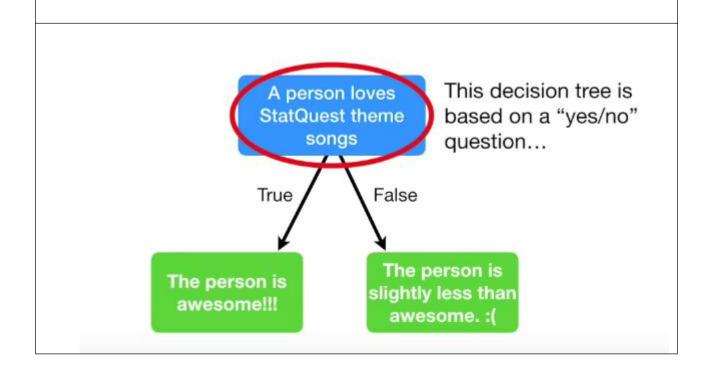
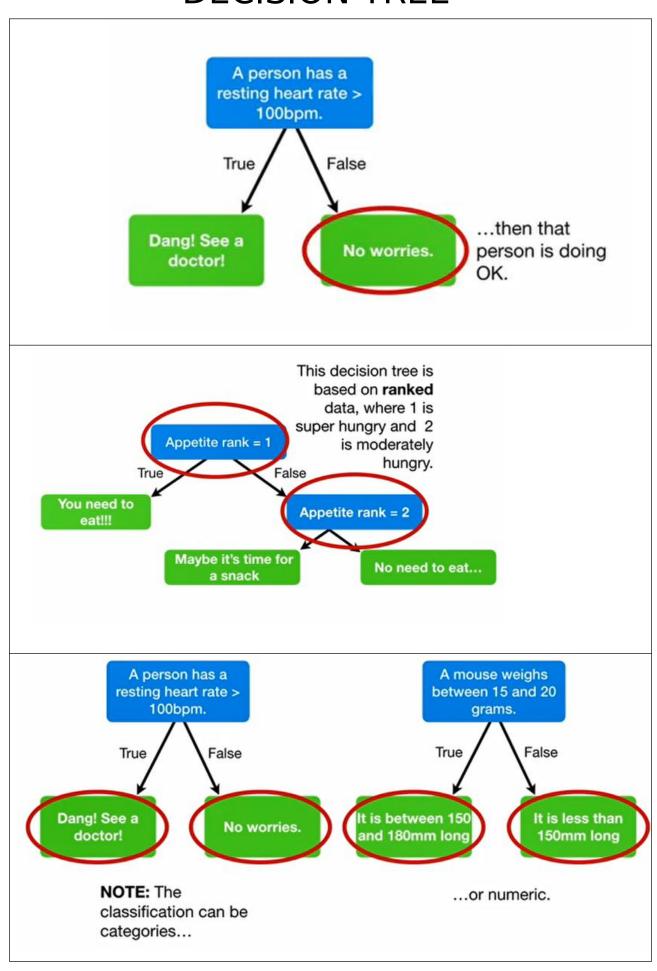
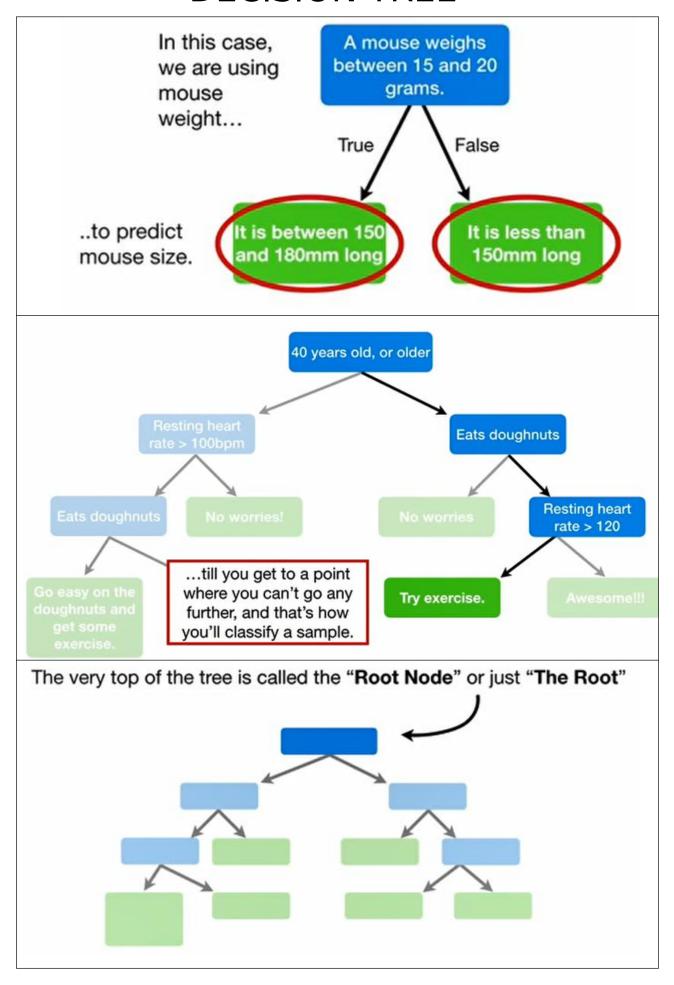
