

R PROGRAMMING

1.Operations on vectors

- Create vectors A and B with multiple element
- Apply arithmetic operators +, -, *, /, and ^ on vectors A and B
- Apply rep(), paste() and cumprod() functions on A and analyse the results.

Code for above program :

```
# Create vectors A and B with multiple elements
A <- c(1, 2, 3, 4, 5)
B <- c(6, 7, 8, 9, 10)

# Arithmetic operations on vectors A and B
sum_A_B <- A + B # Addition
diff_A_B <- A - B # Subtraction
prod_A_B <- A * B # Element-wise multiplication
div_A_B <- A / B # Element-wise division
pow_A_B <- A ^ B # Element-wise exponentiation

# Print the results
cat("Sum of A and B:", sum_A_B, "\n")
cat("Difference of A and B:", diff_A_B, "\n")
cat("Product of A and B:", prod_A_B, "\n")
cat("Division of A and B:", div_A_B, "\n")
cat("Exponentiation of A and B:", pow_A_B, "\n")

# Apply rep(), paste(), and cumprod() functions on vector A
rep_A <- rep(A, times = 2) # Repeat A twice
paste_A <- paste("A:", A, sep = " ") # Concatenate "A:" with each element of A
cumprod_A <- cumprod(A) # Cumulative product of A

# Print the results
cat("Repeated A:", rep_A, "\n")
cat("Concatenated A:", paste_A, "\n")
```

```
cat("Cumulative product of A:", cumprod_A, "\n")
```

Algorithm for the code:

1. Start
2. Create an empty vector A
3. Create an empty vector B
4. Add elements to vector A
5. Add elements to vector B
6. Perform arithmetic operations on vectors A and B:
 - a. Add vector A and vector B, store the result in sum_A_B
 - b. Subtract vector B from vector A, store the result in diff_A_B
 - c. Multiply vector A and vector B element-wise, store the result in prod_A_B
 - d. Divide vector A by vector B element-wise, store the result in div_A_B
 - e. Raise vector A to the power of vector B element-wise, store the result in pow_A_B
7. Print the results of the arithmetic operations
8. Apply functions on vector A:
 - a. Repeat vector A twice using the rep() function, store the result in rep_A
 - b. Concatenate the string "A:" with each element of vector A using the paste() function, store the result in paste_A
 - c. Calculate the cumulative product of vector A using the cumprod() function, store the result in cumprod_A
9. Print the results of the functions applied on vector A
10. End

Methods used in the above program:

Vector Creation: Two empty vectors, A and B, are created using the c() function. Elements can be added to these vectors by assigning values using the assignment operator <-.

Arithmetic Operations:

Various arithmetic operations are performed on vectors A and B:

Addition (+): The + operator is used to add corresponding elements of vectors A and B, resulting in a new vector representing the sum.

Subtraction (-): The - operator subtracts corresponding elements of vector B from vector A, generating a new vector representing the difference.

Multiplication (*): The * operator performs element-wise multiplication between vectors A and B, producing a new vector with the products.

Division (/): The / operator divides corresponding elements of vector A by vector B, resulting in a new vector representing the division.

Exponentiation (^): The ^ operator raises vector A to the power of vector B element-wise, generating a new vector with the exponentiated values.

Functions:

rep(): The rep() function repeats the elements of vector A a specified number of times, generating a new vector. It takes two arguments: the vector to be repeated and the number of repetitions.

paste(): The paste() function concatenates multiple strings or expressions into a single string. In the code, it is used to concatenate the string "A:" with each element of vector A. The sep argument specifies the separator between elements.

cumprod(): The cumprod() function calculates the cumulative product of a vector. It returns a vector where each element represents the product of all preceding elements in the original vector.

Printing Results: The cat() function is used to print the results of the arithmetic operations and functions applied on vector A. It takes one or more arguments and prints them to the console.

These methods enable vector manipulation, mathematical calculations, and analysis in R programming, providing flexibility and efficiency in data processing.

Output :

```
Sum of A and B: 7 9 11 13 15
```

```
Difference of A and B: -5 -5 -5 -5 -5
```

```
Product of A and B: 6 14 24 36 50
```

```
Division of A and B: 0.1666667 0.2857143 0.375 0.4444444 0.5
```

```
Exponentiation of A and B: 1 128 6561 262144 9765625
```

```
Repeated A: 1 2 3 4 5 1 2 3 4 5
```

```
Concatenated A: A: 1 A: 2 A: 3 A: 4 A: 5
```

```
Cumulative product of A: 1 2 6 24 120
```

```
[Execution complete with exit code 0]
```

2. Operations on data frames

1. Create the following data frame, later invert gender for all individuals.

Name	Age	Height	Weight	Gender
Ram	30	177	57	M
Alwin	35	164	48	F
Billy	22	155	45	M
Amera	16	180	60	F
Olive	42	124	52	F
Dora	59	150	55	F

2. Add this data frame column-wise to the previous one (make sure you import the variable Working as character and not factor). How many rows and columns does the new data frame have? What class of data is in each column?

Name	Working
Ram	Y
Alwin	N
Billy	Y
Amera	N
Olive	Y
Dora	N

```
1. Code: Name <- c("Ram", "Alwin", "Billy", "Amera", "Olive",  
  "Dora")  
Age <- c(30, 35, 22, 16, 42, 59)  
Height <- c(177, 164, 155, 180, 124, 150)  
Weight <- c(57, 48, 45, 60, 52, 55)  
Gender <- as.factor(c("M", "F", "M", "F", "F", "F"))  
df <- data.frame(Name, Age, Height, Weight, Gender)  
# Invert gender for all individuals  
levels(df$Gender) <- c("M", "F")  
df$Gender <- as.factor(df$Gender)  
df
```

Output:

Name	Age	Height	Weight	Gender
Ram	30	177	57	F
Alwin	35	164	48	M
Billy	22	155	45	F
Amera	16	180	60	M
Olive	42	124	52	M
Dora	59	150	55	M

<pre>Name <- c("Ram", "Alwin", "Billy", "Amera", "Olive", "Dora") Age <- c(30, 35, 22, 16, 42, 59) Height <- c(177, 164, 155, 180, 124, 150) Weight <- c(57, 48, 45, 60, 52, 55) Gender <- as.factor(c("M", "F", "M", "F", "F", "F")) df <- data.frame(Name, Age, Height, Weight, Gender) # Invert gender for all individuals levels(df\$Gender) <- c("M", "F") df\$Gender <- as.factor(df\$Gender) df</pre>	<pre>Rscript /tmp/08R378qERm.r Name Age Height Weight Gender 1 Ram 30 177 57 F 2 Alwin 35 164 48 M 3 Billy 22 155 45 F 4 Amera 16 180 60 M 5 Olive 42 124 52 M 6 Dora 59 150 55 M</pre>
--	---

Algorithm:

1. Define the variables "Name", "Age", "Height", "Weight", and "Gender" as vectors containing the corresponding data for each individual.
2. Convert the "Gender" variable to a factor using the `as.factor()` function.
3. Combine the variables into a data frame using the `data.frame()` function.
4. Invert the levels of the "Gender" variable using the `levels()` function and reassign the levels to "M" and "F".
5. Assign the inverted "Gender" variable back to the data frame.
6. Print the modified data frame using the `print()` function.

