

# DATA MINING 2

## Gradient Boost

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Slides edited from StatQuest



# Gradient Boosting for Regression

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# Gradient Boost – Main Idea

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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

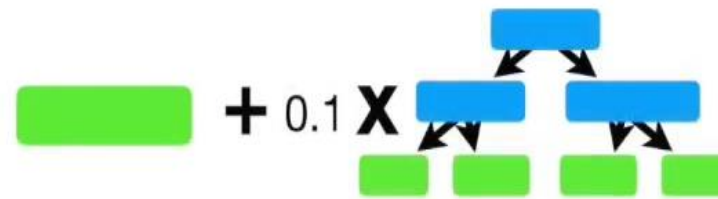
# Gradient Boost – Main Idea

Height (m)	Favorite Color	Gender	Weight (kg)
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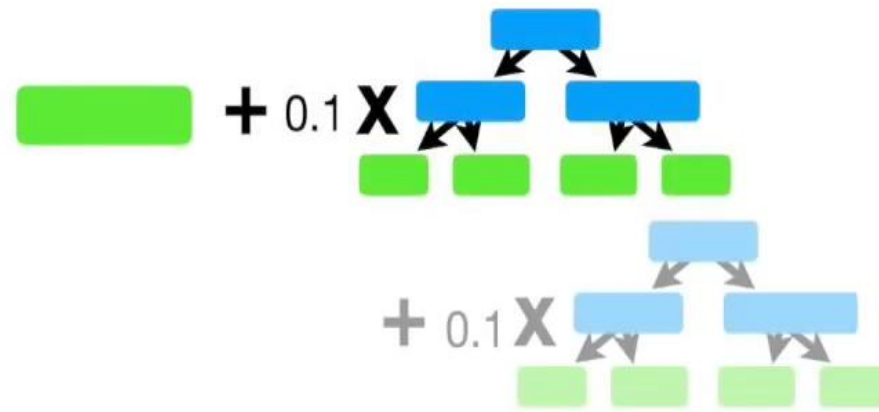
# Gradient Boost – Main Idea

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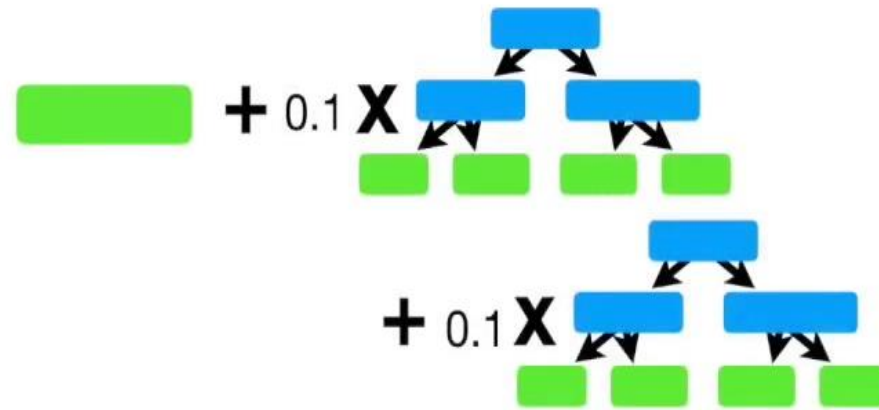
# Gradient Boost – Main Idea

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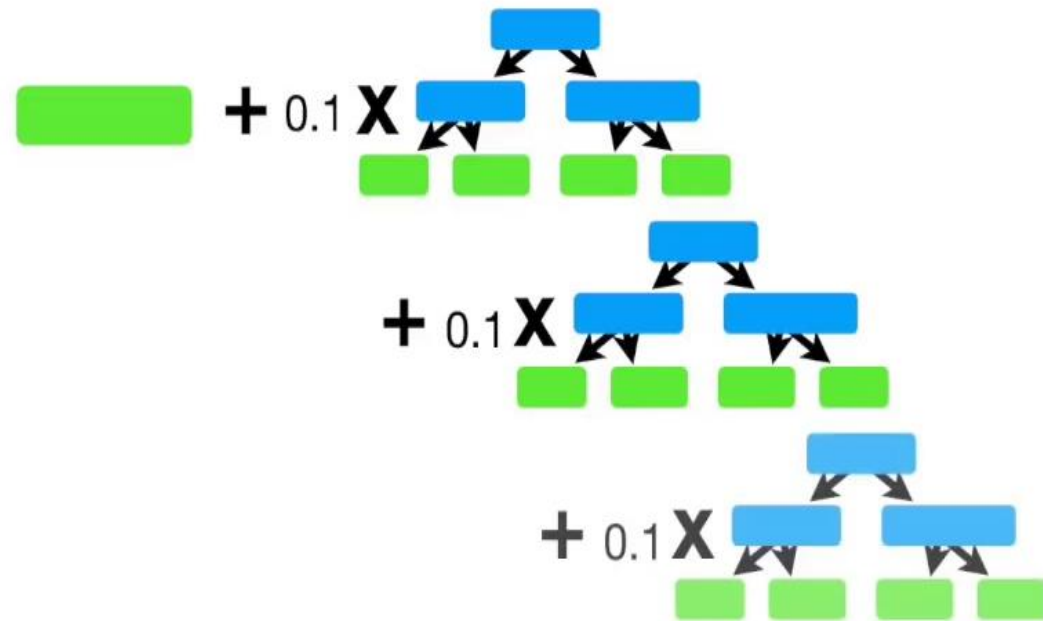
# Gradient Boost – Main Idea

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# Gradient Boost – Main Idea

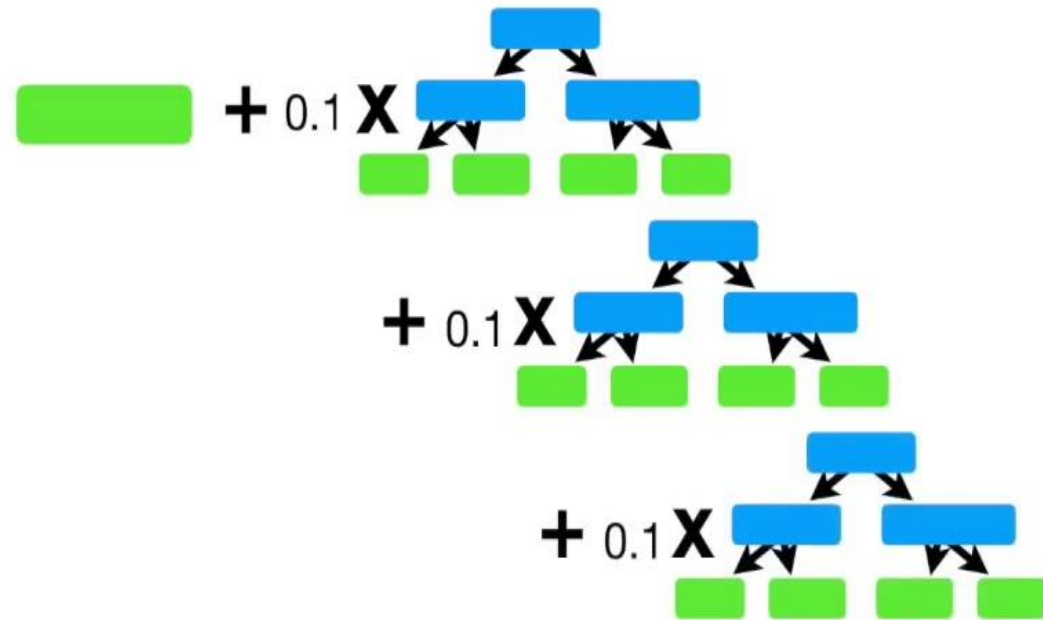
Height (m)	Favorite Color	Gender	Weight (kg)
1.6	Blue	Male	88
1.6	Green	Female	76
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# Gradient Boost – Main Idea

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1.8	Red	Male	73
1.5	Green	Male	77
1.4	Blue	Female	57



# Gradient Boost – Example

Average Weight

71.2

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

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

# Example

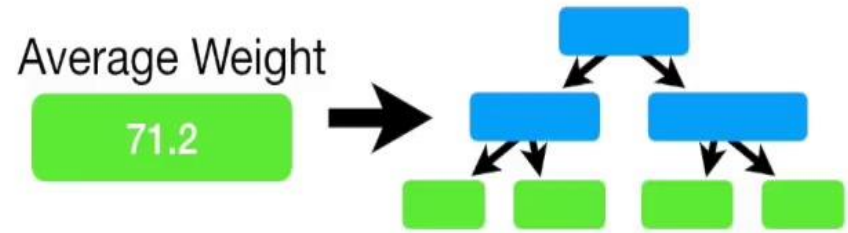
Average Weight

71.2

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

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

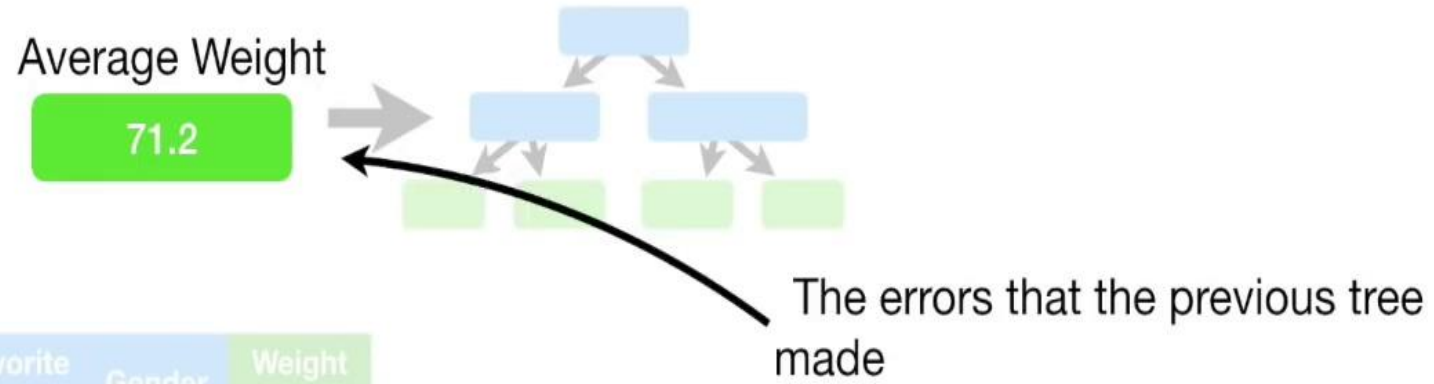
# Example



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

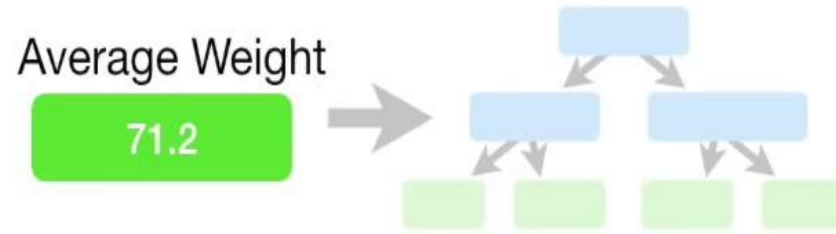
The next thing we do is build a tree based on the errors from the first tree.

# Example



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

# Example

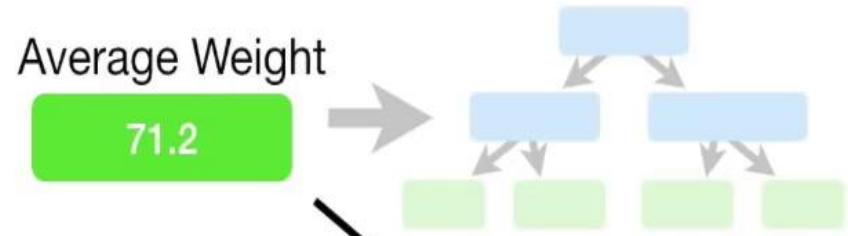


Height (m)	Favorite Color	Gender	Weight (kg)
1.6	Blue	Male	88
1.6	Green	Female	76
1.5	Blue	Female	56
1.8	Red	Male	73
1.5	Green	Male	77
1.4	Blue	Female	57

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

(Observed Weight - Predicted Weight)

# Example



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

The errors that the previous tree made are the differences between the **Observed Weights** and the **Predicted Weight, 71.2**.

(Observed Weight - Predicted Weight)

# Example

Average Weight

71.2

Height (m)	Favorite Color	Gender	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	

...and save the difference, which is called a **Pseudo Residual**, in a new column.

$$(88 - 71.2) = 16.8$$



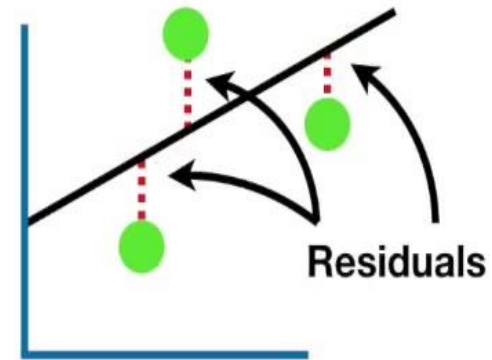
# Example

Average Weight

71.2

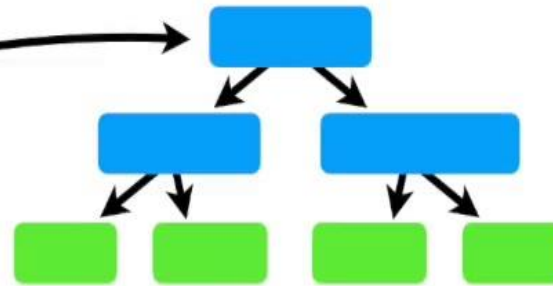
**NOTE:** The term **Pseudo Residual** is based on **Linear Regression**, where the difference between the **Observed** values and the **Predicted** values results in **Residuals**.

Height (m)	Favorite Color	Gender	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	



# Example

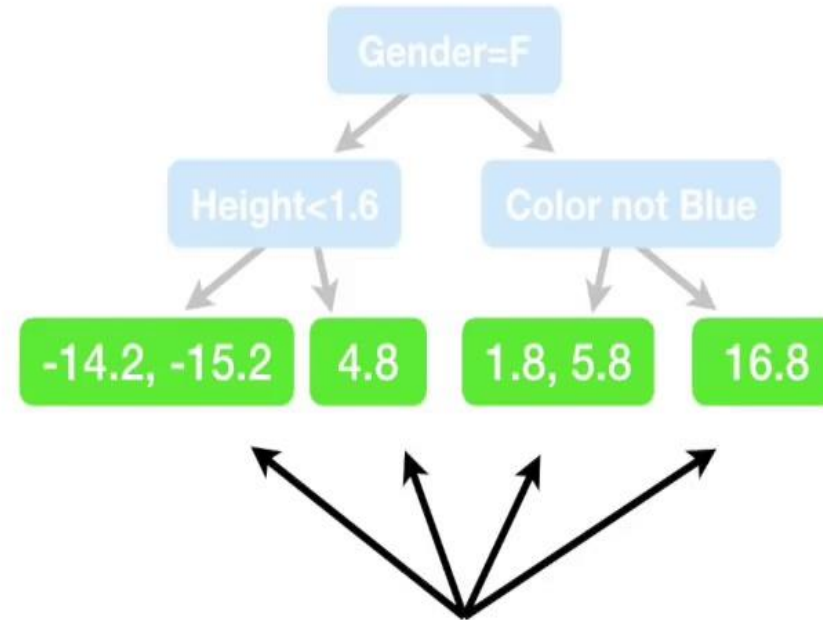
Now we will build a **Tree**



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

# Example

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

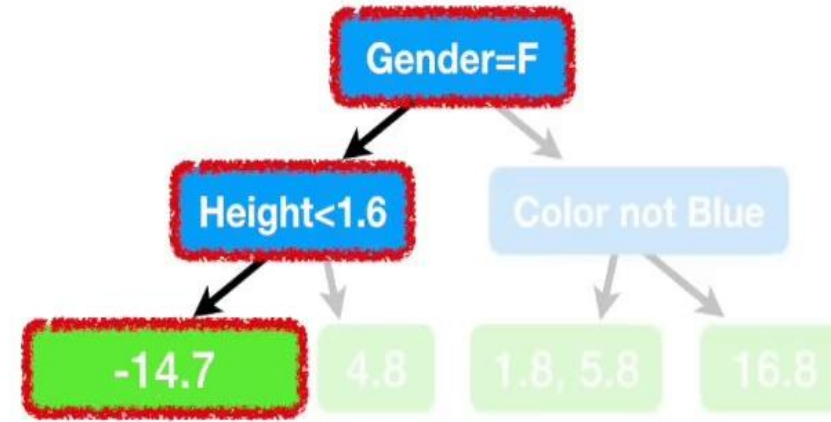


Remember, in this example we are only allowing up to four leaves...

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

# Example

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



So we replace these residuals with their average.

$$\frac{(-14.2 + -15.2)}{2} = -14.7$$

# Example

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



So we replace these residuals with their average.

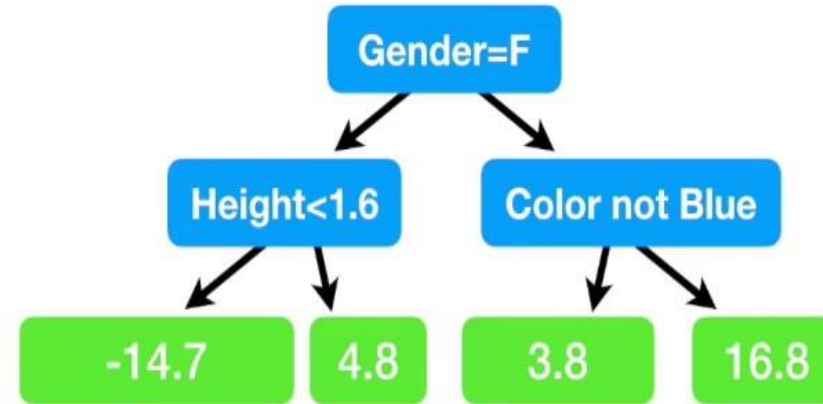
$$\frac{(1.8 + 5.8)}{2}$$

# Example

Average Weight

71.2

+



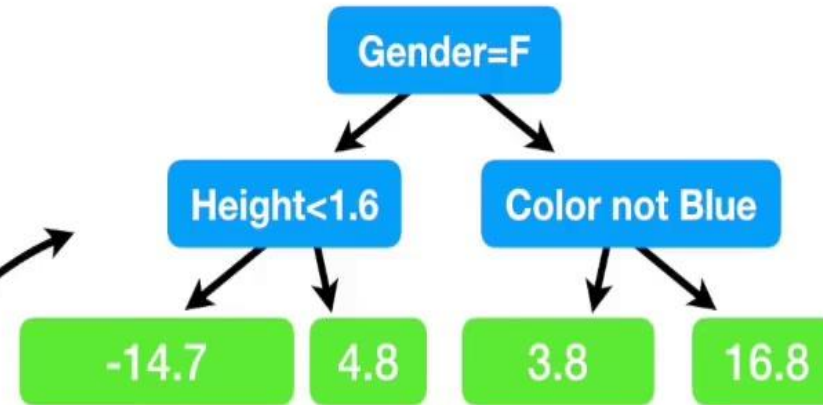
Now we can now combine  
the original leaf...

# Example

Average Weight

71.2

+



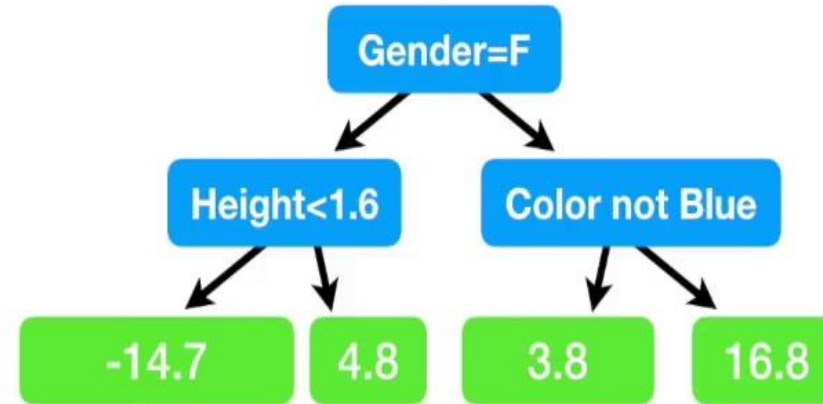
...with the new tree...

# Example

Average Weight

71.2

+



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

...to make a new **Prediction** of an individual's **Weight** from the **Training Data**.

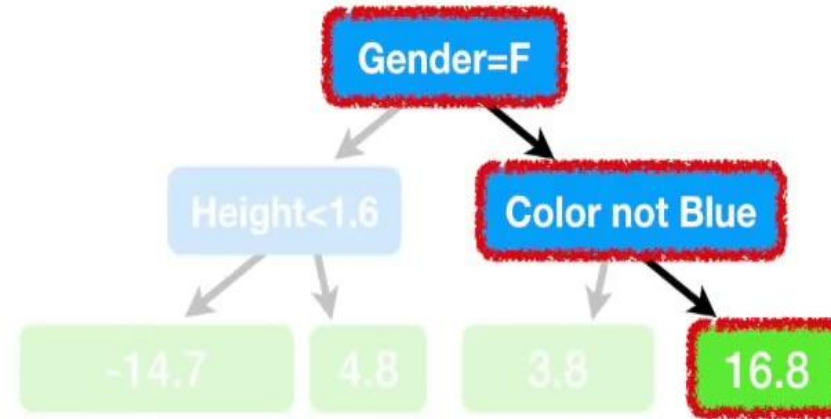


# Example

Average Weight

71.2

+



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

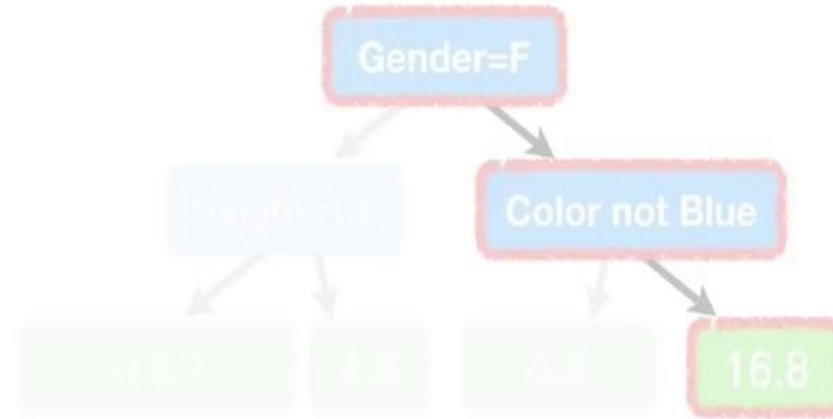
Height (m)	Favorite Color	Gender	Weight (kg)
1.6	Blue	Male	88

# Example

Average Weight

71.2

+

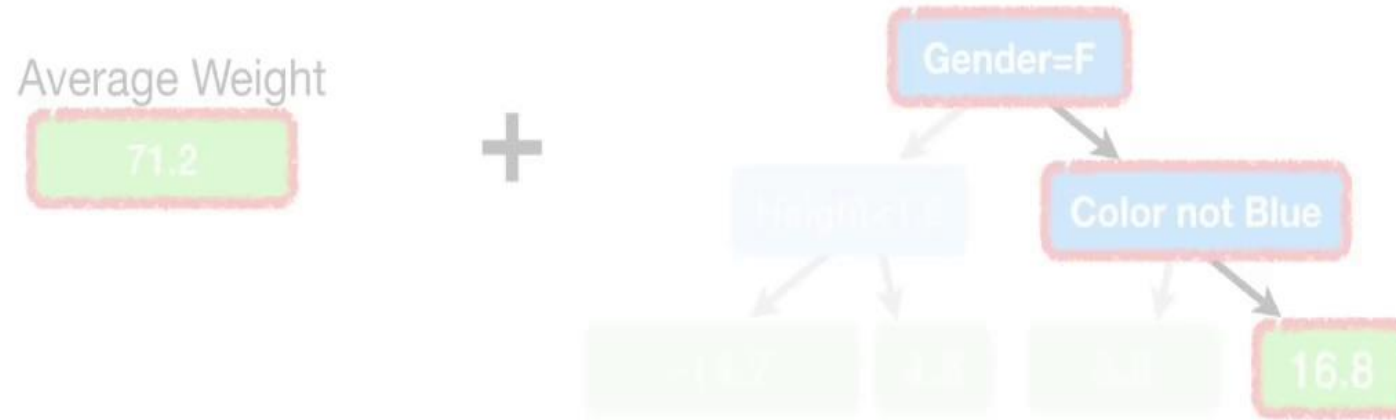


$$\text{Predicted Weight} = 71.2 + 16.8 = 88$$

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

Is this awesome???

# Example



$$\text{Predicted Weight} = 71.2 + 16.8 = 88$$

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

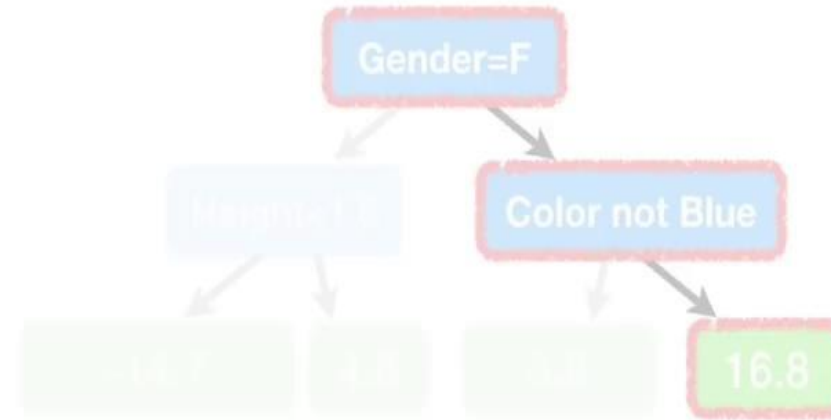
**No.** The model fits the **Training Data** too well.

