

Question 1.

- I. Read the data from "people.txt".

```
> people<-read.csv('people.csv')
> people
  Age agegroup height status yearsmarried
1  21    adult   6.0  single           -1
2   2    child   3.0 married            0
3  18    adult   5.7 married           20
4 221 elderly   5.0 widowed            2
5  34    child  -7.0 married            3
```

- II. Create a ruleset E that contain rules to check for the following conditions:

- a) The age should be in the range 0-150.

```
> E1<- editset(c("Age >=0", "Age<=150"))
> E1

Edit set:
num1 : 0 <= Age
num2 : Age <= 150
```

- b) The age should be greater than yearsmarried.

```
> E2<-editfile("edit.txt")
> E2

Edit set:
num1 : 0 <= Age
num2 : 0 < Height
num3 : Age <= 150
num4 : yearsmarried < Age
```

- c) The status should be married or single or widowed.

```
> E3<-editfile("edit2.txt")
> E3

Data model:
dat1 : Age_Group %in% c('Adult', 'child', 'Elderly')
dat2 : Status %in% c('Married', 'single', 'widowed')

Edit set:
cat1 : if( Age_Group == 'child' ) Status != 'Married'
```

- d) If age is less than 18 the agegroup should be child, if age is between 18 and 65 the agegroup should be adult, if age is more than 65 the agegroup should be elderly.

```
> E4<-editfile("edit3.txt")
> E4

Data model:
dat6 : Age_Group %in% c('Adult', 'child', 'Elderly')
dat7 : Status %in% c('Married', 'widowed')

Edit set:
mix1 : if ( yearsmarried + 17 <= Age ) FALSE
mix2 : if( 18 <= Age ) Age_Group != 'Adult'
mix3 : if( Age < 18 & 65 <= Age ) Age_Group != 'Adult'
mix4 : if( Age < 65 ) Age_Group != 'Adult'
```

III. Check whether ruleset E is violated by the data in the file people.txt.

a)

```
> E <- editset(c("Age >=0", "Age <=150"))
> ve1 <- violatedEdits(E,people)
> ve1
      edit
record num1 num2
      1 FALSE FALSE
      2 FALSE FALSE
      3 FALSE FALSE
      4 FALSE  TRUE
      5 FALSE FALSE
```

b)

```
> ve4 <- violatedEdits(E,people)
> ve4
      edit
record num1 num2 num3 num4
      1 FALSE FALSE FALSE FALSE
      2 FALSE FALSE FALSE FALSE
      3 FALSE FALSE FALSE  TRUE
      4 FALSE FALSE  TRUE FALSE
      5 FALSE  TRUE  FALSE FALSE
```

c)

```
> E <- editfile("edit2.txt")
> ve2 <- violatedEdits(E,people)
> ve2
      edit
record dat1 dat2 cat1
      1 FALSE FALSE FALSE
      2 FALSE FALSE  TRUE
      3 FALSE FALSE FALSE
      4 FALSE FALSE FALSE
      5 FALSE FALSE  TRUE
```

d)

```
> ve3 <- violatedEdits(E,people)
> ve3
      edit
record dat5 mix1 mix2 mix3
1 FALSE  TRUE FALSE  TRUE
2 FALSE FALSE FALSE FALSE
3 FALSE  TRUE FALSE  TRUE
4 FALSE FALSE FALSE FALSE
5 FALSE FALSE FALSE FALSE
```

IV. Summarize the results obtained in part(iii)

```
> summary(ve1)
Edit violations, 5 observations, 0 completely missing (0%):

editname freq rel
      num2    1 20%

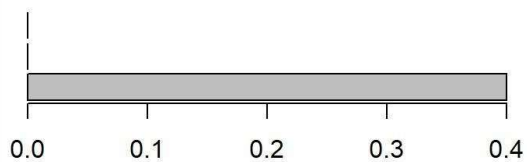
Edit violations per record:

errors freq rel
      0    4 80%
      1    1 20%
```

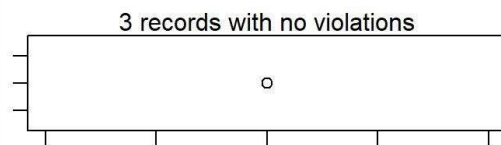
V. Visualize the results obtained in part (iii)

```
>plot(v3)
```

Edit violation frequency of top 3 edits



Edit violations per record



Question 2

Perform the following preprocessing tasks on the dirty_iris dataset

```
dirty_iris<- read.csv('iris_dirty.csv') dirty_iris
```

	X	Sepal.Length	Sepal.Width	Petal.Length	
1	1	5.1	3.5	1.4	
2	2	4.9	3.0	1.4	
3	3	4.7	3.2	1.3	
4	4	4.6	3.1	1.5	
5	5	NA	3.6	1.4	
6	6	5.4	NA	1.7	
7	7	4.6	3.4	1.4	
8	8	5.0	3.4	1.5	
9	9	4.4	2.9	1.4	
10	10	4.9	NA	1.5	
11	11	5.4	3.7	1.5	
12	12	4.8	3.4	1.6	
13	13	4.8	3.0	1.4	
14	14	4.3	3.0	1.1	
15	15	5.8	4.0	1.2	
16	16	5.7	4.4	1.5	
17	17	5.4	3.9	1.3	
18	18	5.1	3.5	1.4	
19	19	5.7	3.8	1.7	
20	20	5.1	3.8	1.5	
21	21	NA	3.4	1.7	
22	22	5.1	3.7	1.5	
23	23	4.6	3.6	1.0	
24	24	5.1	3.3	1.7	
25	25	4.8	3.4	1.9	
26	26	5.0	3.0	1.6	
27	27	5.0	NA	1.6	
28	28	5.2	3.5	1.5	
29	29	5.2	3.4	1.4	
30	30	4.7	3.2	1.6	
31	31	4.8	3.1	1.6	
32	32	5.4	3.4	1.5	
33	33	5.2	4.1	1.5	
34	34	5.5	4.2	1.4	
35	35	4.9	3.1	1.5	
36	36	5.0	3.2	1.2 12
38	38	4.9	3.6	1 12
40	40	5.1	3.4	1 12
42	42	4.5	2.3	1.3	43 43
4.4		3.2	1.344	44	5.0 3.5
1 12
45	45	5.1	3.8	1 12
37	37	5.5	NA	1.3	

39	39	4.4	3.0	1.3
41	41	5.0	3.5	1.3
46	46	NA	3.0	1.4
47	47	5.1	NA	1.6
48	48	4.6	3.2	1.4
49	49	5.3	3.7	1.5
50	50	5.0	3.3	1.4
51	51	7.0	3.2	4.7
52	52	6.4	3.2	4.5
53	53	6.9	NA	4.9
54	54	5.5	2.3	4.0
55	55	6.5	2.8	4.6
56	56	5.7	2.8	4.5
57	57	6.3	3.3	4.7
58	58	4.9	2.4	3.3
59	59	6.6	2.9	4.6
60	60	5.2	2.7	3.9
61	61	5.0	2.0	3.5
62	62	5.9	NA	4.2
63	63	6.0	2.2	4.0
64	64	6.1	2.9	4.7
65	65	5.6	2.9	3.6
66	66	6.7	3.1	4.4
67	67	5.6	3.0	4.5
68	68	5.8	2.7	4.1
69	69	6.2	2.2	4.5
70	70	5.6	2.5	3.9
71	71	5.9	3.2	4.8
72	72	6.1	2.8	4.0
73	73	6.3	2.5	4.9
74	74	6.1	2.8	4.7
75	75	6.4	2.9	4.3
76	76	6.6	3.0	4.4
77	77	6.8	2.8	4.8
78	78	6.7	3.0	5.0
79	79	6.0	2.9	4.5
80	80	5.7	2.6	3.5
81	81	5.5	2.4	3.8
82	82	5.5	2.4	3.7
83	83	5.8	2.7	3.9
84	84	6.0	2.7	5.1
85	85	5.4	3.0	4.5
86	86	6.0	3.4	4.5
87	87	NA	3.1	4.7
88	88	6.3	2.3	4.4
89	89	5.6	NA	4.1
90	90	5.5	2.5	4.0
91	91	5.5	2.6	4.4
92	92	6.1	3.0	4.6

93	93	5.8	2.6	4.0		
94	94	5.0	2.3	3.3	95	95
	5.6	2.7	4.2	96		5.7
	3.0	4.2	97	97	5.7	2.9
	4.2	98	98	6.2	2.9	4.3
99	99	5.1	2.5	3.0		
100	100	5.7	2.8	4.1		
101	101	6.3	3.3	6.0		
102	102	5.8	2.7	5.1		
103	103	7.1	3.0	5.9		
104	104	6.3	2.9	5.6		
105	105	6.5	3.0	5.8		
106	106	7.6	3.0	6.6		
107	107	4.9	2.5	4.5		
108	108	7.3	2.9	6.3		
109	109	6.7	2.5	5.8		
110	110	7.2	3.6	6.1		
111	111	6.5	3.2	5.1		
112	112	6.4	2.7	5.3		
113	113	6.8	3.0	5.5		
114	114	5.7	2.5	5.0		
115	115	5.8	NA	5.1		
116	116	6.4	3.2	5.3		
117	117	6.5	3.0	5.5		
118	118	7.7	3.8	6.7		
119	119	7.7	2.6	6.9		
120	120	6.0	2.2	5.0		
121	121	6.9	3.2	5.7		
122	122	5.6	2.8	4.9		
123	123	7.7	2.8	6.7		
124	124	6.3	2.7	4.9		
125	125	6.7	3.3	5.7		
126	126	7.2	3.2	6.0		
127	127	6.2	2.8	4.8		
128	128	6.1	3.0	4.9		
129	129	6.4	2.8	5.6		
130	130	7.2	3.0	5.8		
131	131	7.4	2.8	6.1		
132	132	7.9	3.8	6.4		
133	133	6.4	2.8	5.6		
134	134	6.3	NA	5.1		
135	135	6.1	2.6	5.6		
136	136	7.7	3.0	6.1		
137	137	6.3	3.4	5.6		
138	138	6.4	3.1	5.5		
139	139	6.0	3.0	4.8		
140	140	NA	3.1	5.4		
141	141	6.7	3.1	5.6		
142	142	6.9	3.1	5.1	143	143
	NA	2.7	5.1			

144	144	6.8	NA	5.9
145	145	NA	3.3	5.7
146	146	6.7	3.0	5.2
147	147	6.3	2.5	5.0
148	148	6.5	3.0	5.2
149	149	6.2	3.4	5.4
	150			150

	NA	3.0	5.1	
	Petal.Width	Species		
1	0.2	Setosa		
2	0.2	setosa		
3	0.2	setosa		
4	0.2	setosa		
5	0.2	setosa		
6	0.4	setosa		
7	0.3	setosa		
8	0.2	SETOSA		
9	0.2	setosa		
10	0.1	setosa		
11	0.2	setosa		
12	0.2	setosa		
13	0.1	setosa		
14	0.1	setosa		
15	0.2	setosa		
16	0.4	setosa		
17	0.4	setosa		
18	0.3	setosa		
19	0.3	setosa		
20	0.3	setosa		
21	0.2	setosa		
22	0.4	setosa		
23	0.2	setosa		
24	0.5	setosa		
25	0.2	setosa		
26	0.2	setosa		
27	0.4	setosa		
28	0.2	setosa		
29	0.2	setosa		
30	0.2	setosa		
31	0.2	setosa		
32	0.4	setosa		
33	0.1	setosa		
34	0.2	setosa		
35	0.2	setosa36	0.2	setosa
37	0.2	setosa		
38	0.1	setosa		
39	0.2	setosa		
40	0.2	setosa		
41	0.3	setosa		
42	0.3	setosa		

43	0.2	setosa		
44	0.6	setosa		
45	0.4	setosa46	0.3	setosa 47
	0.2	setosa		
48	0.2	setosa		
49	0.2	setosa		
50	0.2	setosa		
51	1.4	versicolor		
52	1.5	versicolor		
53	1.5	versicolor		
54	1.3	versicolor		
55	1.5	versicolor		
56	1.3	versicolor		
57	1.6	versicolor		
58	1.0	versicolor		
59	1.3	Versicolor		
60	1.4	versicolor		
61	1.0	versicolor		
62	1.5	versicolor		
63	1.0	versicolor		
64	1.4	VERSICOLOR		
65	1.3	versicolor		
66	1.4	versicolor		
67	1.5	versicolor		
68	1.0	versicolor		
69	1.5	versicolor		
70	1.1	versicolor		
71	1.8	versicolor		
72	1.3	versicolor		
73	1.5	versicolor		
74	1.2	versicolor		
75	1.3	versicolor		
76	1.4	versicolor		
77	1.4	versicolor		
78	1.7	versicolor		
79	1.5	versicolor		
80	1.0	versicolor		
81	1.1	versicolor		
82	1.0	versicolor		
83	1.2	versicolor		
84	1.6	versicolor		
85	1.5	versicolor		
86	1.6	versicolor		
87	1.5	versicolor		
88	1.3	versicolor		
89	1.3	versicolor		
90	1.3	versicolor		
91	1.2	versicolor92	1.4	versicolor 93
	1.2	versicolor 94	1.0	versicolor

95	1.3	versicolor	
96	1.2	versicolor	
97	1.3	versicolor	
98	1.3	versicolor	
99	1.1	versicolor	
100	1.3	versicolor	
101	2.5	virginica	
102	1.9	virginica	
103	2.1	virginica	
104	1.8	virginica	
105	2.2	virginica	
106	2.1	virginica	
107	1.7	virginica	
108	1.8	VIRGINICA	
109	1.8	virginica	
110	2.5	virginica	
111	2.0	virginica	
112	1.9	virginica	
113	2.1	virginica	
114	2.0	virginica	
115	2.4	virginica	
116	2.3	virginica	
117	1.8	virginica	
118	2.2	virginica	
119	2.3	virginica	
120	1.5	virginica	
121	2.3	virginica	
122	2.0	virginica	
123	2.0	virginica	
124	1.8	virginica	
125	2.1	virginica	
126	1.8	virginica	
127	1.8	virginica	
128	1.8	virginica	
129	2.1	virginica	
130	1.6	virginica	
131	1.9	virginica	
132	2.0	virginica	
133	2.2	virginica	
134	1.5	virginica	
135	1.4	virginica	
136	2.3	virginica	
137	2.4	virginica	
138	1.8	virginica	
139	1.8	virginica140	2.1 virginica 141
	2.4	virginica 142	2.3 virginica
143	1.9	virginica	
144	2.3	virginica	
145	2.5	virginica	

```

146      2.3 virginica
147      1.9 virginica
148      2.0 virginica
149      2.3 virginica
150      1.8 virginica

```

1. Calculate the number and percentage of observations that are complete. `c<-sum(complete.cases(dirty_iris)) cat("Number of complete observations:",c,"\n") cat("Percentage of complete observations:",c/(dim(dirty_iris)[1])*100,"\n\n")`

```

Number of complete observations: 131
Percentage of complete observations: 87.33333

```

2. Replace all the special values in data with NA.

```

replace <- function(x){  for (i in
1:length(x)){    if(is.infinite(x[i])
| is.nan(x[i]))    x[i]=NA    }  x }
for(i in 1:length(dirty_iris))
{
dirty_iris[,i]=replace(dirty_iris[,i])
}
dirty_iris

```

```

X Sepal.Length Sepal.Width Petal.Length
1      1      5.1      3.5      1.4
2      2      4.9      3.0      1.4
3      3      4.7      3.2      1.3
4      4      4.6      3.1      1.5
5      5      NA      3.6      1.4
6      6      5.4      NA      1.7
7      7      4.6      3.4      1.4
8      8      5.0      3.4      1.5
9      9      4.4      2.9      1.4
10     10      4.9      NA      1.5
11     11      5.4      3.7      1.5
12     12      4.8      3.4      1.6
13     13      4.8      3.0      1.4
14     14      4.3      3.0      1.1
15     15      5.8      4.0      1.2
16     16      5.7      4.4      1.5
17     17      5.4      3.9      1.3 18      18
      5.1      3.5      1.4
19     19      5.7      3.8      1.7

