Titanic Survivor Classification

Description

This is notebook to build classification for Titanic passengers to classify if those passengers survived or not. Our aim is to get model with the most accuracy possible.

source: https://www.kaggle.com/competitions/titanic/data

Data Overview

import pandas as pd

data.info()

data = pd.read csv('train.csv')

In [1]:

```
<class 'pandas.core.frame.DataFrame'>
       RangeIndex: 891 entries, 0 to 890
       Data columns (total 12 columns):
          Column Non-Null Count Dtype
                       -----
        0
          PassengerId 891 non-null
                                     int64
        1 Survived 891 non-null int64
2 Pclass 891 non-null int64
                      891 non-null object
                      891 non-null object
714 non-null float64
        4 Sex
           Age
        6 SibSp
                      891 non-null int64
        7 Parch
                      891 non-null int64
                      891 non-null object
        8 Ticket
          Fare
                      891 non-null float64
        10 Cabin
                      204 non-null object
        11 Embarked 889 non-null object
       dtypes: float64(2), int64(5), object(5)
       memory usage: 83.7+ KB
In [2]:
       data.describe()
```

| Out[2]: | | PassengerId | Survived | Pclass | Age | SibSp | Parch | Fare |
|---------|-------|-------------|------------|------------|------------|------------|------------|------------|
| | count | 891.000000 | 891.000000 | 891.000000 | 714.000000 | 891.000000 | 891.000000 | 891.000000 |
| | mean | 446.000000 | 0.383838 | 2.308642 | 29.699118 | 0.523008 | 0.381594 | 32.204208 |
| | std | 257.353842 | 0.486592 | 0.836071 | 14.526497 | 1.102743 | 0.806057 | 49.693429 |
| | min | 1.000000 | 0.000000 | 1.000000 | 0.420000 | 0.000000 | 0.000000 | 0.000000 |
| | 25% | 223.500000 | 0.000000 | 2.000000 | 20.125000 | 0.000000 | 0.000000 | 7.910400 |
| | 50% | 446.000000 | 0.000000 | 3.000000 | 28.000000 | 0.000000 | 0.000000 | 14.454200 |
| | 75% | 668.500000 | 1.000000 | 3.000000 | 38.000000 | 1.000000 | 0.000000 | 31.000000 |
| | max | 891.000000 | 1.000000 | 3.000000 | 80.000000 | 8.000000 | 6.000000 | 512.329200 |

```
In [3]: data.head()
```

| Out[3]: | Passengerld | Survived | Pclass | Name | Sex | Age | SibSp | Parch | Ticket | Fare | Cabin | Embarked |
|---------|--|---|--------|--|--------|------|-------|-------|---------------------|---------|-------|----------|
| | 0 1 | 0 | 3 | Braund, Mr. Owen Harris | male | 22.0 | 1 | 0 | A/5 21171 | 7.2500 | NaN | S |
| | 1 2 | 1 | 1 | Cumings, Mrs. John Bradley (Florence Briggs Th | female | 38.0 | 1 | 0 | PC 17599 | 71.2833 | C85 | С |
| | 2 3 | 1 | 3 | Heikkinen, Miss. Laina | female | 26.0 | 0 | 0 | STON/O2. 3101282 | 7.9250 | NaN | S |
| | 3 4 | 1 | 1 | Futrelle, Mrs. Jacques Heath (Lily May Peel) | female | 35.0 | 1 | 0 | 113803 | 53.1000 | C123 | S |
| | 4 5 | 0 | 3 | Allen, Mr. William Henry | male | 35.0 | 0 | 0 | 373450 | 8.0500 | NaN | S |
| In [4]: | data.nuniqu | e() | | | | | | | | | | |
| Out[4]: | PassengerId Survived Pclass Name Sex Age SibSp Parch Ticket Fare Cabin Embarked dtype: int64 | 891 2 3 891 2 88 7 7 681 248 147 3 | | | | | | | | | | |

So, here is Describe for each column

Out[3]:

• Passengerld: id for each passenger for this data

• Survived : if this passenger survived or not (0 = No, 1 = Yes)

• Pclass : Ticket class (1 = 1st, 2 = 2nd, 3 = 3rd)

• Name : name of passenger

• Sex : sex of passenger

• Age : age of passenger

• SibSp: number of siblings / spouses aboard the Titanic

• Parch : number of parents / children aboard the Titanic

• Ticket: Ticket number

• Fare: Passenger fare

• Cabin : Cabin number

• Embarked : Port of Embarkation (C = Cherbourg, Q = Queenstown, S = Southampton)

Clean Up

Remove column that we are not going to use and fill null data.

