titanic

October 31, 2020

SURVIVAL IN TITANIC PREDICTION

```
[0]: import pandas as pd
    from google.colab import drive
    import seaborn as sns
    %matplotlib inline
    from matplotlib import pyplot as plt
    from matplotlib import style
    drive.mount('/content/gdrive')
    #reading datasets from gdrive
    train = pd.read_csv('gdrive/My Drive/Colab Notebooks/titanic/train.csv', __
     x_test = pd.read_csv('gdrive/My Drive/Colab Notebooks/titanic/test.csv',
    y test = pd.read_csv('gdrive/My Drive/Colab Notebooks/titanic/gender_submission.
    test = x_test.merge(y_test, left_index=True,right_index=True,how='inner')
    X = ⊔
     →train[['Pclass','Name','Sex','Age','SibSp','Parch','Ticket','Fare','Cabin','Embarked',]]
    Y = train[['Survived']]
    train.head()
```

Drive already mounted at /content/gdrive; to attempt to forcibly remount, call drive.mount("/content/gdrive", force_remount=True).

```
[0]:
                   Survived Pclass ... Cabin Embarked
     PassengerId
                           0
                                    3
                                            NaN
                                                        S
     1
                                       •••
     2
                                            C85
                                                        C
                           1
                                    1 ...
                                    3 ...
                                                        S
     3
                           1
                                            NaN
     4
                                    1 ... C123
                                                        S
                           1
                                    3 ...
     5
                                           NaN
                                                        S
```

[5 rows x 11 columns]

```
[0]: import numpy as np import matplotlib.pyplot as plt
```

```
from astropy.visualization import hist
#checking the dataset's features for inconsistency
#getting column names
columns = train.columns.values
print(columns)
#here's table with each feature's missing values
total = train.isnull().sum().sort_values(ascending=False)
percent_1 = train.isnull().sum()/train.isnull().count()*100
percent_2 = (round(percent_1, 1)).sort_values(ascending=False)
missing_data = pd.concat([total, percent_2], axis=1, keys=['Total', '%'])
print(missing_data)
#looking at age and sex
male_survived = train[['Survived', 'Sex', 'Age']]
female_survived = train[['Survived','Sex','Age']]
#filtering for respective sex
male_survived = male_survived[male_survived.Sex == 'male']
female_survived = female_survived[female_survived.Sex == 'female']
male_alive = len(male_survived[male_survived.Survived == 1])/len(male_survived.
→Survived)
female_alive = len(female_survived[female_survived.Survived == 1])/
→len(female_survived.Survived)
print('males alive : ',male_alive*100,"%")
print('female alive : ',female_alive*100,"%")
#dropping rows with empty age field
male_survived = male_survived[pd.isnull(male_survived.Age) == False]
female_survived = female_survived[pd.isnull(female_survived.Age) == False]
#plotting the graph
survived = 'survived'
not survived = 'not survived'
fig, axes = plt.subplots(nrows=1, ncols=2,figsize=(15, 6))
ax = sns.distplot(male_survived[male_survived['Survived']==1].Age, bins='auto',__
→label = survived, ax = axes[0], kde =False)
ax = sns.distplot(male survived[male survived['Survived']==0].Age, bins='auto', ___
→label = not_survived, ax = axes[0], kde =False)
ax.legend()
ax.set_title('Males')
ax = sns.distplot(female_survived[female_survived['Survived']==1].Age, __
⇒bins='auto', label = survived, ax = axes[1], kde = False)
ax = sns.distplot(female_survived[female_survived['Survived']==0].Age, __
⇒bins='auto', label = not_survived, ax = axes[1], kde = False)
ax.legend()
ax.set_title('Females')
```

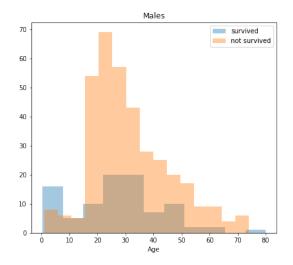
['Survived' 'Pclass' 'Name' 'Sex' 'Age' 'SibSp' 'Parch' 'Ticket' 'Fare'

