Cab Fare Prediction



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Introduction

City taxi rides paint a vibrant picture of life in the city. The millions of rides taken each month can provide insight into traffic patterns, road blockage, or large-scale events that attract many People. With ridesharing apps gaining popularity, it is increasingly important for taxi companies to provide visibility to their estimated fare and ride duration, since the competing apps provide these metrics upfront. Predicting fare and duration of a ride can help passengers decide when is the optimal time to start their commute, or help drivers decide which of two potential rides will be more profitable, for example. Furthermore, this visibility into fare will attract customers during times when ridesharing services are implementing surge pricing.

In order to predict the fare, only data which would be available at the beginning of a ride was used. This includes pickup and drop off coordinates, the date, time and number of passengers. Linear regression, decision tree and random forest models were used to predict the fare amount.

PYTHON

import numpy as np import pandas as pd import seaborn as sns import matplotlib.pyplot as plt from sklearn.decomposition import PCA from sklearn.linear_model import LinearRegression from sklearn.metrics import mean squared error from math import sqrt from sklearn import tree from sklearn.model_selection import train test split sklearn.ensemble import from RandomForestRegressor import os os.chdir(r"E:\MY\Project\Cab Fare Prediction") Train = pd.read csv("train.csv") Test = pd.read_csv("test.csv")

```
R
rm(list=ls())
setwd("E:/MY/Project/Cab Fare
Prediction")
x = c("ggplot2", "corrgram", "DMwR",
"caret", "randomForest", "unbalanced",
"C50", "dummies", "e1071", "Information",
"MASS", "rpart", "gbm", "ROSE", 'sampling',
'DataCombine', 'inTrees', "usdm", "party")
install.packages(x)
lapply(x, require, character.only = TRUE)
rm(x)
Train = read.csv("train.csv", header = T,
na.strings = c(" ", "", "NA"))
Test= read.csv("test.csv", header = T,
na.strings = c(" ", "", "NA"))
```

Exploratory Data Analysis

The train set consists of 16067 obser vations and 7 features containing the target. The missing percentage of "p assenger_count" is 34.23% while that of "fare_amoun"t is 15.55%. The test data does not contain any missing values. There are only numeric features. No features seem to have substantial outliers. The data set is historical dataset containing numeric feature variables. Where, we tried to design a system that predicts the fare amount for a cab ride in the city.



2.1 Data Pre Processing

1. Missing values Imputation

Train["fare_amount"] = pd.to_numeric (Train["fare_amount"],errors='coerce') def missin_val(df): missin_val = pd.DataFrame(df.isnull().sum()) missin_val = missin_val.reset_index() missin_val = missin_val.rename(columns = {'index': 'Variables', 0: 'Missing_percentage'}) missin_val['Missing_percentage'] = (missin_val['Missing_percentage'] /len(df))*100 missin_val = missin_val.sort_values ('Missing_percentage', ascending = False).reset_index(drop = True) return(missin_val) print("The missing value percentage in training data : \n\n",missin_val(Train)) print("The missing value percentage in test data : \n\n",missin_val(Test)) #Impute the missing values Train["passenger_count"] = Train["passenger_count"].fillna(Train["passenger_count"].median()) Train["fare_amount"] = Train["fare_amount"].fillna(Train["fare_amount"].median()) print("Is there still any missing value in the training data:\n\n",Train.isnull().sum())

```
#covert factor to numeric
Train$fare_amount = as.numeric(Train$fare_amount)

#Missing Value Analysis
missing_val = data.frame(apply(Train,2,function(x){sum(is.na(x))}))
missing_val$Columns = row.names(missing_val)
names(missing_val)[1] = "Missing_percentage"
missing_val$Missing_percentage = (missing_val$Missing_percentage/nrow(Train)) * 100
missing_val = missing_val[order(-missing_val$Missing_percentage),]
row.names(missing_val) = NULL
missing_val = missing_val[,c(2,1)]
View(missing_val)

#Median Method
Train$fare_amount[is.na(Train$fare_amount)] = median(Train$fare_amount, na.rm = T)
Train$passenger count[is.na(Train$passenger count)] = median(Train$passenger count, na.rm = T)
```

