Feature Selection By Using The Chi2 Test. • High diamensional data in input usualy not good for classification due to curse of the diamensionality .It significantly increases the time and space complexity for processing the data. Lets suppose you having huge dimension of your data then definetly it will take more time and more space to test and train a model moreover in presence of irrelevant and redudant features learning method tends to overfit and becomes less interpretable. A common way to resolve this problem is a feature selection which reduces the diamensionality by selection the subset of feature from the input feature set and it is often used to reduce the computational cost and remove irrrevant and redudant features for the problem with the high diamensional data. One of the method in filter method is a Fisher Score which can be calculated by doing the chi2(chi sqaure). Fisher score use on numerical and chi2 method is apply only on the categorical data. That categorical data that has to be a finite set of the data so the basical it computes the distributions and its frequency. Distribution and frequncy is basically define accordingly with the help of mode and median. So the Fisher Score is one of the most widely used supervised feature selection method however it select each features independently accordingly their score and the fisher criteria which can lead suboptimal subset of the features. In univaraint selection that the feature are been selected individualy by considering the effect of other features we may endup by selecting a suboptimal feature set. What is the Fisher Score and Chi2(x2) Test. • Chi2(x2) Test: - It is a stastical hypothesis test where the sampling distribution of the stastic of the chi2 distribution and chi2 test is used to determine whether there is significant diffrence between expected frequency and observed frequency in one or more categories thats means one or more features. It is applied only and only categorical datasets. Chi2 test is measure the diffrences between the stochastic variable, so using this function "Weeds out" the features that are most likely to be independed of the class and therefore irrelevant for classifiaction. That we are removing those feature which are the irrevant for the classification or not depended to the target output. Understand With an Example. import pandas as pd import numpy as np import matplotlib.pyplot as plt import seaborn as sns import warnings warnings.filterwarnings('ignore') from sklearn.metrics import accuracy score, classification report, confusion matrix from sklearn.model selection import train test split from sklearn.feature selection import SelectFromModel,chi2,SelectKBest,SelectPercentile from sklearn.ensemble import RandomForestClassifier from sklearn import datasets data = sns.load\_dataset('titanic') In [4]: data.head() embarked class who adult\_male survived pclass age sibsp parch fare deck embark\_town alive alone sex 0 0 22.0 0 7.2500 S Third 3 male True NaN Southampton False man no 1 38.0 0 71.2833 First woman **False** female C Cherbourg False yes 2 7.9250 1 26.0 Third woman False NaN True female S Southampton yes 3 female 35.0 53.1000 First woman **False** Southampton False 8.0500 4 male 35.0 S Third Southampton True NaN True man no from sklearn.preprocessing import LabelEncoder data.info() <class 'pandas.core.frame.DataFrame'> RangeIndex: 891 entries, 0 to 890 Data columns (total 15 columns): # Column Non-Null Count Dtype 0 survived 891 non-null int64 1 pclass 891 non-null int64 3 age 714 non-null float64
4 sibsp 891 non-null int64
5 parch 891 non-null int64
6 fare 891 non-null float64
7 embarked 889 non-null object
8 class 891 non-null category
9 who 891 non-null object 2 891 non-null object sex 10 adult\_male 891 non-null bool
11 deck 203 non-null category
12 cmbark town 889 non-null object 12 embark town 889 non-null object 13 alive 891 non-null object 14 alone 891 non-null bool object dtypes: bool(2), category(2), float64(2), int64(4), object(5) memory usage: 80.6+ KB data.describe() survived pclass age sibsp parch fare **count** 891.000000 891.000000 714.000000 891.000000 891.000000 891.000000 0.383838 2.308642 29.699118 0.523008 0.381594 32.204208 std 0.486592 0.836071 14.526497 1.102743 0.806057 49.693429 min 0.000000 1.000000 0.420000 0.000000 0.000000 0.000000 25% 0.000000 2.000000 20.125000 0.000000 0.000000 7.910400 0.000000 0.000000 **50**% 3.000000 28.000000 0.000000 14.454200 **75**% 1.000000 3.000000 38.000000 