

Dataset Overview:

The Diabetes Prediction Dataset includes patients' medical and demographic information as well as the state of their diabetes (positive or negative). The provided dataset contains various health-related attributes. The information includes gender, age, hypertension, heart disease, smoking history, BMI, HbA1c level, blood glucose level, and diabetes status. This dataset appears to be intended for research or analysis related to health and wellness, with a focus on factors that may impact an individual's health. The project's objective seems to be to investigate the relationships between these factors and their potential impact on developing diabetes. The findings from this analysis could be valuable for understanding how these factors are associated with specific health conditions and for making informed decisions regarding diabetes identification and personalized treatment.

Reference: <https://www.kaggle.com/datasets/iammustafatz/diabetes-prediction-dataset/data>

Data pre-processing:

*Importing the Dataset:

At first the dataset is imported from the correct path. The data is read from a CSV file named "Dataset_Midterm.csv" using the read.csv method in R. The resulting data frame is then assigned to a variable named mydata. mydata is then used to display the contents of the dataset in the console. The output of this code displays the entire contents of the dataset, including all rows and columns.

```
> options(max.print = 1e6)
> mydata<-read.csv("D:/Fall 2023-24 11th sem/Introduction to Data Science/Dataset_Midterm.csv", header = TRUE, sep = ",")
> mydata
```

The output provides a way to verify that the data has been read correctly and allows for a quick visual inspection of the dataset. The output also provides information about the structure of the data, such as the number of columns and rows and the type of data in each column. Overall, this code and its output are a necessary first step in any data analysis project in R. The output is given below:

	gender	age	hypertension	heart_disease	smoking_history	bmi
1	Female	80	0	1	never	25.19
2	Female	54	0	0	No Info	27.32
3	Male	28	0	0	never	-27.32
4	Female	NA	0	0	current	23.45
5	Male	76	1	1	current	20.14
6	Female	20	0	0	never	27.32
7		79	0	0	No Info	23.86
8	Male	42	0	0	never	33.64
9	Female	32	0	0	never	27.32
10	Female	53	0	0	never	27.32
11	Female	54	0	0	former	54.70
12	Female	78	NA	0	former	36.05
13	Female	67	0	0	never	25.69
14	Female	76	0	0	No Info	27.32
15		78	0	0	No Info	27.32
16	Male	15	0	0	never	30.36
17	Female	42	0	0	never	24.48
18	Female	42	0	0	No Info	27.32
19	Male	NA	0	0	ever	25.72
20	Male	40	0	0	current	36.38
21	Male	5	0	0	No Info	18.80
22	Female	69	0	0	never	21.24
23	Female	72	0	1	former	27.94
24	Female	4	0	0	No Info	13.99
25	Male	30	0	0	never	33.76
26	Male	40	0	0	former	27.85
27	Male	45	NA	0	never	26.47
28	Male	43	0	0	never	26.08
29	Female	53	0	0	No Info	31.75
30	Male	50	0	0	No Info	25.15
31	Female	41	0	0	current	22.01
32	Female	20	0	0	never	22.19
33	Female	76	0	0	never	23.55
34	Male	5	0	0	No Info	15.10
35	Female	15	0	0	No Info	21.76
36	Female	26	0	0	never	21.22
37	Male	5	0	0	No Info	27.32
38	Female	77	1	1	never	22.62

	gender	age	hypertension	heart_disease	smoking_history	bmi
39	Female	66	0	0	No Info	29.30
40	Female	67	0	0	No Info	27.32
41	Female	44	0	0	never	24.93
42	Female	29	0	0	never	19.95
43	Female	60	0	0	never	18.03
44	Female	38	0	0	never	28.27
45	Female	3	0	0	No Info	19.27
46	Male	57	0	0	never	27.32
47	Female	43	0	0	No Info	27.32
48	Female	74	0	0	No Info	28.12
49	Female	21	0	0		26.10
50	Female	30	0	0	current	27.32
51	Female	59	0	0	former	27.32
52	Female	290	0	0	not current	30.22
53	Female	59	0	1	ever	23.11
54	Female	19	0	0		27.32
55	Female	NA	0	0	No Info	28.16
56	Male	56	0	0	never	26.78
57	Male	43	0	0	No Info	23.04
58	Male	7	0	0	No Info	15.94
59	Male	3	0	0	No Info	15.80
60	Female	30	0	0	never	27.01
61	Male	43	0	0		27.32
62	Female	76	0	0	never	22.19
63	Female	41	0	0	never	27.45
64	Female	11	0	0	No Info	17.98
65	Female	26	0	0	never	26.45
66	Male	34	0	0	never	31.16
67	Male	80	0	0	former	24.42
68	Female	37	0	0	No Info	30.50
69	Female	44	0	0	never	19.31
70	Male	67	0	1	not current	27.32
71	Male	50	1	0	current	27.32
72	Male	73	0	0	former	25.91
73	Female	53	0	0	former	27.32
74	Male	50	0	0	former	37.16
75	Female	67	0	0	never	63.48
76	Male	57	0	0	No Info	27.32
77	Female	36	0	0	current	32.27
78	Female	60	0	0	never	27.32
79	Female	67	0	0	never	27.32

	gender	age	hypertension	heart_disease	smoking_history	bmi
80	Female	80	1	0	never	27.32
81	Female	NA	0	0	never	31.70
82	Male	80	0	0	never	22.06
83	Female	47	0	0	never	36.49
84	Male	53	0	0	current	30.80
85	Female	61	0	0	not current	39.36
86	Male	76	0	0	never	31.90
87	Female	43	0	0	never	26.71
88	Male	55	0	0	No Info	27.32
89	Male	57	1	1	not current	27.77
90	Female	43	0	0	never	27.32
91	Male	63	1	0	ever	35.06
92	Male	80	0	0	never	23.25
93	Female	70	0	0	current	29.25
94	Female	42	0	0	never	24.81
95	Female	80	0	0	former	36.18
96	Female	52	1	0	never	50.30
97	Male	71	0	0	never	27.09
98	Female	43	0	0	never	27.32
99	Male	71	0	0	never	27.09
100	Male	80	0	1	former	24.36
101	Male	59	0	0	current	29.20
102	Male	29	0	0	current	25.41
103	Female	68	0	0	No Info	40.31
104	Female	52	0	0	No Info	27.32
105	Male	71	0	0	never	26.53
106	Male	48	1	0	current	36.12
107	Female	79	1	0	former	27.32
108	Male	37	0	0	never	37.24
109	Female	73	0	0	never	35.56
110	Female	59	0	0	former	43.41
111	Female	80	0	0	never	27.32
112	Female	64	0	0	ever	49.27
113	Male	43	0	0	never	39.00
114	Male	43	0	0	never	22.43
115	Male	62	0	0	not current	32.19
116	Male	59	1	0	ever	25.94
117	Female	43	0	0	never	27.73
118	Male	43	0	0	ever	19.46
119	Female	280	0	0	No Info	27.32
120	Female	43	0	0	No Info	27.32

	gender	age	hypertension	heart_disease	smoking_history	bmi
1		6.6		140	0	
2		6.6		80	0	
3		5.7		158	0	
4		5.0		155	0	
5		4.8		155	0	
6		6.6		85	0	
7		5.7		85	0	
8		4.8		145	0	
9		5.0		100	0	
10		6.1		85	0	
11		6.0		100	0	
12		5.0		130	0	
13		5.8		200	0	
14		5.0		160	0	
15		6.6		126	0	
16		6.1		200	0	
17		5.7		158	0	
18		5.7		80	0	
19		3.5		159	0	
20		6.0		90	0	
21		6.2		85	0	
22		4.8		85	0	
23		6.5		130	0	
24		4.0		140	0	
25		6.1		126	0	
26		5.8		80	0	
27		4.0		158	0	
28		6.1		155	0	
29		4.0		200	0	
30		4.0		145	0	
31		6.2		126	0	
32		3.5		100	0	
33		5.0		85	0	
34		5.8		85	0	
35		4.5		130	0	
36		6.6		200	0	
37		6.6		130	0	
38		5.0		159	0	
39		4.8		159	0	
40		3.5		160	0	
41		6.1		100	0	

Console	Terminal	Background Jobs					
R 4.3.1 · C:/Users/ASUS/Downloads/				82	9.0	155	1
				83	7.5	155	1
				84	6.6	280	1
42	5.0	90	0	85	9.0	140	1
43	4.0	159	0	86	7.5	155	1
44	6.2	155	0	87	6.5	300	1
45	6.5	100	0	88	6.8	159	1
46	6.1	155	0	89	6.6	160	1
47	4.0	200	0	90	6.2	155	1
48	5.0	100	0	91	5.8	200	1
49	5.8	140	0	92	6.1	159	1
50	6.5	158	0	93	8.2	130	1
51	6.0	159	0	94	9.0	159	1
52	5.7	100	0	95	6.5	200	1
53	6.5	200	0	96	6.6	155	1
54	5.7	145	0	97	8.2	200	1
55	5.0	90	0	98	6.6	130	1
56	4.8	200	0	99	8.2	200	1
57	5.7	160	0	100	7.5	280	1
58	5.8	158	0	101	8.2	220	1
59	6.2	90	0	102	6.1	130	1
60	6.2	145	0	103	7.5	260	1
61	3.5	126	0	104	9.0	140	1
62	6.6	158	0	105	8.8	159	1
63	5.7	130	0	106	6.8	140	1
64	6.5	159	0	107	6.5	159	1
65	5.7	158	0	108	7.0	126	1
66	5.8	90	0	109	5.8	260	1
67	4.0	160	0	110	6.2	160	1
68	5.7	159	0	111	6.0	200	1
69	6.5	200	1	112	8.2	140	1
70	6.5	200	1	113	8.8	220	1
71	5.7	260	1	114	7.0	160	1
72	9.0	160	1	115	5.8	300	1
73	7.0	159	1	116	9.0	140	1
74	9.0	159	1	117	8.8	145	1
75	8.8	155	1	118	9.0	130	1
76	8.2	126	1	119	8.8	159	1
77	6.2	220	1	120	5.8	159	1
78	7.5	300	1				
79	6.2	159	1				
80	6.8	280	1				
81	6.5	280	1				
82	9.0	155	1				

```
> names(mydata)
[1] "gender"      "age"          "hypertension"  "heart_disease"
[5] "smoking_history" "bmi"          "HbA1c_level"   "blood_glucose_level"
[9] "diabetes"
> |
```

* Identifying Missing Values:

To identify missing value “is.na” function has been used to check for missing values in the variable. The is.na function returns a logical value of TRUE for each missing value in the data and FALSE for all other values.

```
Console Terminal Background Jobs
R 4.3.1 · C:/Users/ASUS/Downloads/

> mydata <- mydata %>% mutate(gender = ifelse(grepl("^\\s*$", gender), NA, gender))
> missingdata <- which(is.na(mydata$gender))
> mydata <- mydata %>% mutate(smoking_history = ifelse(grepl("^\\s*$", smoking_history), NA, smoking_history))
> missingdata_sh <- which(is.na(mydata$smoking_history))
> mydata$smoking_history[mydata$smoking_history == "No Info"] <- NA
> is.na(mydata)
```

