Project Report On **Employee Absenteeism**

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1. Introduction

1.1 Problem Statement:

XYZ is a courier company. As we appreciate that human capital plays an important role in collection, transportation and delivery. The company is passing through genuine issue of Absenteeism. The company has shared it dataset and requested to have an answer on the following areas:

- 1. What changes company should bring to reduce the number of absenteeism?
- 2. How much losses every month can we project in 2011 if same trend of absenteeism continues?

1.2 Data:

Dataset Details:

Dataset Characteristics: Timeseries Multivariant

Number of Attributes: 21 Missing Values : Yes

Attribute Information:

- 1. Individual identification (ID)
- 2. Reason for absence (ICD). Absences attested by the International Code of Diseases (ICD) stratified into 21 categories (I to XXI) as follows:
 - I. Certain infectious and parasitic diseases
 - II. Neoplasms
 - III. Diseases of the blood and blood-forming organs and certain disorders involving the immune mechanism
 - IV. Endocrine, nutritional and metabolic diseases
 - V. Mental and behavioural disorders VI Diseases of the nervous system VII Diseases of the eye and adnexa
 - VI. Diseases of the ear and mastoid process
 - VII. Diseases of the circulatory system
 - VIII. Diseases of the respiratory system
 - IX. Diseases of the digestive system
 - X. Diseases of the skin and subcutaneous tissue
 - XI. Diseases of the musculoskeletal system and connective tissue
 - XII. Diseases of the genitourinary system
 - XIII. Pregnancy, childbirth and the puerperium
 - XIV. Certain conditions originating in the perinatal period
 - XV. Congenital malformations, deformations and chromosomal abnormalities

- XVI. Symptoms, signs and abnormal clinical and laboratory findings, not elsewhere classified
- XVII. Injury, poisoning and certain other consequences of external causes
- XVIII. External causes of morbidity and mortality
- XIX. Factors influencing health status and contact with health services

And 7 categories without (CID) patient follow-up (22), medical consultation (23), blood donation (24), laboratory examination (25), unjustified absence (26), physiotherapy (27), dental consultation (28).

- 3. Month of absence
- 4. Day of the week (Monday (2), Tuesday (3), Wednesday (4), Thursday (5), Friday (6))
- 5. Seasons (summer (1), autumn (2), winter (3), spring (4))
- 6. Transportation expense
- 7. Distance from Residence to Work
- 8. Service time
- 9. Age
- 10. Work load Average/day
- 11. Hit target
- 12. Disciplinary failure (yes=1; no=0)
- 13. Education (high school (1), graduate (2), postgraduate (3), master and doctor (4))
- 14. Son (number of children)
- 15. Social drinker (yes=1; no=0)
- 16. Social smoker (yes=1; no=0)
- 17. Pet (number of pet)
- 18. Weight
- 19. Height
- 20. Body mass index
- 21. Absenteeism time in hours (target)

2. Methodology

2.1 Pre-Processing

Data preprocessing is a data science technique that involves transforming raw data into an understandable format. Real-world data is often incomplete, inconsistent, and/or lacking in certain behaviors or trends, and is likely to contain many errors. Data preprocessing is a proven method of resolving such issues. Data preprocessing prepares raw data for further processing.

Data-gathering methods are often loosely controlled, resulting in out-of-range values, impossible data combinations, missing values, etc. This is often called as exploratory data analysis.

