

# Machine Learning Lab

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## Candidate Elimination Algorithm

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### Terms Used:

1. **Concept learning**: Concept learning is basically learning task of the machine (Learn by Train data)
  2. **General Hypothesis**: Not Specifying features to learn the machine.
  3. **G** = { '?', '?', '?', '?', ... } : Number of attributes
  4. **Specific Hypothesis**: Specifying features to learn machine (Specific feature)
  5. **S** = { 'pi', 'pi', 'pi', ... } : Number of pi depends on number of attributes.
  6. **Version Space**: It is intermediate of general hypothesis and Specific hypothesis. It not only just written one hypothesis but a set of all possible hypothesis based on training data-set.
- 

### Algorithm:

```
Step1: Load Data set
Step2: Initialize General Hypothesis and Specific Hypothesis.
Step3: For each training example
Step4: If example is positive example
    if attribute_value == hypothesis_value:
        Do nothing
    else:
        replace attribute value with '?' (Basically generalizing it)
Step5: If example is Negative example
    Make generalize hypothesis more specific.
```

---

In [1]: *#import useful packages*

```
import numpy as np
import pandas as pd
```

In [2]: *#read data set*

```
df = pd.read_csv('data/ce.csv')
```

In [3]: *#preview dataset*

```
df
```

Out[3]:

	sky	airtemp	humidity	wind	water	forecast	enjoysport
0	sunny	warm	normal	strong	warm	same	Yes
1	sunny	warm	high	strong	warm	same	Yes
2	rainy	cold	high	strong	warm	change	No
3	sunny	warm	high	strong	cool	change	Yes

In [4]: *#defining algorithm*

```
def candidate_elimination(con,tar):
    for i,val in enumerate(tar):
        if val == 'Yes':
            specific_h = con[i].copy()
            break
    general_h = [['?' for i in range(len(specific_h))] for i in range(len(specific_h))]

    for i, val in enumerate(con):
        if tar[i] == "Yes":
            for j in range(len(specific_h)):
                if val[j] != specific_h[j]:
                    specific_h[j] = '?'
                    general_h[j][j] = '?'

        if tar[i] == "No":
            for j in range(len(specific_h)):
                if val[j] != specific_h[j]:
                    general_h[j][j] = specific_h[j]
            else:
                general_h[j][j] = '?'
    return specific_h,general_h
```

In [5]: *#run algorithm*

```
con = np.array(df)[:,-1]
tar = np.array(df)[:,-1]

spe_h,gen_h = candidate_elimination(con,tar)
```

```
In [6]: #print specific_h
spe_h
```

```
Out[6]: array(['sunny', 'warm', '?', 'strong', '?', '?'], dtype=object)
```

```
In [7]: #print gen_h
gen_h
```

```
Out[7]: [['sunny', '?', '?', '?', '?', '?'],
         ['?', 'warm', '?', '?', '?', '?'],
         ['?', '?', '?', '?', '?', '?'],
         ['?', '?', '?', '?', '?', '?'],
         ['?', '?', '?', '?', '?', '?'],
         ['?', '?', '?', '?', '?', '?']]
```

---

## Find-S algorithm:

### Important Representation:

1. ? indicates that any value is acceptable for the attribute.
2. specify a single required value ( e.g., Cold ) for the attribute.
3.  $\varnothing$  indicates that no value is acceptable.
4. The most general hypothesis is represented by: {?, ?, ?, ?, ?, ?}
5. The most specific hypothesis is represented by : { $\varnothing$ ,  $\varnothing$ ,  $\varnothing$ ,  $\varnothing$ ,  $\varnothing$ ,  $\varnothing$ }

---

### Algorithm:

1. Initialize h to the most specific hypothesis in H
2. For each positive training instance x For each attribute constraint a, in h
  - If the constraint a, is satisfied by x
  - Then do nothing
  - Else replace a, in h by the next more general constraint that is satisfied by x
3. Output hypothesis h

---

```
In [7]: #import useful packages

import numpy as np
import pandas as pd
```

```
In [8]: #read data set
df = pd.read_csv('data/fs.csv')
```

```
In [9]: #preview dataset
df
```

```
Out[9]:
```