1.000000 0.000000 31.000000 1.000000 3.000000 80.000000 8.000000 6.000000 512.329200 data.isnull().sum() Out[9]: survived pclass 0 sex 0 177 age sibsp parch 0 fare 2 embarked class 0 who 0 0 adult male deck 688 embark town 2 alive alone dtype: int64 In [10]: data['age'].mean() Out[10]: 29.69911764705882 data['age'].median() Out[11]: 28.0 data['age'].fillna(data['age'].median(),inplace=True) data['deck'].value\_counts() Out[13]: C 59 D 33 Ε 32 15 Α F 13 Name: deck, dtype: int64 data['deck'].isnull().sum()/len(data.index)\*100 In [14]: 77.21661054994388 Out[14]: del data['deck'] data.head() fare embarked class survived pclass age sibsp parch who adult\_male embark\_town alive alone 0 7.2500 male 22.0 S Third Southampton False 1 female 38.0 71.2833 Cherbourg First woman yes False 2 7.9250 S Third female 26.0 Southampton yes True 53.1000 3 female 35.0 Southampton False 4 0 male 35.0 8.0500 S Third Southampton True data['embarked'].value counts() S 644 168 77 Name: embarked, dtype: int64 data['embarked'].fillna('S',inplace=True) lable = LabelEncoder() data['sex']=lable.fit transform(data['sex'])  $d = \{ 'S':0, 'C':1, 'Q':2 \}$ data['embarked']=data['embarked'].map(d) xa = {'First':0,'Second':1,'Third':2} data['class']=data['class'].map(xa) lable = LabelEncoder() data['who']=lable.fit\_transform(data['who']) lable = LabelEncoder() data['who']=lable.fit\_transform(data['who']) In [24]: lable = LabelEncoder() data['adult\_male']=lable.fit\_transform(data['adult\_male']) lable = LabelEncoder() data['alive']=lable.fit\_transform(data['alive']) lable = LabelEncoder() data['alone']=lable.fit\_transform(data['alone']) data.head() survived pclass sex age sibsp parch fare embarked class who adult\_male embark\_town alive alone 1 22.0 7.2500 Southampton 38.0 Cherbourg 2 7.9250 Southampton 1 3 35.0 53.1000 Southampton 4 0 35.0 0 8.0500 2 Southampton 0 1 data['embark town'].value counts() 644 Southampton Cherbourg Queenstown 77 Name: embark\_town, dtype: int64 data['embark town'].isnull().sum() Out[29]: 2 data['embark\_town'].fillna('Southampton',inplace=True) data['embark town']=lable.fit transform(data['embark town']) data.head() embarked who adult\_male embark\_town survived pclass sex age sibsp parch class 0 0 1 22.0 0 7.2500 0 2 1 1 2 0 0 0 38.0 71.2833 2 2 2 0 1 3 0 26.0 7.9250 0 2 2 1 1 3 53.1000 35.0 0 4 1 1 35.0 0 8.0500 2 1 0 x = data.drop(columns=['alive']) y = data['alive'] In [34]: x.shape, y.shape ((891, 13), (891,)) x\_train,x\_test,y\_train,y\_test = train\_test\_split(x,y,train\_size=0.80,stratify=y,random\_state=42) x\_train.shape,y\_train.shape ((712, 13), (712,))Chi2\_test f\_score,pvalue = chi2(x\_train,y\_train) #it will return us the f\_score and pvalues In [40]: f\_score Out[40]: array([4.39000000e+02, 2.59119322e+01, 7.63019940e+01, 2.08599242e+01, 1.10666572e+00, 9.06416124e+00, 3.91708928e+03, 1.58304323e+01, 4.57073139e+01, 2.24512976e+01, 8.93946763e+01, 1.06126527e+01, 1.29187387e+01]) pvalue In [41]: Out[41]: array([1.78683855e-97, 3.57353388e-07, 2.43437961e-18, 4.94125395e-06, 2.92807896e-01, 2.60668078e-03, 0.00000000e+00, 6.92792453e-05, 1.37309133e-11, 2.15539829e-06, 3.23405685e-21, 1.12316440e-03, 3.25308982e-04]) In [42]: Pvalues = pd.Series(pvalue,index=x\_train.columns) Pvalues.sort values(ascending=True,inplace=True) In [43]: Pvalues In [44]: Out[44]: fare 0.000000e+00 survived 1.786839e-97 adult\_male 3.234057e-21 2.434380e-18 sex class 1.373091e-11 pclass 3.573534e-07 who 2.155398e-06 4.941254e-06 age embarked 6.927925e-05 alone 3.253090e-04 1.123164e-03 embark\_town 2.606681e-03 parch 2.928079e-01 sibsp dtype: float64

In [45]: plt.figure(figsize=(20,9))

plt.show()

0.30

0.25

0.20

0.15

0.10

0.05

0.00

prediction.

Out[56]: ((712, 10), (179, 10))

Accuracy :- 1.0

Thanks !!!

x\_train\_6.shape,x\_test\_6.shape

clf.fit(x\_train,y\_train)
y pred = clf.predict(x\_test)

def RandomForest(x\_train,x\_test,y\_train,y\_test):

RandomForest(x train 6,x test 6,y train,y test)

clf = RandomForestClassifier(n\_estimators=100,)

print('Accuracy :-',accuracy\_score(y\_test,y\_pred))

plt.bar(Pvalues.index, Pvalues)

embark town

If We see the Pvalues according Feature. Those which having the low pvalus which are highly important for us to get the maximum

x\_train\_6 = x\_train[['fare','survived','adult\_male','sex','class','pclass','age','alone','who','embark\_town']]
x\_test\_6 = x\_test[['fare','survived','adult\_male','sex','class','pclass','age','alone','who','embark\_town']]

plt.xticks(rotation=90)