# **Data Mining Project**

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## THE DATASET LINK

https://www.kaggle.com/datasets/sulianova/cardiovascular-disease-dataset

## **Main Task:**

**Predicting Cardiovascular Disease** 

## How we accomplish this?

Create a predictive model that uses demographic and health data to assess if a person has cardiovascular disease.

Does the individual have cardiovascular disease? (1 = Yes, 0 = No).

# Before starting implementing the tasks..

We adjusted the dataset and made some errors like: Duplicating id, Missing values in age and weight, Value range error for Height and selected a random sample to focus our analysis

## Task 1:

1. Understand the data: Import the data selected "........csv" file into RapidMiner. While importing the data, you should properly choose the types of attributes. Identify the type of each attribute (binary, nominal, ordinal, numeric) and report it. Get statistics about the data to identify the values of the summarizing properties for each attribute, including frequency, location, and spread [e.g., value ranges of the attributes, frequency of values, medians, means, variances, or standard deviation].

# 1.1 Identify the type of each attribute:

Attribute Name	Туре	Description
Age	Numeric	Described as numerical value, presented by days
Weight	Numeric	Described as numerical value ,Weight by kilogram
ID	Nominal	Identifier, unique for everyone
Gender	Numeric	Gender (1=female, 2=male)
Height	Numeric	Described as numerical value, Height by centimeters
Ap_hi	Numeric	Systolic blood pressure
Ap_lo	Numeric	Diastllic blood pressure
Cholesterol	Ordinal	Cholesterol level (1=normal, 2= above normal, 3= High)
Gluc	Ordinal	Glucose level (1=normal, 2= above normal, 3= High)
Smoke	Binary	Smoking status ( 0= doesn't smoke , 1= smoker)
Aloc	Binary	Alcohol user (0= no, 1= yes)
Active	Binary	Physical status (unactive = $0$ , active =1)
Cardio	Binary	Heart disease (0= safe, 1= not safe)

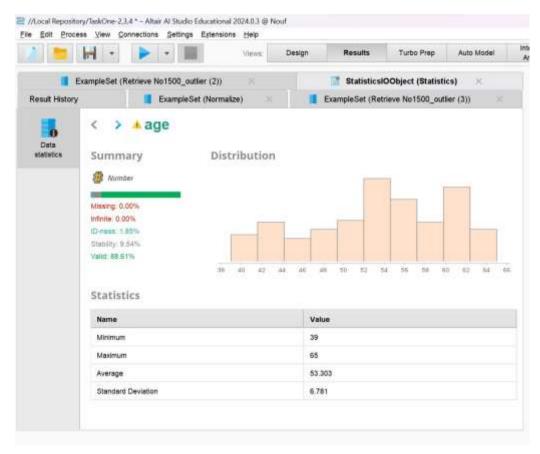
## 1.2 Properties of attributes

	Column	Description
1	Age	Represents the age of the individual in days. It is a numerical attribute used to calculate age in years or categorize individuals into age groups
2	Weight	Represents the weight of the individual in kilograms. It is a continuous variable that can be used for calculating Body Mass Index (BMI).
3	ID	A unique identifier for each individual in the dataset. It is non-numerical and used only for distinguishing individuals.
4	Gender	Represents the gender of the individual
5	Height	Represents the height of the individual in centimeters. This numerical attribute can also be used for BMI calculation.
6	Ap_hi	Represents the systolic blood pressure (the higher value in a blood pressure reading). It is a continuous variable.
7	Ap_lo	Represents the diastolic blood pressure (the lower value in a blood pressure reading). It is a continuous variable.
8	Cholesterol	Indicates the cholesterol level: 0 1  = Normal 0 2 =  Above Normal 0 3 =  High
9	Gluc	Indicates the glucose level:  o 1 = Normal o 2 =  Above Normal o 3 =  High
10	Smoke	Indicates smoking status:  o 0 = Does  not smoke  o 1 =  Smoker
11	Alco	Indicates alcohol usage:  ○ 0 = Does not consume alcohol ○  1 = Consumes alcohol

12	Active	Represents physical activity level:   0 = Not active  1 = Active
13	Cardio	Indicates the presence of cardiovascular disease:   0 = No (safe)  1 = Yes (not safe)

### **Data Exploratory Analysis and Visualization**

#### 1.3 Statistics about some of the data

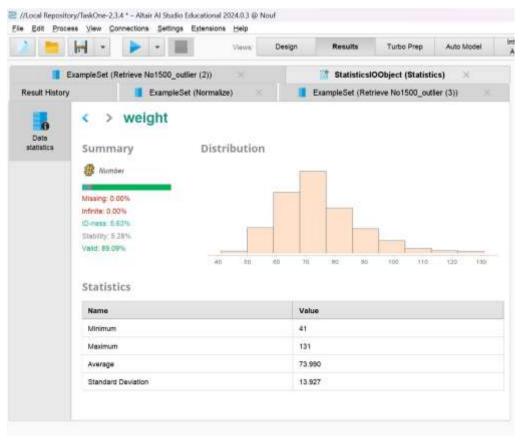


## Brief explaintion About Age:

age (Age Distribution) Description:

The plot shows the age distribution of the dataset's participants, with an average age of 53 and a range of 39 to 65.3.

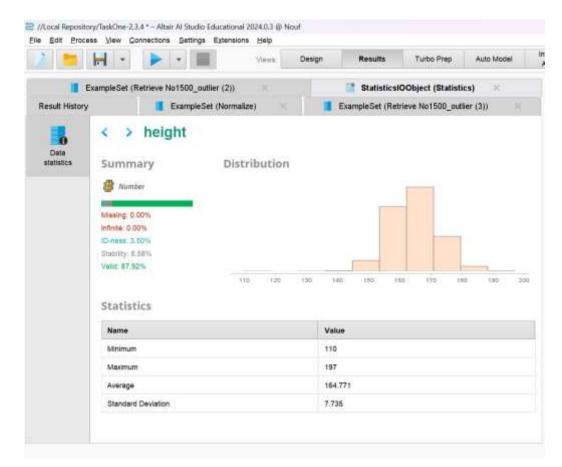
Significance: An important risk factor for cardiovascular disease is age. Since older people are typically at higher risk, this feature is essential.



### Brief explaintion About Weight:

#### distribution of (weight)

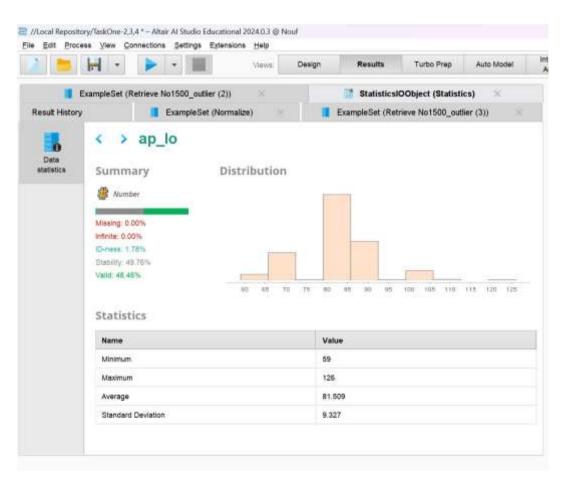
The plot shows the weight distribution, with an average of 73.99 kg and a range of 41 to 131 kg. Relevance: Body mass index (BMI), a recognized risk factor for cardiovascular health problems, is directly correlated with weight.



Brief explaintion About Height: height

#### (Height Distribution)

This plot displays the range of individual heights, with an average of 164.77 cm and a range of 110 to 197 cm. Relevance: Although height by itself may not be a reliable indicator of cardiovascular disease, it may serve as a stand-in when paired with weight (e.g., BMI).

