

#### **GUIDE ON CLASSICAL MACHINE LEARNING**

**About the Tutorial:** Today's Artificial Intelligence (AI) has far surpassed the hype of blockchain and quantum computing. The developers now take advantage of this in creating new Machine Learning models and to re-train the existing models for better performance and results. This tutorial will give an introduction to machine learning and its implementation in Artificial Intelligence.

**Audience:** This tutorial has been prepared for professionals aspiring to learn the complete picture of Machine Learning and Al. This tutorial caters to the learning needs of both the novice learners and experts, to help them understand the concepts and implementation of Al

**Prerequisites:** The learners of this tutorial are expected to know the basics of Python programming. Besides, they need to have a solid understanding of computer programing and fundamentals. If you are new to this arena, we suggest you pick up tutorials based on these concepts first, before you embark on with Machine Learning.

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## **Course Curriculum**

- 1. Introduction to Machine Learning, AI & Data Science
- 2. Use Cases
- 3. Python Basics
- 4. Alternatives to Python for Data Science
- 5. Essential Modules/Libraries
- 6. Types of Learning
- 7. Classification Algorithms
  - a. k-nn
  - b. Decision Trees
  - c. Random Forest
  - d. Naive Bayes
  - e. SVMs
  - f. Logistic Regression
- 8. Regression Algorithms
  - a. Simple Linear Regression
  - b. Multiple Linear Regression
  - c. Polynomial Regression
- 9. Clustering Algorithms
  - a. K-Means Clustering
  - b. Hierarchical Clustering
  - c. Mean-Shift Algorithm
- 10. Association Rule Learning
- 11. Ensemble Techniques
- 12. Time Series Analysis
- 13. Introduction to Deep Learning
- 14. Dimensionality Reduction

Let's have a glance at some of the chapters of this course below:



# **Multiple Linear Regression**

The difference between simple linear regression and multiple linear regression, multiple linear regression has (>1) independent variables, whereas simple linear regression has only 1 independent variable.

Simple Linear Regression  $\rightarrow$  y = bo + b1\*x1

Multiple Linear Regression  $\rightarrow$  y = bo + b1\*x1 + b2\*x2 + .... + bn\*xn

Where, y → Dependent variable

x1, x2, ....xn → Independent variables

### **Assumptions of Linear Regression**

- 1. Linearity
- 2. Homoscedasticity
- 3. Multivariate normality
- 4. Independence of errors
- 5. Lack of multicollinearity

#### Data:

In case we have data as below:

## dataset - DataFrame

Index	R&D Spend	Administration	Marketing Spend	State	Profit
0	165349	136898	471784	New York	192262
1	162598	151378	443899	California	191792
2	153442	101146	407935	Florida	191050
3	144372	118672	383200	New York	182902
4	142107	91391.8	366168	Florida	166188
5	131877	99814.7	362861	New York	156991
6	134615	147199	127717	California	156123

**Goal**: If we have a correlation of Profit with other variables or not.

Profit → Dependent variables

Others → Independent variables



$$Y = bo + b1*x1 + b2*x2 + b3*x3 + b4*x4$$

We can ignore State, as State is a categorical variable. Hence, we need to create dummy variables i.e get\_dummies or one hot encoding.

# **Dummy Variables**

#### ## dataset - DataFrame

Index	R&D Spend	Administration	Marketing Spend	State	Profit
0	165349	136898	471784	New York	192262
1	162598	151378	443899	California	191792
2	153442	101146	407935	Florida	191050
3	144372	118672	383200	New York	182902
4	142107	91391.8	366168	Florida	166188
5	131877	99814.7	362861	New York	156991
6	134615	147199	127717	California	156123

New York	California	Florida	
1	0	0	
0	1	0	
0	0	1	
1	0	0	
0	0	1	
1	0	0	
0	1	0	

$$Y = bo + b1*x1 + b2*x2 + b3*x3 + b4*D1 + b5*D2 + b6*D3$$

Always omit one dummy variable.



