

Source Code Đồ án cuối kỳ

Thông tin chung

- **Môn:** Thiết kế và Phân tích thực nghiệm
- **Lớp:** DS304.L21
- **Nhóm 9**
- **Thành viên nhóm:**

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- **Đề tài báo cáo:** Phân tích ảnh hưởng của các chỉ số sức khỏe đến tiến triển bệnh đái tháo đường

Phân tích thăm dò và trực quan bộ dữ liệu (Python)

In [1]:

```
import pandas as pd
import numpy as np

import seaborn as sns
import matplotlib.pyplot as plt

df = pd.read_csv('/content/drive/MyDrive/#2020-2021 HK2/TK&PTTN/diabetes.tab.tsv', sep='\t')
df
```

Out[1]:

	AGE	SEX	BMI	BP	S1	S2	S3	S4	S5	S6	Y
0	59	2	32.1	101.00	157	93.2	38.0	4.00	4.8598	87	151
1	48	1	21.6	87.00	183	103.2	70.0	3.00	3.8918	69	75
2	72	2	30.5	93.00	156	93.6	41.0	4.00	4.6728	85	141
3	24	1	25.3	84.00	198	131.4	40.0	5.00	4.8903	89	206
4	50	1	23.0	101.00	192	125.4	52.0	4.00	4.2905	80	135
...
437	60	2	28.2	112.00	185	113.8	42.0	4.00	4.9836	93	178
438	47	2	24.9	75.00	225	166.0	42.0	5.00	4.4427	102	104
439	60	2	24.9	99.67	162	106.6	43.0	3.77	4.1271	95	132
440	36	1	30.0	95.00	201	125.2	42.0	4.79	5.1299	85	220
441	36	1	19.6	71.00	250	133.2	97.0	3.00	4.5951	92	57

442 rows × 11 columns

In [2]:

```
import seaborn as sns
import matplotlib.pyplot as plt

#defining colour palette
def custom_palette(custom_colors):
    customPalette = sns.set_palette(sns.color_palette(custom_colors))
    sns.palplot(sns.color_palette(custom_colors), size=0.8)
    plt.tick_params(axis='both', labelsize=0, length = 0)

red = ["#4f000b", "#720026", "#ce4257", "#ff7f51", "#ff9b54"]
bo = ["#6930c3", "#5e60ce", "#0096c7", "#48cae4", "#ade8f4", "#ff7f51", "#ff9b54", "#ffbf69"]
pink = ["#aa4465", "#dd2d4a", "#f26a8d", "#f49cbb", "#ffcbbf", "#e2aaff", "#ff86c8", "#ffa3a5", "#ffbf81", "#e9b827", "#f9e576"]
custom_palette(pink)
```



In [3]:

```
df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 442 entries, 0 to 441
Data columns (total 11 columns):
#   Column  Non-Null Count  Dtype
---  -
```

```

0  AGE      442 non-null    int64
1  SEX      442 non-null    int64
2  BMI      442 non-null    float64
3  BP       442 non-null    float64
4  S1       442 non-null    int64
5  S2       442 non-null    float64
6  S3       442 non-null    float64
7  S4       442 non-null    float64
8  S5       442 non-null    float64
9  S6       442 non-null    int64
10 Y        442 non-null    int64
dtypes: float64(6), int64(5)
memory usage: 38.1 KB

```

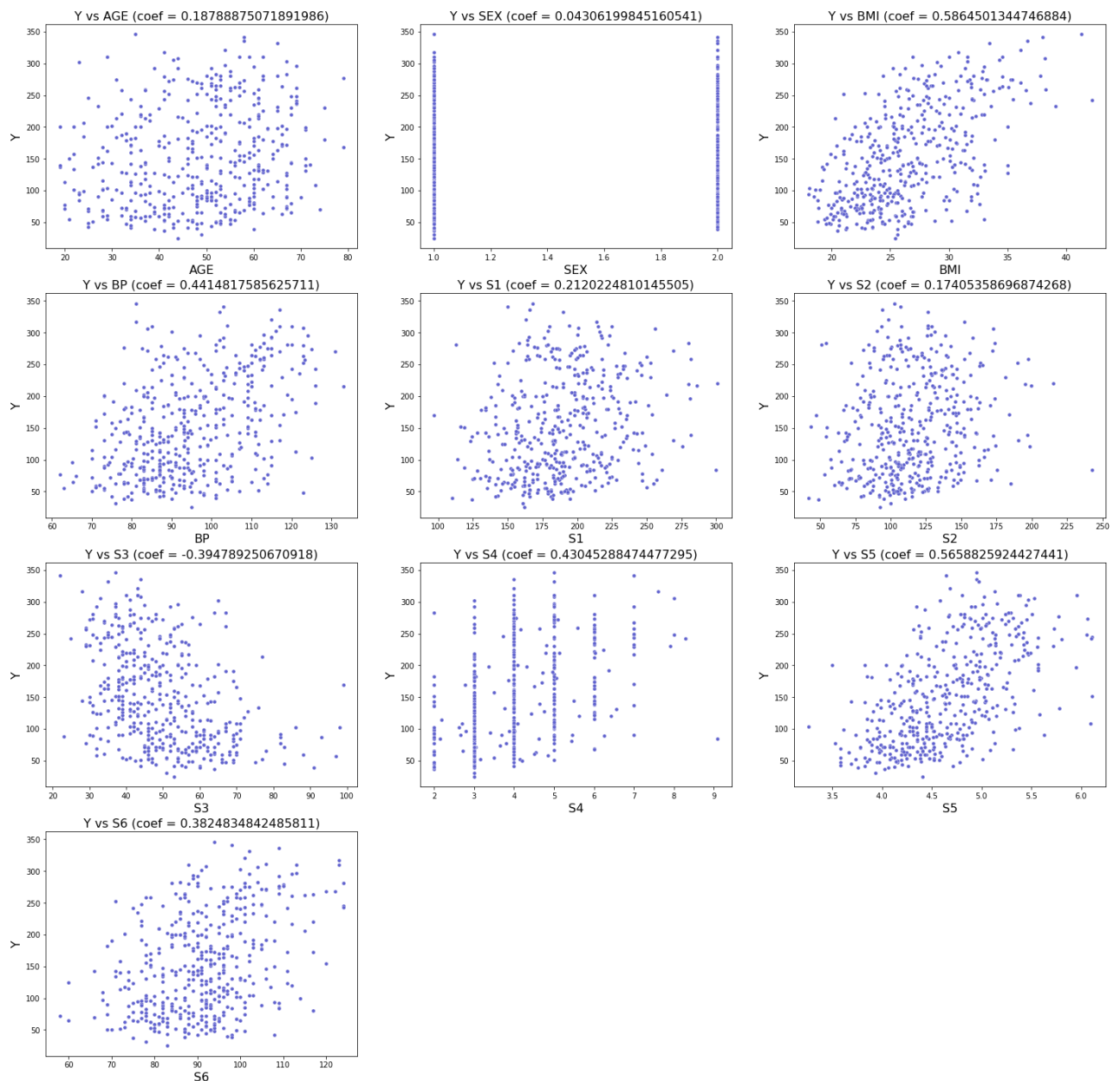
In [4]: `df.describe()`

	AGE	SEX	BMI	BP	S1	S2	S3	S4	S5	S6	Y
count	442.000000	442.000000	442.000000	442.000000	442.000000	442.000000	442.000000	442.000000	442.000000	442.000000	442.000000
mean	48.518100	1.468326	26.375792	94.647014	189.140271	115.439140	49.788462	4.070249	4.641411	91.260181	152.133484
std	13.109028	0.499561	4.418122	13.831283	34.608052	30.413081	12.934202	1.290450	0.522391	11.496335	77.093005
min	19.000000	1.000000	18.000000	62.000000	97.000000	41.600000	22.000000	2.000000	3.258100	58.000000	25.000000
25%	38.250000	1.000000	23.200000	84.000000	164.250000	96.050000	40.250000	3.000000	4.276700	83.250000	87.000000
50%	50.000000	1.000000	25.700000	93.000000	186.000000	113.000000	48.000000	4.000000	4.620050	91.000000	140.500000
75%	59.000000	2.000000	29.275000	105.000000	209.750000	134.500000	57.750000	5.000000	4.997200	98.000000	211.500000
max	79.000000	2.000000	42.200000	133.000000	301.000000	242.400000	99.000000	9.090000	6.107000	124.000000	346.000000

```

In [5]: corr = df.corr()
fig, axs = plt.subplots(4,3,figsize=(25,25))
for i, ax in zip(range(10), axs.flat):
    sns.scatterplot(ax=ax, data=df, x=df.columns[:-1][i] , y='Y',color=bo[1],s=25)
    c = corr['Y'][df.columns[:-1][i]]
    ax.set_title('Y vs {} (coef = {})'.format(df.columns[:-1][i],str(c)), fontsize=16)
    ax.set_xlabel(df.columns[:-1][i], fontsize = 16)
    ax.set_ylabel('Y', fontsize = 16)
fig.delaxes(axs[3][1])
fig.delaxes(axs[3][2])
plt.savefig('Scatter_Y.png',transparent=False,bbox_inches = 'tight',dpi=300)
plt.show()

```



```
In [6]: def hist(col,title,norm):
fig, ax = plt.subplots(1,1, figsize=(16,9))
#plt.figure(figsize = (16,9))
#ax.set_xlim(0,400)
ax = sns.distplot(col,hist=True,bins=35,fit=norm);

values = np.array([patch.get_height() for patch in ax.patches])

#normalizing the values to get a range of colours
norm = plt.Normalize(values.min(), values.max())

#range of colours from colourmap-rainbow
colors = plt.cm.rainbow(norm(values))

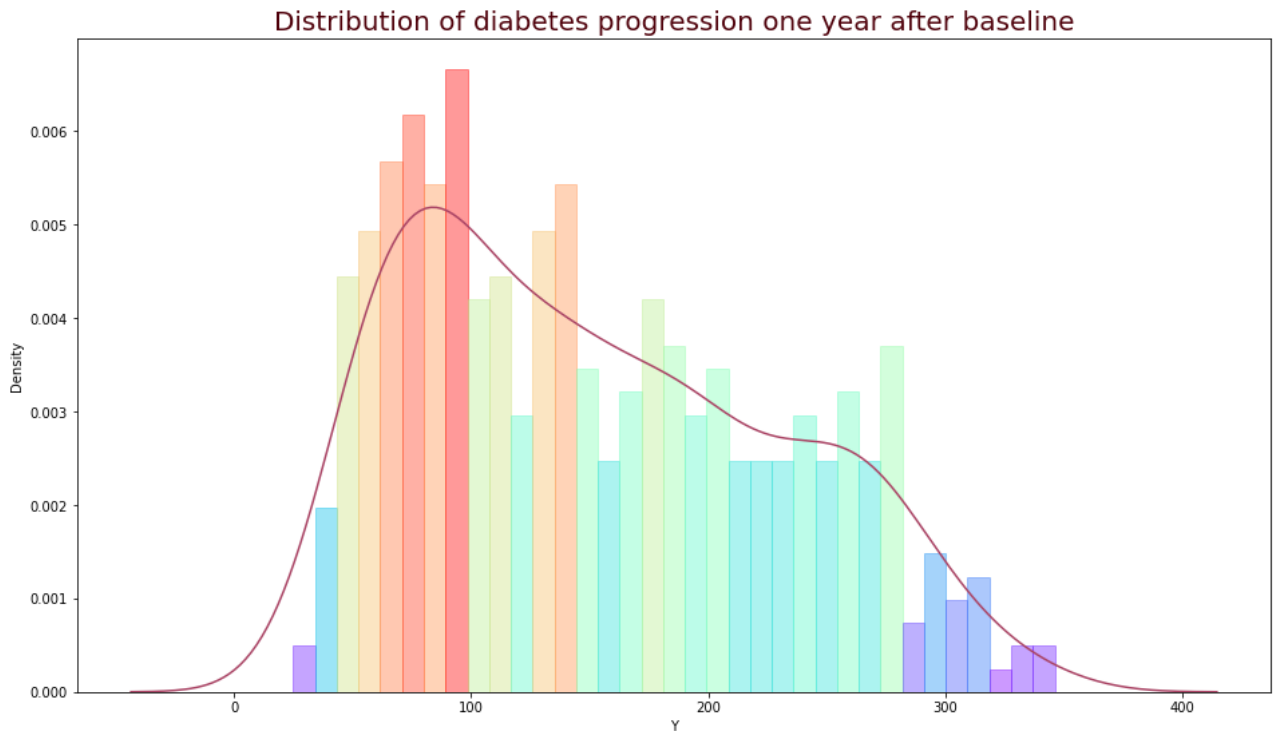
#set colour for each patch
for patch, color in zip(ax.patches, colors):
    patch.set_color(color)

plt.title(title, size = 20, color = red[0])
```

```
In [7]: hist(df['Y'],'Distribution of diabetes progression one year after baseline',None)
plt.savefig('Dist_Y.png',transparent=False,bbox_inches = 'tight',dpi=300)
```

/usr/local/lib/python3.7/dist-packages/seaborn/distributions.py:2557: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

warnings.warn(msg, FutureWarning)



```
In [8]: def get_percent(status):
    if(status=='Underweight'):
        return str(np.round((len(df[df['BMI']<=18.5])/442)*100,2))+'%'
    if(status=='Normal'):
        tmp = df[df['BMI']>18.5]
        return str(np.round((len(tmp[tmp['BMI']<=25])/442)*100,2))+'%'
    if(status=='Overweight'):
        tmp = df[df['BMI']>25]
        return str(np.round((len(tmp[tmp['BMI']<=30])/442)*100,2))+'%'
    tmp = df[df['BMI']>30]
    return str(np.round((len(tmp[tmp['BMI']>30])/442)*100,2))+'%
```

```
In [9]: fig, ax = plt.subplots(1,1, figsize=(20,12))
ax = sns.distplot(df['BMI'], kde=True, bins=24);

values = np.array([patch.get_height() for patch in ax.patches])

norm = plt.Normalize(values.min(), values.max())
colors = plt.cm.rainbow(norm(values))
for patch, color in zip(ax.patches, colors):
    patch.set_color(color)

#colours for different bmi categories
span_color = ['#00a8e8', '#25a18e', '#fb8500', '#ef476f']

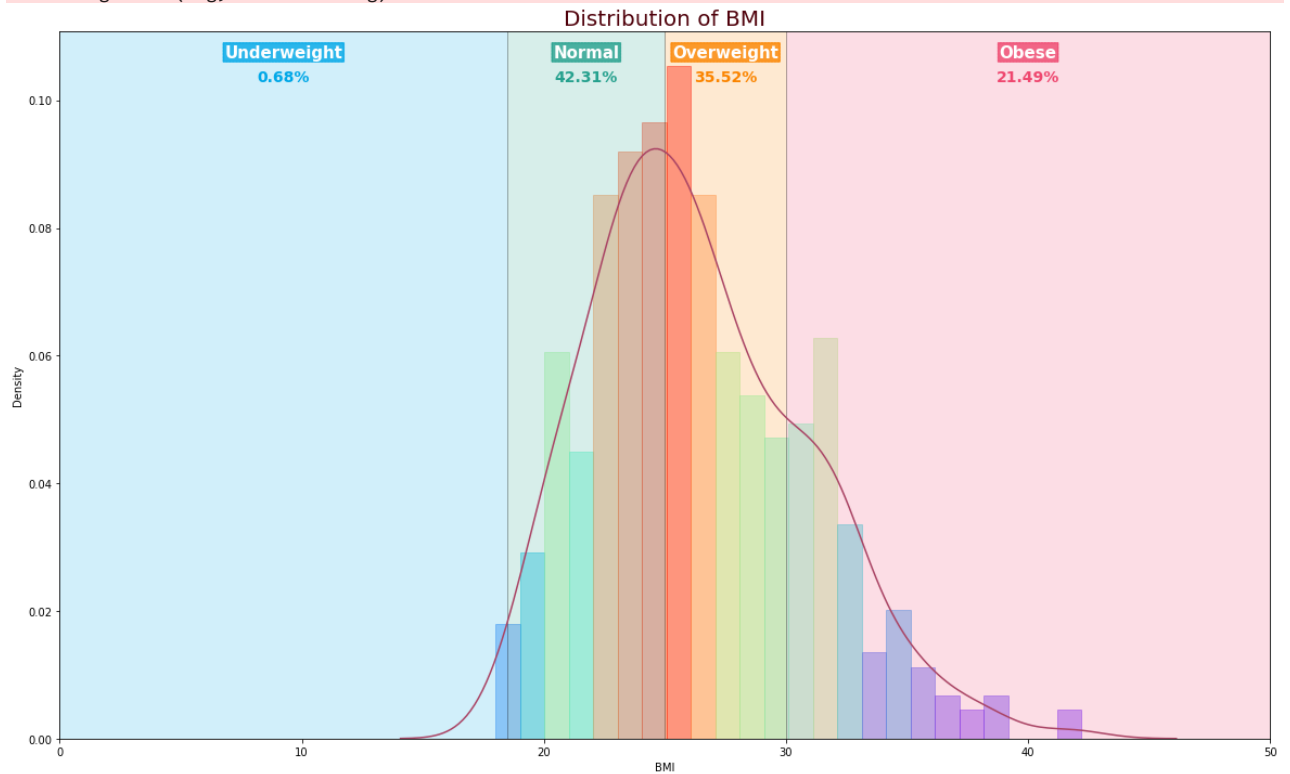
#range of values for different bmi categories
span_range = [[0,18.5], [18.5,25], [25,30], [30,50]]
ax.set_xlim(0,50)
for idx, span_title in enumerate(['Underweight', 'Normal', 'Overweight', 'Obese']):
    ax.annotate(span_title,
                xy=(sum(span_range[idx])/2, 0.029),
                xytext=(0,470), textcoords='offset points',
                va='top', ha="center",
                color="w", fontsize=5, fontweight='bold',
                size=15,
                bbox=dict(boxstyle='sawtooth', pad=0.1, color=span_color[idx], alpha=0.8))

    ax.annotate(get_percent(span_title),
                xy=(sum(span_range[idx])/2, 0.025),
                xytext=(0,470), textcoords='offset points',
                va='top', ha="center",
                color=span_color[idx], fontsize=5, fontweight='bold',
                size=14,
                bbox=dict(boxstyle='sawtooth', pad=0.1, color=span_color[idx], alpha=0))

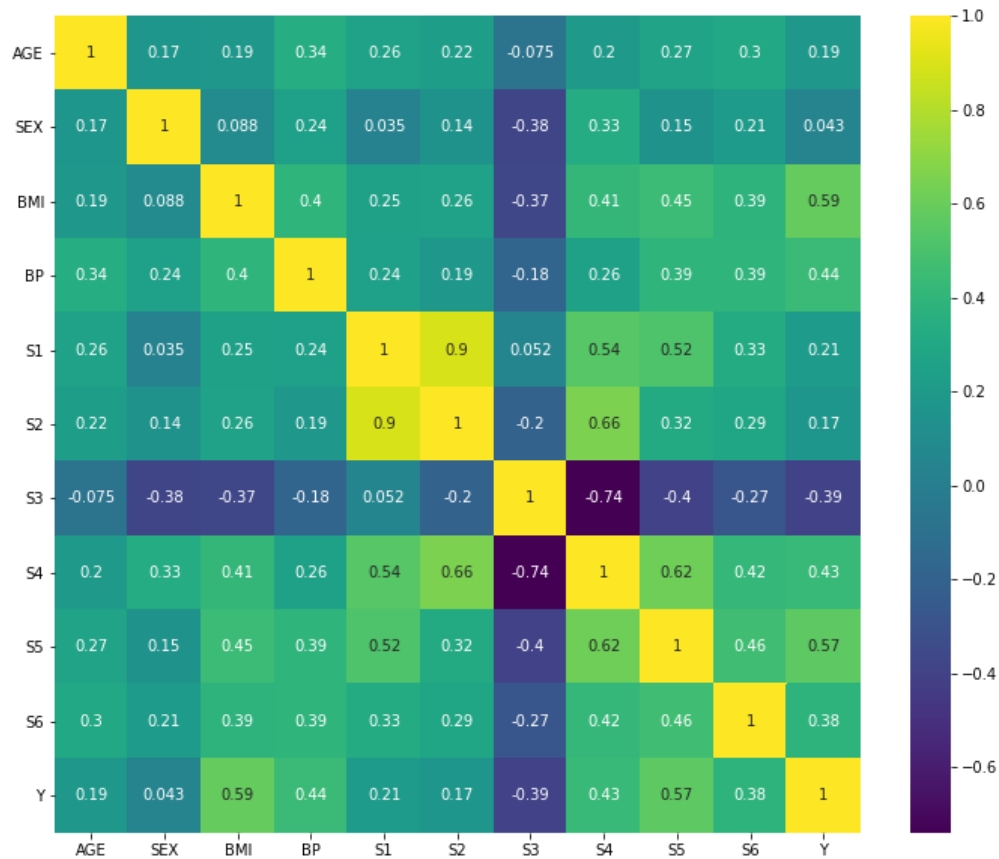
    ax.axvspan(span_range[idx][0], span_range[idx][1], color=span_color[idx], alpha=0.18, ec='black')
plt.title("Distribution of BMI", size = 20, color = red[0])
plt.savefig('Dist_BMI.png', transparent=False, bbox_inches = 'tight', dpi=300)
plt.show()
```