	Color	Toughness	Fungus	Appearance	Poisonous
0	Green	Hard	No	Wrinkled	Yes
1	Green	Hard	Yes	Smooth	No
2	Brown	Soft	No	Wrinkled	No
3	Orange	Hard	No	Wrinkled	Yes
4	Green	Hard	Yes	Wrinkled	Yes
5	Orange	Hard	No	Wrinkled	Yes

```
In [10]: # find s algo
def find_s(con,tar):
    for i,val in enumerate(tar):
        if val == 'Yes':
            specific_h = con[i].copy()
            break

    for i,val in enumerate(con):
        if tar[i] == 'Yes':
            for j,feat in enumerate(val):
                if specific_h[j] != feat:
                    specific_h[j] = '?'
            print(specific_h)
    return specific_h
```

```
In [11]: #run algorithm
con = np.array(df)[:,:-1]
tar = np.array(df)[:,-1]

gen_h = find_s(con,tar)
```

```
['Green' 'Hard' 'No' 'Wrinkled']
['?' 'Hard' 'No' 'Wrinkled']
['?' 'Hard' '?' 'Wrinkled']
['?' 'Hard' '?' 'Wrinkled']
```

```
In [12]: #see result

gen_h
```

```
Out[12]: array(['?', 'Hard', '?', 'Wrinkled'], dtype=object)
```

## A. Simple Linear Analysis (From Scratch)

```
In [1]: #import package
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
```

```
In [2]: #read data
df = pd.read_csv('data/a.csv')
df.head()
```

```
Out[2]:
```

	X	Y
0	3.5	5.1
1	3.0	4.9
2	3.2	4.7
3	3.1	4.6
4	3.6	5.0

Suppose the required regression line equation is :

$$y = m \cdot x + b$$

There are two equations called normal equation from which we can find  $m$  and  $b$  :

$$\Sigma XY = b \cdot \Sigma X + m \cdot \Sigma X^2 \quad \dots\dots\dots(1)$$

$$\Sigma Y = n \cdot b + m \cdot \Sigma X \quad \dots\dots\dots(2)$$

Or we can solve the two equation for  $m$  and  $b$  as:

$$m = \frac{n \cdot \Sigma xy - \Sigma x \cdot \Sigma y}{n \cdot \Sigma x^2 - (\Sigma x)^2}$$

$$b = \bar{y} - m \cdot \bar{x}$$

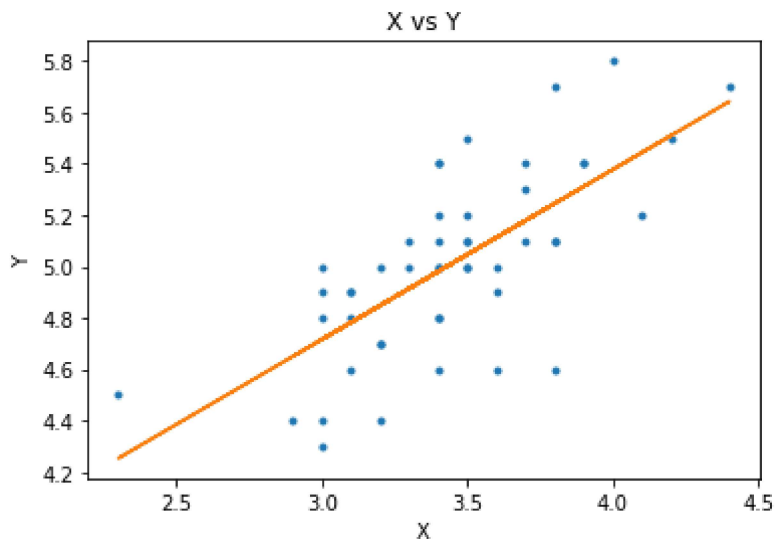
```
In [3]: #Calculate m and b for y = mx+b
X = np.array(df['X'])
Y = np.array(df['Y'])

m = ((len(X)*sum(X*Y)) - sum(X)*sum(Y))/((len(X)*sum(X**2)) - (sum(X)**2))

b = np.mean(Y) - m*np.mean(X)
```

```
In [4]: # append calculated y column in df
df['Y_h'] = m*X + b
```

```
In [5]: #plot data points and y=mx+b line
plt.plot(df['X'],df['Y'],'.')
plt.plot(df['X'],df['Y_h'],'-')
plt.xlabel("X");
plt.ylabel("Y");
plt.title("X vs Y");
```



## B. Multiple Linear Regression (From Scratch)

```
In [6]: df1 = pd.read_csv("data/mlr.csv")
df1.head()
```

```
Out[6]:
```

	X1	X2	X3	X4	X5
0	6.8	225	0.442	0.672	9.2
1	6.3	180	0.435	0.797	11.7
2	6.4	190	0.456	0.761	15.8
3	6.2	180	0.416	0.651	8.6
4	6.9	205	0.449	0.900	23.2

### Data Description

The following data (X1, X2, X3, X4, X5) are for each player.

