

k93lu2nfv

January 30, 2023

1 EDA

1.1 Importing Libraries

```
[1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns

%matplotlib inline

import warnings
warnings.filterwarnings('ignore')

from sklearn.neighbors import KNeighborsClassifier
from sklearn.model_selection import train_test_split, GridSearchCV,
↳RandomizedSearchCV
from sklearn.svm import SVC
from sklearn.naive_bayes import GaussianNB
from sklearn.preprocessing import LabelEncoder
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import (RandomForestClassifier, AdaBoostClassifier,
↳ExtraTreesClassifier, GradientBoostingClassifier)

from sklearn.pipeline import Pipeline
from sklearn.preprocessing import StandardScaler
```

```
[3]: from google.colab import files
files.upload()
```

<IPython.core.display.HTML object>

Saving heart_disease_uci.csv to heart_disease_uci.csv

```
[3]: {'heart_disease_uci.csv': b'id,age,sex,dataset,cp,trestbps,chol,fbs,restecg,thal
ch,exang,oldpeak,slope,ca,thal,num\n1,63,Male,Cleveland,typical
angina,145,233,TRUE,lv hypertrophy,150,FALSE,2.3,downsloping,0,fixed
```

defect,0\n2,67, Male, Cleveland, asymptomatic, 160, 286, FALSE, lv hypertrophy, 108, TRUE, 1.5, flat, 3, normal, 2\n3, 67, Male, Cleveland, asymptomatic, 120, 229, FALSE, lv hypertrophy, 129, TRUE, 2.6, flat, 2, reversable defect, 1\n4, 37, Male, Cleveland, non-anginal, 130, 250, FALSE, normal, 187, FALSE, 3.5, downsloping, 0, normal, 0\n5, 41, Female, Cleveland, atypical angina, 130, 204, FALSE, lv hypertrophy, 172, FALSE, 1.4, upsloping, 0, normal, 0\n6, 56, Male, Cleveland, atypical angina, 120, 236, FALSE, normal, 178, FALSE, 0.8, upsloping, 0, normal, 0\n7, 62, Female, Cleveland, asymptomatic, 140, 268, FALSE, lv hypertrophy, 160, FALSE, 3.6, downsloping, 2, normal, 3\n8, 57, Female, Cleveland, asymptomatic, 120, 354, FALSE, normal, 163, TRUE, 0.6, upsloping, 0, normal, 0\n9, 63, Male, Cleveland, asymptomatic, 130, 254, FALSE, lv hypertrophy, 147, FALSE, 1.4, flat, 1, reversable defect, 2\n10, 53, Male, Cleveland, asymptomatic, 140, 203, TRUE, lv hypertrophy, 155, TRUE, 3.1, downsloping, 0, reversable defect, 1\n11, 57, Male, Cleveland, asymptomatic, 140, 192, FALSE, normal, 148, FALSE, 0.4, flat, 0, fixed defect, 0\n12, 56, Female, Cleveland, atypical angina, 140, 294, FALSE, lv hypertrophy, 153, FALSE, 1.3, flat, 0, normal, 0\n13, 56, Male, Cleveland, non-anginal, 130, 256, TRUE, lv hypertrophy, 142, TRUE, 0.6, flat, 1, fixed defect, 2\n14, 44, Male, Cleveland, atypical angina, 120, 263, FALSE, normal, 173, FALSE, 0, upsloping, 0, reversable defect, 0\n15, 52, Male, Cleveland, non-anginal, 172, 199, TRUE, normal, 162, FALSE, 0.5, upsloping, 0, reversable defect, 0\n16, 57, Male, Cleveland, non-anginal, 150, 168, FALSE, normal, 174, FALSE, 1.6, upsloping, 0, normal, 0\n17, 48, Male, Cleveland, atypical angina, 110, 229, FALSE, normal, 168, FALSE, 1, downsloping, 0, reversable defect, 1\n18, 54, Male, Cleveland, asymptomatic, 140, 239, FALSE, normal, 160, FALSE, 1.2, upsloping, 0, normal, 0\n19, 48, Female, Cleveland, non-anginal, 130, 275, FALSE, normal, 139, FALSE, 0.2, upsloping, 0, normal, 0\n20, 49, Male, Cleveland, atypical angina, 130, 266, FALSE, normal, 171, FALSE, 0.6, upsloping, 0, normal, 0\n21, 64, Male, Cleveland, typical angina, 110, 211, FALSE, lv hypertrophy, 144, TRUE, 1.8, flat, 0, normal, 0\n22, 58, Female, Cleveland, typical angina, 150, 283, TRUE, lv hypertrophy, 162, FALSE, 1, upsloping, 0, normal, 0\n23, 58, Male, Cleveland, atypical angina, 120, 284, FALSE, lv hypertrophy, 160, FALSE, 1.8, flat, 0, normal, 1\n24, 58, Male, Cleveland, non-anginal, 132, 224, FALSE, lv hypertrophy, 173, FALSE, 3.2, upsloping, 2, reversable defect, 3\n25, 60, Male, Cleveland, asymptomatic, 130, 206, FALSE, lv hypertrophy, 132, TRUE, 2.4, flat, 2, reversable defect, 4\n26, 50, Female, Cleveland, non-anginal, 120, 219, FALSE, normal, 158, FALSE, 1.6, flat, 0, normal, 0\n27, 58, Female, Cleveland, non-anginal, 120, 340, FALSE, normal, 172, FALSE, 0, upsloping, 0, normal, 0\n28, 66, Female, Cleveland, typical angina, 150, 226, FALSE, normal, 114, FALSE, 2.6, downsloping, 0, normal, 0\n29, 43, Male, Cleveland, asymptomatic, 150, 247, FALSE, normal, 171, FALSE, 1.5, upsloping, 0, normal, 0\n30, 40, Male, Cleveland, asymptomatic, 110, 167, FALSE, lv hypertrophy, 114, TRUE, 2, flat, 0, reversable defect, 3\n31, 69, Female, Cleveland, typical angina, 140, 239, FALSE, normal, 151, FALSE, 1.8, upsloping, 2, normal, 0\n32, 60, Male, Cleveland, asymptomatic, 117, 230, TRUE, normal, 160, TRUE, 1.4, upsloping, 2, reversable defect, 2\n33, 64, Male, Cleveland, non-anginal, 140, 335, FALSE, normal, 158, FALSE, 0, upsloping, 0, normal, 1\n34, 59, Male, Cleveland, asympt

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defect, 3\n38, 57, Male, Cleveland, asymptomatic, 150, 276, FALSE, lv hypertrophy, 112, TRUE, 0.6, flat, 1, fixed defect, 1\n39, 55, Male, Cleveland, asymptomatic, 132, 353, FALSE, normal, 132, TRUE, 1.2, flat, 1, reversable
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defect, 0\n43, 71, Female, Cleveland, atypical angina, 160, 302, FALSE, normal, 162, FALSE, 0.4, upsloping, 2, normal, 0\n44, 59, Male, Cleveland, non-anginal, 150, 212, TRUE, normal, 157, FALSE, 1.6, upsloping, 0, normal, 0\n45, 61, Female, Cleveland, asymptomatic, 130, 330, FALSE, lv hypertrophy, 169, FALSE, 0, upsloping, 0, normal, 1\n46, 58, Male, Cleveland, non-anginal, 112, 230, FALSE, lv hypertrophy, 165, FALSE, 2.5, flat, 1, reversable
defect, 4\n47, 51, Male, Cleveland, non-anginal, 110, 175, FALSE, normal, 123, FALSE, 0.6, upsloping, 0, normal, 0\n48, 50, Male, Cleveland, asymptomatic, 150, 243, FALSE, lv hypertrophy, 128, FALSE, 2.6, flat, 0, reversable
defect, 4\n49, 65, Female, Cleveland, non-anginal, 140, 417, TRUE, lv hypertrophy, 157, FALSE, 0.8, upsloping, 1, normal, 0\n50, 53, Male, Cleveland, non-anginal, 130, 197, TRUE, lv hypertrophy, 152, FALSE, 1.2, downsloping, 0, normal, 0\n51, 41, Female, Cleveland, atypical angina, 105, 198, FALSE, normal, 168, FALSE, 0, upsloping, 1, normal, 0\n52, 65, Male, Cleveland, asymptomatic, 120, 177, FALSE, normal, 140, FALSE, 0.4, upsloping, 0, reversable defect, 0\n53, 44, Male, Cleveland, asymptomatic, 112, 290, FALSE, lv hypertrophy, 153, FALSE, 0, upsloping, 1, normal, 2\n54, 44, Male, Cleveland, atypical angina, 130, 219, FALSE, lv hypertrophy, 188, FALSE, 0, upsloping, 0, normal, 0\n55, 60, Male, Cleveland, asymptomatic, 130, 253, FALSE, normal, 144, TRUE, 1.4, upsloping, 1, reversable
defect, 1\n56, 54, Male, Cleveland, asymptomatic, 124, 266, FALSE, lv hypertrophy, 109, TRUE, 2.2, flat, 1, reversable defect, 1\n57, 50, Male, Cleveland, non-anginal, 140, 233, FALSE, normal, 163, FALSE, 0.6, flat, 1, reversable
defect, 1\n58, 41, Male, Cleveland, asymptomatic, 110, 172, FALSE, lv hypertrophy, 158, FALSE, 0, upsloping, 0, reversable
defect, 1\n59, 54, Male, Cleveland, non-anginal, 125, 273, FALSE, lv hypertrophy, 152, FALSE, 0.5, downsloping, 1, normal, 0\n60, 51, Male, Cleveland, typical angina, 125, 213, FALSE, lv hypertrophy, 125, TRUE, 1.4, upsloping, 1, normal, 0\n61, 51, Female, Cleveland, asymptomatic, 130, 305, FALSE, normal, 142, TRUE, 1.2, flat, 0, reversable
defect, 2\n62, 46, Female, Cleveland, non-anginal, 142, 177, FALSE, lv hypertrophy, 160, TRUE, 1.4, downsloping, 0, normal, 0\n63, 58, Male, Cleveland, asymptomatic, 128, 216, FALSE, lv hypertrophy, 131, TRUE, 2.2, flat, 3, reversable
defect, 1\n64, 54, Female, Cleveland, non-anginal, 135, 304, TRUE, normal, 170, FALSE, 0, upsloping, 0, normal, 0\n65, 54, Male, Cleveland, asymptomatic, 120, 188, FALSE, normal, 113, FALSE, 1.4, flat, 1, reversable
defect, 2\n66, 60, Male, Cleveland, asymptomatic, 145, 282, FALSE, lv hypertrophy, 142, TRUE, 2.8, flat, 2, reversable defect, 2\n67, 60, Male, Cleveland, non-anginal, 140, 185, FALSE, lv

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 hypertrophy, 140, TRUE, 3.4, downsloping, 0, reversable
 defect, 2\n70, 46, Male, Cleveland, non-anginal, 150, 231, FALSE, normal, 147, FALSE, 3.6, fl
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 , 163, FALSE, 0.2, flat, 2, reversable defect, 3\n73, 62, Male, Cleveland, asymptomatic, 120
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 defect, 1\n74, 65, Male, Cleveland, asymptomatic, 110, 248, FALSE, lv
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 defect, 1\n75, 44, Male, Cleveland, asymptomatic, 110, 197, FALSE, lv
 hypertrophy, 177, FALSE, 0, upsloping, 1, normal, 1\n76, 65, Female, Cleveland, non-
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 hypertrophy, 141, TRUE, 2.8, flat, 1, reversable defect, 1\n78, 51, Female, Cleveland, non-
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 hypertrophy, 142, FALSE, 1.5, upsloping, 1, normal, 0\n79, 48, Male, Cleveland, atypical
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 defect, 3\n81, 45, Male, Cleveland, asymptomatic, 104, 208, FALSE, lv hypertrophy, 148, TRU
 E, 3, flat, 0, normal, 0\n82, 53, Female, Cleveland, asymptomatic, 130, 264, FALSE, lv
 hypertrophy, 143, FALSE, 0.4, flat, 0, normal, 0\n83, 39, Male, Cleveland, non-
 anginal, 140, 321, FALSE, lv
 hypertrophy, 182, FALSE, 0, upsloping, 0, normal, 0\n84, 68, Male, Cleveland, non-
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 defect, 3\n85, 52, Male, Cleveland, atypical angina, 120, 325, FALSE, normal, 172, FALSE, 0.
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 hypertrophy, 156, FALSE, 0, upsloping, 0, normal, 0\n88, 53, Female, Cleveland, non-
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 defect, 1\n97, 59, Male, Cleveland, asymptomatic, 110, 239, FALSE, lv
 hypertrophy, 142, TRUE, 1.2, flat, 1, reversable

defect,2\n98,60,Female,Cleveland,asymptomatic,150,258,FALSE,lv
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hypertrophy,186,FALSE,0,upsloping,0,normal,0\n101,45,Male,Cleveland,asymptomatic,115,260,FALSE,lv
hypertrophy,185,FALSE,0,upsloping,0,normal,0\n102,34,Male,Cleveland,typical angina,118,182,FALSE,lv
hypertrophy,174,FALSE,0,upsloping,0,normal,0\n103,57,Female,Cleveland,asymptomatic,128,303,FALSE,lv
hypertrophy,159,FALSE,0,upsloping,1,normal,0\n104,71,Female,Cleveland,non-anginal,110,265,TRUE,lv
hypertrophy,130,FALSE,0,upsloping,1,normal,0\n105,49,Male,Cleveland,non-anginal,120,188,FALSE,normal,139,FALSE,2,flat,3,reversible
defect,3\n106,54,Male,Cleveland,atypical angina,108,309,FALSE,normal,156,FALSE,0,upsloping,0,reversible defect,0\n107,59, Male,Cleveland,asymptomatic,140,177,FALSE,normal,162,TRUE,0,upsloping,1,reversible defect,2\n108,57, Male,Cleveland,non-anginal,128,229,FALSE,lv
hypertrophy,150,FALSE,0.4,flat,1,reversible defect,1\n109,61, Male,Cleveland,asymptomatic,120,260,FALSE,normal,140,TRUE,3.6,flat,1,reversible defect,2\n110,39, Male,Cleveland,asymptomatic,118,219,FALSE,normal,140,FALSE,1.2,flat,0,reversible defect,3\n111,61, Female,Cleveland,asymptomatic,145,307,FALSE,lv
hypertrophy,146,TRUE,1,flat,0,reversible
defect,1\n112,56, Male,Cleveland,asymptomatic,125,249,TRUE,lv
hypertrophy,144,TRUE,1.2,flat,1,normal,1\n113,52, Male,Cleveland,typical angina,118,186,FALSE,lv
hypertrophy,190,FALSE,0,flat,0,fixed defect,0\n114,43, Female,Cleveland,asymptomatic,132,341,TRUE,lv
hypertrophy,136,TRUE,3,flat,0,reversible defect,2\n115,62, Female,Cleveland,non-anginal,130,263,FALSE,normal,97,FALSE,1.2,flat,1,reversible
defect,2\n116,41, Male,Cleveland,atypical angina,135,203,FALSE,normal,132,FALSE,0,flat,0,fixed defect,0\n117,58, Male,Cleveland,non-anginal,140,211,TRUE,lv
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hypertrophy,132,TRUE,1.8,upsloping,3,reversible defect,3\n120,65, Male,Cleveland,asymptomatic,135,254,FALSE,lv
hypertrophy,127,FALSE,2.8,flat,1,reversible
defect,2\n121,48, Male,Cleveland,asymptomatic,130,256,TRUE,lv
hypertrophy,150,TRUE,0,upsloping,2,reversible
defect,3\n122,63, Female,Cleveland,asymptomatic,150,407,FALSE,lv
hypertrophy,154,FALSE,4,flat,3,reversible defect,4\n123,51, Male,Cleveland,non-anginal,100,222,FALSE,normal,143,TRUE,1.2,flat,0,normal,0\n124,55, Male,Cleveland,asymptomatic,140,217,FALSE,normal,111,TRUE,5.6,downsloping,0,reversible
defect,3\n125,65, Male,Cleveland,typical angina,138,282,TRUE,lv
hypertrophy,174,FALSE,1.4,flat,1,normal,1\n126,45, Female,Cleveland,atypical angina,130,234,FALSE,lv
hypertrophy,175,FALSE,0.6,flat,0,normal,0\n127,56, Female,Cleveland,asymptomatic,200,288,TRUE,lv
hypertrophy,133,TRUE,4,downsloping,2,reversible defect,3\n128,54, Male,Cleveland,

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 defect,3\n129,44,Male,Cleveland,atypical angina,120,220,FALSE,normal,170,FALSE,0
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 anginal,120,258,FALSE,lv hypertrophy,147,FALSE,0.4,flat,0,reversible
 defect,0\n132,51,Male,Cleveland,non-
 anginal,94,227,FALSE,normal,154,TRUE,0,upsloping,1,reversible
 defect,0\n133,29,Male,Cleveland,atypical angina,130,204,FALSE,lv hypertrophy,202
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 lv hypertrophy,186,TRUE,0,upsloping,0,normal,0\n135,43,Female,Cleveland,non-angi
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 atypical angina,135,250,FALSE,lv hypertrophy,161,FALSE,1.4,flat,0,normal,0\n137,
 70,Male,Cleveland,asymptomatic,145,174,FALSE,normal,125,TRUE,2.6,downsloping,0,r
 eversible defect,4\n138,62,Male,Cleveland,atypical angina,120,281,FALSE,lv
 hypertrophy,103,FALSE,1.4,flat,1,reversible defect,3\n139,35,Male,Cleveland,asym
 ptomatic,120,198,FALSE,normal,130,TRUE,1.6,flat,0,reversible
 defect,1\n140,51,Male,Cleveland,non-anginal,125,245,TRUE,lv
 hypertrophy,166,FALSE,2.4,flat,0,normal,0\n141,59,Male,Cleveland,atypical angina
 ,140,221,FALSE,normal,164,TRUE,0,upsloping,0,normal,0\n142,59,Male,Cleveland,typ
 ical angina,170,288,FALSE,lv hypertrophy,159,FALSE,0.2,flat,0,reversible
 defect,1\n143,52,Male,Cleveland,atypical angina,128,205,TRUE,normal,184,FALSE,0,
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 anginal,125,309,FALSE,normal,131,TRUE,1.8,flat,0,reversible
 defect,1\n145,58,Male,Cleveland,non-anginal,105,240,FALSE,lv
 hypertrophy,154,TRUE,0.6,flat,0,reversible defect,0\n146,47,Male,Cleveland,non-a
 nginal,108,243,FALSE,normal,152,FALSE,0,upsloping,0,normal,1\n147,57,Male,Clevel
 and,asymptomatic,165,289,TRUE,lv hypertrophy,124,FALSE,1,flat,3,reversible
 defect,4\n148,41,Male,Cleveland,non-anginal,112,250,FALSE,normal,179,FALSE,0,ups
 loping,0,normal,0\n149,45,Male,Cleveland,atypical angina,128,308,FALSE,lv
 hypertrophy,170,FALSE,0,upsloping,0,normal,0\n150,60,Female,Cleveland,non-angina
 l,102,318,FALSE,normal,160,FALSE,0,upsloping,1,normal,0\n151,52,Male,Cleveland,t
 ypic al angina,152,298,TRUE,normal,178,FALSE,1.2,flat,0,reversible
 defect,0\n152,42,Female,Cleveland,asymptomatic,102,265,FALSE,lv
 hypertrophy,122,FALSE,0.6,flat,0,normal,0\n153,67,Female,Cleveland,non-
 anginal,115,564,FALSE,lv hypertrophy,160,FALSE,1.6,flat,0,reversible
 defect,0\n154,55,Male,Cleveland,asymptomatic,160,289,FALSE,lv
 hypertrophy,145,TRUE,0.8,flat,1,reversible
 defect,4\n155,64,Male,Cleveland,asymptomatic,120,246,FALSE,lv hypertrophy,96,TRU
 E,2.2,downsloping,1,normal,3\n156,70,Male,Cleveland,asymptomatic,130,322,FALSE,l
 v hypertrophy,109,FALSE,2.4,flat,3,normal,1\n157,51,Male,Cleveland,asymptomatic,
 140,299,FALSE,normal,173,TRUE,1.6,upsloping,0,reversible
 defect,1\n158,58,Male,Cleveland,asymptomatic,125,300,FALSE,lv
 hypertrophy,171,FALSE,0,upsloping,2,reversible
 defect,1\n159,60,Male,Cleveland,asymptomatic,140,293,FALSE,lv
 hypertrophy,170,FALSE,1.2,flat,2,reversible defect,2\n160,68,Male,Cleveland,non-
 anginal,118,277,FALSE,normal,151,FALSE,1,upsloping,1,reversible
 defect,0\n161,46,Male,Cleveland,atypical