The output of this code is a matrix that has the same dimensions as the variable. The matrix contains TRUE values in the cells where the corresponding values are missing and FALSE values in all other cells. This output is useful for identifying any missing values in the dataset and determining the extent to which missing values may affect subsequent analysis:

Console	Terminal	Background Jobs
R 4.3.1 - C:/Users/ASUS/Downloads/		
gender age hypertension heart_disease smoking_history bmi_hdlc_level		
[1,]	FALSE	FALSE
[2,]	FALSE	FALSE
[3,]	FALSE	FALSE
[4,]	FALSE	TRUE
[5,]	FALSE	FALSE
[6,]	FALSE	FALSE
[7,]	TRUE	FALSE
[8,]	FALSE	FALSE
[9,]	FALSE	FALSE
[10,]	FALSE	FALSE
[11,]	FALSE	FALSE
[12,]	FALSE	TRUE
[13,]	FALSE	FALSE
[14,]	FALSE	FALSE
[15,]	TRUE	FALSE
[16,]	FALSE	FALSE
[17,]	FALSE	FALSE
[18,]	FALSE	FALSE
[19,]	FALSE	TRUE
[20,]	FALSE	FALSE
[21,]	FALSE	FALSE
[22,]	FALSE	FALSE
[23,]	FALSE	FALSE
[24,]	FALSE	FALSE
[25,]	FALSE	FALSE
[26,]	FALSE	FALSE
[27,]	FALSE	TRUE
[28,]	FALSE	FALSE
[29,]	FALSE	FALSE
[30,]	FALSE	FALSE
[31,]	FALSE	FALSE
[32,]	FALSE	FALSE
[33,]	FALSE	FALSE
[34,]	FALSE	FALSE
[35,]	FALSE	FALSE
[36,]	FALSE	FALSE
[37,]	FALSE	FALSE
[38,]	FALSE	FALSE
[39,]	FALSE	FALSE
[40,]	FALSE	FALSE
[41,]	FALSE	FALSE

Console	Terminal	Background Jobs
R 4.3.1 - C:/Users/ASUS/Downloads/		
[83,]	FALSE	FALSE
[84,]	FALSE	FALSE
[85,]	FALSE	FALSE
[86,]	FALSE	FALSE
[87,]	FALSE	FALSE
[88,]	FALSE	FALSE
[89,]	FALSE	FALSE
[90,]	FALSE	FALSE
[91,]	FALSE	FALSE
[92,]	FALSE	FALSE
[93,]	FALSE	FALSE
[94,]	FALSE	FALSE
[95,]	FALSE	FALSE
[96,]	FALSE	FALSE
[97,]	FALSE	FALSE
[98,]	FALSE	FALSE
[99,]	FALSE	FALSE
[100,]	FALSE	FALSE
[101,]	FALSE	FALSE
[102,]	FALSE	FALSE
[103,]	FALSE	FALSE
[104,]	FALSE	FALSE
[105,]	FALSE	FALSE
[106,]	FALSE	FALSE
[107,]	FALSE	FALSE
[108,]	FALSE	FALSE
[109,]	FALSE	FALSE
[110,]	FALSE	FALSE
[111,]	FALSE	FALSE
[112,]	FALSE	FALSE
[113,]	FALSE	FALSE
[114,]	FALSE	FALSE
[115,]	FALSE	FALSE
[116,]	FALSE	FALSE
[117,]	FALSE	FALSE
[118,]	FALSE	FALSE
[119,]	FALSE	FALSE
[120,]	FALSE	FALSE
blood_glucose_level diabetes		
[1,]	FALSE	FALSE
[2,]	FALSE	FALSE

Console	Terminal	Background Jobs
R 4.3.1 - C:/Users/ASUS/Downloads/		
[42,]	FALSE	FALSE
[43,]	FALSE	FALSE
[44,]	FALSE	FALSE
[45,]	FALSE	FALSE
[46,]	FALSE	FALSE
[47,]	FALSE	FALSE
[48,]	FALSE	FALSE
[49,]	FALSE	FALSE
[50,]	FALSE	FALSE
[51,]	FALSE	FALSE
[52,]	FALSE	FALSE
[53,]	FALSE	FALSE
[54,]	FALSE	FALSE
[55,]	FALSE	FALSE
[56,]	FALSE	FALSE
[57,]	FALSE	FALSE
[58,]	FALSE	FALSE
[59,]	FALSE	FALSE
[60,]	FALSE	FALSE
[61,]	FALSE	FALSE
[62,]	FALSE	FALSE
[63,]	FALSE	FALSE
[64,]	FALSE	FALSE
[65,]	FALSE	FALSE
[66,]	FALSE	FALSE
[67,]	FALSE	FALSE
[68,]	FALSE	FALSE
[69,]	FALSE	FALSE
[70,]	FALSE	FALSE
[71,]	FALSE	FALSE
[72,]	FALSE	FALSE
[73,]	FALSE	FALSE
[74,]	FALSE	FALSE
[75,]	FALSE	FALSE
[76,]	FALSE	FALSE
[77,]	FALSE	FALSE
[78,]	FALSE	FALSE
[79,]	FALSE	FALSE
[80,]	FALSE	FALSE
[81,]	FALSE	FALSE
[82,]	FALSE	FALSE

Console	Terminal	Background Jobs	
R 4.3.1 · C:/Users/ASUS/Downloads/ ↗			
[4,]		FALSE	FALSE
[5,]		FALSE	FALSE
[6,]		FALSE	FALSE
[7,]		FALSE	FALSE
[8,]		FALSE	FALSE
[9,]		FALSE	FALSE
[10,]		FALSE	FALSE
[11,]		FALSE	FALSE
[12,]		FALSE	FALSE
[13,]		FALSE	FALSE
[14,]		FALSE	FALSE
[15,]		FALSE	FALSE
[16,]		FALSE	FALSE
[17,]		FALSE	FALSE
[18,]		FALSE	FALSE
[19,]		FALSE	FALSE
[20,]		FALSE	FALSE
[21,]		FALSE	FALSE
[22,]		FALSE	FALSE
[23,]		FALSE	FALSE
[24,]		FALSE	FALSE
[25,]		FALSE	FALSE
[26,]		FALSE	FALSE
[27,]		FALSE	FALSE
[28,]		FALSE	FALSE
[29,]		FALSE	FALSE
[30,]		FALSE	FALSE
[31,]		FALSE	FALSE
[32,]		FALSE	FALSE
[33,]		FALSE	FALSE
[34,]		FALSE	FALSE
[35,]		FALSE	FALSE
[36,]		FALSE	FALSE
[37,]		FALSE	FALSE
[38,]		FALSE	FALSE
[39,]		FALSE	FALSE
[40,]		FALSE	FALSE
[41,]		FALSE	FALSE
[42,]		FALSE	FALSE
[43,]		FALSE	FALSE
[44,]		FALSE	FALSE
[45,]		FALSE	FALSE

Console	Terminal	Background Jobs	
R 4.3.1 · C:/Users/ASUS/Downloads/ ↗			
[45,]		FALSE	FALSE
[46,]		FALSE	FALSE
[47,]		FALSE	FALSE
[48,]		FALSE	FALSE
[49,]		FALSE	FALSE
[50,]		FALSE	FALSE
[51,]		FALSE	FALSE
[52,]		FALSE	FALSE
[53,]		FALSE	FALSE
[54,]		FALSE	FALSE
[55,]		FALSE	FALSE
[56,]		FALSE	FALSE
[57,]		FALSE	FALSE
[58,]		FALSE	FALSE
[59,]		FALSE	FALSE
[60,]		FALSE	FALSE
[61,]		FALSE	FALSE
[62,]		FALSE	FALSE
[63,]		FALSE	FALSE
[64,]		FALSE	FALSE
[65,]		FALSE	FALSE
[66,]		FALSE	FALSE
[67,]		FALSE	FALSE
[68,]		FALSE	FALSE
[69,]		FALSE	FALSE
[70,]		FALSE	FALSE
[71,]		FALSE	FALSE
[72,]		FALSE	FALSE
[73,]		FALSE	FALSE
[74,]		FALSE	FALSE
[75,]		FALSE	FALSE
[76,]		FALSE	FALSE
[77,]		FALSE	FALSE
[78,]		FALSE	FALSE
[79,]		FALSE	FALSE
[80,]		FALSE	FALSE
[81,]		FALSE	FALSE
[82,]		FALSE	FALSE
[83,]		FALSE	FALSE
[84,]		FALSE	FALSE
[85,]		FALSE	FALSE
[86,]		FALSE	FALSE

[87,]	FALSE	FALSE
[88,]	FALSE	FALSE
[89,]	FALSE	FALSE
[90,]	FALSE	FALSE
[91,]	FALSE	FALSE
[92,]	FALSE	FALSE
[93,]	FALSE	FALSE
[94,]	FALSE	FALSE
[95,]	FALSE	FALSE
[96,]	FALSE	FALSE
[97,]	FALSE	FALSE
[98,]	FALSE	FALSE
[99,]	FALSE	FALSE
[100,]	FALSE	FALSE
[101,]	FALSE	FALSE
[102,]	FALSE	FALSE
[103,]	FALSE	FALSE
[104,]	FALSE	FALSE
[105,]	FALSE	FALSE
[106,]	FALSE	FALSE
[107,]	FALSE	FALSE
[108,]	FALSE	FALSE
[109,]	FALSE	FALSE
[110,]	FALSE	FALSE
[111,]	FALSE	FALSE
[112,]	FALSE	FALSE
[113,]	FALSE	FALSE
[114,]	FALSE	FALSE
[115,]	FALSE	FALSE
[116,]	FALSE	FALSE
[117,]	FALSE	FALSE
[118,]	FALSE	FALSE
[119,]	FALSE	FALSE
[120,]	FALSE	FALSE

> |

The mydata Output after replacing missing values with NA:

Console Terminal Background Jobs											
R 4.3.1 - C:/Users/ASUS/Downloads/											
> mydata	age	hypertension	heart_disease	smoking_history	bmi	HbA1c_level	bb	0	0	<NA>	29.30
1 Female 80	0	1	never	25.19	6.6	40	Female 67	0	0	<NA>	27.32
2 Female 54	0	0	<NA>	27.32	6.6	41	Female 44	0	0	never	24.93
3 Male 28	0	0	never	27.32	5.7	42	Female 29	0	0	never	19.95
4 Female NA	0	0	current	23.45	5.0	43	Female 60	0	0	never	18.03
5 Male 76	1	1	current	20.14	4.8	44	Female 38	0	0	never	28.27
6 Female 20	0	0	never	27.32	6.6	45	Female 3	0	0	<NA>	19.27
7 <NA> 79	0	0	<NA>	23.86	5.7	46	Male 57	0	0	never	27.32
8 Male 42	0	0	never	33.64	4.8	47	Female 43	0	0	<NA>	27.32
9 Female 32	0	0	never	27.32	5.0	48	Female 74	0	0	<NA>	28.12
10 Female 53	0	0	never	27.32	6.1	49	Female 21	0	0	<NA>	26.10
11 Female 54	0	0	former	54.70	6.0	50	Female 30	0	0	current	27.32
12 Female 78	NA	0	former	36.05	5.0	51	Female 59	0	0	former	27.32
13 Female 67	0	0	never	25.69	5.8	52	Female 290	0	0	not current	30.22
14 Female 76	0	0	<NA>	27.32	5.0	53	Female 59	0	1	ever	23.11
15 <NA> 78	0	0	<NA>	27.32	6.6	54	Female 19	0	0	<NA>	27.32
16 Male 15	0	0	never	30.36	6.1	55	Female NA	0	0	<NA>	28.16
17 Female 42	0	0	never	24.48	5.7	56	Male 56	0	0	never	26.78
18 Female 42	0	0	<NA>	27.32	5.7	57	Male 43	0	0	<NA>	23.04
19 Male NA	0	0	ever	25.72	3.5	58	Male 7	0	0	<NA>	15.94
20 Male 40	0	0	current	36.38	6.0	59	Male 3	0	0	<NA>	15.80
21 Male 5	0	0	<NA>	18.80	6.2	60	Female 30	0	0	never	27.01
22 Female 69	0	0	never	21.24	4.8	61	Male 43	0	0	<NA>	27.32
23 Female 72	0	1	former	27.94	6.5	62	Female 76	0	0	never	22.19
24 Female 4	0	0	<NA>	13.99	4.0	63	Female 41	0	0	never	27.45
25 Male 30	0	0	never	33.76	6.1	64	Female 11	0	0	<NA>	17.98
26 Male 40	0	0	former	27.85	5.8	65	Female 26	0	0	never	26.45
27 Male 45	NA	0	never	26.47	4.0	66	Male 34	0	0	never	31.16
28 Male 43	0	0	never	26.08	6.1	67	Male 80	0	0	former	24.42
29 Female 53	0	0	<NA>	31.75	4.0	68	Female 37	0	0	<NA>	30.50
30 Male 50	0	0	<NA>	25.15	4.0	69	Female 44	0	0	never	19.31
31 Female 41	0	0	current	22.01	6.2	70	Male 67	0	1	not current	27.32
32 Female 20	0	0	never	22.19	3.5	71	Male 50	1	0	current	27.32
33 Female 76	0	0	never	23.55	5.0	72	Male 73	0	0	former	25.91
34 Male 5	0	0	<NA>	15.10	5.8	73	Female 53	0	0	former	27.32
35 Female 15	0	0	<NA>	21.76	4.5	74	Male 50	0	0	former	37.16
36 Female 26	0	0	never	21.22	6.6	75	Female 67	0	0	never	63.48
37 Male 5	0	0	<NA>	27.32	6.6	76	Male 57	0	0	<NA>	27.32
38 Female 77	1	1	never	32.02	5.0	77	Female 36	0	0	current	32.27
39 Female 66	0	0	<NA>	29.30	4.8	78	Female 60	0	0	never	27.32
						79	Female 67	0	0	never	27.32
						80	Female 80	1	0	never	27.32

Console Terminal Background Jobs											
R 4.3.1 - C:/Users/ASUS/Downloads/											
81 Female NA	0	0	never	31.70	6.5						
82 Male 80	0	0	never	22.06	9.0						
83 Female 47	0	0	never	36.49	7.5						
84 Male 53	0	0	current	30.80	6.6						
85 Female 61	0	0	not current	39.36	9.0						
86 Male 76	0	0	never	31.90	7.5						
87 Female 43	0	0	never	26.71	6.5						
88 Male 55	0	0	<NA>	27.32	6.8						
89 Male 57	1	1	not current	27.77	6.6						
90 Female 43	0	0	never	27.32	6.2						
91 Male 63	1	0	ever	35.06	5.8						
92 Male 80	0	0	never	23.25	6.1						
93 Female 70	0	0	current	29.25	8.2						
94 Female 42	0	0	never	24.81	9.0						
95 Female 80	0	0	former	36.18	6.5						
96 Female 52	1	0	never	50.30	6.6						
97 Male 71	0	0	never	27.09	8.2						
98 Female 43	0	0	never	27.32	6.6						
99 Male 71	0	0	never	27.09	8.2						
100 Male 80	0	1	former	24.36	7.5						
101 Male 59	0	0	current	29.20	8.2						
102 Male 29	0	0	current	25.41	6.1						
103 Female 68	0	0	<NA>	40.31	7.5						
104 Female 52	0	0	<NA>	27.32	9.0						
105 Male 71	0	0	never	26.53	8.8						
106 Male 48	1	0	current	36.12	6.8						
107 Female 79	1	0	former	27.32	6.5						
108 Male 37	0	0	never	37.24	7.0						
109 Female 73	0	0	never	35.56	5.8						
110 Female 59	0	0	former	43.41	6.2						
111 Female 80	0	0	never	27.32	6.0						
112 Female 64	0	0	ever	49.27	8.2						
113 Male 43	0	0	never	39.00	8.8						
114 Male 43	0	0	never	22.43	7.0						
115 Male 62	0	0	not current	32.19	5.8						
116 Male 59	1	0	ever	25.94	9.0						
117 Female 43	0	0	never	27.73	8.8						
118 Male 43	0	0	ever	19.46	9.0						
119 Female 280	0	0	<NA>	27.32	8.8						
120 Female 43	0	0	<NA>	27.32	5.8						

blood_glucose_level diabetes

```
R 4.3.1 - C:/Users/ASUS/Downloads/
120 R Console
blood_glucose_level diabetes
1      140      0
2       80      0
3      158      0
4      155      0
5      155      0
6       85      0
7       85      0
8      145      0
9      100      0
10      85      0
11      100      0
12      130      0
13      200      0
14      160      0
15      126      0
16      200      0
17      158      0
18       80      0
19      159      0
20       90      0
21       85      0
22       85      0
23      130      0
24      140      0
25      126      0
26       80      0
27      158      0
28      155      0
29      200      0
30      145      0
31      126      0
32      100      0
33       85      0
34       85      0
35      130      0
36      200      0
37      130      0
38      159      0
39      159      0
40      160      0

82      155      1
83      155      1
84      280      1
85      140      1
86      155      1
87      300      1
88      159      1
89      160      1
90      155      1
91      200      1
92      159      1
93      130      1
94      159      1
95      200      1
96      155      1
97      200      1
98      130      1
99      200      1
100     280      1
101     220      1
102     130      1
103     260      1
104     140      1
105     159      1
106     140      1
107     159      1
108     126      1
109     260      1
110     160      1
111     200      1
112     140      1
113     220      1
114     160      1
115     300      1
116     140      1
117     145      1
118     130      1
119     159      1
120     159      1
> |
```

```
Console Terminal Background Jobs
R 4.3.1 - C:/Users/ASUS/Downloads/
40      160      0
41      100      0
42       90      0
43      159      0
44      155      0
45      100      0
46      155      0
47      200      0
48      100      0
49      140      0
50      158      0
51      159      0
52      100      0
53      200      0
54      145      0
55       90      0
56      200      0
57      160      0
58      158      0
59       90      0
60      145      0
61      126      0
62      158      0
63      130      0
64      159      0
65      158      0
66       90      0
67      160      0
68      159      0
69      200      1
70      200      1
71      260      1
72      160      1
73      159      1
74      159      1
75      155      1
76      126      1
77      220      1
78      300      1
79      159      1
80      280      1
81      280      1
```


Also, there are some missing values based on specific columns and rows.
The outputs are:

```
> which(is.na(mydata$gender))
[1] 7 15
> which(is.na(mydata$age))
[1] 4 19 55 81
> which(is.na(mydata$hypertension))
[1] 12 27
> which(is.na(mydata$smoking_history))
[1] 2 7 14 15 18 21 24 29 30 34 35 37 39 40 45 47 48 49 54 55 57 58
[23] 59 61 64 68 76 88 103 104 119 120
> |
```

Data Exploration:

Missing age values, and any negative ages have been converted to their positive counterparts (absolute values):

```
> mydata <- mydata[!is.na(as.numeric(mydata$age)), ]
> unique(mydata$age)
[1] 80 54 28 76 20 79 42 32 53 78 67 15 40 5 69 72 4 30 45 43 50 41
[23] 26 77 66 44 29 60 38 3 57 74 21 59 290 19 56 7 11 34 37 73 36 47
[45] 61 55 63 70 52 71 68 48 64 62 280
> mydata$age[mydata$age < 0] <- abs(mydata$age[mydata$age < 0])
```

Missing bmi values, and any negative ages have been converted to their positive counterparts (absolute values):

```
> mydata <- mydata[!is.na(as.numeric(mydata$bmi)), ]
> unique(mydata$bmi)
[1] 25.19 27.32 -27.32 20.14 23.86 33.64 54.70 36.05 25.69 30.36 24.48 36.38 18.80
[14] 21.24 27.94 13.99 33.76 27.85 26.47 26.08 31.75 25.15 22.01 22.19 23.55 15.10
[27] 21.76 21.22 32.02 29.30 24.93 19.95 18.03 28.27 19.27 28.12 26.10 30.22 23.11
[40] 26.78 23.04 15.94 15.80 27.01 27.45 17.98 26.45 31.16 24.42 30.50 19.31 25.91
[53] 37.16 63.48 32.27 22.06 36.49 30.80 39.36 31.90 26.71 27.77 35.06 23.25 29.25
[66] 24.81 36.18 50.30 27.09 24.36 29.20 25.41 40.31 26.53 36.12 37.24 35.56 43.41
[79] 49.27 39.00 22.43 32.19 25.94 27.73 19.46
> mydata$bmi[mydata$bmi < 0] <- abs(mydata$bmi[mydata$bmi < 0])
```

