## In [1]:

4

0

1

```
import pandas as pd
import numpy as np
#***********
#
#
    Decision Tree, Random Forest & Gradient Boosting
#
#>>> 1> Read file
titanic = pd. read csv('http://biostat.mc. vanderbilt. edu/wiki/pub/Main/DataSets/titanic. txt')
print(titanic.head()) #Pre-look at titanic as a dataframe
  row.names pclass survived
0
         1
             1st
                        1
         2
                        0
1
             1st
2
                        0
         3
             1st
3
                        0
         4
             1st
4
         5
             1st
                        1
                                                       embarked
                                       name
                                                age
                  Allen, Miss Elisabeth Walton
0
                                            29.0000
                                                    Southampton
1
                   Allison, Miss Helen Loraine
                                             2.0000
                                                    Southampton
2
            Allison, Mr Hudson Joshua Creighton
                                            30,0000
                                                    Southampton
  Allison, Mrs Hudson J.C. (Bessie Waldo Daniels)
3
                                            25.0000
                                                    Southampton
```

0.9167

boat

NaN

NaN

11

(135)

2

ticket

NaN

NaN

NaN

NaN

24160 L221

Southampton

sex

female

female

female

male

male

Allison, Master Hudson Trevor

B-5

C26

C26

home.dest room

St Louis, MO

Montreal, PQ / Chesterville, ON

Montreal, PQ / Chesterville, ON

4 Montreal, PQ / Chesterville, ON C22

3 Montreal, PQ / Chesterville, ON

```
print(titanic.info()) #Look at titanic statistic information
```

<class 'pandas.core.frame.DataFrame'> RangeIndex: 1313 entries, 0 to 1312 Data columns (total 11 columns): 1313 non-null int64 row.names 1313 non-null object pclass survived 1313 non-null int64 1313 non-null object name 633 non-null float64 age 821 non-null object embarked 754 non-null object home. dest 77 non-null object room 69 non-null object ticket 347 non-null object boat 1313 non-null object sex dtypes: float64(1), int64(2), object(8)memory usage: 112.9+ KB

None

## In [3]:

```
#>>> 2> Select features and Vectorize
X = titanic[['pclass', 'age', 'sex']]
y = titanic['survived']
print(X.info()) # Check features information
```

<class 'pandas.core.frame.DataFrame'> RangeIndex: 1313 entries, 0 to 1312 Data columns (total 3 columns): pclass 1313 non-null object 633 non-null float64 age 1313 non-null object sex dtypes: float64(1), object(2) memory usage: 30.9+ KB None

```
In [4]:
```

X['age'].fillna(X['age'].mean(),inplace=True)

```
print(X.info()) # Check again after filling out
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1313 entries, 0 to 1312
Data columns (total 3 columns):
pclass
          1313 non-null object
          1313 non-null float64
age
          1313 non-null object
sex
dtypes: float64(1), object(2)
memory usage: 30.9+ KB
None
C:\Program Files\Anaconda3\lib\site-packages\pandas\core\generic.py:3191: SettingWit
hCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/in
dexing. html#indexing-view-versus-copy (http://pandas.pydata.org/pandas-docs/stable/i
ndexing.html#indexing-view-versus-copy)
  self. update inplace (new data)
```

## In [5]:

```
from sklearn.cross_validation import train_test_split
from sklearn.feature_extraction import DictVectorizer

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.25, random_state=33) #Split data

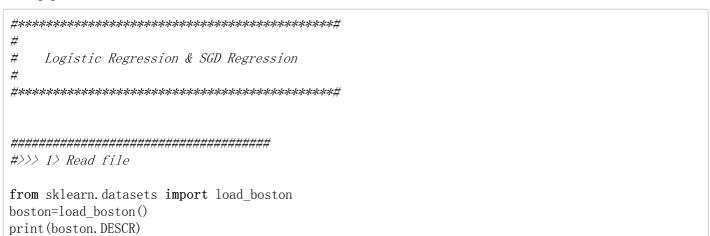
vec = DictVectorizer(sparse=False) #Vectorize
X_train=vec.fit_transform(X_train.to_dict(orient='record'))
print(vec.feature_names_)
X_test=vec.transform(X_test.to_dict(orient='record'))
```