X1 = height in feet

X2 = weight in pounds

X3 = percent of successful field goals (out of 100 attempted)

X4 = percent of successful free throws (out of 100 attempted)

X5 = average points scored per game

```
In [7]: X = np.array(df1.drop('X5',axis = 1))
ones = np.ones((X.shape[0],1))
X = np.append(ones,X,axis = 1)
Y = np.array(df1['X5'])
```

$$y = b_0 + x_1 b_1 + x_2 b_2 + \dots$$

$$Y = Xb$$

$$\mathbf{X} = \begin{bmatrix} 1 & X_1 & X_2 & \dots \\ 1 & \dots & \dots & \cdot \\ \vdots & \vdots & \ddots & \vdots \\ 1 & \dots & \dots & \dots \end{bmatrix}$$

$$\mathbf{Y} = \begin{bmatrix} Y_1 \\ Y_2 \\ \vdots \\ Y_n \end{bmatrix}$$

$$\mathbf{B} = \begin{bmatrix} b_0 & b_1 & b_2 & \dots \end{bmatrix}$$

```
In [8]: b = np.linalg.inv(X.T@X)@X.T@Y
b
```

```
Out[8]: array([ 4.14870671e+00, -3.69049908e+00,  9.45845788e-03,  4.79401992e+01,
 1.13710193e+01])
```

```
In [9]: Y_pred = X@b
Y_pred
```

```
Out[9]: array([10.01235894, 12.51777389, 12.84069605, 10.3157912 , 12.38231366,
 12.18928646, 14.76231003, 12.2882684 , 11.06248103, 13.80102549,
 12.92521215, 13.24396498,  8.64006433, 12.60525851, 12.92430257,
 15.2545908 , 15.58138272,  5.09650334,  9.94940424, 12.49696124,
  9.05637536, 10.25083143,  8.06191203, 10.07255781, 14.74963484,
 12.07610898, 13.95300287,  8.71473239, 11.3419041 , 14.40707455,
 15.27283126, 10.4123043 , 13.64991307, 10.25083143,  8.97445925,
 14.70763753, 14.90726563, 13.253166 ,  7.32888666,  6.6597904 ,
  8.95334041,  5.83073166, 13.82675288, 16.36627839, 12.00928465,
 10.11871685, 17.63471509,  9.70871125,  8.95738083, 14.73644788,
 15.93275941, 12.25972646, 11.34188054, 10.03210379])
```

$$R^2 = \frac{\Sigma(\hat{Y} - \bar{Y})^2}{\Sigma(Y - \bar{Y})^2}$$

```
In [10]: Y_mean = np.mean(Y)

R2 = (np.sum((Y_pred-Y_mean)**2))/(np.sum((Y-Y_mean)**2))
print(f"R^2 value is {R2}")

R^2 value is 0.22225063130014133
```

## Linear Regression Using SKLearn

### 1. Read Data

```
In [11]: df_3 = pd.read_csv("data/insurance.csv")
df_3.head()
```

```
Out[11]:
```

	age	sex	bmi	children	smoker	region	charges
0	19	0	27.900	0	1	3	16884.92400
1	18	1	33.770	1	0	2	1725.55230
2	28	1	33.000	3	0	2	4449.46200
3	33	1	22.705	0	0	1	21984.47061
4	32	1	28.880	0	0	1	3866.85520

```
In [12]: df_3.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 348 entries, 0 to 347
Data columns (total 7 columns):
#   Column      Non-Null Count  Dtype  
---  -
0   age         348 non-null   int64  
1   sex         348 non-null   int64  
2   bmi         348 non-null   float64 
3   children    348 non-null   int64  
4   smoker      348 non-null   int64  
5   region      348 non-null   int64  
6   charges     348 non-null   float64 
dtypes: float64(2), int64(5)
memory usage: 19.2 KB
```

### 2. Import Linear Regressor and Split Data Set



```
In [13]: from sklearn.linear_model import LinearRegression
from sklearn.model_selection import train_test_split
```

```
In [14]: X = df_3.drop('charges', axis = 1)
Y = df_3['charges']
```

```
In [15]: X_train,X_test,Y_train,Y_test = train_test_split(X,Y, test_size = 0.3)
```

### 3. Train Linear Regression Model

```
In [16]: lr_model = LinearRegression()
```

```
In [17]: lr_model.fit(X_train,Y_train)
```

```
Out[17]: LinearRegression()
```

### 4. Predict and check Model

```
In [18]: loc = 10
x = X_test.iloc[loc]
y = Y_test.iloc[loc]
y_pred = lr_model.predict([x])
```

```
In [19]: print(f'x = {x.values}\ny = {y}\nPredicted Value = {y_pred[0]}')
```

```
x = [19.    0.  28.9  0.    0.    3. ]
y = 1743.214
Predicted Value = 1913.0976139823597
```

```
In [20]: print(f'Score of Model = {lr_model.score(X_test,Y_test)}')
```

```
Score of Model = 0.7750219073416568
```

---

## Locally Weight Regression (From Scretch)

```
In [1]: # import useful module
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
```

### Algorithm

Locally weighted Regression is a non parametric Regression Algorithms in which we need data every time we calculate matrix.

