

TDS10 Final Project

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Abstract

IF you wish, you may add here a short abstract of 100 words max.

Introduction

“adsadsds”

Dataset Description

(Write about heart.csv here)

Multinomial Logistic Regression — Theory

Question(1.1):

We are using multinomial Logistic Regression because the response variable can take more than 2 categories. For these categories there is a separate set of coefficients and we choose one as the baseline. The coefficients describe how the predictors(age, sex, chol etc.) affect the probability of belonging to each outcome category.

In this regression the response variable Y can take K -number of different categories. We have to pick one of the categories to be the baseline - category 0, for every other category - $k = 1, 2, \dots, K-1$.

The model shows the probability of an observation belonging to category K using the multinomial logistic regression function:

$$P(Y = k \mid X) = \frac{\exp(\beta_{0k} + \beta_k^T X)}{1 + \sum_{j=1}^{K-1} \exp(\beta_{0j} + \beta_j^T X)}$$

The probability of the baseline category is:

$$P(Y = 0 | X) = \frac{1}{1 + \sum_{j=1}^{K-1} \exp(\beta_{0j} + \beta_j^T X)}$$

Data Preparation

There are several variables in the dataset containing missing values. In order to prepare the data for the multinomial logistic model, we observed how many missing values each variable had and saw that the variables `ca`, `thal`, and `slope` had extremely high numbers of missing values. Because these variables are categorical, and because imputing such a large amount of missing data would introduce strong bias, we decided to remove them from the dataset. For the remaining numeric variables with less missing values- (`trestbps`, `chol`, `thalach`, `oldpeak`) we applied median imputation.

We chose this method because replacing each missing value with the median of the corresponding variable is a robust measure that is not affected by extreme values.

For the categorical variables with very few missing values, we replaced missing entries with the most frequent category (the mode).

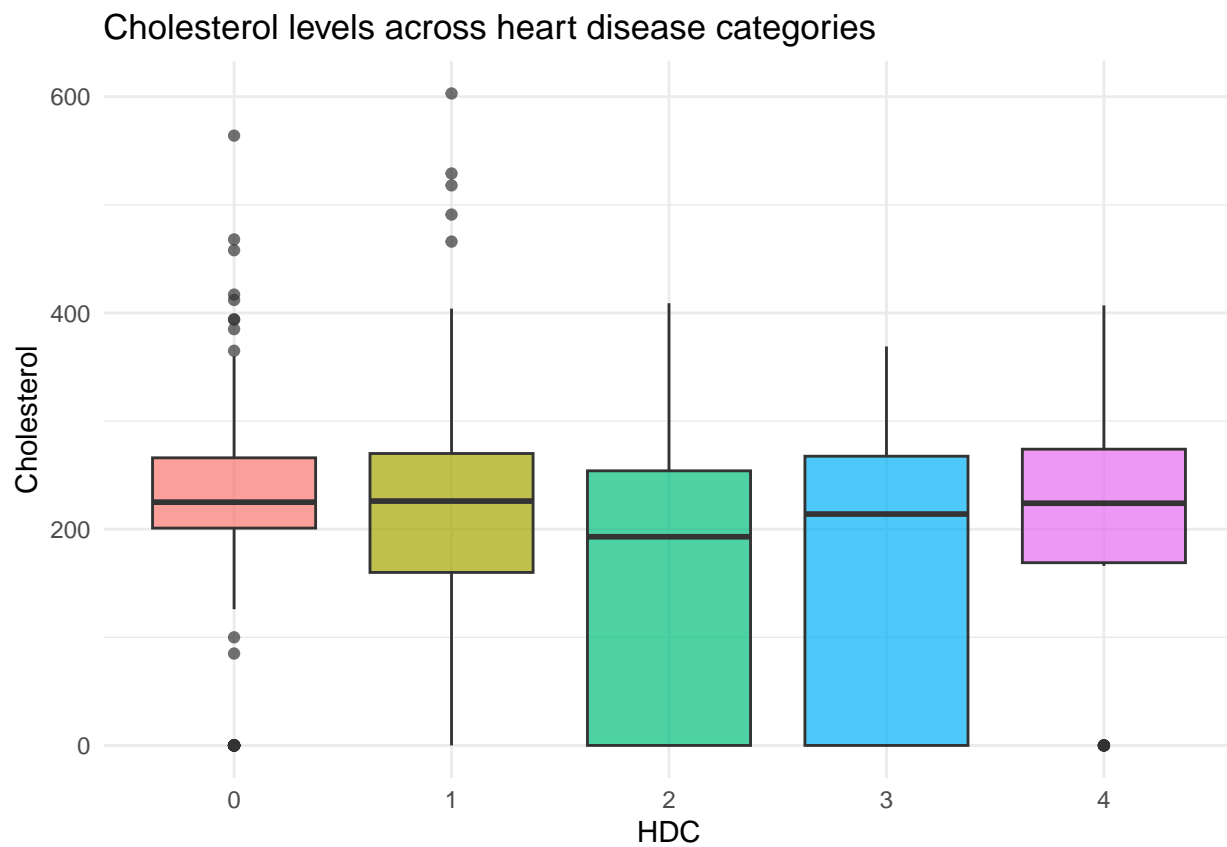
This approach ensures that there are no missing values before using the multinomial logistic regression model.