['age', 'pclass=1st', 'pclass=2nd', 'pclass=3rd', 'sex=female', 'sex=male']

```
#>>> 3> Build model and Predict
# Decision Tree
from sklearn tree import DecisionTreeClassifier
dtc=DecisionTreeClassifier()
dtc.fit(X_train, y_train)
y_predict=dtc.predict(X_test)
# Random Forest
from sklearn.ensemble import RandomForestClassifier
rfc=RandomForestClassifier()
rfc.fit(X_train, y_train)
rfc_y_pred=rfc.predict(X_test)
# Gradient Boosting
from sklearn.ensemble import GradientBoostingClassifier
gbc=GradientBoostingClassifier()
gbc. fit(X_train, y_train)
gbc_y_pred=gbc.predict(X_test)
#>>> 4> Score
from sklearn.metrics import classification_report
# Decision Tree
print('The accuracy of Decision Tree is', dtc.score(X_test, y_test))
print(classification_report(y_predict, y_test))
# Random Forest
print ('The accuracy of Random Forest Classifier is', rfc. score(X_test, y_test))
print(classification_report(rfc_y_pred, y_test))
# Gradient Boosting
print ('The accuracy of Gradient Boosting Classifier is', gbc.score(X_test, y_test))
print(classification_report(gbc_y_pred, y_test))
```

The accuracy of Decision Tree is 0.781155015198 precision recall f1-score support					
0 1	0. 91 0. 58	0. 78 0. 80	0. 84 0. 67	236 93	
avg / total	0.81	0.78	0.79	329	
The accuracy of Random Forest Classifier is 0.781155015198 precision recall fl-score support					
0	0.90	0.78	0.83	234	
1	0. 59	0.79	0.68	95	
avg / total	0.81	0.78	0.79	329	
The accuracy	of Gradient precision		Classifier f1-score	is 0.7902 support	73556231
0	0.92	0.78	0.84	239	
1	0.58	0.82	0.68	90	
avg / total	0.83	0.79	0.80	329	

# In [8]:



Boston House Prices dataset

#### Notes

\_\_\_\_\_

#### Data Set Characteristics:

:Number of Instances: 506

:Number of Attributes: 13 numeric/categorical predictive

:Median Value (attribute 14) is usually the target

:Attribute Information (in order):

- CRIM per capita crime rate by town

- ZN proportion of residential land zoned for lots over 25,000 sq.ft.

- INDUS proportion of non-retail business acres per town

- CHAS Charles River dummy variable (= 1 if tract bounds river; 0 otherw

ise)

- NOX nitric oxides concentration (parts per 10 million)

- RM average number of rooms per dwelling

AGE proportion of owner-occupied units built prior to 1940
 DIS weighted distances to five Boston employment centres

RAD index of accessibility to radial highways
 TAX full-value property-tax rate per \$10,000

- PTRATIO pupil-teacher ratio by town

- B 1000 (Bk - 0.63) 2 where Bk is the proportion of blacks by town

- LSTAT % lower status of the population

- MEDV Median value of owner-occupied homes in \$1000's

:Missing Attribute Values: None

:Creator: Harrison, D. and Rubinfeld, D.L.

This is a copy of UCI ML housing dataset.

http://archive.ics.uci.edu/ml/datasets/Housing (http://archive.ics.uci.edu/ml/datasets/Housing)

This dataset was taken from the StatLib library which is maintained at Carnegie Mell on University.

The Boston house-price data of Harrison, D. and Rubinfeld, D.L. 'Hedonic prices and the demand for clean air', J. Environ. Economics & Management, vol.5, 81-102, 1978. Used in Belsley, Kuh & Welsch, 'Regression diagnostics ...', Wiley, 1980. N.B. Various transformations are used in the table on pages 244-261 of the latter.

