# DATA MINING 2 Gradient Boost

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a.a. 2024/2025



# Gradient Boosting for Regression

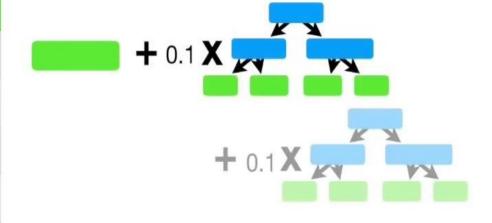
Height (m)	Favorite Color	Gender	Weight (kg)
1.6	Blue	Male	88
1.6	Green	Female	76
1.5	Blue	Female	56
1.8	Red	Male	73
1.5	Green	Male	77
1.4	Blue	Female	57

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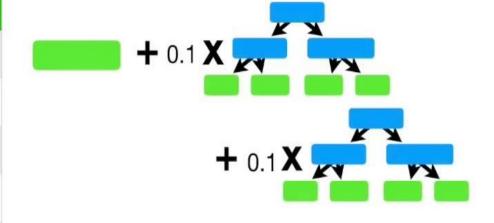
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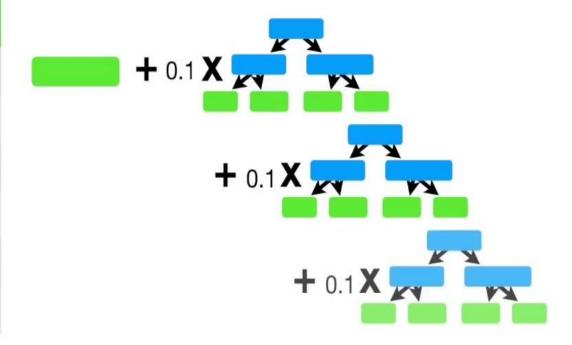
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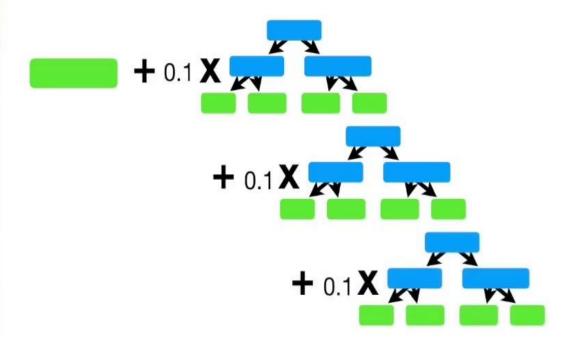
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# Gradient Boost — Example

71.2

		Gender	Weight (kg)
1.6	Blue	Male	88
1.6	Green	Female	76
1.5	Blue	Female	56
1.8	Red	Male	73
1.5	Green	Male	77
1.4	Blue	Female	57

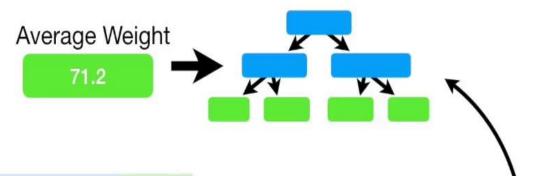
The first thing we do is calculate the average **Weight.** 

Average Weight

71.2

		Gender	Weight (kg)
1.6	Blue	Male	88
1.6	Green	Female	76
1.5	Blue	Female	56
1.8	Red	Male	73
1.5	Green	Male	77
1.4	Blue	Female	57

This is the first attempt at predicting everyone's weight.



1.6	Blue	Male	88
1.6	Green	Female	76
1.5	Blue	Female	56
1.8	Red	Male	73
1.5	Green	Male	77
1.4	Blue	Female	57

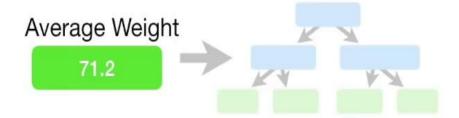
The next thing we do is build a tree based on the errors from the first tree.

Average Weight

71.2

The errors that the previous tree made

1.6	Blue	Male	88
1.6	Green	Female	76
1.5	Blue	Female	56
1.8	Red	Male	73
1.5	Green	Male	77
1.4	Blue	Female	57



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1.6	Blue	Male	88
1.6	Green	Female	76
1.5	Blue	Female	56
1.8	Red	Male	73
1.5	Green	Male	77
1.4	Blue	Female	57

The errors that the previous tree made are the differences between the **Observed Weights** 

(Observed Weight - Predicted Weight)

Average Weight

71.2

		Gender	Weight (kg)
1.6	Blue	Male	88
1.6	Green	Female	76
1.5	Blue	Female	56
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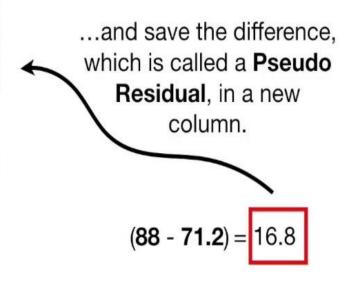
The errors that the previous tree made are the differences between the **Observed Weights** and the **Predicted Weight**, **71.2**.

(Observed Weight - Predicted Weight)

#### Average Weight

71.2

			Weight (kg)	Residual
1.6	Blue	Male	88	16.8
1.6	Green	Female	76	
1.5	Blue	Female	56	
1.8	Red	Male	73	
1.5	Green	Male	77	
1.4	Blue	Female	57	

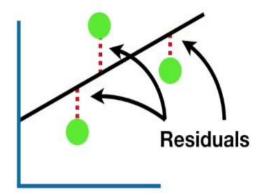


Average Weight

71.2

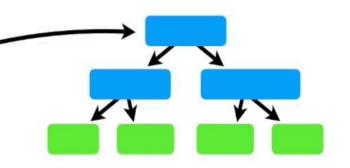
			Weight (kg)	Residual
1.6	Blue	Male	88	16.8
1.6	Green	Female	76	
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1.8	Red	Male	73	
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1.4	Blue	Female	57	

NOTE: The term Pseudo Residual is based on Linear Regression, where the difference between the Observed values and the Predicted values results in Residuals.



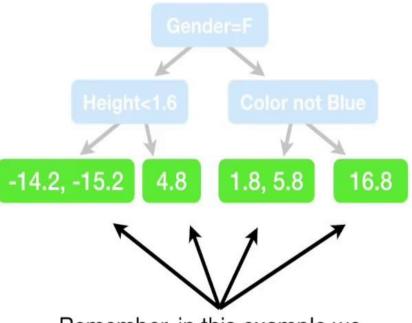
Example

Now we will build a Tree



Height (m)	Favorite Color	Gender	Weight (kg)	Residual
1.6	Blue	Male	88	16.8
1.6	Green	Female	76	4.8
1.5	Blue	Female	56	-15.2
1.8	Red	Male	73	1.8
1.5	Green	Male	77	5.8
1.4	Blue	Female	57	-14.2

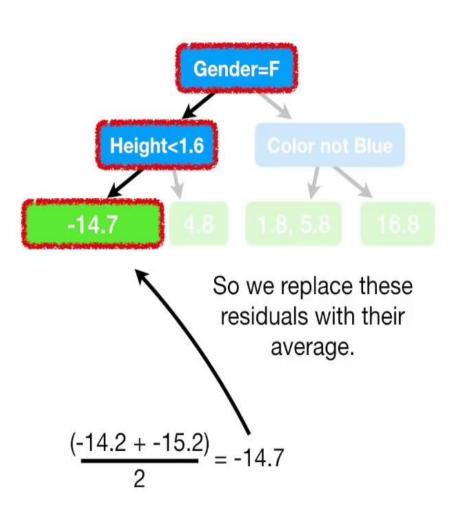
Height (m)			
1.6	Blue	Male	16.8
1.6	Green	Female	4.8
1.5	Blue	Female	-15.2
1.8	Red	Male	1.8
1.5	Green	Male	5.8
1.4	Blue	Female	-14.2



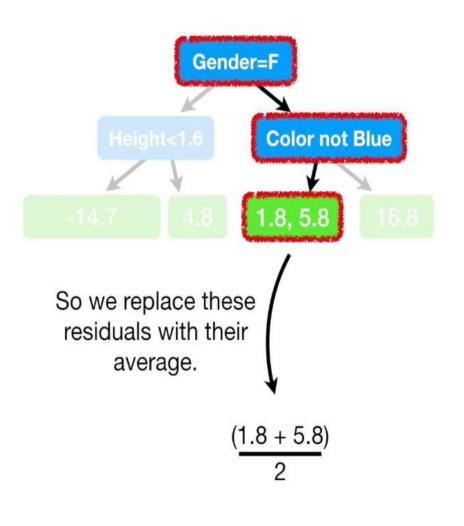
Remember, in this example we are only allowing up to four leaves...

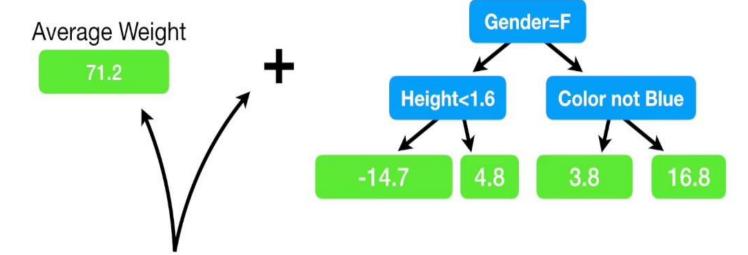
...but when using a larger dataset, it is common to allow anywhere from 8 to 32.

Height (m)	Favorite Color	Gender	Weight (kg)	Residual
1.6	Blue	Male		16.8
1.6	Green	Female		4.8
1.5	Blue	Female		-15.2
1.8	Red	Male		1.8
1.5	Green	Male		5.8
1.4	Blue	Female		-14.2

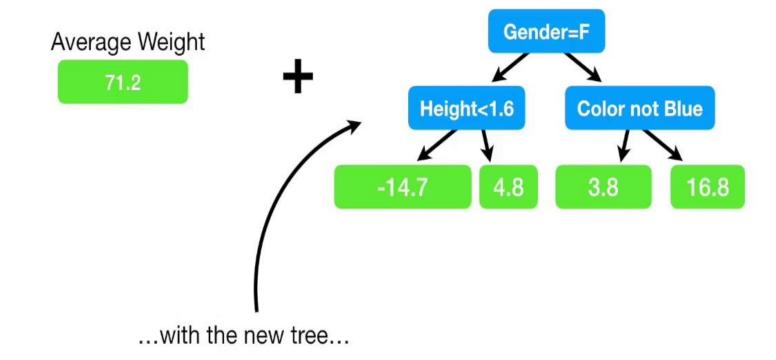


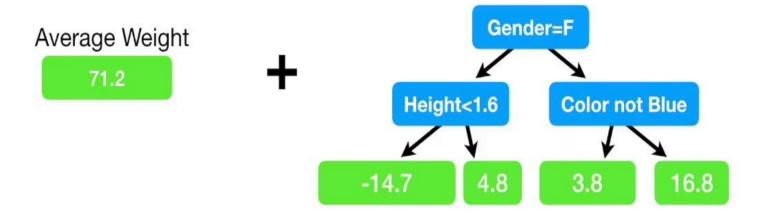
Height (m)	Favorite Color	Gender	Weight (kg)	Residual
1.6	Blue	Male		16.8
1.6	Green	Female		4.8
1.5	Blue	Female		-15.2
1.8	Red	Male		1.8
1.5	Green	Male		5.8
1.4	Blue	Female		-14.2





Now we can now combine the original leaf...



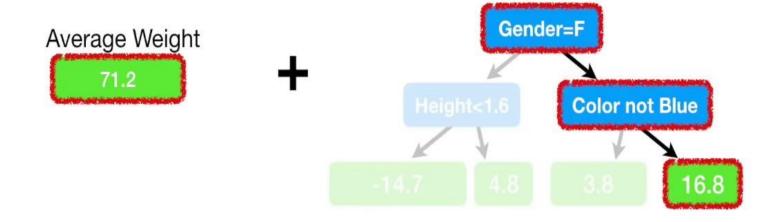


Height	Favorite	Gender	Weight
(m)	Color		(kg)
1.6	Blue	Male	88

...to make a new

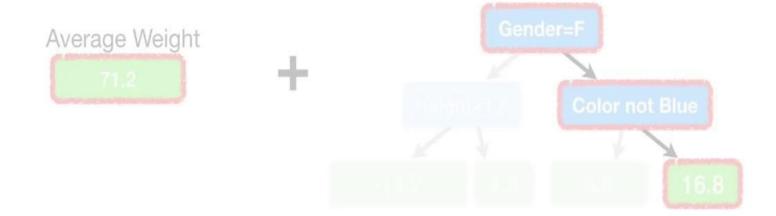
Prediction of an

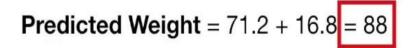
individual's Weight from
the Training Data.



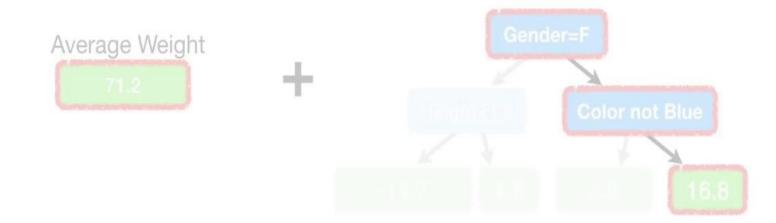
...so the **Predicted Weight** = 71.2 + 16.8 = 88

1.6	Blue	Male	88





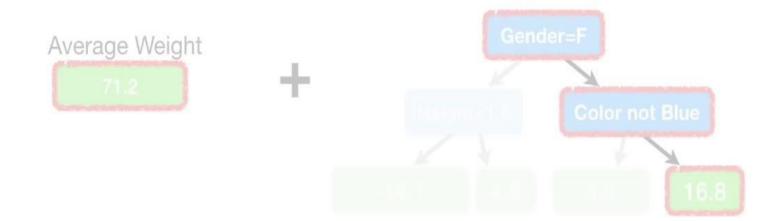
	40.00		Alexander and the	
Height (m)	Favorite Color	Gender	Weight (kg)	Is this awesome???
1.6	Blue	Male	88	<b>←</b>



**Predicted Weight** = 71.2 + 16.8 = 88

Height	Favorite	Gender	Weight
(m)	Color		(kg)
1.6	Blue	Male	88

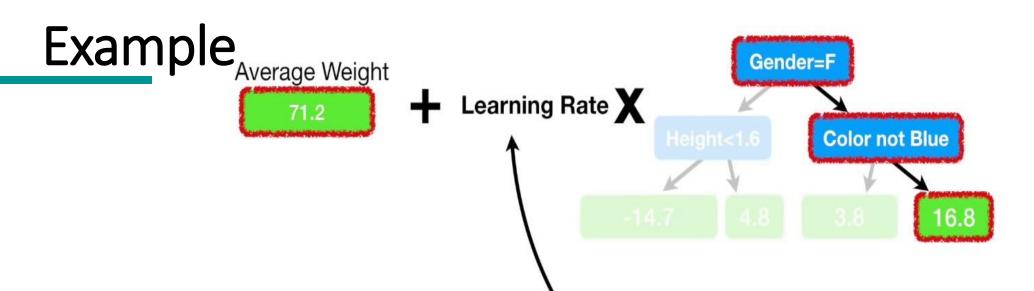
**No.** The model fits the **Training Data** too well.



**Predicted Weight** = 71.2 + 16.8 = 88

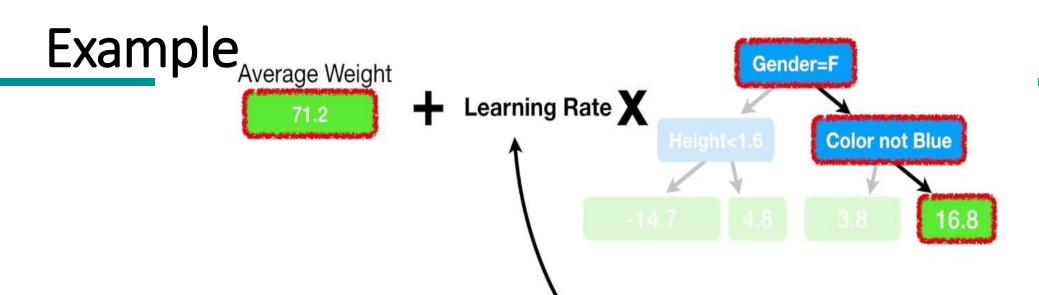
Height	Favorite	Gender	Weight
(m)	Color		(kg)
1.6	Blue	Male	88

In other words, we have low **Bias**, but probably very high **Variance**.



1.6	Blue	Male	88	

Gradient Boost deals with this problem by using a Learning Rate to scale the contribution from the new tree.

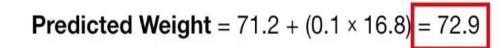


1.6	Blue	Male	88

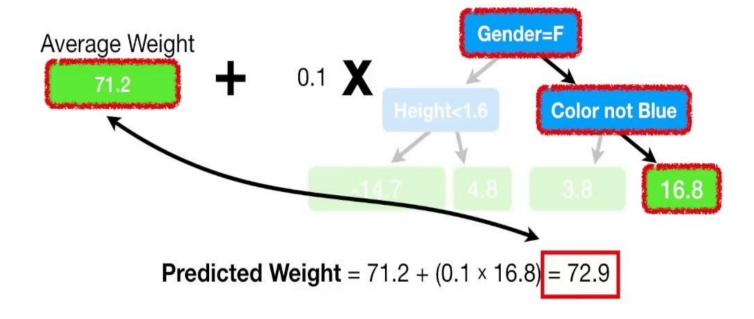
Gradient Boost deals with this problem by using a Learning Rate to scale the contribution from the new tree.

The **Learning Rate** is a value between **0** and **1**.



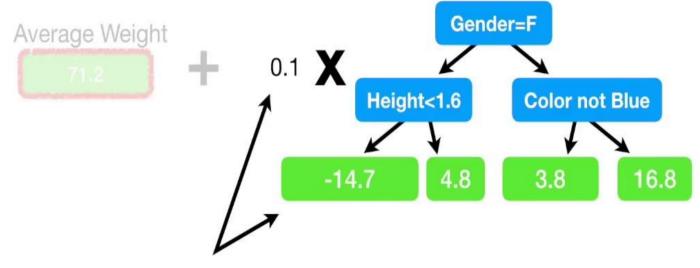


			Weight (kg)	
1.6	Blue	Male	88	With the <b>Learning Rate</b> set to <b>0.1</b> , the new <b>Prediction</b>
				isn't as good as as it was before

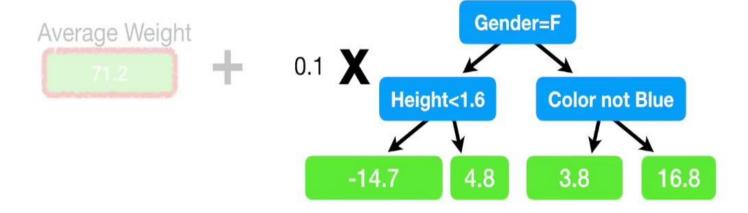


1.6	Blue	Male	88	

...but it's a little bit better than the **Prediction** made with just the original leaf, which predicted that all samples would weigh **71.2**.



In other words, scaling the tree by the **Learning Rate** results in a small step in the right direction.

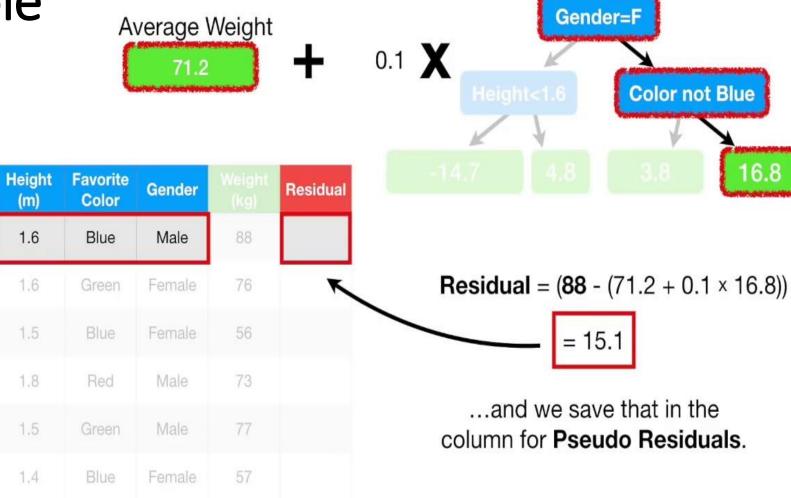


So let's build another tree so we can take another small step in the right direction.

			Weight (kg)	Residual
1.6	Blue	Male	88	
1.6	Green	Female	76	
1.5	Blue	Female	56	
1.8	Red	Male	73	
1.5	Green	Male	77	
1.4	Blue	Female	57	

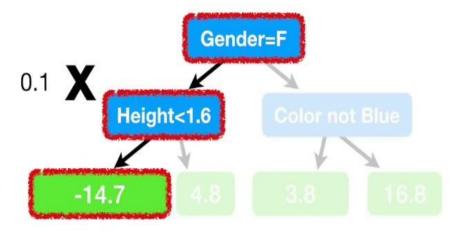
Just like before, we calculate the Pseudo Residuals, the difference between the Observed Weights and our latest Predictions.

Residual = (Observed - Predicted)



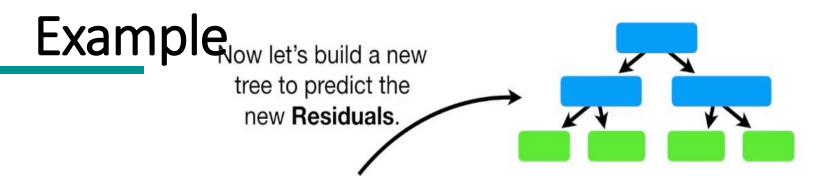


Height (m)	Favorite Color	Gender	Weight (kg)	Residual
1.6	Blue	Male	88	15.1
1.6	Green	Female	76	4.3
1.5	Blue	Female	56	-13.7
1.8	Red	Male	73	1.4
1.5	Green	Male	77	5.4
1.4	Blue	Female	57	-12.7

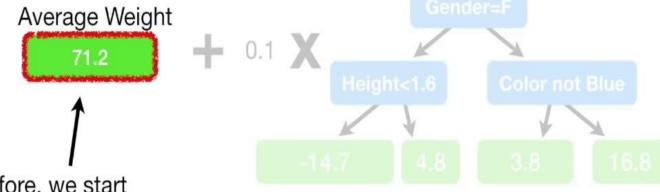


Residual = (Observed - Predicted)

Example Average Weight Gender=F 71.2 Height<1.6 **Color not Blue** -14.73.8 16.8 4.8 Residual Residual 16.8 15.1 The new **Residuals** are all 4.3 4.8 smaller than before, so we've taken a small step in -15.2 -13.7the right direction. 1.4 1.8 5.8 5.4 -14.2 -12.7

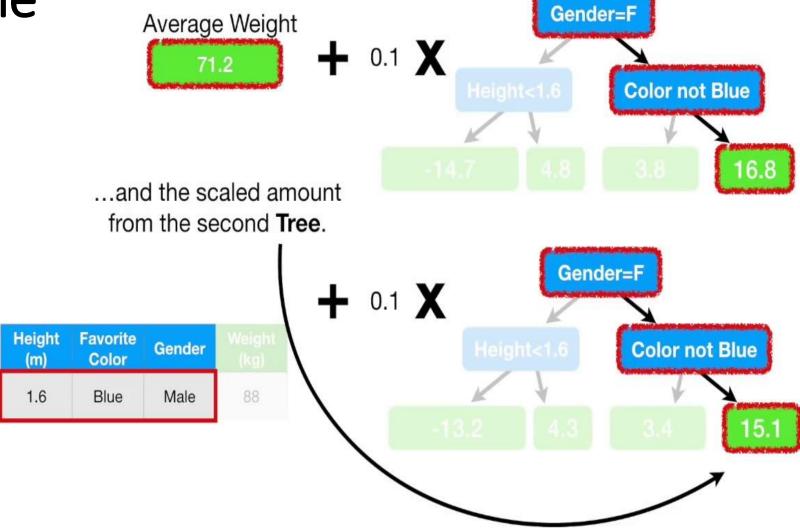


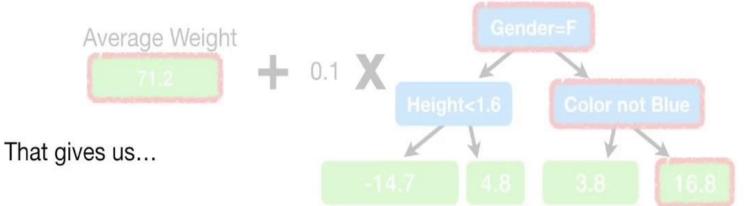
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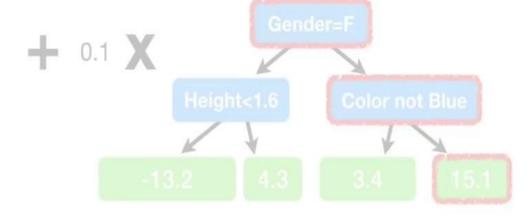
Just like before, we start with the initial **Prediction**...

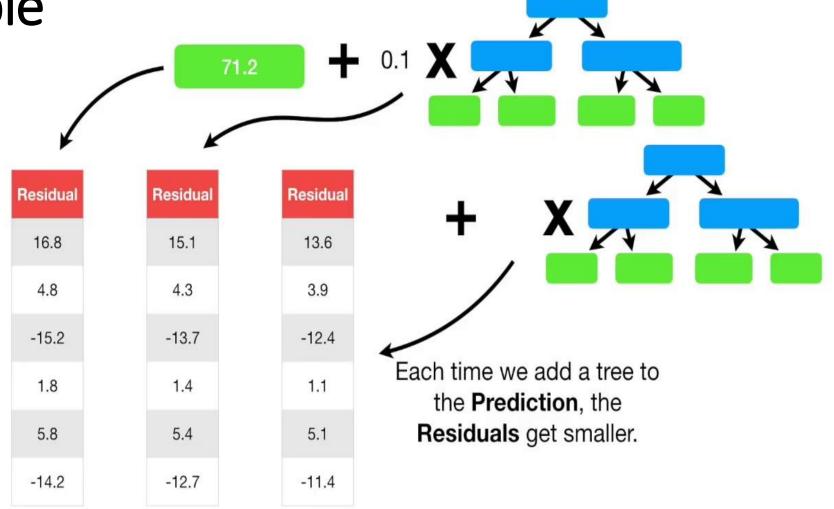
				0.	X				
Height (m)	Favorite Color	Gender	Weight (kg)					Color no	
1.6	Blue	Male	88			K	1		7





$$71.2 + (0.1 \times 16.8) + (0.1 \times 15.1)$$





**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 

**Step 1:** Initialize model with a constant value:  $F_0(x) = \operatorname{argmin} \sum_{i=1}^{n} L(y_i, \gamma)$ 

**Step 2:** for m = 1 to M:

(A) Compute 
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]_{F(x) = F_{m-1}(x)}$$
 for  $i = 1,...,n$ 

- **(B)** Fit a regression tree to the  $r_{im}$  values and create terminal regions  $R_{jm}$ , for  $j = 1...J_m$
- (C) For  $j = 1...J_m$  compute  $\gamma_{jm} = \underset{\gamma}{\operatorname{argmin}} \sum_{x_i \in R_{ij}} L(y_i, F_{m-1}(x_i) + \gamma)$ (D) Update  $F_m(x) = F_{m-1}(x) + \nu \sum_{j=1}^{J_m} \gamma_{jm} I(x \in R_{jm})$

**(D)** Update 
$$F_m(x) = F_{m-1}(x) + \nu \sum_{j=1}^{J_m} \gamma_{jm} I(x \in R_{jm})$$

**Step 3:** Output  $F_M(x)$ 

# GB Algorithm Input: Data $\{(x_i, y_i)\}_{i=1}^n$ and a differentiable Loss Function $L(y_i, F(x))$

The **Loss Function** that is most commonly used when doing Regression with Gradient Boost is...

$$\frac{1}{2}$$
 (Observed - Predicted)<sup>2</sup>

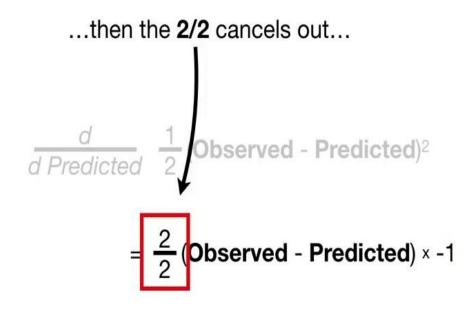
			Weight (kg)
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1.6	Green	Female	76
1.5	Blue	Female	56

**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 

The reason why people choose this **Loss Function** for **Gradient Boost** is that when we differentiate it with respect to "**Predicted**"...

$$\frac{d}{d \text{ Predicted}} \frac{1}{2} (\text{Observed - Predicted})^2$$

**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 



**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 

...and that leaves you with the Observed minus the Predicted multiplied by -1.

$$\frac{d}{d \text{ Predicted}} = \frac{1}{2} (\text{Observed - Predicted})^2$$

$$= \frac{2}{2} (\text{Observed - Predicted}) \times -1$$

= -(Observed - Predicted)

**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 

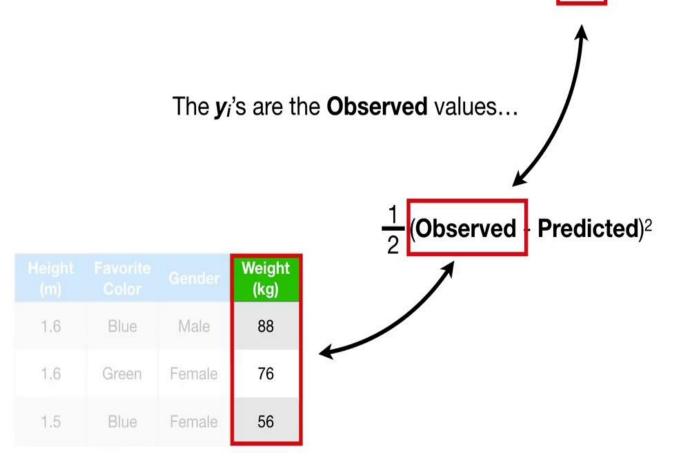
In other words, we are left with the negative **Residual**, and this makes the math easier since **Gradient Boost** uses the derivative a lot.

$$\frac{d}{d \ Predicted} = \frac{1}{2} ( \text{Observed - Predicted} )^2$$

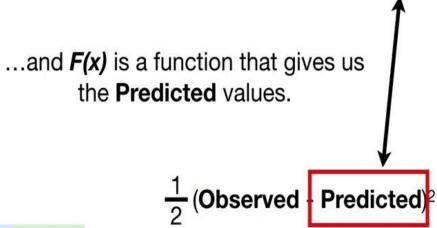
$$=\frac{2}{2}$$
 (Observed - Predicted) × -1

= -(Observed - Predicted)

GB Algorithm Input: Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable Loss Function  $L(y_i, F(x))$ 



GB Algorithm Input: Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable Loss Function  $L(y_i, F(x))$ 



1.6	Blue	Male	88
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**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 

**Step 1:** Initialize model with a constant value:  $F_0(x) = \operatorname{argmin} \sum_{i=1}^{n} L(y_i, \gamma)$ 

$$F_0(x) = \underset{\gamma}{\operatorname{argmin}} \sum_{i=1}^{n} L(y_i, \gamma)$$

We start by initializing the model with a constant value...