```

20	20	5.1	3.8	1.5
21	21	NA	3.4	1.7
22	22	5.1	3.7	1.5
23	23	4.6	3.6	1.0
24	24	5.1	3.3	1.7
25	25	4.8	3.4	1.9
26	26	5.0	3.0	1.6
27	27	5.0	NA	1.6
28	28	5.2	3.5	1.5
29	29	5.2	3.4	1.4
30	30	4.7	3.2	1.6
31	31	4.8	3.1	1.6
32	32	5.4	3.4	1.5
33	33	5.2	4.1	1.5
34	34	5.5	4.2	1.4
35	35	4.9	3.1	1.5
36	36	5.0	3.2	1.2
37	37	5.5	NA	1.3
38	38	4.9	3.6	1.4
39	39	4.4	3.0	1.3
40	40	5.1	3.4	1.5
41	41	5.0	3.5	1.3
42	42	4.5	2.3	1.3
43	43	4.4	3.2	1.3
44	44	5.0	3.5	1.6
45	45	5.1	3.8	1.9
46	46	NA	3.0	1.4
47	47	5.1	NA	1.6
48	48	4.6	3.2	1.4
49	49	5.3	3.7	1.5
50	50	5.0	3.3	1.4
51	51	7.0	3.2	4.7
52	52	6.4	3.2	4.5
53	53	6.9	NA	4.9
54	54	5.5	2.3	4.0
55	55	6.5	2.8	4.6
56	56	5.7	2.8	4.5
57	57	6.3	3.3	4.7
58	58	4.9	2.4	3.3
59	59	6.6	2.9	4.6
60	60	5.2	2.7	3.9
61	61	5.0	2.0	3.5
62	62	5.9	NA	4.2 63 63
	6.0	2.2	4.0	
64	64	6.1	2.9	4.7
65	65	5.6	2.9	3.6
66	66	6.7	3.1	4.4
67	67	5.6	3.0	4.5
68	68	5.8	2.7	4.1

69	69	6.2	2.2	4.5
70	70	5.6	2.5	3.9
71	71	5.9	3.2	4.8
72	72	6.1	2.8	4.0
73	73	6.3	2.5	4.9
74	74	6.1	2.8	4.7
75	75	6.4	2.9	4.3
76	76	6.6	3.0	4.4
77	77	6.8	2.8	4.8
78	78	6.7	3.0	5.0
79	79	6.0	2.9	4.5
80	80	5.7	2.6	3.5
81	81	5.5	2.4	3.8
82	82	5.5	2.4	3.7
83	83	5.8	2.7	3.9
84	84	6.0	2.7	5.1
85	85	5.4	3.0	4.5
86	86	6.0	3.4	4.5
87	87	NA	3.1	4.7
88	88	6.3	2.3	4.4
89	89	5.6	NA	4.1
90	90	5.5	2.5	4.0
91	91	5.5	2.6	4.4
92	92	6.1	3.0	4.6
93	93	5.8	2.6	4.0
94	94	5.0	2.3	3.3
95	95	5.6	2.7	4.2
96	96	5.7	3.0	4.2
97	97	5.7	2.9	4.2
98	98	6.2	2.9	4.3
99	99	5.1	2.5	3.0
100	100	5.7	2.8	4.1
101	101	6.3	3.3	6.0
102	102	5.8	2.7	5.1
103	103	7.1	3.0	5.9
104	104	6.3	2.9	5.6
105	105	6.5	3.0	5.8
106	106	7.6	3.0	6.6
107	107	4.9	2.5	4.5
108	108	7.3	2.9	6.3
109	109	6.7	2.5	5.8
110	110	7.2	3.6	6.1
111	111	6.5	3.2	5.1
112	112	6.4	2.7	5.3
113	113	6.8	3.0	5.5
114	114	5.7	2.5	5.0
115	115	5.8	NA	5.1
116	116	6.4	3.2	5.3
117	117	6.5	3.0	5.5

118	118	7.7	3.8	6.7
119	119	7.7	2.6	6.9
120	120	6.0	2.2	5.0
121	121	6.9	3.2	5.7
122	122	5.6	2.8	4.9
123	123	7.7	2.8	6.7
124	124	6.3	2.7	4.9
125	125	6.7	3.3	5.7
126	126	7.2	3.2	6.0
127	127	6.2	2.8	4.8
128	128	6.1	3.0	4.9
129	129	6.4	2.8	5.6
130	130	7.2	3.0	5.8
131	131	7.4	2.8	6.1
132	132	7.9	3.8	6.4
133	133	6.4	2.8	5.6
134	134	6.3	NA	5.1
135	135	6.1	2.6	5.6
136	136	7.7	3.0	6.1
137	137	6.3	3.4	5.6
138	138	6.4	3.1	5.5
139	139	6.0	3.0	4.8
140	140	NA	3.1	5.4
141	141	6.7	3.1	5.6
142	142	6.9	3.1	5.1
143	143	NA	2.7	5.1
144	144	6.8	NA	5.9
145	145	NA	3.3	5.7
146	146	6.7	3.0	5.2
147	147	6.3	2.5	5.0
148	148	6.5	3.0	5.2
149	149	6.2	3.4	5.4
150	150	NA	3.0	5.1

	Petal.Width	Species		
1	0.2	Setosa		
2	0.2	setosa		
3	0.2	setosa		
4	0.2	setosa		
5	0.2	setosa		
6	0.4	setosa		
7	0.3	setosa		
8	0.2	SETOSA		
9	0.2	setosa		
10	0.1	setosa		
11	0.2	setosa12	0.2	setosa 13
	0.1	setosa		
14	0.1	setosa		
15	0.2	setosa		
16	0.4	setosa		

17	0.4	setosa
18	0.3	setosa
19	0.3	setosa
20	0.3	setosa
21	0.2	setosa
22	0.4	setosa
23	0.2	setosa
24	0.5	setosa
25	0.2	setosa
26	0.2	setosa
27	0.4	setosa
28	0.2	setosa
29	0.2	setosa
30	0.2	setosa
31	0.2	setosa
32	0.4	setosa
33	0.1	setosa
34	0.2	setosa
35	0.2	setosa
36	0.2	setosa
37	0.2	setosa
38	0.1	setosa
39	0.2	setosa
40	0.2	setosa
41	0.3	setosa
42	0.3	setosa
43	0.2	setosa
44	0.6	setosa
45	0.4	setosa
46	0.3	setosa
47	0.2	setosa
48	0.2	setosa
49	0.2	setosa
50	0.2	setosa
51	1.4	versicolor
52	1.5	versicolor
53	1.5	versicolor
54	1.3	versicolor
55	1.5	versicolor
56	1.3	versicolor
57	1.6	versicolor
58	1.0	versicolor59
	1.4	versicolor
61	1.0	versicolor
62	1.5	versicolor
63	1.0	versicolor
64	1.4	VERSICOLOR
65	1.3	versicolor
66	1.4	versicolor

1.3 Versicolor 60

67	1.5 versicolor	
68	1.0 versicolor	
69	1.5 versicolor	
70	1.1 versicolor	
71	1.8 versicolor	
72	1.3 versicolor	
73	1.5 versicolor	
74	1.2 versicolor	
75	1.3 versicolor	
76	1.4 versicolor	
77	1.4 versicolor	
78	1.7 versicolor	
79	1.5 versicolor	
80	1.0 versicolor	
81	1.1 versicolor	
82	1.0 versicolor	
83	1.2 versicolor	
84	1.6 versicolor	
85	1.5 versicolor	
86	1.6 versicolor	
87	1.5 versicolor	
88	1.3 versicolor	
89	1.3 versicolor	
90	1.3 versicolor	
91	1.2 versicolor	
92	1.4 versicolor	
93	1.2 versicolor	
94	1.0 versicolor	
95	1.3 versicolor	
96	1.2 versicolor	
97	1.3 versicolor	
98	1.3 versicolor	
99	1.1 versicolor	
100	1.3 versicolor	
101	2.5 virginica	
102	1.9 virginica	
103	2.1 virginica	
104	1.8 virginica	
105	2.2 virginica106	2.1 virginica 107
	1.7 virginica	
108	1.8 VIRGINICA	
109	1.8 virginica110	2.5 virginica
111	2.0 virginica	
112	1.9 virginica	
113	2.1 virginica	
114	2.0 virginica	
115	2.4 virginica116	2.3 virginica 117
	1.8 virginica	
118	2.2 virginica	
119	2.3 virginica	

120	1.5	virginica
121	2.3	virginica
122	2.0	virginica
123	2.0	virginica
124	1.8	virginica
125	2.1	virginica
126	1.8	virginica
127	1.8	virginica
128	1.8	virginica
129	2.1	virginica
130	1.6	virginica
131	1.9	virginica
132	2.0	virginica
133	2.2	virginica
134	1.5	virginica
135	1.4	virginica
136	2.3	virginica
137	2.4	virginica
138	1.8	virginica
139	1.8	virginica
140	2.1	virginica
141	2.4	virginica
142	2.3	virginica
143	1.9	virginica
144	2.3	virginica
145	2.5	virginica
146	2.3	virginica
147	1.9	virginica
148	2.0	virginica
149	2.3	virginica
150	1.8	virginica

3. Define these rules in a separate text file and read them.

(Use `editfile` function in R (package `editrules`). Use similar function in Python). Print the resulting constraint object.

- Species should be one of the following values: `setosa`, `versicolor` or `virginica`.
- All measured numerical properties of an iris should be positive.
- The petal length of an iris is atleast 2 times its petal width.
- The sepal length of an iris cannot exceed 30cm.
- The sepals of an iris are longer than its petals.


```
> E<-editfile("editQ2.txt")
> E
```

Data model:

```
dat1 : Species %in% c('setosa', 'versicolor', 'virginica')
```

Edit set:

```
num1 : 0 <= Sepal.Length
num2 : 0 <= Sepal.Width
num3 : 0 <= Petal.Length
num4 : 0 <= Petal.Width
num5 : 2*Petal.Width <= Petal.Length
num6 : Sepal.Length <= 30
num7 : Petal.Length < Sepal.Length
num8 : Petal.width < Sepal.Width
```

4. Determine how often each rule is broken (violatedEdits). Also summarize and

plot the result `ve <- violatedEdits(E,dirty_iris) ve`

edit

```
record num1 num2 num3 num4 num5 num6
1 FALSE FALSE FALSE FALSE FALSE FALSE
2 FALSE FALSE FALSE FALSE FALSE FALSE
3 FALSE FALSE FALSE FALSE FALSE FALSE
4 FALSE FALSE FALSE FALSE FALSE FALSE
5 NA FALSE FALSE FALSE FALSE NA
6 FALSE NA FALSE FALSE FALSE FALSE
7 FALSE FALSE FALSE FALSE FALSE FALSE
8 FALSE FALSE FALSE FALSE FALSE FALSE
9 FALSE FALSE FALSE FALSE FALSE FALSE
10 FALSE NA FALSE FALSE FALSE FALSE
11 FALSE FALSE FALSE FALSE FALSE FALSE
12 FALSE FALSE FALSE FALSE FALSE FALSE
13 FALSE FALSE FALSE FALSE FALSE FALSE 14 FALSE
15 FALSE FALSE FALSE FALSE FALSE FALSE
16 FALSE FALSE FALSE FALSE FALSE FALSE
17 FALSE FALSE FALSE FALSE FALSE FALSE
18 FALSE FALSE FALSE FALSE FALSE FALSE
19 FALSE FALSE FALSE FALSE FALSE FALSE
20 FALSE FALSE FALSE FALSE FALSE FALSE
21 NA FALSE FALSE FALSE FALSE NA
22 FALSE FALSE FALSE FALSE FALSE FALSE
23 FALSE FALSE FALSE FALSE FALSE FALSE
24 FALSE FALSE FALSE FALSE FALSE FALSE
25 FALSE FALSE FALSE FALSE FALSE FALSE
26 FALSE FALSE FALSE FALSE FALSE FALSE
27 FALSE NA FALSE FALSE FALSE FALSE
```

28	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
29	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
30	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
31	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
32	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
33	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
34	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
35	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
36	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
37	FALSE	NA	FALSE	FALSE	FALSE	FALSE
38	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
39	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
40	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
41	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
42	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
43	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
44	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
45	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
46	NA	FALSE	FALSE	FALSE	FALSE	NA
47	FALSE	NA	FALSE	FALSE	FALSE	FALSE
48	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
49	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
50	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
51	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
52	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
53	FALSE	NA	FALSE	FALSE	FALSE	FALSE
54	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
55	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
56	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
57	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
58	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
59	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
60	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
61	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
62	FALSE	NA	FALSE	FALSE	FALSE	FALSE
63	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
64	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
65	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
66	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
67	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
68	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
69	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
70	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
71	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
72	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
73	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
74	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
75	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE
76	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE

77	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
78	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
79	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
80	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
81	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
82	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
83	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
84	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
85	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
86	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
87	NA	FALSE	FALSE	FALSE	FALSE	FALSE	NA	
88	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
89	FALSE	NA	FALSE	FALSE	FALSE	FALSE	FALSE	
90	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
91	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
92	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
93	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
94	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
95	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
96	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
97	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
98	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
99	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
100	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
101	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
102	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
103	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
104	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
105	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
106	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
107	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
108	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
109	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	
110	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	111 FALSE
	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	FALSE	edit record num7
	num8	dat1	1	FALSE	FALSE	TRUE		
2	FALSE	FALSE	FALSE					
3	FALSE	FALSE	FALSE4	FALSE	FALSE			
	FALSE	5	NA	FALSE	FALSE			
6	FALSE	NA	FALSE					
7	FALSE	FALSE	FALSE					
8	FALSE	FALSE	TRUE					
9	FALSE	FALSE	FALSE					
10	FALSE	NA	FALSE					
11	FALSE	FALSE	FALSE					
12	FALSE	FALSE	FALSE					
13	FALSE	FALSE	FALSE					
14	FALSE	FALSE	FALSE					
15	FALSE	FALSE	FALSE					
16	FALSE	FALSE	FALSE					