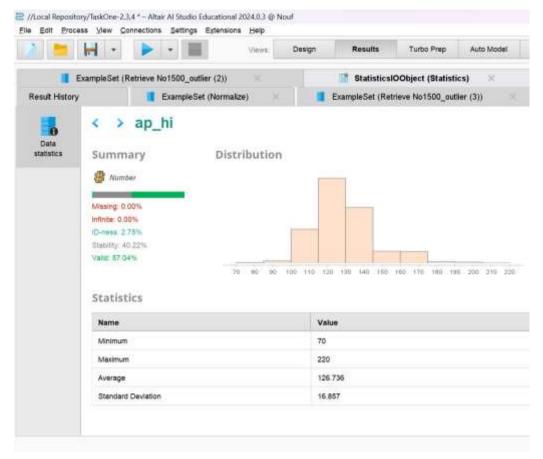


#### Brief explaintion About ap lo:

#### Diastolic Blood Pressure Distribution (ap lo):

This plot displays the dataset's diastolic blood pressure distribution (ap\_lo). It has an average of 81 and ranges from 59 to 126.51.

Relavance: One of the main markers of cardiovascular problems is high or low blood pressure.



Brief explaintion About ap high:

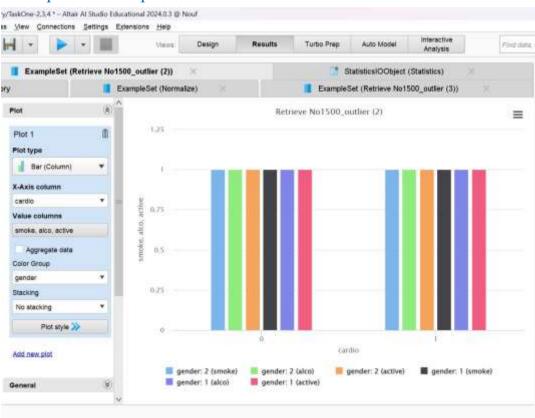
Systolic Blood Pressure Distribution (ap\_hi)

This plot shows the average systolic blood pressure (ap\_hi), which ranges from 70 to 220.

Relevance: Systolic blood pressure is essential for detecting hypertension, just like diastolic blood pressure.

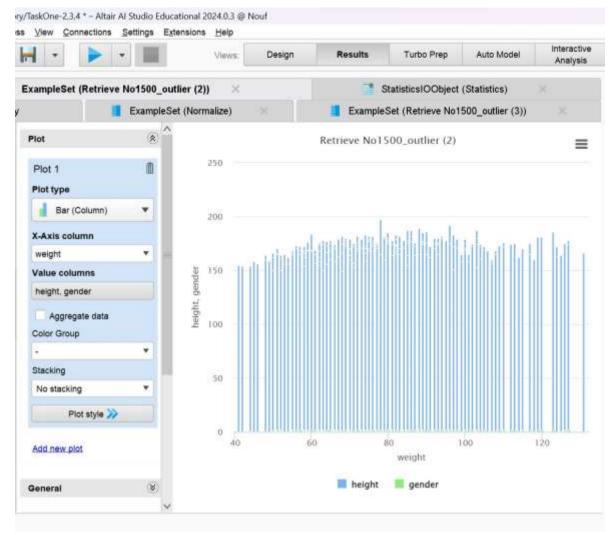
## 2. Analyze the data by using different visualization techniques provided by RapidMiner.

#### 2.1 Explanation of plot one:



#### (Lifestyle Choices and Gender in Relation to Cardiovascular Disease)

The association between gender, physical activity, alcohol use, smoking, and cardiovascular disease is depicted in this plot. Relevance: Modifiable risk factors for cardiovascular disease include lifestyle choices like drinking alcohol and smoking. Risk prediction may also be influenced by gender differences (1 = Female, 2= Male), you can tell that both male and female almost have equal Lifestyle, so the chance for having Cardiovascular Disease is possible for both of them if they don't stop the bad habits like alco drinking, smoking.

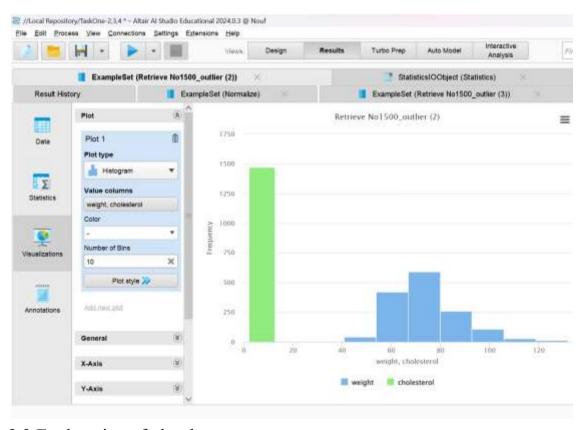


#### 2.2 Explanation of plot two:

### (Weight and Height Overview)

This plot contrasts the distributions of height and weight by gender. It draws attention to individual differences in these parameters.

Relevance: These characteristics affect BMI, which is a measure of obesity, a major risk factor for cardiovascular disease.



### 2.3 Explanation of plot three:

#### (Cholesterol and Weight)

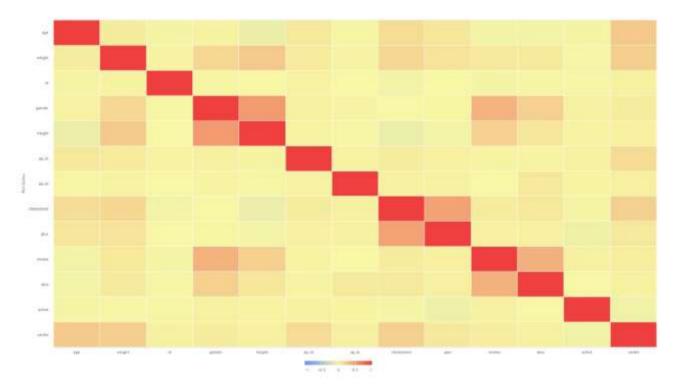
The frequency of weight and cholesterol levels in the dataset is shown visually in this histogram.

Relevance: One of the main cardiovascular risk factors, hyperlipidemia, can be identified by measuring cholesterol levels. The combination of weight and cholesterol yields important information.

3. Find and analyze the correlated attributes (you can use the "Correlation Matrix" operator or Scatter Plot chart).

## 3.1 Correlation matrix

Attribut	age	weight	id	gender	height	ap_hi	ap_lo	cholest	gluc	smoke	alco	active	cardio
198	1	0.057	-0.017	0.022	-0.073	0.077	0.003	0.141	0.089	-0.025	-0.030	-0.017	0.244
weight	0.057	1	0.010	0.171	0.252	0.059	0.014	0.169	0.105	0.079	0.067	-0.008	0.218
id	-0.017	0.010	1.	0.004	-0.002	0.023	-0.001	-0.031	0.000	-0.014	-0.005	-0.006	0.026
gender	0.022	0.171	0.004	1	0.507	0.030	0.012	-0.002	0.007	0.372	0.212	0.028	0.057
height	-0.073	0.252	-0.002	0.507	1	0.014	-0.005	-0.081	-0.025	0.212	0.089	0.015	0.032
ap_hi	0.077	0.059	0.023	0.030	0.014	1	0.024	0.055	0.031	0.009	0.011	0.028	0.148
ap_lo	0.003	0.014	-0.001	0.012	-0.005	0.024	1	0.027	0.029	-0.002	0.079	0.019	0.037
choleste	0.141	0.169	-0.031	-0.002	-0.081	0.055	0.027	(1)	0.458	0.055	0.077	-0.015	0.207
gluc	0.089	0.105	0.000	0.007	-0.025	0.031	0.029	0.458	1	0.033	0.038	-0.055	0.083
smoke	-0.025	0.079	-0.014	0.372	0.212	0.009	-0.002	0.055	0.033	1:	0.378	0.030	0.048
ałco	-0.030	0.067	-0.006	0.212	0.089	0.011	0.079	0.077	0.038	0.378	4	-0.001	0.030
active	-0.017	-0.008	-0.006	0.028	0.015	0.028	0.019	-0.015	-0.055	0.030	-0.001	1	-0.052
cardio	0.244	0.218	0.026	0.057	0.032	0.148	0.037	0.207	0.083	0.048	0.030	-0.052	1



## Explanation:

#### **Positive Correlations:**

height and weight (0.252): Taller individuals tend to weigh more, which is a logical and expected trend.