/usr/local/lib/python3.7/dist-packages/seaborn/distributions.py:2557: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with

th similar flexibility) or `histplot` (an axes-level function for histograms).
 warnings.warn(msg, FutureWarning)



```
In [18]: temp_df = df[df.columns]
corr = temp_df.corr()
plt.figure(figsize=(12,10))
sns.heatmap(corr,
            xticklabels=corr.columns.values,
            yticklabels=corr.columns.values,annot=True,cmap='viridis')
plt.yticks(rotation=360)
plt.savefig('Corr.png',transparent=False,bbox_inches = 'tight',dpi=300)
```

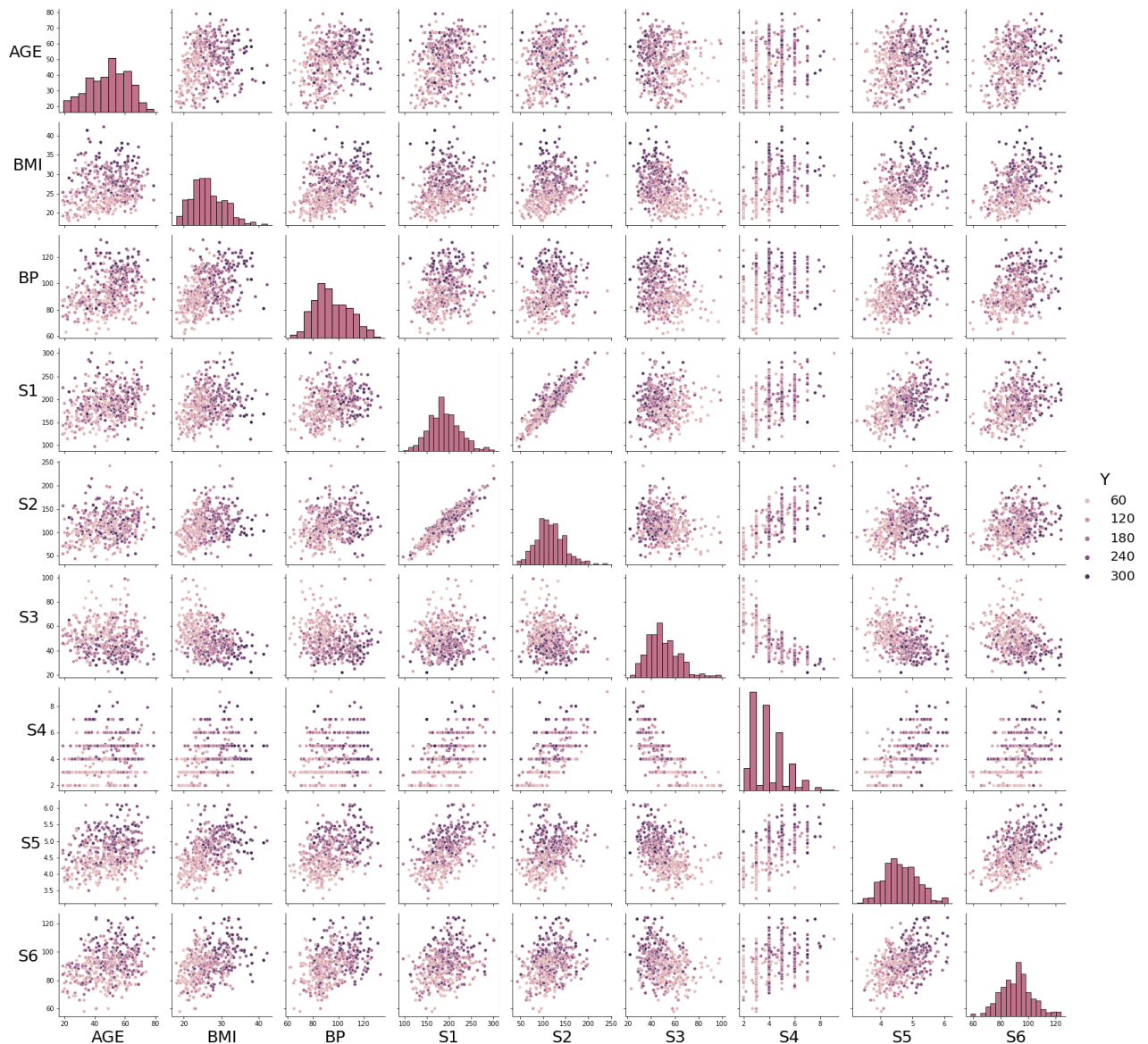


```
In [11]: df2 = df.drop(['SEX'],axis=1)
g = sns.PairGrid(df2,hue='Y')
g.map_diag(sns.histplot,hue=None)
```

```

g.map_offdiag(sns.scatterplot,s=25).add_legend(title='Y', fontsize= 20)
for axes in g.axes.flat:
    axes.set_ylabel(axes.get_ylabel(), rotation=0, horizontalalignment='right')
    axes.xaxis.get_label().set_fontsize(25)
    axes.yaxis.get_label().set_fontsize(25)
plt.setp(g._legend.get_title(), fontsize=25)
plt.savefig('Scatter_factors.png',transparent=False,bbox_inches = 'tight',dpi=300)
plt.show()

```

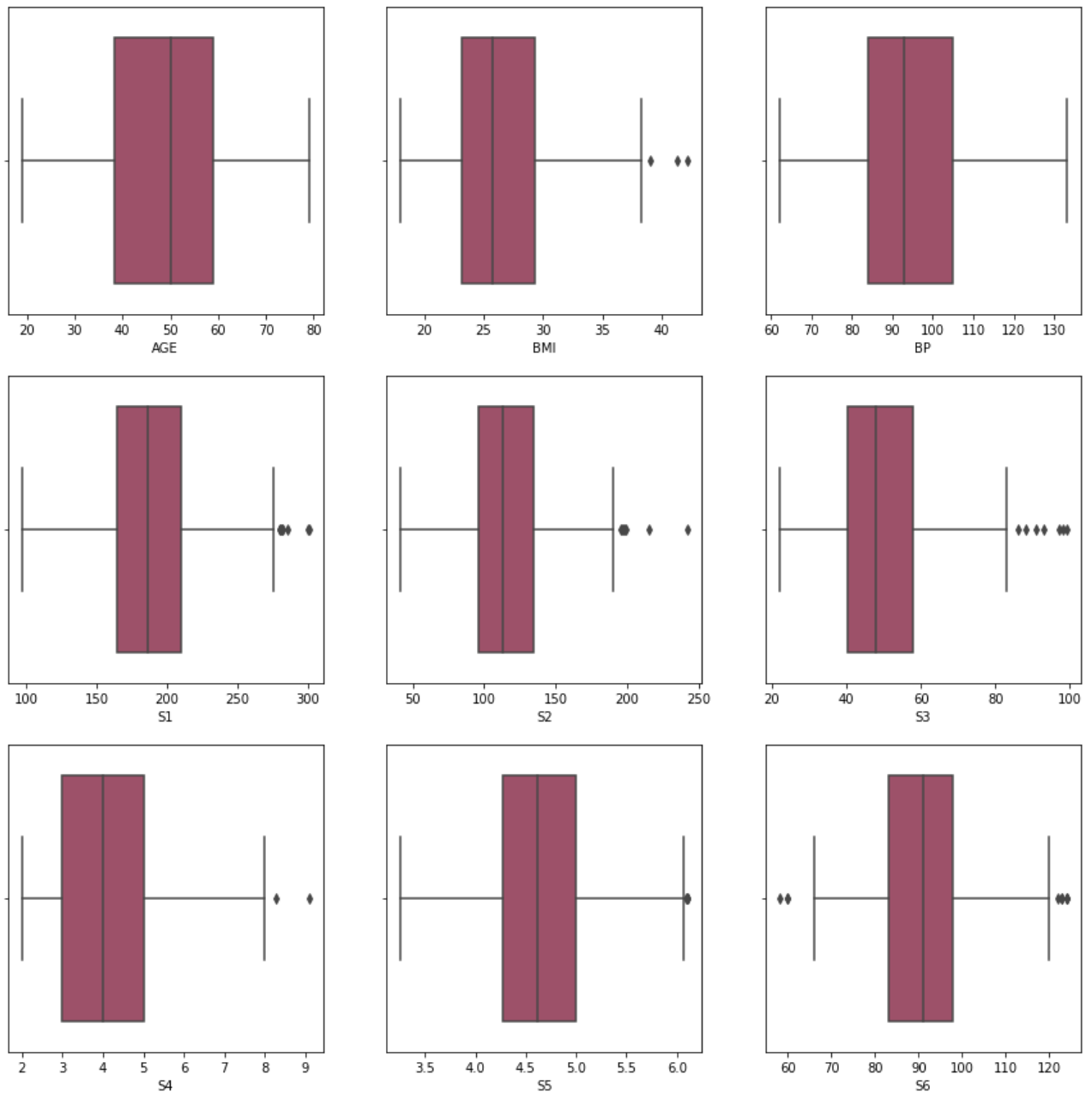


Data Preprocessing

```

In [12]: cols=['AGE', 'BMI', 'BP', 'S1', 'S2', 'S3', 'S4', 'S5', 'S6']
fig, axs = plt.subplots(3,3,figsize=(15,15))
for i, ax in zip(range(10), axs.flat):
    sns.boxplot(x=df[cols[i]],ax=ax)
plt.savefig('box.png',dpi=200)

```



```
In [13]: df_p = df.copy()
cols=['AGE','BMI','BP','S1','S2','S3','S4','S5','S6']
for attr in df_p.columns[:-1]:
    if(attr!='SEX' and attr!='Y'):
        Q1= np.percentile(df_p[attr], 25)
        Q3 = np.percentile(df_p[attr], 75)
        IQR = Q3 - Q1
        olr_up = Q3+1.5*IQR
        olr_low = Q1-1.5*IQR
        df_p = df_p.drop(df_p[df_p[attr]>olr_up].index)
        df_p = df_p.drop(df_p[df_p[attr]<olr_low].index)
```

```
In [14]: import matplotlib as mpl
from scipy.stats import skew
def compare(factor_name):
    i = factor_name

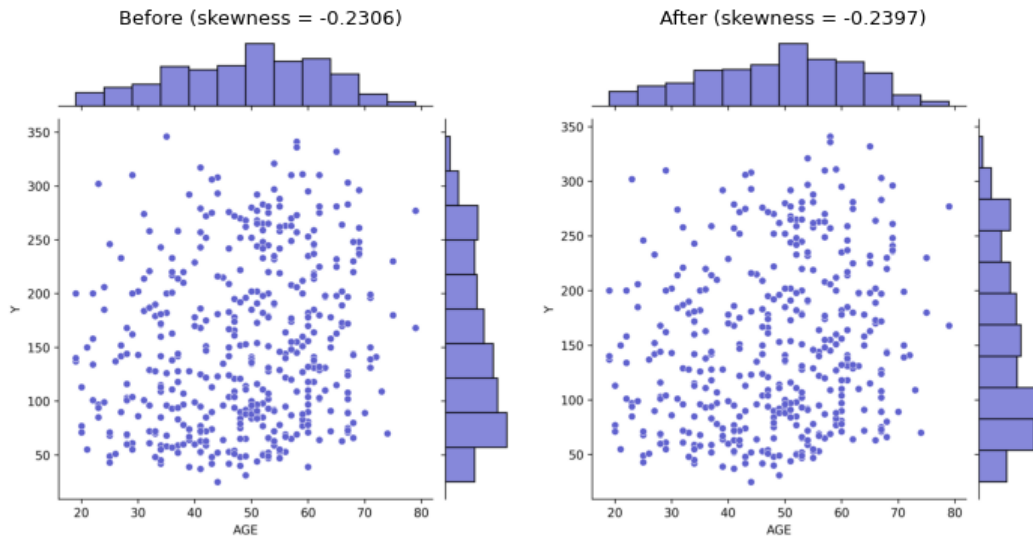
    tmp1 = sns.jointplot(data=df, x= i, y='Y',color=bo[1])
    tmp2 = sns.jointplot(data = df_p, x= i, y='Y',color=bo[1])
    tmp1.savefig('tmp1.png',dpi=300)
    plt.close(tmp1.fig)
    tmp2.savefig('tmp2.png',dpi=300)
    plt.close(tmp2.fig)
    fig, axs = plt.subplots(1,2,figsize=(10,10))
    axs[0].imshow(mpl.image.imread('tmp1.png'))
    axs[0].set_title('Before (skewness = {})'.format(np.round(skew(df[i]),4)),fontsize=13)

    axs[1].imshow(mpl.image.imread('tmp2.png'))
    axs[1].set_title('After (skewness = {})'.format(np.round(skew(df_p[i]),4)),fontsize=13)
    [ax.set_axis_off() for ax in axs.ravel()]
```

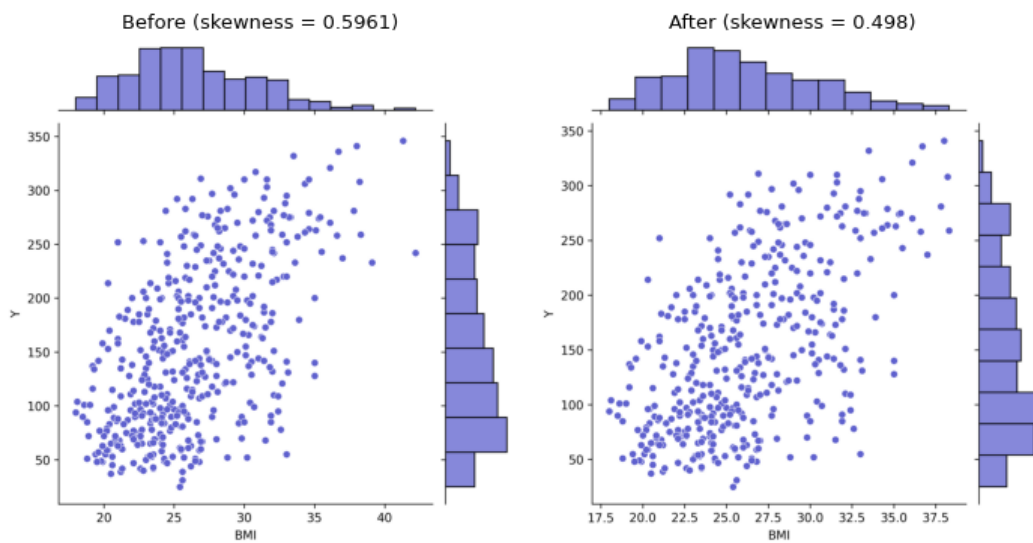
```
plt.tight_layout()
plt.suptitle(factor_name+' Outlier Removal',y=0.8,fontsize=16)
fig.savefig('{}OR.png'.format(factor_name),dpi=100)
plt.show()
```

```
In [15]: for i in cols:
         compare(i)
```

