

Universidad Nacional del Altiplano
Facultad de Ingeniería Estadística e Informática
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Trabajo Encargado -

Mi tareaa

Explicación de la Aplicación Shiny:

"Análisis Estadístico de Datos"

Este código es una aplicación interactiva web creada con el framework Shiny de R.

```
options(shiny.maxRequestSize = 100 * 1024^2) # 100MB max file size

library(shiny)
library(shinythemes)
library(DT)
library(readr)
library(readxl)
library(dplyr)
library(ggplot2)
library(plotly)
library(e1071)
library(nortest)
library(corrplot)
library(shinydashboard)
library(tidyr)

# Interfaz de Usuario
ui <- fluidPage(
  theme = shinytheme("flatly"),

  # Título principal
  titlePanel(
    div(
      h1("Análisis Estadístico de Datos",
        style = "color: #2c3e50; text-align: center; margin-bottom: 30px;"),
      h4("Aplicación para Datos Cualitativos y Cuantitativos",
        style = "color: #7f8c8d; text-align: center; font-weight: 300;")
    )
  ),

  # Layout principal
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sidebarLayout(
  # Panel lateral
  sidebarPanel(
    width = 3,

    # Sección de carga de datos
    div(class = "sidebar-section",
      tags$h4(icon("upload"), "Cargar Datos"),

      fileInput("file", "Seleccione archivo CSV o Excel",
        accept = c(".csv", ".xlsx", ".xls"),
        buttonLabel = "Examinar...",
        placeholder = "Ningún archivo seleccionado"),

      # Opciones para CSV
      conditionalPanel(
        condition = "input.file && input.file.name.endsWith('.csv')",
        checkboxInput("header", "¿Tiene encabezado?", TRUE),
        radioButtons("sep", "Separador:",
          choices = c("Coma" = ",",
            "Punto y coma" = ";",
            "Tabulador" = "\t"),
          selected = ","),
        radioButtons("quote", "Comillas:",
          choices = c("Ninguna" = "",
            "Simple" = "'",
            "Doble" = '"'),
          selected = '"')
      ),

      actionButton("load", "Cargar Datos",
        class = "btn-primary btn-block",
        style = "margin-top: 15px;"),
    ),

  # Información del dataset
  conditionalPanel(
    condition = "output.data_loaded",
    div(class = "sidebar-section",
      tags$h5(icon("info-circle"), "Información del Dataset"),
      verbatimTextOutput("data_info", placeholder = TRUE)
    )
  ),
),

