

## Experiment-1

### Program

@relation exp1

@attribute customerid numeric

@attribute gender{male,female}

@attribute age numeric

@attribute annual\_income numeric

@attribute spending\_score numeric

@data

1,male,19,15,39

2,male,21,15,81

3,female,20,16,6

4,female,23,16,77

5,female,31,17,40

### Output

```
=== Run information ===

Scheme:      weka.clusterers.HierarchicalClusterer -N 2 -L SINGLE -P -A "weka.core.Filterer
Relation:    expl
Instances:    5
Attributes:   5
              customerid
              gender
              age
              annual_income
              spending_score
Test mode:    evaluate on training data

=== Clustering model (full training set) ===

Cluster 0
(((39.0:0.73431,81.0:0.73431):0.90381,77.0:1.63811):0.28319,6.0:1.9213)

Time taken to build model (full training data) : 0 seconds

=== Model and evaluation on training set ===

Clustered Instances

0      4 ( 80%)
1      1 ( 20%)
```

## Experiment-2

### Program

@relation exp1

@attribute emp\_id numeric

@attribute gender{male,female}

@attribute age numeric

@attribute salary numeric

@attribute credit numeric

@data

111,male,28,1500000,39

222,male,25,1500000,27

333,female,26,1600000,42

444,female,25,1600000,40

555,female,30,1700000,64

666,male,29,2000000,72

### Output

```
Initial starting points (random):

Cluster 0: 444,female,25,1600000,40
Cluster 1: 111,male,28,1500000,39
Cluster 2: 222,male,25,1500000,27

Missing values globally replaced with mean/mode

Final cluster centroids:

Attribute      Full Data      Cluster#
              (6.0)      0          1          2
              (3.0)      (1.0)      (2.0)
=====
emp_id         388.5         444         666         166.5
gender         male         female      male         male
age            27.1667        27          29          26.5
salary         1650000 1633333.3333 2000000     1500000
credit         47.3333        48.6667     72          33

Time taken to build model (full training data) : 0 seconds

=== Model and evaluation on training set ===

Clustered Instances

0      3 ( 50%)
1      1 ( 17%)
2      2 ( 33%)
```



Experiment-3 naivebayes in breast\_cancer

Output

Time taken to build model: 0 seconds

=== Evaluation on test split ===

Time taken to test model on test split: 0 seconds

=== Summary ===

Correctly Classified Instances	69	71.134 %
Incorrectly Classified Instances	28	28.866 %
Kappa statistic	0.3274	
Mean absolute error	0.3431	
Root mean squared error	0.4825	
Relative absolute error	79.9872 %	
Root relative squared error	100.9522 %	
Total Number of Instances	97	

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	MCC	ROC Area	PRC Area	Class
	0.828	0.515	0.757	0.828	0.791	0.331	0.676	0.789	no-recurrence-events
	0.485	0.172	0.593	0.485	0.533	0.331	0.676	0.560	recurrence-events
Weighted Avg.	0.711	0.398	0.701	0.711	0.703	0.331	0.676	0.711	

=== Confusion Matrix ===

```

a  b  <-- classified as
53 11 | a = no-recurrence-events
17 16 | b = recurrence-events

