

SWE485: Selected Topics in Software Engineering Software Engineering Department King Saud University 3rd term 1444

Heart Attack Analysis & Prediction using Machine Learning Algorithms



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Phase 1:

1. Introduction

The dataset we have chosen is a heart attack analysis & prediction dataset, We have chosen this dataset since the correct prediction of heart attacks can prevent life threats, and incorrect prediction can prove to be fatal at the same time.

2. The goal of Choosing the dataset

The dataset provides a list of values such as: age, sex, blood pressure, cholesterol level, chest pain and some other attributes. The goal of choosing this dataset is to predict the chance of heart attack by analyzing the relationship between the patient attributes and the target variable, which is binary outcome, so: 0 = less chance of heart attack and 1 = less chance of heart attack by applying machine learning techniques.

3. Machine learning Tasks

Since the class label in the dataset "output" is known, therefore our problem is a supervised machine learning problem. And since some values of the class label are binary values (zero or one), therefore, our problem is a classification problem because the problem requires predicting a target. For that, we will use a supervised machine learning classification algorithm to predict whether it has a chance of a heart attack or not based on the values of some attributes.

Supervised learning

To predict whether the there is a chance of heart attack or not, we will use the following machine learning algorithms:

- Logistic Regression algorithm
- Decision Tree algorithm

4. Data

a. Kind of data:

 Heart Attack Analysis & Prediction Dataset contains information indicate if the person has more chance of heart attack compared with normal person.

b. Data source:

• We got the dataset from Kaggle. Dataset URL: https://www.kaggle.com/datasets/rashikrahmanpritom/heart-attack-analysis-prediction-dataset

- c. Data exploration:
- 1) Number of observations: our data set contains 303 rows and 14 columns
- 2) Describe the meaning of each variable
 - a) Age: Age of the patient
 - b) Sex : Sex of the patient (1 = male; 0 = female)
 - c) exang: exercise induced angina (1 = yes; 0 = no)
 - d) caa: number of major vessels (0-3)
 - e) cp: Chest Pain type chest pain type
 - i) Value 1: typical angina
 - ii) Value 2: atypical angina
 - iii) Value 3: non-anginal pain
 - iv) Value 4: asymptomatic
 - f) trtbps: resting blood pressure (in mm Hg)
 - g) chol: cholesterol in mg/dl fetched via BMI sensor
 - h) fbs: (fasting blood sugar > 120 mg/dl) (1 = true; 0 = false)
 - i) rest_ecg : resting electrocardiographic results
 - i) Value 0: normal.
 - ii) Value 1: having ST-T wave abnormality (T wave inversions and/or ST elevation or depression of > 0.05 mV).
 - iii) Value 2: showing probable or definite left ventricular hypertrophy by Estes' criteria.
 - j) thalach: maximum heart rate achieved
 - k) old peak: ST depression induced by exercise relative to rest
 - thall: thalassemia which is an inherited blood disorder that causes your body to have less hemoglobin than normal.
 - i) 0 = null
 - ii) 1 =fixed defect
 - iii) 2 = normal
 - iv) 3 = reversable defect
 - m) output: 0= less chance of heart attack 1= more chance of heart attack

3) Number of variables and data types:

By using **dtypes** which we can find it in **panda** library we can see the data type of each variable in dataset.

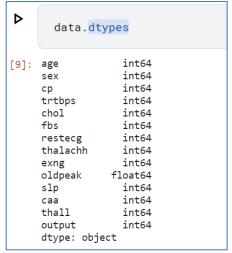


Figure 1: Data type of variable in dataset

d. Sample of raw dataset:

Here we extract some sample of the dataset by using **head()** function in **panda** library



Figure 2: some rows of dataset

e. Variables distribution:

i. Distribution plot

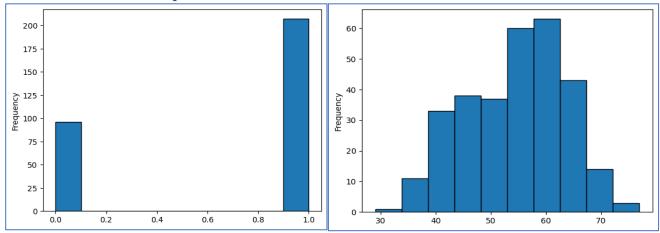


Figure 3: Sex Variable Distribution Plot

Figure 4: Age Variable Distribution plot

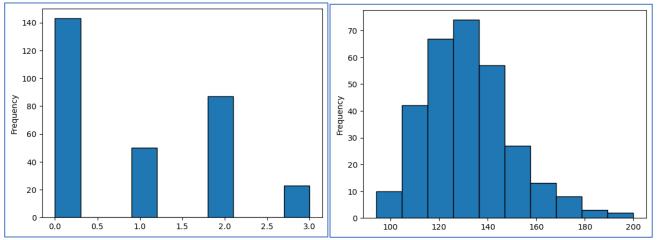


Figure 5: Cp Variable Distribution Plot

Figure 6: Trtbps Variable Distribution plot

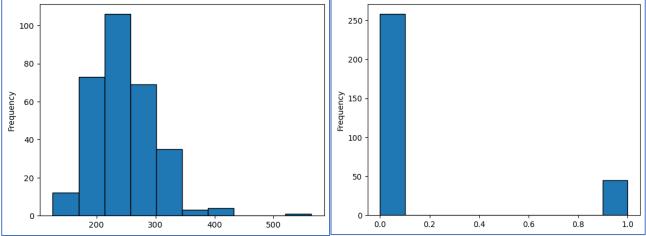
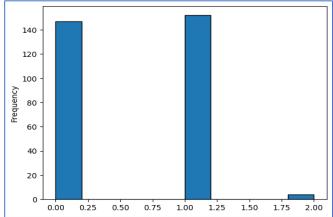


Figure 7: Chol Variable Distribution Plot

Figure 8: Fbs Variable Distribution Plot



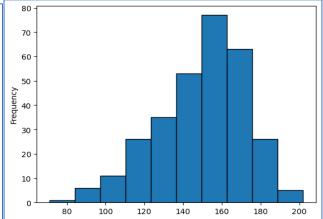


Figure 9: restecg Variable Distribution Plot

Figure 10: thalachh Variable Distribution Plot

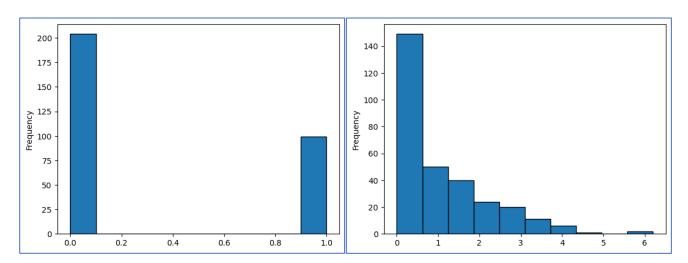


Figure 11: exng Variable Distribution Plot

Figure 12:old peak Variable Distribution Plot

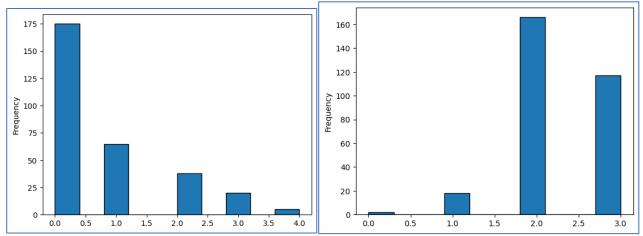


Figure 13: Caa Variable Distribution Plot

Figure 14: thall Variable Distribution Plot

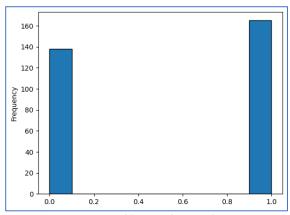


Figure 15: Output Variable Distribution Plot

ii. Pie chart of output types:

*output: 0= less chance of heart attack 1= more chance of heart attack

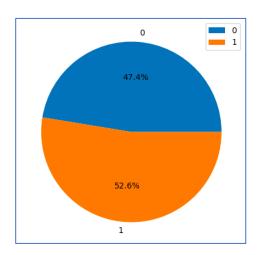


Figure 16: output pie chart

f. Missing values:

We check if the dataset have null or missing values by using isnull() function which is returns the number of missing values in the dataset.

