



भारतीय प्रबंध संस्थान कोयंबटूर
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Globalizing Indian Thought

CAPSTONE PROJECT 3 – PACKAGE PRICING AT MISSION HOSPITAL

Q: The objective is to develop a package pricing strategy for treating ailments of economically disadvantaged individuals, considering market demand and competition. This pricing strategy should be tailored to specific treatments and account for regional variations in pricing due to different state policies.

Furthermore, the package pricing strategy should address the perception of medical tourism and consider its impact on the healthcare sector. It is important to analyze and understand the differences between traditional and conventional strategies in order to support the transition to the conventional mode of package pricing.

Sol:-

```
# Let's read the data;
data_MH <- read.csv("clipboard", sep = "\t", header = T)
# Install some packages to do this analysis;
install.packages("dplyr")
library(dplyr)

install.packages("mice")
library(mice)

install.packages("ggcorrplot")
library(ggcorrplot)

install.packages("fastDummies")
library(fastDummies)

install.packages("lmtest")
library(lmtest)

install.packages("lm.beta")
library(lm.beta)

install.packages("car")
library(car)
```

```

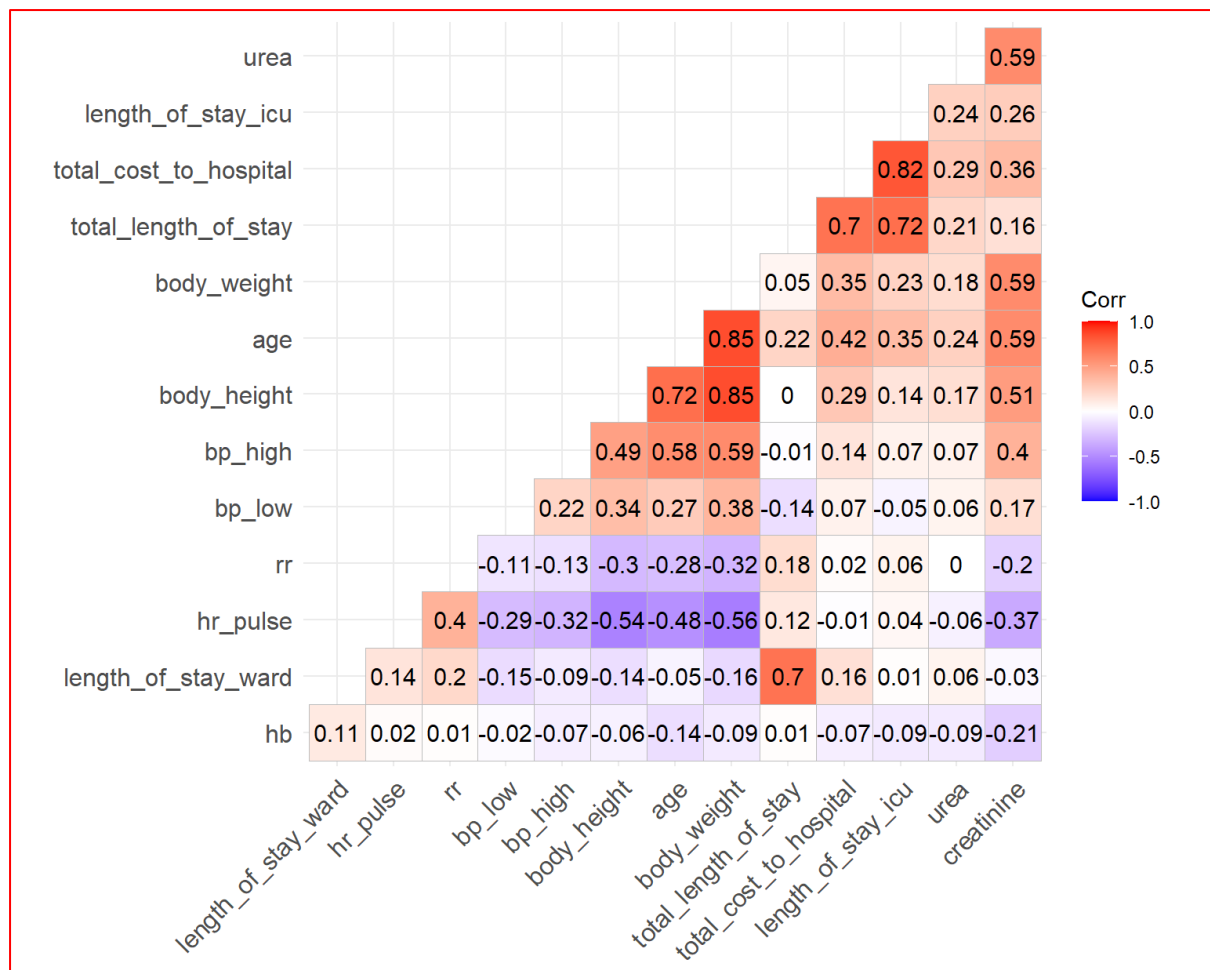
install.packages("janitor")
library(janitor)

# Pre processing the data;

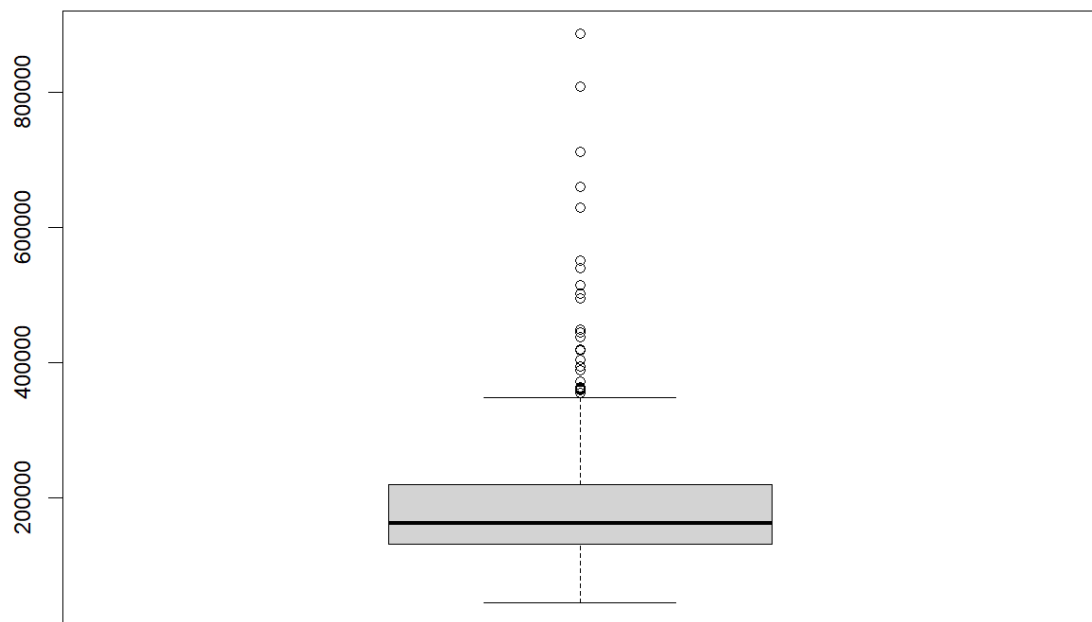
data1_MH <- clean_names(data_MH)
data1_MH$past_medical_history_code[which(is.na(data1_MH$past_medical_histoty_code))]
<- "None"
data2_MH <- data1_MH %>% mutate_if(is.character, as.factor)
data3_MH <- subset(data2_MH, select = -c(cost_of_implant))
# MICE imputation to convert all NA Values
set.seed(1234)
data4_MH <- mice(data3_MH)
# From the imputed values, we have chosen the variable 'Creatinine' for mean comparison
with the imputed values.
mean(data3_MH$creatinine, na.rm = T)
[1] 0.7469767
imp.values <- data4_MH$imp$creatinine
mean(imp.values$"1")
[1] 0.630303
> mean(imp.values$"2")
[1] 0.7242424
> mean(imp.values$"3")
[1] 0.6333333
> mean(imp.values$"4")
[1] 0.6272727
> mean(imp.values$"5")
[1] 0.6969697

# Since the mean value has most proximity to the 1st imputed value, we are selecting that
imputation
data5_MH <- complete(data4_MH, 1)
# Pre-processing of data
names(data5)[23] <- "implant_used"
data5_MH$bp_low <- as.numeric(data5_MH$bp_low)
data6_MH <- subset(data5_MH, select = -c(sl))
# Correlation matrix for creating correlation plot
data6.1 <- data6_MH %>% select_if(is.numeric)
cor.matrix <- cor(data6.1)

