# **Decision Tree**

Regression

# Regression tree vs classification tree

Low pressu re	High Tempe rature	High humidi ty	Wind Speed	Rain
No	No	No	10.0	No
Yes	Yes	Yes	30.0	Yes
Yes	Yes	No	20.0	No
Yes	No	Yes	50.0	No
No	No	Yes	70.0	Yes

classification tree



Discrete data

Low pressu re	High Tempe rature	High humidi ty	Wind Speed	Rain amoun t
No	No	No	10.0	0.1
Yes	Yes	Yes	30.0	0.3
Yes	Yes	No	20.0	1.0
Yes	No	Yes	50.0	2.1
No	No	Yes	70.0	0.7

**Regression tree** 



Continuous data

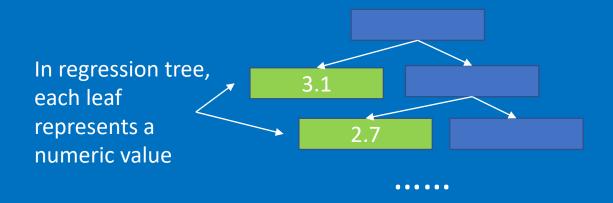
# Regression tree vs classification tree

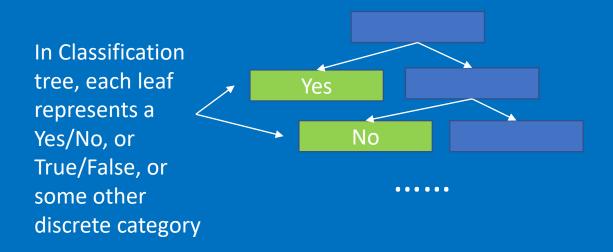
Regression tree is a type of decision tree

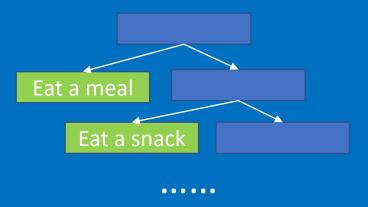


# Regression tree vs classification tree

Regression tree is a type of decision tree







Similar to classification tree, regression tree is built from top down

Let's assume that we have only one predictors, which is drug dosage, and the target is drug effective ratio

Dosage	Drug effect.
10	58
20	60
35	57
5	44
	0 0.0

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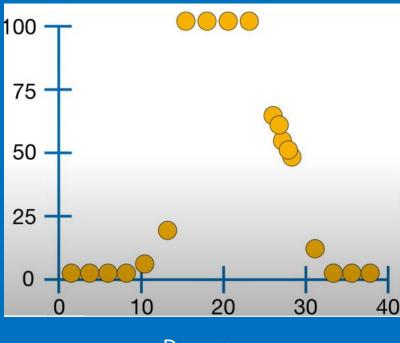
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Dosage	Drug effect.
10	58
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	***

We can plot it out as



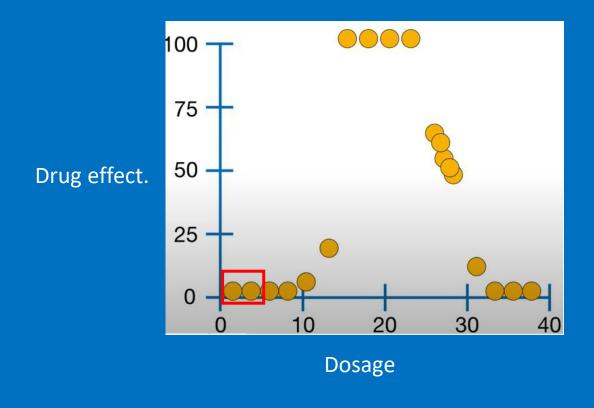
Drug effect.