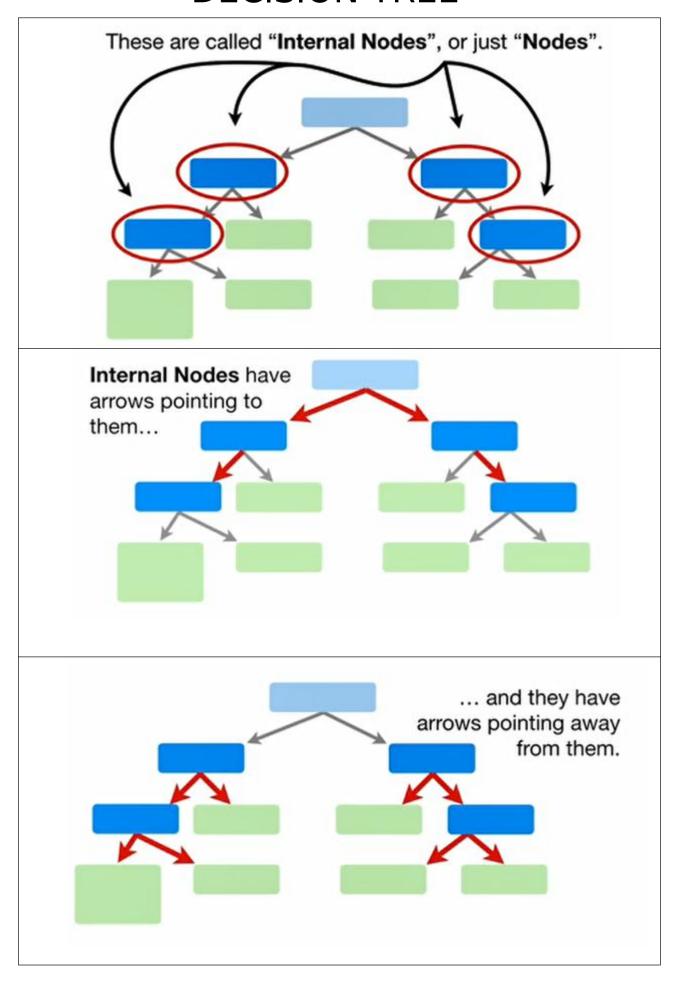
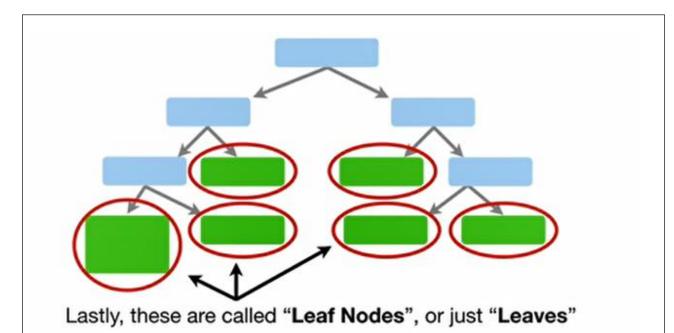
# Big Data Lab Decision tree