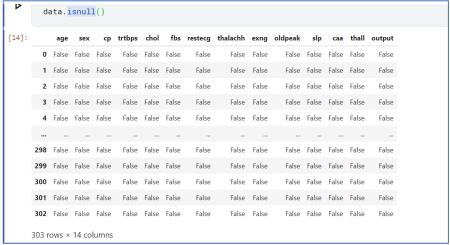


Figure 17: missing values

g. Statistical summaries:

Here we can see the statistical summaries we use describe function is used to get a descriptive statistics summary. This includes mean, count, std deviation, percentiles, and minmax values of all the features. We discover that mean of age is ~ 54 year old.

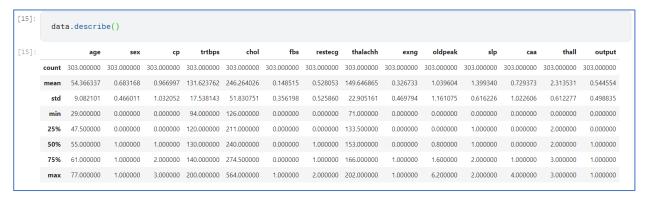


Figure 18: statistical summaries of all except variation

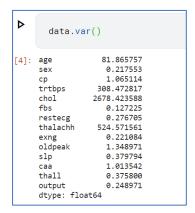


Figure 19: statistical summaries of variation

5. Data preprocessing

We deeply check our dataset to decide what techniques we need to apply. Because all variables in our data are numeral, we didn't need to do the variable transformation. Also, because our data was already classified into categorical attributes, we didn't need to do the discretization. Moreover. Because most of the variables in our data are of type integer, we didn't need to do the normalization.

Data cleaning:

The dataset didn't contain a null value but there is one duplicate in row 164 so we removed it.

```
D
      duplic= data.duplicated()
      print(data[duplic])
                     trtbps
                            chol
                                  fbs restecg thalachh exng
                                                             oldpeak
                                                                      slp
                  2
                        138
                             175
                                                   173
                                                                 0.0
             thall
      + Code
                  + Markdown
```

Figure 20: Check the duplicate in dataset

We use this code to remove duplicate row.

```
data= data.drop_duplicates()
```

Figure 21: remove duplicate row code

And this our data after remove row 164.

```
restecg
                                                thalachh
                                                                   oldpeak
                                                                                                 output
37
                   130
                         250
                                                      187
41
                         204
                  130
                                                      172
                         354
                                                      163
                   140
                         192
                                                      148
                   140
                          294
                                                      153
                   172
                         199
                                                      162
                   150
                         168
                                                      174
                   140
                          239
                                                      160
                   130
                  130
110
                          266
                                                      171
                          211
                                                      144
                          283
                                                      162
                          219
                  120
150
                                                      172
                          226
                                                      114
                                                      171
                                                      151
                   135
                          234
                                                      161
                   130
                          233
                                                      179
```

26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52	59 51 65 53 41 54 54 54 54 55 51 48 45 53 39 52 44 47 53 53 53 53 53 53 53 53 53 53 54 54 54 55 55 56 56 57 57 57 57 57 57 57 57 57 57 57 57 57	1101110011000111000111111000011	2 2 2 2 2 1 0 1 2 3 2 2 2 2 2 1 0 0 2 1 2 2 2 0 2 0 2	150 110 140 130 105 120 130 125 142 135 150 160 140 130 140 140 120 148 138 138 138 120 130	212 175 417 197 198 177 219 273 321 317 304 245 269 360 362 325 235 225 235 257 216 234 256 234 256 234 256 234 256 257 257 257 257 257 257 257 257 257 257	101100000000000000000000000000000000000	110011000010000000000000000000000000000	157 123 157 152 168 140 188 152 125 160 170 165 148 151 142 180 148 143 182 172 180 156 115 160 170 170 170 180 170 180 170 180 170 180 170 170 180 170 180 180 180 180 180 180 180 180 180 18	000000110000010000000000000000000000000	1.6 0.6 0.8 1.2 0.0 0.4 0.5 1.4 0.0 1.6 0.8 0.8 1.5 0.2 0.0 0.4 0.0 0.8 0.8 0.8 0.9 0.9 0.8 0.8 0.9 0.9 0.9 0.9 0.9 0.9 0.9 0.9 0.9 0.9	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	0 0 0 1 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 70 71 72 73 74 75 77 78 80 81 82 83 84 85 88 90 91 92 93 94 95 96 96 97 98 99 99 99 99 99 99 99 99 99 99 99 99	44 63 52 48 45 37 77 54 54 55 45 45 45 45 45 45 45 45 45 45	0 0 1 1 1 1 1 0 0 1 0 1 1 1 1 1 0 0 0 1 1 1 1 1 0 0 1 0 0 1 1 1 1 0 0 1 0 1 1 1 1 1 0 0 1 0 1 1 1 1 1 0 0 1 0 1 1 1 1 1 0 0 1 0 1 1 1 1 1 1 0 1 1 1 0 1	221003021312021102210212112 212302212020211000223312223	108 135 134 122 115 118 128 110 108 118 138 140 130 120 141 120 142 135 140 122 135 140 122 135 140 122 135 140 122 135 140 122 135 140 128 105 112 128 102 115 118 101 110 100 124 130 138 131 112 142 140 128 130 130 148 130 130 148 178 140 129 120 160	141 252 201 182 260 182 265 309 211 183 220 222 234 220 258 227 261 213 250 261 213 250 265 27 214 245 221 245 221 245 221 245 221 245 227 240 255 265 27 27 28 28 29 29 20 20 20 20 20 20 20 20 20 20 20 20 20	000000100000000000000000000000000000000	101000001011011011001001100111001110011100111000111000	175 172 158 186 185 174 159 130 156 190 132 165 182 165 182 143 175 170 163 147 154 202 186 165 164 184 154 179 170 160 178 182 160 178 122 160 151 156 158 122 175 168 169 151 157 147 162 173 178 147 162 173 178 145 179 194	0000000000010000101000101 00000000000010101000000	0.6 0.8 0.0 0.0 0.0 0.0 0.0 0.0 0.0	12222222112211221122111221 2221111221 222111222122221212222020211	001000110000000000000000000000000000000	2 2 2 2 2 2 2 3 1 1 2 2 2 2 2 2 3 3 2 2 2 2	