**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 

**Step 1:** Initialize model with a constant value:  $F_0(x) = \underset{\gamma}{\operatorname{argmin}} \sum_{i=1}^n L(y_i, \gamma)$ 

The summation means that we add up one **Loss Function** for each **Observed** value...

**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 

**Step 1:** Initialize model with a constant value:  $F_0(x) = \underset{\gamma}{\operatorname{argmin}} \sum_{i} L(y_i, \gamma)$ 

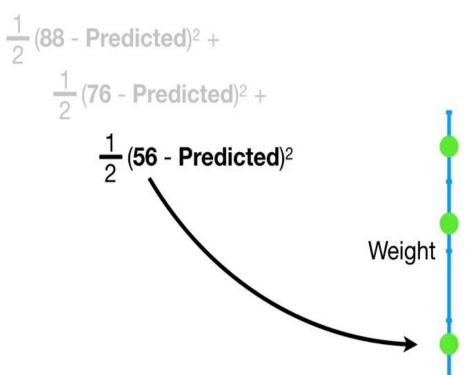
$$\frac{1}{2}$$
 (88 - Predicted)<sup>2</sup> +  $\frac{1}{2}$  (76 - Predicted)<sup>2</sup> +  $\frac{1}{2}$  (56 - Predicted)<sup>2</sup>

Height (m)	Favorite Color	Gender	Weight (kg)
1.6	Blue	Male	88
1.6	Green	Female	76
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...and the "argmin over gamma" means we need to find a Predicted value that minimizes this sum.

**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 

**Step 1:** Initialize model with a constant value:  $F_0(x) = \underset{\gamma}{\operatorname{argmin}} \sum_{i=1}^n L(y_i, \gamma)$ 

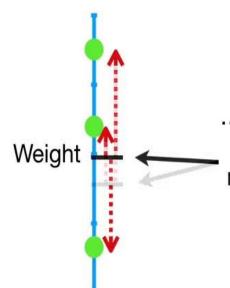


In other words, if we plot the Observed Weights on a number line...

**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 

**Step 1:** Initialize model with a constant value:  $F_0(x) = \underset{\gamma}{\operatorname{argmin}} \sum_{i=1}^n L(y_i, \gamma)$ 

$$\frac{1}{2}$$
 (88 - Predicted)<sup>2</sup> +  $\frac{1}{2}$  (76 - Predicted)<sup>2</sup> +  $\frac{1}{2}$  (56 - Predicted)<sup>2</sup>



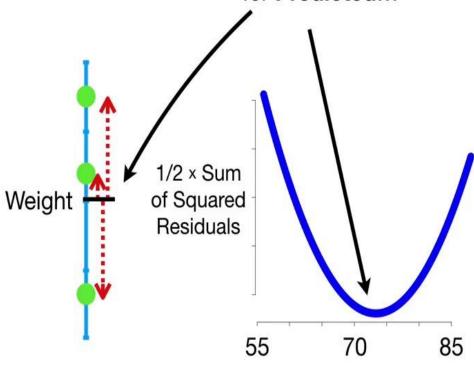
...then we want to find the point on the line that minimizes the sum of the squared residuals...

**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Fu** 

**Step 1:** Initialize model with a constant value:  $F_0(x)$ 

$$\frac{1}{2}$$
 (88 - Predicted)<sup>2</sup> +  $\frac{1}{2}$  (76 - Predicted)<sup>2</sup> +  $\frac{1}{2}$  (56 - Predicted)<sup>2</sup>

NOTE: We could use Gradient Descent to find the optimal value for Predicted...



**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 

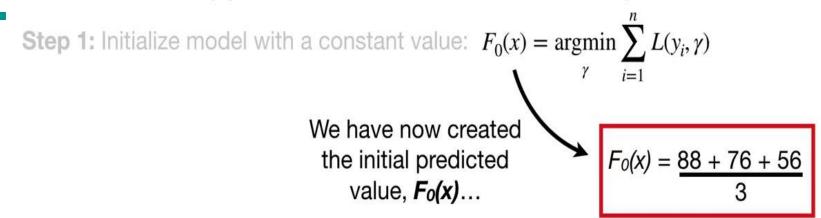
**Step 1:** Initialize model with a constant value:  $F_0(x) = \underset{\gamma}{\operatorname{argmin}} \sum_{i=1}^n L(y_i, \gamma)$ 

**Predicted** = 
$$88 + 76 + 56$$
 3

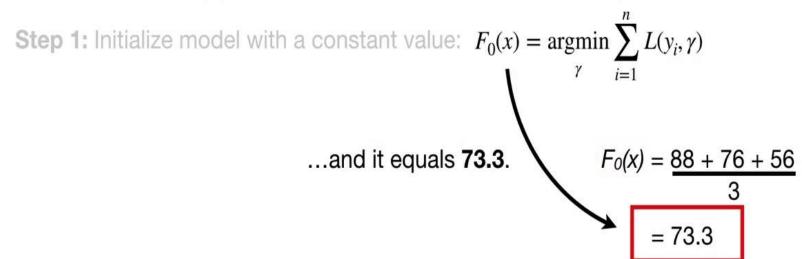
			Weight (kg)
1.6	Blue	Male	88
1.6	Green	Female	76
1.5	Blue	Female	56

...and we end up with the **Average** of the **Observed Weights**.

**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 



**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 



**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 

Step 1: Initialize model with a constant value: 
$$F_0(x) = \underset{\gamma}{\operatorname{argmin}} \sum_{i=1}^n L(y_i, \gamma)$$

That means that the initial predicted value,  $F_0(x)$ , is just a leaf.



73.3

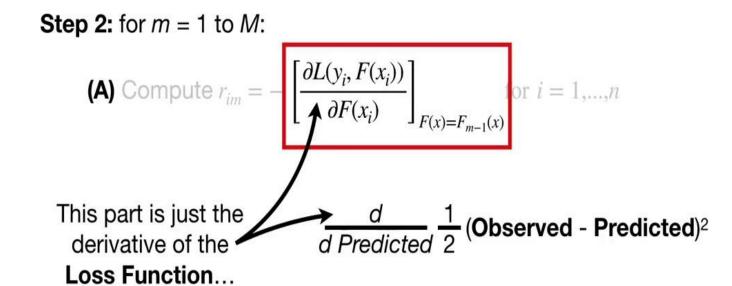
$$F_0(x) = \frac{88 + 76 + 56}{3}$$
$$= 73.3$$

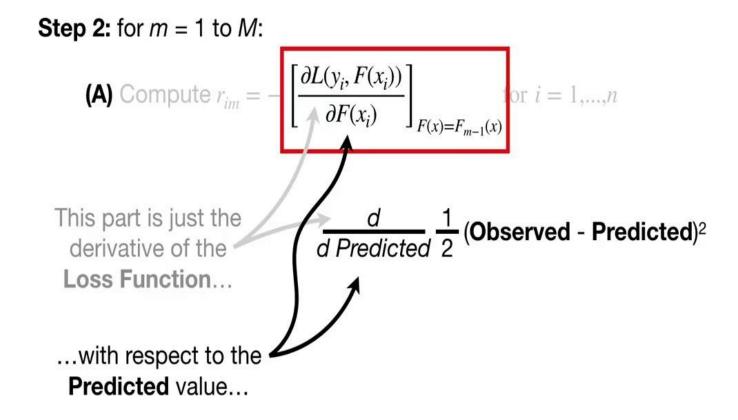
Now we can work on Step 2...

**Step 2:** for m = 1 to M:

**(A)** Compute 
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]_{F(x) = F_{m-1}(x)}$$
 for  $i = 1,...,n$ 

- **(B)** Fit a regression tree to the  $r_{im}$  values and create terminal regions  $R_{jm}$ , for  $j = 1...J_m$
- (C) For  $j = 1...J_m$  compute  $\gamma_{jm} = \underset{\gamma}{\operatorname{argmin}} \sum_{x_i \in R_{ij}} L(y_i, F_{m-1}(x_i) + \gamma)$ (D) Update  $F_m(x) = F_{m-1}(x) + \nu \sum_{i=1}^{J_m} \gamma_m I(x \in R_{jm})$

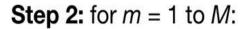


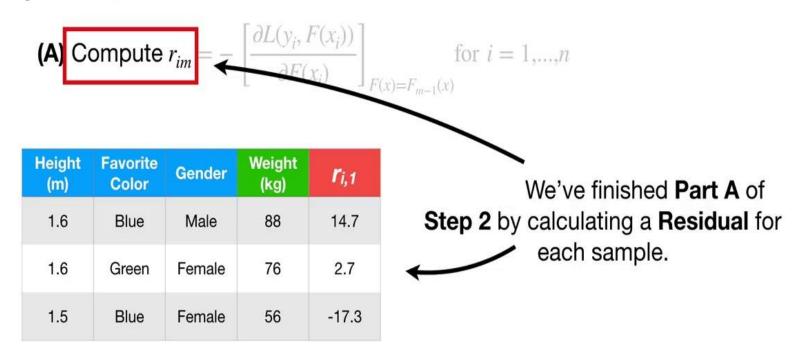


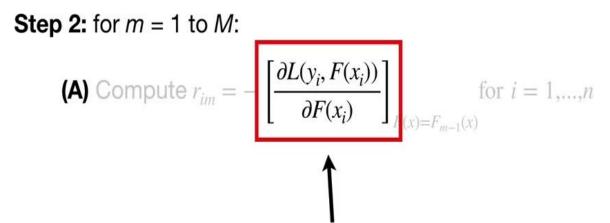
**Step 2:** for m = 1 to M:

**(A)** Compute 
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]_{F(x) = F_{m-1}(x)}$$
 for  $i = 1,...,n$ 

$$\frac{d}{d \ Predicted} \frac{1}{2} ( \textbf{Observed - Predicted} )^2$$
...and we've already 
$$= - ( \textbf{Observed - Predicted} )$$
 calculated this.







NOTE: Before we move on, I just want to point out that this derivative is the Gradient that Gradient Boost is named after.

**Step 2:** for m = 1 to M:

(A) Compute 
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]_{F(x) = F_{m-1}(x)}$$
 for  $i = 1,...,n$ 

I also want to point that the  $r_{i,m}$  values are technically called **Pseudo Residuals**.

**Step 2:** for m = 1 to M:

(A) Compute 
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]_{F(x) = F_{m-1}(x)}$$
 for  $i = 1,...,n$ 

**(B)** Fit a regression tree to the  $r_{im}$  values and create terminal regions  $R_{jm}$ , for  $j = 1...J_m$ 



All this is saying is that we will build a regression tree...

#### Now let's do Part C.

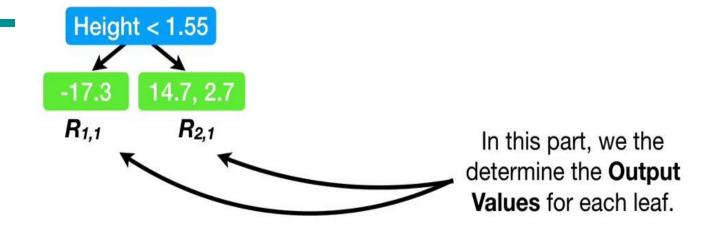
**Step 2:** for m = 1 to M:

**ep 2:** for 
$$m=1$$
 to  $M$ :

**(A)** Compute  $r_{im}=-\left[\frac{\partial L(y_i,F(x_i))}{\partial F(x_i)}\right]_{F(x)=F_{m-1}(x)}$  for  $i=1,...,n$ 

(B) Fit a regression tree to the  $r_{im}$  values and create terminal regions  $R_{jm}$ , for  $j = 1...J_m$ 

(C) For 
$$j = 1...J_m$$
 compute  $\gamma_{jm} = \underset{\gamma}{\operatorname{argmin}} \sum_{x_i \in R_{ij}} L(y_i, F_{m-1}(x_i) + \gamma)$ 



(C) For 
$$j = 1...J_m$$
 compute  $\gamma_{jm} = \underset{\gamma}{\operatorname{argmin}} \sum_{x_i \in R_{ij}} L(y_i, F_{m-1}(x_i) + \gamma)$ 

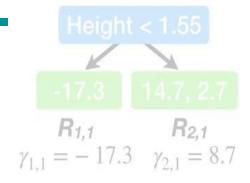
**Step 1:** Initialize model with a constant value:  $F_0(x) = \operatorname{argmin} \sum L(y_i, \gamma)$ 

$$F_0(x) = \underset{\gamma}{\operatorname{argmin}} \sum_{i=1}^n L(y_i, \gamma)$$

**NOTE:** This minimization is like what we did in Step 1.

(C) For 
$$j = 1...J_m$$
 compute

(C) For 
$$j = 1...J_m$$
 compute  $\gamma_{jm} = \underset{\gamma}{\operatorname{argmin}} \sum_{x_i \in R_{ij}} L(y_i, F_{m-1}(x_i) + \gamma)$ 



Given our choice of Loss Function, the Output Values are always the average of the Residuals that end up in the same leaf.

$$\frac{1}{2}$$
 (Observed - Predicted)<sup>2</sup>

#### Now let's do Part D!!!

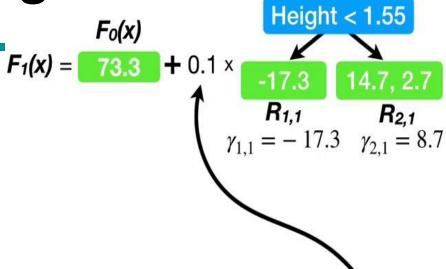
**Step 2:** for m = 1 to M:

(A) Compute 
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]_{F(x) = I_{m-1}(x)}$$
 for  $i = 1, ..., n$ 

(B) Fit a regression tree to the  $r_{im}$  values and create terminal

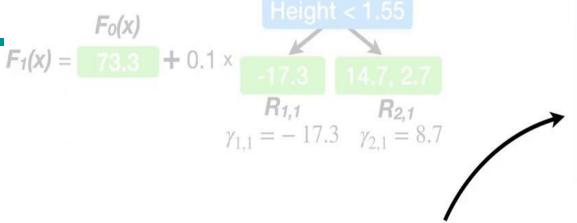
regions 
$$R_{jm}$$
, for  $j=1...J_m$  (C) For  $j=1...J_m$  compute  $\gamma_{jm}=\mathop{\rm argmin} \sum L(y_i,F_{m-1}(x_i)+\gamma)$ 

**(D)** Update 
$$F_m(x) = F_{m-1}(x) + \nu \sum_{j=1}^{J_m} \gamma_{jm} I(x \in R_{jm})$$



In this example, we'll set **nu** to **0.1**.

(**D**) Update 
$$F_m(x) = F_{m-1}(x) + \nu \sum_{j=1}^{J_m} \gamma_{jm} I(x \in R_{jm})$$



Height (m)	Favorite Color	Gender	
1.6	Blue	Male	88
1.6	Green	Female	76
1.5	Blue	Female	56

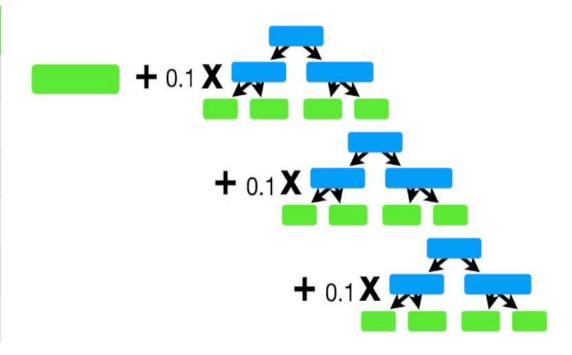
Now we will use  $F_1(x)$  to make new **Predictions** for each sample.

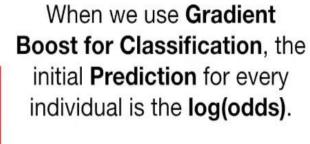
**(D)** Update 
$$F_m(x) = F_{m-1}(x) + \nu \sum_{j=1}^{J_m} \gamma_{jm} I(x \in R_{jm})$$

## Gradient Boosting for Classification

...and walk through, step-by-step, the most common way that **Gradient Boost** fits a model to this **Training Data**.

Likes Popcorn	Age	Favorite Color	Loves Troll 2	
Yes	12	Blue	Yes	
Yes	87	Green Yes		
No	44	Blue	No	
Yes	19	Red No		
No	32	Green	Yes	
No	14	Blue Yes		





		Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
Yes	87	Green	Yes
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes

log(4/2) = 0.7

So this is the **Initial Prediction**.

Yes	12	Blue	Yes
Yes	87	Green	Yes
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes

Just like with Logistic Regression,
the easiest way to use the log(odds)
for Classification is to convert it to a
Probability...

Yes	12	Blue	Yes
Yes	87	Green	Yes
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes

Just like with Logistic Regression,
the easiest way to use the log(odds)
for Classification is to convert it to a
Probability...

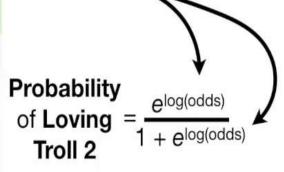
Yes	12	Blue	Yes
Yes	87	Green	Yes
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes

...and we do that with a Logistic Function.

log(4/2) = 0.693

So we plug the log(odds) into the Logistic Function...

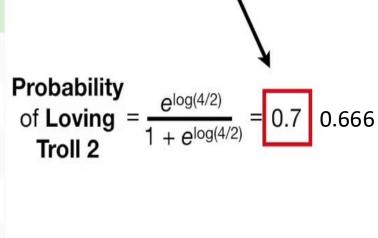
Yes	12	Blue	Yes
Yes	87	Green Yes	
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes



log(4/2) = 0.693

#### ...and we get 0.7 as the Probability of Loving Troll 2.

Yes	12	Blue	Yes
Yes	87	Green	Yes
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes

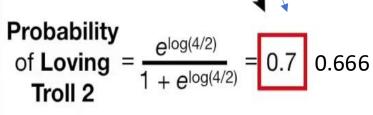


 $\log(4/2) = 0.693$ 

NOTE this are rounded values

...and we get 0.7 as the Probability of Loving Troll 2.

Yes	12	Blue	Yes
Yes	87	Green	Yes
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes



 $\log(4/2) = 0.7$ 

Probability of Loving Troll 2 = 0.7

Since the **Probability** of **Loving Troll 2** is greater than **0.5**, we can **Classify** everyone in the **Training Dataset** as someone who **Loves Troll 2**.

Yes	12	Blue	Yes	
Yes	87 Green		Yes	
No	44	Blue	No	
Yes	19	Red	No	
No	32	Green	Yes	
No	14	Blue	Yes	

log(4/2) = 0.693

Probability of Loving Troll 2 = 0.7 0,666

		Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
Yes	87	Green	Yes
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes

We can measure how bad the initial **Prediction** is by calculating **Pseudo Residuals**, the difference between the **Observed** and the **Predicted** values.

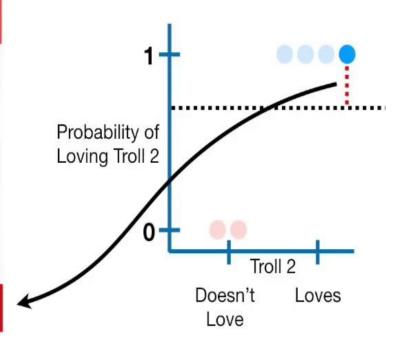
Residual = (Observed - Predicted)

 $\frac{\log(4/2) = 0.7}{\text{Probability of Loving Troll 2}} = 0.7$ 

Then we calculate the rest of the **Residuals**...

Residual = (Observed - Predicted)

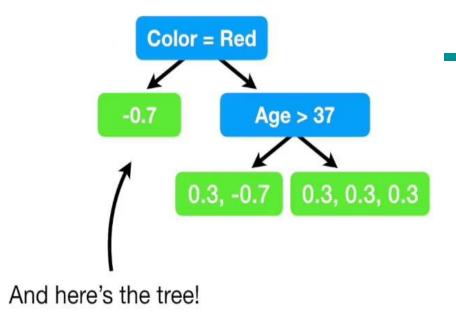
Likes Popcorn	Age	Favorite Color	Loves Troll 2	Residual
Yes	12	Blue	Yes	0.3
Yes	87	Green	Yes	0.3
No	44	Blue	No	-0.7
Yes	19	Red	No	-0.7
No	32	Green	Yes	0.3
No	14	Blue	Yes	0.3



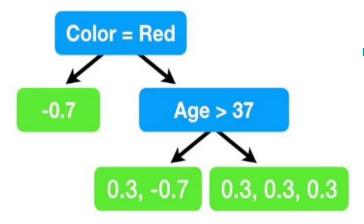
# Example Now we will build a Tree

Likes Popcorn	Age	Favorite Color	Loves Troll 2	Residual
Yes	12	Blue	Yes	0.3
Yes	87	Green	Yes	0.3
No	44	Blue	No	-0.7
Yes	19	Red	No	-0.7
No	32	Green	Yes	0.3
No	14	Blue	Yes	0.3

Likes Popcorn	Age	Favorite Color	Loves Troll 2	Residual
Yes	12	Blue	Yes	0.3
Yes	87	Green	Yes	0.3
No	44	Blue	No	-0.7
Yes	19	Red	No	-0.7
No	32	Green	Yes	0.3
No	14	Blue	Yes	0.3



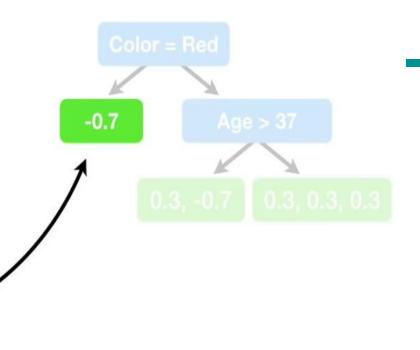
Likes Popcorn	Age	Favorite Color	Loves Troll 2	Residual
Yes	12	Blue	Yes	0.3
Yes	87	Green	Yes	0.3
No	44	Blue	No	-0.7
Yes	19	Red	No	-0.7
No	32	Green	Yes	0.3
No	14	Blue	Yes	0.3

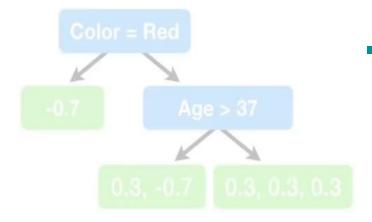


In this simple example, we are limiting the number of leaves to **3**.

In practice people often set the maximum number of leaves to be between 8 and 32

When we used **Gradient Boost** for **Regression**, a leaf with single **Residual** had an **Output Value** equal to that **Residual**.



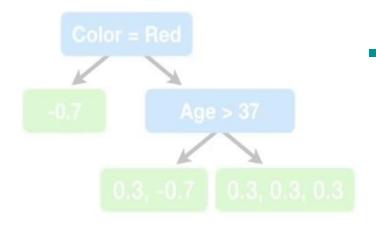


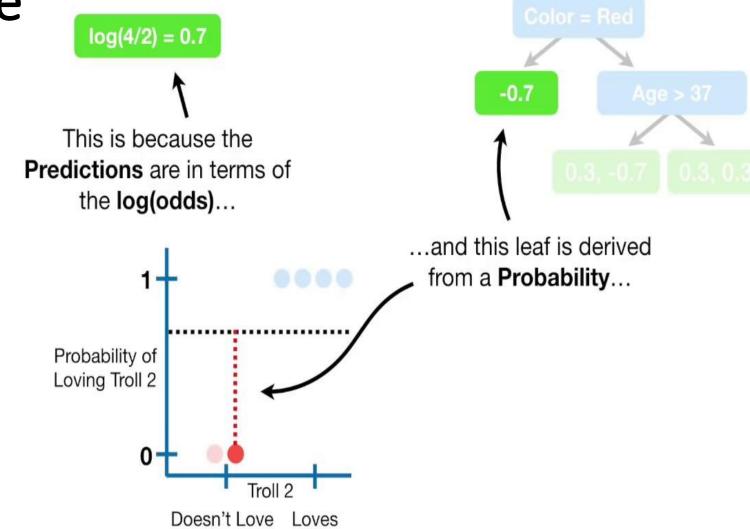
In contrast, when we use **Gradient Boost** for **Classification**, the situation is a little more complex.

log(4/2) = 0.7

1

This is because the **Predictions** are in terms of the **log(odds)**...

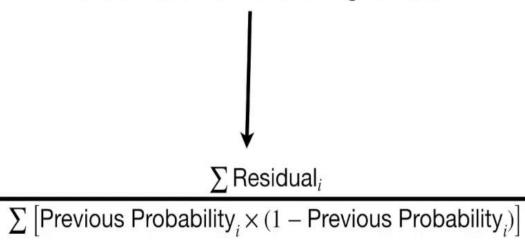


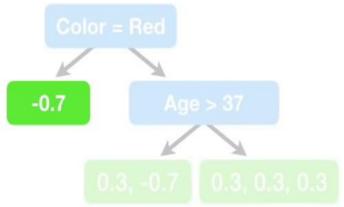


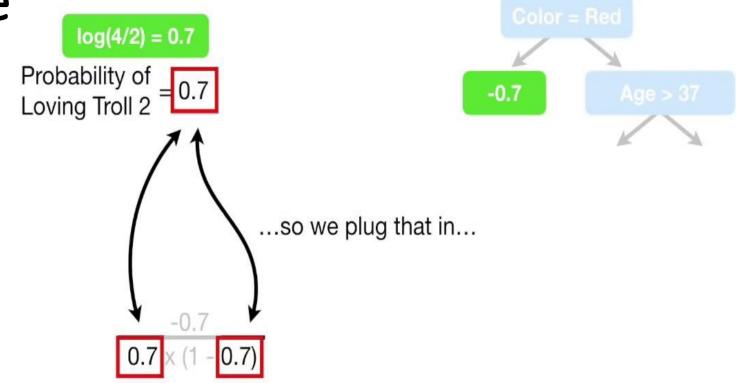


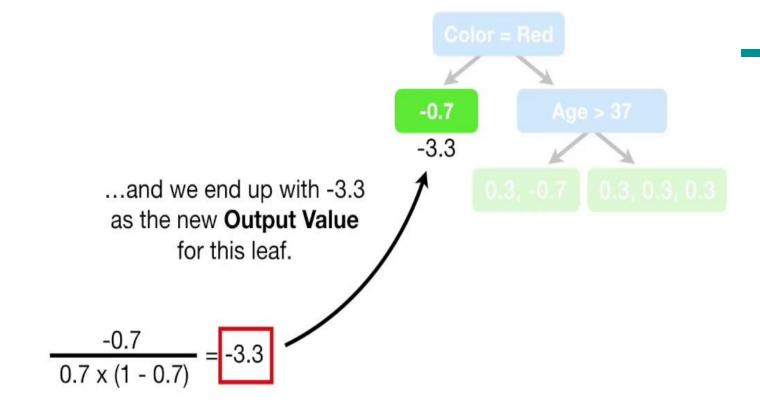
...so we can't just add them together to get a new log(odds) Prediction without some sort of transformation.

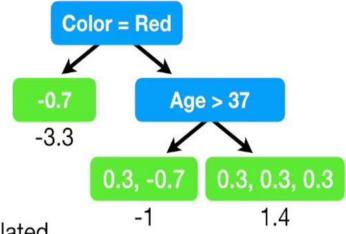
When we use **Gradient Boost** for **Classification**, the most common transformation is the following formula.





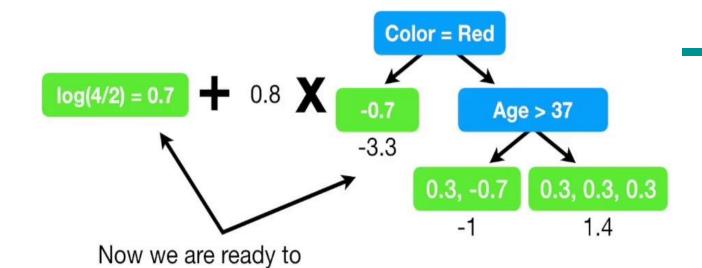






We've calculated

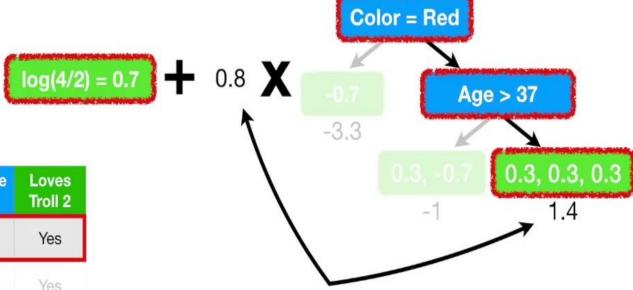
Output Values for all three
leaves in the tree!



update our **Predictions** by

combining the initial leaf

with the new tree.