```



```
# Creating fast dummies of categorical variables
data7_MH <- dummy_cols(data6_MH, remove_most_frequent_dummy = T,
remove_selected_columns = T)
# Outlier detection and removal from dependent variable
outlier <- boxplot(data7_MH$total_cost_to_hospital)$out
> length(outlier)
[1] 24
> outlier.data <- data7_MH[which(data7_MH$total_cost_to_hospital %in% outlier),]
> data8_MH <- data7_MH[-which(data7_MH$total_cost_to_hospital %in% outlier),]
> mode11 <- lm(total_cost_to_hospital ~., data8_MH)
> options(scipen = 100)
```



```
> summary(mode11)
```

Call:

```
lm(formula = total_cost_to_hospital ~ ., data = data8_MH)
```

Residuals:

```
   Min     1Q  Median     3Q    Max
-63942 -16879     0 14833 92114
```

Coefficients:

##	Estimate	Std. Error	t value
## (Intercept)	54750.27	30864.34	1.774
## age	169.28	268.14	0.631
## body_weight	-26.85	289.19	-0.093
## body_height	158.60	109.71	1.446
## hr_pulse	81.23	141.11	0.576
## bp_high	10.65	126.05	0.084
## bp_low	293.45	340.27	0.862
## rr	-1240.58	725.12	-1.711
## hb	-773.45	763.61	-1.013
## urea	-27.06	199.48	-0.136
## creatinine	20219.81	8849.99	2.285

```

## total_length_of_stay          444.89 10240.39 0.043
## length_of_stay_icu            17674.46 10335.92 1.710
## length_of_stay_ward           4977.71 10145.10 0.491
## gender_F                      732.94 4719.94 0.155
## marital_status_MARRIED        -8416.46 10596.59 -0.794
## key_complaints_code_ACHD       -5601.42 7930.75 -0.706
## `key_complaints_code_CAD-DVD`  9255.87 10807.93 0.856
## `key_complaints_code_CAD-SVD` -43194.20 24456.59 -1.766
## `key_complaints_code_CAD-TVD` -85.25 11194.69 -0.008
## `key_complaints_code_CAD-VSD` -10271.08 22180.11 -0.463
## `key_complaints_code_OS-ASD`   240.09 8809.88 0.027
## `key_complaints_code_other- respiratory` 3421.12 7967.69 0.429
## `key_complaints_code_other-general` -39374.88 23354.88 -1.686
## `key_complaints_code_other-nervous` 1187.39 14364.91 0.083
## `key_complaints_code_other-teratology` 31663.00 8392.36 3.773
## `key_complaints_code_PM-VSD` 25371.32 13868.09 1.829
## key_complaints_code_RHD        -4181.06 9791.36 -0.427
## past_medical_history_code_Diabetes1 8471.51 12390.38 0.684
## past_medical_history_code_Diabetes2 46771.15 19210.71 2.435
## past_medical_history_code_hypertension1 2939.12 10837.34 0.271
## past_medical_history_code_Hypertension1 -23568.57 19932.47 -1.182
## past_medical_history_code_hypertension2 -11397.01 10192.43 -1.118
## past_medical_history_code_hypertension3 796.33 17357.20 0.046
## past_medical_history_code_other -11950.97 8606.25 -1.389
## mode_of_arrival_AMBULANCE      47526.71 32619.11 1.457
## mode_of_arrival_TRANSFERRED    -30276.07 16215.94 -1.867
## state_at_the_time_of_arrival_CONFUSED -5647.34 45179.48 -0.125
## type_of_admsn_EMERGENCY        -65286.02 31980.45 -2.041
## implant_used_Y                 78650.77 8473.03 9.282
##                               Pr(>|t|)
## (Intercept)                    0.077733 .
## age                            0.528626
## body_weight                     0.926126
## body_height                     0.149986
## hr_pulse                        0.565557

```

## bp_high	0.932780
## bp_low	0.389598
## rr	0.088789 .
## hb	0.312448
## urea	0.892237
## creatinine	0.023470 *
## total_length_of_stay	0.965394
## length_of_stay_icu	0.088951 .
## length_of_stay_ward	0.624258
## gender_F	0.876766
## marital_status_MARRIED	0.428067
## key_complaints_code_ACHD	0.480900
## `key_complaints_code_CAD-DVD`	0.392892
## `key_complaints_code_CAD-SVD`	0.079028 .
## `key_complaints_code_CAD-TVD`	0.993932
## `key_complaints_code_CAD-VSD`	0.643857
## `key_complaints_code_OS-ASD`	0.978288
## `key_complaints_code_other- respiratory`	0.668153
## `key_complaints_code_other-general`	0.093503 .
## `key_complaints_code_other-nervous`	0.934212
## `key_complaints_code_other-teratology`	0.000217 ***
## `key_complaints_code_PM-VSD`	0.068947 .
## key_complaints_code_RHD	0.669867
## past_medical_history_code_Diabetes1	0.495014
## past_medical_history_code_Diabetes2	0.015861 *
## past_medical_history_code_hypertension1	0.786539
## past_medical_history_code_Hypertension1	0.238564
## past_medical_history_code_hypertension2	0.264946
## past_medical_history_code_hypertension3	0.963457
## past_medical_history_code_other	0.166621
## mode_of_arrival_AMBULANCE	0.146815
## mode_of_arrival_TRANSFERRED	0.063484 .
## state_at_the_time_of_arrival_CONFUSED	0.900662
## type_of_admsn_EMERGENCY	0.042635 *
## implant_used_Y	< 0.0000000000000002 ***

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
# Stepwise regression to identify the significant variables in the model
```

```
> mode12 <- step(mode11, trace = 0)
```

```
> summary(mode12)
```

Call:

```
lm(formula = total_cost_to_hospital ~ body_height + bp_low +  
  rr + hb + creatinine + length_of_stay_icu + length_of_stay_ward +  
  key_complaints_code_ + `key_complaints_code_CAD-DVD` + `key_complaints_code_CAD-  
SVD` +  
  `key_complaints_code_other-general` + `key_complaints_code_other-teratology` +  
  `key_complaints_code_PM-VSD` + past_medical_history_code_Diabetes2 +  
  past_medical_history_code_Hypertension1 + past_medical_history_code_hypertension2 +  
  past_medical_history_code_other + mode_of_arrival_AMBULANCE +  
  mode_of_arrival_TRANSFERRED + type_of_admsn_EMERGENCY + implant_used_y_n_Y,  
  data = data8_MH)
```

Residuals:

```
   Min      1Q  Median      3Q      Max  
-66256 -14930  -120   15521  96471
```

```
## Coefficients:
```

##	Estimate	Std. Error	t value
## (Intercept)	60894.73	20123.04	3.026
## body_height	146.17	66.74	2.190
## rr	-1277.62	659.11	-1.938
## creatinine	19466.47	4845.00	4.018
## length_of_stay_icu	18431.69	1050.34	17.548
## length_of_stay_ward	5319.80	611.65	8.697
## `key_complaints_code_CAD-DVD`	11315.37	7625.61	1.484
## `key_complaints_code_CAD-SVD`	-39429.64	20902.43	-1.886
## `key_complaints_code_other-general`	-34886.66	21159.77	-1.649
## `key_complaints_code_other-teratology`	29709.33	6886.53	4.314
## `key_complaints_code_PM-VSD`	26046.62	12445.49	2.093
## past_medical_history_code_Diabetes2	43754.53	15289.69	2.862
## past_medical_history_code_Hypertension1	-24938.34	17751.26	-1.405

```

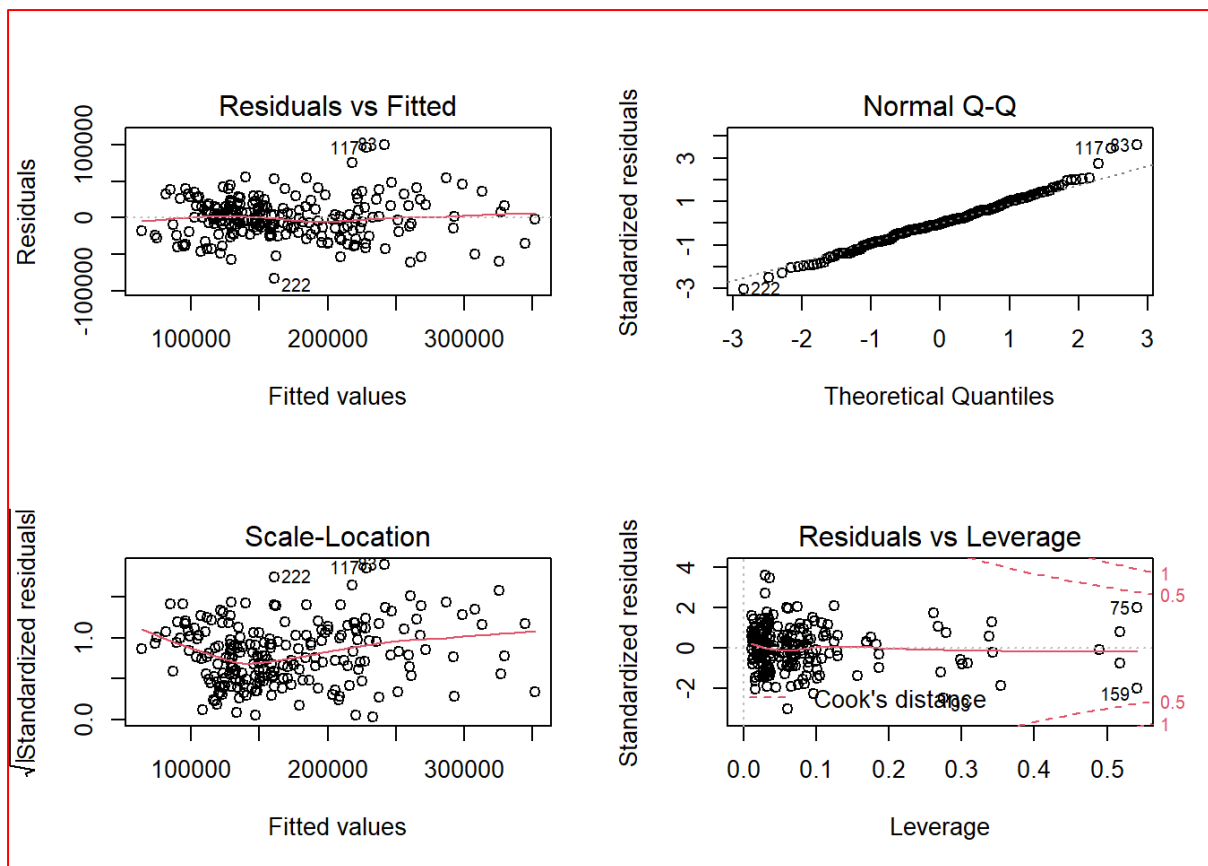
## past_medical_history_code_other      -11872.14  7883.21 -1.506
## mode_of_arrival_AMBULANCE             54838.55  30379.38  1.805
## mode_of_arrival_TRANSFERRED           -31423.99  14729.27 -2.133
## type_of_admsn_EMERGENCY               -72979.91  29536.34 -2.471
## implant_used_Y                        76820.61  5584.35  13.756
##                                     Pr(>|t|)
## (Intercept)                          0.00279 **
## body_height                          0.02964 *
## rr                                    0.05394 .
## creatinine                           0.00008231642922645 ***
## length_of_stay_icu                   < 0.00000000000000002 ***
## length_of_stay_ward                   0.000000000000000108 ***
## `key_complaints_code_CAD-DVD`         0.13937
## `key_complaints_code_CAD-SVD`         0.06065 .
## `key_complaints_code_other-general`   0.10073
## `key_complaints_code_other-teratology` 0.00002485689935067 ***
## `key_complaints_code_PM-VSD`          0.03759 *
## past_medical_history_code_Diabetes2    0.00465 **
## past_medical_history_code_Hypertension1 0.16156
## past_medical_history_code_other        0.13360
## mode_of_arrival_AMBULANCE             0.07252 .
## mode_of_arrival_TRANSFERRED           0.03407 *
## type_of_admsn_EMERGENCY               0.01429 *
## implant_used_Y                        < 0.00000000000000002 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 28140 on 206 degrees of freedom
## Multiple R-squared:  0.8192, Adjusted R-squared:  0.8043
## F-statistic: 54.91 on 17 and 206 DF, p-value: < 0.000000000000000022

```