107	45	0	0	138	236	0	0	152	1	0.2	1	0	2	1
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111	57	1	2	150	126	1	1	173	0	0.2	2	1	3	1
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									1					
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														1
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173		1		132	224	0	0	173	0					
174	60	1	0	130	206	0	0	132	1	2.4	1	2	3	0
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178 179			_		353	0	1	132	1	1.2	1	1	3	0
179	57		a			0			9					0
179 180	57 55	1	0	132		0							2	^
179 180 181	57 55 65	1 0	0	150	225	0	0	114		1.0	1	3	3	0
179 180 181 182	57 55 65 61	1 0 0	0 0	150 130	225 330	0	0	169	0	0.0	2	0	2	0
179 180 181	57 55 65	1 0	0	150	225									
179 180 181 182	57 55 65 61	1 0 0	0 0	150 130	225 330	0	0	169	0	0.0	2	0	2	0
179 180 181 182 183	57 55 65 61 58	1 0 0 1	0 0 2	150 130 112	225 330 230	0 0	0 0	169 165	0 0	0.0 2.5	2 1	0 1	2	0
179 180 181 182 183 184 185	57 55 65 61 58 50 44	1 0 0 1 1	0 0 2 0	150 130 112 150 112	225 330 230 243 290	0 0 0	0 0 0	169 165 128 153	0 0 0	0.0 2.5 2.6 0.0	2 1 1 2	0 1 0 1	2 3 3 2	0 0 0
179 180 181 182 183 184 185 186	57 55 65 61 58 50 44 60	1 0 0 1 1 1	0 2 0 0	150 130 112 150 112 130	225 330 230 243 290 253	0 0 0 0	0 0 0 0	169 165 128 153 144	0 0 0 0	0.0 2.5 2.6 0.0 1.4	2 1 1 2 2	0 1 0 1	2 3 3 2 3	0 0 0 0
179 180 181 182 183 184 185	57 55 65 61 58 50 44	1 0 0 1 1	0 0 2 0	150 130 112 150 112	225 330 230 243 290	0 0 0	0 0 0	169 165 128 153	0 0 0	0.0 2.5 2.6 0.0	2 1 1 2	0 1 0 1	2 3 3 2	0 0 0

