

```
install.packages("tidyverse")
install.packages("e1071")
install.packages("leaps")
install.packages("tree")
library("tidyverse")
library("e1071")
library("leaps")
library("tree")
```

Installing package into ‘/usr/local/lib/R/site-library’
(as ‘lib’ is unspecified)

Installing package into ‘/usr/local/lib/R/site-library’
(as ‘lib’ is unspecified)

also installing the dependency ‘proxy’

Installing package into ‘/usr/local/lib/R/site-library’
(as ‘lib’ is unspecified)

Installing package into ‘/usr/local/lib/R/site-library’
(as ‘lib’ is unspecified)

Warning message in system("timedatectl", intern = TRUE):
“running command ‘timedatectl’ had status 1”

— Attaching packages —

tidyverse 1.3.2 —

✓ ggplot2 3.4.0	✓ purrr 0.3.5
✓ tibble 3.1.8	✓ dplyr 1.0.10
✓ tidyr 1.2.1	✓ stringr 1.4.1
✓ readr 2.1.3	✓ forcats 0.5.2

— Conflicts —

tidyverse_conflicts() —

* dplyr::filter() masks stats::filter()
* dplyr::lag() masks stats::lag()

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

Installing package into ‘/usr/local/lib/R/site-library’
(as ‘lib’ is unspecified)

also installing the dependencies ‘iterators’, ‘foreach’, ‘shape’,
‘Rcpp’, ‘RcppEigen’

Loading required package: Matrix

Attaching package: 'Matrix'

The following objects are masked from 'package:tidyr':

expand, pack, unpack

Loaded glmnet 4.1-6

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

Installing package into '/usr/local/lib/R/site-library'
(as 'lib' is unspecified)

Attaching package: 'MASS'

The following object is masked from 'package:dplyr':

select

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

Installing package into '/usr/local/lib/R/site-library'
(as 'lib' is unspecified)

randomForest 4.7-1.1

Type rfNews() to see new features/changes/bug fixes.

Attaching package: 'randomForest'

The following object is masked from 'package:dplyr':

combine

The following object is masked from 'package:ggplot2':

margin

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

Installing package into ‘/usr/local/lib/R/site-library’
(as ‘lib’ is unspecified)

Attaching package: ‘kernlab’

The following object is masked from ‘package:purrr’:

cross

The following object is masked from ‘package:ggplot2’:

alpha

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

Installing package into ‘/usr/local/lib/R/site-library’
(as ‘lib’ is unspecified)

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

Installing package into ‘/usr/local/lib/R/site-library’
(as ‘lib’ is unspecified)

Attaching package: ‘pls’

The following object is masked from ‘package:stats’:

loadings

```
heart<-read.csv("heart.csv")
```

```
heart<-heart %>%
mutate(sex=as.factor(sex),cp=as.factor(cp),fbs=as.factor(fbs),restecg=
as.factor(restecg),exang=as.factor(exang),slope=as.factor(slope),ca=as
.factor(ca),thal=as.factor(thal),target=as.factor(target))
```

heart

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope
ca thal											
1 2	52	1	0	125	212	0	1	168	0	1.0	2
2 3											
2 0	53	1	0	140	203	1	0	155	1	3.1	0
3 0											
3 0	70	1	0	145	174	0	1	125	1	2.6	0
4 1											
4 1	61	1	0	148	203	0	1	161	0	0.0	2
5 3											
5 3	62	0	0	138	294	1	1	106	0	1.9	1
6 0											
6 0	58	0	0	100	248	0	0	122	0	1.0	1
7 3											
7 3	58	1	0	114	318	0	2	140	0	4.4	0
8 1											
8 1	55	1	0	160	289	0	0	145	1	0.8	1
9 0											
9 0	46	1	0	120	249	0	0	144	0	0.8	2
10 2											
10 2	54	1	0	122	286	0	0	116	1	3.2	1
11 0											
11 0	71	0	0	112	149	0	1	125	0	1.6	1
12 0											
12 0	43	0	0	132	341	1	0	136	1	3.0	1
13 0											
13 0	34	0	1	118	210	0	1	192	0	0.7	2
14 3											
14 3	51	1	0	140	298	0	1	122	1	4.2	1
15 0											
15 0	52	1	0	128	204	1	1	156	1	1.0	1
16 0											
16 0	34	0	1	118	210	0	1	192	0	0.7	2
17 1											
17 1	51	0	2	140	308	0	0	142	0	1.5	2
18 1											
18 1	54	1	0	124	266	0	0	109	1	2.2	1
19 0											
19 0	50	0	1	120	244	0	1	162	0	1.1	2
20 0											
20 0	58	1	2	140	211	1	0	165	0	0.0	2
21 0											
21 0	60	1	2	140	185	0	0	155	0	3.0	1

22 2 2	67	0	0	106	223	0	1	142	0	0.3	2
23 0 2	45	1	0	104	208	0	0	148	1	3.0	1
24 0 2	63	0	2	135	252	0	0	172	0	0.0	2
25 0 2	42	0	2	120	209	0	1	173	0	0.0	1
26 0 3	61	0	0	145	307	0	0	146	1	1.0	1
27 0 2	44	1	2	130	233	0	1	179	1	0.4	2
28 2 2	58	0	1	136	319	1	0	152	0	0.0	2
29 1 1	56	1	2	130	256	1	0	142	1	0.6	1
30 0 2	55	0	0	180	327	0	2	117	1	3.4	1
⋮ ⋮ ⋮	⋮ ⋮ ⋮	⋮ ⋮ ⋮	⋮ ⋮ ⋮	⋮ ⋮ ⋮	⋮ ⋮ ⋮	⋮ ⋮ ⋮	⋮ ⋮ ⋮	⋮ ⋮ ⋮	⋮ ⋮ ⋮	⋮ ⋮ ⋮	⋮ ⋮ ⋮
996 0 3	44	1	1	120	263	0	1	173	0	0.0	2
997 2 3	56	0	0	134	409	0	0	150	1	1.9	1
998 1 3	54	1	0	120	188	0	1	113	0	1.4	1
999 0 1	42	1	0	136	315	0	1	125	1	1.8	1
1000 2 3	67	1	0	125	254	1	1	163	0	0.2	1
1001 2 1	64	1	0	145	212	0	0	132	0	2.0	1
1002 0 2	42	1	0	140	226	0	1	178	0	0.0	2
1003 1 2	66	1	0	112	212	0	0	132	1	0.1	2
1004 3 3	52	1	0	108	233	1	1	147	0	0.1	2
1005 1 2	51	0	2	140	308	0	0	142	0	1.5	2
1006 1 3	55	0	0	128	205	0	2	130	1	2.0	1
1007 0 2	58	1	2	140	211	1	0	165	0	0.0	2
1008 0 3	56	1	3	120	193	0	0	162	0	1.9	1
1009 0 2	42	1	1	120	295	0	1	162	0	0.0	2
1010 0 3	40	1	0	152	223	0	1	181	0	0.0	2

1011	51	1	0	140	299	0	1	173	1	1.6	2
0	3										
1012	45	1	1	128	308	0	0	170	0	0.0	2
0	2										
1013	48	1	1	110	229	0	1	168	0	1.0	0
0	3										
1014	58	1	0	114	318	0	2	140	0	4.4	0
3	1										
1015	44	0	2	108	141	0	1	175	0	0.6	1
0	2										
1016	58	1	0	128	216	0	0	131	1	2.2	1
3	3										
1017	65	1	3	138	282	1	0	174	0	1.4	1
1	2										
1018	53	1	0	123	282	0	1	95	1	2.0	1
2	3										
1019	41	1	0	110	172	0	0	158	0	0.0	2
0	3										
1020	47	1	0	112	204	0	1	143	0	0.1	2
0	2										
1021	59	1	1	140	221	0	1	164	1	0.0	2
0	2										
1022	60	1	0	125	258	0	0	141	1	2.8	1
1	3										
1023	47	1	0	110	275	0	0	118	1	1.0	1
1	2										
1024	50	0	0	110	254	0	0	159	0	0.0	2
0	2										
1025	54	1	0	120	188	0	1	113	0	1.4	1
1	3										

	target
1	0
2	0
3	0
4	0
5	0
6	1
7	0
8	0
9	0
10	0
11	1
12	0
13	1
14	0
15	0
16	1
17	1
18	0
19	1

ca	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	
1	58	0	0	100	248	0	0	122	0	1.0	1	0
2	71	0	0	112	149	0	1	125	0	1.6	1	0