Dosage

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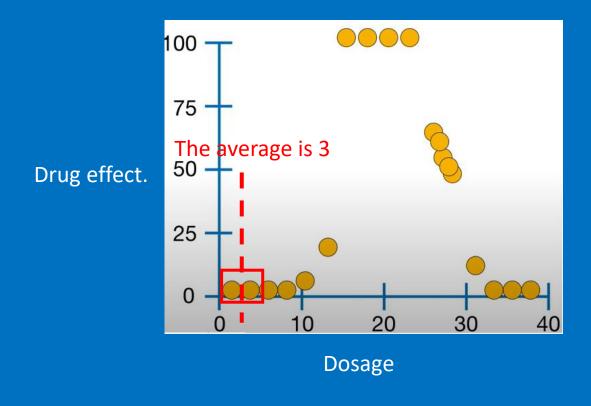
Let's assume that we have only one predictors, which is drug dosage, and the target is drug effective ratio



Step 1: let's look at the first two data points (smallest two values)

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Let's assume that we have only one predictors, which is drug dosage, and the target is drug effective ratio



Step 1: let's look at the first two data points (smallest two values), The average is 3.

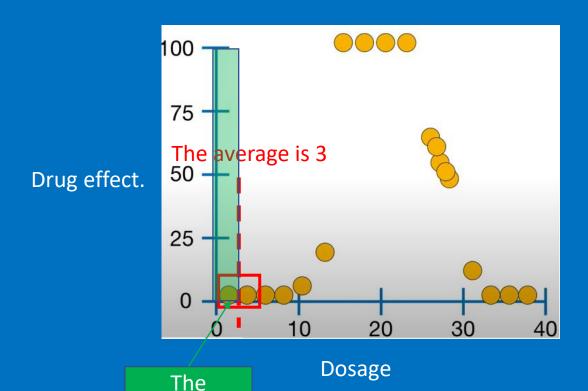
average

drug

effect. is 0

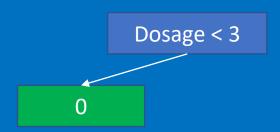
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Let's assume that we have only one predictors, which is drug dosage, and the target is drug effective ratio



Step 1: let's look at the first two data points (smallest two values), The average is 3.

Now we can build a simple tree based on if the dosage is smaller than 3



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Let's assume that we have only one predictors, which is drug dosage, and the target is drug effective ratio

100 75 The average is 3 50 Drug effect. 25 20 30 Dosage The average drug effect. Is 38.8

Step 1: let's look at the first two data points (smallest two values), The average is 3.

Now we can build a simple tree based on if the dosage is smaller than 3



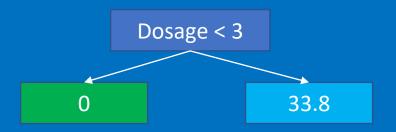
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100 75 The average is 3 50 -Drug effect. 25 20 30 Dosage

Step 1: let's look at the first two data points (smallest two values), The average is 3.

Now we can build a simple tree based on if the dosage is smaller than 3



Step 2: let's estimate how accurate this tree is

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Let's assume that we have only one predictors, which is drug dosage, and the target is drug effective ratio

Drug effect.

The average is 3

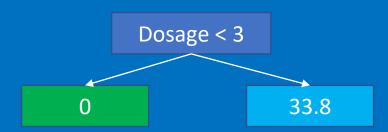
50

25

Dosage

Step 1: let's look at the first two data points (smallest two values), The average is 3.

Now we can build a simple tree based on if the dosage is smaller than 3



Step 2: let's estimate how accurate this tree is

We go through all the points in the graph in the graph with the tree above, the error for this tree is 27468.5