## **Model Building**

In the case of Simple Linear Regression, it's quite easy to build a model, but here we need to decide which variables are important and which are not.

Not necessarily, more variables are better for a good model. We need to construct a model wisely by selecting the perfect set of variables.

#### 5 methods of building models:

- 1. All-in
- 2. Backward Elimination → Stepwise
- 3. Forward Selection → Stepwise
- 4. Bidirectional Elimination → Stepwise
- 5. Score Comparison

**All-in cases** → Domain Knowledge is quite important for these cases, where you already know which variables are going to be useful for your model.

#### Backward Elimination →

Step 1: Select a significance level to stay in the model (SL = 0.05)

Step 2: Fit the full model with all possible predictors

Step 3: Consider the predictor with the highest p-value. If P > SL go to Step 4, otherwise go to END.

Step 4: Remove the predictor

Step 5: Fit the model without this variable → Rebuilding → Coefficients, constant are going to be different

Go to Step 3, keep doing for all variables

Step 6: END → Your model is READY

#### Forward Selection →

Step 1: Select a significance level to stay in the model (SL = 0.05)

Step 2: Fit all simple regression models y~xn. Select the one with lowest p-value

Step 3: Keep this variable and fit all possible models with one extra predictor added to the one(s) you already have.

Step 4: Consider the predictor with the lowest p-value. If P < SL, go to Step 3, otherwise go to END.

Go to Step 3, keep doing till all the variables are used

Step 5: END → Keep the previous MODEL



#### Bidirectional Elimination >

Step 1: Select a significance level to enter and to stay in the model

Eg: SLENTER = 0.05, SLSTAY = 0.05

Step 2: Perform the next step of forward selection (new variable must have P <

SLENTER to enter)

Step 3: Perform all steps of backward elimination (old variables must have P < SLSTAY to stay)

Go to Step 2

Step 4: No new variables can enter no old variables can exit.

Step 5: Model is READY

#### All Possible Models

Step 1: Select a criterion of goodness of fit

Step 2: Construct all possible regression models (2^n -1) total combinations

Step 3: Select the one with the best criterion

Step 4: Model is READY

Example: 10 columns means, 2^10-1 = 1024-1 = **1023 models** 

# **Python Implementation**

The data looks like below:



In the data, we have multiple variables, to make it look easier, let's remove all the categorical variables and just deal with the numerical and proceed with the model building part.

Student's task: Build a multiple linear regression model considering the entire data.

# Multiple Linear Regression

# Importing the libraries import numpy as np



import matplotlib.pyplot as plt import pandas as pd

```
# Importing the dataset
dataset = pd.read_csv('carprice.csv')
#Removing few features to make the solution less complicated
#The below features can be converted into OneHotEncoder and the results can be
enhanced
dataset.pop('fueltype')
dataset.pop('aspiration')
dataset.pop('drivewheel')
dataset.pop('enginelocation')
dataset.pop('enginetype')
dataset.pop('fuelsystem')
dataset.pop('CarName')
#Let's simply convert these categorical to numerical's using LabelEncoder/replace
functionality
dataset['doornumber'].replace(['four','two'],[4,2],inplace=True)
dataset['carbody'].replace(['sedan','hatchback', 'wagon', 'hardtop',
'convertible'],[1,2,3,4,5],inplace=True)
dataset['cylindernumber'].replace(['four','six','five', 'eight', 'two', 'three',
'twelve'],[4,6,5,8,2,3,12],inplace=True)
X = dataset.iloc[:, :-1].values
y = dataset.iloc[:, 18].values
#No use of OneHotEncoder here, as we have removed the categorical fields.
# Encoding categorical data
from sklearn.preprocessing import LabelEncoder, OneHotEncoder
labelencoder = LabelEncoder()
X[:, 2] = labelencoder.fit_transform(X[:, 2])
onehotencoder = OneHotEncoder(categorical_features = [2])
X = onehotencoder.fit_transform(X).toarray()
# Avoiding the Dummy Variable Trap
X = X[:, 1:]
# Splitting the dataset into the Training set and Test set
```



from sklearn.cross\_validation import train\_test\_split
X\_train, X\_test, y\_train, y\_test = train\_test\_split(X, y, test\_size = 0.2, random\_state = 0)