The Boston house-price data has been used in many machine learning papers that address regression problems.

#### \*\*References\*\*

- Belsley, Kuh & Welsch, 'Regression diagnostics: Identifying Influential Data and Sources of Collinearity', Wiley, 1980. 244-261.
- Quinlan, R. (1993). Combining Instance-Based and Model-Based Learning. In Proceedings on the Tenth International Conference of Machine Learning, 236-243, University of Massachusetts, Amherst. Morgan Kaufmann.
- many more! (see http://archive.ics.uci.edu/ml/datasets/Housing) (http://archive.ics.uci.edu/ml/datasets/Housing))

```
#>>> 2> Split data and preprocessing
# Split data
from sklearn.cross_validation import train_test_split
X=boston. data
y=boston. target
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.25, random_state=33) #Split data
print('The max target value is', np.max(boston.target))
print('The min target value is', np.min(boston.target))
print('The average target value is', np.mean(boston.target))
# preprocessing
from sklearn.preprocessing import StandardScaler
ss_X=StandardScaler()
ss_y=StandardScaler()
X_train=ss_X.fit_transform(X_train)
X test=ss X. fit transform(X test)
y_train=ss_y.fit_transform(y train)
y_test=ss_y.fit_transform(y_test)
The max target value is 50.0
```

The average target value is 22.5328063241

C:\Program Files\Anaconda3\lib\site-packages\sklearn\preprocessing\data.py:583: DeprecationWarning: Passing 1d arrays as data is deprecated in 0.17 and will raise Value Error in 0.19. Reshape your data either using X.reshape(-1, 1) if your data has a si

warnings.warn(DEPRECATION\_MSG\_1D, DeprecationWarning)

ngle feature or X. reshape(1, -1) if it contains a single sample.

The min target value is 5.0

C:\Program Files\Anaconda3\lib\site-packages\sklearn\preprocessing\data.py:646: Depr ecationWarning: Passing 1d arrays as data is deprecated in 0.17 and will raise Value Error in 0.19. Reshape your data either using X.reshape(-1, 1) if your data has a single feature or X.reshape(1, -1) if it contains a single sample.

warnings.warn(DEPRECATION\_MSG\_1D, DeprecationWarning)

C:\Program Files\Anaconda3\lib\site-packages\sklearn\preprocessing\data.py:583: Depr ecationWarning: Passing 1d arrays as data is deprecated in 0.17 and will raise Value Error in 0.19. Reshape your data either using X.reshape(-1, 1) if your data has a single feature or X.reshape(1, -1) if it contains a single sample.

warnings.warn(DEPRECATION\_MSG\_1D, DeprecationWarning)

C:\Program Files\Anaconda3\lib\site-packages\sklearn\preprocessing\data.py:646: Depr ecationWarning: Passing 1d arrays as data is deprecated in 0.17 and will raise Value Error in 0.19. Reshape your data either using X.reshape(-1, 1) if your data has a single feature or X.reshape(1, -1) if it contains a single sample.

warnings. warn (DEPRECATION MSG 1D, DeprecationWarning)

```
#>>> 3> Build model and Predict
# Linear Regression
from sklearn.linear model import LinearRegression
1r=LinearRegression()
lr. fit(X_train, y_train)
lr_y_predict=lr.predict(X_test)
# SGDRegressor
from sklearn.linear_model import SGDRegressor
sgdr=SGDRegressor()
sgdr. fit (X_train, y_train)
sgdr_y_pred=sgdr.predict(X_test)
#>>> 4> Score
# Linear Regression
print('The value of default measurement of LinearRegression is', lr. score(X_test, y_test))
from sklearn.metrics import r2_score, mean_squared_error, mean_absolute_error
# R-squared
print('The value of R-squared of LinearRegression is', r2_score(y_test, lr_y_predict))
# Mean squared error
print ('The value of mean squared error of LinearRegression is', mean_squared_error (ss_y. inverse_trans
# Mean absolute error
print ('The value of mean absolute error of LinearRegression is', mean_absolute_error (ss_y.inverse_tra
The value of default measurement of LinearRegression is 0.676930350524
```