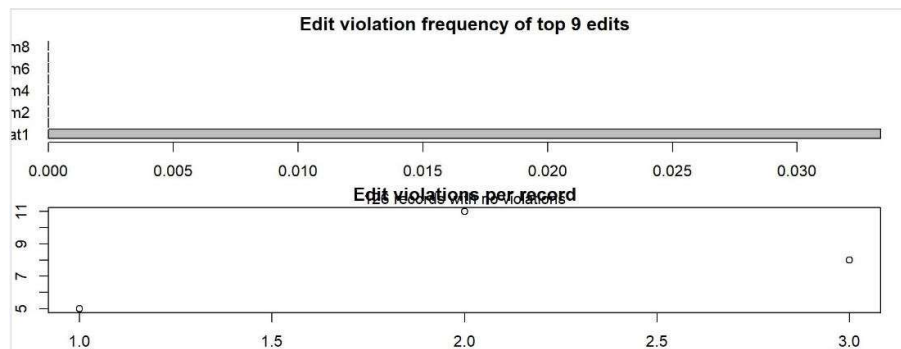
17	FALSE	FALSE	FALSE	
18	FALSE	FALSE	FALSE	
19	FALSE	FALSE	FALSE	
20	FALSE	FALSE	FALSE	
21	NA	FALSE	FALSE	
22	FALSE	FALSE	FALSE	
23	FALSE	FALSE	FALSE	
24	FALSE	FALSE	FALSE	
25	FALSE	FALSE	FALSE	
26	FALSE	FALSE	FALSE	
27	FALSE	NA	FALSE	
28	FALSE	FALSE	FALSE	
29	FALSE	FALSE	FALSE	
30	FALSE	FALSE	FALSE	
31	FALSE	FALSE	FALSE	
32	FALSE	FALSE	FALSE	
33	FALSE	FALSE	FALSE	
34	FALSE	FALSE	FALSE	
35	FALSE	FALSE	FALSE	
36	FALSE	FALSE	FALSE	
37	FALSE	NA	FALSE	
38	FALSE	FALSE	FALSE	
39	FALSE	FALSE	FALSE	
40	FALSE	FALSE	FALSE	
41	FALSE	FALSE	FALSE	
42	FALSE	FALSE	FALSE	
43	FALSE	FALSE	FALSE	
44	FALSE	FALSE	FALSE	
45	FALSE	FALSE	FALSE	
46	NA	FALSE	FALSE	
47	FALSE	NA	FALSE	
48	FALSE	FALSE	FALSE	
49	FALSE	FALSE	FALSE	
50	FALSE	FALSE	FALSE	
51	FALSE	FALSE	FALSE	
52	FALSE	FALSE	FALSE	
53	FALSE	NA	FALSE54	FALSE FALSE FALSE 55 FALSE
	FALSE	FALSE		
56	FALSE	FALSE	FALSE	
57	FALSE	FALSE	FALSE	
58	FALSE	FALSE	FALSE	
59	FALSE	FALSE	TRUE	
60	FALSE	FALSE	FALSE	
61	FALSE	FALSE	FALSE	
62	FALSE	NA	FALSE	
63	FALSE	FALSE	FALSE	
64	FALSE	FALSE	TRUE	
65	FALSE	FALSE	FALSE	
66	FALSE	FALSE	FALSE	

```

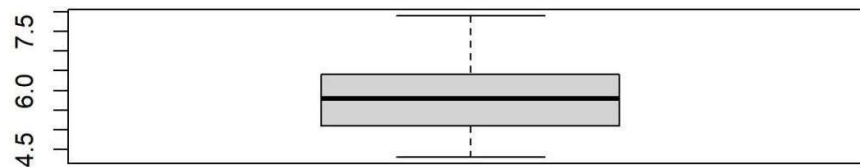
67 FALSE FALSE FALSE
68 FALSE FALSE FALSE
69 FALSE FALSE FALSE
70 FALSE FALSE FALSE
71 FALSE FALSE FALSE
72 FALSE FALSE FALSE
73 FALSE FALSE FALSE
74 FALSE FALSE FALSE
75 FALSE FALSE FALSE
76 FALSE FALSE FALSE
77 FALSE FALSE FALSE
78 FALSE FALSE FALSE
79 FALSE FALSE FALSE
80 FALSE FALSE FALSE
81 FALSE FALSE FALSE
82 FALSE FALSE FALSE
83 FALSE FALSE FALSE
84 FALSE FALSE FALSE
85 FALSE FALSE FALSE
86 FALSE FALSE FALSE
87 NA FALSE FALSE
88 FALSE FALSE FALSE
89 FALSE NA FALSE
90 FALSE FALSE FALSE
91 FALSE FALSE FALSE
92 FALSE FALSE FALSE
93 FALSE FALSE FALSE
94 FALSE FALSE FALSE
95 FALSE FALSE FALSE
96 FALSE FALSE FALSE
97 FALSE FALSE FALSE
98 FALSE FALSE FALSE
99 FALSE FALSE FALSE
100 FALSE FALSE FALSE
101 FALSE FALSE FALSE
102 FALSE FALSE FALSE
103 FALSE FALSE FALSE104 FALSE FALSE FALSE 105 FALSE
    FALSE FALSE
106 FALSE FALSE FALSE
107 FALSE FALSE FALSE
108 FALSE FALSE TRUE
109 FALSE FALSE FALSE
110 FALSE FALSE FALSE
111 FALSE FALSE FALSE

```

Plot(ve)



5. Find outliers in sepal length using boxplot and boxplot.stats
`boxplot(iris$Sepal.Length)`



```
> boxplot.stats((iris$Sepal.Length))
$stats
[1] 4.3 5.1 5.8 6.4 7.9

$n
[1] 150

$conf
[1] 5.632292 5.967708

$out
numeric(0)
```

Question 3

Load the data from wine dataset. Check whether all attributes are standardized or not (mean is 0 and standard deviation is 1). If not, standardize the attributes.

```
wine <- as.data.frame(read.csv("wine.csv"))
fixed_acidity volatile_acidity citric_acid
1          7.4          0.700          0.00
2          7.8          0.880          0.00
```

3	7.8	0.760	0.04
4	11.2	0.280	0.56
5	7.4	0.700	0.00
6	7.4	0.660	0.00
7	7.9	0.600	0.06
8	7.3	0.650	0.00
9	7.8	0.580	0.02
10	7.5	0.500	0.36
11	6.7	0.580	0.08
12	7.5	0.500	0.36
13	5.6	0.615	0.00
14	7.8	0.610	0.29
15	8.9	0.620	0.18
16	8.9	0.620	0.19
17	8.5	0.280	0.56
18	8.1	0.560	0.28
19	7.4	0.590	0.08
20	7.9	0.320	0.51
21	8.9	0.220	0.48
22	7.6	0.390	0.31
23	7.9	0.430	0.21
24	8.5	0.490	0.11
25	6.9	0.400	0.14
26	6.3	0.390	0.16
27	7.6	0.410	0.24
28	7.9	0.430	0.21
29	7.1	0.710	0.00
30	7.8	0.645	0.00
31	6.7	0.675	0.07
32	6.9	0.685	0.00
33	8.3	0.655	0.12
34	6.9	0.605	0.12

35	5.2	0.320	0.25	
36	7.8	0.645	0.00	
37	7.8	0.600	0.14	
38	8.1	0.380	0.28	
39	5.7	1.130	0.09	
40	7.3	0.450	0.36	
41	7.3	0.450	0.36	
42	8.8	0.610	0.30	
43	7.5	0.490	0.20	
44	8.1	0.660	0.22	
45	6.8	0.670	0.02	
46	4.6	0.520	0.15	
47	7.7	0.935	0.43	
48	8.7	0.290	0.52	
49	6.4	0.400	0.23	
50	5.6	0.310	0.37	
51	8.8	0.660	0.26	
52	6.6	0.520	0.04	
53	6.6	0.500	0.04	
54	8.6	0.380	0.36	
55	7.6	0.510	0.15	
56	7.7	0.620	0.04	
57	10.2	0.420	0.57	
58	7.5	0.630	0.12	
59	7.8	0.590	0.18	60
	0.390	0.31		7.3
61	8.8	0.400	0.40	
62	7.7	0.690	0.49	
63	7.5	0.520	0.16	
64	7.0	0.735	0.05	
65	7.2	0.725	0.05	
66	7.2	0.725	0.05	

67	7.5	0.520	0.11	
68	6.6	0.705	0.07	
69	9.3	0.320	0.57	
70	8.0	0.705	0.05	
71	7.7	0.630	0.08	
72	7.7	0.670	0.23	
73	7.7	0.690	0.22	
74	8.3	0.675	0.26	
75	9.7	0.320	0.54	
76	8.8	0.410	0.64	residual_sugar
	chlorides	free_sulfur_dioxide	1	1.90
	0.076		11	
2	2.60	0.098		25
3	2.30	0.092		15
4	1.90	0.075		17
5	1.90	0.076		11
6	1.80	0.075		13
7	1.60	0.069		15
8	1.20	0.065		15
9	2.00	0.073		9
10	6.10	0.071		17
11	1.80	0.097		15
12	6.10	0.071		17
13	1.60	0.089		16
14	1.60	0.114		9 15
	0.176		52	3.80
16	3.90	0.170		51
17	1.80	0.092		35
18	1.70	0.368		16
19	4.40	0.086		6
20	1.80	0.341		17
21	1.80	0.077		29

22	2.30	0.082	23	
23	1.60	0.106	10	
24	2.30	0.084	9	
25	2.40	0.085	21	
26	1.40	0.080	11	
27	1.80	0.080	4	
28	1.60	0.106	10	
29	1.90	0.080	14	
30	2.00	0.082	8	
31	2.40	0.089	17	
32	2.50	0.105	22	
33	2.30	0.083	15	
34	10.70	0.073	40	
35	1.80	0.103	13	
36	5.50	0.086	5	
37	2.40	0.086	3	
38	2.10	0.066	13	
39	1.50	0.172	7	
40	5.90	0.074	12	
41	5.90	0.074	12	
42	2.80	0.088	17	
43	2.60	0.332	8	
44	2.20	0.069	9	
45	1.80	0.050	5	
46	2.10	0.054	8 47	2.20
	0.114		22	
48	1.60	0.113	12	
49	1.60	0.066	5	
50	1.40	0.074	12	
51	1.70	0.074	4	
52	2.20	0.069	8	
53	2.10	0.068	6	

54	3.00	0.081	30
55	2.80	0.110	33
56	3.80	0.084	25
57	3.40	0.070	4
58	5.10	0.111	50
59	2.30	0.076	17
60	2.40	0.074	9
61	2.20	0.079	19
62	1.80	0.115	20
63	1.90	0.085	12
64	2.00	0.081	13
65	4.65	0.086	4
66	4.65	0.086	4
67	1.50	0.079	11
68	1.60	0.076	6
69	2.00	0.074	27
70	1.90	0.074	8
71	1.90	0.076	15
72	2.10	0.088	17
73	1.90	0.084	18
74	2.10	0.084	11
75	2.50	0.094	28
76	2.20	0.093	9

total_sulfur_dioxide density pH sulphates

1	34	0.9978	3.51	0.56
2	67	0.9968	3.20	0.68
3				54
		0.9970	3.26	0.65
4	60	0.9980	3.16	0.58
5	34	0.9978	3.51	0.56
6	40	0.9978	3.51	0.56
7	59	0.9964	3.30	0.46

8	21	0.9946	3.39	0.47
9	18	0.9968	3.36	0.57
10	102	0.9978	3.35	0.80
11	65	0.9959	3.28	0.54
12	102	0.9978	3.35	0.80
13	59	0.9943	3.58	0.52
14	29	0.9974	3.26	1.56
15	145	0.9986	3.16	0.88
16	148	0.9986	3.17	0.93
17	103	0.9969	3.30	0.75
18	56	0.9968	3.11	1.28
19	29	0.9974	3.38	0.50
20	56	0.9969	3.04	1.08
21	60	0.9968	3.39	0.53
22	71	0.9982	3.52	0.65
23	37	0.9966	3.17	0.91
24	67	0.9968	3.17	0.53
25	40	0.9968	3.43	0.63
26	23	0.9955	3.34	0.56
27	11	0.9962	3.28	0.59
28	37	0.9966	3.17	0.91
29	35	0.9972	3.47	0.55
30	16	0.9964	3.38	0.59
31	82	0.9958	3.35	0.54
32	37	0.9966	3.46	0.57
33	113	0.9966	3.17	0.66
	83	0.9993	3.45	0.52
35	50	0.9957	3.38	0.55
36	18	0.9986	3.40	0.55
37	15	0.9975	3.42	0.60
38	30	0.9968	3.23	0.73
39	19	0.9940	3.50	0.48

40	87	0.9978	3.33	0.83
41	87	0.9978	3.33	0.83
42	46	0.9976	3.26	0.51
43	14	0.9968	3.21	0.90
44	23	0.9968	3.30	1.20
45	11	0.9962	3.48	0.52
46	65	0.9934	3.90	0.56
47	114	0.9970	3.25	0.73
48	37	0.9969	3.25	0.58
49	12	0.9958	3.34	0.56
50	96	0.9954	3.32	0.58
51	23	0.9971	3.15	0.74
52	15	0.9956	3.40	0.63
53	14	0.9955	3.39	0.64
54	119	0.9970	3.20	0.56
55	73	0.9955	3.17	0.63
56	45	0.9978	3.34	0.53
57	10	0.9971	3.04	0.63
58	110	0.9983	3.26	0.77
59	54	0.9975	3.43	0.59
60	46	0.9962	3.41	0.54
61	52	0.9980	3.44	0.64
62	112	0.9968	3.21	0.71
63	35	0.9968	3.38	0.62
64	54	0.9966	3.39	0.57
65	11	0.9962	3.41	0.39 66
	11	0.9962	3.41	0.39
67	39	0.9968	3.42	0.58
68	15	0.9962	3.44	0.58
69	65	0.9969	3.28	0.79
70	19	0.9962	3.34	0.95
71	27	0.9967	3.32	0.54

72			96	0.9962	3.32	0.48	
73			94	0.9961	3.31	0.48	
74			43	0.9976	3.31	0.53	
75			83	0.9984	3.28	0.82	76
			42	0.9986	3.54	0.66	alcohol quality
			style 1		9.4	5	red
2	9.8	5	red				
3	9.8	5	red				
4	9.8	6	red				
5	9.4	5	red				
6	9.4	5	red				
7	9.4	5	red				
8	10.0	7	red				
9	9.5	7	red				
10	10.5	5	red				
11	9.2	5	red				
12	10.5	5	red				
13	9.9	5	red				
14	9.1	5	red				
15	9.2	5	red				
16	9.2	5	red				
17	10.5	7	red				
18	9.3	5	red				
19	9.0	4	red				
20	9.2	6	red				
21	9.4	6	red				
22	9.7	5	red				
23	9.5	5	red				
24	9.4	5	red				
25	9.7	6	red				
26	9.3	5	red				
27	9.5	5	red				

28	9.5	5	red			
29	9.4	5	red			
30	9.8	6	red			
31	10.1	5	red			
32	10.6	6	red			
33	9.8	5	red			
34	9.4	6	red			
35	9.2	5	red			
36	9.6	6	red			
37	10.8	6	red			
38	9.7	7	red			
39	9.8	4	red			
40	10.5	5	red			
41	10.5	5	red			
42	9.3	4	red			
43	10.5	6	red			
44	10.3	5	red			
45	9.5	5	red			
46	13.1	4	red			
47	9.2	5	red			
48	9.5	5	red			
49	9.2	5	red			
50	9.2	5	red			
51	9.2	5	red			
52	9.4	6	red	53	9.4	6 red
54	9.4	5	red			
55	10.2	6	red			
56	9.5	5	red			
57	9.6	5	red			
58	9.4	5	red			
59	10.0	5	red			
60	9.4	6	red			

```

61    9.2      5    red
62    9.3      5    red
63    9.5      7    red
64    9.8      5    red
65   10.9      5    red
66   10.9      5    red
67    9.6      5    red
68   10.7      5    red
69   10.7      5    red
70   10.5      6    red
71    9.5      6    red
72    9.5      5    red
73    9.5      5    red
74    9.2      4    red
75    9.6      5    red
76   10.5      5    red

[ reached 'max' / getOption("max.print") -- omitted 6421 rows ]

wine[1:(length(wine)-1)]

flag<-1 j<-1 for(j in
1:length(wine))
{
  if(mean(wine[,j])!=0 || sd(wine[,j])!=1)
  {
    flag<-0
  } j<-j+1
}
if(flag==1)
{
  cat("Dataset is normalized")
} if(flag==0){
  cat("Dataset is not normalized","\n")
cat("Normalizing the dataset:", "\n")  data_std<-
function(X){
  (x-mean(x))/sd(X)
}
}

```


Dataset is not normalized

Normalizing the dataset:

```
wine1<-data.frame(sapply(wine,data_std))
i<-1
for(i in 1:length(wine)-1){
  cat("The mean of ",names(wine[i]),"is ",as.integer(mean(wine1[,i])), " and
  standard
                                deviation is ",as.integer(sd(wine1[,i])))
  cat("\n")  i=i+1
}
```

```
The mean of  fixed_acidity is  0  and standard
deviation is  0 The mean of  volatile_acidity is  0
and standard
                                deviation is
1 The mean of  citric_acid is  0  and standard
deviation is  0 The mean of  residual_sugar is  0
and standard
                                deviation is
1 The mean of  chlorides is  0  and standard
deviation is  1 The mean of  free_sulfur_dioxide is
0  and standard
                                deviation
is  0
The mean of  total_sulfur_dioxide is  0  and standard
                                deviation is  1
The mean of  density is  0  and standard
deviation is  1
The mean of  pH is  0  and standard
                                deviation is  1
The mean of  sulphates is  0  and standard
deviation is  0 The mean of  alcohol is  0
and standard
                                deviation
is  0 The mean of  quality is  0  and
standard
                                deviation is  0
```

Question 4

Run Apriori algorithm to find frequent item sets and association rules

4.1 Use minimum support as 50% and minimum confidence as 75%

4.2 Use minimum support as 60% and minimum confidence as 60%

```
library(arules)
```

```
library(datasets)
data("Groceries")
inspect(Groceries[1:10])
```

```

      items
{citrus      fruit,
semi-finished bread,
margarine,
ready soups}
{tropical    fruit,
yogurt,
coffee}
{whole milk}
{pip         fruit,
yogurt,      cream
cheese ,     meat
spreads}
vegetables,  [5] {other
milk,        whole
milk,        condensed
milk,        long life
bakery product}
milk,        [6] {whole
yogurt,      butter,
abrasive cleaner}
rice,
[7]
{rolls/buns}
[8] {other vegetables,
      UHT-milk,
rolls/buns,
bottled      beer,
liquor       (appetizer)}
[9] {pot plants}
[10] {whole milk,
cereals}
```

```
r<-apriori(Groceries,parameter=list(support=0.05,conf=0.1,minlen=2))
```

```
Apriori
```

```
Parameter specification:
```

```

confidence minval smax arem  aval originalSupport
0.1      0.1      1 none FALSE          TRUE
maxtime support minlen maxlen target  ext
5      0.05      2      10 rules TRUE
```

```
Algorithmic control:
```

```

filter tree heap memopt load sort verbose
0.1 TRUE TRUE  FALSE TRUE    2    TRUE
```

```
Absolute minimum support count: 491
```

```

set item appearances ...[0 item(s)] done [0.00s]. set transactions
...[169 item(s), 9835 transaction(s)] done [0.01s]. sorting and
recoding items ... [28 item(s)] done [0.00s]. creating transaction
tree ... done [0.01s]. checking subsets of size 1 2 done [0.00s].
writing ... [6 rule(s)] done [0.00s]. creating S4 object ... done
[0.00s].