Likes Popcorn	Age	Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
Yes	87	Green	Yes
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes

...plus the **Output Value** from the tree scaled by the **Learning Rate**...

log(odds) Prediction = 
$$0.7 + (0.8 \times 1.4)$$



Likes Popcorn	Age	Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
Yes	87	Green	Yes
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes

...and the new log(odds)

Prediction = 1.8.

log(odds) Prediction = 
$$0.7 + (0.8 \times 1.4) = 1.8$$

Likes Popcorn	Age	Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
Yes	87	Green	Yes
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes

Now we convert the new log(odds) Prediction into a Probability...

Probability = 
$$\frac{e^{\log(\text{odds})}}{1 + e^{\log(\text{odds})}}$$

Likes Popcorn	Age	Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
Yes	87	Green	Yes
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes

Now we convert the new log(odds) Prediction into a Probability...

Probability = 
$$\frac{e^{\log(\text{odds})}}{1 + e^{\log(\text{odds})}}$$
  
 $\log(\text{odds})$  Prediction = 0.7 + (0.8 × 1.4) = 1.8

Likes Popcorn	Age	Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
Yes	87	Green	Yes
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes

Now we convert the new log(odds) Prediction into a Probability...

Probability = 
$$\frac{e^{1.8}}{1 + e^{1.8}} = 0.9$$

log(4/2) = 0.7

Initial Probability of Loving Troll 2 = 0.7

Likes Popcorn	Age	Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
Yes	87	Green	Yes
No	44	Blue	No
Yes	19	Red	No
No	32	Green	Yes
No	14	Blue	Yes

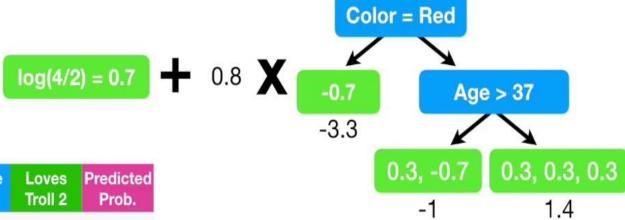
...so we are taking a small step in the right direction since this person **Loves Troll 2**.

Probability = 
$$\frac{e^{1.8}}{1 + e^{1.8}} = 0.9$$

Likes Popcorn	Age	Favorite Color	Loves Troll 2	Predicted Prob.
Yes	12	Blue	Yes	0.9
Yes	87	Green	Yes	
No	44	Blue	No	
Yes	19	Red	No	
No	32	Green	Yes	
No	14	Blue	Yes	

We save the new Predicted Probability here.

Probability = 
$$\frac{e^{1.8}}{1 + e^{1.8}} = 0.9$$



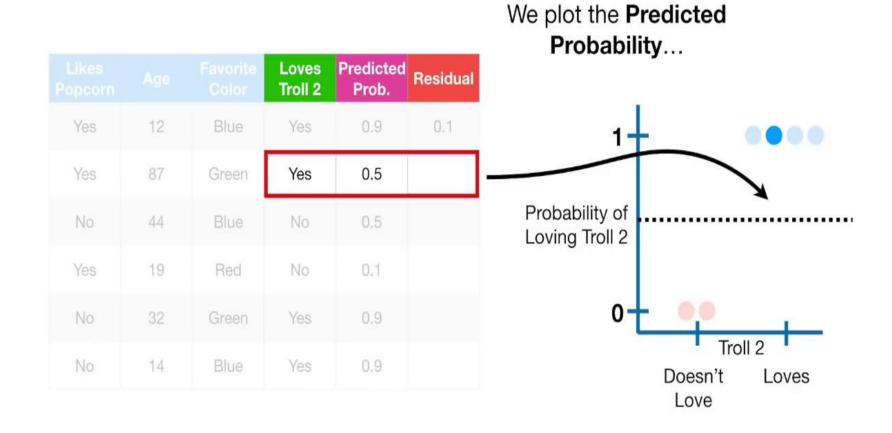
Likes Popcorn	Age	Favorite Color	Loves Troll 2	Predicted Prob.
Yes	12	Blue	Yes	0.9
Yes	87	Green	Yes	0.5
No	44	Blue	No	0.5
Yes	19	Red	No	0.1
No	32	Green	Yes	0.9
No	14	Blue	Yes	0.9

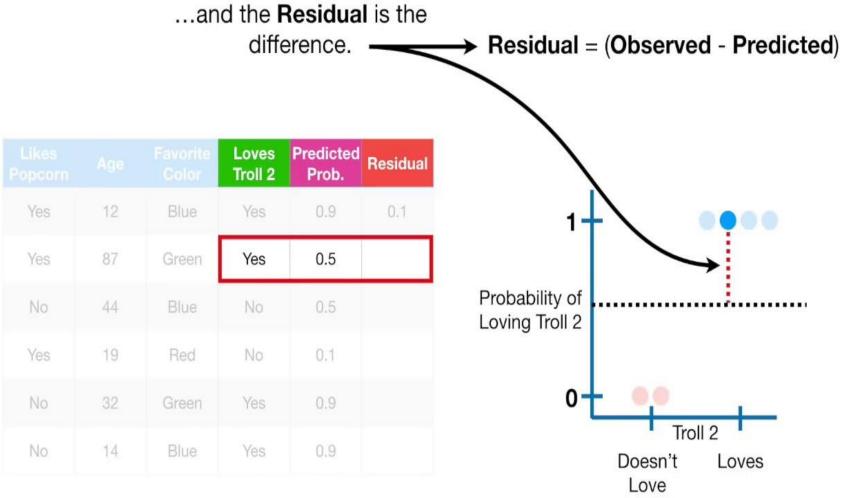
Then we calculate the **Predicted Probabilities** for the remaining people.

And now, just like before, we calculate the new **Residuals**...

-	
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,	4

				Predicted Prob.	Residual
Yes	12	Blue	Yes	0.9	
Yes	87	Green	Yes	0.5	
No	44	Blue	No	0.5	
Yes	19	Red	No	0.1	
No	32	Green	Yes	0.9	
No	14	Blue	Yes	0.9	









Age < 66

0.1, -0.1, 0.1, 0.1

-0.5

0.5

Likes Popcorn	Age	Favorite Color	Loves Troll 2		Residual
Yes	12	Blue	Yes	0.9	0.1
Yes	87	Green	Yes	0.5	0.5
No	44	Blue	No	0.5	-0.5
Yes	19	Red	No	0.1	-0.1
No	32	Green	Yes	0.9	0.1
No	14	Blue	Yes	0.9	0.1

**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 

**Step 1:** Initialize model with a constant value:  $F_0(x) = \operatorname{argmin} \sum_{i=1}^{n} L(y_i, \gamma)$ 

**Step 2:** for m = 1 to M:

(A) Compute 
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]_{F(x) = F_{m-1}(x)}$$
 for  $i = 1,...,n$ 

- **(B)** Fit a regression tree to the  $r_{im}$  values and create terminal regions  $R_{im}$ , for  $j = 1...J_m$
- (C) For  $j = 1...J_m$  compute  $\gamma_{jm} = \underset{\gamma}{\operatorname{argmin}} \sum_{x_i \in R_{ij}} L(y_i, F_{m-1}(x_i) + \gamma)$ (D) Update  $F_m(x) = F_{m-1}(x) + \nu \sum_{j=1}^{J_m} \gamma_{jm} I(x \in R_{jm})$

**(D)** Update 
$$F_m(x) = F_{m-1}(x) + \nu \sum_{j=1}^{J_m} \gamma_{jm} I(x \in R_{jm})$$

**Step 3:** Output  $F_M(x)$ 

# GB Algorithm Input: Data $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable Loss Function $L(y_i, F(x))$

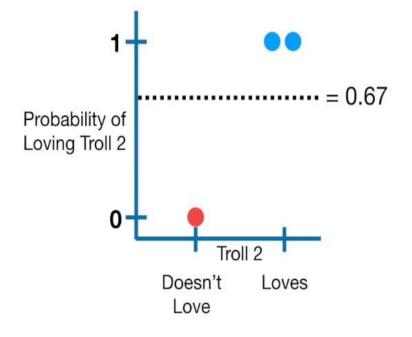


Now we need a differentiable **Loss Function** that will work for Classification.

Log(Likelihood of the Observed Data given the Prediction) =

$$\sum_{i=1}^{N} \mathbf{y}_{i} \times \log(\mathbf{p}) + (1 - \mathbf{y}_{i}) \times \log(1 - \mathbf{p})$$

Likes Popcorn	Age	Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
No	87	Green	Yes
No	44	Blue	No



- Observed 
$$\times \log(\mathbf{p}) + (1 - \mathbf{Observed}) \times \log(1 - \mathbf{p})$$

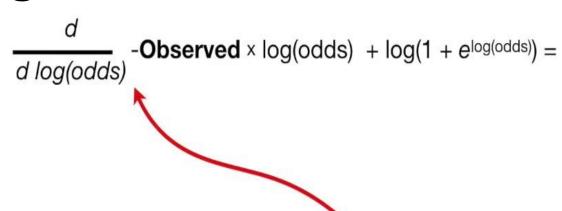
$$\sum_{i=1}^{N} \mathbf{y}_{i} \times \log(\mathbf{p}) + (1 - \mathbf{y}_{i}) \times \log(1 - \mathbf{p})$$

- 1) -Observed  $\times \log(p)$  ( $\mathbf{k}$  Observed)  $\times \log(1 p)$
- 2) -Observed  $\times \log(p) \log(1 \cdot p) + Observed \times \log(1 p)$
- 3) -Observed ×  $[\log(p) \log(1-p)] \log(1-p)$
- 4) -Observed  $\times \log(\text{odds}) \log(1 p)$
- 5) -Observed  $\times \log(\text{odds}) + \log(1 + e^{\log(\text{odds})})$

We converted the **negative Log(likelihood)** of the data, which is a function of the predicted probability, p...

...into a function of the predicted log(odds).

This is the Loss function



So let's take the derivative of the **Loss Function** with respect to the predicted **log(odds)**.

$$\frac{d}{d \log(odds)} - \textbf{Observed} \times \log(odds) + \log(1 + e^{\log(odds)}) =$$

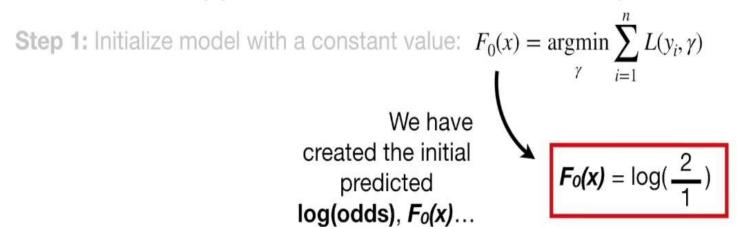
$$= -\mathbf{Observed} + \frac{e^{\log(\text{odds})}}{1 + e^{\log(\text{odds})}}$$

$$=$$
 -Observed +  $p$ 

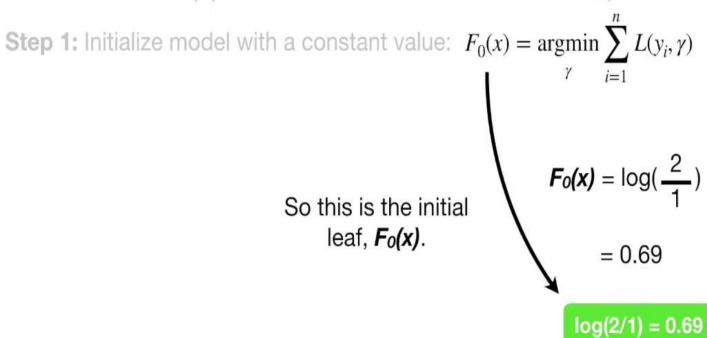
Indeed, in the previous example we performed

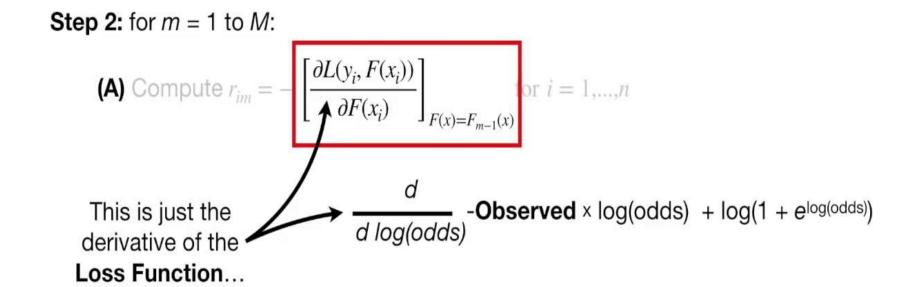
- 1 0.7 = 0.3
- 0 0.7 = -0.7

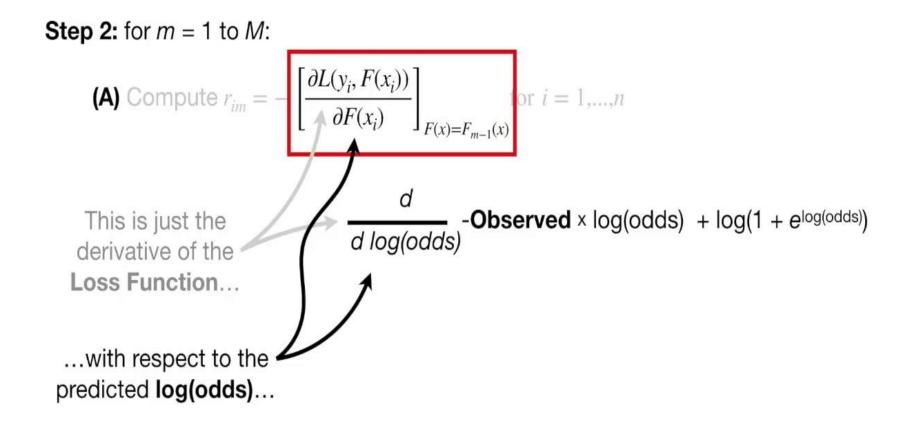
**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 



**Input:** Data  $\{(x_i, y_i)\}_{i=1}^n$ , and a differentiable **Loss Function**  $L(y_i, F(x))$ 







**Step 2:** for m = 1 to M:

**(A)** Compute 
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]_{F(x) = F_{m-1}(x)}$$
 for  $i = 1, ..., n$ 

$$\frac{d}{d \log(odds)} \text{-Observed} \times \log(\text{odds}) + \log(1 + e^{\log(\text{odds})})$$
...and we've already 
$$= (\text{-Observed} + \frac{e^{\log(\text{odds})}}{1 + e^{\log(\text{odds})}})$$
calculated this.

**Step 2:** for m = 1 to M:

**(A)** Compute 
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]_{F(x) = F_{m-1}(x)}$$
 for  $i = 1,...,n$ 

...and that leaves us with this equation for a calculating Pseudo Residuals.

(Observed - 
$$\frac{e^{\log(\text{odds})}}{1 + e^{\log(\text{odds})}}$$
)

**Step 2:** for m = 1 to M:

**(A)** Compute 
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]_{F(x) = F_{m-1}(x)}$$
 for  $i = 1, ..., n$ 

NOTE: As we have seen before, we can replace this term with the predicted probability, 
$$p$$
... (Observed -  $\frac{e^{\log(\text{odds})}}{1 + e^{\log(\text{odds})}}$ )

**Step 2:** for m = 1 to M:

**(A)** Compute 
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]_{F(x) = F_{m-1}(x)}$$
 for  $i = 1, ..., n$ 

**Step 2:** for m = 1 to M:

(A) Compute 
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]_{F(x) = F_{m-1}(x)}$$
 for  $i = 1,...,n$ 

**(B)** Fit a regression tree to the  $r_{im}$  values and create terminal regions  $R_{jm}$ , for  $j = 1...J_m$ 



Now we are ready for **Part B**, where we will build a regression tree.

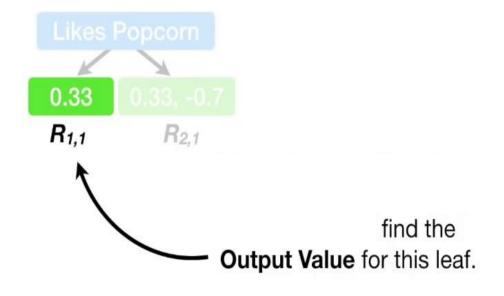
Now let's do **Part C**.

**Step 2:** for m = 1 to M:

**(A)** Compute 
$$r_{im} = -\left[\frac{\partial L(y_i, F(x_i))}{\partial F(x_i)}\right]_{F(x) = F_{m-1}(x)}$$
 for  $i = 1, ..., n$ 

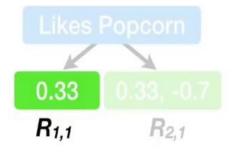
(B) Fit a regression tree to the  $r_{im}$  values and create terminal regions  $R_{jm}$ , for  $j = 1...J_m$ 

(C) For 
$$j = 1...J_m$$
 compute  $\gamma_{jm} = \underset{\gamma}{\operatorname{argmin}} \sum_{x_i \in R_{ij}} L(y_i, F_{m-1}(x_i) + \gamma)$ 



$$\gamma_{1,1} = \underset{\gamma}{\operatorname{argmin}} \sum_{x_i \in R_{ij}} L(y_i, F_{m-1}(x_i) + \gamma)$$

Likes Popcorn	Age	Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
No	87	Green	Yes
No	44	Blue	No

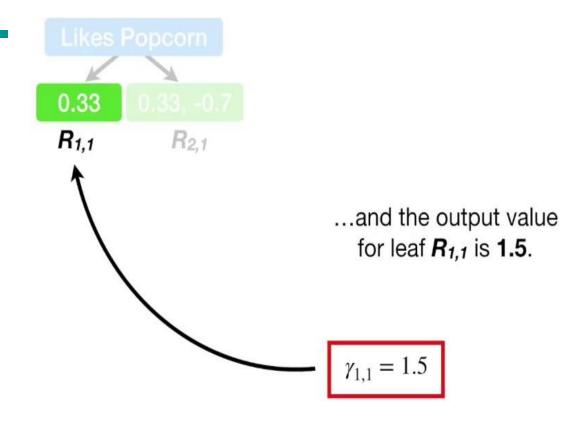


Likes Popcorn	Age	Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
No	87	Green	Yes
No	44	Blue	No

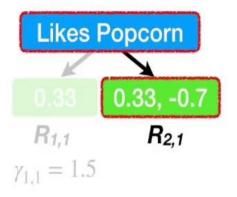
# $\frac{\sum \mathsf{Residual}_i}{\sum \left[\mathsf{Previous\ Probability}_i \times (1 - \mathsf{Previous\ Probability}_i)\right]}$

$$\gamma_{1,1} = \frac{\mathbf{Residual}}{\boldsymbol{p} \times (1 - \boldsymbol{p})}$$

 $0.33/(2/3 \times 1/3) = 1.48$ 



Likes Popcorn	Age	Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
No	87	Green	Yes
No	44	Blue	No

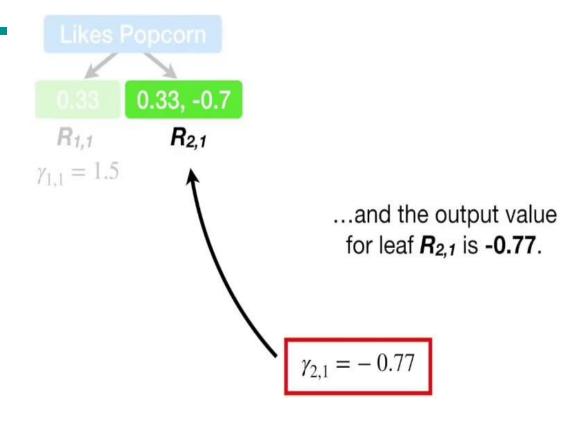


$$\gamma_{2,1} = \frac{\text{Residual}_2 + \text{Residual}_3}{[\boldsymbol{p}_2 \times (1 - \boldsymbol{p}_2)] + [\boldsymbol{p}_3 \times (1 - \boldsymbol{p}_3)]}$$

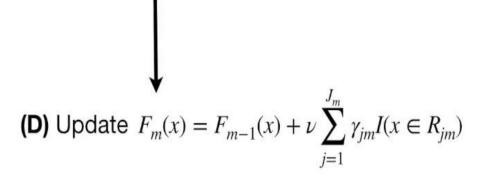
Likes Popcorn	Age	Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
No	87	Green	Yes
No	44	Blue	No

 $\sum$  Residual<sub>i</sub>

 $\sum$  [Previous Probability<sub>i</sub> × (1 – Previous Probability<sub>i</sub>)]



In **Part D**, we make a new prediction for each sample.



$$F_{1}(x) = \log(2/1) = 0.69 + 0.8 \times 0.33 \quad 0.33, -0.7$$

$$R_{1,1} \quad R_{2,1}$$

$$\gamma_{1,1} = 1.5 \quad \gamma_{2,1} = -0.77$$
...the **Output Values** from the first tree we made.

$$\text{(D) Update } F_{m}(x) = F_{m-1}(x) + \nu \sum_{j=1}^{J_{m}} \gamma_{jm} I(x \in R_{jm})$$

# **Gradient Boost Improved**

- XGBoost (Extreme Gradient Boost)
- LightGBM (Light Gradient Boosting Machine)
- Both designed for large complex datasets

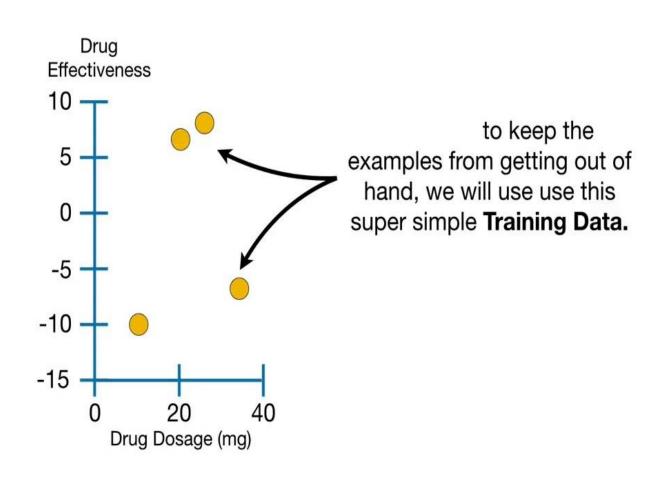
Mass	Age	ВР	Color	Movie	Car	Hair	etc.
120	23	102	Brown	T2	Ford	Long	
150	25	98	Brown	Frozen	Kia	Short	***
165	22	130	Black	Spiderman	Ford	Short	
123	45	98	Red	T2	Kia	Long	•••
156	33`	78	Brown	Frozen 2	Ford	Long	
***		•••	***	***	***	***	***

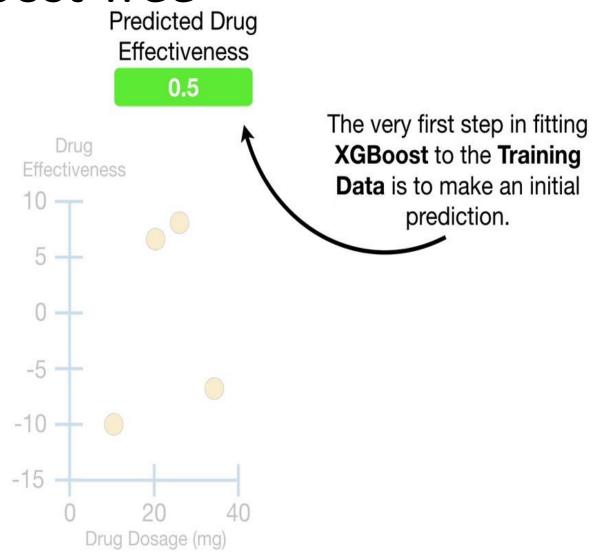
## **XGBoost Characteristics**

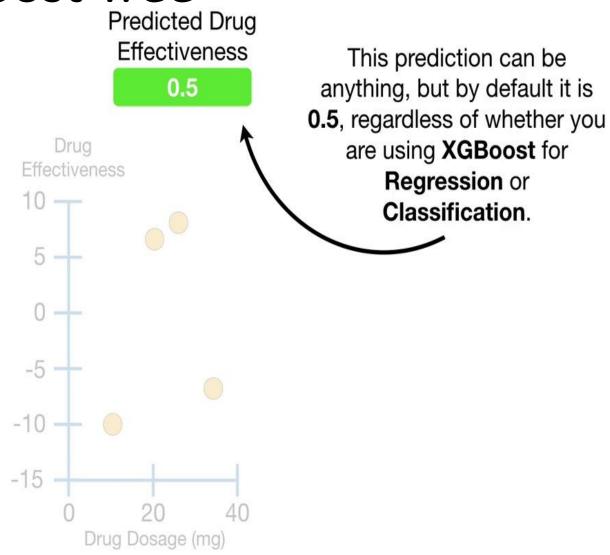
- Gradient Boost
- Regularization
- Approximate Greedy Algorithm
- Weighted Quantile Sketch
- Sparsity-Aware Split Finding
- Parallel Learning
- Cache-Aware Access
- Blocks for Out-of-Core Computation

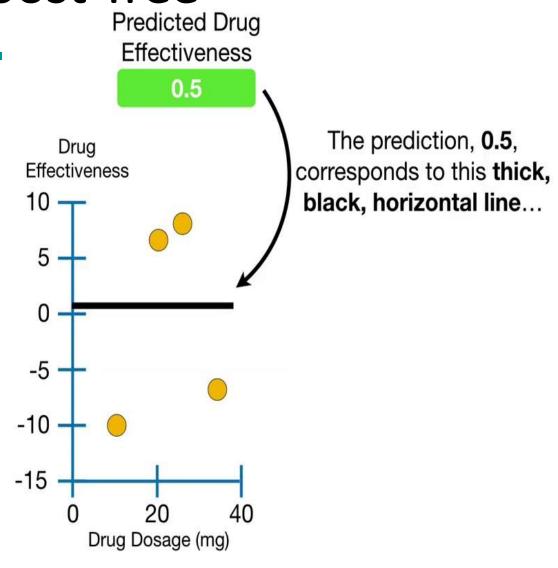
## **XGBoost Characteristics**

- Gradient Boost
- Regularization
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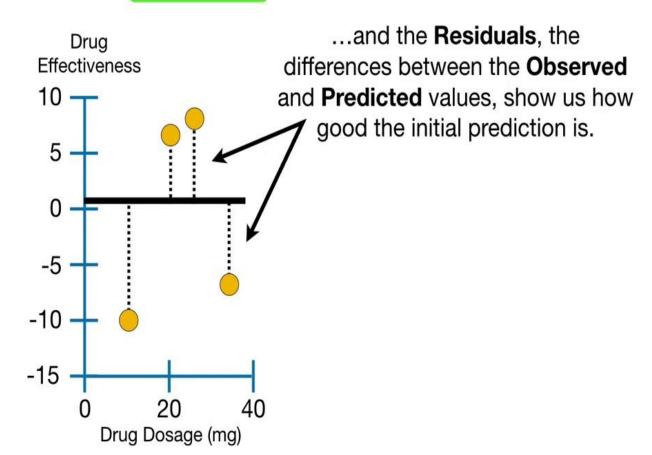




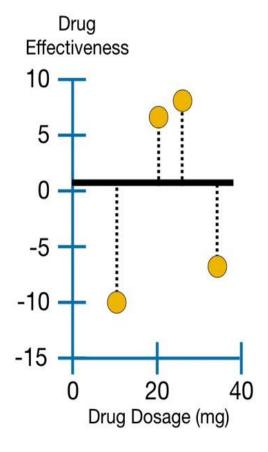


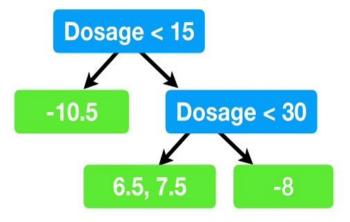
Predicted Drug Effectiveness

0.5



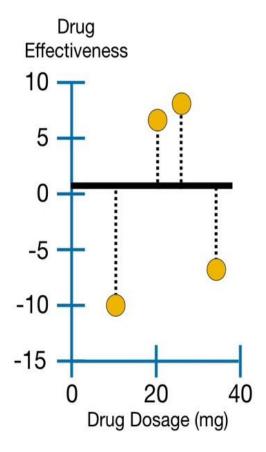
Predicted Drug
Effectiveness
0.5

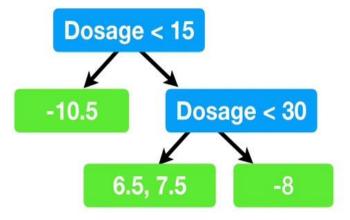




However, unlike unextreme **Gradient Boost**, which typically uses regular,
off-the-shelf, **Regression Trees**...

Predicted Drug
Effectiveness
0.5

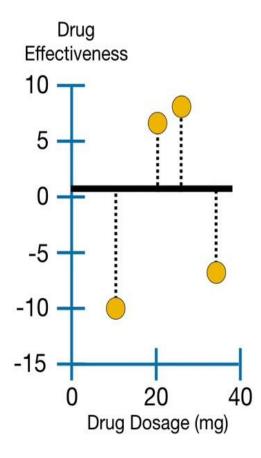


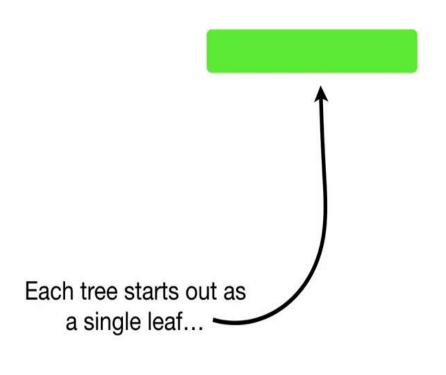


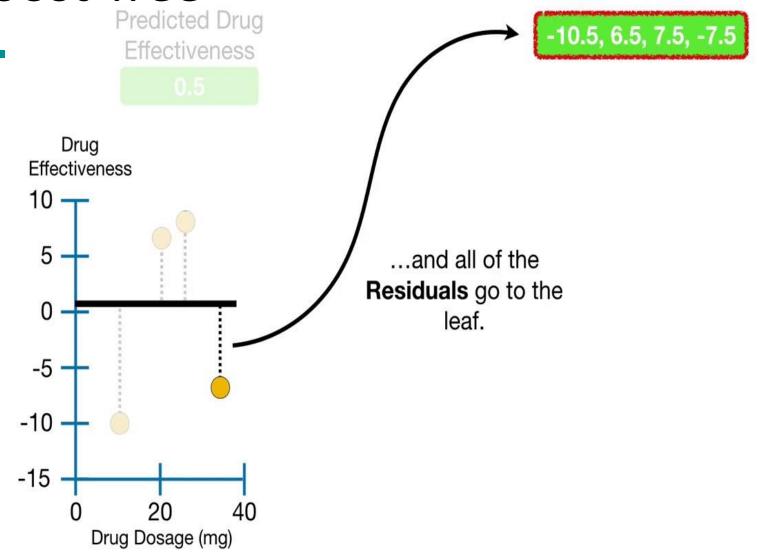
...XGBoost uses a unique Regression Tree that I call an XGBoost Tree.