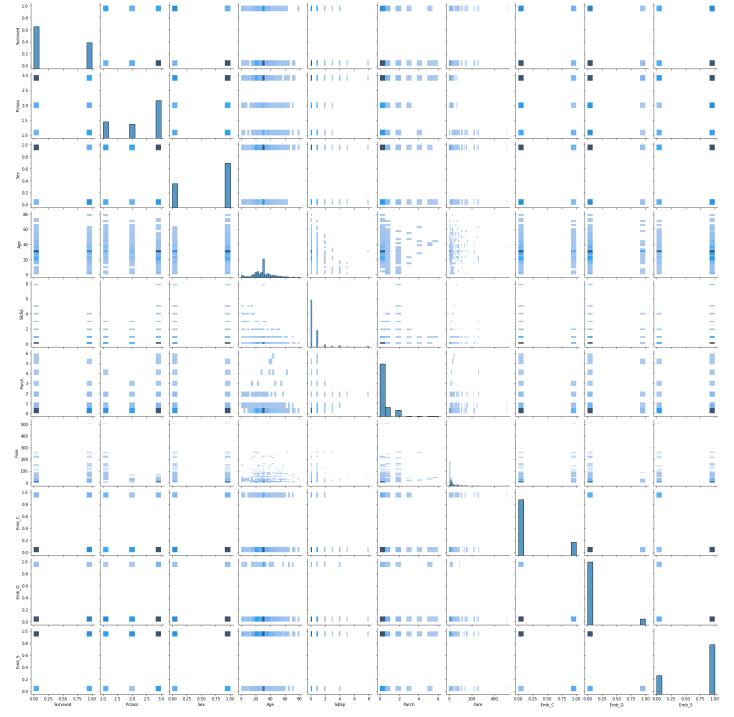
```
# remove cloumns PassengerId, Name, Ticket and Cabin Since it have high cardinality.
In [5]:
        data = data[['Survived','Pclass','Sex','Age','SibSp','Parch','Fare','Embarked']]
In [6]:
        # get mean age
        mean age = data['Age'].mean()
        print('mean age = ', mean age)
         # show histogram for Embarked for impute with the most frequent value
        mode emb = data['Embarked'].mode()
        print('mode Embarked = ', mode emb[0])
        mean age = 29.69911764705882
        mode Embarked = S
In [7]:
         # impute missing data
        values = {'Age' : mean age, 'Embarked': 'S'}
        data = data.fillna(value=values)
In [8]:
         # convert Sex to int and Embarked to 3 int columns
        data['Sex'] = data.Sex.apply(lambda x: int(x == 'male'))
        data['Emb C'] = data.Embarked.apply(lambda x: int(x == 'C'))
        data['Emb Q'] = data.Embarked.apply(lambda x: int(x == 'Q'))
        data['Emb S'] = data.Embarked.apply(lambda x: int(x == 'S'))
        data = data[['Survived', 'Pclass', 'Sex', 'Age', 'SibSp', 'Parch', 'Fare', 'Emb C', 'Emb Q', 'Emk
        data.head()
```

| Out[8]: | | Survived | Pclass | Sex | Age | SibSp | Parch | Fare | Emb_C | Emb_Q | Emb_S |
|---------|---|----------|--------|-----|------|-------|-------|---------|-------|-------|-------|
| | 0 | 0 | 3 | 1 | 22.0 | 1 | 0 | 7.2500 | 0 | 0 | 1 |
| | 1 | 1 | 1 | 0 | 38.0 | 1 | 0 | 71.2833 | 1 | 0 | 0 |
| | 2 | 1 | 3 | 0 | 26.0 | 0 | 0 | 7.9250 | 0 | 0 | 1 |
| | 3 | 1 | 1 | 0 | 35.0 | 1 | 0 | 53.1000 | 0 | 0 | 1 |
| | 4 | 0 | 3 | 1 | 35.0 | 0 | 0 | 8.0500 | 0 | 0 | 1 |

Visual Display

```
import seaborn as sns
import matplotlib.pyplot as plt

sns.pairplot(data, kind="hist")
plt.savefig('pair_plot1.png')
```



In [10]: corr_data = data.corr()
 print(corr_data)