189	41	1	0	110	172	0	0	158	0	0.0	2	0	3	0
		1	0											
190	51	0	0	130	305	0	1	142	1	1.2	1	0	3	0
191	58	1	0	128	216	0	0	131	1	2.2	1	3	3	0
	54													
192		1	0	120	188	0	1	113	0	1.4	1	1	3	0
193	60	1	0	145	282	0	0	142	1	2.8	1	2	3	0
194	60	1	2	140	185	0	0	155	0	3.0	1	0	2	0
195	59	1	0	170	326	0	0	140	1	3.4	0	0	3	0
196	46	1	2	150	231	0	1	147	0	3.6	1	0	2	0
197	67	1	0	125	254	1	1	163	0	0.2	1	2	3	0
198	62	1	0	120	267	0	1	99	1	1.8	1	2	3	0
199	65	1	0	110	248	0	0	158	0	0.6	2	2	1	0
200	44	1	0	110	197	0	0	177	0	0.0	2	1	2	0
201	60	1	0	125	258	0	0	141	1	2.8	1	1	3	0
202	58	1	0	150	270	0	0	111	1	0.8	2	0	3	0
203	68	1	2	180	274	1	0	150	1	1.6	1	0	3	0
204	62	0	0	160	164	0	0	145	0	6.2	0	3	3	0
			0										3	
205	52	1		128	255	0	1	161	1	0.0	2	1		0
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215	43	0	0	132	341	1	0	136	1	3.0	1	0	3	0
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216	62	0	2	130	263	0	1	97	0	1.2	1	1	3	0
217	63	1	0	130	330	1	0	132	1	1.8	2	3	3	0
218	65	1	0	135	254	0	0	127	0	2.8	1	1	3	0
219	48	1	0	130	256	1	0	150	1	0.0	2	2	3	0
													3	
220	63	0	0	150	407	0	0	154	0	4.0	1	3		0
221	55	1	0	140	217	0	1	111	1	5.6	0	0	3	0
222	65	1	3	138	282	1	0	174	0	1.4	1	1	2	0
223	56	0	0	200	288	1	0	133	1	4.0	0	2	3	0
224	54	1	0	110	239	0	1	126	1	2.8	1	1	3	0
225	70	1	0	145	174	0	1	125	1	2.6	0	0	3	0
226	62	1	1	120	281	0	0	103	0	1.4	1	1	3	0
227	35	1	0	120	198	0	1	130	1	1.6	1	0	3	0
228	59	1	3	170	288	0	0	159	0	0.2	1	0	3	0
229	64	1	2	125	309	0	1	131	1	1.8	1	0	3	0
230	47	1	2	108	243	0	1	152	0	0.0	2	0	2	0
231	57	1	0	165	289	1	0	124	0	1.0	1	3	3	0
232	55	1	0	160	289	0	0	145	1	0.8	1	1	3	0
233	64	1	0	120	246	0	0	96	1	2.2	0	1	2	0
234	70	1	0	130	322	0	0	109	0	2.4	1	3	2	0
235	51	1	0	140	299	0	1	173	1	1.6	2	0	3	0
236	58	1	0	125	300	0	0	171	0	0.0	2	2	3	0
237	60	1	0	140	293	0	0	170	0	1.2	1	2	3	0
238	77	1	0	125	304	0	0	162	1	0.0	2	3	2	0
239	35	1	0	126	282	0	0	156	1	0.0	2	0	3	0
240	70	1	2	160	269	0	1	112	1	2.9	1	1	3	0
241	59	0	0	174	249	0	1	143	1	0.0	1	0	2	0
242	64	1	0	145	212	0	0	132	0	2.0	1	2	1	0
		_	_	4-0		_	_		_					_
243	57	1	0	152	274	0	1	88	1	1.2	1	1	3	0
244	56	1	0	132	184	0	0	105	1	2.1	1	1	1	0
245	48	1	0	124	274	0	0	166	0	0.5	1	0	3	0
246	56	0	0	134	409	0	0	150	1	1.9	1	2	3	0
247	66	1	1	160	246	0	1	120	1	0.0	1	3	1	0
248	54	1	1	192	283	0	9	195	0	0.0	2	1	3	0
249	69	1	2	140	254	0	0	146	0	2.0	1	3	3	0
250	51	1	0	140	298	0	1	122	1	4.2	1	3	3	0
												4	3	
251	43	1	0	132	247	1	0	143	1	0.1	1			0
252	62	0	0	138	294	1	1	106	0	1.9	1	3	2	0
253		1	0	100	299	0	0	125	1	0.9	1	2	2	0
	67			700										
0.5	67			4			0	125	0	0.0	2	0	2	
254	59	1	3	160	273	0								0
	59	1					0	147	1			3		
255	59 45	1	0	142	309	0		147	1	0.0	1	3	3	0
255 256	59 45 58	1 1 1	0 0	142 128	309 259	0 0	0	130	1	0.0 3.0	1 1	2	3 3	0 0
255	59 45	1	0	142	309	0				0.0	1		3	0
255 256 257	59 45 58 50	1 1 1	0 0 0	142 128 144	309 259 200	0 0 0	0 0	130 126	1 1	0.0 3.0 0.9	1 1 1	2 0	3 3 3	0 0 0
255 256 257 258	59 45 58 50 62	1 1 1 1 0	9 9 9	142 128 144 150	309 259 200 244	0 0 0	0 0 1	130 126 154	1 1 1	0.0 3.0 0.9 1.4	1 1 1	2 0 0	3 3 3 2	0 0 0
255 256 257	59 45 58 50	1 1 1	0 0 0	142 128 144	309 259 200	0 0 0	0 0	130 126	1 1	0.0 3.0 0.9	1 1 1	2 0	3 3 3	0 0 0
255 256 257 258 259	59 45 58 50 62 38	1 1 1 0 1	0 0 0 0 3	142 128 144 150 120	309 259 200 244 231	0 0 0 0	0 0 1 1	130 126 154 182	1 1 1	0.0 3.0 0.9 1.4 3.8	1 1 1 1	2 0 0	3 3 2 3	0 0 0 0
255 256 257 258 259 260	59 45 58 50 62 38 66	1 1 1 0 1	0 0 0 3 0	142 128 144 150 120 178	309 259 200 244 231 228	0 0 0 0 1	0 0 1 1	130 126 154 182 165	1 1 1 1	0.0 3.0 0.9 1.4 3.8 1.0	1 1 1 1 1	2 0 0 0 2	3 3 2 3 3	0 0 0 0
255 256 257 258 259 260 261	59 45 58 50 62 38 66 52	1 1 1 0 1 0	9 9 9 3 9	142 128 144 150 120 178 112	309 259 200 244 231 228 230	0 0 0 0 1	0 0 1 1 1	130 126 154 182 165 160	1 1 1 1 0	0.0 3.0 0.9 1.4 3.8 1.0 0.0	1 1 1 1 1 2	2 0 0 0 2 1	3 3 2 3 3 2	0 0 0 0 0
255 256 257 258 259 260	59 45 58 50 62 38 66	1 1 1 0 1	0 0 0 3 0	142 128 144 150 120 178	309 259 200 244 231 228	0 0 0 0 1	0 0 1 1	130 126 154 182 165	1 1 1 1	0.0 3.0 0.9 1.4 3.8 1.0	1 1 1 1 1	2 0 0 0 2	3 3 2 3 3	0 0 0 0
255 256 257 258 259 260 261 262	59 45 58 50 62 38 66 52 53	1 1 1 0 1 0 1	0 0 0 3 0 0	142 128 144 150 120 178 112	309 259 200 244 231 228 230 282	0 0 0 0 1 0	0 0 1 1 1 1	130 126 154 182 165 160 95	1 1 1 1 0	0.0 3.0 0.9 1.4 3.8 1.0 0.0 2.0	1 1 1 1 1 2	2 0 0 2 1 2	3 3 2 3 3 2 3	0 0 0 0 0 0
255 256 257 258 259 260 261 262 263	59 45 58 50 62 38 66 52 53 63	1 1 1 0 1 0 1 0	0 0 0 3 0 0	142 128 144 150 120 178 112 123 108	309 259 200 244 231 228 230 282 269	0 0 0 0 1 0 0	0 0 1 1 1 1	130 126 154 182 165 160 95 169	1 1 1 1 0 1	0.0 3.0 0.9 1.4 3.8 1.0 0.0 2.0 1.8	1 1 1 1 1 2 1	2 0 0 2 1 2	3 3 2 3 3 2 3 2	0 0 0 0 0 0
255 256 257 258 259 260 261 262	59 45 58 50 62 38 66 52 53	1 1 1 0 1 0 1	0 0 0 3 0 0	142 128 144 150 120 178 112	309 259 200 244 231 228 230 282	0 0 0 0 1 0	0 0 1 1 1 1	130 126 154 182 165 160 95	1 1 1 1 0	0.0 3.0 0.9 1.4 3.8 1.0 0.0 2.0 1.8	1 1 1 1 1 2	2 0 0 2 1 2 2	3 3 2 3 3 2 3 2 2 2	0 0 0 0 0 0
255 256 257 258 259 260 261 262 263 264	59 45 58 50 62 38 66 52 53 63 54	1 1 1 0 1 0 1 1 0	0 0 0 3 0 0 0 0	142 128 144 150 120 178 112 123 108 110	309 259 200 244 231 228 230 282 269 206	0 0 0 0 1 0 0	0 0 1 1 1 1	130 126 154 182 165 160 95 169 108	1 1 1 1 0 1 1	0.0 3.0 0.9 1.4 3.8 1.0 0.0 2.0 1.8	1 1 1 1 1 2 1 1	2 0 0 2 1 2 2	3 3 2 3 3 2 3 2 2 2	0 0 0 0 0 0
255 256 257 258 259 260 261 262 263 264 265	59 45 58 50 62 38 66 52 53 63 54 66	1 1 1 0 1 0 1 0 1 1	9 9 9 3 9 9 9 9 9	142 128 144 150 120 178 112 123 108 110	309 259 200 244 231 228 230 282 269 206 212	0 0 0 0 1 0 0 0	0 0 1 1 1 1 1 0	130 126 154 182 165 160 95 169 108 132	1 1 1 1 0 1 1	0.0 3.0 0.9 1.4 3.8 1.0 0.0 2.0 1.8 0.0	1 1 1 1 1 2 1 1 1 2	2 0 0 2 1 2 2 1	3 3 2 3 3 2 3 2 2 2 2	0 0 0 0 0 0 0 0
255 256 257 258 259 260 261 262 263 264 265 266	59 45 58 50 62 38 66 52 53 63 54 66 55	1 1 1 0 1 0 1 1 0 1	0 0 0 3 0 0 0 0 0 0	142 128 144 150 120 178 112 123 108 110 112	309 259 200 244 231 228 230 282 269 206 212 327	0 0 0 0 0 0 0 0 0 0 0 0 0	0 0 1 1 1 1 1 0 0	130 126 154 182 165 160 95 169 108 132 117	1 1 1 1 0 1 1 1	0.0 3.0 0.9 1.4 3.8 1.0 0.0 2.0 1.8 0.0	1 1 1 1 1 2 1 1 2	2 0 0 2 1 2 2 1 1 0	3 3 2 3 3 2 3 2 2 2 2 2 2	0 0 0 0 0 0 0 0 0
255 256 257 258 259 260 261 262 263 264 265	59 45 58 50 62 38 66 52 53 63 54 66	1 1 1 0 1 0 1 0 1 1	9 9 9 3 9 9 9 9 9	142 128 144 150 120 178 112 123 108 110	309 259 200 244 231 228 230 282 269 206 212	0 0 0 0 1 0 0 0	0 0 1 1 1 1 1 0	130 126 154 182 165 160 95 169 108 132	1 1 1 1 0 1 1	0.0 3.0 0.9 1.4 3.8 1.0 0.0 2.0 1.8 0.0	1 1 1 1 1 2 1 1 1 2	2 0 0 2 1 2 2 1	3 3 2 3 3 2 3 2 2 2 2	0 0 0 0 0 0 0 0
255 256 257 258 259 260 261 262 263 264 265 266 267	59 45 58 50 62 38 66 52 53 63 54 66 55 49	1 1 1 0 1 0 1 1 0 1 0 1	0 0 0 3 0 0 0 0 0 0 2	142 128 144 150 120 178 112 123 108 110 112 180 118	309 259 200 244 231 228 230 282 269 206 212 327 149	0 0 0 0 1 0 0 0 0	0 0 1 1 1 1 1 0 0	130 126 154 182 165 160 95 169 108 132 117 126	1 1 1 1 0 1 1 1 1	0.0 3.0 0.9 1.4 3.8 1.0 0.0 2.0 1.8 0.0 0.1 3.4	1 1 1 1 1 2 1 1 2 1 2	2 0 0 2 1 2 2 1 0 3	3 3 2 3 2 3 2 2 2 2 2 2	0 0 0 0 0 0 0 0 0
255 256 257 258 259 260 261 262 263 264 265 266	59 45 58 50 62 38 66 52 53 63 54 66 55	1 1 1 0 1 0 1 1 0 1	0 0 0 3 0 0 0 0 0 0	142 128 144 150 120 178 112 123 108 110 112	309 259 200 244 231 228 230 282 269 206 212 327	0 0 0 0 0 0 0 0 0 0 0 0 0	0 0 1 1 1 1 1 0 0	130 126 154 182 165 160 95 169 108 132 117	1 1 1 1 0 1 1 1	0.0 3.0 0.9 1.4 3.8 1.0 0.0 2.0 1.8 0.0	1 1 1 1 1 2 1 1 2	2 0 0 2 1 2 2 1 1 0	3 3 2 3 3 2 3 2 2 2 2 2 2	0 0 0 0 0 0 0 0 0