Predicted Drug Effectiveness

0.5



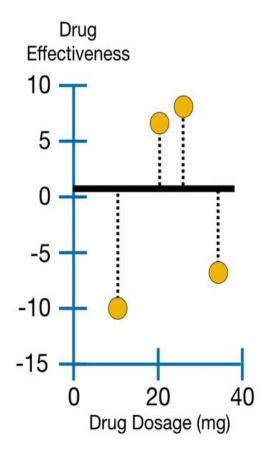




Predicted Drug Effectiveness

0.5

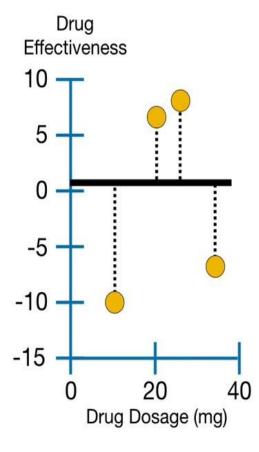
-10.5, 6.5, 7.5, -7.5



Now we calculate a **Quality Score**, or **Similarity Score**, for the **Residuals**.

Predicted Drug Effectiveness

0.5



-10.5, 6.5, 7.5, -7.5

Similarity Score =  $\frac{\text{Sum of Residuals, Squared}}{\text{Number of Residuals } + \lambda}$ 

NOTE: λ (lambda) is a Regularization parameter,

Predicted Drug Effectiveness

0.5

Drug Effectiveness 10 — 5 0 --5 -10 --15 20 40 Drug Dosage (mg)

-10.5, 6.5, 7.5, -7.5

Similarity Score = 
$$\frac{(-10.5 + 6.5 + 7.5 + -7.5)^2}{4 + 0}$$

Predicted Drug Effectiveness

0.5

Drug Effectiveness 10 -5 -5 -10 --15 20 40 Drug Dosage (mg)

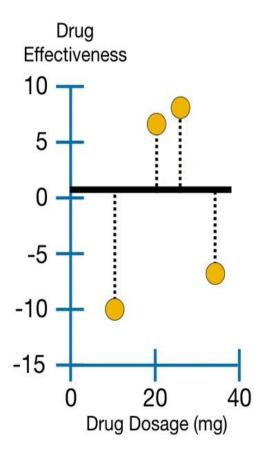
-10.5, 6.5, 7.5, -7.5

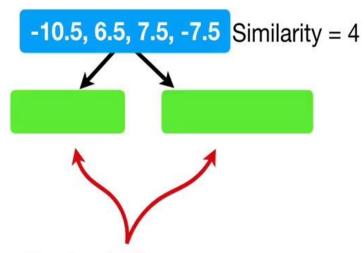
Similarity Score = 
$$\frac{(-4)^2}{4+0}$$
 = 4

Thus, the **Similarity Score** for the **Residuals** in the root = **4**.

Predicted Drug Effectiveness

0.5

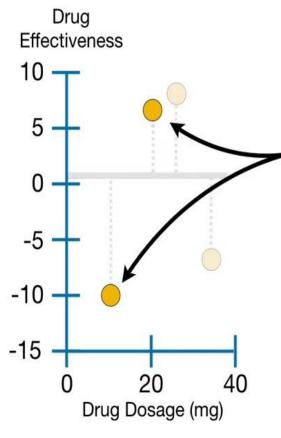




Now the question is whether or not we can do a better job clustering similar **Residuals** if we split them into two groups.

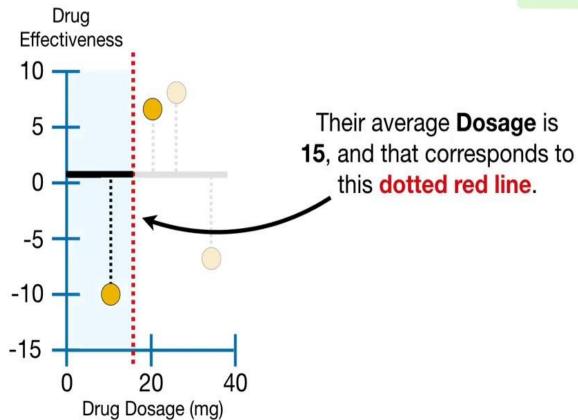
Predicted Drug
Effectiveness

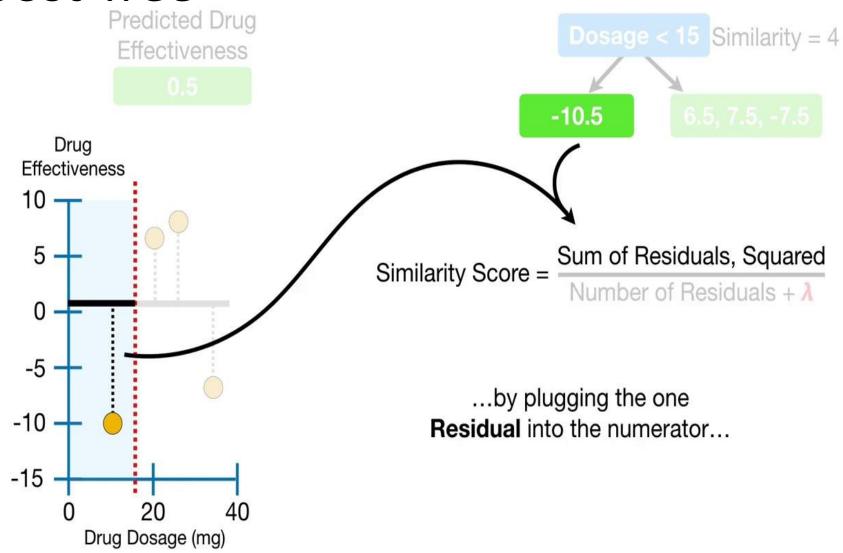


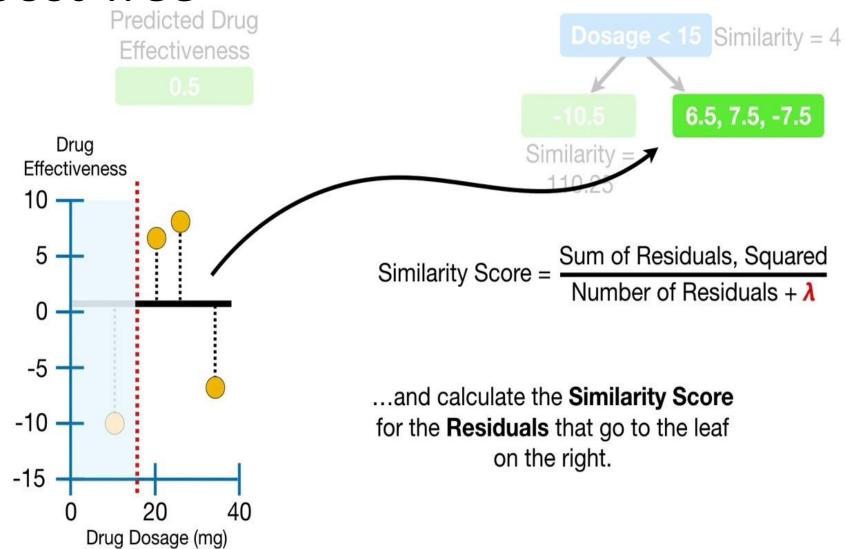


To answer this, we first focus on the two observations with the lowest **Dosages**.

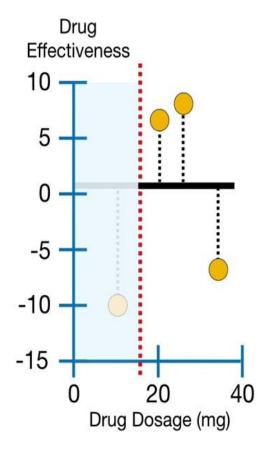


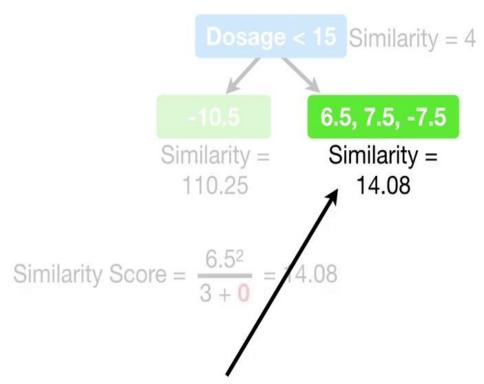




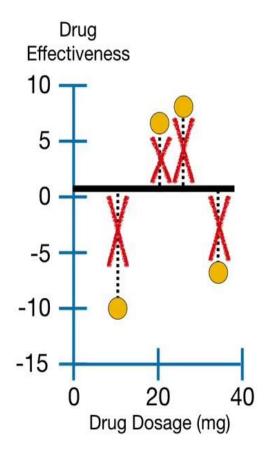


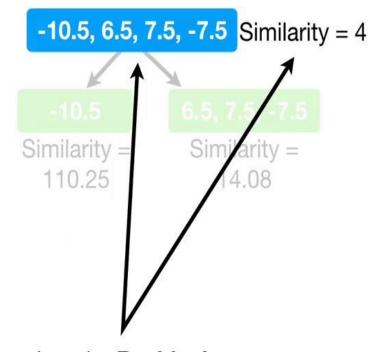
Predicted Drug
Effectiveness
0.5





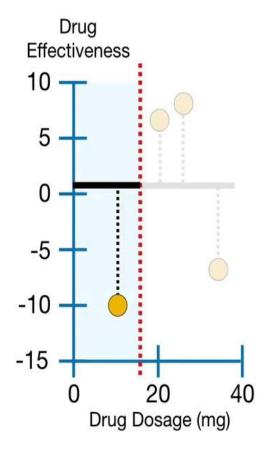
So let's put **Similarity = 14.08** under the leaf so we can keep track of it.

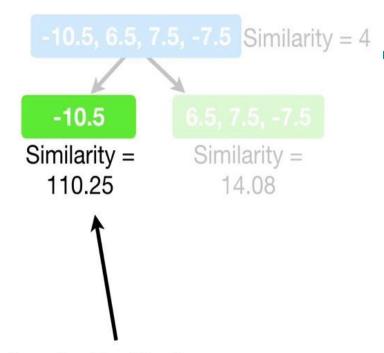




when the **Residuals** in a node are very different, they cancel each other out and the **Similarity Score** is relatively small.

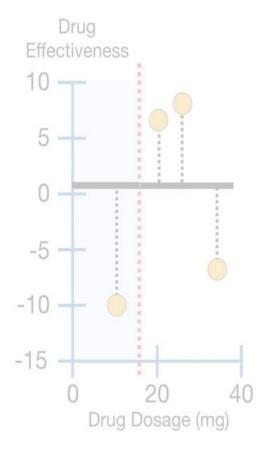
Predicted Drug
Effectiveness
0.5

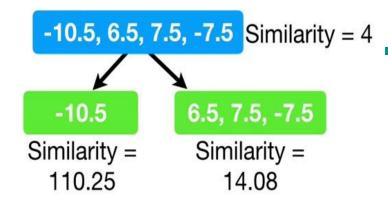




In contrast, when the **Residuals** are similar, or there is just one of them, they do not cancel out and the **Similarity Score** is relatively large.

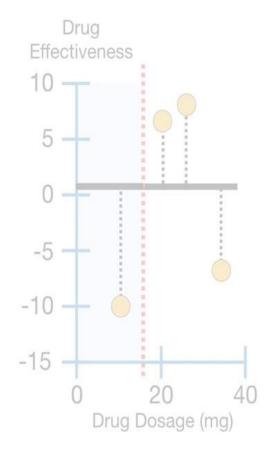
Predicted Drug
Effectiveness
0.5

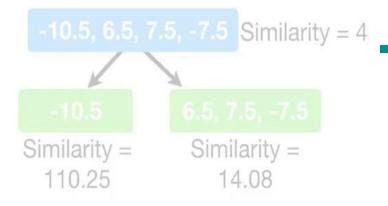


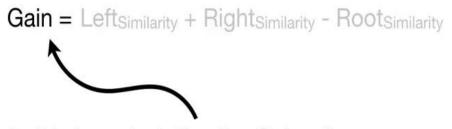


Now we need to quantify how much better the leaves cluster similar **Residuals** than the root.

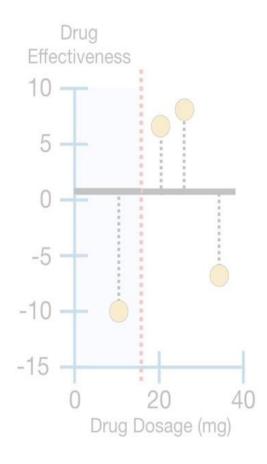
Predicted Drug
Effectiveness
0.5

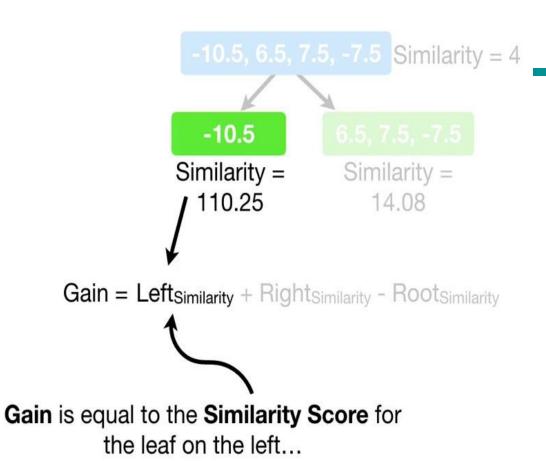


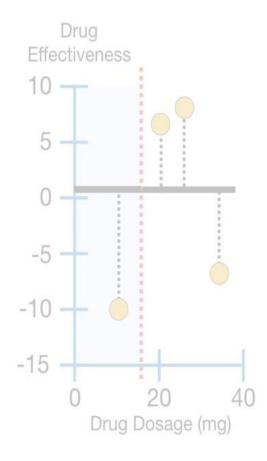


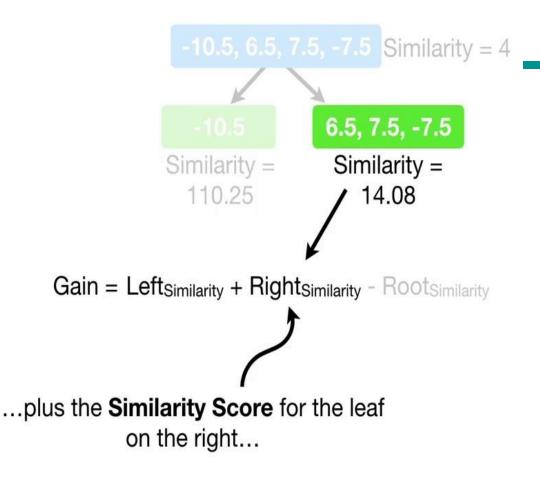


We do this by calculating the **Gain** of splitting the **Residuals** into two groups.

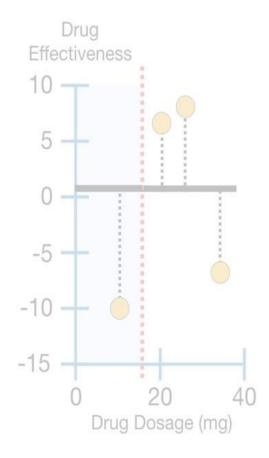


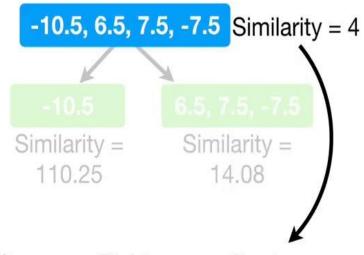






Predicted Drug
Effectiveness

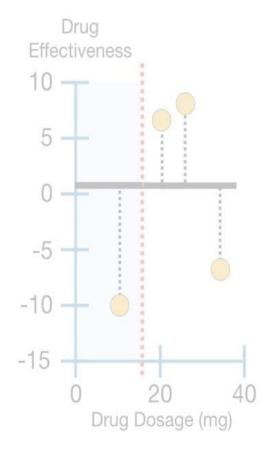


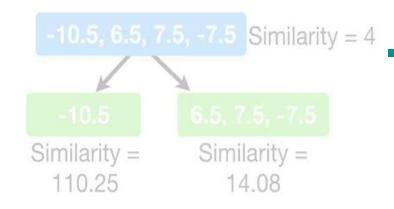


Gain = Left<sub>Similarity</sub> + Right<sub>Similarity</sub> - Root<sub>Similarity</sub>



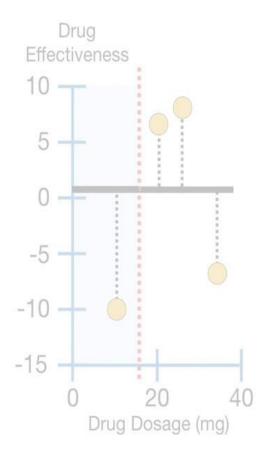
...minus the **Similarity Score** for the root.

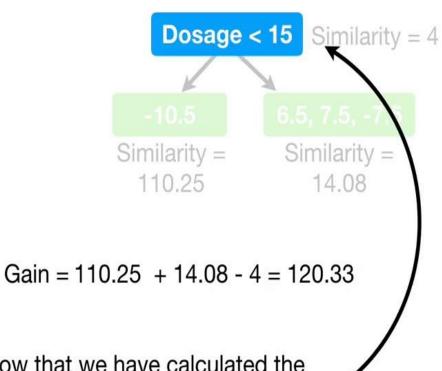




Gain = 
$$110.25 + 14.08 - 4 = 120.33$$

Predicted Drug
Effectiveness

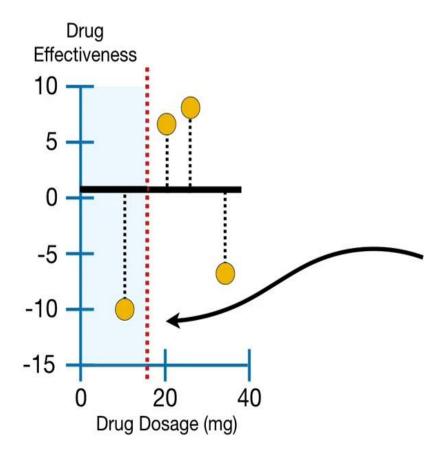




Now that we have calculated the **Gain** for the threshold **Dosage < 15**, we can compare it to the **Gain** calculated for other thresholds.

Predicted Drug Effectiveness

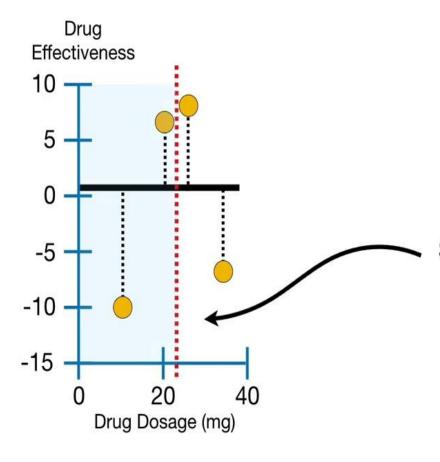
0.5



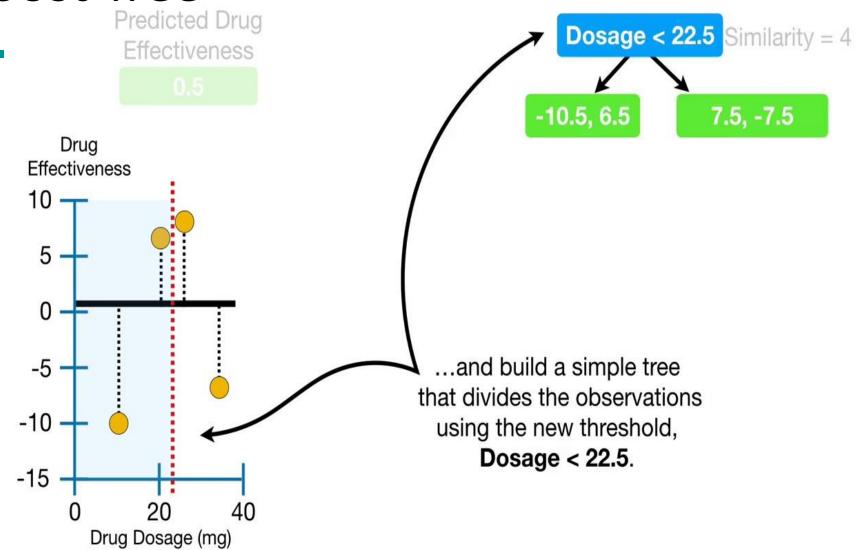
So we shift the threshold over so that it is the average of the next two observations...

Predicted Drug Effectiveness

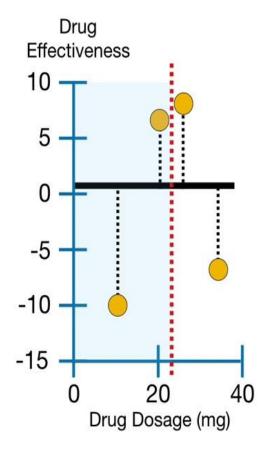
0.5

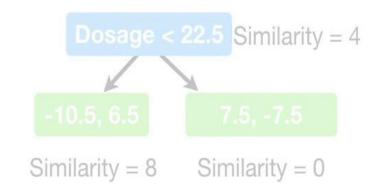


So we shift the threshold over so that it is the average of the next two observations...



Predicted Drug
Effectiveness
0.5

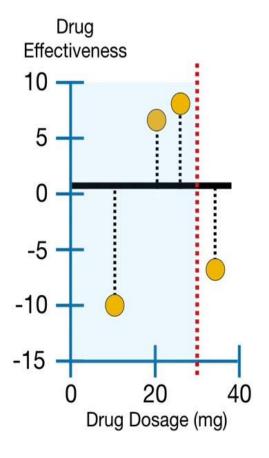


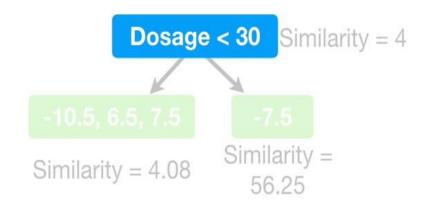


Gain = 
$$8 + 0 - 4 = 4$$

Since the **Gain** for **Dosage** < 22.5 (**Gain** = 4) is less than the **Gain** for **Dosage** < 15 (**Gain** = 120.33), **Dosage** < 15 is better at splitting the **Residuals** into clusters of similar values.

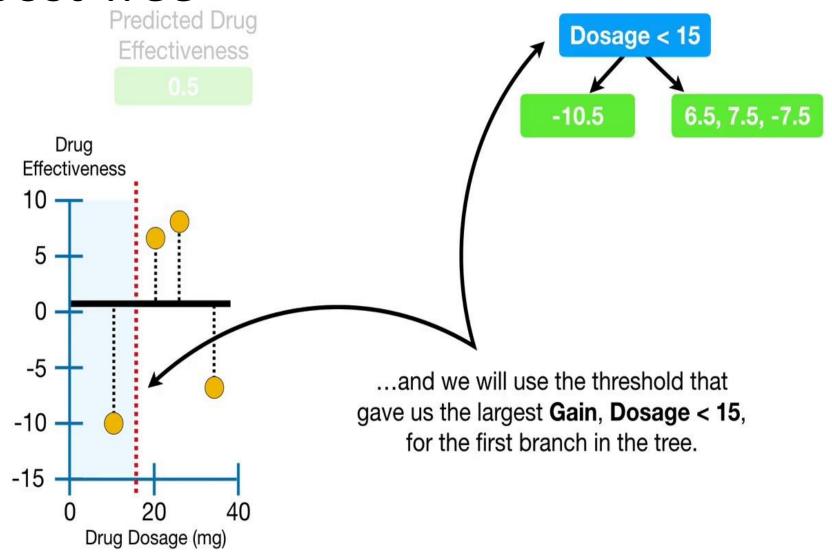
Predicted Drug
Effectiveness
0.5



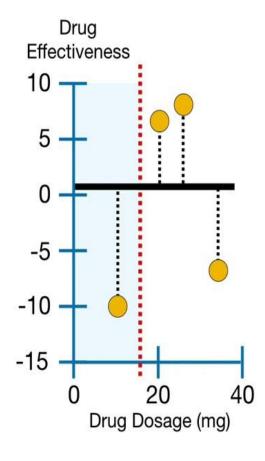


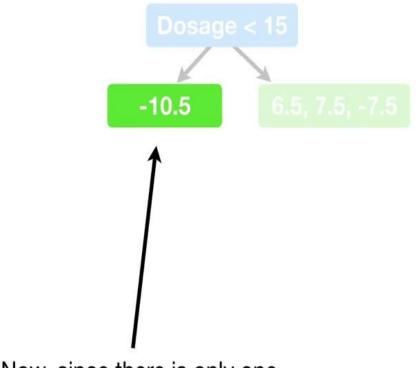
Gain = 
$$4.08 + 56.25 - 4 = 56.33$$

Again, since the **Gain** for **Dosage** < **30** (**Gain** = **56.33**) is less than the **Gain** for **Dosage** < **15** (**Gain** = **120.33**), **Dosage** < **15** is better at splitting the observations.

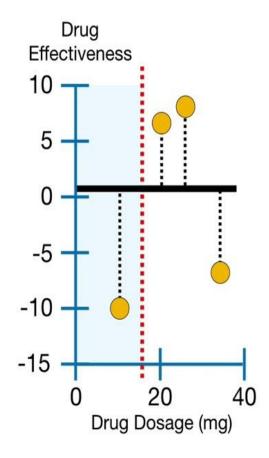


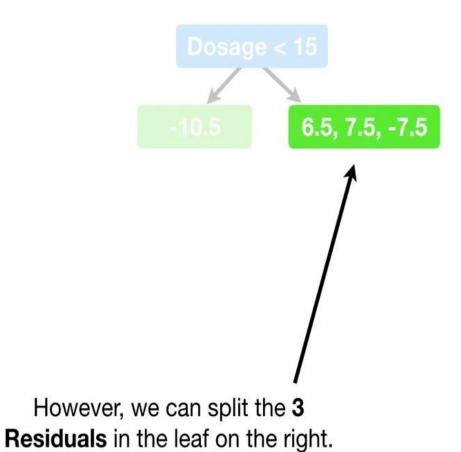
Predicted Drug
Effectiveness
0.5



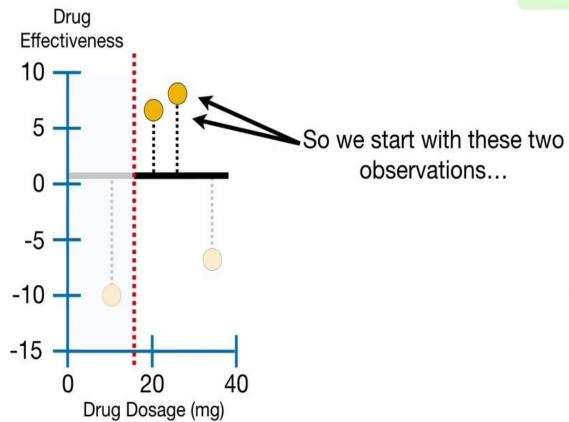


Now, since there is only one **Residual** in the leaf on the left, we can't split it any further.