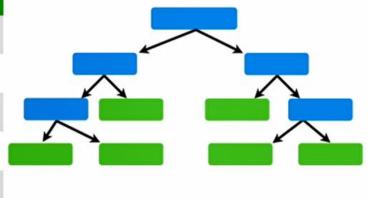
Leaf Nodes have arrows pointing to them...

...but there are no arrows pointing away from them.

Now we are ready to talk about how to go from a raw table of data...

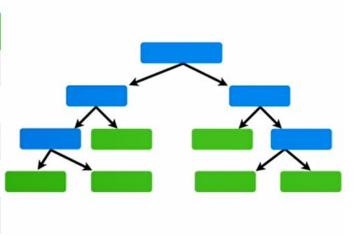
...to a decision tree!!!

Chest Pain	Good Blood Circulation	Blocked Arteries	Heart Disease
No	No	No	No
Yes	Yes	Yes	Yes
Yes	Yes	No	No
Yes	No	???	Yes
etc	etc	etc	etc

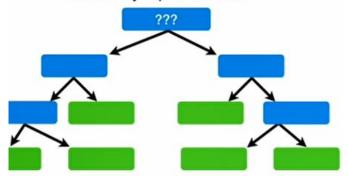


In this example, we want to create a tree that uses chest pain, good blood circulation and blocked artery status to predict...

Chest Pain	Good Blood Circulation	Blocked Arteries	Heart Disease
No	No	No	No
Yes	Yes	Yes	Yes
Yes	Yes	No	No
Yes	No	777	Yes
etc	etc	etc	etc