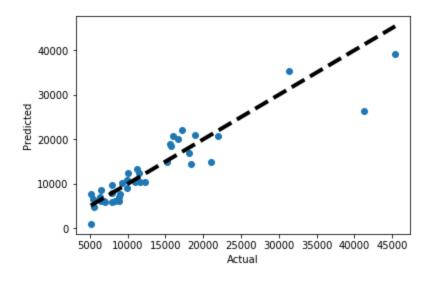
# Fitting Multiple Linear Regression to the Training set from sklearn.linear\_model import LinearRegression regressor = LinearRegression() regressor.fit(X\_train, y\_train)

# Predicting the Test set results
y\_pred = regressor.predict(X\_test)

# **Let's plot Actual vs Predicted results:**

#Let's plot Actual vs Predicted import matplotlib.pyplot as plt

fig, ax = plt.subplots()
ax.scatter(y\_test, y\_pred)
ax.plot([y\_test.min(), y\_test.max()], [y\_test.min(), y\_test.max()], 'k--', lw=4)
ax.set\_xlabel('Actual')
ax.set\_ylabel('Predicted')
plt.show()



Do you think this model is the optimum model with the dataset we have?



When we built the model, we used all the independent variables that were available, but what if among these independent variables, there are some that are highly statistically significant, which has a great impact on the dependent variable (Profit), and some that are not statistically significant variables, which means, if we remove the non statistically significant variables, there won't be much impact on the output of the model.

#### Let's try: Backward Elimination

Remember in the equation of Multiple Linear Regression i.e.

```
y = bo + b1*x1 + b2*x2 + b3*x3 + b4*x4
```

y → Dependent variable bo → Constant

x1, x2, ... xn → Independent variables

bo is basically bo\*xo, where xo = 1.

**Statsmodels** library doesn't take into account bo as constant, so we need to add it to the matrix of the dependent variable, we will add xo = 1.

#Building optimal model using Backward Elimination import statsmodels.api as sm

```
X = np.append(arr = np.ones((205, 1)).astype(int), values = X, axis = 1) #Step 2 of backward elimination
X_opt = X[:, [0,1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17, 18]]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
regressor_OLS.summary()
```

### Output:

Out[94]:

<class 'statsmodels.iolib.summary.Summary'>

OLS Regression Results

\_\_\_\_\_\_

=====

Dep. Variable: y R-squared: 0.872 Model: OLS Adj. R-squared: 0.860 Method: Least Squares F-statistic: 70.41



Mon, 06 Jan 2020 Prob (F-statistic): Date: 6.28e-73

Time: -1921.7 15:14:34 Log-Likelihood: No. Observations: 205 AIC: 3881. Df Residuals: 186 BIC: 3945.

Df Model: 18

Covariance Type: nonrobust

\_\_\_\_\_\_

[0.025 coef std err t P>|t| 0.9751

```
const -4.502e+04 1.55e+04 -2.904
                               0.004 -7.56e+04 -1.44e+04
χl
     -17.8152 4.074 -4.373 0.000 -25.852
                                         -9.778
x2
     65.2123 279.806
                     х3
      224.4821 329.699 0.681 0.497 -425.948 874.912
      321.2178 258.763 1.241 0.216 -189.269 831.705
х4
x5
     115.0941 107.157 1.074 0.284 -96.306 326.494
     -63.1336 55.123 -1.145 0.254 -171.881 45.613
х6
x7
      588.8392 241.943 2.434
                             0.016 111.535 1066.143
      248.6145 135.514 1.835
8x
                           0.068 -18.727 515.956
      0.1585 1.715 0.092 0.926
                                -3.224
                                       3.541
x9
     -1776.7108 669.892 -2.652 0.009 -3098.275 -455.147
x10
x11
     159.8603 24.628 6.491 0.000 111.274 208.446
     -3071.9252 1643.611 -1.869 0.063 -6314.441 170.591
x12
   -4725.0468 910.237 -5.191 0.000 -6520.762 -2929.331
x13
x14
     331.8705 79.060 4.198 0.000 175.900 487.841
x15
      57.1478 16.261 3.514 0.001
                                 25.068
                                        89.228
      1.9091 0.636
x16
                    3.003 0.003
                                0.655
                                        3.163
     -225.4306 171.695 -1.313 0.191 -564.151 113.289
x17
x18
     179.8634 152.851
                      1.177 0.241 -121.682
                                       481.409
______
```