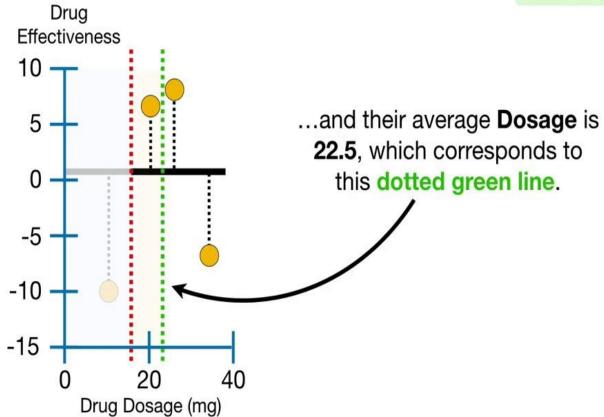








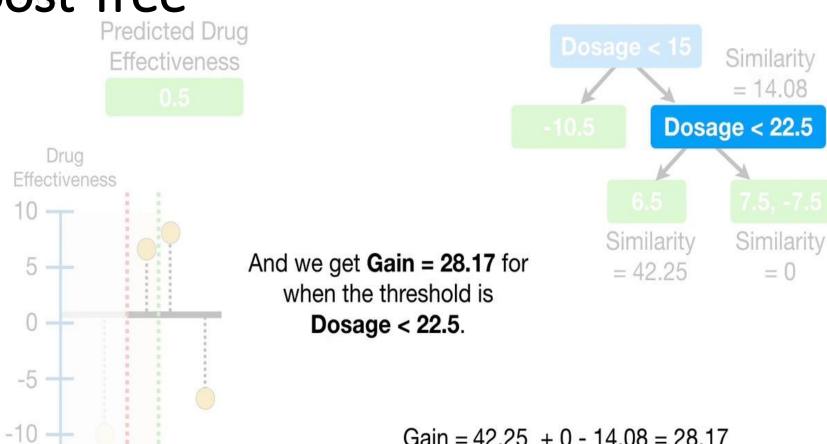




-15

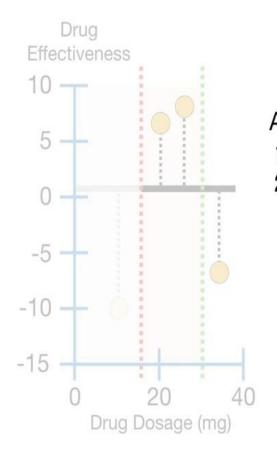
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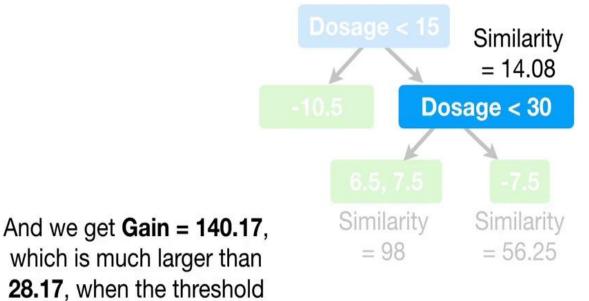
Drug Dosage (mg)



Gain = 
$$42.25 + 0 - 14.08 = 28.17$$

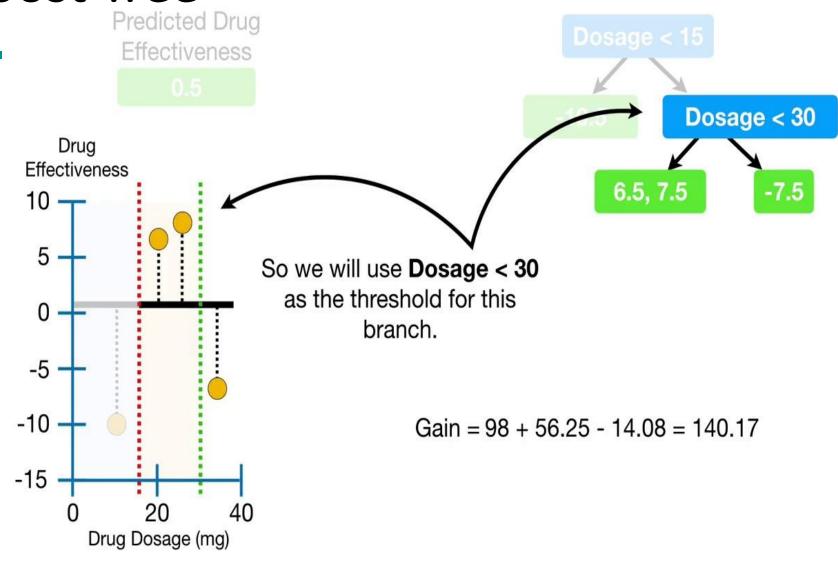
Predicted Drug
Effectiveness
0.5





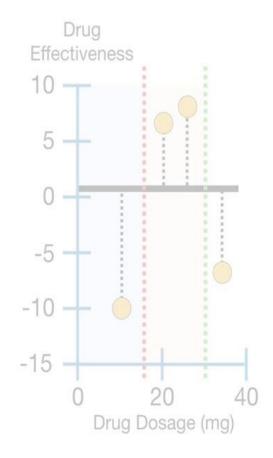
$$Gain = 98 + 56.25 - 14.08 = 140.17$$

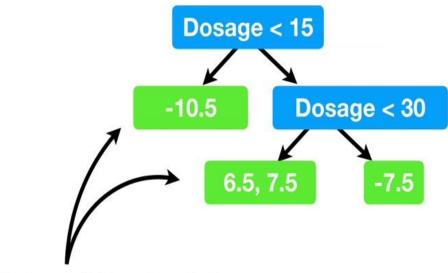
was **Dosage < 22.5**.



#### XGBoost Tree

Predicted Drug
Effectiveness
0.5

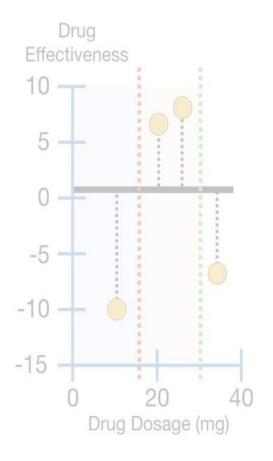


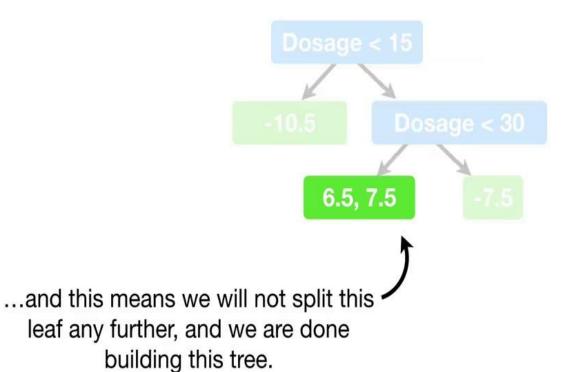


**NOTE:** To keep this example from getting out of hand, I've limited the tree depth to two levels...

#### **XGBoost Tree**

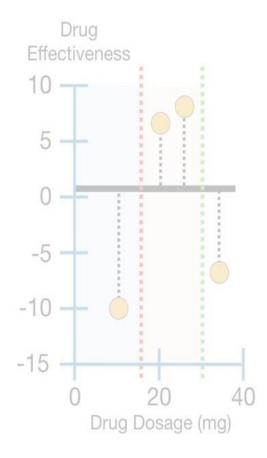
Predicted Drug
Effectiveness
0.5

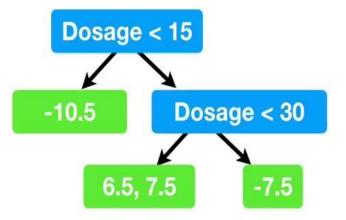




#### XGBoost Tree

Predicted Drug
Effectiveness
0.5

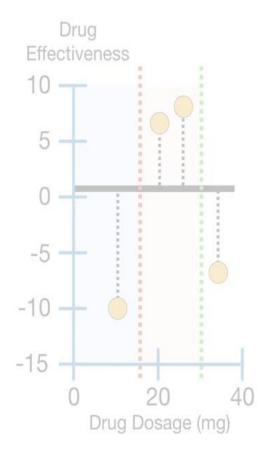


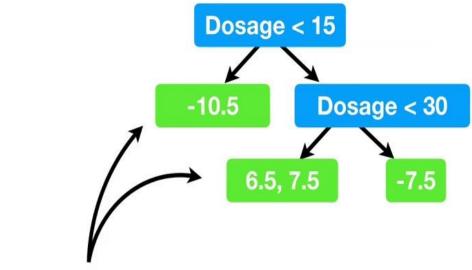


However, the default is to allow up to 6 levels.

Tree Pruning
Predicted Drug

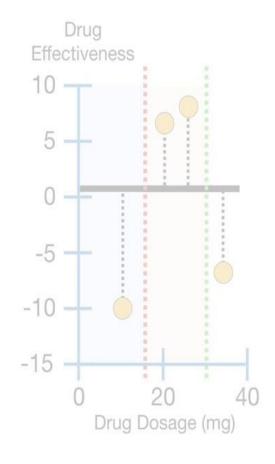
Predicted Drug
Effectiveness
0.5

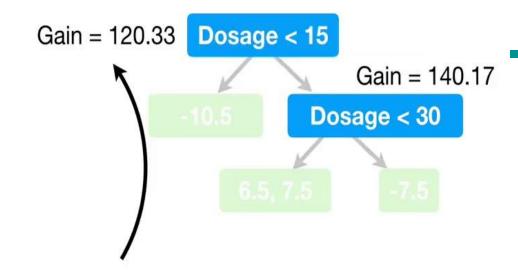




Now we need to talk about how to **Prune** this tree.

Predicted Drug
Effectiveness
0.5





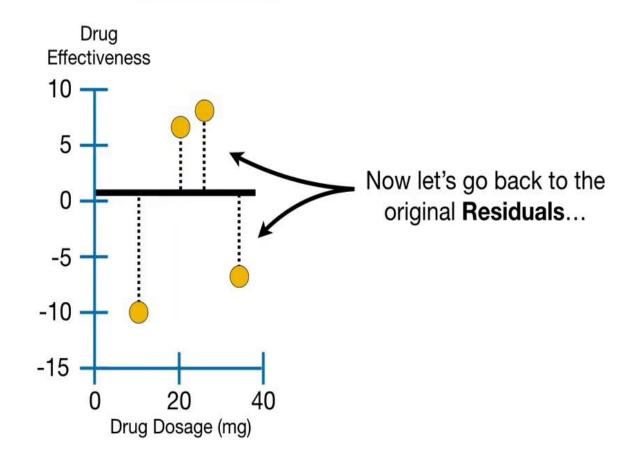
We **Prune** an **XGBoost Tree** based on its **Gain** values.

We set a threshold parameter **gamma**. Then we cut leaves with a gain less than **gamma**.

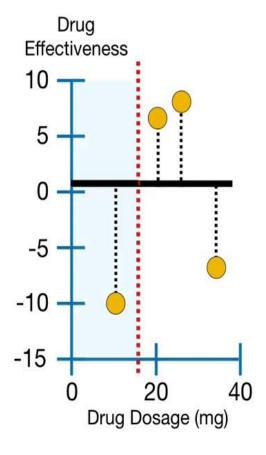
Tree Pruning
Predicted Drug

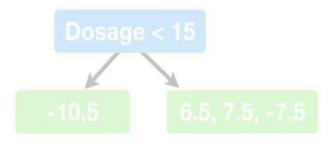
Effectiveness

0.5



Predicted Drug Effectiveness

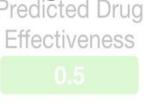


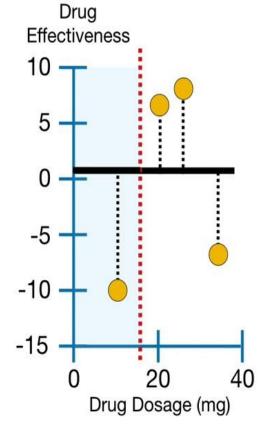


Similarity Score = 
$$\frac{\text{Sum of Residuals, Squared}}{\text{Number of Residuals } + \lambda}$$

...only this time, when we calculate **Similarity Scores**, we will set **λ** (lambda) = 1.

Predicted Drug Effectiveness

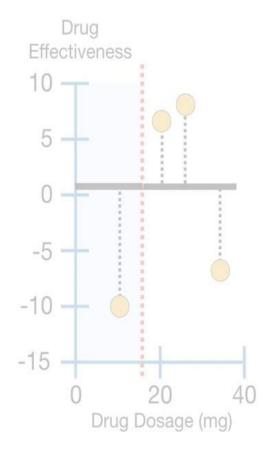


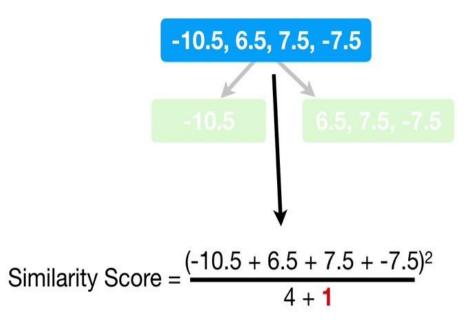




Remember  $\lambda$  (lambda) is a Regularization Parameter, which means that it is intended to reduce the prediction's sensitivity to individual observations.

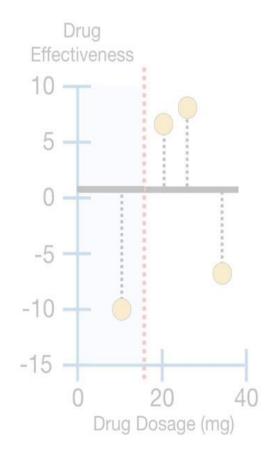
Predicted Drug Effectiveness 0.5

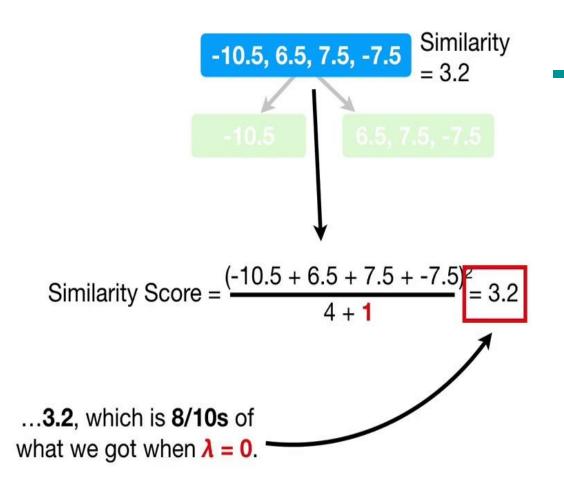




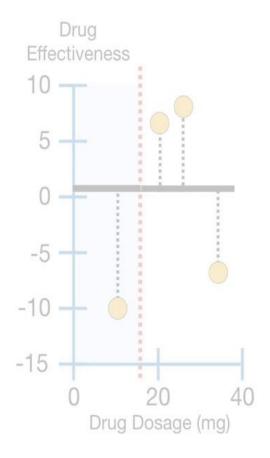
Now the **Similarity Score** for the root is...

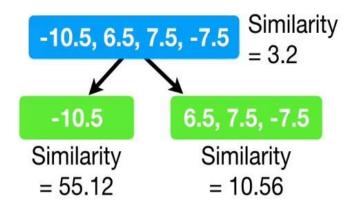
Predicted Drug
Effectiveness
0.5





Predicted Drug
Effectiveness
0.5

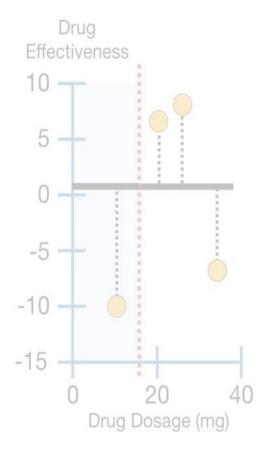


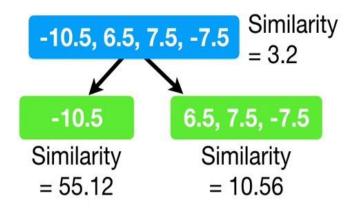


So, one thing we see is that when  $\lambda > 0$ , the **Similarity Scores** are smaller...

Predicted Drug
Effectiveness

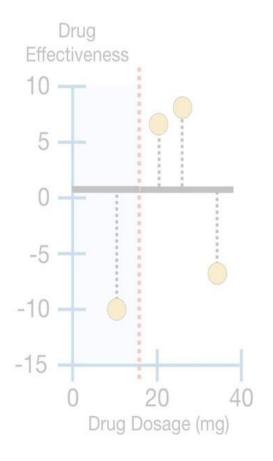
0.5

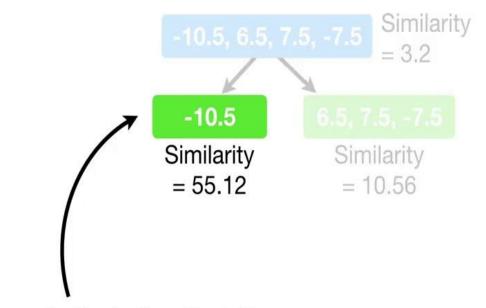




...and the amount of decrease is inversely proportional to the number of **Residuals** in the node.

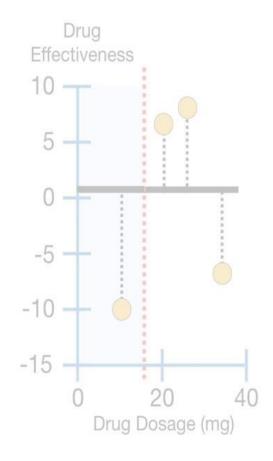
Predicted Drug
Effectiveness
0.5

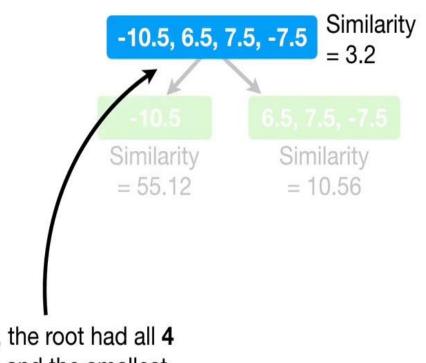




In other words, the leaf on the left had only 1 Residual, and it had the largest decrease in Similarity Score, 50%.

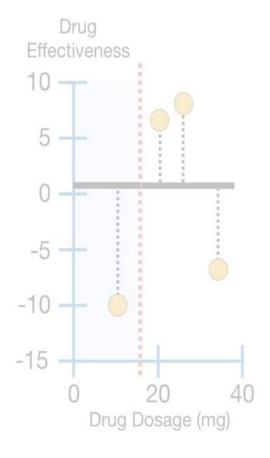
Predicted Drug
Effectiveness
0.5

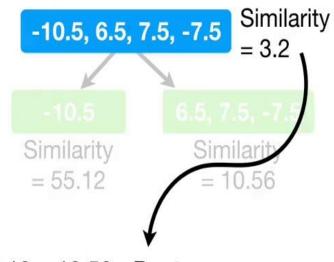




In contrast, the root had all 4 Residuals and the smallest decrease, 20%.

Predicted Drug
Effectiveness
0.5

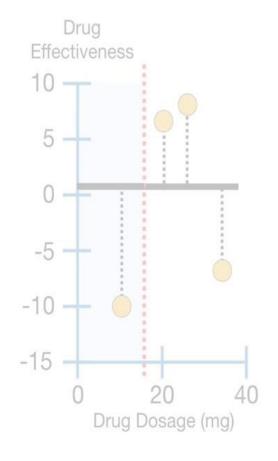


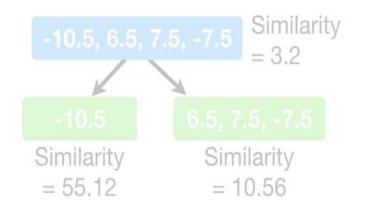


 $Gain = 55.12 + 10.56 - Root_{Similarity}$ 

Now when we calculate the **Gain**...

Predicted Drug
Effectiveness
0.5



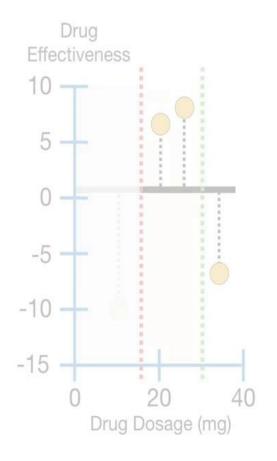


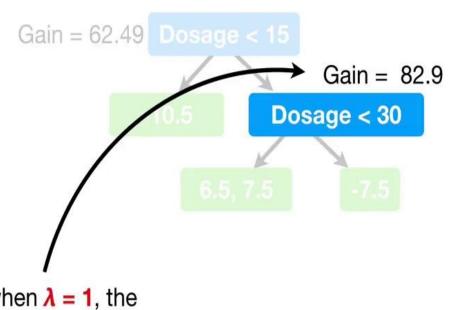
Gain = 
$$55.12 + 10.56 - 3.2 = 62.48$$

...we get **66**, which is a lot less than **120.33**, the value we got when  $\lambda = 0$ .

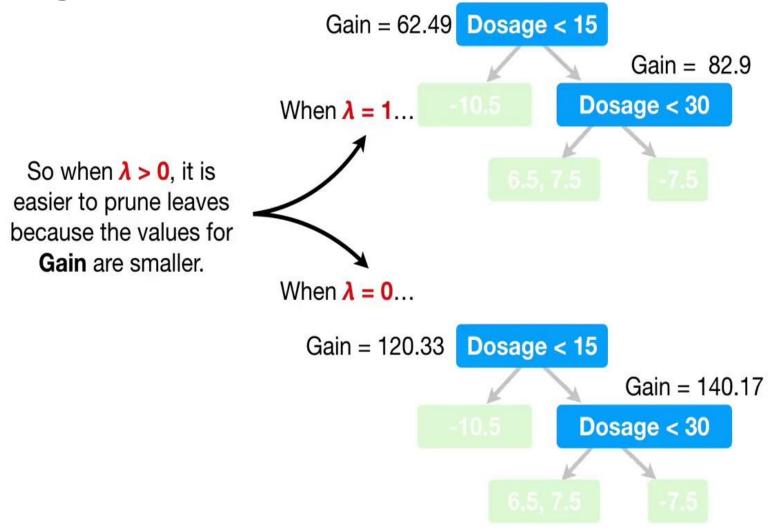
Tree Pruning
Predicted Drug

Predicted Drug Effectiveness 0.5

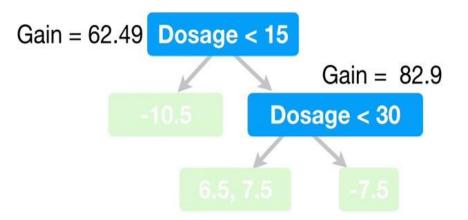


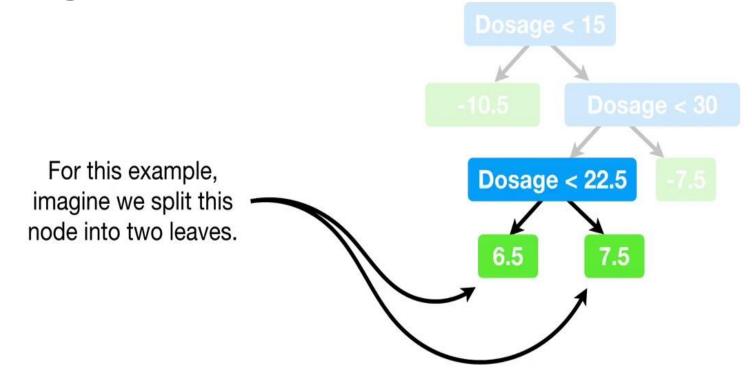


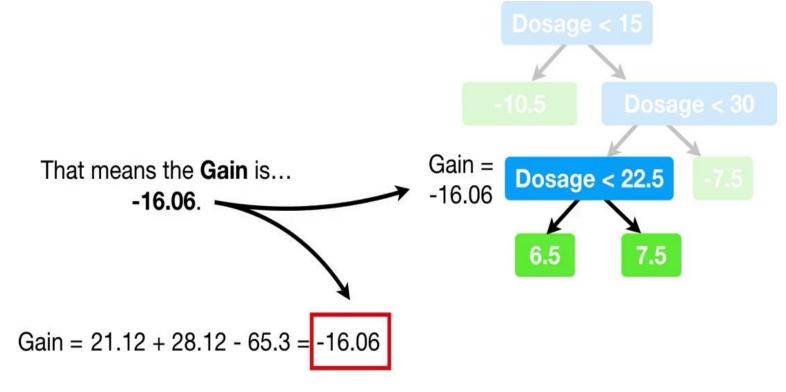
Similarly, when λ = 1, the Gain for the next branch is smaller than before.

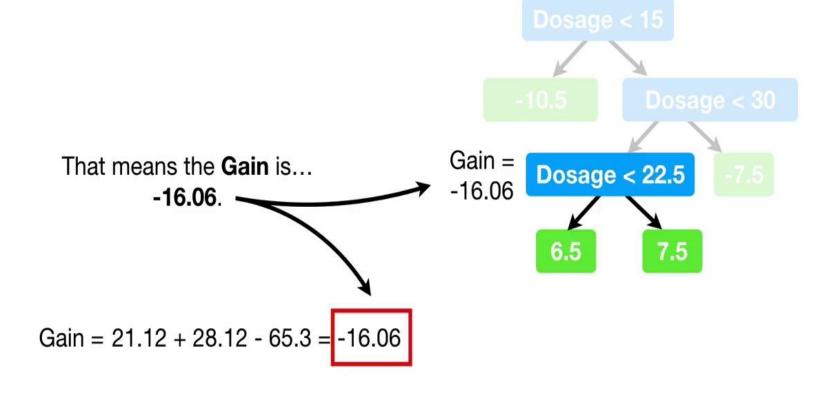


**NOTE:** Before we move on, I want illustrate one last feature of *λ* (lambda).



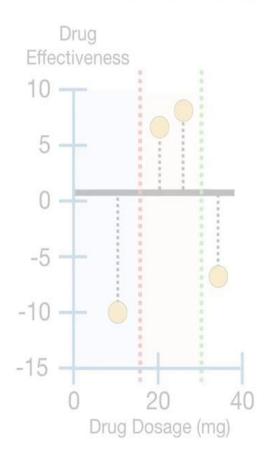


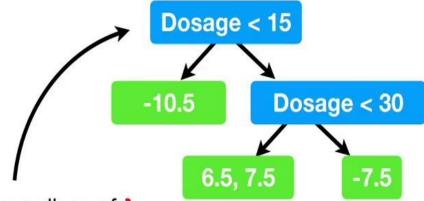




Also if we set **gamma** equals to 0 we prune this branch

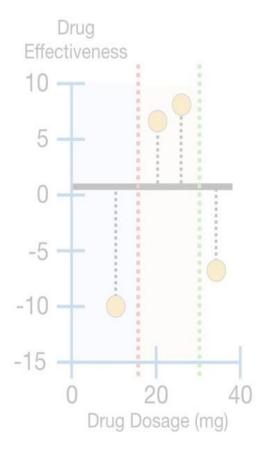
Predicted Drug Effectiveness 0.5

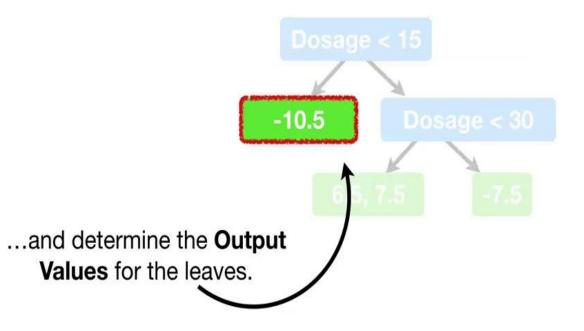




For now, regardless of λ (lambda) and γ (gamma), let's assume this is the tree we are working with...

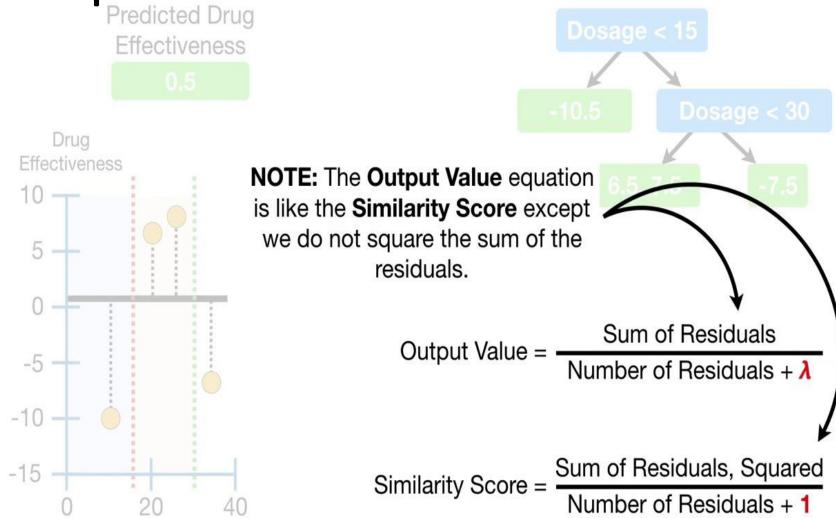
Predicted Drug
Effectiveness
0.5

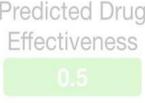


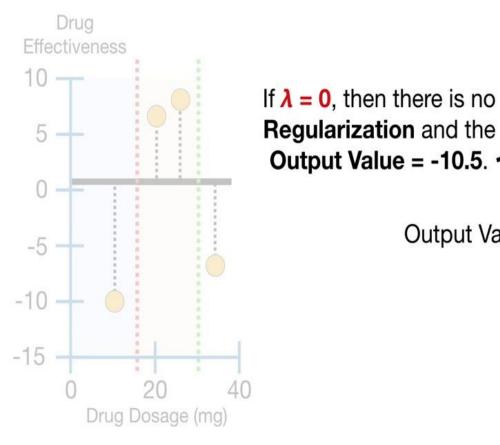


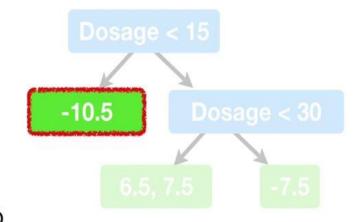
Leaves Update

Drug Dosage (mg)



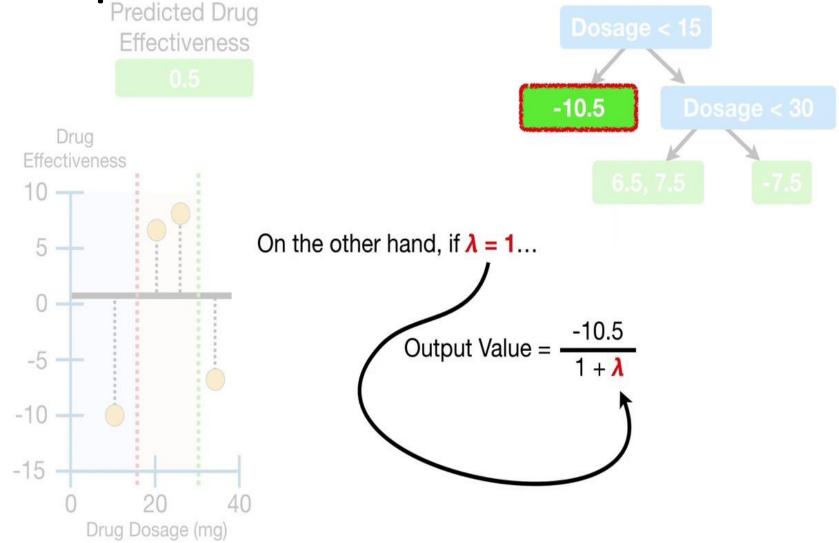


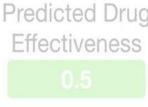


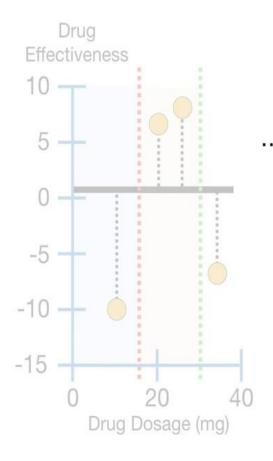


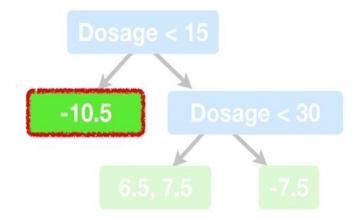
Regularization and the Output Value = -10.5.