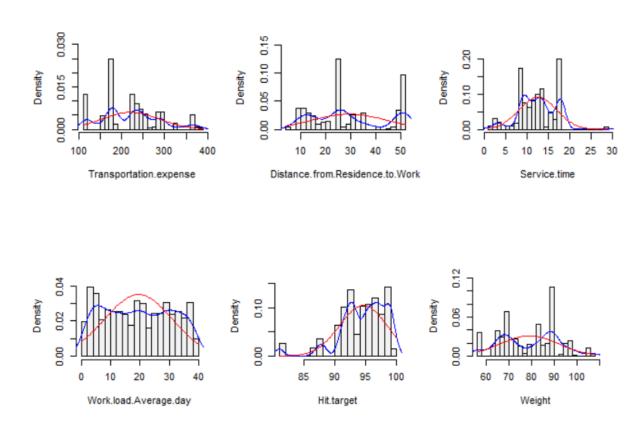
2.2.1 Missing Value Analysis

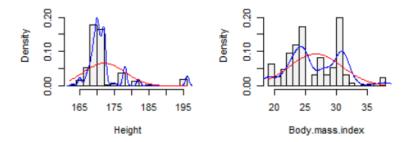
Missing value can arise due to many cases and Proper handling of missing values is important in all statistical analyses. Here Target variable is 'Absenteeism.time.in.hours'. Which itself contains some missing values, in such cases we should neglect the observation. 22 observations from 740 observation has no Target variable. Missing values are imputed using different methods such as Mean, median and KNN imputation. The criterion for imputation is that the variable should have missing values less than 30 percent, In this case no variable has missing values more than 30 percent. To choose the method for imputation we purposefully create a NA and try to impute it using different methods, whichever method gives the closest output, we freeze that method. Method may vary from variable to variable and it mainly depends upon the whether variable is categorical or continuous. Following is the percentage of missing values in each variable

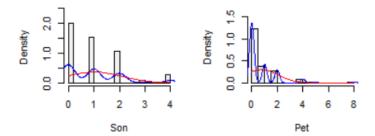
	variable	no of missing value	Missing percentage
1	ID	0	0.0000000
2	Reason.for.absence	3	0.4054054
3	Month.of.absence	1	0.1351351
4	Day.of.the.week	0	0.0000000
5	Seasons	0	0.0000000
6	Transportation.expense	7	0.9459459
7	Distance.from.Residence.to.Work	3	0.4054054
8	Service.time	3	0.4054054
9	Age	3	0.4054054
10	Work.load.Average.day	0	0.0000000
11	Hit.target	6	0.8108108
12	Disciplinary.failure	6	0.8108108
13	Education	10	1.3513514
14	Son	6	0.8108108
15	Social.drinker	3	0.4054054
16	Social.smoker	4	0.5405405
17	Pet	2	0.2702703
18	Weight	1	0.1351351
19	Height	14	1.8918919
20	Body.mass.index	31	4.1891892
21	Absenteeism.time.in.hours	22	2.9729730

2.2.2 Outlier Analysis

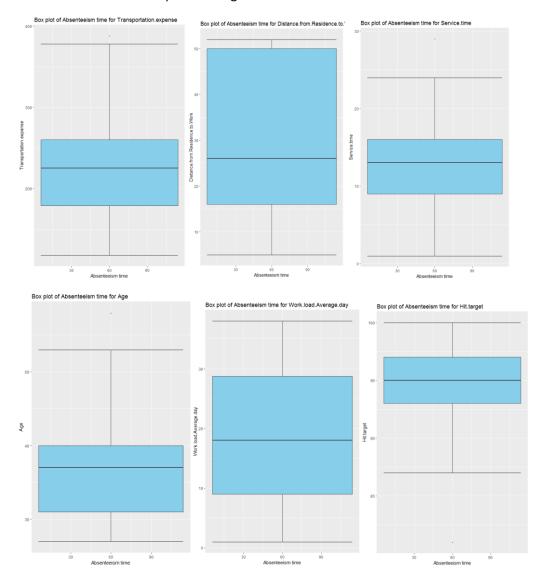
Outliers are extreme values that deviate from other observations on data, they may indicate a variability in a measurement, experimental errors. This data contains outliers in variables such as 'Transportation Expense', 'Service time', 'Hit target' and some personal details. Below are histograms of numerical variables. We can see the distribution is not normal hence we need to perform outlier analysis which will try to impute the outliers with medians or means of the data. Although distribution plots does not help us to find the outliers, we use boxplot method to identify and remove or impute the outliers. We visualize the data using boxplots.



