

SHETH L.U.J.AND SIR M.V. COLLEGE
PRACTICAL(1,2,3,4,5,6) MODULE2
Data Analysis with SAS / SPSS /R

1) AIM:- Generating descriptive statistics using summary() or describe() (R)

INPUT:-

```
library(dplyr)

df <- read.csv("shades.csv")

print("---- 2. Frequency Tables for Shades Dataset ----")

# Automatically select the first categorical column

cat_var <- names(df)[!sapply(df, is.numeric)][1]

print(paste("Using categorical column:", cat_var))

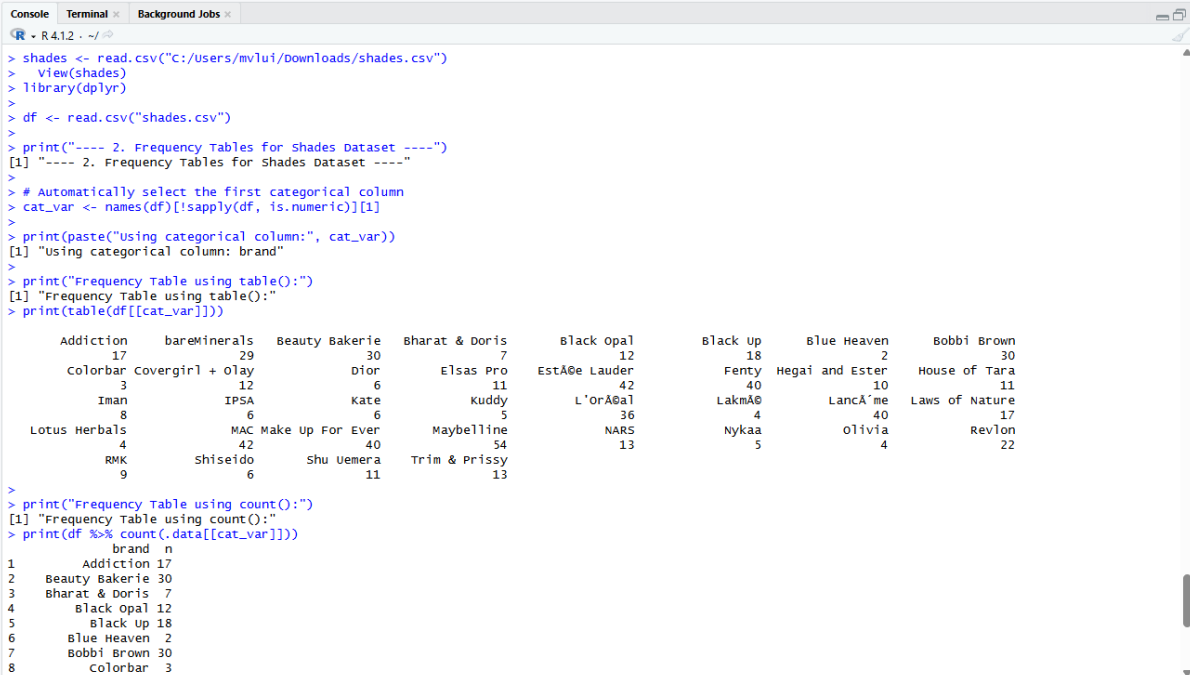
print("Frequency Table using table():")

print(table(df[[cat_var]]))

print("Frequency Table using count():")

print(df %>% count(.data[[cat_var]]))
```

OUTPUT:-



```
Console Terminal Background Jobs
R - R 4.1.2 - ~/...

> shades <- read.csv("C:/Users/mvlu/Downloads/shades.csv")
> View(shades)
> library(dplyr)
> 
> df <- read.csv("shades.csv")
> 
> print("---- 2. Frequency Tables for Shades Dataset ----")
[1] "---- 2. Frequency Tables for Shades Dataset ----"
> 
> # Automatically select the first categorical column
> cat_var <- names(df)[!sapply(df, is.numeric)][1]
> 
> print(paste("Using categorical column:", cat_var))
[1] "Using categorical column: brand"
> 
> print("Frequency Table using table():")
[1] "Frequency Table using table():"
> print(table(df[[cat_var]]))