Code:

```
Name <- c("Ram", "Alwin", "Billy", "Amera", "Olive", "Dora")
Age <- c(30, 35, 22, 16, 42, 59)
Height <- c(177, 164, 155, 180, 124, 150)
Weight <- c(57, 48, 45, 60, 52, 55)
Gender <- as.factor(c("M", "F", "M", "F", "F", "F"))
df <- data.frame(Name, Age, Height, Weight, Gender)
# Invert gender for all individuals
levels(df$Gender) <- c("M", "F")
df$Gender <- as.factor(df$Gender)
df
# Create the new data frame
Working <- c("Y", "N", "Y", "N", "Y", "N")
df2 <- data.frame(Working, stringsAsFactors = FALSE)
# Add the new data frame to the previous one
df <- cbind(df, df2)
# Print the modified data frame
df
```

Output:

Name	Age	Height	Weight	Gender
Ram	30	177	57	F
Alwin	35	164	48	M
Billy	22	155	45	F
Amera	16	180	60	M
Olive	42	124	52	M
Dora	59	150	55	M

Name	Age	Height	Weight	Gender	Working
Ram	30	177	57	F	Y
Alwin	35	164	48	M	N
Billy	22	155	45	F	Y
Amera	16	180	60	M	N
Olive	42	124	52	M	Y

Dora 59 150 55 M N

```
Name <- c("Ram", "Alwin", "Billy", "Amera", "Olive", "Dora")
Age <- c(30, 35, 22, 16, 42, 59)
Height <- c(177, 164, 155, 180, 124, 150)
Weight <- c(57, 48, 45, 60, 52, 55)
Gender <- as.factor(c("M", "F", "M", "F", "F", "F"))
df <- data.frame(Name, Age, Height, Weight, Gender)
# Invert gender for all individuals
levels(df$Gender) <- c("M", "F")
df$Gender <- as.factor(df$Gender)
df
# Create the new data frame
Working <- c("Y", "N", "Y", "N", "Y", "N")
df2 <- data.frame(Working, stringsAsFactors = FALSE)
# Add the new data frame to the previous one
df <- cbind(df, df2)
# Print the modified data frame
df
```

```
Rscript /tmp/08R378qERm.r
Name Age Height Weight Gender
1 Ram 30 177 57 F
2 Alwin 35 164 48 M
3 Billy 22 155 45 F
4 Amera 16 180 60 M
5 Olive 42 124 52 M
6 Dora 59 150 55 M
Name Age Height Weight Gender Working
1 Ram 30 177 57 F Y
2 Alwin 35 164 48 M N
3 Billy 22 155 45 F Y
4 Amera 16 180 60 M N
5 Olive 42 124 52 M Y
6 Dora 59 150 55 M N
```

The new data frame has 6 rows and 6 columns. The class of data in each column is:

1. "Name": character
2. "Age": numeric
3. "Height": numeric
4. "Weight": numeric
5. "Gender": factor
6. "Working": character

Algorithm:

1. Create a new data frame "df2" containing the variable "Working".
2. Add the new data frame "df2" to the previous data frame "df" using the cbind() function.
3. Print the modified data frame using the print() function.

Here are the methods used in the code:

1. `c()` function: This function is used to concatenate the values of the variables "Name", "Age", "Height", "Weight", and "Gender" into vectors.
2. `as.factor()` function: This function is used to convert the "Gender" vector into a factor.

3. ``data.frame()`` function: This function is used to combine the vectors into a data frame.
4. ``levels()`` function: This function is used to access and modify the levels of the "Gender" factor.
5. ``<-`` operator: This operator is used to assign the modified levels of the "Gender" factor back to the data frame.
6. ``print()`` function: This function is used to display the modified data frame.

3. DATASETS

Consider built-in dataset `state.x77`.

- a. Make sure the object is a data frame, if not change it to a data frame.
- b. Find out how many states have an income of less than 5000.
- c. Find out the state with the highest income and lowest income.
- d. Rename the column `state.region` so that only the first 3 letters appear.
- e. Remove the variable `div`, `Life Exp`, `HS Grad`, `Frost`, `abb`
- f. Add a variable to the data frame which should categorize the level of illiteracy: `[0,1)` is low, `[1,2)` is some, `[2, inf]` is high.
- g. Find out which state from the west, with low illiteracy, has the highest income, and what that income

ALGORITHM:

1. Check the class of the `state.x77` object.
2. Convert the `state.x77` object into a data frame named `s77`.
3. Check the class of the `s77` data frame.
4. Generate a summary of the `s77` data frame.

- 5.Count the number of rows in `s77` where the value in the "Income" column is less than 5000.
- 6.Retrieve the row name of the row with the maximum value in the "Income" column of `s77`.
- 7.Retrieve the row name of the row with the minimum value in the "Income" column of `s77`.
- 8.Create a new data frame named `df` by combining specific columns from different objects and using row names from `state.name`.
- 9.Modify the column names of `df` by extracting a substring from the existing column names.
- 10.Combine the `state.x77` and `df` data frames using the `cbind()` function to create a new data frame named `new.df`.
- 11.Remove the "div" column from `new.df`.
- 12.Subset `new.df` to remove columns 4, 6, 7, 9, and 10.
- 13.Create a new column named "Illiteracy.Levels" in `new.df` based on specific conditions using the `ifelse()` function.
- 14.Retrieve the values of the "Illiteracy.Levels" column in `new.df`.
- 15.Subset `new.df` to select rows where the "reg" column is "West" and the "Illiteracy.Levels" column is "Low" using the `subset()` function and store the result in `x`.
- 16.Retrieve the row name of the row with the maximum value in the "Income" column of `x`.
- 17.Retrieve the maximum value in the "Income" column of `x`.

Methods :

1.class() - This function is used to determine the class of an object, such as a data frame. It returns the class or type of the specified object.

2.data.frame() - This function is used to create a data frame from one or more vectors, matrices, or other data frames.

3.summary() - This method generates descriptive statistics of a data frame or specific columns within it. It provides a summary of the data, including minimum, 1st quartile, median, mean, 3rd quartile, and maximum values, as well as counts of missing values.

4.\$ - The dollar sign operator is used to access a specific column or variable within a data frame. It allows you to retrieve or modify values in a specific column.

5.nrow() - This function is used to count the number of rows in a data frame.

6.row.names() - This method is used to retrieve the row names of a data frame or a subset of a data frame.

7.which.max() - This function is used to find the index of the maximum value within a vector or column of a data frame.

8.which.min() - This function is used to find the index of the minimum value within a vector or column of a data frame.

9.substr() - This function is used to extract a substring from a character vector or column names. It allows you to specify the starting and ending positions of the substring.

10.cbind() - This function is used to combine vectors, matrices, or data frames column-wise. It binds the columns of multiple objects together to create a new data frame.

11.NULL - The NULL keyword is used to remove a column from a data frame. Assigning NULL to a column name effectively removes that column from the data frame.

12.subset() - This function is used to create a subset of a data frame based on specified conditions. It allows you to extract rows or columns that meet specific criteria.

13.ifelse() - This function is used to perform conditional operations in R. It allows you to specify a condition and return different values based on the condition.