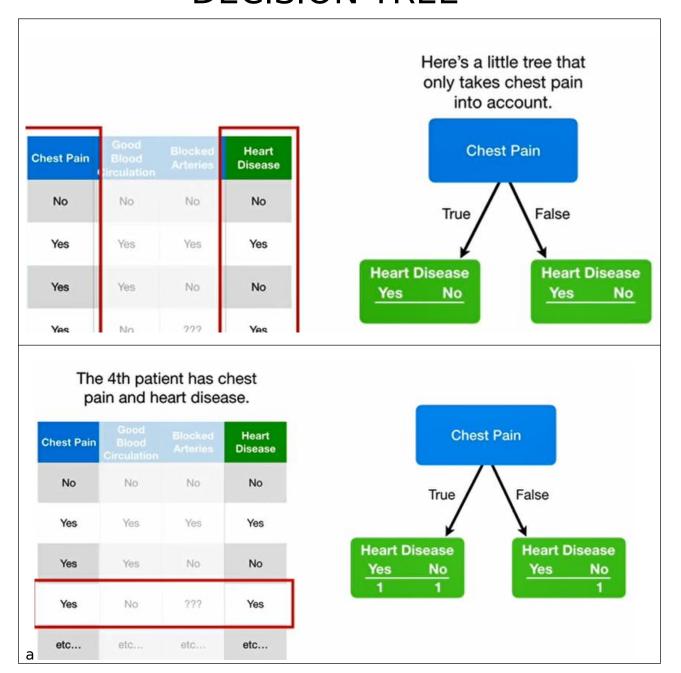


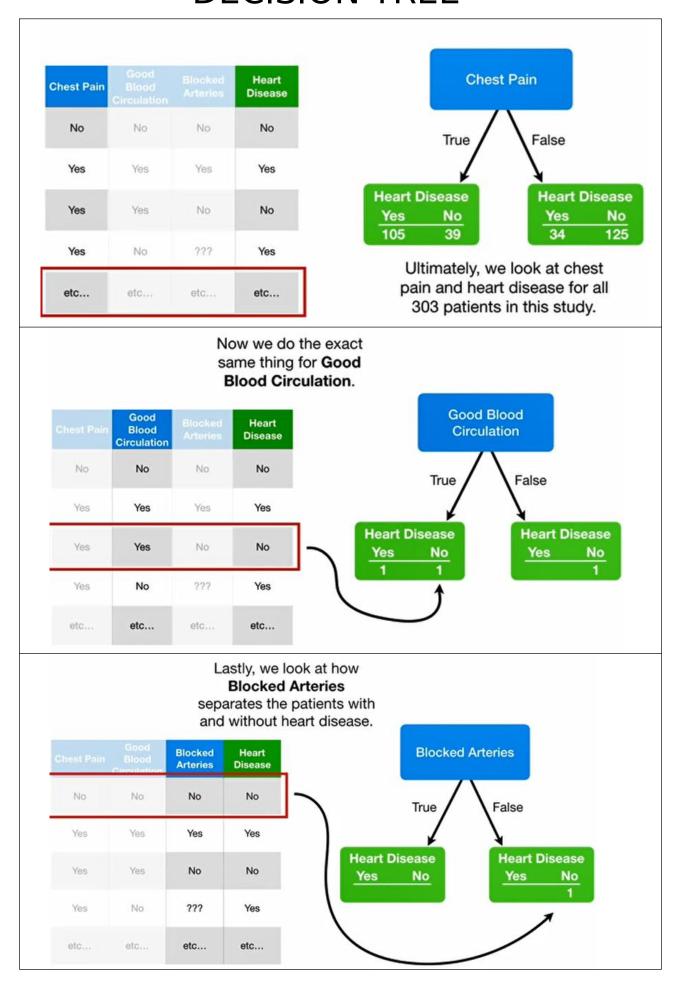
The first thing we want to know is whether Chest Pain, Good Blood Circulation or Blocked Arteries should be at the very top of our tree.