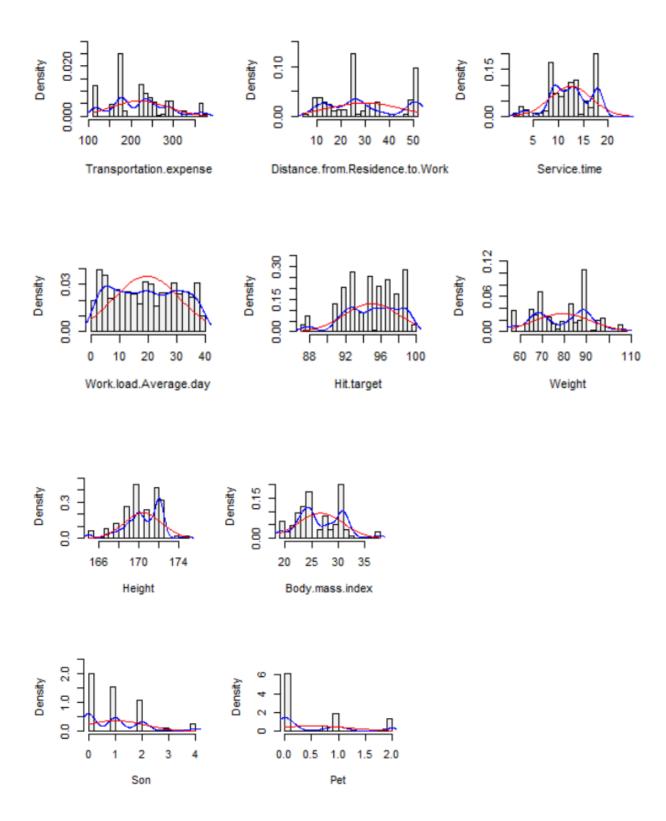




We can clearly see the distribution is not normal hence below are the boxplots for the same. In these boxplots the observations which are above or below 1.5 times the interquartile range are marked as outliers. Interquartile range is shaded as blue while outliers are marked as Red dots.



We used the boxplot method to identify and remove the outliers from the variables and the histogram distribution after removal of outliers is given below



2.2.3 Feature Selection

Before performing any type of modeling we need to assess the importance of each predictor variable in our analysis. There is a possibility that many variables in our analysis are not important at all to the problem of class prediction. There are several methods of doing that. We are here using correlation plot to identify the correlation between numerical variables and we will neglect the variable which are highly correlated to each other so that they does not carry the same information to the model development. In same way we will use Chi-Square test for categorical variable.

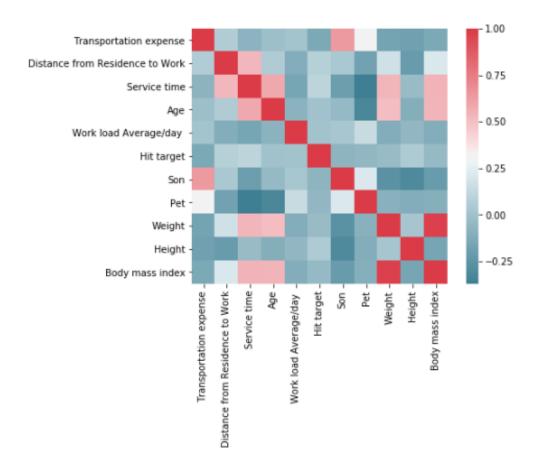
The chi-square test is a statistical test of independence to determine the dependency of two variables. It shares similarities with coefficient of determination, R². However, chi-square test is only applicable to categorical or nominal data while R² is only applicable to numeric data. Below, we can see the output of Chi-square test of independence for categorical variables from given data

1.4954158388291075e-65 Reason for absence 1.5194795569729106e-136 Month of absence 3.432975009623508e-55 Day of the week 1.1353710371238763e-40 Seasons 2.790172171150468e-52 Education 0.999355371240261 Social drinker 6.958462386179855e-17 Social smoker 0.42109821908319545 Disciplinary failure 2.207076113445463e-120

This test resides on P value of the variable. The null hypothesis of this test is that the two variables are independent of each other hence Alternate hypothesis would be that they are dependent on each other hence if the P value is greater than 0.05 which allows null hypothesis to be correct and states that the variables are not dependent on each other.