Code:

```
class(state.x77)
```

```
s77 <- data.frame(state.x77)
```

```
class(s77)
```

#B

```
summary(s77)
```

```
nrow(s77[s77$Income < 5000,])
```

#C

```
row.names(s77 [which.max(s77$Income),])
```

```
row.names(s77 [which.min(s77$Income),])
```

#d

```
df<-
```

```
data.frame(state.abb,state.area,state.division,state.region,row.names=state.name)
```

```
colnames(df)<-substr(colnames(df),7,9)
```

```
new.df<-cbind(state.x77,df)
```

new.df

#e

```
new.df$div<-NULL
```

```
new.df<-subset(new.df,, -c(4,6,7,9,10))
```

```
new.df
```

#f

```
new.df$Illiteracy.Levels<-ifelse(new.df$Illiteracy    >=0    &
```

```
new.df$Illiteracy < 1, "Low",
```

```
ifelse(new.df$Illiteracy >= 1 &
```

```
new.df$Illiteracy < 2,"Some",
```

"High"))

```
new.df$Illiteracy.Levels
```

#g

```
x<-subset(new.df,reg=="West" & Illiteracy.Levels=="Low")
row.names(x[which.max(x$Income),])
```

Output:

```
> class(state.x77)
[1] "matrix" "array"
>
> s77 <- data.frame(state.x77)
> class(s77)
[1] "data.frame"
> #B
> summary(s77)
  Population      Income      Illiteracy      Life.Exp      Murder      HS.Grad      Frost
Min.   : 365   Min.   :3098   Min.   :0.500   Min.   :67.96   Min.   : 1.400   Min.   :37.80   Min.   : 0.00
1st Qu.:1080   1st Qu.:3993   1st Qu.:0.625   1st Qu.:70.12   1st Qu.: 4.350   1st Qu.:48.05   1st Qu.: 66.25
Median :2838   Median :4519   Median :0.950   Median :70.67   Median : 6.850   Median :53.25   Median :114.50
Mean   :4246   Mean   :4436   Mean   :1.170   Mean   :70.88   Mean   : 7.378   Mean   :53.11   Mean   :104.46
3rd Qu.:4968   3rd Qu.:4814   3rd Qu.:1.575   3rd Qu.:71.89   3rd Qu.:10.675   3rd Qu.:59.15   3rd Qu.:139.75
Max.   :21198   Max.   :6315   Max.   :2.800   Max.   :73.60   Max.   :15.100   Max.   :67.30   Max.   :188.00
  Area
Min.   : 1049
1st Qu.: 36985
Median : 54277
Mean   : 70736
3rd Qu.: 81163
Max.   :566432
> nrow(s77[s77$Income < 5000,])
[1] 42
> #C
> row.names(s77 [which.max(s77$Income),])
[1] "Alaska"
>
> row.names(s77 [which.min(s77$Income),])
[1] "Mississippi"
> #d
> df<-data.frame(state.abb,state.area,state.division,state.region,row.names=state.name)
> colnames(df)<-substr(colnames(df),7,9)
> new.df<-cbind(state.x77,df)
> new.df
  Population Income Illiteracy Life.Exp Murder HS.Grad Frost Area abb are div
Alabama    3615   3624      2.1   69.05   15.1   41.3    20  50708 AL  51609 East South Central
Alaska      365   6315      1.5   69.31   11.3   66.7   152 566432 AK  589757 Pacific
Arizona    2212   4530      1.8   70.55    7.8   58.1    15 113417 AZ  113909 Mountain
Arkansas    2110   3378      1.9   70.66   10.1   39.9    65  51945 AR  53104 West South Central
California  21198  5114      1.1   71.71   10.3   62.6    20 156361 CA  158693 Pacific
Colorado    2541  4884      0.7   72.06    6.8   63.9   166 103766 CO  104247 Mountain
Connecticut 3100  5348      1.1   72.48    3.1   56.0   139  4862 CT  5009 New England
Delaware     579  4809      0.9   70.06    6.2   54.6   103  1982 DE  2057 South Atlantic
Florida     8277  4815      1.3   70.66   10.7   52.6    11 54090 FL  58560 South Atlantic
Georgia     4931  4091      2.0   68.54   13.9   40.6    60  58073 GA  58876 South Atlantic
Hawaii       868  4963      1.9   73.60    6.2   61.9    0  6425 HI  6450 Pacific
Idaho        813  4119      0.6   71.87    5.3   59.5   126  82677 ID  83557 Mountain
Illinois    11197  5107      0.9   70.14   10.3   52.6   127  55748 IL  56400 East North Central
```

Ohio	10735	4561	0.8	70.82	7.4	53.2	124	40975	OH	41222	East North Central
Oklahoma	2715	3983	1.1	71.42	6.4	51.6	82	68782	OK	69919	West South Central
Oregon	2284	4660	0.6	72.13	4.2	60.0	44	96184	OR	96981	Pacific
Pennsylvania	11860	4449	1.0	70.43	6.1	50.2	126	44966	PA	45333	Middle Atlantic
Rhode Island	931	4558	1.3	71.90	2.4	46.4	127	1049	RI	1214	New England
South Carolina	2816	3635	2.3	67.96	11.6	37.8	65	30225	SC	31055	South Atlantic
South Dakota	681	4167	0.5	72.08	1.7	53.3	172	75955	SD	77047	West North Central
Tennessee	4173	3821	1.7	70.11	11.0	41.8	70	41328	TN	42244	East South Central
Texas	12237	4188	2.2	70.90	12.2	47.4	35	262134	TX	267339	West South Central
Utah	1203	4022	0.6	72.90	4.5	67.3	137	82096	UT	84916	Mountain
Vermont	472	3907	0.6	71.64	5.5	57.1	168	9267	VT	9609	New England
Virginia	4981	4701	1.4	70.08	9.5	47.8	85	39780	VA	40815	South Atlantic
Washington	3559	4864	0.6	71.72	4.3	63.5	32	66570	WA	68192	Pacific
West Virginia	1799	3617	1.4	69.48	6.7	41.6	100	24070	WV	24181	South Atlantic
Wisconsin	4589	4468	0.7	72.48	3.0	54.5	149	54464	WI	56154	East North Central
Wyoming	376	4566	0.6	70.29	6.9	62.9	173	97203	WY	97914	Mountain
Alabama											South
Alaska											West
Arizona											West
Arkansas											South
California											West
Colorado											West
Connecticut											Northeast
Delaware											South
Florida											South
Georgia											South
Hawaii											West

Idaho											West
Illinois											North Central
Indiana											North Central
Iowa											North Central
Kansas											North Central
Kentucky											South
Louisiana											South
Maine											Northeast
Maryland											South
Massachusetts											Northeast
Michigan											North Central
Minnesota											North Central
Mississippi											South
Missouri											North Central
Montana											West
Nebraska											North Central
Nevada											West
New Hampshire											Northeast
New Jersey											Northeast
New Mexico											West
New York											Northeast
North Carolina											South
North Dakota											North Central
Ohio											North Central
Oklahoma											South
Oregon											West
Pennsylvania											Northeast
Rhode Island											Northeast
South Carolina											South
Wyoming											West
> #e											
> new.df\$div<-NULL											
> new.df<-subset(new.df,, -c(4,6,7,9,10))											
> new.df											
	Population	Income	Illiteracy	Murder	Area		reg				
Alabama	3615	3624	2.1	15.1	50708		South				
Alaska	365	6315	1.5	11.3	566432		West				
Arizona	2212	4530	1.8	7.8	113417		West				
Arkansas	2110	3378	1.9	10.1	51945		South				
California	21198	5114	1.1	10.3	156361		West				
Colorado	2541	4884	0.7	6.8	103766		West				
Connecticut	3100	5348	1.1	3.1	4862		Northeast				
Delaware	579	4809	0.9	6.2	1982		South				
Florida	8277	4815	1.3	10.7	54090		South				
Georgia	4931	4091	2.0	13.9	58073		South				
Hawaii	868	4963	1.9	6.2	6425		West				
Idaho	813	4119	0.6	5.3	82677		West				
Illinois	11197	5107	0.9	10.3	55748		North Central				
Indiana	5313	4458	0.7	7.1	36097		North Central				
Iowa	2861	4628	0.5	2.3	55941		North Central				
Kansas	2280	4669	0.6	4.5	81787		North Central				
Kentucky	3387	3712	1.6	10.6	39650		South				
Louisiana	3806	3545	2.8	13.2	44930		South				
Maine	1058	3694	0.7	2.7	30920		Northeast				
Maryland	4122	5299	0.9	8.5	9891		South				
Massachusetts	5814	4755	1.1	3.3	7826		Northeast				
Michigan	9111	4751	0.9	11.1	56817		North Central				