270	46	1	0	120	249	0	0	144	0	0.8	2	0	3		0	
271	61	1	3	134	234	0	1	145	0	2.6	1	2	2		0	
272	67	1	0	120	237	0	1	71	0	1.0	1	0	2		0	
273	58	1	0	100	234	0	1	156	0	0.1	2	1	3		0	
274	47	1	0	110	275	0	0	118	1	1.0	1	1	2		0	
275	52	1	0	125	212	0	1	168	0	1.0	2	2	3		0	
276	58	1	0	146	218	0	1	105	0	2.0	1	1	3		0	
277	57	1	1	124	261	0	1	141	0	0.3	2	0	3		0	
278	58	0	1	136	319	1	0	152	0	0.0	2	2	2		0	
279	61	1	0	138	166	0	0	125	1	3.6	1	1	2		0	
280	42	1	0	136	315	0	1	125	1	1.8	1	0	1		0	
281	52	1	0	128	204	1	1	156	1	1.0	1	0	0		0	
282	59	1	2	126	218	1	1	134	0	2.2	1	1	1		0	
283	40	1	0	152	223	0	1	181	0	0.0	2	0	3		0	
284	61	1	0	140	207	0	0	138	1	1.9	2	1	3		0	
285	46	1	0	140	311	0	1	120	1	1.8	1	2	3		0	
286	59	1	3	134	204	0	1	162	0	0.8	2	2	2		0	
287	57	1	1	154	232	0	0	164	0	0.0	2	1	2		0	
288	57	1	0	110	335	0	1	143	1	3.0	1	1	3		0	
289	55	0	0	128	205	0	2	130	1	2.0	1	1	3		0	
290	61	1	0	148	203	0	1	161	0	0.0	2	1	3		0	
291	58	1	0	114	318	0	2	140	0	4.4	0	3	1		0	
292	58	0	0	170	225	1	0	146	1	2.8	1	2	1		0	
293	67	1	2	152	212	0	0	150	0	0.8	1	0	3		0	
294	44	1	0	120	169	0	1	144	1	2.8	0	0	1		0	
295	63	1	0	140	187	0	0	144	1	4.0	2	2	3		0	
296	63	0	0	124	197	0	1	136	1	0.0	1	0	2		0	
297	59	1	0	164	176	1	0	90	0	1.0		1	2	1		0
298	57	0	0	140	241	0	1	123	1	0.2		1	0	3		0
299	45	1	3	110	264	0	1	132	0	1.2		1	0	3		0
300	68	1	0	144	193	1	1	141	0	3.4		1	2	3		0
301	57	1	0	130	131	0	1	115	1	1.2		1	1	3		0
302	57	0	1	130	236	0	0	174	0	0.0		1	1	2		0

Phase 2:

1. Supervised Learning

To predict whether it has a chance of a heart attack or not, we will use the following machine learning algorithms:

a. Logistic Regression algorithm

The Logistic Regression is a supervised linear classification algorithm. It can be used to classify objects of binary and multi-class problems.

b. Decision Tree algorithm

A decision tree is a non-parametric supervised learning algorithm, which is utilized for both classification and regression tasks. It has a hierarchical, tree structure, which consists of a root node, branches, internal nodes and leaf nodes.^[1]

To evaluate the performance of logistic regression and decision tree algorithms in predicting the chance of a heart attack, we will divide the dataset into two sets:

Training set: this data will constitute 70% of the dataset and it will be used for training the machine learning models.

Test set: This data constitutes 30% of the data and it will be used for evaluating the performance of the machine learning algorithms. We will use the accuracy, precision, sensitivity, and specificity performance metrics to evaluate the performance of the logistic regression and decision tree algorithms.

2. Implementation of the Algorithms

We implemented the logistic regression and decision tree algorithms in this project using Python programming language. We used Google Colab. ^[2] Colab, or "Collaboratory", allows you to write and execute Python in your browser, with

- Zero configuration required
- Access to GPUs free of charge
- Easy sharing

Loading the dataset

The following figure shows the python code for loading the dataset. We used the function read csv in Panda's library [3] to load the dataset.

```
import pandas as pd
from sklearn import preprocessing as pre
import numpy as np
data = pd.read_csv('heartpreprocessed.csv')
```

Figure 22: import and load the dataset

Preparing the dataset to implement algorithms:

- 1- Separate the dataset into features ('age', 'sex', 'cp', 'trtbps', 'chol', 'fbs', 'restecg', 'thalachh', 'exng', 'oldpeak', 'slp', 'caa', 'thall') and output and then we store the **target variable in variable Y** and store the **features in the matrix X.**
- 2- Divide the dataset into training/testing with ratio 70/30

Figure 23: divide the data set to train and test

Implementation Logistic Regression

In this algorithm, we divided our dataset into training and testing data as shown in the code snapshot where 70% of the data set is training and 30% is testing:

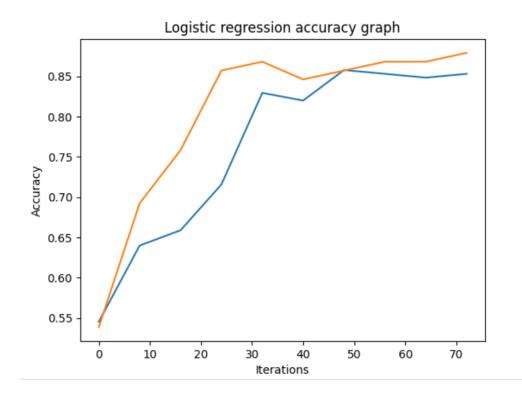
Figure 24: divide dataset for Logistic Regression

Then, we plotted our graph comparing our training and testing sets' accuracies over many iterations to make more accurate result:

```
[44] # Train and validation scores initialized as empty list
     train_score = []
     test_score = []
     #Loop for taking average result
     Array=8*(np.arange(10))
     for i in Array:
         # Create LogisticRegression object and fit
         lr = LogisticRegression( max_iter=i)
         lr.fit(x_train, y_train)
         # Evalueate scores and append to lists
         train_score.append( lr.score(x_train, y_train))
         test_score.append( lr.score(x_test, y_test))
     # Plot results
     plt.ylabel('Accuracy')
     plt.xlabel('Iterations')
     plt.title('Logistic regression accuracy graph')
     plt.plot(Array,train_score)
     plt.plot(Array,test_score)
```

Figure 25: build a Logistic Regression model

In orange is our training accuracy and in blue is our testing accuracy, we can conclude that as the iterations increases both accuracies increases but the training data will have higher accuracy than testing data.