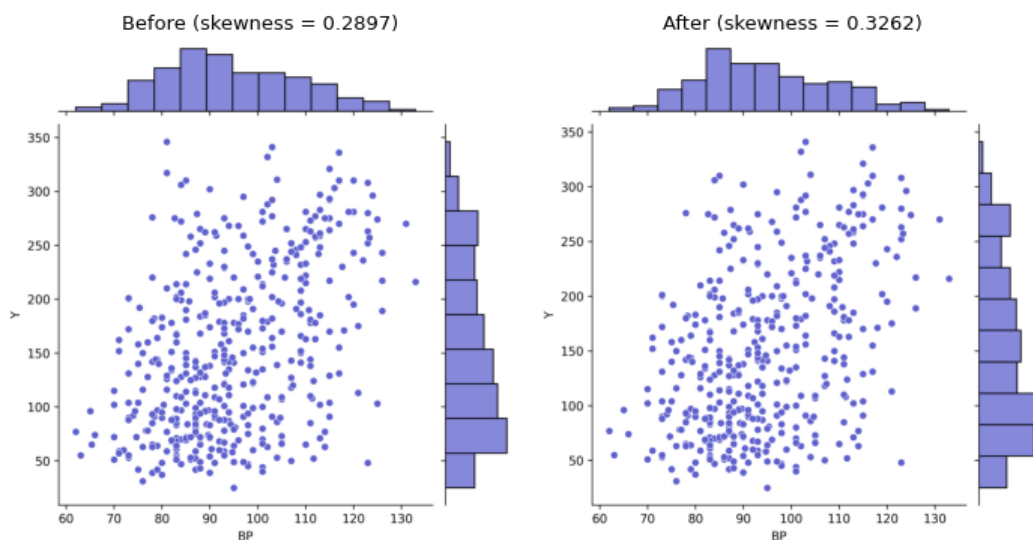
AGE Outlier Removal



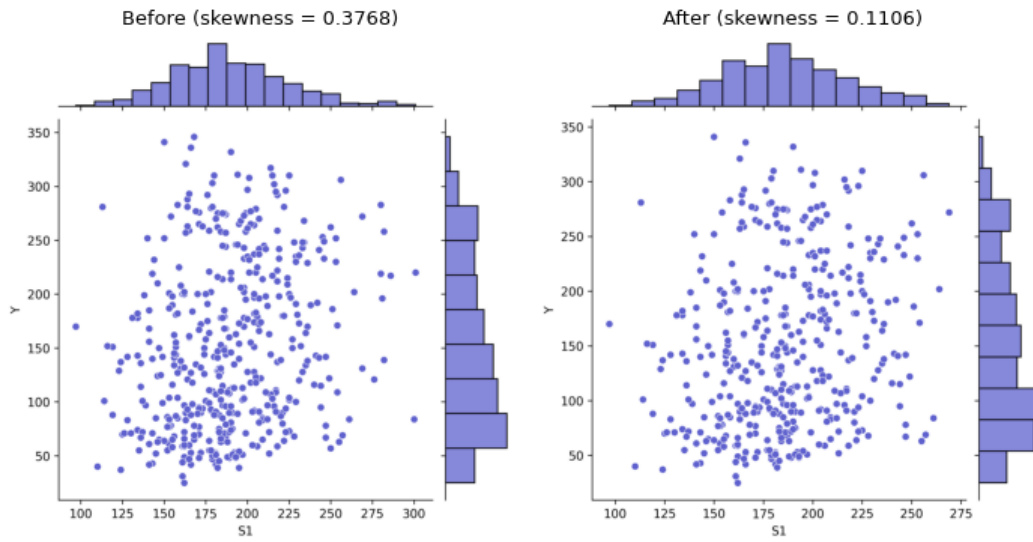
BMI Outlier Removal



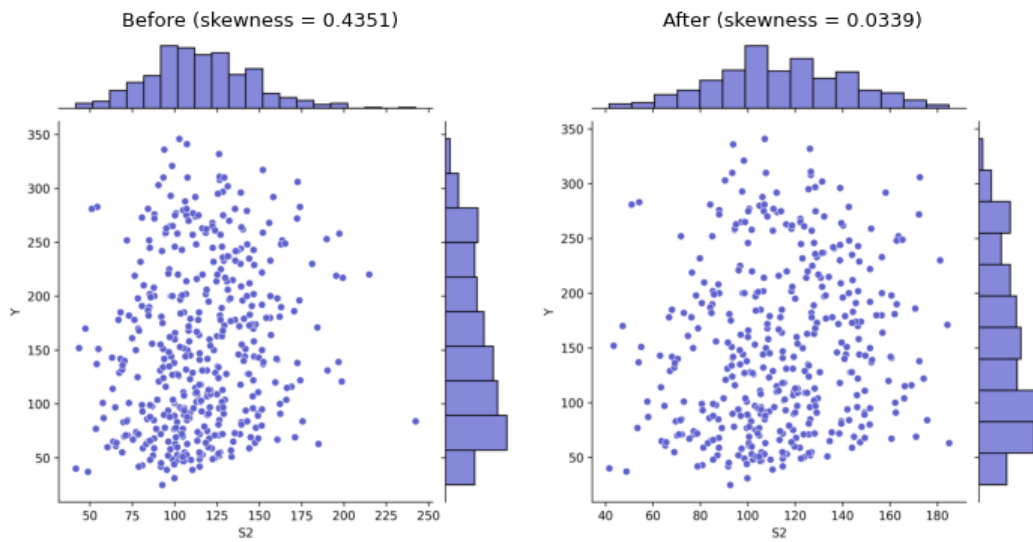
BP Outlier Removal



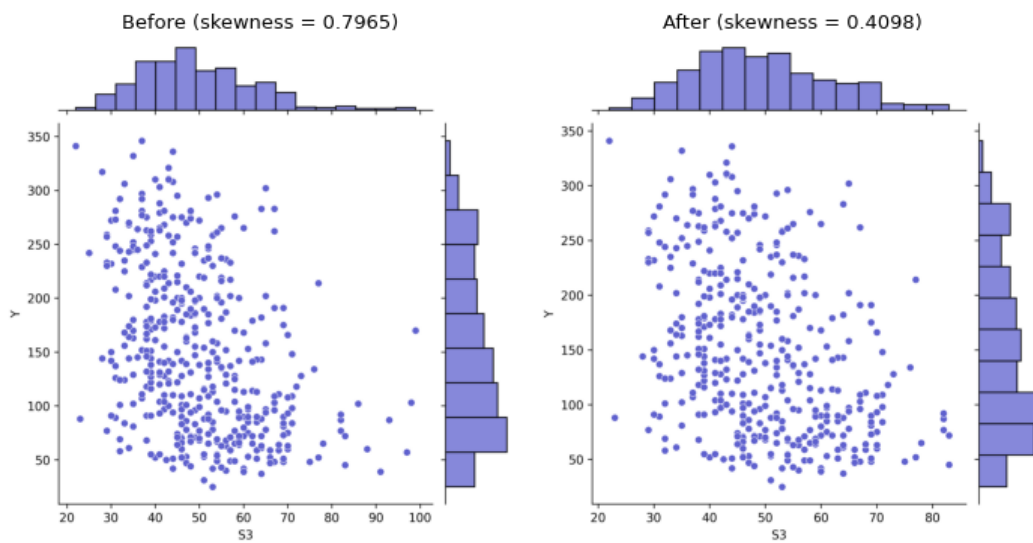
S1 Outlier Removal



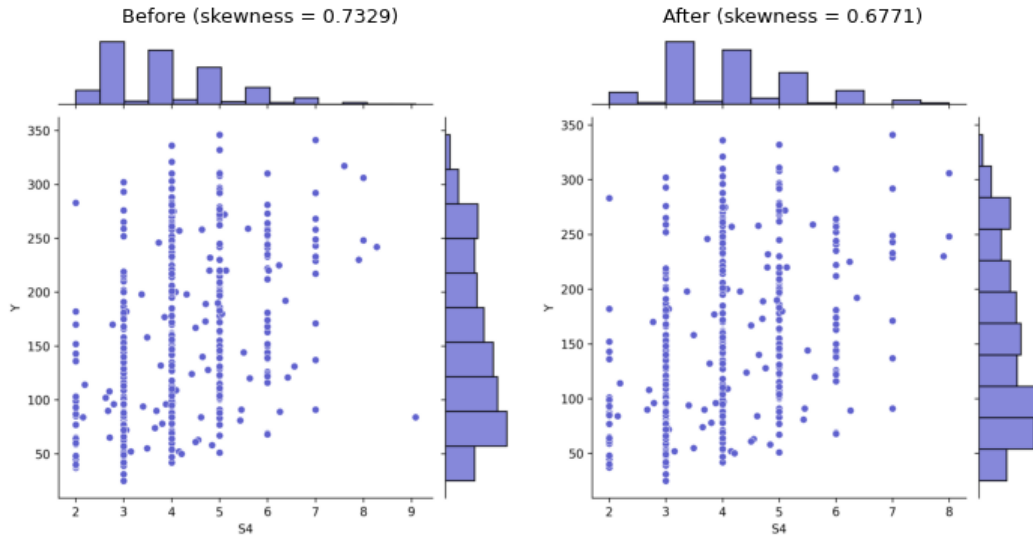
S2 Outlier Removal



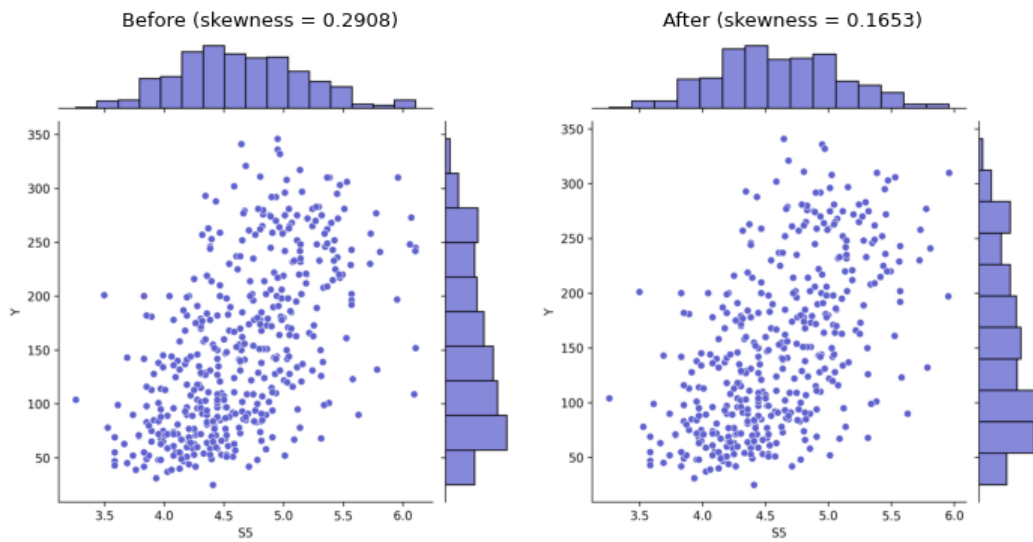
S3 Outlier Removal



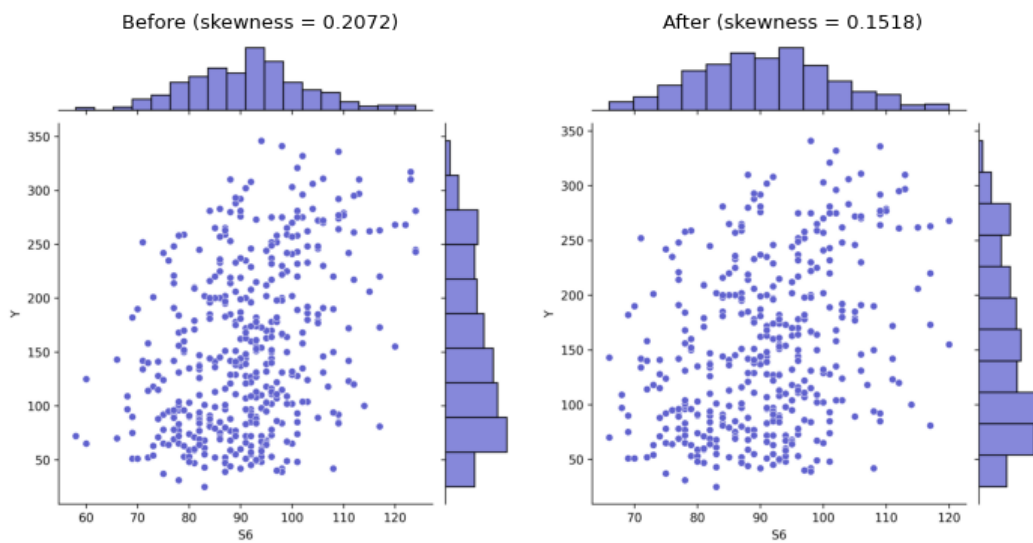
S4 Outlier Removal



S5 Outlier Removal



S6 Outlier Removal



Phân tích ảnh hưởng, tương tác của các yếu tố và xây dựng mô hình hồi quy (R)

1. Trên bộ dữ liệu trước khi xử lý

```
#library
library(kernlab)
library(caret)
```

```
## Loading required package: lattice
```

```
## Loading required package: ggplot2
```

```
##
## Attaching package: 'ggplot2'
```

```
## The following object is masked from 'package:kernlab':
##
##      alpha
```

```
library(glmnet)
```

```
## Loading required package: Matrix
```

```
## Loaded glmnet 4.1-2
```

```
library(moments)
library(knitr)

# Doc du lieu
df <- read.csv('diabetes.tab.tsv', header=TRUE, sep = '\t')

#train-test
set.seed(123)
train.index <- createDataPartition(df$Y, p = .8, list = FALSE)
train <- df[ train.index, ]
test  <- df[ -train.index, ]

x.train <- train[, 1:10]
x.test  <- test[, 1:10]
y.train <- train[, 11]
y.test  <- test[, 11]

RMSE = function(m, o){
  sqrt(mean((m - o)^2))
}
```

```

}

# Effect of Factors no interactions
av <- aov(Y~.,data=train)
summary(av)

```

```

##              Df Sum Sq Mean Sq F value    Pr(>F)
## AGE           1   85855    85855  28.653 1.58e-07 ***
## SEX           1     252      252   0.084   0.772
## BMI           1  591812  591812 197.514 < 2e-16 ***
## BP            1  104605  104605  34.911 8.30e-09 ***
## S1            1    5132     5132   1.713   0.191
## S2            1    4662     4662   1.556   0.213
## S3            1  173465  173465  57.893 2.68e-13 ***
## S4            1    1210     1210   0.404   0.526
## S5            1   50637   50637  16.900 4.93e-05 ***
## S6            1    4458     4458   1.488   0.223
## Residuals    344 1030730    2996
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

# Effect of Factor with interaction
av <- aov(Y~(AGE+BMI+BP+S1+S2+S3+S5+S6)*S4,data=train)
summary(av)

```

```

##              Df Sum Sq Mean Sq F value    Pr(>F)
## AGE           1   85855    85855  27.968 2.22e-07 ***
## BMI           1  591789  591789 192.779 < 2e-16 ***
## BP            1   98099   98099  31.956 3.36e-08 ***
## S1            1    5402     5402   1.760 0.185547
## S2            1    7923     7923   2.581 0.109082
## S3            1  147757  147757  48.133 2.05e-11 ***
## S5            1   47191   47191  15.373 0.000107 ***
## S6            1    3657     3657   1.191 0.275845
## S4            1     599      599   0.195 0.659068
## AGE:S4        1    5122     5122   1.669 0.197321
## BMI:S4        1   10028   10028   3.267 0.071591 .
## BP:S4         1     169      169   0.055 0.814705
## S1:S4         1    1218     1218   0.397 0.529180
## S2:S4         1     386      386   0.126 0.723111
## S3:S4         1     182      182   0.059 0.807668
## S5:S4         1    3702     3702   1.206 0.272916
## S6:S4         1    9221     9221   3.004 0.083984 .
## Residuals    337 1034516    3070
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

av <- aov(Y~(AGE+BMI+BP+S1+S2+S3+S5+S6)*SEX,data=train)
summary(av)

```

```

##              Df Sum Sq Mean Sq F value    Pr(>F)
## AGE           1   85855    85855  29.032 1.34e-07 ***
## BMI           1  591789  591789 200.115 < 2e-16 ***

```

```
## BP          1  98099   98099  33.172 1.90e-08 ***
## S1          1   5402    5402   1.827 0.177416
## S2          1   7923    7923   2.679 0.102591
## S3          1 147757 147757  49.965 9.06e-12 ***
## S5          1  47191   47191  15.958 7.96e-05 ***
## S6          1   3657    3657   1.237 0.266911
## SEX         1  33132   33132  11.204 0.000909 ***
## AGE:SEX     1  16983   16983   5.743 0.017101 *
## BMI:SEX     1   9617    9617   3.252 0.072228 .
## BP:SEX      1   1313    1313   0.444 0.505624
## S1:SEX      1    106     106   0.036 0.850020
## S2:SEX      1   6128    6128   2.072 0.150940
## S3:SEX      1     74     74   0.025 0.874516
## S5:SEX      1    776     776   0.262 0.608866
## S6:SEX      1    425     425   0.144 0.704953
## Residuals   337 996590   2957
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
av <- aov(Y~AGE*BMI*BP*S1*S2*S3*S5*S6+AGE*SEX,data=train)
summary(av)
```

```
##              Df Sum Sq Mean Sq F value    Pr(>F)
## AGE              1  85855   85855  28.663 5.59e-07 ***
## BMI              1 591789  591789 197.571 < 2e-16 ***
## BP              1  98099   98099  32.751 1.12e-07 ***
## S1              1   5402    5402   1.804 0.182355
## S2              1   7923    7923   2.645 0.107038
## S3              1 147757 147757  49.329 2.77e-10 ***
## S5              1  47191   47191  15.755 0.000137 ***
## S6              1   3657    3657   1.221 0.271858
## SEX             1  33132   33132  11.061 0.001238 **
## AGE:BMI         1   9377    9377   3.131 0.079915 .
## AGE:BP          1  11912   11912   3.977 0.048883 *
## BMI:BP          1  21859   21859   7.298 0.008123 **
## AGE:S1          1      2      2    0.001 0.979064
## BMI:S1          1  10422   10422   3.479 0.065098 .
## BP:S1           1   208     208   0.069 0.792619
## AGE:S2          1   8377    8377   2.797 0.097623 .
## BMI:S2          1   523     523   0.175 0.676877
## BP:S2           1    85     85   0.028 0.866446
## S1:S2           1   469     469   0.157 0.693186
## AGE:S3          1   1185    1185   0.396 0.530743
## BMI:S3          1    54     54   0.018 0.893198
## BP:S3           1   2814    2814   0.939 0.334808
## S1:S3           1   1916    1916   0.640 0.425734
## S2:S3           1   3100    3100   1.035 0.311488
## AGE:S5          1   7201    7201   2.404 0.124219
## BMI:S5          1   2705    2705   0.903 0.344256
## BP:S5           1   1357    1357   0.453 0.502542
## S1:S5           1      1      1    0.000 0.984883
## S2:S5           1   3635    3635   1.214 0.273279
## S3:S5           1   1596    1596   0.533 0.467191
## AGE:S6          1   3816    3816   1.274 0.261765
## BMI:S6          1    890     890   0.297 0.586833
```

## BP:S6	1	1259	1259	0.420	0.518276	
## S1:S6	1	895	895	0.299	0.585949	
## S2:S6	1	337	337	0.112	0.738034	
## S3:S6	1	5373	5373	1.794	0.183538	
## S5:S6	1	704	704	0.235	0.628842	
## AGE:SEX	1	18271	18271	6.100	0.015230	*
## AGE:BMI:BP	1	5803	5803	1.937	0.167085	
## AGE:BMI:S1	1	968	968	0.323	0.570925	
## AGE:BP:S1	1	583	583	0.195	0.660098	
## BMI:BP:S1	1	1299	1299	0.434	0.511745	
## AGE:BMI:S2	1	1138	1138	0.380	0.538982	
## AGE:BP:S2	1	7179	7179	2.397	0.124789	
## BMI:BP:S2	1	1414	1414	0.472	0.493618	
## AGE:S1:S2	1	813	813	0.271	0.603555	
## BMI:S1:S2	1	4960	4960	1.656	0.201177	
## BP:S1:S2	1	124	124	0.041	0.839153	
## AGE:BMI:S3	1	375	375	0.125	0.724075	
## AGE:BP:S3	1	1591	1591	0.531	0.467810	
## BMI:BP:S3	1	112	112	0.037	0.847059	
## AGE:S1:S3	1	4336	4336	1.448	0.231792	
## BMI:S1:S3	1	3601	3601	1.202	0.275545	
## BP:S1:S3	1	3811	3811	1.272	0.262034	
## AGE:S2:S3	1	343	343	0.114	0.735880	
## BMI:S2:S3	1	1290	1290	0.431	0.513204	
## BP:S2:S3	1	122	122	0.041	0.840332	
## S1:S2:S3	1	2427	2427	0.810	0.370271	
## AGE:BMI:S5	1	3688	3688	1.231	0.269844	
## AGE:BP:S5	1	11160	11160	3.726	0.056441	.
## BMI:BP:S5	1	1309	1309	0.437	0.510087	
## AGE:S1:S5	1	7515	7515	2.509	0.116400	
## BMI:S1:S5	1	23173	23173	7.737	0.006480	**
## BP:S1:S5	1	3932	3932	1.313	0.254694	
## AGE:S2:S5	1	416	416	0.139	0.710188	
## BMI:S2:S5	1	11092	11092	3.703	0.057179	.
## BP:S2:S5	1	92	92	0.031	0.861310	
## S1:S2:S5	1	9	9	0.003	0.957614	
## AGE:S3:S5	1	13	13	0.004	0.948480	
## BMI:S3:S5	1	3067	3067	1.024	0.314091	
## BP:S3:S5	1	7129	7129	2.380	0.126074	
## S1:S3:S5	1	1311	1311	0.438	0.509794	
## S2:S3:S5	1	91	91	0.030	0.861746	
## AGE:BMI:S6	1	20	20	0.007	0.934821	
## AGE:BP:S6	1	758	758	0.253	0.615936	
## BMI:BP:S6	1	1	1	0.000	0.988486	
## AGE:S1:S6	1	7772	7772	2.595	0.110397	
## BMI:S1:S6	1	128	128	0.043	0.836704	
## BP:S1:S6	1	2193	2193	0.732	0.394283	
## AGE:S2:S6	1	217	217	0.073	0.788285	
## BMI:S2:S6	1	0	0	0.000	0.998612	
## BP:S2:S6	1	207	207	0.069	0.793402	
## S1:S2:S6	1	1446	1446	0.483	0.488765	
## AGE:S3:S6	1	246	246	0.082	0.775180	
## BMI:S3:S6	1	6598	6598	2.203	0.140947	
## BP:S3:S6	1	9932	9932	3.316	0.071631	.
## S1:S3:S6	1	547	547	0.183	0.670023	
## S2:S3:S6	1	3286	3286	1.097	0.297486	
## AGE:S5:S6	1	5080	5080	1.696	0.195853	