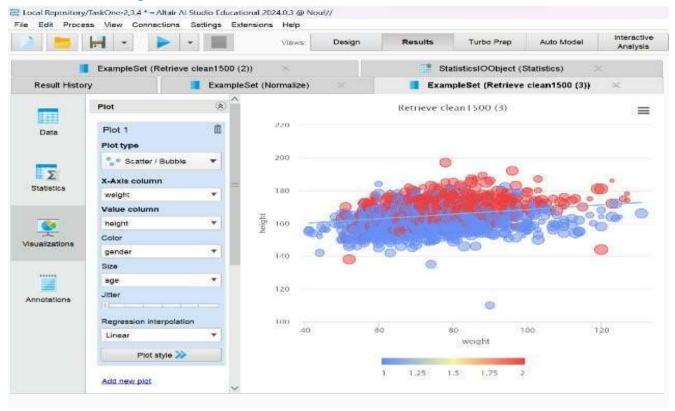
**cholesterol and gluc (0.458):** High cholesterol levels are moderately correlated with elevated glucose levels, indicating potential links to metabolic or cardiovascular health issues. **age and cardio (0.244):** Older age is moderately associated with cardiovascular conditions, reinforcing age as a risk factor.

weight and cardio (0.218): Higher weight is linked to cardiovascular issues, likely due to obesity-related risks.

### **Negative Correlations:**

Attributes like smoke, alco, and active show very low correlations with most variables.

#### 3.2 Scatter plot



#### **Explanation:**

There is a visible positive trend: as height increases, weight also increases.

This matches the correlation value of 0.252 from the matrix.

#### - Gender Based Differences:

Red points and blue points (representing different genders)

Red points (Female) are more concentrated in lower height and weight ranges.

Blue points (Male) are spread across a wider range of height and weight.

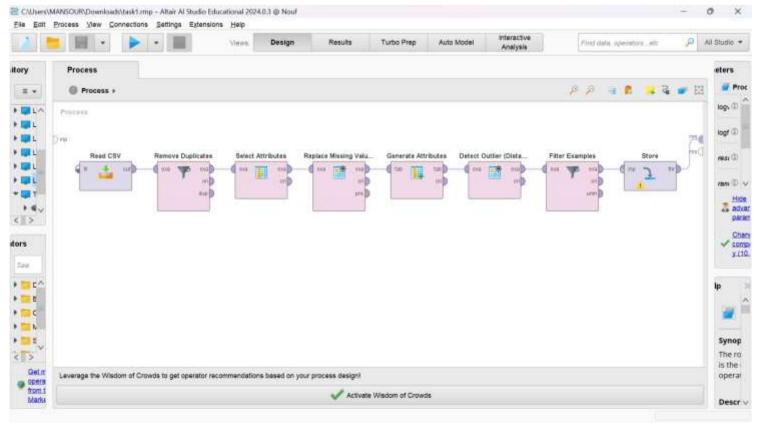
#### - Age Influence:

Larger-sized points (older individuals) are more distributed across the plot, indicating that age influences both weight and height variations.

4. Propose a way of cleaning data: e.g., replacing missing values, doing normalization if needed, corrections or errors, etc. 4.1 Preprocessing

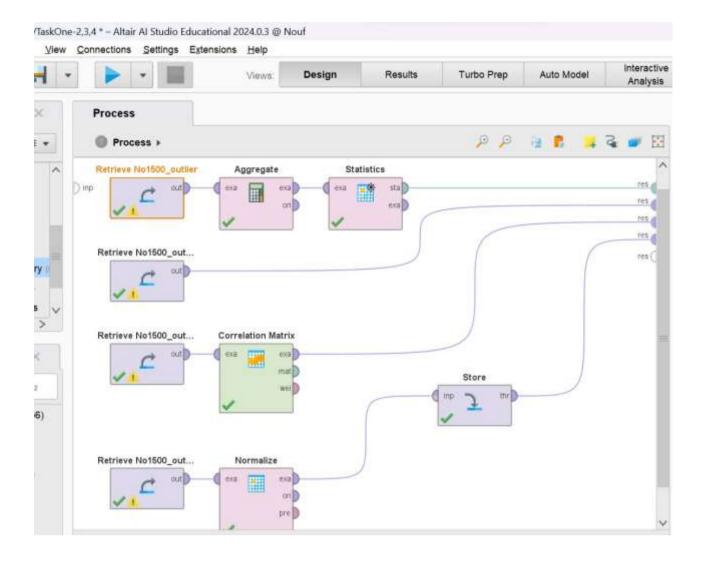
We have fixed those errors Duplicating id, Missing values in age and weight, Value range error for Height by and also did normalization but it will show on the final processes for task one down below:

### Fixing the errors:



## Final Process for Task one:

statistics, visualization, correlation, normalization of (Height, Weight, ap\_lo, ap\_hi) and it has been done by these processes



Task 2:
Evaluation Approach (Performance Measures)

Measure	Description	Definition of Value Function	Weight	Threshold
Accuracy	Percentage of test set tuples that are correctly classified	1 – Test MisClassification Rate	0.60	(>) 0.70
Simplicity	Model should be easy to understand.	Assuming that the ideal DT would have between 3(i.e. $Ideal_{Bot}$ ) through 7 (i.e. $Ideal_{Top}$ ) leaves, and the DT with 2 or less leaves (i.e. $CutOff_{Bot}$ = 2) or 14 or more leaves (i.e. $CutOff_{Top}$ = 14) is unacceptable;	0.05	3 ≤ Leaves < 14
Lift	For a given Target Event, it Measures the relative improvement that the model provides versus a random guess.	Based on the positive target event @ the 3rd decile	0.20	(>) 0
Stability	The model behaves similarly when applied to different datasets. Stability is binary, with 1 indicating a stable model and 0 indicating an unstable model as determined by the visual inspection of the noncumulative % Response Lift Chart for the given model	Based on visual inspection of Non-Cumulative Lift chart	0.15	1=>

1. Generate decision trees for the following parameter combinations and select the 'best' decision tree based on the criteria: Accuracy, Simplicity, Stability, and Lift (based on the positive target event @ the 3rd decile).