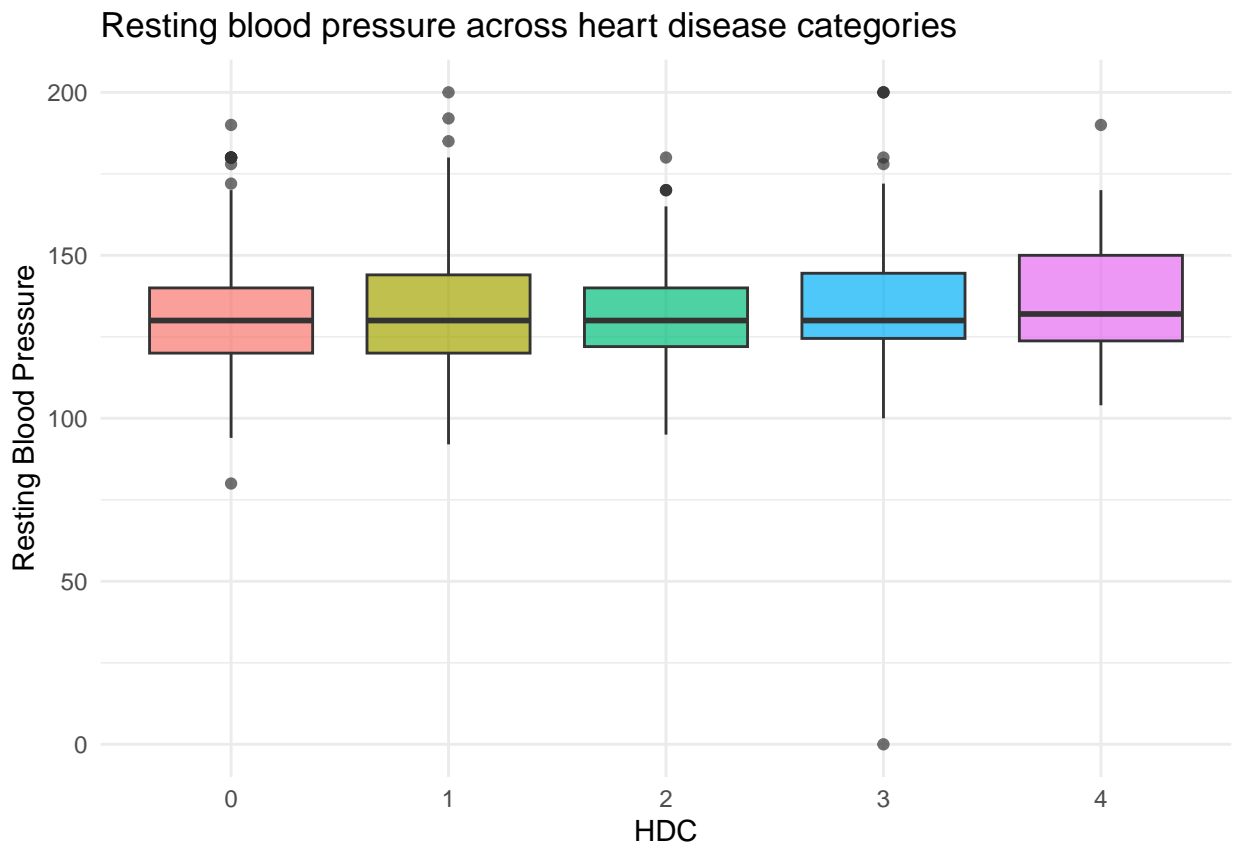
```
##      age      sex    place      cp trestbps      chol      fbs  restecg
##      0        0        0        0        0        0        0        0
##  thalach  exang  oldpeak      hdc
##      0        0        0        0
```

