

# CORRELATION MATRIX

```
In [2]: import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

```
In [4]: # Load the dataset
df1=pd.read_csv(r"C:\Users\bhava\Downloads\cardio_train.csv")
df1
```

```
Out[4]:      id;age;gender;height;weight;ap_hi;ap_lo;cholesterol;gluc;smoke;alco;active;cardio
0      0;18393;2;168;62.0;110;80;1;1;0;0;1;0
1      1;20228;1;156;85.0;140;90;3;1;0;0;1;1
2      2;18857;1;165;64.0;130;70;3;1;0;0;0;1
3      3;17623;2;169;82.0;150;100;1;1;0;0;1;1
4      4;17474;1;156;56.0;100;60;1;1;0;0;0;0
...
69995  99993;19240;2;168;76.0;120;80;1;1;1;0;1;0
69996  99995;22601;1;158;126.0;140;90;2;2;0;0;1;1
69997  99996;19066;2;183;105.0;180;90;3;1;0;1;0;1
69998  99998;22431;1;163;72.0;135;80;1;2;0;0;0;1
69999  99999;20540;1;170;72.0;120;80;2;1;0;0;1;0
```

70000 rows × 1 columns

```
In [8]: df1=pd.read_csv(r"C:\Users\bhava\Downloads\cardio_train.csv",delimiter=';')
df1
```

```
Out[8]:      id  age  gender  height  weight  ap_hi  ap_lo  cholesterol  gluc  smoke  alco  active  cardio
0      0   0  18393      2    168    62.0   110    80           1     1     0     0     1     0
1      1   1  20228      1    156    85.0   140    90           3     1     0     0     1     1
2      2   2  18857      1    165    64.0   130    70           3     1     0     0     0     1
3      3   3  17623      2    169    82.0   150   100           1     1     0     0     1     1
4      4   4  17474      1    156    56.0   100    60           1     1     0     0     0     0
...
69995  99993  19240      2    168    76.0   120    80           1     1     1     0     1     0
69996  99995  22601      1    158   126.0   140    90           2     2     0     0     1     1
69997  99996  19066      2    183   105.0   180    90           3     1     0     1     0     1
69998  99998  22431      1    163    72.0   135    80           1     2     0     0     0     1
69999  99999  20540      1    170    72.0   120    80           2     1     0     0     1     0
```

70000 rows × 13 columns

```
In [16]: # Select relevant features for the correlation matrix
features = ['age', 'gender', 'height', 'weight', 'ap_hi', 'ap_lo', 'cholesterol', 'gluc', 'smoke', 'alco', 'active', 'cardio']
features
```

```
Out[16]: ['age',
'gender',
'height',
'weight',
'ap_hi',
'ap_lo',
'cholesterol',
'gluc',
'smoke',
'alco',
'active',
'cardio']
```

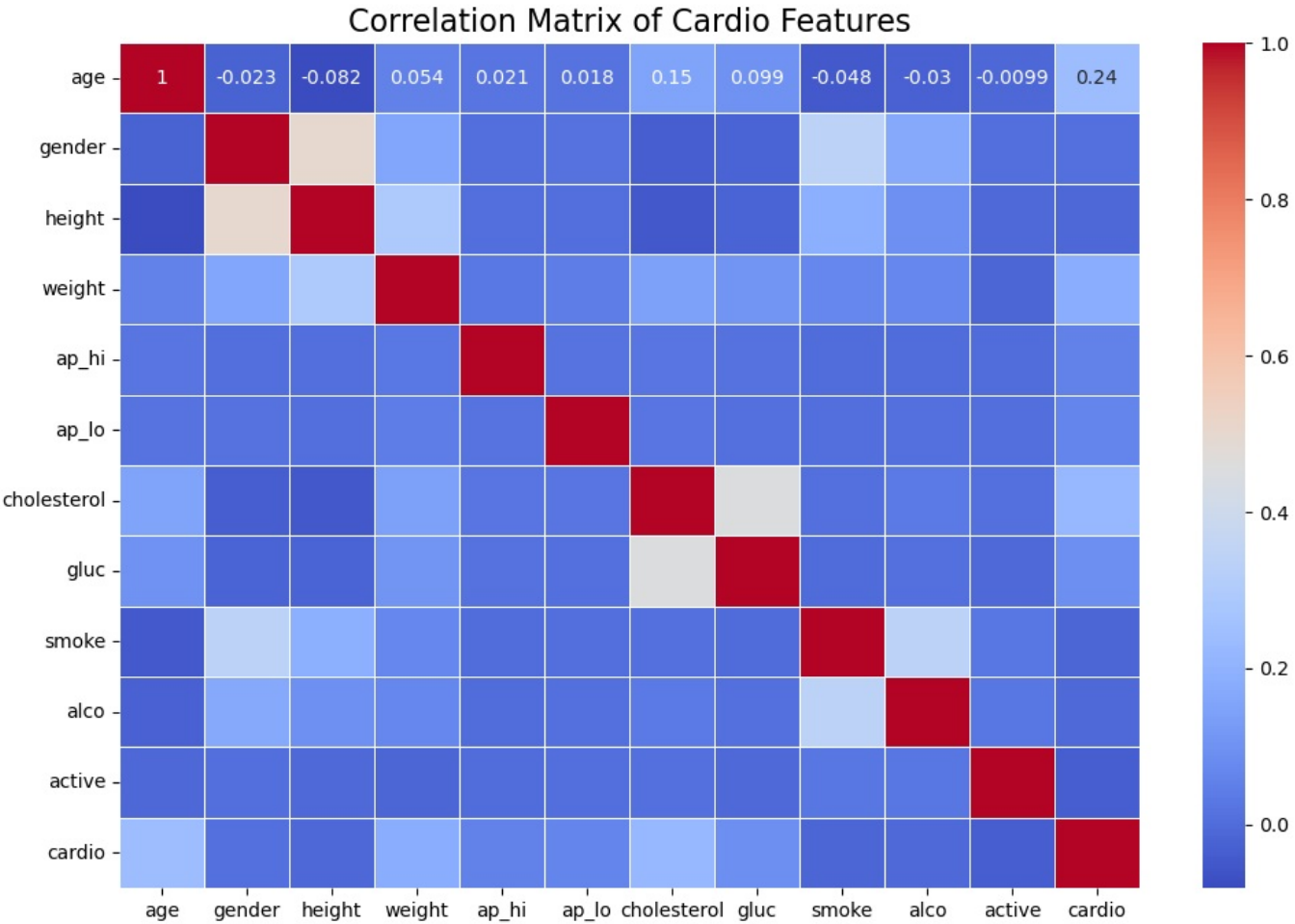
```
In [18]: # Calculate the correlation matrix
correlation_matrix = df1[features].corr()
correlation_matrix
```

Out[18]:

	age	gender	height	weight	ap_hi	ap_lo	cholesterol	gluc	smoke	alco	active
age	1.000000	-0.022811	-0.081515	0.053684	0.020764	0.017647	0.154424	0.098703	-0.047633	-0.029723	-0.009927
gender	-0.022811	1.000000	0.499033	0.155406	0.006005	0.015254	-0.035821	-0.020491	0.338135	0.170966	0.005866
height	-0.081515	0.499033	1.000000	0.290968	0.005488	0.006150	-0.050226	-0.018595	0.187989	0.094419	-0.006570
weight	0.053684	0.155406	0.290968	1.000000	0.030702	0.043710	0.141768	0.106857	0.067780	0.067113	-0.016867
ap_hi	0.020764	0.006005	0.005488	0.030702	1.000000	0.016086	0.023778	0.011841	-0.000922	0.001408	-0.000033
ap_lo	0.017647	0.015254	0.006150	0.043710	0.016086	1.000000	0.024019	0.010806	0.005186	0.010601	0.004780
cholesterol	0.154424	-0.035821	-0.050226	0.141768	0.023778	0.024019	1.000000	0.451578	0.010354	0.035760	0.009911
gluc	0.098703	-0.020491	-0.018595	0.106857	0.011841	0.010806	0.451578	1.000000	-0.004756	0.011246	-0.006770
smoke	-0.047633	0.338135	0.187989	0.067780	-0.000922	0.005186	0.010354	-0.004756	1.000000	0.340094	0.025858
alco	-0.029723	0.170966	0.094419	0.067113	0.001408	0.010601	0.035760	0.011246	0.340094	1.000000	0.025476
active	-0.009927	0.005866	-0.006570	-0.016867	-0.000033	0.004780	0.009911	-0.006770	0.025858	0.025476	1.000000
cardio	0.238159	0.008109	-0.010821	0.181660	0.054475	0.065719	0.221147	0.089307	-0.015486	-0.007330	-0.035653

In [20]:

```
# Plotting the correlation heatmap
plt.figure(figsize=(12, 8))
sns.heatmap(correlation_matrix, annot=True, cmap='coolwarm', linewidths=0.5)
plt.title('Correlation Matrix of Cardio Features', fontsize=16)
plt.show()
```



In [22]:

```
# Display the correlation matrix
print(correlation_matrix)
```

	age	gender	height	weight	ap_hi	ap_lo	\
age	1.000000	-0.022811	-0.081515	0.053684	0.020764	0.017647	
gender	-0.022811	1.000000	0.499033	0.155406	0.006005	0.015254	
height	-0.081515	0.499033	1.000000	0.290968	0.005488	0.006150	
weight	0.053684	0.155406	0.290968	1.000000	0.030702	0.043710	
ap_hi	0.020764	0.006005	0.005488	0.030702	1.000000	0.016086	
ap_lo	0.017647	0.015254	0.006150	0.043710	0.016086	1.000000	
cholesterol	0.154424	-0.035821	-0.050226	0.141768	0.023778	0.024019	
gluc	0.098703	-0.020491	-0.018595	0.106857	0.011841	0.010806	
smoke	-0.047633	0.338135	0.187989	0.067780	-0.000922	0.005186	
alco	-0.029723	0.170966	0.094419	0.067113	0.001408	0.010601	
active	-0.009927	0.005866	-0.006570	-0.016867	-0.000033	0.004780	
cardio	0.238159	0.008109	-0.010821	0.181660	0.054475	0.065719	

	cholesterol	gluc	smoke	alco	active	cardio
age	0.154424	0.098703	-0.047633	-0.029723	-0.009927	0.238159
gender	-0.035821	-0.020491	0.338135	0.170966	0.005866	0.008109
height	-0.050226	-0.018595	0.187989	0.094419	-0.006570	-0.010821
weight	0.141768	0.106857	0.067780	0.067113	-0.016867	0.181660
ap_hi	0.023778	0.011841	-0.000922	0.001408	-0.000033	0.054475
ap_lo	0.024019	0.010806	0.005186	0.010601	0.004780	0.065719
cholesterol	1.000000	0.451578	0.010354	0.035760	0.009911	0.221147
gluc	0.451578	1.000000	-0.004756	0.011246	-0.006770	0.089307
smoke	0.010354	-0.004756	1.000000	0.340094	0.025858	-0.015486
alco	0.035760	0.011246	0.340094	1.000000	0.025476	-0.007330
active	0.009911	-0.006770	0.025858	0.025476	1.000000	-0.035653
cardio	0.221147	0.089307	-0.015486	-0.007330	-0.035653	1.000000

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