3 2	34	0	1	118	210	0	1	192	0	0.7	2	0
4 2	34	0	1	118	210	0	1	192	0	0.7	2	0
5 2	51	0	2	140	308	0	0	142	0	1.5	2	1
6 2	50	0	1	120	244	0	1	162	0	1.1	2	0
7 2	58	1	2	140	211	1	0	165	0	0.0	2	0
8 2	67	0	0	106	223	0	1	142	0	0.3	2	2
9 2	45	1	0	104	208	0	0	148	1	3.0	1	0
10 2	63	0	2	135	252	0	0	172	0	0.0	2	0
11 2	42	0	2	120	209	0	1	173	0	0.0	1	0
12 2	44	1	2	130	233	0	1	179	1	0.4	2	0
13 2	50	0	1	120	244	0	1	162	0	1.1	2	0
14 2	50	1	2	129	196	0	1	163	0	0.0	2	0
15 2	51	1	3	125	213	0	0	125	1	1.4	2	1
16 2	59	1	0	138	271	0	0	182	0	0.0	2	0
17 3	64	1	0	128	263	0	1	105	1	0.2	1	1
18 2	65	0	2	160	360	0	0	151	0	0.8	2	0
19 3	54	1	2	120	258	0	0	147	0	0.4	1	0
20 2	55	0	1	132	342	0	1	166	0	1.2	2	0
21 2	42	1	0	140	226	0	1	178	0	0.0	2	0
22 1	41	1	1	135	203	0	1	132	0	0.0	1	0
23 2	66	0	2	146	278	0	0	152	0	0.0	1	1
24 2	58	0	3	150	283	1	0	162	0	1.0	2	0
25 2	38	1	2	138	175	0	1	173	0	0.0	2	4
26 3	56	1	3	120	193	0	0	162	0	1.9	1	0
27 2	48	1	1	130	245	0	0	180	0	0.2	1	0

282	292	1	1	130	204	0	0	202	0	0.0	2	0
292	66	0	2	146	278	0	0	152	0	0.0	1	1
302	59	1	2	150	212	1	1	157	0	1.6	2	0
⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮
4972	44	0	2	108	141	0	1	175	0	0.6	1	0
4982	76	0	2	140	197	0	2	116	0	1.1	1	0
4992	60	0	2	120	178	1	1	96	0	0.0	2	0
5002	52	1	1	120	325	0	1	172	0	0.2	2	0
5012	38	1	2	138	175	0	1	173	0	0.0	2	4
5023	52	1	2	172	199	1	1	162	0	0.5	2	0
5031	52	1	3	118	186	0	0	190	0	0.0	1	0
5042	51	1	2	125	245	1	0	166	0	2.4	1	0
5053	43	1	0	110	211	0	1	161	0	0.0	2	0
5062	52	1	1	128	205	1	1	184	0	0.0	2	0
5071	57	1	0	140	192	0	1	148	0	0.4	1	0
5082	67	0	0	106	223	0	1	142	0	0.3	2	2
5093	64	1	0	128	263	0	1	105	1	0.2	1	1
5103	59	1	0	135	234	0	1	161	0	0.5	1	0
5113	62	1	2	130	231	0	1	146	0	1.8	1	3
5122	71	0	1	160	302	0	1	162	0	0.4	2	2
5132	56	1	1	120	236	0	1	178	0	0.8	2	0
5142	50	0	0	110	254	0	0	159	0	0.0	2	0
5153	44	1	1	120	263	0	1	173	0	0.0	2	0
5162	42	1	0	140	226	0	1	178	0	0.0	2	0
5173	52	1	0	108	233	1	1	147	0	0.1	2	3

518	51	0	2	140	308	0	0	142	0	1.5	2	1
2												
519	58	1	2	140	211	1	0	165	0	0.0	2	0
2												
520	56	1	3	120	193	0	0	162	0	1.9	1	0
3												
521	42	1	1	120	295	0	1	162	0	0.0	2	0
2												
522	45	1	1	128	308	0	0	170	0	0.0	2	0
2												
523	44	0	2	108	141	0	1	175	0	0.6	1	0
2												
524	47	1	0	112	204	0	1	143	0	0.1	2	0
2												
525	59	1	1	140	221	0	1	164	1	0.0	2	0
2												
526	50	0	0	110	254	0	0	159	0	0.0	2	0
2												

	target
1	1
2	1
3	1
4	1
5	1
6	1
7	1
8	1
9	1
10	1
11	1
12	1
13	1
14	1
15	1
16	1
17	1
18	1
19	1
20	1
21	1
22	1
23	1
24	1
25	1
26	1
27	1
28	1
29	1
30	1
⋮	⋮

497	1
498	1
499	1
500	1
501	1
502	1
503	1
504	1
505	1
506	1
507	1
508	1
509	1
510	1
511	1
512	1
513	1
514	1
515	1
516	1
517	1
518	1
519	1
520	1
521	1
522	1
523	1
524	1
525	1
526	1

```
heart %>% filter(target==0)
```

ca	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	
13	52	1	0	125	212	0	1	168	0	1.0	2	2
23	53	1	0	140	203	1	0	155	1	3.1	0	0
33	70	1	0	145	174	0	1	125	1	2.6	0	0
43	61	1	0	148	203	0	1	161	0	0.0	2	1
52	62	0	0	138	294	1	1	106	0	1.9	1	3
61	58	1	0	114	318	0	2	140	0	4.4	0	3
73	55	1	0	160	289	0	0	145	1	0.8	1	1
83	46	1	0	120	249	0	0	144	0	0.8	2	0

9 2	54	1	0	122	286	0	0	116	1	3.2	1	2
10 3	43	0	0	132	341	1	0	136	1	3.0	1	0
11 3	51	1	0	140	298	0	1	122	1	4.2	1	3
12 0	52	1	0	128	204	1	1	156	1	1.0	1	0
13 3	54	1	0	124	266	0	0	109	1	2.2	1	1
14 2	60	1	2	140	185	0	0	155	0	3.0	1	0
15 3	61	0	0	145	307	0	0	146	1	1.0	1	0
16 2	58	0	1	136	319	1	0	152	0	0.0	2	2
17 1	56	1	2	130	256	1	0	142	1	0.6	1	1
18 2	55	0	0	180	327	0	2	117	1	3.4	1	0
19 1	44	1	0	120	169	0	1	144	1	2.8	0	0
20 3	57	1	0	130	131	0	1	115	1	1.2	1	1
21 3	70	1	2	160	269	0	1	112	1	2.9	1	1
22 2	46	1	2	150	231	0	1	147	0	3.6	1	0
23 3	57	1	2	128	229	0	0	150	0	0.4	1	1
24 2	61	0	0	130	330	0	0	169	0	0.0	2	0
25 3	46	1	0	120	249	0	0	144	0	0.8	2	0
26 3	66	0	0	178	228	1	1	165	1	1.0	1	2
27 3	60	1	0	117	230	1	1	160	1	1.4	2	2
28 3	57	0	0	140	241	0	1	123	1	0.2	1	0
29 3	49	1	2	120	188	0	1	139	0	2.0	1	3
30 3	55	1	0	140	217	0	1	111	1	5.6	0	0
⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮
470 3	53	1	0	140	203	1	0	155	1	3.1	0	0
471 3	39	1	0	118	219	0	1	140	0	1.2	1	0

[illegible]

497	60	1	0	125	258	0	0	141	1	2.8	1	1
3												
498	47	1	0	110	275	0	0	118	1	1.0	1	1
2												
499	54	1	0	120	188	0	1	113	0	1.4	1	1
3												

	target
--	--------

1	0
2	0
3	0
4	0
5	0
6	0
7	0
8	0
9	0
10	0
11	0
12	0
13	0
14	0
15	0
16	0
17	0
18	0
19	0
20	0
21	0
22	0
23	0
24	0
25	0
26	0
27	0
28	0
29	0
30	0
⋮	⋮
470	0
471	0
472	0
473	0
474	0
475	0
476	0
477	0
478	0
479	0
480	0
481	0

482 0
 483 0
 484 0
 485 0
 486 0
 487 0
 488 0
 489 0
 490 0
 491 0
 492 0
 493 0
 494 0
 495 0
 496 0
 497 0
 498 0
 499 0

summary(heart)

age	sex	cp	trestbps	chol	fbs
restecg					
Min. :29.00	0:312	0:497	Min. : 94.0	Min. :126	0:872
1st Qu.:48.00	1:713	1:167	1st Qu.:120.0	1st Qu.:211	1:153
Median :56.00		2:284	Median :130.0	Median :240	
Mean :54.43		3: 77	Mean :131.6	Mean :246	
3rd Qu.:61.00			3rd Qu.:140.0	3rd Qu.:275	
Max. :77.00			Max. :200.0	Max. :564	
thalach	exang	oldpeak	slope	ca	thal
target					
Min. : 71.0	0:680	Min. :0.000	0: 74	0:578	0: 7 0:499
1st Qu.:132.0	1:345	1st Qu.:0.000	1:482	1:226	1: 64 1:526
Median :152.0		Median :0.800	2:469	2:134	2:544
Mean :149.1		Mean :1.072		3: 69	3:410
3rd Qu.:166.0		3rd Qu.:1.800		4: 18	
Max. :202.0		Max. :6.200			

```

normalize <- function(x) {
  return ((x - min(x)) / (max(x) - min(x)))
}
heartmm <- as.data.frame(model.matrix(~.-1,heart))
# we are going to normalize everything
heart <- as.data.frame(lapply(heartmm, normalize))

set.seed(385720)

train<-sample(dim(heart)[1],dim(heart)[1]*0.7)

test<--train

heart.train<-heart[train,]

heart.test<-heart[test,]

head(heart)