angina,101,197,TRUE,normal,156,FALSE,0,upsloping,0,reversable
 defect,0\n162,77,Male,Cleveland,asymptomatic,125,304,FALSE,lv
 hypertrophy,162,TRUE,0,upsloping,3,normal,4\n163,54,Female,Cleveland,non-anginal
 ,110,214,FALSE,normal,158,FALSE,1.6,flat,0,normal,0\n164,58,Female,Cleveland,asy
 mptomatic,100,248,FALSE,lv
 hypertrophy,122,FALSE,1,flat,0,normal,0\n165,48,Male,Cleveland,non-anginal,124,2
 55,TRUE,normal,175,FALSE,0,upsloping,2,normal,0\n166,57,Male,Cleveland,asymptoma
 tic,132,207,FALSE,normal,168,TRUE,0,upsloping,0,reversable
 defect,0\n167,52,Male,Cleveland,non-anginal,138,223,FALSE,normal,169,FALSE,0,ups
 loping,,normal,0\n168,54,Female,Cleveland,atypical angina,132,288,TRUE,lv hypert
 rophy,159,TRUE,0,upsloping,1,normal,0\n169,35,Male,Cleveland,asymptomatic,126,28
 2,FALSE,lv hypertrophy,156,TRUE,0,upsloping,0,reversable
 defect,1\n170,45,Female,Cleveland,atypical angina,112,160,FALSE,normal,138,FALSE
 ,0,flat,0,normal,0\n171,70,Male,Cleveland,non-
 anginal,160,269,FALSE,normal,112,TRUE,2.9,flat,1,reversable
 defect,3\n172,53,Male,Cleveland,asymptomatic,142,226,FALSE,lv
 hypertrophy,111,TRUE,0,upsloping,0,reversable defect,0\n173,59,Female,Cleveland,
 asymptomatic,174,249,FALSE,normal,143,TRUE,0,flat,0,normal,1\n174,62,Female,Clev
 eland,asymptomatic,140,394,FALSE,lv hypertrophy,157,FALSE,1.2,flat,0,normal,0\n1
 75,64,Male,Cleveland,asymptomatic,145,212,FALSE,lv
 hypertrophy,132,FALSE,2,flat,2,fixed defect,4\n176,57,Male,Cleveland,asymptomati
 c,152,274,FALSE,normal,88,TRUE,1.2,flat,1,reversable defect,1\n177,52,Male,Cleve
 land,asymptomatic,108,233,TRUE,normal,147,FALSE,0.1,upsloping,3,reversable
 defect,0\n178,56,Male,Cleveland,asymptomatic,132,184,FALSE,lv
 hypertrophy,105,TRUE,2.1,flat,1,fixed defect,1\n179,43,Male,Cleveland,non-angina
 l,130,315,FALSE,normal,162,FALSE,1.9,upsloping,1,normal,0\n180,53,Male,Cleveland
 ,non-anginal,130,246,TRUE,lv hypertrophy,173,FALSE,0,upsloping,3,normal,0\n181,4
 8,Male,Cleveland,asymptomatic,124,274,FALSE,lv
 hypertrophy,166,FALSE,0.5,flat,0,reversable
 defect,3\n182,56,Female,Cleveland,asymptomatic,134,409,FALSE,lv
 hypertrophy,150,TRUE,1.9,flat,2,reversable
 defect,2\n183,42,Male,Cleveland,typical angina,148,244,FALSE,lv
 hypertrophy,178,FALSE,0.8,upsloping,2,normal,0\n184,59,Male,Cleveland,typical
 angina,178,270,FALSE,lv hypertrophy,145,FALSE,4.2,downsloping,0,reversable
 defect,0\n185,60,Female,Cleveland,asymptomatic,158,305,FALSE,lv
 hypertrophy,161,FALSE,0,upsloping,0,normal,1\n186,63,Female,Cleveland,atypical a
 ngina,140,195,FALSE,normal,179,FALSE,0,upsloping,2,normal,0\n187,42,Male,Clevela
 nd,non-anginal,120,240,TRUE,normal,194,FALSE,0.8,downsloping,0,reversable
 defect,0\n188,66,Male,Cleveland,atypical
 angina,160,246,FALSE,normal,120,TRUE,0,flat,3,fixed
 defect,2\n189,54,Male,Cleveland,atypical angina,192,283,FALSE,lv
 hypertrophy,195,FALSE,0,upsloping,1,reversable
 defect,1\n190,69,Male,Cleveland,non-anginal,140,254,FALSE,lv
 hypertrophy,146,FALSE,2,flat,3,reversable defect,2\n191,50,Male,Cleveland,non-an
 ginal,129,196,FALSE,normal,163,FALSE,0,upsloping,0,normal,0\n192,51,Male,Clevela
 nd,asymptomatic,140,298,FALSE,normal,122,TRUE,4.2,flat,3,reversable
 defect,3\n193,43,Male,Cleveland,asymptomatic,132,247,TRUE,lv

hypertrophy, 143, TRUE, 0.1, flat, , reversable defect, 1\n194, 62, Female, Cleveland, asymptomatic, 138, 294, TRUE, normal, 106, FALSE, 1.9, flat, 3, normal, 2\n195, 68, Female, Cleveland, non-anginal, 120, 211, FALSE, lv hypertrophy, 115, FALSE, 1.5, flat, 0, normal, 0\n196, 67, Male, Cleveland, asymptomatic, 100, 299, FALSE, lv hypertrophy, 125, TRUE, 0.9, flat, 2, normal, 3\n197, 69, Male, Cleveland, typical angina, 160, 234, TRUE, lv hypertrophy, 131, FALSE, 0.1, flat, 1, normal, 0\n198, 45, Female, Cleveland, asymptomatic, 138, 236, FALSE, lv hypertrophy, 152, TRUE, 0.2, flat, 0, normal, 0\n199, 50, Female, Cleveland, atypical angina, 120, 244, FALSE, normal, 162, FALSE, 1.1, upsloping, 0, normal, 0\n200, 59, Male, Cleveland, typical angina, 160, 273, FALSE, lv hypertrophy, 125, FALSE, 0, upsloping, 0, normal, 1\n201, 50, Female, Cleveland, asymptomatic, 110, 254, FALSE, lv hypertrophy, 159, FALSE, 0, upsloping, 0, normal, 0\n202, 64, Female, Cleveland, asymptomatic, 180, 325, FALSE, normal, 154, TRUE, 0, upsloping, 0, normal, 0\n203, 57, Male, Cleveland, non-anginal, 150, 126, TRUE, normal, 173, FALSE, 0.2, upsloping, 1, reversable defect, 0\n204, 64, Female, Cleveland, non-anginal, 140, 313, FALSE, normal, 133, FALSE, 0.2, upsloping, 0, reversable defect, 0\n205, 43, Male, Cleveland, asymptomatic, 110, 211, FALSE, normal, 161, FALSE, 0, upsloping, 0, reversable defect, 0\n206, 45, Male, Cleveland, asymptomatic, 142, 309, FALSE, lv hypertrophy, 147, TRUE, 0, flat, 3, reversable defect, 3\n207, 58, Male, Cleveland, asymptomatic, 128, 259, FALSE, lv hypertrophy, 130, TRUE, 3, flat, 2, reversable defect, 3\n208, 50, Male, Cleveland, asymptomatic, 144, 200, FALSE, lv hypertrophy, 126, TRUE, 0.9, flat, 0, reversable defect, 3\n209, 55, Male, Cleveland, atypical angina, 130, 262, FALSE, normal, 155, FALSE, 0, upsloping, 0, normal, 0\n210, 62, Female, Cleveland, asymptomatic, 150, 244, FALSE, normal, 154, TRUE, 1.4, flat, 0, normal, 1\n211, 37, Female, Cleveland, non-anginal, 120, 215, FALSE, normal, 170, FALSE, 0, upsloping, 0, normal, 0\n212, 38, Male, Cleveland, typical angina, 120, 231, FALSE, normal, 182, TRUE, 3.8, flat, 0, reversable defect, 4\n213, 41, Male, Cleveland, non-anginal, 130, 214, FALSE, lv hypertrophy, 168, FALSE, 2, flat, 0, normal, 0\n214, 66, Female, Cleveland, asymptomatic, 178, 228, TRUE, normal, 165, TRUE, 1, flat, 2, reversable defect, 3\n215, 52, Male, Cleveland, asymptomatic, 112, 230, FALSE, normal, 160, FALSE, 0, upsloping, 1, normal, 1\n216, 56, Male, Cleveland, typical angina, 120, 193, FALSE, lv hypertrophy, 162, FALSE, 1.9, flat, 0, reversable defect, 0\n217, 46, Female, Cleveland, atypical angina, 105, 204, FALSE, normal, 172, FALSE, 0, upsloping, 0, normal, 0\n218, 46, Female, Cleveland, asymptomatic, 138, 243, FALSE, lv hypertrophy, 152, TRUE, 0, flat, 0, normal, 0\n219, 64, Female, Cleveland, asymptomatic, 130, 303, FALSE, normal, 122, FALSE, 2, flat, 2, normal, 0\n220, 59, Male, Cleveland, asymptomatic, 138, 271, FALSE, lv hypertrophy, 182, FALSE, 0, upsloping, 0, normal, 0\n221, 41, Female, Cleveland, non-anginal, 112, 268, FALSE, lv hypertrophy, 172, TRUE, 0, upsloping, 0, normal, 0\n222, 54, Female, Cleveland, non-anginal, 108, 267, FALSE, lv hypertrophy, 167, FALSE, 0, upsloping, 0, normal, 0\n223, 39, Female, Cleveland, non-anginal, 94, 199, FALSE, normal, 179, FALSE, 0, upsloping, 0, normal, 0\n224, 53, Male, Cleveland, asymptomatic, 123, 282, FALSE, normal, 95, TRUE, 2, flat, 2, reversable defect, 3\n225, 63, Female, Cleveland, asymptomatic, 108, 269, FALSE, normal, 169, TRUE, 1.8, flat, 2, normal, 1\n226, 34, Female, Cleveland, atypical angina, 118, 210, FALSE, normal, 192, FALSE, 0.7, upsloping,

ng,0,normal,0\n227,47, Male, Cleveland, asymptomatic, 112, 204, FALSE, normal, 143, FALSE, 0.1, upsloping, 0, normal, 0\n228, 67, Female, Cleveland, non-anginal, 152, 277, FALSE, normal, 172, FALSE, 0, upsloping, 1, normal, 0\n229, 54, Male, Cleveland, asymptomatic, 110, 206, FALSE, lv hypertrophy, 108, TRUE, 0, flat, 1, normal, 3\n230, 66, Male, Cleveland, asymptomatic, 112, 212, FALSE, lv hypertrophy, 132, TRUE, 0.1, upsloping, 1, normal, 2\n231, 52, Female, Cleveland, non-anginal, 136, 196, FALSE, lv hypertrophy, 169, FALSE, 0.1, flat, 0, normal, 0\n232, 55, Female, Cleveland, asymptomatic, 180, 327, FALSE, st-t abnormality, 117, TRUE, 3.4, flat, 0, normal, 2\n233, 49, Male, Cleveland, non-anginal, 118, 149, FALSE, lv hypertrophy, 126, FALSE, 0.8, upsloping, 3, normal, 1\n234, 74, Female, Cleveland, atypical angina, 120, 269, FALSE, lv hypertrophy, 121, TRUE, 0.2, upsloping, 1, normal, 0\n235, 54, Female, Cleveland, non-anginal, 160, 201, FALSE, normal, 163, FALSE, 0, upsloping, 1, normal, 0\n236, 54, Male, Cleveland, asymptomatic, 122, 286, FALSE, lv hypertrophy, 116, TRUE, 3.2, flat, 2, normal, 3\n237, 56, Male, Cleveland, asymptomatic, 130, 283, TRUE, lv hypertrophy, 103, TRUE, 1.6, downsloping, 0, reversable defect, 2\n238, 46, Male, Cleveland, asymptomatic, 120, 249, FALSE, lv hypertrophy, 144, FALSE, 0.8, upsloping, 0, reversable defect, 1\n239, 49, Female, Cleveland, atypical angina, 134, 271, FALSE, normal, 162, FALSE, 0, flat, 0, normal, 0\n240, 42, Male, Cleveland, atypical angina, 120, 295, FALSE, normal, 162, FALSE, 0, upsloping, 0, normal, 0\n241, 41, Male, Cleveland, atypical angina, 110, 235, FALSE, normal, 153, FALSE, 0, upsloping, 0, normal, 0\n242, 41, Female, Cleveland, atypical angina, 126, 306, FALSE, normal, 163, FALSE, 0, upsloping, 0, normal, 0\n243, 49, Female, Cleveland, asymptomatic, 130, 269, FALSE, normal, 163, FALSE, 0, upsloping, 0, normal, 0\n244, 61, Male, Cleveland, typical angina, 134, 234, FALSE, normal, 145, FALSE, 2.6, flat, 2, normal, 2\n245, 60, Female, Cleveland, non-anginal, 120, 178, TRUE, normal, 96, FALSE, 0, upsloping, 0, normal, 0\n246, 67, Male, Cleveland, asymptomatic, 120, 237, FALSE, normal, 71, FALSE, 1, flat, 0, normal, 2\n247, 58, Male, Cleveland, asymptomatic, 100, 234, FALSE, normal, 156, FALSE, 0.1, upsloping, 1, reversable defect, 2\n248, 47, Male, Cleveland, asymptomatic, 110, 275, FALSE, lv hypertrophy, 118, TRUE, 1, flat, 1, normal, 1\n249, 52, Male, Cleveland, asymptomatic, 125, 212, FALSE, normal, 168, FALSE, 1, upsloping, 2, reversable defect, 3\n250, 62, Male, Cleveland, atypical angina, 128, 208, TRUE, lv hypertrophy, 140, FALSE, 0, upsloping, 0, normal, 0\n251, 57, Male, Cleveland, asymptomatic, 110, 201, FALSE, normal, 126, TRUE, 1.5, flat, 0, fixed defect, 0\n252, 58, Male, Cleveland, asymptomatic, 146, 218, FALSE, normal, 105, FALSE, 2, flat, 1, reversable defect, 1\n253, 64, Male, Cleveland, asymptomatic, 128, 263, FALSE, normal, 105, TRUE, 0.2, flat, 1, reversable defect, 0\n254, 51, Female, Cleveland, non-anginal, 120, 295, FALSE, lv hypertrophy, 157, FALSE, 0.6, upsloping, 0, normal, 0\n255, 43, Male, Cleveland, asymptomatic, 115, 303, FALSE, normal, 181, FALSE, 1.2, flat, 0, normal, 0\n256, 42, Female, Cleveland, non-anginal, 120, 209, FALSE, normal, 173, FALSE, 0, flat, 0, normal, 0\n257, 67, Female, Cleveland, asymptomatic, 106, 223, FALSE, normal, 142, FALSE, 0.3, upsloping, 2, normal, 0\n258, 76, Female, Cleveland, non-anginal, 140, 197, FALSE, st-t abnormality, 116, FALSE, 1.1, flat, 0, normal, 0\n259, 70, Male, Cleveland, atypical angina, 156, 245, FALSE, lv hypertrophy, 143, FALSE, 0, upsloping, 0, normal, 0\n260, 57, Male, Cleveland, atypical angina, 124, 261, FALSE, normal, 141, FALSE, 0.3, upsloping, 0, reversable

defect, 1\n261, 44, Female, Cleveland, non-anginal, 118, 242, FALSE, normal, 149, FALSE, 0.3
, flat, 1, normal, 0\n262, 58, Female, Cleveland, atypical angina, 136, 319, TRUE, lv
hypertrophy, 152, FALSE, 0, upsloping, 2, normal, 3\n263, 60, Female, Cleveland, typical an
gina, 150, 240, FALSE, normal, 171, FALSE, 0.9, upsloping, 0, normal, 0\n264, 44, Male, Clevel
and, non-anginal, 120, 226, FALSE, normal, 169, FALSE, 0, upsloping, 0, normal, 0\n265, 61, Ma
le, Cleveland, asymptomatic, 138, 166, FALSE, lv hypertrophy, 125, TRUE, 3.6, flat, 1, norma
l, 4\n266, 42, Male, Cleveland, asymptomatic, 136, 315, FALSE, normal, 125, TRUE, 1.8, flat, 0
, fixed defect, 2\n267, 52, Male, Cleveland, asymptomatic, 128, 204, TRUE, normal, 156, TRUE
, 1, flat, 0, , 2\n268, 59, Male, Cleveland, non-
anginal, 126, 218, TRUE, normal, 134, FALSE, 2.2, flat, 1, fixed defect, 2\n269, 40, Male, Cle
veland, asymptomatic, 152, 223, FALSE, normal, 181, FALSE, 0, upsloping, 0, reversable
defect, 1\n270, 42, Male, Cleveland, non-anginal, 130, 180, FALSE, normal, 150, FALSE, 0, ups
loping, 0, normal, 0\n271, 61, Male, Cleveland, asymptomatic, 140, 207, FALSE, lv
hypertrophy, 138, TRUE, 1.9, upsloping, 1, reversable
defect, 1\n272, 66, Male, Cleveland, asymptomatic, 160, 228, FALSE, lv
hypertrophy, 138, FALSE, 2.3, upsloping, 0, fixed defect, 0\n273, 46, Male, Cleveland, asym
ptomatic, 140, 311, FALSE, normal, 120, TRUE, 1.8, flat, 2, reversable defect, 2\n274, 71, Fe
male, Cleveland, asymptomatic, 112, 149, FALSE, normal, 125, FALSE, 1.6, flat, 0, normal, 0\n
275, 59, Male, Cleveland, typical angina, 134, 204, FALSE, normal, 162, FALSE, 0.8, upslopin
g, 2, normal, 1\n276, 64, Male, Cleveland, typical angina, 170, 227, FALSE, lv
hypertrophy, 155, FALSE, 0.6, flat, 0, reversable
defect, 0\n277, 66, Female, Cleveland, non-anginal, 146, 278, FALSE, lv
hypertrophy, 152, FALSE, 0, flat, 1, normal, 0\n278, 39, Female, Cleveland, non-anginal, 138
, 220, FALSE, normal, 152, FALSE, 0, flat, 0, normal, 0\n279, 57, Male, Cleveland, atypical
angina, 154, 232, FALSE, lv hypertrophy, 164, FALSE, 0, upsloping, 1, normal, 1\n280, 58, Fem
ale, Cleveland, asymptomatic, 130, 197, FALSE, normal, 131, FALSE, 0.6, flat, 0, normal, 0\n2
81, 57, Male, Cleveland, asymptomatic, 110, 335, FALSE, normal, 143, TRUE, 3, flat, 1, reversa
ble defect, 2\n282, 47, Male, Cleveland, non-anginal, 130, 253, FALSE, normal, 179, FALSE, 0
, upsloping, 0, normal, 0\n283, 55, Female, Cleveland, asymptomatic, 128, 205, FALSE, st-t
abnormality, 130, TRUE, 2, flat, 1, reversable
defect, 3\n284, 35, Male, Cleveland, atypical angina, 122, 192, FALSE, normal, 174, FALSE, 0
, upsloping, 0, normal, 0\n285, 61, Male, Cleveland, asymptomatic, 148, 203, FALSE, normal, 1
61, FALSE, 0, upsloping, 1, reversable
defect, 2\n286, 58, Male, Cleveland, asymptomatic, 114, 318, FALSE, st-t
abnormality, 140, FALSE, 4.4, downsloping, 3, fixed
defect, 4\n287, 58, Female, Cleveland, asymptomatic, 170, 225, TRUE, lv
hypertrophy, 146, TRUE, 2.8, flat, 2, fixed defect, 2\n288, 58, Male, Cleveland, atypical
angina, 125, 220, FALSE, normal, 144, FALSE, 0.4, flat, , reversable
defect, 0\n289, 56, Male, Cleveland, atypical angina, 130, 221, FALSE, lv
hypertrophy, 163, FALSE, 0, upsloping, 0, reversable
defect, 0\n290, 56, Male, Cleveland, atypical angina, 120, 240, FALSE, normal, 169, FALSE, 0
, downsloping, 0, normal, 0\n291, 67, Male, Cleveland, non-anginal, 152, 212, FALSE, lv
hypertrophy, 150, FALSE, 0.8, flat, 0, reversable
defect, 1\n292, 55, Female, Cleveland, atypical angina, 132, 342, FALSE, normal, 166, FALSE
, 1.2, upsloping, 0, normal, 0\n293, 44, Male, Cleveland, asymptomatic, 120, 169, FALSE, norm
al, 144, TRUE, 2.8, downsloping, 0, fixed
defect, 2\n294, 63, Male, Cleveland, asymptomatic, 140, 187, FALSE, lv

