

EXPERIMENT - 10

Aim:- Make use of One-way Anova and Two-way Anova techniques in R to explore the sources of variability among different groups.

Introduction:- The aim of this experiment is to utilize One-way ANOVA and Two-way ANOVA techniques in R to explore the sources of variability among different groups. The goal is to understand how these analysis of variance methods can be applied to assess the impact of categorical variables on the variability in a dataset.

Software Required:-

1. R Statistical Software
2. R Studio

Relevance of the Experiment:- Analysis of variance is a powerful statistical technique used to analyze the differences among group means in a sample. This experiment is relevant for individuals involved in experimental design, research and data analysis, as it provides hands-on experience in identifying and understanding the sources of variance variability among different groups.

Description:- The experiment involves applying One-way ANOVA and Two-way ANOVA techniques in R to explore the

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Aim:- Make use of One-way ANOVA and Two-way ANOVA techniques in R to explore the sources of variability among different groups.

```
1 # Load required libraries
2 library(dplyr)
3 library(ggplot2)
4
5 # Load the iris dataset
6 data("iris")
7
8 # Function to perform sampling and estimate mean using a given sampling method
9 perform_sampling <- function(dataset, sample_size, sampling_method) {
10   if (sampling_method == "simple_random") {
11     sampled_data <- dataset[sample(nrow(dataset), sample_size, replace = FALSE), ]
12   } else if (sampling_method == "stratified") {
13     sampled_data <- dataset %>%
14       group_by(Species) %>%
15       sample_n(sample_size, replace = FALSE)
16   } else if (sampling_method == "systematic") {
17     n <- nrow(dataset)
18     k <- sample_size
19     start <- sample(1:k, 1)
20     indices <- seq(start, n, by = k)
21     sampled_data <- dataset[indices, ]
22   } else {
23     stop("Invalid sampling method.")
24   }
25   return(mean(sampled_data$Sepal.Length))
26 }
27
28 # Define parameters
29 sample_size <- 30
30 num_iterations <- 100
31 sampling_methods <- c("simple_random", "stratified", "systematic")
32
33 # Perform sampling and estimate mean using each technique
34 sampling_results <- lapply(sampling_methods, function(method) {
35   replicate(num_iterations, perform_sampling(iris, sample_size, method))
36 })
37
38 # Combine results into a data frame
39 results_df <- data.frame(
```

impact of one or two categorical variables on the variability within a dataset. Participants will learn to formulate hypotheses, conduct ANOVA tests, and interpret the results.

Pseudocode / Algorithm :-

1. Identify the categorical variables of interest.
2. Formulate the null and alternative hypothesis for the ANOVA tests.
3. Conduct One-way ANOVA or Two-way ANOVA using the relevant R functions.
4. Perform post-hoc tests if necessary to identify specific group differences.
5. Interpret the results and draw conclusions based on the ANOVA output.

Learning Outcomes :-

1. Understanding the principles of One-way and Two-way ANOVA.
2. Proficiency in formulating null and alternative hypotheses for ANOVA tests.
3. Ability to conduct One-way & Two-way ANOVA in R.

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```

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35 sampling_results <- lapply(sampling_methods, function(method) {
36   replicate(num_iterations, perform_sampling(iris, sample_size, method))
37 })
38
39 # Combine results into a data frame
40 results_df <- data.frame(
41   sampling_method = rep(sampling_methods, each = num_iterations),
42   mean = unlist(sampling_results)
43 )
44
45 # Plot histograms
46 histogram <- ggplot(results_df, aes(x = mean, fill = sampling_method)) +
47   geom_histogram(position = "identity", alpha = 0.5, bins = 20) +
48   labs(title = "Distribution of Sample Means", x = "Mean Sepal Length", y = "Frequency") +
49   theme_minimal()
50
51 print(histogram)
52
53 # Calculate means
54 mean_results <- sapply(sampling_results, mean)
55 names(mean_results) <- sampling_methods
56
57 # Print results
58 print(mean_results)

```

```

> mean_results <- sapply(sampling_results, mean)
> names(mean_results) <- sampling_methods
>
> # Print results
> print(mean_results)
simple_random stratified systematic
5.832267      5.838022      5.848600
>

```


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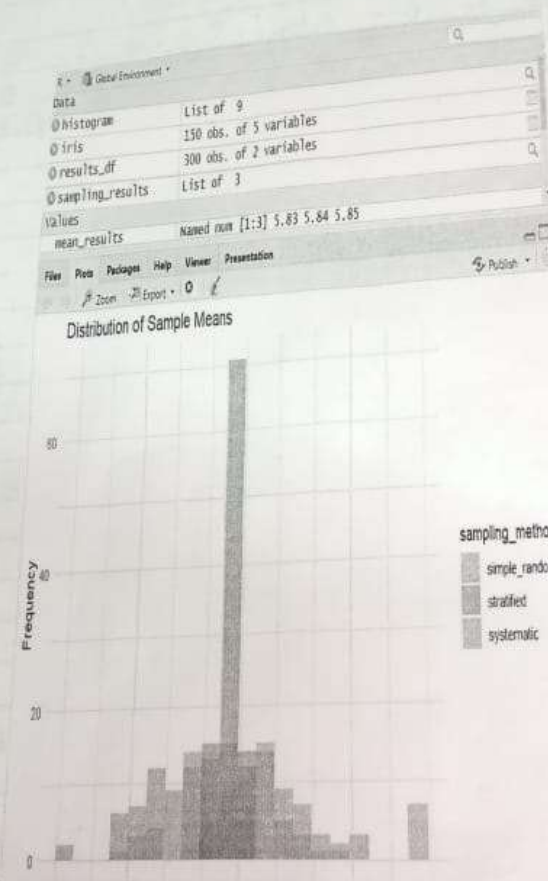
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4. Interpretation of ANOVA results, including F-statistics and p-values.
5. Practical experience in identifying sources of variability among different groups and assessing interaction effects between factors.

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SCREENSHOTS:

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