The value of default measurement of LinearRegression is 0.676930350524 The value of R-squared of LinearRegression is 0.676930350524 The value of mean squared error of LinearRegression is 25.0512388542 The value of mean absolute error of LinearRegression is 3.51371562676

```
In [45]:
```

```
#
#
         SVM
#
#>>> 1> Read file
titanic = pd. read_csv('http://biostat.mc. vanderbilt.edu/wiki/pub/Main/DataSets/titanic.txt')
#>>> 2> Vectorize
X = titanic.drop(['name', 'row.names', 'survived'], axis=1)
y = titanic['survived']
#fill missing value
X['age'].fillna(X['age'].mean(),inplace=True)
X. fillna('UNKNOWN', inplace=True)
from sklearn.cross validation import train test split
from sklearn.feature_extraction import DictVectorizer
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.25, random_state=33) #Split data
vec = DictVectorizer()
X_train=vec.fit_transform(X_train.to_dict(orient='record'))
print(len(vec.feature_names_))
X_test=vec. transform(X_test. to_dict(orient='record'))
```

474

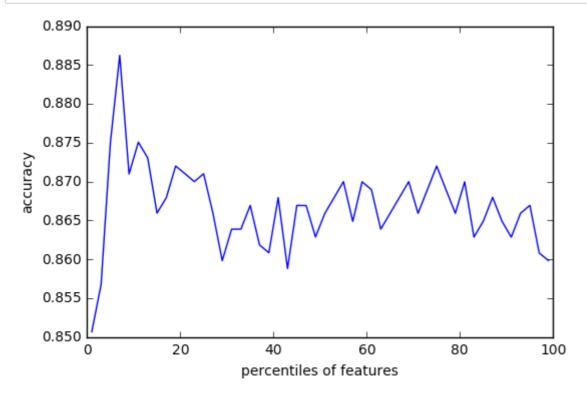
## In [46]:

0.823708206687

```
In [13]:
```

```
#>>> 4> Select feature and predict
from sklearn import feature_selection
fs=feature selection. SelectPercentile (feature selection. chi2, percentile=20)
X_train_fs=fs.fit_transform(X_train, y_train)
dt.fit(X_train_fs, y_train)
X test fs=fs.transform(X test)
print(dt.score(X_test_fs, y_test))
0.817629179331
In \lceil 47 \rceil:
#>>> 5> Validation
from sklearn.cross_validation import cross_val_score
percentiles=range(1, 100, 2)
results=[]
for i in percentiles:
   fs=feature selection. SelectPercentile(feature selection. chi2, percentile=i)
   X_train_fs=fs.fit_transform(X_train, y_train)
   scores=cross_val_score(dt, X_train_fs, y_train, cv=5)
   results=np. append (results, scores. mean ())
print(results)
[ 0.85063904  0.85673057
                       0. 87501546 0. 88622964
                                              0.86590394 0.86998557
 0.86896516 0.87302618 0.86591424
                                   0.87097506
                                              0.86589363 0.86691404
 0.86795506 0.86386312 0.87097506 0.86590394
                                              0.87199546 0.86691404
 0. 86486291 0. 87198516 0. 8597918
                                   0.86791383
                                              0.86792414 0.87096475
 0.86894455
            0.87199546
                       0.87097506
                                   0.87404659
                                              0.87198516 0.86893424
 0. 86996496 0. 86385281
                       0.87098536
                                   0.86998557
                                              0.87096475 0.86996496
 0.86795506
                                              0.87098536 0.86691404
 0.86084313 0.8618223
 0.8598021
            0.85675119
In [48]:
opt=np. where (results==results. max())[0]
print('Optimal number of features %d'%percentiles[opt]) #I still couldn't find out the reason why it
                                       Traceback (most recent call last)
TypeError
<ipython-input-48-3fa7ef434f1d> in <module>()
     1 opt=np. where (results==results. max()) [0]
----> 2 print('Optimal number of features %d'%percentiles[opt])
TypeError: only integer scalar arrays can be converted to a scalar index
```