```

## Experiment-4

### Program

@relation exp1

@attribute person{gopu,babu,baby,gopal,krishna,jai,dev,malini,hema,anu}

@attribute gender{yes,no}

@data

gopu,yes

babu,yes

baby,yes

gopal,no

krishna,yes

jai,no

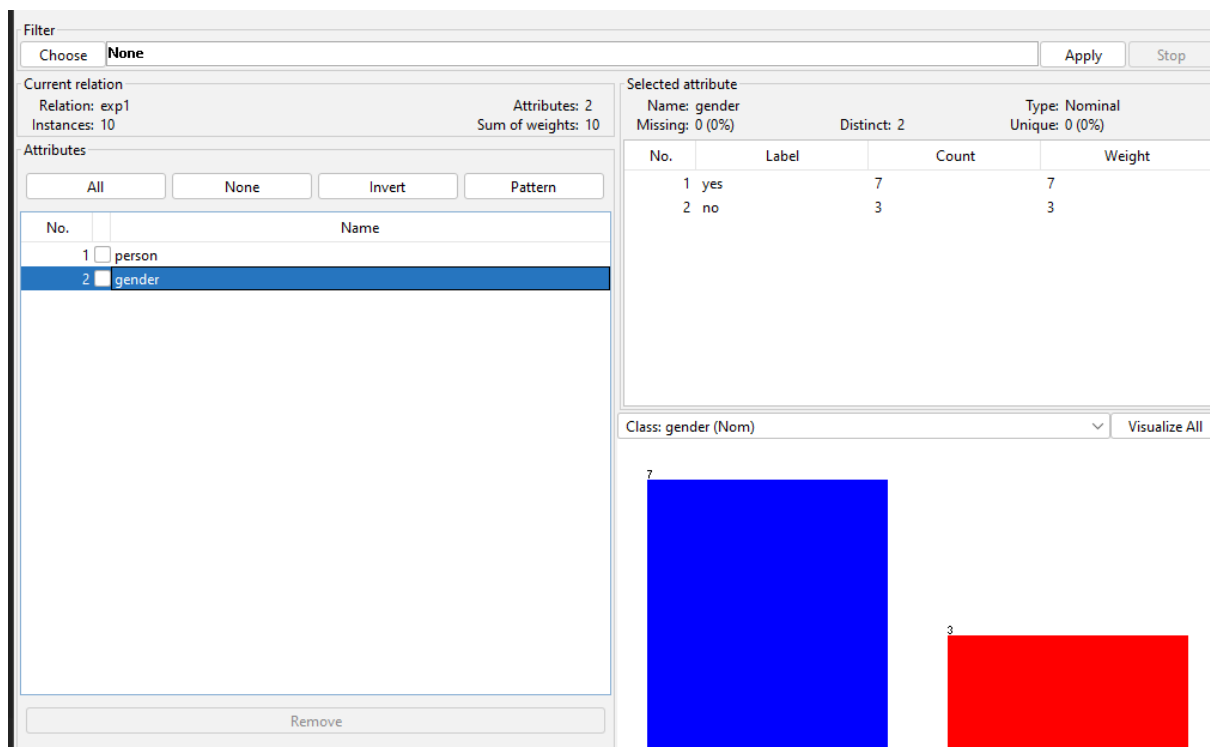
dev,no

malini,yes

hema,yes

anu,yes

output;