Here, 'Education' and 'Social smoker' are two categorical variables which follows null hypothesis and hence the target variable is independent of these two variables. Hence we will drop these two variable and proceed with other for model development

Correlation plot for the numerical variables from data is given below



Here, Dark Red indicates that the two variables are highly positively correlated to each other and Dark Blue indicates that the two variables are highly negatively correlated to each other. As we can see 'Weight' and 'Body mass index' are highly positively correlated to each other so we should drop any one variable from them to avoid multicollinearity.

Using above two techniques to eliminate variables, we have dropped 'Pet', 'Age', 'Son', 'Weight', 'Height', 'Education', 'Social smoker' from the data set.

2.2.4 Feature Scaling

Feature scaling is a method used to standardize the range of independent variables or features of data.

There are two methods of feature scaling viz. Normalization and standardization. Standardization is applied when data is normally distributed and normalization is applied in other cases. Here we have used normalization as there is skewness in some variables

3. Modelling

3.1 Model selection

Model selection depends on the Target variable. In case of given problem statement and dataset the target variable is continuous, hence the model will be regression model.

For regression model there are many models with which we can train our data and test on the same. We will consider some models in here and then depending on error rate we will decide on the same.

3.1.1 Multiple linear Regression

```
#Multiple linear regression
#train, test = train_test_split(df_empabs, test_size=0.2)
#MLR_model = sm.OLS(train.iloc[:,13], train.iloc[:,0:13]).fit()
#MLR_model.summary()
```

Dep. Variable:	Absenteeism t	time in hours		R-equar	ed:	0.308	
Model:		OLS	Adj.	R-equar	ed:	0.282	
Method:	Le	east Squares		F-etatie	tic:	11.83	
Date:	Sun,	12 Aug 2018	Prob (F-statist	ic): 3.6)7e-21	
Time:		05:48:19	Log-	Likeliho	od: -	1431.2	
No. Observations:		359		А	IC:	2888.	
Df Residuals:		346		В	IC:	2939.	
Df Model:		13					
Covariance Type:		nonrobust					
		coef	atd err	t	P> t	[0.025	0.975]
	ID		0.079	1.125	0.262	-0.066	0.243
Resso	on for absence		0.100	-3.288	0.001	-0.527	-0.133
	nth of absence		0.260	1.189	0.235	-0.203	0.822
	ay of the week		0.535	-1.982	0.048	-2.113	-0.008
_	Seasons		0.774	-0.019	0.985	-1.538	1.508
Transport	tation expense		3.346	3.249	0.001	4.291	17.454
Distance from Resi			2.950	-2.059	0.040	-11.877	-0.272
	Service time	13.9477	4.973	2.805	0.005	4.167	23.729
Work loa	d Average/day	2.0219	3.043	0.664	0.507	-3.963	8.007
	Hit target		3.099	2.618	0.009	2.019	14.208
Disc	iplinary fallure		4.141	-3.637	0.000	-23.207	-6.917
	Social drinker		1.857	2.197	0.029	0.426	7.730
Во	dy mass index		3.686	-1.028	0.305	-11.040	3.461
-	_,						
Omnibue: 3	99.438 Dur	bin-Watson:	1	.817			
Prob(Omnibus):	0.000 Jarqu	ie-Bera (JB):	17520	.370			
Skew:	5.028	Prob(JB):		0.00			

As you can see the Adjusted R-squared value, we can explain only about 30% of the data using our multiple linear regression model. This is not very impressive, but at least looking at the F-statistic and combined p-value we can reject the null hypothesis that target variable does not depend on any of the predictor variables.