## In [31]:



## In [34]:

```
#Validation
from sklearn import feature_selection

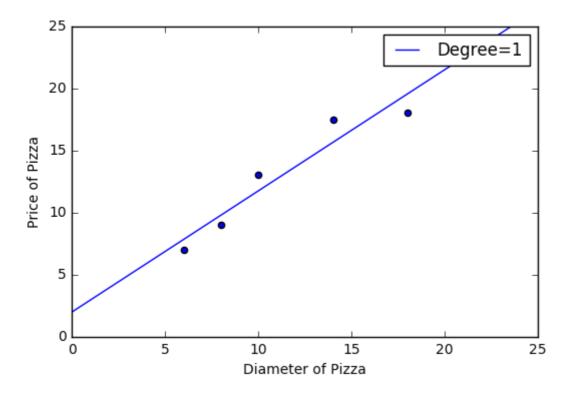
fs=feature_selection. SelectPercentile(feature_selection. chi2, percentile=20)

X_train_fs=fs. fit_transform(X_train, y_train)
dt. fit(X_train_fs, y_train)

X_test_fs=fs. transform(X_test)
print(dt. score(X_test_fs, y_test))
```

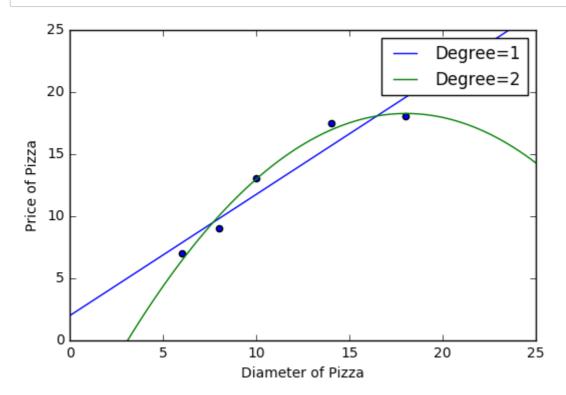
0.823708206687

```
#***********
#
#
         Regularization
#
#************
#linearregression
#>>> 1> Input dataset
X_train=[[6], [8], [10], [14], [18]]
y_train=[[7], [9], [13], [17.5], [18]]
#>>> 2> Build linearregression model
from sklearn. linear model import LinearRegression
regressor=LinearRegression()
regressor.fit(X_train, y_train)
#>>> 3> Predict
xx=np. linspace (0, 26, 100)
xx=xx. reshape (xx. shape [0], 1)
yy=regressor.predict(xx)
#>>> 4> Visualization and Output
import matplotlib.pyplot as plt
plt. scatter(X_train, y_train)
plt1, =plt. plot(xx, yy, label='Degree=1')
plt. axis([0, 25, 0, 25])
plt.xlabel('Diameter of Pizza')
plt.ylabel('Price of Pizza')
plt.legend(handles=[plt1])
plt.show()
print ('The R-squared value of Linear Regressor performing on the training data is', regressor.score(X
```



The R-squared value of Linear Regressor performing on the training data is 0.9100015 96424