Next, we plotted a confusion matrix using test data to asses the accuracy of the classification. The rows correspond to the actual label for which the results were intended. And the columns correspond to predicted label.

```
from sklearn import metrics
     y_pred=lr.predict(x_test)
     cnf_matrix = metrics.confusion_matrix(y_test, y_pred)
     cnf_matrix
[ ] import numpy as np
     import matplotlib.pyplot as plt
     import seaborn as sns
     %matplotlib inline
[42] class_names=[0,1] # name of classes
     fig, ax = plt.subplots()
     tick_marks = np.arange(len(class_names))
     plt.xticks(tick_marks, class_names)
     plt.yticks(tick_marks, class_names)
     # create heatmap
     sns.heatmap(pd.DataFrame(cnf_matrix), annot=True, cmap="YlGnBu" ,fmt='g')
     ax.xaxis.set_label_position("top")
     plt.tight_layout()
     plt.title('Confusion matrix', y=1.1)
     plt.ylabel('Actual label')
     plt.xlabel('Predicted label')
```

The output:

*output: 0 = less chance of heart attack 1 = more chance of heart attack

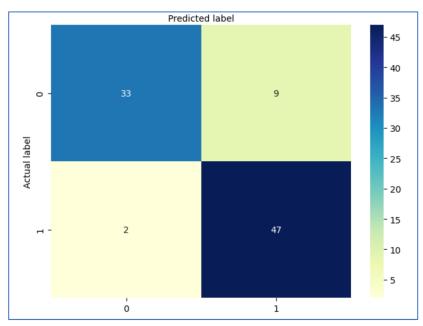


Figure 26: confusion matrix for Logistic Regression

Finally we calculated the CFM evaluation metrics Accuracy, Recall, f1-score using classification_report:

*output: 0 = less chance of heart attack 1 = more chance of heart attack

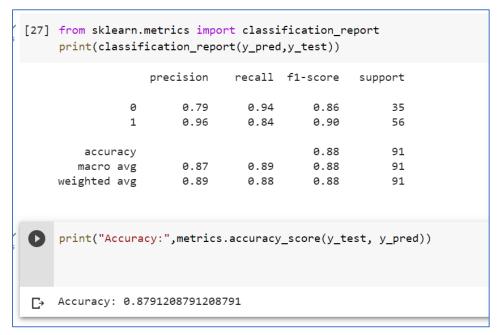


Figure 27: report for Logistic Regression

Implementation Decision Tree

We create the decision tree with entropy and we pass two parameters random_state=100 and max_depth=5 then we trained the model using fit() function then, we performed the test to predict the output(the chance of heart attack) using predict() function passing the test data (x_test) to it. Finally, we calculate the accuracy using accuracy_score() method and generate a report using classification_report() method and cross validation score using cross_val_score() which runs 5-folds cross validation on our dataset to see if the model generalizes to the entire dataset to produce an array of estimator scores for each cross validation run.

*output: 0 = less chance of heart attack 1 = more chance of heart attack

```
from sklearn.metrics import accuracy_score , classification_report
from sklearn.model_selection import cross_val_score
from sklearn.tree import DecisionTreeClassifier
# decision tree with entropy
clf_entropy = DecisionTreeClassifier(criterion="entropy" , random_state = 100 , max_depth=5)
# performing training
clf_entropy.fit(x_train , y_train )
# predicton on test with entropy
y_pred = clf_entropy.predict(x_test)

print("Accuracy :" ,accuracy_score(y_test , y_pred)*100)

print("Report :" ,classification_report(y_test , y_pred))

score= cross_val_score(clf_entropy , X , Y , cv=5)

print('Cross validation score :' , score)
```

Figure 28: build the decision tree model

```
Accuracy : 86.81318681318682
                     precision recall f1-score support
   Report :
                    0.92
                             0.79
                                       0.85
                                                  42
             1
                    0.84
                             0.94
                                                  49
                                       0.88
       accuracy
                                       0.87
                                                  91
                    0.88
                             0.86
      macro avg
                                       0.87
                                                  91
   weighted avg
                    0.87
                            0.87
                                       0.87
                                                  91
   Cross validation score : [0.73770492 0.86885246 0.73333333 0.78333333 0.63333333]
```

Figure 29: report of decision tree model

After that we pass (x train, y train) and (x test, y test) to score() after training the model on the data.

```
# Train Accuracy
data_train_accurcy= clf_entropy.score(x_train , y_train)
print("Training accuracy =" , data_train_accurcy)
# Test Accuracy
data_test_accurcy = clf_entropy.score(x_test , y_test)
print("Testing accuracy =" , data_test_accurcy)

Training accuracy = 0.909952606635071
Testing accuracy = 0.8681318681318682
```

Figure 30: training and test decision tree model with the result of accuracy

Decision tree visual representation:

We represent Decision tree representation using plot_tree() function.

Figure 31: represent Decision tree code

And this the output:

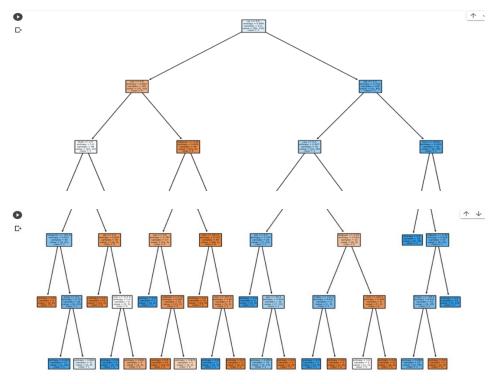


Figure 32: decision tree diagram

Finally, we use confusion_matrix() on the test data to assess the classification's accuracy and identify any model flaws. The rows correspond to the actual courses for which the results were intended. And the columns correspond to the predictions we've made.

```
import seaborn as sns
# confusion matrix
from sklearn.metrics import confusion_matrix
y_pred = clf_entropy.predict(x_test)
y_true = y_test
ab = ['#3b5f91' , '#7daa6a' ]
cm_dt = confusion_matrix(y_true , y_pred)
f , ax = plt.subplots(figsize=(10 , 5))
sns.heatmap(cm_dt , annot = True , linewidth = 0.1 , fmt =".0f" , cmap=ab , ax = ax)
plt.xlabel("Predicted")
plt.ylabel("Actual")
plt.title("Predicted vs actual confusion matrix ")
plt.show()
```

Figure 33: plot the confutation matrix code

The output:

*output: 0 = less chance of heart attack 1 = more chance of heart attack

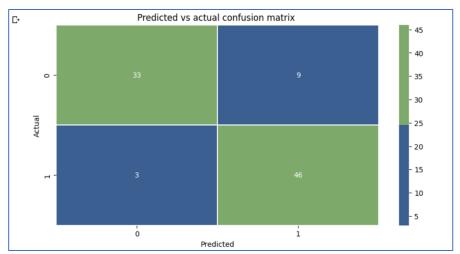


Figure 34: confusion matrix of decision tree

3. Result of Algorithms

Comparison between Logistic Regression and Decision tree in predicting the chance of a heart attack.

Performance Metric	Logistic Regression	Decision tree
Accuracy	87%	86.81%
Precision	89 %	87%
Sensitivity	88%	87%
F1 score	88 %	87%

Based on what is shown in the table, we can conclude that the logistic regression yielded better results than the decision tree.

Phase 3:

1. Unsupervised Learning

In this section, we use unsupervised learning algorithm which is k-means to group the dataset into clusters. To cluster the dataset using k-means algorithm, we perform the following steps:

Step 1: drop the class label

Step 2: Build the K-means model.

Step 3: Train the K-means model.

Step 4: Evaluate the K-means model.