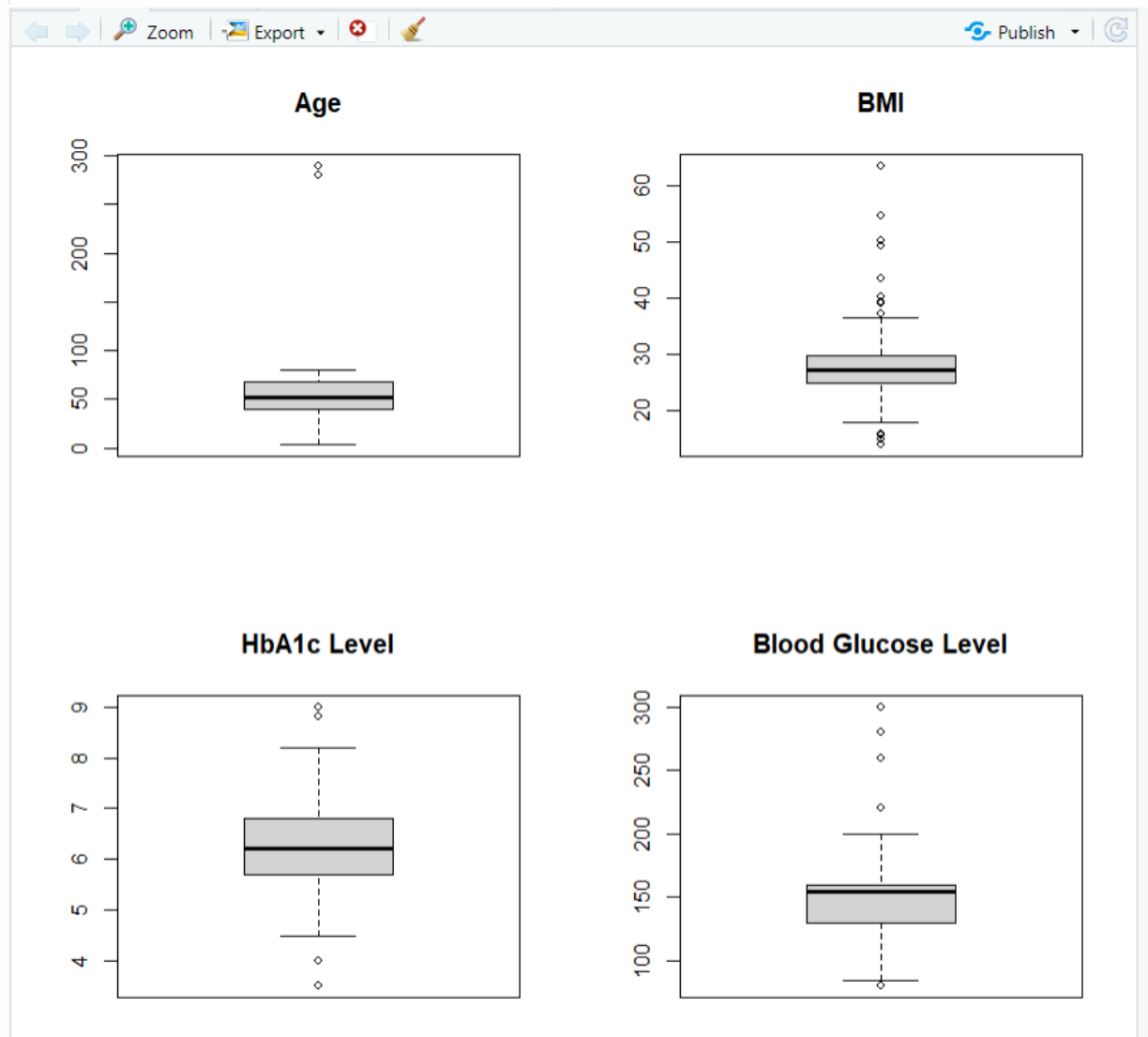
Handling potential outliers using box plots:

This will create a 2x2 grid of boxplots to visualize the distribution and spread of data for four different variables in the mydata dataframe. The output:


```

par(mfrow = c(2, 2))
boxplot(mydata$age, main = "Age")
boxplot(mydata$bmi, main = "BMI")
boxplot(mydata$HbA1c_level, main = "HbA1c Level")
boxplot(mydata$blood_glucose_level, main = "Blood Glucose Level")

```



From the above boxplots it is noticeable that there are few outliers present in the dataset.

Dealing with missing values:

We replaced the main data frame “mydata” each time whenever we performed a specific task on the dataframe so that there would be backup data as mydata. Replacing missing values with the mean value. The output:

```
mydata_mean <- mydata
mean_age <- mean(mydata_mean$age, na.rm = TRUE)
mydata_mean$age[is.na(mydata_mean$age)] <- mean_age
print(mydata_mean)
```

	gender	age	hypertension	heart_disease	smoking_history	bmi	HbA1c_level
1	Female	80	0	1	never	25.19	6.6
2	Female	54	0	0	<NA>	27.32	6.6
3	Male	28	0	0	never	27.32	5.7
5	Male	76	1	1	current	20.14	4.8
6	Female	20	0	0	never	27.32	6.6
7	<NA>	79	0	0	<NA>	23.86	5.7
8	Male	42	0	0	never	33.64	4.8
9	Female	32	0	0	never	27.32	5.0
10	Female	53	0	0	never	27.32	6.1
11	Female	54	0	0	former	54.70	6.0
12	Female	78	NA	0	former	36.05	5.0
13	Female	67	0	0	never	25.69	5.8
14	Female	76	0	0	<NA>	27.32	5.0
15	<NA>	78	0	0	<NA>	27.32	6.6
16	Male	15	0	0	never	30.36	6.1
17	Female	42	0	0	never	24.48	5.7
18	Female	42	0	0	<NA>	27.32	5.7
20	Male	40	0	0	current	36.38	6.0
21	Male	5	0	0	<NA>	18.80	6.2
22	Female	69	0	0	never	21.24	4.8
23	Female	72	0	1	former	27.94	6.5
24	Female	4	0	0	<NA>	13.99	4.0
25	Male	30	0	0	never	33.76	6.1
26	Male	40	0	0	former	27.85	5.8
27	Male	45	NA	0	never	26.47	4.0
28	Male	43	0	0	never	26.08	6.1
29	Female	53	0	0	<NA>	31.75	4.0
30	Male	50	0	0	<NA>	25.15	4.0
31	Female	41	0	0	current	22.01	6.2
32	Female	20	0	0	never	22.19	3.5
33	Female	76	0	0	never	23.55	5.0
34	Male	5	0	0	<NA>	15.10	5.8
35	Female	15	0	0	<NA>	21.76	4.5
36	Female	26	0	0	never	21.22	6.6
37	Male	5	0	0	<NA>	27.32	6.6
38	Female	77	1	1	never	32.02	5.0
39	Female	66	0	0	<NA>	29.30	4.8
40	Female	67	0	0	<NA>	27.32	3.5
41	Female	44	0	0	never	24.93	6.1
42	Female	29	0	0	never	19.95	5.0

	gender	age	hypertension	heart_disease	smoking_history	bmi	HbA1c_level
43	Female	60	0	0	never	18.03	4.0
44	Female	38	0	0	never	28.27	6.2
45	Female	3	0	0	<NA>	19.27	6.5
46	Male	57	0	0	never	27.32	6.1
47	Female	43	0	0	<NA>	27.32	4.0
48	Female	74	0	0	<NA>	28.12	5.0
49	Female	21	0	0	<NA>	26.10	5.8
50	Female	30	0	0	current	27.32	6.5
51	Female	59	0	0	former	27.32	6.0
52	Female	290	0	0	not current	30.22	5.7
53	Female	59	0	1	ever	23.11	6.5
54	Female	19	0	0	<NA>	27.32	5.7
56	Male	56	0	0	never	26.78	4.8
57	Male	43	0	0	<NA>	23.04	5.7
58	Male	7	0	0	<NA>	15.94	5.8
59	Male	3	0	0	<NA>	15.80	6.2
60	Female	30	0	0	never	27.01	6.2
61	Male	43	0	0	<NA>	27.32	3.5
62	Female	76	0	0	never	22.19	6.6
63	Female	41	0	0	never	27.45	5.7
64	Female	11	0	0	<NA>	17.98	6.5
65	Female	26	0	0	never	26.45	5.7
66	Male	34	0	0	never	31.16	5.8
67	Male	80	0	0	former	24.42	4.0
68	Female	37	0	0	<NA>	30.50	5.7
69	Female	44	0	0	never	19.31	6.5
70	Male	67	0	1	not current	27.32	6.5
71	Male	50	1	0	current	27.32	5.7
72	Male	73	0	0	former	25.91	9.0
73	Female	53	0	0	former	27.32	7.0
74	Male	50	0	0	former	37.16	9.0
75	Female	67	0	0	never	63.48	8.8
76	Male	57	0	0	<NA>	27.32	8.2
77	Female	36	0	0	current	32.27	6.2
78	Female	60	0	0	never	27.32	7.5
79	Female	67	0	0	never	27.32	6.2
80	Female	80	1	0	never	27.32	6.8
82	Male	80	0	0	never	22.06	9.0
83	Female	47	0	0	never	36.49	7.5
84	Male	53	0	0	current	30.80	6.6
85	Female	61	0	0	not current	39.36	9.0
86	Male	76	0	0	never	31.90	7.5

	gender	age	hypertension	heart_disease	smoking_history	bmi	HbA1c_level
87	Female	43	0	0	never	26.71	6.5
88	Male	55	0	0	<NA>	27.32	6.8
89	Male	57	1	1	not current	27.77	6.6
90	Female	43	0	0	never	27.32	6.2
91	Male	63	1	0	ever	35.06	5.8
92	Male	80	0	0	never	23.25	6.1
93	Female	70	0	0	current	29.25	8.2
94	Female	42	0	0	never	24.81	9.0
95	Female	80	0	0	former	36.18	6.5
96	Female	52	1	0	never	50.30	6.6
97	Male	71	0	0	never	27.09	8.2
98	Female	43	0	0	never	27.32	6.6
99	Male	71	0	0	never	27.09	8.2
100	Male	80	0	1	former	24.36	7.5
101	Male	59	0	0	current	29.20	8.2
102	Male	29	0	0	current	25.41	6.1
103	Female	68	0	0	<NA>	40.31	7.5
104	Female	52	0	0	<NA>	27.32	9.0
105	Male	71	0	0	never	26.53	8.8
106	Male	48	1	0	current	36.12	6.8
107	Female	79	1	0	former	27.32	6.5
108	Male	37	0	0	never	37.24	7.0
109	Female	73	0	0	never	35.56	5.8
110	Female	59	0	0	former	43.41	6.2
111	Female	80	0	0	never	27.32	6.0
112	Female	64	0	0	ever	49.27	8.2
113	Male	43	0	0	never	39.00	8.8
114	Male	43	0	0	never	22.43	7.0
115	Male	62	0	0	not current	32.19	5.8
116	Male	59	1	0	ever	25.94	9.0
117	Female	43	0	0	never	27.73	8.8
118	Male	43	0	0	ever	19.46	9.0
119	Female	280	0	0	<NA>	27.32	8.8
120	Female	43	0	0	<NA>	27.32	5.8

	blood_glucose_level	diabetes
1	140	0
2	80	0
3	158	0
5	155	0
6	85	0
7	85	0

Console	Terminal	Background Jobs
R 4.3.1 · C:/Users/ASUS/Downloads/		
8	145	0
9	100	0
10	85	0
11	100	0
12	130	0
13	200	0
14	160	0
15	126	0
16	200	0
17	158	0
18	80	0
20	90	0
21	85	0
22	85	0
23	130	0
24	140	0
25	126	0
26	80	0
27	158	0
28	155	0
29	200	0
30	145	0
31	126	0
32	100	0
33	85	0
34	85	0
35	130	0
36	200	0
37	130	0
38	159	0
39	159	0
40	160	0
41	100	0
42	90	0
43	159	0
44	155	0
45	100	0
46	155	0
47	200	0
48	100	0
49	140	0
50	158	0