```

brand	n
Addiction	17
Beauty Bakerie	30
Bharat & Doris	7
Black Opal	12
Black Up	18
Blue Heaven	2
Bobbi Brown	30
Colorbar	3
Covergirl + Olay	12
Dior	6
Elsas Pro	11
Estée Lauder	42
Fenty	40
Hegai and Ester	10
House of Tara	11
Iman	8
IPSA	6
Kate	6
Kuddy	5
L'Oréal	36
Lakmé	4
Lancôme	40
Laws of Nature	17
Lotus Herbals	4
MAC Make Up For Ever	40
Maybelline	54
NARS	13
Nykaa	5
Olivia	4
Revlon	22
RMK	9
Shiseido	6
Shu Uemura	11
Trim & Prissy	13

```
> 
> print("Frequency Table using count():")
[1] "Frequency Table using count():"
> print(df %>% count(.data[[cat_var]]))

```

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```
Console Terminal Background Jobs
R - R 4.1.2 - ~/
> print("Frequency Table using count():")
[1] "Frequency Table using count():"
> print(df %>% count(.data[[cat_var]]))
  brand      n
1  Addition 17
2  Beauty Bakerie 30
3  Bharat & Doris 7
4  Black Opal 12
5  Black Up 18
6  Blue Heaven 2
7  Bobbi Brown 30
8  Colorbar 3
9  Covergirl + Olay 12
10 Dior 6
11 Elsas Pro 11
12 Est  e Lauder 42
13 Fenty 40
14 Hegai and Ester 10
15 House of Tara 11
16 IPSA 6
17 Iman 8
18 Kate 6
19 Kuddy 5
20 L'Or  al 36
21 Lakm   4
22 Lanc  me 40
23 Laws of Nature 17
24 Lotus Herbals 4
25 MAC 42
26 Make Up For Ever 40
27 Maybelline 54
28 NARS 13
29 Nykaa 5
30 Olivia 4
31 RMK 9
32 Revlon 22
33 Shiseido 6
34 Shu Uemura 11
35 Trim & Prissy 13
36 bareMinerals 29
> |
```

2) Generating frequency tables using table() or count() (R).

INPUT:-

```
library(psych)
```

```
df <- read.csv("shades.csv")
```

```
print("---- 1. Descriptive Statistics for Shades Dataset ----")
```

```
print("Summary of a numeric variable:")
```

```
summary(df[sapply(df, is.numeric)][,1])
```

```
print("Detailed Description of numeric variables:")
```

```
describe(df[sapply(df, is.numeric)])
```

OUTPUT:-

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```
> library(psych)
>
> df <- read.csv("shades.csv")
>
> print("---- 1. Descriptive Statistics for Shades Dataset ----")
[1] "---- 1. Descriptive Statistics for Shades Dataset ----"
>
> print("Summary of a numeric variable:")
[1] "Summary of a numeric variable:"
> summary(df[sapply(df, is.numeric)][,1])
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.   NA's
 4.00   23.00   26.00   25.31   29.00   45.00    12
>
> print("Detailed Description of numeric variables:")
[1] "Detailed Description of numeric variables:"
> describe(df[sapply(df, is.numeric)])
      vars  n mean  sd median trimmed  mad min max range skew kurtosis  se
H         1 613 25.31 5.33  26.00   25.58  4.45  4.0 45  41.0 -0.53    1.30 0.22
S         2 613  0.46 0.15  0.44    0.45  0.16  0.1  1  0.9  0.38   -0.17 0.01
V         3 613  0.78 0.17  0.84    0.80  0.13  0.2  1  0.8 -1.06    0.42 0.01
L         4 625 65.92 17.51  71.00   67.56 16.31 11.0 95 84.0 -0.80   -0.11 0.70
group     5 625  3.47  1.98  3.00    3.43  1.48  0.0  7  7.0  0.22   -0.88 0.08
> |
```

3) Creating cross-tabulations and two-way tables using table() (R)

INPUT:-

```
df <- read.csv("shades.csv")
```

```
print("---- 3. Cross-Tabulation for Shades Dataset ----")
```

```
cat_vars <- names(df)[!sapply(df, is.numeric)]
```

```
cross_tab <- table(df[[cat_vars[1]]], df[[cat_vars[2]]])
```

```
print(cross_tab)
```