After changing the test data to 40 percent the model gave output as follow

```
#Multiple linear regression
#train, test = train_test_split(df_empabs, test_size=0.4)
#MLR_model = sm.OLS(train.iloc[:,13], train.iloc[:,0:13]).fit()
#MLR_model.summary()
```

Dep. Variable	: Absent	eeism tir	me in hours	I	R-squar	ed:	0.307	
Model	Model: OLS		Adj.	R-squar	ed:	0.290		
Method	:	Lea	ast Squares		F-statis	tic:	17.90	
Date	:	Sun, 1	2 Aug 2018	Prob (i	F-statist	ic): 1.7	75e-34	
Time	:		07:32:51	Log-	Likeliho	od: -	2092.3	
No. Observations	:		538		А	IC:	4211.	
Df Residuals	:		525		В	IC:	4266.	
Df Model	:		13					
Covariance Type	:		nonrobust					
			coef	atd err	t	P≻ t	[0.025	0.975]
		ID	0.1161	0.055	2.115	0.035	0.008	0.224
Rea	son for at	sence	-0.3420	0.073	-4.696	0.000	-0.485	-0.199
М	onth of at	sence	0.6420	0.185	3.467	0.001	0.278	1.006
	Day of the	e week	-0.7277	0.378	-1.923	0.055	-1.471	0.016
	Se	asons	-0.9000	0.568	-1.584	0.114	-2.016	0.216
Transpo	rtation ex	epense	6.9846	2.402	2.908	0.004	2.266	11.703
Distance from Re	aldence t	Work	-3.2168	2.107	-1.527	0.127	-7.355	0.922
	Servic	e time	9.6767	3.704	2.612	0.009	2.400	16.953
Work Id	ad Avera	ge/day	4.8111	2.265	2.124	0.034	0.361	9.261
	HI	target	6.5114	2.250	2.894	0.004	2.092	10.931
DIS	ciplinary	fallure	-13.4982	2.786	-4.845	0.000	-18.971	-8.025
	Social	drinker	3.2906	1.374	2.395	0.017	0.591	5.990
Е	ody mass	Index	-2.3357	2.688	-0.869	0.385	-7.617	2.945
Omnibus:	625.225	Durb	In-Watson:	2	.117			
Prob(Omnibus):	0.000		-Bera (JB):					
Skew:	5.596		Prob(JB):		0.00			
Kurtosis:	43.965		Cond. No.		239.			

```
MLR model.summary()
# Dep. Variable:
# Absenteeism time in hours
# R-squared: for testsize=0.2
# 0.307
# Adj. R-squared:
# 0.289
# Dep. Variable:
# Absenteeism time in hours
# R-squared: for testsize=0.3
# 0.293
# Adj. R-squared:
# 0.272
# Dep. Variable:
# Absenteeism time in hours
# R-squared: for testsize=0.4
# 0.318
# Adj. R-squared:
# 0.295
# Dep. Variable:
# Absenteeism time in hours
# R-squared: for testsize=0.5
# 0.330
# Adj. R-squared:
# 0.301
```

After changing the test data also it did not change the predictive power of our regression model effectively. Therefore, this is the maximum accuracy that we can get from this model

```
#Calculate MAE
MAE(test.iloc[:,13], predictions_MLR)

#MAE 5.9 for test size=0.3

#MAE 5.8 for test size=0.2

#MAE 5.7 for test size=0.4

#MAE 6.34 for test size=0.5

#MAE 6.83 for test size=0.15
```

3.1.2 KNN Regressor

```
train, test = train_test_split(df_empabs, test_size=0.2)
KNN_model = KNeighborsRegressor(n_neighbors=3).fit(train.iloc[:,0:13], train.iloc[:,13])
predictions_KNN = KNN_model.predict(test.iloc[:,0:13])
#Calculate MAE
MAE(test.iloc[:,13], predictions_KNN)
```

```
#for KNN=3

#MAE 4.19 for test size=0.3

#MAE 3.63 for test size=0.2

#MAE 4.39 for test size=0.4

#MAE 4.14 for test size=0.15

#for KNN=2

#MAE 4.20 for test size=0.3

#MAE 4.53 for test size=0.2

#MAE 3.79 for test size=0.4

#MAE 5.33 for test size=0.15
```