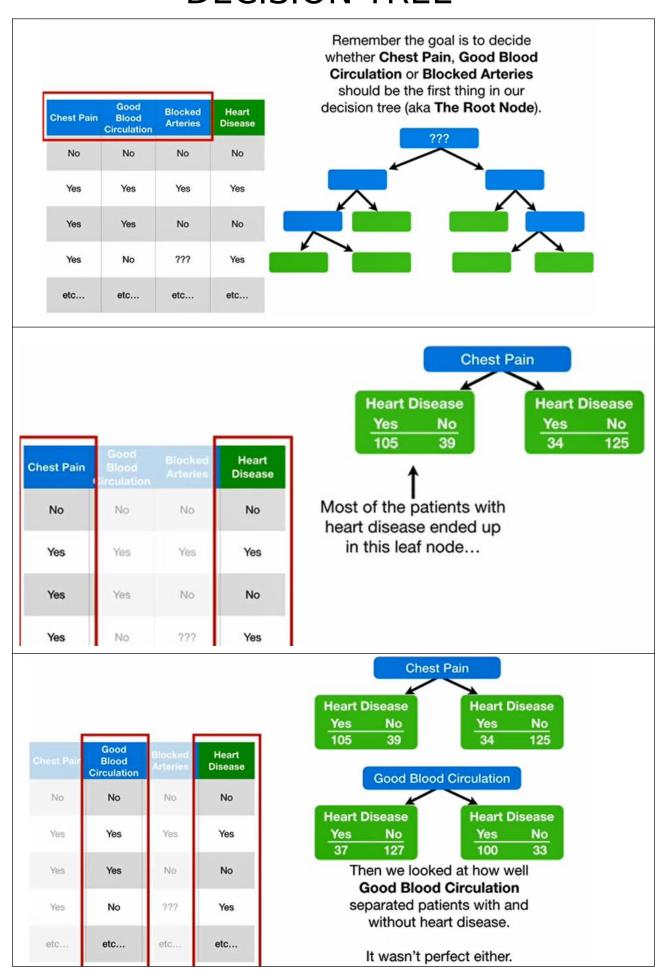


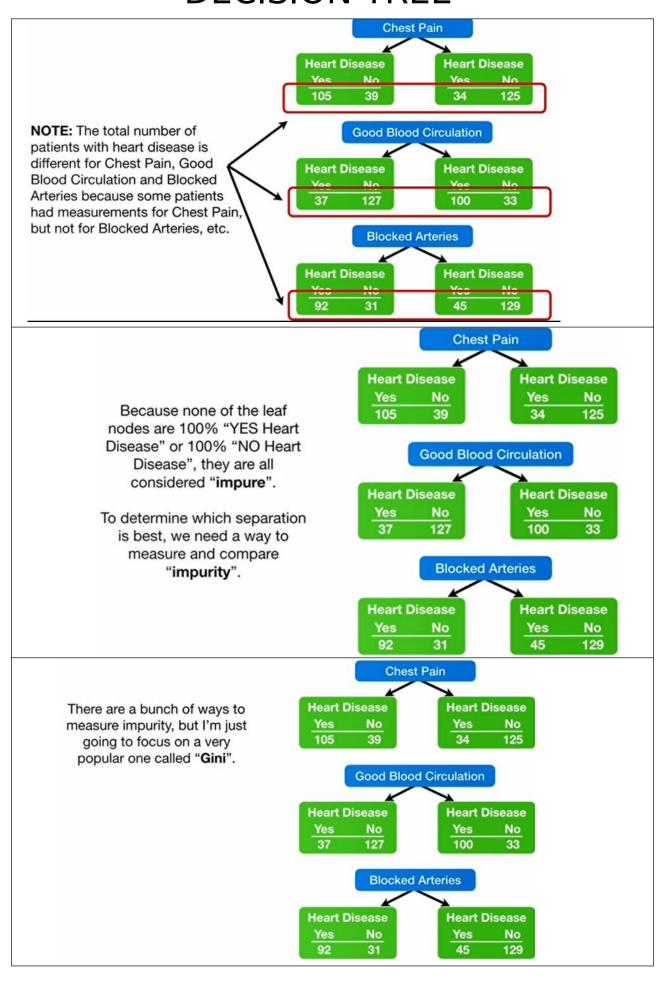
We start by looking at how well **Chest Pain** alone predicts heart disease...

Chest Pain	Good Blood Sirculation	Blocked Arteries	Heart Disease
No	No	No	No
Yes	Yes	Yes	Yes
Yes	Yes	No	No
Yes	No	777	Yes
etc	etc	etc	etc







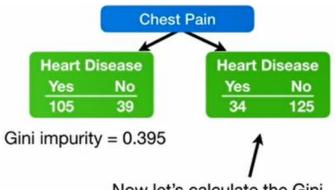




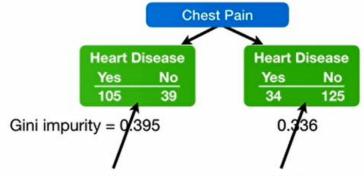
For this leaf, the Gini impurity = 1 - (the probability of "yes")2 - (the probability of "no")2

$$= 1 - (\frac{105}{105 + 39})^2 - (\frac{39}{105 + 39})^2$$

$$= 0.395$$



Now let's calculate the Gini impurity for this leaf node...



Because this leaf node ... and this leaf node represents 144 patients... represents 159 patients...

Thus, the total Gini impurity for using Chest Pain to separate patients with and without heart disease is the weighted average of the leaf node impurities.