OUTPUT:-

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```
> df <- read.csv("shades.csv")
>
> print("---- 3. Cross-Tabulation for Shades Dataset ----")
[1] "---- 3. Cross-Tabulation for Shades Dataset ----"
>
> cat_vars <- names(df)[!sapply(df, is.numeric)]
>
> cross_tab <- table(df[[cat_vars[1]]], df[[cat_vars[2]]])
>
> print(cross_tab)
```

	ad	bb	bd	bh	bm	bo	br	bu	cb	di	el	ep	fe	he	ht	im	ip	ka	ku	la	lc	lh	ln	lo	ma	mb	mu	na	ny	oc	ol	rmk	rv	sh	su	tp
Addiction	17	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
BareMinerals	0	0	0	0	29	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Beauty Bakerie	0	30	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Bharat & Doris	0	0	7	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Black Opal	0	0	0	0	0	12	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Black Up	0	0	0	0	0	0	18	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Blue Heaven	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Bobbi Brown	0	0	0	0	0	30	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Colorbar	0	0	0	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Covergirl + olay	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	12	0	0	0	0	0		
Dior	0	0	0	0	0	0	0	0	6	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Elsas Pro	0	0	0	0	0	0	0	0	0	0	11	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Estée Lauder	0	0	0	0	0	0	0	0	0	42	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Fenty	0	0	0	0	0	0	0	0	0	0	0	40	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Hegal and Ester	0	0	0	0	0	0	0	0	0	0	0	0	10	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
House of Tara	0	0	0	0	0	0	0	0	0	0	0	0	11	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Iman	0	0	0	0	0	0	0	0	0	0	0	0	0	8	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
IPSA	0	0	0	0	0	0	0	0	0	0	0	0	0	6	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Kate	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	6	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Kuddy	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
L'Oréal	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	36	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Lakmé	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Lancôme	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	40	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Laws of Nature	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	17	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
Lotus Herbals	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
MAC	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	42	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Make Up For Ever	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	40	0	0	0	0	0	0	0	0	0	0	0	0

```
[ reached getOption("max.print") -- omitted 9 rows ]
> |
```

4) Performing one-sample t-tests using t.test() (R).

INPUT:-

```
df <- read.csv("shades.csv")
```

```
print("---- 4. One-Sample t-test for Shades Dataset ----")
```

```
num_var <- names(df)[sapply(df, is.numeric)][1]
```

```
test_value <- 100
```

```
t_test_one <- t.test(df[[num_var]], mu = test_value)
```

```
print(t_test_one)
```

OUTPUT:-

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```
> df <- read.csv("shades.csv")
>
> print("---- 4. One-Sample t-test for Shades Dataset ----")
[1] "---- 4. One-Sample t-test for Shades Dataset ----"
>
> num_var <- names(df)[sapply(df, is.numeric)][1]
>
> test_value <- 100
>
> t_test_one <- t.test(df[[num_var]], mu = test_value)
>
> print(t_test_one)

      One Sample t-test

data:  df[[num_var]]
t = -347.07, df = 612, p-value < 2.2e-16
alternative hypothesis: true mean is not equal to 100
95 percent confidence interval:
 24.89224 25.73745
sample estimates:
mean of x
 25.31485

> |
```

5) Performing independent two-sample t-tests using t.test() with grouping (R).

INPUT:-

```
library(dplyr)
```

```
shdes <- read.csv("shades.csv")
```

```
head(shdes)
```

```
print("---- 5. Independent Two-Sample t-test ----")
```

```
num_var <- "H"
```

```
binary_var <- "group"
```

```
if(length(unique(shdes[[binary_var]])) > 2){
```

```
  shdes[[binary_var]] <- ifelse(shdes[[binary_var]] > median(shdes[[binary_var]]), "High",
    "Low")
```