3.1.3 Random Forest Regressor

```
train, test = train_test_split(df_empabs, test_size=0.2)
RFR_model = RandomForestRegressor(n_estimators = 20).fit(train.iloc[:,0:13], train.iloc[:,13])
RFR_Predictions = RFR_model.predict(test.iloc[:,0:13])
#Calculate MAE
MAE(test.iloc[:,13], RFR_Predictions)
```

#MAE 5.86 for test size=0.2 #MAE 5.2 for test size=0.3

4. Conclusion

4.1 Model Evaluation

Now that we have a few models for predicting the target variable, we need to decide which one to choose. There are several criteria that exist for evaluating and comparing models. We can compare models using any of the following criteria:

- 1. Predictive Performance
- 2. Interpretability
- 3. Computational Efficiency

In this case, if we consider Predictive performance of all models then after comparing their MAE (Mean absolute error) we can pick any one model as the MAE for both models are nearly same. This data set contains very less number of observations which affects the model building and predictions hence with high number of observations we might be able to perform well with the same models. The levels in Target variable were also very wide which resulted in average model building, with high number of observations this problem can also be solved.

4.2 What changes company should bring to reduce the number of absenteeism?

As the requirement of problem statement, we can use Random Forest model to predict the contribution of each independent variable resulted in variance of Target variable

```
> RF model = randomForest(Absenteeism.time.in.hours ~ ., df empabs,
                                   importance = TRUE)
> importance(RF model)
                                %IncMSE IncNodePurity
                              4.1255159 7356.1379
TD
                             12.0008526
                                          23818.3700
Reason.for.absence
Month.of.absence
                             3.3613963
                                         8883.9971
                                          7535.9078
Day.of.the.week
                             -0.1893287
                                          4377.5246
Seasons
                              4.8723884
                                         6915.3141
Transportation.expense
                             3.6879140
Distance.from.Residence.to.Work 4.5476416
                                         9642.9045
Service.time
                              5.1999099
                                         4635.8810
Work.load.Average.day
                             2.8401655
                                         12388.5741
                                          6717.5765
Hit.target
                             0.3666597
Disciplinary.failure
                             -0.8092790
                                           922.6676
Social.drinker
                             3.4668085
                                           1602.1034
Body.mass.index
                              5.6755639
                                         5468.1417
```

As we can observe here, Reason.for.absence contributes most than other variables while it comes to node splitting Reason.for.absence as well as Work.load.Average.day contributes more than other variables hence we can treat these two variables as important.

On Analyzing further for Reason.for.absence, we can aggregate the results with reason codes to related Absenteeism hours which gives result as follows

> 1	aggregate(data=df_er	mpabs1,predicted_abs	hrs~Reason.for.absence,sum)
	Reason.for.absence		
1	1	278.08844	
2	2	15.92431	
3	3	17.75265	
4	4	26.11160	
5	5	39.40333	
6	6	106.36139	
7	7	216.68603	
8	8	72.08159	
9	9	57.12723	
10	10	294.06592	
11	11	276.39156	
12		95.07083	
13	13	572.26403	
14		189.26185	
15	15	23.18990	
16		28.76178	
17	17	10.00768	
18	18	197.27571	
19	19	325.53114	
20	20	212.01087	
21		44.95017	
22		232.85517	
23	23	801.70911	
24	24	20.02198	
25		169.53102	
26		162.81082	
27	27	141.98287	
28	28	396.26621	

Here, we can observe that Reason.for.absence code ${\bf 13}$, ${\bf 23}$ are more often responsible for absenteeism. Which are as follows

13 - Pregnancy, childbirth and the puerperium

23 – Medical Consultation

Hence, company should look into these two reasons and workload average per day should be reduced to decrease the rate of absenteeism