```

	age	sex0	sex1	cp1	cp2	cp3	trestbps	chol	fbs1	restecg1	...
slope1											
1	0.4791667	0	1	0	0	0	0.29245283	0.1963470	0	1	...
0											
2	0.5000000	0	1	0	0	0	0.43396226	0.1757991	1	0	...
0											
3	0.8541667	0	1	0	0	0	0.48113208	0.1095890	0	1	...
0											
4	0.6666667	0	1	0	0	0	0.50943396	0.1757991	0	1	...
0											
5	0.6875000	1	0	0	0	0	0.41509434	0.3835616	1	1	...
1											
6	0.6041667	1	0	0	0	0	0.05660377	0.2785388	0	0	...
1											
slope2	ca1	ca2	ca3	ca4	thal1	thal2	thal3	target1			
1	1	0	1	0	0	0	1	0			
2	0	0	0	0	0	0	1	0			
3	0	0	0	0	0	0	1	0			
4	1	1	0	0	0	0	1	0			
5	0	0	0	1	0	0	1	0			
6	0	0	0	0	0	0	1	0			

Logistic Regression

```

logmod <- glm(target1 ~ ., data = heart.train, family = "binomial")
summary(logmod)
logmodselect = step(logmod, direction = "backward")

```

Call:

```
glm(formula = target1 ~ ., family = "binomial", data = heart.train)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-2.87521	-0.29896	0.07339	0.41120	3.10889

Coefficients: (1 not defined because of singularities)

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-1.0625	2.6834	-0.396	0.692141
age	1.0789	0.8296	1.300	0.193439
sex0	2.0465	0.3846	5.320	1.04e-07 ***
sex1	NA	NA	NA	NA
cp1	0.8673	0.3885	2.232	0.025591 *
cp2	1.8740	0.3457	5.421	5.93e-08 ***
cp3	2.6358	0.4723	5.580	2.40e-08 ***
trestbps	-2.9214	0.8638	-3.382	0.000719 ***
chol	-3.1810	1.2239	-2.599	0.009346 **
fbs1	0.1954	0.4047	0.483	0.629268
restecg1	0.2423	0.2727	0.889	0.374227
restecg2	-0.4054	1.8375	-0.221	0.825385
thalach	2.4574	1.0658	2.306	0.021123 *
exang1	-0.7879	0.3101	-2.541	0.011064 *
oldpeak	-3.6010	1.0194	-3.532	0.000412 ***
slope1	-0.9724	0.5796	-1.678	0.093416 .
slope2	0.4652	0.6165	0.755	0.450470
ca1	-2.2040	0.3556	-6.198	5.73e-10 ***
ca2	-3.4790	0.5198	-6.693	2.19e-11 ***
ca3	-2.3177	0.6775	-3.421	0.000624 ***
ca4	0.8779	1.3773	0.637	0.523873
thal1	2.1875	2.5646	0.853	0.393678
thal2	1.9098	2.5170	0.759	0.447994
thal3	0.5836	2.5190	0.232	0.816801

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

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 993.66 on 716 degrees of freedom
 Residual deviance: 409.76 on 694 degrees of freedom
 AIC: 455.76

Number of Fisher Scoring iterations: 6

Start: AIC=455.76

target1 ~ age + sex0 + sex1 + cp1 + cp2 + cp3 + trestbps + chol +
 fbs1 + restecg1 + restecg2 + thalach + exang1 + oldpeak +
 slope1 + slope2 + ca1 + ca2 + ca3 + ca4 + thal1 + thal2 +
 thal3

Step: AIC=455.76

target1 ~ age + sex0 + cp1 + cp2 + cp3 + trestbps + chol + fbs1 +
 restecg1 + restecg2 + thalach + exang1 + oldpeak + slope1 +

slope2 + ca1 + ca2 + ca3 + ca4 + thal1 + thal2 + thal3

	Df	Deviance	AIC
- restecg2	1	409.81	453.81
- thal3	1	409.82	453.82
- fbs1	1	410.00	454.00
- ca4	1	410.17	454.17
- slope2	1	410.32	454.32
- thal2	1	410.39	454.39
- restecg1	1	410.56	454.56
- thal1	1	410.59	454.59
- age	1	411.47	455.47
<none>		409.76	455.76
- slope1	1	412.64	456.64
- cp1	1	414.91	458.91
- thalach	1	415.35	459.35
- exang1	1	416.20	460.20
- chol	1	416.48	460.48
- trestbps	1	421.78	465.78
- oldpeak	1	423.53	467.53
- ca3	1	424.73	468.73
- cp2	1	441.93	485.93
- sex0	1	442.06	486.06
- cp3	1	445.30	489.30
- ca1	1	452.81	496.81
- ca2	1	464.99	508.99

Step: AIC=453.81

target1 ~ age + sex0 + cp1 + cp2 + cp3 + trestbps + chol + fbs1 +
restecg1 + thalach + exang1 + oldpeak + slope1 + slope2 +
ca1 + ca2 + ca3 + ca4 + thal1 + thal2 + thal3

	Df	Deviance	AIC
- thal3	1	409.87	451.87
- fbs1	1	410.05	452.05
- ca4	1	410.22	452.22
- slope2	1	410.37	452.37
- thal2	1	410.44	452.44
- thal1	1	410.63	452.63
- restecg1	1	410.65	452.65
- age	1	411.50	453.50
<none>		409.81	453.81
- slope1	1	412.68	454.68
- cp1	1	414.97	456.97
- thalach	1	415.43	457.43
- exang1	1	416.23	458.23
- chol	1	416.48	458.48
- trestbps	1	421.85	463.85
- oldpeak	1	423.91	465.91
- ca3	1	424.92	466.92

- cp2	1	442.00	484.00
- sex0	1	442.18	484.18
- cp3	1	445.46	487.46
- ca1	1	452.91	494.91
- ca2	1	465.01	507.01

Step: AIC=451.87

target1 ~ age + sex0 + cp1 + cp2 + cp3 + trestbps + chol + fbs1 +
restecg1 + thalach + exang1 + oldpeak + slope1 + slope2 +
ca1 + ca2 + ca3 + ca4 + thal1 + thal2

	Df	Deviance	AIC
- fbs1	1	410.09	450.09
- ca4	1	410.28	450.28
- slope2	1	410.42	450.42
- restecg1	1	410.70	450.70
- age	1	411.56	451.56
<none>		409.87	451.87
- slope1	1	412.74	452.74
- cp1	1	415.05	455.05
- thalach	1	415.50	455.50
- exang1	1	416.31	456.31
- chol	1	416.49	456.49
- thal1	1	416.97	456.97
- trestbps	1	421.87	461.87
- oldpeak	1	423.93	463.93
- ca3	1	424.93	464.93
- thal2	1	431.44	471.44
- cp2	1	442.02	482.02
- sex0	1	442.23	482.23
- cp3	1	445.55	485.55
- ca1	1	452.97	492.97
- ca2	1	465.02	505.02

Step: AIC=450.09

target1 ~ age + sex0 + cp1 + cp2 + cp3 + trestbps + chol + restecg1 +
thalach + exang1 + oldpeak + slope1 + slope2 + ca1 + ca2 +
ca3 + ca4 + thal1 + thal2

	Df	Deviance	AIC
- ca4	1	410.56	448.56
- slope2	1	410.61	448.61
- restecg1	1	410.90	448.90
- age	1	411.75	449.75
<none>		410.09	450.09
- slope1	1	413.02	451.02
- cp1	1	415.56	453.56
- thalach	1	415.84	453.84
- exang1	1	416.36	454.36
- chol	1	416.58	454.58

- thal1	1	417.90	455.90
- trestbps	1	421.87	459.87
- oldpeak	1	424.62	462.62
- ca3	1	424.93	462.93
- thal2	1	431.77	469.77
- sex0	1	442.26	480.26
- cp2	1	444.41	482.41
- cp3	1	446.62	484.62
- ca1	1	453.18	491.18
- ca2	1	465.25	503.25

Step: AIC=448.56

target1 ~ age + sex0 + cp1 + cp2 + cp3 + trestbps + chol + restecg1 +
thalach + exang1 + oldpeak + slope1 + slope2 + ca1 + ca2 +
ca3 + thal1 + thal2

	Df	Deviance	AIC
- slope2	1	411.05	447.05
- restecg1	1	411.39	447.39
- age	1	412.12	448.12
<none>		410.56	448.56
- slope1	1	413.43	449.43
- thalach	1	416.19	452.19
- cp1	1	416.30	452.30
- exang1	1	416.74	452.74
- chol	1	417.12	453.12
- thal1	1	418.22	454.22
- trestbps	1	422.24	458.24
- ca3	1	425.54	461.54
- oldpeak	1	425.66	461.66
- thal2	1	431.98	467.98
- sex0	1	442.36	478.36
- cp2	1	445.01	481.01
- cp3	1	446.88	482.88
- ca1	1	454.06	490.06
- ca2	1	465.46	501.46

Step: AIC=447.05

target1 ~ age + sex0 + cp1 + cp2 + cp3 + trestbps + chol + restecg1 +
thalach + exang1 + oldpeak + slope1 + ca1 + ca2 + ca3 + thal1 +
thal2

	Df	Deviance	AIC
- restecg1	1	412.03	446.03
- age	1	412.59	446.59
<none>		411.05	447.05
- cp1	1	416.78	450.78
- thalach	1	416.88	450.88
- exang1	1	417.20	451.20
- chol	1	417.57	451.57

- thal1	1	418.31	452.31
- trestbps	1	422.61	456.61
- ca3	1	425.76	459.76
- slope1	1	430.45	464.45
- thal2	1	432.47	466.47
- oldpeak	1	433.78	467.78
- sex0	1	442.66	476.66
- cp2	1	445.52	479.52
- cp3	1	447.47	481.47
- ca1	1	454.11	488.11
- ca2	1	465.58	499.58

Step: AIC=446.03

target1 ~ age + sex0 + cp1 + cp2 + cp3 + trestbps + chol + thalach +
exang1 + oldpeak + slope1 + ca1 + ca2 + ca3 + thal1 + thal2

	Df	Deviance	AIC
- age	1	413.43	445.43
<none>		412.03	446.03
- cp1	1	417.64	449.64
- thalach	1	417.87	449.87
- exang1	1	418.14	450.14
- thal1	1	419.23	451.23
- chol	1	420.02	452.02
- trestbps	1	424.15	456.15
- ca3	1	427.32	459.32
- thal2	1	432.61	464.61
- slope1	1	433.14	465.14
- oldpeak	1	434.60	466.60
- sex0	1	445.27	477.27
- cp2	1	446.44	478.44
- cp3	1	447.66	479.66
- ca1	1	457.43	489.43
- ca2	1	466.51	498.51

Step: AIC=445.43

target1 ~ sex0 + cp1 + cp2 + cp3 + trestbps + chol + thalach +
exang1 + oldpeak + slope1 + ca1 + ca2 + ca3 + thal1 + thal2

	Df	Deviance	AIC
<none>		413.43	445.43
- thalach	1	417.94	447.94
- cp1	1	419.53	449.53
- exang1	1	419.92	449.92
- chol	1	420.53	450.53
- thal1	1	421.01	451.01
- trestbps	1	424.23	454.23
- ca3	1	427.71	457.71
- slope1	1	433.69	463.69
- thal2	1	433.87	463.87