=====

19.492 Durbin-Watson: 0.974 Omnibus: Prob(Omnibus): 0.000 Jarque-Bera (JB): 52.169

Skew: 0.336 Prob(JB): 4.69e-12 Kurtosis: 5.378 Cond. No. 4.27e+05

\_\_\_\_\_\_

=====

#### Warnings:

- [1] Standard Errors assume that the covariance matrix of the errors is correctly specified.
- [2] The condition number is large, 4.27e+05. This might indicate that there are strong multicollinearity or other numerical problems.



111111

x9 has the highest p value, hence we remove it in the next run  $\rightarrow$  And the process continues till we have the variables with a p value less than 0.05

```
X_{opt} = X[:, [0,1,2,3,4,5,6,7,8,10,11,12,13,14,15,16,17,18]]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
regressor_OLS.summary()
X_{opt} = X[:, [0,1,3,4,5,6,7,8,10,11,12,13,14,15,16,17,18]]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
regressor_OLS.summary()
X_{opt} = X[:, [0,1,4,5,6,7,8,10,11,12,13,14,15,16,17,18]]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
regressor_OLS.summary()
X_{opt} = X[:, [0,1,4,5,7,8,10,11,12,13,14,15,16,17,18]]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
regressor_OLS.summary()
X_{opt} = X[:, [0,1,4,7,8,10,11,12,13,14,15,16,17,18]]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
regressor_OLS.summary()
X_{opt} = X[:, [0,1,4,7,8,10,11,12,13,14,15,16,17]]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
regressor_OLS.summary()
X_{opt} = X[:, [0,1,4,7,8,10,11,12,13,14,15,16]]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
regressor_OLS.summary()
X_{opt} = X[:, [0,1,7,8,10,11,12,13,14,15,16]]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
regressor_OLS.summary()
X_{opt} = X[:, [0,1,7,8,10,11,13,14,15,16]]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
regressor_OLS.summary()
X_{opt} = X[:, [0,1,7,10,11,13,14,15,16]]
regressor_OLS = sm.OLS(endog = y, exog = X_opt).fit()
```



#### regressor\_OLS.summary()

Warnings:

#### Output Out[104]: <class 'statsmodels.iolib.summary.Summary'> **OLS Regression Results** \_\_\_\_\_ ===== Dep. Variable: y R-squared: 0.861 OLS Adj. R-squared: Least Squares F-statistic: Model: 0.856 Method: 152.3 Date: Mon, 06 Jan 2020 Prob (F-statistic): 8.15e-80 Time: 15:18:43 Log-Likelihood: -1929.9 No. Observations: 205 AIC: 3878. Df Residuals: 196 BIC: 3908. Df Model: 8 Covariance Type: nonrobust \_\_\_\_\_\_ ===== coef std err t P>|t| [0.025 0.975] \_\_\_\_\_\_ const -4.985e+04 1.06e+04 -4.697 0.000 -7.08e+04 -2.89e+04 x1 -17.7694 3.842 -4.625 0.000 -25.346 -10.193 724.7305 158.059 4.585 0.000 413.016 1036.445 x2 x3 -1211.8347 409.088 -2.962 0.003 -2018.614 -405.055 142.7100 16.585 8.605 0.000 110.001 175.419 x4 x5 -4230.4288 770.725 -5.489 0.000 -5750.407 -2710.451 x6 314.4289 64.536 4.872 0.000 187.154 441.704 44.8901 12.055 3.724 0.000 21.116 68.664 x7 2.0100 0.620 3.241 0.001 0.787 3.233 8x \_\_\_\_\_\_ ===== Omnibus: 15.874 Durbin-Watson: 0.943 Prob(Omnibus): 0.000 Jarque-Bera (JB): 33.383 0.335 Prob(JB): 5.64e-08 Skew: Kurtosis: 4.860 Cond. No. 2.58e+05 \_\_\_\_\_\_