# Example

Average Weight

71.2

+



$$\text{Predicted Weight} = 71.2 + 16.8 = 88$$

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

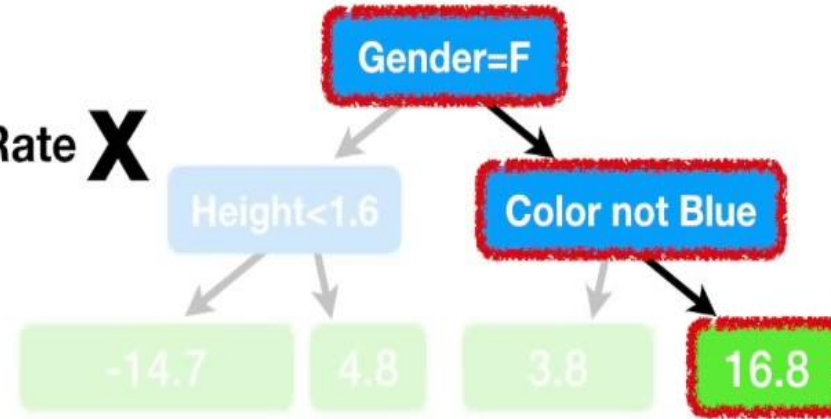
In other words, we have low **Bias**, but probably very high **Variance**.

# Example

Average Weight

71.2

+ Learning Rate  $\times$



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

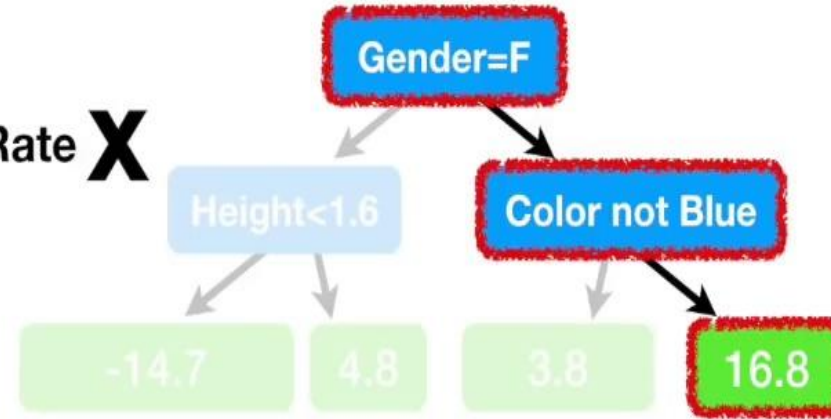
Height (m)	Favorite Color	Gender	Weight (kg)
1.6	Blue	Male	88

# Example

Average Weight

71.2

+ Learning Rate  $\times$

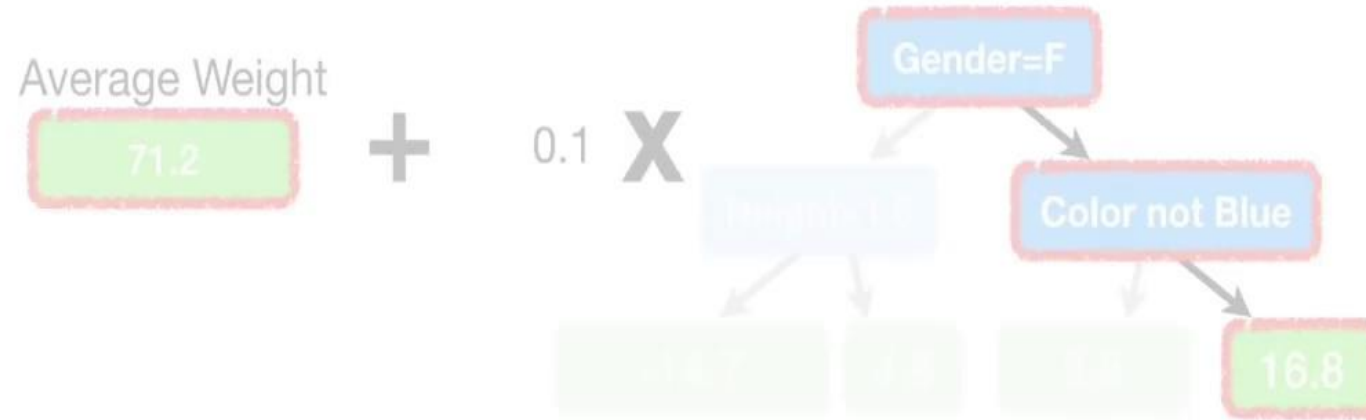


Height (m)	Favorite Color	Gender	Weight (kg)
1.6	Blue	Male	88

**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**.

# Example

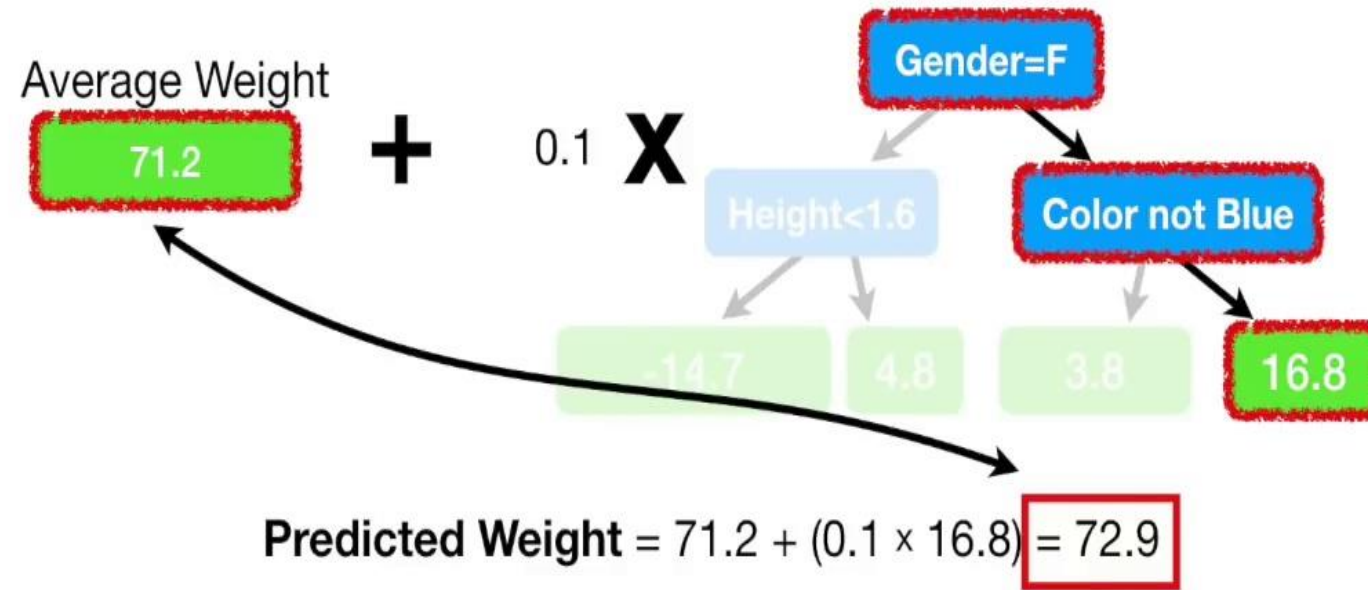


$$\text{Predicted Weight} = 71.2 + (0.1 \times 16.8) = 72.9$$

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

With the **Learning Rate** set to **0.1**, the new **Prediction** isn't as good as it was before...

# Example

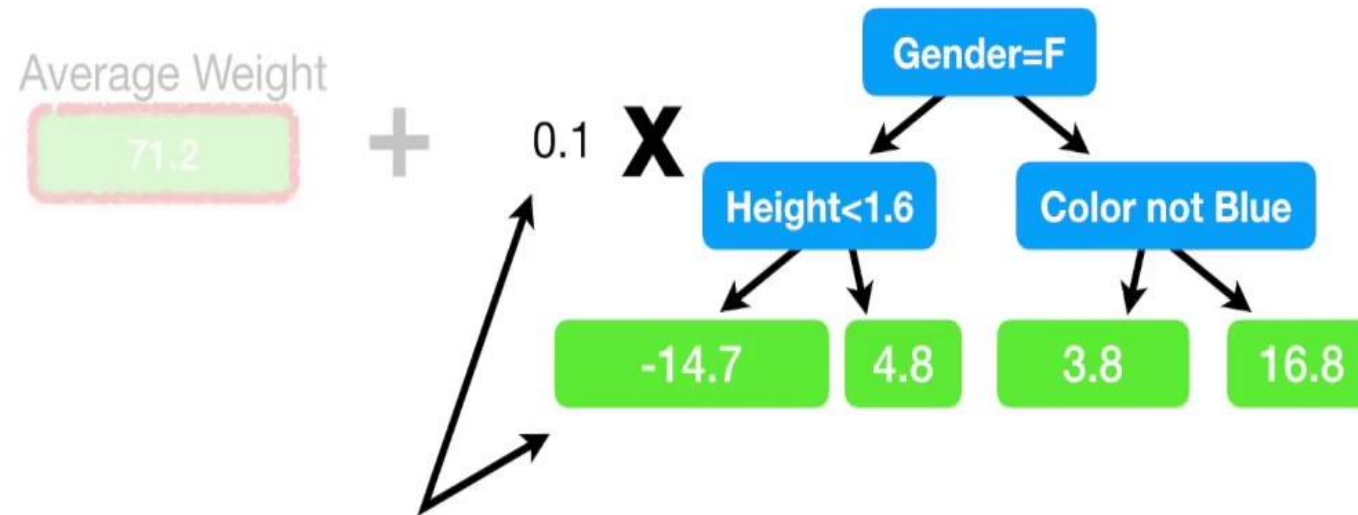


Height (m)	Favorite Color	Gender	Weight (kg)
1.6	Blue	Male	88

...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**.

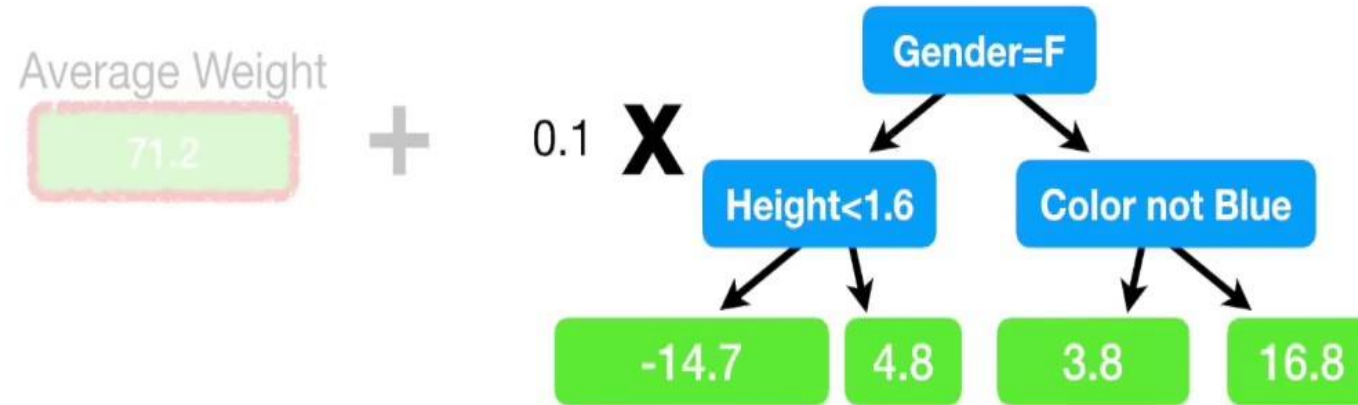


# Example



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

# Example



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

# Example

Height (m)	Favorite Color	Gender	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)**

# Example

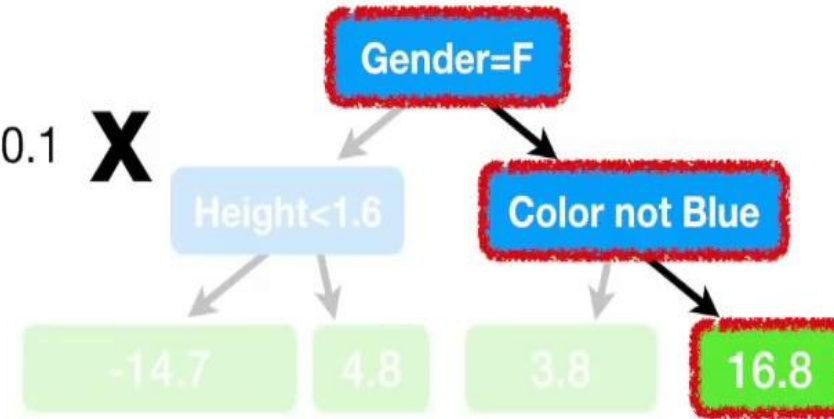
Average Weight

71.2

+

0.1

**X**



Height (m)	Favorite Color	Gender	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	

$$\text{Residual} = (88 - (71.2 + 0.1 \times 16.8))$$

= 15.1

...and we save that in the column for **Pseudo Residuals**.

# Example

Average Weight

71.2

+

0.1 X

Height < 1.6

Gender = F

Color not Blue

-14.7

4.8

3.8

16.8

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

71.2

+

0.1

X

Gender=F

Height<1.6

Color not Blue

-14.7

4.8

3.8

16.8

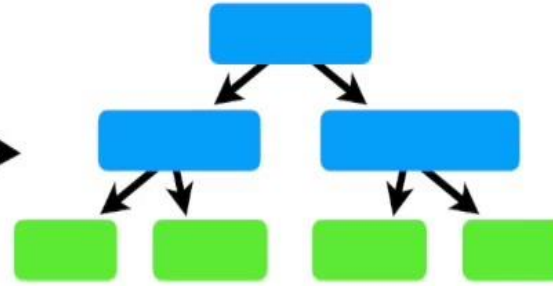
Residual
16.8
4.8
-15.2
1.8
5.8
-14.2

Residual
15.1
4.3
-13.7
1.4
5.4
-12.7

The new **Residuals** are all smaller than before, so we've taken a small step in the right direction.

# Example

Now let's build a new tree to predict the new **Residuals**.



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

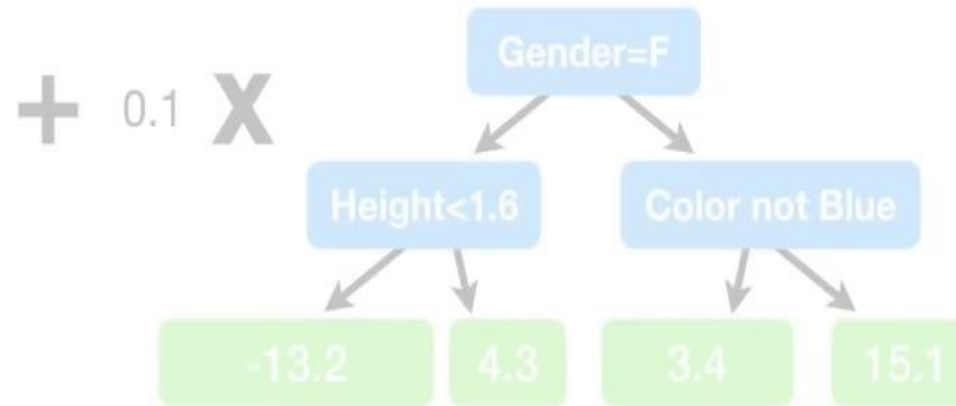
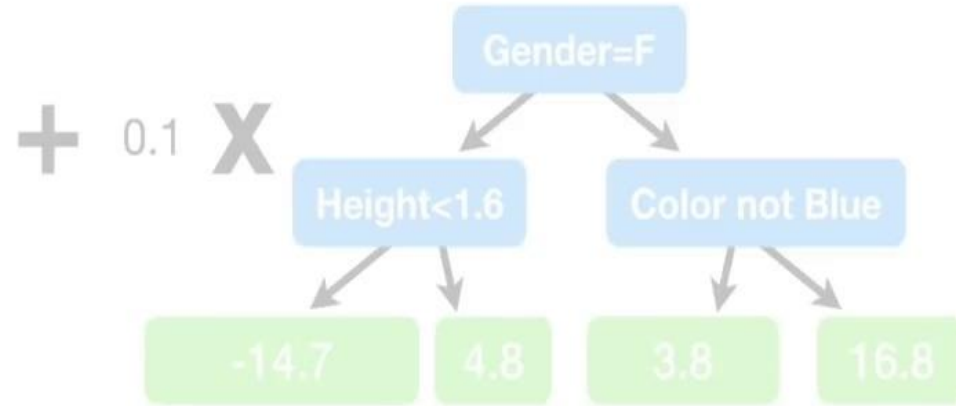
# Example

Average Weight

71.2

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

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



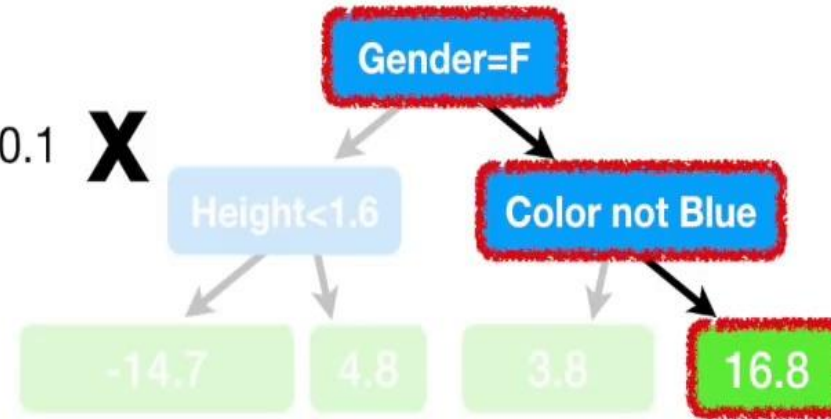


# Example

Average Weight

71.2

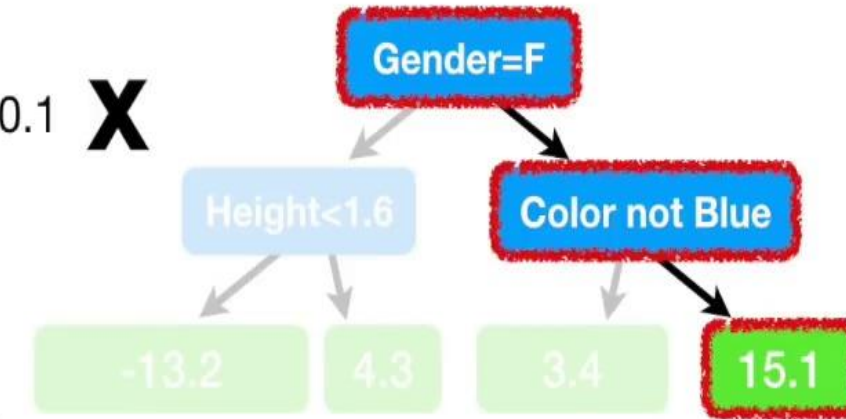
+ 0.1 X



...and the scaled amount from the second **Tree**.

+ 0.1 X

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



# Example

Average Weight

71.2

+ 0.1 X



That gives us...

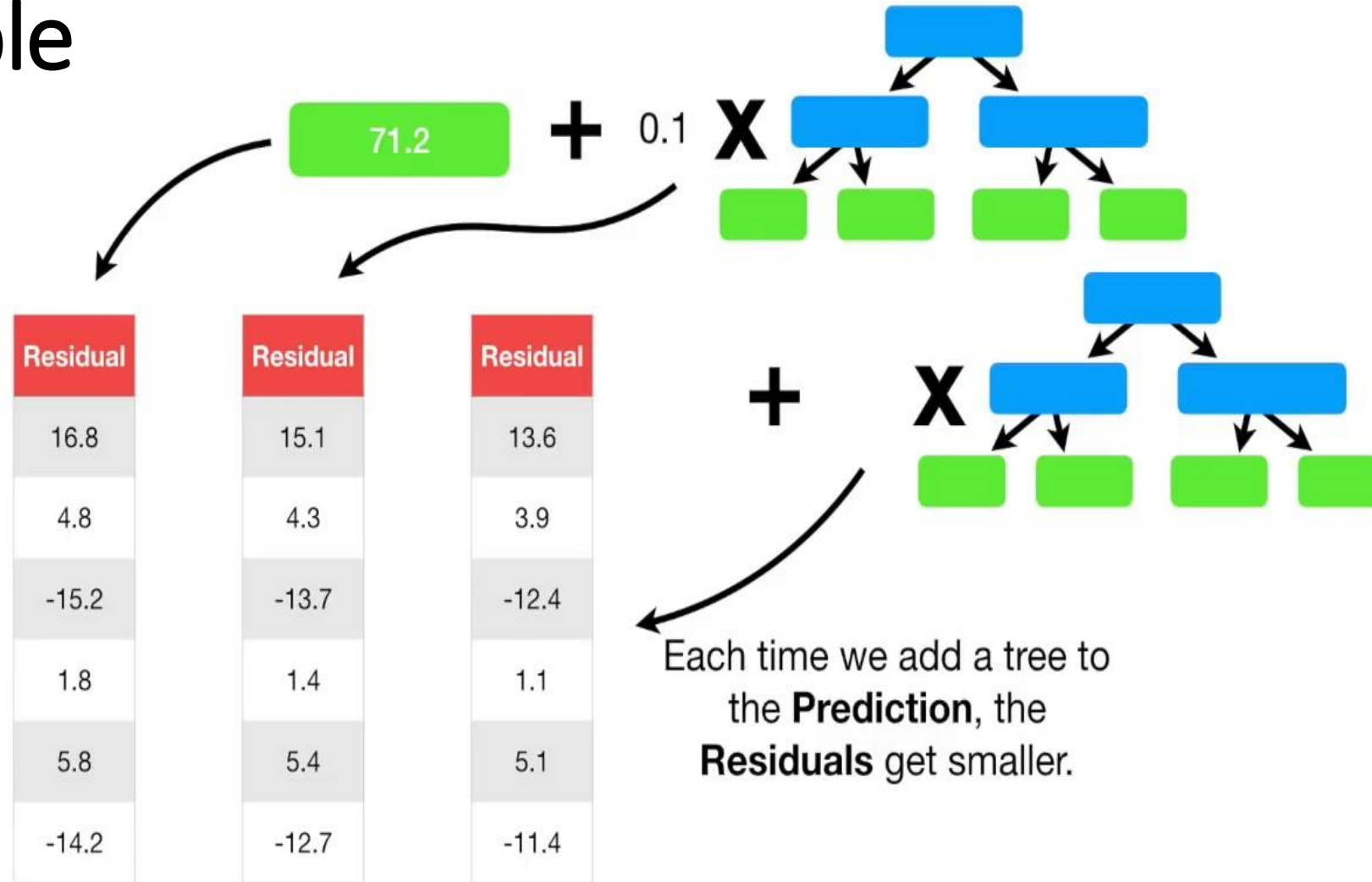
$$71.2 + (0.1 \times 16.8) + (0.1 \times 15.1)$$

= 74.4

+ 0.1 X



# Example



# GB Algorithm

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

**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  $i = 1, \dots, n$

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

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

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

**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...

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

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

# GB Algorithm

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**”...



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

# GB Algorithm

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

...then the **2/2** cancels out...

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

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

# GB Algorithm

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**.

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

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

$$\boxed{= -(\text{Observed} - \text{Predicted})}$$



# GB Algorithm

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.

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

$$= -(\text{Observed} - \text{Predicted})$$

# GB Algorithm

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

The  $y_i$ 's are the **Observed** values...

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

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

# GB Algorithm

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

...and  $F(x)$  is a function that gives us the **Predicted** values.

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

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

# GB Algorithm

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

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

# GB Algorithm

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

$$\frac{1}{2} (88 - \text{Predicted})^2 +$$
$$\frac{1}{2} (76 - \text{Predicted})^2 +$$
$$\frac{1}{2} (56 - \text{Predicted})^2$$

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

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

# GB Algorithm

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

$$\frac{1}{2} (88 - \text{Predicted})^2 + \\ \frac{1}{2} (76 - \text{Predicted})^2 + \\ \frac{1}{2} (56 - \text{Predicted})^2$$

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

...and the “**argmin over gamma**” means we need to find a **Predicted** value that minimizes this sum.

# GB Algorithm

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

$$\frac{1}{2} (88 - \text{Predicted})^2 +$$

$$\frac{1}{2} (76 - \text{Predicted})^2 +$$

$$\frac{1}{2} (56 - \text{Predicted})^2$$



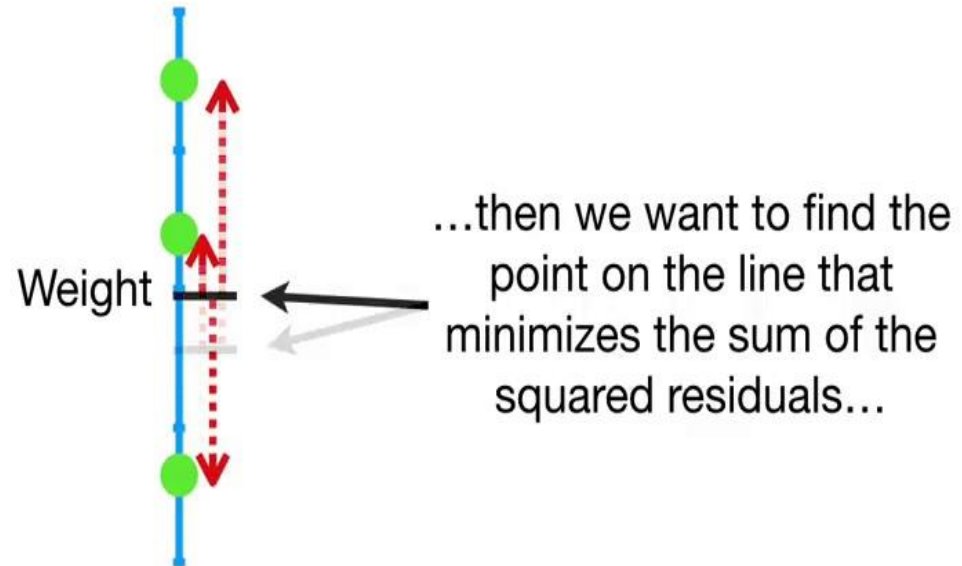
In other words, if we plot the **Observed Weights** on a number line...