2. Data Alignment:

```
Python
#Split our Datetime into individual columns for ease of data processing and modelling
def align datetime(df):
  df["pickup_datetime"] = df["pickup_datetime"].map(lambda x: str(x)[:-3])
  df["pickup datetime"] = pd.to datetime(df["pickup datetime"], format='%Y-%m-%d %H:%M:%S')
  df['year'] = df.pickup datetime.dt.year
  df['month'] = df.pickup_datetime.dt.month
  df['day'] = df.pickup_datetime.dt.day
  df['weekday'] = df.pickup_datetime.dt.weekday
  df['hour'] = df.pickup datetime.dt.hour
  return(df["pickup_datetime"].head())
align_datetime(Train)
align datetime(Test)
#Remove the datetime column
Train.drop('pickup_datetime', axis=1, inplace=True)
Test.drop('pickup_datetime', axis=1, inplace=True)
```

```
R
#covert Train datetime
Train$pickup datetime1 = as.Date(Train$pickup datetime)
Train$year = as.numeric(format(Train$pickup_datetime1, format = "%Y"))
Train$month = as.numeric(format(Train$pickup_datetime1, format = "%m"))
Train$day = as.numeric(format(Train$pickup_datetime1, format = "%d"))
Train$pickup_time <- sapply(strsplit(as.character(Train$pickup_datetime), " "), "[", 2)
Train$Hour <- sapply(strsplit(as.character(Train$pickup_time), ":"), "[", 1)</pre>
Train$pickup_datetime = NULL
Train$pickup_datetime1 = NULL
Train$pickup_time =NULL
View(Train)
#covert Test datetime
Test$pickup datetime1 = as.Date(Test$pickup datetime)
Test$year = as.numeric(format(Test$pickup_datetime1, format = "%Y"))
Test$month = as.numeric(format(Test$pickup_datetime1, format = "%m"))
Test$day = as.numeric(format(Test$pickup_datetime1, format = "%d"))
Test$pickup_time <- sapply(strsplit(as.character(Test$pickup_datetime), " "), "[", 2)
Test$Hour <- sapply(strsplit(as.character(Test$pickup_time), ":"), "[", 1)</pre>
Test$pickup_datetime = NULL
Test$pickup_datetime1 = NULL
Test$pickup_time =NULL
```

3. Convert data in their proper data type.

```
#Setting proper data type for each columns
Train=
Train.astype({"fare_amount":float,"pickup_longitude":float,"pickup_latitude":float,"dropoff_longitude":float,"dropoff_latitude":float,"passenger_count":int,"year":int,"month":int ,"day"
:int,"weekday":int,"hour":int})
Train.dtypes
```

<u>R</u>

Train\$Hour = as.numeric(Train\$Hour)
Test\$Hour = as.numeric(Test\$Hour)

4. Individual Distribution:

```
import gc
gc.collect();

print('Distributions columns')
plt.figure(figsize=(30, 185))
for i, col in enumerate(numerical_features):
    plt.subplot(50, 4, i + 1)
    plt.hist(Train[col])
    plt.title(col)
gc.collect();
```