Console	Terminal	Background Jobs	
R 4.3.1 · C:/Users/ASUS/Downloads/			
51		159	0
52		100	0
53		200	0
54		145	0
56		200	0
57		160	0
58		158	0
59		90	0
60		145	0
61		126	0
62		158	0
63		130	0
64		159	0
65		158	0
66		90	0
67		160	0
68		159	0
69		200	1
70		200	1
71		260	1
72		160	1
73		159	1
74		159	1
75		155	1
76		126	1
77		220	1
78		300	1
79		159	1
80		280	1
82		155	1
83		155	1
84		280	1
85		140	1
86		155	1
87		300	1
88		159	1
89		160	1
90		155	1
91		200	1
92		159	1
93		130	1
94		159	1

95	200	1
96	155	1
97	200	1
98	130	1
99	200	1
100	280	1
101	220	1
102	130	1
103	260	1
104	140	1
105	159	1
106	140	1
107	159	1
108	126	1
109	260	1
110	160	1
111	200	1
112	140	1
113	220	1
114	160	1
115	300	1
116	140	1
117	145	1
118	130	1
119	159	1
120	159	1
>		

Check for missing values and count:

```

> missing_count <- colSums(is.na(mydata))
> missing_count
      gender      age hypertension      heart_disease
      2      0      2      0
smoking_history      bmi      HbA1c_level blood_glucose_level
      31      0      0      0
      diabetes
      0

```

Data Visualization:

ggplot2 library, which was used for creating data visualizations. We replaced mydata data frame with mydata_remove. Then we removed rows with missing values, ensuring that the dataset used for analysis and visualization is complete and doesn't introduce bias or inaccuracies due to missing data. The output:

```
> library(ggplot2)
> mydata_remove <- mydata
> mydata_remove <- mydata_remove[complete.cases(mydata_remove), ]
> mydata_remove
```

	gender	age	hypertension	heart_disease	smoking_history	bmi	HbA1c_level
1	Female	80	0	1	never	25.19	6.6
3	Male	28	0	0	never	27.32	5.7
5	Male	76	1	1	current	20.14	4.8
6	Female	20	0	0	never	27.32	6.6
8	Male	42	0	0	never	33.64	4.8
9	Female	32	0	0	never	27.32	5.0
10	Female	53	0	0	never	27.32	6.1
11	Female	54	0	0	former	54.70	6.0
13	Female	67	0	0	never	25.69	5.8
16	Male	15	0	0	never	30.36	6.1
17	Female	42	0	0	never	24.48	5.7
20	Male	40	0	0	current	36.38	6.0
22	Female	69	0	0	never	21.24	4.8
23	Female	72	0	1	former	27.94	6.5
25	Male	30	0	0	never	33.76	6.1
26	Male	40	0	0	former	27.85	5.8
28	Male	43	0	0	never	26.08	6.1
31	Female	41	0	0	current	22.01	6.2
32	Female	20	0	0	never	22.19	3.5
33	Female	76	0	0	never	23.55	5.0
36	Female	26	0	0	never	21.22	6.6
38	Female	77	1	1	never	32.02	5.0
41	Female	44	0	0	never	24.93	6.1
42	Female	29	0	0	never	19.95	5.0
43	Female	60	0	0	never	18.03	4.0
44	Female	38	0	0	never	28.27	6.2
46	Male	57	0	0	never	27.32	6.1
50	Female	30	0	0	current	27.32	6.5
51	Female	59	0	0	former	27.32	6.0
52	Female	290	0	0	not current	30.22	5.7
53	Female	59	0	1	ever	23.11	6.5
56	Male	56	0	0	never	26.78	4.8
60	Female	30	0	0	never	27.01	6.2
62	Female	76	0	0	never	22.19	6.6
63	Female	41	0	0	never	27.45	5.7
65	Female	26	0	0	never	26.45	5.7

	gender	age	hypertension	heart_disease	smoking_history	bmi	HbA1c_level
65	Female	26	0	0	never	26.45	5.7
66	Male	34	0	0	never	31.16	5.8
67	Male	80	0	0	former	24.42	4.0
69	Female	44	0	0	never	19.31	6.5
70	Male	67	0	1	not current	27.32	6.5
71	Male	50	1	0	current	27.32	5.7
72	Male	73	0	0	former	25.91	9.0
73	Female	53	0	0	former	27.32	7.0
74	Male	50	0	0	former	37.16	9.0
75	Female	67	0	0	never	63.48	8.8
77	Female	36	0	0	current	32.27	6.2
78	Female	60	0	0	never	27.32	7.5
79	Female	67	0	0	never	27.32	6.2
80	Female	80	1	0	never	27.32	6.8
82	Male	80	0	0	never	22.06	9.0
83	Female	47	0	0	never	36.49	7.5
84	Male	53	0	0	current	30.80	6.6
85	Female	61	0	0	not current	39.36	9.0
86	Male	76	0	0	never	31.90	7.5
87	Female	43	0	0	never	26.71	6.5
89	Male	57	1	1	not current	27.77	6.6
90	Female	43	0	0	never	27.32	6.2
91	Male	63	1	0	ever	35.06	5.8
92	Male	80	0	0	never	23.25	6.1
93	Female	70	0	0	current	29.25	8.2
94	Female	42	0	0	never	24.81	9.0
95	Female	80	0	0	former	36.18	6.5
96	Female	52	1	0	never	50.30	6.6
97	Male	71	0	0	never	27.09	8.2
98	Female	43	0	0	never	27.32	6.6
99	Male	71	0	0	never	27.09	8.2
100	Male	80	0	1	former	24.36	7.5
101	Male	59	0	0	current	29.20	8.2
102	Male	29	0	0	current	25.41	6.1
105	Male	71	0	0	never	26.53	8.8
106	Male	48	1	0	current	36.12	6.8
107	Female	79	1	0	former	27.32	6.5
108	Male	37	0	0	never	37.24	7.0
109	Female	73	0	0	never	35.56	5.8
110	Female	59	0	0	former	43.41	6.2
111	Female	80	0	0	never	27.32	6.0
112	Female	64	0	0	ever	49.27	8.2
113	Male	43	0	0	never	39.00	8.8
114	Male	43	0	0	never	22.43	7.0
115	Male	62	0	0	not current	32.19	5.8
116	Male	59	1	0	ever	25.94	9.0
117	Female	43	0	0	never	27.73	8.8
118	Male	43	0	0	ever	19.46	9.0

```
Console Terminal Background Jobs
R 4.3.1 - C:/Users/ASUS/Downloads/
112 Female 64 0 0 ever 49.27 8.2
113 Male 43 0 0 never 39.00 8.8
114 Male 43 0 0 never 22.43 7.0
115 Male 62 0 0 not current 32.19 5.8
116 Male 59 1 0 ever 25.94 9.0
117 Female 43 0 0 never 27.73 8.8
118 Male 43 0 0 ever 19.46 9.0
blood_glucose_level diabetes
1 140 0
3 158 0
5 155 0
6 85 0
8 145 0
9 100 0
10 85 0
11 100 0
13 200 0
16 200 0
17 158 0
20 90 0
22 85 0
23 130 0
25 126 0
26 80 0
28 155 0
31 126 0
32 100 0
33 85 0
36 200 0
38 159 0
41 100 0
42 90 0
43 159 0
44 155 0
46 155 0
50 158 0
51 159 0
52 100 0
53 200 0
56 200 0
60 145 0
```

```
Console Terminal Background Jobs
R 4.3.1 - C:/Users/ASUS/Downloads/
62 158 0
63 130 0
65 158 0
66 90 0
67 160 0
69 200 1
70 200 1
71 260 1
72 160 1
73 159 1
74 159 1
75 155 1
77 220 1
78 300 1
79 159 1
80 280 1
82 155 1
83 155 1
84 280 1
85 140 1
86 155 1
87 300 1
89 160 1
90 155 1
91 200 1
92 159 1
93 130 1
94 159 1
95 200 1
96 155 1
97 200 1
98 130 1
99 200 1
100 280 1
101 220 1
102 130 1
105 159 1
106 140 1
107 159 1
108 126 1
109 260 1
```

```

110          160      1
111          200      1
112          140      1
113          220      1
114          160      1
115          300      1
116          140      1
117          145      1
118          130      1

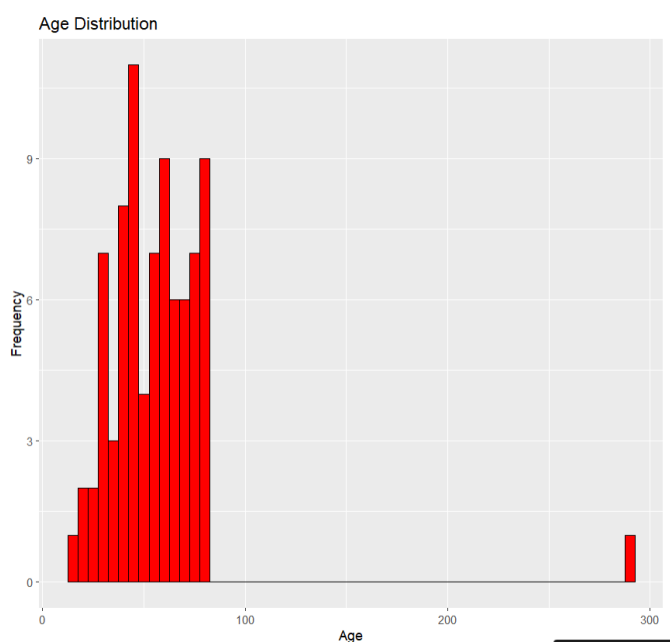
```

To indentify pattern trend in the dataset .The output:

```

> ggplot(mydata_remove, aes(x = age)) +
+ geom_histogram(binwidth = 5, fill = "red", color = "black") +
+ labs(x = "Age", y = "Frequency", title = "Age Distribution")
> |

```



Checking for missing values and count.The output:

```

> missing_count <- colSums(is.na(mydata))
> missing_count
      gender      age hypertension      heart_disease
      2         0         2             0
smoking_history      bmi  HbA1c_level blood_glucose_level
      31         0         0             0
      diabetes
      0