4.3 How much losses every month can we project in 2011 if same trend of absenteeism continues?

We can answer this question by feeding whole independent variable's observation to our model and compare the resulted absenteeism hours to the month of absence, which will give us the pattern of monthly absenteeism for further coming year (provided same trend continues)

```
> predictions LR = predict(lm model, df empabs[,1:13])
> df empabs1$predicted abs hrs=predictions LR
> aggregate(data=df empabs1, predicted abs hrs~Month.of.absence,sum)
  Month.of.absence predicted abs hrs
                           227.1016
1
                1
2
                2
                           393.3676
3
                3
                          633.4353
4
                4
                          335.6969
5
                5
                          566.5677
6
                6
                          371.7813
7
                7
                          574.0996
8
                8
                          358.5721
9
                9
                          320.9492
10
                10
                          444.2815
11
               11
                          455.6757
12
               12
                           341.9666
```

These many number of hours are predicted by the model if the same trend continues for upcoming year

5. Appendix

5.1 R Code

```
rm(list=ls())
x = c("ggplot2", "corrgram", "DMwR", "caret", "randomForest", "unbalanced", "C50",
"dummies", "e1071", "Information",
   "MASS", "rpart", "gbm", "ROSE", 'sampling', 'DataCombine', 'inTrees', "usdm", "class")
lapply(x, require, character.only = TRUE)
df empabs=read.csv("Absenteeism at work Project.xls")
sum(is.na(df_empabs))
#first remove the observations for which target variable is null
df_empabs = df_empabs[which(!df_empabs$Absenteeism.time.in.hours %in% NA),]
dim(df empabs)
#data preprocessing
df_empabs$Work.load.Average.day=as.numeric(df_empabs$Work.load.Average.day)
#Change Reason code 0 to 20
df empabs$Reason.for.absence[which(df empabs$Reason.for.absence %in% 0)] = 20
#for month = 0, make it 12
df empabs$Month.of.absence[which(df empabs$Month.of.absence %in% 0)] = 12
#missing value analysis
df_empabs$Work.load.Average.day[is.na(df_empabs$Work.load.Average.daye)] =
median(df empabs$Work.load.Average.day, na.rm = T)
df_empabs$Month.of.absence[is.na(df_empabs$Month.of.absencee)] =
median(df empabs$Month.of.absence, na.rm = T)
df_empabs$Reason.for.absence[is.na(df_empabs$Reason.for.absence)] =
median(df empabs$Reason.for.absence, na.rm = T)
#other variables missing values imputation
df_empabs=knnImputation(df_empabs,k=3)
# df=subset(df empabs,select=
c(Transportation.expense, Distance.from.Residence.to.Work, Service.time, Work.load.Average.da
y, Hit. target,
#
                 Weight, Height, Body.mass.index))
```

```
#outlier analysis
multi.hist(df, main = NA, dcol = c("blue", "red"),
      dlty = c("solid", "solid"), bcol = "grey95")
#boxplot analysis
numeric var=c("Transportation.expense","Distance.from.Residence.to.Work","Service.time","A
ge","Work.load.Average.day","Hit.target",
       "Son","Pet","Weight","Height","Body.mass.index")
for (i in 1:length(numeric var))
{
 assign(pasteO("gn",i), ggplot(aes_string(y = (numeric_var[i]), x = "Absenteeism.time.in.hours"),
data = df_empabs)+
     stat boxplot(geom = "errorbar", width = 0.5) +
     geom_boxplot(outlier.colour="red", fill = "skyblue",outlier.shape=20,
            outlier.size=1, notch=FALSE) +
     theme(legend.position="bottom")+