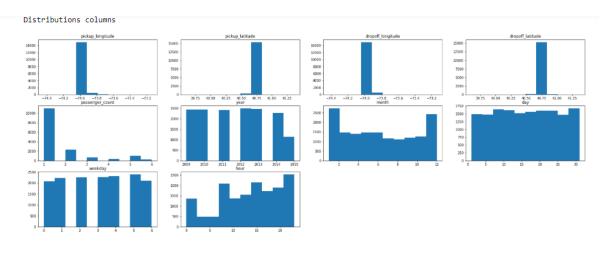
R

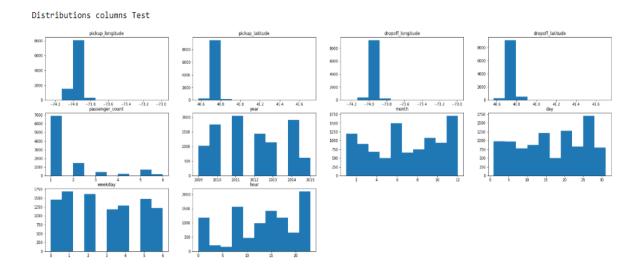
#Distribution of train attributes

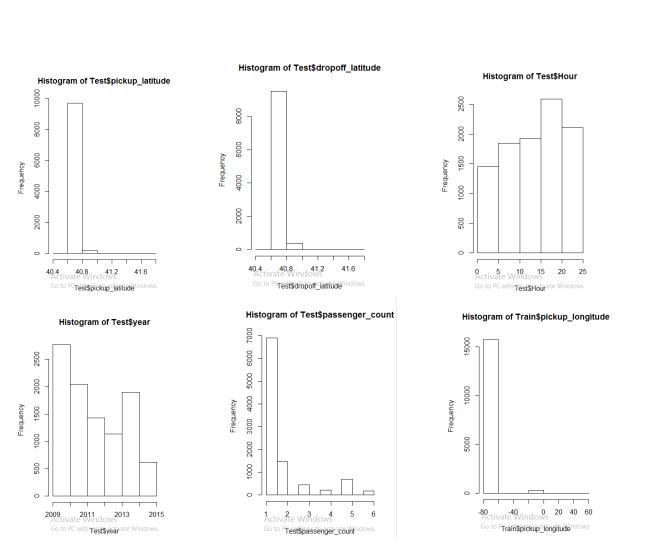
hist(Train\$pickup_longitude, breaks=7) hist(Train\$dropoff_longitude, breaks=7) hist(Train\$pickup_latitude, breaks=7) hist(Train\$dropoff_latitude, breaks=7) hist(Train\$Hour, breaks=7) hist(Train\$year, breaks=7) hist(Train\$passenger_count)

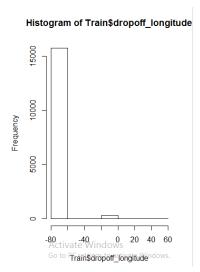
hist(Test\$pickup_longitude, breaks=7)
hist(Test\$dropoff_longitude, breaks=7)
hist(Test\$pickup_latitude, breaks=7)
hist(Test\$dropoff_latitude, breaks=7)
hist(Test\$Hour, breaks=7)

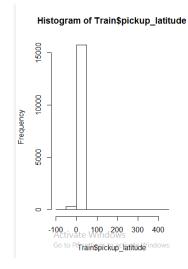
hist(Test\$year, breaks=7) hist(Test\$passenger_count)