## Experiment-6

### Program

@relation exp1

@attribute bread{true,false}

@attribute cheese{true,false}

@attribute egg{true,false}

@attribute juice{true,false}

@attribute milk{true,false}

@attribute yogurt{true,false}

@data

true,true,true,true,false,false

true,true,false,true,false,false

true,false,false,false,true,true

true,false,false,true,true,false

false,true,false,true,true,false

output;

```
=== Run information ===

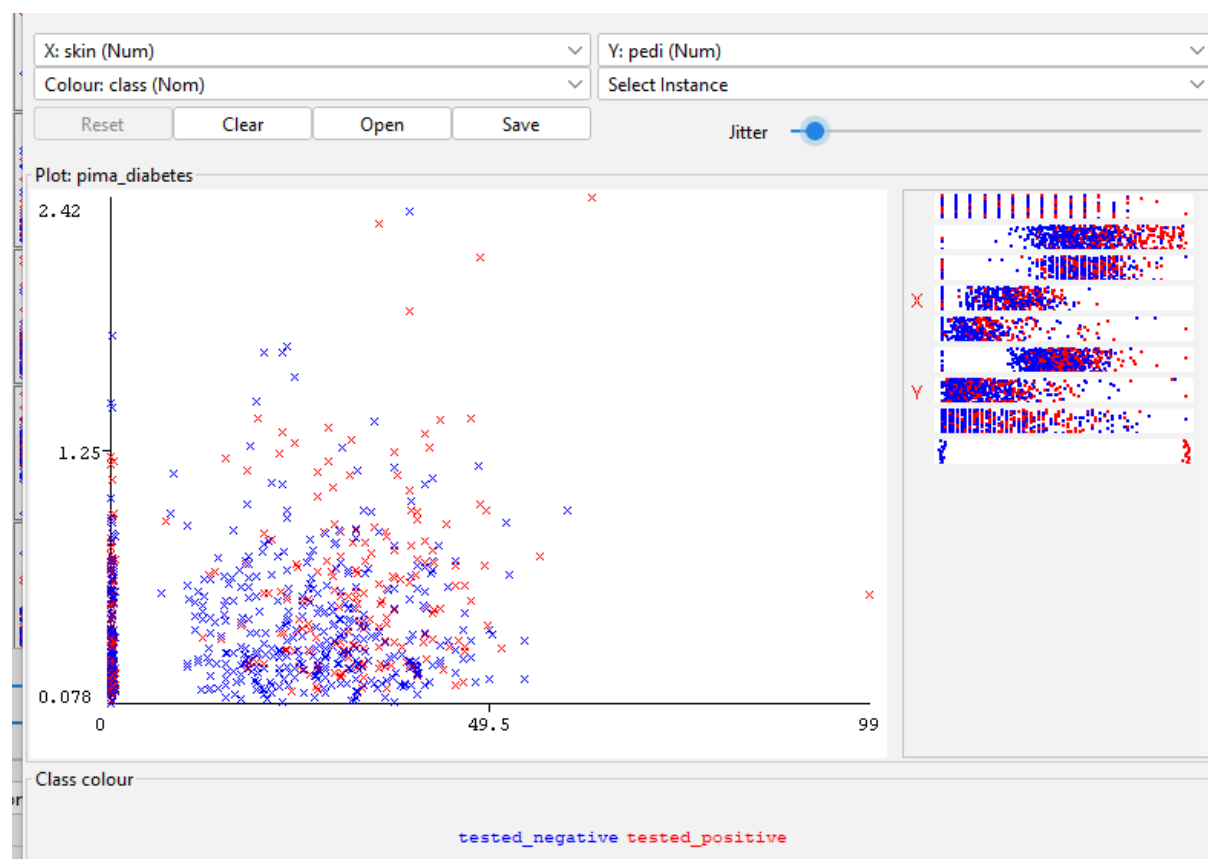
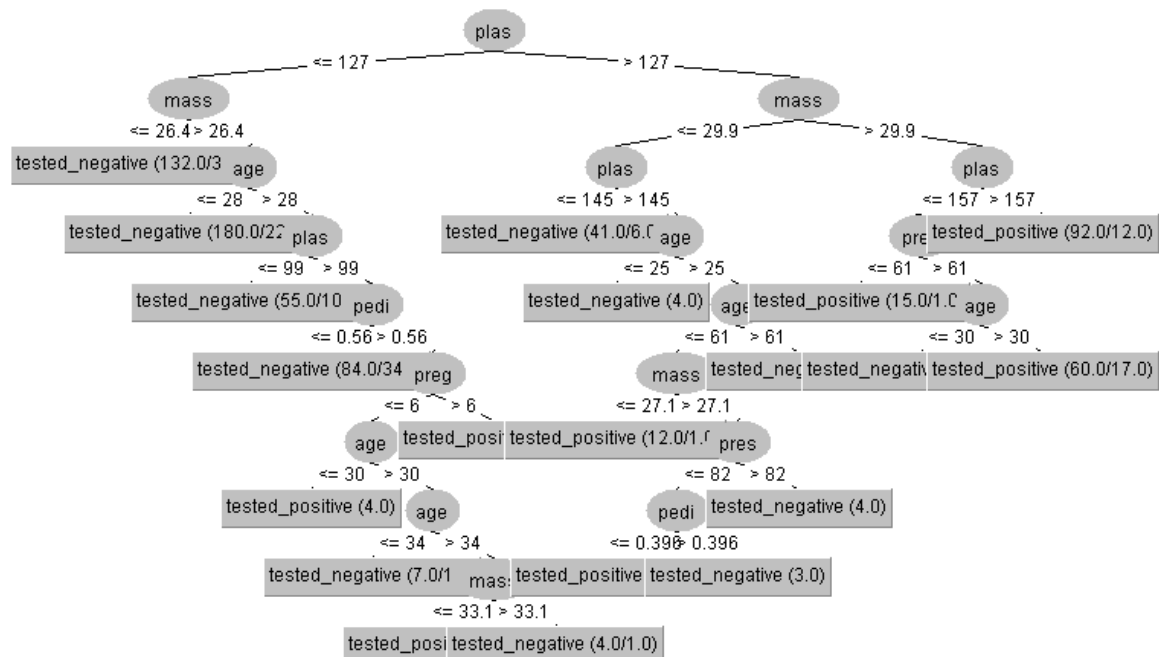
Scheme:      weka.associations.FPGrowth -P 2 -I -1 -N 10 -T 0 -C 0.75 -D 0.05 -U 2.0 -M 0.1
Relation:    exp1
Instances:   5
Attributes:  6
             bread
             cheese
             egg
             juice
             milk
             yogurt

=== Associator model (full training set) ===

FPGrowth found 14 rules (displaying top 10)

1. [milk=false]: 2 ==> [yogurt=false]: 2   <conf:(1)> lift:(1.25) lev:(0.08) conv:(0.4)
2. [bread=false]: 1 ==> [yogurt=false]: 1   <conf:(1)> lift:(1.25) lev:(0.04) conv:(0.2)
3. [cheese=false]: 2 ==> [egg=false]: 2     <conf:(1)> lift:(1.25) lev:(0.08) conv:(0.4)
4. [juice=false]: 1 ==> [egg=false]: 1     <conf:(1)> lift:(1.25) lev:(0.04) conv:(0.2)
5. [bread=false]: 1 ==> [egg=false]: 1     <conf:(1)> lift:(1.25) lev:(0.04) conv:(0.2)
6. [juice=false]: 1 ==> [cheese=false]: 1   <conf:(1)> lift:(2.5) lev:(0.12) conv:(0.6)
7. [egg=false, milk=false]: 1 ==> [yogurt=false]: 1   <conf:(1)> lift:(1.25) lev:(0.04) conv:(0.2)
8. [yogurt=false, cheese=false]: 1 ==> [egg=false]: 1   <conf:(1)> lift:(1.25) lev:(0.04) conv:(0.2)
9. [bread=false]: 1 ==> [yogurt=false, egg=false]: 1   <conf:(1)> lift:(1.67) lev:(0.08) conv:(0.4)
10. [yogurt=false, bread=false]: 1 ==> [egg=false]: 1   <conf:(1)> lift:(1.25) lev:(0.04) conv:(0.2)
```