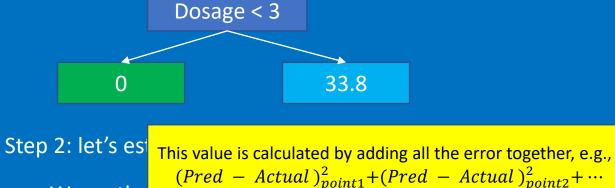
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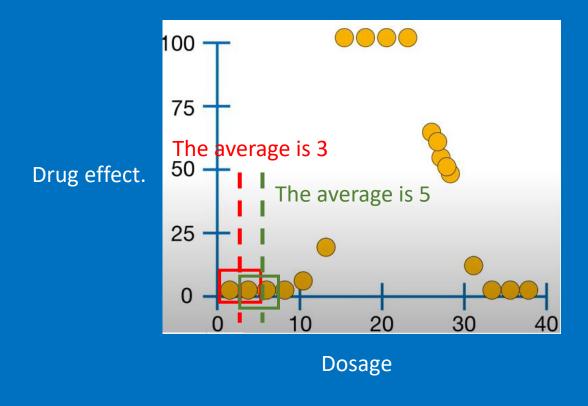
Now we can build a simple tree based on if the dosage is smaller than 3



We go thr graph with  $t^{\prime}$  tree above, the error for this tree is 27468.5

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Let's assume that we have only one predictors, which is drug dosage, and the target is drug effective ratio



Step 3: then we can move to the next 2 points, and use the average there as the new root threshold, which is 5

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Drug effect.

The average is 3

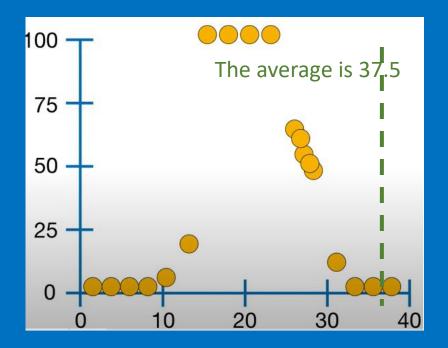
The average is 5

25

Dosage

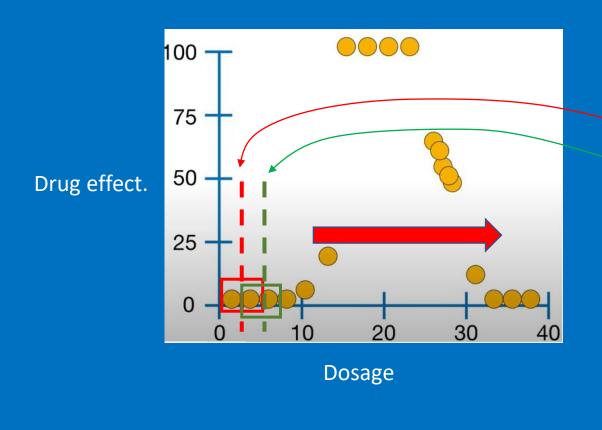
Step 3: then we can move to the next 2 points, and use the average there as the new root threshold, which is 5

Then we can get a new error, which is 23763.2



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By repeating the above steps, we can get a series of root node threshold, and the corresponding errors

#### For example

Threshold 3 27468.5

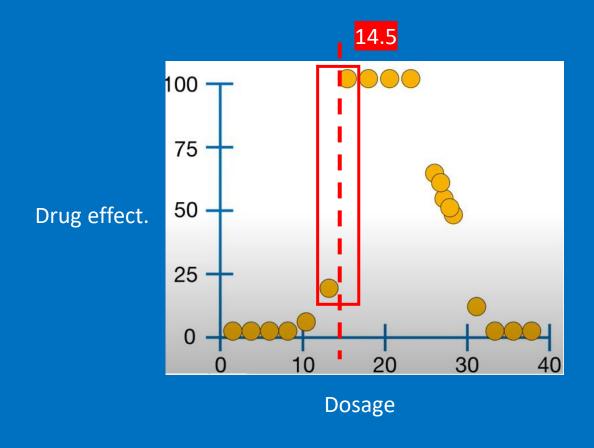
Threshold 5 23763.2

Threshold 8 .....

.....