Output Value = 
$$\frac{-10.5}{1+0}$$
 = -10.5





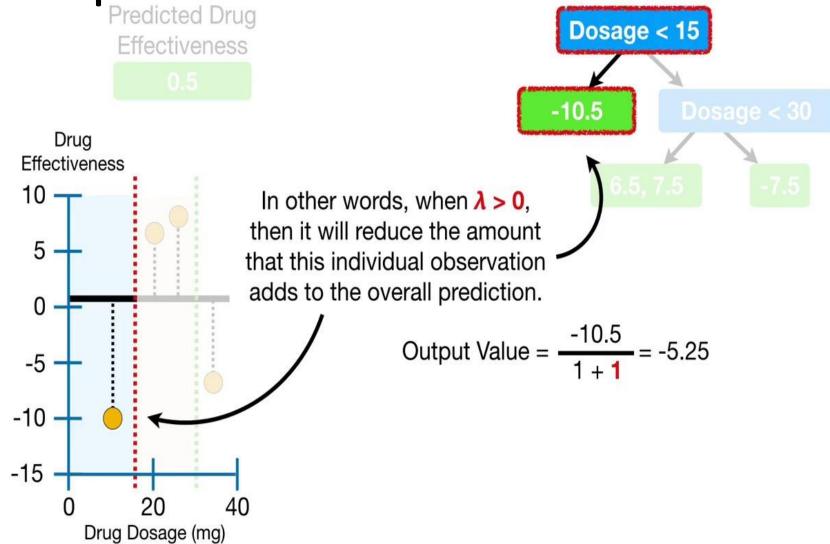


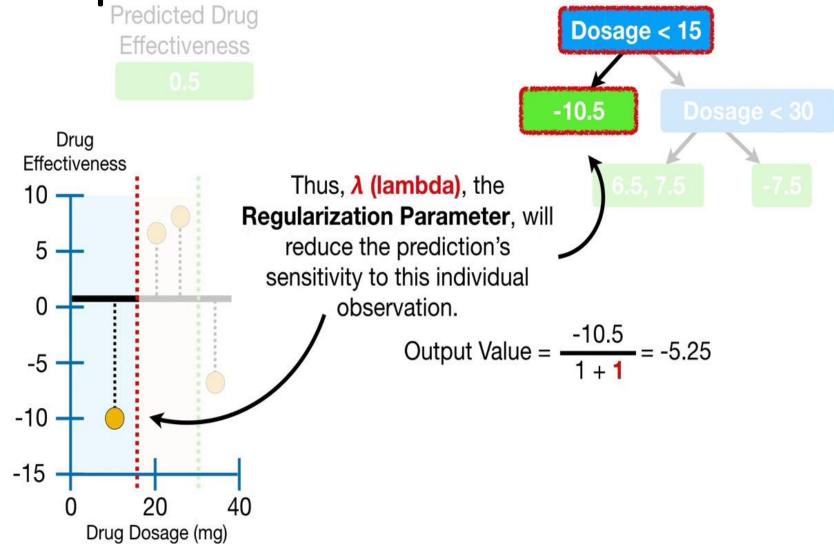


...the Output Value = -5.25.

Output Value = 
$$\frac{-10.5}{1+1}$$
 = -5.25

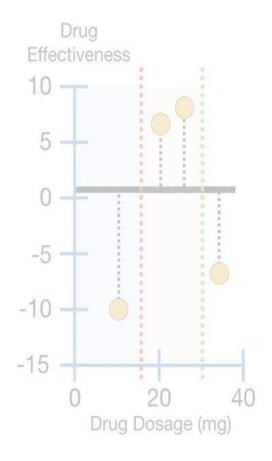
**Leaves Update** 

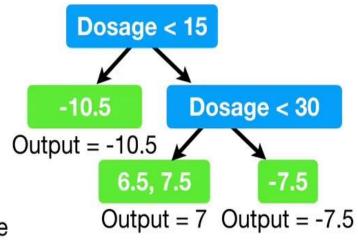




**Leaves Update** 

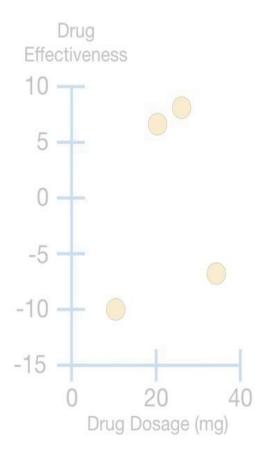
Predicted Drug Effectiveness 0.5

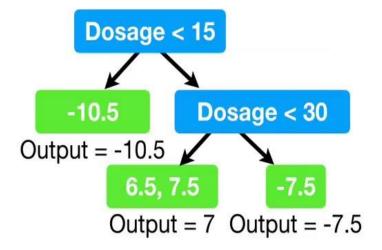




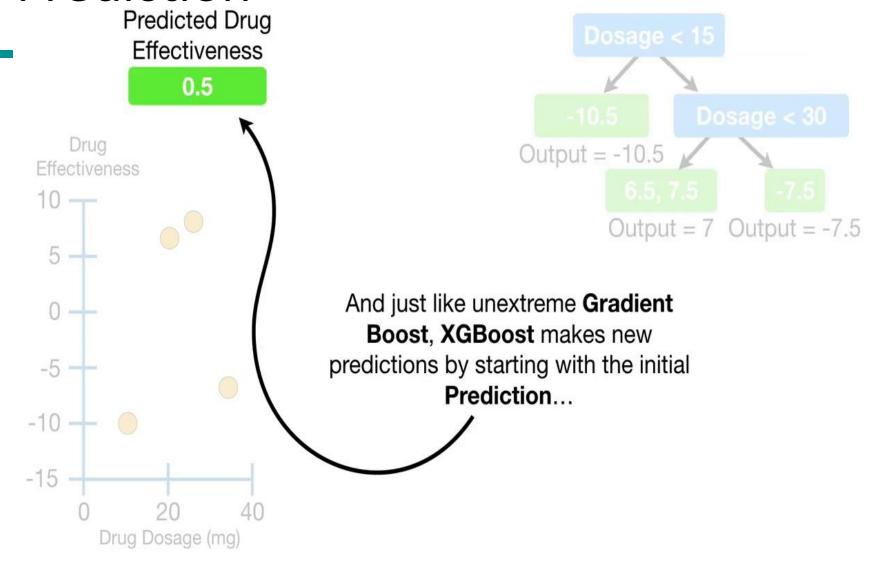
Now, at long last, the first tree is complete!

Predicted Drug
Effectiveness
0.5

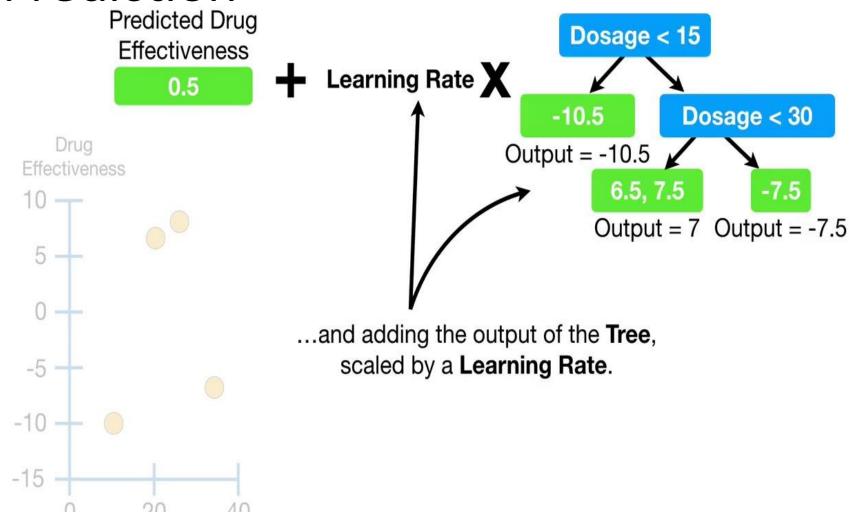




Since we have built our first tree, we can make new **Predictions**.

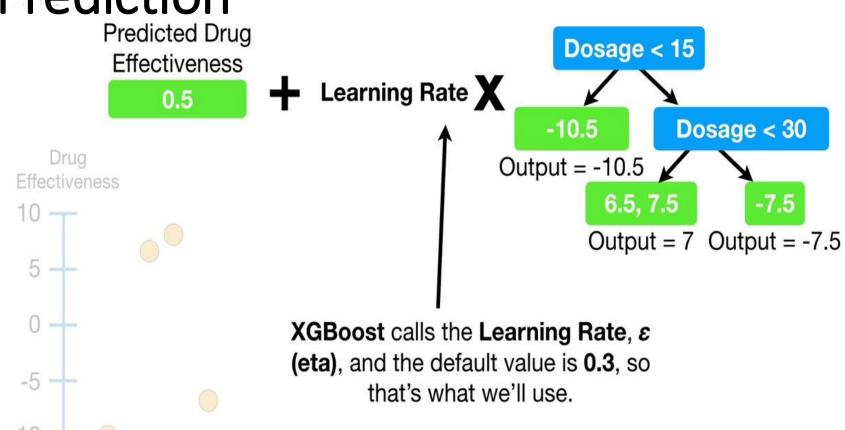


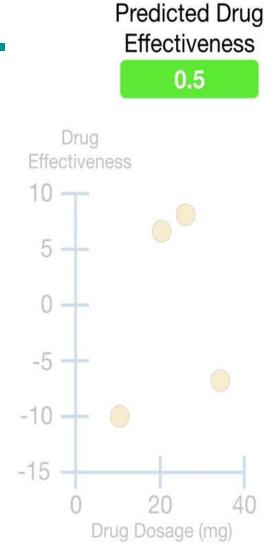
Drug Dosage (mg)

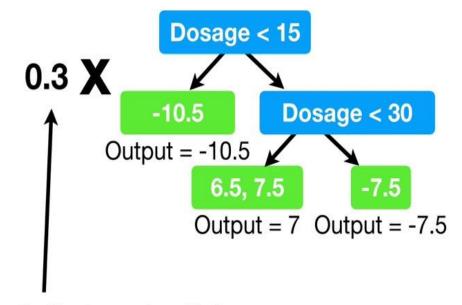


-7.5

Drug Dosage (mg)

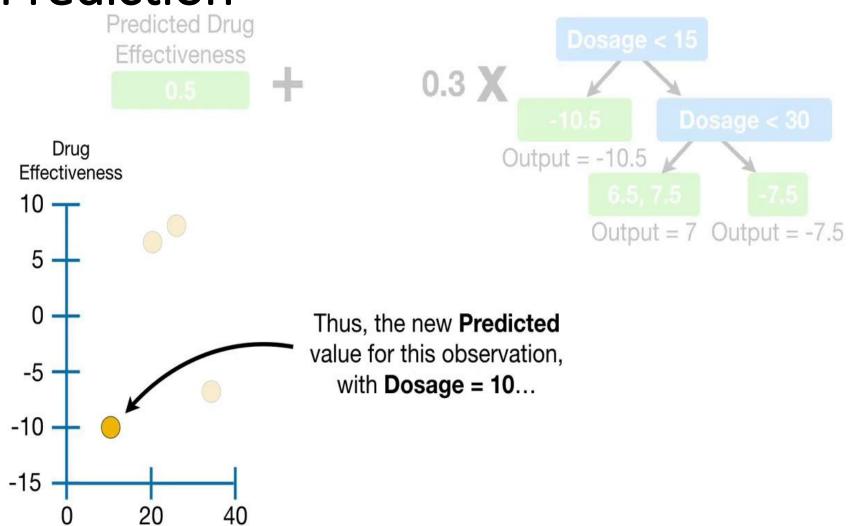


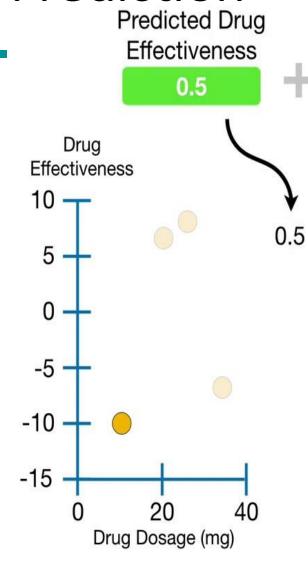


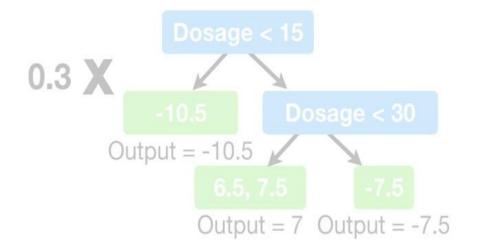


XGBoost calls the Learning Rate, ε (eta), and the default value is 0.3, so that's what we'll use.

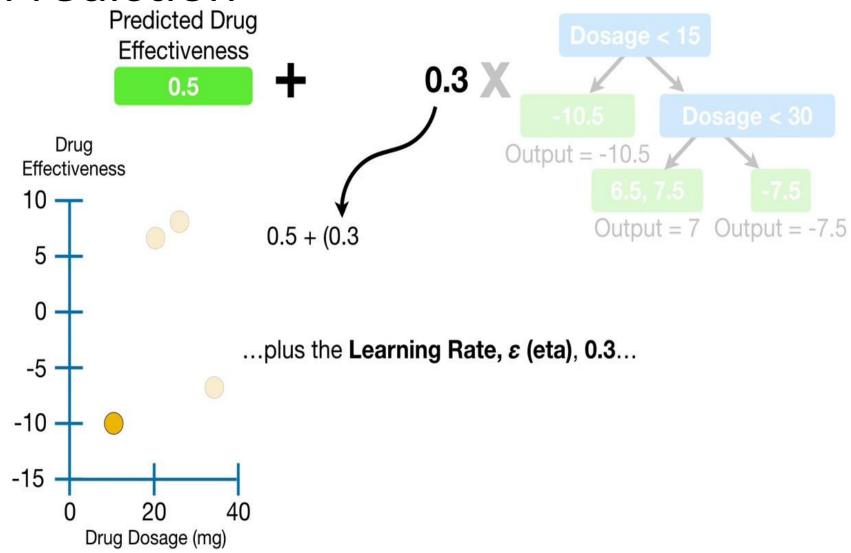
Drug Dosage (mg)







...is the original prediction, **0.5**...

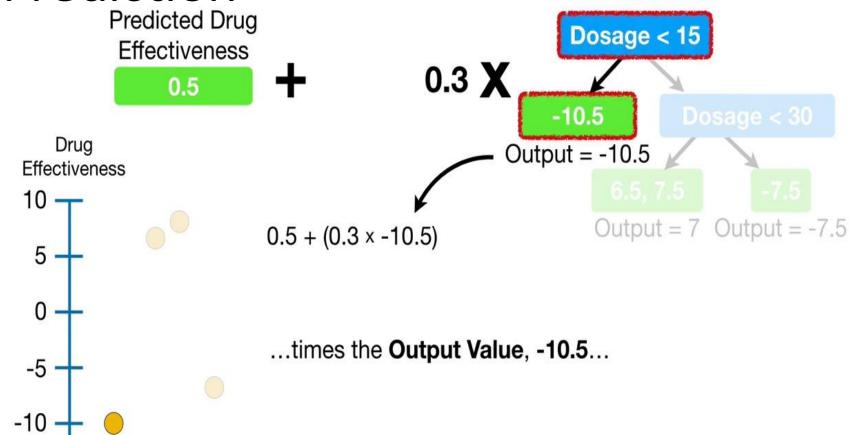


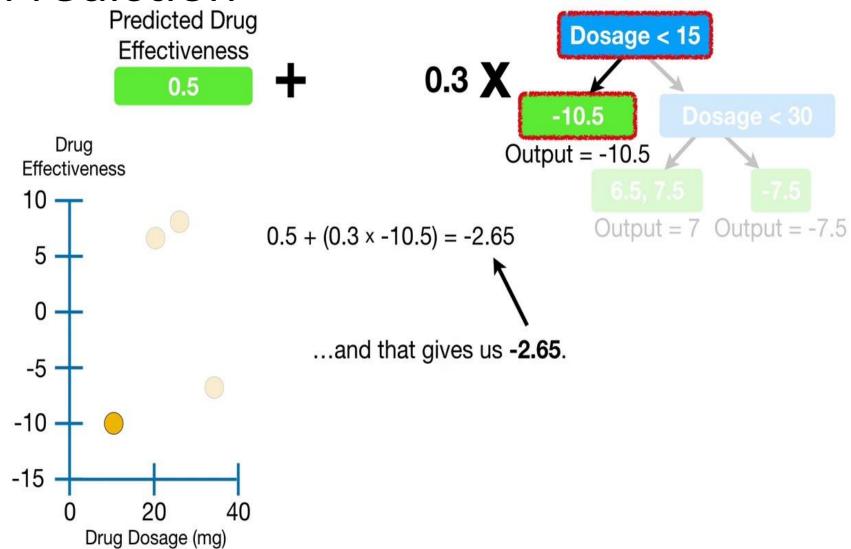
-15

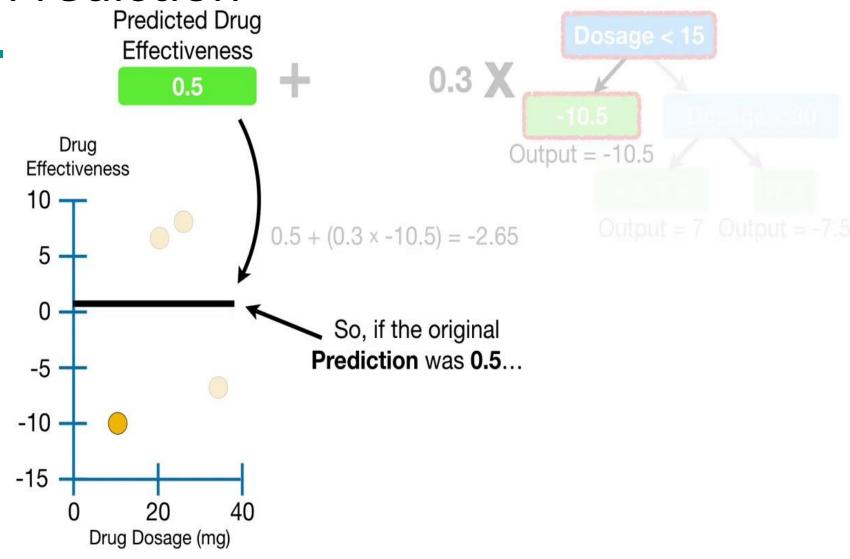
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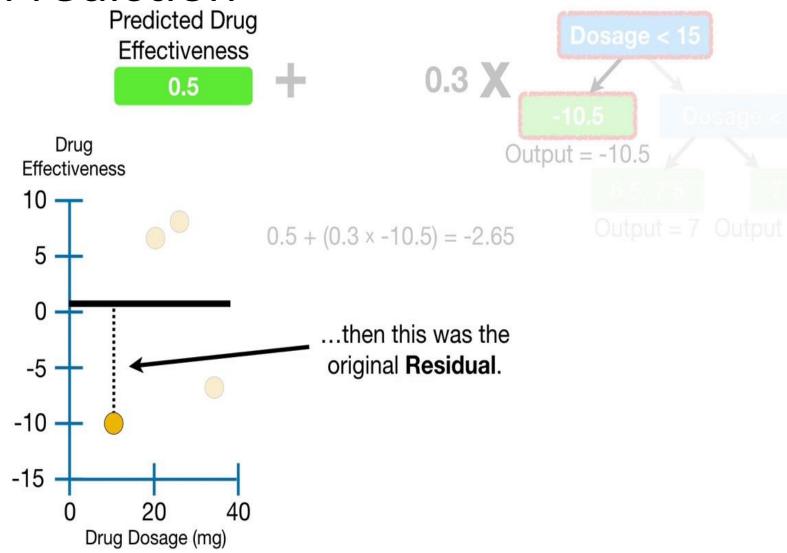
Drug Dosage (mg)

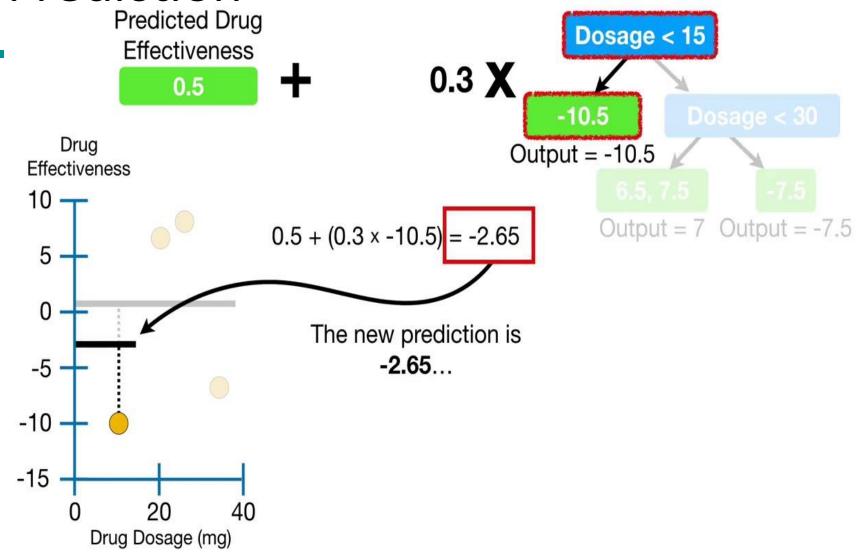
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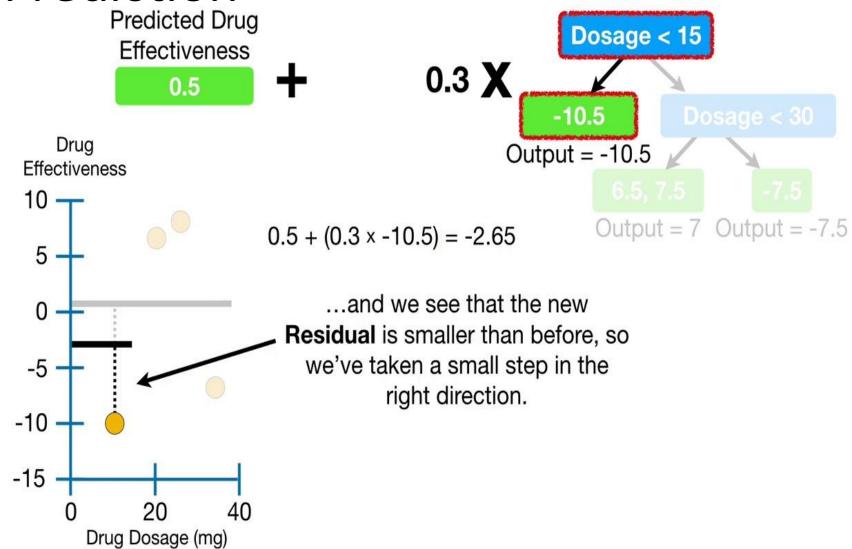










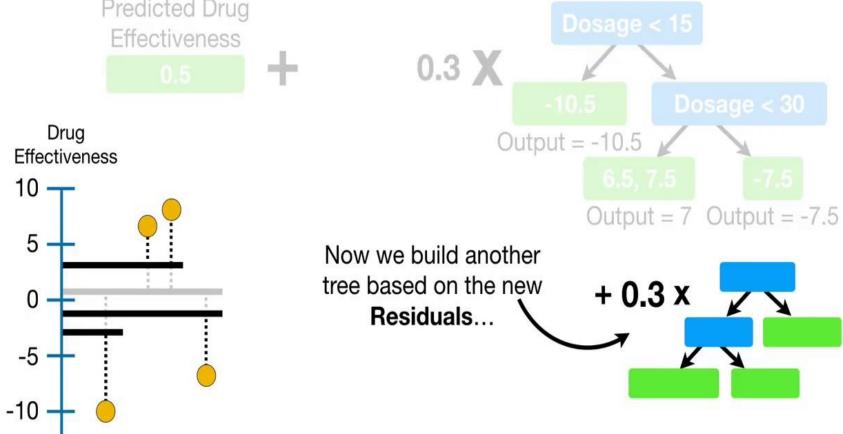


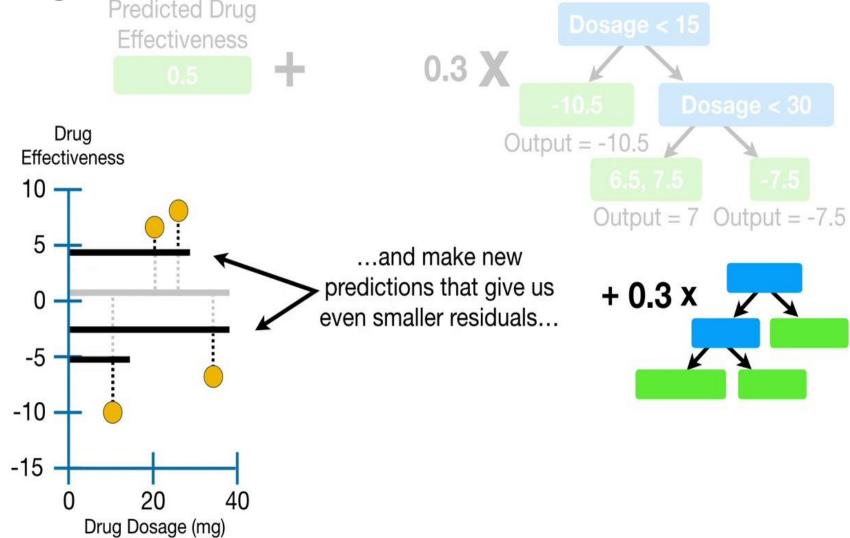
-15

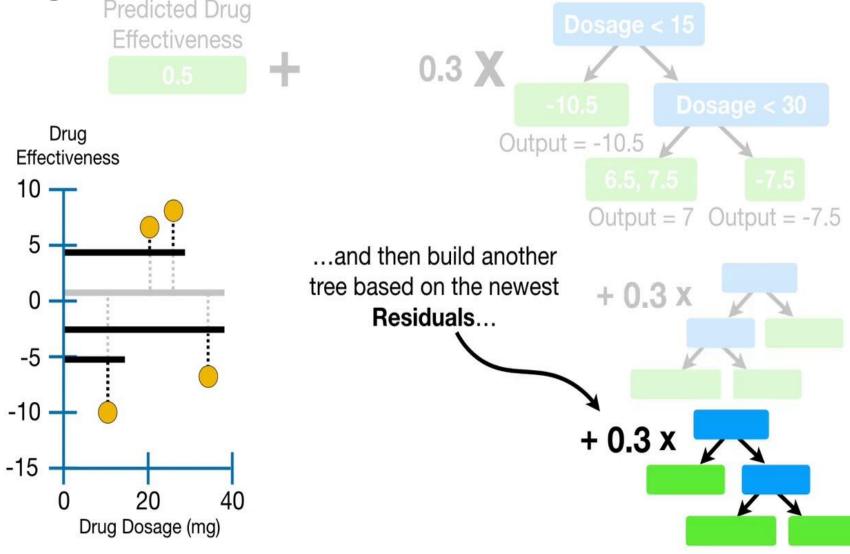
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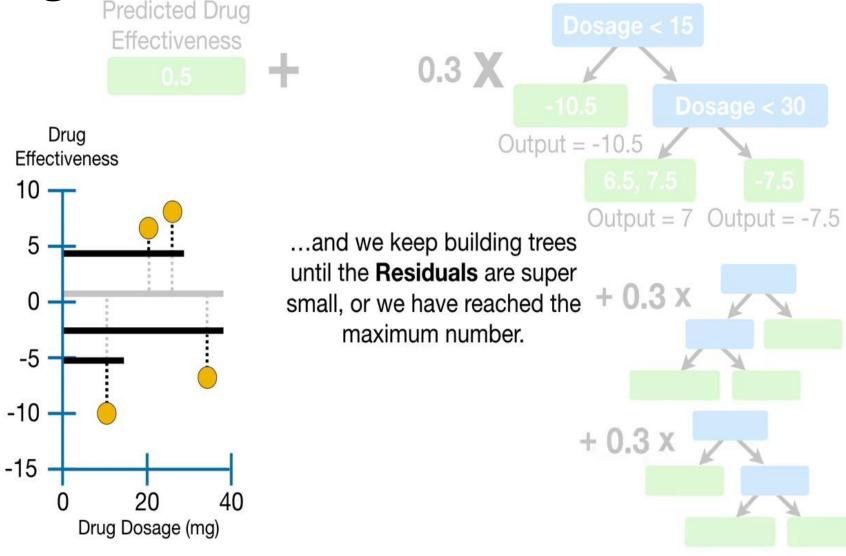
Drug Dosage (mg)

