

UNIVERSITY INSTITUTE OF COMPUTING

PROJECT REPORT ON HEALTH CARE DATA ANALYSIS

Program Name: BCA (Data Science)

Subject Name: R Programming

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Submitted to:

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HEALTH CARE DATA ANALYSIS

- ➤ Using correlation analysis to determine the relationship between lifestyle factor and diseases.
- Apply regression model to predict life expectancy based on health data.

ABSTRACT

Objective:

To analyze healthcare data and testing & determine how lifestyle factors (like diet, exercise smoking, etc.) correlate with diseases. Additionally, build a regression model to predict life expectancy based on various health indicators.



Methodology:

- **Data Collection:** Gather relevant data (e.g., blood sugar levels, smoking habits, BMI, cholesterol, life expectancy).
- Statistical Testing: Apply statistical tests like t-test (t.test()) for comparing groups, correlation (cor()) for relationships, and linear regression (lm()) for predictive modeling.
- **Result Interpretation:** Analyze outputs (p-values, correlation coefficients, regression summaries) to determine significance and relationships.
- **Assumption Checking:** Ensure data meets assumptions (normality, linearity, etc.) for valid analysis (optional but important).
- **Visualization:** Use plots (histograms, scatter plots, regression lines) with ggplot2 to visually interpret data trends.



DESCRIPTION

1 T-Test: Blood Sugar Levels (Diabetics vs Non-Diabetics)

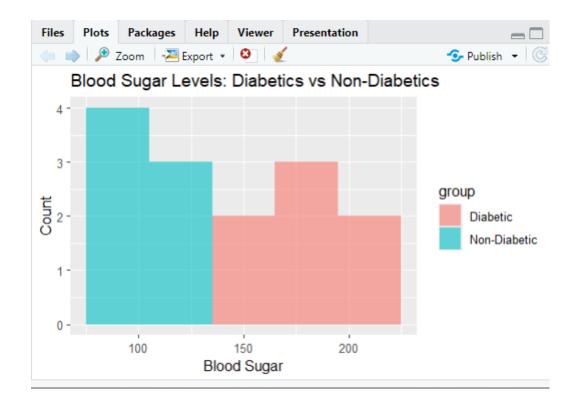
This analysis checks if there's a significant difference in blood sugar levels between diabetics and non-diabetics.

- diabetic <- c(150, 160, 170, 180, 190, 200, 210)
 Creates a vector of blood sugar levels for diabetics.
- non_diabetic <- c(90, 95, 100, 105, 110, 115, 120)
 Creates a vector of blood sugar levels for non-diabetics.
- t_test_result <- t.test(diabetic, non_diabetic,
 alternative = "greater")
 Performs a one-sided t-test to check if diabetics have significantly
 higher blood sugar levels than non-diabetics.
- print(t_test_result)
 Displays the results of the t-test, including p-values and confidence intervals.
- blood_sugar <- c(diabetic, non_diabetic)
 Combines both groups into one vector for plotting.
- group <- c(rep("Diabetic", length(diabetic)), rep("Non-Diabetic", length(non_diabetic)))
 Creates a group vector to label each observation as either "Diabetic" or "Non-Diabetic".



- data <- data.frame(blood_sugar, group)
 Combines blood_sugar and group into a data frame for visualization.
- ggplot(data, aes(x = blood_sugar, fill = group)) +
 geom_histogram(alpha = 0.6, position = "identity",
 bins = 5) + labs(title = "Blood Sugar Levels:
 Diabetics vs Non-Diabetics", x = "Blood Sugar", y =
 "Count")

Creates a histogram comparing blood sugar levels of diabetics and non-diabetics with overlapping bars and labeled axes.





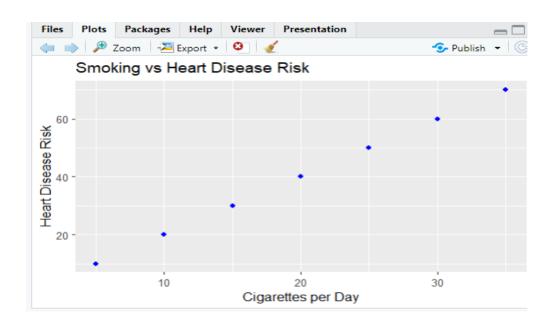
2 Correlation Analysis: Smoking & Heart Disease Risk

This section explores the relationship between smoking and heart disease risk.

