



Cluster Sampling – Population Mean Estimation

Subject:

Sampling Techniques

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1. Introduction

Cluster sampling is a probability-based survey technique where the population is divided into mutually exclusive groups called clusters. Instead of sampling individual units from the entire population, a subset of clusters is randomly selected, and all units within these clusters are included in the survey. This approach is especially useful when the population is large, geographically dispersed, or when surveying every unit individually is costly or time-consuming.

This report demonstrates a Shiny-based application for designing a cluster sampling survey, estimating population mean, and visualizing the selected clusters.

2. Objective

The main objectives of this cluster sampling exercise are:

- To determine the **number of clusters** to include in the survey based on user-defined parameters such as allowable error and confidence level.
- To calculate **expected cost and time** for surveying the selected clusters.
- To provide a **visual representation** of the survey design for better understanding and planning.
- To create an **interactive tool** that allows users to experiment with different survey parameters and immediately see the effect on cluster selection.

3. Methodology

1. Population and Clusters Setup:

- The total population is divided into a chosen number of clusters.
- Each cluster contains a number of units, and each unit has an associated observed value (simulated in the app for demonstration purposes).
- Cost and time per cluster are defined to allow estimation of survey resources.

2. Cluster Summary:

- For each cluster, the app computes:
 - Cluster size (number of units)
 - Mean of observed values
 - Variance of observed values
 - Cost and time

3. Cluster Selection:

- Based on the allowable error (sampling bias) and confidence level, the app determines how many clusters should be selected.
- Clusters are selected randomly to ensure unbiased sampling.

4. Cost & Time Calculation:

- Total expected cost and survey time are calculated by summing the cost and time across the selected clusters.

5. Visualization:

- The design of experiment is plotted using a scatter plot faceted by cluster.
- Units within selected clusters are highlighted in a distinctive color, while non-selected clusters are shown in a neutral color.
- This visual representation helps quickly understand which clusters are included and the overall survey coverage.