hypertrophy,144,TRUE,4,upsloping,2,reversable defect,2\n295,63,Female,Cleveland,
 asymptomatic,124,197,FALSE,normal,136,TRUE,0,flat,0,normal,1\n296,41,Male,Clevel
 and,atypical angina,120,157,FALSE,normal,182,FALSE,0,upsloping,0,normal,0\n297,5
 9,Male,Cleveland,asymptomatic,164,176,TRUE,lv
 hypertrophy,90,FALSE,1,flat,2,fixed defect,3\n298,57,Female,Cleveland,asymptomat
 ic,140,241,FALSE,normal,123,TRUE,0.2,flat,0,reversable
 defect,1\n299,45,Male,Cleveland,typical
 angina,110,264,FALSE,normal,132,FALSE,1.2,flat,0,reversable defect,1\n300,68,Mal
 e,Cleveland,asymptomatic,144,193,TRUE,normal,141,FALSE,3.4,flat,2,reversable def
 ect,2\n301,57,Male,Cleveland,asymptomatic,130,131,FALSE,normal,115,TRUE,1.2,flat
 ,1,reversable defect,3\n302,57,Female,Cleveland,atypical angina,130,236,FALSE,lv
 hypertrophy,174,FALSE,0,flat,1,normal,1\n303,38,Male,Cleveland,non-anginal,138,1
 75,FALSE,normal,173,FALSE,0,upsloping,,normal,0\n304,28,Male,Cleveland,atypical
 angina,130,132,FALSE,lv
 hypertrophy,185,FALSE,0,,,0\n305,29,Male,Hungary,atypical
 angina,120,243,FALSE,normal,160,FALSE,0,,,0\n306,29,Male,Hungary,atypical
 angina,140,,FALSE,normal,170,FALSE,0,,,0\n307,30,Female,Hungary,typical
 angina,170,237,FALSE,st-t abnormality,170,FALSE,0,,,fixed
 defect,0\n308,31,Female,Hungary,atypical angina,100,219,FALSE,st-t
 abnormality,150,FALSE,0,,,0\n309,32,Female,Hungary,atypical
 angina,105,198,FALSE,normal,165,FALSE,0,,,0\n310,32,Male,Hungary,atypical
 angina,110,225,FALSE,normal,184,FALSE,0,,,0\n311,32,Male,Hungary,atypical
 angina,125,254,FALSE,normal,155,FALSE,0,,,0\n312,33,Male,Hungary,non-
 anginal,120,298,FALSE,normal,185,FALSE,0,,,0\n313,34,Female,Hungary,atypical
 angina,130,161,FALSE,normal,190,FALSE,0,,,0\n314,34,Male,Hungary,atypical
 angina,150,214,FALSE,st-t
 abnormality,168,FALSE,0,,,0\n315,34,Male,Hungary,atypical
 angina,98,220,FALSE,normal,150,FALSE,0,,,0\n316,35,Female,Hungary,typical
 angina,120,160,FALSE,st-t abnormality,185,FALSE,0,,,0\n317,35,Female,Hungary,as
 ymptomatic,140,167,FALSE,normal,150,FALSE,0,,,0\n318,35,Male,Hungary,atypical
 angina,120,308,FALSE,lv
 hypertrophy,180,FALSE,0,,,0\n319,35,Male,Hungary,atypical
 angina,150,264,FALSE,normal,168,FALSE,0,,,0\n320,36,Male,Hungary,atypical
 angina,120,166,FALSE,normal,180,FALSE,0,,,0\n321,36,Male,Hungary,non-anginal,11
 2,340,FALSE,normal,184,FALSE,1,flat,,normal,0\n322,36,Male,Hungary,non-
 anginal,130,209,FALSE,normal,178,FALSE,0,,,0\n323,36,Male,Hungary,non-
 anginal,150,160,FALSE,normal,172,FALSE,0,,,0\n324,37,Female,Hungary,atypical
 angina,120,260,FALSE,normal,130,FALSE,0,,,0\n325,37,Female,Hungary,non-anginal,
 130,211,FALSE,normal,142,FALSE,0,,,0\n326,37,Female,Hungary,asymptomatic,130,17
 3,FALSE,st-t abnormality,184,FALSE,0,,,0\n327,37,Male,Hungary,atypical
 angina,130,283,FALSE,st-t abnormality,98,FALSE,0,,,0\n328,37,Male,Hungary,non-a
 nginal,130,194,FALSE,normal,150,FALSE,0,,,0\n329,37,Male,Hungary,asymptomatic,1
 20,223,FALSE,normal,168,FALSE,0,,,normal,0\n330,37,Male,Hungary,asymptomatic,130
 ,315,FALSE,normal,158,FALSE,0,,,0\n331,38,Female,Hungary,atypical
 angina,120,275,,normal,129,FALSE,0,,,0\n332,38,Male,Hungary,atypical
 angina,140,297,FALSE,normal,150,FALSE,0,,,0\n333,38,Male,Hungary,non-
 anginal,145,292,FALSE,normal,130,FALSE,0,,,0\n334,39,Female,Hungary,non-

anginal,110,182,FALSE,st-t
 abnormality,180,FALSE,0,,,0\n335,39,Male,Hungary,atypical
 angina,120,,FALSE,st-t
 abnormality,146,FALSE,2,upsloping,,,0\n336,39,Male,Hungary,atypical
 angina,120,200,FALSE,normal,160,TRUE,1,flat,,,0\n337,39,Male,Hungary,atypical
 angina,120,204,FALSE,normal,145,FALSE,0,,,0\n338,39,Male,Hungary,atypical
 angina,130,,FALSE,normal,120,FALSE,0,,,0\n339,39,Male,Hungary,atypical
 angina,190,241,FALSE,normal,106,FALSE,0,,,0\n340,39,Male,Hungary,non-
 anginal,120,339,FALSE,normal,170,FALSE,0,,,0\n341,39,Male,Hungary,non-anginal,1
 60,147,TRUE,normal,160,FALSE,0,,,0\n342,39,Male,Hungary,asymptomatic,110,273,FA
 LSE,normal,132,FALSE,0,,,0\n343,39,Male,Hungary,asymptomatic,130,307,FALSE,norm
 al,140,FALSE,0,,,0\n344,40,Male,Hungary,atypical
 angina,130,275,FALSE,normal,150,FALSE,0,,,0\n345,40,Male,Hungary,atypical
 angina,140,289,FALSE,normal,172,FALSE,0,,,0\n346,40,Male,Hungary,non-
 anginal,130,215,FALSE,normal,138,FALSE,0,,,0\n347,40,Male,Hungary,non-
 anginal,130,281,FALSE,normal,167,FALSE,0,,,0\n348,40,Male,Hungary,non-
 anginal,140,,FALSE,normal,188,FALSE,0,,,0\n349,41,Female,Hungary,atypical
 angina,110,250,FALSE,st-t
 abnormality,142,FALSE,0,,,0\n350,41,Female,Hungary,atypical
 angina,125,184,FALSE,normal,180,FALSE,0,,,0\n351,41,Female,Hungary,atypical
 angina,130,245,FALSE,normal,150,FALSE,0,,,0\n352,41,Male,Hungary,atypical
 angina,120,291,FALSE,st-t
 abnormality,160,FALSE,0,,,0\n353,41,Male,Hungary,atypical
 angina,120,295,FALSE,normal,170,FALSE,0,,,0\n354,41,Male,Hungary,atypical
 angina,125,269,FALSE,normal,144,FALSE,0,,,0\n355,41,Male,Hungary,asymptomatic,112,25
 0,FALSE,normal,142,FALSE,0,,,0\n356,42,Female,Hungary,non-
 anginal,115,211,FALSE,st-t
 abnormality,137,FALSE,0,,,0\n357,42,Male,Hungary,atypical
 angina,120,196,FALSE,normal,150,FALSE,0,,,0\n358,42,Male,Hungary,atypical
 angina,120,198,FALSE,normal,155,FALSE,0,,,0\n359,42,Male,Hungary,atypical
 angina,150,268,FALSE,normal,136,FALSE,0,,,0\n360,42,Male,Hungary,non-
 anginal,120,228,FALSE,normal,152,TRUE,1.5,flat,,,0\n361,42,Male,Hungary,non-angi
 nal,160,147,FALSE,normal,146,FALSE,0,,,0\n362,42,Male,Hungary,asymptomatic,140,
 358,FALSE,normal,170,FALSE,0,,,0\n363,43,Female,Hungary,typical
 angina,100,223,FALSE,normal,142,FALSE,0,,,0\n364,43,Female,Hungary,atypical
 angina,120,201,FALSE,normal,165,FALSE,0,,,0\n365,43,Female,Hungary,atypical
 angina,120,215,FALSE,st-t
 abnormality,175,FALSE,0,,,0\n366,43,Female,Hungary,atypical
 angina,120,249,FALSE,st-t
 abnormality,176,FALSE,0,,,0\n367,43,Female,Hungary,atypical
 angina,120,266,FALSE,normal,118,FALSE,0,,,0\n368,43,Female,Hungary,atypical
 angina,150,186,FALSE,normal,154,FALSE,0,,,0\n369,43,Female,Hungary,non-
 anginal,150,,FALSE,normal,175,FALSE,0,,,normal,0\n370,43,Male,Hungary,atypical
 angina,142,207,FALSE,normal,138,FALSE,0,,,0\n371,44,Female,Hungary,asymptomatic,
 120,218,FALSE,st-t abnormality,115,FALSE,0,,,0\n372,44,Male,Hungary,atypical
 angina,120,184,FALSE,normal,142,FALSE,1,flat,,,0\n373,44,Male,Hungary,atypical
 angina,130,215,FALSE,normal,135,FALSE,0,,,0\n374,44,Male,Hungary,asymptomatic,15

0,412,FALSE,normal,170,FALSE,0,,,0\n375,45,Female,Hungary,atypical
 angina,130,237,FALSE,normal,170,FALSE,0,,,0\n376,45,Female,Hungary,atypical
 ina,180,,FALSE,normal,180,FALSE,0,,,0\n377,45,Female,Hungary,asymptomatic,132,2
 97,FALSE,normal,144,FALSE,0,,,0\n378,45,Male,Hungary,atypical
 angina,140,224,TRUE,normal,122,FALSE,0,,,0\n379,45,Male,Hungary,non-anginal,135
 ,,FALSE,normal,110,FALSE,0,,,0\n380,45,Male,Hungary,asymptomatic,120,225,FALSE,
 normal,140,FALSE,0,,,0\n381,45,Male,Hungary,asymptomatic,140,224,FALSE,normal,1
 44,FALSE,0,,,0\n382,46,Female,Hungary,asymptomatic,130,238,FALSE,normal,90,FALS
 E,0,,,0\n383,46,Male,Hungary,atypical
 angina,140,275,FALSE,normal,165,TRUE,0,,,0\n384,46,Male,Hungary,non-
 anginal,120,230,FALSE,normal,150,FALSE,0,,,0\n385,46,Male,Hungary,non-anginal,1
 50,163,,normal,116,FALSE,0,,,0\n386,46,Male,Hungary,asymptomatic,110,238,FALSE,
 st-t abnormality,140,TRUE,1,flat,,normal,0\n387,46,Male,Hungary,asymptomatic,110
 ,240,FALSE,st-t abnormality,140,FALSE,0,,,normal,0\n388,46,Male,Hungary,asymptom
 atic,180,280,FALSE,st-t
 abnormality,120,FALSE,0,,,0\n389,47,Female,Hungary,atypical angina,140,257,FALS
 E,normal,135,FALSE,1,upsloping,,,0\n390,47,Female,Hungary,non-
 anginal,130,,FALSE,normal,145,FALSE,2,flat,,,0\n391,47,Male,Hungary,typical
 angina,110,249,FALSE,normal,150,FALSE,0,,,0\n392,47,Male,Hungary,atypical angin
 a,160,263,FALSE,normal,174,FALSE,0,,,0\n393,47,Male,Hungary,asymptomatic,140,27
 6,TRUE,normal,125,TRUE,0,,,0\n394,48,Female,Hungary,atypical
 angina,,308,FALSE,st-t
 abnormality,,,2,upsloping,,,0\n395,48,Female,Hungary,atypical
 angina,120,,TRUE,st-t
 abnormality,148,FALSE,0,,,0\n396,48,Female,Hungary,atypical
 angina,120,284,FALSE,normal,120,FALSE,0,,,0\n397,48,Female,Hungary,non-anginal,
 120,195,FALSE,normal,125,FALSE,0,,,0\n398,48,Female,Hungary,asymptomatic,108,16
 3,FALSE,normal,175,FALSE,2,upsloping,,,0\n399,48,Female,Hungary,asymptomatic,120
 ,254,FALSE,st-t abnormality,110,FALSE,0,,,0\n400,48,Female,Hungary,asymptomatic
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 angina,100,,FALSE,normal,100,FALSE,0,,,0\n402,48,Male,Hungary,atypical
 angina,130,245,FALSE,normal,160,FALSE,0,,,0\n403,48,Male,Hungary,atypical
 angina,140,238,FALSE,normal,118,FALSE,0,,,0\n404,48,Male,Hungary,non-
 anginal,110,211,FALSE,normal,138,FALSE,0,,,fixed
 defect,0\n405,49,Female,Hungary,atypical
 angina,110,,FALSE,normal,160,FALSE,0,,,0\n406,49,Female,Hungary,atypical
 angina,110,,FALSE,normal,160,FALSE,0,,,0\n407,49,Female,Hungary,atypical
 angina,124,201,FALSE,normal,164,FALSE,0,,,0\n408,49,Female,Hungary,non-
 anginal,130,207,FALSE,st-t
 abnormality,135,FALSE,0,,,0\n409,49,Male,Hungary,atypical
 angina,100,253,FALSE,normal,174,FALSE,0,,,0\n410,49,Male,Hungary,non-anginal,14
 0,187,FALSE,normal,172,FALSE,0,,,0\n411,49,Male,Hungary,asymptomatic,120,297,,n
 ormal,132,FALSE,1,flat,,,0\n412,49,Male,Hungary,asymptomatic,140,,FALSE,normal,1
 30,FALSE,0,,,0\n413,50,Female,Hungary,atypical angina,110,202,FALSE,normal,145,
 FALSE,0,,,0\n414,50,Female,Hungary,asymptomatic,120,328,FALSE,normal,110,TRUE,1
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 angina,120,168,FALSE,normal,160,FALSE,0,,0,,0\n416,50,Male,Hungary,atypical

angina,140,216,FALSE,normal,170,FALSE,0,,normal,0\n417,50,Male,Hungary,atypical
 angina,170,209,FALSE,st-t abnormality,116,FALSE,0,,0\n418,50,Male,Hungary,asym
 ptomatic,140,129,FALSE,normal,135,FALSE,0,,0\n419,50,Male,Hungary,asymptomatic
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 anginal,110,190,FALSE,normal,120,FALSE,0,,0\n422,51,Female,Hungary,non-anginal
 ,130,220,FALSE,normal,160,TRUE,2,upsloping,,0\n423,51,Female,Hungary,non-angina
 l,150,200,FALSE,normal,120,FALSE,0.5,upsloping,,0\n424,51,Male,Hungary,atypical
 angina,125,188,FALSE,normal,145,FALSE,0,,0\n425,51,Male,Hungary,atypical angin
 a,130,224,FALSE,normal,150,FALSE,0,,0\n426,51,Male,Hungary,asymptomatic,130,17
 9,FALSE,normal,100,FALSE,0,,reversable defect,0\n427,52,Female,Hungary,atypical
 angina,120,210,FALSE,normal,148,FALSE,0,,0\n428,52,Female,Hungary,atypical
 angina,140,,FALSE,normal,140,FALSE,0,,0\n429,52,Female,Hungary,non-anginal,125
 ,272,FALSE,normal,139,FALSE,0,,0\n430,52,Female,Hungary,asymptomatic,130,180,F
 ALSE,normal,140,TRUE,1.5,flat,,0\n431,52,Male,Hungary,atypical
 angina,120,284,FALSE,normal,118,FALSE,0,,0\n432,52,Male,Hungary,atypical
 angina,140,100,FALSE,normal,138,TRUE,0,,0\n433,52,Male,Hungary,atypical
 angina,160,196,FALSE,normal,165,FALSE,0,,0\n434,52,Male,Hungary,non-
 anginal,140,259,FALSE,st-t
 abnormality,170,FALSE,0,,0\n435,53,Female,Hungary,atypical
 angina,113,468,,normal,127,FALSE,0,,0\n436,53,Female,Hungary,atypical
 angina,140,216,FALSE,normal,142,TRUE,2,flat,,0\n437,53,Female,Hungary,non-
 anginal,120,274,FALSE,normal,130,FALSE,0,,0\n438,53,Male,Hungary,atypical
 angina,120,,FALSE,normal,132,FALSE,0,,0\n439,53,Male,Hungary,atypical
 angina,140,320,FALSE,normal,162,FALSE,0,,0\n440,53,Male,Hungary,non-anginal,12
 0,195,FALSE,normal,140,FALSE,0,,0\n441,53,Male,Hungary,asymptomatic,124,260,FA
 LSE,st-t abnormality,112,TRUE,3,flat,,0\n442,53,Male,Hungary,asymptomatic,130,1
 82,FALSE,normal,148,FALSE,0,,0\n443,53,Male,Hungary,asymptomatic,140,243,FALSE
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 rmal,138,FALSE,1,upsloping,,0\n445,54,Female,Hungary,atypical
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 na,120,273,FALSE,normal,150,FALSE,1.5,flat,,0\n447,54,Female,Hungary,atypical
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 abnormality,155,FALSE,0,,0\n448,54,Female,Hungary,atypical
 angina,140,309,,st-t
 abnormality,140,FALSE,0,,0\n449,54,Female,Hungary,atypical
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 angina,110,208,FALSE,normal,142,FALSE,0,,0\n453,54,Male,Hungary,atypical
 angina,120,238,FALSE,normal,154,FALSE,0,,0\n454,54,Male,Hungary,atypical
 angina,120,246,FALSE,normal,110,FALSE,0,,0\n455,54,Male,Hungary,atypical
 angina,160,195,FALSE,st-t
 abnormality,130,FALSE,1,upsloping,,0\n456,54,Male,Hungary,atypical
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E, st-t abnormality, 134, FALSE, 1, upsloping, , , 0\n460, 55, Female, Hungary, atypical
 angina, 110, 344, FALSE, st-t
 abnormality, 160, FALSE, 0, , , 0\n461, 55, Female, Hungary, atypical
 angina, 122, 320, FALSE, normal, 155, FALSE, 0, , , 0\n462, 55, Female, Hungary, atypical
 angina, 130, 394, FALSE, lv
 hypertrophy, 150, FALSE, 0, , , 0\n463, 55, Male, Hungary, atypical
 angina, 120, 256, TRUE, normal, 137, FALSE, 0, , , reversable
 defect, 0\n464, 55, Male, Hungary, atypical
 angina, 140, 196, FALSE, normal, 150, FALSE, 0, , , reversable
 defect, 0\n465, 55, Male, Hungary, atypical
 angina, 145, 326, FALSE, normal, 155, FALSE, 0, , , 0\n466, 55, Male, Hungary, non-
 anginal, 110, 277, FALSE, normal, 160, FALSE, 0, , , 0\n467, 55, Male, Hungary, non-
 anginal, 120, 220, FALSE, lv hypertrophy, 134, FALSE, 0, , , 0\n468, 55, Male, Hungary, asymp-
 tomatic, 120, 270, FALSE, normal, 140, FALSE, 0, , , 0\n469, 55, Male, Hungary, asymptomatic,
 140, 229, FALSE, normal, 110, TRUE, 0.5, flat, , , 0\n470, 56, Female, Hungary, non-
 anginal, 130, 219, , st-t abnormality, 164, FALSE, 0, , , reversable
 defect, 0\n471, 56, Male, Hungary, atypical
 angina, 130, 184, FALSE, normal, 100, FALSE, 0, , , 0\n472, 56, Male, Hungary, non-
 anginal, 130, , FALSE, normal, 114, FALSE, 0, , , 0\n473, 56, Male, Hungary, non-
 anginal, 130, 276, FALSE, normal, 128, TRUE, 1, upsloping, , fixed defect, 0\n474, 56, Male, H
 ungary, asymptomatic, 120, 85, FALSE, normal, 140, FALSE, 0, , , 0\n475, 57, Female, Hungary,
 typical angina, 130, 308, FALSE, normal, 98, FALSE, 1, flat, , , 0\n476, 57, Female, Hungary, a
 symptomatic, 180, 347, FALSE, st-t
 abnormality, 126, TRUE, 0.8, flat, , , 0\n477, 57, Male, Hungary, atypical
 angina, 140, 260, TRUE, normal, 140, FALSE, 0, , , fixed
 defect, 0\n478, 58, Male, Hungary, atypical
 angina, 130, 230, FALSE, normal, 150, FALSE, 0, , , 0\n479, 58, Male, Hungary, atypical
 angina, 130, 251, FALSE, normal, 110, FALSE, 0, , , 0\n480, 58, Male, Hungary, non-anginal, 14
 0, 179, FALSE, normal, 160, FALSE, 0, , , 0\n481, 58, Male, Hungary, asymptomatic, 135, 222, FA
 LSE, normal, 100, FALSE, 0, , , 0\n482, 59, Female, Hungary, atypical
 angina, 130, 188, FALSE, normal, 124, FALSE, 1, flat, , , 0\n483, 59, Male, Hungary, atypical
 angina, 140, 287, FALSE, normal, 150, FALSE, 0, , , 0\n484, 59, Male, Hungary, non-
 anginal, 130, 318, FALSE, normal, 120, TRUE, 1, flat, , normal, 0\n485, 59, Male, Hungary, non-
 anginal, 180, 213, FALSE, normal, 100, FALSE, 0, , , 0\n486, 59, Male, Hungary, asymptomatic,
 140, , FALSE, normal, 140, FALSE, 0, , 0, 0\n487, 60, Male, Hungary, non-
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 mptomatic, 130, 294, FALSE, st-t abnormality, 120, TRUE, 1, flat, , , 0\n489, 61, Male, Hungar
 y, asymptomatic, 125, 292, FALSE, st-t
 abnormality, 115, TRUE, 0, , , 0\n490, 62, Female, Hungary, typical
 angina, 160, 193, FALSE, normal, 116, FALSE, 0, , , 0\n491, 62, Male, Hungary, atypical angin
 a, 140, 271, FALSE, normal, 152, FALSE, 1, upsloping, , , 0\n492, 31, Male, Hungary, asymptomat
 ic, 120, 270, FALSE, normal, 153, TRUE, 1.5, flat, , , 1\n493, 33, Female, Hungary, asymptomati
 c, 100, 246, FALSE, normal, 150, TRUE, 1, flat, , , 1\n494, 34, Male, Hungary, typical
 angina, 140, 156, FALSE, normal, 180, FALSE, 0, , , 1\n495, 35, Male, Hungary, atypical
 angina, 110, 257, FALSE, normal, 140, FALSE, 0, , , 1\n496, 36, Male, Hungary, atypical angin
 a, 120, 267, FALSE, normal, 160, FALSE, 3, flat, , , 1\n497, 37, Male, Hungary, asymptomatic, 14
 0, 207, FALSE, normal, 130, TRUE, 1.5, flat, , , 1\n498, 38, Male, Hungary, asymptomatic, 110, 1