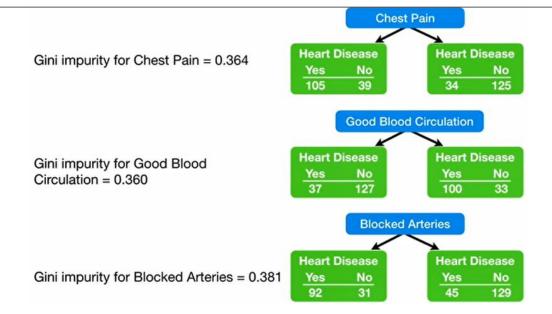
Gini impurity for Chest Pain = weighted average of Gini impurities for the leaf nodes

$$=(\frac{144}{144+159})\ 0.395 + (\frac{159}{144+159})\ 0.336$$

$$= 0.364$$

**Chest Pain Heart Disease Heart Disease** Gini impurity for Chest Pain = 0.364 Yes Yes No No 105 39 34 125 **Good Blood Circulation Heart Disease Heart Disease** Yes No Yes No 37 127 100 33 **Blocked Arteries Heart Disease Heart Disease** Yes No Yes No 92 45 31 129

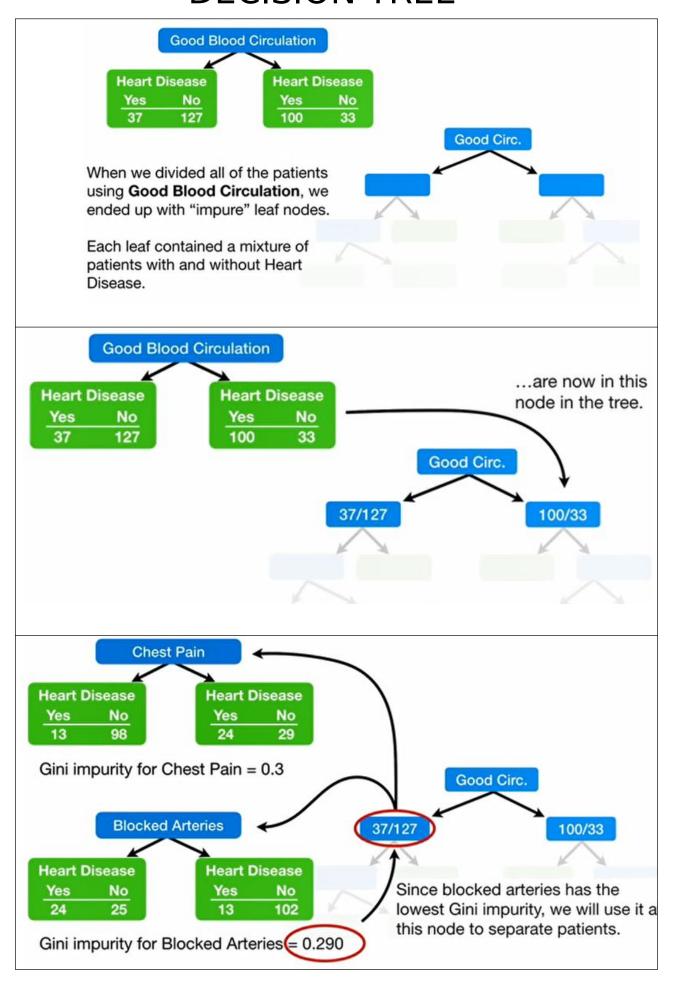


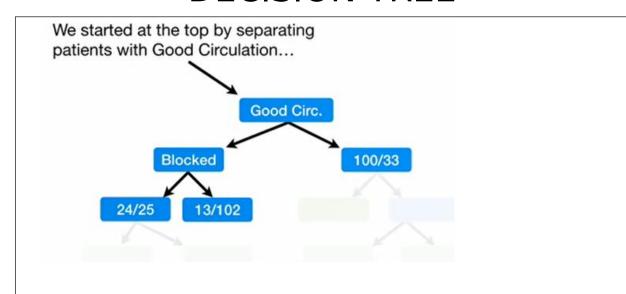


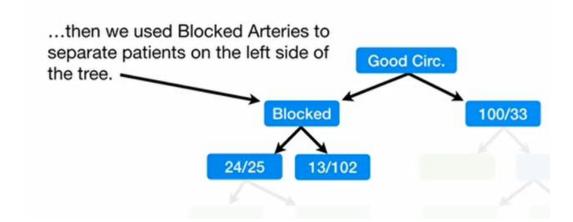
Gini impurity for Chest Pain = 0.364

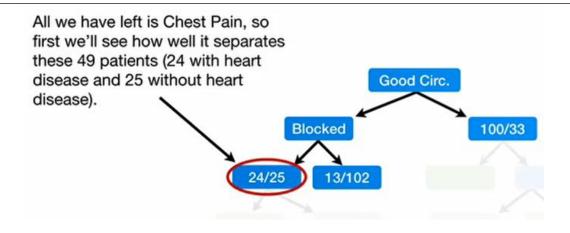