# GB Algorithm

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

$$\begin{aligned} & \frac{1}{2} (88 - \text{Predicted})^2 + \\ & \frac{1}{2} (76 - \text{Predicted})^2 + \\ & \frac{1}{2} (56 - \text{Predicted})^2 \end{aligned}$$





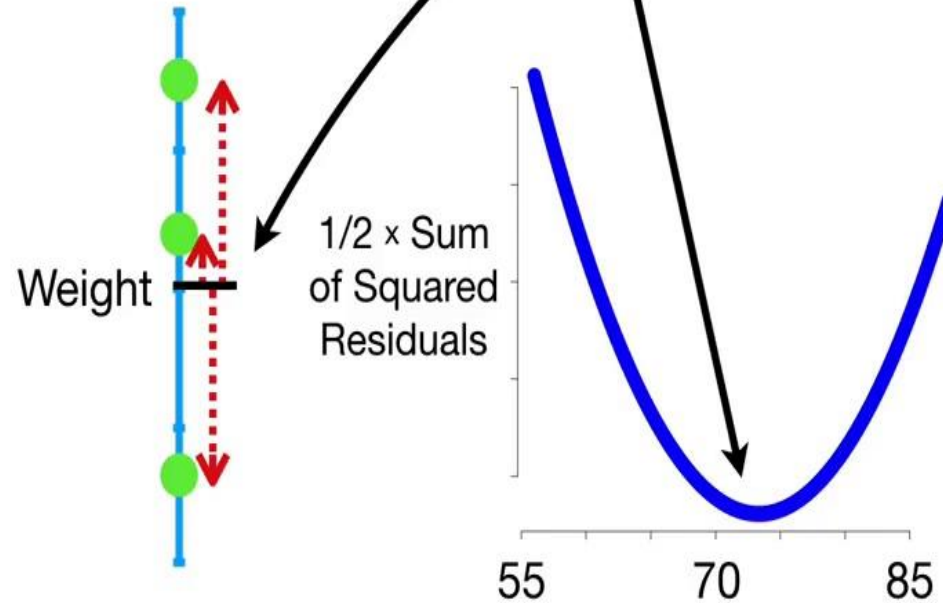
# GB Algorithm

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

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

$$\frac{1}{2} (88 - \text{Predicted})^2 +$$
$$\frac{1}{2} (76 - \text{Predicted})^2 +$$
$$\frac{1}{2} (56 - \text{Predicted})^2$$

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



# GB Algorithm

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

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

$$\text{Predicted} = \frac{88 + 76 + 56}{3}$$

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

# GB Algorithm

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

We have now created  
the initial predicted  
value,  $F_0(x)$ ...

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

# GB Algorithm

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

...and it equals **73.3**.

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

$$= 73.3$$

# GB Algorithm

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

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

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



73.3

# GB Algorithm

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  $i = 1, \dots, n$

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

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

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

# GB Algorithm

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  $i = 1, \dots, n$

This part is just the  
derivative of the  
**Loss Function...**

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

# GB Algorithm

**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  $i = 1, \dots, n$

This part is just the  
derivative of the  
**Loss Function...**

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

...with respect to the  
**Predicted** value...



# GB Algorithm

**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  $i = 1, \dots, n$

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

...and we've already  
calculated this.


$$= -(\text{Observed} - \text{Predicted})$$

# GB Algorithm

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  $i = 1, \dots, n$

Height (m)	Favorite Color	Gender	Weight (kg)	$r_{i,1}$
1.6	Blue	Male	88	14.7
1.6	Green	Female	76	2.7
1.5	Blue	Female	56	-17.3

We've finished **Part A** of **Step 2** by calculating a **Residual** for each sample.

# GB Algorithm

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

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

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

# GB Algorithm

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  $i = 1, \dots, n$

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


# GB Algorithm

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  $i = 1, \dots, n$

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

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



# GB Algorithm

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

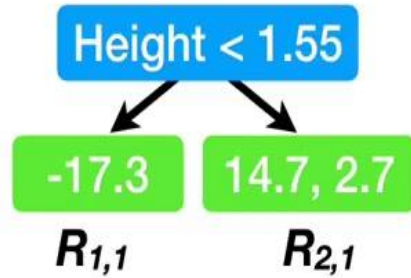
Step 2: for  $m = 1$  to  $M$ :

(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  $i = 1, \dots, n$

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

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

# GB Algorithm



In this part, we determine the **Output Values** for each leaf.

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

# GB Algorithm

**Step 1:** Initialize model with a constant value:

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

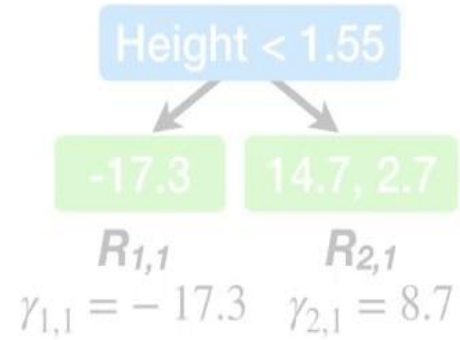
**NOTE:** This minimization is like what we did in **Step 1**.

(C) For  $j = 1 \dots J_m$  compute

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



# GB Algorithm



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

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

# GB Algorithm

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

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

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

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

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

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

# GB Algorithm

$$F_1(x) = \overbrace{F_0(x)}^{73.3} + 0.1 \times \begin{cases} -17.3 & R_{1,1} \\ 14.7, 2.7 & R_{2,1} \end{cases}$$

$\gamma_{1,1} = -17.3 \quad \gamma_{2,1} = 8.7$

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

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

# GB Algorithm

$$F_1(x) = F_0(x) + 0.1 \times \begin{cases} -17.3 & \text{Height} < 1.55 \\ 14.7, 2.7 & \text{Height} \geq 1.55 \end{cases}$$

$R_{1,1}$        $R_{2,1}$   
 $\gamma_{1,1} = -17.3$        $\gamma_{2,1} = 8.7$

Height (m)	Favorite Color	Gender	Weight (kg)
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  $F_m(x) = F_{m-1}(x) + \nu \sum_{j=1}^{J_m} \gamma_{jm} I(x \in R_{jm})$

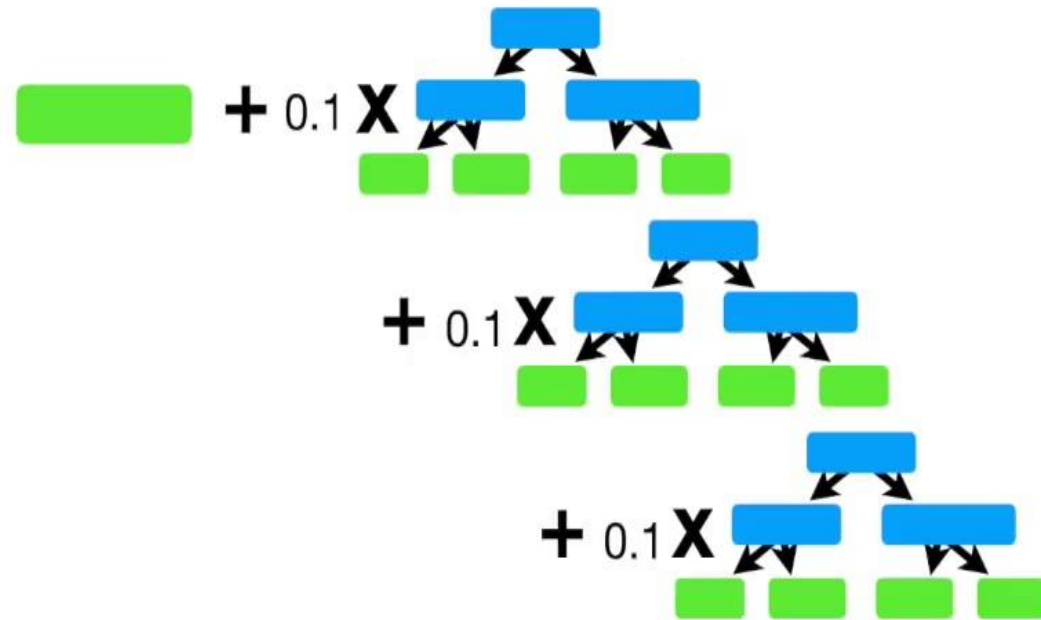
# Gradient Boosting for Classification

---

# Example

...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



# Example



When we use **Gradient Boost for Classification**, the initial **Prediction** for every individual is the  **$\log(\text{odds})$** .

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

# Example

$$\log(4/2) = 0.7$$

← So this is the **Initial Prediction**.

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



# Example

$$\log(4/2) = 0.693$$

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

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

# Example

$$\log(4/2) = 0.693$$

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

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

# Example

$$\log(4/2) = 0.693$$

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

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

**Probability of Loving Troll 2** =  $\frac{e^{\log(\text{odds})}}{1 + e^{\log(\text{odds})}}$

# Example

$$\log(4/2) = 0.693$$

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

Probability of Loving Troll 2 =  $\frac{e^{\log(4/2)}}{1 + e^{\log(4/2)}} = \boxed{0.7}$  0.666

# Example

$$\log(4/2) = 0.693$$

NOTE this are rounded values

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

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

$$\text{Probability of Loving Troll 2} = \frac{e^{\log(4/2)}}{1 + e^{\log(4/2)}} = \boxed{0.7} \quad 0.666$$

# Example

$$\log(4/2) = 0.7$$

Probability of  
Loving Troll 2 = 0.7

0.666

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**.

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

# Example

$$\log(4/2) = 0.693$$

Probability of  
**Loving Troll 2** = 0.7 <sup>0.666</sup>

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

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

$$\text{Residual} = (\text{Observed} - \text{Predicted})$$

# Example

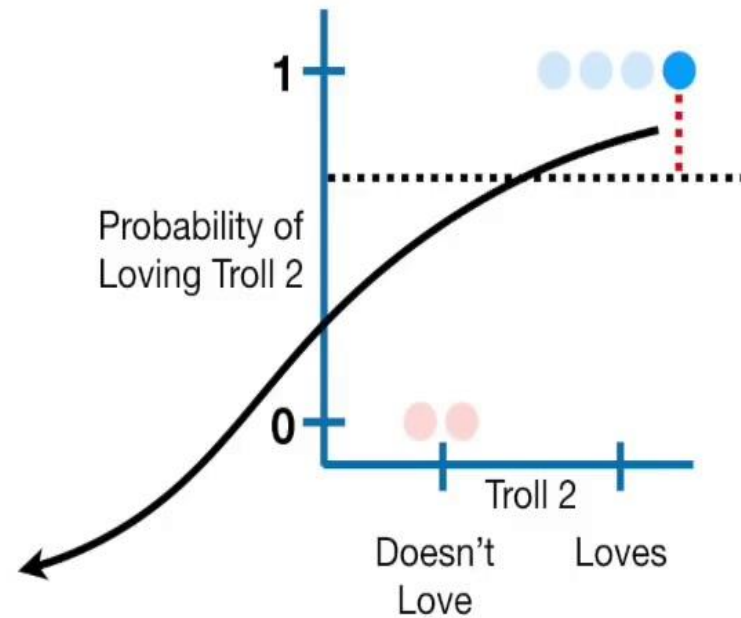
$$\log(4/2) = 0.7$$

Probability of  
Loving Troll 2 = 0.7

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

$$\text{Residual} = (\text{Observed} - \text{Predicted})$$

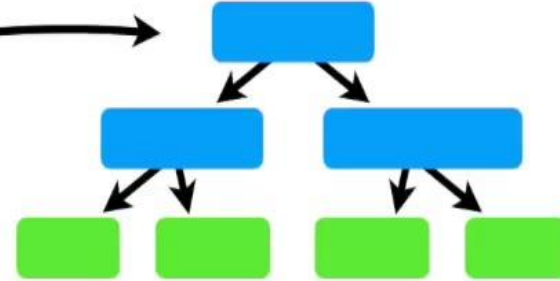
Likes Popcorn	Age	Favorite Color	Loves Troll 2	Residual
Yes	12	Blue	Yes	0.3
Yes	87	Green	Yes	0.3
No	44	Blue	No	-0.7
Yes	19	Red	No	-0.7
No	32	Green	Yes	0.3
No	14	Blue	Yes	0.3





# Example

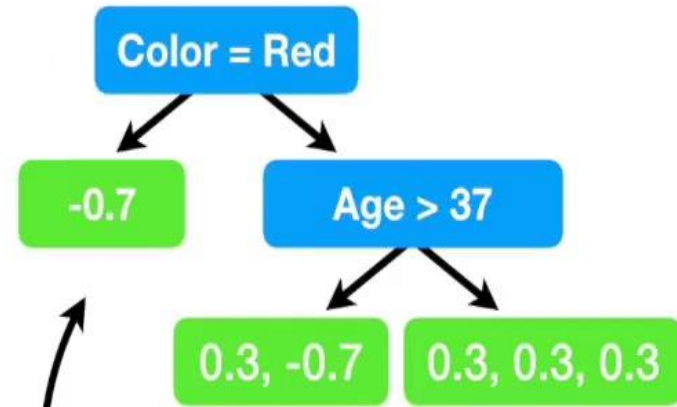
Now we will build a **Tree**



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

# Example

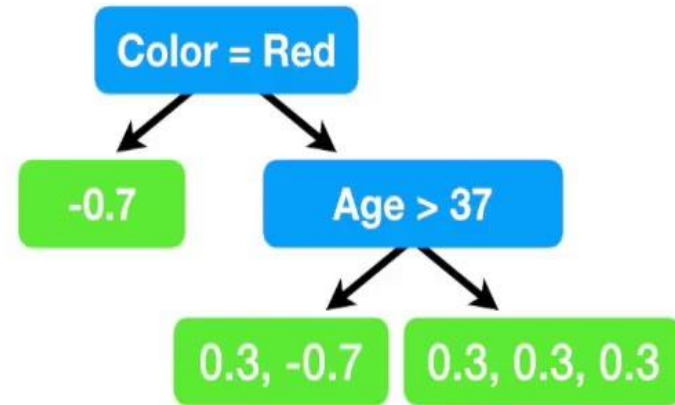
Likes Popcorn	Age	Favorite Color	Loves Troll 2	Residual
Yes	12	Blue	Yes	0.3
Yes	87	Green	Yes	0.3
No	44	Blue	No	-0.7
Yes	19	Red	No	-0.7
No	32	Green	Yes	0.3
No	14	Blue	Yes	0.3



And here's the tree!

# Example

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**

# Example



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

# Example



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

# Example

$$\log(4/2) = 0.7$$



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



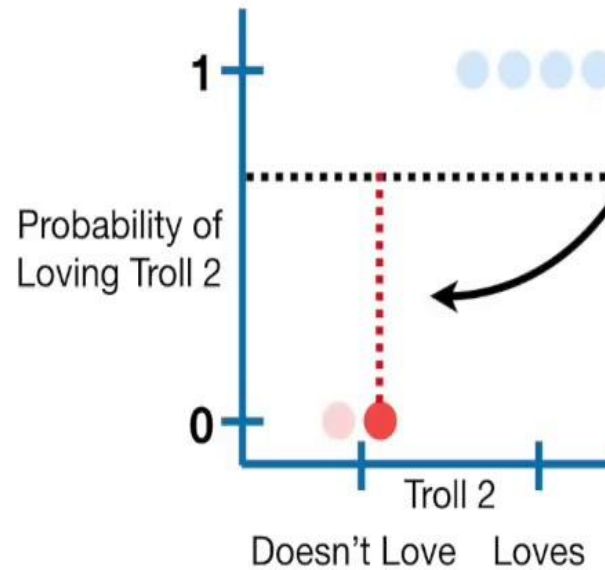
# Example

$$\log(4/2) = 0.7$$

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



...and this leaf is derived from a **Probability**...



# Example

$$\log(4/2) = 0.7$$



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





# Example

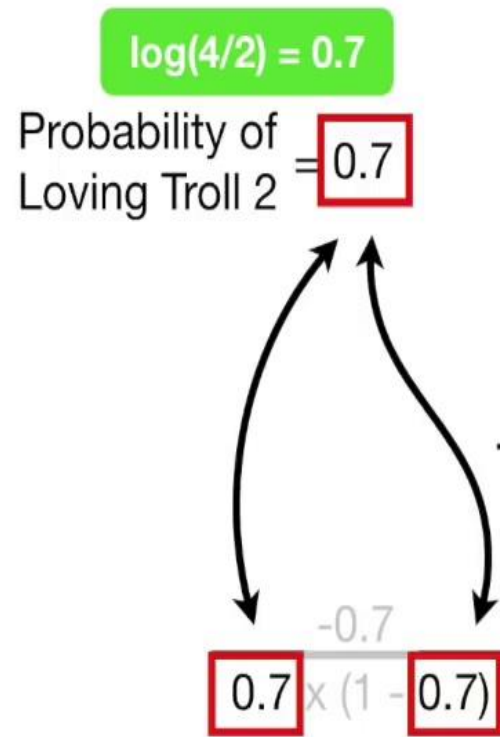
When we use **Gradient Boost** for **Classification**, the most common transformation is the following formula.



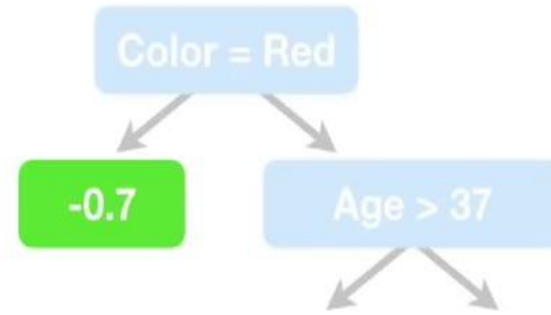
$$\frac{\sum \text{Residual}_i}{\sum [\text{Previous Probability}_i \times (1 - \text{Previous Probability}_i)]}$$



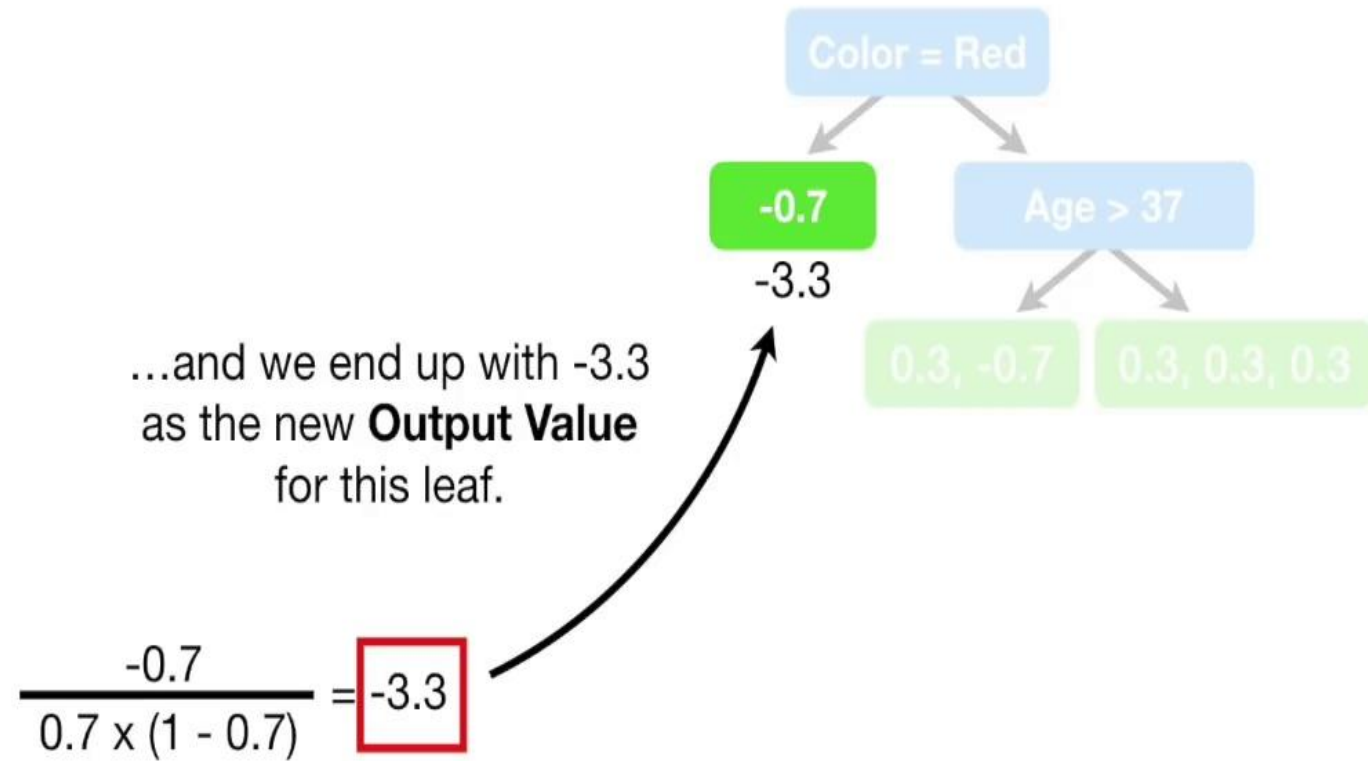
# Example



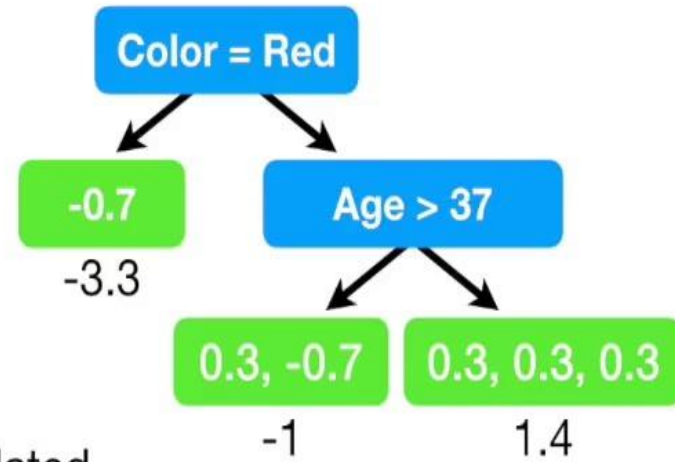
...so we plug that in...



# Example

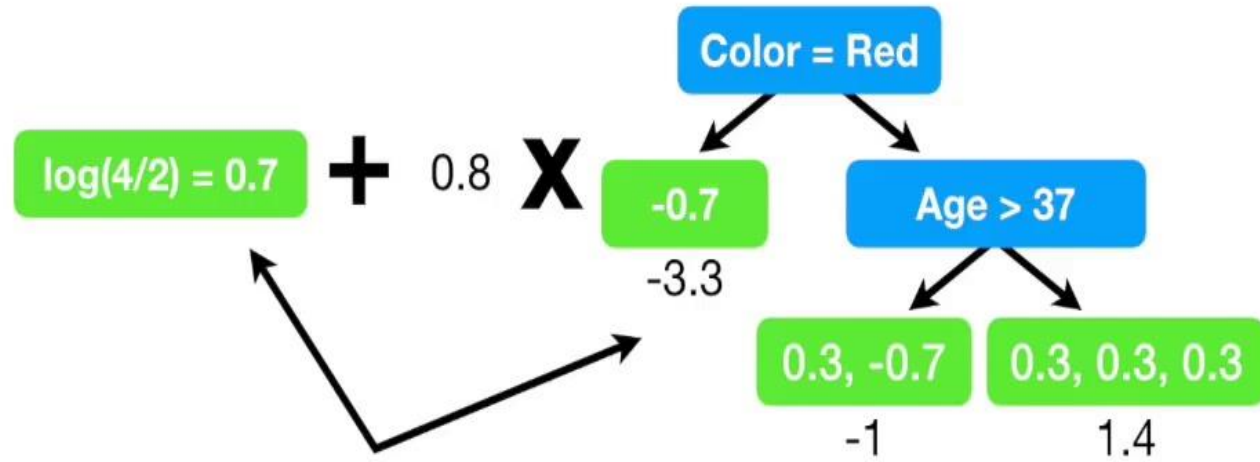


# Example



We've calculated **Output Values** for all three leaves in the tree!

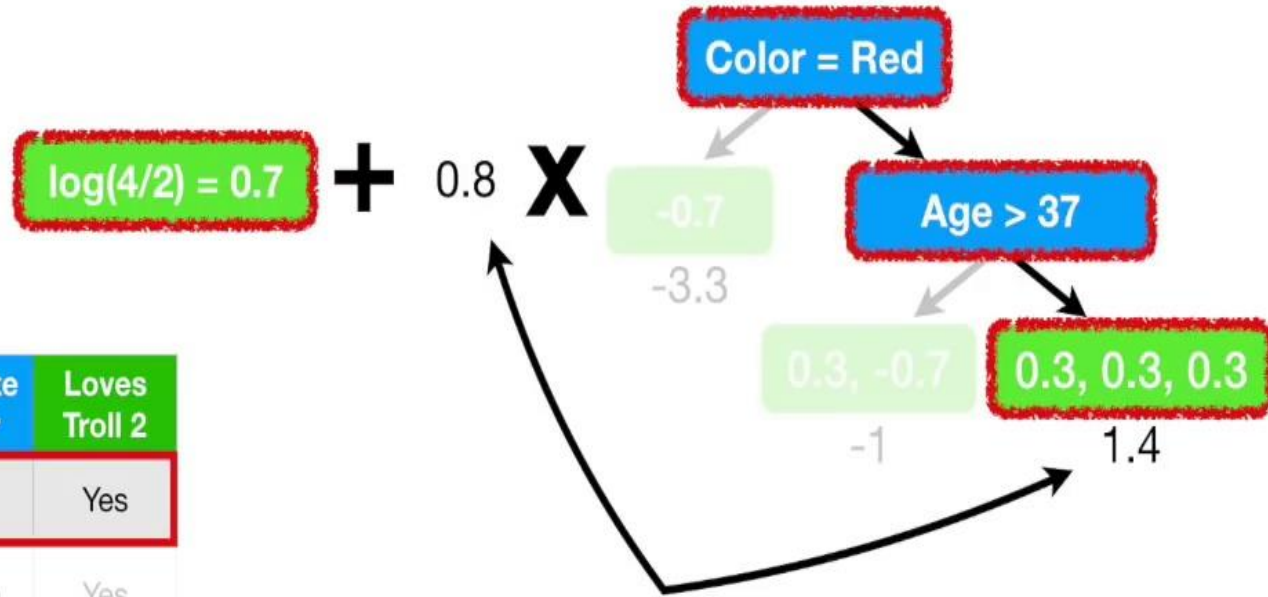
# Example



Now we are ready to update our **Predictions** by combining the initial leaf with the new tree.