- [1] Standard Errors assume that the covariance matrix of the errors is correctly specified.
- [2] The condition number is large, 2.58e+05. This might indicate that there are strong multicollinearity or other numerical problems.



# **Ensemble Techniques**

# **Ensemble Learning**

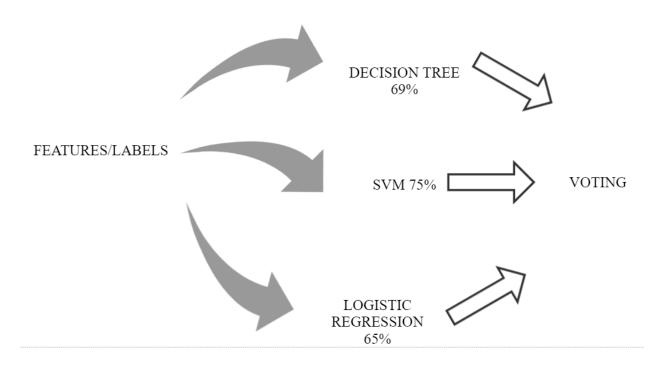
Ensemble Learning is the mechanism to use multiple algorithms together to have a better prediction than the individual models.

Let say, we extract the features from a use case, and try to check the model's accuracy or behaviour individually.

Features → Decision Trees → 69%

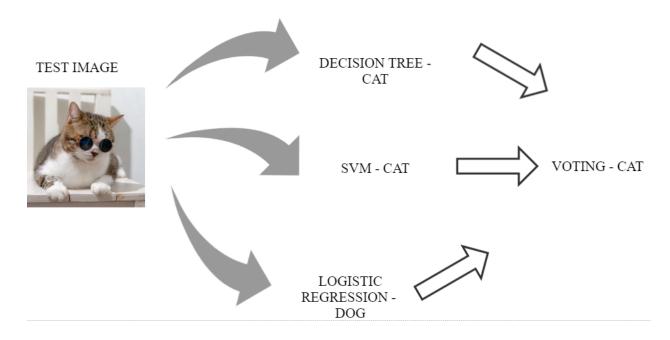
Features → SVM → 75%

Features → Logistic Regression → 65%



In the above picture, we tried different algorithms and got a maximum accuracy of 75%, so what if we combine 3 models, and take a vote on each output, will it increase the accuracy of the entire model?





Let's say, we have a test image of a cat, and we train different classifiers, where Decision Tree & SVN identified it as CAT, but Logistic Regression identified it as a DOG, and considering it as a whole model, we take the voting and predict the image as a CAT. Obviously, the efficiency of ensemble models will be better than individual models, but the computational power will increase, and training a model will take more time as compared to the individual models.

# So why use Ensemble Learning?

- 1. Better Accuracy (Low Error)
- 2. Higher Consistency (Avoids Overfitting)
- 3. Reduces Bias & Variance Frrors

# When and Where do we use Ensemble Learning?

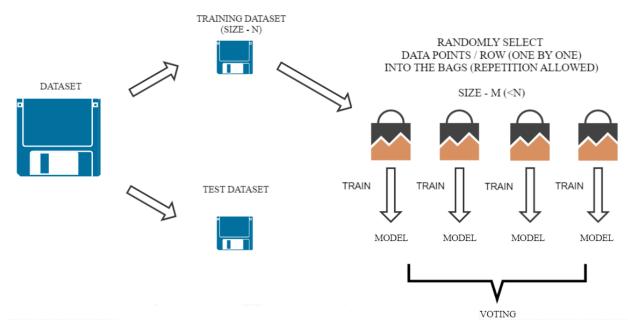
- 1. Single model overfits
- 2. Results worth the extra training
- 3. Can be used for classification & regression both.