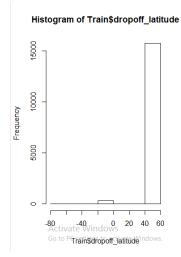


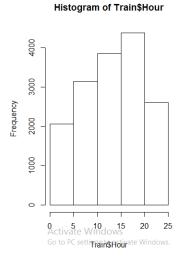


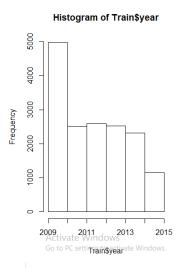


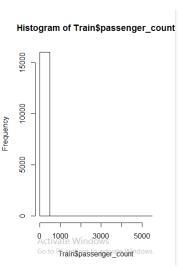


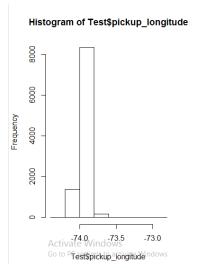


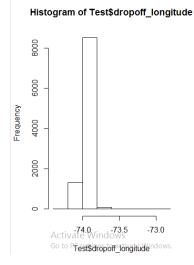






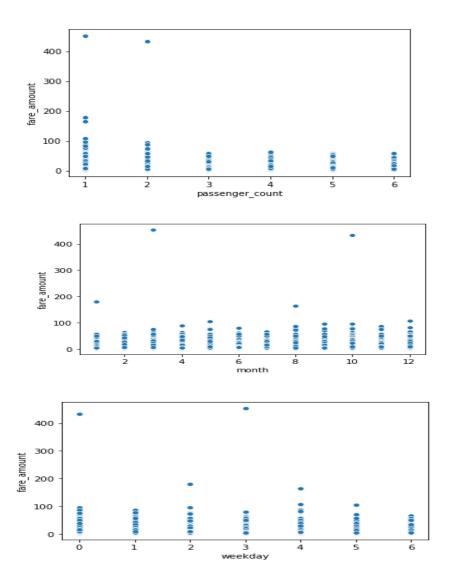






5. Scatter Plot analysis.

```
#Scatter plots
cmap = sns.cubehelix_palette(dark=.3, light=.8, as_cmap=True)
sns.scatterplot(x="passenger_count", y="fare_amount", data= Train, palette="Set2")
sns.scatterplot(x="month", y="fare_amount", data= Train, palette="Set2")
sns.scatterplot(x="weekday", y="fare_amount", data= Train, palette="Set2")
sns.scatterplot(x="hour", y="fare_amount", data= Train, palette="Set2")
```

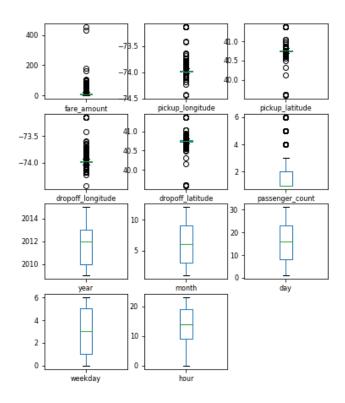


6. Outlier Analysis:

```
Python
#Outlier analysis
Train.plot(kind='box', subplots=True, layout=(8,3), sharex=False, sharey=False, fontsize=8)
plt.subplots_adjust(left=0.125, bottom=0.1, right=0.9, top= 3,wspace=0.2, hspace=0.2)
plt.show()
def outliers_analysis(df):
  for i in df.columns:
    print(i)
    q75, q25 = np.percentile(df.loc[:,i], [75,25])
    iqr = q75 - q25
    min = q25 - (iqr*1.5)
    max = q75 + (iqr*1.5)
    print(min)
    print(max)
    df = df.drop(df[df.loc[:,i] < min].index)</pre>
    df = df.drop(df[df.loc[:,i] > max].index)
    return(df)
train = outliers_analysis(Train)
test = outliers_analysis(Test)
```

```
#remove outlier from test and Train
for(i in cnames){
    print(i)
    val = Train[,i][Train[,i] %in% boxplot.stats(Train[,i])$out]
    #print(length(val))
    Train = Train[which(!Train[,i] %in% val),]
}

for(i in cnames1){
    print(i)
    val = Test[,i][Test[,i] %in% boxplot.stats(Test[,i])$out]
    #print(length(val))
    Test = Test[which(!Test[,i] %in% val),]
}
```



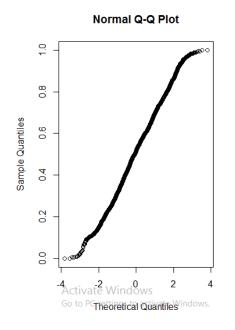
7. Feature scaling with the help of Normalisation:

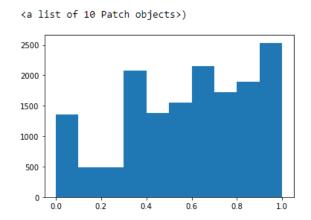
```
##Normalisation
def Normalisation(df):
    for i in df.columns:
        df[i] = (df[i] - df[i].min())/(df[i].max() - df[i].min())

Normalisation(X_train)
Normalisation(Test)
```

```
#for Train
for(i in cnames){
    print(i)
    Train[,i] = (Train[,i] - min(Train[,i]))/(max(Train[,i] - min(Train[,i])))
}

#for Test
for(i in cnames1){
    print(i)
    Test[,i] = (Test[,i] - min(Test[,i]))/(max(Test[,i] - min(Test[,i])))
}
```



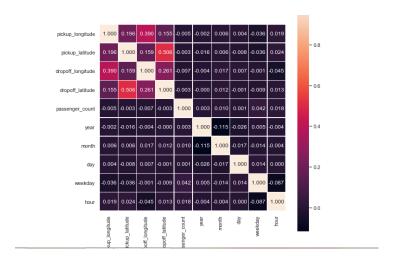


8. Feature selection using correlation analysis.

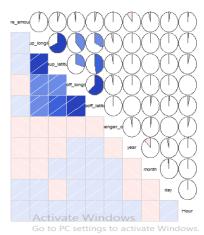
```
corrgram(Train[,numeric_index], order = F,
    upper.panel=panel.pie, text.panel=panel.txt, main = "Correlation Plot Train")

corrgram(Test[,numeric_index1], order = F,
    upper.panel=panel.pie, text.panel=panel.txt, main = "Correlation Plot Test")

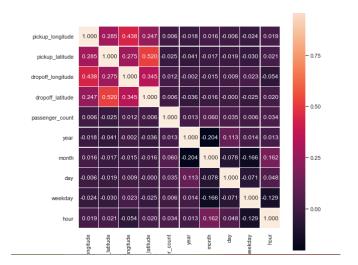
## Dimension Reduction
Train = subset(Train, select = -c(Hour,day,month, year))
View(Train)
Test = subset(Test, select = -c(Hour,day,month, year))
View(Test)
```

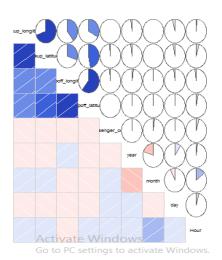


Correlation Plot



Correlation Plot Test





9. Dimensionality Reduction using PCA:

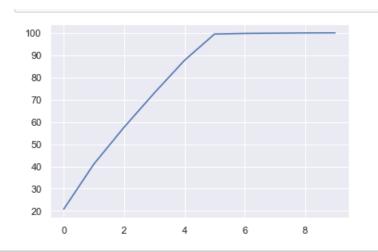
Python

pca = PCA(n_components=10)
pca.fit(X_train)
var= pca.explained_variance_ratio_
var1=np.cumsum(np.round(pca.explained_variance_ratio_, decimals=4)*100)
plt.plot(var1)
plt.show()

#lets reduce our no. of variables to 5 as it explains 100% features of our Data pca = PCA(n_components=5)

X = pca.fit(X_train).transform(X_train)

Test = pca.fit(Test).transform(Test)



Solution Methods:



We use 3 different methods to come to a solution: Logistic Regression, Decision Tree and Random Forest from Scikit Learn.

1 Logistic Regression:

Logistic regression is a relatively 'simple' machine learning algorithm and we expect fast, but not great results. It tries to attach the best constant values to how features interact with the target, based on the train set, minimizing an error term. It then applies this same formula to the test data set. It is pursued here as a baseline in order to compare to more sophisticated models. For more details see (Bishop, 2006).

```
####### Linear Regression #########

import statsmodels.api as sm

# Train the model using the training sets
model = sm.OLS(y_train, X_train).fit()
model.summary()

predictions_LR = model.predict(X_test)

def rmse(y, y_pred):
    return np.sqrt(np.mean(np.square(y - y_pred)))

rmse(y_test, predictions_LR)
```

