

```
# Roll no. 33140
# Batch: L9
# P.S.: Application of Linear regression on Heart disease dataset to predict
        the fate (prob. of heart disease)
```

```
R version 3.6.2 (2019-12-12) -- "Dark and Stormy Night"
Copyright (C) 2019 The R Foundation for Statistical Computing
Platform: x86_64-w64-mingw32/x64 (64-bit)
```

```
R is free software and comes with ABSOLUTELY NO WARRANTY.
You are welcome to redistribute it under certain conditions.
Type 'license()' or 'licence()' for distribution details.
```

```
Natural language support but running in an English locale
```

```
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```

```
Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.
```

```
[workspace loaded from G:/College/SL6/Assignment6/.RData]
```

```
> # Set working directory
```

```
> setwd("G:/College/SL6/Assignment6/")
```

```
> # Read the CSV file and analyse
```

```
> hdata <- read.csv("../..\\SI-VI DataSets/HeartDisease/Cleveland.csv", header=TRUE,
                    sep=",")
```

```
> names(hdata)
```

```
[1] "x63.0" "x1.0" "x1.0.1" "x145.0" "x233.0" "x1.0.2" "x2.0" "x150.0" "x0.0"
[2] "x2.3" "x3.0" "x0.0.1" "x6.0"
[14] "x0"
```

```
> str(hdata)
```

```
'data.frame': 302 obs. of 14 variables:
 $ x63.0 : num 67 67 37 41 56 62 57 63 53 57 ...
 $ x1.0 : num 1 1 1 0 1 0 0 1 1 1 ...
 $ x1.0.1: num 4 4 3 2 2 4 4 4 4 4 ...
 $ x145.0: num 160 120 130 130 120 140 120 130 140 140 ...
 $ x233.0: num 286 229 250 204 236 268 354 254 203 192 ...
 $ x1.0.2: num 0 0 0 0 0 0 0 0 1 0 ...
 $ x2.0 : num 2 2 0 2 0 2 0 2 2 0 ...
 $ x150.0: num 108 129 187 172 178 160 163 147 155 148 ...
 $ x0.0 : num 1 1 0 0 0 0 1 0 1 0 ...
 $ x2.3 : num 1.5 2.6 3.5 1.4 0.8 3.6 0.6 1.4 3.1 0.4 ...
 $ x3.0 : num 2 2 3 1 1 3 1 2 3 2 ...
 $ x0.0.1: Factor w/ 5 levels "?","0.0","1.0",...: 5 4 2 2 2 4 2 3 2 2 ...
 $ x6.0 : Factor w/ 4 levels "?","3.0","6.0",...: 2 4 2 2 2 2 2 4 4 3 ...
 $ x0 : int 2 1 0 0 0 3 0 2 1 0 ...
```

```
> dim(hdata)
```

```
[1] 302 14
```

```
> # Change the headers
```

```
> names(hdata)[1] <- "age"
```

```

> names(hdata)[2] <- "sex"
> names(hdata)[3] <- "cp"
> names(hdata)[4] <- "trestbps"
> names(hdata)[5] <- "chol"
> names(hdata)[6] <- "fbs"
> names(hdata)[7] <- "restecg"
> names(hdata)[8] <- "thalach"
> names(hdata)[9] <- "exang"
> names(hdata)[10] <- "oldpeak"
> names(hdata)[11] <- "slope"
> names(hdata)[12] <- "ca"
> names(hdata)[13] <- "thal"
> names(hdata)[14] <- "num"