maine	1038	3034	0.7	2.7	30320	Northeast
Maryland	4122	5299	0.9	8.5	9891	South
Massachusetts	5814	4755	1.1	3.3	7826	Northeast
Michigan	9111	4751	0.9	11.1	56817	North Central
Minnesota	3921	4675	0.6	2.3	79289	North Central
Mississippi	2341	3098	2.4	12.5	47296	South
Missouri	4767	4254	0.8	9.3	68995	North Central
Montana	746	4347	0.6	5.0	145587	West
Nebraska	1544	4508	0.6	2.9	76483	North Central
Nevada	590	5149	0.5	11.5	109889	West
New Hampshire	812	4281	0.7	3.3	9027	Northeast
New Jersey	7333	5237	1.1	5.2	7521	Northeast
New Mexico	1144	3601	2.2	9.7	121412	West
New York	18076	4903	1.4	10.9	47831	Northeast
North Carolina	5441	3875	1.8	11.1	48798	South
North Dakota	637	5087	0.8	1.4	69273	North Central
Ohio	10735	4561	0.8	7.4	40975	North Central
Oklahoma	2715	3983	1.1	6.4	68782	South
Oregon	2284	4660	0.6	4.2	96184	West
Pennsylvania	11860	4449	1.0	6.1	44966	Northeast
Rhode Island	931	4558	1.3	2.4	1049	Northeast
South Carolina	2816	3635	2.3	11.6	30225	South
South Dakota	681	4167	0.5	1.7	75955	North Central
Tennessee	4173	3821	1.7	11.0	41328	South
Texas	12237	4188	2.2	12.2	262134	South
Utah	1203	4022	0.6	4.5	82096	West
Vermont	472	3907	0.6	5.5	9267	Northeast
Virginia	4981	4701	1.4	9.5	39780	South
Washington	3559	4864	0.6	4.3	66570	West

```

Console terminal background jobs
R 4.3.0 · C:/Users/vishu/Desktop/R/ ↗
Oregon 2284 4660 0.6 4.2 96184 West
Pennsylvania 11860 4449 1.0 6.1 44966 Northeast
Rhode Island 931 4558 1.3 2.4 1049 Northeast
South Carolina 2816 3635 2.3 11.6 30225 South
South Dakota 681 4167 0.5 1.7 75955 North Central
Tennessee 4173 3821 1.7 11.0 41328 South
Texas 12237 4188 2.2 12.2 262134 South
Utah 1203 4022 0.6 4.5 82096 West
Vermont 472 3907 0.6 5.5 9267 Northeast
Virginia 4981 4701 1.4 9.5 39780 South
Washington 3559 4864 0.6 4.3 66570 West
West Virginia 1799 3617 1.4 6.7 24070 South
Wisconsin 4589 4468 0.7 3.0 54464 North Central
Wyoming 376 4566 0.6 6.9 97203 West
> #f
> new.df$illiteracy.Levels<-ifelse(new.df$illiteracy >=0 & new.df$illiteracy < 1, "Low",
+ ifelse(new.df$illiteracy >= 1 & new.df$illiteracy < 2,"Some",
+ "High"))
> new.df$illiteracy.Levels
[1] "High" "Some" "Some" "Some" "Some" "Low" "Some" "Low" "Some" "High" "Some" "Low" "Low" "Low" "Low" "Low" "Some"
[18] "High" "Low" "Low" "Some" "Low" "Low" "High" "Low" "Low" "Low" "Low" "Some" "High" "Some" "Some" "Some" "Low"
[35] "Low" "Some" "Low" "Some" "Some" "High" "Low" "Some" "High" "Low" "Low" "Some" "Low" "Some" "Low" "Low"
>
> #g
> x<-subset(new.df,reg=="West" & illiteracy.Levels=="Low")
> row.names(x[which.max(x$Income),])
[1] "Nevada"
>

```

4. Loop implementation

- Write a nested loop, where the outer for loop increments "a" 3 times, and the inner for loop increments "b" 3 times. The break statement exits the inner for loop after 2 incrementations. The nested loop prints the values of variables, "a" and "b".
- The next statement is used within loops in order to skip the current evaluation, and instead proceed to the next evaluation. Therefore, write a while loop that prints the variable "; that is incremented from 2 - 5, and uses the next statement, to skip the printing of the number.

ALGORITHM: A: 1. Start

2. Set the initial value of "a" to 1.
 3. Start the outer loop:
 4. Repeat the following steps 3 times:
 - a. Set the initial value of "b" to 1.
 - b. Start the inner loop:
 - c. Print the values of "a" and "b".
 - d. Increment the value of "b" by 1.
 - e. If the value of "b" is equal to 2, break out of the inner loop.
 - f. Increment the value of "a" by 1.
 5. End the outer loop.
 6. End B.
-
1. Start
 2. Set the initial value of the variable "x" to 2.
 3. Start the while loop:
 4. Repeat the following steps as long as the value of "x" is less than or equal to
 - 5: a. If the value of "x" is equal to 4, use the "next" statement to skip the current iteration.
 - b. Print the value of "x".
 - c. Increment the value of "x" by 1.
 5. End the while loop
 - . 6. End

METHODS USED IN THE CODE: 1. for()- used to repeat a specific block of code a known number of times. Syntax:
 for(expression1;expression2;expression3({ //code to be executed }

2. break()- a loop control statement that is used to terminate the loop.
Syntax: break;

3. print()- used to print the statement given within the double codes.
Syntax: print(" ");

4. if()- a decision making statement that is used to execute a block of code based on the value of the given expression. Syntax:
if(expression){ //code to be executed if condition is true }else{ //code to be executed if the condition is false }

5. while()- allows user to repeat a statement until a specified expression becomes false. Syntax: while(condition){ //code to be executed }

6. next()- a continue statement causes the iteration to go to the next statement of the loop, skipping the code in between. Syntax: continue;
CODE: a. for (a in 1:3)

```
{  
for (b in 1:3)  
{  
  print(paste("a =", a, "b =", b))  
  if (b == 2)  
  {  
    break  
  }  
}  
}
```

b.

```
i <- 2 while (i <= 5)  
{
```



```

if (i == 4)
{
i <- i + 1 next
}
print(i)
i <- i + 1
}

```

5. Create a function that given an integer will calculate how many divisors it has (other than 1 and itself). Print all the divisors and its count.

ALGORITHM

1. Start with an empty list of divisors.
2. Iterate through the numbers from 2 to (num - 1).
3. For each number, check if it divides the given integer evenly (i.e., the remainder of the division is 0).
4. If the number is a divisor, add it to the list of divisors.
5. After iterating through all the numbers, count the number of divisors in the list.
6. Print the list of divisors and the count.

PROGRAM CODE:

R program that calculates the number of divisors for a given integer and prints all the divisors along with their count:

```

countDivisors <- function(num) {
  divisors <- c()
  # Find divisors
  for (i in 2:(num-1)) {
    if (num %% i == 0) {
      divisors <- c(divisors, i)
    }
  }
}

```

```

divisorCount <- length(divisors)

# Print divisors and count

cat("Divisors:", divisors, "\n")

cat("Number of Divisors:", divisorCount, "\n")

}

# Read input from user

num <- as.integer(readline("Enter an integer: "))

# Call the function with user input

countDivisors(num)

```

FUNCTIONS EXPLANATION:

1. `countDivisors(num)`: This is the main function that calculates the number of divisors for a given integer. It takes an input parameter `num`, representing the integer for which we want to find the divisors. Here's a detailed explanation of the steps within this function:

- `divisors`: This is an empty vector that will store the divisors of the given integer `num`.
- `for` loop: It iterates through the numbers from 2 to `(num - 1)`. This loop will check each number to see if it divides `num` evenly.
- `if` statement: Within the loop, it checks if the current number `i` divides `num` evenly, which is determined by `num %% i == 0`. If the condition is true, it means that `i` is a divisor of `num`.
- Adding divisors: If `i` is a divisor, it is added to the `divisors` vector using the `c()` function. This appends the current divisor to the existing list of divisors.