96,FALSE,normal,166,FALSE,0,,,1\n499,38,Male,Hungary,asymptomatic,120,282,FALSE,normal,170,FALSE,0,,,1\n500,38,Male,Hungary,asymptomatic,92,117,FALSE,normal,134,TRUE,2.5,flat,,,1\n501,40,Male,Hungary,asymptomatic,120,466,,normal,152,TRUE,1,flat,,fixed defect,1\n502,41,Male,Hungary,asymptomatic,110,289,FALSE,normal,170,FALSE,0,,,fixed defect,1\n503,41,Male,Hungary,asymptomatic,120,237,,normal,138,TRUE,1,flat,,,1\n504,43,Male,Hungary,asymptomatic,150,247,FALSE,normal,130,TRUE,2,flat,,,1\n505,46,Male,Hungary,asymptomatic,110,202,FALSE,normal,150,TRUE,0,,,1\n506,46,Male,Hungary,asymptomatic,118,186,FALSE,normal,124,FALSE,0,,,reversible defect,1\n507,46,Male,Hungary,asymptomatic,120,277,FALSE,normal,125,TRUE,1,flat,,,1\n508,47,Male,Hungary,non-anginal,140,193,FALSE,normal,145,TRUE,1,flat,,,1\n509,47,Male,Hungary,asymptomatic,150,226,FALSE,normal,98,TRUE,1.5,flat,0,reversible defect,1\n510,48,Male,Hungary,asymptomatic,106,263,TRUE,normal,110,FALSE,0,,,1\n511,48,Male,Hungary,asymptomatic,120,260,FALSE,normal,115,FALSE,2,flat,,,1\n512,48,Male,Hungary,asymptomatic,160,268,FALSE,normal,103,TRUE,1,flat,,,1\n513,49,Female,Hungary,non-anginal,160,180,FALSE,normal,156,FALSE,1,flat,,,1\n514,49,Male,Hungary,non-anginal,115,265,FALSE,normal,175,FALSE,0,,,1\n515,49,Male,Hungary,asymptomatic,130,206,FALSE,normal,170,FALSE,0,,,1\n516,50,Female,Hungary,non-anginal,140,288,FALSE,normal,140,TRUE,0,,,reversible defect,1\n517,50,Male,Hungary,asymptomatic,145,264,FALSE,normal,150,FALSE,0,,,1\n518,51,Female,Hungary,asymptomatic,160,303,FALSE,normal,150,TRUE,1,flat,,,1\n519,52,Male,Hungary,asymptomatic,130,225,FALSE,normal,120,TRUE,2,flat,,,1\n520,54,Male,Hungary,asymptomatic,125,216,FALSE,normal,140,FALSE,0,,,1\n521,54,Male,Hungary,asymptomatic,125,224,FALSE,normal,122,FALSE,2,flat,,,1\n522,55,Male,Hungary,asymptomatic,140,201,FALSE,normal,130,TRUE,3,flat,,,1\n523,57,Male,Hungary,atypical angina,140,265,FALSE,st-t abnormality,145,TRUE,1,flat,,,1\n524,58,Male,Hungary,non-anginal,130,213,FALSE,st-t abnormality,140,FALSE,0,,,fixed defect,1\n525,59,Female,Hungary,asymptomatic,130,338,TRUE,st-t abnormality,130,TRUE,1.5,flat,,,1\n526,60,Male,Hungary,asymptomatic,100,248,FALSE,normal,125,FALSE,1,flat,,,1\n527,63,Male,Hungary,asymptomatic,150,223,FALSE,normal,115,FALSE,0,,,1\n528,65,Male,Hungary,asymptomatic,140,306,TRUE,normal,87,TRUE,1.5,flat,,,1\n529,32,Male,Hungary,asymptomatic,118,529,FALSE,normal,130,FALSE,0,,,1\n530,38,Male,Hungary,asymptomatic,110,,FALSE,normal,150,TRUE,1,flat,,,1\n531,39,Male,Hungary,asymptomatic,110,280,FALSE,normal,150,FALSE,0,,,fixed defect,1\n532,40,Female,Hungary,asymptomatic,150,392,FALSE,normal,130,FALSE,2,flat,,fixed defect,1\n533,43,Male,Hungary,typical angina,120,291,FALSE,st-t abnormality,155,FALSE,0,,,1\n534,45,Male,Hungary,asymptomatic,130,219,FALSE,st-t abnormality,130,TRUE,1,flat,,,1\n535,46,Male,Hungary,asymptomatic,120,231,FALSE,normal,115,TRUE,0,,,1\n536,46,Male,Hungary,asymptomatic,130,222,FALSE,normal,112,FALSE,0,,,1\n537,48,Male,Hungary,asymptomatic,122,275,TRUE,st-t abnormality,150,TRUE,2,down sloping,,,1\n538,48,Male,Hungary,asymptomatic,160,193,FALSE,normal,102,TRUE,3,flat,,,1\n539,48,Male,Hungary,asymptomatic,160,329,FALSE,normal,92,TRUE,1.5,flat,,,1\n540,48,Male,Hungary,asymptomatic,160,355,FALSE,normal,99,TRUE,2,flat,,,1\n541,50,Male,Hungary,asymptomatic,130,233,FALSE,normal,121,TRUE,2,flat,,reversible defect,1\n542,52,Male,Hungary,asymptomatic,120,182,FALSE,normal,150,FALSE,0,,,1\n543,52,Male,Hungary,asymptomatic,170,,FALSE,normal,126,TRUE,1.5,flat,,,1\n544,

53, Male, Hungary, asymptomatic, 120, 246, FALSE, normal, 116, TRUE, 0, , , 1\n545, 54, Male, Hungary, non-anginal, 120, 237, FALSE, normal, 150, TRUE, 1.5, , , reversable defect, 1\n546, 54, Male, Hungary, asymptomatic, 130, 242, FALSE, normal, 91, TRUE, 1, flat, , , 1\n547, 54, Male, Hungary, asymptomatic, 130, 603, TRUE, normal, 125, TRUE, 1, flat, , , 1\n548, 54, Male, Hungary, asymptomatic, 140, , FALSE, normal, 118, TRUE, 0, , , 1\n549, 54, Male, Hungary, asymptomatic, 200, 198, FALSE, normal, 142, TRUE, 2, flat, , , 1\n550, 55, Male, Hungary, asymptomatic, 140, 268, FALSE, normal, 128, TRUE, 1.5, flat, , , 1\n551, 56, Male, Hungary, asymptomatic, 150, 213, TRUE, normal, 125, TRUE, 1, flat, , , 1\n552, 57, Male, Hungary, asymptomatic, 150, 255, FALSE, normal, 92, TRUE, 3, flat, , , 1\n553, 58, Male, Hungary, non-anginal, 160, 211, TRUE, st-t abnormality, 92, FALSE, 0, , , 1\n554, 58, Male, Hungary, asymptomatic, 130, 263, FALSE, normal, 140, TRUE, 2, flat, , , 1\n555, 41, Male, Hungary, asymptomatic, 130, 172, FALSE, st-t abnormality, 130, FALSE, 2, flat, , , 1\n556, 43, Male, Hungary, asymptomatic, 120, 175, FALSE, normal, 120, TRUE, 1, flat, , , reversable defect, 1\n557, 44, Male, Hungary, atypical angina, 150, 288, FALSE, normal, 150, TRUE, 3, flat, , , 1\n558, 44, Male, Hungary, asymptomatic, 130, 290, FALSE, normal, 100, TRUE, 2, flat, , , 1\n559, 46, Male, Hungary, typical angina, 140, 272, TRUE, normal, 175, FALSE, 2, flat, , , 1\n560, 47, Female, Hungary, non-anginal, 135, 248, TRUE, normal, 170, FALSE, 0, , , 1\n561, 48, Female, Hungary, asymptomatic, 138, 214, FALSE, normal, 108, TRUE, 1.5, flat, , , 1\n562, 49, Male, Hungary, asymptomatic, 130, 341, FALSE, normal, 120, TRUE, 1, flat, , , 1\n563, 49, Male, Hungary, asymptomatic, 140, 234, FALSE, normal, 140, TRUE, 1, flat, , , 1\n564, 51, Male, Hungary, non-anginal, 135, 160, FALSE, normal, 150, FALSE, 2, flat, , , 1\n565, 52, Male, Hungary, asymptomatic, 112, 342, FALSE, st-t abnormality, 96, TRUE, 1, flat, , , 1\n566, 52, Male, Hungary, asymptomatic, 130, 298, FALSE, normal, 110, TRUE, 1, flat, , , 1\n567, 52, Male, Hungary, asymptomatic, 140, 404, FALSE, normal, 124, TRUE, 2, flat, , , 1\n568, 52, Male, Hungary, asymptomatic, 160, 246, FALSE, st-t abnormality, 82, TRUE, 4, flat, , , 1\n569, 53, Male, Hungary, non-anginal, 145, 518, FALSE, normal, 130, FALSE, 0, , , 1\n570, 53, Male, Hungary, asymptomatic, 180, 285, FALSE, st-t abnormality, 120, TRUE, 1.5, flat, , , 1\n571, 54, Male, Hungary, asymptomatic, 140, 216, FALSE, normal, 105, FALSE, 1.5, flat, , , 1\n572, 55, Male, Hungary, typical angina, 140, 295, FALSE, , 136, FALSE, 0, , , 1\n573, 55, Male, Hungary, atypical angina, 160, 292, TRUE, normal, 143, TRUE, 2, flat, , , 1\n574, 55, Male, Hungary, asymptomatic, 145, 248, FALSE, normal, 96, TRUE, 2, flat, , , 1\n575, 56, Female, Hungary, atypical angina, 120, 279, FALSE, normal, 150, FALSE, 1, flat, , , 1\n576, 56, Male, Hungary, asymptomatic, 150, 230, FALSE, st-t abnormality, 124, TRUE, 1.5, flat, , , 1\n577, 56, Male, Hungary, asymptomatic, 170, 388, FALSE, st-t abnormality, 122, TRUE, 2, flat, , , 1\n578, 58, Male, Hungary, atypical angina, 136, 164, FALSE, st-t abnormality, 99, TRUE, 2, flat, , , 1\n579, 59, Male, Hungary, asymptomatic, 130, , FALSE, normal, 125, FALSE, 0, , , 1\n580, 59, Male, Hungary, asymptomatic, 140, 264, TRUE, lv hypertrophy, 119, TRUE, 0, , , 1\n581, 65, Male, Hungary, asymptomatic, 170, 263, TRUE, normal, 112, TRUE, 2, flat, , , 1\n582, 66, Male, Hungary, asymptomatic, 140, , FALSE, normal, 94, TRUE, 1, flat, , , 1\n583, 41, Male, Hungary, asymptomatic, 120, 336, FALSE, normal, 118, TRUE, 3, flat, , , 1\n584, 43, Male, Hungary, asymptomatic, 140, 288, FALSE, normal, 135, TRUE, 2, flat, , , 1\n585, 44, Male, Hungary, asymptomatic, 135, 491, FALSE, normal, 135, FALSE, 0, , , 1\n586, 47, Female, Hungary, asymptomatic, 120, 205, FALSE, normal, 98, TRUE, 2, flat, , , fixed defect, 1\n587, 47, Male, Hungary, asymptomatic, 160, 291, FALSE, st-t abnormality, 158, TRUE, 3, flat, , , 1\n588, 49, Male, Hungary, asymptomatic, 128, 212, FALSE, normal, 96, TRUE, 0, , , 1\n589, 49, Male, Hungary, asymptomatic, 150, 222, FALSE, normal, 122, FALSE, 2, flat, , , 1\n590, 50, Male, Hungary, asymptomatic, 140, 231, FALSE, st-t abnormality, 140, T

RUE,5,flat,,1\n591,50, Male, Hungary, asymptomatic, 140, 341, FALSE, st-t abnormality, 125, TRUE, 2.5, flat,,1\n592,52, Male, Hungary, asymptomatic, 140, 266, FALSE, normal, 134, TRUE, 2, flat,,1\n593,52, Male, Hungary, asymptomatic, 160, 331, FALSE, normal, 94, TRUE, 2.5,,1\n594,54, Female, Hungary, non-anginal, 130, 294, FALSE, st-t abnormality, 100, TRUE, 0, flat,,1\n595,56, Male, Hungary, asymptomatic, 155, 342, TRUE, normal, 150, TRUE, 3, flat,,1\n596,58, Female, Hungary, atypical
 angina, 180, 393, FALSE, normal, 110, TRUE, 1, flat,, reversable
 defect, 1\n597, 65, Male, Hungary, asymptomatic, 130, 275, FALSE, st-t
 abnormality, 115, TRUE, 1, flat,,1\n598, 32, Male, Switzerland, typical angina, 95, 0,, normal, 127, FALSE, 0.7, upsloping,,1\n599, 34, Male, Switzerland, asymptomatic, 115, 0,, 154, FALSE, 0.2, upsloping,,1\n600, 35, Male, Switzerland, asymptomatic,,0,, normal, 130, TRUE,, , reversable defect, 3\n601, 36, Male, Switzerland, asymptomatic, 110, 0,, normal, 125, TRUE, 1, flat,, fixed defect, 1\n602, 38, Female, Switzerland, asymptomatic, 105, 0,, normal, 166, FALSE, 2.8, upsloping,, 2\n603, 38, Female, Switzerland, asymptomatic, 110, 0, FALSE, normal, 156, FALSE, 0, flat,, normal, 1\n604, 38, Male, Switzerland, non-anginal, 100, 0,, normal, 179, FALSE, -1.1, upsloping,, 0\n605, 38, Male, Switzerland, non-anginal, 115, 0, FALSE, normal, 128, TRUE, 0, flat,, reversable defect, 1\n606, 38, Male, Switzerland, asymptomatic, 135, 0,, normal, 150, FALSE, 0,, normal, 2\n607, 38, Male, Switzerland, asymptomatic, 150, 0,, normal, 120, TRUE,, , normal, 1\n608, 40, Male, Switzerland, asymptomatic, 95, 0,, st-t abnormality, 144, FALSE, 0, upsloping,, 2\n609, 41, Male, Switzerland, asymptomatic, 125, 0,, normal, 176, FALSE, 1.6, upsloping,, 2\n610, 42, Male, Switzerland, asymptomatic, 105, 0,, normal, 128, TRUE, -1.5, downsloping,, 1\n611, 42, Male, Switzerland, asymptomatic, 145, 0, FALSE, normal, 99, TRUE, 0, flat,, 2\n612, 43, Male, Switzerland, asymptomatic, 100, 0,, normal, 122, FALSE, 1.5, downsloping,, 3\n613, 43, Male, Switzerland, asymptomatic, 115, 0, FALSE, normal, 145, TRUE, 2, flat,, reversable
 defect, 4\n614, 43, Male, Switzerland, asymptomatic, 140, 0, FALSE, st-t
 abnormality, 140, TRUE, 0.5, upsloping,, reversable
 defect, 2\n615, 45, Male, Switzerland, non-anginal, 110, 0,, normal, 138, FALSE, -0.1, upsloping,, 0\n616, 46, Male, Switzerland, asymptomatic, 100, 0,, st-t abnormality, 133, FALSE, -2.6, flat,, 1\n617, 46, Male, Switzerland, asymptomatic, 115, 0, FALSE, normal, 113, TRUE, 1.5, flat,, reversable defect, 1\n618, 47, Male, Switzerland, non-anginal, 110, 0,, normal, 120, TRUE, 0,, normal, 1\n619, 47, Male, Switzerland, non-anginal, 155, 0, FALSE, normal, 118, TRUE, 1, flat,, normal, 3\n620, 47, Male, Switzerland, asymptomatic, 110, 0,, st-t abnormality, 149, FALSE, 2.1, upsloping,, 2\n621, 47, Male, Switzerland, asymptomatic, 160, 0, FALSE, normal, 124, TRUE, 0, flat,, reversable defect, 1\n622, 48, Male, Switzerland, asymptomatic, 115, 0,, normal, 128, FALSE, 0, flat,, fixed defect, 2\n623, 50, Female, Switzerland, asymptomatic, 160, 0,, normal, 110, FALSE, 0,, normal, 1\n624, 50, Male, Switzerland, asymptomatic, 115, 0, FALSE, normal, 120, TRUE, 0.5, flat,, fixed
 defect, 3\n625, 50, Male, Switzerland, asymptomatic, 120, 0, FALSE, st-t
 abnormality, 156, TRUE, 0, upsloping,, fixed defect, 3\n626, 50, Male, Switzerland, asymptomatic, 145, 0,, normal, 139, TRUE, 0.7, flat,, 1\n627, 51, Female, Switzerland, asymptomatic, 120, 0,, normal, 127, TRUE, 1.5, upsloping,, 2\n628, 51, Male, Switzerland, asymptomatic, 110, 0,, normal, 92, FALSE, 0, flat,, 4\n629, 51, Male, Switzerland, asymptomatic, 120, 0, TRUE, normal, 104, FALSE, 0, flat,, normal, 3\n630, 51, Male, Switzerland, asymptomatic, 130, 0,, normal, 170, FALSE, -0.7, upsloping,, 2\n631, 51, Male, Switzerland, asymptomatic, 130, 0, 0,, st-t abnormality, 163, FALSE, 0,, reversable defect, 1\n632, 51, Male, Switzerland, asymptomatic, 140, 0, FALSE, normal, 60, FALSE, 0, flat,, normal, 2\n633, 51, Male, Switzerland

and, asymptomatic, 95, 0, , normal, 126, FALSE, 2.2, flat, , , 2\n634, 52, Male, Switzerland, asymptomatic, 130, 0, , normal, 120, FALSE, 0, flat, , reversable defect, 2\n635, 52, Male, Switzerland, asymptomatic, 135, 0, , normal, 128, TRUE, 2, flat, , reversable defect, 2\n636, 52, Male, Switzerland, asymptomatic, 165, 0, , normal, 122, TRUE, 1, upsloping, , reversable defect, 2\n637, 52, Male, Switzerland, asymptomatic, 95, 0, , normal, 82, TRUE, , , , 2\n638, 53, Male, Switzerland, atypical angina, 120, 0, FALSE, normal, 95, FALSE, 0, flat, , normal, 3\n639, 53, Male, Switzerland, atypical angina, 130, 0, , st-t abnormality, 120, FALSE, 0.7, downsloping, , , 0\n640, 53, Male, Switzerland, non-anginal, 105, 0, FALSE, normal, 115, FALSE, 0, flat, , reversable defect, 1\n641, 53, Male, Switzerland, non-anginal, 160, 0, , lv hypertrophy, 122, TRUE, 0, , , reversable defect, 1\n642, 53, Male, Switzerland, asymptomatic, 120, 0, , normal, 120, FALSE, 0, flat, , reversable defect, 1\n643, 53, Male, Switzerland, asymptomatic, 125, 0, , normal, 120, FALSE, 1.5, upsloping, , , 4\n644, 53, Male, Switzerland, asymptomatic, 130, 0, FALSE, lv hypertrophy, 135, TRUE, 1, flat, , reversable defect, 2\n645, 53, Male, Switzerland, asymptomatic, 80, 0, , normal, 141, TRUE, 2, downsloping, , , 0\n646, 54, Male, Switzerland, asymptomatic, 120, 0, FALSE, normal, 155, FALSE, 0, flat, , reversable defect, 2\n647, 54, Male, Switzerland, asymptomatic, 130, 0, , normal, 110, TRUE, 3, flat, , reversable defect, 3\n648, 54, Male, Switzerland, asymptomatic, 180, 0, , normal, 150, FALSE, 1.5, flat, , reversable defect, 1\n649, 55, Male, Switzerland, atypical angina, 140, 0, , st-t abnormality, 150, FALSE, 0.2, upsloping, , , 0\n650, 55, Male, Switzerland, asymptomatic, 115, 0, , normal, 155, FALSE, 0.1, flat, , , 1\n651, 55, Male, Switzerland, asymptomatic, 120, 0, FALSE, st-t abnormality, 92, FALSE, 0.3, upsloping, , reversable defect, 4\n652, 55, Male, Switzerland, asymptomatic, 140, 0, FALSE, normal, 83, FALSE, 0, flat, , reversable defect, 2\n653, 56, Male, Switzerland, non-anginal, 120, 0, FALSE, normal, 97, FALSE, 0, flat, , reversable defect, 0\n654, 56, Male, Switzerland, non-anginal, 125, 0, , normal, 98, FALSE, -2, flat, , reversable defect, 2\n655, 56, Male, Switzerland, non-anginal, 155, 0, FALSE, st-t abnormality, 99, FALSE, 0, flat, , normal, 2\n656, 56, Male, Switzerland, asymptomatic, 115, 0, , st-t abnormality, 82, FALSE, -1, upsloping, , , 1\n657, 56, Male, Switzerland, asymptomatic, 120, 0, FALSE, st-t abnormality, 100, TRUE, -1, downsloping, , reversable defect, 2\n658, 56, Male, Switzerland, asymptomatic, 120, 0, FALSE, st-t abnormality, 148, FALSE, 0, flat, , , 2\n659, 56, Male, Switzerland, asymptomatic, 125, 0, TRUE, normal, 103, TRUE, 1, flat, , reversable defect, 3\n660, 56, Male, Switzerland, asymptomatic, 140, 0, , normal, 121, TRUE, 1.8, upsloping, , , 1\n661, 57, Male, Switzerland, non-anginal, 105, 0, , normal, 148, FALSE, 0.3, flat, , , 1\n662, 57, Male, Switzerland, asymptomatic, 110, 0, , st-t abnormality, 131, TRUE, 1.4, upsloping, 1, , 3\n663, 57, Male, Switzerland, asymptomatic, 140, 0, FALSE, normal, 120, TRUE, 2, flat, , fixed defect, 2\n664, 57, Male, Switzerland, asymptomatic, 140, 0, , normal, 100, TRUE, 0, , , fixed defect, 3\n665, 57, Male, Switzerland, asymptomatic, 160, 0, , normal, 98, TRUE, 2, flat, , reversable defect, 2\n666, 57, Male, Switzerland, asymptomatic, 95, 0, , normal, 182, FALSE, 0.7, downsloping, , , 1\n667, 58, Male, Switzerland, asymptomatic, 115, 0, , normal, 138, FALSE, 0.5, upsloping, , , 1\n668, 58, Male, Switzerland, asymptomatic, 130, 0, FALSE, st-t abnormality, 100, TRUE, 1, flat, , fixed defect, 4\n669, 58, Male, Switzerland, asymptomatic, 170, 0, , st-t abnormality, 105, TRUE, 0, , , normal, 1\n670, 59, Male, Switzerland, non-anginal, 125, 0, , normal, 175, FALSE, 2.6, flat, , , 1\n671, 59, Male, Switzerland, asymptomatic, 110, 0, , normal,