```

```
# Panel principal
mainPanel(
  width = 9,

  tabsetPanel(
    id = "main_tabs",

    # Tab 1: Vista Previa de Datos
    tabPanel("Vista Previa",
      value = "preview",
      icon = icon("table"),

      fluidRow(
        column(12,
          div(class = "content-section",
            h3("Resumen General de los Datos"),
            verbatimTextOutput("summary")
          )
        )
      ),

      fluidRow(
        column(6,
          div(class = "content-section",
            h4("Tipos de Variables"),
            DT::dataTableOutput("data_types")
          )
        ),
        column(6,
          div(class = "content-section",
            h4("Valores Faltantes"),
            DT::dataTableOutput("missing_values")
          )
        )
      ),

      fluidRow(
        column(12,
          div(class = "content-section",
            h4("Vista de Datos (Primeras 50 filas)"),
            DT::dataTableOutput("preview")
          )
        )
      )
    )
  ),
```

```
# Tab 2: Estadística Descriptiva
tabPanel("Estadística Descriptiva",
  value = "descriptive",
  icon = icon("chart-bar"),

  sidebarLayout(
    sidebarPanel(
      width = 4,

      selectInput("var_desc", "Seleccione variable:",
        choices = NULL),

      checkboxGroupInput("stats", "Estadísticos a mostrar:",
        choices = c("Media" = "mean",
                    "Mediana" = "median",
                    "Moda" = "mode",
                    "Desviación estándar" = "sd",
                    "Varianza" = "var",
                    "Rango" = "range",
                    "Cuartiles" = "quartiles",
                    "Asimetría" = "skewness",
                    "Curtosis" = "kurtosis"),
        selected = c("mean", "median", "sd")),

      radioButtons("plot_type", "Tipo de gráfico:",
        choices = c("Histograma" = "hist",
                    "Boxplot" = "box",
                    "Densidad" = "density",
                    "Q-Q Plot" = "qq"),
        selected = "hist"),

      # Opciones adicionales para gráficos
      conditionalPanel(
        condition = "input.plot_type == 'hist'",
        sliderInput("bins", "Número de bins:",
          min = 5, max = 50, value = 20)
      ),

      actionButton("calc_desc", "Calcular",
        class = "btn-primary btn-block")
    ),

    mainPanel(
      width = 8,
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    fluidRow(
      column(12,
        div(class = "content-section",
          h4("Resultados Estadísticos"),
          verbatimTextOutput("desc_results")
        )
      )
    ),

    fluidRow(
      column(12,
        div(class = "content-section",
          h4("Visualización"),
          plotlyOutput("desc_plot", height = "400px")
        )
      )
    )
  ),

  # Tab 3: Análisis de Normalidad
  tabPanel("Normalidad",
    value = "normality",
    icon = icon("wave-square"),

    sidebarLayout(
      sidebarPanel(
        width = 4,

        selectInput("var_norm", "Variable para análisis:",
          choices = NULL),

        checkboxGroupInput("norm_tests", "Pruebas de normalidad:",
          choices = c("Shapiro-Wilk" = "shapiro",
            "Kolmogorov-Smirnov" = "ks",
            "Anderson-Darling" = "ad",
            "Lilliefors" = "lillie"),
          selected = c("shapiro", "ks")),

        numericInput("alpha", "Nivel de significancia:",
          value = 0.05, min = 0.01, max = 0.1, step = 0.01),

        actionButton("calc_norm", "Realizar Pruebas",

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        class = "btn-primary btn-block")
    ),

    mainPanel(
        width = 8,

        fluidRow(
            column(12,
                div(class = "content-section",
                    h4("Resultados de Pruebas de Normalidad"),
                    verbatimTextOutput("norm_results")
                )
            )
        ),

        fluidRow(
            column(6,
                div(class = "content-section",
                    h5("Q-Q Plot"),
                    plotlyOutput("qq_plot", height = "300px")
                )
            ),
            column(6,
                div(class = "content-section",
                    h5("Histograma con Curva Normal"),
                    plotlyOutput("norm_hist", height = "300px")
                )
            )
        )
    ),

    # Tab 4: Correlaciones (nueva funcionalidad)
    tabPanel("Correlaciones",
        value = "correlation",
        icon = icon("project-diagram"),

        fluidRow(
            column(4,
                div(class = "content-section",
                    h4("Configuración"),

                    selectInput("corr_vars", "Seleccionar variables numéricas:",
                        choices = NULL,

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        multiple = TRUE),

        radioButtons("corr_method", "Método de correlación:",
                      choices = c("Pearson" = "pearson",
                                   "Spearman" = "spearman",
                                   "Kendall" = "kendall"),
                      selected = "pearson"),

        actionButton("calc_corr", "Calcular Correlaciones",
                      class = "btn-primary btn-block")
      )
    ),

    column(8,
      div(class = "content-section",
        h4("Matriz de Correlación"),
        plotOutput("corr_plot", height = "400px"),

        h5("Tabla de Correlaciones"),
        DT::dataTableOutput("corr_table")
      )
    )
  )
)
),

# CSS personalizado
tags$head(
  tags$style(HTML("
    .content-section {
      background-color: #ffffff;
      border-radius: 8px;
      padding: 20px;
      margin-bottom: 20px;
      box-shadow: 0 2px 4px rgba(0,0,0,0.1);
    }

    .sidebar-section {
      background-color: #f8f9fa;
      border-radius: 8px;
      padding: 15px;
      margin-bottom: 20px;
      border-left: 4px solid #3498db;

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    }

    .btn-primary {
      background-color: #3498db;
      border-color: #3498db;
    }

    .btn-primary:hover {
      background-color: #2980b9;
      border-color: #2980b9;
    }
  ")))
)
)

# Servidor
server <- function(input, output, session) {

  # Valores reactivos
  rv <- reactiveValues(
    data = NULL,
    data_types = NULL,
    numeric_vars = NULL
  )

  # Indicador de datos cargados
  output$data_loaded <- reactive({
    return(!is.null(rv$data))
  })
  outputOptions(output, "data_loaded", suspendWhenHidden = FALSE)

  # Cargar datos
  observeEvent(input$load, {
    req(input$file)

    tryCatch({
      ext <- tools::file_ext(input$file$name)

      if (ext == "csv") {
        rv$data <- read.csv(input$file$datapath,
                           header = input$header,
                           sep = input$sep,
                           quote = input$quote)
      } else if (ext %in% c("xlsx", "xls")) {
        rv$data <- read_excel(input$file$datapath)
```



```
}

# Clasificar tipos de variables
rv$data_types <- sapply(rv$data, function(x) {
  if (is.numeric(x)) {
    unique_vals <- length(unique(x[!is.na(x)]))
    if (unique_vals <= 10) {
      return("Cuantitativa Discreta")
    } else {
      return("Cuantitativa Continua")
    }
  } else {
    unique_vals <- length(unique(x[!is.na(x)]))
    if (unique_vals <= 10) {
      return("Cualitativa Nominal")
    } else {
      return("Cualitativa Ordinal")
    }
  }
})

# Variables numéricas
rv$numeric_vars <- names(rv$data)[sapply(rv$data, is.numeric)]

# Actualizar opciones de selección
updateSelectInput(session, "var_desc", choices = names(rv$data))
updateSelectInput(session, "var_norm", choices = rv$numeric_vars)
updateSelectInput(session, "corr_vars", choices = rv$numeric_vars)

showNotification("Datos cargados correctamente", type = "success")

}, error = function(e) {
  showNotification(paste("Error al cargar datos:", e$message), type = "error")
})
})

# Información del dataset
output$data_info <- renderText({
  req(rv$data)
  paste(
    paste("Filas:", nrow(rv$data)),
    paste("Columnas:", ncol(rv$data)),
    paste("Variables numéricas:", length(rv$numeric_vars)),
    sep = "\n"
  )
})
```

