### **Machine Learning Package:**

### **Package Used: Scikit Learn**

I have used **Scikit Learn** for implementing the given task chosen because of the following reason:

- It's easy to use even for beginners and a great choice for simpler data analysis tasks.
- Can understand how the data is being analysed and users have great control over data to get different outcomes.
- Also, it will be good start for me to work on ML tasks with python coding involved in it.

### Features [a]:

- **Consistency:** All objects share a common interface drawn from a limited set of methods, with consistent documentation.
- **Inspection:** All specified parameter values are exposed as public attributes.
- Limited object hierarchy: Only algorithms are represented by Python classes; datasets are represented in standard formats (NumPy arrays, Pandas DataFrames, SciPy sparse matrices) and parameter names use standard Python strings.
- **Composition:** Many machine learning tasks can be expressed as sequences of more fundamental algorithms, and Scikit-Learn makes use of this wherever possible.
- **Sensible defaults:** When models require user-specified parameters, the library defines an appropriate default value

### **Data Preparation:**

#### Data preparation before feeding data into algorithm:

- 1. Copy hazelnut.txt data into hazelnut.csv. Transpose the data (to correctly arrange rows and columns for the headers provided)
- Add headers to each column in the order as specified in the assignment1.pdf document.

### Data preparation after feeding data into algorithm:

- 1. **Read csv** file and store it in a data frame (using pandas).
- 2. **Drop 'sample\_id'** (since it contains only unique ID and doesn't contribute in deciding the target) and **'variety'** (since it the target) column, and store the remaining data into X (now X contains attributes)
- 3. Load the values of column 'variety' into 'y' (y contains the target data)
- 4. Code Implementation:

```
import pandas as pd
hazelnut = pd.read_csv('./hazelnut.csv')
print(hazelnut.head())
X = hazelnut.drop(['sample_id','variety'],axis=1)
y = hazelnut['variety'].values
```

References: [a]: https://jakevdp.github.io/PythonDataScienceHandbook/05.02-introducing-scikit-learn.html

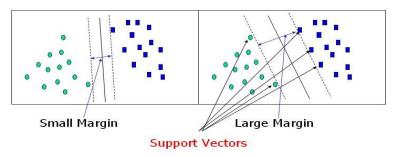
## **Classification Algorithms:**

### Algorithm 1: K Nearest Neighbors (k-NN) [1]

- The principle behind nearest neighbor method is to find a predefined number of training samples closest in distance to the new point and predict the label from these.
- k-NN is a type of instance-based learning, or lazy learning, where the function is only approximated locally, and all computation is deferred until classification.
- The **input consists of the k closest training examples** in the feature space.
- The output is a class membership.
- An object is classified by a plurality vote of its neighbors, with the object being assigned to the class most common among its k nearest neighbors (k is a positive integer, typically small).
- If k = 1, then the object is simply assigned to the class of that single nearest neighbor.
- Choosing the optimal value for K is best done by first inspecting the data.
- In general, a large K value is more precise as it reduces the overall noise but there is no guarantee.
- Cross-validation is another way to retrospectively determine a good K value by using an independent dataset to validate the K value.
- Historically, the optimal K for most datasets has been between 3-21.

## Algorithm 2: Support Vector Machines (SVM) [2][3]

- The objective of the support vector machine algorithm is to find a hyperplane in an N-dimensional space(N - the number of features) that distinctly classifies the data points.
- An SVM model is a **representation of the examples as points in space**, mapped so that the examples of the separate categories are **divided by a clear gap that is as wide as possible**.
- A good separation is achieved by the hyperplane that has the largest distance to the nearest training-data point of any class (so-called functional margin), since in general the larger the margin, the lower the generalization error of the classifier.
- New examples are then mapped into that same space and predicted to belong to a category based on the side of the gap on which they fall.
- An SVM is a discriminative classifier formally defined by a separating hyperplane.
- In other words, given labelled training data (supervised learning), the algorithm outputs an optimal hyperplane which categorizes new examples.
- In **two-dimensional space this hyperplane is a line** dividing a plane in two parts where in each class lay in either side.



https://towardsdatascience.com/support-vector-machine-introduction-to-machine-learning-algorithms-934a444fca47