NumOfObsInModelSet =1471
TrainingPercent = 0.70
Test rows = 441
Train rows = 1030
MinObservationsPerLeaf = Ceiling(NumOfObsInModelSet * TrainingPercent * MinPercentOfTrainingObsPerLeaf)
Minimal Size for Split = 2* Leaf Size

DT No.	Splitting Criterion	MinPercentOfTrainingObsPerLeaf	Leaf Size	Minimal Size for Split	Confidence
1	Information Gain	3.50%	37	74	0.1
2	Information Gain	3.50%	37	74	0.2
3	Information Gain	6.00%	62	124	0.1
4	Information Gain	6.00%	62	124	0.2
5	Information Ratio	3.50%	37	74	0.1
6	Information Ratio	3.50%	37	74	0.2
7	Information Ratio	6.00%	62	124	0.1
8	Information Ratio	6.00%	62	124	0.2
9	Gini	3.50%	37	74	0.1
10	Gini	3.50%	37	74	0.2
11	Gini	6.00%	62	124	0.1
12	Gini	6.00%	62	124	0.2

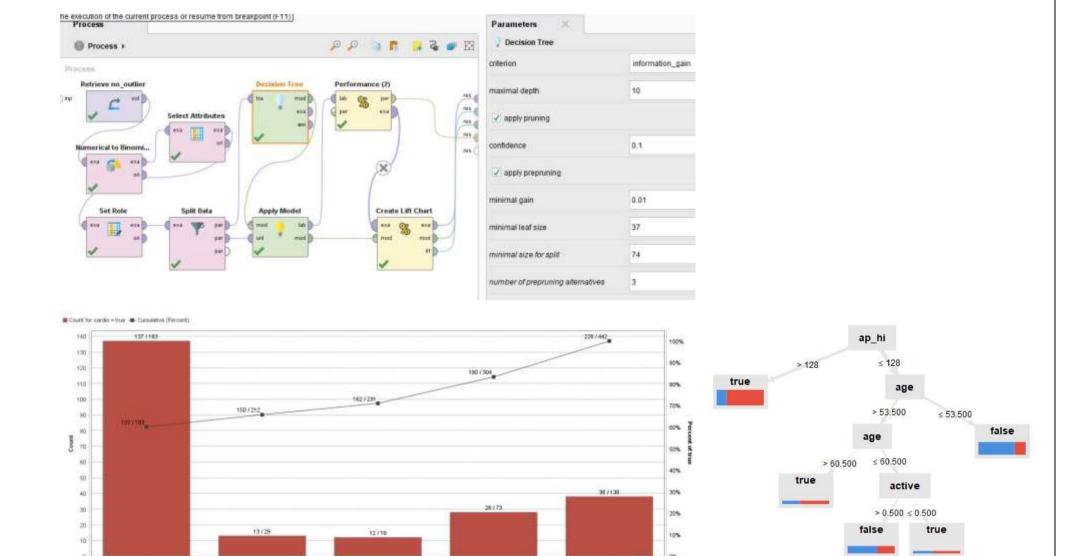
2. Specify combination function including weights that is to be used to obtain the Composite Performance Score for each DT:

$$\begin{aligned} Overall\_Score &= w_{Accuracy}*Score_{Accuracy} & + \\ & w_{Simplicity}*Score_{Simplicity} & + \\ w_{Lift}*Score_{Lift} & + \\ w_{Stability}*Score_{Stability} & \end{aligned}$$

Combination Function (overall score) =0.60\*Accuracy+0.05\*Simplicity+0.20\*Lift+0.15\*Stability 2.

# **Summary of Results**

3. Evidence of Experimentation 1. DT 1



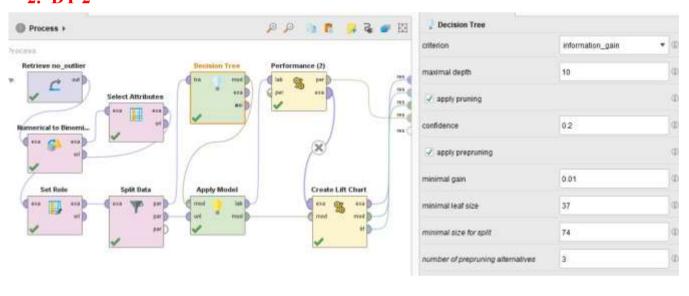
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#### 2. DT 2

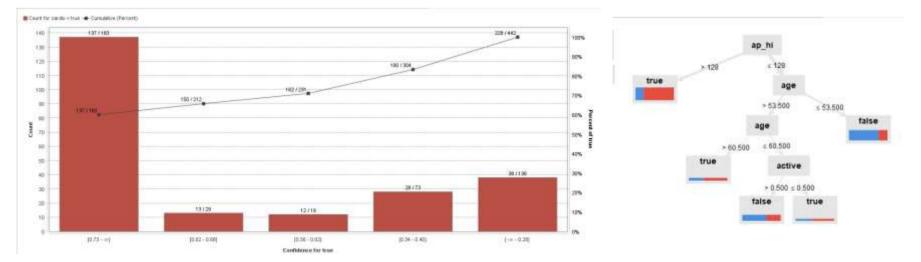
 $j \equiv 73 - -j$ 

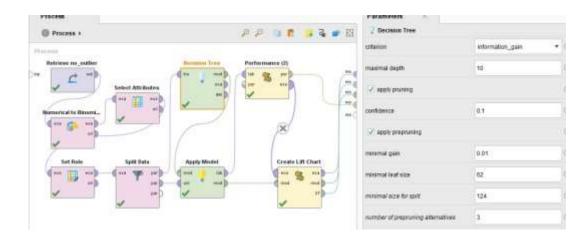
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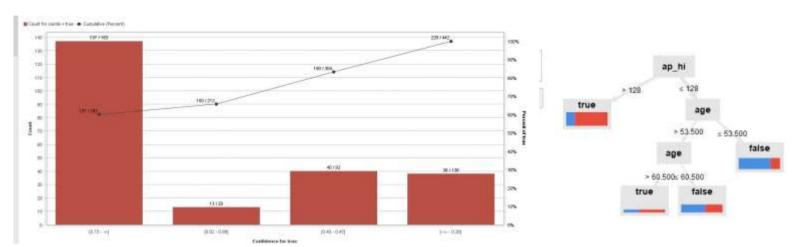


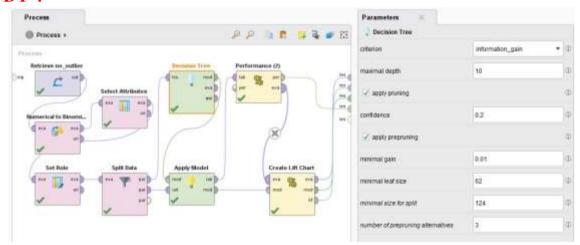
10:56-0:62]