4. Features of the Shiny App



The screenshot shows the RStudio interface with the R code for the shiny app. The code defines the library imports, UI components, and server logic for generating a cluster sampling design.

```
1 library(shiny)
2 library(ggplot2)
3 library(dplyr)
4 library(DT)
5
6 ui <- fluidPage(
7   titlePanel("Cluster Sampling - Population Mean Estimation"),
8   sidebarLayout(
9     sidebarPanel(
10       numericInput("N", "Population Size (N):", value = 300, min = 50),
11       numericInput("C", "Total Number of Clusters:", value = 10, min = 2),
12       numericInput("B", "Allowable Error / Sampling Bias (B):", value = 4, min = 1),
13       selectInput("Z", "Confidence Level:",
14                  choices = c("90%" = 1.645,
15                             "95%" = 1.96,
16                             "99%" = 2.576),
17                  selected = 1.96),
18       numericInput("cost", "Cost per Cluster:", value = 100, min = 1),
19       numericInput("time", "Time per Cluster (hours):", value = 2, min = 0.1),
20       actionButton("calc", "Generate Cluster Design")
21     ),
22   )
23 )
```

```
29 mainPanel(
30   tabsetPanel(
31     tabPanel("Cluster Information",
32       DTOutput("cluster_table")
33     ),
34     tabPanel("Sampling Summary",
35       tableOutput("summary")
36     ),
37     tabPanel("Design of Experiment",
38       plotOutput("cluster_plot", height = "450px")
39     )
40   )
41 )
42 )
43 )
44 )
45 )
46 )
47
48 server <- function(input, output) {
49
50   observeEvent(input$calc, {
51     set.seed(123)
52
53     # POPULATION
54     units_per_cluster <- ceiling(input$N / input$C)
55
56     population <- data.frame(
57       Unit = 1:input$N,
58       ... = ...
59
60     y = rnorm(input$N, mean = 60, sd = 15),
61     Cluster = rep(paste0("C", 1:input$C), each = units_per_cluster)[1:input$N],
62     Cost = rep(input$cost, input$N),
63     Time = rep(input$time, input$N)
64   )
65
66   # Cluster-level summaries
67   cluster_info <- population %>%
68     group_by(Cluster) %>%
69     summarise(
70       Cluster_Size = n(),
71       Mean = mean(y),
72       Variance = var(y),
73       Cost = first(Cost),
74       Time = first(Time),
75       .groups = "drop"
76     )
77
78   M <- nrow(cluster_info)           # total clusters
79   S2 <- var(cluster_info$Mean)      # between-cluster variance
80   Z <- as.numeric(input$Z)
81   B <- input$B
82
83   # REQUIRED NUMBER OF CLUSTERS
84   m <- ceiling((Z^2 * S2) / (B^2))
85   m <- min(m, M)
86
87   # Select clusters (SRS of clusters)
88   selected_clusters <- sample(cluster_info$Cluster, m)
```

```
89     population$Selected <- ifelse(
90         population$Cluster %in% selected_clusters,
91         "Selected", "Not Selected"
92     )
93
94     # COST & TIME CALCULATION
95     cost_time <- cluster_info %>%
96         filter(Cluster %in% selected_clusters) %>%
97         summarise(
98             Total_Clusters_Selected = m,
99             Expected_Cost = sum(Cost),
100            Expected_Time = sum(Time)
101        )
102
103     # OUTPUTS
104     output$cluster_table <- renderDT({
105         datatable(cluster_info, options = list(pageLength = 10))
106     })
107
108     output$summary <- renderTable({
109         data.frame(
110             Parameter = c("Total Clusters",
111                         "Clusters Selected",
112                         "Allowable Error (B)",
113                         "Confidence Level (Z)",
114                         "Expected Cost",
115                         "Expected Time"),
116             Value = c(M,
117                         m,
118                         B,
119                         z
120
121             )
122
123     output$cluster_plot <- renderPlot({
124         ggplot(population,
125             aes(x = Unit, y = y, color = Selected)) +
126             geom_point(size = 2) +
127             facet_wrap(~Cluster, scales = "free_x") +
128             scale_color_manual(
129                 values = c("Selected" = "#1F78B4",
130                           "Not Selected" = "#D9D9D9")
131             ) +
132             tabs(
133                 title = "Cluster Sampling Design",
134                 subtitle = paste("Selected Clusters:", paste(selected_clusters, collapse = ", ")),
135                 x = "Population Units",
136                 y = "Observation Value",
137                 color = "Cluster Status"
138             ) +
139             theme_minimal(base_size = 14) +
140             theme(
141                 plot.title = element_text(face = "bold", size = 16),
142                 plot.subtitle = element_text(size = 12),
143                 legend.position = "bottom",
144                 legend.title = element_text(face = "bold")
145             )
146         )
147     })
148
149
150 })
151
152 }
153
154 shinyApp(ui, server)
```

- **Interactive Inputs:** Allows users to define population size, number of clusters, allowable error, confidence level, cost, and time per cluster.
- **Cluster Information Tab:** Displays detailed summary for each cluster in an interactive table.
- **Sampling Summary Tab:** Provides overall information including total clusters, clusters selected, allowable error, confidence level, expected cost, and time.
- **Design of Experiment Plot:** Shows a clear visualization of selected vs. non-selected clusters, facilitating planning and communication.
- **Reactivity:** Changing any input updates all outputs immediately, allowing users to experiment and observe effects dynamically.

5. Output screenshots:

Screenshot of the "Cluster Sampling – Population Mean Estimation" application interface. The application is titled "Data Mining clustering-sampling - Microsoft Edge" and shows a table of cluster data.