Gini impurity for Good Blood Circulation = 0.360 Good Blood Circulation has the lowest impurity (it separates patients with and without heart disease the best)...

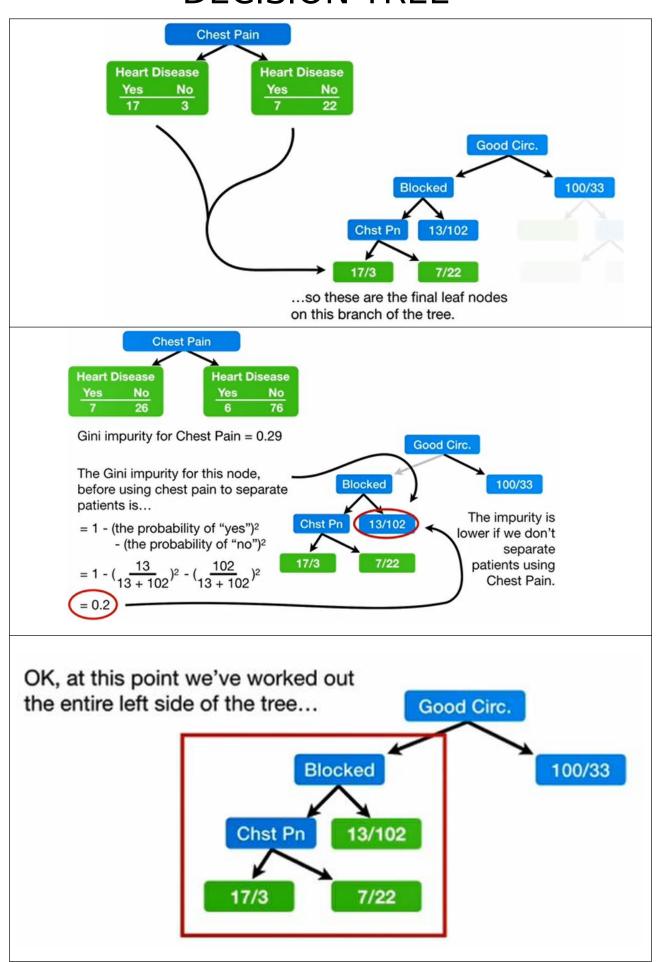
Gini impurity for Blocked Arteries = 0.381





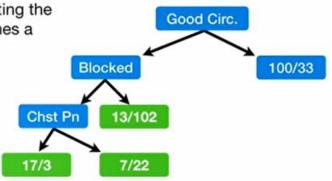






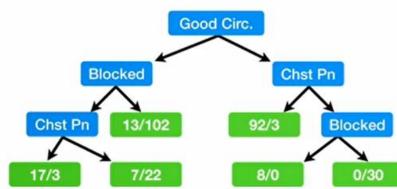
The good news is that we follow the exact same steps as we did on the left side:

- 1) Calculate all of the Gini impurity scores.
- If the node itself has the lowest score, than there is no point in separating the patients any more and it becomes a leaf node.
- If separating the data results in an improvement, than pick the separation with the lowest impurity value.