2 Decision Tree:

This section roughly describes what a decision tree is. The concept is fairly simple. If a prediction needs to be made, go from the top of the tree to the bottom. This tree is constructed by at each depth level greedily finding the best feature to split on, maximizing information gain via the Gini impurity. Let p0 be the proportion of observations that belong to class 0 and let p1 be the proportion of observations that belong to class 1 out of all observations.

$$GINI = 1 - (P_0^2 + P_1^2)$$

This metric is an example of how to measure how pure a split is. It is more pure the lower it is with minimum 0 and maximum 0.5. A split is purer if in the branches the ratio of positive to negative examples is close to 0 or 1 or in other words the split is highly discriminatory. The maximum depth of this tree is set at 2 to keep it tractable and this effectively also keeps it from over fitting. A decision tree can otherwise perfectly match a training set, which does not generalize well.

DecisionTree Modelling ########## from sklearn.tree import DecisionTreeRegressor fit_DT = DecisionTreeRegressor(max_depth=2).fit(X_train, y_train) predictions_DT = fit_DT.predict(X_test) rmse(y_test, predictions_DT)

```
########Decion Tree#########

fit = rpart(fare_amount ~ ., data = train, method = "anova")
predictions_DT = predict(fit, test[, -1])

#RMSE

RMSE(test[,1], predictions_DT)
```

3 Random Forest

The Random Forest model generates multiple decision trees. (Breiman, 2001) Only a subset of predictive features is now considered during each split that is randomly selected. The decision trees lead to one single prediction together by averaging all the predictions they give individually. The parameters of a Random Forest model are really important and need to be tuned appropriately. N_ESTIMATORS determines the number of trees in the forest.

MAX_FEATURES controls the maximum random amount of features to consider when determining a best split during the algorithm. MAX_DEPTH limits the depth of a tree in the random forest. MIN_SAMPLES_SPLIT determines the minimum amount of observations that need to be in a node for it to be considered for splits. MIN_SAMPLES_LEAF constrains that leaf nodes have at least this number of observations. N_JOBS controls the number of processors. Trees in a random forest can be made in parallel, so the more cores working, the less computation time needed. CLASS_WEIGHT can be set to 'balanced' to deal with imbalanced datasets.

PYTHON

Random Forest Modelling

```
rf_model = RandomForestRegressor(max_depth= 5, n_estimators = 100).fit(X_train, y_train)
rf_pred_train = rf_model.predict(X_train)
rf_pred= rf_model.predict(X_test)

rmse(y_test, rf_pred)
```

R


```
rm_model <- randomForest(fare_amount ~., data = train)
print(rm_model)
predictions_RF = predict(rm_model, test[, -1])
RMSE(test[,1], predictions_RF)</pre>
```

Result

S.No	ALGORITHM	RMSE
1	Linear Regression	17.26
2	Decision Tree	12.87
3	Random Forest	12.83

Why RMSE?

When we fit linear regression model in time series and if in case we need to use dummy variable for observed but unpredictable spike(this is know as episodic events) then there could be possibility that model is over fit in forecast periods. It means that instead of switching off episodic events, our model also captured the error terms (this is the one of the difficult task to differentiate random error and episodic events). So in forecast, if we want to check the accuracy of the model and avoid over fit and any episodic event in future; we can square root the MSE value and due to square root, model avoids outliers (robust against extreme values) in the forecast period. As we cannot model spike in the forecast period. Also, RMSE more aggressively punishes big errors than small ones, whereas MAD/MAPE are more linear

Conclusion

Considering what is and what is not accounted for in the models built in this study, their predicting results are fairly accurate. To further improve the prediction accuracy, more variability's need to be considered and modelled. Although the rides in hour and average speed in hour work as proxies for traffic, more modelling on the effect of location is needed. These quantities could be calculated for different areas to further model local effects of traffic. Also, modelling traffic and the effect of location in between pickup and drop-off points should be considered as well as difference in drivers' speed. These further steps could be taken both by analysing larger sets of the data to infer relationships and effects of location and traffic at different times, as well as aggregation with other datasets, as data on traffic, speed limitations, etc.