> hdata$ca
[1] 3.0 2.0 0.0 0.0 0.0 2.0 0.0 1.0 0.0 0.0 0.0 1.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0
[30] 2.0 2.0 0.0 0.0 0.0 0.0 0.0 1.0 1.0 0.0 3.0 0.0 2.0 0.0 0.0 1.0 0.0 0.0 1.0 0.0
[59] 1.0 0.0 0.0 3.0 0.0 1.0 2.0 0.0 0.0 0.0 0.0 0.0 2.0 2.0 2.0 1.0 0.0 1.0 1.0 0.0
[88] 0.0 0.0 0.0 3.0 3.0 0.0 0.0 1.0 1.0 2.0 1.0 0.0 0.0 0.0 1.0 1.0 3.0 0.0 1.0 1.0
[117] 0.0 3.0 1.0 2.0 3.0 0.0 0.0 1.0 0.0 2.0 1.0 0.0 0.0 0.0 1.0 0.0 0.0 0.0 0.0 0.0
[146] 3.0 0.0 0.0 1.0 0.0 0.0 0.0 1.0 1.0 3.0 0.0 2.0 2.0 1.0 0.0 3.0 0.0 0.0 2.0 0.0
[175] 1.0 3.0 1.0 1.0 3.0 0.0 2.0 2.0 0.0 0.0 2.0 0.0 3.0 1.0 3.0 0.0 3.0 2.0 3.0 0.0
[204] 0.0 3.0 2.0 0.0 0.0 0.0 0.0 0.0 0.0 2.0 1.0 0.0 0.0 0.0 2.0 0.0 0.0 0.0 0.0 2.0
[233] 1.0 1.0 2.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 2.0 0.0 0.0 1.0 1.0 2.0 0.0 0.0 1.0 1.0
[262] 0.0 0.0 1.0 0.0 0.0 1.0 0.0 0.0 1.0 0.0 2.0 0.0 2.0 0.0 1.0 0.0 1.0 0.0 1.0 0.0
[291] 0.0 0.0 2.0 0.0 0.0 2.0 0.0 0.0 2.0 1.0 1.0 ?