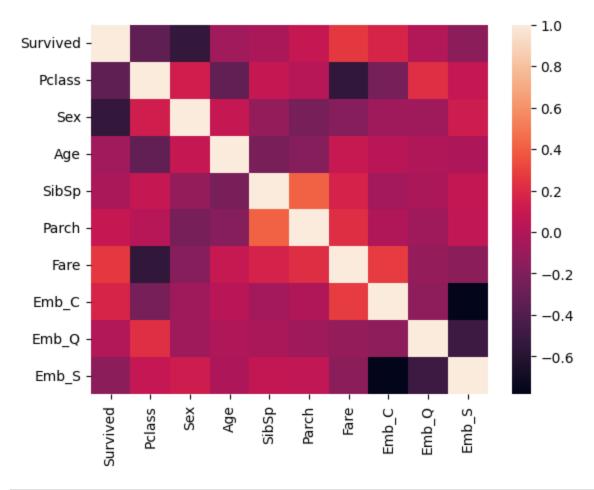
| | Survived | Pclass | Sex | Age | SibSp | Parch | \ |
|----------|-----------|-----------|-----------|-----------|-----------|-----------|---|
| Survived | 1.000000 | -0.338481 | -0.543351 | -0.069809 | -0.035322 | 0.081629 | |
| Pclass | -0.338481 | 1.000000 | 0.131900 | -0.331339 | 0.083081 | 0.018443 | |
| Sex | -0.543351 | 0.131900 | 1.000000 | 0.084153 | -0.114631 | -0.245489 | |
| Age | -0.069809 | -0.331339 | 0.084153 | 1.000000 | -0.232625 | -0.179191 | |
| SibSp | -0.035322 | 0.083081 | -0.114631 | -0.232625 | 1.000000 | 0.414838 | |
| Parch | 0.081629 | 0.018443 | -0.245489 | -0.179191 | 0.414838 | 1.000000 | |
| Fare | 0.257307 | -0.549500 | -0.182333 | 0.091566 | 0.159651 | 0.216225 | |
| Emb_C | 0.168240 | -0.243292 | -0.082853 | 0.032024 | -0.059528 | -0.011069 | |
| Emb_Q | 0.003650 | 0.221009 | -0.074115 | -0.013855 | -0.026354 | -0.081228 | |
| Emb_S | -0.149683 | 0.074053 | 0.119224 | -0.019336 | 0.068734 | 0.060814 | |
| | | | | | | | |
| | Fare | Emb C | Emb Q | Emb S | | | |

Survived 0.257307 0.168240 0.003650 -0.149683 Pclass -0.549500 -0.243292 0.221009 0.074053 Sex -0.182333 -0.082853 -0.074115 0.119224

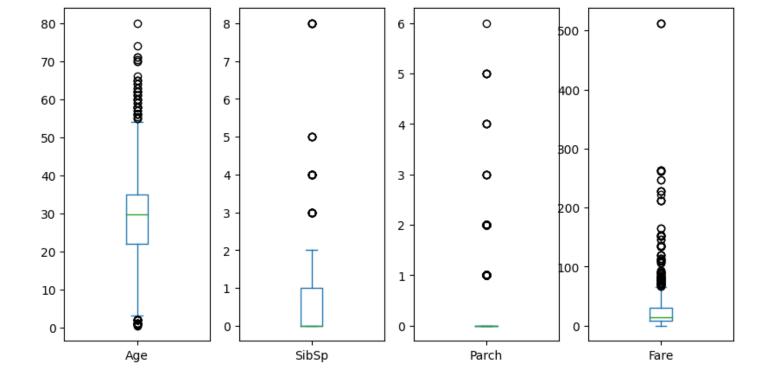
```
0.091566 0.032024 -0.013855 -0.019336
Age
          0.159651 -0.059528 -0.026354
                                      0.068734
SibSp
Parch
          0.216225 -0.011069 -0.081228
                                       0.060814
          1.000000 0.269335 -0.117216 -0.162184
Fare
Emb C
          0.269335 1.000000 -0.148258 -0.782742
Emb Q
         -0.117216 -0.148258 1.000000 -0.499421
         -0.162184 -0.782742 -0.499421 1.000000
Emb S
```

```
In [11]: sns.heatmap(corr data)
```

Out[11]: <AxesSubplot:>



```
In [12]: plot_data = data[['Age','SibSp','Parch','Fare']]
    plot_data.plot(kind="box", subplots=True, figsize =(10, 5))
```



Analysis

From information above, seems there is no strong correlation between features as you and see from correlation heatmap or raw data (>0.7). You may see Emb_C and Emb_S have high value but those columns was extracted from Embarked feature. There are lot of outiler data as you can see. But, I think those are all valid. Age are in reasonable range (more than 0 and less than 90). SibSp and Parch also resonable because most of people board alone. Fare also reasonbale for me.