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  })

  # Resumen de datos
  output$summary <- renderPrint({
    req(rv$data)
    summary(rv$data)
  })

  # Tipos de variables
  output$data_types <- renderDataTable({
    req(rv$data_types)
    data.frame(
      Variable = names(rv$data_types),
      Tipo = rv$data_types,
      stringsAsFactors = FALSE
    )
  }, options = list(pageLength = 10, searching = FALSE))

  # Valores faltantes
  output$missing_values <- renderDataTable({
    req(rv$data)
    missing_data <- rv$data %>%
      summarise_all(~sum(is.na(.))) %>%
      gather(key = "Variable", value = "Valores_Faltantes") %>%
      mutate(Porcentaje = round(Valores_Faltantes / nrow(rv$data) * 100, 2)) %>%
      arrange(desc(Valores_Faltantes))

    missing_data
  }, options = list(pageLength = 10, searching = FALSE))

  # Vista previa de datos
  output$preview <- renderDataTable({
    req(rv$data)
    head(rv$data, 50)
  }, options = list(scrollX = TRUE, pageLength = 15))

  # Estadística descriptiva
  output$desc_results <- renderPrint({
    req(rv$data, input$var_desc, input$calc_desc)

    var_data <- rv$data[[input$var_desc]]
    var_data <- var_data[!is.na(var_data)]