Levels: ? 0.0 1.0 2.0 3.0

```

```

> levels(hdata$ca)[levels(hdata$ca) == "?"] <- "0.0"

```

```

> hdata
  age sex cp trestbps chol fbs restecg thalach exang oldpeak slope ca thal num
1  67   1  4     160  286   0         2     108    1     1.5     2 3.0  3.0   2
2  67   1  4     120  229   0         2     129    1     2.6     2 2.0  7.0   1
3  37   1  3     130  250   0         0     187    0     3.5     3 0.0  3.0   0
4  41   0  2     130  204   0         2     172    0     1.4     1 0.0  3.0   0
5  56   1  2     120  236   0         0     178    0     0.8     1 0.0  3.0   0
6  62   0  4     140  268   0         2     160    0     3.6     3 2.0  3.0   3
7  57   0  4     120  354   0         0     163    1     0.6     1 0.0  3.0   0
8  63   1  4     130  254   0         2     147    0     1.4     2 1.0  7.0   2
9  53   1  4     140  203   1         2     155    1     3.1     3 0.0  7.0   1

```

10	57	1	4	140	192	0	0	148	0	0.4	2	0.0	6.0	0
11	56	0	2	140	294	0	2	153	0	1.3	2	0.0	3.0	0
12	56	1	3	130	256	1	2	142	1	0.6	2	1.0	6.0	2
13	44	1	2	120	263	0	0	173	0	0.0	1	0.0	7.0	0
14	52	1	3	172	199	1	0	162	0	0.5	1	0.0	7.0	0
15	57	1	3	150	168	0	0	174	0	1.6	1	0.0	3.0	0
16	48	1	2	110	229	0	0	168	0	1.0	3	0.0	7.0	1
17	54	1	4	140	239	0	0	160	0	1.2	1	0.0	3.0	0
18	48	0	3	130	275	0	0	139	0	0.2	1	0.0	3.0	0
19	49	1	2	130	266	0	0	171	0	0.6	1	0.0	3.0	0
20	64	1	1	110	211	0	2	144	1	1.8	2	0.0	3.0	0
21	58	0	1	150	283	1	2	162	0	1.0	1	0.0	3.0	0
22	58	1	2	120	284	0	2	160	0	1.8	2	0.0	3.0	1
23	58	1	3	132	224	0	2	173	0	3.2	1	2.0	7.0	3
24	60	1	4	130	206	0	2	132	1	2.4	2	2.0	7.0	4
25	50	0	3	120	219	0	0	158	0	1.6	2	0.0	3.0	0
26	58	0	3	120	340	0	0	172	0	0.0	1	0.0	3.0	0
27	66	0	1	150	226	0	0	114	0	2.6	3	0.0	3.0	0
28	43	1	4	150	247	0	0	171	0	1.5	1	0.0	3.0	0
29	40	1	4	110	167	0	2	114	1	2.0	2	0.0	7.0	3
30	69	0	1	140	239	0	0	151	0	1.8	1	2.0	3.0	0
31	60	1	4	117	230	1	0	160	1	1.4	1	2.0	7.0	2
32	64	1	3	140	335	0	0	158	0	0.0	1	0.0	3.0	1
33	59	1	4	135	234	0	0	161	0	0.5	2	0.0	7.0	0
34	44	1	3	130	233	0	0	179	1	0.4	1	0.0	3.0	0
35	42	1	4	140	226	0	0	178	0	0.0	1	0.0	3.0	0
36	43	1	4	120	177	0	2	120	1	2.5	2	0.0	7.0	3
37	57	1	4	150	276	0	2	112	1	0.6	2	1.0	6.0	1
38	55	1	4	132	353	0	0	132	1	1.2	2	1.0	7.0	3
39	61	1	3	150	243	1	0	137	1	1.0	2	0.0	3.0	0
40	65	0	4	150	225	0	2	114	0	1.0	2	3.0	7.0	4
41	40	1	1	140	199	0	0	178	1	1.4	1	0.0	7.0	0
42	71	0	2	160	302	0	0	162	0	0.4	1	2.0	3.0	0
43	59	1	3	150	212	1	0	157	0	1.6	1	0.0	3.0	0
44	61	0	4	130	330	0	2	169	0	0.0	1	0.0	3.0	1
45	58	1	3	112	230	0	2	165	0	2.5	2	1.0	7.0	4
46	51	1	3	110	175	0	0	123	0	0.6	1	0.0	3.0	0
47	50	1	4	150	243	0	2	128	0	2.6	2	0.0	7.0	4
48	65	0	3	140	417	1	2	157	0	0.8	1	1.0	3.0	0
49	53	1	3	130	197	1	2	152	0	1.2	3	0.0	3.0	0
50	41	0	2	105	198	0	0	168	0	0.0	1	1.0	3.0	0
51	65	1	4	120	177	0	0	140	0	0.4	1	0.0	7.0	0
52	44	1	4	112	290	0	2	153	0	0.0	1	1.0	3.0	2
53	44	1	2	130	219	0	2	188	0	0.0	1	0.0	3.0	0
54	60	1	4	130	253	0	0	144	1	1.4	1	1.0	7.0	1
55	54	1	4	124	266	0	2	109	1	2.2	2	1.0	7.0	1
56	50	1	3	140	233	0	0	163	0	0.6	2	1.0	7.0	1
57	41	1	4	110	172	0	2	158	0	0.0	1	0.0	7.0	1
58	54	1	3	125	273	0	2	152	0	0.5	3	1.0	3.0	0
59	51	1	1	125	213	0	2	125	1	1.4	1	1.0	3.0	0
60	51	0	4	130	305	0	0	142	1	1.2	2	0.0	7.0	2
61	46	0	3	142	177	0	2	160	1	1.4	3	0.0	3.0	0
62	58	1	4	128	216	0	2	131	1	2.2	2	3.0	7.0	1
63	54	0	3	135	304	1	0	170	0	0.0	1	0.0	3.0	0
64	54	1	4	120	188	0	0	113	0	1.4	2	1.0	7.0	2
65	60	1	4	145	282	0	2	142	1	2.8	2	2.0	7.0	2
66	60	1	3	140	185	0	2	155	0	3.0	2	0.0	3.0	1
67	54	1	3	150	232	0	2	165	0	1.6	1	0.0	7.0	0
68	59	1	4	170	326	0	2	140	1	3.4	3	0.0	7.0	2
69	46	1	3	150	231	0	0	147	0	3.6	2	0.0	3.0	1
70	65	0	3	155	269	0	0	148	0	0.8	1	0.0	3.0	0
71	67	1	4	125	254	1	0	163	0	0.2	2	2.0	7.0	3

[reached 'max' / getOption("max.print") -- omitted 231 rows]