# **Popular Ensemble Methods**

# 1. Bootstrap Aggregation (Bagging)

Multiple models of the same learning algorithm trained with subsets of dataset randomly picked from the training dataset.



# Bagging Algorithms:

- a. Bagged Decision Trees
- b. Random Forest
- c. Extra Trees

Code Snippet to be used for each of the below examples like Bagged Decision Trees, Random Forest, Extra Trees, and other Boost examples:

#Importing Libraries import pandas as pd from sklearn.tree import DecisionTreeClassifier from sklearn import model\_selection from sklearn.ensemble import BaggingClassifier

#Reading data
data = pd.read\_csv(r"Social\_Network\_Ads.csv")
data.head()



#Removing User ID, as it's an incremental value that doesn't add to our classification prediction data.pop('User ID')

#Replacing categorical values to numericals data['Gender'].replace(['Male', Female'], [1,0], inplace=True)

#Using features: Gender, Age for prediction of Purchased label feature\_cols = ['Gender', 'Age'] X = data[feature\_cols] # Features Y = data.Purchased # Target variable

## **Bagged Decision Trees**

Bagging performs best with algorithms with high variance. Decision tree is one of the most popular examples.

In this example, we will see the usage of BaggingClassifier along with DecisionTreeClassifier. A total of 100 trees are used.

seed = 7
kfold = model\_selection.KFold(n\_splits=10, random\_state=seed)
cart = DecisionTreeClassifier()
num\_trees = 100
model = BaggingClassifier(base\_estimator=cart, n\_estimators=num\_trees, random\_state=seed)
results = model\_selection.cross\_val\_score(model, X, Y, cv=kfold)
print("Accuracy for BaggingClassifier:", results.mean())

Output: Accuracy for BaggingClassifier: 0.8

#### **Random Forest**

RF is an extension of Bagged decision trees.

Samples of the training dataset are taken with replacement, but the trees are constructed in a way that reduces the correlation between individual classifiers. Specifically, rather than greedily choosing the best split point in the construction of the tree, only a random subset of features are considered for each split.

You can construct a Random Forest model for classification using the RandomForestClassifier class.



The example below provides an example of Random Forest for classification with 100 trees.

from sklearn.ensemble import RandomForestClassifier

```
seed = 7
num_trees = 100
max_features = 3
kfold = model_selection.KFold(n_splits=10, random_state=seed)
model = RandomForestClassifier(n_estimators=num_trees)
results = model_selection.cross_val_score(model, X, Y, cv=kfold)
print("Accuracy for RandomForestClassifier:", results.mean())
```

Output: Accuracy for RandomForestClassifier: 0.7975

#### **Extra Trees**

Extra Trees are another modification of bagging where random trees are constructed from samples of the training dataset.

You can construct an Extra Trees model for classification using the ExtraTreesClassifier class.

The example below provides a demonstration of extra trees with the number of trees set to 100 and splits chosen from 7 random features.

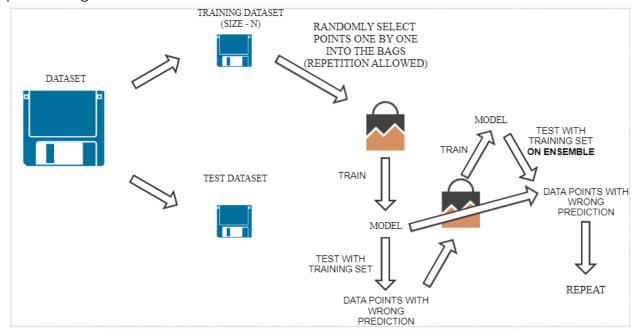
from sklearn.ensemble import ExtraTreesClassifier

```
seed = 7
num_trees = 100
max_features = 7
kfold = model_selection.KFold(n_splits=10, random_state=seed)
model = ExtraTreesClassifier(n_estimators=num_trees)
results = model_selection.cross_val_score(model, X, Y, cv=kfold)
print("Accuracy for ExtraTreesClassifier:", results.mean())
```