## BMI:S5:S6	1	327	327	0.109	0.741668	
## BP:S5:S6	1	224	224	0.075	0.785122	
## S1:S5:S6	1	1429	1429	0.477	0.491414	
## S2:S5:S6	1	5046	5046	1.685	0.197343	
## S3:S5:S6	1	25	25	0.008	0.927733	
## AGE:BMI:BP:S1	1	174	174	0.058	0.810231	
## AGE:BMI:BP:S2	1	4509	4509	1.506	0.222736	
## AGE:BMI:S1:S2	1	56	56	0.019	0.891595	
## AGE:BP:S1:S2	1	452	452	0.151	0.698440	
## BMI:BP:S1:S2	1	202	202	0.067	0.795564	
## AGE:BMI:BP:S3	1	694	694	0.232	0.631409	
## AGE:BMI:S1:S3	1	3934	3934	1.313	0.254560	
## AGE:BP:S1:S3	1	459	459	0.153	0.696184	
## BMI:BP:S1:S3	1	3389	3389	1.131	0.290048	
## AGE:BMI:S2:S3	1	473	473	0.158	0.691930	
## AGE:BP:S2:S3	1	8922	8922	2.979	0.087493	.
## BMI:BP:S2:S3	1	299	299	0.100	0.752688	
## AGE:S1:S2:S3	1	4464	4464	1.490	0.225069	
## BMI:S1:S2:S3	1	8022	8022	2.678	0.104910	
## BP:S1:S2:S3	1	1029	1029	0.344	0.559078	
## AGE:BMI:BP:S5	1	210	210	0.070	0.791508	
## AGE:BMI:S1:S5	1	2977	2977	0.994	0.321222	
## AGE:BP:S1:S5	1	2802	2802	0.936	0.335790	
## BMI:BP:S1:S5	1	3643	3643	1.216	0.272745	
## AGE:BMI:S2:S5	1	96	96	0.032	0.858016	
## AGE:BP:S2:S5	1	4189	4189	1.399	0.239789	
## BMI:BP:S2:S5	1	1560	1560	0.521	0.472130	
## AGE:S1:S2:S5	1	3633	3633	1.213	0.273459	
## BMI:S1:S2:S5	1	3148	3148	1.051	0.307754	
## BP:S1:S2:S5	1	424	424	0.141	0.707683	
## AGE:BMI:S3:S5	1	268	268	0.089	0.765510	
## AGE:BP:S3:S5	1	645	645	0.215	0.643662	
## BMI:BP:S3:S5	1	3993	3993	1.333	0.251055	
## AGE:S1:S3:S5	1	3683	3683	1.230	0.270164	
## BMI:S1:S3:S5	1	51	51	0.017	0.896038	
## BP:S1:S3:S5	1	108	108	0.036	0.850044	
## AGE:S2:S3:S5	1	4179	4179	1.395	0.240363	
## BMI:S2:S3:S5	1	7586	7586	2.533	0.114705	
## BP:S2:S3:S5	1	6304	6304	2.105	0.150023	
## S1:S2:S3:S5	1	12891	12891	4.304	0.040624	*
## AGE:BMI:BP:S6	1	4623	4623	1.544	0.217032	
## AGE:BMI:S1:S6	1	8599	8599	2.871	0.093337	.
## AGE:BP:S1:S6	1	410	410	0.137	0.712268	
## BMI:BP:S1:S6	1	2995	2995	1.000	0.319810	
## AGE:BMI:S2:S6	1	180	180	0.060	0.806952	
## AGE:BP:S2:S6	1	34	34	0.011	0.915741	
## BMI:BP:S2:S6	1	9743	9743	3.253	0.074342	.
## AGE:S1:S2:S6	1	3366	3366	1.124	0.291665	
## BMI:S1:S2:S6	1	1383	1383	0.462	0.498411	
## BP:S1:S2:S6	1	22	22	0.007	0.931540	
## AGE:BMI:S3:S6	1	201	201	0.067	0.796198	
## AGE:BP:S3:S6	1	4877	4877	1.628	0.204948	
## BMI:BP:S3:S6	1	6383	6383	2.131	0.147528	
## AGE:S1:S3:S6	1	148	148	0.050	0.824281	
## BMI:S1:S3:S6	1	1352	1352	0.451	0.503319	
## BP:S1:S3:S6	1	10654	10654	3.557	0.062231	.
## AGE:S2:S3:S6	1	8005	8005	2.672	0.105275	

## BMI:S2:S3:S6	1	3519	3519	1.175	0.281033	
## BP:S2:S3:S6	1	1496	1496	0.499	0.481444	
## S1:S2:S3:S6	1	1997	1997	0.667	0.416119	
## AGE:BMI:S5:S6	1	467	467	0.156	0.693922	
## AGE:BP:S5:S6	1	2167	2167	0.724	0.397036	
## BMI:BP:S5:S6	1	25	25	0.008	0.927013	
## AGE:S1:S5:S6	1	12395	12395	4.138	0.044600	*
## BMI:S1:S5:S6	1	2869	2869	0.958	0.330102	
## BP:S1:S5:S6	1	12000	12000	4.006	0.048071	*
## AGE:S2:S5:S6	1	13	13	0.004	0.948533	
## BMI:S2:S5:S6	1	5392	5392	1.800	0.182764	
## BP:S2:S5:S6	1	1925	1925	0.643	0.424635	
## S1:S2:S5:S6	1	4975	4975	1.661	0.200494	
## AGE:S3:S5:S6	1	52	52	0.017	0.895350	
## BMI:S3:S5:S6	1	657	657	0.219	0.640482	
## BP:S3:S5:S6	1	1263	1263	0.422	0.517577	
## S1:S3:S5:S6	1	2787	2787	0.930	0.337095	
## S2:S3:S5:S6	1	3948	3948	1.318	0.253713	
## AGE:BMI:BP:S1:S2	1	12	12	0.004	0.948950	
## AGE:BMI:BP:S1:S3	1	3183	3183	1.063	0.305121	
## AGE:BMI:BP:S2:S3	1	4269	4269	1.425	0.235386	
## AGE:BMI:S1:S2:S3	1	88	88	0.029	0.864395	
## AGE:BP:S1:S2:S3	1	10	10	0.003	0.955105	
## BMI:BP:S1:S2:S3	1	438	438	0.146	0.702982	
## AGE:BMI:BP:S1:S5	1	915	915	0.305	0.581777	
## AGE:BMI:BP:S2:S5	1	2448	2448	0.817	0.368182	
## AGE:BMI:S1:S2:S5	1	32	32	0.011	0.917599	
## AGE:BP:S1:S2:S5	1	1250	1250	0.417	0.519768	
## BMI:BP:S1:S2:S5	1	3417	3417	1.141	0.288086	
## AGE:BMI:BP:S3:S5	1	2949	2949	0.985	0.323483	
## AGE:BMI:S1:S3:S5	1	1469	1469	0.490	0.485388	
## AGE:BP:S1:S3:S5	1	2293	2293	0.766	0.383707	
## BMI:BP:S1:S3:S5	1	1549	1549	0.517	0.473692	
## AGE:BMI:S2:S3:S5	1	4066	4066	1.358	0.246754	
## AGE:BP:S2:S3:S5	1	1	1	0.000	0.986737	
## BMI:BP:S2:S3:S5	1	2910	2910	0.972	0.326703	
## AGE:S1:S2:S3:S5	1	9545	9545	3.186	0.077312	.
## BMI:S1:S2:S3:S5	1	14151	14151	4.724	0.032122	*
## BP:S1:S2:S3:S5	1	351	351	0.117	0.732986	
## AGE:BMI:BP:S1:S6	1	8841	8841	2.952	0.088920	.
## AGE:BMI:BP:S2:S6	1	364	364	0.122	0.728126	
## AGE:BMI:S1:S2:S6	1	2743	2743	0.916	0.340930	
## AGE:BP:S1:S2:S6	1	6501	6501	2.170	0.143859	
## BMI:BP:S1:S2:S6	1	60	60	0.020	0.888023	
## AGE:BMI:BP:S3:S6	1	9216	9216	3.077	0.082508	.
## AGE:BMI:S1:S3:S6	1	134	134	0.045	0.832823	
## AGE:BP:S1:S3:S6	1	0	0	0.000	0.994720	
## BMI:BP:S1:S3:S6	1	6837	6837	2.282	0.134035	
## AGE:BMI:S2:S3:S6	1	6121	6121	2.044	0.156002	
## AGE:BP:S2:S3:S6	1	367	367	0.123	0.727079	
## BMI:BP:S2:S3:S6	1	2170	2170	0.725	0.396713	
## AGE:S1:S2:S3:S6	1	4793	4793	1.600	0.208860	
## BMI:S1:S2:S3:S6	1	2374	2374	0.793	0.375477	
## BP:S1:S2:S3:S6	1	2223	2223	0.742	0.391102	
## AGE:BMI:BP:S5:S6	1	1971	1971	0.658	0.419194	
## AGE:BMI:S1:S5:S6	1	2908	2908	0.971	0.326882	
## AGE:BP:S1:S5:S6	1	1285	1285	0.429	0.514077	

## BMI:BP:S1:S5:S6	1	8467	8467	2.827	0.095862	.
## AGE:BMI:S2:S5:S6	1	90	90	0.030	0.862521	
## AGE:BP:S2:S5:S6	1	3310	3310	1.105	0.295715	
## BMI:BP:S2:S5:S6	1	3289	3289	1.098	0.297272	
## AGE:S1:S2:S5:S6	1	2275	2275	0.759	0.385606	
## BMI:S1:S2:S5:S6	1	1116	1116	0.373	0.543011	
## BP:S1:S2:S5:S6	1	1984	1984	0.662	0.417714	
## AGE:BMI:S3:S5:S6	1	5573	5573	1.860	0.175662	
## AGE:BP:S3:S5:S6	1	14094	14094	4.705	0.032465	*
## BMI:BP:S3:S5:S6	1	471	471	0.157	0.692654	
## AGE:S1:S3:S5:S6	1	1977	1977	0.660	0.418520	
## BMI:S1:S3:S5:S6	1	3149	3149	1.051	0.307712	
## BP:S1:S3:S5:S6	1	216	216	0.072	0.788850	
## AGE:S2:S3:S5:S6	1	4627	4627	1.545	0.216834	
## BMI:S2:S3:S5:S6	1	2493	2493	0.832	0.363782	
## BP:S2:S3:S5:S6	1	125	125	0.042	0.838393	
## S1:S2:S3:S5:S6	1	75	75	0.025	0.874342	
## AGE:BMI:BP:S1:S2:S3	1	3923	3923	1.310	0.255236	
## AGE:BMI:BP:S1:S2:S5	1	655	655	0.219	0.640987	
## AGE:BMI:BP:S1:S3:S5	1	870	870	0.290	0.591185	
## AGE:BMI:BP:S2:S3:S5	1	3895	3895	1.300	0.256897	
## AGE:BMI:S1:S2:S3:S5	1	3525	3525	1.177	0.280623	
## AGE:BP:S1:S2:S3:S5	1	6	6	0.002	0.965009	
## BMI:BP:S1:S2:S3:S5	1	3021	3021	1.008	0.317730	
## AGE:BMI:BP:S1:S2:S6	1	850	850	0.284	0.595373	
## AGE:BMI:BP:S1:S3:S6	1	1430	1430	0.477	0.491196	
## AGE:BMI:BP:S2:S3:S6	1	850	850	0.284	0.595404	
## AGE:BMI:S1:S2:S3:S6	1	151	151	0.050	0.822883	
## AGE:BP:S1:S2:S3:S6	1	2366	2366	0.790	0.376257	
## BMI:BP:S1:S2:S3:S6	1	1426	1426	0.476	0.491756	
## AGE:BMI:BP:S1:S5:S6	1	8223	8223	2.745	0.100708	
## AGE:BMI:BP:S2:S5:S6	1	313	313	0.104	0.747254	
## AGE:BMI:S1:S2:S5:S6	1	1794	1794	0.599	0.440781	
## AGE:BP:S1:S2:S5:S6	1	1286	1286	0.429	0.513875	
## BMI:BP:S1:S2:S5:S6	1	1020	1020	0.341	0.560784	
## AGE:BMI:BP:S3:S5:S6	1	4	4	0.001	0.970736	
## AGE:BMI:S1:S3:S5:S6	1	1828	1828	0.610	0.436524	
## AGE:BP:S1:S3:S5:S6	1	125	125	0.042	0.838408	
## BMI:BP:S1:S3:S5:S6	1	4108	4108	1.372	0.244343	
## AGE:BMI:S2:S3:S5:S6	1	157	157	0.052	0.819297	
## AGE:BP:S2:S3:S5:S6	1	1029	1029	0.343	0.559195	
## BMI:BP:S2:S3:S5:S6	1	612	612	0.204	0.652249	
## AGE:S1:S2:S3:S5:S6	1	844	844	0.282	0.596789	
## BMI:S1:S2:S3:S5:S6	1	1024	1024	0.342	0.560139	
## BP:S1:S2:S3:S5:S6	1	4609	4609	1.539	0.217761	
## AGE:BMI:BP:S1:S2:S3:S5	1	1048	1048	0.350	0.555534	
## AGE:BMI:BP:S1:S2:S3:S6	1	1086	1086	0.362	0.548547	
## AGE:BMI:BP:S1:S2:S5:S6	1	31	31	0.010	0.918828	
## AGE:BMI:BP:S1:S3:S5:S6	1	723	723	0.241	0.624329	
## AGE:BMI:S1:S2:S3:S5:S6	1	1466	1466	0.490	0.485757	
## AGE:BP:S1:S2:S3:S5:S6	1	7632	7632	2.548	0.113632	
## BMI:BP:S1:S2:S3:S5:S6	1	3459	3459	1.155	0.285175	
## Residuals	99	296537	2995			
## ---						
## Signif. codes:	0	'***'	0.001	'**'	0.01	'*' 0.05
					'.'	0.1 ' ' 1

```
#Build Models
#SLR
#train
fit <- lm(formula = Y~BMI,data=train)
y.train.pred <- predict(fit,newdata=x.train)
summary(fit)
```

```
##
## Call:
## lm(formula = Y ~ BMI, data = train)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -162.809  -43.569   -7.261   48.156  152.338
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -107.096     20.376  -5.256 2.55e-07 ***
## BMI           9.846       0.764  12.887 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 62.89 on 353 degrees of freedom
## Multiple R-squared:  0.3199, Adjusted R-squared:  0.318
## F-statistic: 166.1 on 1 and 353 DF,  p-value: < 2.2e-16
```

```
RMSE(y.train.pred,y.train) #62.7097
```

```
## [1] 62.7097
```

```
#test
y.test.pred <- predict(fit,newdata=x.test)
predict <- lm(y.test~y.test.pred)
summary(predict)
```

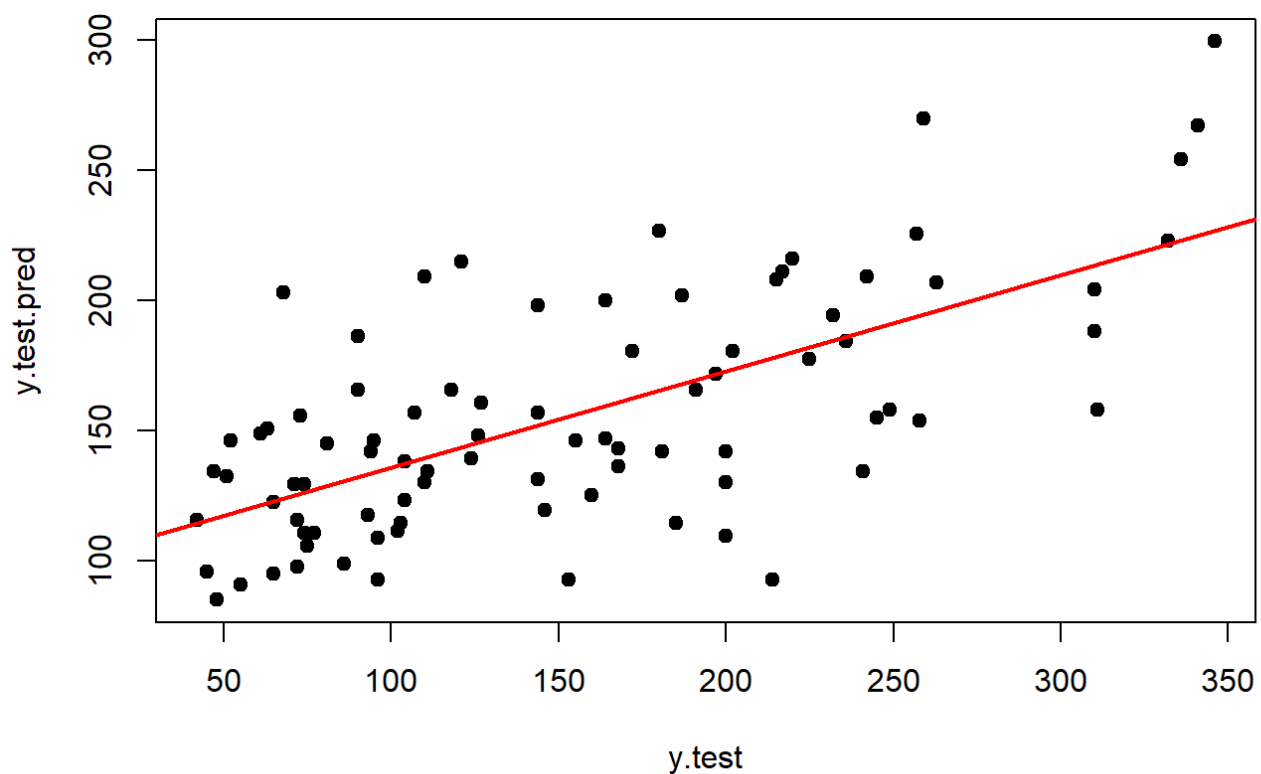
```
##
## Call:
## lm(formula = y.test ~ y.test.pred)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -141.74  -42.70  -10.73   36.79  155.10
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -31.6173     23.5179  -1.344   0.182
## y.test.pred   1.1887      0.1455   8.172 2.55e-12 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 61.18 on 85 degrees of freedom
```

```
## Multiple R-squared:  0.44, Adjusted R-squared:  0.4334
## F-statistic: 66.78 on 1 and 85 DF,  p-value: 2.551e-12
```

```
RMSE(y.test.pred,y.test) #61.11294
```

```
## [1] 61.11294
```

```
plot(y.test,y.test.pred, pch = 19, cex = 1, col = "black")
abline(lm(y.test.pred~y.test),col='red',lwd=2)
```



```
#multiple linear regression
#train
fit <- lm(formula = Y~SEX+BMI+BP+S1+S2+S5+S6,data=train)
y.train.pred <- predict(fit,newdata=x.train)
summary(fit)
```

```
##
## Call:
## lm(formula = Y ~ SEX + BMI + BP + S1 + S2 + S5 + S6, data = train)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -152.851  -38.319   -0.824   36.942  149.420
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
##
```

```
## (Intercept) -320.7058      29.6318 -10.823 < 2e-16 ***
## SEX          -21.9590       6.5182  -3.369  0.00084 ***
## BMI           4.9575       0.8188   6.055  3.65e-09 ***
## BP            1.0507       0.2497   4.208  3.29e-05 ***
## S1           -1.0232       0.2523  -4.055  6.19e-05 ***
## S2            0.8703       0.2622   3.319  0.00100 **
## S5           71.9213       8.7834   8.188  5.12e-15 ***
## S6            0.3715       0.2944   1.262  0.20784
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 54.54 on 347 degrees of freedom
## Multiple R-squared:  0.4972, Adjusted R-squared:  0.4871
## F-statistic: 49.03 on 7 and 347 DF,  p-value: < 2.2e-16
```

```
RMSE(y.train.pred,y.train) #53.91863
```

```
## [1] 53.91863
```