```

```
inspect(r)
```

```

lhs                      rhs                      [1]
{yogurt}                  => {whole milk}
[2] {whole milk}          => {yogurt}
[3] {rolls/buns}          => {whole milk}
[4] {whole milk}          => {rolls/buns}
[5] {other vegetables} => {whole milk} [6] {whole milk}
=> {other vegetables}      support    confidence coverage
lift      count [1] 0.05602440 0.4016035 0.1395018 1.571735
551
[2] 0.05602440 0.2192598 0.2555160 1.571735 551
[3] 0.05663447 0.3079049 0.1839349 1.205032 557
[4] 0.05663447 0.2216474 0.2555160 1.205032 557
[5] 0.07483477 0.3867578 0.1934926 1.513634 736
[6] 0.07483477 0.2928770 0.2555160 1.513634 736

```

```
data("Adult") inspect(Adult[1:10])
```

```

rul<-apriori(Adult,parameter=list(support=0.6,conf=0.6,minlen=2))
inspect(rul)

```

```
Parameter specification:
```

```

confidence minval smax arem  aval originalSupport
0.6      0.1      1 none FALSE      TRUE
maxtime support minlen maxlen target  ext
5       0.6       2      10  rules TRUE

```

```
Algorithmic control:
```

```

filter tree heap memopt load sort verbose
0.1 TRUE TRUE  FALSE TRUE    2    TRUE

```

```
Absolute minimum support count: 29305
```

```

set item appearances ...[0 item(s)] done [0.00s]. set transactions
...[115 item(s), 48842 transaction(s)] done [0.09s]. sorting and
recoding items ... [6 item(s)] done [0.01s]. creating transaction
tree ... done [0.03s]. checking subsets of size 1 2 3 4 done
[0.00s]. writing ... [41 rule(s)] done [0.00s]. creating S4 object
... done [0.00s].

```

```
> inspect(rul)
```

```

lhs                      rhs
support confidence  coverage    lift count

```

```

[1] {sex=Male} => {capital-gain=None}
0.6050735 0.9051455 0.6684820 0.9866565 29553
[2] {capital-gain=None} => {sex=Male}
0.6050735 0.6595621 0.9173867 0.9866565 29553
[3] {sex=Male} => {capital-loss=None}
0.6331027 0.9470750 0.6684820 0.9934931 30922
[4] {capital-loss=None} => {sex=Male}
0.6331027 0.6641323 0.9532779 0.9934931 30922
[5] {workclass=Private} => {native-country=United-
States} 0.6171942 0.8890757 0.6941976 0.9906971 30145
[6] {native-country=United-States} => {workclass=Private}
0.6171942 0.6877396 0.8974243 0.9906971 30145
[7] {workclass=Private} => {capital-gain=None}
0.6413742 0.9239073 0.6941976 1.0071078 31326
[8] {capital-gain=None} => {workclass=Private}
0.6413742 0.6991318 0.9173867 1.0071078 31326
[9] {workclass=Private} => {capital-loss=None}
0.6639982 0.9564974 0.6941976 1.0033773 32431
[10] {capital-loss=None} => {workclass=Private}
0.6639982 0.6965421 0.9532779 1.0033773 32431
[11] {race=White} => {native-country=United-
States}
0.7881127 0.9217231 0.8550428 1.0270761 38493
[12] {native-country=United-States} => {race=White}
0.7881127 0.8781940 0.8974243 1.0270761 38493
[13] {race=White} => {capital-gain=None}
0.7817862 0.9143240 0.8550428 0.9966616 38184
[14] {capital-gain=None} => {race=White}
0.7817862 0.8521883 0.9173867 0.9966616 38184
[15] {race=White} => {capital-loss=None}
0.8136849 0.9516307 0.8550428 0.9982720 39742
[16] {capital-loss=None} => {race=White}
0.8136849 0.8535653 0.9532779 0.9982720 39742
[17] {native-country=United-States} => {capital-gain=None}
0.8219565 0.9159062 0.8974243 0.9983862 40146
[18] {capital-gain=None} => {native-country=United-
States} 0.8219565 0.8959761 0.9173867 0.9983862 40146
[19] {native-country=United-States} => {capital-loss=None}
0.8548380 0.9525461 0.8974243 0.9992323 41752
[20] {capital-loss=None} => {native-country=United-
States} 0.8548380 0.8967354 0.9532779 0.9992323 41752
[21] {capital-gain=None} => {capital-loss=None}
0.8706646 0.9490705 0.9173867 0.9955863 42525
[22] {capital-loss=None} => {capital-gain=None}
0.8706646 0.9133376 0.9532779 0.9955863 42525
[23] {workclass=Private,
capital-gain=None} => {capital-loss=None}
0.6111748 0.9529145 0.6413742 0.9996188 29851
[24] {workclass=Private,
capital-loss=None} => {capital-gain=None}

```

```

0.6111748 0.9204465 0.6639982 1.0033354 29851
[25] {capital-gain=None,
      capital-loss=None}          => {workclass=Private}
0.6111748 0.7019636 0.8706646 1.0111869 29851
[26] {race=White,
      native-country=United-States} => {capital-gain=None}
0.7194628 0.9128933 0.7881127 0.9951019 35140
[27] {race=White,
      capital-gain=None}          => {native-country=United-
      States}
0.7194628 0.9202807 0.7817862 1.0254689 35140
[28] {capital-gain=None,
      native-country=United-States} => {race=White}
0.7194628 0.8753051 0.8219565 1.0236975 35140
[29] {race=White,
      native-country=United-States} => {capital-loss=None}
0.7490480 0.9504325 0.7881127 0.9970152 36585
[30] {race=White,
      capital-loss=None}          => {native-country=United-
      States}
0.7490480 0.9205626 0.8136849 1.0257830 36585
[31] {capital-loss=None,
      native-country=United-States} => {race=White}
0.7490480 0.8762454 0.8548380 1.0247972 36585
[32] {race=White,
      capital-gain=None}          => {capital-loss=None}
0.7404283 0.9470983 0.7817862 0.9935175 36164
[33] {race=White,
      capital-loss=None}          => {capital-gain=None}
0.7404283 0.9099693 0.8136849 0.9919147 36164
[34] {capital-gain=None,
      capital-loss=None}          => {race=White}
0.7404283 0.8504174 0.8706646 0.9945905 36164
[35] {capital-gain=None,
      native-country=United-States} => {capital-loss=None}
0.7793702 0.9481891 0.8219565 0.9946618 38066
[36] {capital-loss=None,
      native-country=United-States} => {capital-gain=None}
0.7793702 0.9117168 0.8548380 0.9938195 38066
[37] {capital-gain=None,
      capital-loss=None}          => {native-country=United-
      States}
0.7793702 0.8951440 0.8706646 0.9974590 38066
[38] {race=White,
      capital-gain=None,
      native-country=United-States} => {capital-loss=None}
0.6803980 0.9457029 0.7194628 0.9920537 33232
[39] {race=White,
      capital-loss=None,
      native-country=United-States} => {capital-gain=None}

```

```

0.6803980 0.9083504 0.7490480 0.9901500 33232
[40] {race=White,
      capital-gain=None,
      capital-loss=None} => {native-country=United-
      States}
0.6803980 0.9189249 0.7404283 1.0239581 33232
[41] {capital-gain=None,
      capital-loss=None,
      native-country=United-States} => {race=White}
0.6803980 0.8730100 0.7793702 1.0210133 33232

```

Question 5

Use Naive bayes, K-nearest, and Decision tree classification algorithms and build classifiers.

Divide the dataset into training and test set. Compare the accuracy of the different classifiers under the following situations:

5.1

- a) Training set=75% Test set= 25%
- b) Training set = 66.6% (2/3rd of total), Test set = 33.3%

5.2 Training set is chosen by

- i) holdout method
- ii) Random subsampling
- iii) Cross-Validation.

Compare the accuracy of the classifiers obtained. 5.3 Data is scaled to standard format.

NAÏVE BAYES

```

library(caret) library(klaR)
library(rpart)
library(rpart.plot)
library(class) library(plyr)

```

a) TRAINING SET 75%

```

s<-createDataPartition(iris$Species,p =
0.75,list=F) iris_train<- iris[s,] iris_train

```

OUTPUT:

```

Sepal.Length Sepal.Width Petal.Length
1           5.1           3.5           1.4
2           4.9           3.0           1.4
3           4.7           3.2           1.3
6           5.4           3.9           1.7
7           4.6           3.4           1.4

```

8	5.0	3.4	1.5
9	4.4	2.9	1.4
10	4.9	3.1	1.5
12	4.8	3.4	1.6
14	4.3	3.0	1.1
15	5.8	4.0	1.2
16	5.7	4.4	1.5
17	5.4	3.9	1.3
18	5.1	3.5	1.4
19	5.7	3.8	1.7
21	5.4	3.4	1.7
22	5.1	3.7	1.5
23	4.6	3.6	1.0
24	5.1	3.3	1.7
25	4.8	3.4	1.9
27	5.0	3.4	1.6
28	5.2	3.5	1.5
29	5.2	3.4	1.4
30	4.7	3.2	1.6
32	5.4	3.4	1.5
34	5.5	4.2	1.4
35	4.9	3.1	1.5
36	5.0	3.2	1.2
39	4.4	3.0	1.3
40	5.1	3.4	1.5
42	4.5	2.3	1.3
43	4.4	3.2	1.3
44	5.0	3.5	1.6
45	5.1	3.8	1.9
46	4.8	3.0	1.4
48	4.6	3.2	1.4
49	5.3	3.7	1.5
50	5.0	3.3	1.4
51	7.0	3.2	4.7
52	6.4	3.2	4.5
53	6.9	3.1	4.9
54	5.5	2.3	4.0
56	5.7	2.8	4.5
61	5.0	2.0	3.5
62	5.9	3.0	4.2
63	6.0	2.2	4.0
64	6.1	2.9	4.7
65	5.6	2.9	3.6
66	6.7	3.1	4.4
67	5.6	3.0	4.5
68	5.8	2.7	4.1
69	6.2	2.2	4.5
70	5.6	2.5	3.9
71	5.9	3.2	4.8

72	6.1	2.8	4.0
73	6.3	2.5	4.9
74	6.1	2.8	4.7
75	6.4	2.9	4.3
76	6.6	3.0	4.4
77	6.8	2.8	4.8
79	6.0	2.9	4.5
80	5.7	2.6	3.5
81	5.5	2.4	3.8
82	5.5	2.4	3.7
83	5.8	2.7	3.9
84	6.0	2.7	5.1
87	6.7	3.1	4.7
89	5.6	3.0	4.1
90	5.5	2.5	4.0
93	5.8	2.6	4.0
94	5.0	2.3	3.3
95	5.6	2.7	4.2
97	5.7	2.9	4.2
98	6.2	2.9	4.3
99	5.1	2.5	3.0
100	5.7	2.8	4.1
101	6.3	3.3	6.0
102	5.8	2.7	5.1
103	7.1	3.0	5.9
104	6.3	2.9	5.6
105	6.5	3.0	5.8
106	7.6	3.0	6.6
107	4.9	2.5	4.5
108	7.3	2.9	6.3
109	6.7	2.5	5.8
110	7.2	3.6	6.1
111	6.5	3.2	5.1
112	6.4	2.7	5.3
115	5.8	2.8	5.1
116	6.4	3.2	5.3
119	7.7	2.6	6.9
120	6.0	2.2	5.0
121	6.9	3.2	5.7
122	5.6	2.8	4.9
123	7.7	2.8	6.7
124	6.3	2.7	4.9
125	6.7	3.3	5.7
127	6.2	2.8	4.8
128	6.1	3.0	4.9
130	7.2	3.0	5.8
131	7.4	2.8	6.1
132	7.9	3.8	6.4
133	6.4	2.8	5.6

136	7.7	3.0	6.1
138	6.4	3.1	5.5
140	6.9	3.1	5.4
142	6.9	3.1	5.1
143	5.8	2.7	5.1
144	6.8	3.2	5.9
146	6.7	3.0	5.2
147	6.3	2.5	5.0
148	6.5	3.0	5.2
149	6.2	3.4	5.4
150	5.9	3.0	5.1

	Petal.Width	Species
1	0.2	setosa
2	0.2	setosa
3	0.2	setosa
6	0.4	setosa
7	0.3	setosa
8	0.2	setosa
9	0.2	setosa
10	0.1	setosa
12	0.2	setosa
14	0.1	setosa
15	0.2	setosa
16	0.4	setosa
17	0.4	setosa
18	0.3	setosa
19	0.3	setosa
21	0.2	setosa
22	0.4	setosa
23	0.2	setosa
24	0.5	setosa
25	0.2	setosa
27	0.4	setosa
28	0.2	setosa
29	0.2	setosa
30	0.2	setosa
32	0.4	setosa
34	0.2	setosa
35	0.2	setosa
36	0.2	setosa
39	0.2	setosa
40	0.2	setosa
42	0.3	setosa
43	0.2	setosa
44	0.6	setosa
45	0.4	setosa
46	0.3	setosa
48	0.2	setosa
49	0.2	setosa
50	0.2	setosa