`divisorCount`: After the loop completes, we determine the number of divisors by calculating the length of the `divisors` vector using the `length()` function.

Printing divisors and count: Finally, the function prints the list of divisors using the `cat()` function, followed by the count of divisors.

2. `readline("Enter an integer: ")`: This function is used to read input from the user. It prompts the user to enter an integer and waits for the user's input. The input provided by the user is returned as a string.

3. `as.integer()`: This function is used to convert the user input (which is initially a string) into an integer data type. It takes a value as an argument and attempts to convert it into an integer. In this code, it is used to convert the user input string into an integer and store it in the variable `num`.

4. `cat("Divisors:", divisors, "\n")`: This function is used to print the divisors of the given integer. It takes three arguments: a string `"Divisors:"`, the `divisors` vector containing the divisors, and `"\n"` to add a newline character after printing the divisors.

5. `cat("Number of Divisors:", divisorCount, "\n")`: This function is used to print the count of divisors. It takes three arguments: a string `"Number of Divisors:"`, the `divisorCount` variable storing the count, and `"\n"` to add a newline character after printing the count.

These functions work together to calculate the divisors of a given integer, read user input, convert the input to an integer, and display the divisors along with their count.

OUTPUT:

```
```R
```

```
Enter an integer: 24
```

```
Divisors: 2 3 4 6 8 12
```

```
Number of Divisors: 6
```

```
```
```

In this example, the user inputs the number 24. The code then calculates the divisors of 24, which are 2, 3, 4, 6, 8, and 12. The code prints the divisors and the count, indicating that there are 6 divisors for the number 24.

6. Create a function that given a data frame, and a number or character. The function will return the data frame with the character or number changed to NA.

ALGORITHM:-

The algorithm used in the `replace_with_na` function is as follows:

The function `replace_with_na` takes two arguments: `data_frame`, which represents the input data frame, and `value`, which is the number or character to be replaced with NA.

The function loops through each column of the data frame using `for (col in colnames(data_frame))`.

Within the loop, the function checks if the elements in the current column (`data_frame[[col]]`) are equal to the specified value (`value`) using `data_frame[[col]] == value`.

If an element in the column matches the specified value, the function assigns NA to that element using the assignment operator `<- NA`.

After iterating through all the columns, the modified data frame is returned using `return(data_frame)`.

METHODS USED IN THE CODE:-

1. `for()`- used to repeat a specific block of code a known number of times.

Syntax: `for(expression1;expression2;expression3(`

```
{  
//code to be executed  
}
```

CODE:-

Here we have created sample data frame

```
df <- data.frame(  
  col1 = c(1, 2, 3, 4),  
  col2 = c("a", "b", "c", "d"),  
  col3 = c(5, 6, 7, 8)  
)
```

Print the original data frame

```
print(df);
```

OUTPUT:-

```
col1 col2 col3
```

```
1  1  a  5
```

```
2  2  b  6
```

```
3  3  c  7
```

```
4  4  d  8
```

After Using Replace using NA Function

Here's a function in R that takes a data frame and a number or character and replaces all occurrences of that number or character with NA

CODE:-

```
replace_with_na <- function(data_frame, value) {  
  # Loop through each column of the data frame  
  for (col in colnames(data_frame)) {  
    # Replace the specified value with NA in the column  
    data_frame[[col]][data_frame[[col]] == value] <- NA  
  }  
  
  return(data_frame)
```

```

}

# Create a sample data frame

df <- data.frame(
  col1 = c(1, 2, 3, 4),
  col2 = c("a", "b", "c", "d"),
  col3 = c(5, 6, 7, 8)
)

# Print the original data frame

print(df)

# Call the replace_with_na function to replace "b" with NA

df_new <- replace_with_na(df, "b")

# Print the modified data frame

print(df_new)

```

OUTPUT:-

```
col1 col2 col3
```

```
1 1 a 5
```

```
2 2 b 6
```

```
3 3 c 7
```

```
4 4 d 8
```

```
col1 col2 col3
```

```
1 1 a 5
```

```
2 2 <NA> 6
```

```
3 3 c 7
```

```
4 4 d 8
```

Here the function `replace_with_na` is called with the data frame `df` and the character "b". It replaces all occurrences of "b" in the data frame with NA and returns the modified data frame `df_new`.

7. Consider the file `table0.txt` containing the following contents.

| Name | Age | Height | Weight | Gender |
|-------|-----|--------|--------|--------|
| Ram | 30 | 177 | 57 | M |
| Alwin | 35 | 164 | 48 | F |
| Billy | -- | 155 | 45 | M |
| Amera | 16 | * | 60 | F |
| Olive | 42 | 124 | ** | F |
| Dora | 59 | 150 | 55 | F |

- Read the contents of the file.
- Find the missing values.
- Replace the missing values on column 4, with the average of the column values.

Step 1: Prepare the file

Create a file named `table0.txt` with the provided contents:

| Name | Age | Height | Weight | Gender |
|-------|-----|--------|--------|--------|
| Ram | 30 | 177 | 57 | M |
| Alwin | 35 | 164 | 48 | F |
| Billy | -- | 155 | 45 | M |
| Amera | 16 | * | 60 | F |
| Olive | 42 | 124 | ** | F |
| Dora | 59 | 150 | 55 | F |

Step 2: Set up R environment

We have installed 'RStudio' for the implementation.

Step 3: Algorithm

- Read the contents of the file "table0.txt" using the `'read.delim()'` function and store it in the 'contents' data frame.
- Find the missing values:
 - Create a subset of 'contents' using the `'subset()'` function, filtering rows where 'Age', 'Height', or 'Weight' values are '--', '*', or '**'.
 - Store the subset in 'missing_values'.
 - Print "Missing values" and 'missing_values'.
- Replace the missing values with the average:
 - Convert the 'Weight' column of 'contents' to numeric values using `'as.numeric(as.character())'`.

- b. Calculate the average of the numeric values using the 'mean()' function, excluding any missing values ('NA') by specifying 'na.rm = TRUE'.
- c. Assign the average value to the rows in the 'Weight' column where the value is '--' or '**' using logical indexing.
- d. Print "Modified contents" and contents.

Step 4: Implementation of the code

```
find_missing.r
1 # Read the content of the file:
2 file_path <- "table0.txt"
3 contents <- read.delim(file_path, header = TRUE, sep = "\t")
4 print(contents)
5
6
7 #Find the missing values:
8 missing_values <- subset(contents, Age == "--" | Height == "--" | Weight == "--" | Age
9 print("Missing values:")
10 print(missing_values)
11
12
13 # Replace the missing values in column 4 with the average of the column values:
14 numeric_values <- as.numeric(as.character(contents$Weight))
15 average <- mean(numeric_values, na.rm = TRUE)
16
17 contents$Weight[contents$Weight == "--" | contents$Weight == "**"] <- average
18
19 print("Modified contents:")
20 print(contents)
```

Explanation:

a. Reading the contents of the file:

We use the read.table() function to read the contents of the file "table0.txt". We specify header = TRUE to indicate that the first row contains column names, and sep = "\t" to specify that the columns are separated by a tab character.

b. Finding the missing values:

We create a logical matrix missing_values to identify missing values in the data. In this case, we consider "--", "**", and "*" as missing values. The expression is.na(data) checks for NA values, and data == "--" checks for values equal to "--", and so on.

c. Replacing missing values in column 4:

We calculate the average of the values in the "Weight" column using the mean() function, and store it in the variable average_weight. Then, we replace the missing values in the "Weight" column with the calculated average by assigning average_weight to the corresponding elements of data\$Weight.

