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vectorA <- c(1, 2, 3, 4, 5) matrix_result <- matrix(0, nrow = 5, ncol = 5)
for (i in 1:5) { for (j in 1:5) { matrix_result[i, j] <- abs(vectorA[i] - vectorA[j]) } }
print(matrix_result)
for (i in 1:5) { output <- " " for (j in 1:i) { output <- paste0(output, "*") } print(output) }
fib_sequence <- function() { start <- as.integer(readline(prompt = "Enter starting number:"))
if (start <= 0) { print("Please enter a positive number") return() }
a <- 0 b <- 1 count <- 0 result <- c()
repeat { if (count >= start) { if (b > 500) { break } result <- c(result, b) }
temp <- a + b
a <- b
b <- temp
count <- count + 1
} print(result) }
household_data <- read.csv("import_march.csv") head(household_data, 6)
male_subset <- household_data[household_data$Gender == "Male", ] female_subset <- household_data[household_data$Gender == "Female", ]
print(paste("Number of male observations:", nrow(male_subset))) print(paste("Number of female observations:", nrow(female_subset)))
gender_counts <- table(household_data$Gender)
barplot(gender_counts, main = "Number of Males and Females in Household", xlab = "Gender", ylab = "Count", col = c("pink", "lightblue"), legend.text = c("Female", "Male"), beside = TRUE)
expenses <- c(60, 10, 5, 25) expense_labels <- c("Food", "Electricity", "Savings", "Miscellaneous") percentages <- paste(expense_labels, expenses, "%")
pie(expenses, labels = percentages, main = "Dela Cruz Family Monthly Expenses", col = c("skyblue", "yellow", "green", "orange"))
data(iris) str(iris)
iris_means <- data.frame( Sepal.Length = mean(iris$Sepal.Length), Sepal.Width = mean(iris$Sepal.Width), Petal.Length = mean(iris$Petal.Length), Petal.Width = mean(iris$Petal.Width) ) print(iris_means)
species_counts <- table(iris$Species) pie(species_counts, main = "Iris Species Distribution", col = c("red", "blue", "green"), labels = paste(names(species_counts), species_counts))
setosa <- iris[iris$Species == "setosa", ] versicolor <- iris[iris$Species == "versicolor", ] virginica <- iris[iris$Species == "virginica", ]
print("Last 6 rows of Setosa:") print(tail(setosa, 6))
print("Last 6 rows of Versicolor:") print(tail(versicolor, 6))
print("Last 6 rows of Virginica:") print(tail(virginica, 6))
irisSpecies <- as.factor(iris$Species)
plot(iris$Sepal.Length, iris$Sepal.Width, main = "Iris Dataset", sub = "Sepal width and length", xlab = "Sepal Length", ylab = "Sepal Width", pch = c(16, 17, 18)[as.numeric(iris$Species)], col = c("red", "blue", "green")[as.numeric(iris$Species)])
legend("topright", legend = levels(iris$Species), pch = c(16, 17, 18), col = c("red", "blue", "green"))

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library(readxl) library(dplyr)
alexa <- read_xlsx("alexa-file.xlsx")

alexavariation <- gsub("BlackDot", "BlackDot", alexavariation) alexavariation <- gsub("BlackPlus", "BlackPlus", alex
alexavariation <- gsub("BlackShow", "BlackShow", alexavariation) alexavariation <- gsub("BlackSpot", "BlackSpot", a
alexavariation <- gsub("WhiteDot", "WhiteDot", alexavariation) alexavariation <- gsub("WhitePlus", "WhitePlus", ale
alexavariation <- gsub("WhiteShow", "WhiteShow", alexavariation) alexavariation <- gsub("WhiteSpot", "WhiteSpot",

head(alexa, 10)

variations <- alexa %>% count(variation) print(variations)

save(variations, file = "variations.RData")

barplot(variationsn, names.arg = variationsvariation, main = "Total Number of Alexa Variations", xlab =
"Variation", ylab = "Count", col = rainbow(nrow(variations)), las = 2, cex.names = 0.7)

black_vars <- variations[grepl("Black", variationsvariation),] white_vars <- variations[grepl("White", variationsvariation),
]

par(mfrow = c(1, 2))

barplot(black_vars, names.arg = black_varsvariation, main = "Black Variations", xlab = "Variation", ylab
= "Count", col = "black", las = 2, cex.names = 0.8)

barplot(white_vars, names.arg = white_varsvariation, main = "White Variations", xlab = "Variation", ylab
= "Count", col = "gray", las = 2, cex.names = 0.8)

par(mfrow = c(1, 1))

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