## 2. Boosting



Boosting is used to create a collection of predictors. In this technique, learners are learned sequentially with early learners fitting simple models to the data and then analysing data for errors. Consecutive trees (random sample) are fit and at every step, the goal is to improve the accuracy from the prior tree. When an input is misclassified by a hypothesis, its weight is increased so that the next hypothesis is more likely to classify it correctly. This process converts weak learners into better performing model.



Most common Boosting techniques are:

- a. AdaBoost
- b. Stochastic Gradient Boosting

#### AdaBoost

AdaBoost was perhaps the first successful boosting ensemble algorithm. It generally works by weighting instances in the dataset by how easy or difficult they are to classify, allowing the algorithm to pay or or less attention to them in the construction of subsequent models.

You can construct an AdaBoost model for classification using the AdaBoostClassifier class.

The example below demonstrates the construction of 30 decision trees in sequence using the AdaBoost algorithm.



from sklearn.ensemble import AdaBoostClassifier

seed = 7
num\_trees = 30
kfold = model\_selection.KFold(n\_splits=10, random\_state=seed)
model = AdaBoostClassifier(n\_estimators=num\_trees, random\_state=seed)
results = model\_selection.cross\_val\_score(model, X, Y, cv=kfold)
print("Accuracy for AdaBoostClassifier:", results.mean())

Output: Accuracy for AdaBoostClassifier: 0.835

### **Stochastic Gradient Boosting**

Stochastic Gradient Boosting (also called Gradient Boosting Machines) are one of the most sophisticated ensemble techniques. It is also a technique that is proving to be perhaps of the best techniques available for improving performance via ensembles.

You can construct a Gradient Boosting model for classification using the GradientBoostingClassifier class.

The example below demonstrates Stochastic Gradient Boosting for classification with 100 trees.

from sklearn.ensemble import GradientBoostingClassifier

```
seed = 7
num_trees = 100
kfold = model_selection.KFold(n_splits=10, random_state=seed)
model = GradientBoostingClassifier(n_estimators=num_trees, random_state=seed)
results = model_selection.cross_val_score(model, X, Y, cv=kfold)
print("Accuracy for GradientBoostingClassifier:", results.mean())
```

**Output**: Accuracy for GradientBoostingClassifier: 0.814999999999999

#### **Voting Ensemble**

To create a **VotingClassifier**, simply aggregate the predictions of each classifier and predict the class that gets the most votes. This majority-vote classifier is called a *hard voting* classifier.



Here's how it's done in Scikit-Learn:

```
# Voting Ensemble for Classification
from sklearn import model_selection
from sklearn.linear_model import LogisticRegression
from sklearn.tree import DecisionTreeClassifier
from sklearn.svm import SVC
from sklearn.ensemble import VotingClassifier
seed = 7
kfold = model_selection.KFold(n_splits=10, random_state=seed)
# create the sub models
estimators = []
model1 = LogisticRegression()
estimators.append(('logistic', model1))
model2 = DecisionTreeClassifier()
estimators.append(('cart', model2))
model3 = SVC()
estimators.append(('svm', model3))
# create the ensemble model
ensemble = VotingClassifier(estimators)
results = model_selection.cross_val_score(ensemble, X, Y, cv=kfold)
print("Accuracy for VotingClassifier:", results.mean())
```

Output: Accuracy for VotingClassifier: 0.8324999999999999

### Conclusion:

In this module, you learnt about various Ensemble algorithms, and using an example, we found that **AdaBoost** gave us the best results, but it doesn't mean that boosting provides us with the best results every time. It's all about hit and trial, these algorithms have to be tried separately for each use case, and then we can come to a conclusion. In some cases, standalone classifiers also outperforms as compared to the ensemble techniques.