We evaluate the k-means model for different number of clusters: from 2 clusters to 9 clusters. To evaluate the k-means model, we used two evaluation metrics: total within-cluster sum of square and Silhouette coefficient. The following figure shows python code for clustering using K-means algorithm and evaluating the performance of the algorithm.

Step 1:

While clustering is an unsupervised machine learning task, so we are going to give the algorithm a lot of input data with no class labels, so the first step will be dropping the class label, we also drop the first column, which is the index of the rows (not beneficial data).

```
import pandas as pd
from sklearn import preprocessing as pre
import numpy as np
data = pd.read_csv('heartpreprocessed.csv')
data.drop(['output', data.columns[0]], axis=1)
```

Figure 35: drop the class label

	age	sex	ср	trtbps	chol	fbs	restecg	thalachh	exng	oldpeak	slp	caa	thall
0	63	1	3	145	233	1	0	150	0	2.3	0	0	1
1	37	1	2	130	250	0	1	187	0	3.5	0	0	2
2	41	0	1	130	204	0	0	172	0	1.4	2	0	2
3	56	1	1	120	236	0	1	178	0	8.0	2	0	2
4	57	0	0	120	354	0	1	163	1	0.6	2	0	2
297	57	0	0	140	241	0	1	123	1	0.2	1	0	3
298	45	1	3	110	264	0	1	132	0	1.2	1	0	3
299	68	1	0	144	193	1	1	141	0	3.4	1	2	3
300	57	1	0	130	131	0	1	115	1	1.2	1	1	3
301	57	0	1	130	236	0	0	174	0	0.0	1	1	2

302 rows x 13 columns

Figure 36: drop the class label result

Step 2, 3, 4:

In this step we will build a k-means model which is technique used to identify clusters of data objects in a dataset. [4]

First, we will import the needed library for build, train, evaluate the k-means model.

```
from traitlets.config import List
from sklearn.metrics.cluster import silhouette_score
import matplotlib.pyplot as plt
from sklearn.cluster import KMeans
from sklearn.preprocessing import StandardScaler
from yellowbrick.cluster import SilhouetteVisualizer
```

Figure 37: library used for build, Train, Evaluate the k-means model

Second, we build our model then we train and evaluate the model

```
# Extract features: Feature extraction is one of the crucial
# steps to obtain an efficient representation of input patterns
X = data.iloc[:, :].values
WCSS=[]
for k in range(2,10):
  #create K-means model
  Kmeans=KMeans(n clusters=k, max iter=1000)
  #train the model using the dataset
  Kmeans.fit(X)
  #evaluate the model
  lables =Kmeans.predict(X)
  #claculate Silhouette coefficient score for each number of clustrs
  score =silhouette score(X,lables, metric='euclidean')
  print("K"+str(k)+" Silhouette coefficient score: " +str(score))
  #claculate WCSS for each number of clusters
  WCSS.append(Kmeans.inertia )
```

Figure 38: build, train and evaluate the model

Third, as we see in above code figure 38, we calculate the Silhouette score for each number of clusters, and as we know the silhouette ranges from -1 to +1, where a high value indicates that the object is well matched to its own cluster and poorly matched to neighboring clusters. If most objects have a high value, then the clustering configuration is appropriate.

The following figure shows the Silhouette score for different number of clusters. The results show that the highest Silhouette score was 0.4328835 for two clusters. This implies that the optimal number of clusters are two clusters

```
K2 Silhouette coefficient score: 0.4328835170484906
K3 Silhouette coefficient score: 0.3072623091388496
K4 Silhouette coefficient score: 0.295036719835046
K5 Silhouette coefficient score: 0.3037831612983324
K6 Silhouette coefficient score: 0.2830645031989993
K7 Silhouette coefficient score: 0.2757697191237553
K8 Silhouette coefficient score: 0.2503332935005701
K9 Silhouette coefficient score: 0.23767388915462362
```

Figure 39: Silhouette score for each cluster

```
K2 WCSS: 1630264.624818801
K3 WCSS: 1288613.1833022707
K4 WCSS: 1068553.1340408612
K5 WCSS: 871773.8790552414
K6 WCSS: 779569.3397454319
K7 WCSS: 711998.9920311256
K8 WCSS: 655760.4541068628
K9 WCSS: 602258.2916314325
```

Figure 40: calculate SSE for each cluster

to justify why we choice [2, 3, 5] size for K, for that we will plot the elbow diagram to find the best value of K, by using number of cluster and WCSS which we have collect it in Figure 38 and displayed in figure 40. ^[6]

```
# plot the elbow diagram by using number of cluster and WCSS
plt.figure()
plt.plot(range(1, 10), WCSS)
plt.xlabel("Number of clusters")
plt.ylabel("WCSS")
plt.show()
```

Figure 41: plot the elbow diagram code

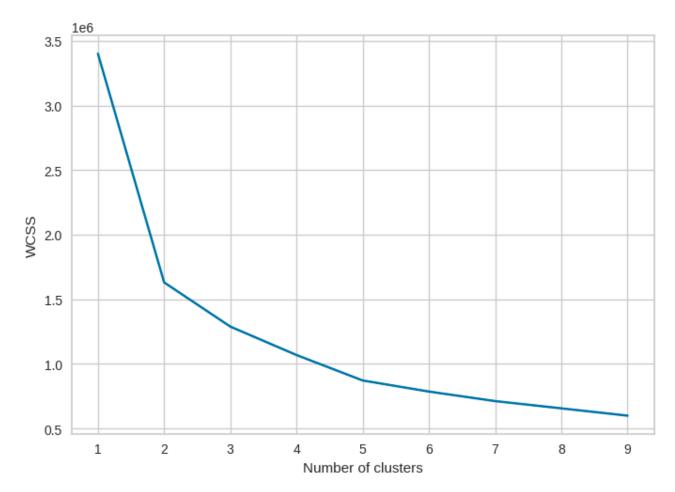


Figure 42: elbow diagram

So, after that we have find the optimal value of clusters which is looks like an Elbow we can find it clearly in cluster 2 and 5 but 3 is difficult to find, and now we will find plot of the optimal values of the Silhouette Visualizer. [5]

```
fig, ax = plt.subplots(2,2, figsize=(6,4))
for i in [2,3,5]:
    #create KMeans instanc for different number of clusters
    km= KMeans(n_clusters=i, init="k-means++", n_init=10, max_iter=1000, random_state=1)
    q, mod= divmod(i, 2)
    #create SilhouetteVisualizer instance with KMeans instane and Fit the visualizer
    visualizer= SilhouetteVisualizer(km, colors='yellowbrick', ax=ax[q-1][mod])
    visualizer.fit(X)
```

Figure 43: plot the Silhouette Visualizer code

*x-axis: represent the Silhouette score and y-axis: cluster label

The result of following figure is the best result we have implemented.