```

Replacing missing values in the column with the mode. The output:

```
R 4.3.1 ~ C:/Users/ASUS/Downloads/ ↵
```

```
> mydata_mode <- mydata
> mode_gender <- names(sort(table(mydata_mode$gender), decreasing = TRUE))[1]
> mydata_mode$gender[is.na(mydata_mode$gender)] <- mode_gender
> mode_age <- names(sort(table(mydata_mode$age), decreasing = TRUE))[1]
> mydata_mode$age[is.na(mydata_mode$age)] <- mode_age
> mode_hypertension <- names(sort(table(mydata_mode$hypertension), decreasing = TRUE))[1]
> mydata_mode$hypertension[is.na(mydata_mode$hypertension)] <- mode_hypertension
> mode_hsmoking <- names(sort(table(mydata_mode$smoking_history), decreasing = TRUE))[1]
> mydata_mode$smoking_history[is.na(mydata_mode$smoking_history)] <- mode_hsmoking
> mydata_mode
```

	gender	age	hypertension	heart_disease	smoking_history	bmi	HbA1c_level
1	Female	80	0	1	never	25.19	6.6
2	Female	54	0	0	never	27.32	6.6
3	Male	28	0	0	never	27.32	5.7
5	Male	76	1	1	current	20.14	4.8
6	Female	20	0	0	never	27.32	6.6
7	Female	79	0	0	never	23.86	5.7
8	Male	42	0	0	never	33.64	4.8
9	Female	32	0	0	never	27.32	5.0
10	Female	53	0	0	never	27.32	6.1
11	Female	54	0	0	former	54.70	6.0
12	Female	78	0	0	former	36.05	5.0
13	Female	67	0	0	never	25.69	5.8
14	Female	76	0	0	never	27.32	5.0
15	Female	78	0	0	never	27.32	6.6
16	Male	15	0	0	never	30.36	6.1
17	Female	42	0	0	never	24.48	5.7
18	Female	42	0	0	never	27.32	5.7
20	Male	40	0	0	current	36.38	6.0
21	Male	5	0	0	never	18.80	6.2
22	Female	69	0	0	never	21.24	4.8
23	Female	72	0	1	former	27.94	6.5
24	Female	4	0	0	never	13.99	6.0
25	Male	30	0	0	never	33.76	4.1
26	Male	40	0	0	former	27.85	5.8
27	Male	45	0	0	never	26.47	4.0
28	Male	43	0	0	never	26.08	6.1
29	Female	53	0	0	never	31.75	4.0
30	Male	50	0	0	never	25.15	4.0
31	Female	41	0	0	current	22.01	6.2

[illegible]

74	Male	50	0	0	former	37.16	9.0
75	Female	67	0	0	never	63.48	8.8
76	Male	57	0	0	never	27.32	8.2
77	Female	36	0	0	current	32.27	6.2
78	Female	60	0	0	never	27.32	7.5
79	Female	67	0	0	never	27.32	6.2
80	Female	80	1	0	never	27.32	6.8
82	Male	80	0	0	never	22.06	9.0
83	Female	47	0	0	never	36.49	7.5
84	Male	53	0	0	current	30.80	6.6
85	Female	61	0	0	not current	39.36	9.0
86	Male	76	0	0	never	31.90	7.5
87	Female	43	0	0	never	26.71	6.5
88	Male	55	0	0	never	27.32	6.8
89	Male	57	1	1	not current	27.77	6.6
90	Female	43	0	0	never	27.32	6.2
91	Male	63	1	0	ever	35.06	5.8
92	Male	80	0	0	never	23.25	6.1
93	Female	70	0	0	current	29.25	8.2
94	Female	42	0	0	never	24.81	9.0
95	Female	80	0	0	former	36.18	6.5
96	Female	52	1	0	never	50.30	6.6
97	Male	71	0	0	never	27.09	8.2
98	Female	43	0	0	never	27.32	6.6
99	Male	71	0	0	never	27.09	8.2
100	Male	80	0	1	former	24.36	7.5
101	Male	59	0	0	current	29.20	8.2
102	Male	29	0	0	current	25.41	6.1
103	Female	68	0	0	never	40.31	7.5
104	Female	52	0	0	never	27.32	9.0
105	Male	71	0	0	never	26.53	8.8
106	Male	48	1	0	current	36.12	6.8
107	Female	79	1	0	former	27.32	6.5
108	Male	37	0	0	never	37.24	7.0
109	Female	73	0	0	never	35.56	5.8
110	Female	59	0	0	former	43.41	6.2
111	Female	80	0	0	never	27.32	6.0
112	Female	64	0	0	ever	49.27	8.2
113	Male	43	0	0	never	39.00	8.8
114	Male	43	0	0	never	22.43	7.0
115	Male	62	0	0	not current	32.19	5.8

116	Male	59	1	0	ever	25.94	9.0
117	Female	43	0	0	never	27.73	8.8
118	Male	43	0	0	ever	19.46	9.0
119	Female	280	0	0	never	27.32	8.8
120	Female	43	0	0	never	27.32	5.8
blood_glucose_level diabetes							
1		140	0				
2		80	0				
3		158	0				
5		155	0				
6		85	0				
7		85	0				
8		145	0				
9		100	0				
10		85	0				
11		100	0				
12		130	0				
13		200	0				
14		160	0				
15		126	0				
16		200	0				
17		158	0				
18		80	0				
20		90	0				
21		85	0				
22		85	0				
23		130	0				
24		140	0				
25		126	0				
26		80	0				
27		158	0				
28		155	0				
29		200	0				
30		145	0				
31		126	0				
32		100	0				
33		85	0				
34		85	0				
35		130	0				
36		200	0				

Console	Terminal	Background Jobs	
R 4.3.1 · C:/Users/ASUS/Downloads/			
38	159	0	
39	159	0	
40	160	0	
41	100	0	
42	90	0	
43	159	0	
44	155	0	
45	100	0	
46	155	0	
47	200	0	
48	100	0	
49	140	0	
50	158	0	
51	159	0	
52	100	0	
53	200	0	
54	145	0	
56	200	0	
57	160	0	
58	158	0	
59	90	0	
60	145	0	
61	126	0	
62	158	0	
63	130	0	
64	159	0	
65	158	0	
66	90	0	
67	160	0	
68	159	0	
69	200	1	
70	200	1	
71	260	1	
72	160	1	
73	159	1	
74	159	1	
75	155	1	
76	126	1	
77	220	1	
78	300	1	
79	159	1	
80	280	1	

82	155	1
83	155	1
84	280	1
85	140	1
86	155	1
87	300	1
88	159	1
89	160	1
90	155	1
91	200	1
92	159	1
93	130	1
94	159	1
95	200	1
96	155	1
97	200	1
98	130	1
99	200	1
100	280	1
101	220	1
102	130	1
103	260	1
104	140	1
105	159	1
106	140	1
107	159	1
108	126	1
109	260	1
110	160	1
111	200	1
112	140	1
113	220	1
114	160	1
115	300	1
116	140	1
117	145	1
118	130	1
119	159	1
120	159	1

```

> mydata_median <- mydata
> median(mydata_median$age)
[1] 52.5
> mydata_median
  gender age hypertension heart_disease smoking_history   bmi HbA1c_level
1  Female 80           0             1      never 25.19      6.6
2  Female 54           0             1      <NA> 27.32      6.6
3   Male 28           0             0      never 27.32      5.7
5   Male 76           1             1    current 20.14      4.8
6  Female 20           0             0      never 27.32      6.6
7   <NA> 79           0             0      <NA> 23.86      5.7
8   Male 42           0             0      never 33.64      4.8
9  Female 32           0             0      never 27.32      5.0
10 Female 53           0             0      never 27.32      6.1
11 Female 54           0             0    former 54.70      6.0
12 Female 78          NA             0    former 36.05      5.0
13 Female 67           0             0      never 25.69      5.8
14 Female 76           0             0      <NA> 27.32      5.0
15   <NA> 78           0             0      <NA> 27.32      6.6
16   Male 15           0             0      never 30.36      6.1
17 Female 42           0             0      never 24.48      5.7
18 Female 42           0             0      <NA> 27.32      5.7
20   Male 40           0             0    current 36.38      6.0
21   Male 5            0             0      <NA> 18.80      6.2
22 Female 69           0             0      never 21.24      4.8
23 Female 72           0             1    former 27.94      6.5
24 Female 4            0             0      <NA> 13.99      4.0
25   Male 30           0             0      never 33.76      6.1
26   Male 40           0             0    former 27.85      5.8
27   Male 45          NA             0      never 26.47      4.0
28   Male 43           0             0      never 26.08      6.1
29 Female 53           0             0      <NA> 31.75      4.0
30   Male 50           0             0      <NA> 25.15      4.0
31 Female 41           0             0    current 22.01      6.2
32 Female 20           0             0      never 22.19      3.5
33 Female 76           0             0      never 23.55      5.0
34   Male 5            0             0      <NA> 15.10      5.8
35 Female 15           0             0      <NA> 21.76      4.5
36 Female 26           0             0      never 21.22      6.6
37   Male 5            0             0      <NA> 27.32      6.6
38 Female 77           1             1      never 32.02      5.0

```

39	Female	66	0	0	<NA>	29.30	4.8
40	Female	67	0	0	<NA>	27.32	3.5
41	Female	44	0	0	never	24.93	6.1
42	Female	29	0	0	never	19.95	5.0
43	Female	60	0	0	never	18.03	4.0
44	Female	38	0	0	never	28.27	6.2
45	Female	3	0	0	<NA>	19.27	6.5
46	Male	57	0	0	never	27.32	6.1
47	Female	43	0	0	<NA>	27.32	4.0
48	Female	74	0	0	<NA>	28.12	5.0
49	Female	21	0	0	<NA>	26.10	5.8
50	Female	30	0	0	current	27.32	6.5
51	Female	59	0	0	former	27.32	6.0
52	Female	290	0	0	not current	30.22	5.7
53	Female	59	0	1	ever	23.11	6.5
54	Female	19	0	0	<NA>	27.32	5.7
56	Male	56	0	0	never	26.78	4.8
57	Male	43	0	0	<NA>	23.04	5.7
58	Male	7	0	0	<NA>	15.94	5.8
59	Male	3	0	0	<NA>	15.80	6.2
60	Female	30	0	0	never	27.01	6.2
61	Male	43	0	0	<NA>	27.32	3.5
62	Female	76	0	0	never	22.19	6.6
63	Female	41	0	0	never	27.45	5.7
64	Female	11	0	0	<NA>	17.98	6.5
65	Female	26	0	0	never	26.45	5.7
66	Male	34	0	0	never	31.16	5.8
67	Male	80	0	0	former	24.42	4.0
68	Female	37	0	0	<NA>	30.50	5.7
69	Female	44	0	0	never	19.31	6.5
70	Male	67	0	1	not current	27.32	6.5
71	Male	50	1	0	current	27.32	5.7
72	Male	73	0	0	former	25.91	9.0
73	Female	53	0	0	former	27.32	7.0
74	Male	50	0	0	former	37.16	9.0
75	Female	67	0	0	never	63.48	8.8
76	Male	57	0	0	<NA>	27.32	8.2
77	Female	36	0	0	current	32.27	6.2
78	Female	60	0	0	never	27.32	7.5
79	Female	67	0	0	never	27.32	6.2
80	Female	80	1	0	never	27.32	6.8
82	Male	80	0	0	never	22.06	9.0