94,FALSE,0,,fixed defect,3\n672,59,Male,Switzerland,asymptomatic,120,0,FALSE,normal,115,FALSE,0,flat,,normal,2\n673,59,Male,Switzerland,asymptomatic,125,0,,normal,119,TRUE,0.9,upsloping,,1\n674,59,Male,Switzerland,asymptomatic,135,0,FALSE,normal,115,TRUE,1,flat,,reversible defect,1\n675,60,Male,Switzerland,non-anginal,115,0,,normal,143,FALSE,2.4,upsloping,,1\n676,60,Male,Switzerland,asymptomatic,125,0,,normal,110,FALSE,0.1,upsloping,2,,3\n677,60,Male,Switzerland,asymptomatic,130,0,,st-t abnormality,130,TRUE,1.1,downsloping,1,,1\n678,60,Male,Switzerland,asymptomatic,135,0,FALSE,normal,63,TRUE,0.5,upsloping,,reversible defect,3\n679,60,Male,Switzerland,asymptomatic,160,0,FALSE,st-t abnormality,99,TRUE,0.5,flat,,reversible defect,3\n680,60,Male,Switzerland,asymptomatic,160,0,,normal,149,FALSE,0.4,flat,,1\n681,61,Male,Switzerland,non-anginal,200,0,,st-t abnormality,70,FALSE,0,,normal,3\n682,61,Male,Switzerland,asymptomatic,105,0,,normal,110,TRUE,1.5,upsloping,,1\n683,61,Male,Switzerland,asymptomatic,110,0,,normal,113,FALSE,1.4,flat,,1\n684,61,Male,Switzerland,asymptomatic,125,0,FALSE,normal,105,TRUE,0,downsloping,,reversible defect,3\n685,61,Male,Switzerland,asymptomatic,130,0,FALSE,lv hypertrophy,115,FALSE,0,flat,,reversible defect,3\n686,61,Male,Switzerland,asymptomatic,130,0,,normal,77,FALSE,2.5,flat,,3\n687,61,Male,Switzerland,asymptomatic,150,0,FALSE,normal,105,TRUE,0,flat,,reversible defect,1\n688,61,Male,Switzerland,asymptomatic,150,0,FALSE,normal,117,TRUE,2,flat,,reversible defect,2\n689,61,Male,Switzerland,asymptomatic,160,0,TRUE,st-t abnormality,145,FALSE,1,flat,,reversible defect,2\n690,62,Female,Switzerland,typical angina,140,0,,normal,143,FALSE,0,,normal,2\n691,62,Female,Switzerland,asymptomatic,120,0,,st-t abnormality,123,TRUE,1.7,downsloping,,1\n692,62,Male,Switzerland,typical angina,120,0,,lv hypertrophy,134,FALSE,-0.8,flat,2,,1\n693,62,Male,Switzerland,non-anginal,160,0,FALSE,normal,72,TRUE,0,flat,,normal,3\n694,62,Male,Switzerland,asymptomatic,115,0,,normal,128,TRUE,2.5,downsloping,,2\n695,62,Male,Switzerland,asymptomatic,115,0,,normal,72,TRUE,-0.5,flat,,normal,1\n696,62,Male,Switzerland,asymptomatic,150,0,,st-t abnormality,78,FALSE,2,flat,,reversible defect,3\n697,63,Male,Switzerland,asymptomatic,100,0,,normal,109,FALSE,-0.9,flat,,1\n698,63,Male,Switzerland,asymptomatic,140,0,,lv hypertrophy,149,FALSE,2,upsloping,,2\n699,63,Male,Switzerland,asymptomatic,150,0,FALSE,normal,86,TRUE,2,flat,,3\n700,63,Male,Switzerland,asymptomatic,150,0,,st-t abnormality,154,FALSE,3.7,upsloping,,3\n701,63,Male,Switzerland,asymptomatic,185,0,FALSE,normal,98,TRUE,0,upsloping,,reversible defect,1\n702,64,Female,Switzerland,asymptomatic,200,0,FALSE,normal,140,TRUE,1,flat,,normal,3\n703,64,Female,Switzerland,asymptomatic,95,0,,normal,145,FALSE,1.1,downsloping,,1\n704,64,Male,Switzerland,asymptomatic,110,0,,normal,114,TRUE,1.3,downsloping,,1\n705,65,Male,Switzerland,asymptomatic,115,0,FALSE,normal,93,TRUE,0,flat,,reversible defect,1\n706,65,Male,Switzerland,asymptomatic,145,0,,st-t abnormality,67,FALSE,,fixed defect,3\n707,65,Male,Switzerland,asymptomatic,155,0,,normal,154,FALSE,1,upsloping,,0\n708,65,Male,Switzerland,asymptomatic,160,0,TRUE,st-t abnormality,122,FALSE,,reversible defect,3\n709,66,Female,Switzerland,asymptomatic,155,0,,normal,90,FALSE,0,,reversible defect,1\n710,66,Male,Switzerland,asymptomatic,150,0,FALSE,normal,108,TRUE,2,flat,,reversible defect,3\n711,67,Male,Switzerland,typical angina,145,0,FALSE,lv hypertrophy,125,

FALSE,0,flat,,normal,2\n712,68,Male,Switzerland,asymptomatic,135,0,FALSE,st-t
abnormality,120,TRUE,0,upsloping,,reversable defect,3\n713,68,Male,Switzerland,a
symptomatic,145,0,,normal,136,FALSE,1.8,upsloping,,1\n714,69,Male,Switzerland,a
symptomatic,135,0,FALSE,normal,130,FALSE,0,flat,,fixed
defect,1\n715,69,Male,Switzerland,asymptomatic,,0,FALSE,st-t
abnormality,,,,,reversable
defect,3\n716,70,Male,Switzerland,asymptomatic,115,0,FALSE,st-t
abnormality,92,TRUE,0,flat,,reversable defect,1\n717,70,Male,Switzerland,asympto
matic,140,0,TRUE,normal,157,TRUE,2,flat,,reversable
defect,3\n718,72,Male,Switzerland,non-anginal,160,0,,lv
hypertrophy,114,FALSE,1.6,flat,2,,0\n719,73,Female,Switzerland,non-
anginal,160,0,FALSE,st-t
abnormality,121,FALSE,0,upsloping,,normal,1\n720,74,Male,Switzerland,atypical
angina,145,0,,st-t abnormality,123,FALSE,1.3,upsloping,,1\n721,63,Male,VA Long
Beach,asymptomatic,140,260,FALSE,st-t
abnormality,112,TRUE,3,flat,,2\n722,44,Male,VA Long
Beach,asymptomatic,130,209,FALSE,st-t
abnormality,127,FALSE,0,,,0\n723,60,Male,VA Long
Beach,asymptomatic,132,218,FALSE,st-t
abnormality,140,TRUE,1.5,downsloping,,2\n724,55,Male,VA Long
Beach,asymptomatic,142,228,FALSE,st-t
abnormality,149,TRUE,2.5,upsloping,,1\n725,66,Male,VA Long Beach,non-
anginal,110,213,TRUE,lv hypertrophy,99,TRUE,1.3,flat,,0\n726,66,Male,VA Long
Beach,non-anginal,120,0,FALSE,st-t
abnormality,120,FALSE,-0.5,upsloping,,0\n727,65,Male,VA Long
Beach,asymptomatic,150,236,TRUE,st-t abnormality,105,TRUE,0,,,3\n728,60,Male,VA
Long Beach,non-anginal,180,0,FALSE,st-t
abnormality,140,TRUE,1.5,flat,,0\n729,60,Male,VA Long Beach,non-
anginal,120,0,,normal,141,TRUE,2,upsloping,,3\n730,60,Male,VA Long
Beach,atypical angina,160,267,TRUE,st-t
abnormality,157,FALSE,0.5,flat,,1\n731,56,Male,VA Long Beach,atypical
angina,126,166,FALSE,st-t abnormality,140,FALSE,0,,,0\n732,59,Male,VA Long
Beach,asymptomatic,140,0,FALSE,st-t
abnormality,117,TRUE,1,flat,,1\n733,62,Male,VA Long Beach,asymptomatic,110,0,FA
LSE,normal,120,TRUE,0.5,flat,,normal,1\n734,63,Male,VA Long Beach,non-
anginal,,0,FALSE,lv hypertrophy,,,,,1\n735,57,Male,VA Long
Beach,asymptomatic,128,0,TRUE,st-t
abnormality,148,TRUE,1,flat,,1\n736,62,Male,VA Long
Beach,asymptomatic,120,220,FALSE,st-t
abnormality,86,FALSE,0,,,0\n737,63,Male,VA Long Beach,asymptomatic,170,177,FALS
E,normal,84,TRUE,2.5,downsloping,,4\n738,46,Male,VA Long
Beach,asymptomatic,110,236,FALSE,normal,125,TRUE,2,flat,,1\n739,63,Male,VA Long
Beach,asymptomatic,126,0,FALSE,st-t
abnormality,120,FALSE,1.5,downsloping,,0\n740,60,Male,VA Long
Beach,asymptomatic,152,0,FALSE,st-t abnormality,118,TRUE,0,,,reversable
defect,0\n741,58,Male,VA Long
Beach,asymptomatic,116,0,FALSE,normal,124,FALSE,1,upsloping,,2\n742,64,Male,VA

Long Beach, asymptomatic, 120, 0, TRUE, st-t
 abnormality, 106, FALSE, 2, flat, , , 1\n743, 63, Male, VA Long Beach, non-
 anginal, 130, 0, FALSE, st-t abnormality, 111, TRUE, 0, , , 3\n744, 74, Male, VA Long
 Beach, non-anginal, , 0, FALSE, normal, , , , , 0\n745, 52, Male, VA Long Beach, non-
 anginal, 128, 0, FALSE, st-t abnormality, 180, FALSE, 3, upsloping, , , 2\n746, 69, Male, VA
 Long Beach, asymptomatic, 130, 0, TRUE, st-t abnormality, 129, FALSE, 1, flat, , fixed
 defect, 2\n747, 51, Male, VA Long Beach, asymptomatic, , 0, TRUE, st-t
 abnormality, , , , , 2\n748, 60, Male, VA Long Beach, asymptomatic, 130, 186, TRUE, st-t
 abnormality, 140, TRUE, 0.5, flat, , , 1\n749, 56, Male, VA Long
 Beach, asymptomatic, 120, 100, FALSE, normal, 120, TRUE, 1.5, flat, 0, reversable
 defect, 1\n750, 55, Male, VA Long Beach, non-anginal, , 228, FALSE, st-t
 abnormality, , , , , 3\n751, 54, Male, VA Long Beach, asymptomatic, , 0, FALSE, st-t
 abnormality, , , , , 3\n752, 77, Male, VA Long Beach, asymptomatic, 124, 171, FALSE, st-t
 abnormality, 110, TRUE, 2, upsloping, , , 3\n753, 63, Male, VA Long
 Beach, asymptomatic, 160, 230, TRUE, normal, 105, TRUE, 1, flat, , , 3\n754, 55, Male, VA Long
 Beach, non-anginal, 0, 0, FALSE, normal, 155, FALSE, 1.5, flat, , , 3\n755, 52, Male, VA Long
 Beach, non-anginal, 122, 0, FALSE, normal, 110, TRUE, 2, downsloping, , , 2\n756, 64, Male, VA
 Long Beach, asymptomatic, 144, 0, FALSE, st-t
 abnormality, 122, TRUE, 1, flat, , , 3\n757, 60, Male, VA Long
 Beach, asymptomatic, , 281, FALSE, st-t abnormality, , , , , 2\n758, 60, Male, VA Long
 Beach, asymptomatic, 120, 0, FALSE, normal, 133, TRUE, 2, upsloping, , reversable
 defect, 0\n759, 58, Male, VA Long
 Beach, asymptomatic, , 203, TRUE, normal, , , , , 1\n760, 59, Male, VA Long
 Beach, asymptomatic, 154, 0, FALSE, st-t
 abnormality, 131, TRUE, 1.5, , 0, , 0\n761, 61, Male, VA Long Beach, non-
 anginal, 120, 0, FALSE, normal, 80, TRUE, 0, flat, , , 3\n762, 40, Male, VA Long
 Beach, asymptomatic, 125, 0, TRUE, normal, 165, FALSE, 0, , , reversable
 defect, 1\n763, 61, Male, VA Long Beach, asymptomatic, , 0, TRUE, st-t
 abnormality, 86, FALSE, 1.5, flat, , reversable defect, 3\n764, 41, Male, VA Long
 Beach, asymptomatic, 104, 0, FALSE, st-t abnormality, 111, FALSE, 0, , , , 0\n765, 57, Male, VA
 Long Beach, asymptomatic, , 277, TRUE, st-t abnormality, , , , , 4\n766, 63, Male, VA Long
 Beach, asymptomatic, 136, 0, FALSE, normal, 84, TRUE, 0, , , reversable
 defect, 2\n767, 59, Male, VA Long Beach, asymptomatic, 122, 233, FALSE, normal, 117, TRUE, 1
 .3, downsloping, , , 1\n768, 51, Male, VA Long
 Beach, asymptomatic, 128, 0, FALSE, normal, 107, FALSE, 0, , , , 0\n769, 59, Male, VA Long
 Beach, non-anginal, , 0, FALSE, normal, 128, TRUE, 2, downsloping, , , 2\n770, 42, Male, VA
 Long Beach, non-anginal, 134, 240, , normal, 160, FALSE, 0, , , , 0\n771, 55, Male, VA Long
 Beach, non-anginal, 120, 0, FALSE, st-t abnormality, 125, TRUE, 2.5, flat, , reversable
 defect, 1\n772, 63, Female, VA Long Beach, atypical
 angina, , 0, FALSE, normal, , , , , 0\n773, 62, Male, VA Long
 Beach, asymptomatic, 152, 153, FALSE, st-t
 abnormality, 97, TRUE, 1.6, upsloping, , reversable defect, 2\n774, 56, Male, VA Long
 Beach, atypical angina, 124, 224, TRUE, normal, 161, FALSE, 2, flat, , , 0\n775, 53, Male, VA
 Long Beach, asymptomatic, 126, 0, FALSE, normal, 106, FALSE, 0, , , , 1\n776, 68, Male, VA Long
 Beach, asymptomatic, 138, 0, FALSE, normal, 130, TRUE, 3, flat, , , 2\n777, 53, Male, VA Long
 Beach, asymptomatic, 154, 0, , st-t abnormality, 140, TRUE, 1.5, flat, , , 2\n778, 60, Male, VA
 Long Beach, non-anginal, , 316, TRUE, st-t abnormality, , , , , , 3\n779, 62, Male, VA Long

Beach, atypical angina, ,0, FALSE, normal, , , , , ,0\n780, 59, Male, VA Long
 Beach, asymptomatic, 178, 0, TRUE, lv hypertrophy, 120, TRUE, 0, , , reversible
 defect, 1\n781, 51, Male, VA Long Beach, asymptomatic, , 218, TRUE, lv
 hypertrophy, , , , , ,0\n782, 61, Male, VA Long
 Beach, asymptomatic, 110, 0, , normal, 108, TRUE, 2, downsloping, , , 2\n783, 57, Male, VA Long
 Beach, asymptomatic, 130, 311, , st-t abnormality, 148, TRUE, 2, flat, , , 1\n784, 56, Male, VA
 Long Beach, non-anginal, 170, 0, FALSE, lv
 hypertrophy, 123, TRUE, 2.5, , , 4\n785, 58, Male, VA Long Beach, atypical
 angina, 126, 0, TRUE, normal, 110, TRUE, 2, flat, , , 2\n786, 69, Male, VA Long Beach, non-
 anginal, 140, 0, , st-t abnormality, 118, FALSE, 2.5, downsloping, , , 2\n787, 67, Male, VA
 Long Beach, typical
 angina, 142, 270, TRUE, normal, 125, FALSE, 2.5, upsloping, , , 3\n788, 58, Male, VA Long
 Beach, asymptomatic, 120, 0, FALSE, lv
 hypertrophy, 106, TRUE, 1.5, downsloping, , reversible defect, 1\n789, 65, Male, VA Long
 Beach, asymptomatic, , 0, FALSE, normal, , , , , , 1\n790, 63, Male, VA Long Beach, atypical
 angina, , 217, TRUE, st-t abnormality, , , , , , 1\n791, 55, Male, VA Long Beach, atypical
 angina, 110, 214, TRUE, st-t abnormality, 180, FALSE, , , , 0\n792, 57, Male, VA Long
 Beach, asymptomatic, 140, 214, FALSE, st-t abnormality, 144, TRUE, 2, flat, , fixed
 defect, 2\n793, 65, Male, VA Long Beach, typical
 angina, , 252, FALSE, normal, , , , , , 0\n794, 54, Male, VA Long
 Beach, asymptomatic, 136, 220, FALSE, normal, 140, TRUE, 3, flat, , , 3\n795, 72, Male, VA Long
 Beach, non-anginal, 120, 214, FALSE, normal, 102, TRUE, 1, flat, , , 3\n796, 75, Male, VA Long
 Beach, asymptomatic, 170, 203, TRUE, st-t abnormality, 108, FALSE, 0, , , reversible
 defect, 1\n797, 49, Male, VA Long Beach, typical angina, 130, 0, FALSE, st-t
 abnormality, 145, FALSE, 3, flat, , , 2\n798, 51, Male, VA Long Beach, non-
 anginal, , 339, FALSE, normal, , , , , , 3\n799, 60, Male, VA Long
 Beach, asymptomatic, 142, 216, FALSE, normal, 110, TRUE, 2.5, flat, , , 2\n800, 64, Female, VA
 Long Beach, asymptomatic, 142, 276, FALSE, normal, 140, TRUE, 1, flat, , reversible
 defect, 1\n801, 58, Male, VA Long Beach, asymptomatic, 132, 458, TRUE, normal, 69, FALSE, 1,
 downsloping, , , 0\n802, 61, Male, VA Long Beach, asymptomatic, 146, 241, FALSE, normal, 148
 , TRUE, 3, downsloping, , , 2\n803, 67, Male, VA Long
 Beach, asymptomatic, 160, 384, TRUE, st-t
 abnormality, 130, TRUE, 0, flat, , , 2\n804, 62, Male, VA Long
 Beach, asymptomatic, 135, 297, FALSE, normal, 130, TRUE, 1, flat, , , 2\n805, 65, Male, VA Long
 Beach, asymptomatic, 136, 248, FALSE, normal, 140, TRUE, 4, downsloping, , , 4\n806, 63, Male,
 VA Long
 Beach, asymptomatic, 130, 308, FALSE, normal, 138, TRUE, 2, flat, , , 2\n807, 69, Male, VA Long
 Beach, asymptomatic, 140, 208, FALSE, st-t
 abnormality, 140, TRUE, 2, , , , 3\n808, 51, Male, VA Long
 Beach, asymptomatic, , 227, TRUE, st-t abnormality, , , , , , 0\n809, 62, Male, VA Long Beach
 , asymptomatic, 158, 210, TRUE, normal, 112, TRUE, 3, downsloping, , , 1\n810, 55, Male, VA
 Long Beach, non-anginal, , 245, TRUE, st-t abnormality, , , , , , 1\n811, 75, Male, VA Long
 Beach, asymptomatic, 136, 225, FALSE, normal, 112, TRUE, 3, flat, , , 3\n812, 40, Male, VA Long
 Beach, non-anginal, 106, 240, FALSE, normal, 80, TRUE, 0, , , reversible
 defect, 0\n813, 67, Male, VA Long Beach, asymptomatic, 120, 0, TRUE, normal, 150, FALSE, 1.5
 , downsloping, , , 3\n814, 58, Male, VA Long
 Beach, asymptomatic, 110, 198, FALSE, normal, 110, FALSE, 0, , , , 1\n815, 60, Male, VA Long