Silhouette Plot of KMeans Clustering for 302 Samples in 2 Centers

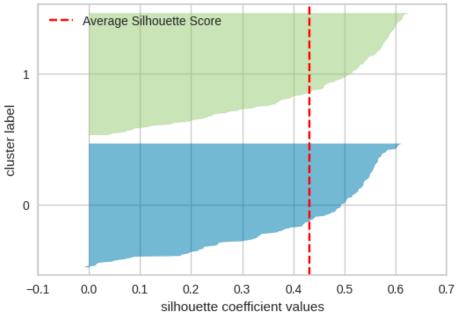


Figure 44: Silhouette Visualizer for two cluster

Silhouette Plot of KMeans Clustering for 302 Samples in 3 Centers

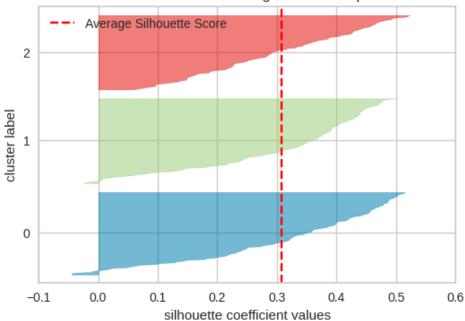


Figure 45: Silhouette Visualizer for three cluster



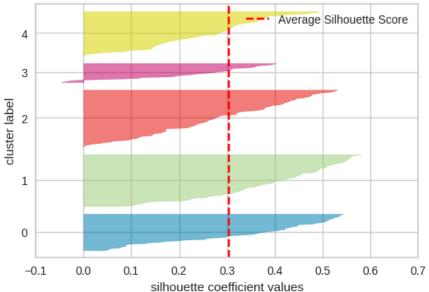
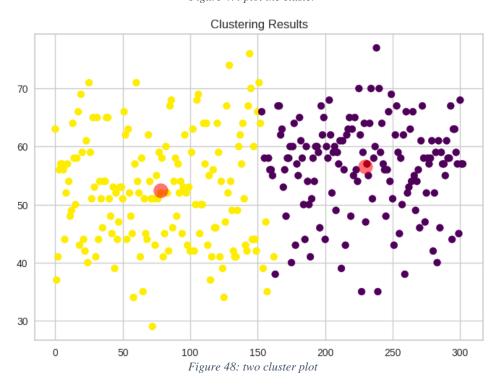


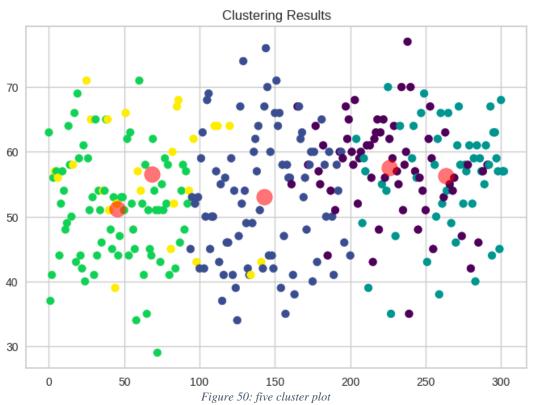
Figure 46: Silhouette Visualizer for five cluster

```
#plot the result of each clustr
plt.scatter(X[:, 0], X[:, 1], c= lables, s=50, cmap='viridis')
plt.scatter(Kmeans.cluster_centers_[:, 0], Kmeans.cluster_centers_[:, 1], c='red', s=200, alpha=0.5)
plt.title('Clustering Results')
plt.show()
```

Figure 47: plot the cluster







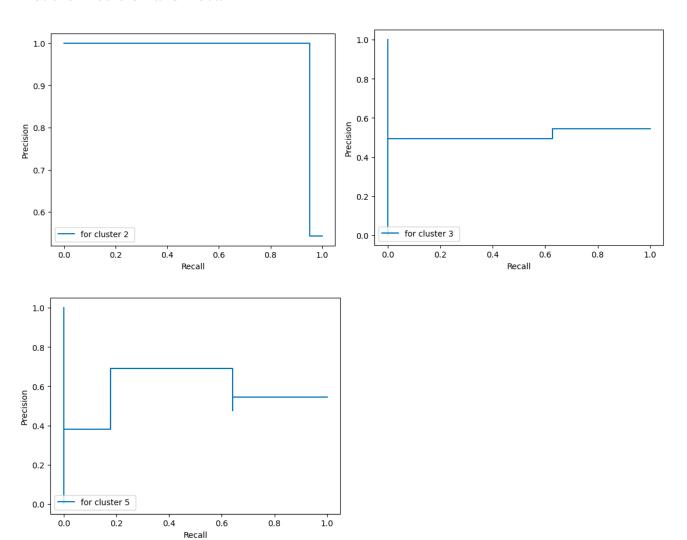
Plot Precision Recall Curve

Precision can be seen as a measure of quality and recall as a measure of quantity. Higher precision means that an algorithm returns more relevant results

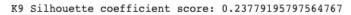
Precision and Recall Formulas

$$precision = \frac{tp}{tp + fp}$$
 $recall = \frac{tp}{tp + fn}$

Plot the Precision and Recall



And to see the different between the optimal values of K and bad ones we have taken k=9 as example which is the worst value we have implement it, we can see that the Silhouette score have value of 0.23779 which considered not good score, also in the figure 51 we can clearly see the unacceptable result.



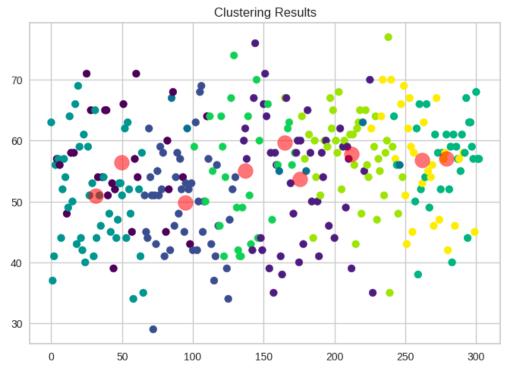


Figure 51: nine cluster plot

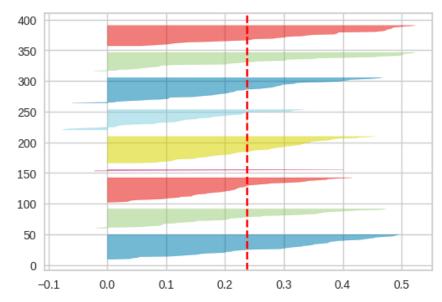


Figure 52: Silhouette Visualizer for nine cluster

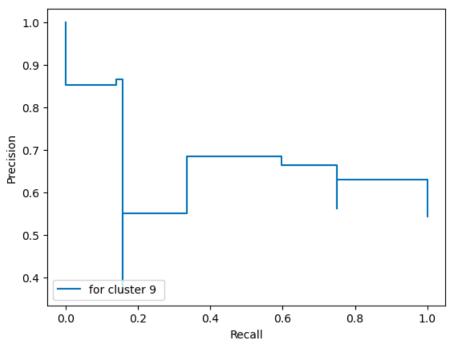
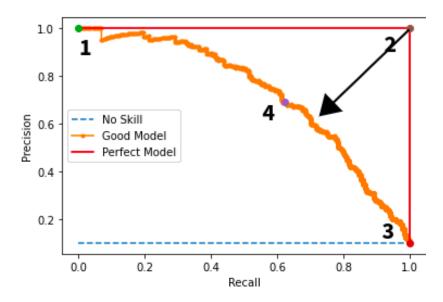


Figure 53: Precision and Recall for nine cluster

So, as we know the perfict fit of Precision and Recall is to fit the border of right and up. Ex:



We see that the k=3.5 have a good fit while the k=2 have properly perfect fit.

Conclusion

In this project, we applied two supervised machine learning classification algorithms, namely Logistic Regression and DT to predicting the chance of a heart attack using measurements of mineral elements. We also used K-means clustering algorithm to group the dataset into different number of clusters. The dataset was cleaned before data mining tasks. Results showed that data mining classification algorithms can successfully predict the chance of a heart attack with a classification accuracy near to 90%. Results of unsupervised learning showed that the optimal number of clusters were two clusters.

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