```
- oldpeak    1    437.45 467.45
- sex0       1    447.91 477.91
- cp2        1    448.62 478.62
- cp3        1    450.00 480.00
- ca1        1    457.43 487.43
- ca2        1    467.52 497.52
```

```
colnames(heart)
```

```
[1] "age"      "sex0"     "sex1"     "cp1"      "cp2"      "cp3"
[7] "trestbps" "chol"     "fbs1"     "restecg1" "restecg2" "thalach"
[13] "exang1"   "oldpeak"  "slope1"   "slope2"   "ca1"      "ca2"
[19] "ca3"      "ca4"      "thal1"    "thal2"    "thal3"    "target1"
```

```
bestmod <- glm(target1 ~ sex0 + cp1 + cp2 + cp3 + trestbps + chol +
  thalach +
  exang1 + oldpeak + slope1 + ca1 + ca2 + ca3 + thal1 + thal2, data
= heart.train, family = "binomial")
```

```
logpred = predict(bestmod, newdata = heart.test, type = "response")
logpred = ifelse(logpred > 0.5, 1, 0)
table(logpred, heart.test$target1)
logRegaccuracy<-mean(logpred == heart.test$target1)
```

```
logpred    0    1
          0 119  21
          1  29 139
```

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

```
Installing package into '/usr/local/lib/R/site-library'
(as 'lib' is unspecified)
```

```
also installing the dependencies 'listenv', 'parallelly', 'future',
'globals', 'future.apply', 'numDeriv', 'progressr', 'SQUAREM', 'lava',
'prodlim', 'clock', 'gower', 'hardhat', 'ipred', 'timeDate',
'ModelMetrics', 'plyr', 'pROC', 'recipes', 'reshape2'
```

```
Loading required package: lattice
```

```
Attaching package: 'caret'
```

```
The following object is masked from 'package:pls':
```

```
R2
```

The following object is masked from 'package:purrr':

lift

```
cm <- confusionMatrix(data = as.factor(logpred), reference =
as.factor(heart.test$target1),positive='1')

draw_confusion_matrix <- function(cm) {

  total <- sum(cm$table)
  res <- as.numeric(cm$table)

  # Generate color gradients. Palettes come from RColorBrewer.
  greenPalette <-
c("#F7FCF5", "#E5F5E0", "#C7E9C0", "#A1D99B", "#74C476", "#41AB5D", "#238B45",
", "#006D2C", "#00441B")
  redPalette <-
c("#FFF5F0", "#FEE0D2", "#FCBBA1", "#FC9272", "#FB6A4A", "#EF3B2C", "#CB181D",
", "#A50F15", "#67000D")
  getColor <- function (greenOrRed = "green", amount = 0) {
    if (amount == 0)
      return("#FFFFFF")
    palette <- greenPalette
    if (greenOrRed == "red")
      palette <- redPalette
    colorRampPalette(palette)(100)[10 + ceiling(90 * amount / total)]
  }

  # set the basic layout
  layout(matrix(c(1,1,2)))
  par(mar=c(2,2,2,2))
  plot(c(100, 345), c(300, 450), type = "n", xlab="", ylab="",
xaxt='n', yaxt='n')
  title('CONFUSION MATRIX', cex.main=2)

  # create the matrix
  classes = colnames(cm$table)
  rect(150, 430, 240, 370, col=getColor("green", res[1]))
  text(195, 435, classes[1], cex=1.2)
  rect(250, 430, 340, 370, col=getColor("red", res[3]))
  text(295, 435, classes[2], cex=1.2)
  text(125, 370, 'Predicted', cex=1.3, srt=90, font=2)
  text(245, 450, 'Actual', cex=1.3, font=2)
  rect(150, 305, 240, 365, col=getColor("red", res[2]))
  rect(250, 305, 340, 365, col=getColor("green", res[4]))
  text(140, 400, classes[1], cex=1.2, srt=90)
  text(140, 335, classes[2], cex=1.2, srt=90)
```

```

# add in the cm results
text(195, 400, res[1], cex=1.6, font=2, col='white')
text(195, 335, res[2], cex=1.6, font=2, col='white')
text(295, 400, res[3], cex=1.6, font=2, col='white')
text(295, 335, res[4], cex=1.6, font=2, col='white')

# add in the specifics
plot(c(100, 0), c(100, 0), type = "n", xlab="", ylab="", main =
"DETAILS", xaxt='n', yaxt='n')
text(10, 85, names(cm$byClass[1]), cex=1.2, font=2)
text(10, 70, round(as.numeric(cm$byClass[1]), 3), cex=1.2)
text(30, 85, names(cm$byClass[2]), cex=1.2, font=2)
text(30, 70, round(as.numeric(cm$byClass[2]), 3), cex=1.2)
text(50, 85, names(cm$byClass[5]), cex=1.2, font=2)
text(50, 70, round(as.numeric(cm$byClass[5]), 3), cex=1.2)
text(70, 85, names(cm$byClass[6]), cex=1.2, font=2)
text(70, 70, round(as.numeric(cm$byClass[6]), 3), cex=1.2)
text(90, 85, names(cm$byClass[7]), cex=1.2, font=2)
text(90, 70, round(as.numeric(cm$byClass[7]), 3), cex=1.2)