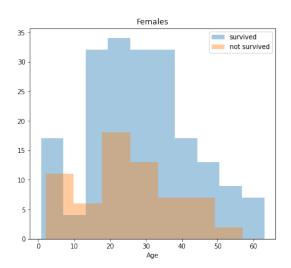
'Cabin'	'Embarked']			
	Total	%		
Cabin	687	77.1		
Age	177	19.9		
Embarked	2	0.2		
Fare	0	0.0		
Ticket	0	0.0		
Parch	0	0.0		
SibSp	0	0.0		
Sex	0	0.0		
Name	0	0.0		
Pclass	0	0.0		
Survived	0	0.0		
males ali	ve :	18.8908	314558058924	%

female alive: 74.20382165605095 %

[0]: Text(0.5, 1.0, 'Females')

[0]:





Age and sex are pretty relevant features since the data is distributed. Above data shows that males only 19% males survived. Most fatalities were among young (16-55 yrs) were the people who died mostly. In Females there wasn't much fatalities as 74% of them survived. Thus we can use these features in the training dataset.

```
[0]: #Embarking represents which passenger entered from which gate, which could_

differ based on passenger class

#checking embarking's correlation with survival rate and Passenger class

print(train.Embarked.unique())

g = sns.FacetGrid(train, col="Embarked")

g.map(sns.pointplot, 'Pclass', 'Survived', 'Sex', palette=None, order=None, 

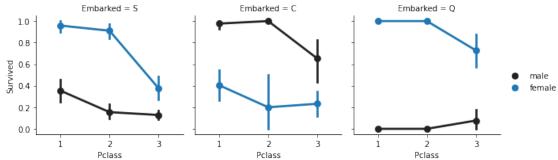
hue_order=None)

g.add_legend()
```

['S' 'C' 'Q' nan]

[0]: <seaborn.axisgrid.FacetGrid at 0x7fc9157a84a8>

[0]:



Here we compare passenger class and survival rate based on their port of entry which is defined in the embarked class.

This data suggests that people who entered the ship from port S, among them first and second class females had a fairly high rate of survival rate. Men apart from first class passgengers barely survived.

In the second graph, people who entered the ship from port C, among them male passengers had a fairly high rate of survival rate. Woman apart from first class passgengers barely survived.

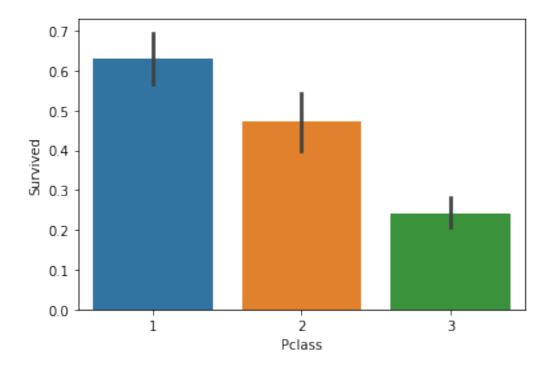
In the thrid graph, people who entered the ship from port Q, among them female passengers had a fairly high rate of survival rate. Man apart from third class passgengers barely survived.

```
[0]: print(train.Pclass.unique())
sns.barplot(x='Pclass', y='Survived', data=train)
```

[3 1 2]

[0]: <matplotlib.axes. subplots.AxesSubplot at 0x7fc9157a02e8>

[0]:

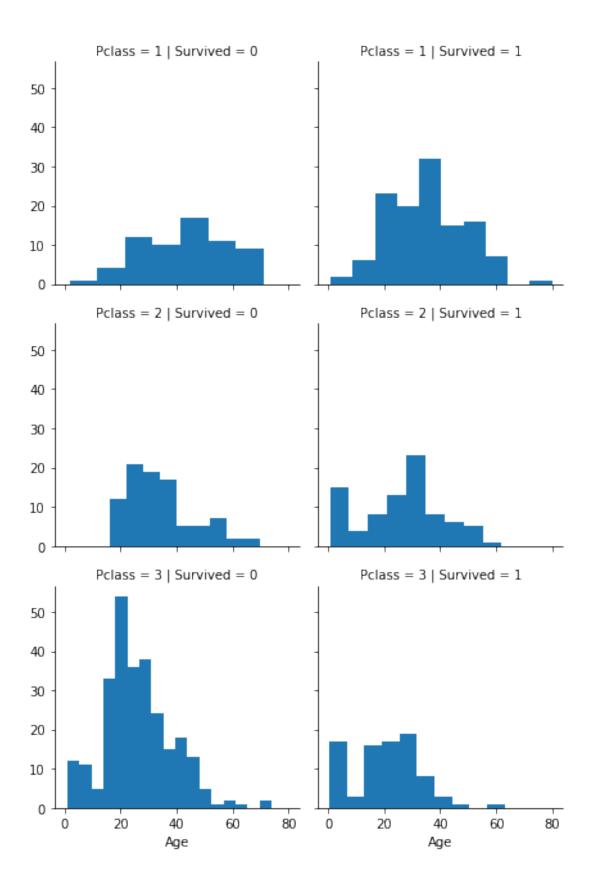


In the embarked feature test, we can see that the passenger class plays an important role in the determining the survival rate. Thus, we plot a graph to see the correlation of passenger class with survival rate.

As we can see, people with higher passenger class had higher survival rate. Thus, its safe to say that passenger class plays an important role in survival.

```
[0]: g = sns.FacetGrid(train, col='Survived', row='Pclass')
g.map(plt.hist, 'Age', bins='auto')
g.add_legend();
```

[0]:



To further elaborate on passenger class, we relate it to age class and see who survived. Turns out maximum, fatalities were among passengers from third class and most survivers were first class passengers.

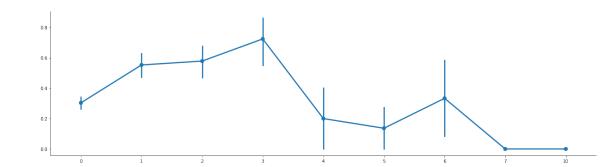
```
[0]: print(train.Parch.unique())
     print(train.SibSp.unique())
     data = [train, test]
     for dataset in data:
         #dataset['relatives'] = dataset['SibSp'] + dataset['Parch']
         dataset.loc[(dataset['SibSp'] + dataset['Parch']) > 0, 'not_alone'] = 1
         dataset.loc[(dataset['SibSp'] + dataset['Parch']) == 0, 'not_alone'] = 0
         dataset['not alone'] = dataset['not alone'].astype(int)
     print(train['not_alone'].value_counts())
     g = sns.factorplot((train.Parch+train.SibSp), 'Survived', data=train , aspect =
      \rightarrow3.5)
    [0 1 2 5 3 4 6]
    [1 0 3 4 2 5 8]
    0
         537
         354
    Name: not_alone, dtype: int64
    /usr/local/lib/python3.6/dist-packages/seaborn/categorical.py:3669: UserWarning:
    The `factorplot` function has been renamed to `catplot`. The original name will
    be removed in a future release. Please update your code. Note that the default
    `kind` in `factorplot` (`'point'`) has changed `'strip'` in `catplot`.
      warnings.warn(msg)
    /usr/local/lib/python3.6/dist-packages/pandas/core/ops/array_ops.py:253:
    FutureWarning: elementwise comparison failed; returning scalar instead, but in
    the future will perform elementwise comparison
      res_values = method(rvalues)
```

```
ValueError
                                           Traceback (most recent call last)
<ipython-input-23-1d35ca25c525> in <module>()
            dataset['not_alone'] = dataset['not_alone'].astype(int)
      9 print(train['not alone'].value counts())
---> 10 g = sns.factorplot((train.Parch+train.SibSp), 'Survived', data=train, __
\rightarrowaspect = 3.5)
/usr/local/lib/python3.6/dist-packages/seaborn/categorical.py in_
→factorplot(*args, **kwargs)
   3677
            kwargs.setdefault("kind", "point")
   3678
-> 3679
            return catplot(*args, **kwargs)
   3680
   3681
```

```
/usr/local/lib/python3.6/dist-packages/seaborn/categorical.py in catplot(x, y, u
→hue, data, row, col, col_wrap, estimator, ci, n_boot, units, seed, order, →hue_order, row_order, col_order, kind, height, aspect, orient, color, palette
→legend, legend_out, sharex, sharey, margin_titles, facet_kws, **kwargs)
   3763