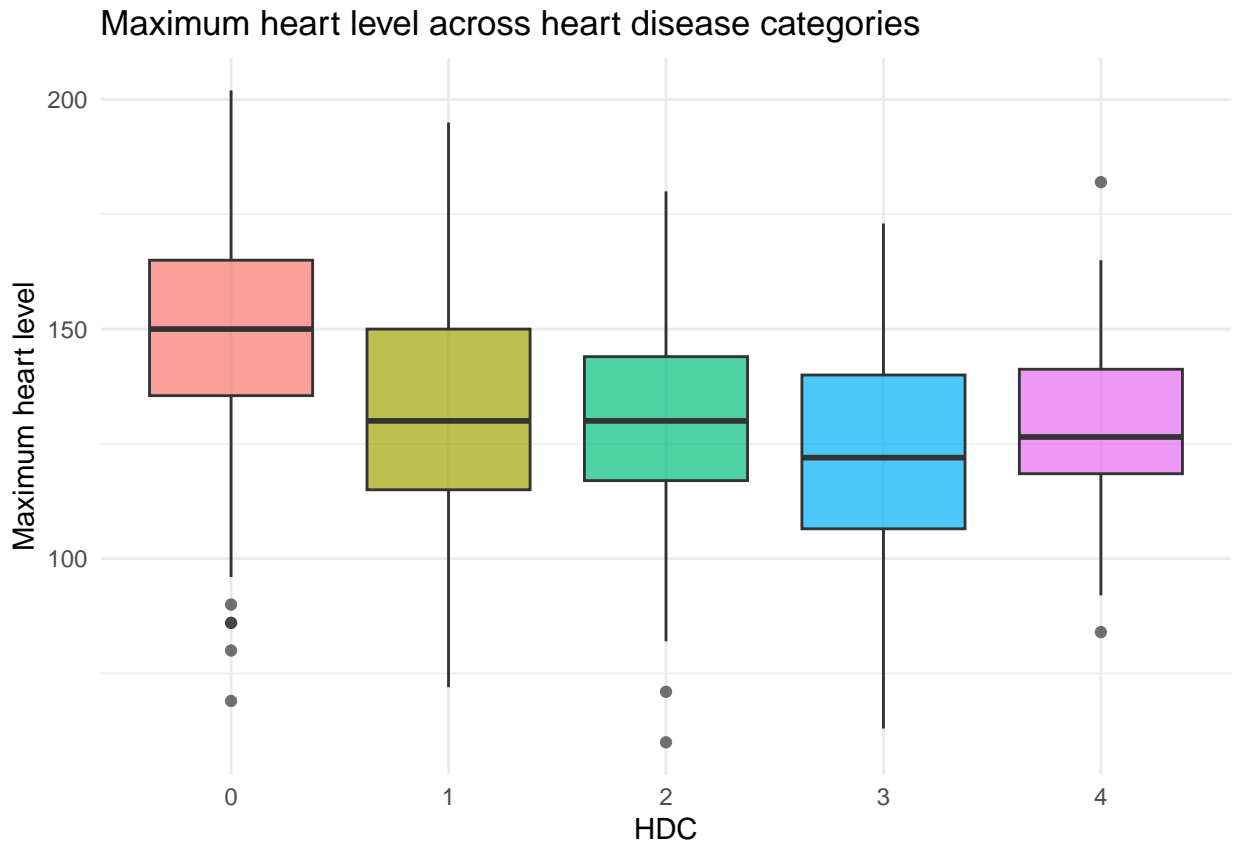
```
##  age sex    place      cp trestbps  chol fbs      restecg thalach
## 1  63  1 Cleveland  typical angina    145  233  1 lv hypertrophy    150
## 2  67  1 Cleveland  asymptomatic    160  286  0 lv hypertrophy    108
## 3  67  1 Cleveland  asymptomatic    120  229  0 lv hypertrophy    129
## 4  37  1 Cleveland  non-anginal    130  250  0      normal    187
## 5  41  0 Cleveland  atypical angina    130  204  0 lv hypertrophy    172
## 6  56  1 Cleveland  atypical angina    120  236  0      normal    178
##  exang oldpeak  hdc
## 1     0     2.3   0
## 2     1     1.5   2
## 3     1     2.6   1
## 4     0     3.5   0
## 5     0     1.4   0
## 6     0     0.8   0
```

