

Team Presentation





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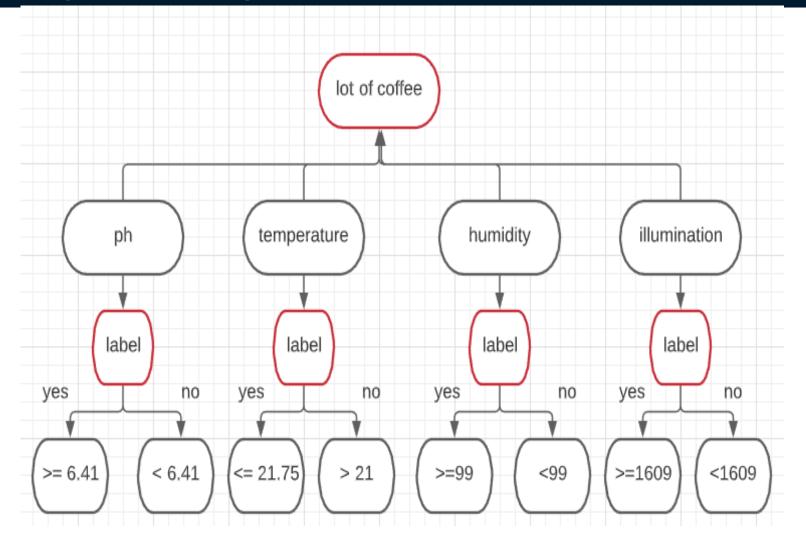
Mauricio Toro





Algorithm Design





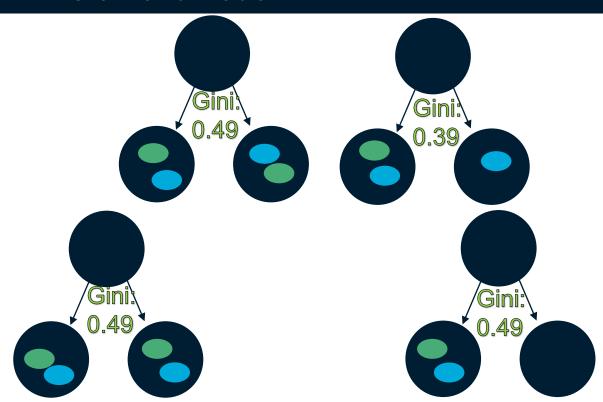
As we know, the main function of the tree is to decide if any of the variables considered that are fundamental for the root, such as temperature, humidity, etc. contain the virus in label and according to the standards of each variable it is indicated whether or not it has a label.

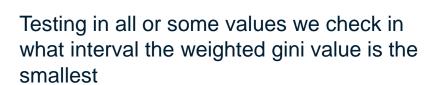
here we use the decision tree id3 to identify that the most influential values are the soil humidity and illuminance.

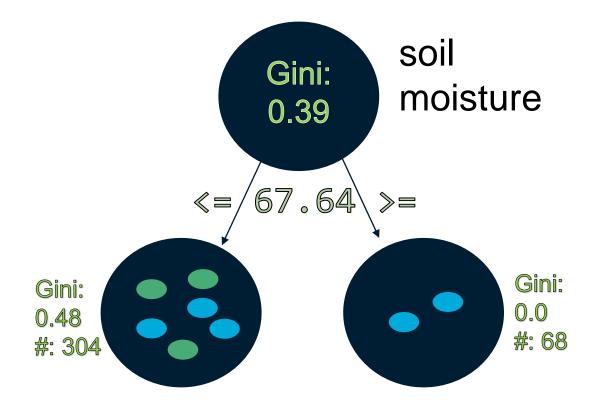


Division of a node









Here we can confirm that when moisture is higher than 68 then its guaranteed that the plant doesn't have the Roya disease and we can separate the tree



Complexity of the Algorithm



	Time complexity	Memory complexity
Train the decision tree	O(N*M)	O(N*M)
Validate the decision tree	O(N)	O(N)
Reader.reader()	O(N)	
Trainer.main()	O(N*M)	
Trainer.trainer()	O(N*M)	
Tree.main()	O(N)	
Tree.tree()	O(N)	

The time and memory complexity of our algorithm is performed with CART. The n and m terms are representations of the complexity of the algorithm.

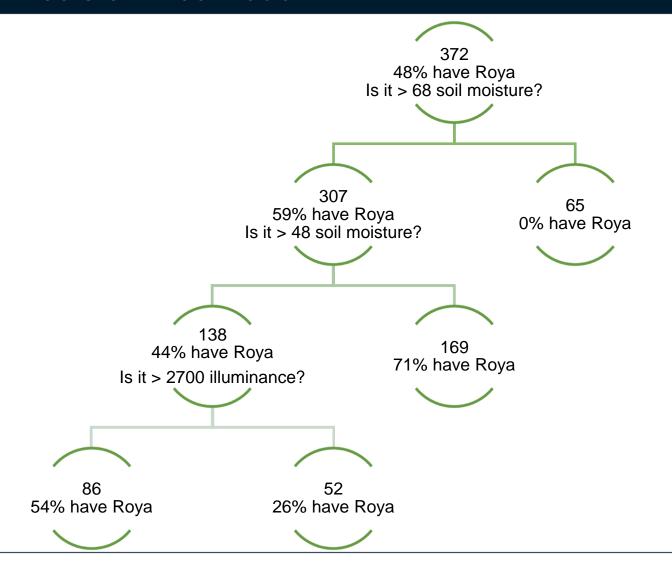
- . N is the amount of data in the file.
- . M is the number of times the cycle of searching for the best division number is repeated.





Decision Tree Model





Relevant characteristics:



Soil moisture



Illuminance

The CART decision tree, it tries multiple values and selects the one that leaves the purest node possible, then it divides the impure one



Evaluation Metrics

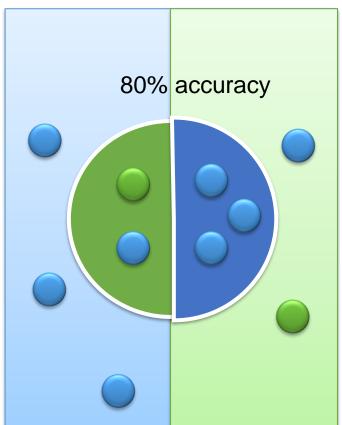


Precision, the percentage of values classified correctly in the field

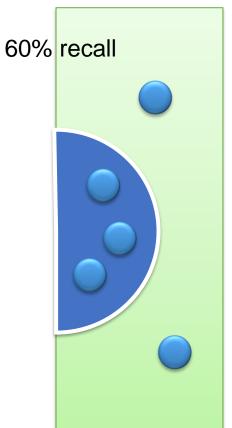
66% precision

of in

Accuracy, the percentage of values classified accordingly



Recall, the percentage of values classified correctly from the whole data





Evaluation Metrics



	Data set 1	Data set 2	Data set 3
Accuracy	85%	81%	62%
Precision	90%	66%	52%
Recall	96%	80%	84%

Evaluation metrics obtained with the training data sets of size 300, 600 and 900





Time and memory consumption