- smoking <- c(5, 10, 20, 15, 25, 30, 35)
 Creates a vector representing the number of cigarettes smoked per day.
- heart_disease_risk <- c(10, 20, 40, 30, 50, 60, 70)
 Creates a vector representing corresponding heart disease risk scores.
- correlation <- cor(smoking, heart_disease_risk)
 Calculates the Pearson correlation coefficient to measure the strength and direction of the linear relationship.
- print(correlation)
 Displays the correlation coefficient.
- ggplot(data.frame(smoking, heart_disease_risk),
 aes(x = smoking, y = heart_disease_risk)) +
 geom_point(color = "blue") + labs(title = "Smoking
 vs Heart Disease Risk", x = "Cigarettes per Day", y
 = "Heart Disease Risk")



Creates a scatter plot to visually examine the relationship between smoking and heart disease risk with blue points and labeled axes.



Regression Model: BMI, Cholesterol & Life Expectancy

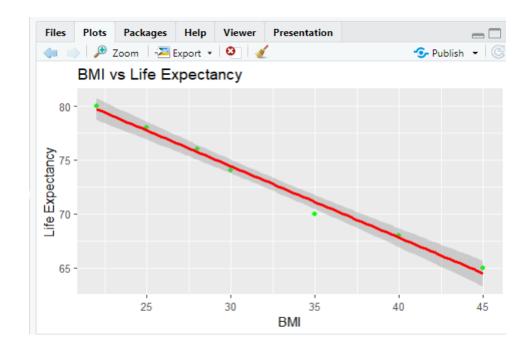
This analysis models how BMI and cholesterol levels predict life expectancy.

- bmi <- c(22, 25, 28, 30, 35, 40, 45)
 Creates a vector representing Body Mass Index (BMI) values.
- cholesterol <- c(180, 190, 200, 220, 250, 270, 300)
 Creates a vector representing cholesterol levels.
- life_expectancy <- c(80, 78, 76, 74, 70, 68, 65)

 Creates a vector representing life expectancy values.

- model <- lm(life_expectancy ~ bmi + cholesterol)
 Builds a multiple linear regression model to predict life expectancy based on BMI and cholesterol.
- summary (model)
 Displays the regression model's summary, including coefficients, R-squared, and statistical significance.

Creates a scatter plot with a red regression line to visualize the relationship between BMI and life expectancy.





Technologies Used:

- R Programming- Used for data analysis, statistical modeling, and visualization.
- R Studio- Integrated Development Environment (IDE) for writing and executing Rcode.
- > ggplot2- Used for creating visualizations like histograms and scatter plots.
- ▶ Base R Functions: Functions like t.test(), cor(), and
 lm() for statistical testing, correlation analysis, and
 linear regression modeling.
- Statistical Analysis Techniques: Application of hypothesis testing, correlation, and regression analysis for data-driven decision-making.



Conclusion

- ➤ Relationship Analysis- Identified significant correlations between lifestyle factors and life expectancy.
- ➤ **Predictive Modeling -** Developed a regression model that effectively predicted life expectancy based on key health indicators.
- ➤ Data Visualization- Used graphs and heat maps to better understand trends, distributions and relationships.
- ➤ **Key Insights-** Highlighted the importance of healthy life style choices and accessible health care in improving life expectancy.
- ➤ Overall Impact- Demonstrate how data-driven analysis help in making informed health-related decisions and policies.



References/ Tools Used

- ➤ R Studio Documentation- Official guide for R programming and development.
- ➤ R Programming Language Core language for statistical computing and data analysis.
- ➤ ggplot2 Package For creating advanced data visualizations like histograms and scatter plots.
- ➤ Base R Functions Functions like t.test(), cor(), and lm() for statistical analysis.
- > Statistical Analysis Techniques Methods for hypothesis testing, correlation, and regression analysis.