# add in the accuracy information
text(30, 35, names(cm$overall[1]), cex=1.5, font=2)
text(30, 20, round(as.numeric(cm$overall[1]), 3), cex=1.4)
text(70, 35, names(cm$overall[2]), cex=1.5, font=2)
text(70, 20, round(as.numeric(cm$overall[2]), 3), cex=1.4)
}

draw_confusion_matrix(cm)

```




logRegaccuracy

```
[1] 0.8376623
```

Linear Discriminant Analysis

```
lda.fit <- lda(target1 ~ ., data = heart.train)
```

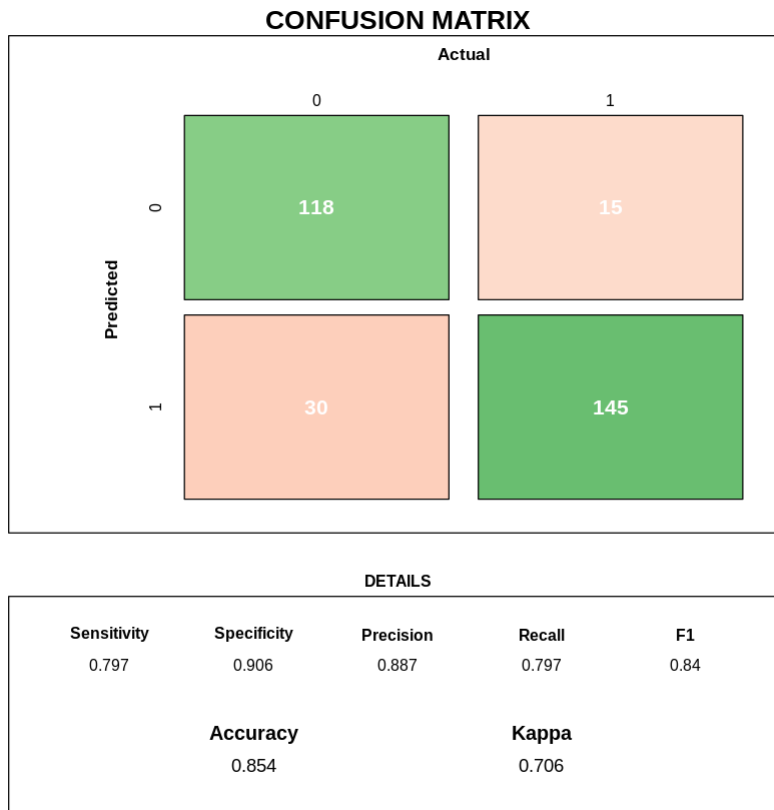
```
Warning message in lda.default(x, grouping, ...):  
"variables are collinear"
```

```
lda.pred <- predict(lda.fit, heart.test)  
names(lda.pred)
```

```
[1] "class"      "posterior" "x"
```

```
lda.class <- lda.pred$class  
cm<-confusionMatrix(as.factor(lda.class),  
as.factor(heart.test$target1))
```

```
draw_confusion_matrix(cm)
```



```
ldaaccuracy<-mean(lda.class==heart.test$target1)
ldaaccuracy
```

```
[1] 0.8538961
```

Naive Bayes

```
nb.fit <- naiveBayes(target1~., data = heart.train)
```

```
nb.class <- predict(nb.fit, heart.test)
cm<-confusionMatrix(as.factor(nb.class),
as.factor(heart.test$target1))
```

```
draw_confusion_matrix(cm)
```



```
naivebayesaccuracy<-mean(nb.class==heart.test$target1)
naivebayesaccuracy
```

```
[1] 0.8019481
```

SVM

```
svm.linear <- tune(svm, target1 ~ ., data = heart.train, kernel =
"linear",
  ranges = list(cost = c(0.001, 0.01, 0.1, 1, 5, 10, 100)))
```

```
svm.linear$best.model
```

Call:

```
best.tune(METHOD = svm, train.x = target1 ~ ., data = heart.train,
  ranges = list(cost = c(0.001, 0.01, 0.1, 1, 5, 10, 100)), kernel =
"linear")
```

Parameters:

```
SVM-Type: eps-regression
SVM-Kernel: linear
  cost: 0.01
  gamma: 0.04347826
  epsilon: 0.1
```

Number of Support Vectors: 576

```
pred.linear<-predict(svm.linear$best.model,heart.test)
pred.linear<-ifelse(pred.linear>0.5,1,0)
```

```
cm<-
confusionMatrix(data=as.factor(pred.linear),reference=as.factor(heart.
test$target1))
```

```
draw_confusion_matrix(cm)
```



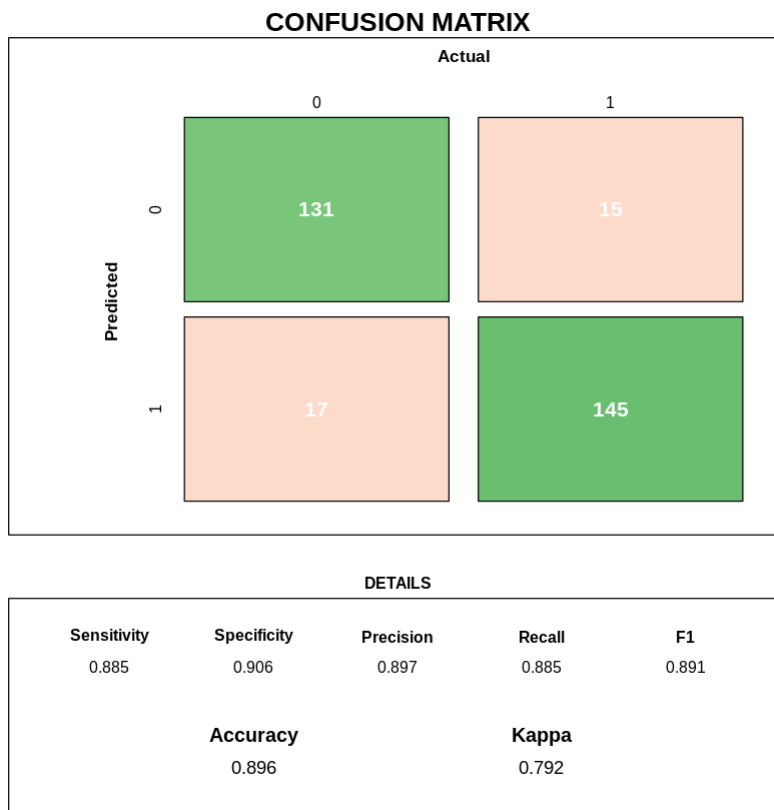
```
linearsvmaccuracy<-mean(pred.linear==heart.test$target1)
```

```
linearsvmaccuracy
```

```
[1] 0.8344156
```

```
svm.radial <- ksvm(target1 ~ ., data = heart.train, kernel = "rbfdot")
pred.radial <- predict(svm.radial, newdata = heart.test)
pred.radial <- ifelse(pred.radial > 0.5, 1, 0)
cm<-
confusionMatrix(data=as.factor(pred.radial),reference=as.factor(heart.
test$target1))
```

```
draw_confusion_matrix(cm)
```



```
radialsvmaccuracy<-mean(pred.radial==heart.test$target1)
```

```
radialsvmaccuracy
```

```
[1] 0.8961039
```

```
KNN
```

```
knnttrain <- heart.train
knntesting <- heart.test
trainlabels <- knnttrain$target1
testing_labels <- knntesting$target1
knnttrain$target1 <- NULL
knntesting$target1 <- NULL
knnprediction <- knn(train = knnttrain, test = knntesting,
                     cl = trainlabels, k=14)
```

```
knnprediction
```

```
[1] 0 0 1 1 1 0 0 1 0 1 1 1 1 0 1 1 1 0 0 0 1 0 1 0 1 1 0 1 1
1 1 0 1 0
[38] 1 0 0 0 0 0 0 1 1 1 1 0 1 0 1 0 0 0 1 1 0 0 1 1 0 0 1 1 0 1 0 1
0 0 1 1 1
[75] 0 1 1 1 0 0 1 1 0 0 0 1 0 1 0 1 1 1 0 1 1 0 1 0 1 1 1 0 1 1 1 1
```

```

1 0 0 1 1
[112] 1 1 1 0 0 0 1 1 0 0 1 0 1 1 1 0 1 0 1 1 1 0 1 1 1 0 1 0
0 1 1 1 1
[149] 1 0 1 0 1 0 1 1 1 1 1 1 1 0 1 1 1 1 1 0 1 1 1 0 0 1 1 0 1 0 0 0
1 1 0 0 1
[186] 1 0 1 0 1 1 1 0 1 1 1 1 0 1 0 0 0 0 0 0 0 0 1 1 1 1 1 0 0 0 1 0
1 1 0 1 1
[223] 1 1 0 1 1 1 0 0 1 0 0 0 0 1 1 1 0 0 0 1 0 0 0 0 0 1 1 1 0 0 0 1
1 1 1 1 1
[260] 1 1 0 0 0 0 1 0 1 0 0 1 1 0 0 0 1 1 0 0 0 0 0 1 0 1 0 1 1 0 0 1
1 1 1 0 0
[297] 1 1 1 0 0 0 0 0 1 0 0 1
Levels: 0 1

```