```

> par(mfrow = c(2,2))
> plot(mode12)
72, 135, 142

```

```
> bptest(mode12)
```

studentized Breusch-Pagan test

data: mode12

BP = 43.247, df = 21, p-value = 0.002923

Conducting Breusch-Pagan test for checking heteroscedasticity

```
bptest(model2)
```

studentized Breusch-Pagan test

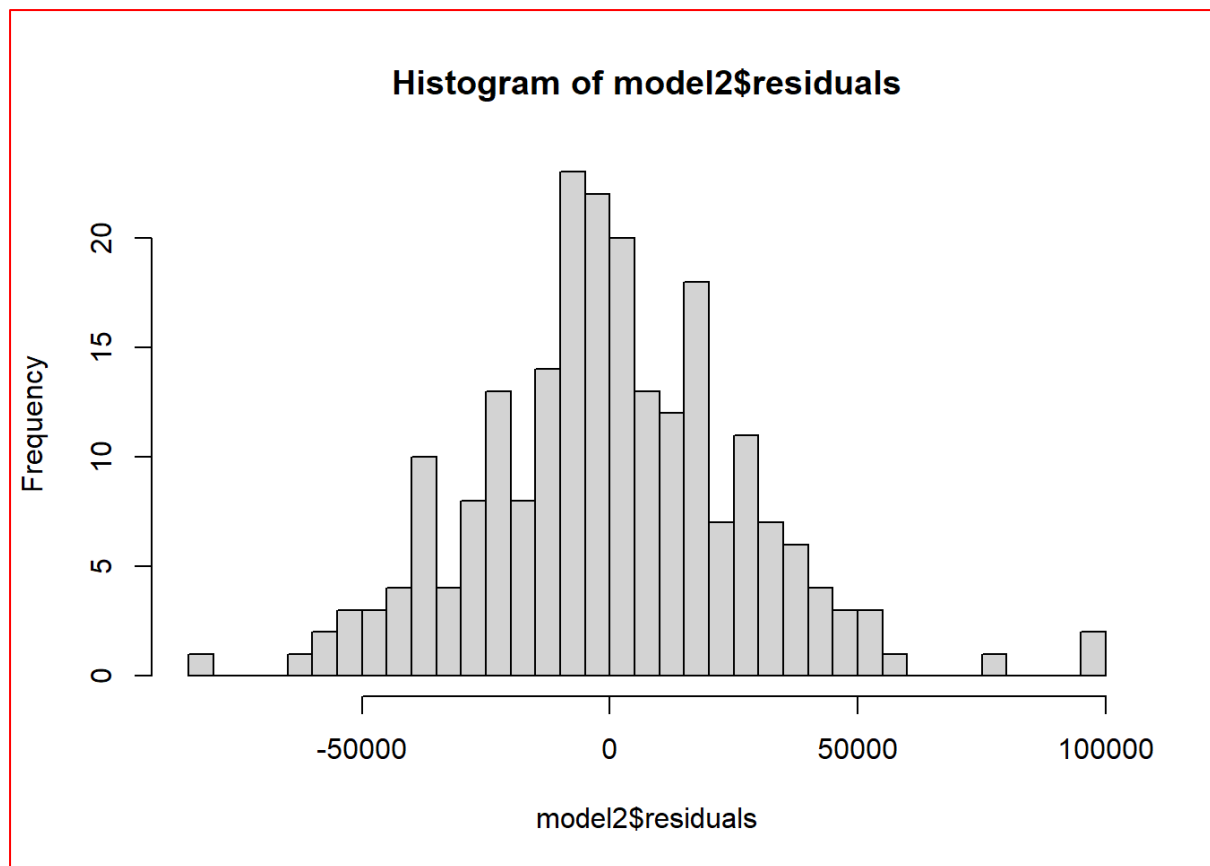
data: model2

BP = 30.557, df = 17, p-value = 0.0226

Getting the standard normal values of regression coefficients to identify the order of significance