   3764
            # Draw the plot onto the facets
-> 3765
            g.map_dataframe(plot_func, x, y, hue, **plot_kws)
   3766
   3767
            # Special case axis labels for a count type plot
/usr/local/lib/python3.6/dist-packages/seaborn/axisgrid.py in_
→map_dataframe(self, func, *args, **kwargs)
    834
    835
                 # Finalize the annotations and layout
--> 836
                 self. finalize grid(args[:2])
    837
    838
                 return self
/usr/local/lib/python3.6/dist-packages/seaborn/axisgrid.py in_
→ finalize_grid(self, axlabels)
            def _finalize_grid(self, axlabels):
    857
                 """Finalize the annotations and layout."""
    858
                 self.set axis labels(*axlabels)
--> 859
    860
                 self.set_titles()
    861
                 self.fig.tight_layout()
/usr/local/lib/python3.6/dist-packages/seaborn/axisgrid.py in_
 ⇒set_axis_labels(self, x_var, y_var)
    883
                 if x_var is not None:
    884
                     self. x var = x var
--> 885
                     self.set xlabels(x var)
    886
                 if y_var is not None:
    887
                     self. y var = y var
/usr/local/lib/python3.6/dist-packages/seaborn/axisgrid.py in set_xlabels(self,
 →label, **kwargs)
    894
                     label = self._x_var
                 for ax in self._bottom_axes:
    895
--> 896
                     ax.set_xlabel(label, **kwargs)
    897
                 return self
    898
/usr/local/lib/python3.6/dist-packages/matplotlib/axes/_axes.py in_
→set_xlabel(self, xlabel, fontdict, labelpad, **kwargs)
    246
                 if labelpad is not None:
                     self.xaxis.labelpad = labelpad
    247
--> 248
                 return self.xaxis.set label text(xlabel, fontdict, **kwargs)
```

```
249
    250
            def get_ylabel(self):
/usr/local/lib/python3.6/dist-packages/matplotlib/axis.py in_
→set label text(self, label, fontdict, **kwargs)
   1611
   1612
                self.isDefault label = False
                self.label.set text(label)
-> 1613
   1614
                if fontdict is not None:
                    self.label.update(fontdict)
   1615
/usr/local/lib/python3.6/dist-packages/matplotlib/text.py in set_text(self, s)
   1163
                if s is None:
                    s = ''
   1164
                if s != self._text:
-> 1165
   1166
                    self._text = str(s)
   1167
                    self.stale = True
/usr/local/lib/python3.6/dist-packages/pandas/core/generic.py in_
→__nonzero__(self)
  1477
            def __nonzero__(self):
                raise ValueError(
   1478
                    f"The truth value of a {type(self).__name__} is ambiguous.
-> 1479
                    "Use a.empty, a.bool(), a.item(), a.any() or a.all()."
   1480
   1481
                )
ValueError: The truth value of a Series is ambiguous. Use a.empty, a.bool(), a.
\rightarrowitem(), a.any() or a.all().
```