### Cost Function in Linear Regression :

$$\text{costfn} = \sum (y_i - \theta x_i)^2$$

### Cost Function for Locally Weighted Regression:

$$\text{costfn} = \sum w_i (y_i - \theta x_i)^2$$

where  $x_i, y_i$  is the  $i^{\text{th}}$  example and  $w_i$  is weight.

Generally

$$w_i = e^{-\frac{(x_i - x)^2}{2\tau^2}}$$

by minimize costfn for  $\theta$  and solving expression we get

$$\theta = (X^T W X)^{-1} (X^T W Y)$$

And prediction is  $y = X\theta$ .

```
In [2]: def w_i(point,X,tau):  
    m,n = X.shape  
    w = np.mat(np.eye(m))  
  
    for j in range(m):  
        diff = X[j]-point  
        w[j, j] = np.exp(diff * diff.T / (-2.0 * tau**2))  
    return w
```

```
In [3]: def theta(X,Y,point,tau):  
    W = w_i(point,X,tau)  
    th_x = X.T@W@X  
    th_y = X.T@W@Y.T  
    th = th_x.I@th_y  
    return th
```

```
In [4]: def predict(X,Y,tau):  
    m,n = np.shape(X)  
    ypred = np.zeros(m)  
  
    for i in range(m):  
        ypred[i] = X[i] * theta( X, Y, X[i],tau)  
    return ypred
```

---

## Part 1

Generate X and Y and apply

```
In [5]: n = 100
xs = np.linspace(0, np.pi, n)
ys = 1 + np.sin(xs) + np.cos(xs**2) + np.random.normal(0, 0.1, n)
```

```
In [6]: X = np.mat(xs)
Y = np.mat(ys)

#add 1 with each entry in X
m = X.shape[1]
one = np.ones((1,m))
X = np.hstack((one.T,X.T))

print(X[:5])
```

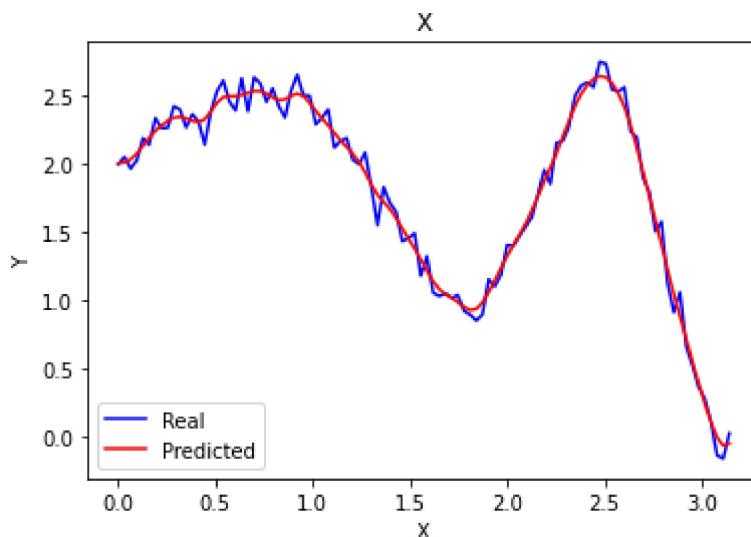
```
[[1.      0.      ]
 [1.      0.03173326]
 [1.      0.06346652]
 [1.      0.09519978]
 [1.      0.12693304]]
```

```
In [7]: Y_prediction_05 = predict(X,Y,0.05)
```

```
In [8]: X_l = xs
Y_l = ys
Y_p = Y_prediction_05

plt.plot(X_l,Y_l,'b-')
plt.plot(X_l,Y_p,'r-')
plt.xlabel('X')
plt.ylabel('Y')
plt.title('X')
plt.legend(['Real', 'Predicted'])
```

Out[8]: <matplotlib.legend.Legend at 0x2202eed070>



## Part 2

Use on DataSet

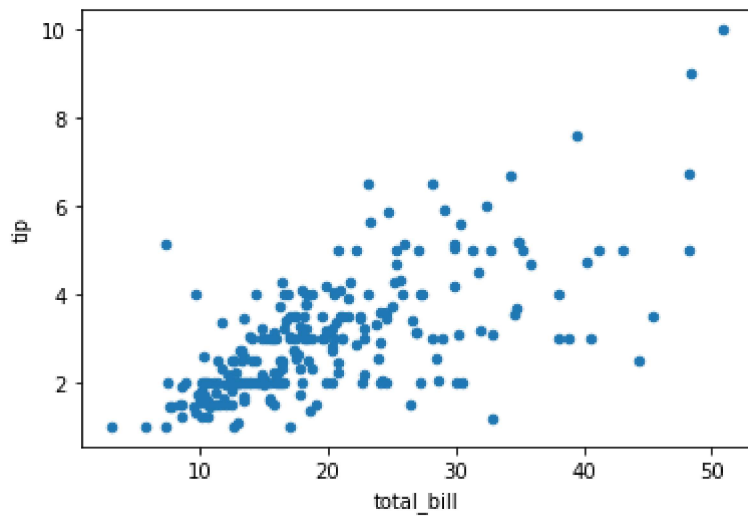
```
In [9]: # Read Data and make DataFrame
df = pd.read_csv('data/tips.csv')
df.head()
```

```
Out[9]:
```