Classification

I want to use Random Forest Tree to solve this problem. Because we have only fews features.

First, split data in to train and test (test for 20%).

```
In [23]: from sklearn.model_selection import train_test_split

X = data[['Pclass','Sex','Age','SibSp','Parch','Fare','Emb_C', 'Emb_Q', 'Emb_S']]
Y = data['Survived']
x_train, x_test, y_train, y_test = train_test_split(X, Y, test_size=0.2, random_state=1992
print(len(x_train), len(x_test))
712 179
```

Then, Build first model to be our base line.

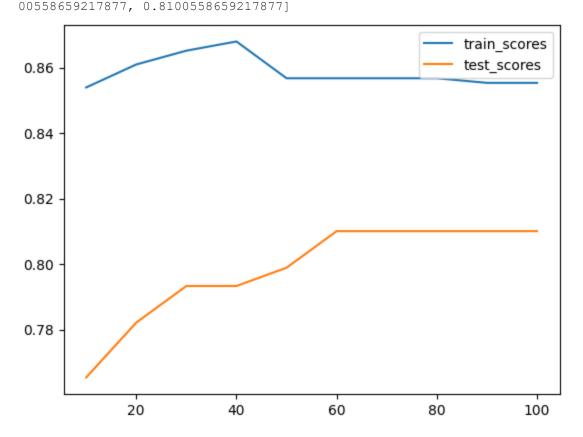
```
In [24]: from sklearn.ensemble import RandomForestClassifier

bl_model = RandomForestClassifier(n_estimators=10, max_depth=5, random_state=1992).fit(x_t
print('score with training data = ',bl_model.score(x_train,y_train))
print('score with test data = ',bl_model.score(x_test,y_test))

score with training data = 0.8539325842696629
score with test data = 0.7653631284916201
```

try to play with n_estimators and max_depth

train_scores = [0.8539325842696629, 0.8609550561797753, 0.8651685393258427, 0.86797752808 98876, 0.8567415730337079, 0.8567415730337079, 0.8567415730337079, 0.8567415730337079, 0.8567415730337079, 0.8553370786516854, 0.8553370786516854]
test_scores = [0.7653631284916201, 0.7821229050279329, 0.7932960893854749, 0.793296089385 4749, 0.7988826815642458, 0.8100558659217877, 0.8100558659217877, 0.81



Try with max_depth=10

```
In [28]:
    es = list(range(10,110,10))
    train_scores = []
    test_scores = []
    for e in es:
        model = RandomForestClassifier(n_estimators=e, max_depth=10, random_state=1992).fit(x_train_scores.append(model.score(x_train,y_train))
        test_scores.append(model.score(x_test,y_test))

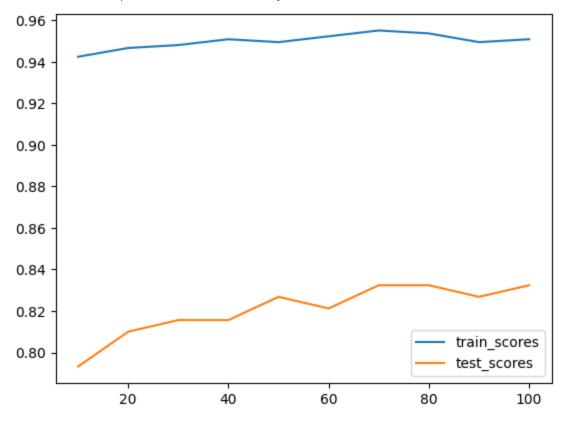
    print('train_scores = ', train_scores)
    print('test_scores = ', test_scores)

plt.plot(es, train_scores)
plt.plot(es, test_scores)
```

```
plt.legend(['train_scores', 'test_scores'])
plt.show()
```

train_scores = [0.9424157303370787, 0.9466292134831461, 0.9480337078651685, 0.95084269662 92135, 0.949438202247191, 0.952247191011236, 0.9550561797752809, 0.9536516853932584, 0.949 438202247191, 0.9508426966292135]

test_scores = [0.7932960893854749, 0.8100558659217877, 0.8156424581005587, 0.8156424581006, 0.8212290502793296, 0.8324022346368715, 0.8324022346368715, 0.8268156424581006, 0.8324022346368715]



From graphs we can see that using max_depth=10 is better. And the reasonal n_estimators is 50.