```
par(mfrow= c(1,1))
```

```
hist(model2$residuals, breaks = 30)
```



```
library(lm.beta)
```

```
coeff <- lm.beta(model2)
```

```
std.coef <- data.frame(coeff$standardized.coefficients)
```

```
std.coef
```

##	coeff.standardized.coefficients
## (Intercept)	NA
## body_height	0.09025878
## rr	-0.06988629
## creatinine	0.15915995
## length_of_stay_icu	0.59033228
## length_of_stay_ward	0.27803070
## `key_complaints_code_CAD-DVD`	0.05411276
## `key_complaints_code_CAD-SVD`	-0.05843636
## `key_complaints_code_other-general`	-0.05170347
## `key_complaints_code_other-teratology`	0.13347249
## `key_complaints_code_PM-VSD`	0.06625582
## past_medical_history_code_Diabetes2	0.09129211

## past_medical_history_code_Hypertension1	-0.04516408
## past_medical_history_code_other	-0.04675378
## mode_of_arrival_AMBULANCE	0.24071796
## mode_of_arrival_TRANSFERRED	-0.06556492
## type_of_admsn_EMERGENCY	-0.32787041
## implant_used_Y	0.45894933

Checking multicollinearity

```
vif <- data.frame(vif(model2))
```

```
vif
```

##	vif.model2.
## body_height	1.935469
## rr	1.481242
## creatinine	1.788177
## length_of_stay_icu	1.289602
## length_of_stay_ward	1.164468
## `key_complaints_code_CAD-DVD`	1.515444
## `key_complaints_code_CAD-SVD`	1.093562
## `key_complaints_code_other-general`	1.120655
## `key_complaints_code_other-teratology`	1.090756
## `key_complaints_code_PM-VSD`	1.142085
## past_medical_history_code_Diabetes2	1.159703
## past_medical_history_code_Hypertension1	1.177712
## past_medical_history_code_other	1.098273
## mode_of_arrival_AMBULANCE	20.264324
## mode_of_arrival_TRANSFERRED	1.076247
## type_of_admsn_EMERGENCY	20.065060
## implant_used_Y	1.268380

Creating Training and test data