51	1.4 versicolor	
52	1.5 versicolor	
53	1.5 versicolor	
54	1.3 versicolor	
56	1.3 versicolor	
61	1.0 versicolor	
62	1.5 versicolor	
63	1.0 versicolor	
64	1.4 versicolor	
65	1.3 versicolor	
66	1.4 versicolor	
67	1.5 versicolor	
68	1.0 versicolor	
69	1.5 versicolor	
70	1.1 versicolor	
71	1.8 versicolor	
72	1.3 versicolor	
73	1.5 versicolor74	1.2 versicolor 75
	1.3 versicolor	
76	1.4 versicolor	
77	1.4 versicolor	
79	1.5 versicolor 80	1.0 versicolor
81	1.1 versicolor	
82	1.0 versicolor	
83	1.2 versicolor	
84	1.6 versicolor	
87	1.5 versicolor	
89	1.3 versicolor	
90	1.3 versicolor	
93	1.2 versicolor	
94	1.0 versicolor	
95	1.3 versicolor	
97	1.3 versicolor	
98	1.3 versicolor	
99	1.1 versicolor	
100	1.3 versicolor	
101	2.5 virginica	
102	1.9 virginica	
103	2.1 virginica	
104	1.8 virginica	
105	2.2 virginica	
106	2.1 virginica	
107	1.7 virginica	
108	1.8 virginica	
109	1.8 virginica	
110	2.5 virginica	
111	2.0 virginica	
112	1.9 virginica	
115	2.4 virginica	

```

116      2.3 virginica
119      2.3 virginica
120      1.5 virginica
121      2.3 virginica
122      2.0 virginica
123      2.0 virginica
124      1.8 virginica
125      2.1 virginica
127      1.8 virginica
128      1.8 virginica
130      1.6 virginica
131      1.9 virginica
132      2.0 virginica
133      2.2 virginica
136      2.3 virginica
138      1.8 virginica
140      2.1 virginica
142      2.3 virginica
143      1.9 virginica
144      2.3 virginica
146      2.3 virginica
147      1.9 virginica148      2.0 virginica
149      2.3 virginica
150      1.8 virginica

```

```
iris_test<- iris[-s,] iris_test
```

OUTPUT:

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
4	4.6	3.1	1.5	0.2	setosa
5	5.0	3.6	1.4	0.2	setosa
11	5.4	3.7	1.5	0.2	setosa
13	4.8	3.0	1.4	0.1	setosa
20	5.1	3.8	1.5	0.3	setosa
26	5.0	3.0	1.6	0.2	setosa
31	4.8	3.1	1.6	0.2	setosa
33	5.2	4.1	1.5	0.1	setosa
37	5.5	3.5	1.3	0.2	setosa
38	4.9	3.6	1.4	0.1	setosa
41	5.0	3.5	1.3	0.3	setosa
47	5.1	3.8	1.6	0.2	setosa
55	6.5	2.8	4.6	1.5	versicolor
57	6.3	3.3	4.7	1.6	versicolor
58	4.9	2.4	3.3	1.0	versicolor
59	6.6	2.9	4.6	1.3	versicolor
60	5.2	2.7	3.9	1.4	versicolor
78	6.7	3.0	5.0	1.7	versicolor
85	5.4	3.0	4.5	1.5	versicolor
86	6.0	3.4	4.5	1.6	versicolor

88	6.3	2.3	4.4	1.3	versicolor
91	5.5	2.6	4.4	1.2	versicolor
92	6.1	3.0	4.6	1.4	versicolor
96	5.7	3.0	4.2	1.2	versicolor
113	6.8	3.0	5.5	2.1	virginica
114	5.7	2.5	5.0	2.0	virginica
117	6.5	3.0	5.5	1.8	virginica
118	7.7	3.8	6.7	2.2	virginica
126	7.2	3.2	6.0	1.8	virginica
129	6.4	2.8	5.6	2.1	virginica
134	6.3	2.8	5.1	1.5	virginica
135	6.1	2.6	5.6	1.4	virginica
137	6.3	3.4	5.6	2.4	virginica
139	6.0	3.0	4.8	1.8	virginica
141	6.7	3.1	5.6	2.4	virginica
145	6.7	3.3	5.7	2.5	virginica

```
model<- NaiveBayes(Species~., data=iris_train)
```

```
model
```

```
apriori
```

```
grouping
```

```
      setosa versicolor  virginica
0.3333333  0.3333333  0.3333333
```

```
$tables
```

```
$tables$Sepal.Length
```

```
 [,1]      [,2] setosa
4.997368 0.3809567 versicolor
5.936842 0.5004976 virginica
6.605263 0.6685795
```

```
$tables$Sepal.Width
```

```
 [,1]      [,2] setosa
3.410526 0.3881971 versicolor
2.739474 0.3071574 virginica
2.952632 0.3125629
```

```
$tables$Petal.Length
```

```
 [,1]      [,2] setosa
1.460526 0.1910702 versicolor
4.218421 0.4780904 virginica
5.550000 0.5764523
```

```
$tables$Petal.Width
```

```
 [,1]      [,2] setosa
0.2631579 0.1100885 versicolor
1.3052632 0.1944464 virginica
2.0342105 0.2507031
```

```
$levels  
[1] "setosa"      "versicolor" "virginica"
```

```
$call  
NaiveBayes.default(x = X, grouping = Y)
```

```
$x  
      Sepal.Length Sepal.Width Petal.Length Petal.Width  
1           5.1           3.5           1.4           0.2  
2           4.9           3.0           1.4           0.2  
3           4.7           3.2           1.3           0.2 6  
           5.4           3.9           1.7           0.4  
7           4.6           3.4           1.4           0.3  
8           5.0           3.4           1.5           0.2  
9           4.4           2.9           1.4           0.2  
10          4.9           3.1           1.5           0.1  
12          4.8           3.4           1.6           0.2 14  
4.3         3.0           1.1           0.1  
15          5.8           4.0           1.2           0.2  
16          5.7           4.4           1.5           0.4  
17          5.4           3.9           1.3           0.4 18  
           5.1           3.5           1.4           0.3  
19          5.7           3.8           1.7           0.3  
21          5.4           3.4           1.7           0.2  
22          5.1           3.7           1.5           0.4  
23          4.6           3.6           1.0           0.2  
24          5.1           3.3           1.7           0.5  
25          4.8           3.4           1.9           0.2  
27          5.0           3.4           1.6           0.4  
28          5.2           3.5           1.5           0.2  
29          5.2           3.4           1.4           0.2  
30          4.7           3.2           1.6           0.2  
32          5.4           3.4           1.5           0.4  
34          5.5           4.2           1.4           0.2  
35          4.9           3.1           1.5           0.2  
36          5.0           3.2           1.2           0.2  
39          4.4           3.0           1.3           0.2  
40          5.1           3.4           1.5           0.2  
42          4.5           2.3           1.3           0.3  
43          4.4           3.2           1.3           0.2  
44          5.0           3.5           1.6           0.6  
45          5.1           3.8           1.9           0.4  
46          4.8           3.0           1.4           0.3  
48          4.6           3.2           1.4           0.2  
49          5.3           3.7           1.5           0.2  
50          5.0           3.3           1.4           0.2  
51          7.0           3.2           4.7           1.4  
52          6.4           3.2           4.5           1.5  
53          6.9           3.1           4.9           1.5  
54          5.5           2.3           4.0           1.3
```

56	5.7	2.8	4.5	1.3
61	5.0	2.0	3.5	1.0
62	5.9	3.0	4.2	1.5
63	6.0	2.2	4.0	1.0
64	6.1	2.9	4.7	1.4
65	5.6	2.9	3.6	1.3
66	6.7	3.1	4.4	1.4
67	5.6	3.0	4.5	1.5
68	5.8	2.7	4.1	1.0
69	6.2	2.2	4.5	1.5
70	5.6	2.5	3.9	1.1
71	5.9	3.2	4.8	1.8
72	6.1	2.8	4.0	1.3
73	6.3	2.5	4.9	1.5 74
	6.1	2.8	4.7	1.2 75
	6.4	2.9	4.3	1.3
76	6.6	3.0	4.4	1.4 77
6.8	2.8	4.8	1.4 79	6.0
2.9	4.5	1.5		
80	5.7	2.6	3.5	1.0
81	5.5	2.4	3.8	1.1
82	5.5	2.4	3.7	1.0
83	5.8	2.7	3.9	1.2
84	6.0	2.7	5.1	1.6
87	6.7	3.1	4.7	1.5
89	5.6	3.0	4.1	1.3
90	5.5	2.5	4.0	1.3
93	5.8	2.6	4.0	1.2
94	5.0	2.3	3.3	1.0
95	5.6	2.7	4.2	1.3
97	5.7	2.9	4.2	1.3
98	6.2	2.9	4.3	1.3
99	5.1	2.5	3.0	1.1
100	5.7	2.8	4.1	1.3
101	6.3	3.3	6.0	2.5
102	5.8	2.7	5.1	1.9
103	7.1	3.0	5.9	2.1
104	6.3	2.9	5.6	1.8
105	6.5	3.0	5.8	2.2
106	7.6	3.0	6.6	2.1
107	4.9	2.5	4.5	1.7
108	7.3	2.9	6.3	1.8
109	6.7	2.5	5.8	1.8
110	7.2	3.6	6.1	2.5
111	6.5	3.2	5.1	2.0
112	6.4	2.7	5.3	1.9
115	5.8	2.8	5.1	2.4
116	6.4	3.2	5.3	2.3
119	7.7	2.6	6.9	2.3
120	6.0	2.2	5.0	1.5

121	6.9	3.2	5.7	2.3
122	5.6	2.8	4.9	2.0
123	7.7	2.8	6.7	2.0
124	6.3	2.7	4.9	1.8
125	6.7	3.3	5.7	2.1
127	6.2	2.8	4.8	1.8
128	6.1	3.0	4.9	1.8
130	7.2	3.0	5.8	1.6
131	7.4	2.8	6.1	1.9
132	7.9	3.8	6.4	2.0
133	6.4	2.8	5.6	2.2 136
	7.7	3.0	6.1	2.3
138	6.4	3.1	5.5	1.8
140	6.9	3.1	5.4	2.1
142	6.9	3.1	5.1	2.3
143	5.8	2.7	5.1	1.9 144
	6.8	3.2	5.9	2.3 146
	6.7	3.0	5.2	2.3
147	6.3	2.5	5.0	1.9
148	6.5	3.0	5.2	2.0
149	6.2	3.4	5.4	2.3
150	5.9	3.0	5.1	1.8

```
$usekernel
```

```
[1] FALSE
```

```
$varnames
```

```
[1] "Sepal.Length" "Sepal.Width" "Petal.Length" "Petal.Width"
attr("class")
```

```
[1] "NaiveBayes"
```

```
pred<- predict(model, iris_test) pred$class
```

4	5	11	13	20	26	31
33	37					
setosa	setosa	setosa	setosa	setosa	setosa	setosa
setosa	setosa	setosa				
	38	41	47	55	57	58
59	60	78				
setosa	setosa	setosa	versicolor	versicolor	versicolor	
versicolor	versicolor	virginica				
	85	86	88	91	92	96
113	114	117				
versicolor	versicolor	versicolor	versicolor	versicolor	versicolor	
virginica	virginica	virginica				
	118	126	129	134	135	137
139	141	145				
virginica	virginica	virginica	versicolor	versicolor	virginica	
virginica	virginica	virginica				

Levels: setosa versicolor virginica

```
conf<- confusionMatrix(pred$class ,iris_test$Species)$table
conf
```

```
Reference
Prediction  setosa versicolor virginica
setosa      12         0         0
versicolor   0        11         2
virginica    0         1        10
```

```
conf1<-as.matrix(conf)
d<-diag(conf1) s<-sum(d)
s1<-sum(conf1)
accuracy1<-(s/s1)*100
accuracy1
91.66667
```

b) TRAINING SET 66.6%(2/3)

```
s<-createDataPartition(iris$Species,p = 0.66,list=F)
iris_train<- iris[s,] iris_train
```

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
2	4.9	3.0	1.4	0.2	setosa
3	4.7	3.2	1.3	0.2	setosa
4	4.6	3.1	1.5	0.2	setosa
5	5.0	3.6	1.4	0.2	setosa
6	5.4	3.9	1.7	0.4	setosa
7	4.6	3.4	1.4	0.3	setosa
11	5.4	3.7	1.5	0.2	setosa
12	4.8	3.4	1.6	0.2	setosa
13	4.8	3.0	1.4	0.1	setosa
14	4.3	3.0	1.1	0.1	setosa
15	5.8	4.0	1.2	0.2	setosa
16	5.7	4.4	1.5	0.4	setosa
17	5.4	3.9	1.3	0.4	setosa
19	5.7	3.8	1.7	0.3	setosa
20	5.1	3.8	1.5	0.3	setosa
21	5.4	3.4	1.7	0.2	setosa
24	5.1	3.3	1.7	0.5	setosa
26	5.0	3.0	1.6	0.2	setosa
28	5.2	3.5	1.5	0.2	setosa
29	5.2	3.4	1.4	0.2	setosa
30	4.7	3.2	1.6	0.2	setosa
31	4.8	3.1	1.6	0.2	setosa
32	5.4	3.4	1.5	0.4	setosa
33	5.2	4.1	1.5	0.1	setosa
35	4.9	3.1	1.5	0.2	setosa
37	5.5	3.5	1.3	0.2	setosa

39	4.4	3.0	1.3	0.2	setosa
40	5.1	3.4	1.5	0.2	setosa
41	5.0	3.5	1.3	0.3	setosa
42	4.5	2.3	1.3	0.3	
	setosa45		5.1	3.8	1.9
	0.4	setosa			
46	4.8	3.0	1.4	0.3	setosa
48	4.6	3.2	1.4	0.2	setosa
51	7.0	3.2	4.7	1.4	versicolor
52	6.4	3.2	4.5	1.5	versicolor
53	6.9	3.1	4.9	1.5	versicolor
54	5.5	2.3	4.0	1.3	
	versicolor55		6.5	2.8	4.6
	1.5 versicolor 57		6.3	3.3	
	4.7	1.6 versicolor 58		4.9	
	2.4	3.3	1.0 versicolor		
61	5.0	2.0	3.5	1.0	versicolor
64	6.1	2.9	4.7	1.4	versicolor
65	5.6	2.9	3.6	1.3	versicolor
66	6.7	3.1	4.4	1.4	versicolor
67	5.6	3.0	4.5	1.5	versicolor
68	5.8	2.7	4.1	1.0	versicolor
69	6.2	2.2	4.5	1.5	versicolor
70	5.6	2.5	3.9	1.1	versicolor
73	6.3	2.5	4.9	1.5	versicolor
74	6.1	2.8	4.7	1.2	versicolor
75	6.4	2.9	4.3	1.3	versicolor
77	6.8	2.8	4.8	1.4	versicolor
78	6.7	3.0	5.0	1.7	versicolor
79	6.0	2.9	4.5	1.5	versicolor
81	5.5	2.4	3.8	1.1	versicolor
82	5.5	2.4	3.7	1.0	versicolor
86	6.0	3.4	4.5	1.6	versicolor
87	6.7	3.1	4.7	1.5	versicolor
89	5.6	3.0	4.1	1.3	versicolor
90	5.5	2.5	4.0	1.3	versicolor
92	6.1	3.0	4.6	1.4	versicolor
93	5.8	2.6	4.0	1.2	versicolor
94	5.0	2.3	3.3	1.0	versicolor
95	5.6	2.7	4.2	1.3	versicolor
98	6.2	2.9	4.3	1.3	versicolor
100	5.7	2.8	4.1	1.3	versicolor
102	5.8	2.7	5.1	1.9	virginica
103	7.1	3.0	5.9	2.1	virginica
104	6.3	2.9	5.6	1.8	virginica
106	7.6	3.0	6.6	2.1	virginica
109	6.7	2.5	5.8	1.8	virginica
110	7.2	3.6	6.1	2.5	virginica
111	6.5	3.2	5.1	2.0	virginica
112	6.4	2.7	5.3	1.9	virginica

```

113      6.8      3.0      5.5      2.1 virginica
114      5.7      2.5      5.0      2.0 virginica
115      5.8      2.8      5.1      2.4 virginica
116      6.4      3.2      5.3      2.3 virginica
117      6.5      3.0      5.5      1.8 virginica
118      7.7      3.8      6.7      2.2 virginica
119      7.7      2.6      6.9      2.3
      virginica120      6.0      2.2      5.0
      1.5 virginica
121      6.9      3.2      5.7      2.3 virginica
122      5.6      2.8      4.9      2.0 virginica
125      6.7      3.3      5.7      2.1 virginica
129      6.4      2.8      5.6      2.1 virginica
130      7.2      3.0      5.8      1.6 virginica
131      7.4      2.8      6.1      1.9 virginica
132      7.9      3.8      6.4      2.0 virginica
133      6.4      2.8      5.6      2.2 virginica
134      6.3      2.8      5.1      1.5 virginica
137      6.3      3.4      5.6      2.4 virginica
138      6.4      3.1      5.5      1.8
      virginica139      6.0      3.0      4.8
      1.8 virginica 144      6.8      3.2
      5.9      2.3 virginica
146      6.7      3.0      5.2      2.3 virginica
148      6.5      3.0      5.2      2.0 virginica
149      6.2      3.4      5.4      2.3 virginica
150      5.9      3.0      5.1      1.8 virginica