Finally, we print the updated data using `print(data)`.

We have to make sure that the file "table0.txt" is in the working directory of R environment or else we need to provide the full path to the file in the `read.table()` function.

Functions used in the Program:

1. read.delim(): This function is used to read tabular data from a file with delimiter-separated values. It is similar to `read.table()`, but specifically designed to read tab-separated files. In the code, it reads the contents of the file "table0.txt" and returns a data frame, which is stored in the variable `contents`.

2. subset(): This function is used to extract a subset of rows from a data frame based on specified conditions. In the code, it is used to create a subset `missing_values` by filtering rows where the values in the 'Age', 'Height', or 'Weight' columns are '--', '*', or '**'.

3. print(): This function is used to display output on the console. In the code, it is used to print the messages and the contents of the data frames (`missing_values` and `contents`).

4. as.numeric(): This function is used to convert a vector or a variable to a numeric type. In the code, it is used to convert the 'Weight' column of the `contents` data frame to numeric values. `as.character()` is used to convert the values to character type before converting them to numeric.

5. mean(): This function is used to calculate the mean (average) of a numeric vector or values. In the code, it is used to calculate the average of the numeric values in the 'Weight' column, excluding any missing values (NA). The `na.rm = TRUE` parameter is used to handle missing values and ignore them in the calculation.

6. Logical Indexing: Logical indexing is used to select specific rows in a data frame based on a condition. In the code, it is used to identify the rows in the 'Weight' column where the value is '--' or '**', and replace those values with the calculated average.

Step 5: Run the code.

```
> file_path <- "table0.txt"
> contents <- read.delim(file_path, header = TRUE, sep = "\t")
> print(contents)
  Name Age Height Weight Gender
1  Ram  30   177    57      M
2 Alwin  35   164    48      F
3 Billy --   155    45      M
4 Amera 16     *    60      F
5 Olive 42   124    **      F
6 Dora  59   150    55      F
> missing_values <- subset(contents, Age == "--" | Height == "--" | Weight == "--" | Age == "*" | Height == "*" | Weight == "**")
> print("Missing values:")
[1] "Missing values:"
> print(missing_values)
  Name Age Height Weight Gender
3 Billy --   155    45      M
4 Amera 16     *    60      F
5 Olive 42   124    **      F
> numeric_values <- as.numeric(as.character(contents$Weight))
Warning message:
NAs introduced by coercion
> average <- mean(numeric_values, na.rm = TRUE)
>
> contents$Weight[contents$Weight == "--" | contents$Weight == "**"] <- average
>
> print("Modified contents:")
[1] "Modified contents:"
> print(contents)
  Name Age Height Weight Gender
1  Ram  30   177    57      M
2 Alwin  35   164    48      F
3 Billy --   155    45      M
4 Amera 16     *    60      F
5 Olive 42   124    53      F
6 Dora  59   150    55      F
```


8.. Read the file Table1.txt

| | | | | |
|-------|----|-----|----|---|
| Ram | 30 | 177 | 57 | M |
| Alwin | 35 | 164 | 48 | F |
| Billy | 28 | 155 | 45 | M |
| Amera | 16 | 168 | 60 | F |
| Olive | 42 | 124 | 70 | F |
| Dora | 59 | 150 | 55 | F |

a. Change the names of the columns to Name, Age, Height, Weight and Gender

b. Change the row names so that they are the same as Name, and remove the variable Name.

Program code:

Read the file

```
data <- read.table("Table1.txt", header = FALSE)
```

Set the column names

```
colnames(data) <- c("Name", "Age", "Height", "Weight", "Gender")
```

Set the row names to the values in the first column (Name)

```
rownames(data) <- data$Name
```

Remove the Name variable

```
data <- data[, -1]
```

Print the modified data

```
print(data)
```

Title: R Program to Modify Column and Row Names in Table Data

Program Steps:

1. Reading the File:

The program starts by reading the contents of the "Table1.txt" file using the ``read.table()`` function. The ``read.table()`` function is a versatile tool in R used to read tabular data from files. In this case, we set the ``header`` parameter to ``FALSE`` since the file does not contain a header row.

2. Changing Column Names:

To change the column names, the program uses the ``colnames()`` function. By assigning a character vector with the desired column names to ``colnames(data)``, the program modifies the column names to "Name," "Age," "Height," "Weight," and "Gender." This step ensures that the column names accurately represent the corresponding variables in the data.

3. Setting Row Names:

Next, the program sets the row names of the data using the ``rownames()`` function. To achieve this, the program assigns the values from the "Name" column to ``rownames(data)``. By doing so, the row names become identical to the values in the "Name" column, enabling easier identification and indexing of rows.

4. Removing the Name Variable:

To remove the "Name" variable from the data, the program uses R's subsetting capabilities. By excluding the first column (Name) from the data, the program retains only the "Age," "Height," "Weight," and "Gender" variables. This step ensures that the modified data no longer contains the "Name" variable.

5. Printing the Modified Data:

Finally, the program prints the modified data to the console using the ``print()`` function. The resulting output displays the table with updated column names and row names, providing a clear representation of the modified data.

Algorithms:

1. Read the file:

1. Start

2. Read the file "Table1.txt" using the `read.table()` function and assign the data to a variable (e.g., `data`)

3. Return the data

4. Stop

2. Set the column names:
 1. Start
 2. Set the column names of the data using the `colnames()` function
 3. Assign the desired column names to `colnames(data)` as a character vector
 4. Return the modified data
 5. Stop
3. Set the Row names:
 1. Start
 2. Set the row names of the data using the `rownames()` function
 3. Assign the values from the "Name" column of the data to `rownames(data)`
 4. Return the modified data
 5. Stop
4. Removing the Name Variable:
 1. Start
 2. Remove the first column (Name) from the data using subsetting
 3. Assign the modified data to a new variable (e.g., `data_modified`)
 4. Return the `data_modified`
 5. Stop
5. Printing the modified data:
 1. Start
 2. Print the modified data using the `print()` function
 3. Stop

Outputs:

- **dataset**

V1 V2 V3 V4 V5

1 Ram 30 177 57 M

2 Alwin 35 164 48 F

```

3 Billy 28 155 45 M
4 Amera 16 168 60 F
5 Olive 42 124 70 F
6 Dora 59 150 55 F

```

- Set the column names

```

> colnames(data) <- c("Name", "Age", "Height", "Weight", "Gender")
> data

```

| | Name | Age | Height | Weight | Gender |
|---|-------|-----|--------|--------|--------|
| 1 | Ram | 30 | 177 | 57 | M |
| 2 | Alwin | 35 | 164 | 48 | F |
| 3 | Billy | 28 | 155 | 45 | M |
| 4 | Amera | 16 | 168 | 60 | F |
| 5 | Olive | 42 | 124 | 70 | F |
| 6 | Dora | 59 | 150 | 55 | F |

- Set the row names to the values in the first column (Name)

```

> rownames(data) <- data$Name
> data

```

| | Name | Age | Height | Weight | Gender |
|-------|-------|-----|--------|--------|--------|
| Ram | Ram | 30 | 177 | 57 | M |
| Alwin | Alwin | 35 | 164 | 48 | F |
| Billy | Billy | 28 | 155 | 45 | M |
| Amera | Amera | 16 | 168 | 60 | F |
| Olive | Olive | 42 | 124 | 70 | F |
| Dora | Dora | 59 | 150 | 55 | F |

- **Remove the Name variable**

```
> data <- data[, -1]
```

```
> # Print the modified data
```

```
> print(data)
```

| | Age | Height | Weight | Gender |
|-------|-----|--------|--------|--------|
| Ram | 30 | 177 | 57 | M |
| Alwin | 35 | 164 | 48 | F |
| Billy | 28 | 155 | 45 | M |
| Amera | 16 | 168 | 60 | F |
| Olive | 42 | 124 | 70 | F |
| Dora | 59 | 150 | 55 | F |

9. Plot Data

- Plot Miles/(US) gallon versus Rear axle ratio by plot(mpg,drat).
- Change type of visualisation to scatterplot.