```

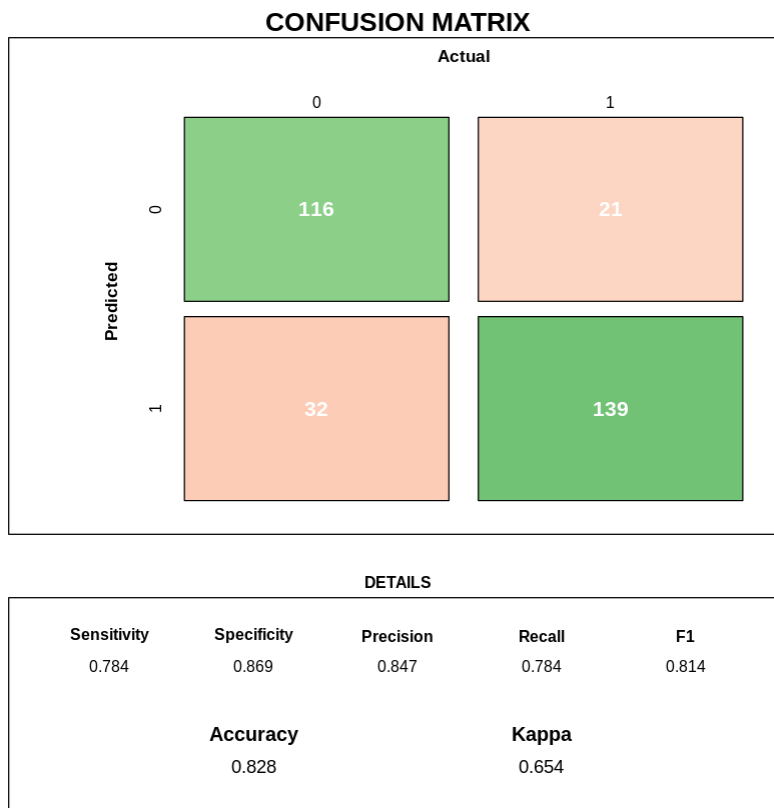
cm<-confusionMatrix(as.factor(knnprediction),
as.factor(heart.test$target1))

```

```

draw_confusion_matrix(cm)

```



```

knnaccuracy<-mean(knnprediction==heart.test$target1)
knnaccuracy

```

```

[1] 0.8279221

```

```

accuracy<-list()
for(i in 1:50)

```

```
{
  knntrain<-heart.train
  knntesting<-heart.test
  trainlabels<-knntrain$target1
  testing_labels<-knntesting$target1
  knntrain$target1<-NULL
  knntesting$target1<-NULL
  knnprediction<-
knn(train=knntrain,test=knntesting,cl=trainlabels,k=i)
  knnaccuracy<-mean(knnprediction==heart.test$target1)
  accuracy<-append(accuracy,knnaccuracy)
}
```

```
valuesofk<-1:50
```

```
accuracy
```

```
[[1]]
[1] 0.9902597
```

```
[[2]]
[1] 0.8993506
```

```
[[3]]
[1] 0.8246753
```

```
[[4]]
[1] 0.8214286
```

```
[[5]]
[1] 0.8084416
```

```
[[6]]
[1] 0.8051948
```

```
[[7]]
[1] 0.8019481
```

```
[[8]]
[1] 0.8214286
```

```
[[9]]
[1] 0.8051948
```

```
[[10]]
[1] 0.8279221
```

```
[[11]]
[1] 0.8246753
```

[[12]]
[1] 0.8376623

[[13]]
[1] 0.8116883

[[14]]
[1] 0.8246753

[[15]]
[1] 0.8246753

[[16]]
[1] 0.8116883

[[17]]
[1] 0.7987013

[[18]]
[1] 0.8051948

[[19]]
[1] 0.7954545

[[20]]
[1] 0.8019481

[[21]]
[1] 0.8019481

[[22]]
[1] 0.7954545

[[23]]
[1] 0.7987013

[[24]]
[1] 0.8019481

[[25]]
[1] 0.788961

[[26]]
[1] 0.7922078

[[27]]
[1] 0.7824675

[[28]]

[1] 0.788961

[[29]]

[1] 0.7922078

[[30]]

[1] 0.7922078

[[31]]

[1] 0.788961

[[32]]

[1] 0.788961

[[33]]

[1] 0.7857143

[[34]]

[1] 0.7857143

[[35]]

[1] 0.7987013

[[36]]

[1] 0.7954545

[[37]]

[1] 0.7954545

[[38]]

[1] 0.7857143

[[39]]

[1] 0.7824675

[[40]]

[1] 0.7824675

[[41]]

[1] 0.7824675

[[42]]

[1] 0.788961

[[43]]

[1] 0.788961

[[44]]

[1] 0.788961

```
[[45]]  
[1] 0.788961
```

```
[[46]]  
[1] 0.788961
```

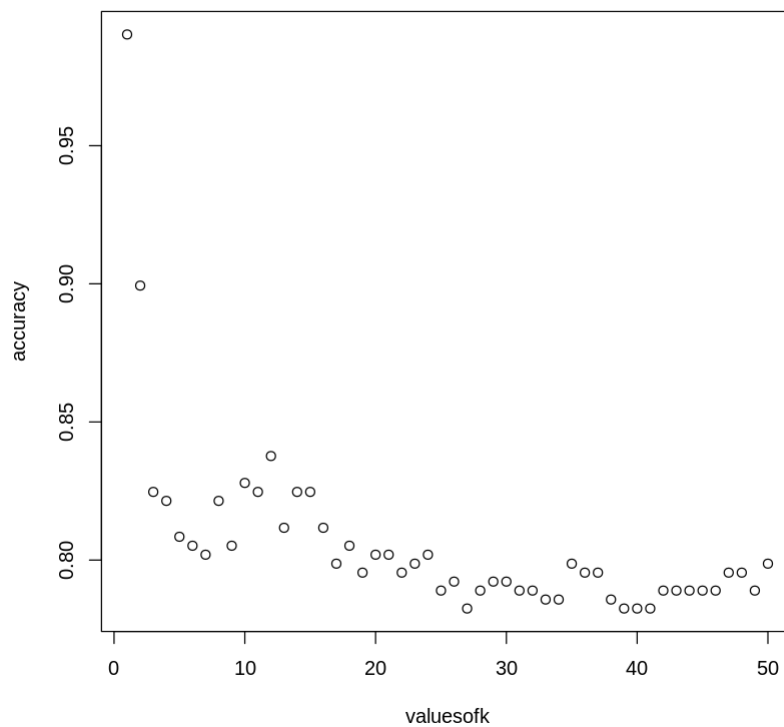
```
[[47]]  
[1] 0.7954545
```

```
[[48]]  
[1] 0.7954545
```

```
[[49]]  
[1] 0.788961
```

```
[[50]]  
[1] 0.7987013
```

```
plot(valuesofk,accuracy)
```



Tree

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

```
Installing package into '/usr/local/lib/R/site-library'
(as 'lib' is unspecified)
```

```
Loaded gbm 2.1.8.1
```

```
heart.tree<-tree(target1~.,data=heart.train)
summary(heart.tree)
```

Regression tree:

```
tree(formula = target1 ~ ., data = heart.train)
```

Variables actually used in tree construction:

```
[1] "thal2"      "oldpeak"    "trestbps"   "chol"       "restecg1"   "age"
[7] "cp3"        "cp2"        "thalach"    "exang1"     "ca3"        "slope1"
```

Number of terminal nodes: 24

Residual mean deviance: 0.06084 = 42.16 / 693

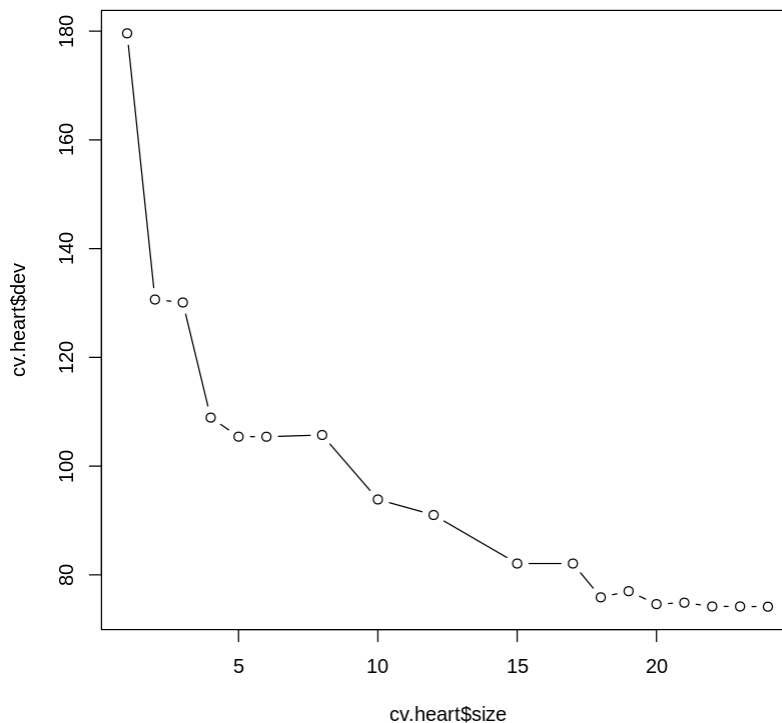
Distribution of residuals:

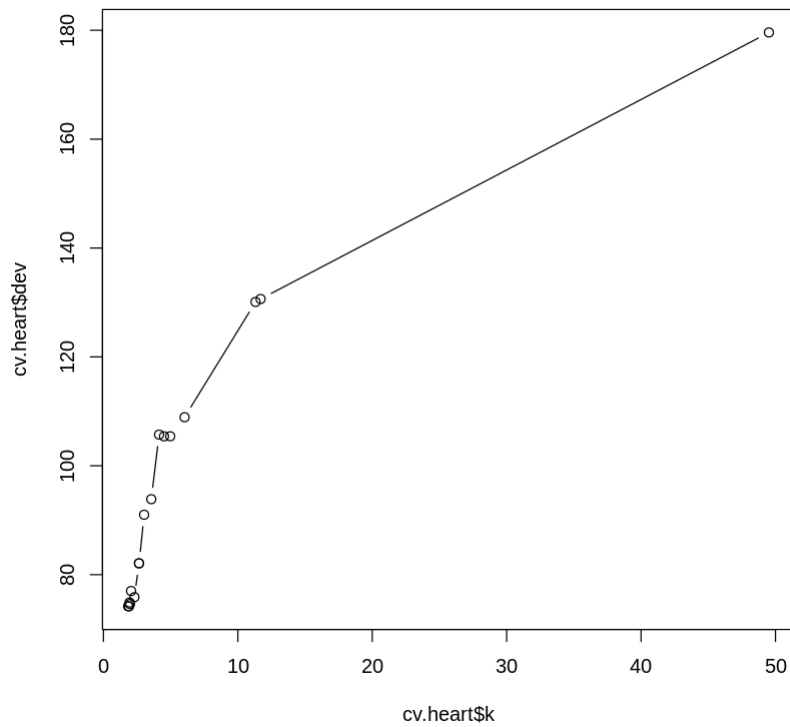
Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
-0.91410	-0.02235	0.00000	0.00000	0.08594	0.97770

```
cv.heart <- cv.tree(heart.tree, FUN = prune.tree,K=10)
```

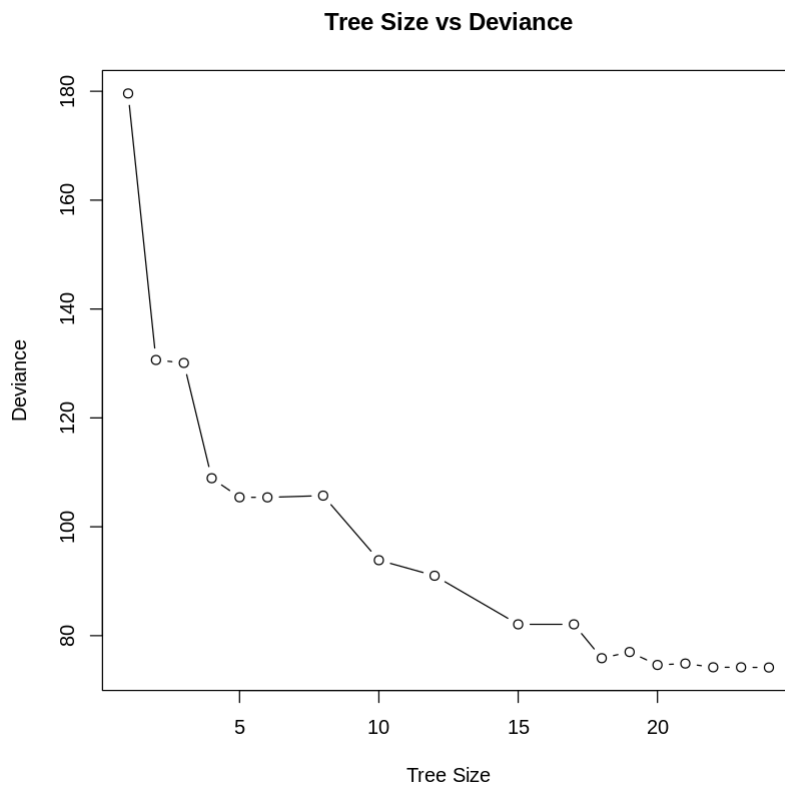
```
plot(cv.heart$size, cv.heart$dev, type = "b")
```

```
plot(cv.heart$k, cv.heart$dev, type = "b")
```





```
plot(cv.heart$size, cv.heart$dev, type = "b", xlab = "Tree Size", ylab = "Deviance", main = "Tree Size vs Deviance")
```



```
cv.heart$size
```

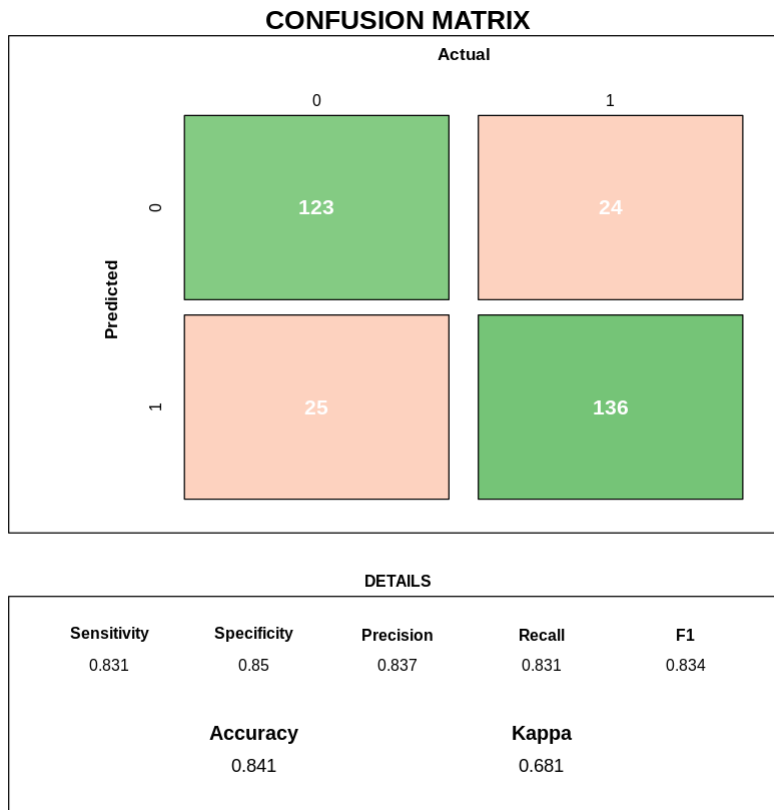
```
[1] 24 23 22 21 20 19 18 17 15 12 10 8 6 5 4 3 2 1
```

```
cv.heart$dev
```

```
[1] 74.15202 74.19153 74.19153 74.88207 74.62077 76.99959
75.86681
[8] 82.08400 82.08400 91.00989 93.86512 105.72492 105.40436
105.41887
[15] 108.92160 130.09359 130.63983 179.59392
```

```
heart.pruned<-prune.tree(heart.tree,best=23)
```

```
plot(heart.pruned)
text(heart.pruned,pretty=0)
```

```
table(predheart.pruned,heart.test$target1)
```

```
predheart.pruned  0   1
                  0 123  24
                  1  25 136
```

```
treeaccuracy<-mean(predheart.pruned==heart.test$target1)
treeaccuracy
```

```
[1] 0.8409091
```

Random Forest

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

```
Installing package into ‘/usr/local/lib/R/site-library’
(as ‘lib’ is unspecified)
```

```
bag.heart<-randomForest(target1~.,data = heart.train,mtry = 5,ntree =
100,importance = T)
```

```
Warning message in randomForest.default(m, y, ...):
"The response has five or fewer unique values. Are you sure you want
to do regression?"