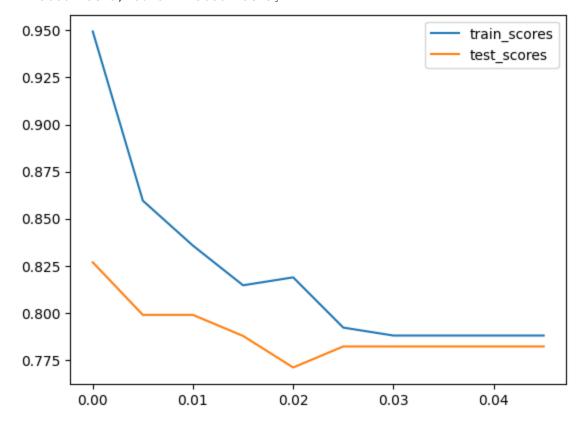
I will try to tune ccp_alpha to reduce data overfitted.

```
In [29]:
         alphas = []
         for i in range(10):
             alphas.append(i*0.005)
         train scores = []
         test scores = []
         for a in alphas:
             model = RandomForestClassifier(n estimators=50, max depth=10, ccp alpha=a, random stat
             train scores.append(model.score(x train, y train))
              test scores.append(model.score(x test,y test))
         print('train scores = ', train scores)
         print('test scores = ', test scores)
         plt.plot(alphas, train scores)
         plt.plot(alphas, test scores)
         plt.legend(['train scores', 'test scores'])
         plt.show()
```

0337, 0.8188202247191011, 0.7921348314606742, 0.7879213483146067, 0.7879213483146067, 0.7879213483146067, 0.7879213483146067, 0.7879213483146067, 0.7879213483146067]
test_scores = [0.8268156424581006, 0.7988826815642458, 0.7988826815642458, 0.787709497206]

train scores = [0.949438202247191, 0.8595505617977528, 0.8356741573033708, 0.814606741573

7039, 0.770949720670391, 0.7821229050279329, 0.7821229050279329, 0.7821229050279329, 0.7821229050279329, 0.7821229050279329]



Incrase ccp_alpha helps reduce over fitted problem. But, I don't want to sacrifice score fro this.

So, the best model for me is n_estimators=50, max_depth=10 and ccp_alpha=0

Use this setting and train with all data we have to predict test data from Kaggle.

In [31]:

```
print('score with training data = ',bl_model.score(X,Y))

score with training data = 0.8361391694725028

In [32]:
    org_test_data = pd.read_csv('test.csv')
    test_data = org_test_data[['Pclass','Sex','Age','SibSp','Parch','Fare','Embarked']]
    mean_fare = data['Fare'].mean()

values = {'Age' : mean_age, 'Fare': mean_fare, 'Embarked': 'S'}
    test_data = test_data.fillna(value=values)
```

best model = RandomForestClassifier(n estimators=50, max depth=10, ccp alpha=0, random sta

```
In [33]:
    test_data['Sex'] = test_data.Sex.apply(lambda x: int(x =='male'))
    test_data['Emb_C'] = test_data.Embarked.apply(lambda x: int(x =='C'))
    test_data['Emb_Q'] = test_data.Embarked.apply(lambda x: int(x =='Q'))
    test_data['Emb_S'] = test_data.Embarked.apply(lambda x: int(x =='S'))

    test_data = test_data[['Pclass','Sex','Age','SibSp','Parch','Fare','Emb_C', 'Emb_Q', 'Emb_
```

```
In [34]: predictions = best_model.predict(test_data)
  output = pd.DataFrame({'PassengerId': org_test_data.PassengerId, 'Survived': predictions})
  output.to_csv('submission.csv', index=False)
  print("Your submission was successfully saved!")
```

Your submission was successfully saved!

Score for submit this prediction to Kaggle = 0.77751

Conclusion

For this problem first we look at overview of data we have. Then, we clean it up (select only useful feature fill null data). After that we do analysis about corrlation and outlier. And, we build classification model with random forest tree and tune it to get the best model. Last, we use our best model to do prediction and submit it to Kaggle.

Actually, I also try Adaboost but it worst than this. But I still think that there are better approach to this problem which I havn't try it.

github: https://github.com/Satjarporn/Titanic

| In []: |]: | |
|---------|----|--|
| | | |