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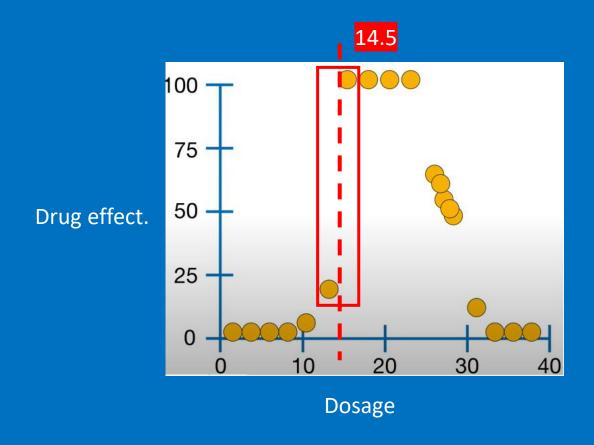


By repeating the above steps, we can get a series of root node threshold, and the corresponding errors

The dosage threshold of 14.5 has the smallest error, and it will be used to build our actual tree

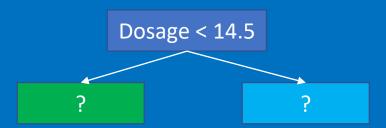
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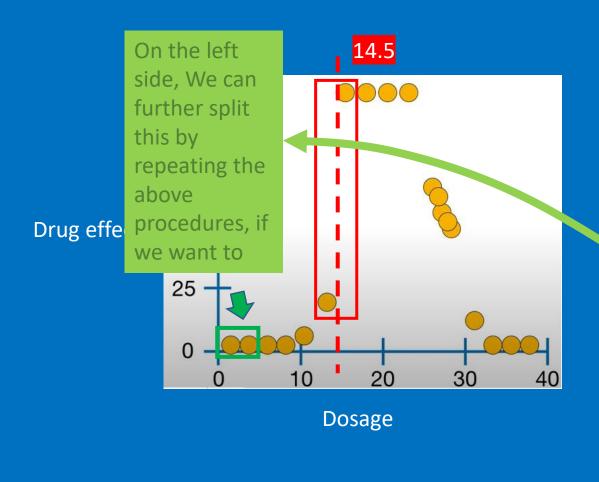
The dosage threshold of 14.5 has the smallest error, and it will be used to build our actual tree



In summary, we split the data into two groups by finding the threshold that give us the smallest sum of squared residuals (errors)

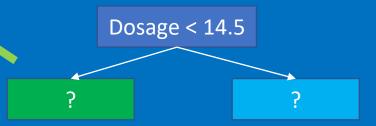
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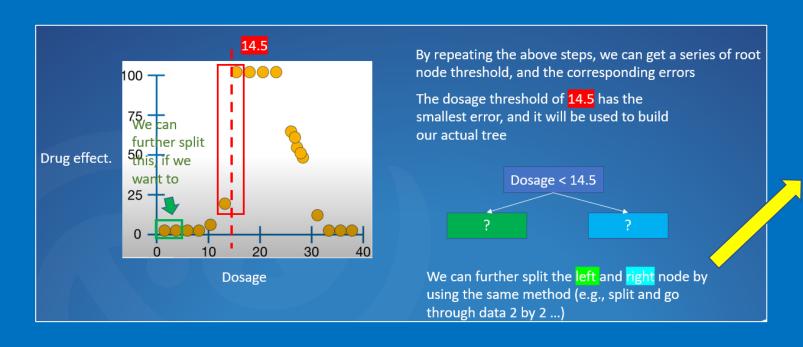
The dosage threshold of 14.5 has the smallest error, and it will be used to build our actual tree



We can further split the left and right node by using the same method (e.g., split and go through data 2 by 2 ...)

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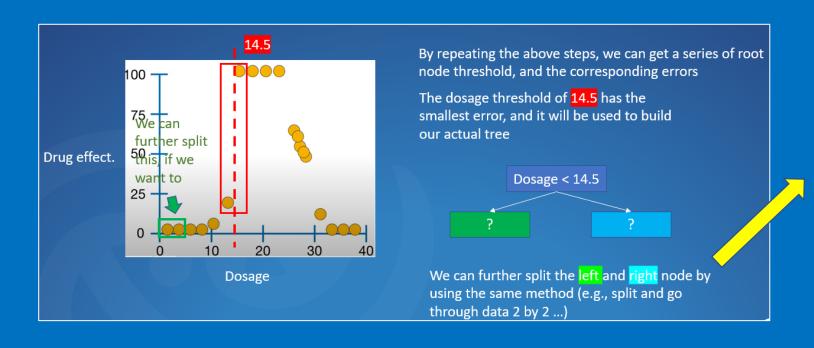
Let's assume that we have only one predictors, which is drug dosage, and the target is drug effective ratio



However, we may not want to do so (in order to avoid over-fitting). Usually we can constrain how many data points still being left and decide if we still want to further split the tree ...