[0]:



Lastly we look at Sibsp(# of siblings present on the ship) and parch(# of parent present on the ship) classes. The above graph shows that people with relatives had a higher chances. The chances seems higher when someone has 1-3 relatives on board.

```
[0]: #Pre Processing data #Filling the missing values
```

```
import re
     #make class deck out of cabin and then drop it
     deck = {"A": 1, "B": 2, "C": 3, "D": 4, "E": 5, "F": 6, "G": 7, "N": 8}
     data = [train, test]
     for dataset in data:
         dataset['Cabin'] = dataset['Cabin'].fillna("NO")
         dataset['Deck'] = dataset['Cabin'].map(lambda x: re.compile("([a-zA-Z]+)").
      \rightarrowsearch(x).group())
         dataset['Deck'] = dataset['Deck'].map(deck)
         dataset['Deck'] = dataset['Deck'].fillna(0)
         dataset['Deck'] = dataset['Deck'].astype(int)# we can now drop the cabin_
      \rightarrow feature
     train = train.drop(columns='Cabin')
     test = test.drop(columns='Cabin')
     #filling the values of Embarked with the most common ones and then mapping them_
     → to their numeric values
     print(train['Embarked'].describe())
     train['Embarked'] = train['Embarked'].fillna('S')
     test['Embarked'] = test['Embarked'].fillna('S')
     ports = {"S": 0, "C": 1, "Q": 2}
     train['Embarked'] = train['Embarked'].map(ports)
     test['Embarked'] = test['Embarked'].map(ports)
    count
              889
    unique
                3
                S
    top
              644
    freq
    Name: Embarked, dtype: object
[0]: #filling the missing values in age
     mean = train["Age"].mean()
     std = test["Age"].std()
     is_null = train["Age"].isnull().sum()
     print(mean,std,is_null)
     # compute random numbers between the mean, std and is_null
     rand_age = np.random.randint(mean - std, mean + std, size = is_null)
     print(rand_age)
     # fill NaN values in Age column with random values generated
     new_age = train["Age"].copy()
     new_age[np.isnan(new_age)] = rand_age
     train["Age"] = new_age
     train["Age"] = train["Age"].astype(int)
```

```
mean = train["Age"].mean()
std = test["Age"].std()
is_null = test["Age"].isnull().sum()
print(mean,std,is_null)
# compute random numbers between the mean, std and is_null
rand_age = np.random.randint(mean - std, mean + std, size = is_null)
print(rand_age)
# fill NaN values in Age column with random values generated
new_age = test["Age"].copy()
new age[np.isnan(new age)] = rand age
test["Age"] = new_age
test["Age"] = test["Age"].astype(int)
print("Number of NaN's in the training dataset : ",train["Age"].isnull().sum())
print("Number of NaN's in the testing dataset : ",test["Age"].isnull().sum())
# Changing the age to categories of age
train['Age'] = train['Age'].astype(int)
train.loc[ train['Age'] <= 10, 'Age'] = 0</pre>
train.loc[(train['Age'] > 10) & (train['Age'] <= 20), 'Age'] = 1</pre>
train.loc[(train['Age'] > 20) & (train['Age'] <= 30), 'Age'] = 2</pre>
train.loc[(train['Age'] > 30) & (train['Age'] <= 40), 'Age'] = 3
train.loc[(train['Age'] > 40) & (train['Age'] <= 50), 'Age'] = 4</pre>
train.loc[(train['Age'] > 50) & (train['Age'] <= 60), 'Age'] = 5</pre>
train.loc[(train['Age'] > 60) & (train['Age'] <= 70), 'Age'] = 6
train.loc[ train['Age'] > 70, 'Age'] = 7
test['Age'] = test['Age'].astype(int)
test.loc[ test['Age'] <= 10, 'Age'] = 0
test.loc[(test['Age'] > 10) & (test['Age'] <= 20), 'Age'] = 1
test.loc[(test['Age'] > 20) & (test['Age'] <= 30), 'Age'] = 2
test.loc[(test['Age'] > 30) & (test['Age'] <= 40), 'Age'] = 3
test.loc[(test['Age'] > 40) & (test['Age'] <= 50), 'Age'] = 4
test.loc[(test['Age'] > 50) & (test['Age'] <= 60), 'Age'] = 5
test.loc[(test['Age'] > 60) & (test['Age'] <= 70), 'Age'] = 6
test.loc[ test['Age'] > 70, 'Age'] = 7
# let's see how it's distributed
print('Values distribution in training set : ',train['Age'].value_counts())
print('Values distribution in testing set : ',test['Age'].value_counts())
29.69911764705882 14.18120923562442 177
[16 27 20 24 20 35 23 22 37 17 31 34 19 40 41 28 29 40 33 21 24 28 42 35
30 38 26 39 36 39 22 26 21 22 32 41 16 29 25 41 42 22 31 24 22 19 40 40
40 42 27 37 27 42 29 21 31 41 33 28 27 19 28 19 38 23 15 36 34 37 33 26
 24 15 41 41 36 40 20 42 38 17 33 19 29 23 21 22 33 42 37 32 36 35 28 35
```

```
37 36 36 30 22 17 16 18 37 23 15 19 27 40 39 37 30 23 40 34 22 36 26 42
     27 32 29 20 31 35 18 28 38 19 30 24 19 22 26 39 37 20 21 41 21 26 22 17
     32 26 25 34 17 34 40 23 42]
    29.60830527497194 14.18120923562442 86
    [25 38 34 37 18 28 22 29 42 19 15 24 38 20 35 30 16 24 17 26 28 25 39 31
     22 37 20 42 21 15 27 25 32 21 36 25 26 42 18 16 40 21 27 33 28 15 42 27
     30 28 23 35 29 33 28 40 39 42 39 40 20 26 21 41 27 23 24 40 33 38 36 37
     22 39 28 42 35 16 18 27 22 15 29 30 16 18]
    Number of NaN's in the training dataset: 0
    Number of NaN's in the testing dataset : 0
    Values distribution in training set : 2
                                                 299
    3
         217
         146
    1
    4
         101
    0
          64
    5
          42
    6
          18
    7
           4
    Name: Age, dtype: int64
    Values distribution in testing set : 2
                                                168
    3
          81
    1
          64
    4
          52
    0
          22
    5
          21
    6
           9
    7
           1
    Name: Age, dtype: int64
[0]: train['Fare'] = train['Fare'].fillna(0)
     test['Fare'] = test['Fare'].fillna(0)
     train['Fare'] = train['Fare'].astype(int)
     test['Fare'] = test['Fare'].astype(int)
     print("Number of NaN's in the training dataset : ",train["Fare"].isnull().sum())
     print("Number of NaN's in the testing dataset : ",test["Fare"].isnull().sum())
     train.loc[ train['Fare'] <= 7.91, 'Fare'] = 0</pre>
     train.loc[(train['Fare'] > 7.91) & (train['Fare'] <= 14.454), 'Fare'] = 1
     train.loc[(train['Fare'] > 14.454) & (train['Fare'] <= 31), 'Fare']</pre>
     train.loc[(train['Fare'] > 31) & (train['Fare'] <= 99), 'Fare']</pre>
     train.loc[(train['Fare'] > 99) & (train['Fare'] <= 250), 'Fare']</pre>
     train.loc[ train['Fare'] > 250, 'Fare'] = 5
     train['Fare'] = train['Fare'].astype(int)
     test.loc[ test['Fare'] <= 7.91, 'Fare'] = 0
     test.loc[(test['Fare'] > 7.91) & (test['Fare'] <= 14.454), 'Fare'] = 1
```