```

```
iris_test<- iris[-s,] iris_test
```

```

      Sepal.Length Sepal.Width Petal.Length Petal.Width Species
1          5.1          3.5          1.4          0.2    setosa
8          5.0          3.4          1.5          0.2    setosa
9          4.4          2.9          1.4          0.2    setosa
10         4.9          3.1          1.5          0.1    setosa
18         5.1          3.5          1.4          0.3    setosa
22         5.1          3.7          1.5          0.4    setosa
23         4.6          3.6          1.0          0.2    setosa
25         4.8          3.4          1.9          0.2    setosa
27         5.0          3.4          1.6          0.4    setosa
34         5.5          4.2          1.4          0.2    setosa
36         5.0          3.2          1.2          0.2    setosa
38         4.9          3.6          1.4          0.1    setosa
43         4.4          3.2          1.3          0.2    setosa
44         5.0          3.5          1.6          0.6    setosa
47         5.1          3.8          1.6          0.2    setosa
49         5.3          3.7          1.5          0.2    setosa
50         5.0          3.3          1.4          0.2    setosa
56         5.7          2.8          4.5          1.3 versicolor
59         6.6          2.9          4.6          1.3 versicolor

```

60	5.2	2.7	3.9	1.4	versicolor
62	5.9	3.0	4.2	1.5	versicolor
63	6.0	2.2	4.0	1.0	versicolor
71	5.9	3.2	4.8	1.8	versicolor
72	6.1	2.8	4.0	1.3	versicolor
76	6.6	3.0	4.4	1.4	versicolor
80	5.7	2.6	3.5	1.0	versicolor
83	5.8	2.7	3.9	1.2	versicolor
84	6.0	2.7	5.1	1.6	
	versicolor85		5.4	3.0	4.5
	1.5				versicolor
88	6.3	2.3	4.4	1.3	versicolor
91	5.5	2.6	4.4	1.2	versicolor
96	5.7	3.0	4.2	1.2	versicolor
97	5.7	2.9	4.2	1.3	versicolor
99	5.1	2.5	3.0	1.1	versicolor
101	6.3	3.3	6.0	2.5	virginica
105	6.5	3.0	5.8	2.2	virginica
107	4.9	2.5	4.5	1.7	virginica
108	7.3	2.9	6.3	1.8	virginica
123	7.7	2.8	6.7	2.0	virginica
124	6.3	2.7	4.9	1.8	
	virginica126		7.2	3.2	6.0
	1.8				virginica
127	6.2	2.8	4.8	1.8	virginica
128	6.1	3.0	4.9	1.8	virginica
135	6.1	2.6	5.6	1.4	virginica
136	7.7	3.0	6.1	2.3	virginica
140	6.9	3.1	5.4	2.1	virginica
141	6.7	3.1	5.6	2.4	virginica
142	6.9	3.1	5.1	2.3	virginica
143	5.8	2.7	5.1	1.9	virginica
145	6.7	3.3	5.7	2.5	virginica
147	6.3	2.5	5.0	1.9	virginica

```
model<- NaiveBayes(Species~., data=iris_train)
```

```
model $apriori grouping
```

```
setosa versicolor virginica
0.3333333 0.3333333 0.3333333
```

```
$tables
```

```
$tables$Sepal.Length
```

```
[,1] [,2] setosa
```

```
5.033333 0.3845994 versicolor
```

```
5.987879 0.5577620 virginica
```

```
6.600000 0.6093029
```

```
$tables$Sepal.Width
```

```
[,1] [,2] setosa
```

```
3.406061 0.4167879 versicolor
```

```
2.775758 0.3400646 virginica
3.003030 0.3513233
```

```
$tables$Petal.Length
[,1]      [,2] setosa
1.469697 0.1667424 versicolor
4.284848 0.4657797 virginica
5.578788 0.5295975
```

```
$tables$Petal.Width
      [,1]      [,2] setosa
0.2484848 0.09721501 versicolor
1.3303030 0.19603069 virginica
2.0333333 0.25819889
```

```
$levels
[1] "setosa"      "versicolor" "virginica"
```

```
$call
NaiveBayes.default(x = X, grouping = Y)
```

```
$x
      Sepal.Length Sepal.Width Petal.Length Petal.Width
2           4.9         3.0         1.4         0.2
3           4.7         3.2         1.3         0.2
4           4.6         3.1         1.5         0.2
5           5.0         3.6         1.4         0.2
6           5.4         3.9         1.7         0.4
7           4.6         3.4         1.4         0.3
11          5.4         3.7         1.5         0.2
12          4.8         3.4         1.6         0.2
13          4.8         3.0         1.4         0.1
14          4.3         3.0         1.1         0.1
15          5.8         4.0         1.2         0.2
16          5.7         4.4         1.5         0.4
17          5.4         3.9         1.3         0.4
19          5.7         3.8         1.7         0.3
20          5.1         3.8         1.5         0.3
21          5.4         3.4         1.7         0.2
24          5.1         3.3         1.7         0.5
26          5.0         3.0         1.6         0.2
28          5.2         3.5         1.5         0.2
29          5.2         3.4         1.4         0.2
30          4.7         3.2         1.6         0.2
31          4.8         3.1         1.6         0.2
32          5.4         3.4         1.5         0.4
33          5.2         4.1         1.5         0.1
35          4.9         3.1         1.5         0.2
37          5.5         3.5         1.3         0.2
```

39	4.4	3.0	1.3	0.2
40	5.1	3.4	1.5	0.2
41	5.0	3.5	1.3	0.3
42	4.5	2.3	1.3	0.3
45	5.1	3.8	1.9	0.4
46	4.8	3.0	1.4	0.3
48	4.6	3.2	1.4	0.2
51	7.0	3.2	4.7	1.4
52	6.4	3.2	4.5	1.5
53	6.9	3.1	4.9	1.5
54	5.5	2.3	4.0	1.3
55	6.5	2.8	4.6	1.5
57	6.3	3.3	4.7	1.6
58	4.9	2.4	3.3	1.0
61	5.0	2.0	3.5	1.0
64	6.1	2.9	4.7	1.4
65	5.6	2.9	3.6	1.3
66	6.7	3.1	4.4	1.4
67	5.6	3.0	4.5	1.5
68	5.8	2.7	4.1	1.0
69	6.2	2.2	4.5	1.5
70	5.6	2.5	3.9	1.1
73	6.3	2.5	4.9	1.5
74	6.1	2.8	4.7	1.2
75	6.4	2.9	4.3	1.3
77	6.8	2.8	4.8	1.4
78	6.7	3.0	5.0	1.7
79	6.0	2.9	4.5	1.5
81	5.5	2.4	3.8	1.1
82	5.5	2.4	3.7	1.0
86	6.0	3.4	4.5	1.6
87	6.7	3.1	4.7	1.5
89	5.6	3.0	4.1	1.3
90	5.5	2.5	4.0	1.3
92	6.1	3.0	4.6	1.4
93	5.8	2.6	4.0	1.2
94	5.0	2.3	3.3	1.0
95	5.6	2.7	4.2	1.3
98	6.2	2.9	4.3	1.3
100	5.7	2.8	4.1	1.3
102	5.8	2.7	5.1	1.9
103	7.1	3.0	5.9	2.1
104	6.3	2.9	5.6	1.8
106	7.6	3.0	6.6	2.1
109	6.7	2.5	5.8	1.8
110	7.2	3.6	6.1	2.5
111	6.5	3.2	5.1	2.0
112	6.4	2.7	5.3	1.9
113	6.8	3.0	5.5	2.1

```

114      5.7      2.5      5.0      2.0
115      5.8      2.8      5.1      2.4
116      6.4      3.2      5.3      2.3
117      6.5      3.0      5.5      1.8
118      7.7      3.8      6.7      2.2
119      7.7      2.6      6.9      2.3
120      6.0      2.2      5.0      1.5
121      6.9      3.2      5.7      2.3
122      5.6      2.8      4.9      2.0
125      6.7      3.3      5.7      2.1
129      6.4      2.8      5.6      2.1
130      7.2      3.0      5.8      1.6
131      7.4      2.8      6.1      1.9
132      7.9      3.8      6.4      2.0 133
      6.4      2.8      5.6      2.2
134      6.3      2.8      5.1      1.5
137      6.3      3.4      5.6      2.4
138      6.4      3.1      5.5      1.8
139      6.0      3.0      4.8      1.8
144      6.8      3.2      5.9      2.3
146      6.7      3.0      5.2      2.3
148      6.5      3.0      5.2      2.0
149      6.2      3.4      5.4      2.3
150      5.9      3.0      5.1      1.8

$usekernel
[1] FALSE

$varnames
[1] "Sepal.Length" "Sepal.Width" "Petal.Length" "Petal.Width"
attr(,"class")
[1] "NaiveBayes"

pred<- predict(model, iris_test) pred$class
      1      8      9      10      18      22
23      25      27
      setosa      setosa      setosa      setosa      setosa      setosa
setosa      setosa      setosa
      34      36      38      43      44      47
49      50      56
      setosa      setosa      setosa      setosa      setosa      setosa
setosa      setosa versicolor
      59      60      62      63      71      72
76      80      83
versicolor versicolor versicolor versicolor virginica versicolor
versicolor versicolor versicolor
      84      85      88      91      96      97
99      101      105
versicolor versicolor versicolor versicolor versicolor versicolor
versicolor virginica virginica

```

```

      107      108      123      124      126      127
128      135      136
versicolor virginica virginica virginica virginica virginica
virginica versicolor virginica
      140      141      142      143      145      147
virginica virginica virginica virginica virginica virginica
Levels: setosa versicolor virginica

```

```
conf<- confusionMatrix(pred$class ,iris_test$Species)$table
```

```
conf Reference
```

```

Prediction   setosa versicolor virginica
setosa        17         0         0
versicolor    0         16         2
virginica     0          1        15

```

```

conf1<-
as.matrix(conf) d<-
diag(conf1) s<-sum(d)
s1<-sum(conf1)
accuracy2<-(s/s1)*100
accuracy2
94.11765

```

```

if(accuracy1>accuracy2)
{
  cat("Training set of 75% records is better")
}
if(accuracy2>accuracy1)
{
  cat("Training set of 66.6% records is better")
}

```

```
Training set of 66.6% records is better
```

HOLD OUT METHOD

```

s<-sample(150,75)
iris_train<-iris[s,]
iris_test<-iris[-s,]
model<-NaiveBayes(Species~.,data=iris_train) pred<-
predict(model,iris_test)
conf<-confusionMatrix(pred$class ,iris_test$Species)$table
conf1<-as.matrix(conf)
acchld<-((sum(diag(conf1)))/sum(conf1))*100 cat("Accuracy
of holdout method is ",acchld)

```

```
Accuracy of holdout method is 94.66667
```

RANDOM SUBSAMPLING

```

i<-75 j<-1
acc<-c() for(i
in 75:100)
{   s<-sample(150,i)
iris_train<-iris[s,]
iris_test<-iris[-s,]
  model<-NaiveBayes(Species~.,data=iris_train)  pred<-
predict(model,iris_test)
  conf<-confusionMatrix(pred$class,iris_test$Species)$table
conf1<-as.matrix(conf)
  acc[j]<-c((sum(diag(conf1))/sum(conf1))*100,acc)
j=j+1 } accrs<-mean(acc)
cat("Accuracy of random subsampling is ",accrs)

```

Accuracy of random subsampling is 95.5321

CROSS VALIDATION METHOD

```

x=iris[,-5]
y=iris$Species
model = train(x,y,'nb',trControl=trainControl(method='cv',number=10))
cons<-table(predict(model$finalModel,x)$class,y) cons1<-as.matrix(cons)
accv<-(sum(diag(cons1))/sum(cons1))*100 cat("Accuracy
of cross validation is ",accv)

```

Accuracy of cross validation is 96

COMPARISON

```

greatest<-max(acchld,accrs,accv) if(greatest==acchld)
{
  cat("Holdout method does best classification")
}
if(greatest==accrs)
{
  cat("Random subsampling method does best classification")
}
if(greatest==accv)
{
  cat("Cross validation method does best classification")
}

```

Cross validation method does best classification

DATA SCALING

```

data_std <- function(x)
{
  (x-mean(x))/sd(x)
}
sapply(iris[,-5],data_std)

```


	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
[1,]	-0.89767388	1.01560199	-1.33575163	-1.3110521482
[2,]	-1.13920048	-0.13153881	-1.33575163	-1.3110521482
[3,]	-1.38072709	0.32731751	-1.39239929	-1.3110521482
[4,]	-1.50149039	0.09788935	-1.27910398	-1.3110521482
[5,]	-1.01843718	1.24503015	-1.33575163	-1.3110521482
[6,]	-0.53538397	1.93331463	-1.16580868	-1.0486667950
[7,]	-1.50149039	0.78617383	-1.33575163	-1.1798594716
[8,]	-1.01843718	0.78617383	-1.27910398	-1.3110521482
[9,]	-1.74301699	-0.36096697	-1.33575163	-1.3110521482
[10,]	-1.13920048	0.09788935	-1.27910398	-1.4422448248
[11,]	-0.53538397	1.47445831	-1.27910398	-1.3110521482
[12,]	-1.25996379	0.78617383	-1.22245633	-1.3110521482
[13,]	-1.25996379	-0.13153881	-1.33575163	-1.4422448248
[14,]	-1.86378030	-0.13153881	-1.50569459	-1.4422448248
[15,]	-0.05233076	2.16274279	-1.44904694	-1.3110521482
[16,]	-0.17309407	3.08045544	-1.27910398	-1.0486667950
[17,]	-0.53538397	1.93331463	-1.39239929	-1.0486667950
[18,]	-0.89767388	1.01560199	-1.33575163	-1.1798594716
[19,]	-0.17309407	1.70388647	-1.16580868	-1.1798594716
[20,]	-0.89767388	1.70388647	-1.27910398	-1.1798594716
[21,]	-0.53538397	0.78617383	-1.16580868	-1.3110521482
[22,]	-0.89767388	1.47445831	-1.27910398	-1.0486667950
[23,]	-1.50149039	1.24503015	-1.56234224	-1.3110521482
[24,]	-0.89767388	0.55674567	-1.16580868	-0.9174741184
[25,]	-1.25996379	0.78617383	-1.05251337	-1.3110521482
[26,]	-1.01843718	-0.13153881	-1.22245633	-1.3110521482
[27,]	-1.01843718	0.78617383	-1.22245633	-1.0486667950
[28,]	-0.77691058	1.01560199	-1.27910398	-1.3110521482
[29,]	-0.77691058	0.78617383	-1.33575163	-1.3110521482
[30,]	-1.38072709	0.32731751	-1.22245633	-1.3110521482
[31,]	-1.25996379	0.09788935	-1.22245633	-1.3110521482
[32,]	-0.53538397	0.78617383	-1.27910398	-1.0486667950
[33,]	-0.77691058	2.39217095	-1.27910398	-1.4422448248
[34,]	-0.41462067	2.62159911	-1.33575163	-1.3110521482
[35,]	-1.13920048	0.09788935	-1.27910398	-1.3110521482
[36,]	-1.01843718	0.32731751	-1.44904694	-1.3110521482
[37,]	-0.41462067	1.01560199	-1.39239929	-1.3110521482
[38,]	-1.13920048	1.24503015	-1.33575163	-1.4422448248
[39,]	-1.74301699	-0.13153881	-1.39239929	-1.3110521482
[40,]	-0.89767388	0.78617383	-1.27910398	-1.3110521482
[41,]	-1.01843718	1.01560199	-1.39239929	-1.1798594716
[42,]	-1.62225369	-1.73753594	-1.39239929	-1.1798594716
[43,]	-1.74301699	0.32731751	-1.39239929	-1.3110521482
[44,]	-1.01843718	1.01560199	-1.22245633	-0.7862814418
[45,]	-0.89767388	1.70388647	-1.05251337	-1.0486667950
[46,]	-1.25996379	-0.13153881	-1.33575163	-1.1798594716
[47,]	-0.89767388	1.70388647	-1.22245633	-1.3110521482