#### References:

[1]: https://en.wikipedia.org/wiki/K-nearest\_neighbors\_algorithm

[2]: https://en.wikipedia.org/wiki/Support-vector machine

[3]: https://medium.com/machine-learning-101/chapter-2-svm-support-vector-machine-theory-f0812effc72

### 10-fold cross-validation:

### **Algorithm 1:** K Nearest Neighbors (k-NN)

- To perform 10-fold cross validation, the value of k is taken as 3
- For each value fold in 10-fold cross validation, and the prediction accuracy is stored in list.
- Below code shows the implementation of 10-fold cross validation and the corresponding graph.
- Implementation takes the complete data prepared in previous step.

```
from sklearn.model_selection import cross_val_score
#create a KNN model
knn_cv = KNeighborsClassifier(n_neighbors=3)
#train model with cv of 10
cv_scores = cross_val_score(knn_cv, X, y, cv=10)
print("Cross Validation scores: \n",cv_scores)
# plot the graph for each fold in knn
cv_range = (1,2,3,4,5,6,7,8,9,10)
plt.plot(cv_range,cv_scores)
plt.xlabel('Number of folds')
plt.ylabel('10-fold cross validation accuracy')
```

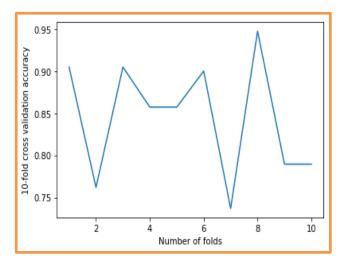


image source:

#### 10-fold Cross Validation score for k=3:

```
print("Cross Validation scores: \n",cv_scores)
```

```
Cross Validation scores:

[0.9047619 0.76190476 0.9047619 0.85714286 0.85714286 0.9

0.73684211 0.94736842 0.78947368 0.78947368]
```

By taking the mean of 10-fold Cross Validation score, accuracy of the model can be measured as below:

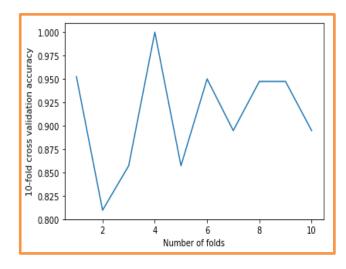
```
mean = format(np.mean(cv_scores))
mean = float(mean)*int(100)
print("Accuracy of the model after 10-fold cross validation: %0.2f"%mean,"%")
```

```
Accuracy of the model after 10-fold cross validation: 84.49 %
```

### **Algorithm 2: Support Vector Machines (SVM)**

- To perform 10-fold cross validation, linear kernel is used.
- A penalty parameter is introduced for the error term
- Below code shows the implementation of 10-fold cross validation and the corresponding graph.
- Implementation takes the complete data prepared in previous step.

```
import matplotlib.pyplot as plt
from sklearn.svm import SVC
from sklearn.model_selection import cross_val_score
# select kernel as linear with penalty parameter C
clf = SVC(kernel='linear', C=1)
# train model with cv of 10
cv_scores = cross_val_score(clf, X, y, cv=10)
# plot the graph for each fold in knn
cv_range = (1,2,3,4,5,6,7,8,9,10)
plt.plot(cv_range,cv_scores)
plt.xlabel('Number of folds')
plt.ylabel('10-fold cross validation accuracy')
```



#### 10-fold Cross Validation score:

```
print("Cross Validation scores: \n",cv_scores)
```

```
Cross Validation scores:

[0.95238095 0.80952381 0.85714286 1. 0.85714286 0.95

0.89473684 0.94736842 0.94736842 0.89473684]
```

By taking the mean of 10-fold Cross Validation score, accuracy of the model can be measured as below:

```
mean = cv_scores.mean()
mean = float(mean) * int(100)
print("Accuracy: %0.2f" %mean,"%")

Accuracy: 91.10 %
```

### **Conclusion:**

The results and observations show that **SVMs are a more reliable classifier** than k-NN.

Algorithm	Training data	Accuracy
k-NN	201	8445
SVM	201	91.10

- KNN is less computationally intensive than SVM.
- Since, k-NN is easy to implement, the classification of Multi-class data should be done with k-NN.
- If data is homogenous to look at, one might be able to classify better by putting in a kernel into the SVM.
- For most practical problems, KNN is a bad choice because it scales badly, if there are a million labelled examples, it would take a long time (linear to the number of examples) to find K nearest neighbours