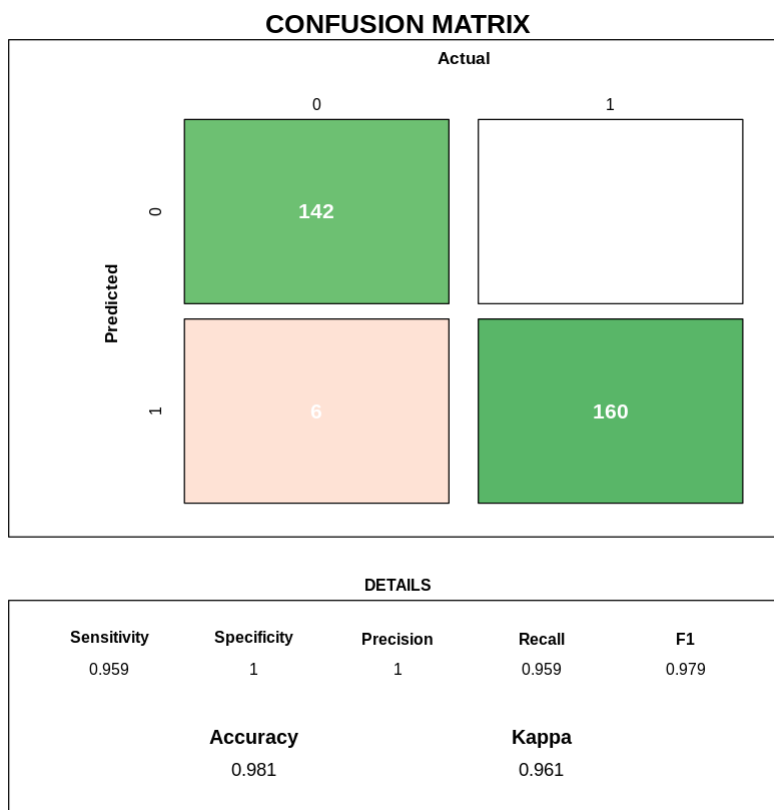
randomforestpredict<-predict(bag.heart,heart.test)
randomforestpredict <- ifelse(randomforestpredict > 0.5, 1, 0)
cm<-
confusionMatrix(data=as.factor(randomforestpredict),reference=as.factor(heart.test$target1))

randomforestaccuracy<-mean(randomforestpredict==heart.test$target1)

randomforestaccuracy

[1] 0.9805195

draw_confusion_matrix(cm)
```



```
summary(bag.heart)
```

	Length	Class	Mode
call	6	-none-	call
type	1	-none-	character
predicted	717	-none-	numeric
mse	100	-none-	numeric
rsq	100	-none-	numeric

oob.times	717	-none-	numeric
importance	46	-none-	numeric
importanceSD	23	-none-	numeric
localImportance	0	-none-	NULL
proximity	0	-none-	NULL
ntree	1	-none-	numeric
mtry	1	-none-	numeric
forest	11	-none-	list
coefs	0	-none-	NULL
y	717	-none-	numeric
test	0	-none-	NULL
inbag	0	-none-	NULL
terms	3	terms	call

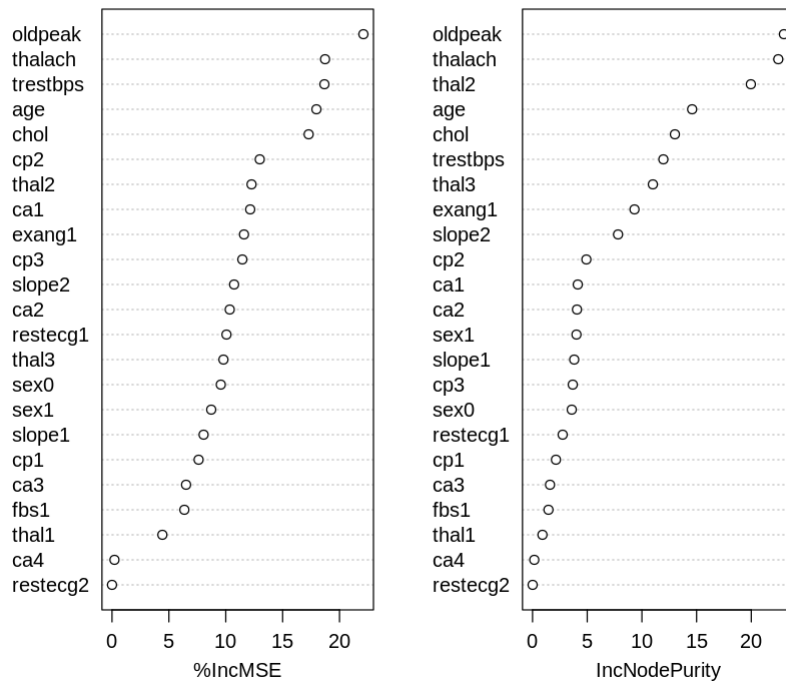
```
importance<-importance(bag.heart)
```

```
importance
```

	%IncMSE	IncNodePurity
age	17.9692447	14.59223130
sex0	9.5635295	3.58466263
sex1	8.7209044	4.01616520
cp1	7.6136227	2.13949528
cp2	12.9890674	4.92208486
cp3	11.4586969	3.67854049
trestbps	18.6731028	11.96179824
chol	17.2859821	13.01639675
fbs1	6.3607371	1.45607112
restecg1	10.0625733	2.75499606
restecg2	0.0000000	0.01004385
thalach	18.7299357	22.47016559
exang1	11.6032109	9.32221798
oldpeak	22.1003386	22.98636683
slope1	8.0518571	3.80983385
slope2	10.7328526	7.81790120
ca1	12.1487972	4.13621118
ca2	10.3503962	4.06272349
ca3	6.5106885	1.59802012
ca4	0.2171856	0.15954128
thal1	4.4287194	0.90696603
thal2	12.2661648	19.96232085
thal3	9.7964155	11.00353702

```
varImpPlot(bag.heart,main="Predictor Importance")
```

Predictor Importance



Artificial Neural Network

```
install.packages('neuralnet')
library('neuralnet')
```

Installing package into '/usr/local/lib/R/site-library'
(as 'lib' is unspecified)

also installing the dependency 'Deriv'

Attaching package: 'neuralnet'

The following object is masked from 'package:dplyr':

compute

```
trainlabels <- heart.train$target1
testing_labels <- heart.test$target1
neuralmod = neuralnet(target1~ ., data = heart.train,hidden=6)
neuralpred = predict(neuralmod, newdata = heart.test)
```

```
neuralpred <- ifelse(neuralpred > 0.5, 1, 0)
cm<-
confusionMatrix(data=as.factor(neuralpred),reference=as.factor(testing_
_labels))
draw_confusion_matrix(cm)
```



```
neuralnetworkaccuracy<-mean(neuralpred==testing_labels)
```

```
neuralnetworkaccuracy
```

```
[1] 0.9480519
```

```
tribble(
  ~Model,~Accuracy,
  "Logistic Regression", logRegaccuracy,
  "LDA", ldaaccuracy,
  "Naive Bayes", naivebayesaccuracy,
  "Linear SVM", linearsvmaccuracy,
  "Radial SVM", radialsvmaccuracy,
  "KNN", knnaccuracy,
  "Tree", treeaccuracy,
  "ANN", neuralnetworkaccuracy,
)
```

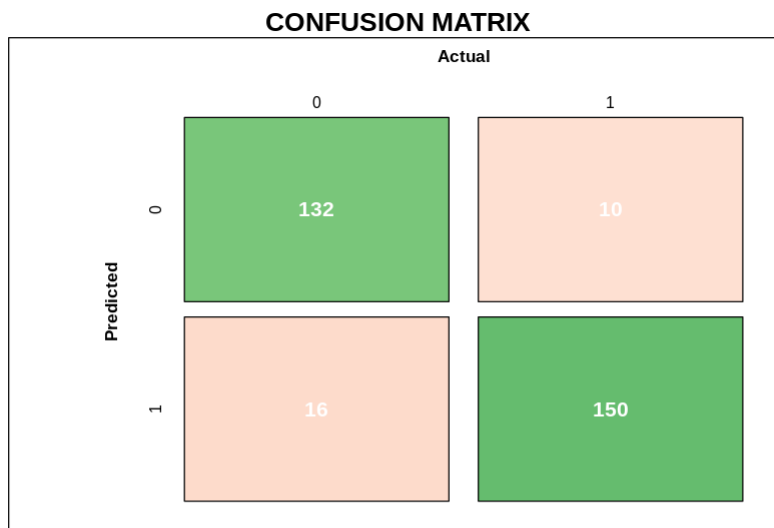
Model	Accuracy
1 Logistic Regression	0.8376623
2 LDA	0.8538961
3 Naive Bayes	0.8019481
4 Linear SVM	0.8344156
5 Radial SVM	0.8961039
6 KNN	0.7987013
7 Tree	0.8409091
8 ANN	0.9480519

```

trainlabels <- heart.train$target1
testing_labels <- heart.test$target1
neuralmod = neuralnet(target1~ ., data = heart.train)
neuralpred = predict(neuralmod, newdata = heart.test)
neuralpred <- ifelse(neuralpred > 0.5, 1, 0)
cm<-
confusionMatrix(data=as.factor(neuralpred),reference=as.factor(testing_labels))

draw_confusion_matrix(cm)

```



DETAILS

Sensitivity	Specificity	Precision	Recall	F1
0.892	0.938	0.93	0.892	0.91
Accuracy		Kappa		
0.916		0.831		