[48,] -1.50149039 0.32731751 -1.33575163 -1.3110521482 [49,]
-0.65614727 1.47445831 -1.27910398 -1.3110521482
[50,] -1.01843718 0.55674567 -1.33575163 -1.3110521482
[51,] 1.39682886 0.32731751 0.53362088 0.2632599711
[52,] 0.67224905 0.32731751 0.42032558 0.3944526477
[53,] 1.27606556 0.09788935 0.64691619 0.3944526477
[54,] -0.41462067 -1.73753594 0.13708732 0.1320672944
[55,] 0.79301235 -0.59039513 0.47697323 0.3944526477
[56,] -0.17309407 -0.59039513 0.42032558 0.1320672944
[57,] 0.55148575 0.55674567 0.53362088 0.5256453243
[58,] -1.13920048 -1.50810778 -0.25944625 -0.2615107354
[59,] 0.91377565 -0.36096697 0.47697323 0.1320672944
[60,] -0.77691058 -0.81982329 0.08043967 0.2632599711
[61,] -1.01843718 -2.42582042 -0.14615094 -0.2615107354
[62,] 0.06843254 -0.13153881 0.25038262 0.3944526477
[63,] 0.18919584 -1.96696410 0.13708732 -0.2615107354
[64,] 0.30995914 -0.36096697 0.53362088 0.2632599711
[65,] -0.29385737 -0.36096697 -0.08950329 0.1320672944
[66,] 1.03453895 0.09788935 0.36367793 0.2632599711
[67,] -0.29385737 -0.13153881 0.42032558 0.3944526477
[68,] -0.05233076 -0.81982329 0.19373497 -0.2615107354
[69,] 0.43072244 -1.96696410 0.42032558 0.3944526477
[70,] -0.29385737 -1.27867961 0.08043967 -0.1303180588
[71,] 0.06843254 0.32731751 0.59026853 0.7880306775
[72,] 0.30995914 -0.59039513 0.13708732 0.1320672944
[73,] 0.55148575 -1.27867961 0.64691619 0.3944526477
[74,] 0.30995914 -0.59039513 0.53362088 0.0008746178
[75,] 0.67224905 -0.36096697 0.30703027 0.1320672944
[76,] 0.91377565 -0.13153881 0.36367793 0.2632599711
[77,] 1.15530226 -0.59039513 0.59026853 0.2632599711
[78,] 1.03453895 -0.13153881 0.70356384 0.6568380009
[79,] 0.18919584 -0.36096697 0.42032558 0.3944526477
[80,] -0.17309407 -1.04925145 -0.14615094 -0.2615107354
[81,] -0.41462067 -1.50810778 0.02379201 -0.1303180588
[82,] -0.41462067 -1.50810778 -0.03285564 -0.2615107354
[83,] -0.05233076 -0.81982329 0.08043967 0.0008746178
[84,] 0.18919584 -0.81982329 0.76021149 0.5256453243
[85,] -0.53538397 -0.13153881 0.42032558 0.3944526477
[86,] 0.18919584 0.78617383 0.42032558 0.5256453243
[87,] 1.03453895 0.09788935 0.53362088 0.3944526477
[88,] 0.55148575 -1.73753594 0.36367793 0.1320672944
[89,] -0.29385737 -0.13153881 0.19373497 0.1320672944
[90,] -0.41462067 -1.27867961 0.13708732 0.1320672944
[91,] -0.41462067 -1.04925145 0.36367793 0.0008746178
[92,] 0.30995914 -0.13153881 0.47697323 0.2632599711
[93,] -0.05233076 -1.04925145 0.13708732 0.0008746178
[94,] -1.01843718 -1.73753594 -0.25944625 -0.2615107354
[95,] -0.29385737 -0.81982329 0.25038262 0.1320672944
[96,] -0.17309407 -0.13153881 0.25038262 0.0008746178
[97,] -0.17309407 -0.36096697 0.25038262 0.1320672944

[98,]	0.43072244	-0.36096697	0.30703027	0.1320672944	[99,]
-0.89767388	-1.27867961	-0.42938920	-0.1303180588	[100,]	-
0.17309407	-0.59039513	0.19373497	0.1320672944	[101,]	
0.55148575	0.55674567	1.27004036	1.7063794137		
[102,]	-0.05233076	-0.81982329	0.76021149	0.9192233541	
[103,]	1.51759216	-0.13153881	1.21339271	1.1816087073	
[104,]	0.55148575	-0.36096697	1.04344975	0.7880306775	
[105,]	0.79301235	-0.13153881	1.15674505	1.3128013839	
[106,]	2.12140867	-0.13153881	1.60992627	1.1816087073	
[107,]	-1.13920048	-1.27867961	0.42032558	0.6568380009	
[108,]	1.75911877	-0.36096697	1.43998331	0.7880306775	
[109,]	1.03453895	-1.27867961	1.15674505	0.7880306775	
[110,]	1.63835547	1.24503015	1.32668801	1.7063794137	
[111,]	0.79301235	0.32731751	0.76021149	1.0504160307	
[112,]	0.67224905	-0.81982329	0.87350679	0.9192233541	
[113,]	1.15530226	-0.13153881	0.98680210	1.1816087073	
[114,]	-0.17309407	-1.27867961	0.70356384	1.0504160307	
[115,]	-0.05233076	-0.59039513	0.76021149	1.5751867371	
[116,]	0.67224905	0.32731751	0.87350679	1.4439940605	
[117,]	0.79301235	-0.13153881	0.98680210	0.7880306775	
[118,]	2.24217198	1.70388647	1.66657392	1.3128013839	
[119,]	2.24217198	-1.04925145	1.77986923	1.4439940605	
[120,]	0.18919584	-1.96696410	0.70356384	0.3944526477	
[121,]	1.27606556	0.32731751	1.10009740	1.4439940605	
[122,]	-0.29385737	-0.59039513	0.64691619	1.0504160307	
[123,]	2.24217198	-0.59039513	1.66657392	1.0504160307	
[124,]	0.55148575	-0.81982329	0.64691619	0.7880306775	
[125,]	1.03453895	0.55674567	1.10009740	1.1816087073	
[126,]	1.63835547	0.32731751	1.27004036	0.7880306775	
[127,]	0.43072244	-0.59039513	0.59026853	0.7880306775	
[128,]	0.30995914	-0.13153881	0.64691619	0.7880306775	
[129,]	0.67224905	-0.59039513	1.04344975	1.1816087073	
[130,]	1.63835547	-0.13153881	1.15674505	0.5256453243	
[131,]	1.87988207	-0.59039513	1.32668801	0.9192233541	
[132,]	2.48369858	1.70388647	1.49663097	1.0504160307	
[133,]	0.67224905	-0.59039513	1.04344975	1.3128013839	
[134,]	0.55148575	-0.59039513	0.76021149	0.3944526477	
[135,]	0.30995914	-1.04925145	1.04344975	0.2632599711	
[136,]	2.24217198	-0.13153881	1.32668801	1.4439940605	
[137,]	0.55148575	0.78617383	1.04344975	1.5751867371	
[138,]	0.67224905	0.09788935	0.98680210	0.7880306775	
[139,]	0.18919584	-0.13153881	0.59026853	0.7880306775	
[140,]	1.27606556	0.09788935	0.93015445	1.1816087073	
[141,]	1.03453895	0.09788935	1.04344975	1.5751867371	
[142,]	1.27606556	0.09788935	0.76021149	1.4439940605	
[143,]	-0.05233076	-0.81982329	0.76021149	0.9192233541	
[144,]	1.15530226	0.32731751	1.21339271	1.4439940605	
[145,]	1.03453895	0.55674567	1.10009740	1.7063794137	
[146,]	1.03453895	-0.13153881	0.81685914	1.4439940605	
[147,]	0.55148575	-1.27867961	0.70356384	0.9192233541	

```

[148,]  0.79301235 -0.13153881  0.81685914  1.0504160307 [149,]
0.43072244  0.78617383   0.93015445  1.4439940605 [150,]
0.06843254 -0.13153881   0.76021149  0.7880306775
s<-sample(150,90)
iris_train<-iris[s,]
iris_test<-iris[-s,]
model<-NaiveBayes(Species~.,data=iris_train) pred<-
predict(model,iris_test)
conf<-confusionMatrix(pred$class ,iris_test$Species)$table
conf1<-as.matrix(conf)
accds<-((sum(diag(conf1)))/sum(conf1))*100
cat("Accuracy of standardized dataset is",accds,"%")

```

Accuracy of standardized dataset is 98.33333 %

K-NEAREST NEIGHBOUR

a) TRAINING SET IS 75%

```

iris_train<-iris_norm[1:113,]
iris_test<-iris_norm[114:150,]
iris_pred<-knn(iris_train,iris_test,iris[1:113,5],k=13)
con<-table(iris_pred,iris[114:150,5]) accuracy<-
((sum(diag(con)))/sum(con))*100
cat("Accuracy with 75% training set is ",accuracy,"%")

```

Accuracy with 75% training set is 51.35135 %

b) TRAINING SET IS 66.6%

```

iris_train<-iris_norm[1:100,] iris_test<-
iris_norm[101:150,]
iris_pred<-knn(iris_train,iris_test,iris[1:100,5],k=13)
con<-table(iris_pred,iris[101:150,5]) accu66<-
((sum(diag(con)))/sum(con))*100
cat("Accuracy with 66.6% training set is ",accu66,"%")

```

Accuracy with 66.6% training set is 0 %

HOLD OUT METHOD

```

iris_train<-iris_norm[1:75,] iris_test<-iris_norm[76:150,]
iris_pred<-knn(iris_train,iris_test,iris[1:75,5],k=13)
con<-table(iris_pred,iris[76:150,5]) acchld<-

```

```
((sum(diag(con)))/sum(con))*100 cat("Accuracy with hold  
out is ",acchld,"%")
```

Accuracy with hold out is 33.33333 %

RANDOM SUBSAMPLING

```
a) TRAINING SET IS 80%  
iris_train<-iris_norm[1:80,]  
iris_test<-iris_norm[81:150,]  
iris_pred<-knn(iris_train,iris_test,iris[1:80,5],k=13)  
con<-table(iris_pred,iris[81:150,5]) accu1<-  
((sum(diag(con)))/sum(con))*100
```

```
b) TRAINING SET IS 90%  
iris_train<-iris_norm[1:90,]  
iris_test<-iris_norm[91:150,]  
iris_pred<-knn(iris_train,iris_test,iris[1:90,5],k=13)  
con<-table(iris_pred,iris[91:150,5]) accu2<-  
((sum(diag(con)))/sum(con))*100 accrs<-max(accu1,accu2)  
cat("Accuracy with random subsampling is",accrs)
```

Accuracy with random subsampling is 28.57143

CROSS VALIDATION

```
x=iris[, -5]  
y=as.factor(iris$Species)  
res<-knn.cv(x,y,1:length(y))  
con<-table(res,y)  
accucs<-((sum(diag(con)))/sum(con))*100 cat("Accuracy  
with cross validation is",accucs)
```

Accuracy with cross validation is 96

COMPARISON

```
greatest<-max(acchld,accrs,accucs) if(greatest==acchld)  
{  
  cat("Holdout method does best classification")  
}  
if(greatest==accrs)  
{  
  cat("Random subsampling method does best classification")  
}  
if(greatest==accucs)  
{  
  cat("cross validation method does best classification")  
}
```

cross validation method does best classification

DATA SCALING

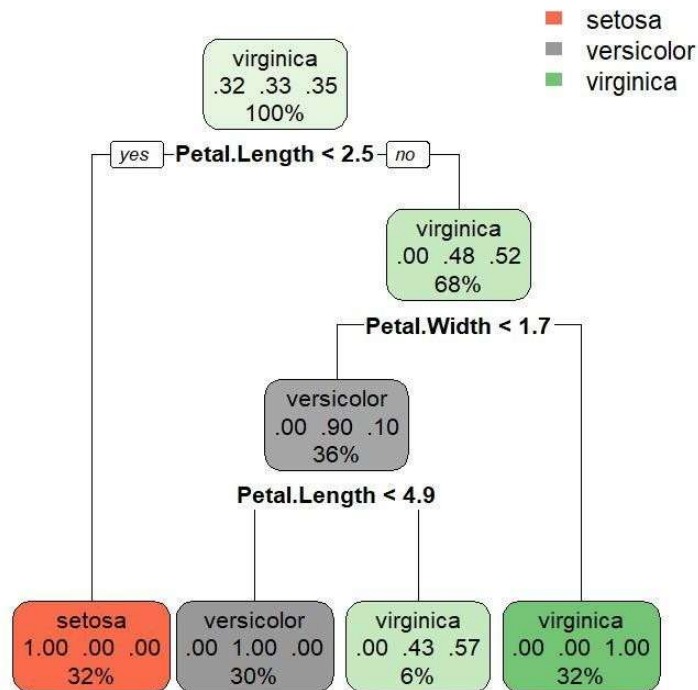
```
data_std <- function(x)
{
  (x-mean(x))/sd(x)
}
sapply(iris[,-5],data_std)
iris_train<-iris_norm[1:100,]
iris_test<-iris_norm[101:150,]
iris_pred<-knn(iris_train,iris_test,iris[1:100,5],k=13)
con<-table(iris_pred,iris[101:150,5]) accursd<-
((sum(diag(con)))/sum(con))*100 cat("Accuracy of sd
is",accurd)
```

Accuracy of sd is 0

DECISION TREE

a) TRAINING SET IS 75%

```
s<-sample(150,113) iris_train<-iris[s,]
iris_test<-iris[-s,]
dtm<-rpart(Species~.,iris_train,method="class")
rpart.plot(dtm)
```



```

p<-predict(dtm,iris_test,type="class")
cn<-confusionMatrix(iris_test[,5],p)$table
print(cn)

```

Reference

Prediction	setosa	versicolor	virginica
setosa	14	0	0
versicolor	0	11	2
virginica	0	0	10

```

cn1<-as.matrix(cn) accu<-
(sum(diag(cn1))/sum(cn1))*100
cat("The accuracy with 75% training data is ",accu,"%")

```

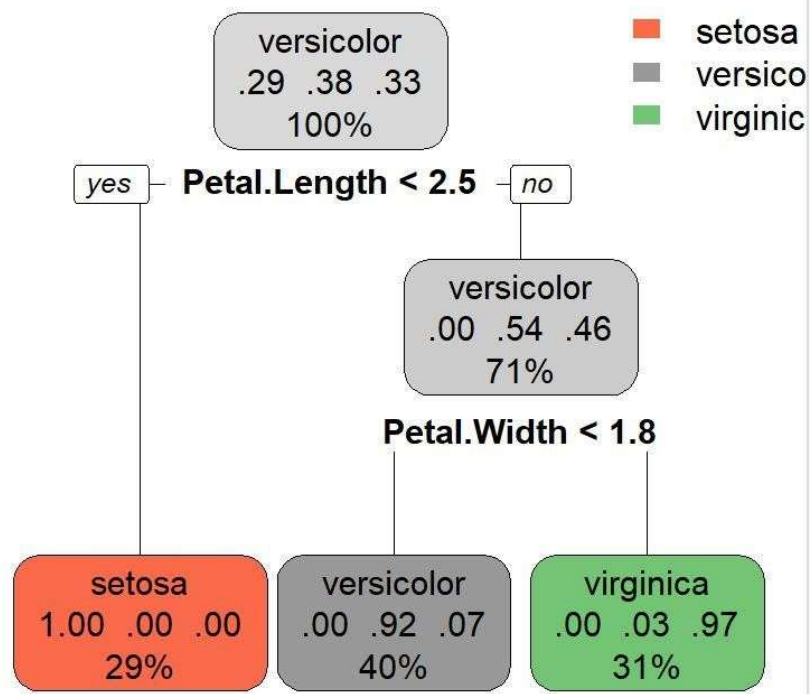
The accuracy with 75% training data is 94.59459 %

b) TRAINING SET IS 66.6%

```

s<-sample(150,100) iris_train<-iris[s,]
iris_test<-iris[-s,]
dtm<-rpart(Species~.,iris_train,method="class")
rpart.plot(dtm)