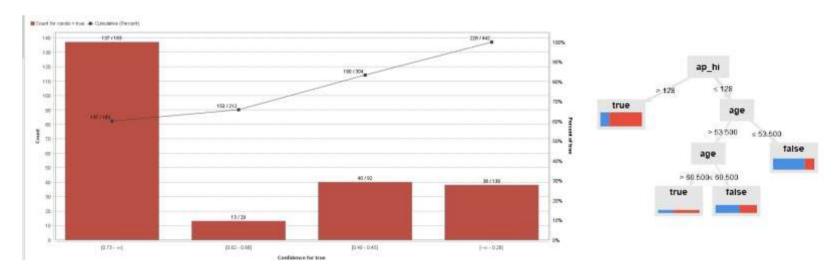
Confidence for true

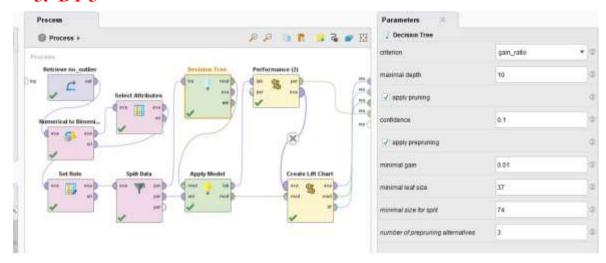


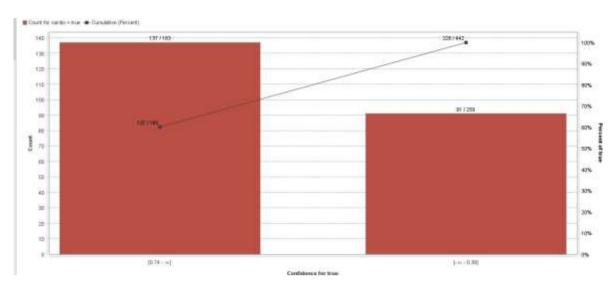


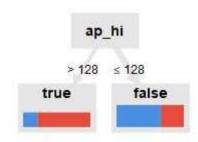


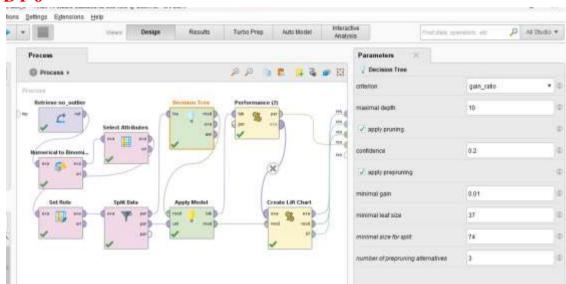


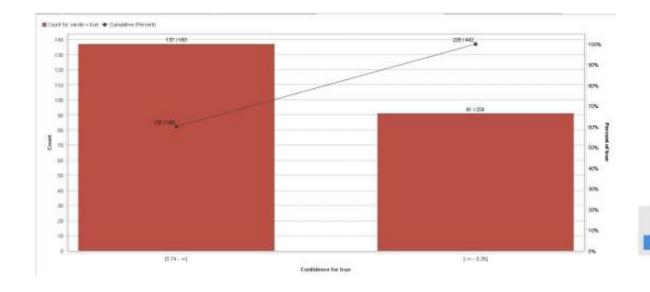


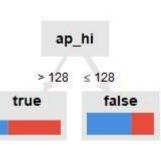


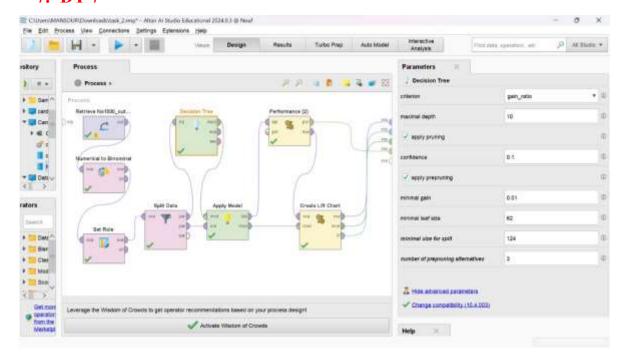


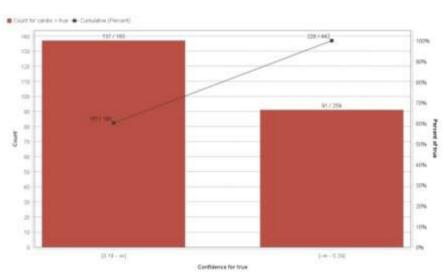


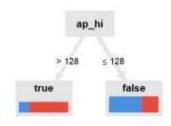


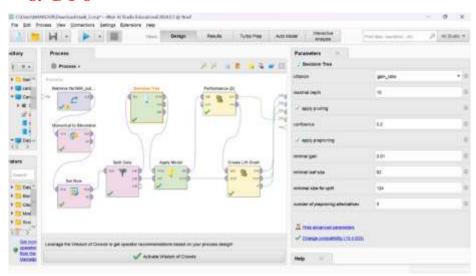


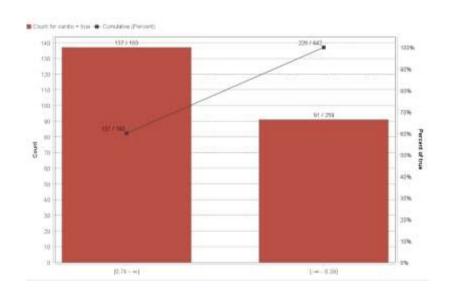


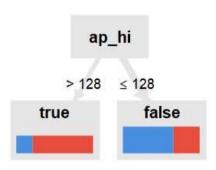




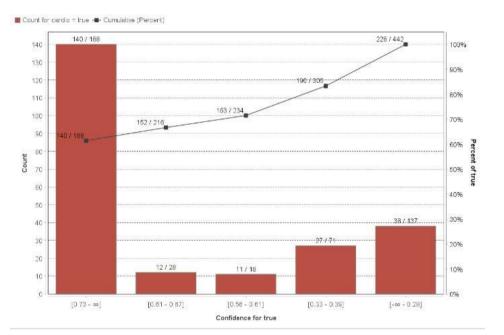


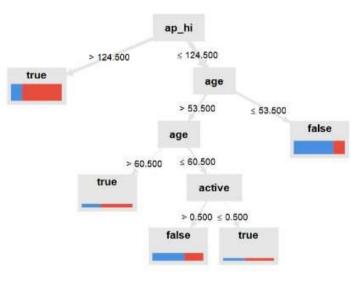


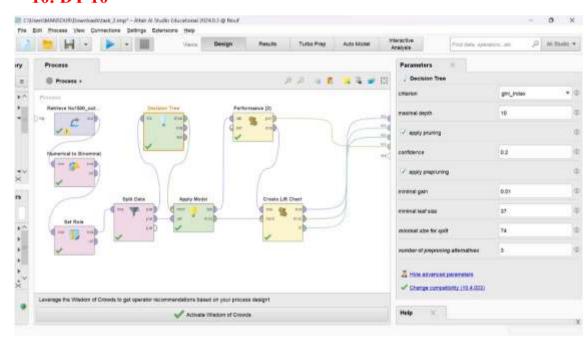


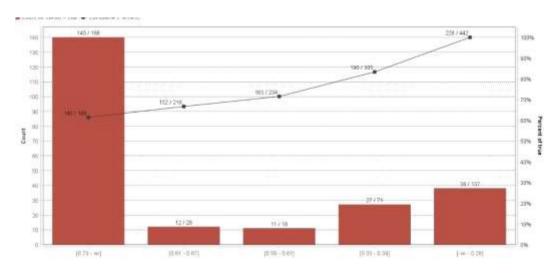


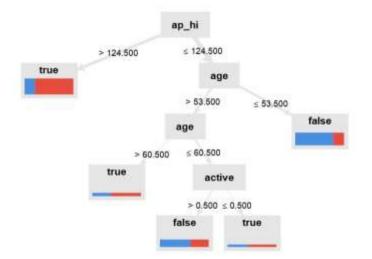




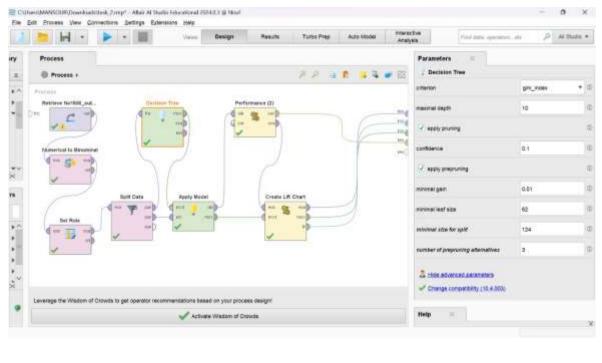


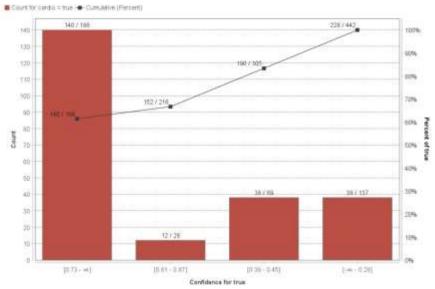