```
index <- sample(1:nrow(data8), 0.80*(nrow(data8)))
```

```
train_data <- data8[index,]
```

```
test_data <- data8[-index,]
```

Creating model using training data

```
model3 <- lm(total_cost_to_hospital ~., train_data)
```

```
summary(model3)
```

```
##
```

```
## Call:
```

```
## lm(formula = total_cost_to_hospital ~ ., data = train_data)
```

```
##
```

```
## Residuals:
```

```
##   Min     1Q  Median     3Q    Max
```

```
## -74361 -16277 -1021  16750  98322
```

```
##
```

```
## Coefficients:
```

```
##
```

```
Estimate Std. Error t value
```

```
## (Intercept)      63658.99  37427.60  1.701
```

```
## age              179.00   331.45  0.540
```

```
## body_weight      112.12   342.78  0.327
```

```
## body_height      134.46   135.38  0.993
```

```
## hr_pulse         120.41   165.49  0.728
```

```
## bp_high          84.75   161.20  0.526
```

```
## bp_low           149.15   428.55  0.348
```

```
## rr               -2112.67  899.73 -2.348
```

```
## hb               -743.55   903.77 -0.823
```

```
## urea             -40.22   238.80 -0.168
```

```
## creatinine       20591.93 11018.67  1.869
```

```
## total_length_of_stay      1801.76 11095.76  0.162
```

```
## length_of_stay_icu        16019.11 11266.66  1.422
```

```
## length_of_stay_ward       4091.76 10991.51  0.372
```

```
## gender_F            -425.35  5733.43 -0.074
```

```
## marital_status_MARRIED    -15808.27 13072.06 -1.209
```

```
## key_complaints_code_ACHD   -2888.34  9608.64 -0.301
```

```
## `key_complaints_code_CAD-DVD`    12171.25 12967.32  0.939
```

```
## `key_complaints_code_CAD-SVD`   -48579.52 26482.56 -1.834
```

```
## `key_complaints_code_CAD-TVD`    1866.71 13541.02  0.138
```

```
## `key_complaints_code_CAD-VSD`  -11804.45 23759.70 -0.497
```

```
## `key_complaints_code_OS-ASD`    -3623.44 10449.68 -0.347
```

```
## `key_complaints_code_other- respiratory`  3953.95  9988.24  0.396
```

```

## `key_complaints_code_other-general`    -35389.31  25279.64 -1.400
## `key_complaints_code_other-nervous`     1482.35  16901.96  0.088
## `key_complaints_code_other-teratology`  32526.94  10070.20  3.230
## `key_complaints_code_PM-VSD`           30238.11  18038.72  1.676
## key_complaints_code_RHD                 -5806.00  11711.22 -0.496
## past_medical_history_code_Diabetes1     -1283.97  16255.29 -0.079
## past_medical_history_code_Diabetes2     58785.13  24715.09  2.379
## past_medical_history_code_hypertension1  3732.02  12800.26  0.292
## past_medical_history_code_Hypertension1 -24411.24  22023.71 -1.108
## past_medical_history_code_hypertension2 -14788.93  13160.41 -1.124
## past_medical_history_code_hypertension3  77.17  18868.69  0.004
## past_medical_history_code_other         -12209.15  9754.71 -1.252
## mode_of_arrival_AMBULANCE               55586.89  35531.88  1.564
## mode_of_arrival_TRANSFERRED             -44336.48  24991.05 -1.774
## state_at_the_time_of_arrival_CONFUSED   -16002.21  52088.37 -0.307
## type_of_admsn_EMERGENCY                 -74634.31  34485.79 -2.164
## implant_used_Y                          81968.11  9973.07  8.219
##                                         Pr(>|t|)
## (Intercept)                            0.09121 .
## age                                    0.59001
## body_weight                            0.74409
## body_height                            0.32234
## hr_pulse                              0.46809
## bp_high                               0.59989
## bp_low                                0.72834
## rr                                    0.02028 *
## hb                                    0.41207
## urea                                  0.86648
## creatinine                            0.06375 .
## total_length_of_stay                   0.87124
## length_of_stay_icu                     0.15732
## length_of_stay_ward                     0.71026
## gender_F                              0.94097
## marital_status_MARRIED                 0.22859
## key_complaints_code_ACHD               0.76417

```

```

## `key_complaints_code_CAD-DVD`          0.34956
## `key_complaints_code_CAD-SVD`          0.06873 .
## `key_complaints_code_CAD-TVD`          0.89055
## `key_complaints_code_CAD-VSD`          0.62010
## `key_complaints_code_OS-ASD`           0.72930
## `key_complaints_code_other- respiratory` 0.69281
## `key_complaints_code_other-general`     0.16377
## `key_complaints_code_other-nervous`     0.93024
## `key_complaints_code_other-teratology`  0.00155 **
## `key_complaints_code_PM-VSD`           0.09593 .
## key_complaints_code_RHD                0.62084
## past_medical_history_code_Diabetes1     0.93716
## past_medical_history_code_Diabetes2     0.01874 *
## past_medical_history_code_hypertension1 0.77106
## past_medical_history_code_Hypertension1 0.26960
## past_medical_history_code_hypertension2 0.26306
## past_medical_history_code_hypertension3 0.99674
## past_medical_history_code_other         0.21281
## mode_of_arrival_AMBULANCE              0.11999
## mode_of_arrival_TRANSFERRED            0.07824 .
## state_at_the_time_of_arrival_CONFUSED  0.75914
## type_of_admsn_EMERGENCY                0.03216 *
## implant_used_Y                        0.000000000000129 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 30810 on 139 degrees of freedom
## Multiple R-squared:  0.8239, Adjusted R-squared:  0.7745
## F-statistic: 16.68 on 39 and 139 DF,  p-value: < 0.00000000000000022

```

```

> mode14 <- step(mode13, trace = 0)
> summary(mode14)

```

Call:

```

lm(formula = total_cost_to_hospital ~ body_height + hr_pulse +
    rr + hb + creatinine + length_of_stay_icu + length_of_stay_ward +

```

```
key_complaints_code_ + key_complaints_code_ACHD + `key_complaints_code_CAD-SVD`
+
`key_complaints_code_other-teratology` + `key_complaints_code_PM-VSD` +
past_medical_history_code_hypertension1 + mode_of_arrival_AMBULANCE +
mode_of_arrival_TRANSFERRED + state_at_the_time_of_arrival_CONFUSED +
type_of_admsn_EMERGENCY + implant_used_y_n_Y, data = train_data)
```

Residuals:

```
Min    1Q  Median    3Q   Max
-62297 -14711   471  14209 103666
```

Coefficients:

##	Estimate	Std. Error	t value
## (Intercept)	77505.98	23332.76	3.322
## body_height	180.91	74.66	2.423
## rr	-2273.86	764.80	-2.973
## creatinine	18362.56	5444.29	3.373
## length_of_stay_icu	18482.24	1209.90	15.276
## length_of_stay_ward	5692.97	687.81	8.277
## `key_complaints_code_CAD-SVD`	-47367.06	21820.58	-2.171
## `key_complaints_code_other-general`	-35057.81	22131.67	-1.584
## `key_complaints_code_other-teratology`	30585.50	8150.25	3.753
## `key_complaints_code_PM-VSD`	32859.88	15969.00	2.058
## past_medical_history_code_Diabetes2	54088.27	18379.04	2.943
## mode_of_arrival_AMBULANCE	55114.65	31645.29	1.742
## mode_of_arrival_TRANSFERRED	-43879.89	22437.95	-1.956
## type_of_admsn_EMERGENCY	-74398.91	30758.42	-2.419
## implant_used_Y	77777.92	6288.27	12.369
##	Pr(> t)		
## (Intercept)	0.001103	**	
## body_height	0.016471	*	
## rr	0.003392	**	
## creatinine	0.000928	***	
## length_of_stay_icu	< 0.000000000000000002	***	
## length_of_stay_ward	0.00000000000000419	***	
## `key_complaints_code_CAD-SVD`	0.031388	*	
## `key_complaints_code_other-general`	0.115107		

```
## `key_complaints_code_other-teratology`      0.000242 ***
## `key_complaints_code_PM-VSD`                0.041199 *
## past_medical_history_code_Diabetes2         0.003723 **
## mode_of_arrival_AMBULANCE                   0.083447 .
## mode_of_arrival_TRANSFERRED                 0.052210 .
## type_of_admsn_EMERGENCY                     0.016665 *
## implant_used_Y                             < 0.0000000000000002 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 29320 on 164 degrees of freedom
## Multiple R-squared:  0.8119, Adjusted R-squared:  0.7959
## F-statistic: 50.57 on 14 and 164 DF, p-value: < 0.00000000000000022
```

Using the model to predict values of test data and comparing the results

```
test_data$Predicted <- predict(model4, test_data)
comparison <- data.frame(Actual_cost =test_data$total_cost_to_hospital, Predicted_cost=
test_data$Predicted)
comparison
```

```
##   Actual_cost Predicted_cost
## 1   341109.0    273140.85
## 2   144037.2    126197.74
## 3   164962.0    107180.31
## 4   120131.0    133664.99
## 5   138923.0    156152.22
## 6   122892.0    111192.02
## 7   142552.0    134051.78
## 8   109085.8    106938.44
## 9   125643.0    118862.96
## 10  128196.0     95529.98
## 11  109085.8    106938.44
## 12  125643.0    120518.31
## 13  294615.9    296296.33
## 14  156576.9    154020.51
## 15  109575.6    167453.97
```



```
## 16 201219.0 202625.43
## 17 214679.0 270628.50
## 18 189701.5 178381.03
## 19 139723.0 152435.75
## 20 119685.6 117874.87
## 21 276458.0 260702.84
## 22 150337.0 130609.74
## 23 139067.0 146234.87
## 24 127899.0 110775.28
## 25 146355.0 170204.38
## 26 97060.8 119870.27
## 27 106070.0 111186.56
## 28 140372.0 155320.25
## 29 138769.4 128346.19
## 30 77241.0 74704.68
## 31 49700.0 73587.04
## 32 137273.0 100162.45
## 33 193543.0 228469.83
## 34 191102.0 234498.35
## 35 132585.0 160565.91
## 36 170654.0 162985.88
## 37 174074.0 128594.76
## 38 210622.0 210393.38
## 39 46093.0 50167.98
## 40 188824.0 196148.22
## 41 146700.0 144152.96
## 42 149462.0 158009.78
## 43 186450.0 227407.55
## 44 132997.0 132721.96
## 45 248112.0 265147.35
```

```
# Checking the validity of model using MAPE and RMSE
```

```
mape <- mean(abs(comparison$Actual_cost-
comparison$Predicted_cost)/comparison$Actual_cost)
```

```
mape
```

```
## [1] 0.1253493
```

```

1-mape
## [1] 0.8746507
rmse <- sqrt(mean(comparison$Actual_cost- comparison$Predicted_cost)^2)
rmse
## [1] 1226.584

```

Conclusion:-

The MAPE value of 0.1253493 indicates that, on average, the absolute percentage difference between the predicted values and the actual values is approximately 12.53%. This value represents the average relative error in percentage terms.

1. The MAPE value suggests that, on average, the model's predictions deviate by around 12.53% from the actual values.
2. The RMSE value indicates that the average prediction error, in absolute terms, is approximately 1226.584. This value gives an idea of the magnitude of the errors made by the model.

Presented by;

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17.06.2023

*******Thank You*******