```



```

p<-predict(dtm,iris_test,type="class")
cn<-confusionMatrix(iris_test[,5],p)$table
print(cn)

```

```

Reference
Prediction  setosa versicolor virginica
setosa      21         0         0
versicolor  0         12        0
virginica   0         2        15

```

```

cn1<-as.matrix(cn)
accu6<-(sum(diag(cn1))/sum(cn1))*100
cat("The accuracy with 66.6% training data is ",accu6,"%") The
    accuracy with 66.6% training data is 96 %

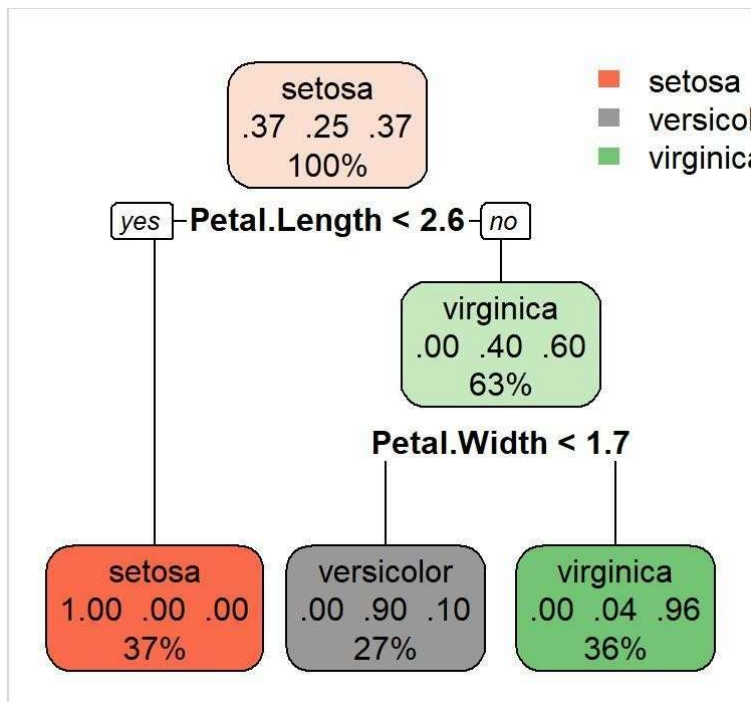
```

HOLD-OUT METHOD

```

s<-sample(150,75) iris_train<-iris[s,]
iris_test<-iris[-s,]
dtm<-rpart(Species~.,iris_train,method="class")
rpart.plot(dtm)

```

```

p<-predict(dtm,iris_test,type="class")
cn<-confusionMatrix(iris_test[,5],p)$table
cn1<-as.matrix(cn)
acchld<-(sum(diag(cn1))/sum(cn1))*100
cat("The accuracy with hold out method is ",acchld,"%")

```

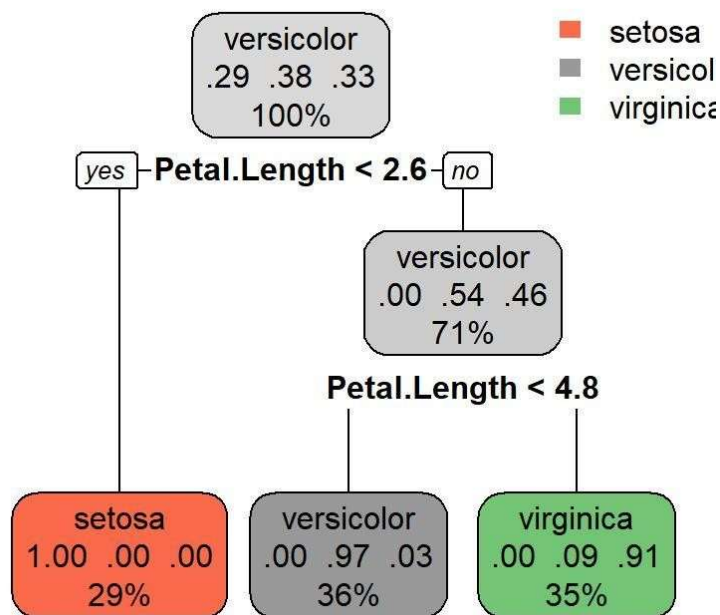
The accuracy with hold out method is 96 %

RANDOM SUBSAMPLING

```

i<-75 j<-1
acc<-c() for(i
in 75:100)
{ s<-sample(150,i)
iris_train<-iris[s,]
iris_test<-iris[-s,]
dtm<-rpart(Species~.,iris_train,method="class")
rpart.plot(dtm)
p<-predict(dtm,iris_test,type="class") cn<-
confusionMatrix(iris_test[,5],p)$table cn1<-as.matrix(cn)
acc[j]<-c((sum(diag(cn1))/sum(cn1))*100,acc)
j=j+1 }
acrs<-mean(acc)
cat("The accuracy with random subsampling method is ",acrs,"%")

```



The accuracy with random subsampling method is 94.11932 %

CROSS VALIDATION

```

set.seed(123)
form <- "Species ~ Sepal.Length + Sepal.Width + Petal.Length + Petal.Width"
folds <- split(iris, cut(sample(1:nrow(iris)),10))
errs <- rep(NA, length(folds))
for (i in 1:length(folds)) {
  test <- ldply(folds[i], data.frame)
  train <- ldply(folds[-i], data.frame)
  tmp.model <- rpart(form, train, method = "class")
  tmp.predict <- predict(tmp.model, newdata = test, type = "class")
  conf.mat <- table(test$Species, tmp.predict)
  errs[i] <- 1 - sum(diag(conf.mat))/sum(conf.mat)
}
accv <- (1 - mean(errs)) * 100
cat("The average accuracy using k-fold cross validation is", accv, "%")
  
```

The average accuracy using k-fold cross validation is 94 %

COMPARISON

```

greatest <- max(acchld, acrs, accv)
if (greatest == acchld) {
  cat("Holdout method does best classification")
}
if (greatest == acrs)
  
```

```

{
  cat("Random subsampling method does best classification")
}
if(greatest==accv)
{
  cat("Cross validation method does best classification")
}

```

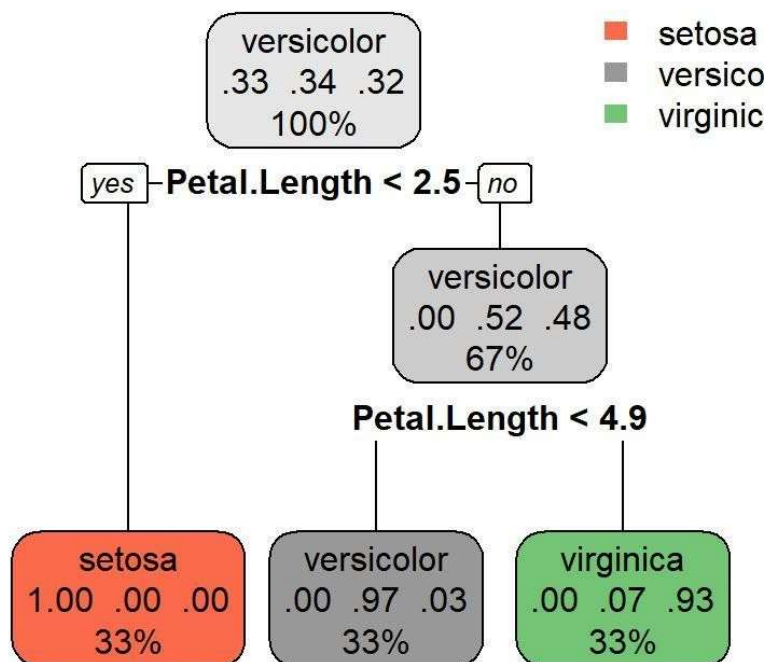
Holdout method does best classification

DATA SCALING

```

data_std <- function(x)
{
  (x-mean(x))/sd(x)
}
sapply(iris[, -5], data_std)
s<-sample(150,90)
iris_train<-iris[s,]
iris_test<-iris[-s,]
dtm<-rpart(Species~.,iris_train,method="class")
rpart.plot(dtm)

```



```

p<-predict(dtm,iris_test,type="class")
cn<-confusionMatrix(iris_test[,5],p)$table
cn1<- as.matrix(cn)
accsd<-(sum(diag(cn1))/sum(cn1))*100
cat("The accuracy in standardised iris data is ",accsd,"%")

```

The accuracy in standardised iris data is 95 %

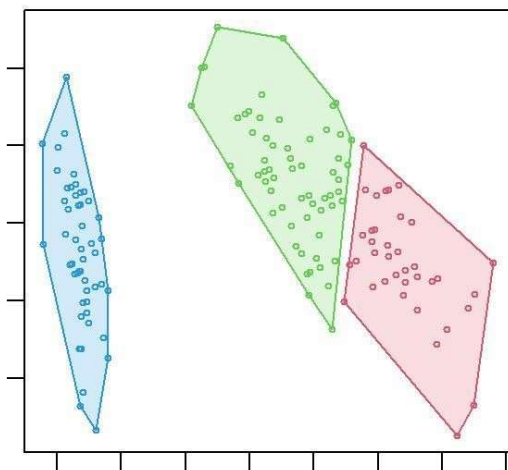
Question 6

Use Simple Kmeans, DBScan, Hierarchical clustering algorithms for clustering. Compare the performance of clusters by changing the parameters involved in the algorithm

K-means Clustering

```
library(cluster) library(dbSCAN)
ir <- iris k<-
kmeans(ir[1:4],3)
clusplot(ir[1:4], k$cluster, labels=2, lines=T)
hullplot(ir[1:4],k$cluster)
```

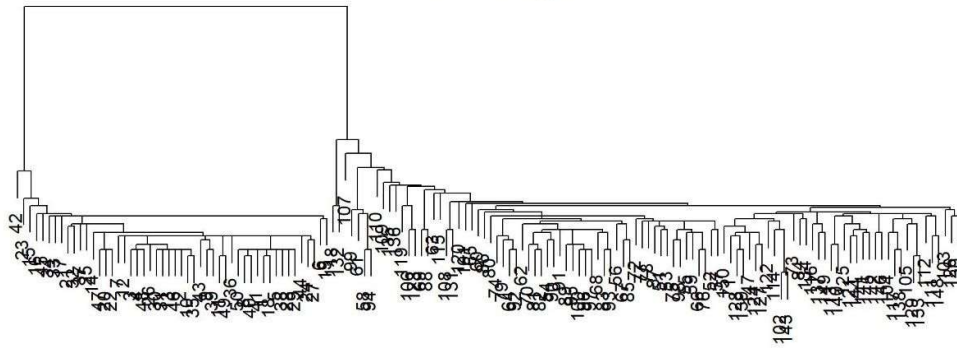
Convex Cluster Hulls



Hierarchical Clustering

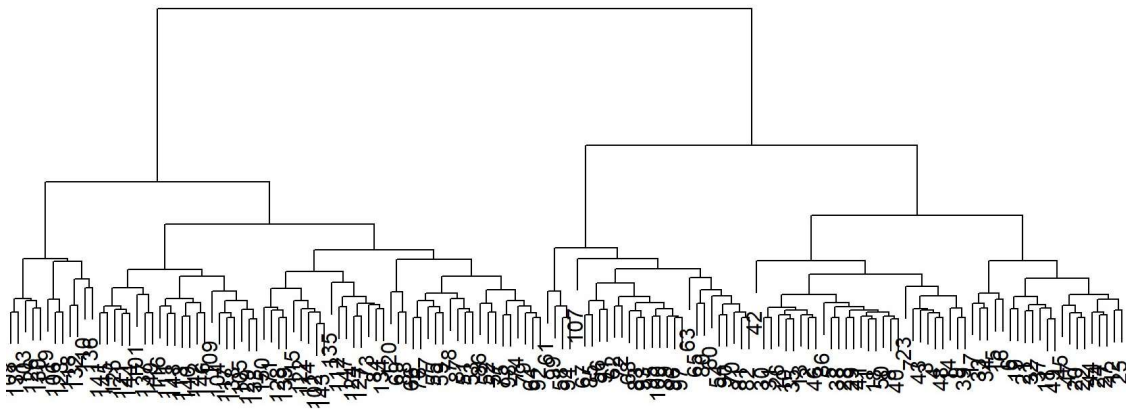
```
d<- dist(ir[1:4],method =
"euclidean") h <- hclust(d,method =
"single") plot(h)
```

Cluster Dendrogram



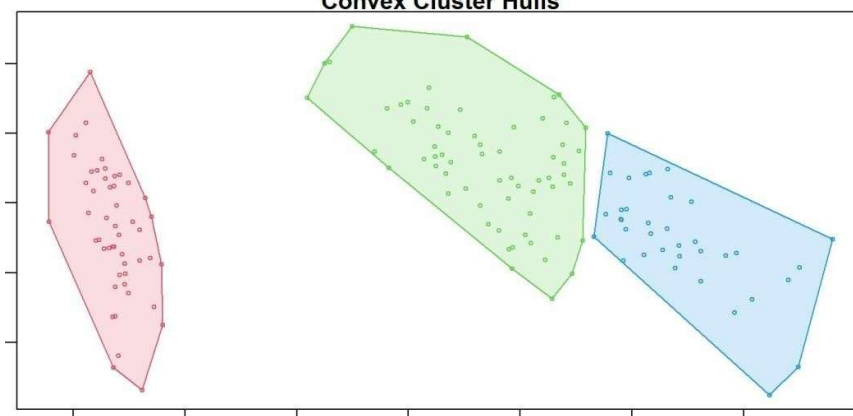
```
h1<-hclust(d,method="complete") plot(h1)
```

Cluster Dendrogram



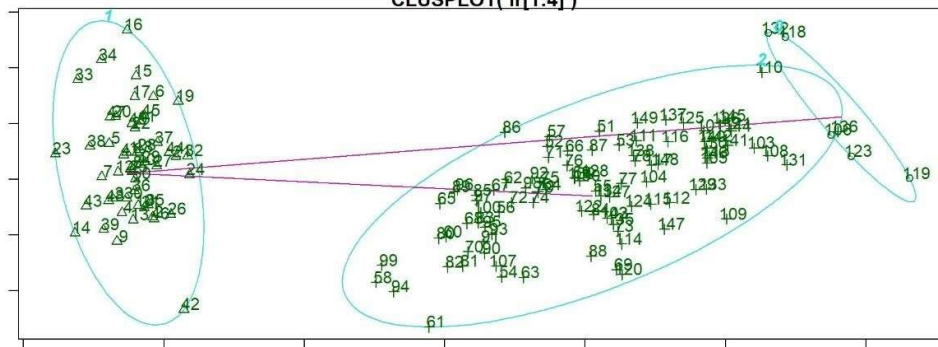
```
h2 <- hclust(d,method="average")
h2=cutree(h2,k=3) hullplot(ir[1:4],h2)
```

Convex Cluster Hulls



DB scan

```
d <- dist(ir[1:4],method ="euclidean") dbc<-
dbscan(ir[1:4],eps=0.8,minPts=10) dbc<-
dbscan(d,eps=0.8,minPts=10)
clusplot(ir[1:4],dbc$cluster, labels=2,lines=T)
```



```
hullplot(ir[1:4],dbc$cluster)
```