Algorithm:

Load the mtcars dataset.

Plot the Miles/(US) gallon versus Rear axle ratio using the plot() function with 'mpg' and 'drat' columns.

Customize the plot by setting the title, x-axis label, and y-axis label.

Display the line plot.

Modify the plot by changing the visualization type to a scatter plot.

Display the scatter plot.

Methods Used in the Code:

- plot() - Used to create a basic line or scatter plot.
- data() - Used to load the mtcars dataset.
- main, xlab, ylab - Arguments used to set the title, x-axis label, and y-axis label in the plot.

Code:

```
# Load the mtcars dataset
data(mtcars)

# Plot Miles/(US) gallon versus Rear axle ratio as a line plot
plot(mtcars$mpg, mtcars$drat, main = "Miles/(US) gallon versus Rear axle
ratio",
      xlab = "Miles/(US) gallon", ylab = "Rear axle ratio")

# Change the visualization to a scatter plot
plot(mtcars$mpg, mtcars$drat, main = "Miles/(US) gallon versus Rear axle ratio
(Scatter Plot)",
      xlab = "Miles/(US) gallon", ylab = "Rear axle ratio", type = "p")
```

Step 1: Load the dataset

To begin, we load the mtcars dataset into our R environment. The following code accomplishes this task:

```
data(mtcars)
```

Step 2: Plotting Miles/(US) gallon versus Rear axle ratio as a line plot

For the initial line plot, we make use of the plot() function, considering the 'mpg' and 'drat' columns from the mtcars dataset. Our code implementation is as follows:

```
plot(mtcars$mpg, mtcars$drat, main = "Miles/(US) gallon versus Rear axle
ratio",
      xlab = "Miles/(US) gallon", ylab = "Rear axle ratio")
```

This code generates a line plot where the x-axis represents Miles/(US) gallon, while the y-axis represents Rear axle ratio. The main, xlab, and ylab arguments are utilized to set the title, x-axis label, and y-axis label, respectively.

Output:

The output of the above code is a line plot showing the relationship between Miles/(US) gallon and Rear axle ratio.

Step 3: Changing the visualization to a scatter plot

To transform the line plot into a scatter plot, we modify the code by including the type argument within the plot() function. Here is the updated code:

```
plot(mtcars$mpg, mtcars$drat, main = "Miles/(US) gallon versus Rear axle ratio  
(Scatter Plot)",
```

```
      xlab = "Miles/(US) gallon", ylab = "Rear axle ratio", type = "p")
```

This code produces a scatter plot where each data point corresponds to a car from the mtcars dataset. The title, x-axis label, and y-axis label are appropriately specified.

Output:

The output of the modified code is a scatter plot illustrating the relationship between Miles/(US) gallon and Rear axle ratio.

Explanation:

In our solution, we followed a step-by-step approach to plot the Miles/(US) gallon versus Rear axle ratio using the mtcars dataset. Firstly, we loaded the mtcars dataset using the data(mtcars) command.

Then, we created a line plot by utilizing the plot() function with the 'mpg' and 'drat' columns from the mtcars dataset. The resulting plot demonstrated the relationship between the two variables.

To change the visualization to a scatter plot, we updated the code by incorporating the type = "p" argument within the plot() function. This alteration transformed the line plot into a scatter plot, providing a different representation of the data.

10. Select four different continuous variables from the mtcars dataset.

a. Plot with lines and points for different variables with different colors.

b. Choose one variable from each and highlight it with different color.

Function to create the employee data frame

```
createEmployeeDataFrame <- function() {  
  # Read the number of employees from the user  
  numEmployees <- readline("Enter the number of employees: ")  
  numEmployees <- as.integer(numEmployees)  
  # Initialize empty vectors for each field  
  empID <- vector("integer", numEmployees)  
  empName <- vector("character", numEmployees)  
  doj <- vector("character", numEmployees)  
  dept <- vector("character", numEmployees)  
  desig <- vector("character", numEmployees)
```

Loop to read employee details

```
for (i in 1:numEmployees) {  
  cat(paste("Enter details for Employee", i, ":\n"))
```

Read employee details from the user

```
empID[i] <- readline("EmpID: ")  
empName[i] <- readline("EmpName: ")  
doj[i] <- readline("DOJ: ")  
dept[i] <- readline("Dept: ")  
desig[i] <- readline("Desig: ")  
}
```

Create the data frame

```
employeeData <- data.frame(EmpID = empID, EmpName = empName, DOJ = doj, Dept = dept,  
Desig = desig)
```



```
    return(employeeData)
}
```

Function to write the employee data frame to CSV

```
writeDataFrameToCSV <- function(employeeData, fileName) {
  write_csv(employeeData, fileName)
  cat(paste("Employee data written to", fileName, "successfully!\n"))
}
```

Function to read and display the contents of the CSV file

```
readAndDisplayCSV <- function(fileName) {
  employeeData <- read_csv(fileName)
  cat(paste("Contents of", fileName, ":\n"))
  print(employeeData)
}
```

Function to append a new row to the CSV file

```
appendRowToCSV <- function(employeeData, fileName) {
  newEmployeeData <- createEmployeeDataFrame()
  appendedData <- rbind(employeeData, newEmployeeData)
  write_csv(appendedData, fileName)
  cat(paste("New row appended to", fileName, "successfully!\n"))
}
```

Main program

```
csvFileName <- "check.csv"
```

Step 1: Create the employee data frame

```
employeeData <- createEmployeeDataFrame()
```

Step 2: Store the data frame in the CSV file

```
writeDataFrameToCSV(employeeData, csvFileName)
```

```
# Step 3: Read and display the contents of the CSV file
```

```
readAndDisplayCSV(csvFileName)
```

```
# Step 4: Append a new row to the CSV file
```

```
appendRowToCSV(employeeData, csvFileName)
```

ALGORITHM

1. The ``createEmployeeDataFrame`` function prompts the user to enter the number of employees. It then initializes empty vectors for each employee field (EmpID, EmpName, DOJ, Dept, Desig). It uses a loop to read employee details from the user and stores them in the respective vectors. Finally, it creates a data frame using the collected employee data and returns it.

2. The ``writeDataFrameToCSV`` function takes an employee data frame and a file name as inputs. It writes the data frame to a CSV file using the ``write_csv`` function from the appropriate library. It displays a success message after the operation.

3. The ``readAndDisplayCSV`` function takes a file name as an input. It reads the contents of the CSV file using the ``read_csv`` function and displays them using the ``print`` function.

4. The ``appendRowToCSV`` function takes an existing employee data frame and a file name as inputs. It calls the ``createEmployeeDataFrame`` function to collect details for a new employee. It then appends the new employee data to the existing data frame using the ``rbind`` function. The updated data frame is written to the CSV file using the ``write_csv`` function, and a success message is displayed.

METHODS USED

- 'readline': Used to read user input for the number of employees and employee details.

- 'vector': Used to initialize empty vectors for each employee field.

- 'for' loop: Used to iterate over the employee details input process.

- 'rbind': Used to append a new row to the employee data frame.