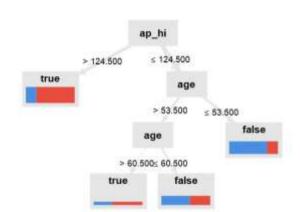




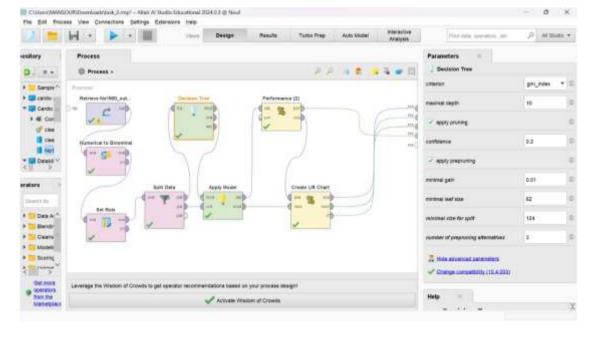
#### 11.DT 11

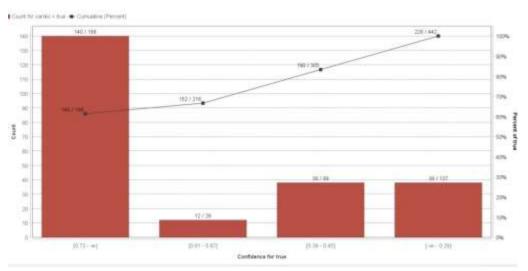


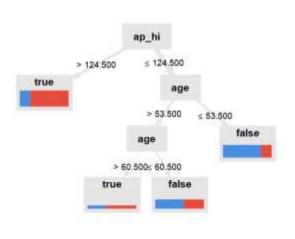




### 12.DT 12

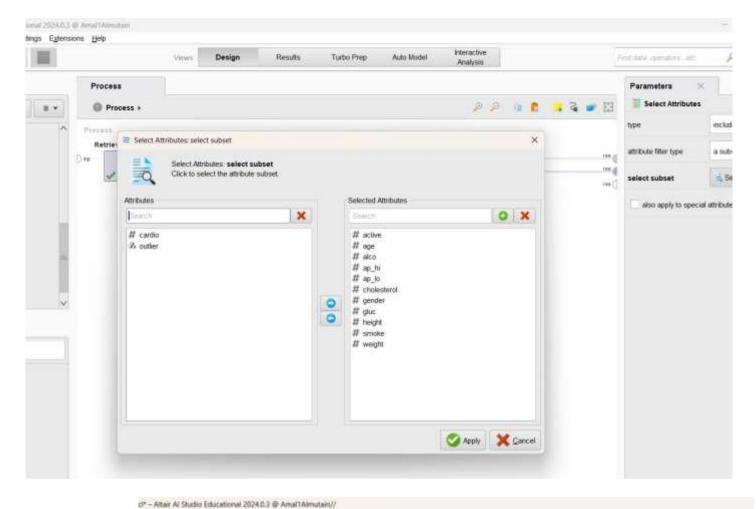


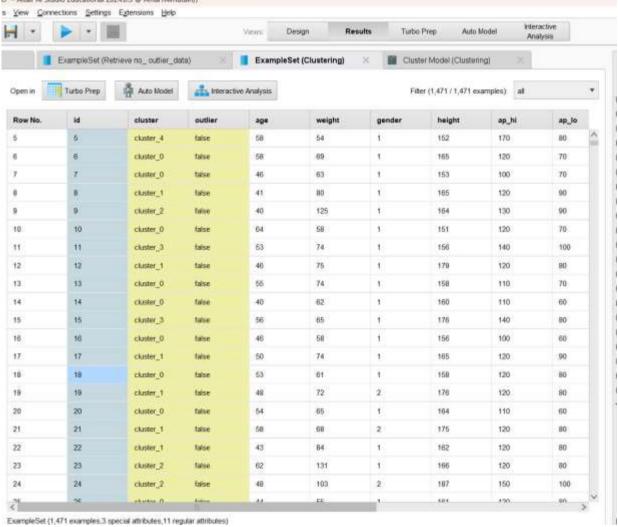




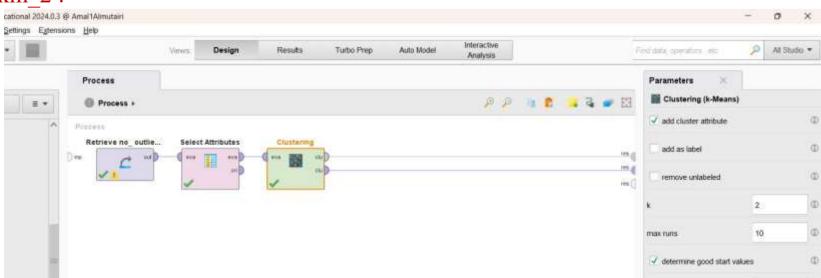
4. Try to find a combination different from the above combinations that gives the best results.

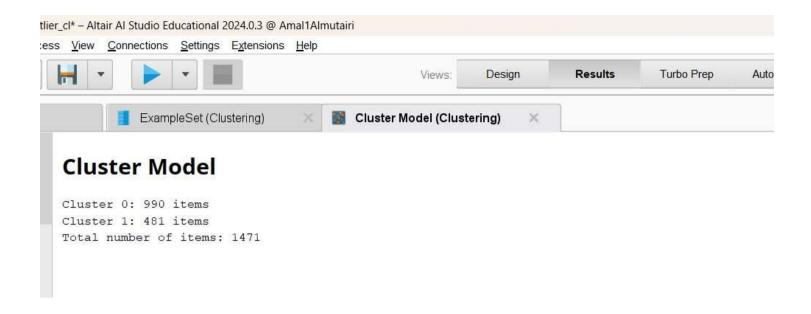
<b>Task 3:</b> 1. Use k-m	eans to generate 4 segmentations with the parameter Number of Clusters described in the table below. You
	gnore the following variables: ID variable, and the target variable.
	Attributes operator in RapidMiner to exclude the ID and target attributes from the analysis. The target attribute in our dataset, ed as it is not relevant to the clustering process