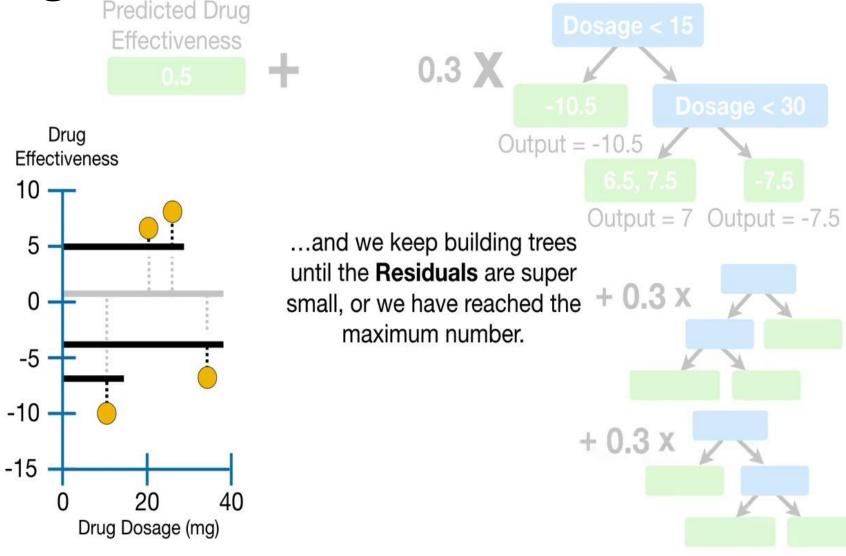
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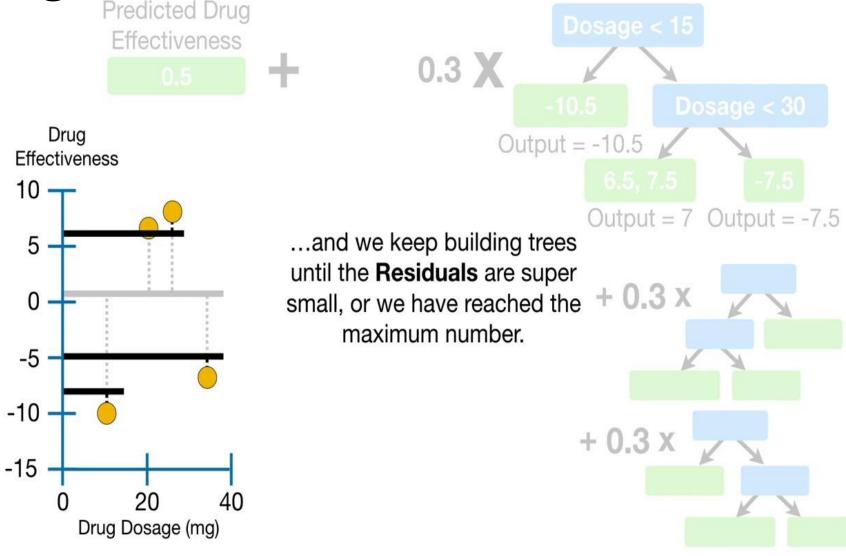


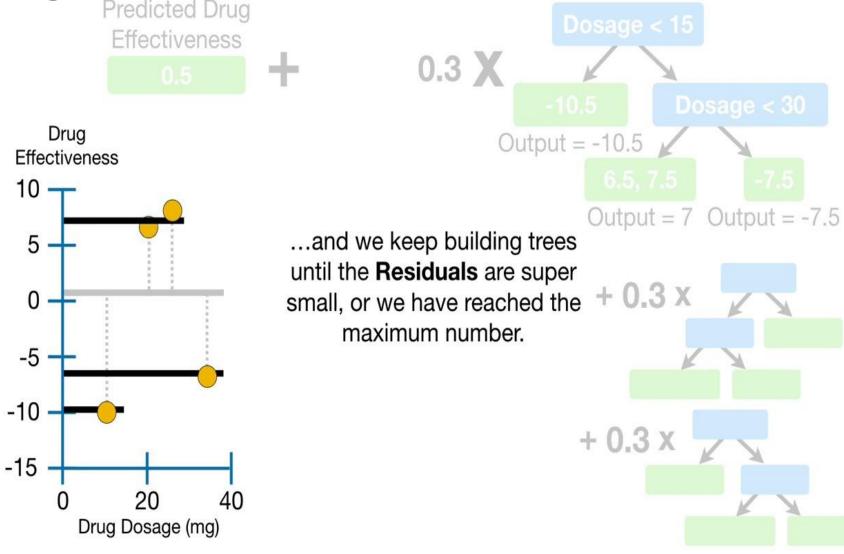




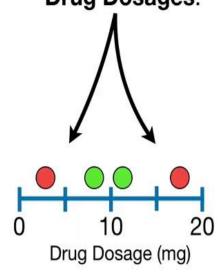


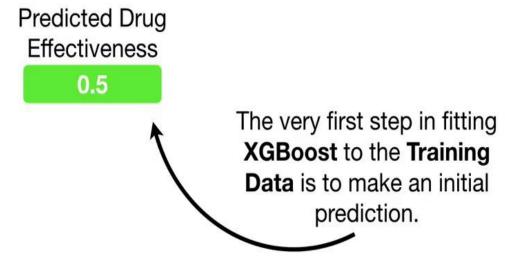


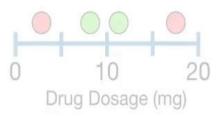




to keep the examples from getting out of hand, we will use this super simple **Training Data** consisting of **4** different **Drug Dosages**.

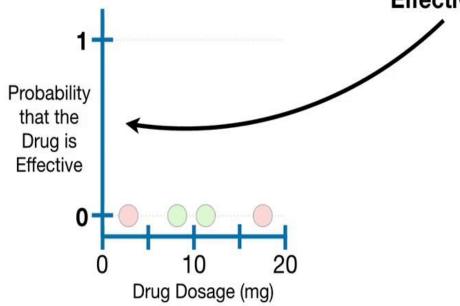


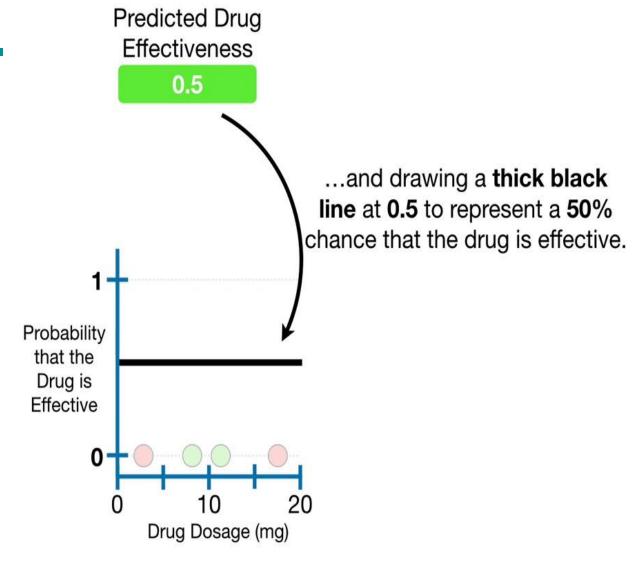






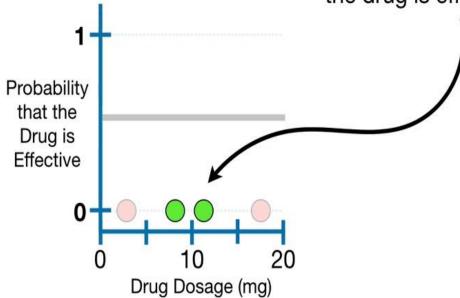
We can illustrate the initial prediction by adding a *y*-axis to our graph to represent the **Probability that the Drug is Effective...** 





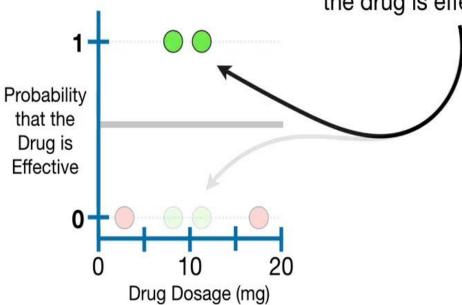
Predicted Drug Effectiveness

Since these two **Green Dots** represent effective dosages, we will move them to the top of the graph, where the probability that the drug is effective is **1**.



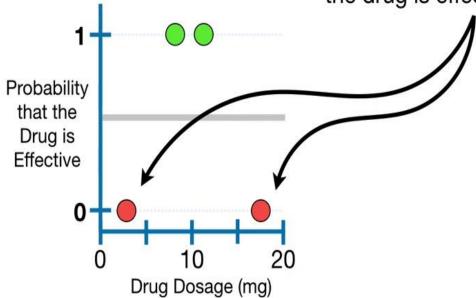
Predicted Drug
Effectiveness
0.5

Since these two **Green Dots** represent effective dosages, we will move them to the top of the graph, where the probability that the drug is effective is **1**.



Predicted Drug
Effectiveness

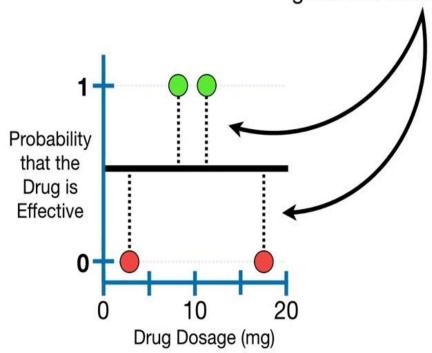
These two **Red Dots** represent ineffective dosages, so we will leave them at the bottom of the graph, where the probability that the drug is effective is **0**.

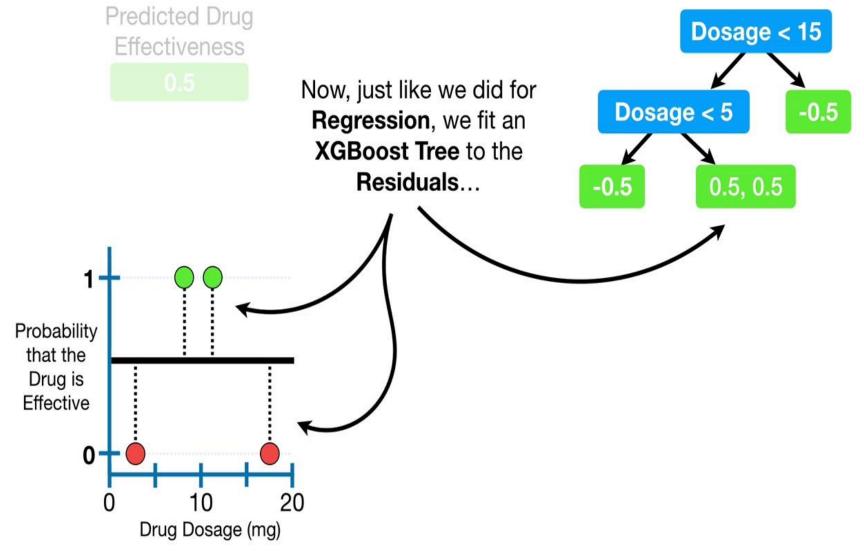


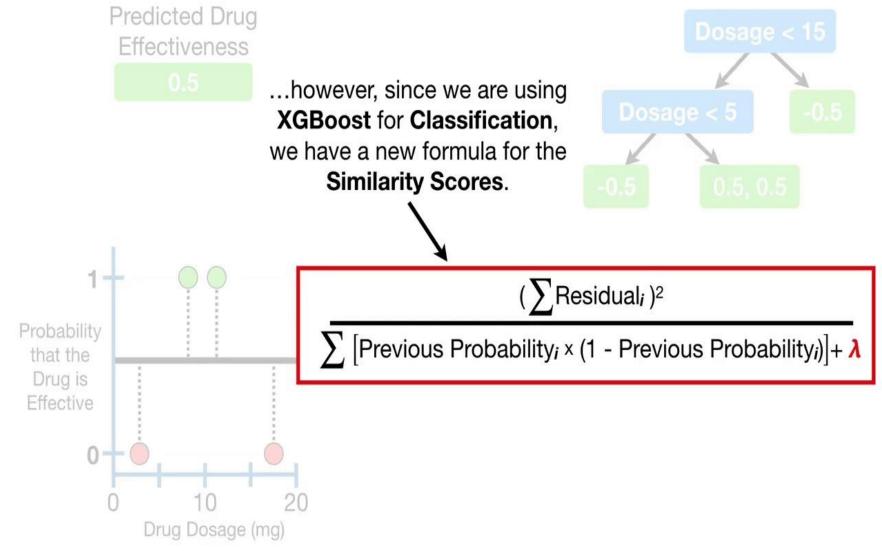
Predicted Drug Effectiveness

0.5

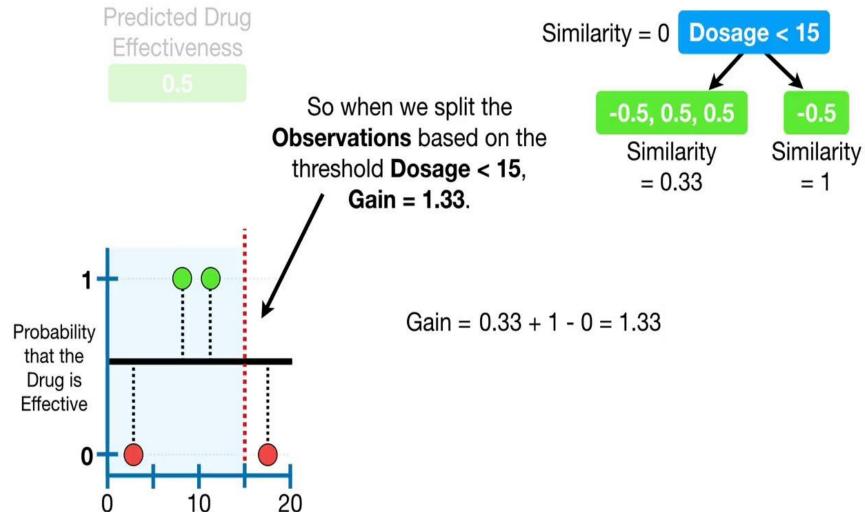
The **Residuals**, the differences between the **Observed** and **Predicted** values, show us how good the initial prediction is.







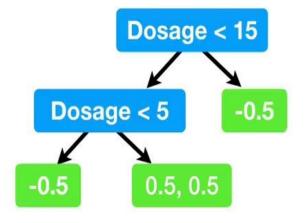
Drug Dosage (mg)

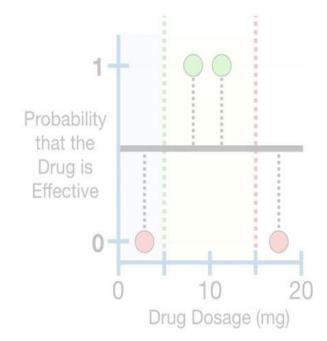


Predicted Drug
Effectiveness

0.5

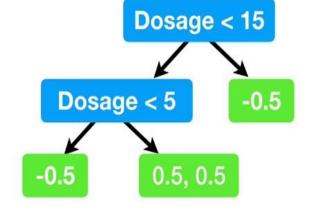
Now, since I'm limiting trees to 2 levels, we will not split this leaf any further, and we are done building this tree.

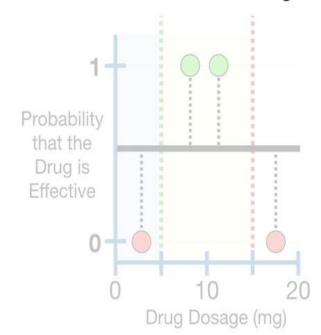


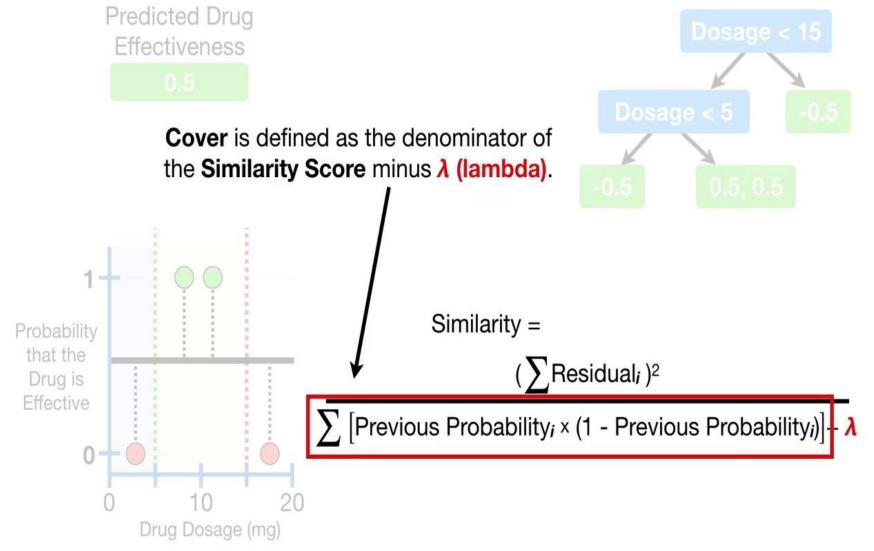


Predicted Drug
Effectiveness
0.5

The minimum number of **Residuals** in each leaf is determined by calculating something called **Cover**.

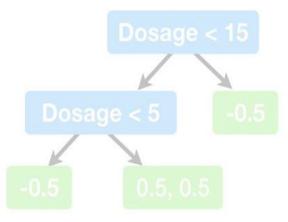


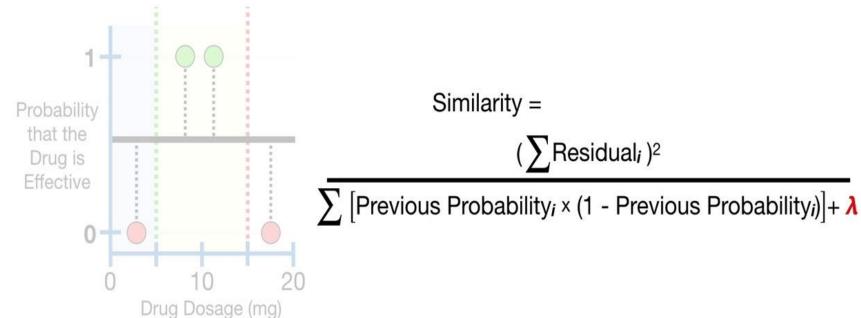


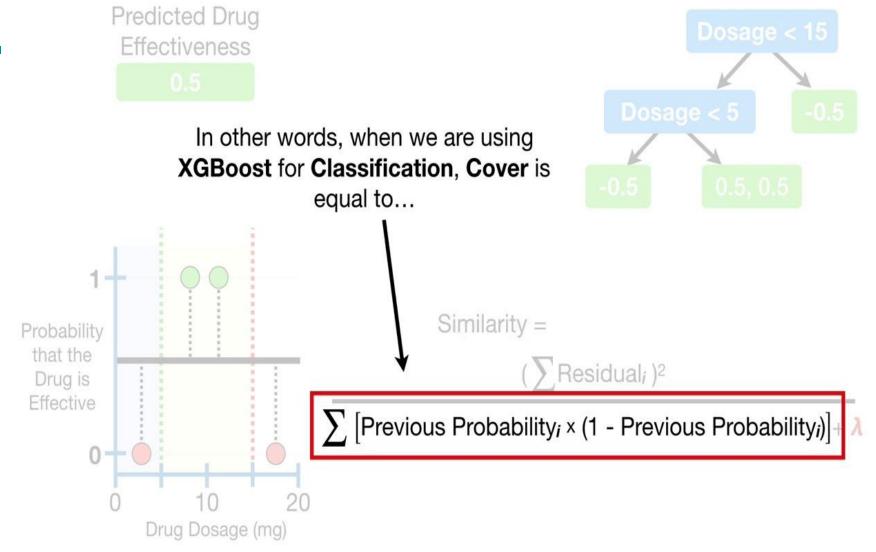


Predicted Drug
Effectiveness
0.5

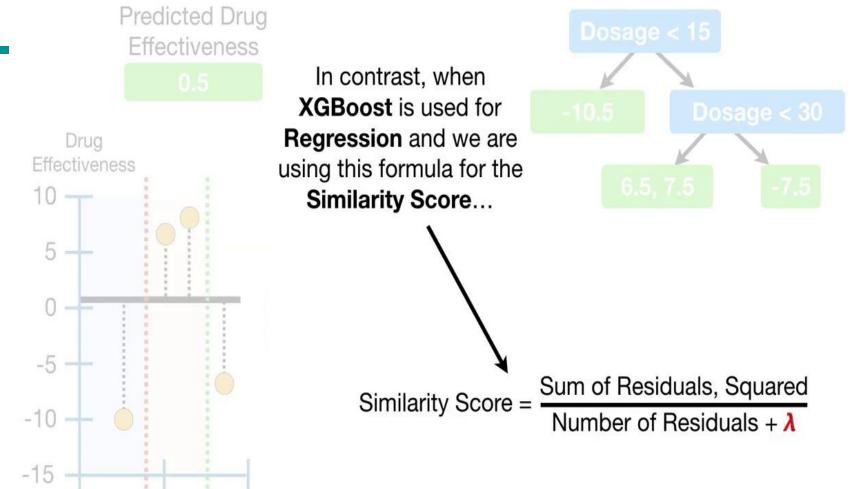
In other words, when we are using XGBoost for Classification, Cover is equal to...

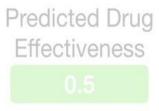


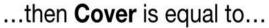


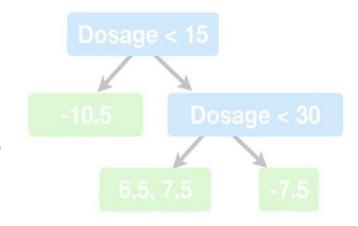


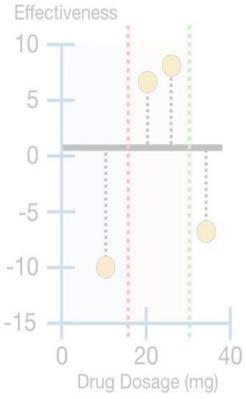
Drug Dosage (mg)





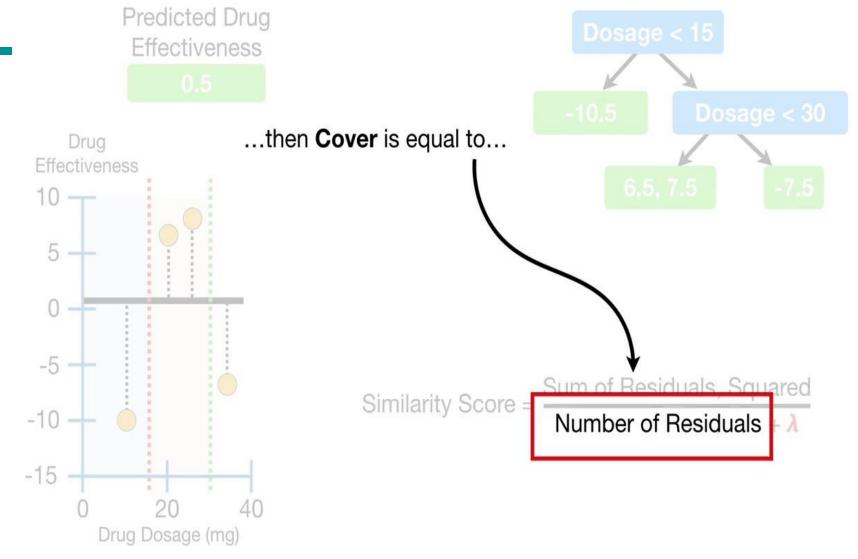


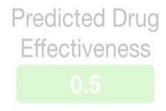


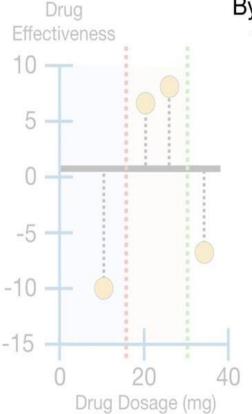


Drug

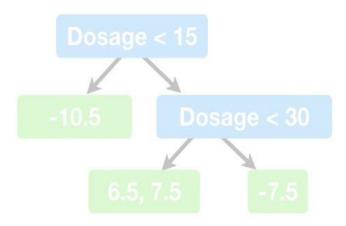
Similarity Score = 
$$\frac{\text{Sum of Residuals, Squared}}{\text{Number of Residuals } + \lambda}$$







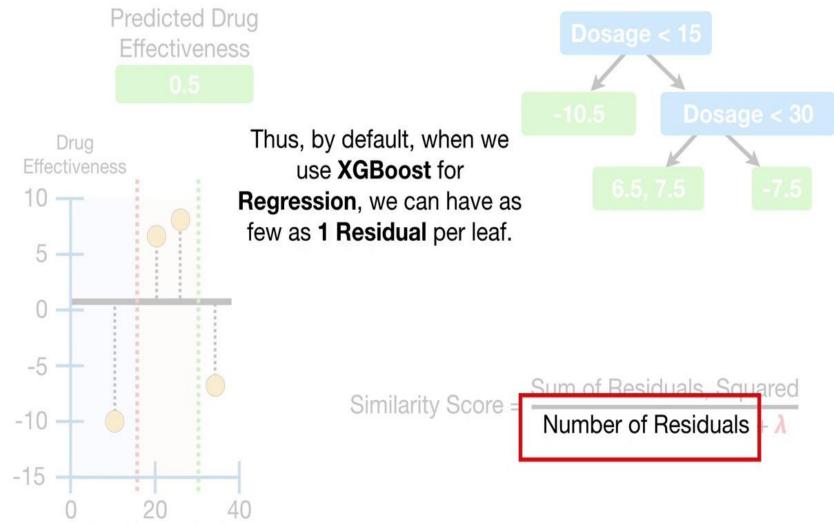
By default, the minimum value for **Cover** is **1**.

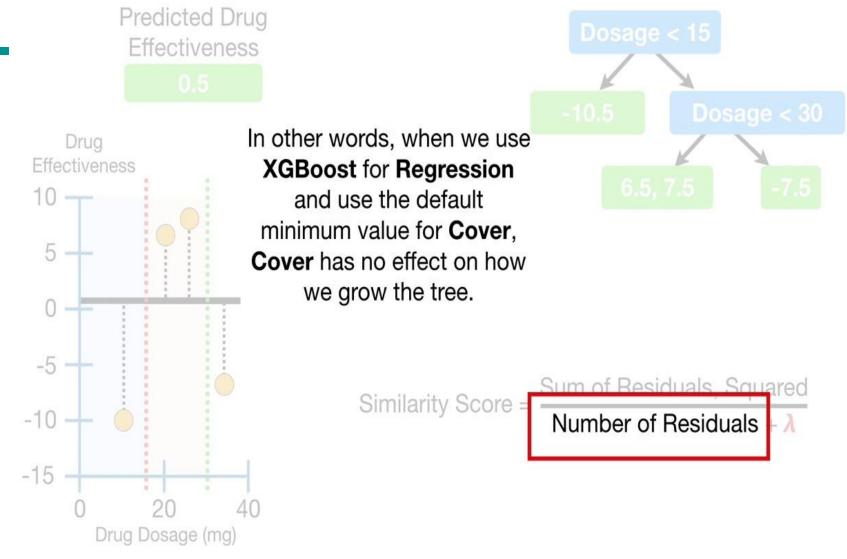


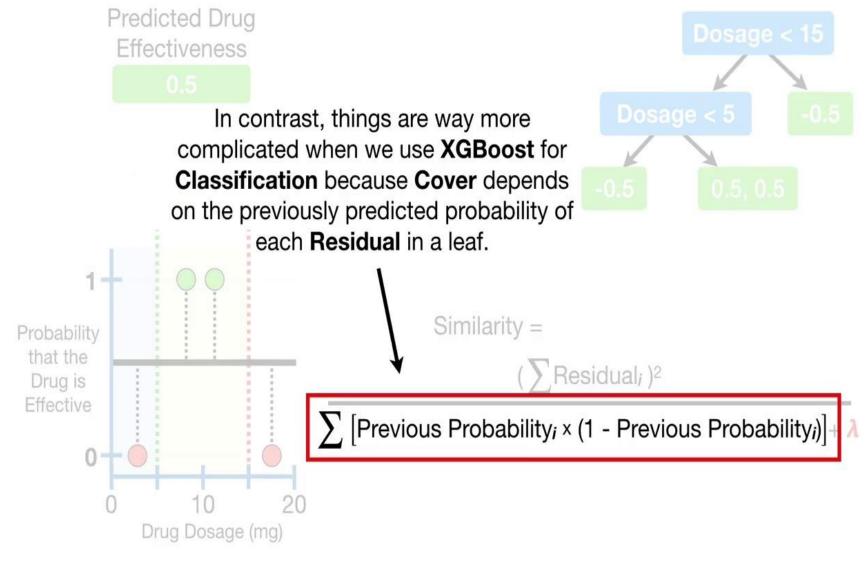
Similarity Score = Sum of Residuals, Squared

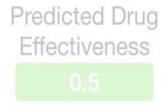
Number of Residuals + \(\lambda\)

Drug Dosage (mg)

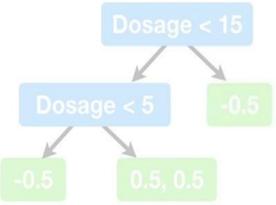


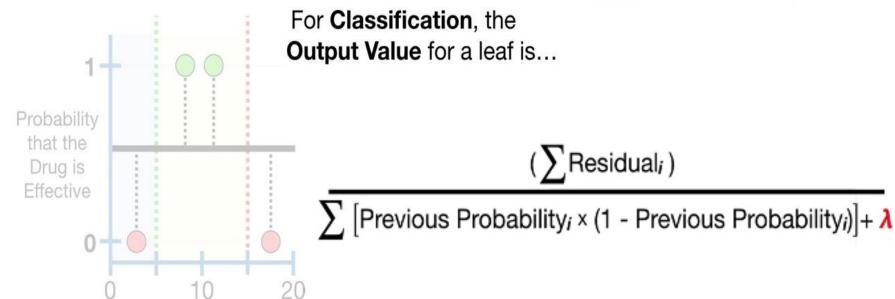


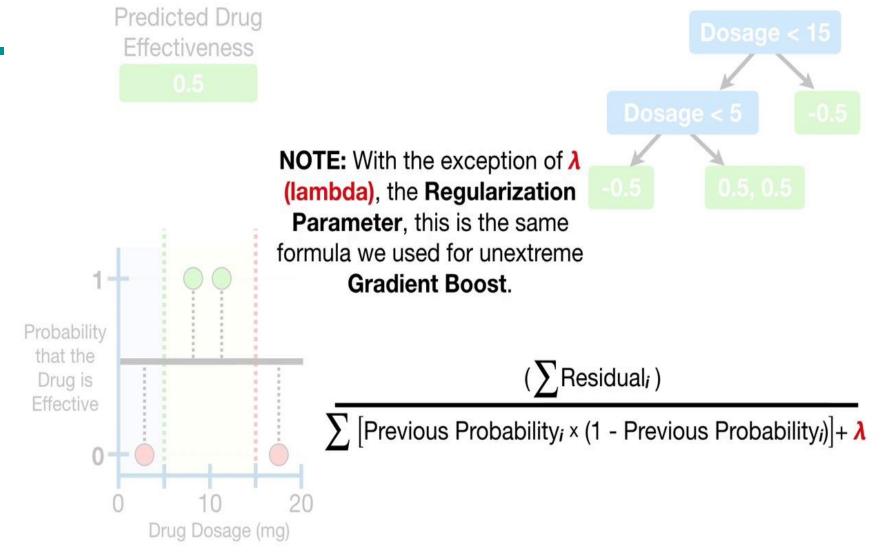


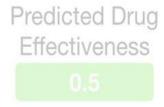


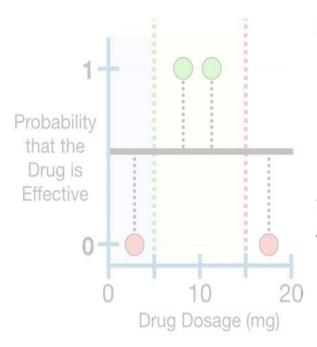
Drug Dosage (mg)



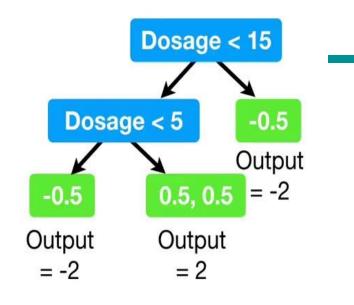








The first tree is complete!