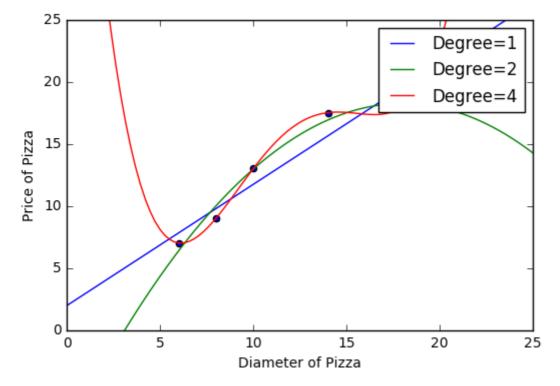
```
#polynominal featrues in degree 2
#>>> 2&3> Build model and Predict
from sklearn.preprocessing import PolynomialFeatures
poly2=PolynomialFeatures(degree=2)
X_train_poly2=poly2.fit_transform(X_train)
regressor_poly2=LinearRegression()
regressor_poly2.fit(X_train_poly2, y_train)
xx poly2=poly2. transform(xx)
yy_poly2=regressor_poly2.predict(xx_poly2)
#>>> 4> Visualization and Output
plt. scatter(X_train, y_train)
plt1, =plt. plot(xx, yy, label='Degree=1')
plt2, =plt. plot(xx, yy_poly2, label='Degree=2')
plt. axis([0, 25, 0, 25])
plt.xlabel('Diameter of Pizza')
plt.ylabel('Price of Pizza')
plt. legend (handles=[plt1, plt2])
plt. show()
print ('The R-squared value of Polynominal Regressor (Degree=2) performing on the training data is', re
```



The R-squared value of Polynominal Regressor(Degree=2) performing on the training da ta is 0.98164216396

```
#polynominal featrues in degree 4
#>>> 2&3> Build model and Predict
from sklearn.preprocessing import PolynomialFeatures
poly4=PolynomialFeatures (degree=4)
X_train_poly4=poly4.fit_transform(X_train)
regressor_poly4=LinearRegression()
regressor poly4. fit (X train poly4, y train)
xx poly4=poly4. transform(xx)
yy_poly4=regressor_poly4.predict(xx_poly4)
print(regressor_poly4.coef_)
print(np. sum(regressor_poly4.coef_**2))
#>>> 4> Visualization and Output
plt. scatter(X_train, y_train)
plt1, =plt. plot(xx, yy, label='Degree=1')
plt2, =plt. plot(xx, yy_poly2, label='Degree=2')
plt4, =plt. plot(xx, yy_poly4, label='Degree=4')
plt. axis([0, 25, 0, 25])
plt.xlabel('Diameter of Pizza')
plt.ylabel('Price of Pizza')
plt.legend(handles=[plt1, plt2, plt4])
plt.show()
print ('The R-squared value of Polynominal Regressor (Degree=4) performing on the training data is', re
   0.00000000e+00 -2.51739583e+01
                                    3. 68906250e+00 -2. 12760417e-01
```





The R-squared value of Polynominal Regressor (Degree=4) performing on the training da ta is  $1.0\,$ 

# In [40]: #>>> 5> Validation #Prepare test dataset X test=[[6], [8], [11], [16]] y\_test=[[8], [12], [15], [18]] #linear regression print ('The R-squared value of Linear Regressor performing on the test data is', regressor.score(X\_tes #polynominal 2 X test poly2=poly2.fit\_transform(X\_test) print ('The R-squared value of Polynominal Regressor (Degree=2) performing on the test data is', regres #polynominal 4 X test poly4=poly4.fit transform(X test) print ('The R-squared value of Polynominal Regressor (Degree=4) performing on the test data is', regres The R-squared value of Linear Regressor performing on the test data is 0.80972683246 The R-squared value of Polynominal Regressor(Degree=2) performing on the test data i s 0.867544365635 The R-squared value of Polynominal Regressor(Degree=4) performing on the test data i s 0.809588079577 In [42]: #lasso #>>> 2&3> Build model and Predict

# 0.83889268736

- [ 0.00000000e+00 0.00000000e+00 1.17900534e-01 5.42646770e-05 -2.23027128e-04]
- C:\Program Files\Anaconda3\lib\site-packages\sklearn\linear\_model\coordinate\_descen t.py:466: ConvergenceWarning: Objective did not converge. You might want to increase the number of iterations ConvergenceWarning)