Beach, asymptomatic, ,195,FALSE,normal,,,,,0\n816,63, Male, VA Long
 Beach, asymptomatic, 160,267,TRUE,st-t abnormality,88,TRUE,2,,,3\n817,35, Male, VA
 Long Beach, non-anginal, ,161,FALSE,st-t abnormality,,,,,0\n818,62, Male, VA Long
 Beach, typical angina, 112,258,FALSE,st-t
 abnormality,150,TRUE,,,,,1\n819,43, Male, VA Long Beach, asymptomatic, 122,0,FALSE,n
 ormal,120,FALSE,0.5,upsloping,,,1\n820,63, Male, VA Long Beach, non-
 anginal, 130,0,TRUE,st-t abnormality,160,FALSE,3,flat,,,0\n821,68, Male, VA Long
 Beach, non-anginal, 150,195,TRUE,normal,132,FALSE,0,,,fixed
 defect,1\n822,65, Male, VA Long
 Beach, asymptomatic, 150,235,FALSE,normal,120,TRUE,1.5,flat,,,3\n823,48, Male, VA
 Long Beach, non-anginal, 102,0,,st-t
 abnormality,110,TRUE,1,downsloping,,,1\n824,63, Male, VA Long
 Beach, asymptomatic, 96,305,FALSE,st-t
 abnormality,121,TRUE,1,upsloping,,,1\n825,64, Male, VA Long
 Beach, asymptomatic, 130,223,FALSE,st-t
 abnormality,128,FALSE,0.5,flat,,,0\n826,61, Male, VA Long
 Beach, asymptomatic, 120,282,FALSE,st-t abnormality,135,TRUE,4,downsloping,,fixed
 defect,3\n827,50, Male, VA Long Beach, asymptomatic, 144,349,FALSE,lv
 hypertrophy,120,TRUE,1,upsloping,,reversible defect,1\n828,59, Male, VA Long
 Beach, asymptomatic, 124,,FALSE,normal,117,TRUE,1,flat,,,1\n829,55, Male, VA Long
 Beach, asymptomatic, 150,160,FALSE,st-t
 abnormality,150,FALSE,0,,,0\n830,45, Male, VA Long Beach, non-
 anginal,,236,FALSE,normal,,,,,0\n831,65, Male, VA Long
 Beach, asymptomatic,,312,FALSE,lv hypertrophy,,,,,3\n832,61, Male, VA Long
 Beach, atypical angina,,283,FALSE,normal,,,,,0\n833,49, Male, VA Long Beach, non-
 anginal,,142,FALSE,normal,,,,,3\n834,72, Male, VA Long
 Beach, asymptomatic,,211,FALSE,normal,,,,,1\n835,50, Male, VA Long
 Beach, asymptomatic,,218,FALSE,normal,,,,,1\n836,64, Male, VA Long
 Beach, asymptomatic,,306,TRUE,st-t abnormality,,,,,3\n837,55, Male, VA Long
 Beach, asymptomatic, 116,186,TRUE,st-t
 abnormality,102,FALSE,0,,,2\n838,63, Male, VA Long
 Beach, asymptomatic, 110,252,FALSE,st-t
 abnormality,140,TRUE,2,flat,,,2\n839,59, Male, VA Long Beach, asymptomatic, 125,222,
 FALSE,normal,135,TRUE,2.5,downsloping,,,3\n840,56, Male, VA Long
 Beach, asymptomatic,,0,FALSE,lv hypertrophy,,,,,1\n841,62, Male, VA Long
 Beach, non-anginal,,0,TRUE,st-t abnormality,,,,,2\n842,74, Male, VA Long
 Beach, asymptomatic, 150,258,TRUE,st-t
 abnormality,130,TRUE,4,downsloping,,,3\n843,54, Male, VA Long
 Beach, asymptomatic, 130,202,TRUE,normal,112,TRUE,2,flat,,,1\n844,57, Male, VA Long
 Beach, asymptomatic, 110,197,FALSE,lv hypertrophy,100,FALSE,0,,,0\n845,62, Male, VA
 Long Beach, non-anginal,,204,FALSE,st-t abnormality,,,,,1\n846,76, Male, VA Long
 Beach, non-anginal, 104,,FALSE,lv
 hypertrophy,120,FALSE,3.5,downsloping,,,4\n847,54, Female, VA Long
 Beach, asymptomatic, 138,274,FALSE,normal,105,TRUE,1.5,flat,,,1\n848,70, Male, VA
 Long Beach, asymptomatic, 170,192,FALSE,st-t
 abnormality,129,TRUE,3,downsloping,,,2\n849,61, Female, VA Long Beach, atypical
 angina, 140,298,TRUE,normal,120,TRUE,0,,,reversible defect,0\n850,48, Male, VA Long

Beach, asymptomatic, ,272, FALSE, st-t abnormality, , , , , 0\n851, 48, Male, VA Long
 Beach, non-anginal, 132, 220, TRUE, st-t abnormality, 162, FALSE, 0, , , fixed
 defect, 1\n852, 61, Male, VA Long Beach, typical angina, 142, 200, TRUE, st-t
 abnormality, 100, FALSE, 1.5, downsloping, , , 3\n853, 66, Male, VA Long Beach, asymptomati
 c, 112, 261, FALSE, normal, 140, FALSE, 1.5, upsloping, , , 1\n854, 68, Male, VA Long
 Beach, typical angina, , 181, TRUE, st-t abnormality, , , , , 0\n855, 55, Male, VA Long
 Beach, asymptomatic, 172, 260, FALSE, normal, 73, FALSE, 2, , , 3\n856, 62, Male, VA Long
 Beach, non-anginal, 120, 220, FALSE, lv hypertrophy, 86, FALSE, 0, , , 0\n857, 71, Male, VA
 Long Beach, non-anginal, , 221, FALSE, normal, , , , , 3\n858, 74, Male, VA Long
 Beach, typical angina, , 216, TRUE, normal, , , , , 3\n859, 53, Male, VA Long Beach, non-
 anginal, 155, 175, TRUE, st-t abnormality, 160, FALSE, , , , fixed
 defect, 0\n860, 58, Male, VA Long Beach, non-anginal, 150, 219, FALSE, st-t
 abnormality, 118, TRUE, 0, , , 2\n861, 75, Male, VA Long
 Beach, asymptomatic, 160, 310, TRUE, normal, 112, TRUE, 2, downsloping, , reversable
 defect, 0\n862, 56, Male, VA Long Beach, non-anginal, , 208, TRUE, st-t
 abnormality, , , , , 4\n863, 58, Male, VA Long Beach, non-anginal, , 232, FALSE, st-t
 abnormality, , , , , 2\n864, 64, Male, VA Long Beach, asymptomatic, 134, 273, FALSE, normal
 , 102, TRUE, 4, downsloping, , , 4\n865, 54, Male, VA Long Beach, non-
 anginal, , 203, FALSE, st-t abnormality, , , , , 0\n866, 54, Male, VA Long Beach, atypical
 angina, , 182, FALSE, st-t abnormality, , , , , 0\n867, 59, Male, VA Long
 Beach, asymptomatic, 140, 274, FALSE, normal, 154, TRUE, 2, flat, , , 0\n868, 55, Male, VA Long
 Beach, asymptomatic, , 204, TRUE, st-t abnormality, , , , , 1\n869, 57, Male, VA Long
 Beach, asymptomatic, 144, 270, TRUE, st-t
 abnormality, 160, TRUE, 2, flat, , , 3\n870, 61, Male, VA Long
 Beach, asymptomatic, , 292, FALSE, st-t abnormality, , , , , 3\n871, 41, Male, VA Long
 Beach, asymptomatic, 150, 171, FALSE, normal, 128, TRUE, 1.5, flat, , , 0\n872, 71, Male, VA
 Long Beach, asymptomatic, 130, 221, FALSE, st-t
 abnormality, 115, TRUE, 0, , , 3\n873, 38, Male, VA Long Beach, asymptomatic, 110, 289, FALS
 E, normal, 105, TRUE, 1.5, downsloping, , , 1\n874, 55, Male, VA Long
 Beach, asymptomatic, 158, 217, FALSE, normal, 110, TRUE, 2.5, flat, , , 1\n875, 56, Male, VA
 Long Beach, asymptomatic, 128, 223, FALSE, st-t
 abnormality, 119, TRUE, 2, downsloping, , , 2\n876, 69, Male, VA Long
 Beach, asymptomatic, , TRUE, normal, , , , , 2\n877, 64, Male, VA Long
 Beach, asymptomatic, 150, 193, FALSE, st-t
 abnormality, 135, TRUE, 0.5, flat, , , 2\n878, 72, Male, VA Long
 Beach, asymptomatic, 160, , TRUE, lv hypertrophy, 130, FALSE, 1.5, , , 2\n879, 69, Male, VA
 Long Beach, asymptomatic, , 210, TRUE, st-t abnormality, , , , , 2\n880, 56, Male, VA Long
 Beach, asymptomatic, , 282, TRUE, normal, , , , , 1\n881, 62, Male, VA Long
 Beach, asymptomatic, , 170, FALSE, st-t abnormality, 120, TRUE, 3, , , 4\n882, 67, Male, VA
 Long Beach, asymptomatic, , 369, FALSE, normal, , , , , 3\n883, 57, Male, VA Long
 Beach, asymptomatic, 156, 173, FALSE, lv
 hypertrophy, 119, TRUE, 3, downsloping, , , 3\n884, 69, Male, VA Long
 Beach, asymptomatic, , 289, TRUE, st-t abnormality, , , , , 3\n885, 51, Male, VA Long
 Beach, asymptomatic, , TRUE, lv hypertrophy, , , , , reversable
 defect, 1\n886, 48, Male, VA Long
 Beach, asymptomatic, 140, , FALSE, normal, 159, TRUE, 1.5, upsloping, , , 3\n887, 69, Male, VA
 Long Beach, asymptomatic, 122, 216, TRUE, lv hypertrophy, 84, TRUE, 0, , , reversable

defect,2\n888,69,Male,VA Long Beach,non-anginal,,271,FALSE,lv
hypertrophy,,,,,0\n889,64,Male,VA Long Beach,asymptomatic,,244,TRUE,st-t
abnormality,,,,,2\n890,57,Male,VA Long Beach,atypical angina,180,285,TRUE,st-t
abnormality,120,FALSE,0.8,,,1\n891,53,Male,VA Long
Beach,asymptomatic,124,243,FALSE,normal,122,TRUE,2,flat,,reversable
defect,1\n892,37,Male,VA Long Beach,non-anginal,118,240,FALSE,lv
hypertrophy,165,FALSE,1,flat,,normal,0\n893,67,Male,VA Long
Beach,asymptomatic,140,219,FALSE,st-t abnormality,122,TRUE,2,flat,,reversable
defect,3\n894,74,Male,VA Long Beach,non-
anginal,140,237,TRUE,normal,94,FALSE,0,,,1\n895,63,Male,VA Long Beach,atypical
angina,,165,FALSE,st-t abnormality,,,,,0\n896,58,Male,VA Long
Beach,asymptomatic,100,213,FALSE,st-t
abnormality,110,FALSE,0,,,0\n897,61,Male,VA Long
Beach,asymptomatic,190,287,TRUE,lv
hypertrophy,150,TRUE,2,downsloping,,,4\n898,64,Male,VA Long
Beach,asymptomatic,130,258,TRUE,lv hypertrophy,130,FALSE,0,,,fixed
defect,2\n899,58,Male,VA Long Beach,asymptomatic,160,256,TRUE,lv
hypertrophy,113,TRUE,1,upsloping,,,3\n900,60,Male,VA Long
Beach,asymptomatic,130,186,TRUE,lv
hypertrophy,140,TRUE,0.5,flat,,,1\n901,57,Male,VA Long
Beach,asymptomatic,122,264,FALSE,lv hypertrophy,100,FALSE,0,,,1\n902,55,Male,VA
Long Beach,non-anginal,,,FALSE,st-t abnormality,,,,,0\n903,55,Male,VA Long
Beach,asymptomatic,120,226,FALSE,lv
hypertrophy,127,TRUE,1.7,downsloping,,reversable defect,1\n904,56,Male,VA Long
Beach,asymptomatic,130,203,TRUE,normal,98,FALSE,1.5,flat,,reversable
defect,1\n905,57,Male,VA Long Beach,asymptomatic,130,207,FALSE,st-t
abnormality,96,TRUE,1,flat,,,0\n906,61,Male,VA Long Beach,non-
anginal,,284,FALSE,normal,,,,,1\n907,61,Male,VA Long Beach,non-
anginal,120,337,FALSE,normal,98,TRUE,0,,,3\n908,58,Male,VA Long Beach,non-
anginal,150,219,FALSE,st-t abnormality,118,TRUE,0,,,2\n909,74,Male,VA Long Beac
h,asymptomatic,155,310,FALSE,normal,112,TRUE,1.5,downsloping,,,2\n910,68,Male,VA
Long Beach,non-
anginal,134,254,TRUE,normal,151,TRUE,0,,,normal,0\n911,51,Female,VA Long
Beach,asymptomatic,114,258,TRUE,lv
hypertrophy,96,FALSE,1,upsloping,,,0\n912,62,Male,VA Long
Beach,asymptomatic,160,254,TRUE,st-t
abnormality,108,TRUE,3,flat,,,4\n913,53,Male,VA Long
Beach,asymptomatic,144,300,TRUE,st-t
abnormality,128,TRUE,1.5,flat,,,3\n914,62,Male,VA Long
Beach,asymptomatic,158,170,FALSE,st-t
abnormality,138,TRUE,0,,,1\n915,46,Male,VA Long
Beach,asymptomatic,134,310,FALSE,normal,126,FALSE,0,,,normal,2\n916,54,Female,VA
Long Beach,asymptomatic,127,333,TRUE,st-t
abnormality,154,FALSE,0,,,1\n917,62,Male,VA Long Beach,typical
angina,,139,FALSE,st-t abnormality,,,,,0\n918,55,Male,VA Long
Beach,asymptomatic,122,223,TRUE,st-t abnormality,100,FALSE,0,,,fixed
defect,2\n919,58,Male,VA Long Beach,asymptomatic,,385,TRUE,lv

```
hypertrophy,,,,,,0\n920,62,Male,VA Long Beach,atypical angina,120,254,FALSE,lv
hypertrophy,93,TRUE,0,,,,,1\n'}
```

1.2 Reading Dataset

```
[4]: hd = pd.read_csv('/content/heart_disease_uci.csv')
hd.head()
```

```
[4]:   id  age  sex  dataset          cp  trestbps  chol  fbs  \
0   1   63  Male  Cleveland  typical angina    145.0  233.0  True
1   2   67  Male  Cleveland  asymptomatic    160.0  286.0  False
2   3   67  Male  Cleveland  asymptomatic    120.0  229.0  False
3   4   37  Male  Cleveland  non-anginal    130.0  250.0  False
4   5   41  Female  Cleveland  atypical angina    130.0  204.0  False

      restecg  thalach  exang  oldpeak    slope  ca  \
0  lv hypertrophy    150.0  False     2.3  downsloping  0.0
1  lv hypertrophy    108.0   True     1.5     flat  3.0
2  lv hypertrophy    129.0   True     2.6     flat  2.0
3           normal    187.0  False     3.5  downsloping  0.0
4  lv hypertrophy    172.0  False     1.4   upsloping  0.0

      thal  num
0     fixed defect    0
1           normal    2
2  reversable defect    1
3           normal    0
4           normal    0
```

1.3 Columns Info

- id : (Unique id for each patient)
- age : (Age of the patient in years)
- origin : (place of study)
- sex : (Male/Female)
- cp : chest pain type ([typical angina, atypical angina, non-anginal, asymptomatic])
- trestbps : resting blood pressure (resting blood pressure (in mm Hg on admission to the hospital))
- chol : (serum cholesterol in mg/dl)
- fbs : (if fasting blood sugar > 120 mg/dl)
- restecg : (resting electrocardiographic results)
- – Values: [normal, stt abnormality, lv hypertrophy]
- thalach: maximum heart rate achieved
- exang: exercise-induced angina (True/ False)
- oldpeak: ST depression induced by exercise relative to rest
- slope: the slope of the peak exercise ST segment
- ca: number of major vessels (0-3) colored by fluoroscopy

- thal: [normal; fixed defect; reversible defect]
- num: the predicted attribute

```
[5]: hd.shape
```

```
[5]: (920, 16)
```

```
[6]: hd.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 920 entries, 0 to 919
Data columns (total 16 columns):
#   Column      Non-Null Count  Dtype
---  -
0   id           920 non-null    int64
1   age          920 non-null    int64
2   sex          920 non-null    object
3   dataset      920 non-null    object
4   cp           920 non-null    object
5   trestbps     861 non-null    float64
6   chol         890 non-null    float64
7   fbs          830 non-null    object
8   restecg      918 non-null    object
9   thalch       865 non-null    float64
10  exang        865 non-null    object
11  oldpeak      858 non-null    float64
12  slope        611 non-null    object
13  ca           309 non-null    float64
14  thal         434 non-null    object
15  num          920 non-null    int64
dtypes: float64(5), int64(3), object(8)
memory usage: 115.1+ KB
```

Dropping Column ‘dataset’ as it is not required for visualization. ‘id’ and ‘dataset’ column is not important to us as ‘id’ is a unique value and ‘dataset’ column contains the place where applicant has studied. ‘ca’ column has more than 60% of Nan values if we fill that it could mislead us, so it’s better that we drop that column.

```
[7]: hd = hd.drop(['id', 'dataset', 'ca'], axis=1)
```

Renaming Columns Renaming some columns for our better understanding.

```
[8]: hd.rename(columns = {'cp': 'Chest_Pain_Type', 'trestbps':
    ↪ 'Resting_Blood_Pressure', 'chol': 'serum_cholesterol', 'fbs':
    ↪ 'Fasting_Blood_Sugar', 'restecg': 'Resting_Electrocardiographic', 'thalch':
    ↪ 'Maximum_Heart_Rate', 'exang': 'Exercise_Induced_Angina', 'oldpeak':
    ↪ 'ST_Depression_Induced', 'thal': 'Heart_Condition'}, inplace = True)
```

1.4 Columns After Renaming

- id : (Unique id for each patient)
- age : (Age of the patient in years)
- origin : (place of study)
- sex : (Male/Female)
- Chest_Pain_Type : chest pain type ([typical angina, atypical angina, non-anginal, asymptomatic])
- Resting_Blood_Pressure : resting blood pressure (resting blood pressure (in mm Hg on admission to the hospital))
- serum_cholesterol : (serum cholesterol in mg/dl)
- Fasting_Blood_Sugar : (if fasting blood sugar > 120 mg/dl)
- Resting_Electrocardiographic : (resting electrocardiographic results)
- – Values: [normal, stt abnormality, lv hypertrophy]
- Maximum_Heart_Rate: maximum heart rate achieved
- Exercise_Induced_Angina: exercise-induced angina (True/ False)
- ST_Depression_Induced: ST depression induced by exercise relative to rest
- slope: the slope of the peak exercise ST segment
- Heart_Condition: [normal; fixed defect; reversible defect]
- num: the predicted attribute

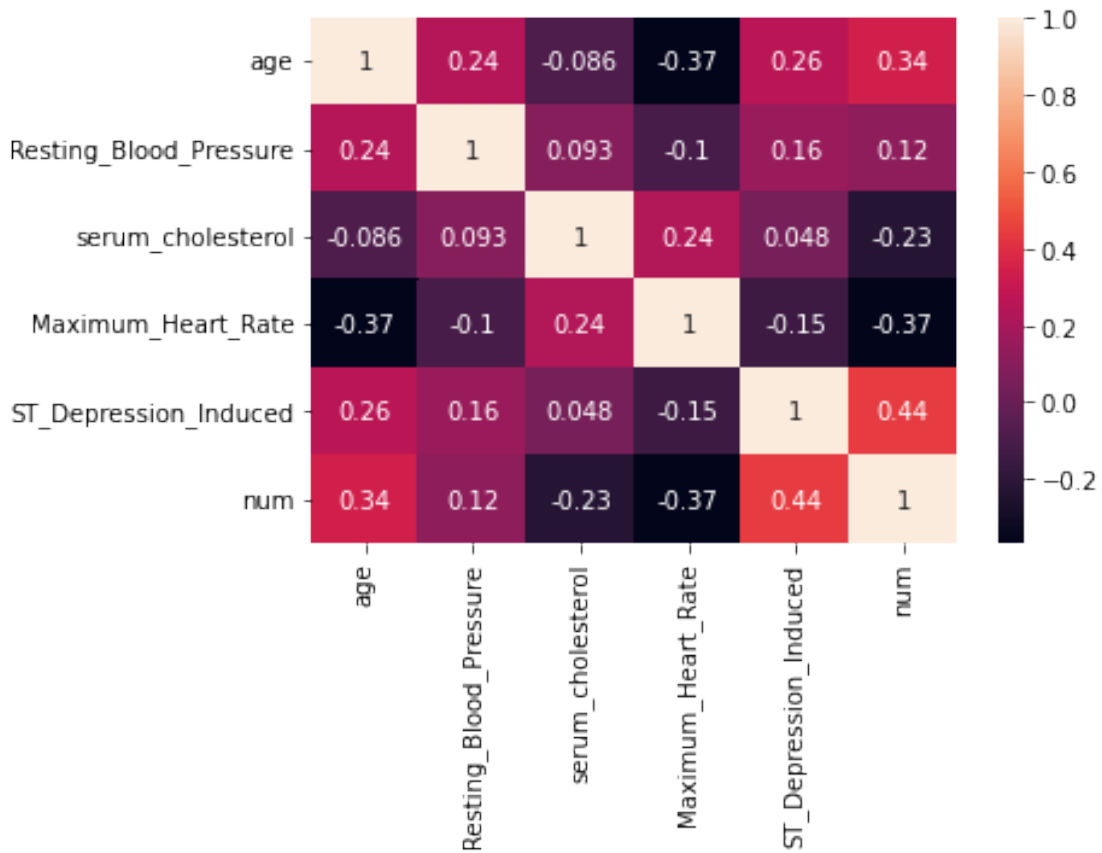
```
[9]: hd.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 920 entries, 0 to 919
Data columns (total 13 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   age                                    920 non-null    int64
1   sex                                    920 non-null    object
2   Chest_Pain_Type                       920 non-null    object
3   Resting_Blood_Pressure                 861 non-null    float64
4   serum_cholesterol                     890 non-null    float64
5   Fasting_Blood_Sugar                   830 non-null    object
6   Resting_Electrocardiographic          918 non-null    object
7   Maximum_Heart_Rate                    865 non-null    float64
8   Exercise_Induced_Angina               865 non-null    object
9   ST_Depression_Induced                  858 non-null    float64
10  slope                                  611 non-null    object
11  Heart_Condition                        434 non-null    object
12  num                                    920 non-null    int64
dtypes: float64(4), int64(2), object(7)
memory usage: 93.6+ KB
```

It shows that there are null values in the dataset. Columns ‘Vessels_Colored’ and ‘Heart_Condition’ have more than 50% null values.

```
[10]: sns.heatmap(hd.corr(),annot=True)
```

[10]: <matplotlib.axes._subplots.AxesSubplot at 0x7f70a281c670>



1.5 Cleaning Dataset

Now we need to remove Null values from the Dataset.