# Example

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

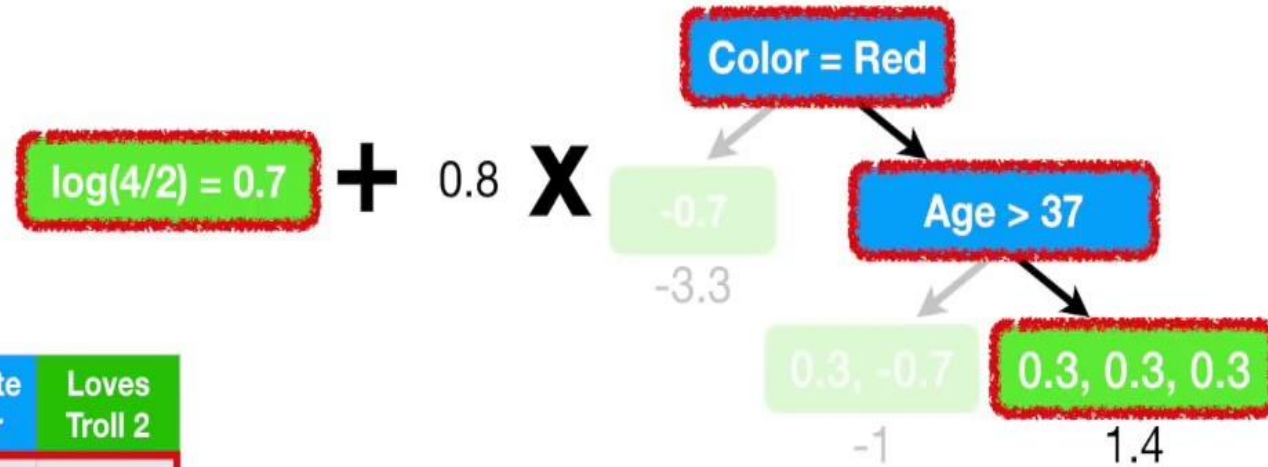


...plus the **Output Value** from the tree scaled by the **Learning Rate**...

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

# Example

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.**

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

# Example

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...**

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


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



# Example

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**...

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


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

# Example

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**...

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

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

# Example

$$\log(4/2) = 0.7$$

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**.

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

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

# Example

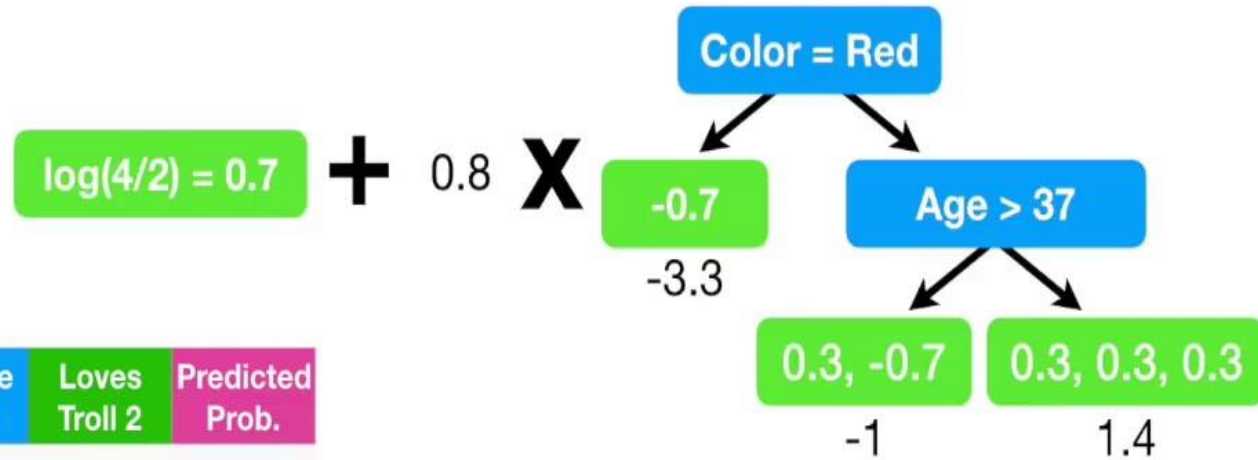
Likes Popcorn	Age	Favorite Color	Loves Troll 2	Predicted Prob.
Yes	12	Blue	Yes	0.9
Yes	87	Green	Yes	
No	44	Blue	No	
Yes	19	Red	No	
No	32	Green	Yes	
No	14	Blue	Yes	

We save the new **Predicted Probability** here.

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

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

# Example



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.

# Example

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

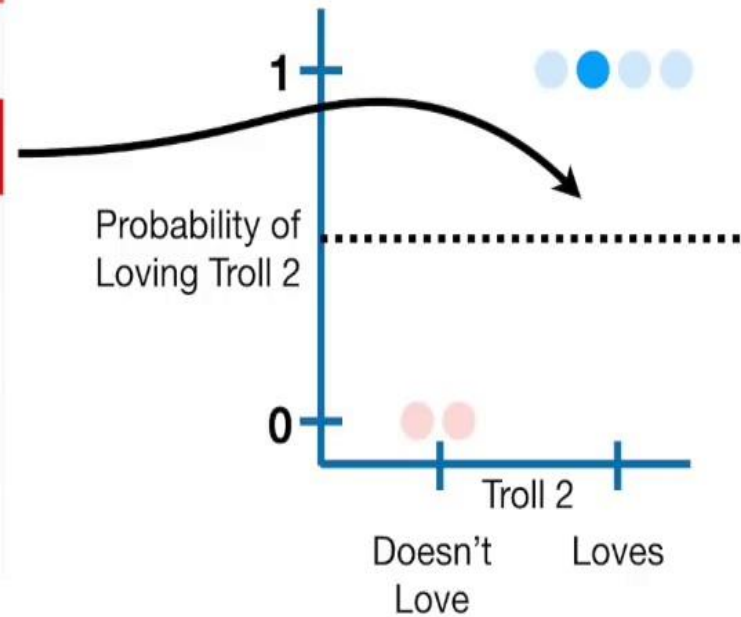


Likes Popcorn	Age	Favorite Color	Loves Troll 2	Predicted Prob.	Residual
Yes	12	Blue	Yes	0.9	
Yes	87	Green	Yes	0.5	
No	44	Blue	No	0.5	
Yes	19	Red	No	0.1	
No	32	Green	Yes	0.9	
No	14	Blue	Yes	0.9	

# Example

Likes Popcorn	Age	Favorite Color	Loves Troll 2	Predicted Prob.	Residual
Yes	12	Blue	Yes	0.9	0.1
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	

We plot the **Predicted Probability...**

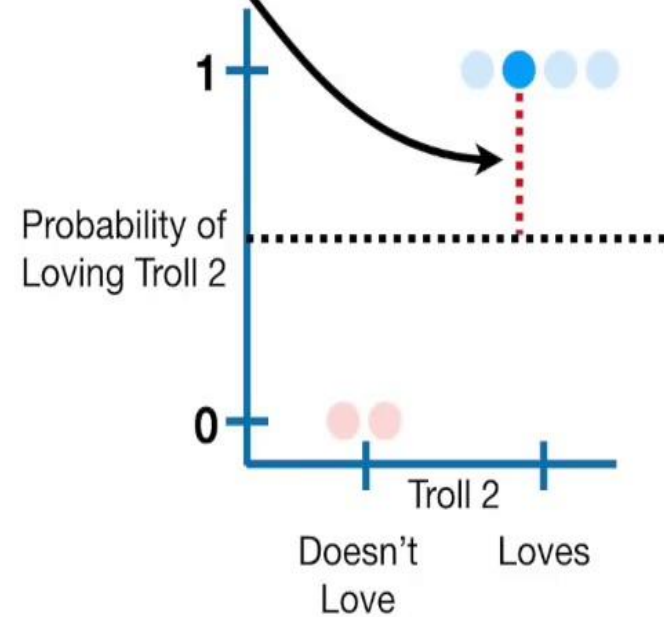


# Example

...and the **Residual** is the difference.

$$\text{Residual} = (\text{Observed} - \text{Predicted})$$

Likes Popcorn	Age	Favorite Color	Loves Troll 2	Predicted Prob.	Residual
Yes	12	Blue	Yes	0.9	0.1
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	

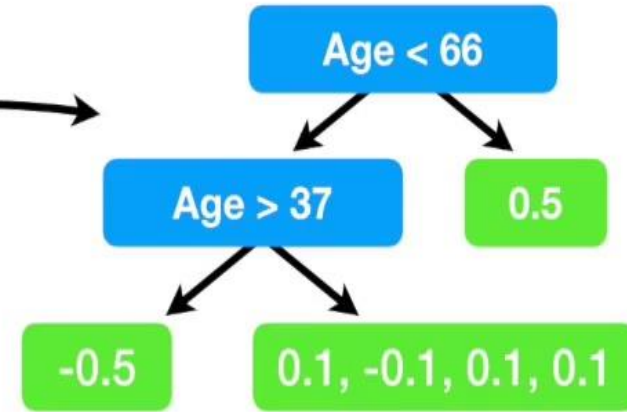




# Example

Now that we have the **Residuals**, we can build a new tree...

Likes Popcorn	Age	Favorite Color	Loves Troll 2	Predicted Prob.	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



# GB Algorithm

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

**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  $i = 1, \dots, n$

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

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

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

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

# GB Algorithm

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



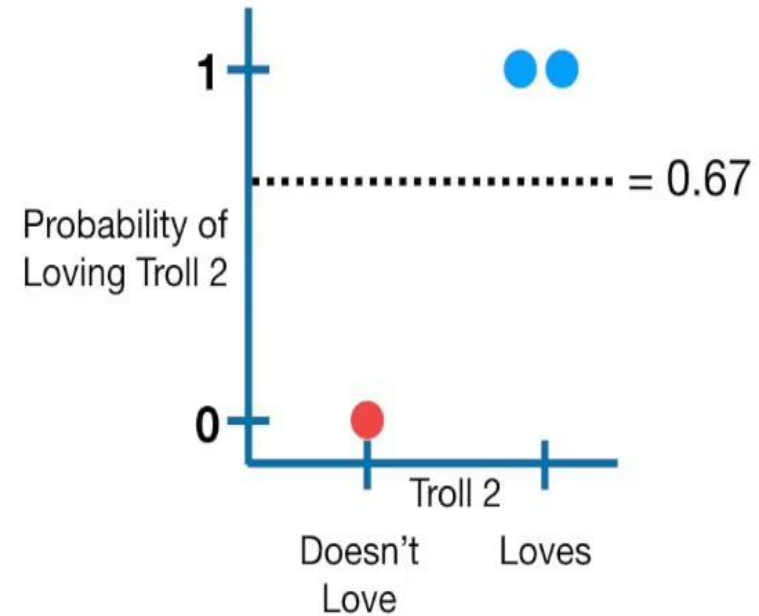
Now we need a differentiable  
**Loss Function** that will work  
for **Classification**.

# GB Algorithm

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

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

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



$$-\left[ \text{Observed} \times \log(p) + (1 - \text{Observed}) \times \log(1 - p) \right]$$

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

- 1)  $-\text{Observed} \times \log(p) - (1 - \text{Observed}) \times \log(1 - p)$
- 2)  $-\text{Observed} \times \log(p) - \log(1 - p) + \text{Observed} \times \log(1 - p)$
- 3)  $-\text{Observed} \times [\log(p) - \log(1 - p)] - \log(1 - p)$
- 4)  $-\text{Observed} \times \log(\text{odds}) - \log(1 - p)$

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


$$5) -\text{Observed} \times \log(\text{odds}) + \log(1 + e^{\log(\text{odds})})$$

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

This is the Loss function

# GB Algorithm

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



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

# GB Algorithm

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

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

$$= -\text{Observed} + p$$

Indeed, in the previous example we performed

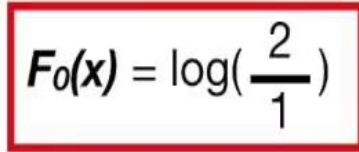
- $1 - 0.7 = 0.3$
- $0 - 0.7 = -0.7$

# GB Algorithm

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

We have  
created the initial  
predicted  
**log(odds),  $F_0(x)$ ...**


$$F_0(x) = \log\left(\frac{2}{1}\right)$$



# GB Algorithm

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

So this is the initial  
leaf,  $F_0(x)$ .

$$F_0(x) = \log\left(\frac{2}{1}\right) \\ = 0.69$$

$$\log(2/1) = 0.69$$

# GB Algorithm

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  $i = 1, \dots, n$

This is just the  
derivative of the  
**Loss Function...**

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

# GB Algorithm

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  $i = 1, \dots, n$

This is just the  
derivative of the  
**Loss Function...**

...with respect to the  
predicted **log(odds)**...

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

# GB Algorithm

**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  $i = 1, \dots, n$

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

...and we've already  
calculated this.

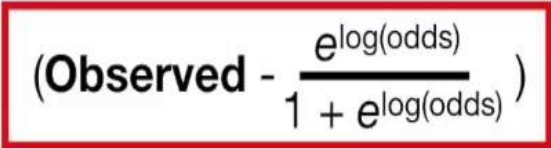

$$= \left( -\mathbf{Observed} + \frac{e^{\log(\text{odds})}}{1 + e^{\log(\text{odds})}} \right)$$

# GB Algorithm

**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  $i = 1, \dots, n$

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


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

# GB Algorithm

**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  $i = 1, \dots, n$

**NOTE:** As we have seen before, we can replace this term with the predicted probability,  $p$ ...

$(\text{Observed} - p)$

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

# GB Algorithm

**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  $i = 1, \dots, n$

...so we can think of the **Pseudo Residuals** as the **Observed** probability minus the **Predicted** probability...  $\rightarrow$  **(Observed -  $p$ )**

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

# GB Algorithm

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  $i = 1, \dots, n$

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



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



# GB Algorithm

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

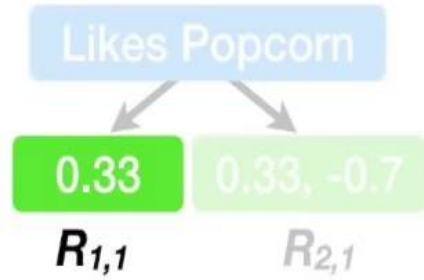
Step 2: for  $m = 1$  to  $M$ :

(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  $i = 1, \dots, n$

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

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

# GB Algorithm

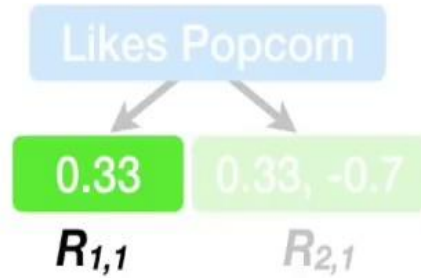


find the  
**Output Value** for this leaf.

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

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

# GB Algorithm



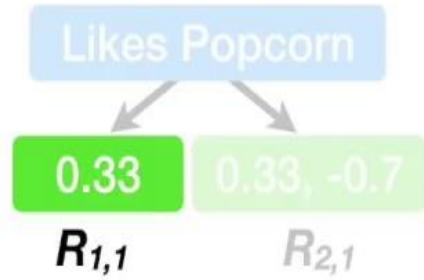
Likes Popcorn	Age	Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
No	87	Green	Yes
No	44	Blue	No

$$\frac{\sum \text{Residual}_i}{\sum [\text{Previous Probability}_i \times (1 - \text{Previous Probability}_i)]}$$

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

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

# GB Algorithm

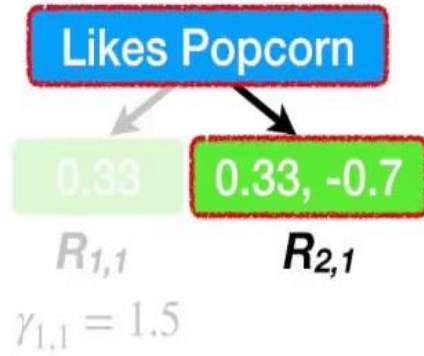


...and the output value for leaf  $R_{1,1}$  is **1.5**.

$$\gamma_{1,1} = 1.5$$

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

# GB Algorithm

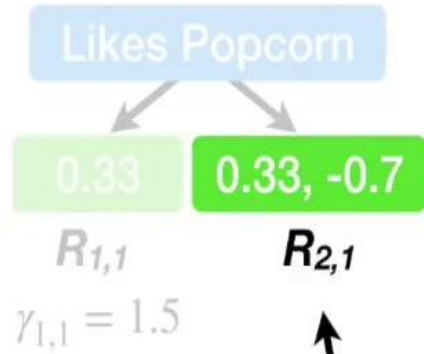


Likes Popcorn	Age	Favorite Color	Loves Troll 2
Yes	12	Blue	Yes
No	87	Green	Yes
No	44	Blue	No

$$\gamma_{2,1} = \frac{\text{Residual}_2 + \text{Residual}_3}{[p_2 \times (1 - p_2)] + [p_3 \times (1 - p_3)]}$$

$$\frac{\sum \text{Residual}_i}{\sum [\text{Previous Probability}_i \times (1 - \text{Previous Probability}_i)]}$$

# GB Algorithm



...and the output value  
for leaf  $R_{2,1}$  is **-0.77**.

$$\gamma_{2,1} = -0.77$$

# GB Algorithm

---

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



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

# GB Algorithm

$$F_1(x) = \overbrace{\log(2/1) = 0.69}^{F_0(x)} + 0.8 \times \begin{matrix} \text{Likes Popcorn} \\ \swarrow \quad \searrow \\ \boxed{0.33} \quad \boxed{0.33, -0.7} \\ R_{1,1} \quad R_{2,1} \end{matrix}$$

$$\gamma_{1,1} = 1.5 \quad \gamma_{2,1} = -0.77$$

...the **Output Values**  
from the first tree we  
made.

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





# 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

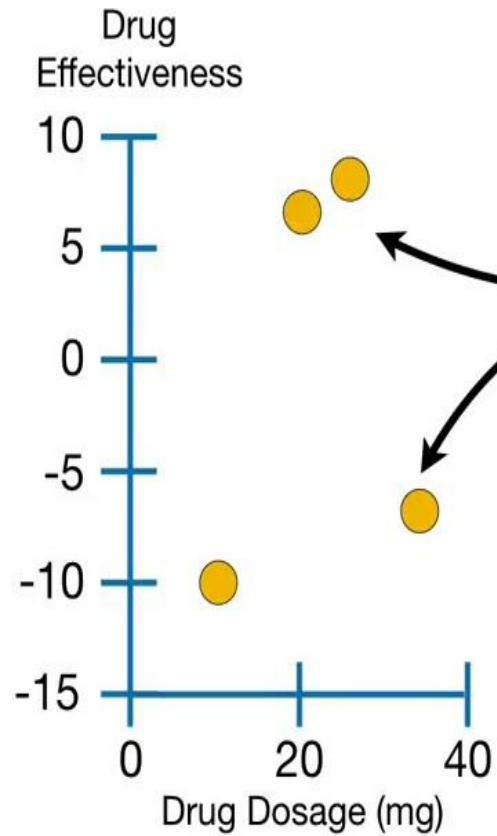
# XGBoost Characteristics

---

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

# XGBoost Tree

---

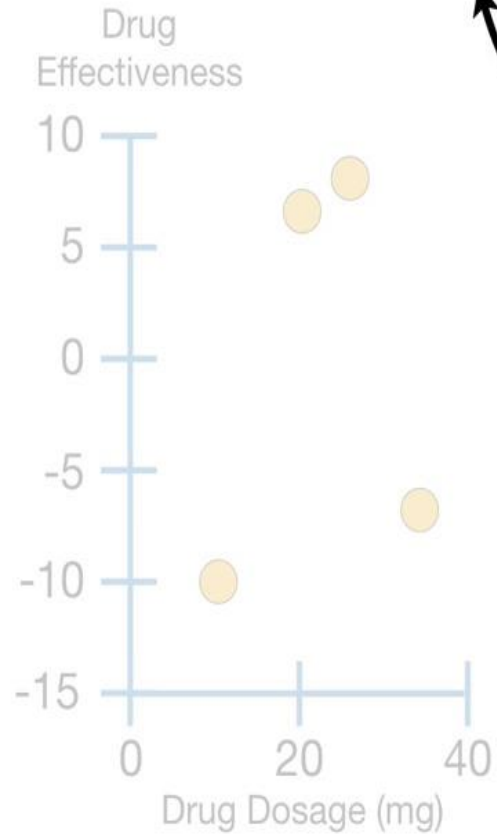


to keep the examples from getting out of hand, we will use use this super simple **Training Data**.

# XGBoost Tree

Predicted Drug  
Effectiveness

0.5



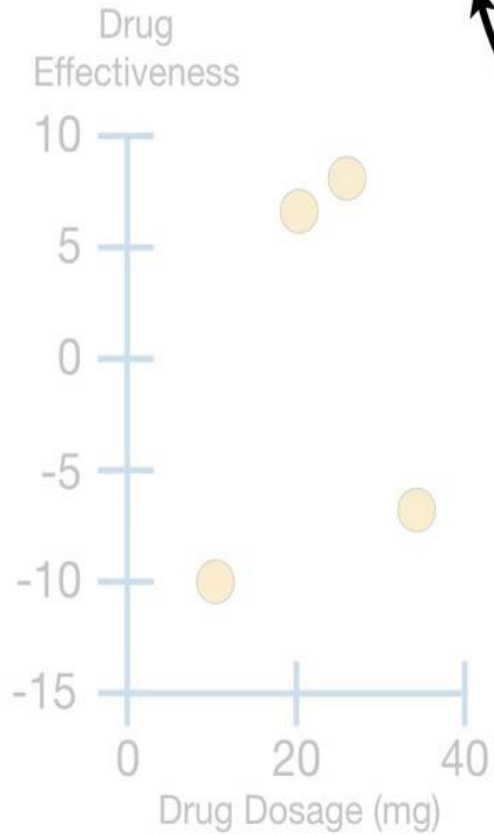
The very first step in fitting  
**XGBoost** to the **Training  
Data** is to make an initial  
prediction.

# XGBoost Tree

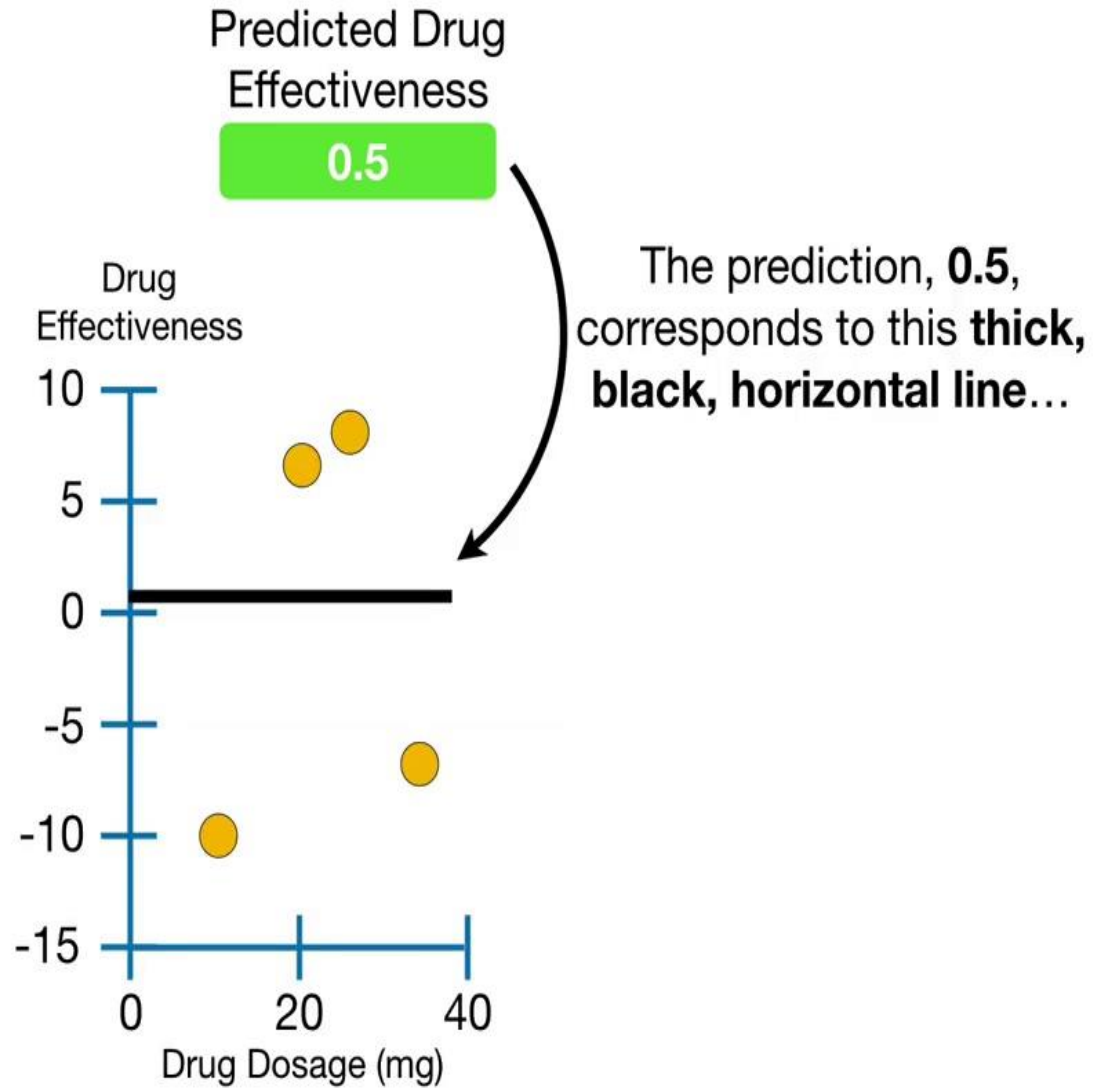
Predicted Drug  
Effectiveness

0.5

This prediction can be anything, but by default it is **0.5**, regardless of whether you are using **XGBoost** for **Regression** or **Classification**.