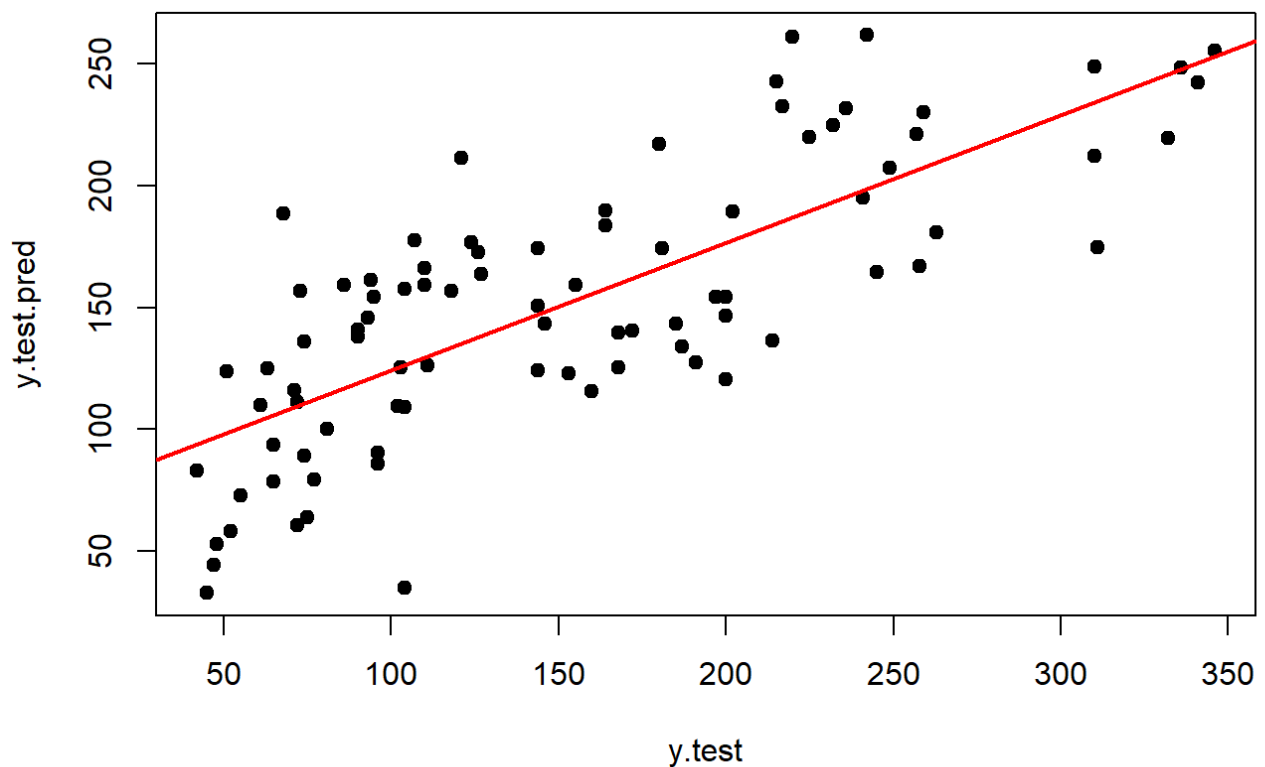
```
#test
y.test.pred <- predict(fit,newdata=x.test)
predict <- lm(y.test~y.test.pred)
summary(predict)
```

```
##
## Call:
## lm(formula = y.test ~ y.test.pred)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -125.766  -41.293   -3.789   36.917  132.700
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  -16.311     16.485  -0.989   0.325
## y.test.pred    1.115      0.102  10.929 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 52.71 on 85 degrees of freedom
## Multiple R-squared:  0.5843, Adjusted R-squared:  0.5794
## F-statistic: 119.5 on 1 and 85 DF,  p-value: < 2.2e-16
```

```
RMSE(y.test.pred,y.test) #52.50263
```

```
## [1] 52.50263
```

```
plot(y.test,y.test.pred, pch = 19, cex = 1, col = "black")
abline(lm(y.test.pred~y.test),col='red',lwd=2)
```



```
#polynomial regression
#train
fit <- lm(formula = Y~AGE+BP+BMI+S3+SEX +
            I (AGE^2)+I (S5^2)+I (S3^2)+
            I (AGE*SEX) +
            I (BMI*BP) + I (S1*S2*S3*S5)+
            I (BMI*S1*S5) + I (BMI*S2*S5)+
            I (AGE*S2*S3*S6)+
            I (BMI*S1*S2*S3*S5) + I (AGE*BP*S3*S5*S6),
            data=train)
y.train.pred <- predict(fit,newdata=x.train)
summary(fit)
```

```
##
## Call:
## lm(formula = Y ~ AGE + BP + BMI + S3 + SEX + I(AGE^2) + I(S5^2) +
##     I(S3^2) + I(AGE * SEX) + I(BMI * BP) + I(S1 * S2 * S3 * S5) +
##     I(BMI * S1 * S5) + I(BMI * S2 * S5) + I(AGE * S2 * S3 * S6) +
##     I(BMI * S1 * S2 * S3 * S5) + I(AGE * BP * S3 * S5 * S6),
##     data = train)
##
## Residuals:
```

	Min	1Q	Median	3Q	Max
	-141.774	-37.239	-4.278	34.056	142.793

```
##
## Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	5.822e+02	1.603e+02	3.631	0.000326 ***

```
## AGE -4.194e+00 1.497e+00 -2.802 0.005378 **
## BP -3.800e+00 1.333e+00 -2.851 0.004630 **
## BMI -9.096e+00 4.660e+00 -1.952 0.051797 .
## S3 -4.029e+00 1.788e+00 -2.254 0.024850 *
## SEX -7.898e+01 2.348e+01 -3.363 0.000858 ***
## I (AGE^2) 1.928e-02 1.490e-02 1.294 0.196462
## I (S5^2) 6.541e+00 2.762e+00 2.368 0.018431 *
## I (S3^2) 2.170e-02 1.296e-02 1.674 0.095126 .
## I (AGE * SEX) 1.103e+00 4.586e-01 2.405 0.016708 *
## I (BMI * BP) 1.473e-01 4.599e-02 3.203 0.001491 **
## I (S1 * S2 * S3 * S5) -1.270e-05 9.678e-06 -1.312 0.190352
## I (BMI * S1 * S5) -8.057e-03 3.735e-03 -2.157 0.031720 *
## I (BMI * S2 * S5) 7.455e-03 4.012e-03 1.858 0.064044 .
## I (AGE * S2 * S3 * S6) -1.541e-06 1.210e-06 -1.273 0.203986
## I (BMI * S1 * S2 * S3 * S5) 6.639e-07 3.585e-07 1.852 0.064908 .
## I (AGE * BP * S3 * S5 * S6) 8.688e-07 3.069e-07 2.831 0.004916 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 52.69 on 338 degrees of freedom
## Multiple R-squared: 0.543, Adjusted R-squared: 0.5213
## F-statistic: 25.1 on 16 and 338 DF, p-value: < 2.2e-16
```

```
RMSE(y.train.pred,y.train) #51.4093
```

```
## [1] 51.4093
```

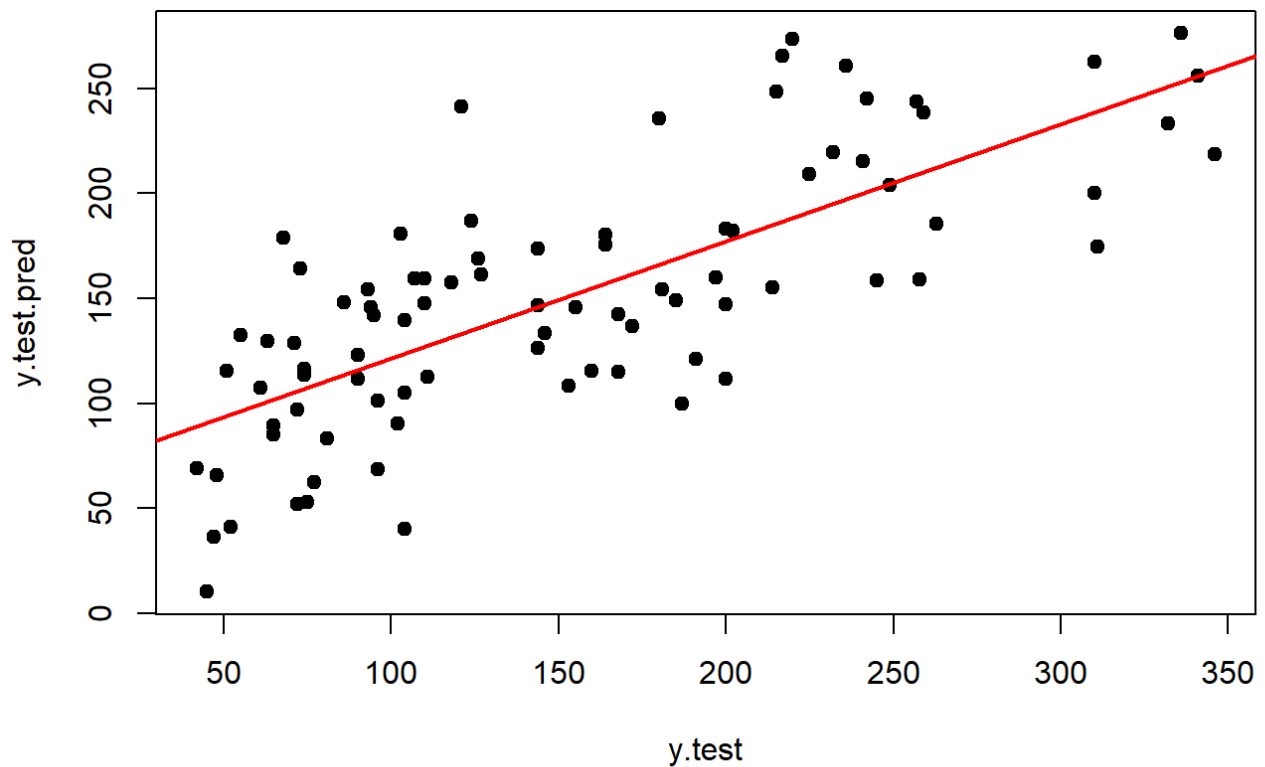
```
#test
y.test.pred <- predict(fit,newdata=x.test)
predict <- lm(y.test~y.test.pred)
summary(predict)
```

```
##
## Call:
## lm(formula = y.test ~ y.test.pred)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -122.067  -41.691   -3.283   33.429  134.537
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  2.83478    15.70374   0.181   0.857
## y.test.pred  0.99529     0.09664  10.299 <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 54.53 on 85 degrees of freedom
## Multiple R-squared: 0.5551, Adjusted R-squared: 0.5499
## F-statistic: 106.1 on 1 and 85 DF, p-value: < 2.2e-16
```

```
RMSE(y.test.pred,y.test) #53.94088
```

```
## [1] 53.94088
```

```
plot(y.test,y.test.pred, pch = 19, cex = 1, col = "black")  
abline(lm(y.test.pred~y.test),col='red',lwd=2)
```



```
#ridge (alpha = 0)  
set.seed(123)  
ridge.fit <- cv.glmnet(as.matrix(x.train),y.train,type.measure = 'mse',alpha=0,family='gaussian')  
ridge.fit$lambda.1se
```

```
## [1] 63.87225
```

```
coef(ridge.fit)
```

```
## 11 x 1 sparse Matrix of class "dgCMatrix"  
##              s1  
## (Intercept) -148.67220469  
## AGE          0.12349837  
## SEX          -8.51972205  
## BMI          3.27742361  
## BP           0.72694871  
## S1           0.02000633  
## S2          -0.03983784  
## S3          -0.54474454
```

```
## S4          4.07225690
## S5          25.39873274
## S6          0.49832477
```

```
#train
ridge.predict.train <- predict(ridge.fit,s=ridge.fit$lambda.1se,newx = as.matri
ix(x.train))
fit <- lm(y.train~ridge.predict.train)
summary(fit)
```

```
##
## Call:
## lm(formula = y.train ~ ridge.predict.train)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -130.410  -41.102   -0.567   40.395  160.876
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   -50.35408    11.66405   -4.317 2.06e-05 ***
## ridge.predict.train    1.33142     0.07431   17.917 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 55.19 on 353 degrees of freedom
## Multiple R-squared:  0.4763, Adjusted R-squared:  0.4748
## F-statistic: 321 on 1 and 353 DF, p-value: < 2.2e-16
```

```
RMSE(ridge.predict.train,y.train) #56.56072
```

```
## [1] 56.56072
```

```
#test
ridge.predict.test <- predict(ridge.fit,s=ridge.fit$lambda.1se,newx = as.matri
x(x.test))
predict <- lm(y.test~ridge.predict.test)
summary(predict)
```

```
##
## Call:
## lm(formula = y.test ~ ridge.predict.test)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -125.346  -44.308   -3.681   39.165  123.582
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   -71.9704    23.2521   -3.095  0.00266 **
## ridge.predict.test    1.4742     0.1473   10.005 5.03e-16 ***
## ---
```

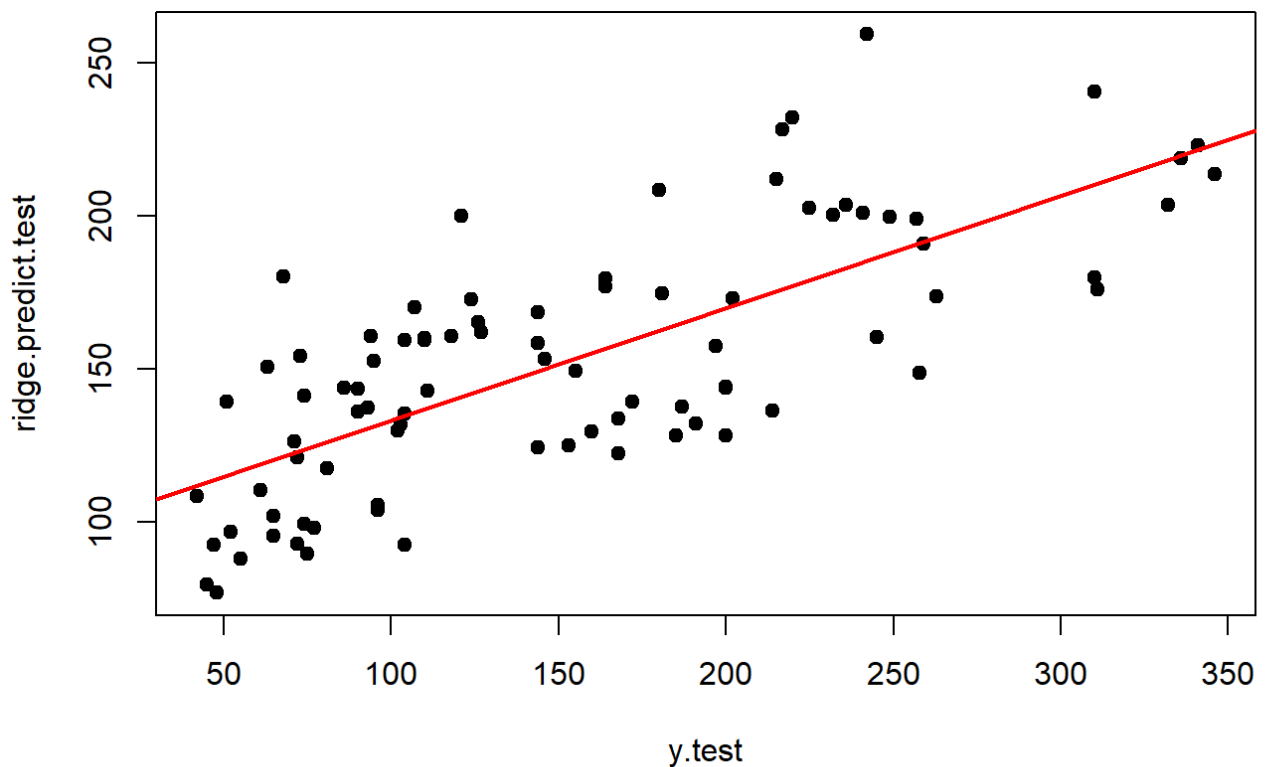


```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 55.4 on 85 degrees of freedom
## Multiple R-squared:  0.5408, Adjusted R-squared:  0.5354
## F-statistic: 100.1 on 1 and 85 DF,  p-value: 5.028e-16
```

```
RMSE(ridge.predict.test,y.test) #58.00265
```

```
## [1] 58.00265
```

```
#plot
plot(y.test,ridge.predict.test, pch = 19, cex = 1, col = "black")
abline(lm(ridge.predict.test~y.test),col='red',lwd=2)
```



```
#lasso (alpha = 1)
set.seed(123)
lasso.fit <- cv.glmnet(as.matrix(x.train),y.train,type.measure = 'mse',alpha=1,family='gaussian')
lasso.fit$lambda.1se
```

```
## [1] 8.845581
```

```
coef(lasso.fit)
```

```
## 11 x 1 sparse Matrix of class "dgCMatrix"
##              s1
## (Intercept) -198.2166275
## AGE          .
## SEX          .
## BMI          4.7320556
## BP           0.5720361
## S1           .
## S2           .
## S3          -0.2404157
## S4           .
## S5          39.4848485
## S6           .
```

```
#summary train
lasso.predict.train <- predict(lasso.fit,s=lasso.fit$lambda.1se,newx = as.matrix(x.train))
fit <- lm(y.train~lasso.predict.train)
summary(fit)
```

```
##
## Call:
## lm(formula = y.train ~ lasso.predict.train)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -139.201  -39.521   -0.881   40.409  144.993
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -42.0800     11.3590  -3.705 0.000246 ***
## lasso.predict.train  1.2770      0.0722  17.686 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 55.53 on 353 degrees of freedom
## Multiple R-squared:  0.4698, Adjusted R-squared:  0.4683
## F-statistic: 312.8 on 1 and 353 DF,  p-value: < 2.2e-16
```

```
RMSE(lasso.predict.train,y.train) #56.51291
```

```
## [1] 56.51291
```

```
#summary test
lasso.predict.test <- predict(lasso.fit,s=lasso.fit$lambda.1se,newx = as.matrix(x.test))
predict <- lm(y.test~lasso.predict.test)
summary(predict)
```

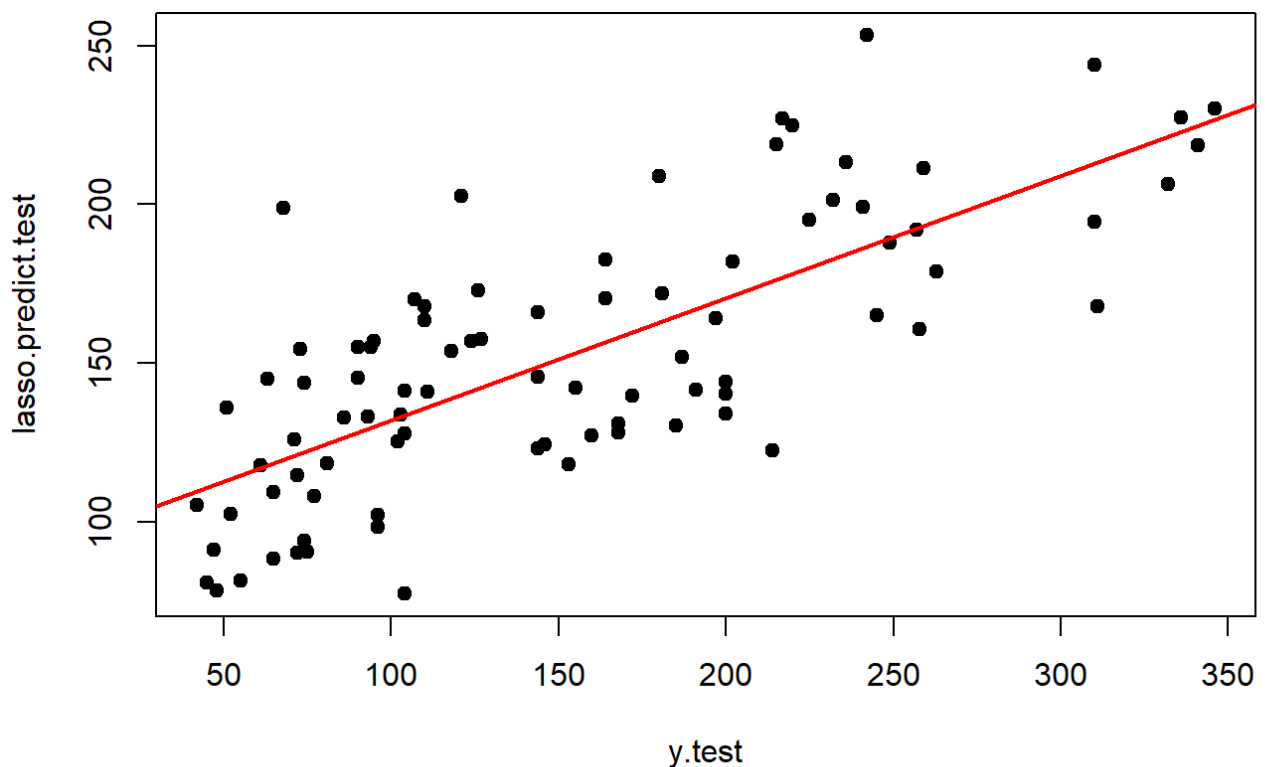
```
##
## Call:
## lm(formula = y.test ~ lasso.predict.test)
```

```
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -151.854  -36.884   -0.261   40.122  135.796
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -66.1576     22.1202  -2.991  0.00364 **
## lasso.predict.test  1.4374      0.1399  10.271  < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 54.61 on 85 degrees of freedom
## Multiple R-squared:  0.5538, Adjusted R-squared:  0.5486
## F-statistic: 105.5 on 1 and 85 DF,  p-value: < 2.2e-16
```

```
RMSE(lasso.predict.test,y.test) #56.99875
```

```
## [1] 56.99875
```

```
#plot
plot(y.test,lasso.predict.test, pch = 19, cex = 1, col = "black")
abline(lm(lasso.predict.test~y.test),col='red',lwd=2)
```