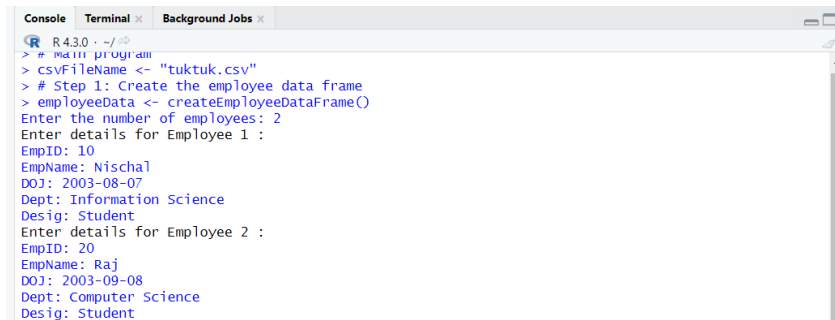
- 'write_csv': Used to write the employee data frame to a CSV file.

- 'read_csv': Used to read the contents of the CSV file.

- 'print': Used to display the contents of the data frame.

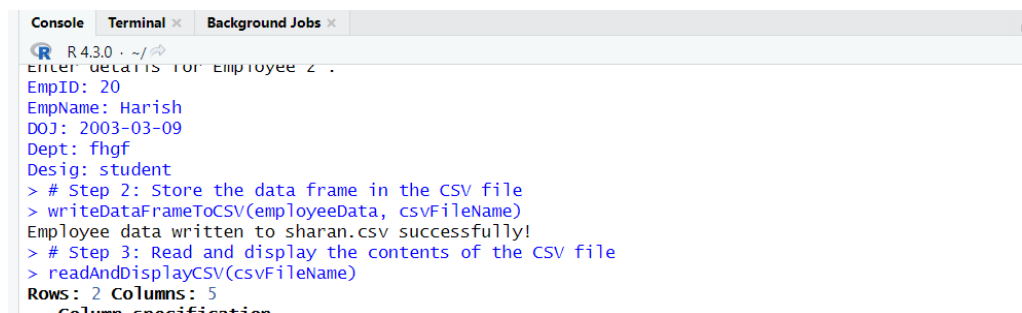
OUTPUT OF THE PROGRAM

- Upon execution, the program prompts the user to enter the number of employees and employee details for each employee.



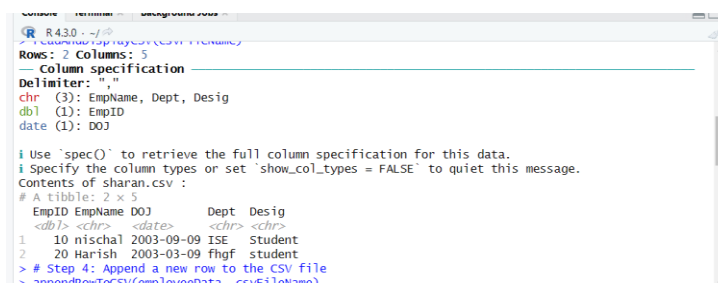
```
R 4.3.0 ~ / > # main program
> csvFileName <- "tuktuk.csv"
> # Step 1: Create the employee data frame
> employeeData <- createEmployeeDataFrame()
Enter the number of employees: 2
Enter details for Employee 1 :
EmpID: 10
EmpName: Nischal
DOJ: 2003-08-07
Dept: Information Science
Desig: Student
Enter details for Employee 2 :
EmpID: 20
EmpName: Raj
DOJ: 2003-09-08
Dept: Computer Science
Desig: Student
```

- It writes the employee data to a CSV file and displays a success message.



```
R 4.3.0 ~ / > # Step 2: Store the data frame in the CSV file
> writeDataFrameToCSV(employeeData, csvFileName)
Employee data written to sharan.csv successfully!
> # Step 3: Read and display the contents of the CSV file
> readAndDisplayCSV(csvFileName)
Rows: 2 Columns: 5
Column specification
-----
Delimiter: ,
chr (3): EmpName, Dept, Desig
dbl (1): EmpID
date (1): DOJ
```

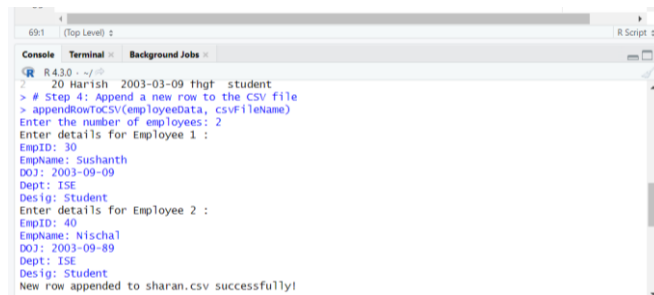
- It reads the contents of the CSV file and displays them.



```
R 4.3.0 ~ / > # Step 4: Append a new row to the CSV file
> appendRowToCSV(employeeData, csvFileName)
Rows: 2 Columns: 5
Column specification
-----
Delimiter: ,
chr (3): EmpName, Dept, Desig
dbl (1): EmpID
date (1): DOJ

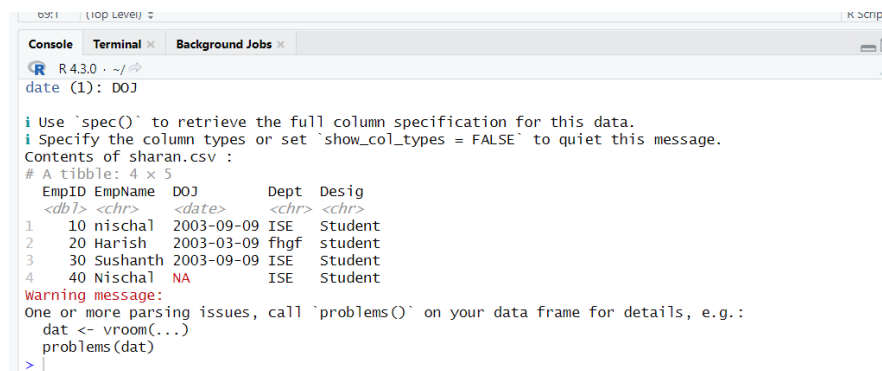
i Use 'spec()' to retrieve the full column specification for this data.
i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
Contents of sharan.csv :
# A tibble: 2 x 5
  EmpID EmpName DOJ      Dept Desig
  <dbl> <chr> <date> <chr> <chr>
1    10 nischal 2003-09-09 ISE Student
2    20 Harish 2003-03-09 fhgf student
```

- It prompts the user to enter details for a new employee and appends the new row to the CSV file.



```
R 4.3.0 ~/> 20 Harish 2003-03-09 fhgf student
> # Step 4: Append a new row to the CSV file
> appendRowToCSV(employeeData, csvFileName)
Enter the number of employees: 2
Enter details for Employee 1 :
EmpID: 30
EmpName: Sushanth
DOJ: 2003-09-09
Dept: ISE
Desig: Student
Enter details for Employee 2 :
EmpID: 40
EmpName: Nischal
DOJ: 2003-09-09
Dept: ISE
Desig: Student
New row appended to sharan.csv successfully!
```

After each operation, appropriate success messages or contents of the data frame are displayed.



```
R 4.3.0 ~/> date (1): DOJ
i Use 'spec()' to retrieve the full column specification for this data.
i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
Contents of sharan.csv :
# A tibble: 4 x 5
  EmpID EmpName DOJ      Dept Desig
  <dbl> <chr>   <date>   <chr> <chr>
1     10 nischal 2003-09-09 ISE    Student
2     20 Harish 2003-03-09 fhgf    student
3     30 Sushanth 2003-09-09 ISE    Student
4     40 Nischal NA       ISE    Student
Warning message:
One or more parsing issues, call 'problems()' on your data frame for details, e.g.:
  dat <- vroom(...)
  problems(dat)
> |
```