Columns 'Resting_Blood_Pressure', 'serum_cholesterol', 'Fasting_Blood_Sugar', 'Resting_Electrocardiographic', and 'Heart_Condition' has Nan values.

Column = 'Resting_Blood_Pressure'

```
[11]: hd['Resting_Blood_Pressure'].unique()
```

```
[11]: array([145., 160., 120., 130., 140., 172., 150., 110., 132., 117., 135.,  
112., 105., 124., 125., 142., 128., 170., 155., 104., 180., 138.,  
108., 134., 122., 115., 118., 100., 200., 94., 165., 102., 152.,  
101., 126., 174., 148., 178., 158., 192., 129., 144., 123., 136.,  
146., 106., 156., 154., 114., 164., 98., 190., nan, 113., 92.,  
95., 80., 185., 116., 0., 96., 127.]
```

```
[12]: drop_row = hd[hd['Resting_Blood_Pressure']==0].index
hd.drop(drop_row, inplace=True)
```

```
[13]: hd['Resting_Blood_Pressure'].isnull().value_counts()
```

```
[13]: False    860
      True     59
      Name: Resting_Blood_Pressure, dtype: int64
```

There are 59 Nan values, Now we will fill these by mean of the 'Resting_Blood_Pressure' column by grouping them with age.

```
[14]: mean_value1 = hd.groupby('age')['Resting_Blood_Pressure'].mean()
```

```
[15]: for i,j in zip(mean_value1.index,mean_value1):
      hd.loc[hd['age']==i,'Resting_Blood_Pressure']= hd.
      ↪loc[hd['age']==i,'Resting_Blood_Pressure'].fillna(j)
```

```
[16]: hd['Resting_Blood_Pressure'].isnull().sum()
```

```
[16]: 0
```

We have filled the Nan values with mean of grouping 'Resting_Blood_Pressure' with 'age' column. It is clean now with no Nan values. And we will do similar process for 'serum_cholesterol'.

Column = 'serum_cholesterol'

```
[17]: hd['serum_cholesterol'].unique()
```

```
[17]: array([233., 286., 229., 250., 204., 236., 268., 354., 254., 203., 192.,
        294., 256., 263., 199., 168., 239., 275., 266., 211., 283., 284.,
        224., 206., 219., 340., 226., 247., 167., 230., 335., 234., 177.,
        276., 353., 243., 225., 302., 212., 330., 175., 417., 197., 198.,
        290., 253., 172., 273., 213., 305., 216., 304., 188., 282., 185.,
        232., 326., 231., 269., 267., 248., 360., 258., 308., 245., 270.,
        208., 264., 321., 274., 325., 235., 257., 164., 141., 252., 255.,
        201., 222., 260., 182., 303., 265., 309., 307., 249., 186., 341.,
        183., 407., 217., 288., 220., 209., 227., 261., 174., 281., 221.,
        205., 240., 289., 318., 298., 564., 246., 322., 299., 300., 293.,
        277., 214., 207., 223., 160., 394., 184., 315., 409., 244., 195.,
        196., 126., 313., 259., 200., 262., 215., 228., 193., 271., 210.,
        327., 149., 295., 306., 178., 237., 218., 242., 319., 166., 180.,
        311., 278., 342., 169., 187., 157., 176., 241., 131., 132., nan,
        161., 173., 194., 297., 292., 339., 147., 291., 358., 412., 238.,
        163., 280., 202., 328., 129., 190., 179., 272., 100., 468., 320.,
        312., 171., 365., 344., 85., 347., 251., 287., 156., 117., 466.,
        338., 529., 392., 329., 355., 603., 404., 518., 285., 279., 388.,
        336., 491., 331., 393., 0., 153., 316., 458., 384., 349., 142.,
```

```
181., 310., 170., 369., 165., 337., 333., 139., 385.]])
```

```
[18]: hd['serum_cholesterol'].isnull().sum()
```

```
[18]: 30
```

```
[19]: mean_value2 = hd.groupby('age')['serum_cholesterol'].mean()
```

```
[20]: for i,j in zip(mean_value2.index,mean_value2.values):  
      hd.loc[hd['age']==i,'serum_cholesterol']= hd.  
      ↪loc[hd['age']==i,'serum_cholesterol'].fillna(j)
```

```
[21]: for i,j in zip(mean_value2.index,mean_value2.values):  
      hd.loc[(hd['age']==i)& (hd['serum_cholesterol']==0.0),'serum_cholesterol']=j
```

```
[22]: hd['serum_cholesterol'].isnull().sum()
```

```
[22]: 0
```

Column = 'Fasting_Bood_Sugar'

```
[23]: hd['Fasting_Blood_Sugar'].isnull().sum()
```

```
[23]: 90
```

```
[24]: hd['Fasting_Blood_Sugar'].unique()
```

```
[24]: array([True, False, nan], dtype=object)
```

Here we can see that data is not even as male count is way more than females, so we can't fill the Nan values on the basis of mode and as this values are categorical so mean is not possible. So we have drop these values and as there are 90 Nan values it will not affect our visualization.

```
[25]: hd.groupby('Fasting_Blood_Sugar')['sex'].value_counts()
```

```
[25]: Fasting_Blood_Sugar  sex  
False                Male    527  
                   Female   164  
True                 Male    119  
                   Female    19  
Name: sex, dtype: int64
```

Column = 'Resting_Electrocardiographic'

```
[26]: hd['Resting_Electrocardiographic'].unique()
```

```
[26]: array(['lv hypertrophy', 'normal', 'st-t abnormality', nan], dtype=object)
```

Column = 'Maximum_Heart_Rate'


```
[27]: hd['Maximum_Heart_Rate'].unique()
```

```
[27]: array([150., 108., 129., 187., 172., 178., 160., 163., 147., 155., 148.,
        153., 142., 173., 162., 174., 168., 139., 171., 144., 132., 158.,
        114., 151., 161., 179., 120., 112., 137., 157., 169., 165., 123.,
        128., 152., 140., 188., 109., 125., 131., 170., 113., 99., 177.,
        141., 180., 111., 143., 182., 156., 115., 149., 145., 146., 175.,
        186., 185., 159., 130., 190., 136., 97., 127., 154., 133., 126.,
        202., 103., 166., 164., 184., 124., 122., 96., 138., 88., 105.,
        194., 195., 106., 167., 95., 192., 117., 121., 116., 71., 118.,
        181., 134., 90., 98., 176., 135., 110., nan, 100., 87., 102.,
        92., 91., 82., 119., 94., 104., 60., 83., 63., 70., 77.,
        72., 78., 86., 93., 67., 84., 80., 107., 69., 73.]
```

There are 55 Nan values. We are going fill these with mean of these values of that age.

```
[28]: mean_value3 = hd.groupby('age')['Maximum_Heart_Rate'].mean()
```

```
[29]: for i,j in zip(mean_value3.index,mean_value3):
        hd.loc[hd['age']==i,'Maximum_Heart_Rate']= hd.
        ↪loc[hd['age']==i,'Maximum_Heart_Rate'].fillna(j)
```

```
[30]: hd['Maximum_Heart_Rate'].isnull().sum()
```

```
[30]: 0
```

Column = 'Exercise_Induced_Angina'

```
[ ]: # hd['Exercise_Induced_Angina'].unique()
```

Column = 'ST_Depression_Induced'

```
[31]: hd['ST_Depression_Induced'].unique()
```

```
[31]: array([ 2.3,  1.5,  2.6,  3.5,  1.4,  0.8,  3.6,  0.6,  3.1,  0.4,  1.3,
         0. ,  0.5,  1.6,  1. ,  1.2,  0.2,  1.8,  3.2,  2.4,  2. ,  2.5,
         2.2,  2.8,  3. ,  3.4,  6.2,  4. ,  5.6,  2.9,  0.1,  2.1,  1.9,
         4.2,  0.9,  1.1,  3.8,  0.7,  0.3,  4.4,  5. , nan, -1.1, -1.5,
        -0.1, -2.6, -0.7, -2. , -1. ,  1.7, -0.8, -0.5, -0.9,  3.7])
```

```
[32]: hd['ST_Depression_Induced'].isnull().sum()
```

```
[32]: 62
```

```
[33]: mean_value4 = hd.groupby('age')['ST_Depression_Induced'].mean()
```

```
[34]: for i,j in zip(mean_value4.index,mean_value4):
```

```
hd.loc[hd['age']==i, 'ST_Depression_Induced']= hd.  
↳loc[hd['age']==i, 'ST_Depression_Induced'].fillna(j)
```

```
[35]: hd['ST_Depression_Induced'].isnull().sum()
```

```
[35]: 0
```

Column = 'slope'

```
[36]: hd['slope'].unique()
```

```
[36]: array(['downsloping', 'flat', 'upsloping', nan], dtype=object)
```

```
[37]: hd['slope'].isnull().sum()
```

```
[37]: 309
```

```
[38]: mode_value = hd.groupby('ST_Depression_Induced')['slope'].agg(pd.Series.mode)  
mode_value
```

```
[38]: ST_Depression_Induced  
-2.6          flat  
-2.0          flat  
-1.5      downsloping  
-1.1          upsloping  
-1.0  [downsloping, upsloping]  
...  
4.2  [downsloping, flat]  
4.4      downsloping  
5.0          flat  
5.6      downsloping  
6.2      downsloping  
Name: slope, Length: 76, dtype: object
```

In some columns we can't fill values with mean or mode of that variable because at last it's a human body and every body is different from one another and these variables depend on many other aspects also, so its better to drop these rows instead of filling them.

```
[39]: hd = hd.dropna()
```

```
[40]: hd.shape
```

```
[40]: (372, 13)
```

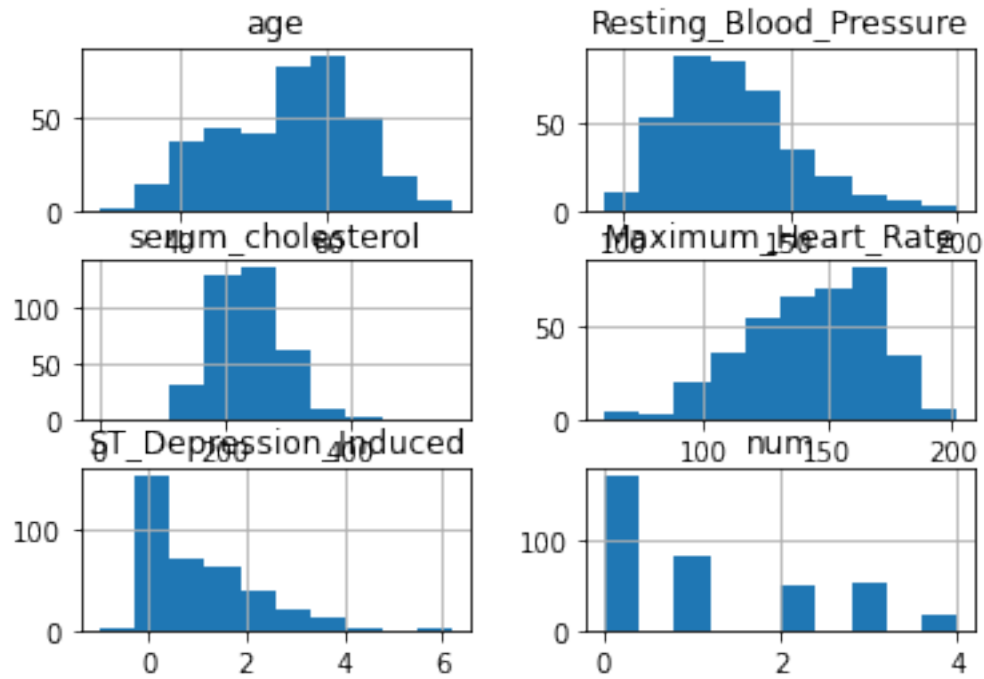
2 Data Visualization

```
[41]: hd.head()
```

```
[41]:   age      sex Chest_Pain_Type Resting_Blood_Pressure  serum_cholesterol  \  
0    63    Male  typical angina                145.0             233.0  
1    67    Male  asymptomatic                160.0             286.0  
2    67    Male  asymptomatic                120.0             229.0  
3    37    Male  non-anginal                  130.0             250.0  
4    41  Female  atypical angina                130.0             204.0  
  
      Fasting_Blood_Sugar Resting_Electrocardiographic  Maximum_Heart_Rate  \  
0                True                lv hypertrophy                150.0  
1                False                lv hypertrophy                108.0  
2                False                lv hypertrophy                129.0  
3                False                normal                    187.0  
4                False                lv hypertrophy                172.0  
  
      Exercise_Induced_Angina  ST_Depression_Induced      slope  \  
0                False                2.3  downsloping  
1                True                1.5    flat  
2                True                2.6    flat  
3                False                3.5  downsloping  
4                False                1.4  upsloping  
  
      Heart_Condition  num  
0      fixed defect    0  
1                normal  2  
2  reversable defect    1  
3                normal  0  
4                normal  0
```

```
[42]: hd.hist()
```

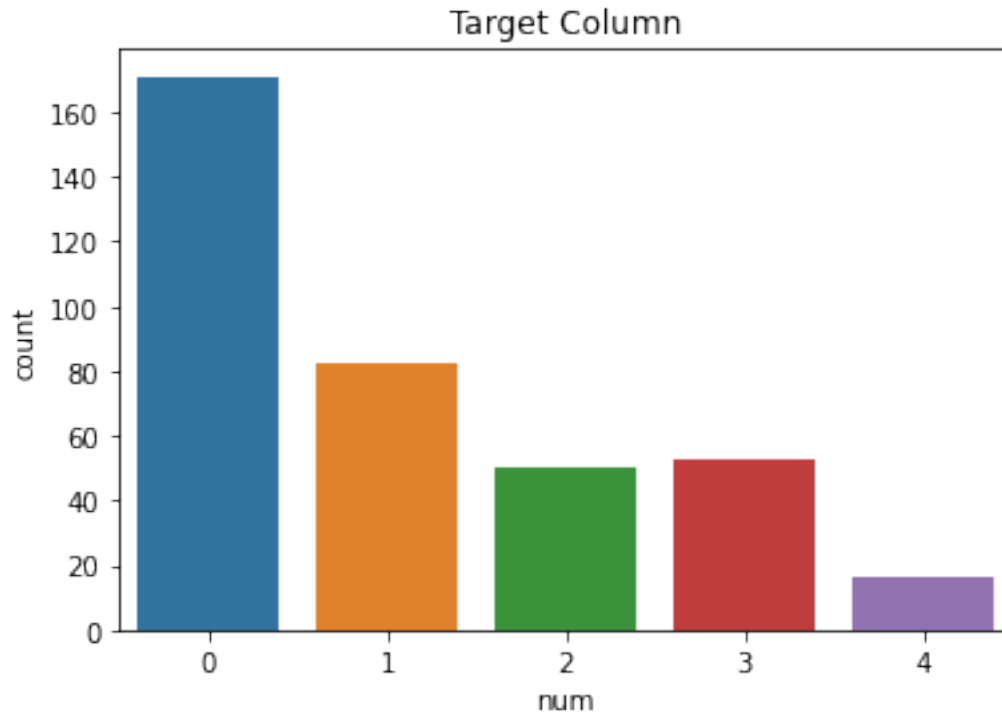
```
[42]: array([[<matplotlib.axes._subplots.AxesSubplot object at 0x7f709f9d50d0>,  
          <matplotlib.axes._subplots.AxesSubplot object at 0x7f709f985520>],  
        [<matplotlib.axes._subplots.AxesSubplot object at 0x7f709f92f940>,  
          <matplotlib.axes._subplots.AxesSubplot object at 0x7f709f95ad30>],  
        [<matplotlib.axes._subplots.AxesSubplot object at 0x7f709f911190>,  
          <matplotlib.axes._subplots.AxesSubplot object at 0x7f709f8ba4c0>]],  
      dtype=object)
```



2.0.1 Target Column

```
[43]: sns.countplot(x='num',data = hd)
plt.title('Target Column')
```

```
[43]: Text(0.5, 1.0, 'Target Column')
```



2.0.2 Independent Variable (Categorical)

```
[44]: plt.figure(1)
plt.figure(figsize=(20,10))

# Column (sex)
plt.subplot(221)
hd['sex'].value_counts().plot(kind='bar',color='crimson',rot=0)
plt.title('sex')

# Column (Chest_Pain_Type)
plt.subplot(222)
hd['Chest_Pain_Type'].value_counts().plot(kind='bar',color='green',rot=0)
plt.title('Chest_Pain_Type')

# Column (Fasting_Blood_Sugar)
plt.subplot(223)
hd['Fasting_Blood_Sugar'].value_counts().plot(kind='bar',color='orange',rot=0)
plt.title('Fasting_Blood_Sugar')

# Column (Resting_Electrocardiographic)
plt.subplot(224)
```

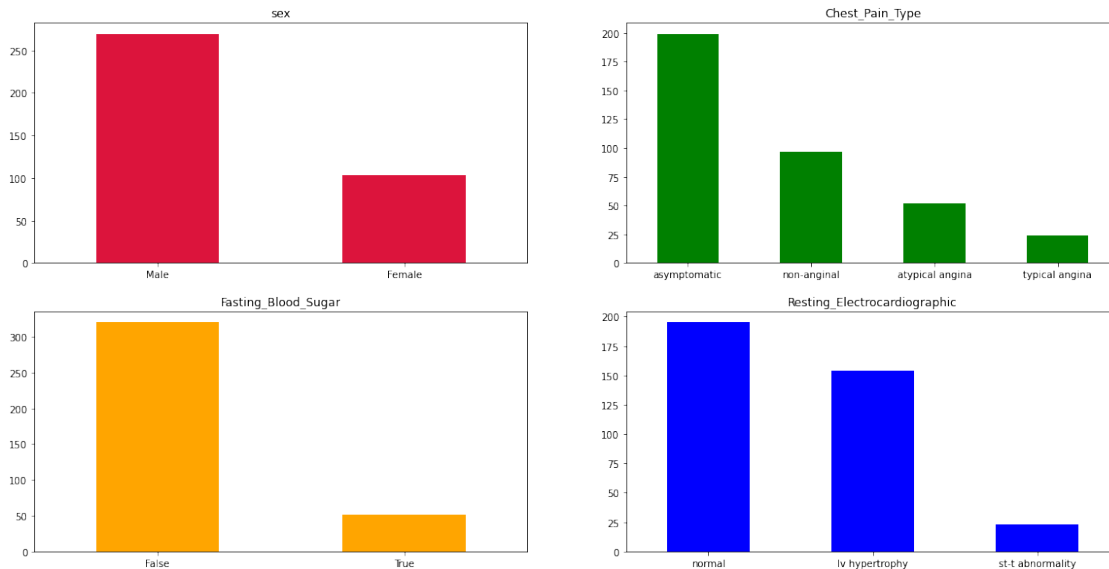
```

hd['Resting_Electrocardiographic'].value_counts().
    plot(kind='bar',color='blue',rot=0)
plt.title('Resting_Electrocardiographic')

plt.show()