24 28 21 35 28 28 32 28 41 17 37 15 42 26 39 40 34 16 28 19 40 22 28 24

```
test.loc[(test['Fare'] > 14.454) & (test['Fare'] <= 31), 'Fare'] = 2
test.loc[(test['Fare'] > 31) & (test['Fare'] <= 99), 'Fare'] = 3
test.loc[(test['Fare'] > 99) & (test['Fare'] <= 250), 'Fare'] = 4
test.loc[ test['Fare'] > 250, 'Fare'] = 5
test['Fare'] = test['Fare'].astype(int)
```

Number of NaN's in the training dataset : 0 Number of NaN's in the testing dataset : 0

```
[0]: #extracting titles out of the names and ranking them
    titles = {"Mr": 1, "Miss": 2, "Mrs": 3, "Master": 4, "Rare": 5}
    # extract titles
    train['Title'] = train.Name.str.extract(' ([A-Za-z]+)\.', expand=False)
    print(train.Title.unique())
    # replace titles with a more common title or as Rare

→ 'Col', 'Don', 'Dr', 'Major', 'Rev', 'Sir', 'Jonkheer', 'Dona'], 'Rare')
    train['Title'] = train['Title'].replace('Mlle', 'Miss')
    train['Title'] = train['Title'].replace('Ms', 'Miss')
    train['Title'] = train['Title'].replace('Mme', 'Mrs')
    # convert titles into numbers
    train['Title'] = train['Title'].map(titles)
    # filling NaN with O, to get safe
    train['Title'] = train['Title'].fillna(0)
    # extract titles
    test['Title'] = test.Name.str.extract(' ([A-Za-z]+)\.', expand=False)
    print(test.Title.unique())
    # replace titles with a more common title or as Rare
    test['Title'] = test['Title'].replace(['Lady', 'Countess', 'Capt', 'Col', 'Don', __
     →'Dr','Major', 'Rev', 'Sir', 'Jonkheer', 'Dona'], 'Rare')
    test['Title'] = test['Title'].replace('Mlle', 'Miss')
    test['Title'] = test['Title'].replace('Ms', 'Miss')
    test['Title'] = test['Title'].replace('Mme', 'Mrs')
    # convert titles into numbers
    test['Title'] = test['Title'].map(titles)
    # filling NaN with O, to get safe
    test['Title'] = test['Title'].fillna(0)
    train = train.drop(columns='Name')
    test = test.drop(columns='Name')
```

```
['Mr' 'Mrs' 'Miss' 'Master' 'Don' 'Rev' 'Dr' 'Mme' 'Ms' 'Major' 'Lady'
'Sir' 'Mlle' 'Col' 'Capt' 'Countess' 'Jonkheer']
['Mr' 'Mrs' 'Miss' 'Master' 'Ms' 'Col' 'Rev' 'Dr' 'Dona']
```

```
[0]: #changing gender field
     genders = {"male": 0, "female": 1}
     train['Sex'] = train['Sex'].map(genders)
     test['Sex'] = test['Sex'].map(genders)
[0]: #ticket id is unique and isnt useful while predicting thus we drop it
     train = train.drop(columns='Ticket')
     test = test.drop(columns = 'Ticket')
[0]: print('Train sets')
     print(train.columns.values)
     print(test.columns.values)
     Y_TRAIN = train[['Survived']]
     #Y_TRAIN.reset_index(drop = True)
     X TRAIN = train
     X_TRAIN = X_TRAIN.drop(['Survived'],axis =1)
     print(X_TRAIN.columns.values)
     print(Y_TRAIN.columns.values)
     print('Test sets')
     print(test.columns.values)
     Y_Test = train[['Survived']]
     X Test = train
     X_Test = X_Test.drop(['Survived'],axis =1)
     print(X Test.columns.values)
     print(Y_Test.columns.values)
     train.head()
    Train sets
    ['Survived' 'Pclass' 'Sex' 'Age' 'SibSp' 'Parch' 'Fare' 'Embarked' 'Deck'
     'Title'
    ['Pclass' 'Sex' 'Age' 'SibSp' 'Parch' 'Fare' 'Embarked' 'Survived' 'Deck'
     'Title']
    ['Pclass' 'Sex' 'Age' 'SibSp' 'Parch' 'Fare' 'Embarked' 'Deck' 'Title']
    ['Survived']
    Test sets
    ['Pclass' 'Sex' 'Age' 'SibSp' 'Parch' 'Fare' 'Embarked' 'Survived' 'Deck'
     'Title']
    ['Pclass' 'Sex' 'Age' 'SibSp' 'Parch' 'Fare' 'Embarked' 'Deck' 'Title']
    ['Survived']
[0]:
                  Survived Pclass Sex Age ... Fare Embarked Deck Title
    PassengerId
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```