```
#elastic net
results.train <-data.frame()
for (i in 0:20)
```

```

{
  set.seed(123)
  fit <- cv.glmnet(as.matrix(x.train), y.train, type.measure="mse", alpha=i/20,
                  family="gaussian")
  y.pred <- predict(fit, s=fit$lambda.1se, newx=as.matrix(x.train))
  predict <- lm(y.train~y.pred)

  temp <- data.frame(alpha=i/20,R2= summary(predict)$r.squared,Adj_R2=summary(pr
edict)$adj.r.squared,rmse=RMSE(y.pred,y.train),lambda=fit$lambda.1se)
  results.train <- rbind(results.train, temp)
}
#alpha = 0.05 (best adj R2)
set.seed(123)
elastic.fit <- cv.glmnet(as.matrix(x.train), y.train, type.measure="mse", alpha=
0.05,
                        family="gaussian")
elastic.fit$lambda.1se

```

```
## [1] 48.09499
```

```
coef(fit)
```

```

## 11 x 1 sparse Matrix of class "dgCMatrix"
##              s1
## (Intercept) -198.2166275
## AGE          .
## SEX          .
## BMI          4.7320556
## BP           0.5720361
## S1           .
## S2           .
## S3          -0.2404157
## S4           .
## S5          39.4848485
## S6           .

```

```

#train
elastic.predict.train <- predict(elastic.fit, s=elastic.fit$lambda.1se, newx=a
s.matrix(x.train))
fit <- lm(y.train~ elastic.predict.train)
summary(fit)

```

```

##
## Call:
## lm(formula = y.train ~ elastic.predict.train)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -131.996  -41.093   -0.335   41.231  155.550
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -50.98749    11.67363   -4.368 1.65e-05 ***

```

```
## elastic.predict.train    1.33559    0.07438   17.956   < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 55.13 on 353 degrees of freedom
## Multiple R-squared:  0.4774, Adjusted R-squared:  0.4759
## F-statistic: 322.4 on 1 and 353 DF,  p-value: < 2.2e-16
```

```
RMSE(elastic.predict.train,y.train) #56.53718
```

```
## [1] 56.53718
```

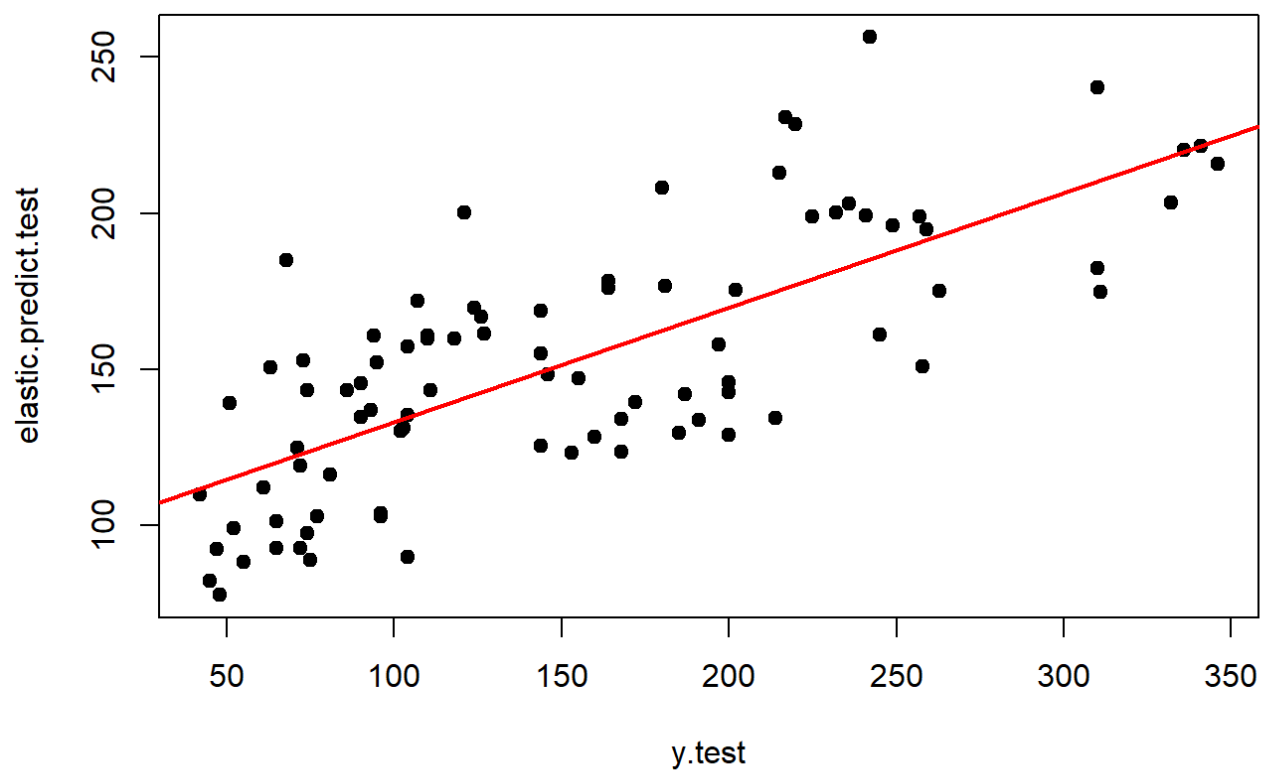
```
#test
elastic.predict.test <- predict(elastic.fit, s=elastic.fit$lambda.1se, newx=a
s.matrix(x.test))
predict <- lm(y.test~elastic.predict.test)
summary(predict)
```

```
##
## Call:
## lm(formula = y.test ~ elastic.predict.test)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -132.744  -43.152   -2.361   40.418  125.093
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -73.1442    23.1805  -3.155  0.00222 **
## elastic.predict.test    1.4820     0.1469  10.087 3.43e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 55.16 on 85 degrees of freedom
## Multiple R-squared:  0.5448, Adjusted R-squared:  0.5395
## F-statistic: 101.7 on 1 and 85 DF,  p-value: 3.432e-16
```

```
RMSE(elastic.predict.test,y.test) #57.86804
```

```
## [1] 57.86804
```

```
plot(y.test, elastic.predict.test, pch = 19, cex = 1, col = "black")
abline(lm(elastic.predict.test~y.test),col='red',lwd=2)
```



Phân tích ảnh hưởng, tương tác của các yếu tố và xây dựng mô hình hồi quy (R)

2. Trên bộ dữ liệu sau khi xử lý

```
#library
library(kernlab)
library(caret)
```

```
## Loading required package: lattice
```

```
## Loading required package: ggplot2
```

```
##
## Attaching package: 'ggplot2'
```

```
## The following object is masked from 'package:kernlab':
##
##      alpha
```

```
library(glmnet)
```

```
## Loading required package: Matrix
```

```
## Loaded glmnet 4.1-2
```

```
# Doc du lieu
df <- read.csv('diabetes.tab.tsv', header=TRUE, sep = '\t')

#preprocess

df_p = df
for( i in names(df_p))
{
  if(i!='SEX' & i!='Y')
  {
    Q3 = quantile(df_p[,i], 0.75)
    Q1 = quantile(df_p[,i], 0.25)
    IQR = Q3 - Q1
    olr_up = Q3+1.5*IQR
    olr_low = Q1-1.5*IQR

    df_p<-df_p[!(df_p[,i] > olr_up), ]
    df_p<-df_p[!(df_p[,i] < olr_low), ]
  }
}
```

```

}

#chia train-test
set.seed(123)
train.index <- createDataPartition(df_p$Y, p = .8, list = FALSE)

train_p <- df_p[ train.index,]
test_p <- df_p[-train.index,]

x.train_p <- train_p[,1:10]
x.test_p <- test_p[,1:10]
y.train_p <- train_p[,11]
y.test_p <- test_p[,11]

RMSE = function(m, o){
  sqrt(mean((m - o)^2))
}

# Effect of Factors
av <- aov(Y~.,data=train_p)
summary(av)

```

```

##              Df Sum Sq Mean Sq F value    Pr(>F)
## AGE              1   70622     70622   24.021 1.53e-06 ***
## SEX              1         2         2    0.001 0.98169
## BMI              1  464953   464953  158.147 < 2e-16 ***
## BP               1   91331    91331   31.065 5.35e-08 ***
## S1               1    1124     1124    0.382 0.53685
## S2               1     746      746    0.254 0.61481
## S3               1  223592   223592   76.052 < 2e-16 ***
## S4               1     358      358    0.122 0.72744
## S5               1   20759    20759    7.061 0.00828 **
## S6               1     661      661    0.225 0.63580
## Residuals      316  929043     2940
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

av <- aov(Y~(AGE+BMI+BP+S1+S2+S3+S5+S6)*S4,data=train_p)
summary(av)

```

```

##              Df Sum Sq Mean Sq F value    Pr(>F)
## AGE              1   70622     70622   23.338 2.14e-06 ***
## BMI              1  464743   464743  153.583 < 2e-16 ***
## BP               1   86284    86284   28.514 1.81e-07 ***
## S1               1    1359     1359    0.449 0.50329
## S2               1    2053     2053    0.678 0.41079
## S3               1  192796   192796   63.713 2.85e-14 ***
## S5               1   22091    22091    7.301 0.00727 **
## S6               1     369      369    0.122 0.72731
## S4               1     345      345    0.114 0.73595
## AGE:S4           1    6128     6128    2.025 0.15574
## BMI:S4           1    5028     5028    1.662 0.19836
## BP:S4            1     268      268    0.088 0.76637
## S1:S4            1      14       14    0.005 0.94559

```



```
## S2:S4      1      10      10    0.003  0.95391
## S3:S4      1    1404    1404    0.464  0.49630
## S5:S4      1    4644    4644    1.535  0.21635
## S6:S4      1    9999    9999    3.304  0.07007 .
## Residuals  309 935035    3026
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
av <- aov(Y~(AGE+BMI+BP+S1+S2+S3+S5+S6)*SEX,data=train_p)
summary(av)
```

```
##              Df Sum Sq Mean Sq F value    Pr(>F)
## AGE           1  70622   70622  24.333 1.33e-06 ***
## BMI           1 464743  464743 160.132 < 2e-16 ***
## BP            1  86284   86284  29.730 1.02e-07 ***
## S1            1   1359    1359   0.468 0.494335
## S2            1   2053    2053   0.707 0.401000
## S3            1 192796  192796  66.430 9.13e-15 ***
## S5            1  22091   22091   7.612 0.006145 **
## S6            1    369     369   0.127 0.721790
## SEX           1  32870   32870  11.326 0.000861 ***
## AGE:SEX       1  18617   18617   6.415 0.011813 *
## BMI:SEX       1   6800    6800   2.343 0.126872
## BP:SEX        1    519     519   0.179 0.672724
## S1:SEX        1    614     614   0.212 0.645764
## S2:SEX        1   1687    1687   0.581 0.446379
## S3:SEX        1   2718    2718   0.936 0.333951
## S5:SEX        1    157     157   0.054 0.816196
## S6:SEX        1   2096    2096   0.722 0.396099
## Residuals    309 896796    2902
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
# Effect of interaction
```

```
av <- aov(Y~AGE*BMI*BP*S1*S2*S3*S5*S6+AGE*SEX,data=train_p)
summary(av)
```

```
##              Df Sum Sq Mean Sq F value    Pr(>F)
## AGE           1  70622   70622  23.261 7.81e-06 ***
## BMI           1 464743  464743 153.076 < 2e-16 ***
## BP            1  86284   86284  28.420 1.10e-06 ***
## S1            1   1359    1359   0.448 0.50567
## S2            1   2053    2053   0.676 0.41368
## S3            1 192796  192796  63.503 1.91e-11 ***
## S5            1  22091   22091   7.276 0.00872 **
## S6            1    369     369   0.121 0.72853
## SEX           1  32870   32870  10.827 0.00156 **
## AGE:BMI       1   4782    4782   1.575 0.21357
## AGE:BP        1    147     147   0.048 0.82652
## BMI:BP        1  21990   21990   7.243 0.00887 **
## AGE:S1        1    293     293   0.096 0.75700
## BMI:S1        1   1316    1316   0.433 0.51244
## BP:S1         1    129     129   0.043 0.83726
## AGE:S2        1    875     875   0.288 0.59298
```

## BMI:S2	1	5232	5232	1.723	0.19351
## BP:S2	1	347	347	0.114	0.73628
## S1:S2	1	159	159	0.052	0.81972
## AGE:S3	1	92	92	0.030	0.86254
## BMI:S3	1	728	728	0.240	0.62597
## BP:S3	1	3345	3345	1.102	0.29743
## S1:S3	1	25	25	0.008	0.92727
## S2:S3	1	226	226	0.075	0.78568
## AGE:S5	1	10836	10836	3.569	0.06294 .
## BMI:S5	1	4246	4246	1.399	0.24091
## BP:S5	1	2439	2439	0.803	0.37317
## S1:S5	1	2	2	0.001	0.97820
## S2:S5	1	1431	1431	0.471	0.49461
## S3:S5	1	3165	3165	1.043	0.31068
## AGE:S6	1	10106	10106	3.329	0.07229 .
## BMI:S6	1	245	245	0.081	0.77732
## BP:S6	1	2123	2123	0.699	0.40582
## S1:S6	1	3629	3629	1.195	0.27798
## S2:S6	1	540	540	0.178	0.67438
## S3:S6	1	980	980	0.323	0.57175
## S5:S6	1	6	6	0.002	0.96502
## AGE:SEX	1	27985	27985	9.218	0.00335 **
## AGE:BMI:BP	1	3784	3784	1.246	0.26801
## AGE:BMI:S1	1	1301	1301	0.428	0.51485
## AGE:BP:S1	1	889	889	0.293	0.59019
## BMI:BP:S1	1	88	88	0.029	0.86528
## AGE:BMI:S2	1	2205	2205	0.726	0.39694
## AGE:BP:S2	1	4533	4533	1.493	0.22580
## BMI:BP:S2	1	40	40	0.013	0.90874
## AGE:S1:S2	1	989	989	0.326	0.57005
## BMI:S1:S2	1	283	283	0.093	0.76114
## BP:S1:S2	1	167	167	0.055	0.81545
## AGE:BMI:S3	1	1	1	0.000	0.98235
## AGE:BP:S3	1	5	5	0.002	0.96802
## BMI:BP:S3	1	34	34	0.011	0.91585
## AGE:S1:S3	1	1150	1150	0.379	0.54023
## BMI:S1:S3	1	262	262	0.086	0.76981
## BP:S1:S3	1	5144	5144	1.694	0.19725
## AGE:S2:S3	1	8	8	0.003	0.96014
## BMI:S2:S3	1	116	116	0.038	0.84538
## BP:S2:S3	1	1472	1472	0.485	0.48858
## S1:S2:S3	1	5631	5631	1.855	0.17754
## AGE:BMI:S5	1	2467	2467	0.813	0.37037
## AGE:BP:S5	1	2191	2191	0.722	0.39849
## BMI:BP:S5	1	1838	1838	0.605	0.43913
## AGE:S1:S5	1	277	277	0.091	0.76332
## BMI:S1:S5	1	7448	7448	2.453	0.12174
## BP:S1:S5	1	394	394	0.130	0.71958
## AGE:S2:S5	1	1360	1360	0.448	0.50545
## BMI:S2:S5	1	8731	8731	2.876	0.09430 .
## BP:S2:S5	1	1375	1375	0.453	0.50308
## S1:S2:S5	1	3371	3371	1.110	0.29555
## AGE:S3:S5	1	534	534	0.176	0.67631
## BMI:S3:S5	1	6886	6886	2.268	0.13651
## BP:S3:S5	1	5528	5528	1.821	0.18150
## S1:S3:S5	1	2012	2012	0.663	0.41838
## S2:S3:S5	1	7123	7123	2.346	0.13004

## AGE:BMI:S6	1	328	328	0.108	0.74341
## AGE:BP:S6	1	131	131	0.043	0.83624
## BMI:BP:S6	1	582	582	0.192	0.66290
## AGE:S1:S6	1	3894	3894	1.282	0.26125
## BMI:S1:S6	1	680	680	0.224	0.63748
## BP:S1:S6	1	1473	1473	0.485	0.48842
## AGE:S2:S6	1	324	324	0.107	0.74501
## BMI:S2:S6	1	33	33	0.011	0.91735
## BP:S2:S6	1	286	286	0.094	0.75986
## S1:S2:S6	1	526	526	0.173	0.67844
## AGE:S3:S6	1	17	17	0.006	0.94089
## BMI:S3:S6	1	15662	15662	5.159	0.02616 *
## BP:S3:S6	1	476	476	0.157	0.69340
## S1:S3:S6	1	70	70	0.023	0.88011
## S2:S3:S6	1	551	551	0.181	0.67146
## AGE:S5:S6	1	74	74	0.024	0.87636
## BMI:S5:S6	1	452	452	0.149	0.70087
## BP:S5:S6	1	645	645	0.213	0.64622
## S1:S5:S6	1	2583	2583	0.851	0.35943
## S2:S5:S6	1	90	90	0.029	0.86413
## S3:S5:S6	1	1490	1490	0.491	0.48585
## AGE:BMI:BP:S1	1	1733	1733	0.571	0.45250
## AGE:BMI:BP:S2	1	2922	2922	0.962	0.32993
## AGE:BMI:S1:S2	1	351	351	0.115	0.73498
## AGE:BP:S1:S2	1	1650	1650	0.544	0.46340
## BMI:BP:S1:S2	1	364	364	0.120	0.73034
## AGE:BMI:BP:S3	1	49	49	0.016	0.89894
## AGE:BMI:S1:S3	1	14307	14307	4.712	0.03329 *
## AGE:BP:S1:S3	1	1643	1643	0.541	0.46437
## BMI:BP:S1:S3	1	4256	4256	1.402	0.24035
## AGE:BMI:S2:S3	1	2836	2836	0.934	0.33705
## AGE:BP:S2:S3	1	238	238	0.078	0.78024
## BMI:BP:S2:S3	1	2183	2183	0.719	0.39935
## AGE:S1:S2:S3	1	483	483	0.159	0.69111
## BMI:S1:S2:S3	1	2180	2180	0.718	0.39968
## BP:S1:S2:S3	1	173	173	0.057	0.81225
## AGE:BMI:BP:S5	1	900	900	0.296	0.58790
## AGE:BMI:S1:S5	1	450	450	0.148	0.70140
## AGE:BP:S1:S5	1	2425	2425	0.799	0.37450
## BMI:BP:S1:S5	1	2912	2912	0.959	0.33074
## AGE:BMI:S2:S5	1	2675	2675	0.881	0.35112
## AGE:BP:S2:S5	1	1168	1168	0.385	0.53700
## BMI:BP:S2:S5	1	222	222	0.073	0.78756
## AGE:S1:S2:S5	1	116	116	0.038	0.84576
## BMI:S1:S2:S5	1	5	5	0.001	0.96923
## BP:S1:S2:S5	1	7879	7879	2.595	0.11162
## AGE:BMI:S3:S5	1	5180	5180	1.706	0.19571
## AGE:BP:S3:S5	1	2504	2504	0.825	0.36689
## BMI:BP:S3:S5	1	3453	3453	1.137	0.28982
## AGE:S1:S3:S5	1	1548	1548	0.510	0.47756
## BMI:S1:S3:S5	1	1447	1447	0.477	0.49219
## BP:S1:S3:S5	1	1484	1484	0.489	0.48668
## AGE:S2:S3:S5	1	331	331	0.109	0.74236
## BMI:S2:S3:S5	1	69	69	0.023	0.88058
## BP:S2:S3:S5	1	1962	1962	0.646	0.42417
## S1:S2:S3:S5	1	4808	4808	1.584	0.21238
## AGE:BMI:BP:S6	1	16	16	0.005	0.94238

## AGE:BMI:S1:S6	1	4171	4171	1.374	0.24507	
## AGE:BP:S1:S6	1	30	30	0.010	0.92057	
## BMI:BP:S1:S6	1	234	234	0.077	0.78216	
## AGE:BMI:S2:S6	1	7740	7740	2.550	0.11477	
## AGE:BP:S2:S6	1	2	2	0.001	0.97775	
## BMI:BP:S2:S6	1	7763	7763	2.557	0.11426	
## AGE:S1:S2:S6	1	367	367	0.121	0.72901	
## BMI:S1:S2:S6	1	726	726	0.239	0.62642	
## BP:S1:S2:S6	1	193	193	0.064	0.80151	
## AGE:BMI:S3:S6	1	790	790	0.260	0.61163	
## AGE:BP:S3:S6	1	2957	2957	0.974	0.32704	
## BMI:BP:S3:S6	1	1357	1357	0.447	0.50594	
## AGE:S1:S3:S6	1	942	942	0.310	0.57934	
## BMI:S1:S3:S6	1	10686	10686	3.520	0.06475	.
## BP:S1:S3:S6	1	14426	14426	4.752	0.03259	*
## AGE:S2:S3:S6	1	9321	9321	3.070	0.08406	.
## BMI:S2:S3:S6	1	666	666	0.219	0.64090	
## BP:S2:S3:S6	1	67	67	0.022	0.88274	
## S1:S2:S3:S6	1	9781	9781	3.222	0.07692	.
## AGE:BMI:S5:S6	1	7751	7751	2.553	0.11452	
## AGE:BP:S5:S6	1	810	810	0.267	0.60719	
## BMI:BP:S5:S6	1	165	165	0.054	0.81654	
## AGE:S1:S5:S6	1	7214	7214	2.376	0.12764	
## BMI:S1:S5:S6	1	0	0	0.000	0.99403	
## BP:S1:S5:S6	1	6580	6580	2.167	0.14539	
## AGE:S2:S5:S6	1	758	758	0.250	0.61882	
## BMI:S2:S5:S6	1	15618	15618	5.144	0.02637	*
## BP:S2:S5:S6	1	4558	4558	1.501	0.22454	
## S1:S2:S5:S6	1	512	512	0.169	0.68260	
## AGE:S3:S5:S6	1	1262	1262	0.416	0.52116	
## BMI:S3:S5:S6	1	1295	1295	0.427	0.51574	
## BP:S3:S5:S6	1	29	29	0.009	0.92289	
## S1:S3:S5:S6	1	15143	15143	4.988	0.02868	*
## S2:S3:S5:S6	1	9144	9144	3.012	0.08699	.
## AGE:BMI:BP:S1:S2	1	1032	1032	0.340	0.56166	
## AGE:BMI:BP:S1:S3	1	2229	2229	0.734	0.39436	
## AGE:BMI:BP:S2:S3	1	410	410	0.135	0.71424	
## AGE:BMI:S1:S2:S3	1	3810	3810	1.255	0.26640	
## AGE:BP:S1:S2:S3	1	26	26	0.009	0.92669	
## BMI:BP:S1:S2:S3	1	1290	1290	0.425	0.51656	
## AGE:BMI:BP:S1:S5	1	6598	6598	2.173	0.14485	
## AGE:BMI:BP:S2:S5	1	1304	1304	0.429	0.51443	
## AGE:BMI:S1:S2:S5	1	1759	1759	0.579	0.44911	
## AGE:BP:S1:S2:S5	1	416	416	0.137	0.71249	
## BMI:BP:S1:S2:S5	1	174	174	0.057	0.81173	
## AGE:BMI:BP:S3:S5	1	2282	2282	0.752	0.38889	
## AGE:BMI:S1:S3:S5	1	503	503	0.166	0.68508	
## AGE:BP:S1:S3:S5	1	14	14	0.005	0.94631	
## BMI:BP:S1:S3:S5	1	4485	4485	1.477	0.22824	
## AGE:BMI:S2:S3:S5	1	2631	2631	0.866	0.35509	
## AGE:BP:S2:S3:S5	1	721	721	0.237	0.62762	
## BMI:BP:S2:S3:S5	1	3519	3519	1.159	0.28530	
## AGE:S1:S2:S3:S5	1	664	664	0.219	0.64140	
## BMI:S1:S2:S3:S5	1	6208	6208	2.045	0.15712	
## BP:S1:S2:S3:S5	1	1953	1953	0.643	0.42516	
## AGE:BMI:BP:S1:S6	1	8588	8588	2.829	0.09699	.
## AGE:BMI:BP:S2:S6	1	116	116	0.038	0.84578	

## AGE:BMI:S1:S2:S6	1	2042	2042	0.673	0.41489	
## AGE:BP:S1:S2:S6	1	580	580	0.191	0.66351	
## BMI:BP:S1:S2:S6	1	959	959	0.316	0.57587	
## AGE:BMI:BP:S3:S6	1	52	52	0.017	0.89653	
## AGE:BMI:S1:S3:S6	1	553	553	0.182	0.67070	
## AGE:BP:S1:S3:S6	1	7192	7192	2.369	0.12821	
## BMI:BP:S1:S3:S6	1	5564	5564	1.833	0.18009	
## AGE:BMI:S2:S3:S6	1	933	933	0.307	0.58101	
## AGE:BP:S2:S3:S6	1	1249	1249	0.411	0.52329	
## BMI:BP:S2:S3:S6	1	363	363	0.120	0.73035	
## AGE:S1:S2:S3:S6	1	616	616	0.203	0.65390	
## BMI:S1:S2:S3:S6	1	5100	5100	1.680	0.19915	
## BP:S1:S2:S3:S6	1	23514	23514	7.745	0.00690	**
## AGE:BMI:BP:S5:S6	1	9093	9093	2.995	0.08787	.
## AGE:BMI:S1:S5:S6	1	8536	8536	2.811	0.09799	.
## AGE:BP:S1:S5:S6	1	5864	5864	1.931	0.16894	
## BMI:BP:S1:S5:S6	1	2526	2526	0.832	0.36482	
## AGE:BMI:S2:S5:S6	1	2691	2691	0.886	0.34968	
## AGE:BP:S2:S5:S6	1	6561	6561	2.161	0.14598	
## BMI:BP:S2:S5:S6	1	73	73	0.024	0.87729	
## AGE:S1:S2:S5:S6	1	8866	8866	2.920	0.09184	.
## BMI:S1:S2:S5:S6	1	3263	3263	1.075	0.30342	
## BP:S1:S2:S5:S6	1	1918	1918	0.632	0.42934	
## AGE:BMI:S3:S5:S6	1	3296	3296	1.086	0.30097	
## AGE:BP:S3:S5:S6	1	2571	2571	0.847	0.36056	
## BMI:BP:S3:S5:S6	1	2200	2200	0.725	0.39746	
## AGE:S1:S3:S5:S6	1	6816	6816	2.245	0.13846	
## BMI:S1:S3:S5:S6	1	4357	4357	1.435	0.23492	
## BP:S1:S3:S5:S6	1	4313	4313	1.421	0.23725	
## AGE:S2:S3:S5:S6	1	1179	1179	0.388	0.53512	
## BMI:S2:S3:S5:S6	1	4899	4899	1.614	0.20812	
## BP:S2:S3:S5:S6	1	58	58	0.019	0.89018	
## S1:S2:S3:S5:S6	1	3042	3042	1.002	0.32023	
## AGE:BMI:BP:S1:S2:S3	1	130	130	0.043	0.83650	
## AGE:BMI:BP:S1:S2:S5	1	1380	1380	0.455	0.50234	
## AGE:BMI:BP:S1:S3:S5	1	2118	2118	0.698	0.40641	
## AGE:BMI:BP:S2:S3:S5	1	808	808	0.266	0.60762	
## AGE:BMI:S1:S2:S3:S5	1	2674	2674	0.881	0.35121	
## AGE:BP:S1:S2:S3:S5	1	32	32	0.011	0.91863	
## BMI:BP:S1:S2:S3:S5	1	384	384	0.126	0.72326	
## AGE:BMI:BP:S1:S2:S6	1	1962	1962	0.646	0.42419	
## AGE:BMI:BP:S1:S3:S6	1	2763	2763	0.910	0.34332	
## AGE:BMI:BP:S2:S3:S6	1	146	146	0.048	0.82709	
## AGE:BMI:S1:S2:S3:S6	1	2933	2933	0.966	0.32897	
## AGE:BP:S1:S2:S3:S6	1	1331	1331	0.438	0.51007	
## BMI:BP:S1:S2:S3:S6	1	5452	5452	1.796	0.18448	
## AGE:BMI:BP:S1:S5:S6	1	4169	4169	1.373	0.24518	
## AGE:BMI:BP:S2:S5:S6	1	5742	5742	1.891	0.17337	
## AGE:BMI:S1:S2:S5:S6	1	21394	21394	7.047	0.00980	**
## AGE:BP:S1:S2:S5:S6	1	9	9	0.003	0.95580	
## BMI:BP:S1:S2:S5:S6	1	7851	7851	2.586	0.11226	
## AGE:BMI:BP:S3:S5:S6	1	435	435	0.143	0.70631	
## AGE:BMI:S1:S3:S5:S6	1	577	577	0.190	0.66429	
## AGE:BP:S1:S3:S5:S6	1	664	664	0.219	0.64153	
## BMI:BP:S1:S3:S5:S6	1	1477	1477	0.486	0.48782	
## AGE:BMI:S2:S3:S5:S6	1	477	477	0.157	0.69310	
## AGE:BP:S2:S3:S5:S6	1	276	276	0.091	0.76384	