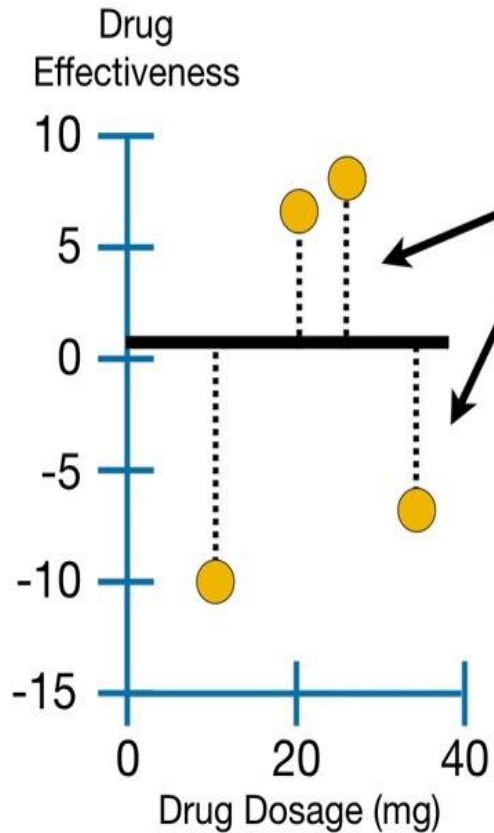
# XGBoost Tree



# XGBoost Tree

Predicted Drug  
Effectiveness

0.5

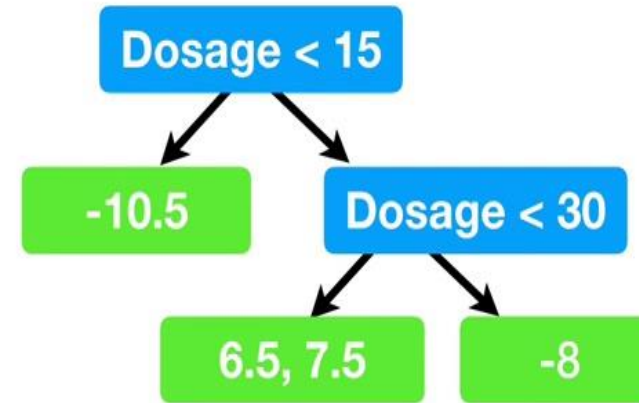
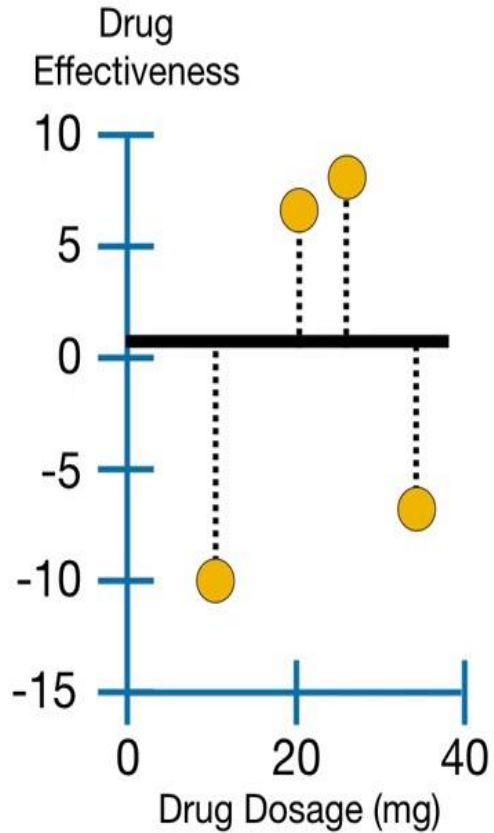


...and the **Residuals**, the differences between the **Observed** and **Predicted** values, show us how good the initial prediction is.



# XGBoost Tree

Predicted Drug Effectiveness  
0.5

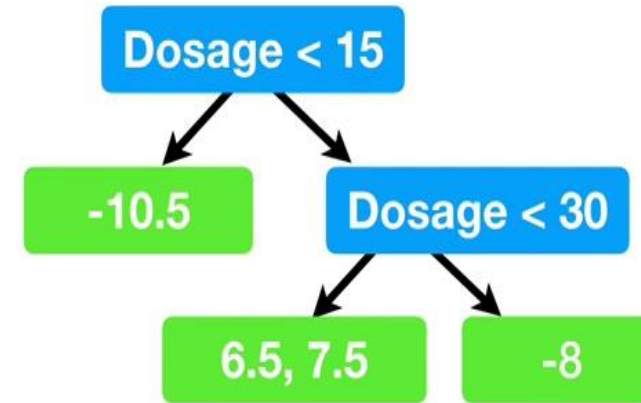
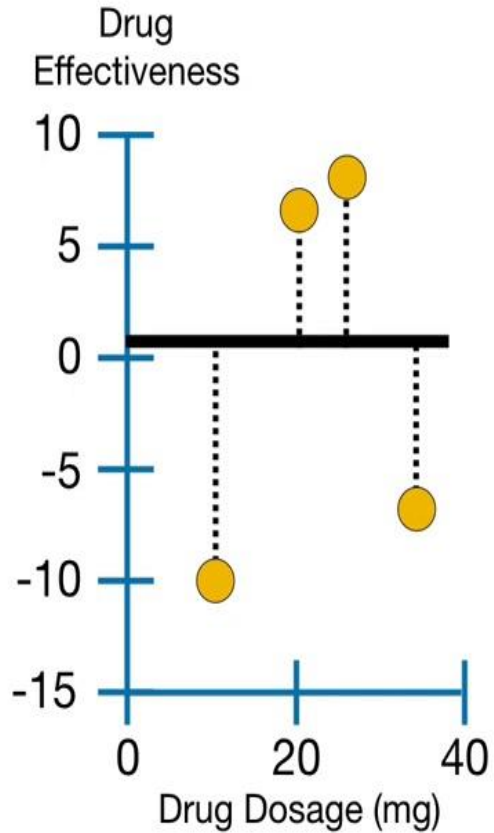


However, unlike unextreme **Gradient Boost**, which typically uses regular, off-the-shelf, **Regression Trees**...

# XGBoost Tree

Predicted Drug Effectiveness

0.5

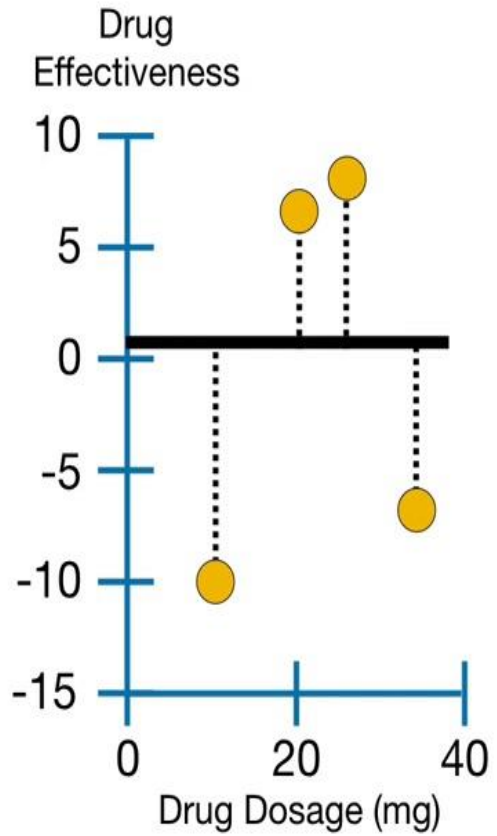


...XGBoost uses a unique Regression Tree that I call an XGBoost Tree.

# XGBoost Tree

Predicted Drug  
Effectiveness

0.5



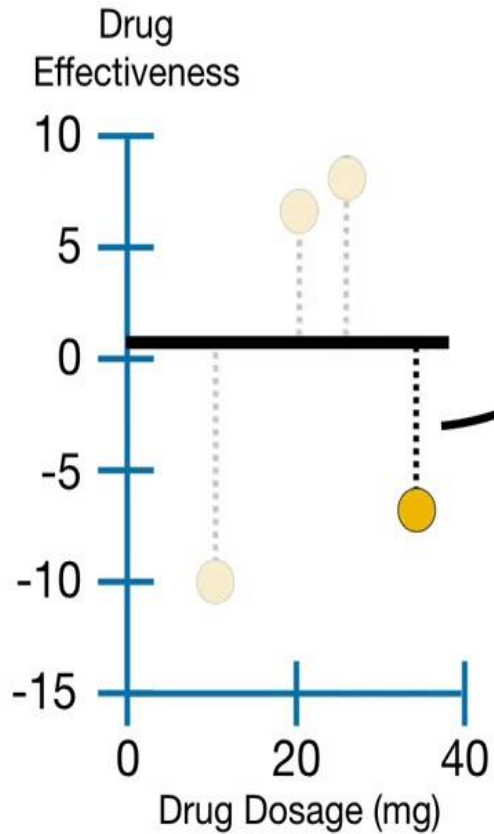
Each tree starts out as  
a single leaf...



# XGBoost Tree

Predicted Drug  
Effectiveness

0.5



-10.5, 6.5, 7.5, -7.5

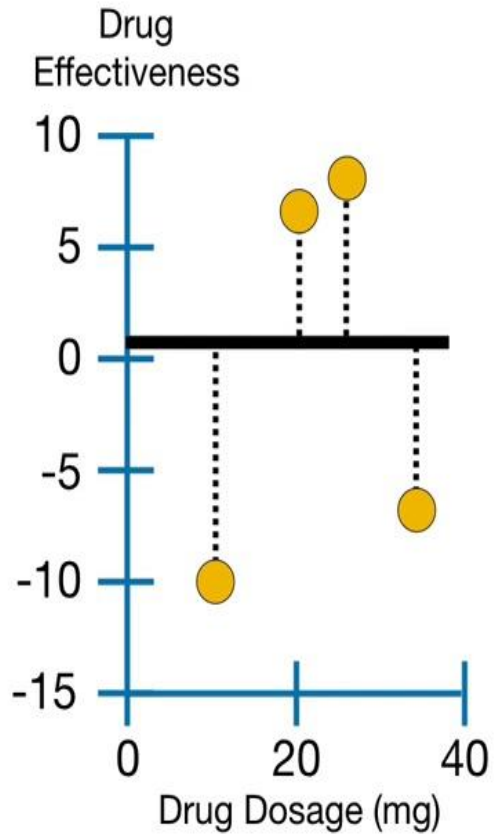
...and all of the  
**Residuals** go to the  
leaf.

# XGBoost Tree

Predicted Drug  
Effectiveness

0.5

-10.5, 6.5, 7.5, -7.5



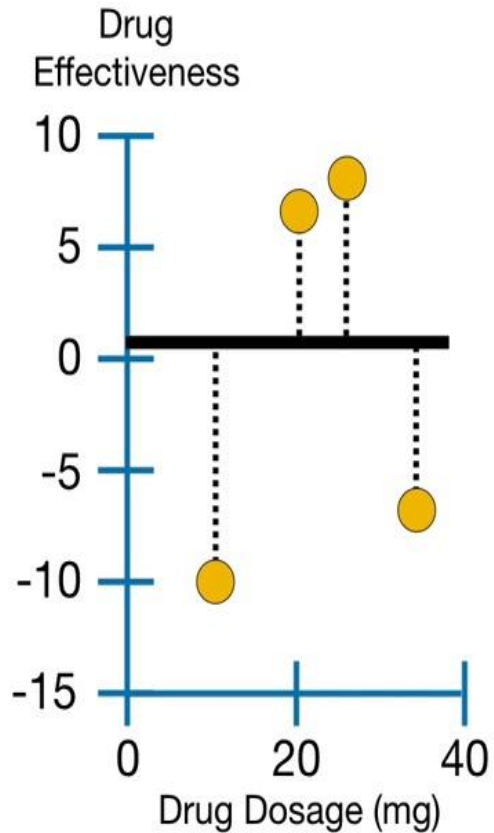
Now we calculate a **Quality Score**, or **Similarity Score**, for the **Residuals**.

# XGBoost Tree

Predicted Drug  
Effectiveness

0.5

-10.5, 6.5, 7.5, -7.5



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

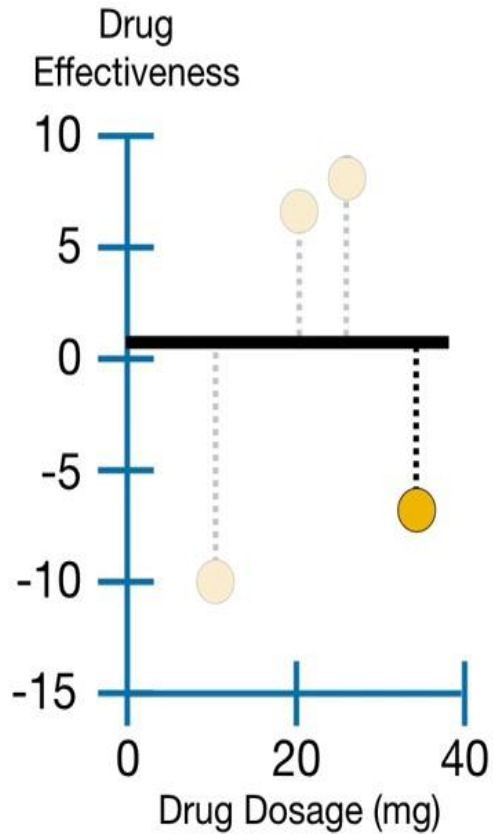
NOTE:  $\lambda$  (lambda) is a  
Regularization parameter,

# XGBoost Tree

Predicted Drug  
Effectiveness

0.5

-10.5, 6.5, 7.5, -7.5



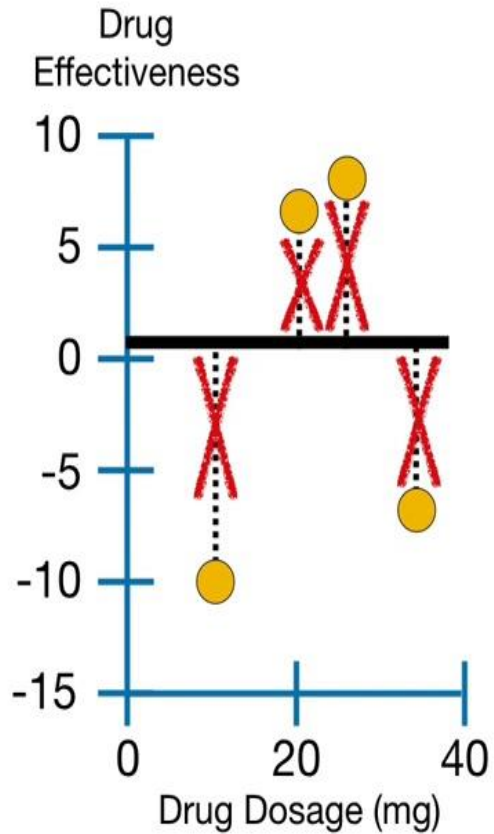
$$\text{Similarity Score} = \frac{(-10.5 + 6.5 + 7.5 + -7.5)^2}{4 + 0}$$

# XGBoost Tree

Predicted Drug Effectiveness

0.5

-10.5, 6.5, 7.5, -7.5



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

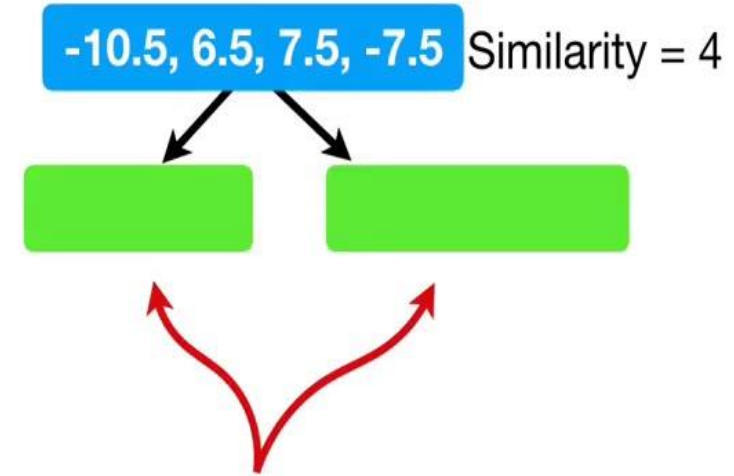
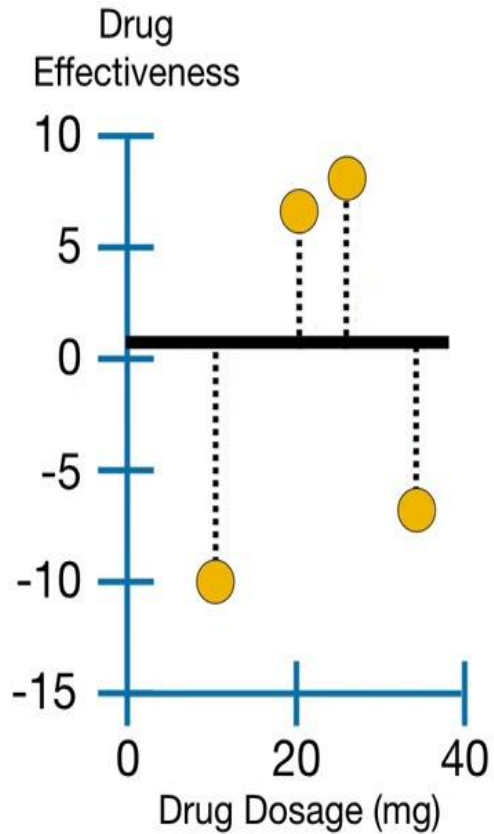
Thus, the **Similarity Score** for the **Residuals** in the root = 4.



# XGBoost Tree

Predicted Drug Effectiveness

0.5



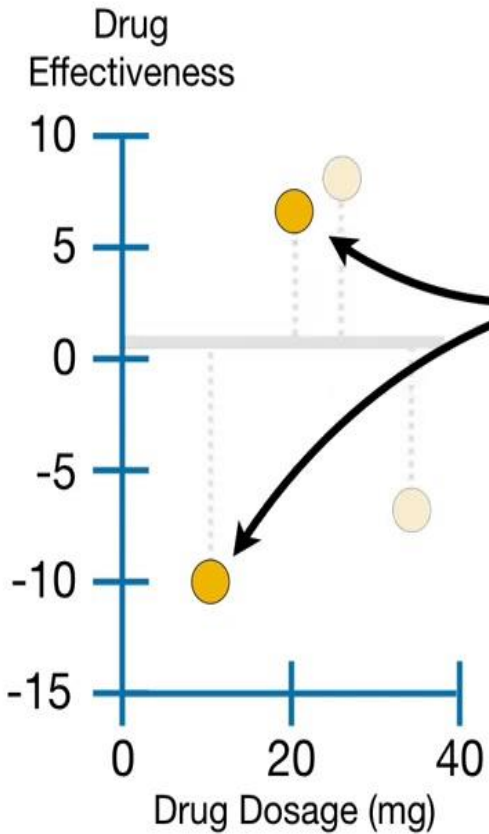
Now the question is whether or not we can do a better job clustering similar **Residuals** if we split them into two groups.

# XGBoost Tree

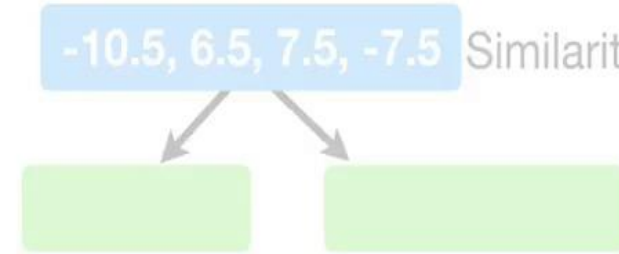
Predicted Drug Effectiveness

0.5

-10.5, 6.5, 7.5, -7.5 Similarity = 4



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

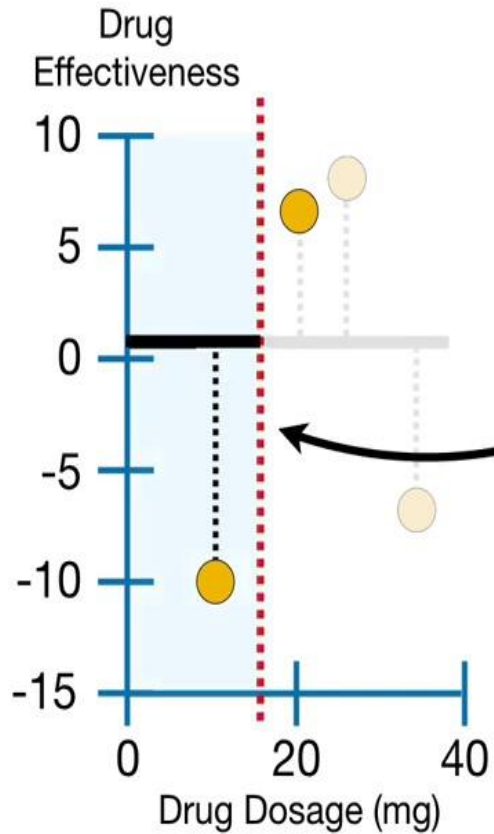


# XGBoost Tree

Predicted Drug Effectiveness

0.5

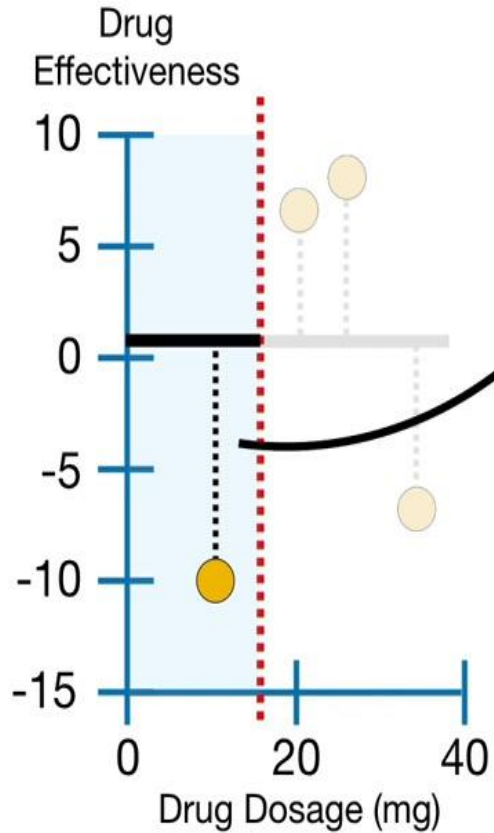
-10.5, 6.5, 7.5, -7.5 Similarity = 4



Their average **Dosage** is **15**, and that corresponds to this **dotted red line**.

# XGBoost Tree

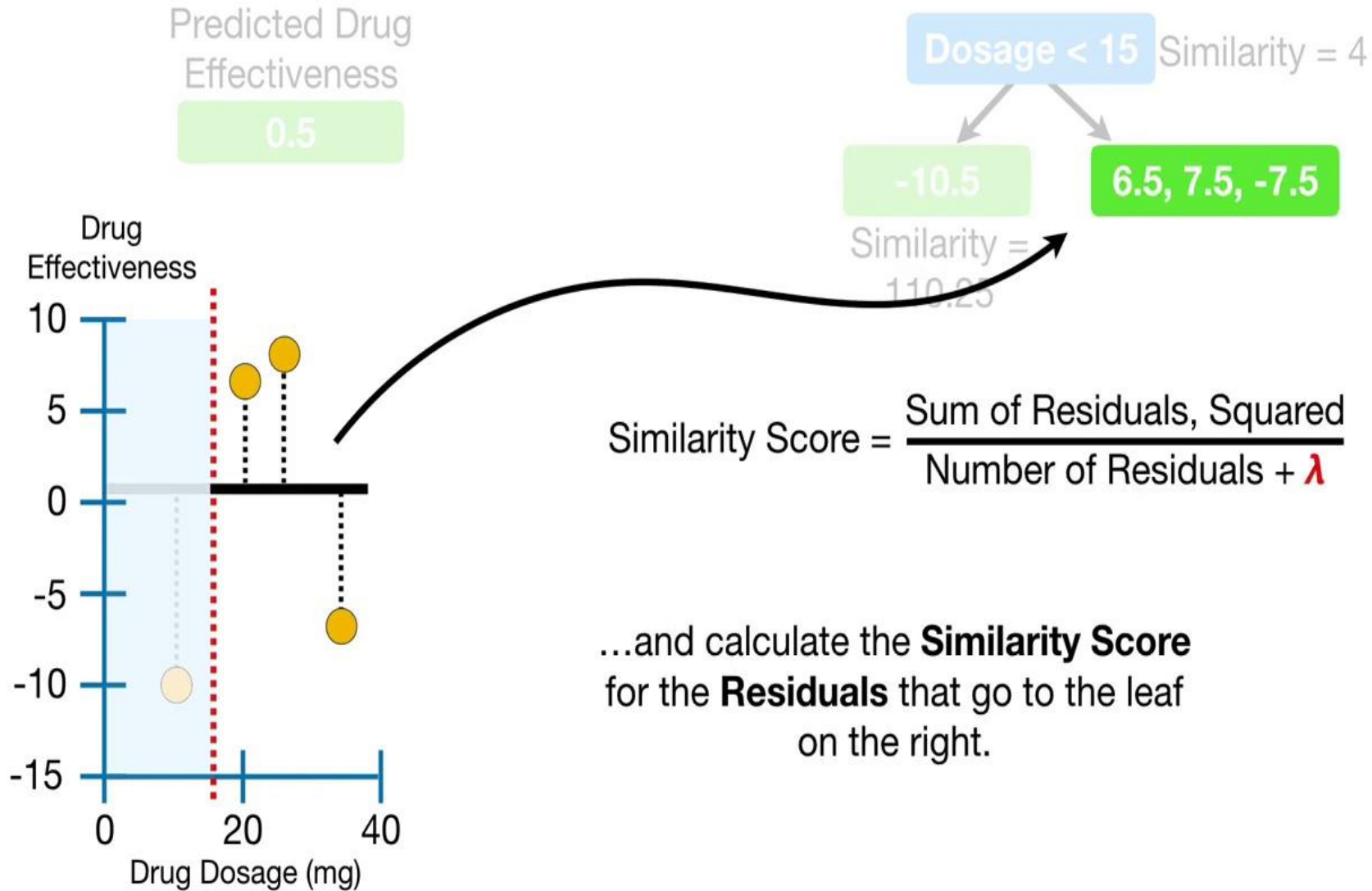
Predicted Drug Effectiveness  
0.5



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

...by plugging the one **Residual** into the numerator...