	total_bill	tip	sex	smoker	day	time	size
0	16.99	1.01	Female	No	Sun	Dinner	2
1	10.34	1.66	Male	No	Sun	Dinner	3
2	21.01	3.50	Male	No	Sun	Dinner	3
3	23.68	3.31	Male	No	Sun	Dinner	2
4	24.59	3.61	Female	No	Sun	Dinner	4

```
In [10]: #plot total_bill vs tip
df.plot(x = 'total_bill',y = 'tip',kind = 'scatter')
```

```
Out[10]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



```
In [11]: # select columns regression
X = np.mat(df['total_bill'])
Y = np.mat(df['tip'])

#add 1 with each entry in X
m = X.shape[1]
one = np.ones((1,m))
X = np.hstack((one.T,X.T))

print(X[:5])
```

```
[[ 1.  16.99]
 [ 1.  10.34]
 [ 1.  21.01]
 [ 1.  23.68]
 [ 1.  24.59]]
```

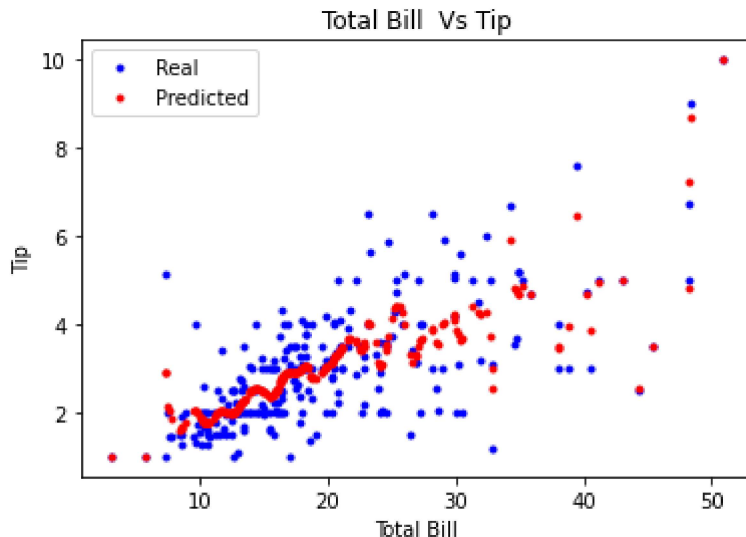
```
In [12]: Y_prediction = predict(X,Y,0.4)
```

## Plot the predicted result

```
In [13]: X_l = df['total_bill']
Y_l = df['tip']
Y_p = Y_prediction

plt.plot(X_l,Y_l,'b.')
plt.plot(X_l,Y_p,'r.')
plt.xlabel('Total Bill')
plt.ylabel('Tip')
plt.title('Total Bill Vs Tip')
plt.legend(['Real', 'Predicted'])
```

```
Out[13]: <matplotlib.legend.Legend at 0x2202f37ef70>
```



# Decision Tree

```
In [1]: #import useful package
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier, plot_tree
import matplotlib.pyplot as plt
from sklearn.metrics import confusion_matrix, accuracy_score
```

```
In [2]: #read data
df = pd.read_csv('data/Iris.csv')
df.head()
```

Out[2]:

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	1	5.1	3.5	1.4	0.2	Iris-setosa
1	2	4.9	3.0	1.4	0.2	Iris-setosa
2	3	4.7	3.2	1.3	0.2	Iris-setosa
3	4	4.6	3.1	1.5	0.2	Iris-setosa
4	5	5.0	3.6	1.4	0.2	Iris-setosa

```
In [3]: df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 6 columns):
 #   Column          Non-Null Count  Dtype
---  -
 0   Id              150 non-null   int64
 1   SepalLengthCm   150 non-null   float64
 2   SepalWidthCm    150 non-null   float64
 3   PetalLengthCm   150 non-null   float64
 4   PetalWidthCm    150 non-null   float64
 5   Species         150 non-null   object
dtypes: float64(4), int64(1), object(1)
memory usage: 7.2+ KB
```

```
In [4]: #Splitting data in to train and test part

X = df.drop(['Species', 'Id'], axis=1)
class_name = list(df.Species.unique())
feat_name = list(X.columns)
Y = df.Species.apply(lambda x: class_name.index(x))
#Y = df.Species
```