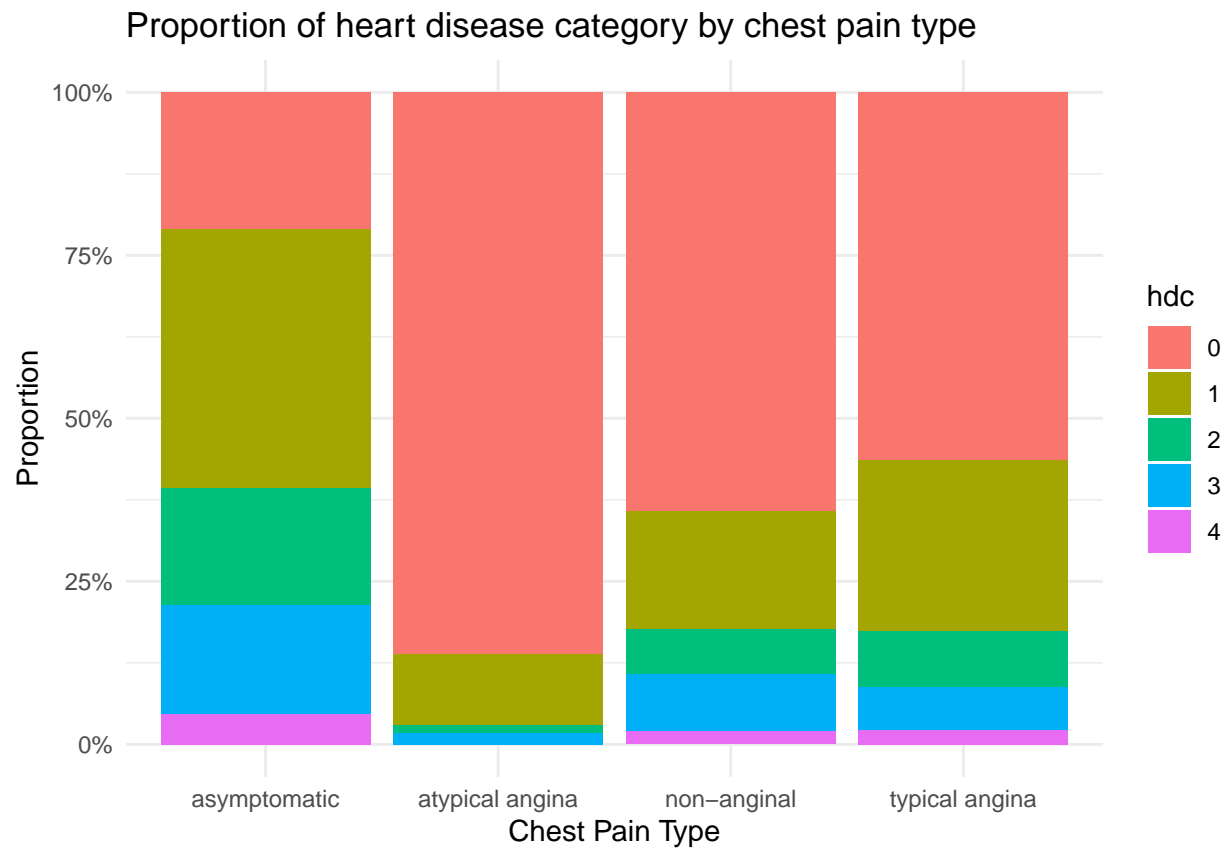
Exploratory Data Analysis