## Experiment-7



## Experiment-8

```
library(dplyr)

library(tibble)

library(ggplot2)

data <- c(55, 60, 71, 63, 55, 65, 50, 55, 58, 59, 61, 63, 65, 67, 71, 72, 75)

equi_depth <- quantile(data, probs = c(0, 1/3, 2/3, 1))

equi_depth_partitioned <- cut(data, breaks = equi_depth, labels = c("Low", "Medium", "High"),
include.lowest = TRUE)

min_value <- min(data)

max_value <- max(data)

width <- (max_value - min_value)/3

equal_width <- seq(min_value, max_value, by = width)

equal_width_partitioned <- cut(data, breaks = equal_width, labels = c("Low", "Medium", "High"),
include.lowest = TRUE)

kmeans_model <- kmeans(data, centers = 3)

cluster_assignments <- as.factor(kmeans_model$cluster)

levels(cluster_assignments) <- c("Low", "Medium", "High")


data_tibble <- tibble(data = data, equi_depth_partitioned = equi_depth_partitioned,
equal_width_partitioned = equal_width_partitioned, cluster_assignments = cluster_assignments)


ggplot(data_tibble, aes(x = data)) +
  geom_histogram(binwidth = 5) +
  facet_wrap(~equi_depth_partitioned, ncol = 1, scales = "free_x") +
  ggtitle("Histogram using Equi-Depth Partitioning")

ggplot(data_tibble, aes(x = data)) +
  geom_histogram(binwidth = 5) +
  facet_wrap(~equal_width_partitioned, ncol = 1, scales = "free_x") +
  ggtitle("Histogram using Equal-Width Partitioning")

ggplot(data_tibble, aes(x = data)) +
  geom_histogram(binwidth = 5) +
  facet_wrap(~cluster_assignments, ncol = 1, scales = "free_x") +
```