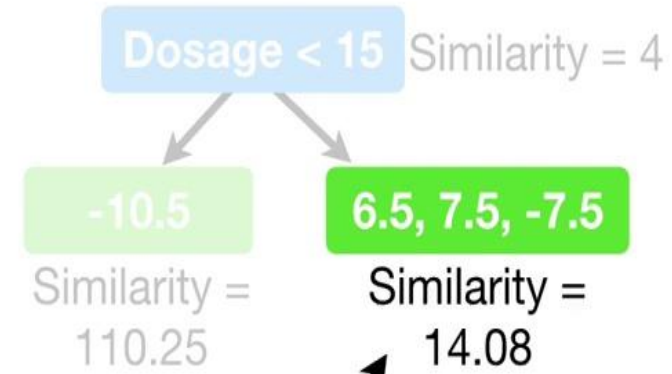
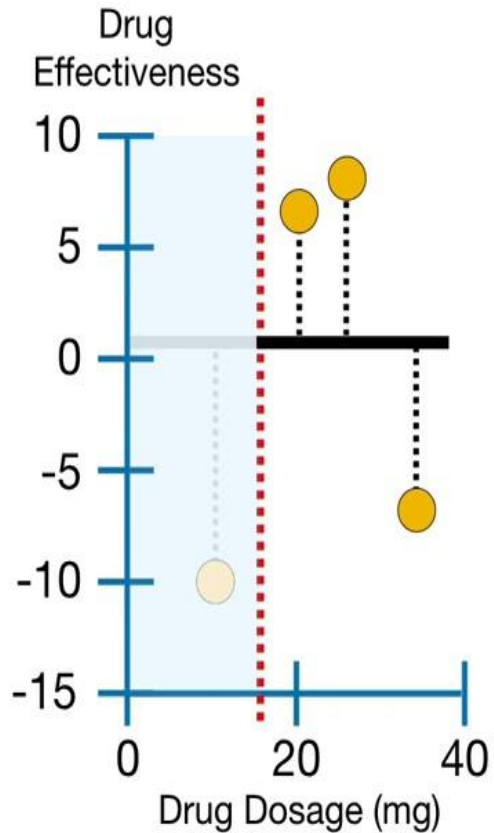
# XGBoost Tree



# XGBoost Tree

Predicted Drug Effectiveness

0.5



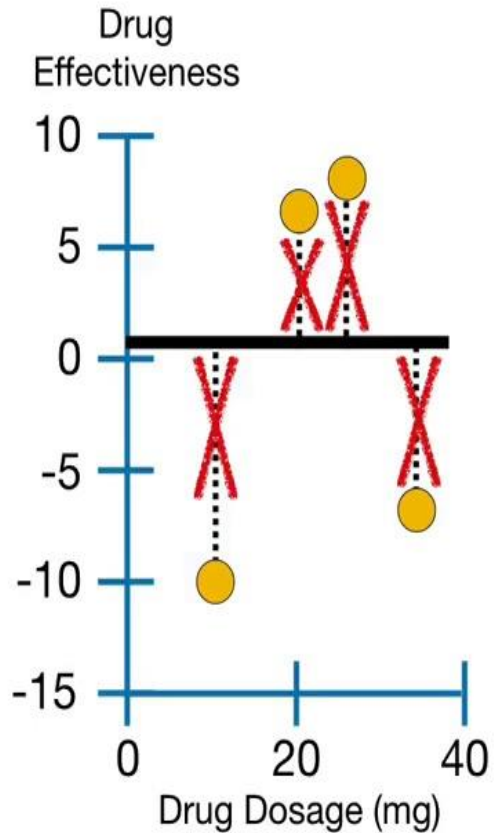
$$\text{Similarity Score} = \frac{6.5^2}{3 + 0} = 14.08$$

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

# XGBoost Tree

Predicted Drug Effectiveness

0.5



-10.5, 6.5, 7.5, -7.5 Similarity = 4

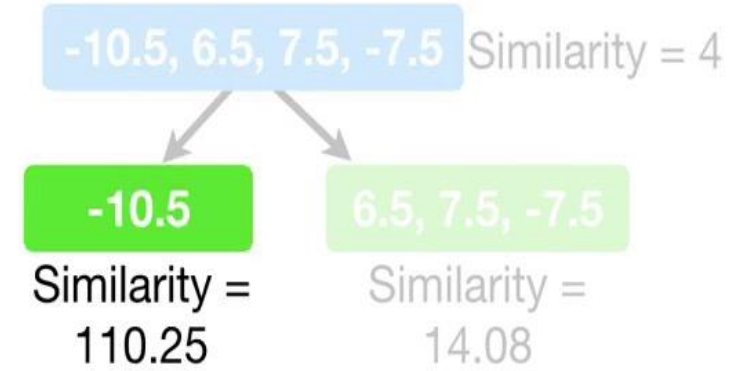
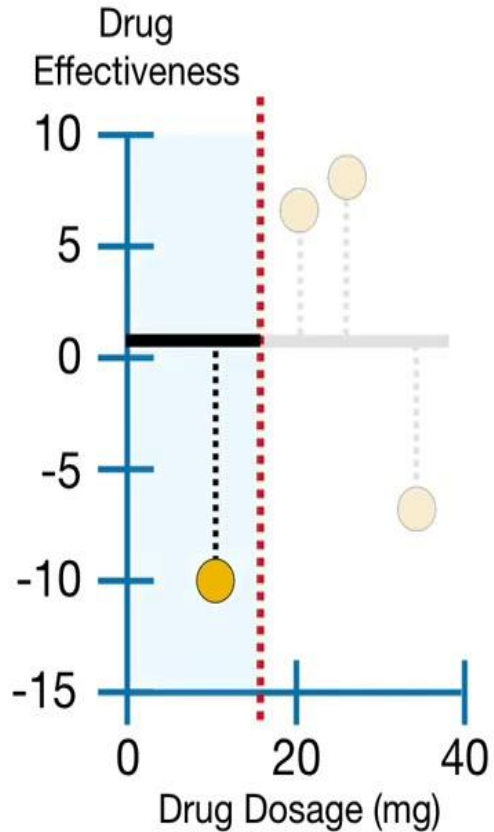
-10.5  
Similarity =  
110.25

6.5, 7.5, -7.5  
Similarity =  
14.08

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

# XGBoost Tree

Predicted Drug Effectiveness  
0.5



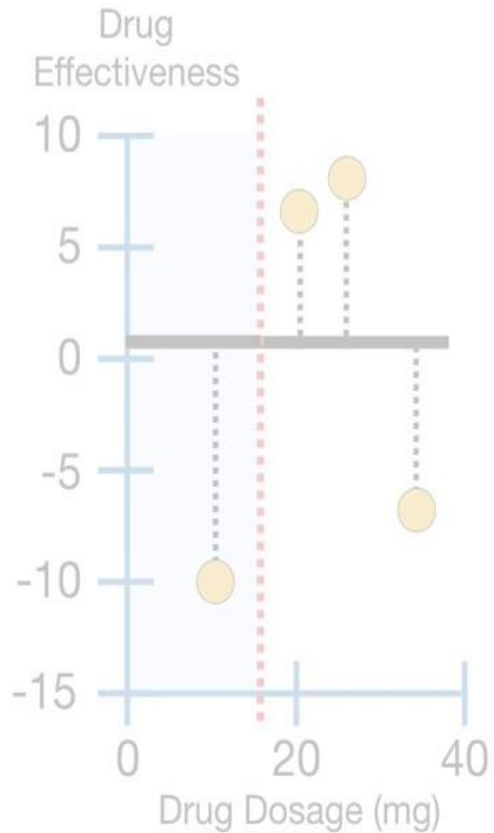
In contrast, when the **Residuals** are similar, or there is just one of them, they do not cancel out and the **Similarity Score** is relatively large.



# XGBoost Tree

Predicted Drug Effectiveness

0.5



-10.5, 6.5, 7.5, -7.5 Similarity = 4

-10.5

Similarity =  
110.25

6.5, 7.5, -7.5

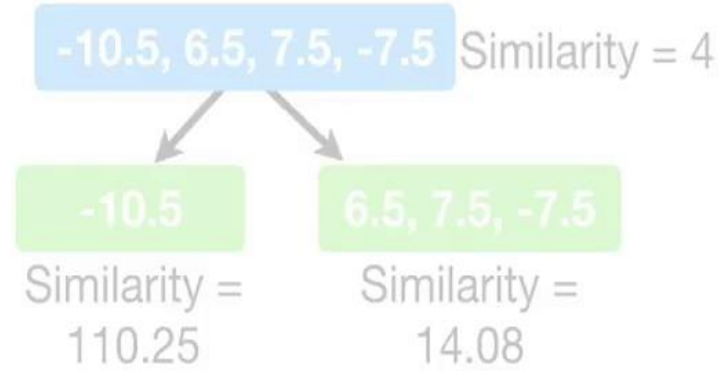
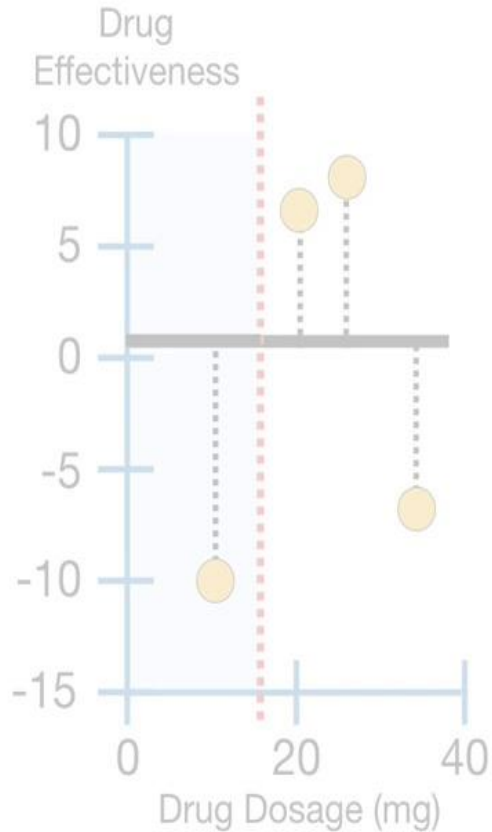
Similarity =  
14.08

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

# XGBoost Tree

Predicted Drug Effectiveness

0.5



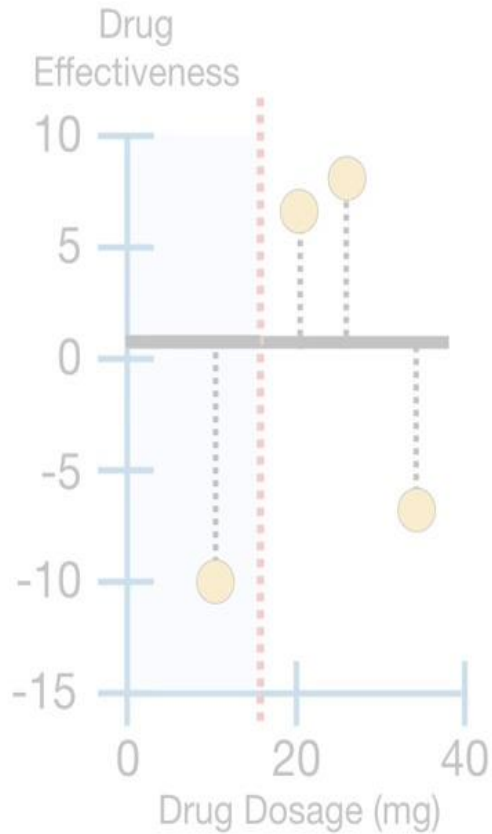
$$\text{Gain} = \text{Left}_{\text{Similarity}} + \text{Right}_{\text{Similarity}} - \text{Root}_{\text{Similarity}}$$

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

# XGBoost Tree

Predicted Drug Effectiveness

0.5



-10.5, 6.5, 7.5, -7.5 Similarity = 4

-10.5

Similarity =  
110.25

6.5, 7.5, -7.5

Similarity =  
14.08

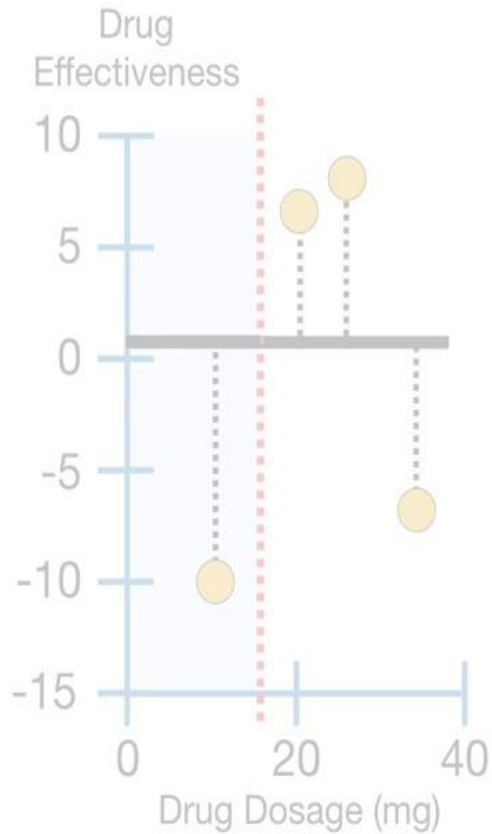
$$\text{Gain} = \text{Left}_{\text{Similarity}} + \text{Right}_{\text{Similarity}} - \text{Root}_{\text{Similarity}}$$

**Gain** is equal to the **Similarity Score** for the leaf on the left...

# XGBoost Tree

Predicted Drug Effectiveness

0.5



-10.5, 6.5, 7.5, -7.5 Similarity = 4

-10.5

Similarity =  
110.25

6.5, 7.5, -7.5

Similarity =  
14.08

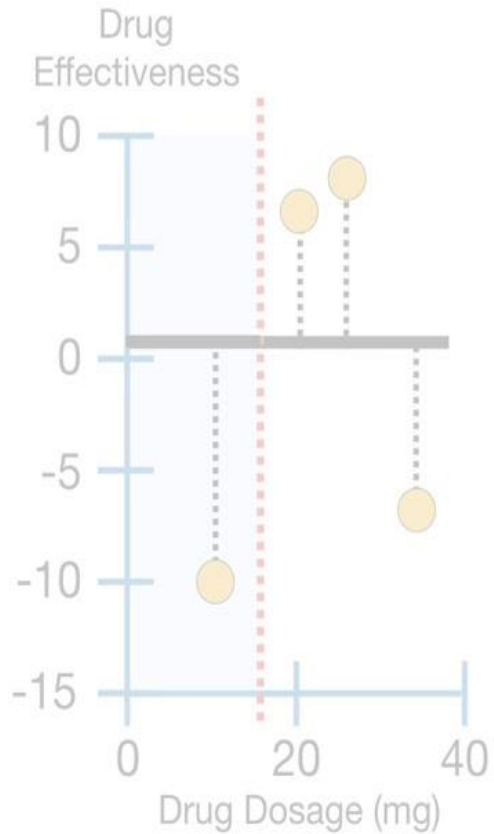
$$\text{Gain} = \text{Left}_{\text{Similarity}} + \text{Right}_{\text{Similarity}} - \text{Root}_{\text{Similarity}}$$

...plus the **Similarity Score** for the leaf on the right...

# XGBoost Tree

Predicted Drug Effectiveness

0.5



-10.5, 6.5, 7.5, -7.5 Similarity = 4

-10.5

Similarity =  
110.25

6.5, 7.5, -7.5

Similarity =  
14.08

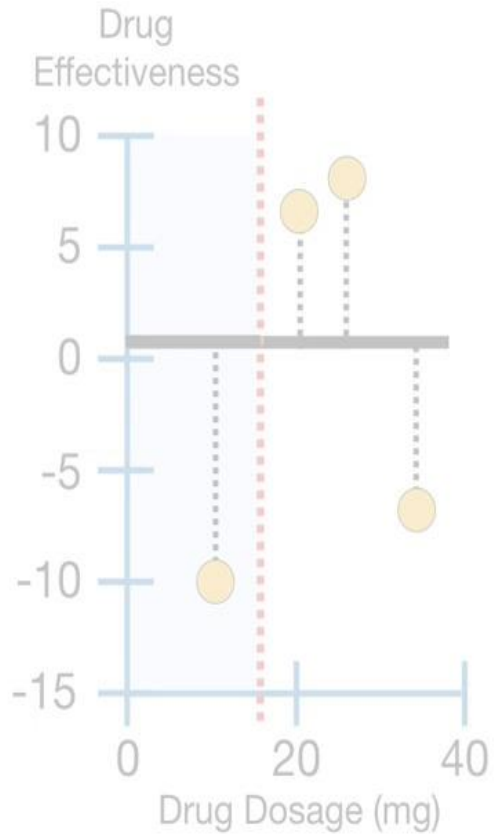
$$\text{Gain} = \text{Left}_{\text{Similarity}} + \text{Right}_{\text{Similarity}} - \text{Root}_{\text{Similarity}}$$

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

# XGBoost Tree

Predicted Drug Effectiveness

0.5



-10.5, 6.5, 7.5, -7.5 Similarity = 4

-10.5

Similarity =  
110.25

6.5, 7.5, -7.5

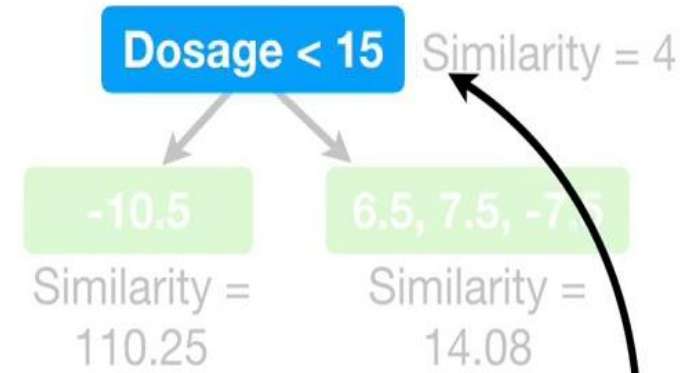
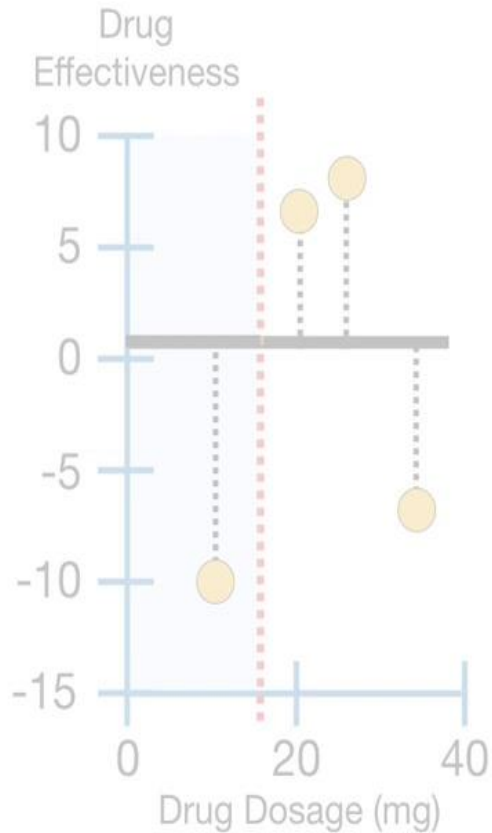
Similarity =  
14.08

$$\text{Gain} = 110.25 + 14.08 - 4 = 120.33$$

# XGBoost Tree

Predicted Drug Effectiveness

0.5



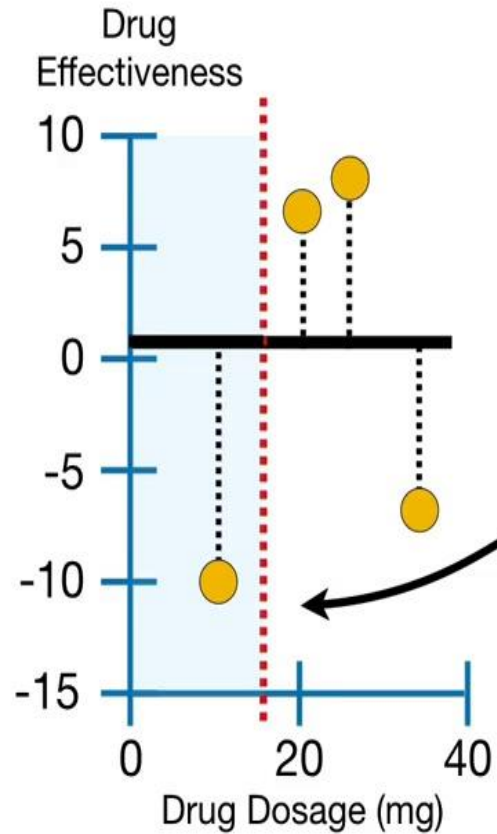
$$\text{Gain} = 110.25 + 14.08 - 4 = 120.33$$

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

# XGBoost Tree

Predicted Drug  
Effectiveness

0.5



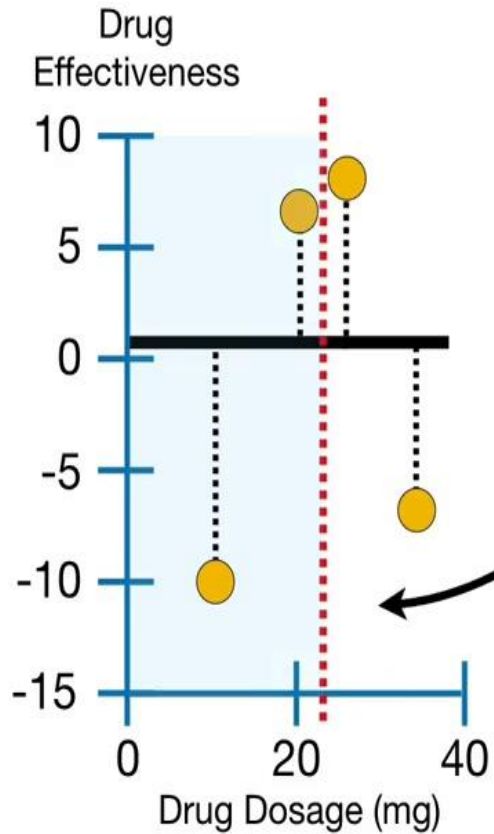
So we shift the threshold  
over so that it is the  
average of the next two  
observations...



# XGBoost Tree

Predicted Drug  
Effectiveness

0.5

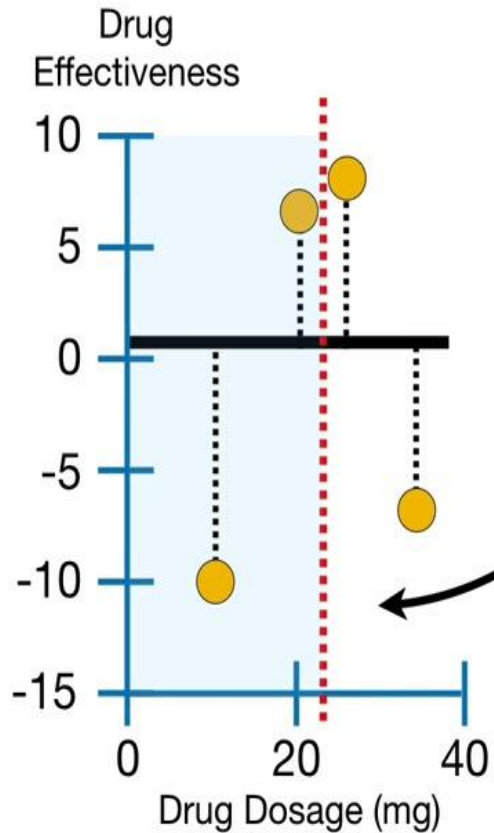


So we shift the threshold over so that it is the average of the next two observations...

# XGBoost Tree

Predicted Drug Effectiveness

0.5



Dosage < 22.5 Similarity = 4

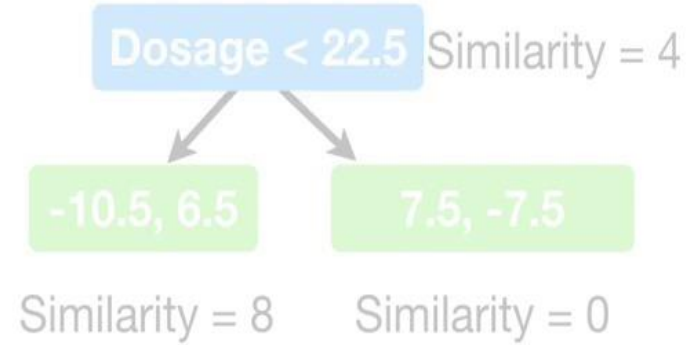
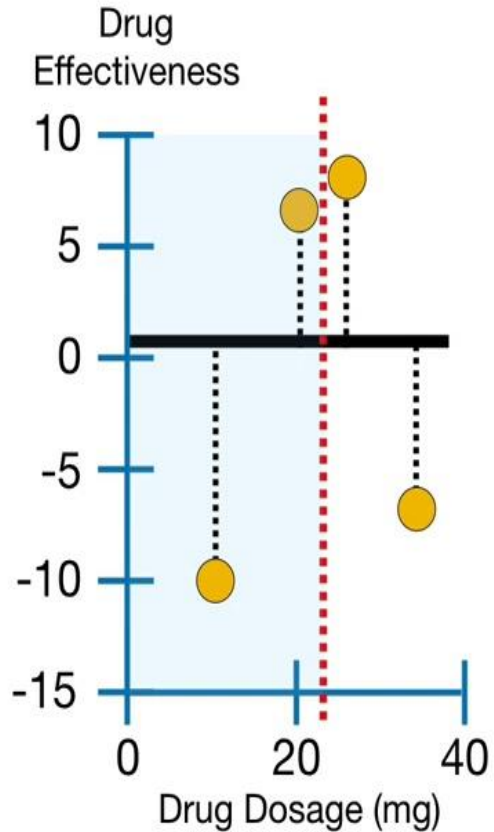
-10.5, 6.5

7.5, -7.5

...and build a simple tree that divides the observations using the new threshold, **Dosage < 22.5.**

# XGBoost Tree

Predicted Drug Effectiveness  
0.5



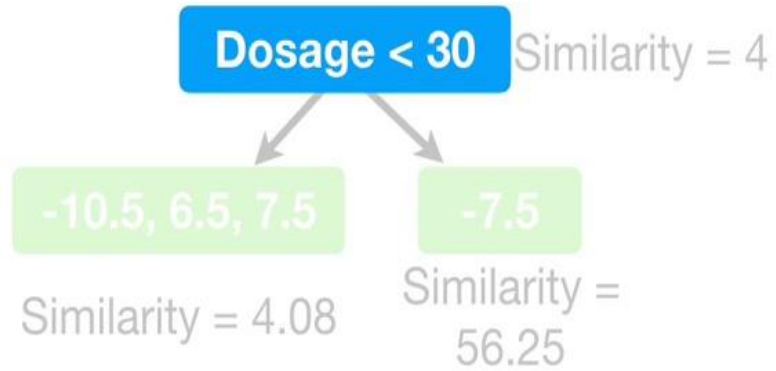
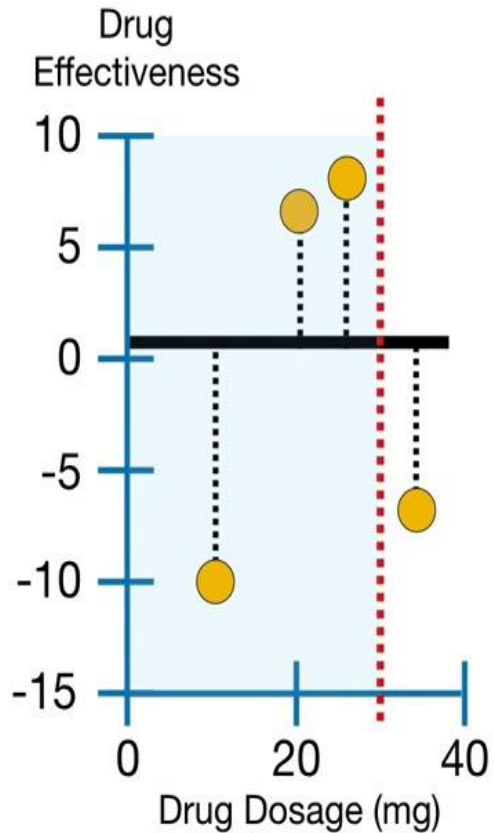
$$\text{Gain} = 8 + 0 - 4 = 4$$

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.