     labs(y=numeric var[i],x="Absenteeism time")+
     ggtitle(paste("Box plot of Absenteeism time for",numeric_var[i])))
}
#gn1
#boxplot analysis
boxplot.stats(df empabs$Transportation.expense)$out
val = df empabs$Transportation.expense[df empabs$Transportation.expense %in%
boxplot.stats(df empabs$Transportation.expense)$out]
df empabs$Transportation.expense[df empabs$Transportation.expense %in% val] =
mean(df_empabs$Transportation.expense, na.rm = T)
boxplot.stats(df empabs$Hit.target)$out
val = df_empabs$Hit.target[df_empabs$Hit.target %in%
boxplot.stats(df_empabs$Hit.target)$out]
df empabs$Hit.target[df empabs$Hit.target %in% val] = mean(df empabs$Hit.target, na.rm =
T)
boxplot.stats(df_empabs$Service.time)$out
val = df empabs$Service.time[df empabs$Service.time %in%
boxplot.stats(df_empabs$Service.time)$out]
df_empabs$Service.time[df_empabs$Service.time %in% val] = mean(df_empabs$Service.time,
na.rm = T
boxplot.stats(df empabs$Age)$out
val = df_empabs$Age[df_empabs$Age %in% boxplot.stats(df_empabs$Age)$out]
df_empabs$Age[df_empabs$Age %in% val] = mean(df_empabs$Age, na.rm = T)
boxplot.stats(df_empabs$Work.load.Average.day)$out
```

```
val = df empabs$Work.load.Average.day[df empabs$Work.load.Average.day %in%
boxplot.stats(df_empabs$Work.load.Average.day)$out]
df empabs$Work.load.Average.day[df empabs$Work.load.Average.day %in% val] =
mean(df empabs$Work.load.Average.day, na.rm = T)
#Feature selection
numeric_index=c("Transportation.expense","Distance.from.Residence.to.Work","Service.time","
Age","Work.load.Average.day","Hit.target","Son","Pet","Weight","Height","Body.mass.index")
corrgram(df empabs[,numeric index], order = F,
     upper.panel=panel.pie, text.panel=panel.txt, main = "Correlation Plot")
## Chi-squared Test of Independence
factor_index = c("ID","Reason.for.absence", "Month.of.absence"
,"Day.of.the.week","Seasons","Disciplinary.failure","Education","Social.drinker","Social.smoker"
factor data = df empabs[,factor index]
for (i in 1:9)
{
print(names(factor data)[i])
print(chisq.test(table(df_empabs$Absenteeism.time.in.hours,factor_data[,i]),simulate.p.value
= TRUE))
}
df_empabs = subset(df_empabs,select = -
c(Age,Son,Pet,Weight,Height,Education,Social.smoker))
#feature scaling
#Normalisation
cnames = c("Transportation.expense"
,"Distance.from.Residence.to.Work","Service.time","Work.load.Average.day","Hit.target","Body
.mass.index")
for(i in cnames){
print(i)
df empabs[,i] = (df empabs[,i] - min(df empabs[,i]))/
  (max(df_empabs[,i] - min(df_empabs[,i])))
}
```

```
#sampling
train_index = sample(1:nrow(df_empabs), 0.8 * nrow(df_empabs))
train = df empabs[train index,]
test = df_empabs[-train_index,]
#Linear Regression
vif(df empabs[,-14])
lm_model = lm(Absenteeism.time.in.hours ~., data = train)
summary(Im model)
predictions_LR = predict(Im_model, test[,1:13])
#MAE
MAE = function(y, yhat){
mean(abs((y - yhat)))
}
#Calculate MAE
MAE(test[,14], predictions_LR)
#KNN regressor model
k= knn(train[,1:13],test[,1:13],train$Absenteeism.time.in.hours, k=3)
MAE(test[,14], as.numeric(k))
library("randomForest")
RF_model = randomForest(Absenteeism.time.in.hours ~ ., train, importance = TRUE)
importance(RF model)
predictions_RF = predict(RF_model, test[,1:13])
MAE(test[,14], predictions_RF)
#Predict for problem statement
predictions LR = predict(lm model, df empabs[,1:13])
df_empabs1=df_empabs
df_empabs1$predicted_abs_hrs=predictions_LR
aggregate(data=df_empabs1,predicted_abs_hrs~Reason.for.absence,sum)
aggregate(data=df_empabs1,predicted_abs_hrs~Month.of.absence,sum)
```