experiment-9

program

@relation exp1

@attribute SONY{true,false}

@attribute BPL{true,false}

@attribute LG{true,false}

@attribute SAMSUNG{true,false}

@attribute ONIDA{true,false}

@data

true,true,true,false,false

false,true,false,true,false

false,true,false,false,true

true,true,false,true,false

true,false,false,false,true

false,true,false,false,true

true,false,false,false,true

true,true,true,false,true

true,true,false,false,true

output;

```
=== Run information ===

Scheme:      weka.associations.FPGrowth -P 2 -I -1 -N 10 -T 0 -C 0.5 -D 0.05 -U 2.0 -M 0.1
Relation:    expl
Instances:    9
Attributes:   5
              SONY
              BPL
              LG
              SAMSUNG
              ONIDA
=== Associator model (full training set) ===

FPGrowth found 10 rules (displaying top 10)

1. [BPL=false]: 2 ==> [SAMSUNG=false]: 2 <conf:(1)> lift:(1.29) lev:(0.05) conv:(0.44)
2. [BPL=false]: 2 ==> [LG=false]: 2 <conf:(1)> lift:(1.29) lev:(0.05) conv:(0.44)
3. [SAMSUNG=false, SONY=false]: 2 --> [LG=false]: 2 <conf:(1)> lift:(1.29) lev:(0.05) conv:(0.44)
4. [BPL=false]: 2 ==> [SAMSUNG=false, LG=false]: 2 <conf:(1)> lift:(1.8) lev:(0.1) conv:(0.89)
5. [SAMSUNG=false, BPL=false]: 2 ==> [LG=false]: 2 <conf:(1)> lift:(1.29) lev:(0.05) conv:(0.44)
6. [LG=false, BPL=false]: 2 ==> [SAMSUNG=false]: 2 <conf:(1)> lift:(1.29) lev:(0.05) conv:(0.44)
7. [SONY=false]: 3 ==> [SAMSUNG=false]: 2 <conf:(0.67)> lift:(0.86) lev:(-0.04) conv:(0.33)
8. [ONIDA=false]: 3 ==> [LG=false]: 2 <conf:(0.67)> lift:(0.86) lev:(-0.04) conv:(0.33)
9. [SONY=false]: 3 ==> [SAMSUNG=false, LG=false]: 2 <conf:(0.67)> lift:(1.2) lev:(0.04) conv:(0.67)
10. [LG=false, SONY=false]: 3 ==> [SAMSUNG=false]: 2 <conf:(0.67)> lift:(0.86) lev:(-0.04) conv:(0.33)
```



```

Apriori
=====

Minimum support: 0.6 (5 instances)
Minimum metric <confidence>: 0.5
Number of cycles performed: 8

Generated sets of large itemsets:

Size of set of large itemsets L(1): 5

Size of set of large itemsets L(2): 6

Size of set of large itemsets L(3): 1

Best rules found:

1. ONIDA=true 6 ==> SAMSUNG=false 6    <conf:(1)> lift:(1.29) lev:(0.15) [1] conv:(1.33)
2. LG=false ONIDA=true 5 ==> SAMSUNG=false 5    <conf:(1)> lift:(1.29) lev:(0.12) [1] conv:(1.11)
3. LG=false SAMSUNG=false 5 ==> ONIDA=true 5    <conf:(1)> lift:(1.5) lev:(0.19) [1] conv:(1.67)
4. SAMSUNG=false 7 ==> ONIDA=true 6    <conf:(0.86)> lift:(1.29) lev:(0.15) [1] conv:(1.17)
5. SONY=true 6 ==> SAMSUNG=false 5    <conf:(0.83)> lift:(1.07) lev:(0.04) [0] conv:(0.67)
6. ONIDA=true 6 ==> LG=false 5    <conf:(0.83)> lift:(1.07) lev:(0.04) [0] conv:(0.67)
7. SAMSUNG=false ONIDA=true 6 ==> LG=false 5    <conf:(0.83)> lift:(1.07) lev:(0.04) [0] conv:(0.67)
8. ONIDA=true 6 ==> LG=false SAMSUNG=false 5    <conf:(0.83)> lift:(1.5) lev:(0.19) [1] conv:(1.33)
9. SAMSUNG=false 7 ==> SONY=true 5    <conf:(0.71)> lift:(1.07) lev:(0.04) [0] conv:(0.78)
10. LG=false 7 ==> BPL=true 5    <conf:(0.71)> lift:(0.92) lev:(-0.05) [0] conv:(0.52)

```

#### Experiment-10 Program

```
speed<-c(78,81,82,74,83,82,77,80,70)
```

```
time<-c(39,37,36,42,35,36,40,38,46)
```

```
a<-var(speed)
```

```
b<-var(time)
```

```
ss<-sqrt(var(speed))
```

```
st<-sqrt(var(time))
```

```
a
```

```
b
```

```
ss
```

```
st
```

output

```

> speed<-c(78,81,82,74,83,82,77,80,70)
> time<-c(39,37,36,42,35,36,40,38,46)
> a<-var(speed)
> b<-var(time)
> ss<-sqrt(var(speed))
> st<-sqrt(var(time))
> a
[1] 18.52778
> b
[1] 12.19444
> ss
[1] 4.304391
> st
[1] 3.492054
~ |