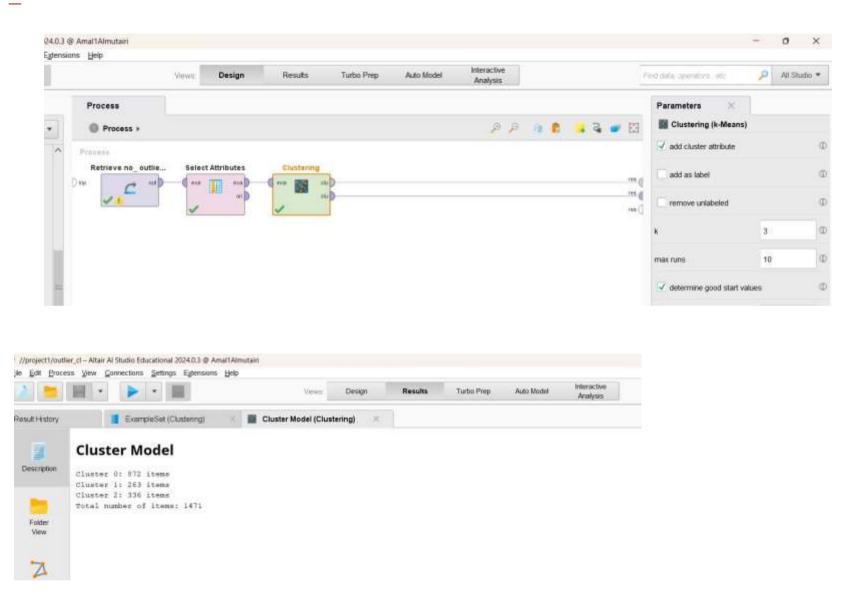


## km 2:





# km\_3:



# $km_4$

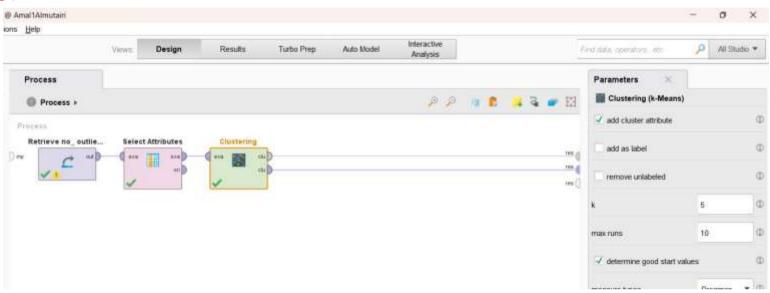
0.3 @ AmalTAlr ensions Help Interactive Views Design All Studio \* Results Turbo Prep Auto Model First mite, openitors...etc. Process Parameters E 💌 🗗 👢 🐧 😰 🔍 🔍 Clustering (k-Means) Process + ✓ add cluster attribute add as label remove unlabeled 4 10 ✓ determine good start values



#### **Cluster Model**

Cluster 0: 719 items Cluster 1: 129 items Cluster 2: 290 items Cluster 3: 333 items Total number of items: 1471

## km 5:





#### **Cluster Model**

Cluster 0: 453 items Cluster 1: 433 items Cluster 2: 162 items Cluster 3: 300 items Cluster 4: 123 items Total number of items: 1471

S	Segmentation Label	Number of Clusters	Cluster Sizes	Size of Smallest Cluster	Outlier (Yes or No)
1.	km_2	2	Cluster 0: 990 Cluster 1: 481	Cluster 1: 481	NO
2.	km_3	3	Cluster 0: 872 Cluster 1: 263 Cluster 2: 336	Cluster 1: 263	NO
3.	km_4	4	Cluster 0: 719 Cluster 1: 129 Cluster 2: 290 Cluster 3: 333	Cluster 1: 129	Yes
4.	km_5	5	Cluster 0: 453 Cluster 1: 433 Cluster 2: 162 Cluster 3: 300 Cluster 4: 123	Cluster 4: 123	Yes

2. Describe the clusters in segmentation km 3 using the descriptive statistics from the results.

Cluster 0: Contains 872 observations, the largest group.

- Age: The mean age is 52.525, making this group middle-aged.
- Weight: The mean weight is 67.655, the lowest among the groups, indicating that this group is lighter.
- Gender: The mean value for gender is 1.334, indicating that this group is mostly female.
- Height: The mean height is 164.347 cm, shorter compared to Group 1 but close to Group 2.
- Blood pressure:
  - ☐ Systolic (ap\_hi): 117.366, the lowest among the groups, indicating better blood pressure health ☐ Diastolic (ap\_lo): 77.462, also the lowest, indicating good cardiovascular health.
- Cholesterol and glucose: Both values are low (cholesterol: 1.258, glucose: 1.164), indicating healthier metabolic conditions.
- Smoking: Very low at 0.078.
- Alcohol consumption: Very low at 0.052.
- Physical activity: High at 0.776, indicating that this group is active.

#### Cluster 1: Contains 263 observations, the smallest group.

- Age: Mean age 53.388, slightly older than Group 0.
- Weight: Mean weight 94.859, the heaviest of the groups, indicating a high BMI.
- Gender: The mean value for gender is 1.502, closer to 1.5, indicating that there are females and males, but the proportion of males is slightly higher than females.
- Height: Mean height 167.707 cm, the tallest of the groups.
- Blood pressure:
  - ☐ Systolic (ap\_hi): 128.384, higher than group 0 but lower than group 2. ☐ Diastolic (ap\_lo): 82.856, average.
- Cholesterol and glucose: slightly higher (cholesterol: 1.586, glucose: 1.308) compared to group 0, showing early signs of metabolic risk.
- Smoking: slightly higher at 0.148.
- Alcohol consumption: average at 0.091.
- Physical activity: high at 0.810, similar to group 2.

#### Cluster 2: Contains 336 observations, middle sized group.

- Age: The average age is 55.357, making this group the oldest.
- Weight: The mean weight is 73.765, higher than group 0 but lower than group 1.
- Gender: The mean gender value (1.381) shows that the group includes both males and females, with a higher proportion of females.
- Height: The mean height is 163.583 cm, the shortest among the groups.
- Blood pressure:
  - Systolic (ap\_hi): 149.601, the highest, indicating a high prevalence of hypertension.
  - Diastolic (ap\_lo): 90.926, also the highest, a potential indicator of cardiovascular risk.
- Cholesterol and glucose: the highest levels (cholesterol: 1.554, glucose: 1.310), reflecting a high metabolic risk.
- Smoking: slightly higher at 0.119.
- Alcohol consumption: Low at 0.068.
- Physical activity: High at 0.812, similar to the other groups.
- 3. If the domain expert was interested in evaluating outlier clusters, which segmentations (i.e., sets of clusters) would you provide to the expert for evaluation? Provide justification for your answer. You may assume that a cluster is an outlier if it contains less than 10% of the observations.

To evaluate outlier clusters, We will provide the following segmentation to the domain expert:

Segmentations with Outliers: km 4

(4 clusters):

Cluster 1 contains 129 observations, which is less than 10% of the total (1471 observations).

Justification: 10% of 1471 is approximately equal to 147 observations. Cluster 1 in km\_4 contains less than this threshold, making it eligible to be an outlier.

#### km\_5 (5 clusters):

Cluster 4 contains 123 observations, which is less than 10% of the total (1471 observations).

Justification: Cluster 4 in km\_5 also falls below the 10% threshold and is considered an outlier.

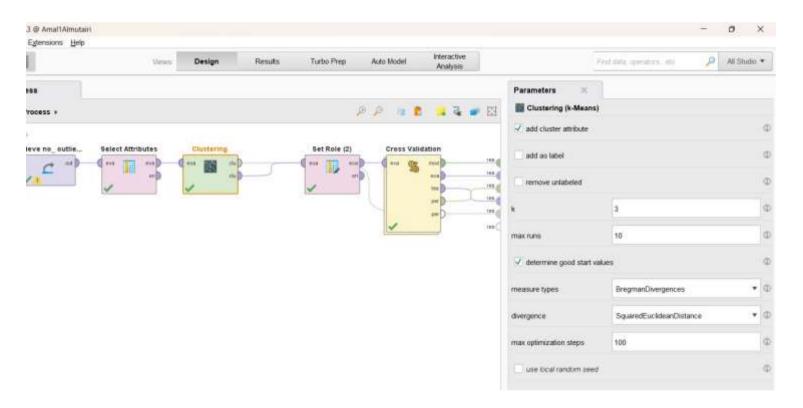
#### Excluded segmentation:

km\_2 (2 clusters): Both clusters contain more than 10% of the total observations. km\_3 (3 clusters): All clusters contain more than 10% of the total observations.

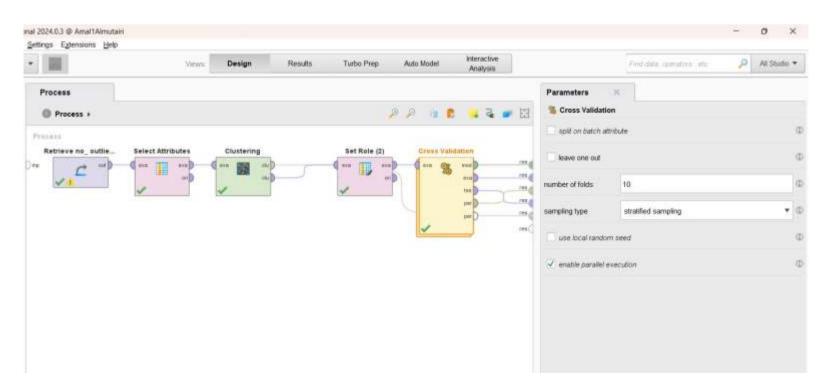
A domain expert should evaluate km\_4 and km\_5, because they contain clusters that meet the outlier criteria based on the 10% threshold.