```
In [5]: x_train, x_test, y_train, y_test = train_test_split(X, Y, test_size = 0.3)
```

# Train Model :- ID3

## Algorithm

```
ID3 (Examples, Target_Attribute, Attributes)
    Create a root node for the tree
    If all examples are positive, Return the single-node tree Root, with label = +.
    If all examples are negative, Return the single-node tree Root, with label = -.
    If number of predicting attributes is empty, then Return the single node tree Root,
    with label = most common value of the target attribute in the examples.
    Otherwise Begin
        A ← The Attribute that best classifies examples.
        Decision Tree attribute for Root = A.
        For each possible value, vi, of A,
            Add a new tree branch below Root, corresponding to the test A = vi.
            Let Examples(vi) be the subset of examples that have the value vi for A
            If Examples(vi) is empty
                Then below this new branch add a leaf node with label = most common target value in the examples
            Else below this new branch add the subtree ID3 (Examples(vi), Target_Attribute, Attributes - {A})
        End
    Return Root
```

ID3 algorithm is based on entropy and information gain calculation.

Entropy is calculated as

$$\text{Entropy} = -\sum p(X) \log p(X)$$

Where  $p(X)$  is a Fraction of example in given class

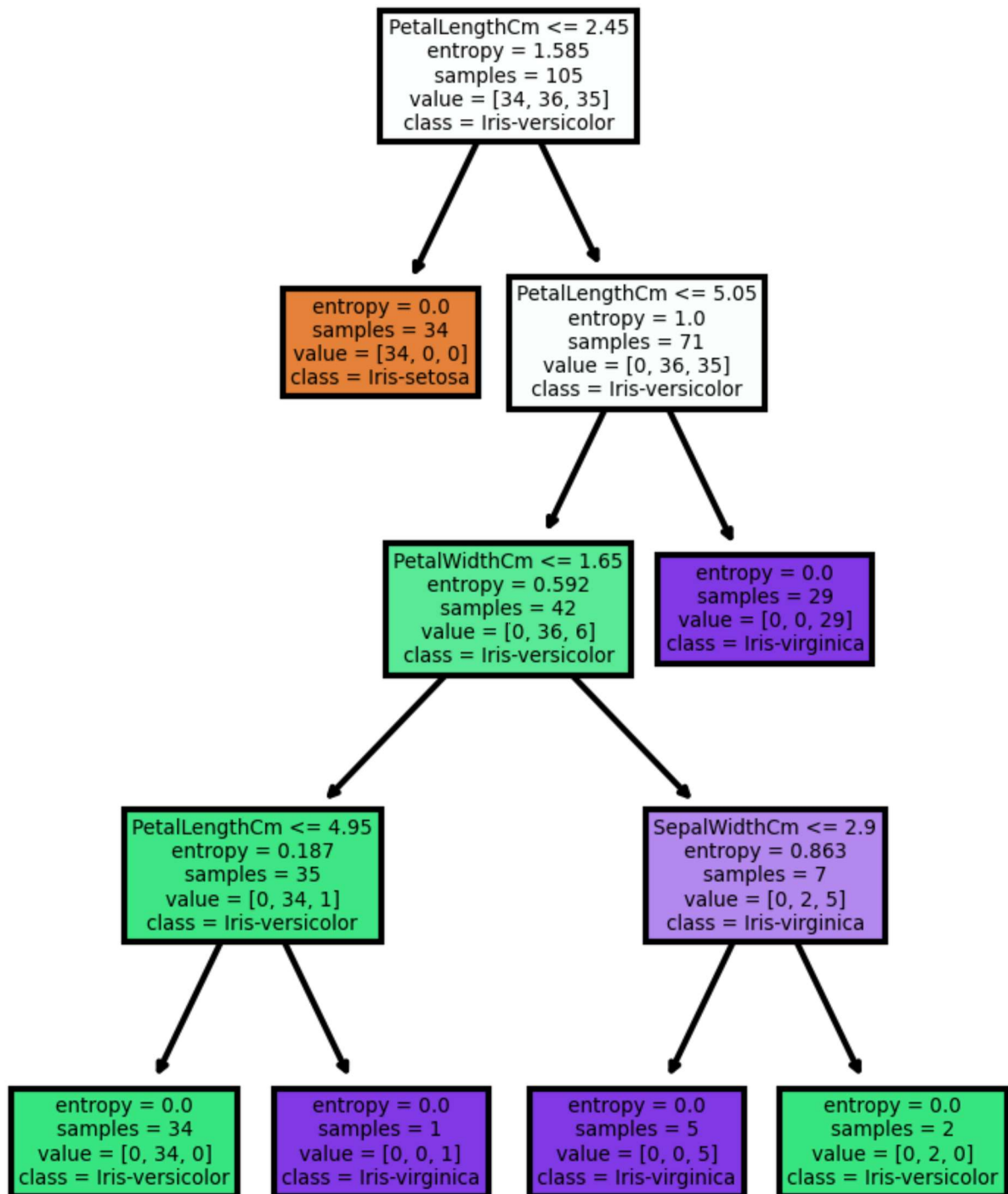
and information gain is calculated as-

$$\text{GAIN} = \text{Entropy}(p) - \sum \frac{n_i}{n} \text{Entropy}(i)$$