```
## BMI:BP:S2:S3:S5:S6      1      930      930      0.306      0.58177
## AGE:S1:S2:S3:S5:S6      1     4483     4483     1.476     0.22835
## BMI:S1:S2:S3:S5:S6      1       42       42     0.014     0.90643
## BP:S1:S2:S3:S5:S6       1     5983     5983     1.971     0.16474
## AGE:BMI:BP:S1:S2:S3:S5  1     5757     5757     1.896     0.17282
## AGE:BMI:BP:S1:S2:S3:S6  1     1170     1170     0.385     0.53681
## AGE:BMI:BP:S1:S2:S5:S6  1       747       747     0.246     0.62151
## AGE:BMI:BP:S2:S3:S5:S6  1     3722     3722     1.226     0.27191
## AGE:BMI:S1:S2:S3:S5:S6  1     2288     2288     0.754     0.38822
## AGE:BP:S1:S2:S3:S5:S6   1     5154     5154     1.698     0.19682
## BMI:BP:S1:S2:S3:S5:S6   1        21        21     0.007     0.93424
## Residuals                71 215558      3036
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
#Build Models
#SLR
#train
fit <- lm(formula = Y~BMI,data=train_p)
y.train.pred <- predict(fit,newdata=x.train_p)
summary(fit)
```

```
##
## Call:
## lm(formula = Y ~ BMI, data = train_p)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -156.418  -45.526   -7.454   47.302  157.173
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -100.1405     21.8255  -4.588 6.39e-06 ***
## BMI           9.4412      0.8184   11.536 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 62.74 on 325 degrees of freedom
## Multiple R-squared:  0.2905, Adjusted R-squared:  0.2883
## F-statistic: 133.1 on 1 and 325 DF, p-value: < 2.2e-16
```

```
RMSE(y.train.pred,y.train_p)
```

```
## [1] 62.54935
```

```
#test
y.test.pred <- predict(fit,newdata=x.test_p)
predict <- lm(y.test_p~y.test.pred)
summary(predict)
```

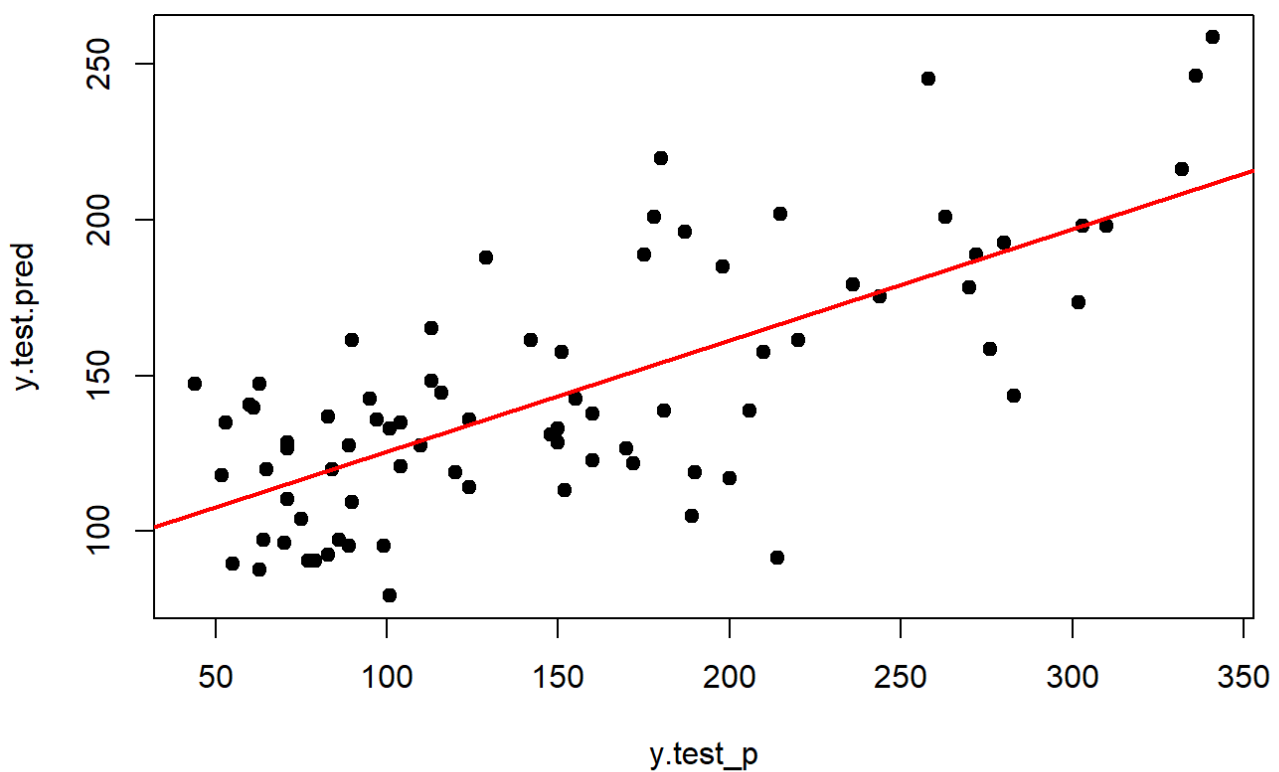
```
##
## Call:
```

```
## lm(formula = y.test_p ~ y.test.pred)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -112.682  -40.283   -3.243   39.606  137.618
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -55.5430    23.6648  -2.347   0.0215 *
## y.test.pred   1.4416     0.1581   9.120 6.31e-14 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 56.46 on 78 degrees of freedom
## Multiple R-squared:  0.5161, Adjusted R-squared:  0.5099
## F-statistic: 83.18 on 1 and 78 DF,  p-value: 6.311e-14
```

```
RMSE(y.test.pred,y.test_p)
```

```
## [1] 59.04361
```

```
plot(y.test_p,y.test.pred, pch = 19, cex = 1, col = "black")
abline(lm(y.test.pred~y.test_p),col='red',lwd=2)
```



```
#multiple linear regression
#train
fit <- lm(formula = Y~SEX+BMI+BP+S1+S3+S5,data=train_p)
```

```
y.train.pred <- predict(fit,newdata=x.train_p)
summary(fit)
```

```
##
## Call:
## lm(formula = Y ~ SEX + BMI + BP + S1 + S3 + S5, data = train_p)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -149.543  -37.501   -0.557   35.273  141.916
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  -215.9629    44.5227  -4.851 1.93e-06 ***
## SEX           -22.8206     6.8001  -3.356 0.000886 ***
## BMI             4.0890     0.8783   4.655 4.74e-06 ***
## BP              1.3396     0.2709   4.945 1.23e-06 ***
## S1             -0.1542     0.1181  -1.306 0.192529
## S3             -1.1371     0.3262  -3.486 0.000560 ***
## S5              54.1603     8.6748   6.243 1.36e-09 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 53.93 on 320 degrees of freedom
## Multiple R-squared:  0.4838, Adjusted R-squared:  0.4741
## F-statistic: 49.98 on 6 and 320 DF,  p-value: < 2.2e-16
```

```
RMSE(y.train.pred,y.train_p)
```

```
## [1] 53.3535
```

```
#test
y.test.pred <- predict(fit,newdata=x.test_p)
predict <- lm(y.test_p~y.test.pred)
summary(predict)
```

```
##
## Call:
## lm(formula = y.test_p ~ y.test.pred)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -110.539  -42.363   -5.933   35.771  158.181
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   -6.9275    18.9331  -0.366   0.715
## y.test.pred    1.0656     0.1192   8.941 1.4e-13 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 57.04 on 78 degrees of freedom
```

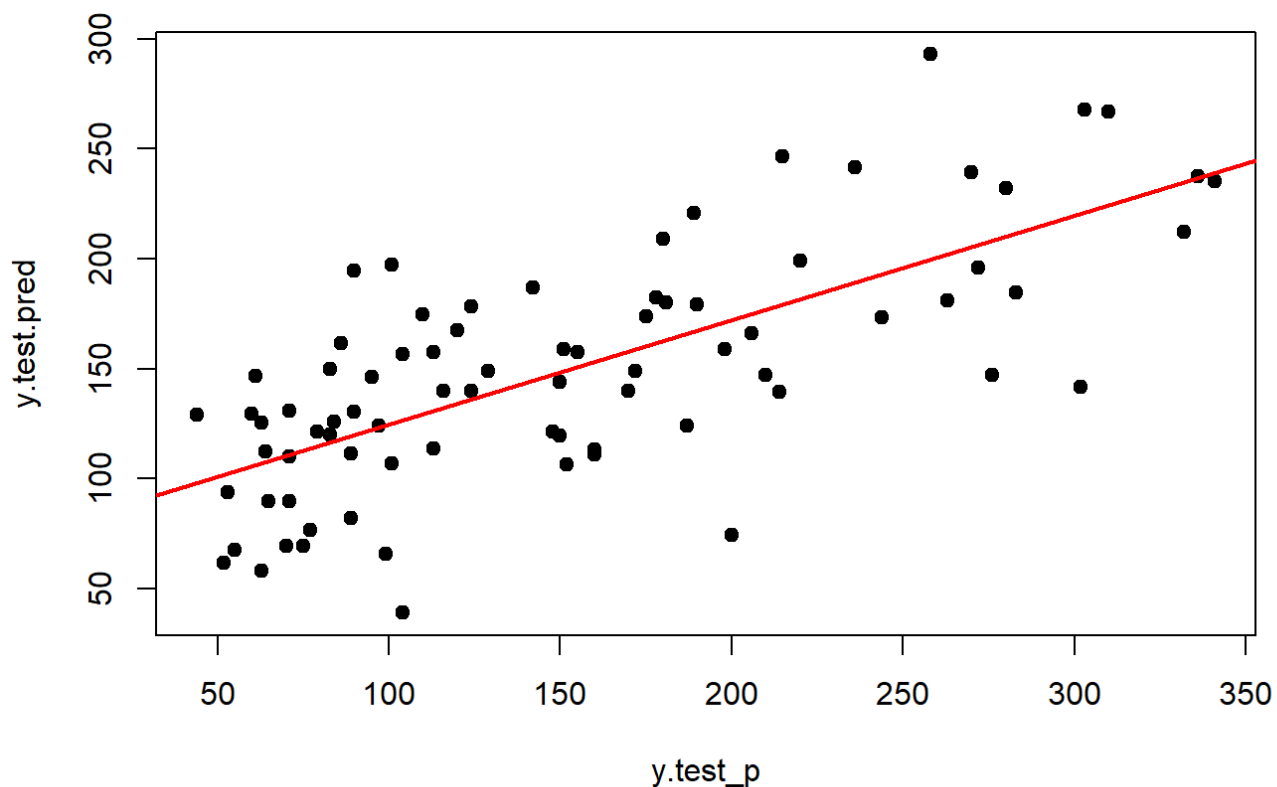


```
## Multiple R-squared:  0.5061, Adjusted R-squared:  0.4998
## F-statistic: 79.94 on 1 and 78 DF,  p-value: 1.404e-13
```

```
RMSE(y.test.pred,y.test_p)
```

```
## [1] 56.50382
```

```
plot(y.test_p,y.test.pred, pch = 19, cex = 1, col = "black")
abline(lm(y.test.pred~y.test_p),col='red',lwd=2)
```



```
#polynomial regression
#train
fit <- lm(formula = Y~AGE+BMI+S3+SEX+S5+I(BP^2)+I(AGE*SEX)+I(BMI*BP)+I(BP*S1*S
3*S6)+I(AGE*BMI*S1*S3)+
          I(BMI*S2*S5*S6)+I(BP*S1*S2*S3*S6),
          data=train_p)
y.train.pred <- predict(fit,newdata=x.train_p)
summary(fit)
```

```
##
## Call:
## lm(formula = Y ~ AGE + BMI + S3 + SEX + S5 + I(BP^2) + I(AGE *
##     SEX) + I(BMI * BP) + I(BP * S1 * S3 * S6) + I(AGE * BMI *
##     S1 * S3) + I(BMI * S2 * S5 * S6) + I(BP * S1 * S2 * S3 *
##     S6), data = train_p)
##
```