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Let's assume that we have only one predictors, which is drug dosage, and the target is drug effective ratio

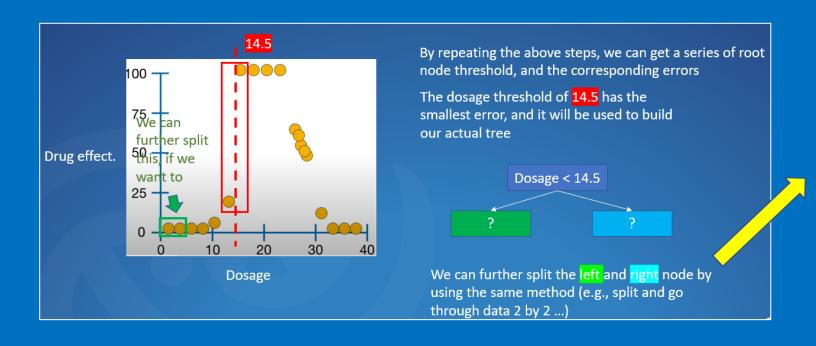


However, we may not want to do so (in order to avoid over-fitting). Usually we can constrain how many datapoints still being left and decide if we still want to further split the tree ...

For example, on the left side of "dosage < 14.5", we still have 6 points

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Let's assume that we have only one predictors, which is drug dosage, and the target is drug effective ratio



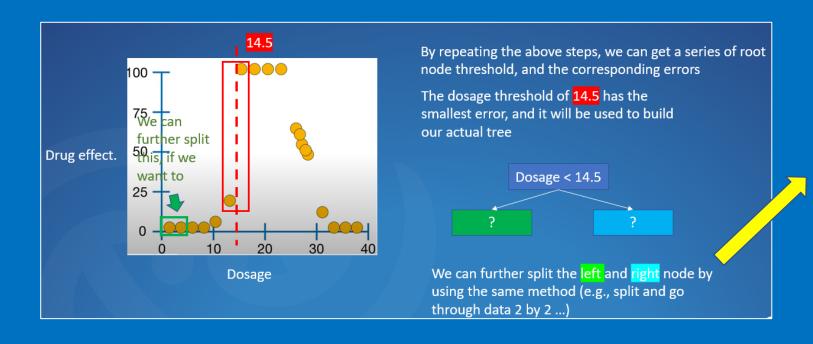
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For example, on the left side of "dosage < 14.5", we still have 6 points

 If the condition says that we stop splitting when the "minimum data points == 6", then we stop splitting

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Let's assume that we have only one predictors, which is drug dosage, and the target is drug effective ratio



However, we may not want to do so (in order to avoid over-fitting). Usually we can constrain how many datapoints still being left and decide if we still want to further split the tree ...

For example, on the left side of "dosage < 14.5", we still have 6 points

- If the condition says that we stop splitting when the "minimum data points == 6", then we stop splitting
- If the condition says that we stop splitting when the "minimum data points == 4", then we can keep splitting further (since we still have more than 4 data points ...) ...

Similar to classification tree, regression tree is built from top down

For multiple predictors, it is the similar process, we basically go through the predictor one by one, and calculate their best threshold, and compare the errors, respectively. Then decide which one to be used as the root node ...

Dosage	Drug effect.
10	58
20	60
35	57
5	44
•••	•••



Dosage		Drug effect.
10	•••	58
20	•••	60
35	•••	57
5	•••	44
•••		