```

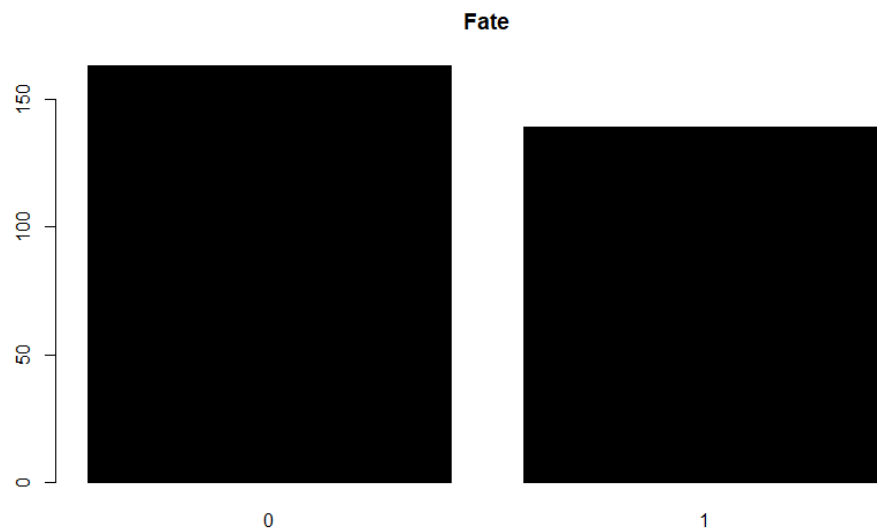
> hdata$ca[hdata$ca == 1.0]
factor(0)
Levels: 0.0 1.0 2.0 3.0

> typeof(hdata$ca)
[1] "integer"

> nrow(hdata)
[1] 302

> # Plotting Fate vs number of records
> hdata$num[hdata$num >= 1] <- 1 # Edit the fate to 0 and 1
> barplot(table(hdata$num), main="Fate", col="black")

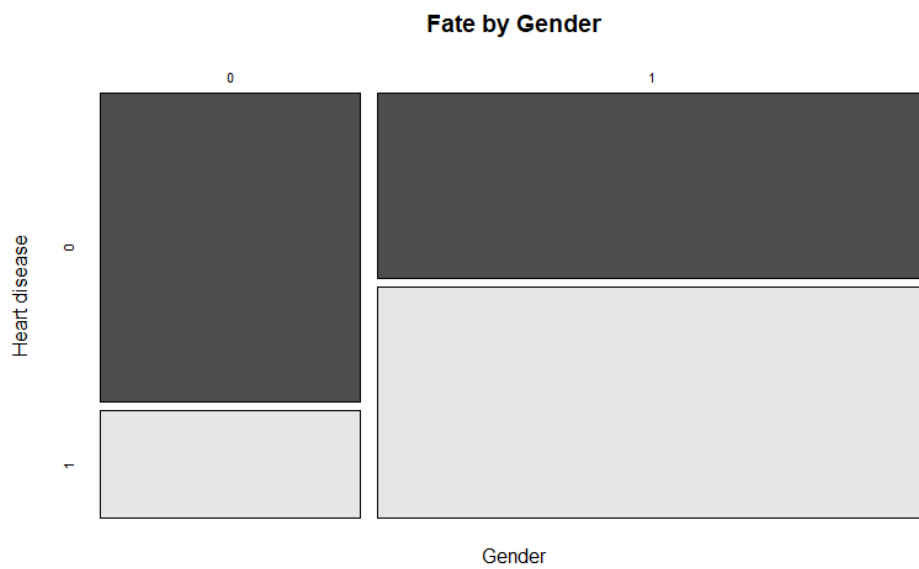
```



```

> # Plot Fate vs gender
> mosaicplot(hdata$sex ~ hdata$num, main="Fate by Gender",
+           shade=FALSE, color=TRUE, xlab="Gender", ylab="Heart disease")

```



```
> # Plot Fate vs Age
> mosaicplot(hdata$age ~ hdata$num,main="Fate by Age",
+           shade=FALSE,color=TRUE,xlab="Age", ylab="Heart disease")
```



```
> # Most important step, change the values of NA
> levels(hdata$thal)[levels(hdata$thal)=="?"]<-"3.0"
> # removal of additional NA
```

