

Coronary Heart Disease Prediction

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Abstract

The early diagnostic of cardiovascular diseases can aid healthcare practitioners in making decisions on lifestyle changes in high risk patients with the purpose of reducing patient complications. This research intends to pinpoint the most relevant/risk factors of heart disease as well as predict the overall risk using logistic regression.

Data: The data set is called “heart” and was obtain from the following website:
<https://web.stanford.edu/~hastie/ElemStatLearn//datasets/SAheart.data>

Methods: logistic regression model was built to capture the risk of having a heart disease.

Stepwise model selection method was used to find the optimal model.

The prediction accuracy of the best model (classification error rate in predicting the class of new test data) was 72.17%.

Exploratory Data Analysis

The data set contains 462 observations and 10 variables. The response is “chd”, the presence (chd=1) or absence (chd=0) of coronary heart disease. The list of variables is shown in table 1.

In the data exploration analysis showed that the median Age of those who have a Coronary Heart Disease (chd=1) is higher than in those who do not have the condition (chd=0). Moreover, can be noted a more symmetrical distribution and higher Interquartile Range in the group with no coronary heart disease (chd=0) than on those with the condition (chd=1) Fig.1.

Table 1. Variables in the Heart DataSet

Name	Description	Type	Missing (%)
sbp	Systolic Blood Pressure	Integer	0
tobacco	Cumulative tobacco (kg)	Numeric	0
ldl	Low Density Lipoprotein Cholesterol	Numeric	0
adiposity	Measure in fat percentage	Numeric	0
famhist	Family history of heart disease (Present, Absent)	Factor	0
typea	Type-A behavior	Integer	0
obesity	Measure in bmi	Numeric	0
alcohol	Current Alcohol Consumption	Numeric	0
age	Age at onset	Integer	0
chd	The presence (chd=1) or absence (chd=0) of Coronary Heart Disease.	Factor	0

The corresponding mean and standard deviation of the quantitative continuous variables in the data table 2.

Table 2. Continuous Variables – Data Description

Variable	Mean	Standard Deviation
sbp	138.33	20.49
tobacco	3.64	4.59
ldl	4.74	2.07
adiposity	25.41	7.78
typea	53.10	9.82
obesity	26.04	4.21
alcohol	17.04	24.48
age	42.82	14.61

The distribution of alcohol between the two groups of individuals with Coronary Heart Disease (chd=1) and individuals with no Coronary Heart Disease (chd=0). looks almost identical (Fig 8). Similarly, no changes can be notice in the distribution of obesity between the two groups (Fig 7). The median consumption of Tobacco in the group with Coronary Heart Disease (chd=1) is slightly higher than in those with no Coronary Heart Disease (chd=0) (Fig. 2). Similarly, the median ldl, typea behavior, adiposity and sbp is slightly higher than those without the condition (chd=0) (Fig 3-6).

Figure 1. Distribution of Age by Coronary Heart Disease Status

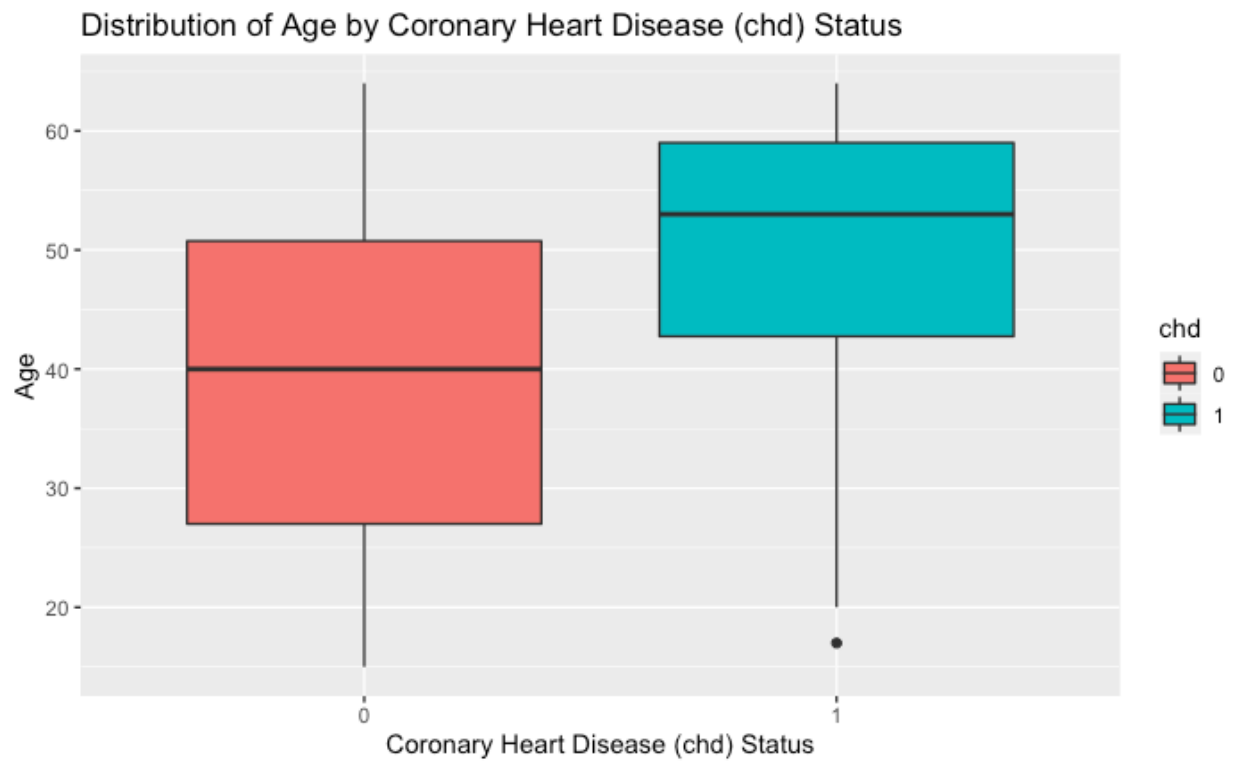


Figure 2. Tobacco by CHD Status

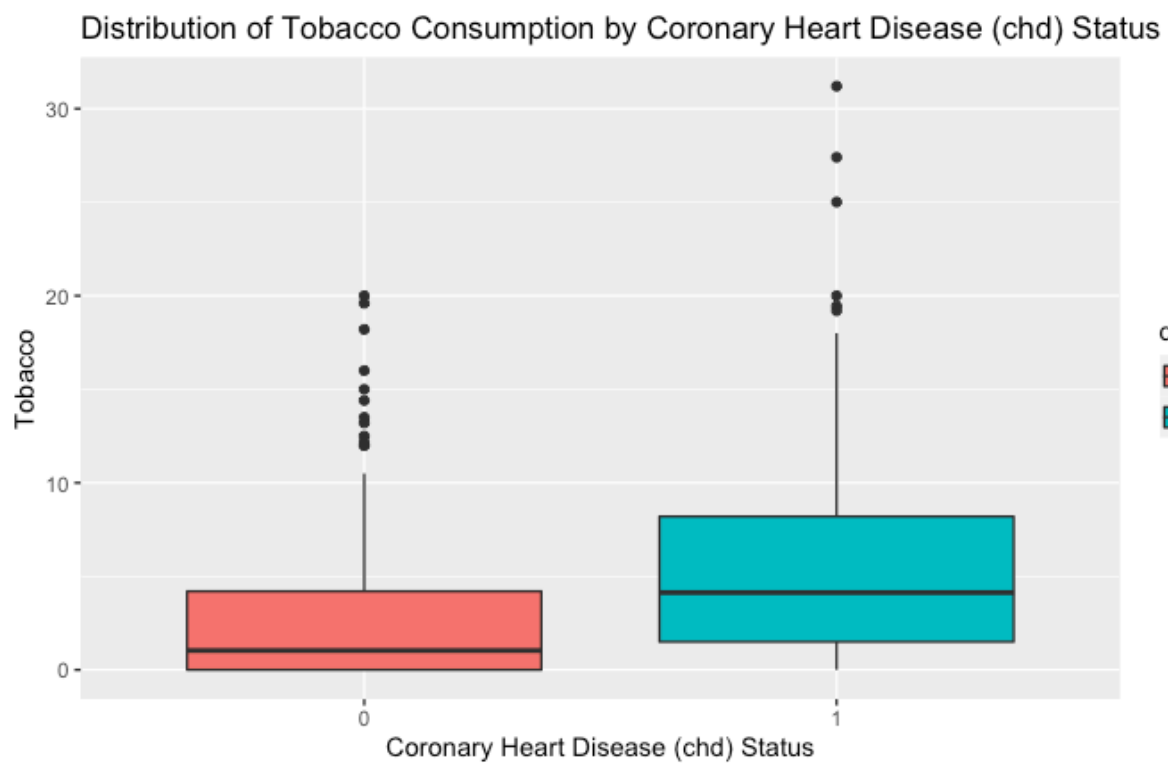


Figure 3.LDL Cholesterol by CHD Status

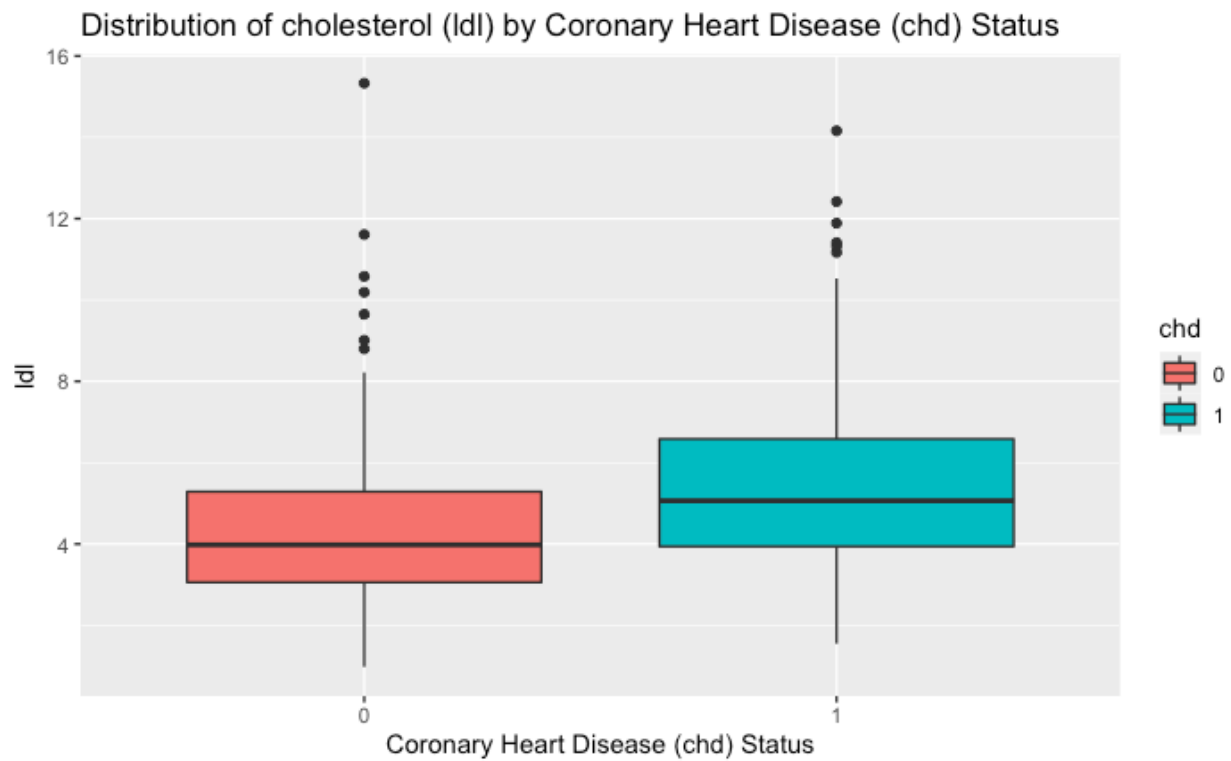


Figure 4.Adiposity by CHD Status

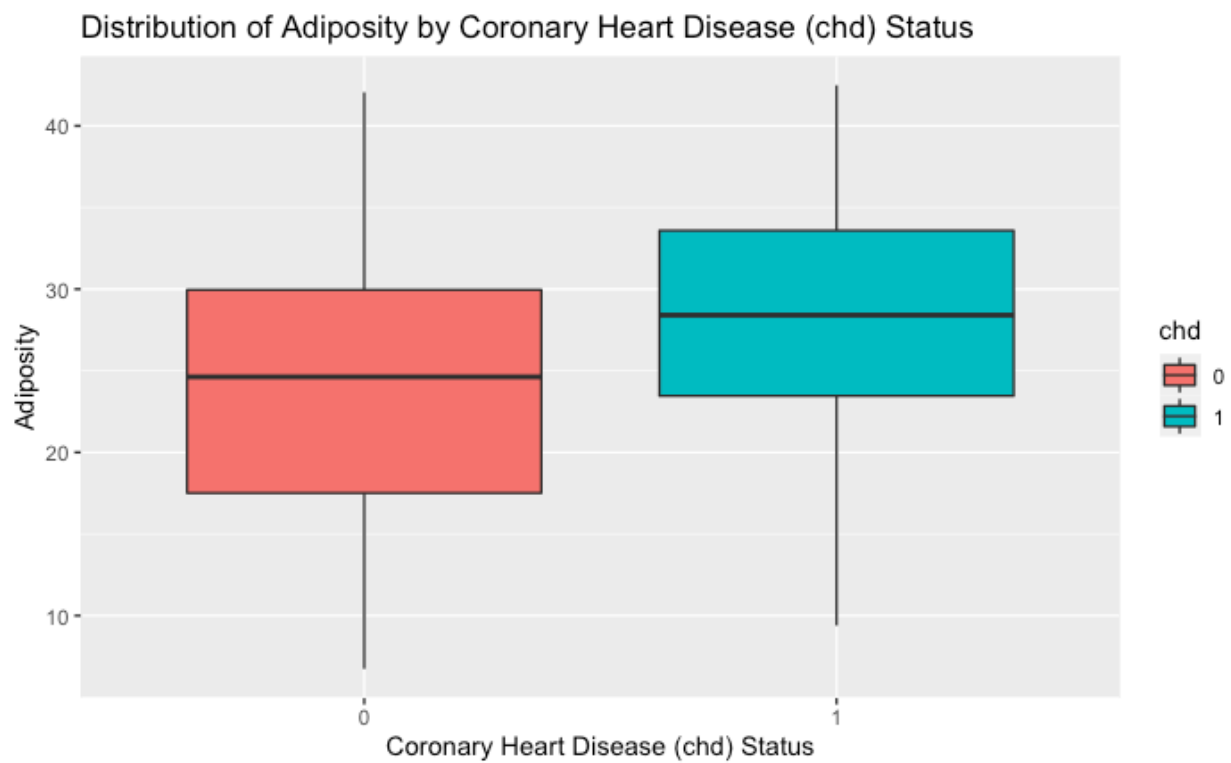


Figure 5. Type a Behavior by CHD Status

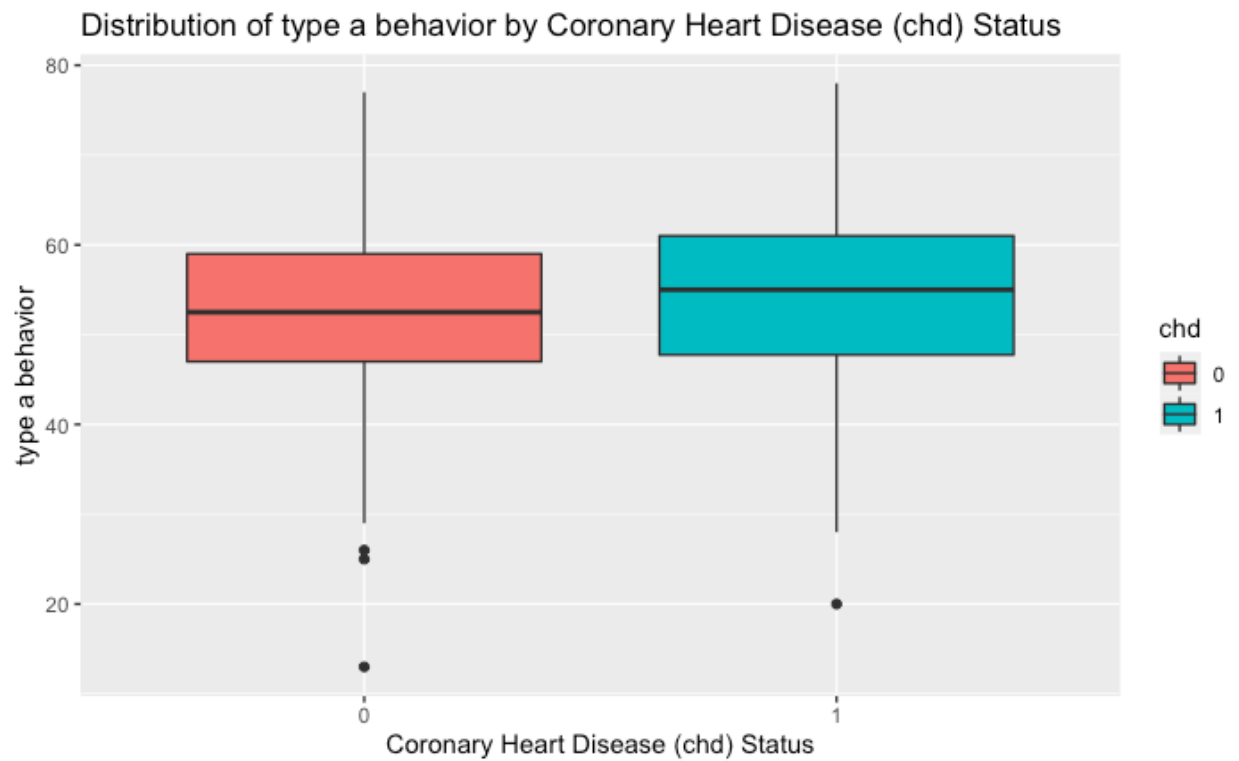


Figure 6. SBP by CHD Status

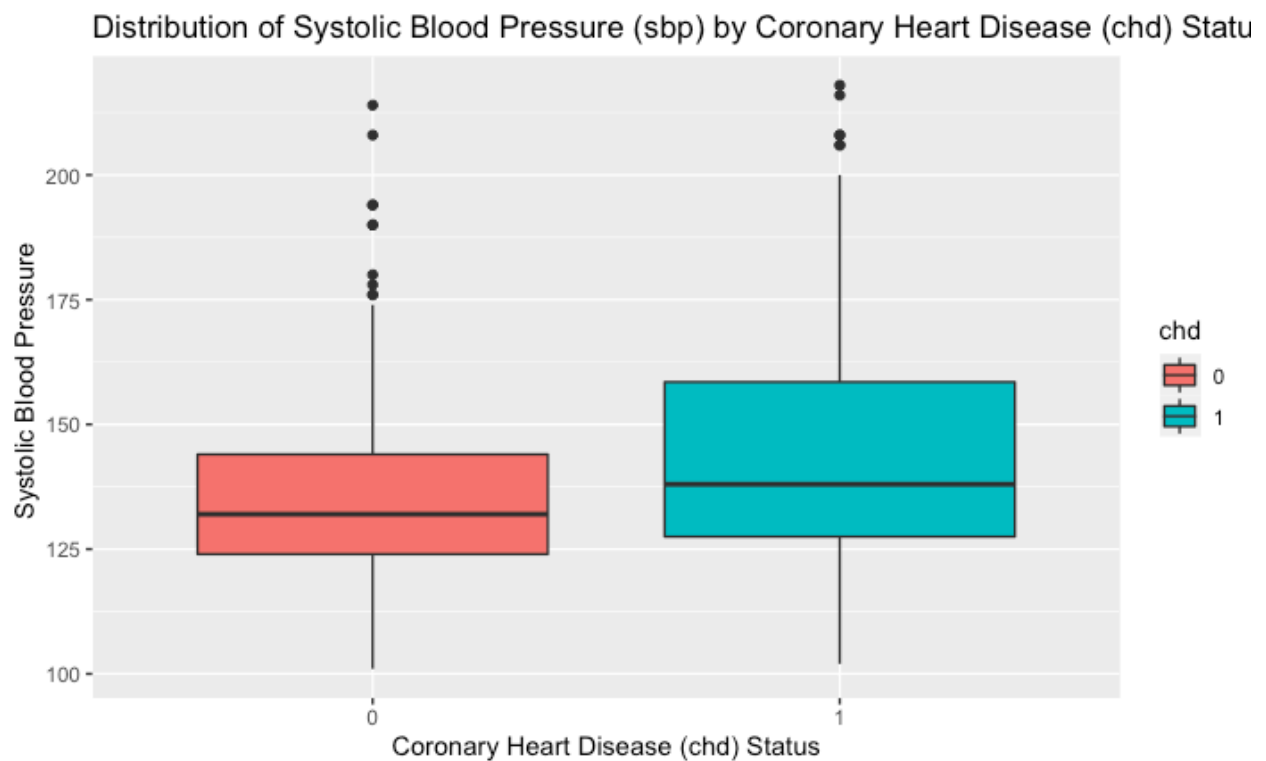


Figure 7 Obesity by CHD Status

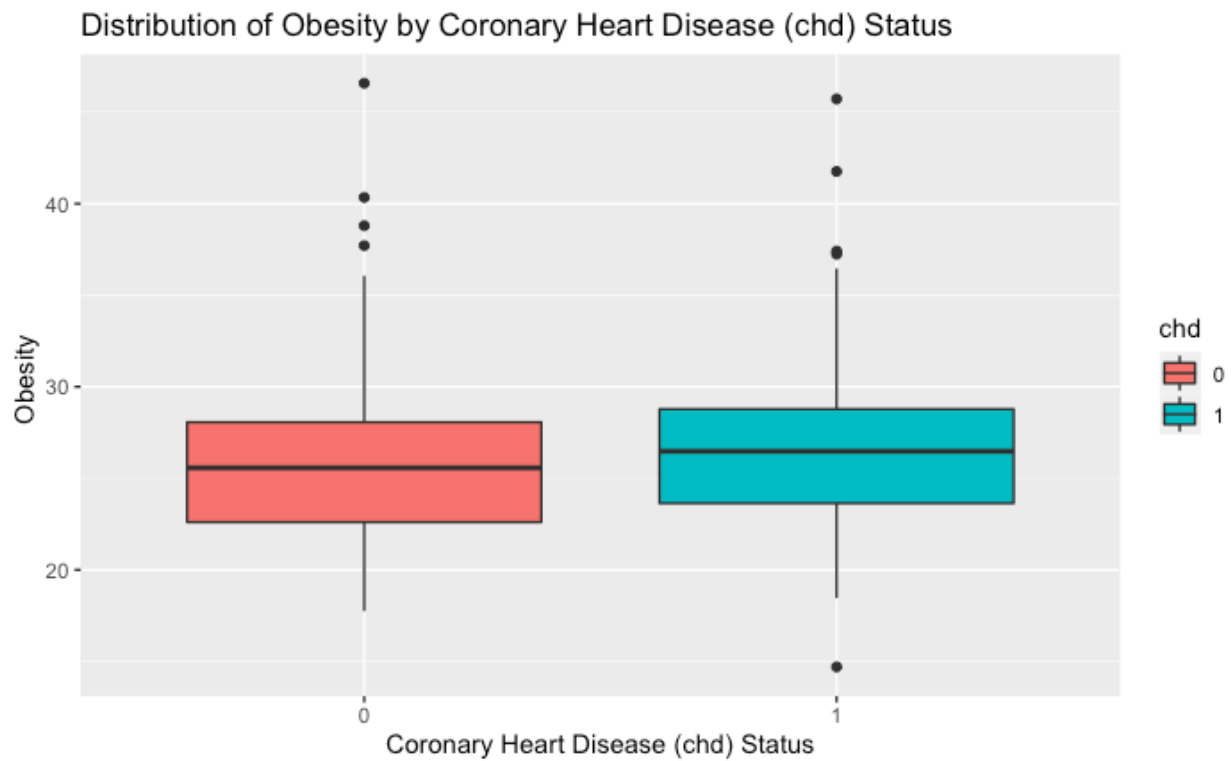
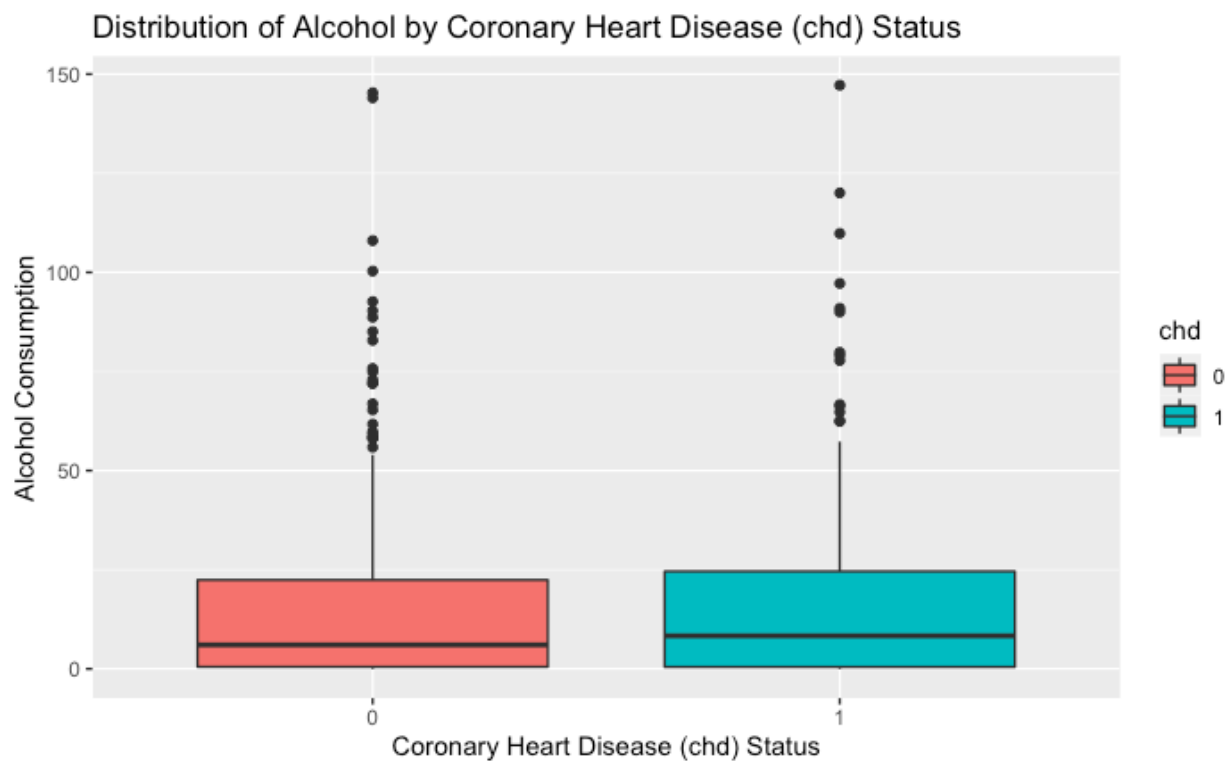


Figure 8. Alcohol by CHD Status

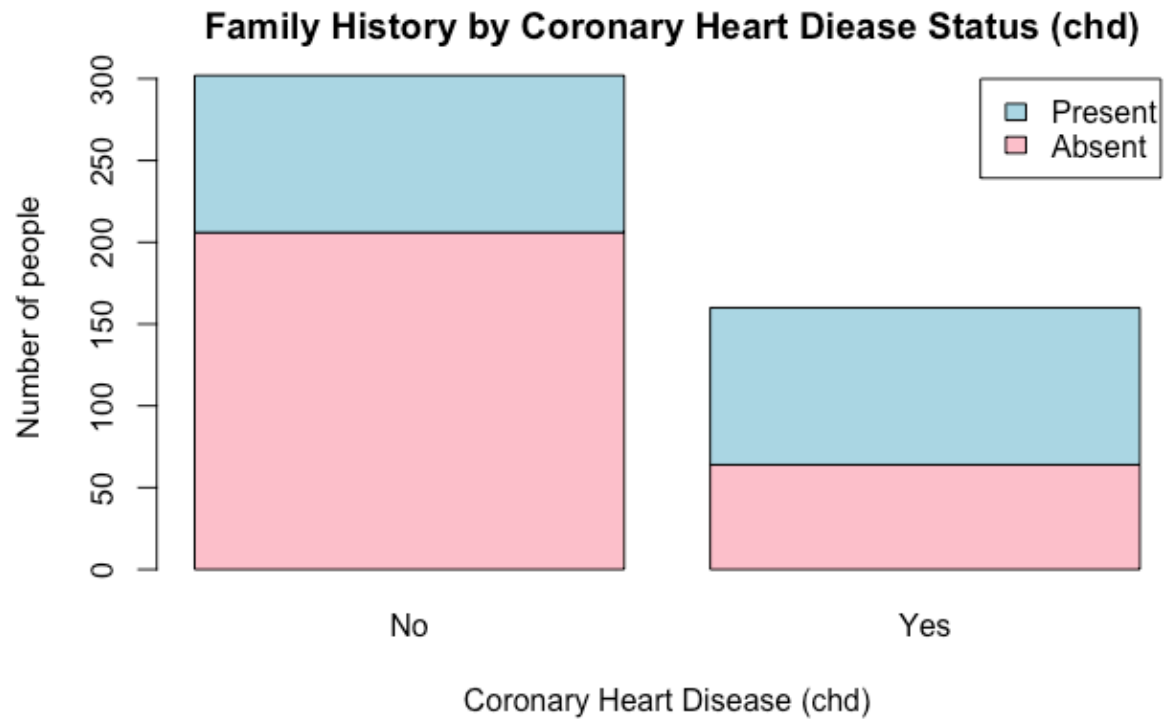


The exploration of the categorical variable Family History of heart disease is summarized in table 3. The proportion of a present family history on individuals with coronary heart disease is higher than in the individuals without the condition fig 9.

Table 3. Family History of Heart Disease vs Coronary Heart Disease Status

Fam History	Coronary Heart Disease (chd)		Totals
	No	Yes	
Absent	206	64	270
Present	96	96	192
Totals	302	160	462

Figure 9. Family History by Coronary Heart Disease Status (chd)



Setting up a training and testing data set

A training data set was created called "train" which contain 75% of the observation from the original data set. The rest of the data was allocated for a test data set "Test".

```
set.seed(3456)
trainIndex <- createDataPartition(heart$chd, p = .75, list = FALSE)
```

```
Train <- heart[ trainIndex,]
Test <- heart[ -trainIndex,]
```

Full logistic regression model

A full logistic regression model was built which includes all the predictors and fitted using the training data.

```
mylogitfull <- glm (chd ~., data = Train, family = "binomial")
summary(mylogitfull)
```

Call:
glm(formula = chd ~ ., family = "binomial", data = Train)

Deviance Residuals:

Min	1Q	Median	3Q	Max
-1.7928	-0.7847	-0.4213	0.8599	2.5008

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-5.148747	1.480688	-3.477	0.000507	***
sbp	0.003835	0.006449	0.595	0.552105	
tobacco	0.082248	0.030124	2.730	0.006326	**
ldl	0.204374	0.075954	2.691	0.007129	**
adiposity	0.031659	0.034743	0.911	0.362178	
famhistPresent	0.902090	0.269484	3.347	0.000816	***
typea	0.040143	0.014145	2.838	0.004542	**
obesity	-0.109172	0.054591	-2.000	0.045522	*
alcohol	-0.001097	0.005377	-0.204	0.838374	
age	0.046491	0.014065	3.306	0.000948	***

```
Null deviance: 447.51 on 346 degrees of freedom
Residual deviance: 346.37 on 337 degrees of freedom
AIC: 366.37
```

Stepwise variable selection

Select the most contributive variables:

```
#Stepwise model selection
```

```
library(MASS)
```

```
library(dplyr)
```

```
step.model<-mylogitfull %>% stepAIC(trace=FALSE)
```

```
coef(step.model)
```

(Intercept)	tobacco	ldl	famhistPresent	typea
-5.20391216	0.08171878	0.22508537	0.89113868	0.03885501
obesity	age			
-0.07082302	0.05464758			

```
summary(step.model)
```

Call:

```
glm(formula = chd ~ tobacco + ldl + famhist + typea + obesity +  
    age, family = "binomial", data = Train)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-1.7810	-0.7714	-0.4065	0.8623	2.5230

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-5.20391	1.24004	-4.197	2.71e-05	***
tobacco	0.08172	0.02943	2.777	0.005485	**
ldl	0.22509	0.07310	3.079	0.002075	**
famhistPresent	0.89114	0.26825	3.322	0.000893	***
typea	0.03886	0.01407	2.762	0.005744	**
obesity	-0.07082	0.03577	-1.980	0.047686	*
age	0.05465	0.01204	4.541	5.61e-06	***

```
Null deviance: 447.51 on 346 degrees of freedom
Residual deviance: 347.60 on 340 degrees of freedom
AIC: 361.6
```

After stepwise variable selection the reduced model named “step. model” removed the predictors adiposity and alcohol from the full model. Both models were used to predict the response on the test data set.

Full vs Reduced model

The performance of the full and the stepwise (reduced) logistic model were made. The best model is defined as the model that has the lowest classification error rate in predicting the class of new test data. The results are summarized in table 4.

```
#Prediction accuracy of the full logistic regression model:

# Make predictions
probabilities <- mylogitfull %>% predict(Test, type = "response")

predicted.classes <- ifelse(probabilities > 0.5, "1", "0")

# Prediction accuracy

observed.classes <- Test$chd
mean(predicted.classes == observed.classes)

[1] 0.6956522

#Prediction accuracy of the stepwise logistic regression model:

# Make predictions
probabilitiesstepmodel <- step.model %>% predict(Test, type = "response")

predicted.classessstepmodel <- ifelse(probabilitiesstepmodel > 0.5, "1", "0")

# Prediction accuracy

observed.classessstepmodel <- Test$chd
mean(predicted.classessstepmodel == observed.classessstepmodel)

[1] 0.7217391
```

Table 4. Comparison Full vs Reduced Model

	Full Model	Reduced Model
Prediction Accuracy	69.56%	72.17%
AIC	366.37	361.6

Results

The stepwise logistic regression model was chosen.

$$\text{logit}(p) = \log(p/(1-p)) = \beta_0 + \beta_1 \cdot \text{tobacco} + \beta_2 \cdot \text{ldl} + \beta_3 \cdot \text{famhistPresent} + \beta_4 \cdot \text{typea} + \beta_5 \cdot \text{obesity} + \beta_6 \cdot \text{age}$$

The result of the model chosen can be seen below in the table 5:

Table 5. Results from logistic regression

Term	β estimate	Standard error	P value
Intercept (β_0)	-5.20391	1.24004	<0.001
tobacco (β_1)	0.08172	0.02943	0.005485
ldl (β_2)	0.22509	0.07310	0.002075
famhistPresent (β_3)	0.89114	0.26825	<0.001
Type (β_4)	0.03886	0.01407	0.005744
obesity (β_5)	-0.07082	0.03577	0.047686
age (β_6)	0.05465	0.01204	<0.001

Individuals who do not have a family history of coronary heart disease have an 8.5 % increase in the odds of having a Coronary Heart Disease for each one -unit increase in cumulative tobacco (kg). Since $\exp(0.08172) = 1.08515$. Similarly, those individuals have an 25.24 % increase in the odds of having a Coronary Heart Disease for each one unit increase in ldl. Since, $\exp(0.22509) = 1.25243$. Similarly, Individuals who do not have a family history of coronary heart disease have an % 3.96 % increase in the odds of having a Coronary Heart Disease for each one -unit increase in typea behavior. Since $\exp(0.03886) = 1.03962$. Individuals who do not have a family history of coronary heart disease have an 5.6% increase in the odds of having a Coronary Heart Disease for each one -unit increase in age. Since $\exp(0.05465) = 1.05617$.

While holding the other predictors at a fixed value, the odds of having a Coronary Heart Disease is 143.79 % higher for individuals with a family history of Coronary Heart Disease than for individuals who do not have a family history. Since $\exp(0.89114) = 2.4379$.

Surprisingly, in this population individuals with no family history of heart disease the odds of having a Coronary Heart Disease decreases by 6.8 % for each one unit increase in obesity. Since $\exp(-0.07082) = 0.9312$.

Conclusion

In the analysis was found that the factors that increase the risk of having a Coronary Heart Disease in this population are a family history of Coronary Heart Disease, Cumulative tobacco (kg), Low Density Lipoprotein Cholesterol (ldl), type A Behavior and Age at onset.