]: clf = DecisionTreeClassifier(criterion="gini")
     clf.fit(X_train, y_train)
[]: clf.__dict__
[]: clf.__dict__["tree_"]
[]: type(clf)
[]: clf.__dict__["classes_"]
[]: from sklearn.tree import export_graphviz
     from graphviz import Digraph, Source
     def view_tree(model, X, save_tree_img=False):
         estimators = clf
         dot_graph = export_graphviz(
             estimators,
             feature_names=X.columns,
             class_names=model.__dict__["classes_"],
             rounded=True,
             proportion=False,
             precision=2,
             filled=True,
         )
               with open(f'tree{tree_index}.dot') as f:
                   dot graph = f.read()
         g = Source(dot_graph)
         if save_tree_img:
             g.render(f"tree", format="png", view=False, cleanup=True)
```

return g

[]: view\_tree(clf, X\_test, save\_tree\_img=False)

Gini impurity is a measure of how often a randomly chosen element from the set would be incorrectly labeled if it was randomly labeled according to the distribution of labels in the subset. ... It reaches its minimum (zero) when all cases in the node fall into a single target category. (https://en.wikipedia.org/wiki/Decision\_tree\_learning#Gini\_impurity)

$$\mathbf{I}_G(p) = \sum_{i=1}^J \left( p_i \sum_{k \neq i} p_k \right) = \sum_{i=1}^J p_i (1-p_i) = \sum_{i=1}^J (p_i - p_i^{\ 2}) = \sum_{i=1}^J p_i - \sum_{i=1}^J p_i^{\ 2} = 1 - \sum_{i=1}^J p_i^{\ 2}$$

gini value at the top level

$$gini = 1 - \left( \left( \frac{7}{56} \right)^2 + \left( \frac{23}{56} \right)^2 + \left( \frac{26}{56} \right)^2 \right) = .6$$

## 1.2 3D scatter

%matplotlib inline

# # switch back to inline mode

# %pylab qt

```
ax.set_ylabel("Total Fat (g)")
ax.set_zlabel("Sugars (g)")

[]: predicted = model.predict(X_test)
predicted

[]: pd.crosstab(y_test, predicted)
```