    results <- list()
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    if ("mean" %in% input$stats && is.numeric(var_data)) {
      results$Media <- round(mean(var_data), 4)
    }
    if ("median" %in% input$stats && is.numeric(var_data)) {
      results$Mediana <- round(median(var_data), 4)
    }
    if ("mode" %in% input$stats) {
      mode_val <- names(sort(table(var_data), decreasing = TRUE))[1]
      results$Moda <- mode_val
    }
    if ("sd" %in% input$stats && is.numeric(var_data)) {
      results$'Desviación Estándar' <- round(sd(var_data), 4)
    }
    if ("var" %in% input$stats && is.numeric(var_data)) {
      results$Varianza <- round(var(var_data), 4)
    }
    if ("range" %in% input$stats && is.numeric(var_data)) {
      results$Rango <- paste(round(range(var_data), 4), collapse = " - ")
    }
    if ("quartiles" %in% input$stats && is.numeric(var_data)) {
      q <- quantile(var_data, probs = c(0.25, 0.5, 0.75))
      results$'Q1, Q2, Q3' <- paste(round(q, 4), collapse = ", ")
    }
    if ("skewness" %in% input$stats && is.numeric(var_data)) {
      results$Asimetría <- round(e1071::skewness(var_data), 4)
    }
    if ("kurtosis" %in% input$stats && is.numeric(var_data)) {
      results$Curtosis <- round(e1071::kurtosis(var_data), 4)
    }

    results
  })

# Gráfico descriptivo
output$desc_plot <- renderPlotly({
  req(rv$data, input$var_desc, input$calc_desc)

  var_data <- rv$data[[input$var_desc]]
  var_name <- input$var_desc

  p <- switch(input$plot_type,
    "hist" = {
      if (is.numeric(var_data)) {
        ggplot(data.frame(x = var_data), aes(x = x)) +
          geom_histogram(bins = input$bins, fill = "#3498db", alpha = 0.7, c

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        labs(title = paste("Histograma de", var_name), x = var_name, y = " ")
        theme_minimal()
    } else {
        ggplot(data.frame(x = var_data), aes(x = x)) +
        geom_bar(fill = "#3498db", alpha = 0.7) +
        labs(title = paste("Gráfico de barras de", var_name), x = var_name) +
        theme_minimal() +
        theme(axis.text.x = element_text(angle = 45, hjust = 1))
    }
},
"box" = {
    if (is.numeric(var_data)) {
        ggplot(data.frame(x = var_data), aes(y = x)) +
        geom_boxplot(fill = "#e74c3c", alpha = 0.7) +
        labs(title = paste("Boxplot de", var_name), y = var_name) +
        theme_minimal()
    } else {
        ggplot(data.frame(x = "Variable", y = 1), aes(x = x, y = y)) +
        geom_text(aes(label = "No aplicable para variables categóricas"),
        theme_void()
    }
},
"density" = {
    if (is.numeric(var_data)) {
        ggplot(data.frame(x = var_data), aes(x = x)) +
        geom_density(fill = "#2ecc71", alpha = 0.7) +
        labs(title = paste("Densidad de", var_name), x = var_name, y = "Densidad") +
        theme_minimal()
    } else {
        ggplot(data.frame(x = "Variable", y = 1), aes(x = x, y = y)) +
        geom_text(aes(label = "No aplicable para variables categóricas"),
        theme_void()
    }
},
"qq" = {
    if (is.numeric(var_data)) {
        ggplot(data.frame(sample = var_data), aes(sample = sample)) +
        stat_qq() + stat_qq_line(color = "red") +
        labs(title = paste("Q-Q Plot de", var_name)) +
        theme_minimal()
    } else {
        ggplot(data.frame(x = "Variable", y = 1), aes(x = x, y = y)) +
        geom_text(aes(label = "No aplicable para variables categóricas"),
        theme_void()
    }
}

```

```
    }
  )

  ggplotly(p)
})