```
> hdata$thal
```

```
 [1] 3.0 7.0 3.0 3.0 3.0 3.0 3.0 7.0 7.0 6.0 3.0 6.0 7.0 7.0 3.0 7.0 3.0 3.0 3.0 3.0
 [30] 3.0 7.0 3.0 7.0 3.0 3.0 7.0 6.0 7.0 3.0 7.0 7.0 3.0 3.0 3.0 7.0 3.0 7.0 3.0 3.0
 [59] 3.0 7.0 3.0 7.0 3.0 7.0 7.0 3.0 7.0 7.0 3.0 3.0 7.0 7.0 6.0 3.0 3.0 7.0 3.0 3.0
 [88] 3.0 3.0 3.0 7.0 7.0 3.0 3.0 7.0 7.0 7.0 3.0 3.0 3.0 3.0 3.0 3.0 7.0 7.0 7.0 7.0
 [117] 3.0 7.0 7.0 7.0 7.0 3.0 7.0 3.0 3.0 7.0 7.0 3.0 3.0 7.0 7.0 3.0 3.0 3.0 3.0 7.0
 [146] 7.0 3.0 3.0 3.0 7.0 3.0 7.0 7.0 3.0 3.0 7.0 7.0 7.0 7.0 7.0 3.0 3.0 3.0 3.0 7.0
 [175] 7.0 7.0 6.0 3.0 3.0 7.0 7.0 3.0 7.0 3.0 7.0 3.0 7.0 6.0 7.0 7.0 3.0 7.0 7.0 3.0
 [204] 7.0 7.0 7.0 7.0 3.0 3.0 3.0 7.0 3.0 7.0 3.0 7.0 3.0 3.0 3.0 3.0 3.0 3.0 3.0 7.0
 [233] 3.0 3.0 3.0 7.0 7.0 3.0 3.0 3.0 3.0 3.0 3.0 3.0 3.0 7.0 3.0 7.0 3.0 6.0 7.0 7.0
 [262] 3.0 3.0 3.0 6.0 3.0 6.0 7.0 3.0 7.0 6.0 7.0 3.0 3.0 7.0 3.0 3.0 3.0 3.0 7.0 3.0
 [291] 3.0 6.0 7.0 3.0 3.0 6.0 7.0 7.0 7.0 7.0 3.0 3.0
```

```
Levels: 3.0 6.0 7.0
```

```
> table(hdata$thal)
```

```
3.0 6.0 7.0
168 17 117
```

```
> table(hdata$ca)
```

```
0.0 1.0 2.0 3.0
179 65 38 20
```

```
> library(caTools) # import library caTools
```

```
Warning message:
package 'caTools' was built under R version 3.6.3
```

```
> n<- sapply(hdata[, c(1)], mean) # get the average values
```

```
> set.seed(123) # generate a pseudo-random number
```

```
> v3 <- hdata[c(11:14),c(2,7:9)]
```

```
> v3
```

	sex	restecg	thalach	exang
11	0	2	153	0
12	1	2	142	1
13	1	0	173	0
14	1	0	162	0

```
> m<- sapply(v3,max)
```

```
> m
```

sex	restecg	thalach	exang
1	2	173	1

```
> set.seed(121)
```

```
> # Divide the dataset into 2/3 for training, and 1/3 for testing
```

```
> split = sample.split(hdata$num, SplitRatio = 2/3)
```

```

> train_hdata = subset(hdata, split == TRUE)
> test_hdata = subset(hdata, split == FALSE)
> # Apply linear regression for Fate vs age
> regressor=lm(formula = num~age, data=train_hdata)
> View(regressor)
> regressor

```

```

Call:
lm(formula = num ~ age, data = train_hdata)

```

```

Coefficients:
(Intercept)      age
   -0.33038      0.01453

```

```

> # Apply regression on test data
> hd_age_predict = predict(regressor, newdata=test_hdata)

```

```

> hd_age_predict

```

25	2	4	7	8	10	18	19	20
0.6430055	0.2652722	0.4977234	0.5848927	0.4977234	0.3669696	0.3814978	0.5994209	
0.3960260	0.2507440	0.6720619	0.5413081					
	33	34	42	43	45	47	52	59
61	65	68	70					
0.5267799	0.3088568	0.7011183	0.5267799	0.5122517	0.3960260	0.3088568	0.4105542	
0.3379132	0.5413081	0.5267799	0.6139491					
	73	75	78	79	83	84	85	86
	108	109	110	111				
0.6139491	0.6139491	0.3669696	0.5122517	0.6575337	0.4250824	0.3088568	0.3524414	
0.5558363	0.2362158	0.5558363	0.4831952					
	115	118	122	129	134	136	137	139
	141	144	147	149				
0.2652722	0.5848927	0.4105542	0.5703645	0.2943286	0.6865901	0.5703645	0.4105542	
0.5267799	0.5122517	0.2652722	0.5413081					
	155	156	158	168	169	173	175	176
	179	184	185	186				
0.6865901	0.4105542	0.5413081	0.1781030	0.3233850	0.5703645	0.4977234	0.4250824	
0.4396106	0.5413081	0.5848927	0.2798004					
	187	192	194	197	201	204	206	207
	214	217	219	220				
0.6284773	0.2943286	0.6575337	0.3233850	0.5994209	0.2943286	0.5122517	0.3960260	
0.4250824	0.3379132	0.5267799	0.2652722					
	225	226	229	230	232	236	237	240
	241	247	248	252				
0.1635748	0.3524414	0.6284773	0.4250824	0.3814978	0.4831952	0.3379132	0.2652722	
0.2652722	0.3524414	0.4250824	0.5994209					
	255	258	262	265	267	273	276	282
	284	285	288	292				
0.2798004	0.6865901	0.5413081	0.2798004	0.5267799	0.7011183	0.6284773	0.4686670	
0.5558363	0.5122517	0.4831952	0.3088568					
	293	296	301	302				
0.5848927	0.5267799	0.4977234	0.2216876					