```
In [6]: cd3 = DecisionTreeClassifier(criterion='entropy')
cd3.fit(x_train,y_train)
```

```
Out[6]: DecisionTreeClassifier(criterion='entropy')
```

```
In [15]: #plot tree
fig, axes = plt.subplots(nrows = 1,ncols = 1,figsize = (3,4), dpi=300)
plot_tree(cd3,
          feature_names = feat_name,
          class_names=class_name,
          filled = True);
```





```
In [8]: #check accurecy
y_pred = cd3.predict(x_test)
print(f'Accuracy Score of model is : {accuracy_score(y_test,y_pred)}')
```

Accuracy Score of model is : 0.9333333333333333

```
In [9]: print('Confusion Metrics :')
print(confusion_matrix(y_test,y_pred))
```

Confusion Metrics :

```
[[16  0  0]
 [ 0 13  1]
 [ 0  2 13]]
```

---

## Train Model :- CART

### Algorithm

- Step 1: Start at the root node with all training instances
- Step 2: Select an attribute on the basis of splitting criteria
- Step 3: Partition instances according to selected attribute recursively

ID3 algorithm is based on GINI Impurity.

GINI is calculated as

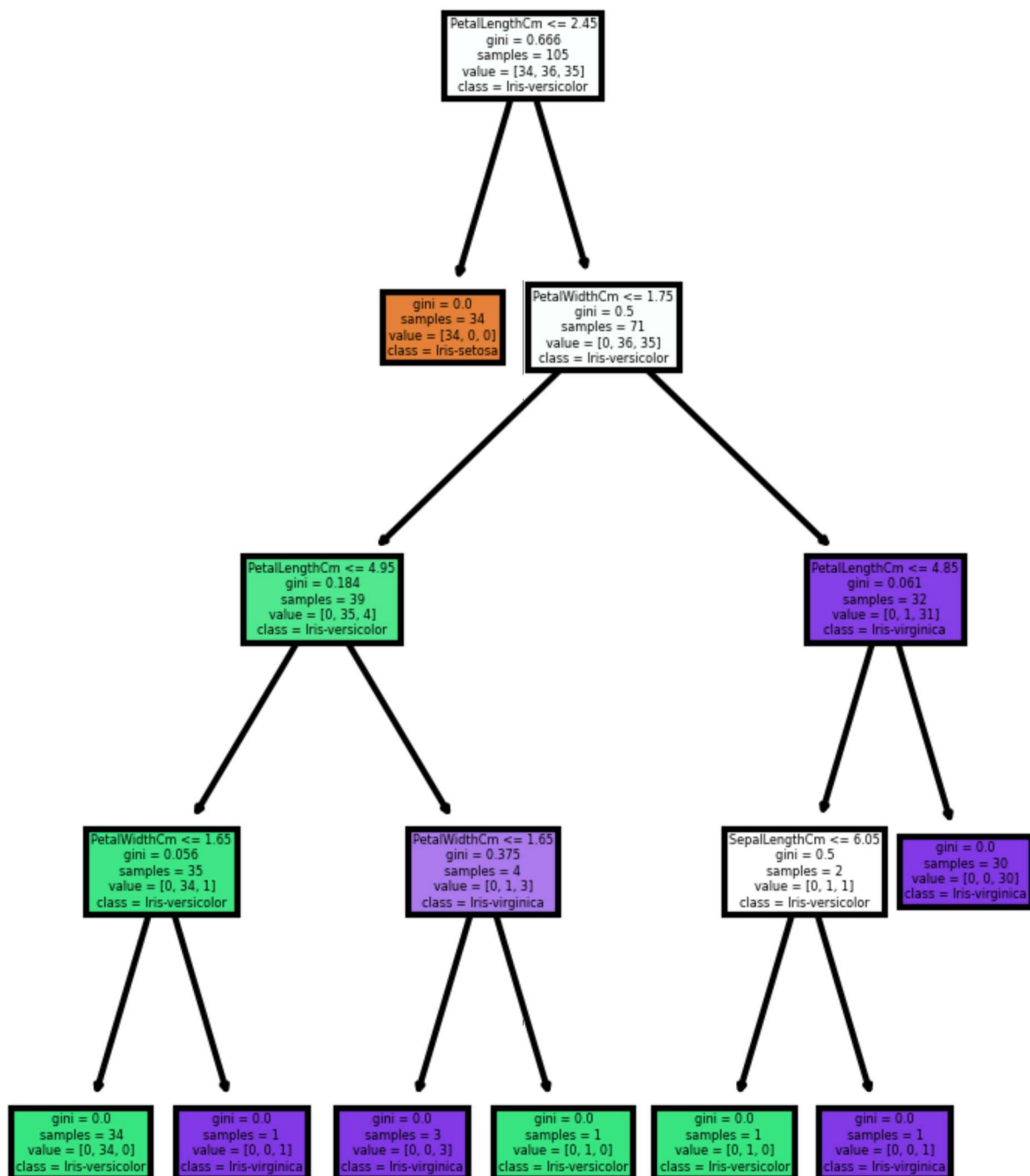
$$\mathbf{GINI} = 1 - \Sigma(p_i)^2$$

Where  $p_i$  is the probability that a tuple in  $D$  belongs to the class  $C$