# Pruebas de normalidad
output$norm_results <- renderPrint({
  req(rv$data, input$var_norm, input$calc_norm)

  var_data <- na.omit(rv$data[[input$var_norm]])

  results <- list()

  for (test in input$norm_tests) {
    test_result <- switch(test,
      "shapiro" = {
        if (length(var_data) <= 5000) {
          shapiro.test(var_data)
        } else {
          list(method = "Shapiro-Wilk",
               p.value = NA,
               statistic = NA,
               note = "Muestra muy grande (>5000). Use otras pruebas")
        }
      },
      "ks" = ks.test(var_data, "pnorm", mean(var_data), sd(var_data)),
      "ad" = nortest::ad.test(var_data),
      "lillie" = nortest::lillie.test(var_data)
    )

    results[[test]] <- test_result
  }

  # Interpretación
  cat("INTERPRETACIÓN DE RESULTADOS:\n")
  cat("H0: Los datos siguen una distribución normal\n")
  cat("H1: Los datos NO siguen una distribución normal\n")
  cat(paste("Nivel de significancia:", input$alpha, "\n\n"))

  for (i in seq_along(results)) {
    test_name <- names(results)[i]
    test_result <- results[[i]]

    cat(paste("===", toupper(test_name), "===\n"))
  }
}
```

```
    if (!is.na(test_result$p.value)) {
      cat(paste("Estadístico:", round(test_result$statistic, 6), "\n"))
      cat(paste("p-valor:", format(test_result$p.value, scientific = TRUE), "\n"))

      if (test_result$p.value < input$alpha) {
        cat("CONCLUSIÓN: Se RECHAZA H0. Los datos NO siguen distribución normal.\n")
      } else {
        cat("CONCLUSIÓN: NO se rechaza H0. Los datos podrían seguir distribución normal.\n")
      }
    } else if (!is.null(test_result$note)) {
      cat(test_result$note, "\n")
    }
    cat("\n")
  }

  results
})

# Q-Q Plot para normalidad
output$qq_plot <- renderPlotly({
  req(rv$data, input$var_norm, input$calc_norm)

  var_data <- na.omit(rv$data[[input$var_norm]])

  p <- ggplot(data.frame(sample = var_data), aes(sample = sample)) +
    stat_qq(color = "#3498db") +
    stat_qq_line(color = "red", linewidth = 1) +
    labs(title = "Q-Q Plot Normal",
         x = "Cuantiles teóricos",
         y = "Cuantiles observados") +
    theme_minimal()

  ggplotly(p)
})

# Histograma con curva normal
output$norm_hist <- renderPlotly({
  req(rv$data, input$var_norm, input$calc_norm)

  var_data <- na.omit(rv$data[[input$var_norm]])

  p <- ggplot(data.frame(x = var_data), aes(x = x)) +
    geom_histogram(aes(y = after_stat(density)), bins = 30,
                   fill = "#3498db", alpha = 0.7, color = "white") +
    stat_function(fun = dnorm,
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```
      args = list(mean = mean(var_data), sd = sd(var_data)),
      color = "red", linewidth = 1) +
  labs(title = "Histograma vs Distribución Normal",
       x = input$var_norm, y = "Densidad") +
  theme_minimal()

  ggplotly(p)
})

# Análisis de correlación
output$corr_plot <- renderPlot({
  req(rv$data, input$corr_vars, input$calc_corr)
  req(length(input$corr_vars) >= 2)

  corr_data <- rv$data[input$corr_vars]
  corr_matrix <- cor(corr_data, use = "complete.obs", method = input$corr_method)

  corrploth(corr_matrix, method = "color", type = "upper",
            order = "hclust", tl.cex = 0.8, tl.col = "black",
            addCoef.col = "black", number.cex = 0.7)
})

# Tabla de correlaciones
output$corr_table <- renderDataTable({
  req(rv$data, input$corr_vars, input$calc_corr)
  req(length(input$corr_vars) >= 2)

  corr_data <- rv$data[input$corr_vars]
  corr_matrix <- cor(corr_data, use = "complete.obs", method = input$corr_method)

  # Convertir matriz a formato largo
  corr_df <- expand.grid(Var1 = rownames(corr_matrix), Var2 = colnames(corr_matrix))
  corr_df$Correlacion <- as.vector(corr_matrix)
  corr_df <- corr_df[corr_df$Var1 != corr_df$Var2, ]
  corr_df$Correlacion <- round(corr_df$Correlacion, 4)

  corr_df
}, options = list(pageLength = 10))
}

# Ejecutar aplicación
shinyApp(ui = ui, server = server)
```