# XGBoost Tree

Predicted Drug Effectiveness

0.5

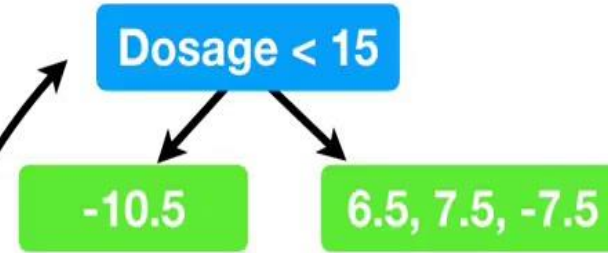
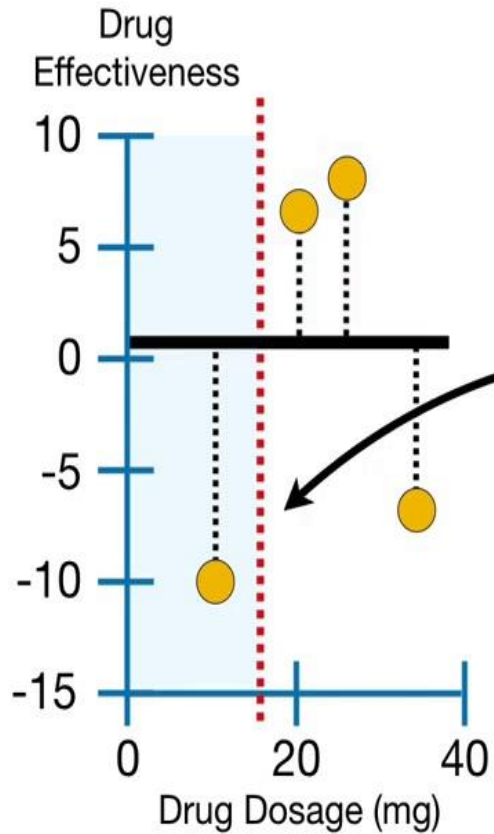


$$\text{Gain} = 4.08 + 56.25 - 4 = 56.33$$

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.

# XGBoost Tree

Predicted Drug Effectiveness  
0.5

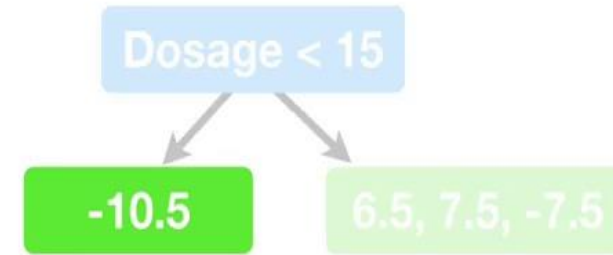
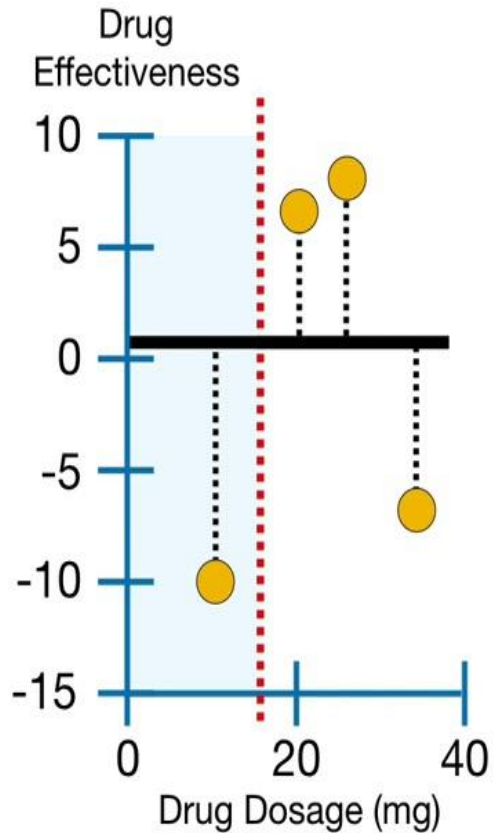


...and we will use the threshold that gave us the largest **Gain**, **Dosage < 15**, for the first branch in the tree.

# XGBoost Tree

Predicted Drug Effectiveness

0.5

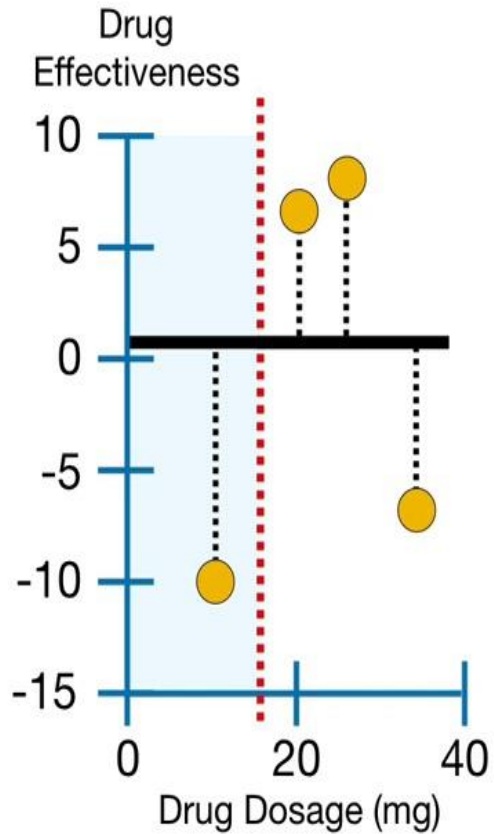


Now, since there is only one **Residual** in the leaf on the left, we can't split it any further.

# XGBoost Tree

Predicted Drug Effectiveness

0.5

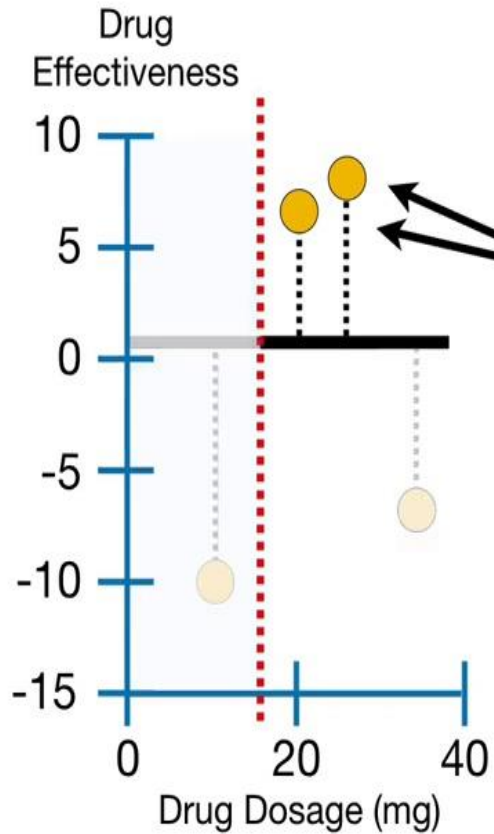


However, we can split the **3 Residuals** in the leaf on the right.

# XGBoost Tree

Predicted Drug Effectiveness

0.5



So we start with these two observations...

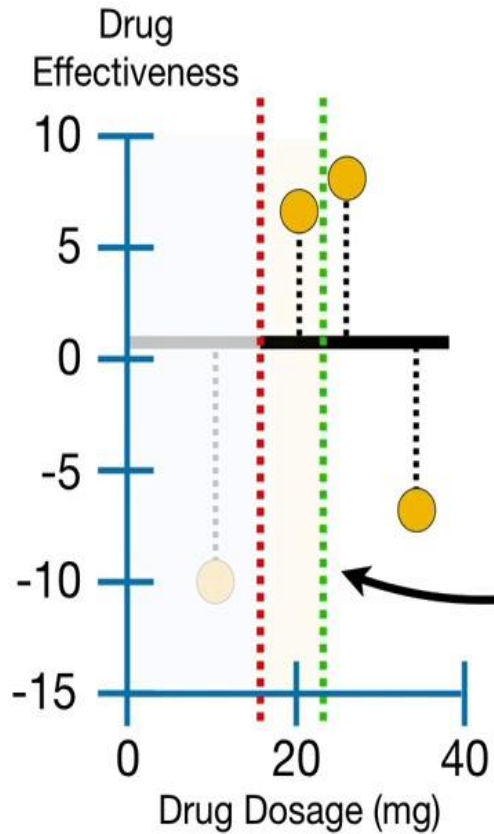




# XGBoost Tree

Predicted Drug Effectiveness

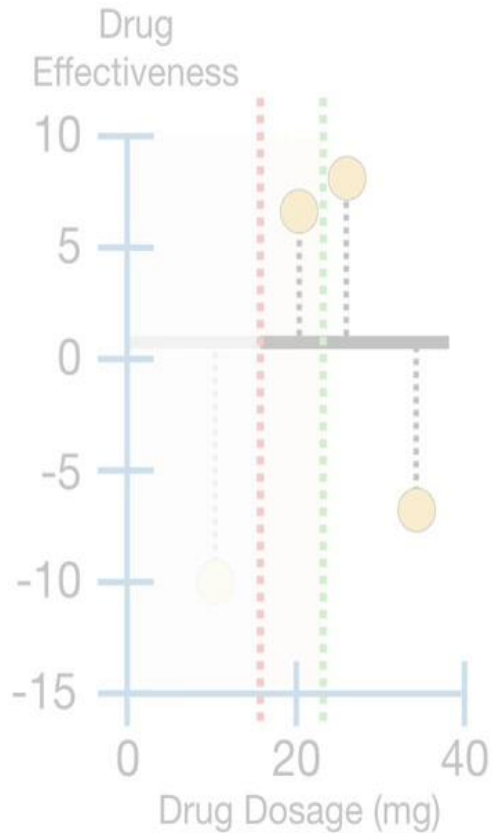
0.5



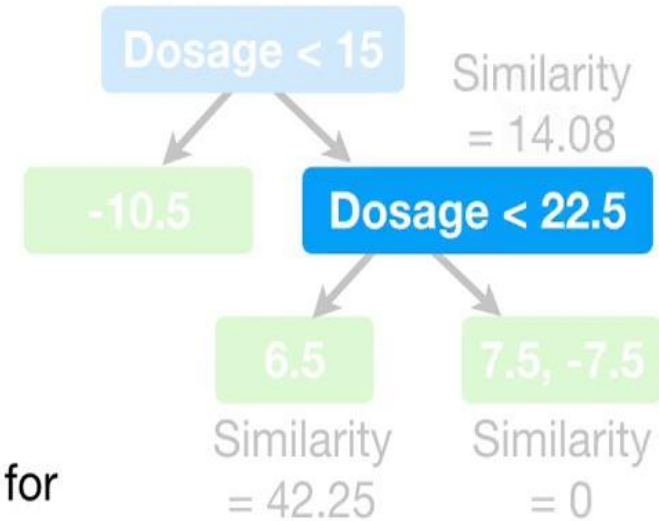
...and their average **Dosage** is **22.5**, which corresponds to this **dotted green line**.

# XGBoost Tree

Predicted Drug Effectiveness  
0.5



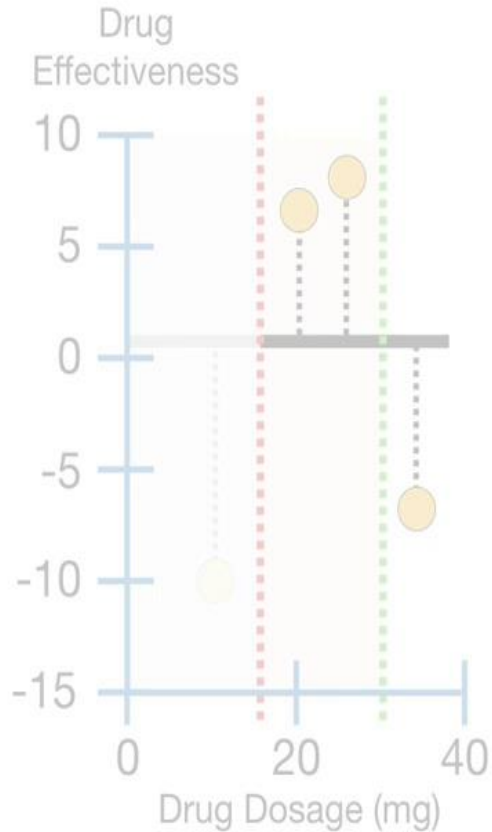
And we get **Gain = 28.17** for when the threshold is **Dosage < 22.5**.



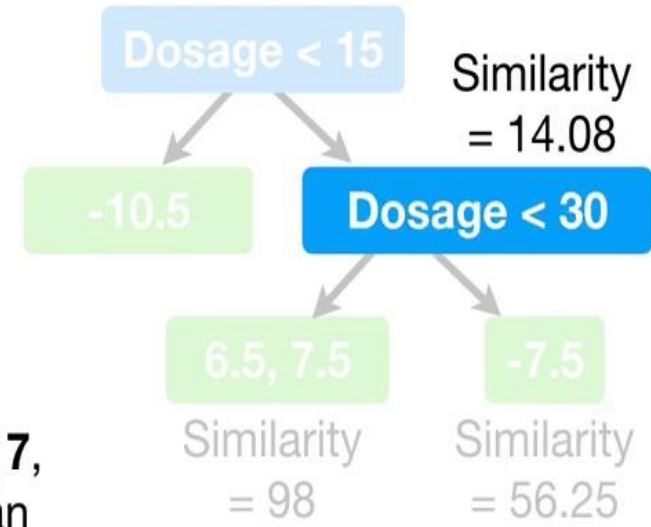
$$\text{Gain} = 42.25 + 0 - 14.08 = 28.17$$

# XGBoost Tree

Predicted Drug Effectiveness  
0.5



And we get **Gain = 140.17**, which is much larger than **28.17**, when the threshold was **Dosage < 22.5**.

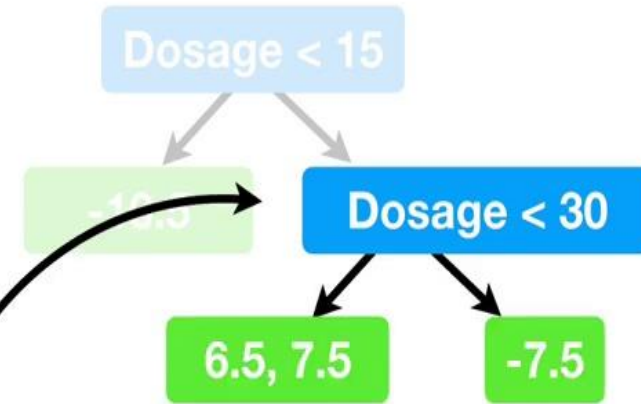
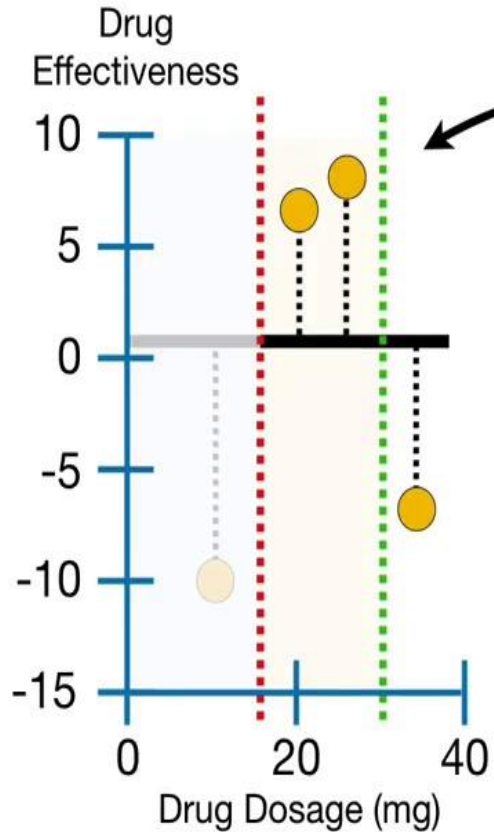


$$\text{Gain} = 98 + 56.25 - 14.08 = 140.17$$

# XGBoost Tree

Predicted Drug Effectiveness

0.5

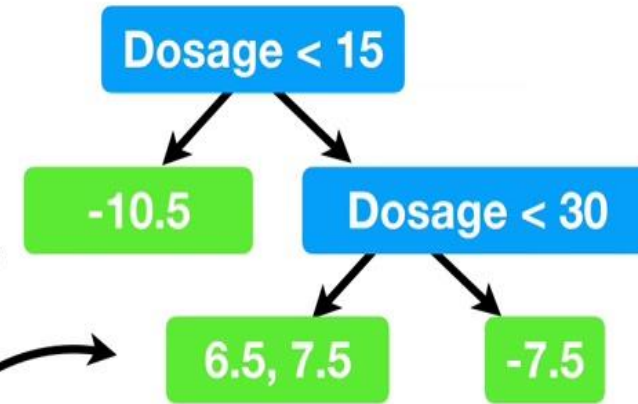
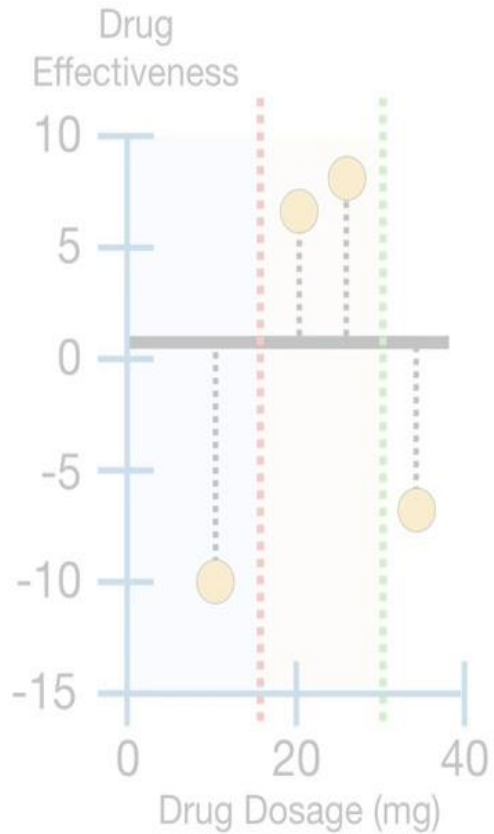


So we will use **Dosage < 30** as the threshold for this branch.

$$\text{Gain} = 98 + 56.25 - 14.08 = 140.17$$

# XGBoost Tree

Predicted Drug Effectiveness  
0.5

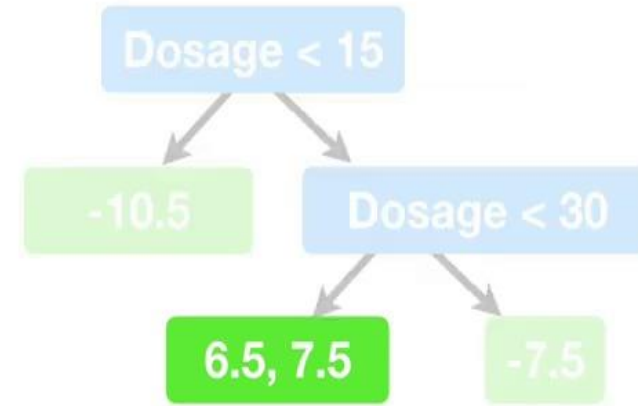
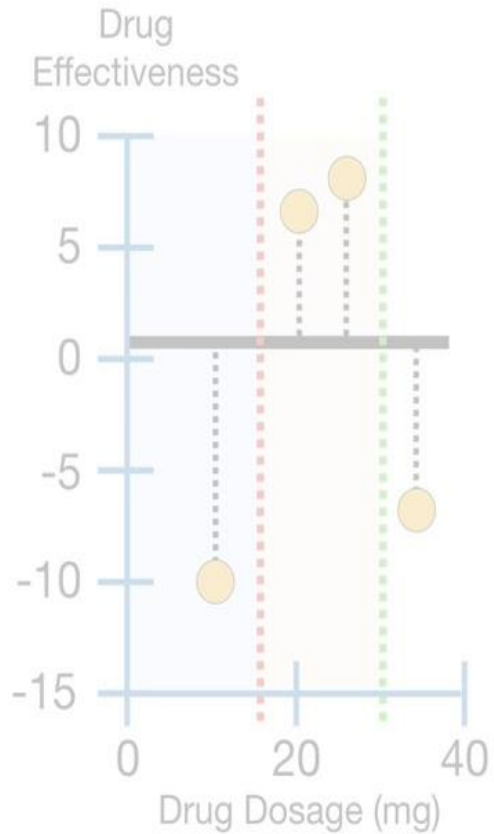


**NOTE:** To keep this example from getting out of hand, I've limited the tree depth to two levels...

# XGBoost Tree

Predicted Drug Effectiveness

0.5

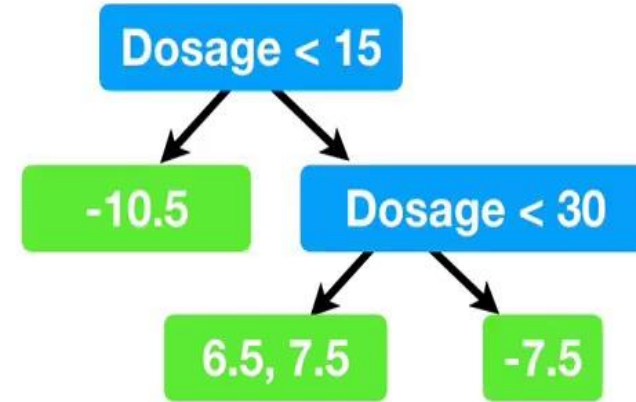
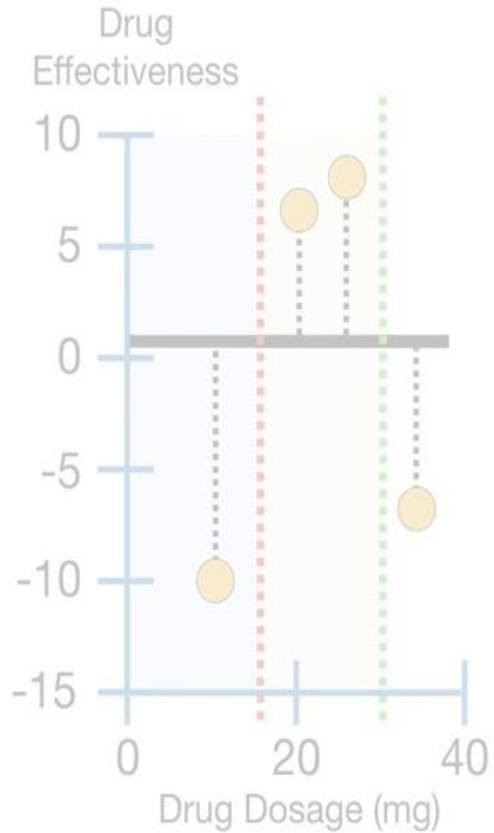


...and this means we will not split this leaf any further, and we are done building this tree.