```

## Experiment-11

```
a<-c(100,70,60,90,90)
```

```
min=0
```

```
max=1
```

```
v<-((a-min(a)/max(a)-min(a))*max+min)/100
```

```
summary(v,method=c("range"))
```

```
v
```

```
#zscore
```

```
w=mean(a)
```

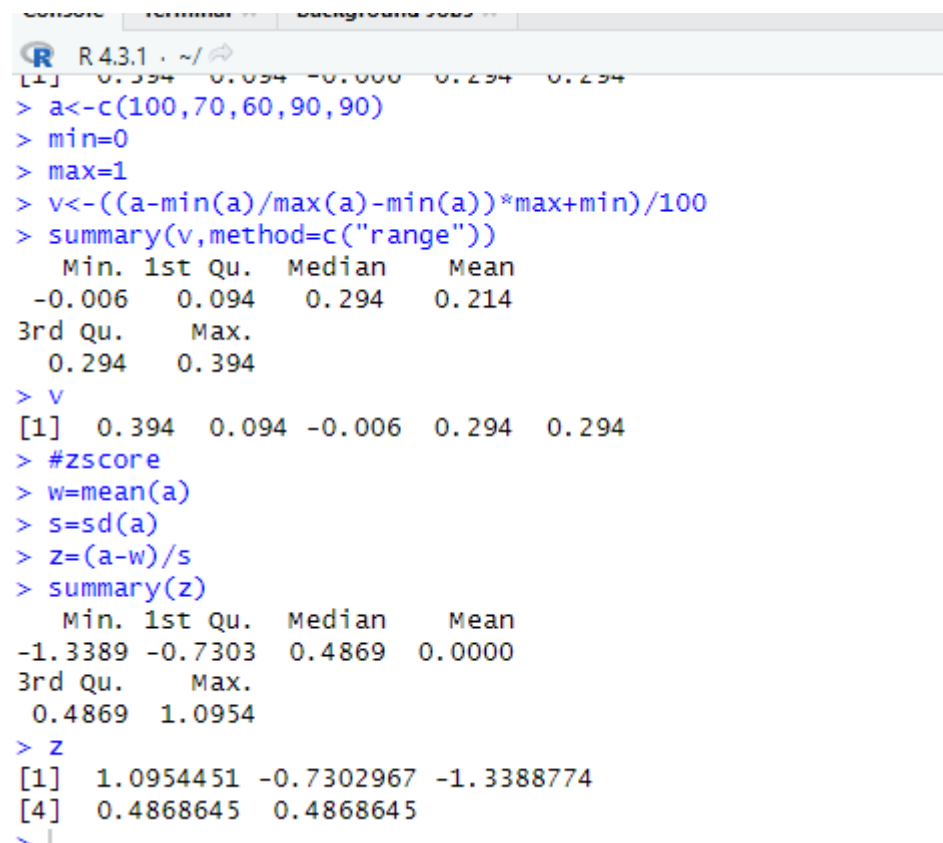
```
s=sd(a)
```

```
z=(a-w)/s
```

```
summary(z)
```

```
z
```

output



```
R 4.3.1 ~/  
[1] 0.394 0.094 -0.006 0.294 0.294  
> a<-c(100,70,60,90,90)  
> min=0  
> max=1  
> v<-((a-min(a)/max(a)-min(a))*max+min)/100  
> summary(v,method=c("range"))  
      Min. 1st Qu.  Median    Mean  
-0.006   0.094   0.294   0.214  
3rd Qu.    Max.  
 0.294   0.394  
> v  
[1] 0.394 0.094 -0.006 0.294 0.294  
> #zscore  
> w=mean(a)  
> s=sd(a)  
> z=(a-w)/s  
> summary(z)  
      Min. 1st Qu.  Median    Mean  
-1.3389 -0.7303  0.4869  0.0000  
3rd Qu.    Max.  
 0.4869  1.0954  
> z  
[1] 1.0954451 -0.7302967 -1.3388774  
[4] 0.4868645  0.4868645  
~ |
```