```

<Figure size 432x288 with 0 Axes>



```

[45]: plt.figure(1)
plt.figure(figsize=(24,6))

# Column (Exercise_Induced_Angina)
plt.subplot(131)
hd['Exercise_Induced_Angina'].value_counts().
    plot(kind='bar',color='crimson',rot=0)
plt.title('Exercise_Induced_Angina')

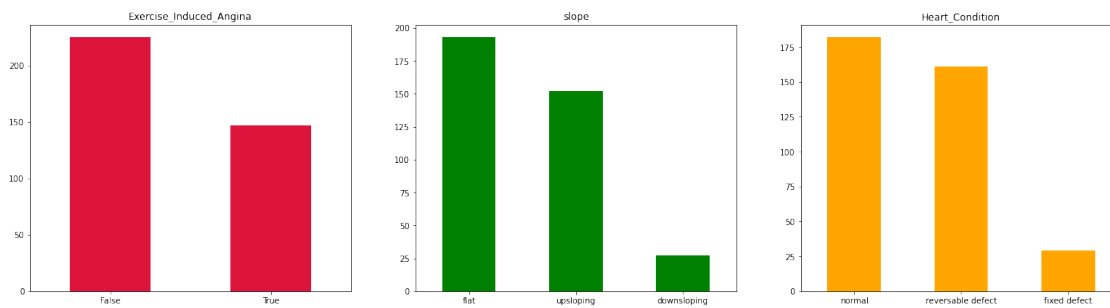
# Column (slope)
plt.subplot(132)
hd['slope'].value_counts().plot(kind='bar',color='green',rot=0)
plt.title('slope')

# Column (Heart_Condition)
plt.subplot(133)
hd['Heart_Condition'].value_counts().plot(kind='bar',color='orange',rot=0)
plt.title('Heart_Condition')

plt.show()

```

<Figure size 432x288 with 0 Axes>



2.0.3 Independent Variable (Continuous)

```
[46]: plt.figure(1)
plt.figure(figsize=(20,10))
plt.subplots_adjust(left=0.1,bottom=0.1,right=0.9,top=0.9,wspace=0.2,hspace=0.4)

# Column (age)
plt.subplot(321)
sns.distplot(hd['age'])
plt.title('age')

# Column (Resting_Blood_Pressure)
plt.subplot(322)
sns.distplot(hd['Resting_Blood_Pressure'])
plt.title('Resting_Blood_Pressure')

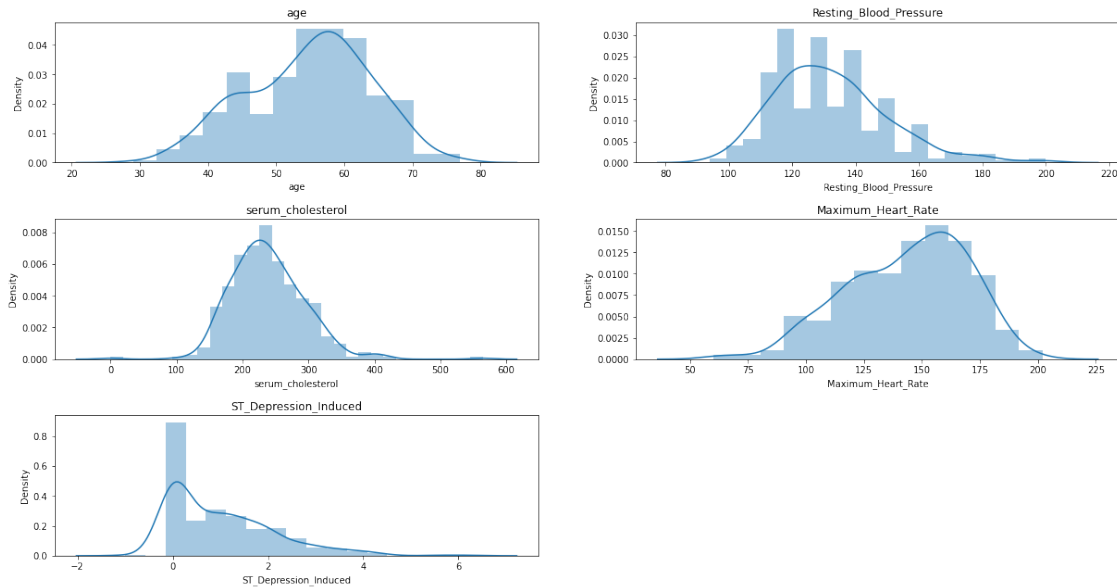
# Column (serum_cholesterol)
plt.subplot(323)
sns.distplot(hd['serum_cholesterol'])
plt.title('serum_cholesterol')

# Column (Maximum_Heart_Rate)
plt.subplot(324)
sns.distplot(hd['Maximum_Heart_Rate'])
plt.title('Maximum_Heart_Rate')

# Column (Maximum_Heart_Rate)
plt.subplot(325)
sns.distplot(hd['ST_Depression_Induced'])
plt.title('ST_Depression_Induced')

plt.show()
```

<Figure size 432x288 with 0 Axes>



2.0.4 Target Variable vs Independent Variable (Categorical)

```
[47]: plt.figure(1)
plt.figure(figsize = (20, 12))
plt.subplots_adjust(left=0.1,bottom=0.1,right=0.9,top=0.9,wspace=0.2,hspace=0.4)

# Column (num vs sex)
plt.subplot(221)
sns.barplot(x = hd['sex'], y = hd['num'], palette = "nipy_spectral")
plt.title('sex vs num')

# Column (num vs Chest_Pain_Type)
plt.subplot(222)
sns.barplot(x = hd['Chest_Pain_Type'], y = hd['num'], palette = "nipy_spectral")
plt.title('Chest_Pain_Type vs num')

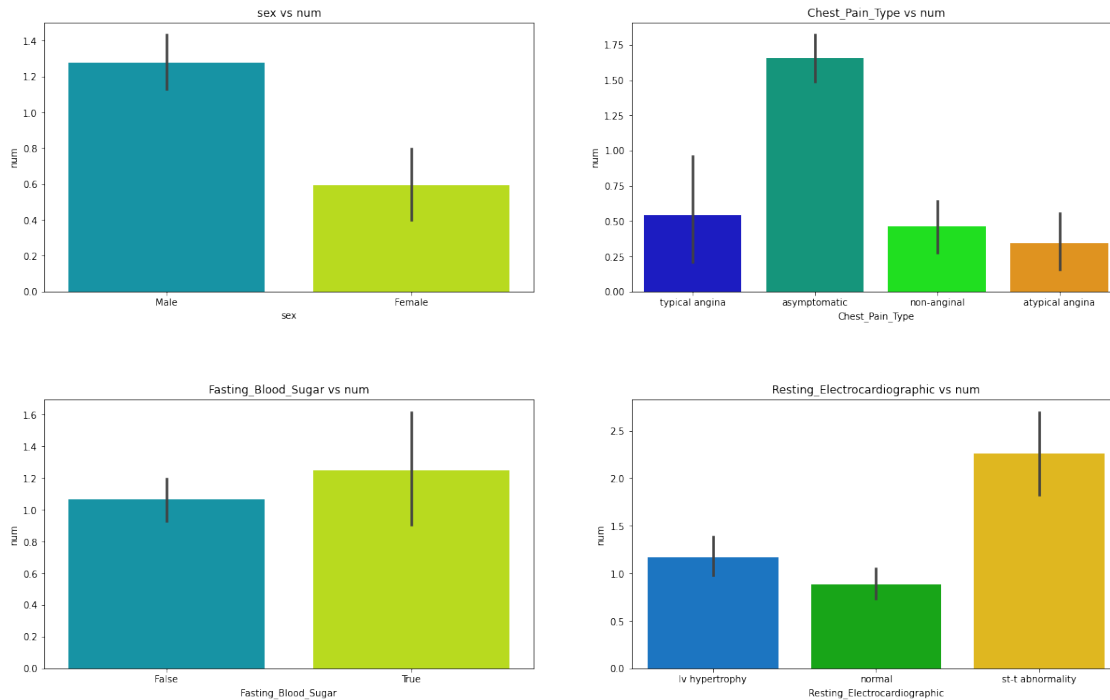
# Column (num vs Fasting_Blood_Sugar)
plt.subplot(223)
sns.barplot(x = hd['Fasting_Blood_Sugar'], y = hd['num'], palette = "nipy_spectral")
plt.title('Fasting_Blood_Sugar vs num')

# Column (num vs Resting_Electrocardiographic)
plt.subplot(224)
sns.barplot(x = hd['Resting_Electrocardiographic'], y = hd['num'], palette = "nipy_spectral")
plt.title('Resting_Electrocardiographic vs num')
```



```
plt.show()
```

<Figure size 432x288 with 0 Axes>



```
[48]: plt.figure(1)
plt.figure(figsize = (24, 6))

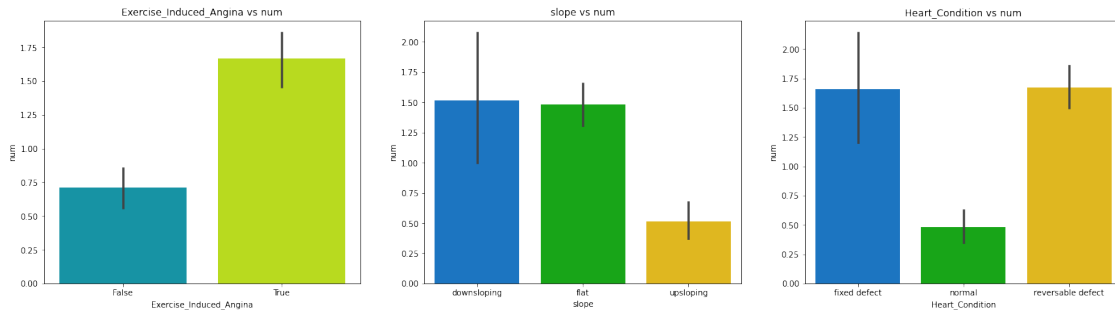
# Column (num vs Exercise_Induced_Angina)
plt.subplot(131)
sns.barplot(x = hd['Exercise_Induced_Angina'], y = hd['num'], palette = "nipy_spectral")
plt.title('Exercise_Induced_Angina vs num')

# Column (num vs slope)
plt.subplot(132)
sns.barplot(x = hd['slope'], y = hd['num'], palette = "nipy_spectral")
plt.title('slope vs num')

# Column (num vs Heart_Condition)
plt.subplot(133)
sns.barplot(x = hd['Heart_Condition'], y = hd['num'], palette = "nipy_spectral")
plt.title('Heart_Condition vs num')
```

```
plt.show()
```

<Figure size 432x288 with 0 Axes>



2.0.5 Target Variable vs Independent Variable (Continuous)

```
[49]: plt.figure(1)
plt.figure(figsize = (20, 10))

# Column (age vs num)
plt.subplot(321)
plt.ticklabel_format(style = 'plain')
plt.scatter(hd['age'], hd['num'])
plt.xlabel('age')
plt.ylabel('num')

# Column (Resting_Blood_Pressure vs num)
plt.subplot(322)
plt.ticklabel_format(style = 'plain')
plt.scatter(hd['Resting_Blood_Pressure'], hd['num'])
plt.xlabel('Resting_Blood_Pressure')
plt.ylabel('num')

# Column (serum_cholesterol vs num)
plt.subplot(323)
plt.ticklabel_format(style = 'plain')
plt.scatter(hd['serum_cholesterol'], hd['num'])
plt.xlabel('serum_cholesterol')
plt.ylabel('num')

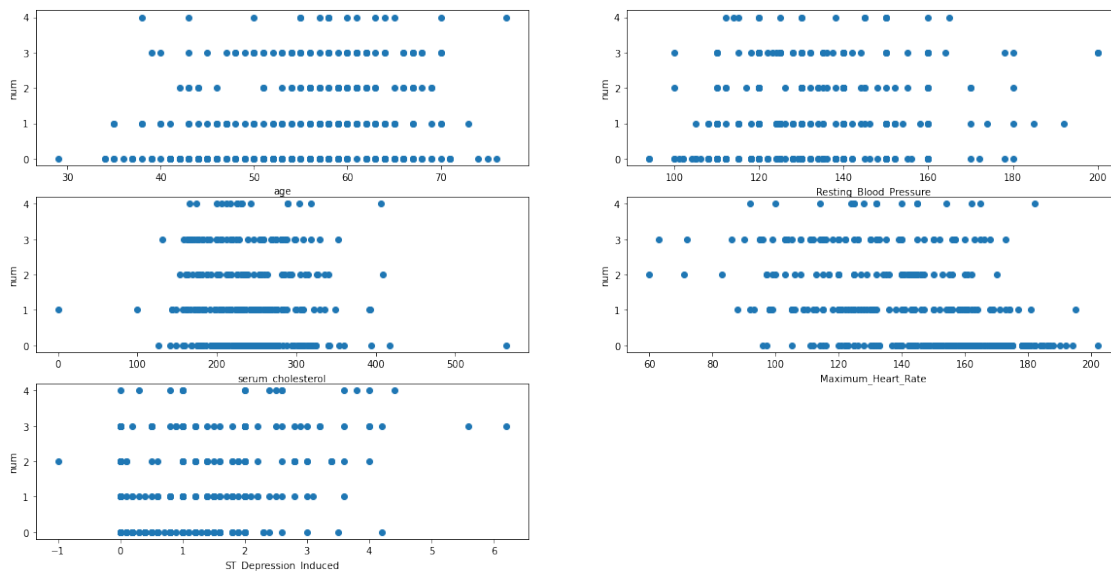
# Column (Maximum_Heart_Rate vs num)
plt.subplot(324)
plt.ticklabel_format(style = 'plain')
plt.scatter(hd['Maximum_Heart_Rate'], hd['num'])
plt.xlabel('Maximum_Heart_Rate')
```

```
plt.ylabel('num')

# Column (ST_Depression_Induced vs num)
plt.subplot(325)
plt.ticklabel_format(style = 'plain')
plt.scatter(hd['ST_Depression_Induced'], hd['num'])
plt.xlabel('ST_Depression_Induced')
plt.ylabel('num')
```

[49]: Text(0, 0.5, 'num')

<Figure size 432x288 with 0 Axes>



3 ML Models

```
[50]: def replace_fun(column):
    l = hd[column].unique()
    for i,j in enumerate(l):
        hd[column] = hd[column].replace(j,i)

    return hd[column]
```

```
[51]: replace_fun('sex')
replace_fun('Chest_Pain_Type')
replace_fun('Fasting_Blood_Sugar')
replace_fun('Resting_Electrocardiographic')
replace_fun('Exercise_Induced_Angina')
```

```
replace_fun('slope')
replace_fun('Heart_Condition')
hd.head()
```

```
[51]:
```

	age	sex	Chest_Pain_Type	Resting_Blood_Pressure	serum_cholesterol	\
0	63	0	0	145.0	233.0	
1	67	0	1	160.0	286.0	
2	67	0	1	120.0	229.0	
3	37	0	2	130.0	250.0	
4	41	1	3	130.0	204.0	

	Fasting_Blood_Sugar	Resting_Electrocardiographic	Maximum_Heart_Rate	\
0	1	0	150.0	
1	1	0	108.0	
2	1	0	129.0	
3	1	1	187.0	
4	1	0	172.0	

	Exercise_Induced_Angina	ST_Depression_Induced	slope	Heart_Condition	num
0	0	2.3	0	0	0
1	1	1.5	1	1	2
2	1	2.6	1	2	1
3	0	3.5	0	1	0
4	0	1.4	2	1	0

```
[52]: X = hd.drop('num', axis=1)
      Y = hd['num']
```

```
[53]: X_train, X_test, Y_train, Y_test = train_test_split(X,Y)
```

3.0.1 KNN Model

```
[54]: gs_knn = {'n_neighbors': [5,21,31], 'weights': ['uniform', 'distance'], 'p':
             ↪ [1,2], 'leaf_size': [10,15,40]}
```

```
[56]: knn = GridSearchCV(KNeighborsClassifier(), param_grid=gs_knn, cv=15)
      knn.fit(X_train,Y_train)
```

```
[56]: GridSearchCV(cv=15, estimator=KNeighborsClassifier(),
                  param_grid={'leaf_size': [10, 15, 40], 'n_neighbors': [5, 21, 31],
                              'p': [1, 2], 'weights': ['uniform', 'distance']})
```

```
[57]: knn.best_score_, knn.best_estimator_
```

```
[57]: (0.5046783625730994,
      KNeighborsClassifier(leaf_size=10, n_neighbors=21, p=1, weights='distance'))
```

```
[58]: knn.score(X_train, Y_train),knn.score(X_test, Y_test)
```

```
[58]: (1.0, 0.5053763440860215)
```

3.0.2 SVM Model

We can use Label Encoding but i have just tried by making a function above, so will continue with that.

```
[ ]: # X_1 = X1.copy()
# for col in X1.columns:
#     lb = LabelEncoder()
#     X_1[col] = lb.fit_transform(X1[col].values)

# X_1.head()
```

```
[59]: svm = SVC(kernel = 'poly', C =1500)
svm.fit(X_train, Y_train)
```

```
[59]: SVC(C=1500, kernel='poly')
```

```
[60]: svm.score(X_train,Y_train), svm.score(X_test, Y_test)
```

```
[60]: (0.6415770609318996, 0.5161290322580645)
```

3.0.3 Naive Bayes Model

```
[61]: nb = GaussianNB()
nb.fit(X_train,Y_train)
```

```
[61]: GaussianNB()
```

```
[62]: nb.score(X_train, Y_train), nb.score(X_test, Y_test)
```

```
[62]: (0.5663082437275986, 0.4838709677419355)
```

3.0.4 Decision Tree with Grid Search CV

```
[63]: dt = DecisionTreeClassifier()
dt.fit(X_train, Y_train)
```

```
[63]: DecisionTreeClassifier()
```

```
[64]: param_grid = {"max_depth": [5, 8, None],
                  "max_features": [5, 8, 12],
                  "min_samples_split": [4, 7, 10],
                  "criterion": ["gini", "entropy"]}
```

```
[65]: gs = GridSearchCV(dt, param_grid=param_grid, cv=20)
      gs.fit(X, Y)
```

```
[65]: GridSearchCV(cv=20, estimator=DecisionTreeClassifier(),
                  param_grid={'criterion': ['gini', 'entropy'],
                              'max_depth': [5, 8, None], 'max_features': [5, 8, 12],
                              'min_samples_split': [4, 7, 10]})
```

```
[66]: gs.score(X_train, Y_train), gs.score(X_test, Y_test)
```

```
[66]: (0.6379928315412187, 0.5913978494623656)
```

3.0.5 Random Forest

```
[67]: rfc = RandomForestClassifier()
      rfc.fit(X_train, Y_train)
```

```
[67]: RandomForestClassifier()
```

```
[68]: rfc.score(X_train, Y_train), rfc.score(X_test, Y_test)
```

```
[68]: (1.0, 0.4838709677419355)
```

```
[69]: rfc.feature_importances_
```

```
[69]: array([0.11766391, 0.02921877, 0.06966261, 0.12636389, 0.13877517,
          0.          , 0.04814355, 0.18848835, 0.04584362, 0.11553022,
          0.04465302, 0.07565688])
```

```
[70]: feature_importance = pd.DataFrame({'importance': rfc.feature_importances_},
      ↪ index= X.columns).sort_values('importance')
      feature_importance
```

```
[70]:
```

	importance
Fasting_Blood_Sugar	0.000000
sex	0.029219
slope	0.044653
Exercise_Induced_Angina	0.045844
Resting_Electrocardiographic	0.048144
Chest_Pain_Type	0.069663
Heart_Condition	0.075657
ST_Depression_Induced	0.115530
age	0.117664
Resting_Blood_Pressure	0.126364
serum_cholesterol	0.138775
Maximum_Heart_Rate	0.188488

```
[71]: feature_importance[feature_importance.importance > 0.06]
```

```
[71]:
```

	importance
Chest_Pain_Type	0.069663
Heart_Condition	0.075657
ST_Depression_Induced	0.115530
age	0.117664
Resting_Blood_Pressure	0.126364
serum_cholesterol	0.138775
Maximum_Heart_Rate	0.188488

3.0.6 AdaBoost

```
[72]: ada = AdaBoostClassifier()  
ada.fit(X, Y)
```

```
[72]: AdaBoostClassifier()
```

```
[73]: ada.score(X_train, Y_train),ada.score(X_test, Y_test)
```

```
[73]: (0.5698924731182796, 0.5591397849462365)
```

```
[ ]:
```