```

> # Round the values of fate in prediction
> round_age=hd_age_predict
> r=round(round_age)
> r

```

```

  2   4   7   8  10  18  19  20  25  29  30  31  33  34  42  43  45  47  52  59  61
65  68  70  73  75  78  79  83  84
1   1   0   0   1   0   0   1   0   0   1   1   1   0   1   1   1   0   0   0   0
85  86 108 109 110 111 115 118 122 129 134 136 137 139 141 144 147 149 155 156 158
168 169 173 175 176 179 184 185 186
0   0   1   0   0   0   1   1   0   1   0   1   1   0   1   1   0   1   1   0   1
187 192 194 197 201 204 206 207 214 217 219 220 225 226 229 230 232 236 237 240 241
247 248 252 255 258 262 265 267 273
1   0   1   0   1   0   1   0   0   0   1   0   0   0   1   0   0   0   0   0   0
0   0   1   0   1   1   0   1   1   1   1   0   0   0   1   0   0   0   0   0   0
276 282 284 285 288 292 293 296 301 302
1   0   1   1   0   0   1   1   0   0

```

```
> table(r,test_hdata$num)
```

```

r      0  1
0 34 20
1 20 26

```

```
> library(e1071)
```

```
> library(caret)
```

```
> typeof(r)
[1] "double"
```

```
> levels(r)
NULL
```

```
> levels(test_hdata$num)
NULL
```

```
> str(r)
Named num [1:100] 1 0 0 1 0 0 0 1 0 0 ...
- attr(*, "names")= chr [1:100] "2" "4" "7" "8" ...
```

```
> r1 = as.data.frame(r)
```

```

> r1
  r
2  1
4  0
7  0
8  1
10 0
18 0
19 0
20 1
25 0
29 0
30 1
31 1
33 1
34 0
42 1
43 1
45 1
47 0
52 0
59 0

```


61	0
65	1
68	1
70	1
73	1
75	1
78	0
79	1
83	1
84	0
85	0
86	0
108	1
109	0
110	1
111	0
115	0
118	1
122	0
129	1
134	0
136	1
137	1
139	0
141	1
144	1
147	0
149	1
155	1
156	0
158	1
168	0
169	0
173	1
175	0
176	0
179	0
184	1
185	1
186	0
187	1
192	0
194	1
197	0
201	1
204	0
206	1
207	0
214	0
217	0
219	1
220	0
225	0
226	0
229	1
230	0
232	0
236	0
237	0
240	0
241	0
247	0
248	0
252	1

```

255 0
258 1
262 1
265 0
267 1
273 1
276 1
282 0
284 1
285 1
288 0
292 0
293 1
296 1
301 0
302 0
> df1=confusionMatrix(as.factor(r1$r),as.factor(test_hdata$num))

```

```

> df1
Confusion Matrix and Statistics

```

```

      Reference
Prediction 0  1
      0 34 20
      1 20 26

      Accuracy : 0.6
      95% CI : (0.4972, 0.6967)
No Information Rate : 0.54
P-Value [Acc > NIR] : 0.1347

      Kappa : 0.1948

McNemar's Test P-Value : 1.0000

      Sensitivity : 0.6296
      Specificity : 0.5652
      Pos Pred Value : 0.6296
      Neg Pred Value : 0.5652
      Prevalence : 0.5400
      Detection Rate : 0.3400
      Detection Prevalence : 0.5400
      Balanced Accuracy : 0.5974

      'Positive' Class : 0

```

```

>

```