```
## Residuals:
##      Min       1Q   Median       3Q      Max
## -142.038  -37.410   -0.199   34.083  133.681
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    2.947e+02  1.329e+02   2.218 0.027300 *
## AGE           -2.731e+00  1.053e+00  -2.594 0.009927 **
## BMI           -1.548e+01  5.445e+00  -2.842 0.004776 **
## S3            -2.081e+00  1.026e+00  -2.028 0.043409 *
## SEX           -9.272e+01  2.422e+01  -3.829 0.000155 ***
## S5             4.012e+01  1.002e+01   4.005 7.76e-05 ***
## I(BP^2)        -1.931e-02  8.467e-03  -2.281 0.023212 *
## I(AGE * SEX)    1.431e+00  4.810e-01   2.976 0.003147 **
## I(BMI * BP)     1.841e-01  5.534e-02   3.326 0.000985 ***
## I(BP * S1 * S3 * S6) 6.006e-07  5.372e-07   1.118 0.264405
## I(AGE * BMI * S1 * S3) 2.929e-06  2.529e-06   1.158 0.247655
## I(BMI * S2 * S5 * S6) 2.103e-05  1.936e-05   1.086 0.278230
## I(BP * S1 * S2 * S3 * S6) -5.426e-09  3.016e-09  -1.799 0.072951 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 52.78 on 314 degrees of freedom
## Multiple R-squared:  0.5149, Adjusted R-squared:  0.4964
## F-statistic: 27.78 on 12 and 314 DF,  p-value: < 2.2e-16
```

```
RMSE(y.train.pred,y.train_p)
```

```
## [1] 51.71859
```

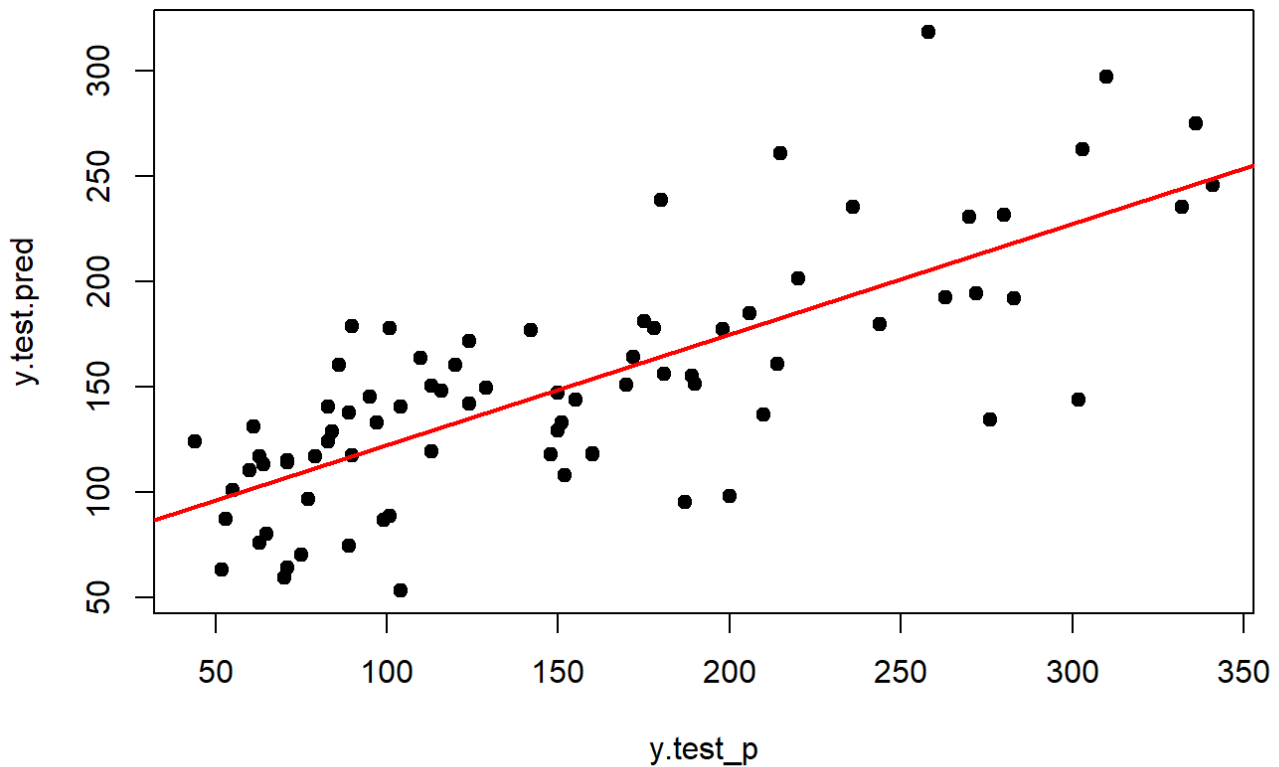
```
#test
y.test.pred <- predict(fit,newdata=x.test_p)
predict <- lm(y.test_p~y.test.pred)
summary(predict)
```

```
##
## Call:
## lm(formula = y.test_p ~ y.test.pred)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
##  -93.94  -43.09   -4.08   30.47  155.92
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  -10.6151    17.0283  -0.623   0.535
## y.test.pred   1.0891     0.1066  10.217 4.83e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 53.08 on 78 degrees of freedom
## Multiple R-squared:  0.5724, Adjusted R-squared:  0.5669
## F-statistic: 104.4 on 1 and 78 DF,  p-value: 4.83e-16
```

```
RMSE(y.test.pred,y.test_p)
```

```
## [1] 52.71497
```

```
plot(y.test_p,y.test.pred, pch = 19, cex = 1, col = "black")  
abline(lm(y.test.pred~y.test_p),col='red',lwd=2)
```



```
#ridge (alpha = 0)  
set.seed(123)  
ridge.fit <- cv.glmnet(as.matrix(x.train_p),y.train_p,type.measure = 'mse',alpha=  
0,family='gaussian')  
ridge.fit$lambda.1se
```

```
## [1] 56.88734
```

```
coef(ridge.fit)
```

```
## 11 x 1 sparse Matrix of class "dgCMatrix"  
##              s1  
## (Intercept) -1.447451e+02  
## AGE         1.147955e-01  
## SEX        -9.351097e+00  
## BMI         3.051928e+00  
## BP          8.154471e-01  
## S1          2.035922e-03
```

```
## S2          -6.766365e-02
## S3          -6.811350e-01
## S4           4.990239e+00
## S5           2.890386e+01
## S6           3.586102e-01
```

```
#summary train
ridge.predict.train <- predict(ridge.fit,s=ridge.fit$lambda.1se,newx = as.matri
x(x.train_p))
fit <- lm(y.train_p~ridge.predict.train)
summary(fit)
```

```
##
## Call:
## lm(formula = y.train_p ~ ridge.predict.train)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -133.567  -39.101   -1.942   40.989  128.080
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -47.28045     11.94939   -3.957 9.33e-05 ***
## ridge.predict.train  1.31854      0.07792  16.921 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 54.31 on 325 degrees of freedom
## Multiple R-squared:  0.4684, Adjusted R-squared:  0.4667
## F-statistic: 286.3 on 1 and 325 DF,  p-value: < 2.2e-16
```

```
RMSE(ridge.predict.train,y.train_p)
```

```
## [1] 55.51858
```

```
#summary test
ridge.predict.test <- predict(ridge.fit,s=ridge.fit$lambda.1se,newx = as.matri
x(x.test_p))
predict <- lm(y.test_p~ridge.predict.test)
summary(predict)
```

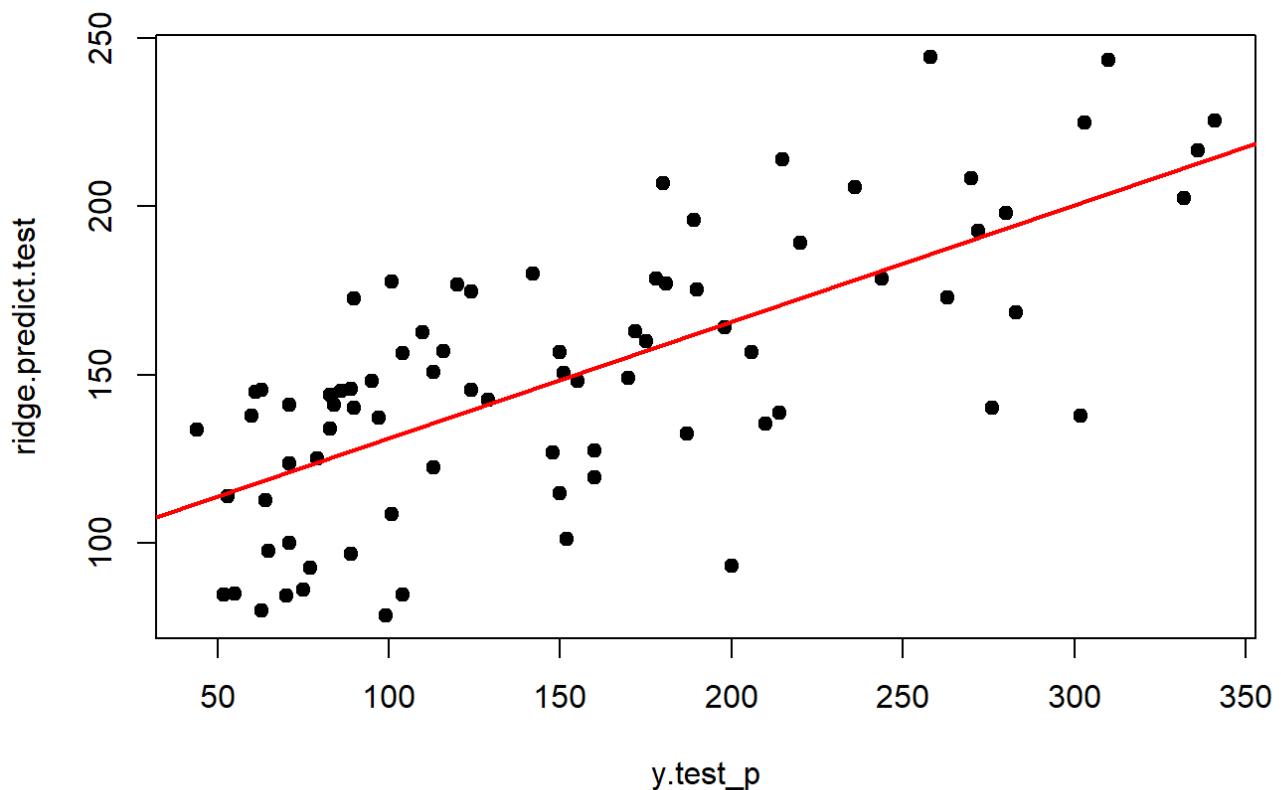
```
##
## Call:
## lm(formula = y.test_p ~ ridge.predict.test)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
##  -95.51  -48.11   -2.48   42.97  165.79
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -58.6149     25.2386   -2.322  0.0228 *
```

```
## ridge.predict.test    1.4134      0.1633    8.654 5.08e-13 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 57.97 on 78 degrees of freedom
## Multiple R-squared:  0.4898, Adjusted R-squared:  0.4833
## F-statistic: 74.88 on 1 and 78 DF,  p-value: 5.078e-13
```

```
RMSE(ridge.predict.test,y.test_p)
```

```
## [1] 59.6312
```

```
#plot
plot(y.test_p,ridge.predict.test, pch = 19, cex = 1, col = "black")
abline(lm(ridge.predict.test~y.test_p),col='red',lwd=2)
```



```
#lasso (alpha = 1)
set.seed(123)
lasso.fit <- cv.glmnet(as.matrix(x.train_p),y.train_p,type.measure ='mse',alpha=
1,family='gaussian')
lasso.fit$lambda.1se
```

```
## [1] 7.87825
```

```
coef(lasso.fit)
```

```
## 11 x 1 sparse Matrix of class "dgCMatrix"
##              s1
## (Intercept) -192.5409138
## AGE          .
## SEX          .
## BMI          3.9613444
## BP           0.6776335
## S1           .
## S2           .
## S3          -0.5086763
## S4           .
## S5          43.0942120
## S6           .
```

```
#summary train
lasso.predict.train <- predict(lasso.fit,s=lasso.fit$lambda.1se,newx = as.matrix(x.train_p))
fit <- lm(y.train_p~lasso.predict.train)
summary(fit)
```

```
##
## Call:
## lm(formula = y.train_p ~ lasso.predict.train)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -142.971  -38.190   -1.197   41.732  135.871
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -38.37113     11.64643   -3.295  0.00109 **
## lasso.predict.train  1.25852      0.07576  16.611 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 54.78 on 325 degrees of freedom
## Multiple R-squared:  0.4592, Adjusted R-squared:  0.4575
## F-statistic: 275.9 on 1 and 325 DF,  p-value: < 2.2e-16
```

```
RMSE(lasso.predict.train,y.train_p)
```

```
## [1] 55.58046
```

```
#summary test
lasso.predict.test <- predict(lasso.fit,s=lasso.fit$lambda.1se,newx = as.matrix(x.test_p))
predict <- lm(y.test_p~lasso.predict.test)
summary(predict)
```

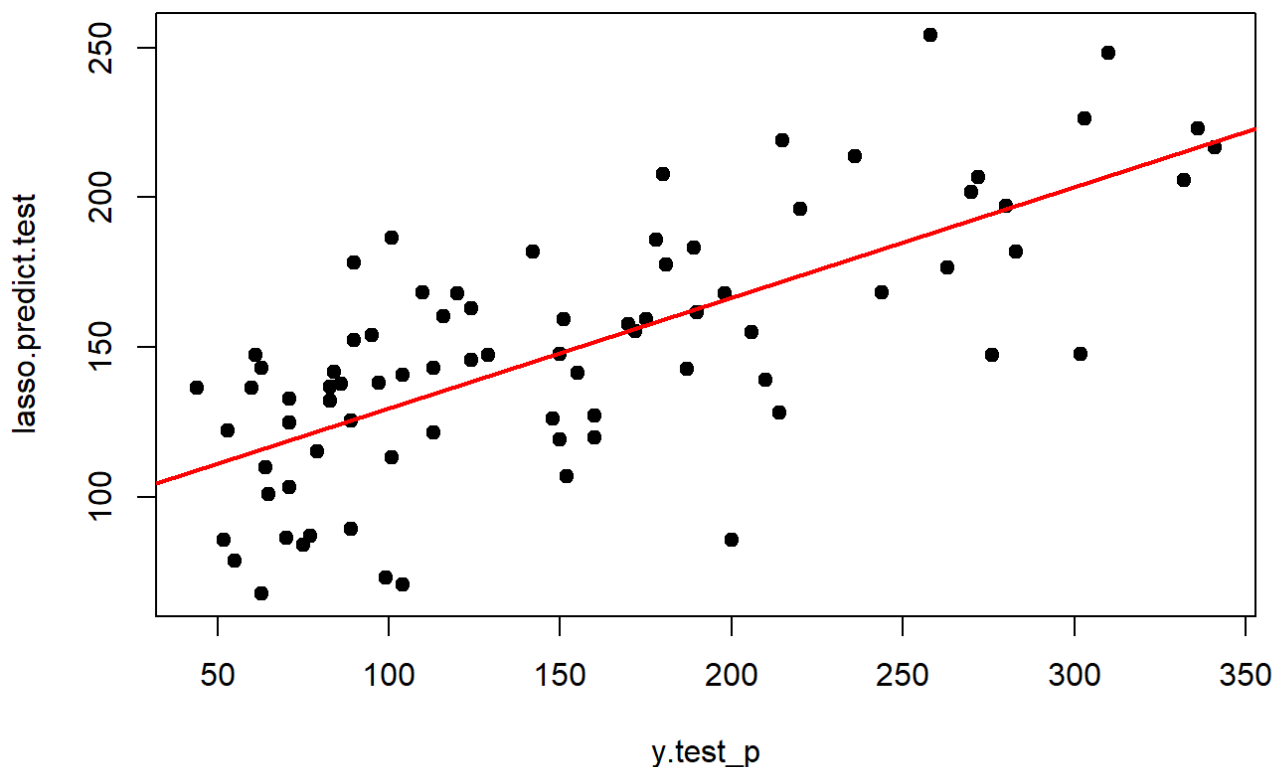
```
##
## Call:
## lm(formula = y.test_p ~ lasso.predict.test)
```

```
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -102.830  -46.847   -2.457   41.180  151.067
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -50.8834     23.7395  -2.143   0.0352 *
## lasso.predict.test  1.3650      0.1535   8.894 1.73e-13 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 57.19 on 78 degrees of freedom
## Multiple R-squared:  0.5035, Adjusted R-squared:  0.4972
## F-statistic: 79.11 on 1 and 78 DF,  p-value: 1.73e-13
```

```
RMSE(lasso.predict.test,y.test_p)
```

```
## [1] 58.58606
```

```
#plot
plot(y.test_p,lasso.predict.test, pch = 19, cex = 1, col = "black")
abline(lm(lasso.predict.test~y.test_p),col='red',lwd=2)
```



```
#elastic net
results.train <-data.frame()
for (i in 0:20)
```

```

{
  set.seed(123)
  fit <- cv.glmnet(as.matrix(x.train_p), y.train_p, type.measure="mse", alpha=i/
20,
                  family="gaussian")
  y.pred <- predict(fit, s=fit$lambda.1se, newx=as.matrix(x.train_p))
  predict <- lm(y.train_p~y.pred)

  temp <- data.frame(alpha=i/20,R2= summary(predict)$r.squared,Adj_R2=summary(pr
edict)$adj.r.squared,rmse=RMSE(y.pred,y.train_p),lambda=fit$lambda.1se)
  results.train <- rbind(results.train, temp)
}
#best alpha = 0.05 (best adj R2)

set.seed(123)
elastic.fit <- cv.glmnet(as.matrix(x.train_p), y.train_p, type.measure="mse", al
pha=0.05,
                        family="gaussian")
elastic.fit$lambda.1se

```

```
## [1] 39.03005
```

```
coef(elastic.fit)
```

```

## 11 x 1 sparse Matrix of class "dgCMatrix"
##              s1
## (Intercept) -156.68332510
## AGE          0.00625415
## SEX          -8.01131052
## BMI          3.28575464
## BP           0.84842402
## S1            .
## S2          -0.01193017
## S3          -0.69496170
## S4           3.61105763
## S5           32.11917544
## S6           0.26372880

```

```

#train
elastic.predict.train <- predict(elastic.fit, s=elastic.fit$lambda.1se, newx=a
s.matrix(x.train_p))
fit <- lm(y.train_p~elastic.predict.train)
summary(fit)

```

```

##
## Call:
## lm(formula = y.train_p ~ elastic.predict.train)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -135.223  -38.024   -1.198   40.713  130.613
##
## Coefficients:

```



```
##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)      -43.03977    11.69242  -3.681 0.000272 ***
## elastic.predict.train  1.28997     0.07614  16.943 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 54.28 on 325 degrees of freedom
## Multiple R-squared:  0.469, Adjusted R-squared:  0.4674
## F-statistic: 287.1 on 1 and 325 DF, p-value: < 2.2e-16
```

```
RMSE(elastic.predict.train,y.train_p)
```

```
## [1] 55.30564
```

```
#test
elastic.predict.test <- predict(elastic.fit, s=elastic.fit$lambda.1se, newx=a
s.matrix(x.test_p))
predict <- lm(y.test_p~elastic.predict.test)
summary(predict)
```

```
##
## Call:
## lm(formula = y.test_p ~ elastic.predict.test)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -95.690 -49.031  -2.624   41.652  158.960
##
## Coefficients:
##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)      -53.7196    24.5495  -2.188   0.0316 *
## elastic.predict.test  1.3812     0.1587   8.706 4.02e-13 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 57.8 on 78 degrees of freedom
## Multiple R-squared:  0.4928, Adjusted R-squared:  0.4863
## F-statistic: 75.79 on 1 and 78 DF, p-value: 4.021e-13
```

```
RMSE(elastic.predict.test,y.test_p)
```

```
## [1] 59.23687
```

```
#plot
plot(y.test_p, elastic.predict.test, pch = 19, cex = 1, col = "black")
abline(lm(elastic.predict.test~y.test_p),col='red',lwd=2)
```