83	Female	47	0	0	never	36.49	7.5	R 4.3.1 · C:/Users/ASUS/Downloads/	3	158	0
84	Male	53	0	0	current	30.80	6.6		5	155	0
85	Female	61	0	0	not current	39.36	9.0		6	85	0
86	Male	76	0	0	never	31.90	7.5		7	85	0
87	Female	43	0	0	never	26.71	6.5		8	145	0
88	Male	55	0	0	<NA>	27.32	6.8		9	100	0
89	Male	57	1	1	not current	27.77	6.6		10	85	0
90	Female	43	0	0	never	27.32	6.2		11	100	0
91	Male	63	1	0	ever	35.06	5.8		12	130	0
92	Male	80	0	0	never	23.25	6.1		13	200	0
93	Female	70	0	0	current	29.25	8.2		14	160	0
94	Female	42	0	0	never	24.81	9.0		15	126	0
95	Female	80	0	0	former	36.18	6.5		16	200	0
96	Female	52	1	0	never	50.30	6.6		17	158	0
97	Male	71	0	0	never	27.09	8.2		18	80	0
98	Female	43	0	0	never	27.32	6.6		20	90	0
99	Male	71	0	0	never	27.09	8.2		21	85	0
100	Male	80	0	1	former	24.36	7.5		22	85	0
101	Male	59	0	0	current	29.20	8.2		23	130	0
102	Male	29	0	0	current	25.41	6.1		24	140	0
103	Female	68	0	0	<NA>	40.31	7.5		25	126	0
104	Female	52	0	0	<NA>	27.32	9.0		26	80	0
105	Male	71	0	0	never	26.53	8.8		27	158	0
106	Male	48	1	0	current	36.12	6.8		28	155	0
107	Female	79	1	0	former	27.32	6.5		29	200	0
108	Male	37	0	0	never	37.24	7.0		30	145	0
109	Female	73	0	0	never	35.56	5.8		31	126	0
110	Female	59	0	0	former	43.41	6.2		32	100	0
111	Female	80	0	0	never	27.32	6.0		33	85	0
112	Female	64	0	0	ever	49.27	8.2		34	85	0
113	Male	43	0	0	never	39.00	8.8		35	130	0
114	Male	43	0	0	never	22.43	7.0		36	200	0
115	Male	62	0	0	not current	32.19	5.8		37	130	0
116	Male	59	1	0	ever	25.94	9.0		38	159	0
117	Female	43	0	0	never	27.73	8.8		39	159	0
118	Male	43	0	0	ever	19.46	9.0		40	160	0
119	Female	280	0	0	<NA>	27.32	8.8		41	100	0
120	Female	43	0	0	<NA>	27.32	5.8		42	90	0
blood_glucose_level diabetes									43	159	0
1			140	0					44	155	0
2									45	100	0
									46	155	0

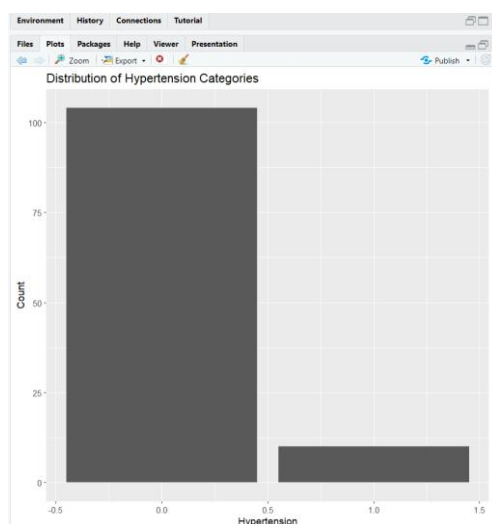
47	200	0	90	155	1
48	100	0	91	200	1
49	140	0	92	159	1
50	158	0	93	130	1
51	159	0	94	159	1
52	100	0	95	200	1
53	200	0	96	155	1
54	145	0	97	200	1
56	200	0	98	130	1
57	160	0	99	200	1
58	158	0	100	280	1
59	90	0	101	220	1
60	145	0	102	130	1
61	126	0	103	260	1
62	158	0	104	140	1
63	130	0	105	159	1
64	159	0	106	140	1
65	158	0	107	159	1
66	90	0	108	126	1
67	160	0	109	260	1
68	159	0	110	160	1
69	200	1	111	200	1
70	200	1	112	140	1
71	260	1	113	220	1
72	160	1	114	160	1
73	159	1	115	300	1
74	159	1	116	140	1
75	155	1	117	145	1
76	126	1	118	130	1
77	220	1	119	159	1
78	300	1	120	159	1
79	159	1			
80	280	1			
82	155	1			
83	155	1			
84	280	1			
85	140	1			
86	155	1			
87	300	1			
88	159	1			
89	159	1			

Data cleaning by counting the total missing column. The output:

```
> missing_count <- colSums(is.na(mydata))
> missing_count
      gender      age      hypertension      heart_disease
      2        0        2                0
smoking_history      bmi      HbA1c_level blood_glucose_level
      31        0        0                0
      diabetes
      0
```

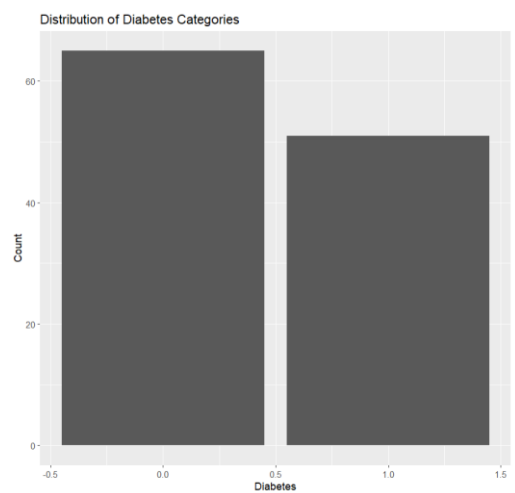
A visual representation of the counts of different hypertension categories, that will help to understand the distribution of this variable in mydata. The output:

```
C:\Users\ASUS\AppData\Local\Temp\RtmpyHtELK\downloaded_packages
> library(ggplot2)
Warning message:
package 'ggplot2' was built under R version 4.3.2
> ggplot(mydata, aes(x = hypertension)) +
+   geom_bar() +
+   labs(x = "Hypertension", y = "Count", title = "Distribution of Hypertension Categories")
```



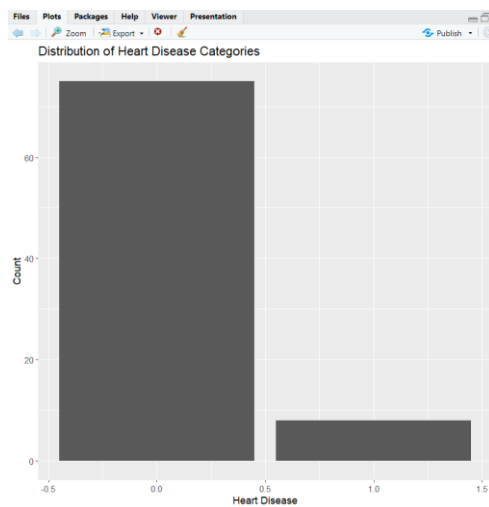
A visual representation of the counts of different diabetes categories, that will help to understand the distribution of this variable in my data. The output:

```
> ggplot(mydata, aes(x = diabetes)) +
+   geom_bar() +
+   labs(x = "Diabetes", y = "Count", title = "Distribution of Diabetes Categories")
> |
```



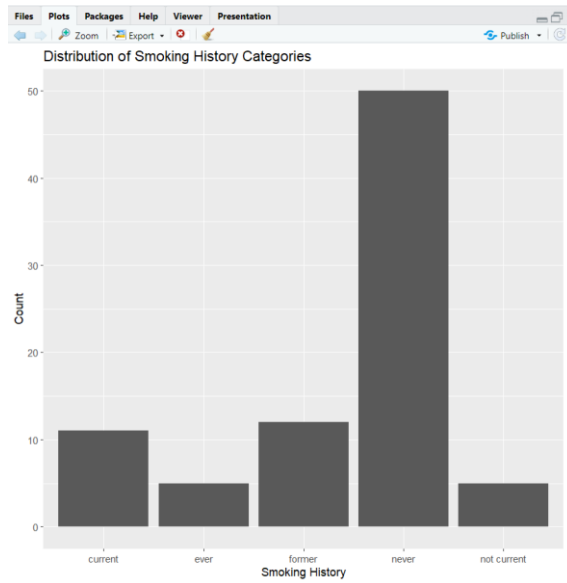
A visual representation of the counts of different heart disease categories.
The output:

```
ggplot(mydata_remove, aes(x = heart_disease)) +  
  geom_bar() +  
  labs(x = "Heart Disease", y = "Count", title = "Distribution of Heart Disease Categories")
```



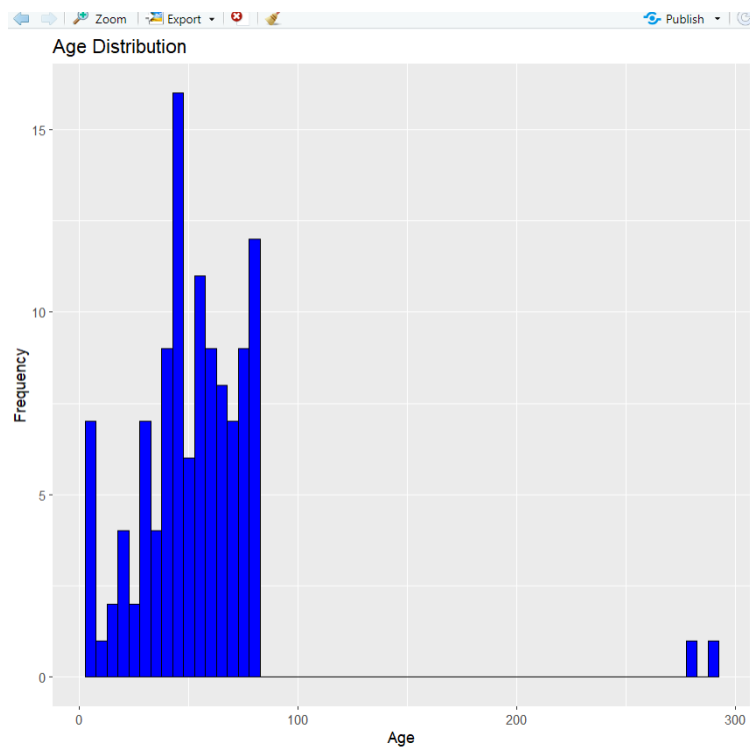
A visual representation of the counts of different smoking histories categories. The output:

```
> ggplot(mydata_remove, aes(x = smoking_history)) +  
+   geom_bar() +  
+   labs(x = "Smoking History", y = "Count", title = "Distribution of Smoking History Categories")
```