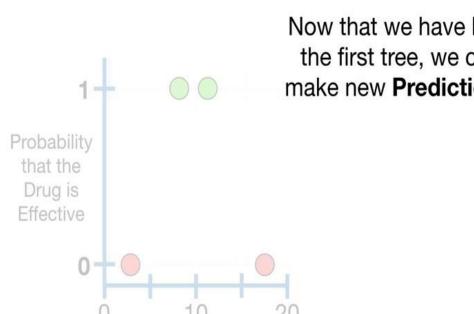
$$-0.5/(0.5 \times (1-0.5) + 0) = -2$$

 $(\sum \mathsf{Residual}_i)$ 

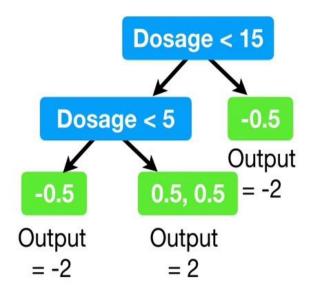
 $\sum [Previous Probability_i \times (1 - Previous Probability_i)] + \lambda$ 

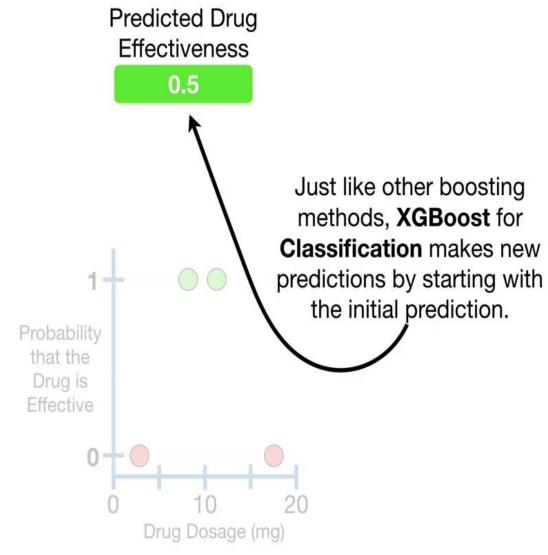
Predicted Drug Effectiveness

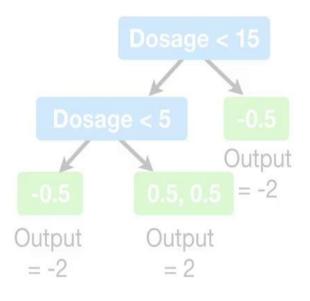
> Now that we have built the first tree, we can make new Predictions.

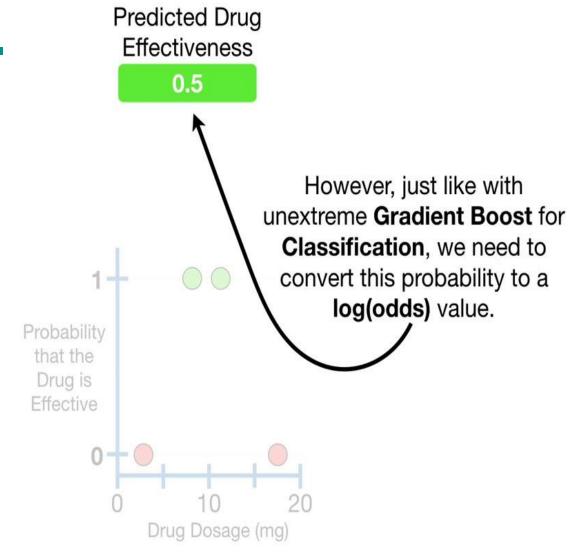


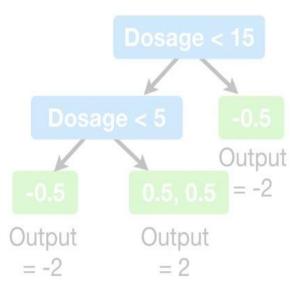
Drug Dosage (mg)

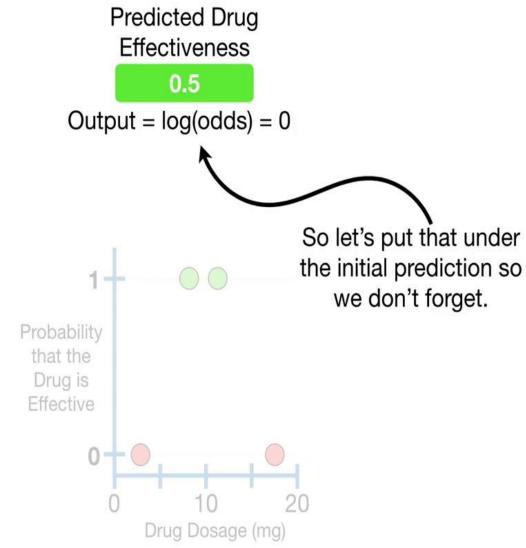


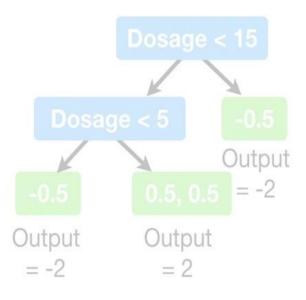


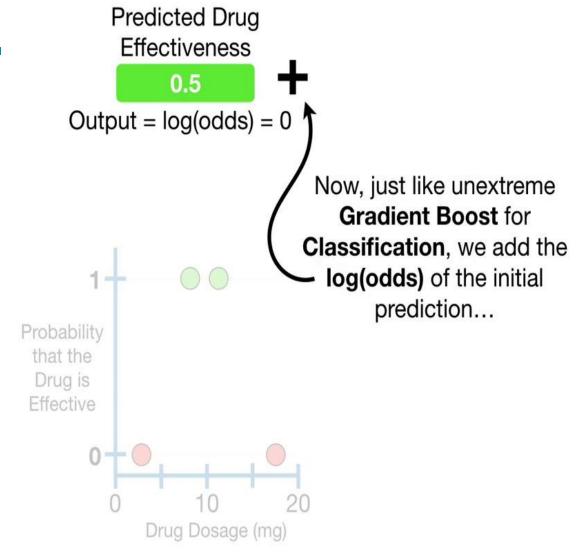


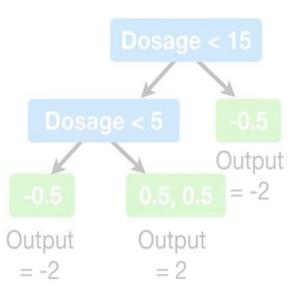


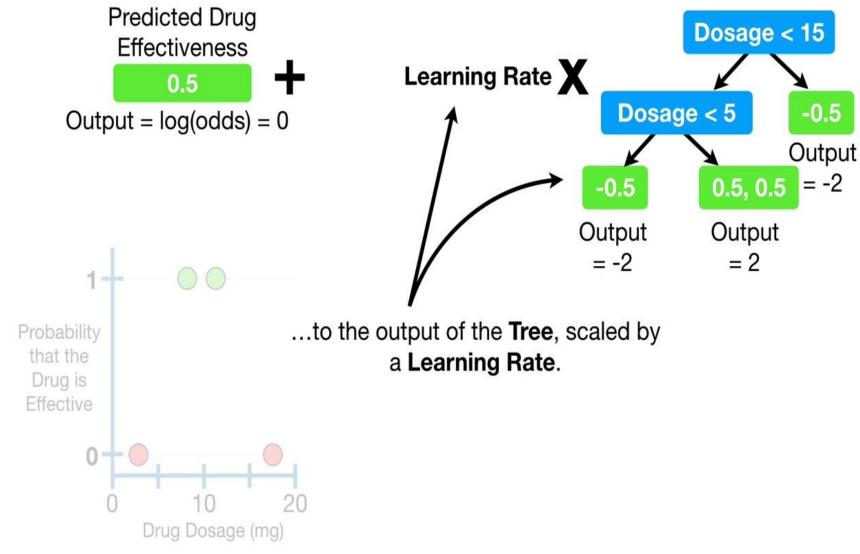




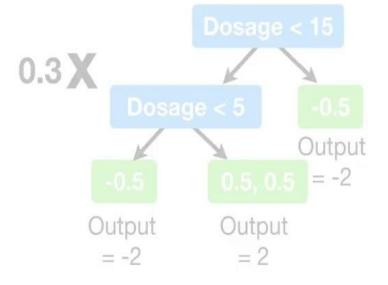


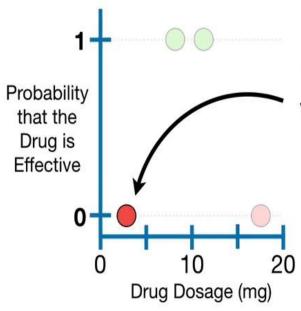




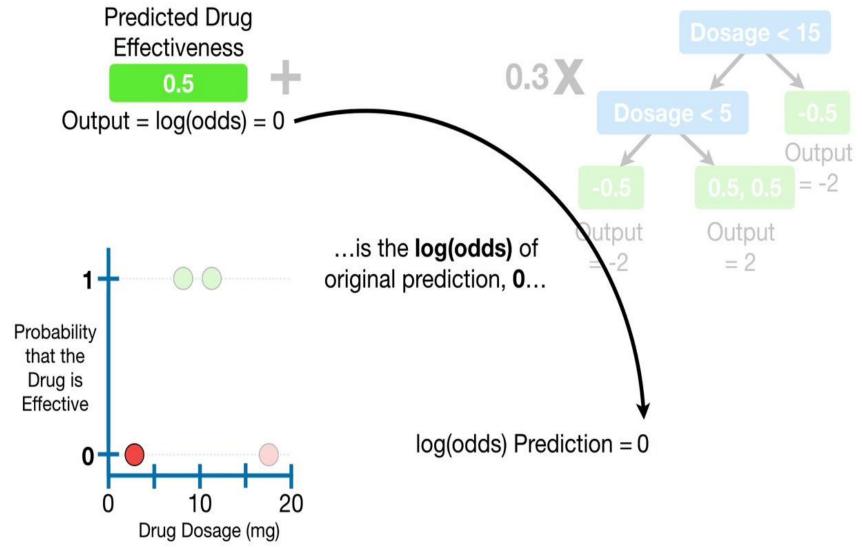


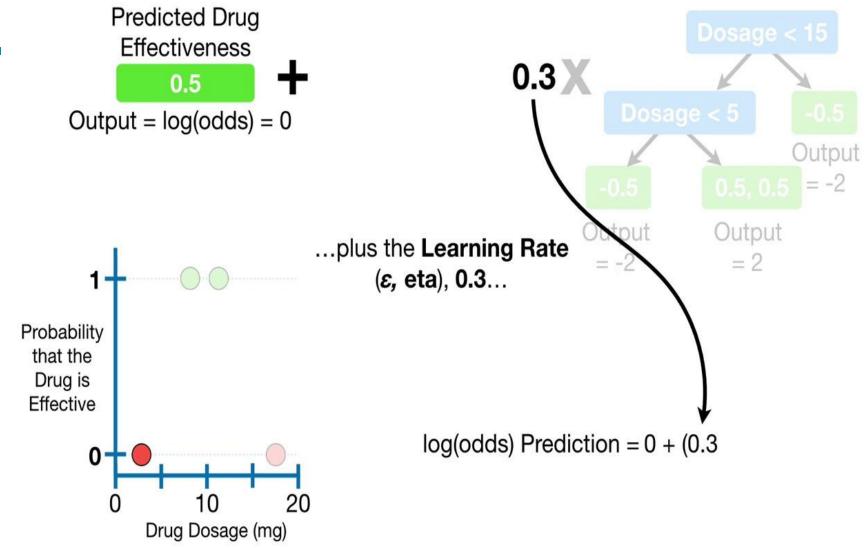


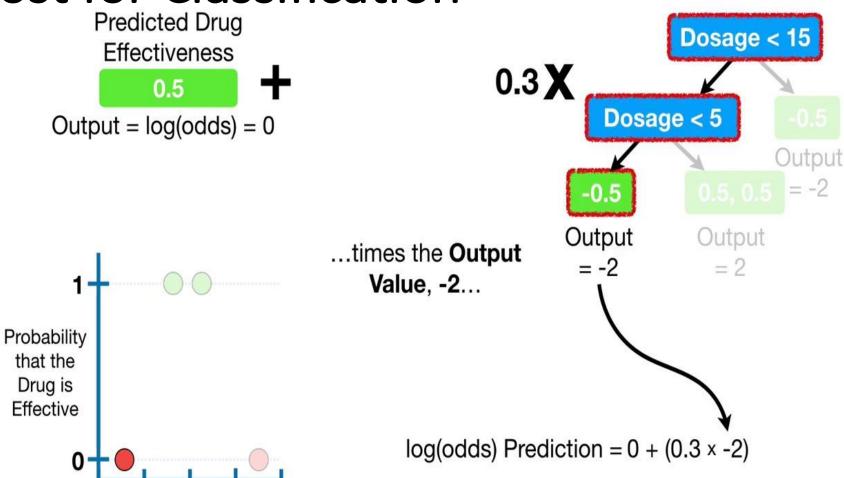




Thus, the new **Predicted** value for this observation, with **Dosage = 2...** 



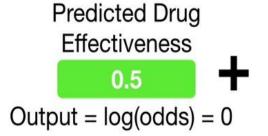


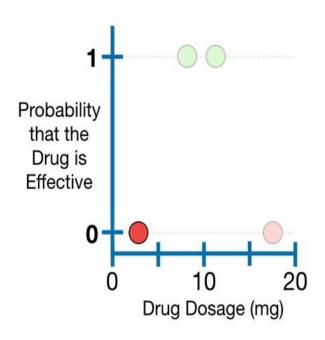


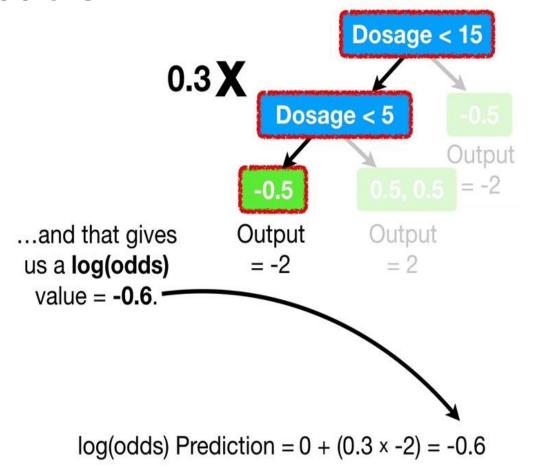
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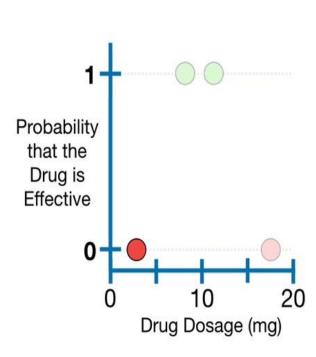
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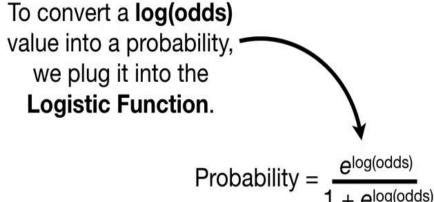
Drug Dosage (mg)



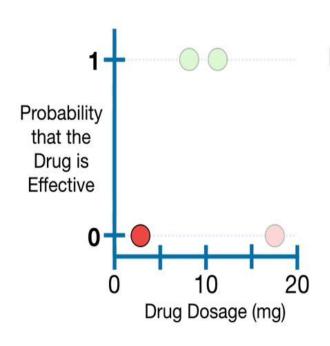


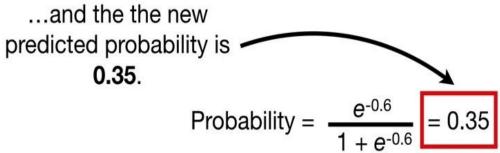




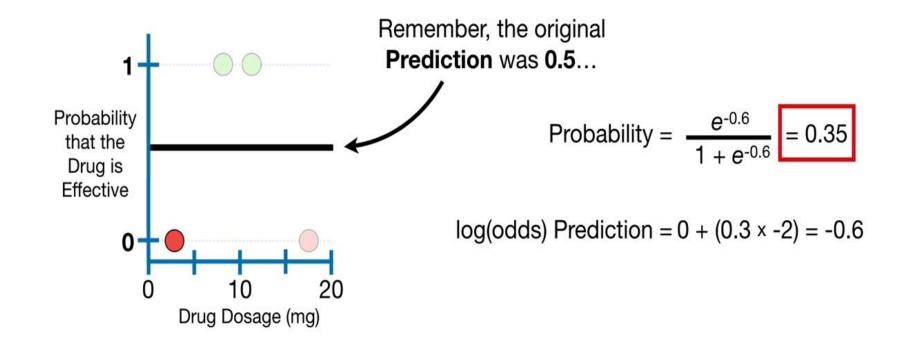


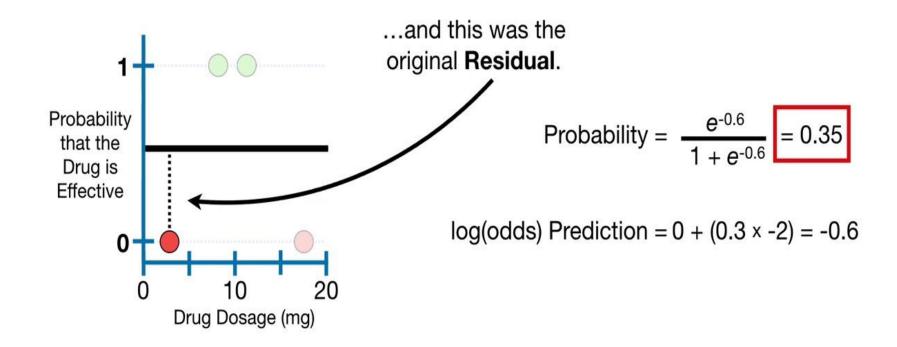
$$log(odds)$$
 Prediction = 0 +  $(0.3 \times -2)$  = -0.6

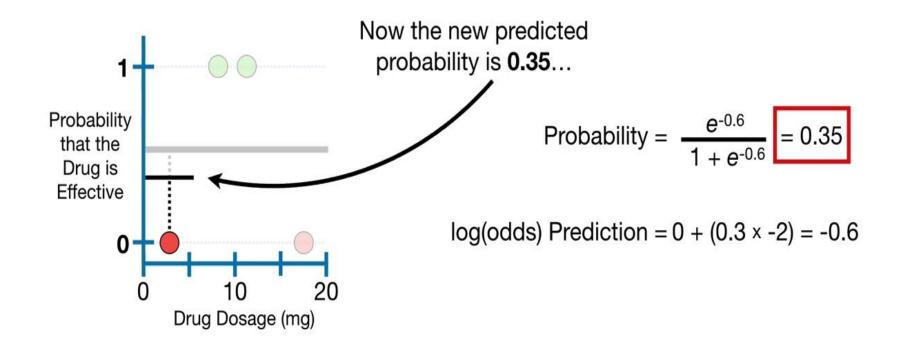


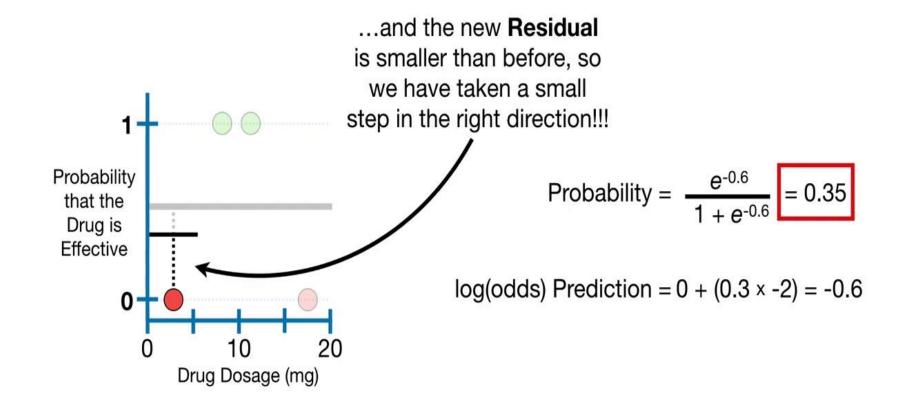


$$log(odds)$$
 Prediction = 0 +  $(0.3 \times -2)$  = -0.6







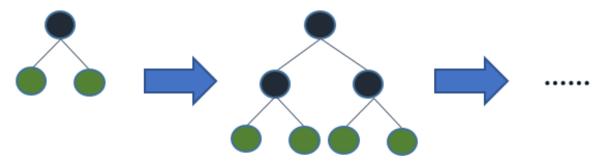


## LightGBM

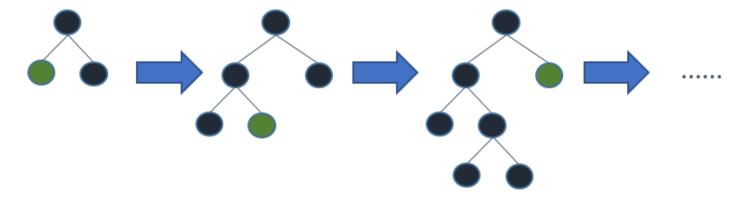
- LightGBM (by Microsoft) is a distributed high-performance framework that uses decision trees for classification, and regression tasks.
- Advantage w.r.t. XGBoost
  - Faster training speed and accuracy resulting from LightGBM being a histogram-based algorithm that performs bucketing of values (also requires lesser memory)
  - Compatible with large and complex datasets but is much faster during training
  - Support for both parallel learning and GPU learning

## LightGBM vs XGBoost

• XGBoost: level-wise (horizontal) growth



• LightGBM: out leaf-wise (vertical) growth



• LightGBM is significantly faster than XGBoost but delivers almost equivalent performance

# Gradient-Based One-Side Sampling (GOSS)

- In Gradient Boosted Decision Trees, the data instances have no native weight which is leveraged by GOSS.
- Data instances with larger gradients contribute more towards information gain.
- To maintain the accuracy of the information, GOSS retains instances with larger gradients and performs random sampling on instances with smaller gradients.

# **Exclusive Feature Bundling (EFB)**

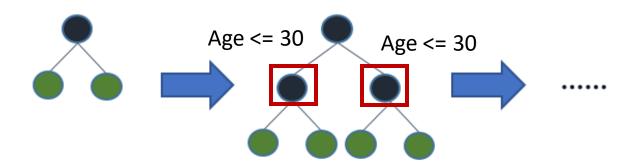
- Exclusive Feature Bundling is a near lossless method to reduce the number of effective features.
- Just like One-Hot encoded features, in the sparse space, many features rarely take non-zero values simultaneously.

# LightGBM and XGBoost

- Handling Categorical Features
- Handling Missing Values

#### **CatBoost**

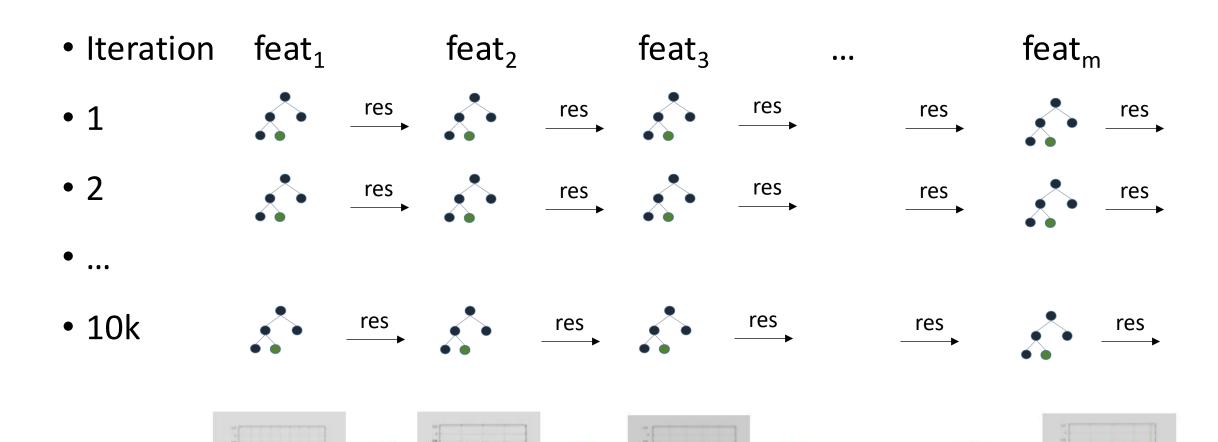
- Categorial Boosting
- Gradient Boosting Machine
- Adopts Ordered Target Encoding procedure for categorical attributes
- Rely on Symmetric Decision Trees, i.e., dedicion trees with the same splitting conditions for nodes at the same lavel



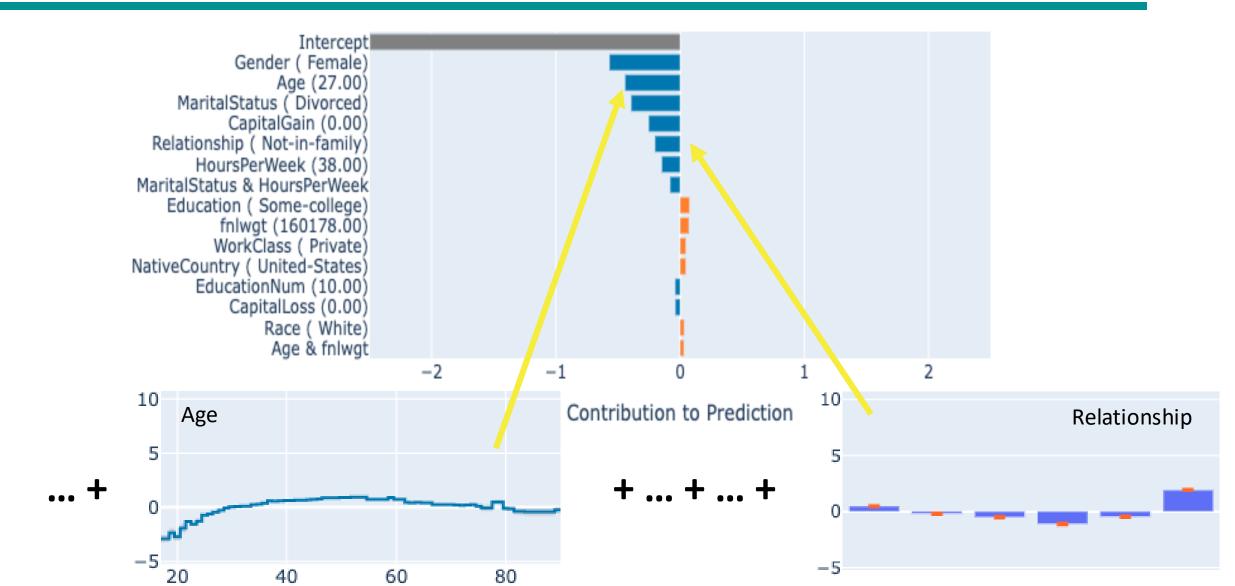
# Explainable Boosting Machines (EMB)

- Type of Generalized Additive Models (GAMS)
- Additive Model:  $y = f_1(x_1) + f_2(x_2) + ... + f_m(x_m)$
- Linear Model:  $y = \beta_1 x_1 + \beta_2 x_2 + ... + \beta_m x_m$
- Complex Model:  $y = f(x_1, x_2, ..., x_m)$

## EBM Algorithm Sketch



## **EBM Prediction Time**



#### References

- Mason, L., Baxter, J., Bartlett, P., & Frean, M. (1999). Boosting algorithms as gradient descent. Advances in neural information processing systems, 12.
- Friedman, J. H. (2002). Stochastic gradient boosting. *Computational statistics & data analysis*, 38(4), 367-378.
- Chen, T., & Guestrin, C. (2016, August). Xgboost: A scalable tree boosting system. In *Proceedings of the 22nd acm sigkdd international conference on knowledge discovery and data mining* (pp. 785-794).
- Ke, G., Meng, Q., Finley, T., Wang, T., Chen, W., Ma, W., ... & Liu, T. Y. (2017). Lightgbm: A highly efficient gradient boosting decision tree. *Advances in neural information processing systems*, 30.