4. For each cluster in km\_3, generate 3 DTs (i.e., Information Gain, Gini, Gain Ratio) & select the DT that has the best test accuracy & good simplicity. Assume that the weights are: 0.60 for Accuracy & 0.40 for Simplicity. Also assume that there are no thresholds on these measures. Compare these segmentations based on the associated DTs. Note that for each of these DTs, the target variable is the Cluster ID.

We used the km 3 segmentation to generate 3 Decision Trees



We specified 10 folds in Cross Validation and also used the Stratified method.



We calculated the smallest cluster, which is Cluster 1 with a size of 263 and then multiplied it by 0.3 to get the number of observations per leaf, and multiplied the leaf size by 2 to get the minimal partition size. These values were applied to all the trees.

### The Frist Decision Tree is Information Gain:

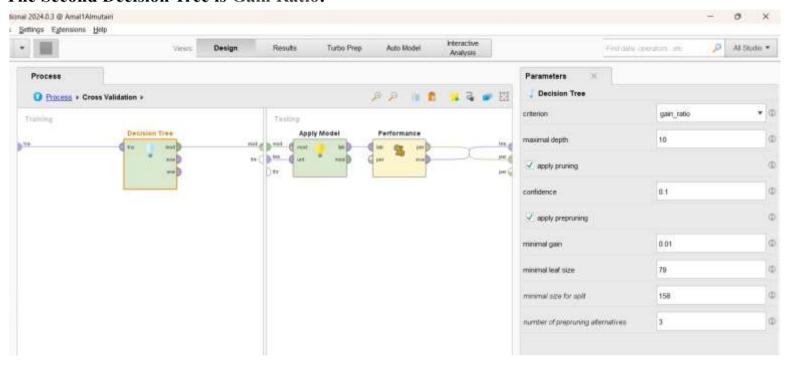
#### **Confusion Matrix Table:**



#### accuracy: 92.93% +/- 2.64% (micro average: 92.93%)

	true cluster_1	true cluster_0	true cluster_2	class precision
pred. cluster_1	247	32	42	76.95%
pred. cluster_0	12	839	13	97.11%
pred. cluster_2	4	1	281	98.25%
class recall	93.92%	96.22%	83.63%	

#### The Second Decision Tree is Gain Ratio:



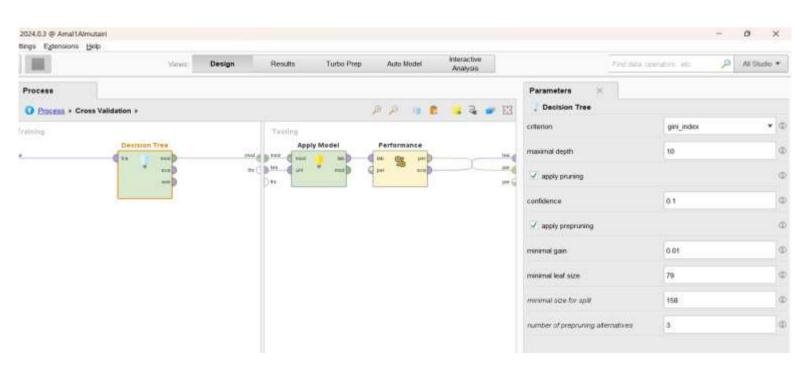


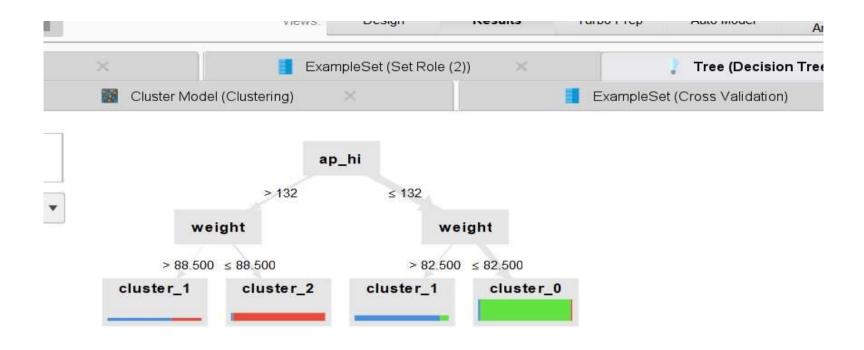
### **The Confusion Matrix Table:**

#### accuracy: 91.98% +/- 2.64% (micro average: 91.98%)

	true cluster_1	true cluster_0	true cluster_2	class precision
ored. cluster_1	186	16	11	87.32%
pred. cluster_0	32	855	13	95.00%
pred. cluster_2	45	1	312	87.15%
class recall	70.72%	98.05%	92.86%	

## The third Decision Tree is Gini:





#### **The Confusion Matrix Table:**

accuracy: 92.32% +/- 2.65% (micro average: 92.32%)

	true cluster_1	true cluster_0	true cluster_2	class precision
pred. cluster_1	199	27	8	85.04%
pred. cluster_0	19	844	13	96.35%
pred. cluster_2	45	1	315	87.26%
class recall	75.67%	96.79%	93.75%	

### combination function (overall score )=0.60\*Accuracy+0.40\*Simplicity

	Pe	rformance Me	Overall Score		
DT	Accuracy		Simplicity		
Label/Descripti on			No of		
	% Error	Score	Leaves	Score	
1-Information					
Gain	1-0.9293=0.0707	92.93%	4	1	0.60*0.9293+0.40*1=95758
2-Gain Ratio	1-0.9198=0.0802	91.98%	4	1	0.60*0.9198+0.40*1=0.95188
3-Gini	1-0.9232=0.0768	92.32%	4	1	0.60*0.9232+0.40*1=0.95392

We assumed in Task 2 that the ideal DT has 3-7 leaves (IdealBot = 3, IdealTop = 7). DTs with  $\leq 2$  or  $\geq 14$  leaves are considered unacceptable. Since all trees have 4 leaves, which falls within the ideal range, the Simplicity Score is 1 for all.

The formula used is:

ScoreSimplicity(NoOfLeaves) = 1 if IdealBot  $\leq$  NoOfLeaves  $\leq$  IdealTop.

The Information Gain decision tree achieves the highest overall score of 0.95758, making it the most suitable option as it offers the best balance between accuracy and simplicity.

## 5. Based on the results of the best DT, explain the produced clusters:

## Importance of Variables:

Variable	Cluster_0	Cluster_1	Cluster_2	Explanation
ap_hi	≤ 132	> 132	> 132	The patients in Cluster_0 have a systolic blood pressure (ap_hi) of ≤ 132, while those in Cluster_1 and Cluster_2 have

				higher blood pressure (> 132).
weight	≤ 81.500	> 81.500 & ≤ 86.500	> 86.500	Cluster_0 consists of patients with lower weight (≤ 81.500), while Cluster_1 contains patients with moderate weight (between 81.500 and 86.500), and Cluster_2 has patients with a weight greater than 86.500

## **Narrative on Comparison:**

This table explains how patients were grouped based on their ap\_hi (systolic blood pressure) and weight. Cluster\_0 includes patients with both lower blood pressure and weight, while Cluster\_1 covers those with moderate blood pressure and moderate weight. Cluster\_2 represents patients with high blood pressure and higher weight.