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```
}
```

```
shdes[[binary_var]] <- as.factor(shdes[[binary_var]])
```

```
t_test_two <- t.test(shdes[[num_var]] ~ shdes[[binary_var]])
```

```
print(t_test_two)
```

OUTPUT:-

```
> library(dplyr)
>
> shdes <- read.csv("shades.csv")
>
> head(shdes)
  brand brand_short product product_short hex H S V L group
1 Maybelline      mb Fit Me      fmf f3c3b3 26 0.26 0.95 86 2
2 Maybelline      mb Fit Me      fmf ffe3c2 32 0.24 1.00 92 2
3 Maybelline      mb Fit Me      fmf ffe0cd 23 0.20 1.00 91 2
4 Maybelline      mb Fit Me      fmf ffd3be 19 0.25 1.00 88 2
5 Maybelline      mb Fit Me      fmf bd9584 18 0.30 0.74 65 2
6 Maybelline      mb Fit Me      fmf eabda6 20 0.29 0.92 80 2
>
> print("---- 5. Independent Two-Sample t-test ----")
[1] "---- 5. Independent Two-Sample t-test ----"
>
> num_var <- "H"
> binary_var <- "group"
>
>
> if(length(unique(shdes[[binary_var]])) > 2){
+   shdes[[binary_var]] <- ifelse(shdes[[binary_var]] > median(shdes[[binary_var]]), "High", "Low")
+ }
> shdes[[binary_var]] <- as.factor(shdes[[binary_var]])
>
> t_test_two <- t.test(shdes[[num_var]] ~ shdes[[binary_var]])
>
> print(t_test_two)

Welch Two Sample t-test

data: shdes[[num_var]] by shdes[[binary_var]]
t = 3.3099, df = 610.62, p-value = 0.0009882
alternative hypothesis: true difference in means between group High and group Low is not equal to 0
95 percent confidence interval:
```

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```

R > R 4.1.2 ~ /
> shdes <- read.csv("shades.csv")
> head(shdes)
  brand brand_short product product_short hex H S V L group
1 Maybelline      mb Fit Me      fmf f3cfb3 26 0.26 0.95 86 2
2 Maybelline      mb Fit Me      fmf ffe3c2 32 0.24 1.00 92 2
3 Maybelline      mb Fit Me      fmf ffe0cd 23 0.20 1.00 91 2
4 Maybelline      mb Fit Me      fmf ffd3be 19 0.25 1.00 88 2
5 Maybelline      mb Fit Me      fmf bd9584 18 0.30 0.74 65 2
6 Maybelline      mb Fit Me      fmf eabda6 20 0.29 0.92 80 2
>
> print("---- 5. Independent Two-Sample t-test ----")
[1] "---- 5. Independent Two-Sample t-test ----"
>
> num_var <- "H"
> binary_var <- "group"
>
> if(length(unique(shdes[[binary_var]])) > 2){
+   shdes[[binary_var]] <- ifelse(shdes[[binary_var]] > median(shdes[[binary_var]]), "High", "Low")
+ }
> shdes[[binary_var]] <- as.factor(shdes[[binary_var]])
>
> t_test_two <- t.test(shdes[[num_var]] ~ shdes[[binary_var]])
> print(t_test_two)

welch Two Sample t-test

data: shdes[[num_var]] by shdes[[binary_var]]
t = 3.3099, df = 610.62, p-value = 0.0009882
alternative hypothesis: true difference in means between group High and group Low is not equal to 0
95 percent confidence interval:
 0.5736462 2.2474929
sample estimates:
mean in group High mean in group Low
      26.05119      24.64062
>

```

6) Performing paired t-tests using t.test(paired=TRUE) (R).

INPUT:-

```
df <- read.csv("shades.csv")
```

```
print("---- 6. Paired t-test for Shades Dataset ----")
```

```
num_var <- names(df)[sapply(df, is.numeric)][1]
```

```
set.seed(123)
```

```
before_values <- df[[num_var]] - runif(nrow(df), min = 0.1, max = 0.5)
```

```
after_values <- df[[num_var]]
```

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```
t_test_paired <- t.test(  
  before_values,  
  after_values,  
  paired = TRUE  
)
```

```
print(t_test_paired)
```

OUTPUT:-

```
> df <- read.csv("shades.csv")  
>  
> print("---- 6. Paired t-test for Shades Dataset ----")  
[1] "---- 6. Paired t-test for Shades Dataset ----"  
>  
> num_var <- names(df)[sapply(df, is.numeric)][1]  
>  
> set.seed(123)  
>  
> before_values <- df[[num_var]] - runif(nrow(df), min = 0.1, max = 0.5)  
> after_values <- df[[num_var]]  
>  
> t_test_paired <- t.test(  
+   before_values,  
+   after_values,  
+   paired = TRUE  
+ )  
>  
> print(t_test_paired)  
  
Paired t-test  
  
data: before_values and after_values  
t = -64.168, df = 612, p-value < 2.2e-16  
alternative hypothesis: true difference in means is not equal to 0  
95 percent confidence interval:  
 -0.3095536 -0.2911685  
sample estimates:  
mean of the differences  
 -0.300361  
  
> |
```