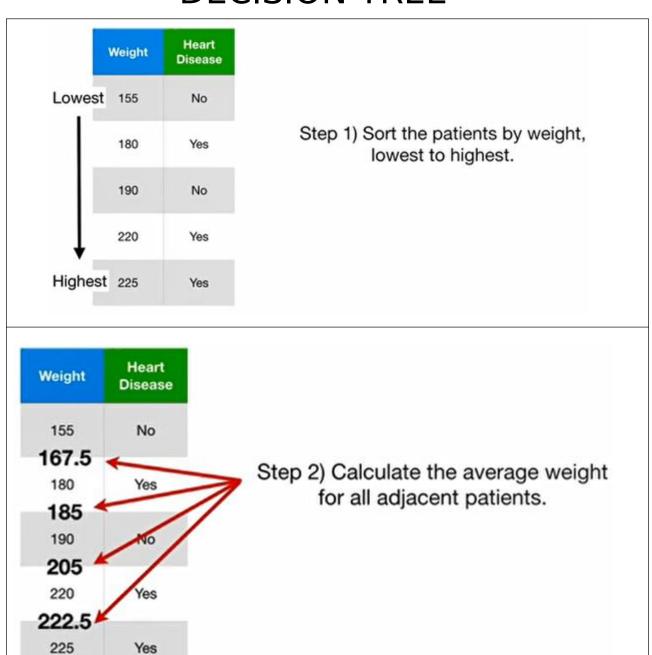
So far we've seen how to build a tree with "yes/no" questions at each step...

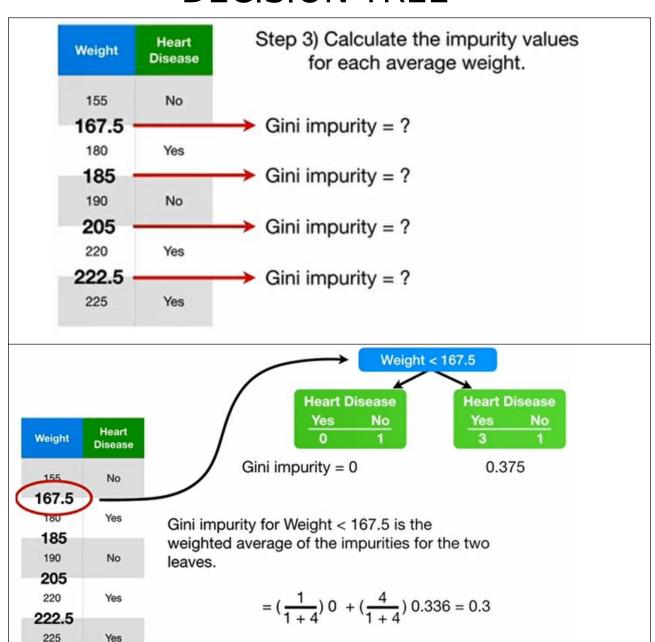
...but what if we have numeric data, like patient weight?

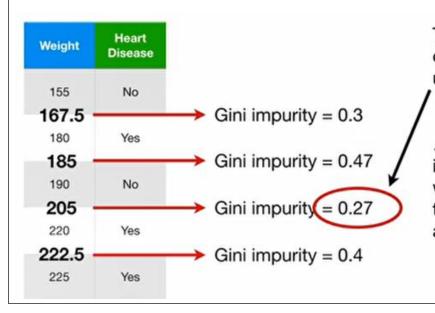


Weight	Heart Disease	
220	Yes	
180	Yes	
225	Yes	
190	No	
155	No	

Imagine if this were our data...







The lowest impurity occurs when we separate using weight < 205...

...so this is the cutoff and impurity value we will use when we compare weight to chest pain or blocked arteries.