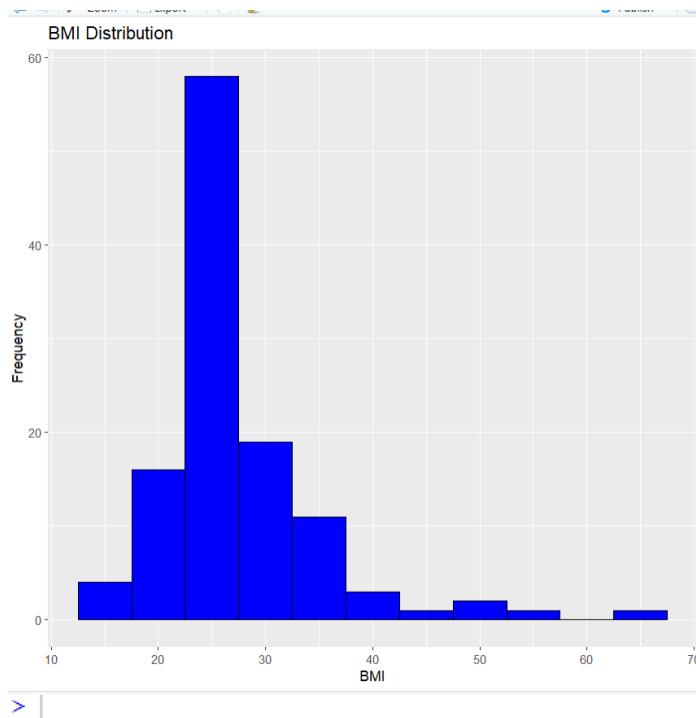
Ages are distributed across different ranges. The output:

```
> ggplot(mydata, aes(x = age)) +  
+   geom_histogram(binwidth = 5, fill = "blue", color = "black") +  
+   labs(x = "Age", y = "Frequency", title = "Age Distribution")  
>
```



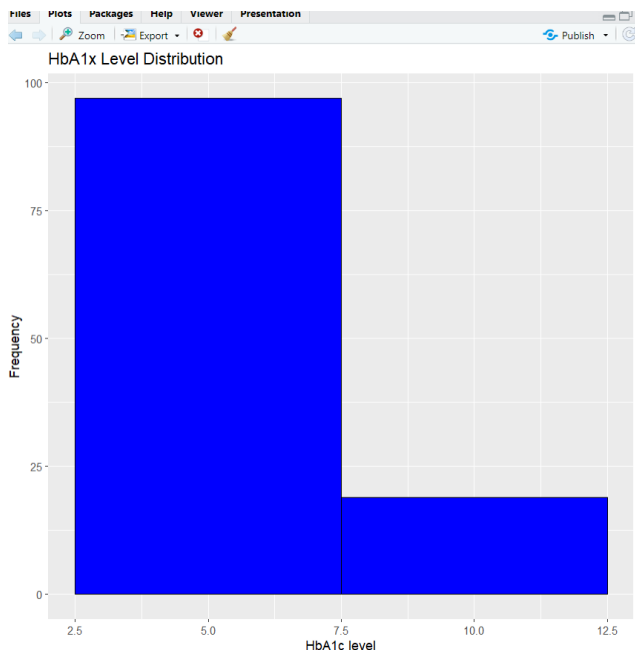
The BMIs are distributed across different ranges. The output:

```
ggplot(mydata, aes(x = bmi)) +
  geom_histogram(binwidth = 5, fill = "blue", color = "black") +
  labs(x = "BMI", y = "Frequency", title = "BMI Distribution")
```



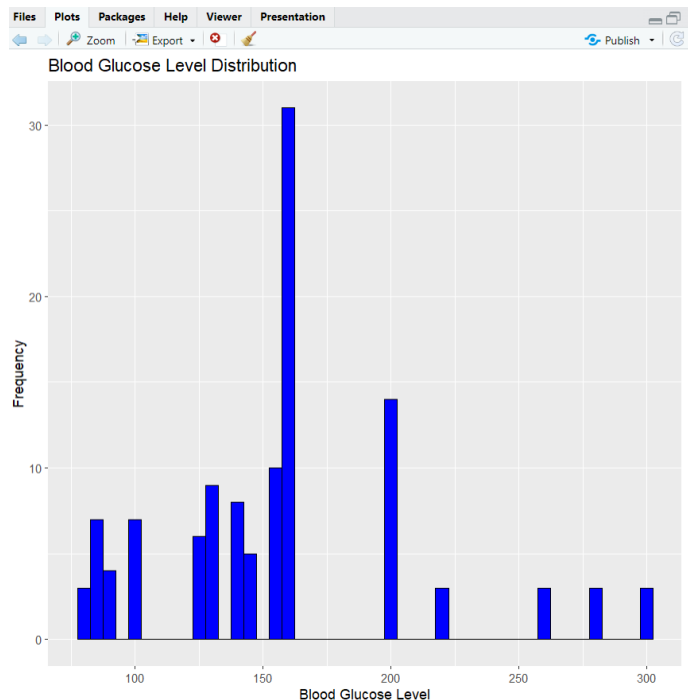
nges. The output:

```
lor = "black") +
  = "HbA1c Level Distribution")
```



The Blood glucose levels are distributed across different ranges. The output:

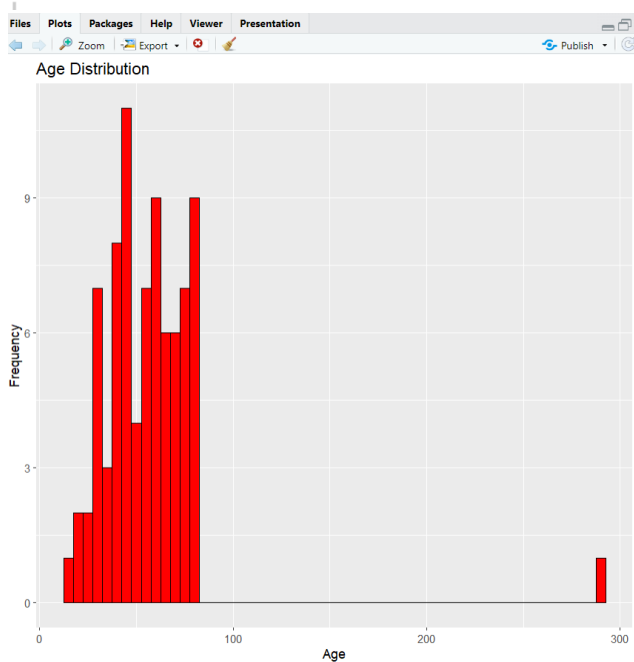
```
ggplot(mydata, aes(x = blood_glucose_level)) +
  geom_histogram(binwidth = 5, fill = "blue", color = "black") +
  labs(x = "Blood Glucose Level", y = "Frequency", title = "Blood Glucose Level Distribution")
```



After missing values handling:

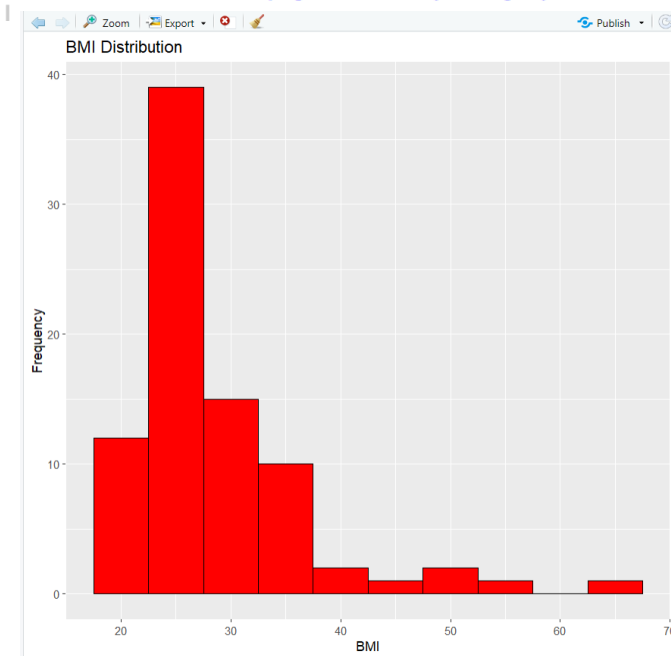
Numerical values of ages are distributed across different age ranges in 'mydata_remove' dataset. The output:

```
ggplot(mydata_remove, aes(x = age)) +
  geom_histogram(binwidth = 5, fill = "red", color = "black") +
  labs(x = "Age", y = "Frequency", title = "Age Distribution")
```



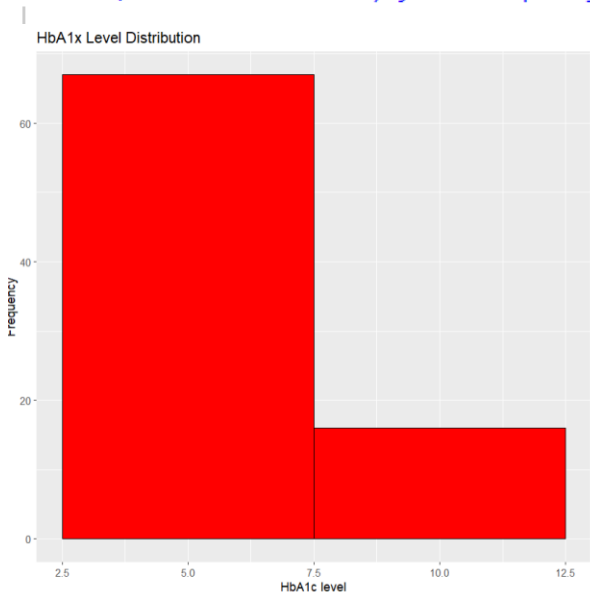
Numerical values of BMIs are distributed across different bmi ranges in my 'mydata_remove' dataset. The output:

```
ggplot(mydata_remove, aes(x = bmi)) +  
  geom_histogram(binwidth = 5, fill = "red", color = "black") +  
  labs(x = "BMI", y = "Frequency", title = "BMI Distribution")
```



How HbA1cs are distributed between two different ranges in my 'mydata_remove' dataset. The output:

```
ggplot(mydata_remove, aes(x = HbA1c_level)) +  
  geom_histogram(binwidth = 5, fill = "red", color = "black") +  
  labs(x = "HbA1c level", y = "Frequency", title = "HbA1x Level Distribution")
```



This is how an effective comparison can be identified by performing data exploration, outliers' detection, missing values detection and handling the missing values by implementing mean, mode, median and row wise removing techniques.