```
In [10]: cart = DecisionTreeClassifier(criterion='gini')
cart.fit(x_train,y_train)
```

```
Out[10]: DecisionTreeClassifier()
```

```
In [14]: #plot tree
fig, axes = plt.subplots(nrows = 1,ncols = 1,figsize = (3,4), dpi=300)
plot_tree(cart,
          feature_names = feat_name,
          class_names=class_name,
          filled = True);
```



```
In [12]: #check accurecy
y_pred = cart.predict(x_test)
print(f'Accuracy Score of model is : {accuracy_score(y_test,y_pred)}')
```

Accuracy Score of model is : 0.9555555555555556

```
In [13]: print('Confusion Metrics :')
print(confusion_matrix(y_test,y_pred))
```

Confusion Metrics :  
[[16 0 0]  
 [ 0 13 1]  
 [ 0 1 14]]

## Logistic Regression

Logistic Regression is a Machine Learning algorithm which is used for the classification problems, it is a predictive analysis algorithm and based on the concept of probability.

**For Logistic Regression Output Function :**

$$\sigma(z) = \text{sigmoid}(z)$$

Where

$$z = \beta_0 + \beta_1 x + \dots$$

And

$$\text{sigmoid}(z) = \frac{1}{1 + e^{-z}} = \frac{1}{1 + e^{-\beta_0 + \beta_1 x + \dots}}$$

$\sigma(z)$  gives values between 0 and 1.

and Cost Function is :

$$J(\theta) = -\frac{1}{m} \sum [y \log \sigma(z) + (1 - y) \log (1 - \sigma(z))]$$

To minimise our cost Function we use Gradient Descent .

$$\theta_j := \theta_j - \alpha \sum_{i=1}^m (\sigma(x^{(i)}) - y^{(i)}) x_j^{(i)}$$

```
In [10]: #import useful package
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
import matplotlib.pyplot as plt
from sklearn.metrics import confusion_matrix, accuracy_score
```

```
In [11]: df = pd.read_csv('data/Social_Network_Ads.csv')
df.head()
```

```
Out[11]:
```

	User ID	Gender	Age	EstimatedSalary	Purchased
0	15624510	Male	19	19000	0
1	15810944	Male	35	20000	0
2	15668575	Female	26	43000	0
3	15603246	Female	27	57000	0
4	15804002	Male	19	76000	0

```
In [12]: df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 400 entries, 0 to 399
Data columns (total 5 columns):
#   Column                Non-Null Count  Dtype
---  -
0   User ID                400 non-null   int64
1   Gender                 400 non-null   object
2   Age                    400 non-null   int64
3   EstimatedSalary        400 non-null   int64
4   Purchased              400 non-null   int64
dtypes: int64(4), object(1)
memory usage: 15.8+ KB
```

```
In [13]: #convert class feature in numerical
g_list = ['Male', 'Female']
df.Gender = df.Gender.apply(lambda x: g_list.index(x))
```

```
In [15]: #Splitting Data in train and test
X = df.drop(['User ID', 'Purchased'], axis = 1)
Y = df['Purchased']

x_train, x_test, y_train, y_test = train_test_split(X, Y, test_size = 0.3)
```

```
In [16]: #Training Logistic Regression Model
LogReg = LogisticRegression()
LogReg.fit(X, Y)
```

```
Out[16]: LogisticRegression()
```

```
In [26]: print("Coefficient for Regression:")  
print(*LogReg.coef_[0], sep = '\n')
```

```
Coefficient for Regression:  
-9.322295567048426e-11  
-2.1041517809748262e-09  
-2.693014040541572e-06
```

```
In [28]: #Test accurecy  
y_pred = LogReg.predict(x_test)  
  
print(f'Accurecy = {accuracy_score(y_test,y_pred)}')
```

```
Accurecy = 0.6416666666666667
```

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