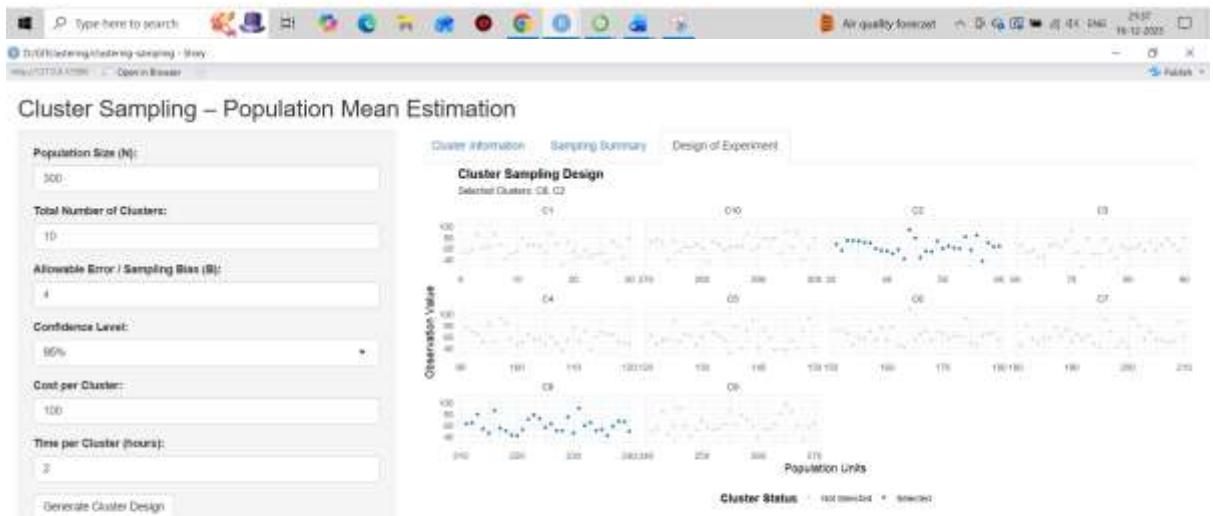
Cluster	Cluster_Size	Mean	Variance	Cost	Time
1 C1	30	60.2934438885238	218.5447821556587	100	2
2 C10	30	60.09033461762997	179.974323301260	100	2
3 C2	30	62.67507564500631	156.923881504166	100	2
4 C3	30	60.386302587541929	170.22177388657250	100	2
5 C4	30	68.991886652754431	188.0439243334386	100	2
6 C5	30	67.24629380157513	298.7758354135769	100	2
7 C6	30	62.305743641020616	189.40630233600459	100	2
8 C7	30	60.22089564044069	210.5868886470662	100	2
9 C8	30	58.65525338029037	171.6937729279165	100	2
10 C9	30	60.79147573181689	238.2794844591589	100	2

The application also includes tabs for "Cluster Information", "Sampling Summary", and "Design of Experiment". A search bar and navigation buttons (Previous, Next) are visible at the bottom.



Cluster Sampling – Population Mean Estimation

Population Size (N): 300	Cluster Information	Sampling Summary	Design of Experiment
Total Number of Clusters: 10	Parameter	Value	
Allowable Error / Sampling Bias (E): 4	Total Clusters	10.00	
Confidence Level: 95%	Clusters Selected	2.00	
Cost per Cluster: 100	Allowable Error (E)	4.00	
Time per Cluster (hours): 2	Confidence Level (Z)	1.96	
<input type="button" value="Generate Cluster Design"/>	Expected Cost	300.00	
	Expected Time	4.00	



6. Conclusion

- The Shiny app provides an **efficient and interactive tool** for designing cluster-based surveys.
- It allows users to **estimate required clusters, calculate expected resources, and visualize survey design** quickly.
- The approach ensures **efficient survey planning**, minimizing cost and time while maintaining statistical reliability.
- The visualization component makes it easier for survey designers and stakeholders to interpret and communicate the survey plan.

7.Professional github link

Including the git repository link : <https://github.com/ann-maria-anil/clustering-sampling-r-shiny>