```
4 1 1 1 3 ... 3 0 3 3
5 0 3 0 3 ... 1 0 8 1
```

[5 rows x 10 columns]

```
[0]: from sklearn.model_selection import cross_val_score
     from sklearn.ensemble import RandomForestClassifier
     from sklearn.ensemble import ExtraTreesClassifier
     from sklearn.tree import DecisionTreeClassifier
     from sklearn.metrics import accuracy_score
     from sklearn.ensemble import BaggingClassifier
     #training the algorithms before dropping the least important class
     clf = RandomForestClassifier(n estimators=1000, max depth=None,
     →min_samples_split=2)
     clf = clf.fit(X_TRAIN, Y_TRAIN.values.ravel())
     print('before')
     accuracy_clf_before = accuracy_score(Y_Test,clf.predict(X_Test))
     print('accuracy random forest before : ',accuracy_clf_before)
     clf2 = ExtraTreesClassifier(n_estimators=1000, max_depth=None, random_state=0)
     clf2.fit(X_TRAIN,Y_TRAIN.values.ravel())
     accuracy_extra_tree = accuracy_score(Y_Test,clf2.predict(X_Test))
     print('accuracy extra tree before : ',accuracy_extra_tree)
     #plotting a graph to visualize each feature's importance
     feature_imp = pd.DataFrame({'feature':X_TRAIN.columns,'importance':np.round(clf.

→feature_importances_,3)})
     feature_imp = feature_imp.sort_values('importance',ascending=False).
     ⇔set_index('feature')
     feature imp.plot.bar()
     #dropping the least important class
     X_TRAIN = X_TRAIN.drop(columns='Parch')
     X_Test = X_Test.drop(columns='Parch')
     #retraining the algorithms to compare
     clf = clf.fit(X_TRAIN, Y_TRAIN.values.ravel())
     print('after removing parch')
     accuracy_clf_after = accuracy_score(Y_Test,clf.predict(X_Test))
     print('accuracy random forest after : ',accuracy clf after)
     scores_random_forest = cross_val_score(clf, X_TRAIN, Y_TRAIN.values.ravel(),_
     \rightarrowcv=10)
     print(' random forest cross val score: ',scores_random_forest.mean())
```

before

accuracy random forest before : 0.920314253647587 accuracy extra tree before : 0.920314253647587

after removing parch

accuracy random forest after: 0.9180695847362514 random forest cross val score: 0.8485018726591761 extra tree cross val score: 0.8439950062421973 accuracy extra tree after: 0.9180695847362514

[0]:

