

# Final\_programming

Biswajit Chowdhury

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## Description:

This is a primer for missing value imputation. Here, I used the ‘heart data’ from UCI depository. The dataset does not include missing values. Therefore, I introduced 20% missing value using introduced missing values of the total observation using the prodNA function from the missForest package. The prodNA function incorporates missing values completely at random. Therefore, the characteristics of missing values in the given dataset are missing completely at random (MCAR). The missing values were imputed by four different ways using Multiple imputation using chained equations (MICE).

Dataset 1: Imputed all variables using default MICE function. In this function, numerical variables were imputed by predictive mean matching (PMM) and categorical variables either by logreg, or polyreg. This dataset has been defined as ‘MICE’ in this analysis.

Dataset 2: Imputed all variables by PMM. This is a semi-parametric imputation approach, which draws imputed values from an observed empirical distribution. This has been defined as “PMM” in this article.

Dataset 3: Changed numerical variable with mean and kept categorical as dataset one and defined this as ‘Mean’.

Dataset 4: Introduced missing values only in the numerical variables (age, resting blood pressure, cholesterol level, max\_heart rate and oldpeak) and keep others unchanged (as train data). Then replaced the missing values only for five numerical variables with mean imputation and labeled it as ‘mean\_numeric’.

## Load library

## Reading data file

```
##   age sex cp trestbps chol fbs restecg thalach exang oldpeak slope ca thal
## 1  63   1  3     145  233    1      0    150     0    2.3    0  0    1
## 2  37   1  2     130  250    0      1    187     0    3.5    0  0    2
## 3  41   0  1     130  204    0      0    172     0    1.4    2  0    2
##   target
## 1      1
## 2      1
## 3      1

## 'data.frame': 303 obs. of 14 variables:
## $ age           : int  63 37 41 56 57 57 56 44 52 57 ...
## $ sex           : int  1 1 0 1 0 1 0 1 1 1 ...
## $ chest_pain    : int  3 2 1 1 0 0 1 1 2 2 ...
```

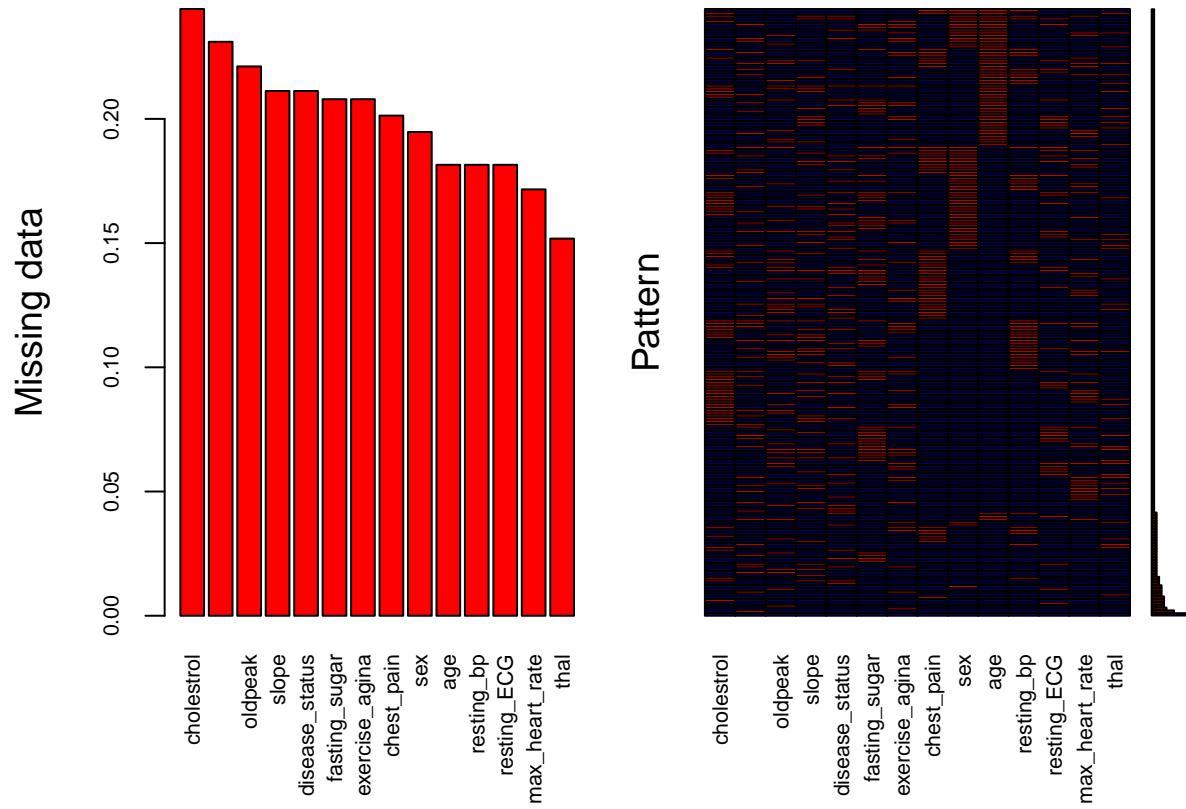
```

## $ resting_bp      : int  145 130 130 120 120 140 140 120 172 150 ...
## $ cholesterol    : int  233 250 204 236 354 192 294 263 199 168 ...
## $ fasting_sugar  : int  1 0 0 0 0 0 0 1 0 ...
## $ resting_ECG     : int  0 1 0 1 1 1 0 1 1 1 ...
## $ max_heart_rate : int  150 187 172 178 163 148 153 173 162 174 ...
## $ exercise_agina : int  0 0 0 0 1 0 0 0 0 0 ...
## $ oldpeak         : num  2.3 3.5 1.4 0.8 0.6 0.4 1.3 0 0.5 1.6 ...
## $ slope           : int  0 0 2 2 2 1 1 2 2 2 ...
## $ number_major_vessels: int  0 0 0 0 0 0 0 0 0 0 ...
## $ thal             : int  1 2 2 2 2 1 2 3 3 2 ...
## $ disease_status   : int  1 1 1 1 1 1 1 1 1 1 ...

```

## Preprocess the variables

### Create missing values in the data set



```

##
## Variables sorted by number of missings:
##          Variable      Count
##          cholesterol 0.2442244
##  number_major_vessels 0.2310231
##          oldpeak 0.2211221
##          slope 0.2112211

```

```

##      disease_status 0.2112211
##      fasting_sugar 0.2079208
##      exercise_agina 0.2079208
##      chest_pain 0.2013201
##      sex 0.1947195
##      age 0.1815182
##      resting_bp 0.1815182
##      resting_ECG 0.1815182
##      max_heart_rate 0.1716172
##      thal 0.1518152

```

Figure 1. Aggregation plot for missing data on the train data set. The missing values were introduced using the prodNA function from the missForest package. The visualization of the missing values patterns was generated using the VIM package in R. (A) represents the proportion of the missing and (B) distribution patterns in each variable. The red in graph representing missing values along with observed values (blue).

## 1. Multiple imputation

impute missing values using multivariate imputation by chained equations (MICE)

```

## Class: mids
## Number of multiple imputations: 5
## Imputation methods:
##      age          sex          chest_pain
##      "pmm"        "logreg"     "polyreg"
##      resting_bp   cholestrol   fasting_sugar
##      "pmm"        "pmm"       "logreg"
##      resting_ECG  max_heart_rate exercise_agina
##      "polyreg"    "pmm"       "logreg"
##      oldpeak      slope       number_major_vessels
##      "pmm"        "polyreg"    "polyreg"
##      thal         disease_status
##      "polyreg"    "logreg"
## PredictorMatrix:
##      age sex chest_pain resting_bp cholestrol fasting_sugar
## age  0   1      1           1      1           1
## sex  1   0      1           1      1           1
## chest_pain 1   1      0           1      1           1
## resting_bp  1   1      1           0      1           1
## cholestrol 1   1      1           1      0           1
## fasting_sugar 1   1      1           1      1           0
##      resting_ECG max_heart_rate exercise_agina oldpeak slope
## age            1              1              1      1      1
## sex            1              1              1      1      1
## chest_pain    1              1              1      1      1
## resting_bp    1              1              1      1      1
## cholestrol   1              1              1      1      1
## fasting_sugar 1              1              1      1      1
##      number_major_vessels thal disease_status
## age                  1   1           1
## sex                  1   1           1

```

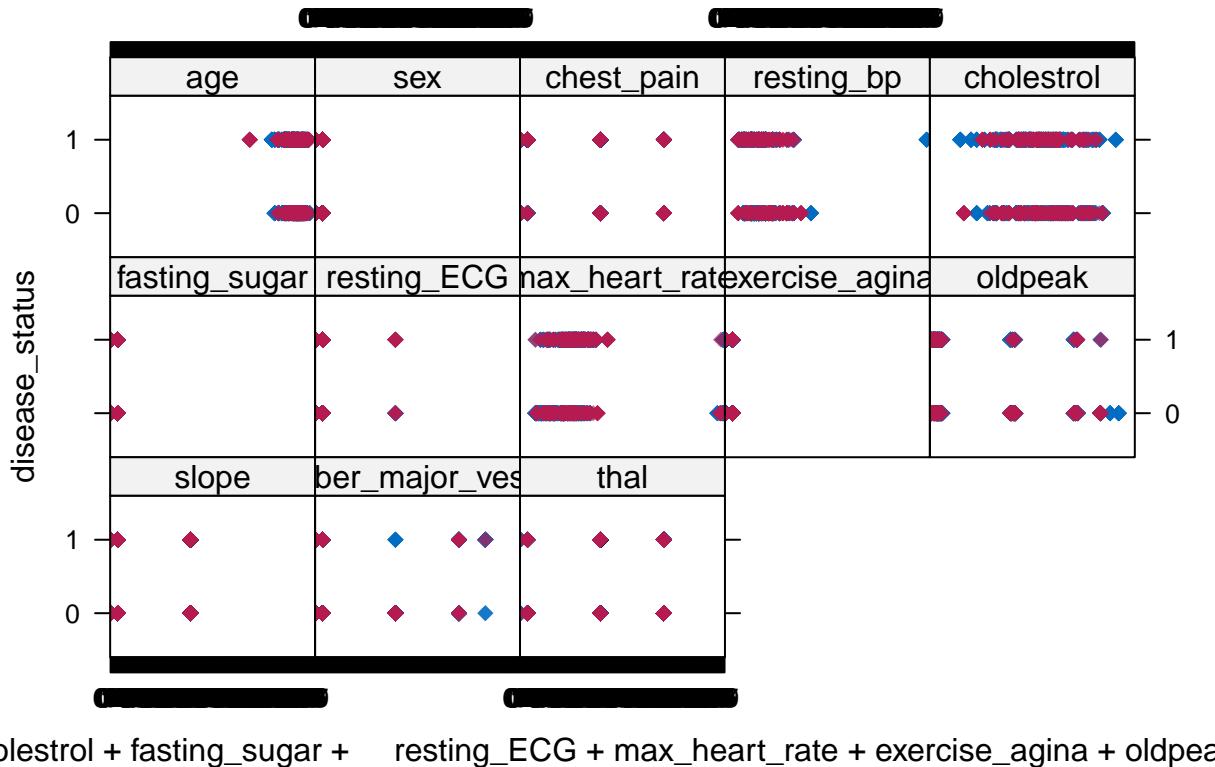
```

## chest_pain           1   1   1
## resting_bp          1   1   1
## cholesterol         1   1   1
## fasting_sugar       1   1   1

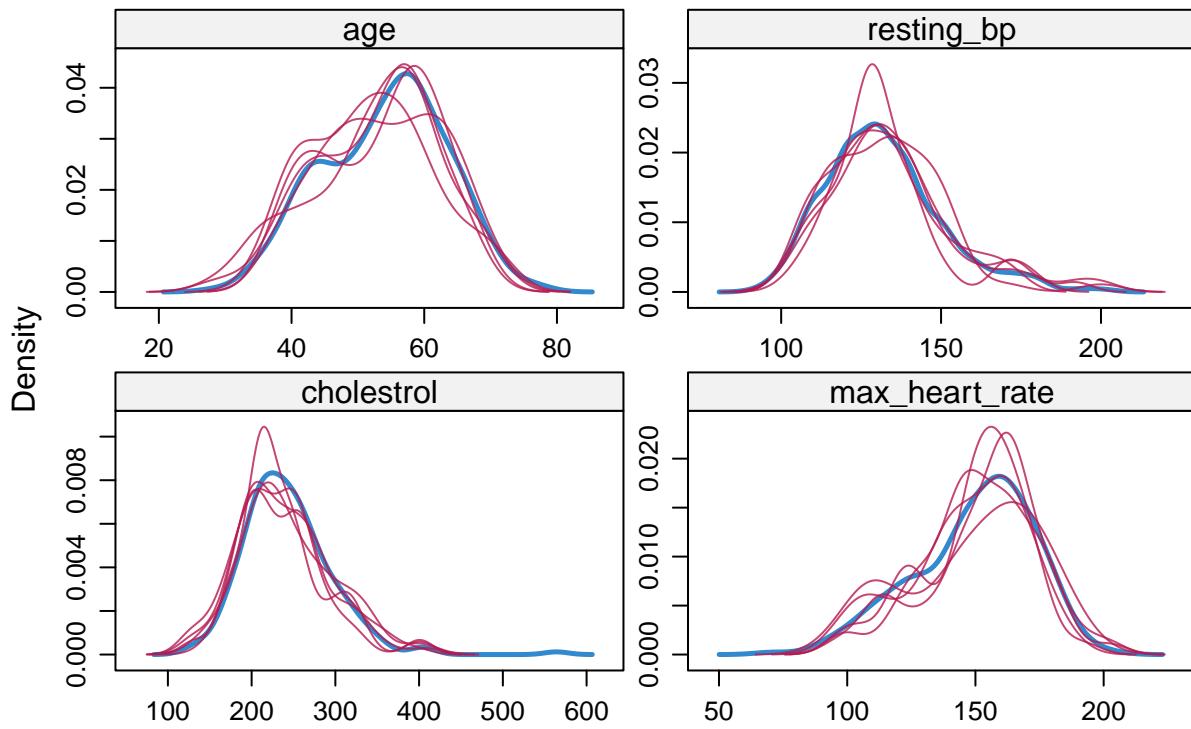
## [1] 1515 16

```

#Inspecting the distribution of original and imputed data



## MICE



## 2. Multiple imputation with PMM

```

## Class: mids
## Number of multiple imputations:  5
## Imputation methods:
##           age          sex      chest_pain
## "pmm"     "pmm"      "pmm"      "pmm"
##           resting_bp   cholesterol  fasting_sugar
## "pmm"     "pmm"      "pmm"      "pmm"
##           resting_ECG  max_heart_rate exercise_agina
## "pmm"     "pmm"      "pmm"      "pmm"
##           oldpeak      slope number_major_vessels
## "pmm"     "pmm"      "pmm"      "pmm"
##           thal        disease_status
## "pmm"     "pmm"      "pmm"
## PredictorMatrix:
##           age sex chest_pain resting_bp cholesterol fasting_sugar
## ## age       0   1         1         1         1         1
## ## sex       1   0         1         1         1         1
## ## chest_pain 1   1         0         1         1         1
## ## resting_bp 1   1         1         0         1         1
## ## cholesterol 1   1         1         1         0         1
## ## fasting_sugar 1   1         1         1         1         0
## ##             resting_ECG max_heart_rate exercise_agina oldpeak slope

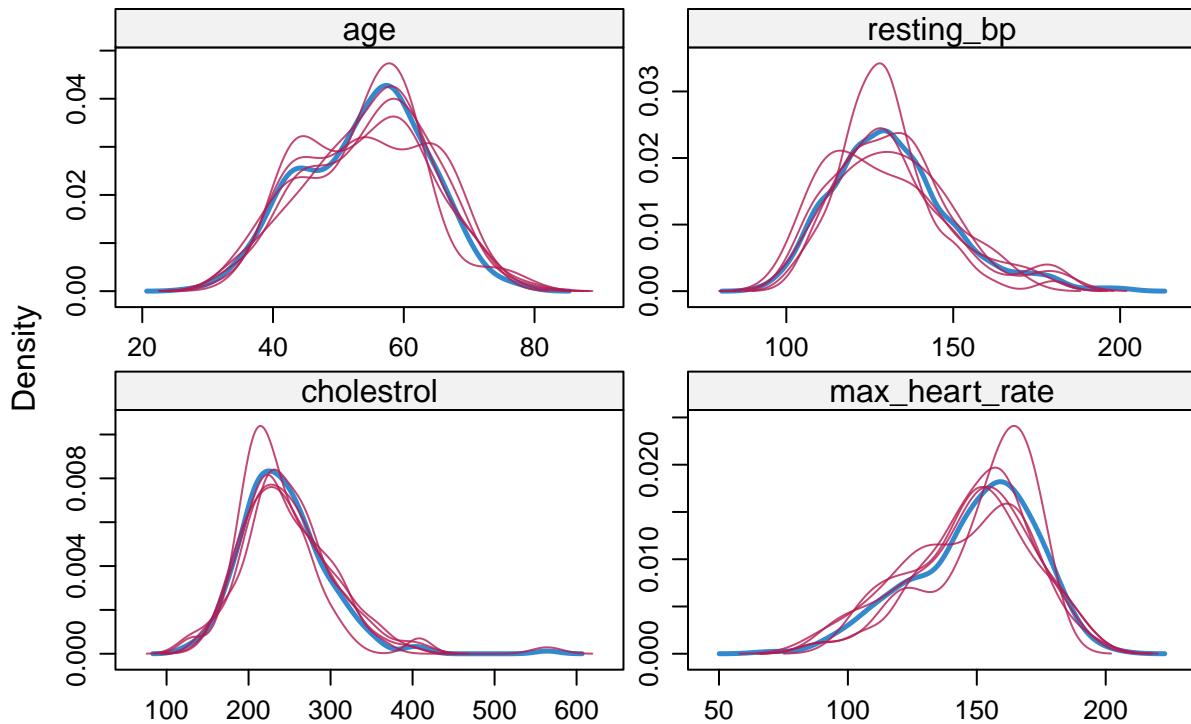
```

```

## age          1           1           1           1           1
## sex          1           1           1           1           1
## chest_pain   1           1           1           1           1
## resting_bp   1           1           1           1           1
## cholesterol  1           1           1           1           1
## fasting_sugar 1           1           1           1           1
##               number_major_vessels thal disease_status
## age              1           1           1
## sex              1           1           1
## chest_pain      1           1           1
## resting_bp      1           1           1
## cholesterol    1           1           1
## fasting_sugar  1           1           1

```

## PMM



### 3. Multiple imputation with Mean

```

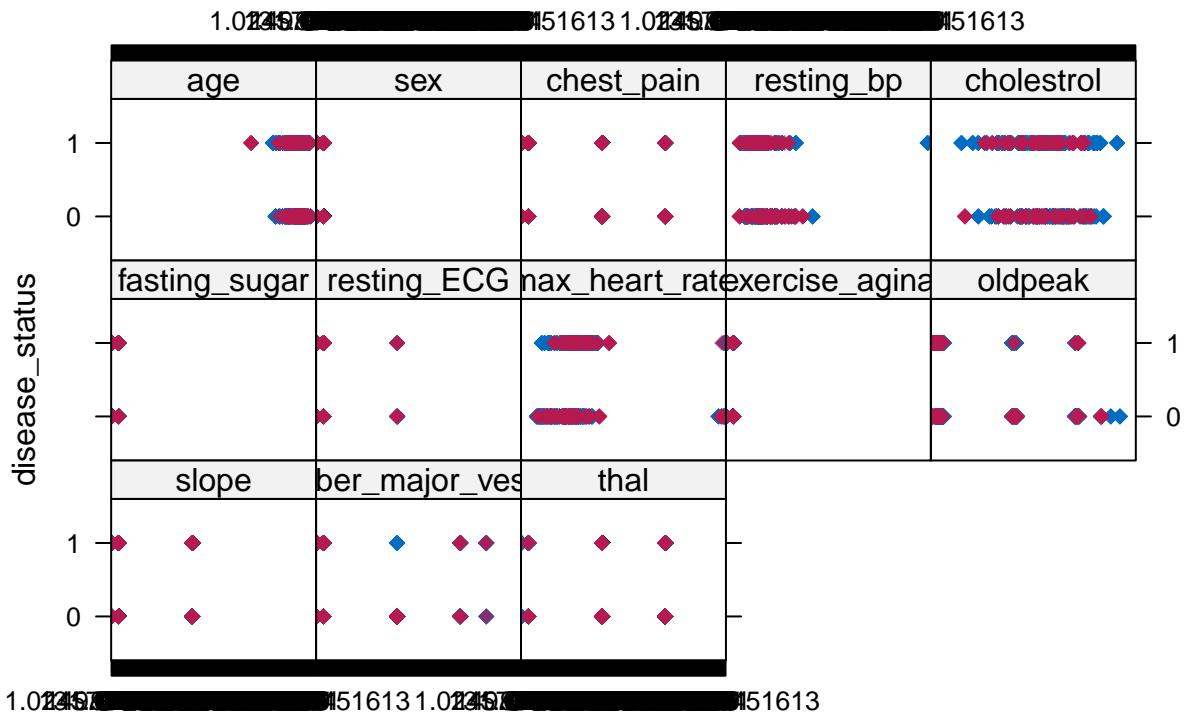
## Class: mids
## Number of multiple imputations:  5
## Imputation methods:
##               age             sex           chest_pain
##       "mean"        "logreg"        "polyreg"
##               resting_bp      cholesterol      fasting_sugar
##       "mean"        "mean"         "logreg"
##               resting_ECG     max_heart_rate exercise_agina

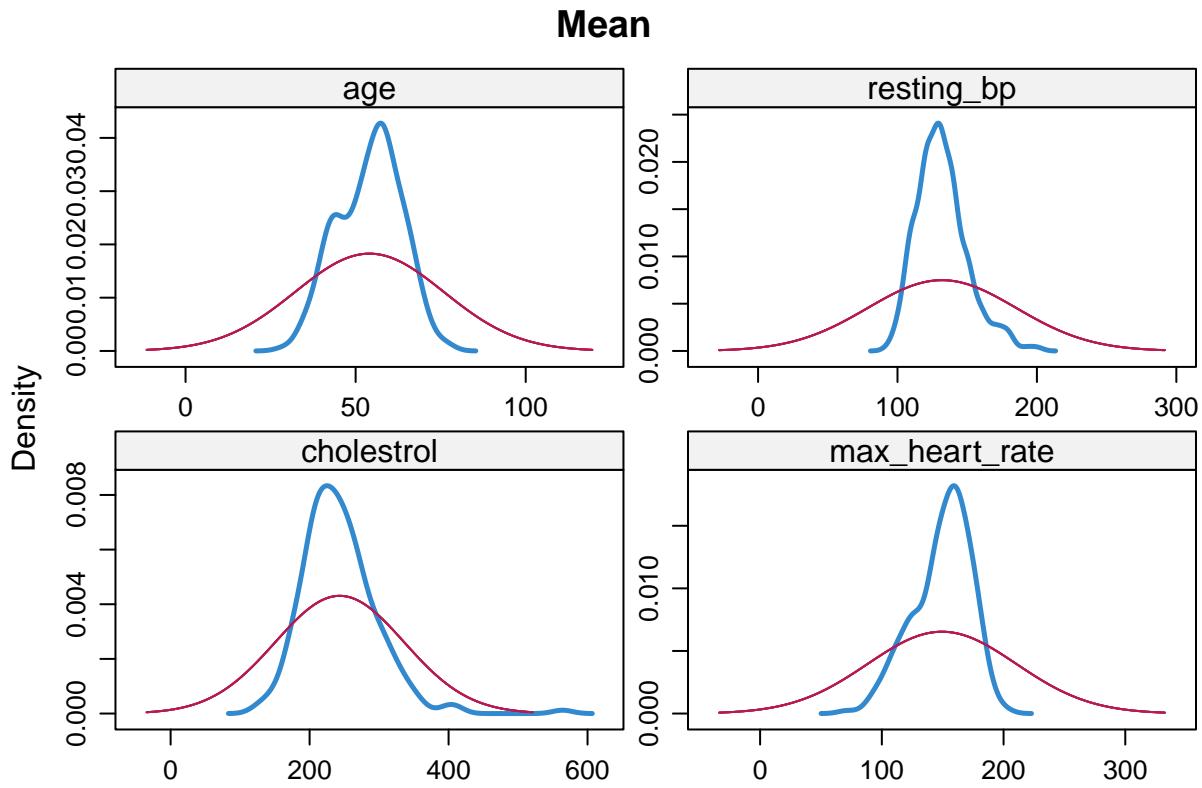
```

```

##          "polyreg"           "mean"           "logreg"
##      oldpeak           slope number_major_vessels
##      "mean"           "polyreg"           "polyreg"
##      thal       disease_status
##      "polyreg"           "logreg"
## PredictorMatrix:
##      age sex chest_pain resting_bp cholesterol fasting_sugar
## age      0   1         1         1         1         1
## sex      1   0         1         1         1         1
## chest_pain 1   1         0         1         1         1
## resting_bp 1   1         1         0         1         1
## cholesterol 1   1         1         1         0         1
## fasting_sugar 1   1         1         1         1         0
##      resting_ECG max_heart_rate exercise_agina oldpeak slope
## age            1             1             1         1         1
## sex            1             1             1         1         1
## chest_pain     1             1             1         1         1
## resting_bp     1             1             1         1         1
## cholesterol    1             1             1         1         1
## fasting_sugar  1             1             1         1         1
##      number_major_vessels thal disease_status
## age            1             1             1
## sex            1             1             1
## chest_pain     1             1             1
## resting_bp     1             1             1
## cholesterol    1             1             1
## fasting_sugar  1             1             1

```





#### 4. Adding missing values only in numerical variables

```

##      age      resting_bp      cholesterol      max_heart_rate
##  Min.   :29.00   Min.   :94.0   Min.   :131.0   Min.   : 71.0
##  1st Qu.:47.00  1st Qu.:120.0  1st Qu.:212.0  1st Qu.:135.5
##  Median :55.00  Median :130.0  Median :243.0  Median :154.0
##  Mean   :54.35  Mean   :131.5  Mean   :248.2  Mean   :149.7
##  3rd Qu.:61.00  3rd Qu.:140.0  3rd Qu.:277.0  3rd Qu.:166.0
##  Max.   :77.00  Max.   :200.0  Max.   :564.0  Max.   :202.0
##  NA's    :66     NA's    :68     NA's    :62     NA's    :59
##      oldpeak
##  Min.   :0.000
##  1st Qu.:0.000
##  Median :0.800
##  Mean   :1.034
##  3rd Qu.:1.600
##  Max.   :6.200
##  NA's    :48

```

**Impute numerical missing values using mean imputation**

```

## Class: mids
## Number of multiple imputations: 5

```

```

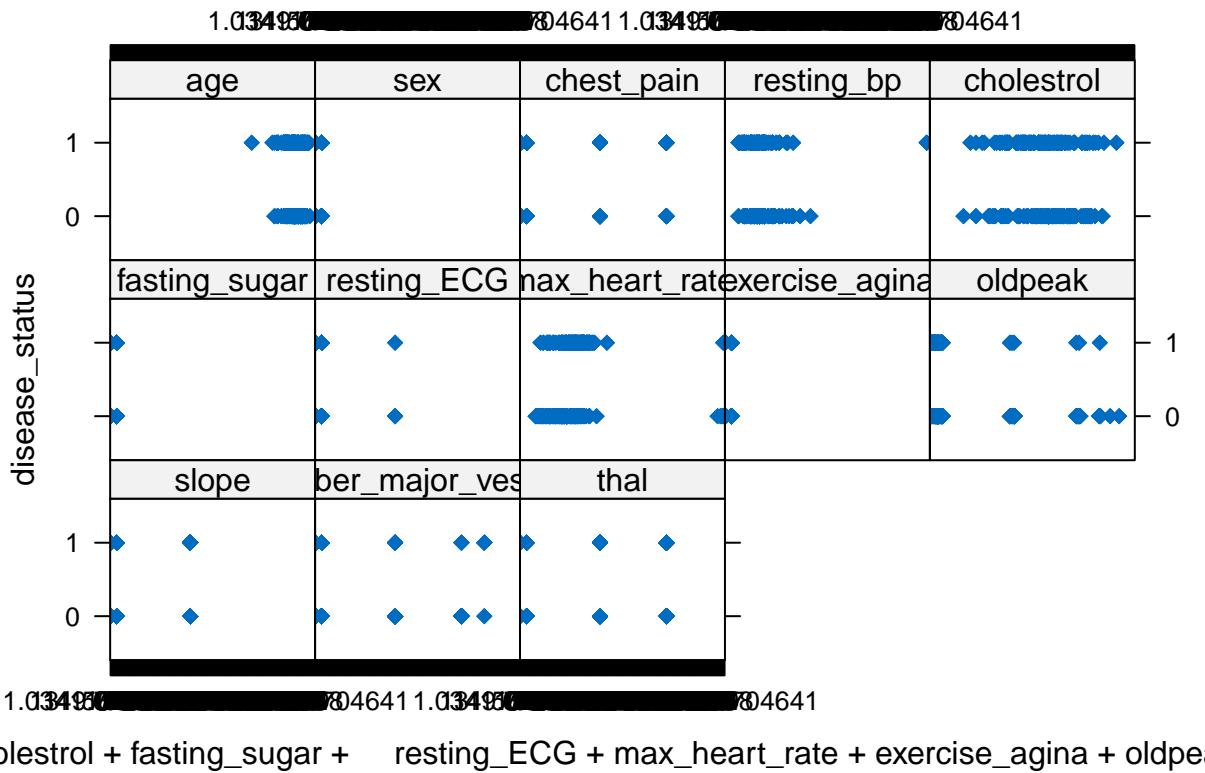
## Imputation methods:
##      age          sex        chest_pain
##      "mean"       ""         ""
##      resting_bp   cholesterol   fasting_sugar
##      "mean"       "mean"     ""
##      resting_ECG  max_heart_rate exercise_agina
##      ""           "mean"     ""
##      oldpeak      slope    number_major_vessels
##      "mean"       ""         ""
##      thal         disease_status
##      ""           ""

## PredictorMatrix:
##      age sex chest_pain resting_bp cholesterol fasting_sugar
##  age    0   1       1       1       1       1
##  sex    1   0       1       1       1       1
##  chest_pain 1   1       0       1       1       1
##  resting_bp 1   1       1       0       1       1
##  cholesterol 1   1       1       1       0       1
##  fasting_sugar 1   1       1       1       1       0
##      resting_ECG max_heart_rate exercise_agina oldpeak slope
##  age            1           1           1       1       1
##  sex            1           1           1       1       1
##  chest_pain    1           1           1       1       1
##  resting_bp    1           1           1       1       1
##  cholesterol   1           1           1       1       1
##  fasting_sugar 1           1           1       1       1
##      number_major_vessels thal disease_status
##  age             1   1       1
##  sex             1   1       1
##  chest_pain     1   1       1
##  resting_bp     1   1       1
##  cholesterol   1   1       1
##  fasting_sugar 1   1       1

##      .imp      .id      age      sex      chest_pain      resting_bp
##  Min.   :1  Min.   : 1  Min.   :29.00  0: 480  0:715      Min.   : 94.0
##  1st Qu.:2  1st Qu.: 76  1st Qu.:50.00  1:1035  1:250      1st Qu.:123.0
##  Median :3  Median :152  Median :54.35   2:435   Median :131.5
##  Mean   :3  Mean   :152  Mean   :54.35   3:115   Mean   :131.5
##  3rd Qu.:4  3rd Qu.:228 3rd Qu.:59.00   4:100   3rd Qu.:138.0
##  Max.   :5  Max.   :303  Max.   :77.00   5: 10   Max.   :200.0
##      cholesterol      fasting_sugar      resting_ECG      max_heart_rate      exercise_agina
##  Min.   :131.0  0:1290      0:735      Min.   : 71.0  0:1020
##  1st Qu.:221.0  1: 225      1:760      1st Qu.:142.0  1: 495
##  Median :248.2           2: 20      Median :149.7
##  Mean   :248.2           3: 20      Mean   :149.7
##  3rd Qu.:268.0           4: 20      3rd Qu.:163.0
##  Max.   :564.0           5: 20      Max.   :202.0
##      oldpeak      slope      number_major_vessels      thal      disease_status
##  Min.   :0.000  0:105      0:875      0: 10      0:690
##  1st Qu.:0.000  1:700      1:325      1: 90      1:825
##  Median :1.034  2:710      2:190      2:830
##  Mean   :1.034  3:100      3:585
##  3rd Qu.:1.400  4: 25

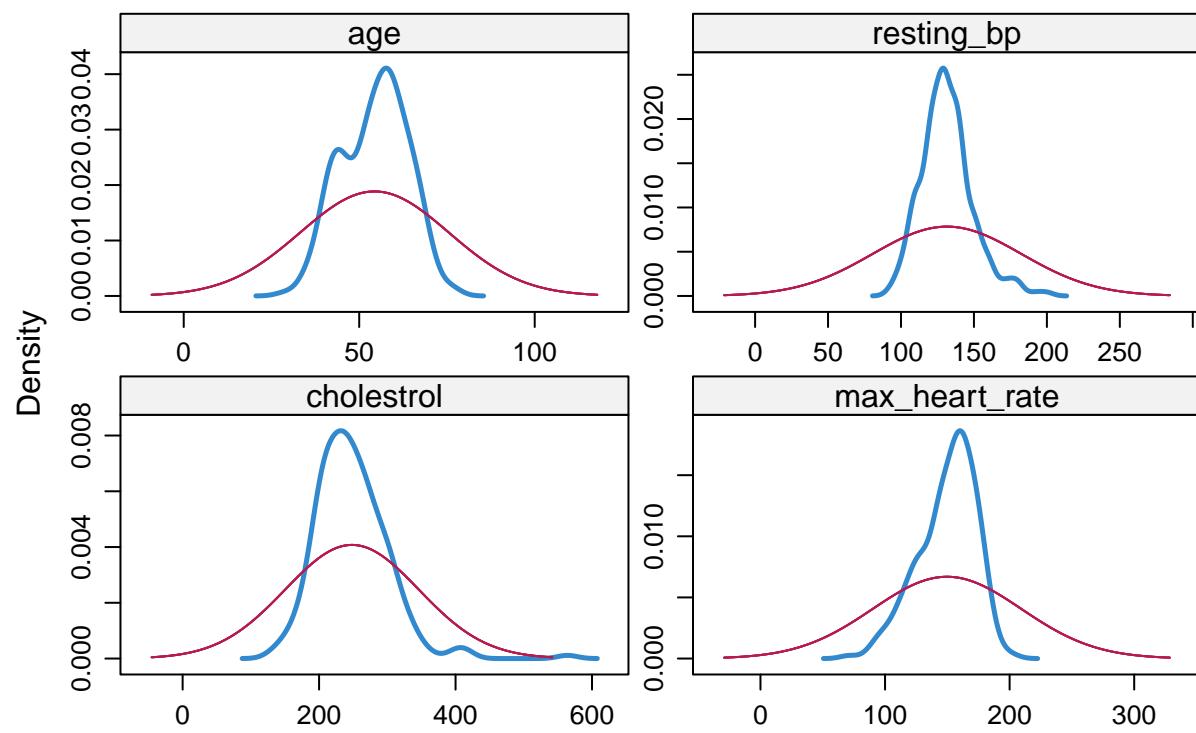
```

## Max. : 6.200



cholesterol + fasting\_sugar + resting\_ECG + max\_heart\_rate + exercise\_agina + oldpeak

## Mean\_numeric



## Summary table for original and imputed data using gtsummary

**Characteristic**	**Mean**, N = 303	**Original**, N = 303	**PMM**, N = 303
age	54 (8)	54 (9)	54 (9)
sex			
0	303 (29%)	303 (32%)	303 (32%)
1	303 (71%)	303 (68%)	303 (68%)
chest_pain			
0	303 (48%)	303 (47%)	303 (48%)
1	303 (17%)	303 (17%)	303 (17%)
2	303 (27%)	303 (29%)	303 (28%)
3	303 (7.3%)	303 (7.6%)	303 (7.3%)
resting_bp	132 (16)	132 (18)	132 (18)
cholesterol	243 (46)	246 (52)	245 (53)
fasting_sugar			
0	303 (84%)	303 (85%)	303 (84%)
1	303 (16%)	303 (15%)	303 (16%)
resting_ECG			
0	303 (48%)	303 (49%)	303 (49%)
1	303 (49%)	303 (50%)	303 (50%)
2	303 (2.6%)	303 (1.3%)	303 (1.0%)
max_heart_rate	149 (21)	150 (23)	149 (23)
exercise_agina			
0	303 (64%)	303 (67%)	303 (65%)
1	303 (36%)	303 (33%)	303 (35%)
oldpeak	1.02 (1.04)	1.04 (1.16)	1.00 (1.17)
slope			
0	303 (8.3%)	303 (6.9%)	303 (7.9%)
1	303 (47%)	303 (46%)	303 (48%)
2	303 (45%)	303 (47%)	303 (45%)
number_major_vessels			
0	303 (58%)	303 (58%)	303 (59%)
1	303 (22%)	303 (21%)	303 (23%)
2	303 (12%)	303 (13%)	303 (11%)
3	303 (6.6%)	303 (6.6%)	303 (5.9%)
4	303 (1.7%)	303 (1.7%)	303 (1.0%)
thal			
0	303 (1.3%)	303 (0.7%)	303 (1.0%)
1	303 (7.3%)	303 (5.9%)	303 (5.9%)
2	303 (53%)	303 (55%)	303 (54%)
3	303 (38%)	303 (39%)	303 (39%)
disease_status			
0	303 (47%)	303 (46%)	303 (48%)
1	303 (53%)	303 (54%)	303 (52%)