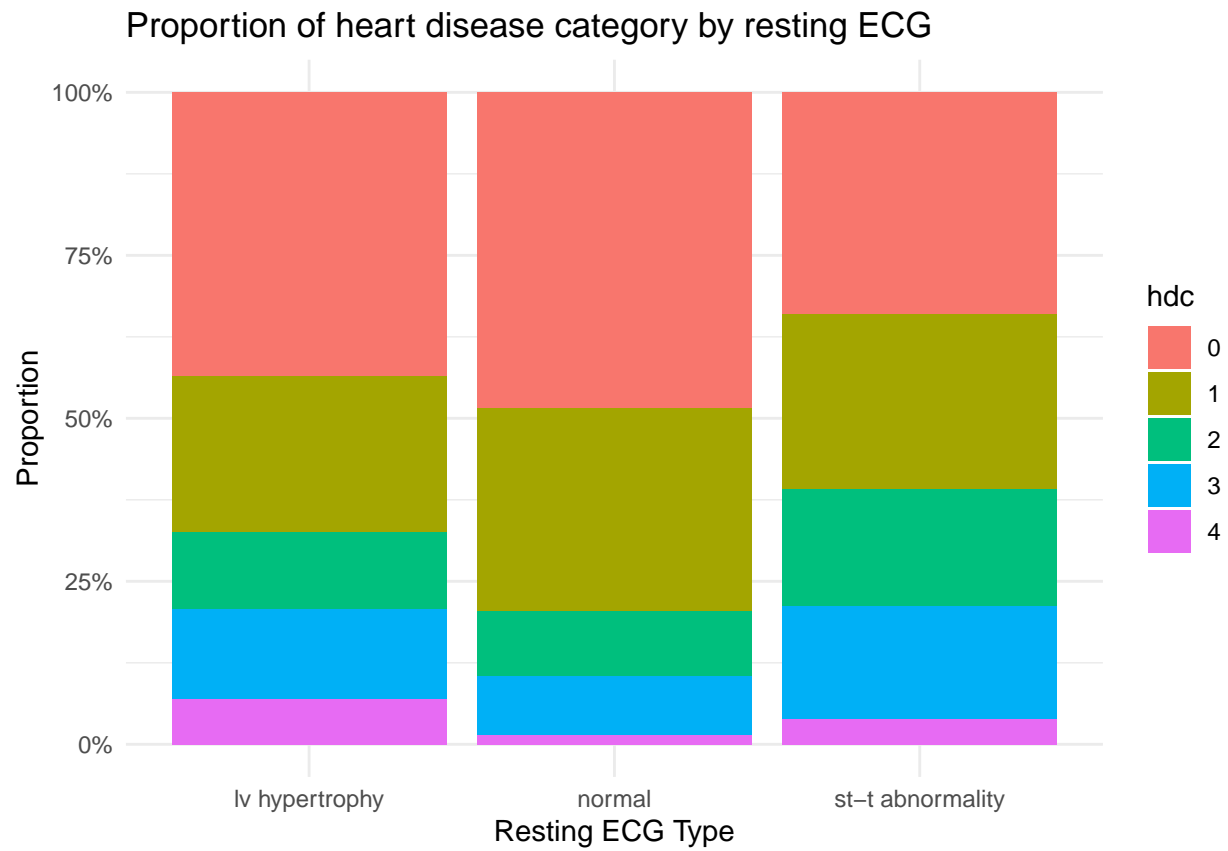
Warning: package 'ggplot2' was built under R version 4.5.2

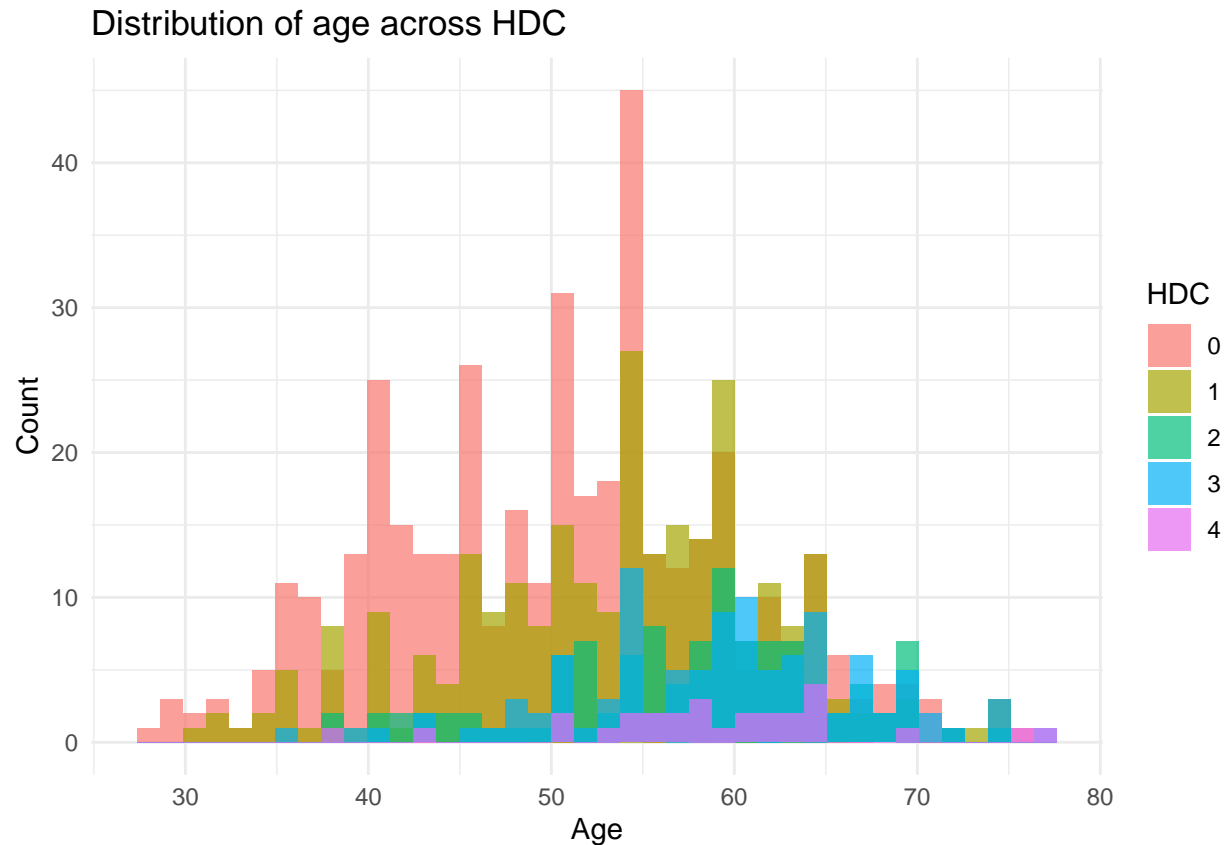












Multinomial Logistic Regression

(Fit model + interpretation)

Model Evaluation

(Cross-validation)

Model Improvement

(Stepwise model / alternative model)

Binary Logistic Regression

(Create hdc01 + logistic model)

Model Comparison

(Compare multinomial vs binary)

Conclusion

(Brief summary)