# XGBoost Tree

Predicted Drug  
Effectiveness

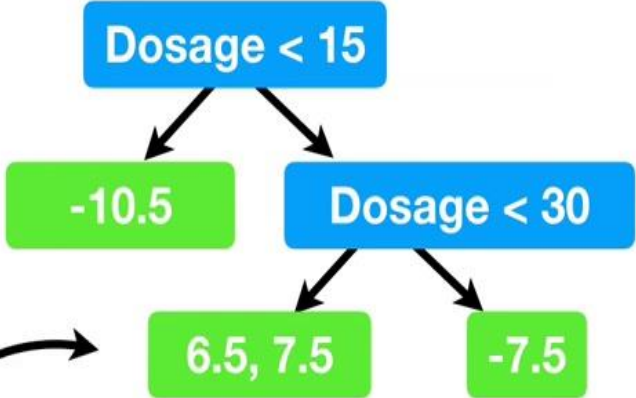
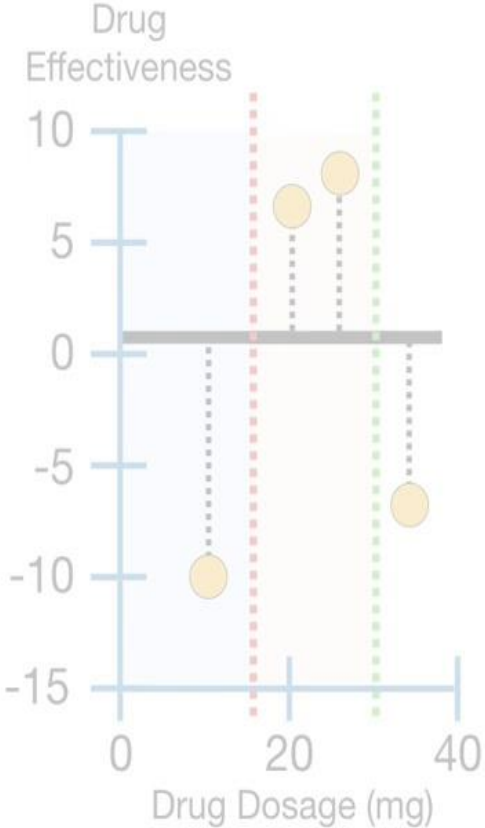
0.5



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

# Tree Pruning

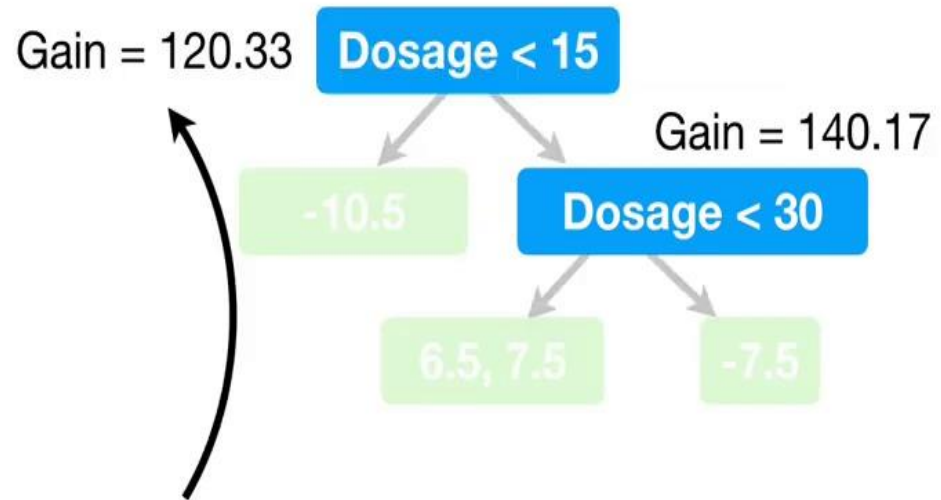
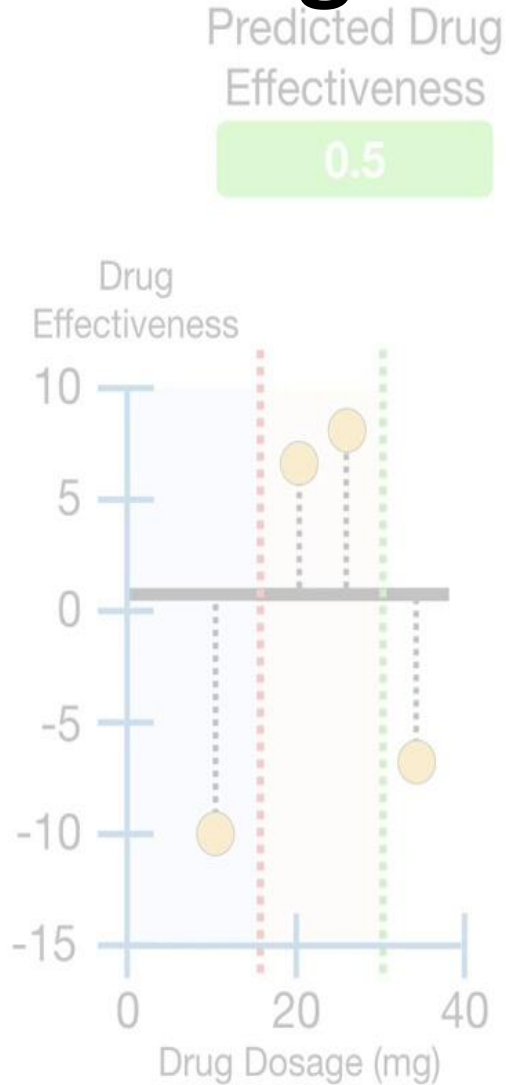
Predicted Drug Effectiveness  
0.5



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



# Tree Pruning



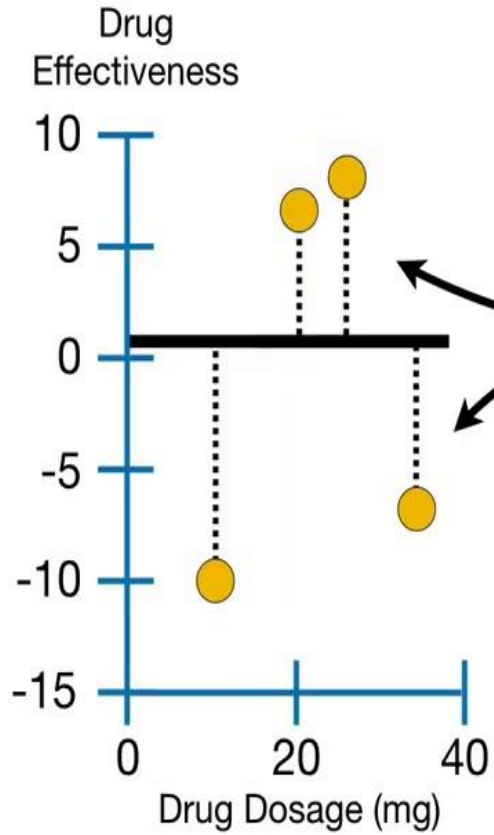
We **Prune** an **XGBoost Tree** based on its **Gain** values.

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

# Tree Pruning

Predicted Drug  
Effectiveness

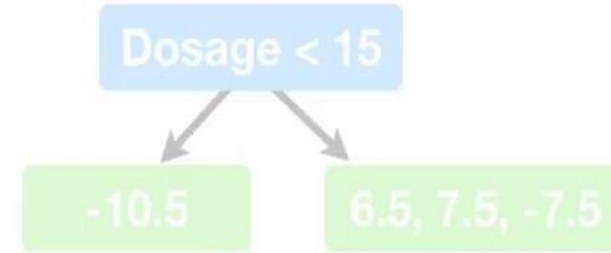
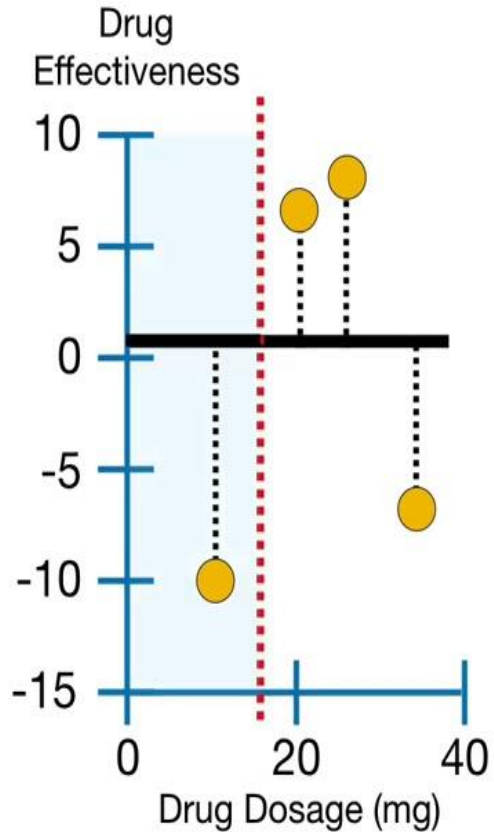
0.5



Now let's go back to the  
original **Residuals**...

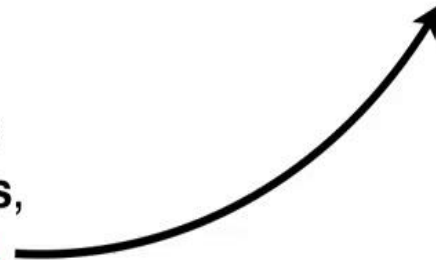
# Tree Pruning

Predicted Drug Effectiveness  
0.5



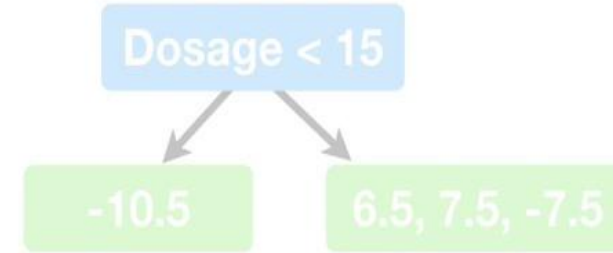
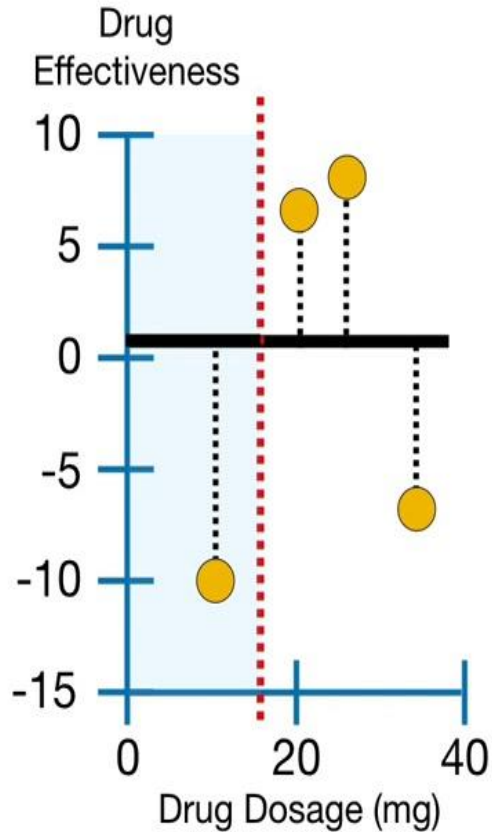
$$\text{Similarity Score} = \frac{\text{Sum of Residuals, Squared}}{\text{Number of Residuals} + \lambda}$$

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



# Tree Pruning

Predicted Drug Effectiveness  
0.5



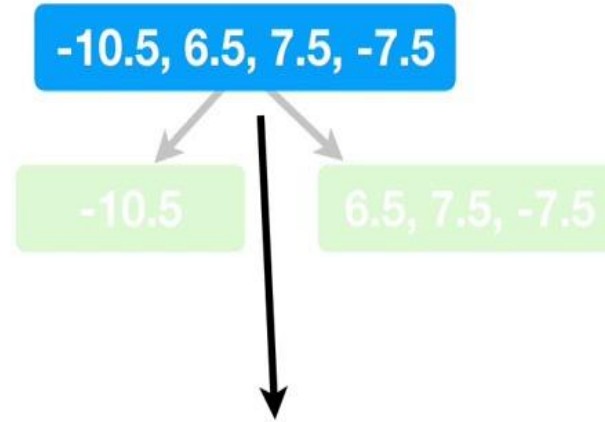
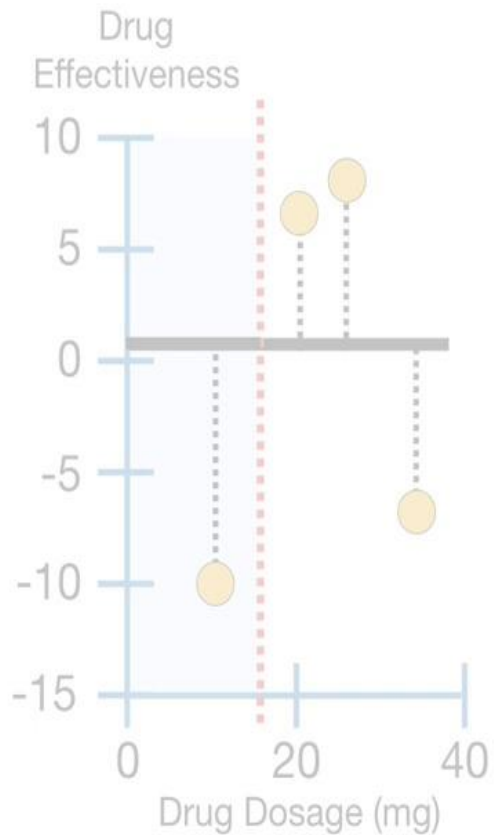
$$\text{Similarity Score} = \frac{\text{Sum of Residuals, Squared}}{\text{Number of Residuals} + 1}$$

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

# Tree Pruning

Predicted Drug Effectiveness

0.5



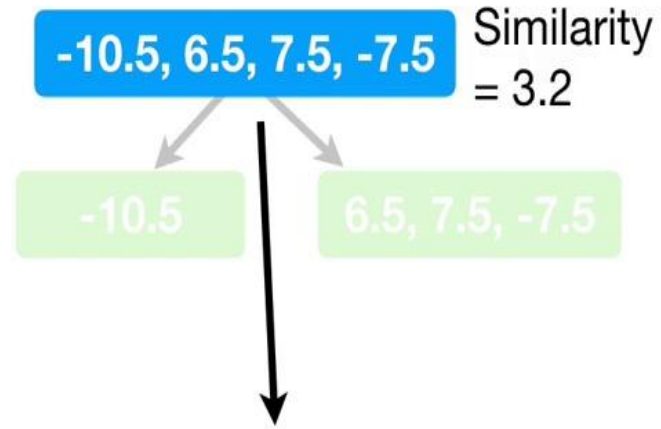
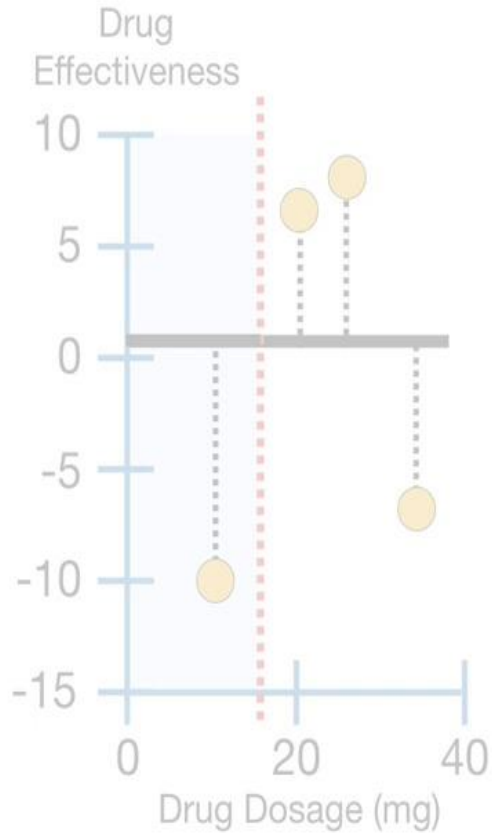
$$\text{Similarity Score} = \frac{(-10.5 + 6.5 + 7.5 + -7.5)^2}{4 + 1}$$

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

# Tree Pruning

Predicted Drug Effectiveness

0.5

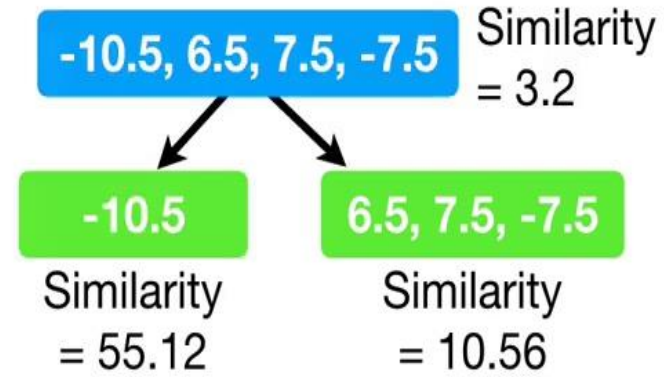
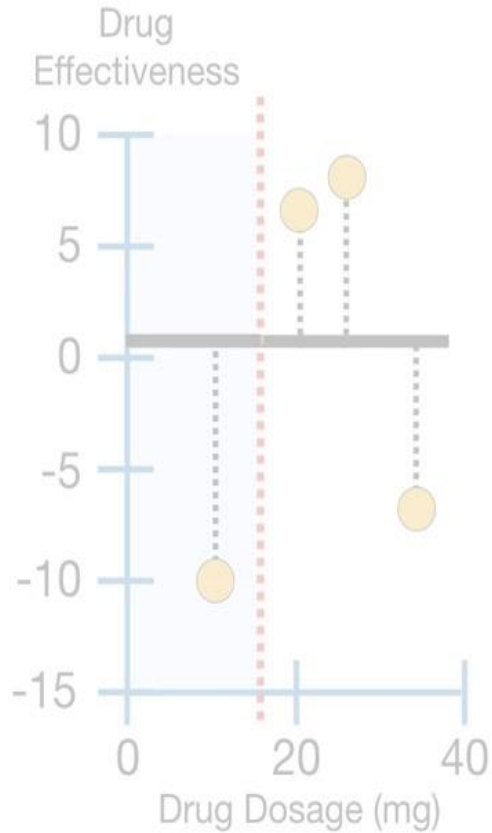


$$\text{Similarity Score} = \frac{(-10.5 + 6.5 + 7.5 + -7.5)^2}{4 + 1} = 3.2$$

...**3.2**, which is **8/10s** of what we got when  $\lambda = 0$ .

# Tree Pruning

Predicted Drug Effectiveness  
0.5

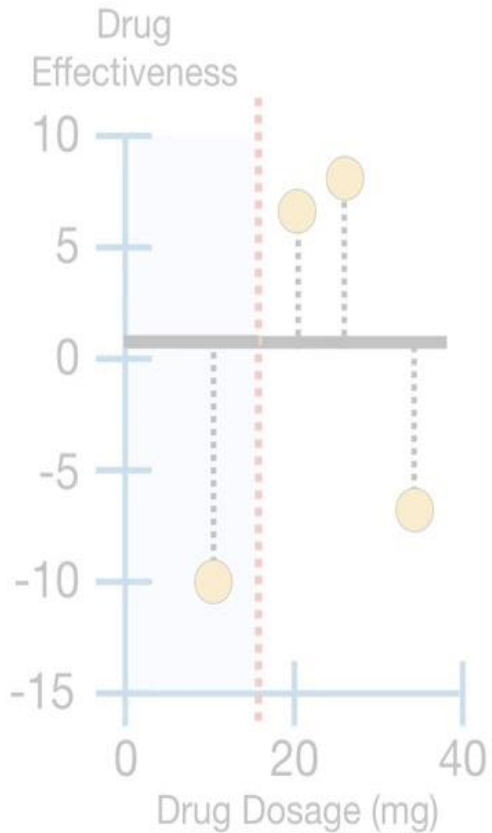


So, one thing we see is that when  $\lambda > 0$ , the **Similarity Scores** are smaller...

# Tree Pruning

Predicted Drug Effectiveness

0.5



-10.5, 6.5, 7.5, -7.5 Similarity = 3.2

-10.5

Similarity = 55.12

6.5, 7.5, -7.5

Similarity = 10.56

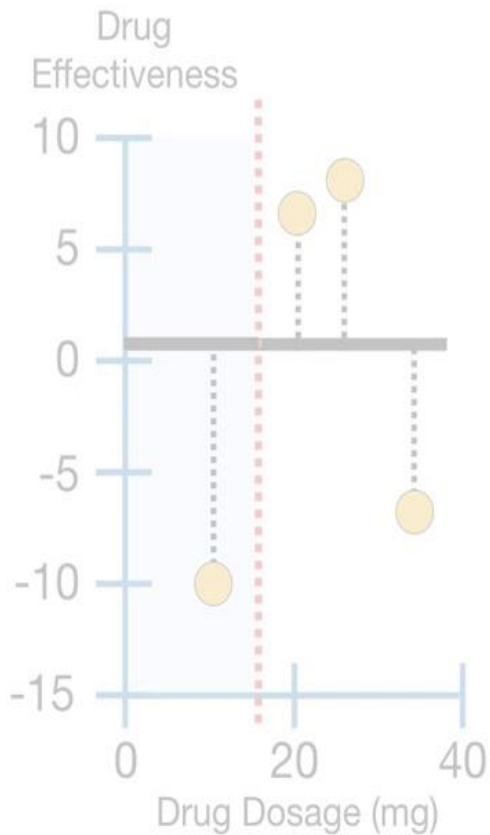
...and the amount of decrease is **inversely proportional** to the number of **Residuals** in the node.



# Tree Pruning

Predicted Drug Effectiveness

0.5



-10.5, 6.5, 7.5, -7.5 Similarity = 3.2

-10.5

Similarity = 55.12

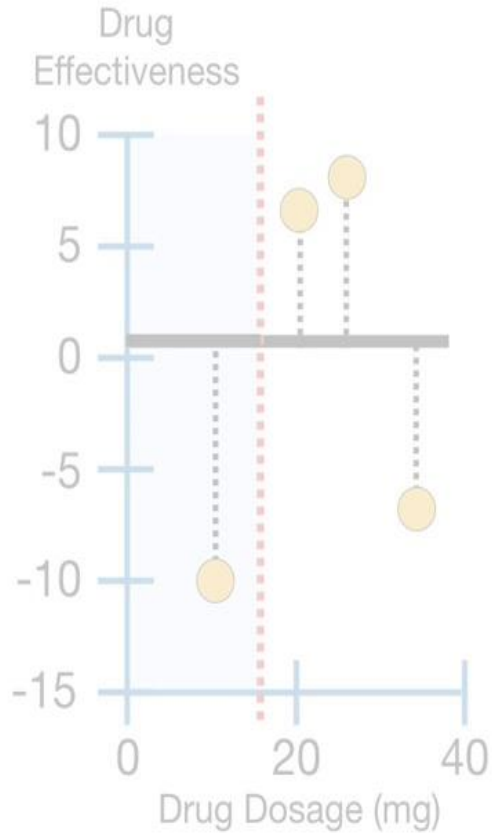
6.5, 7.5, -7.5

Similarity = 10.56

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

# Tree Pruning

Predicted Drug Effectiveness  
0.5



-10.5, 6.5, 7.5, -7.5 Similarity = 3.2

-10.5 Similarity = 55.12

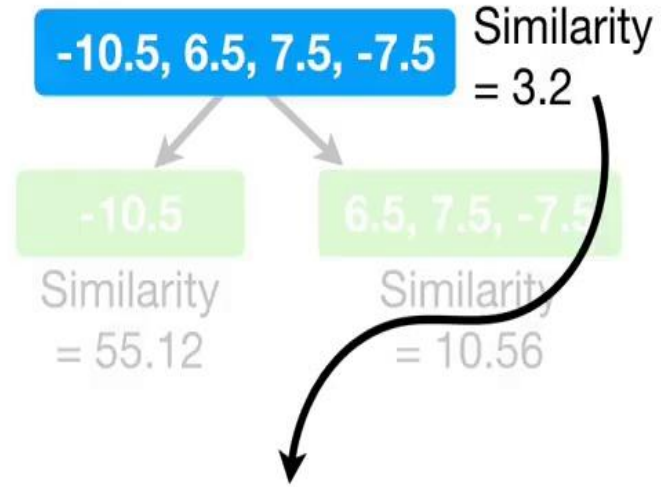
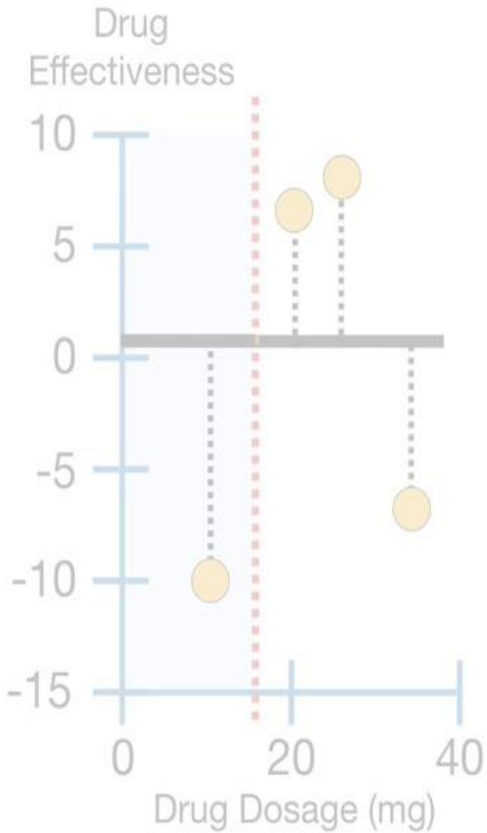
6.5, 7.5, -7.5 Similarity = 10.56

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

# Tree Pruning

Predicted Drug Effectiveness

0.5



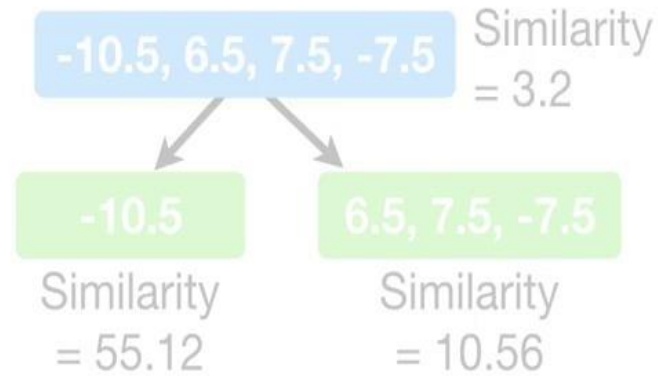
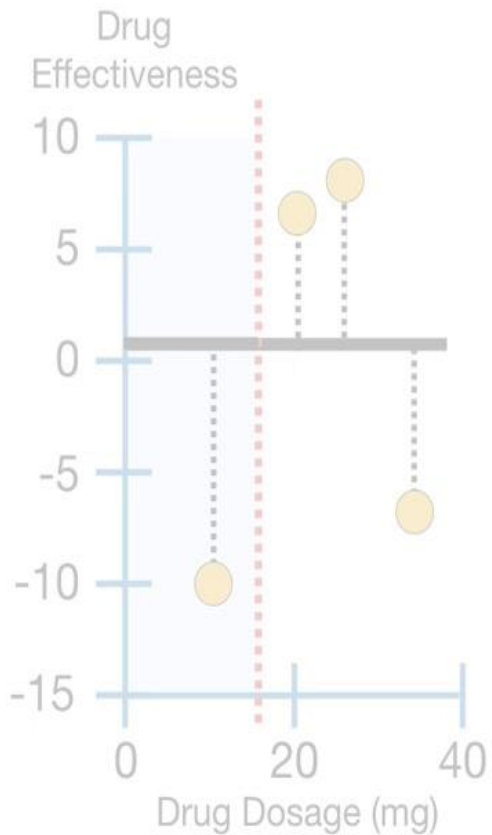
$$\text{Gain} = 55.12 + 10.56 - \text{RootSimilarity}$$

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

# Tree Pruning

Predicted Drug Effectiveness

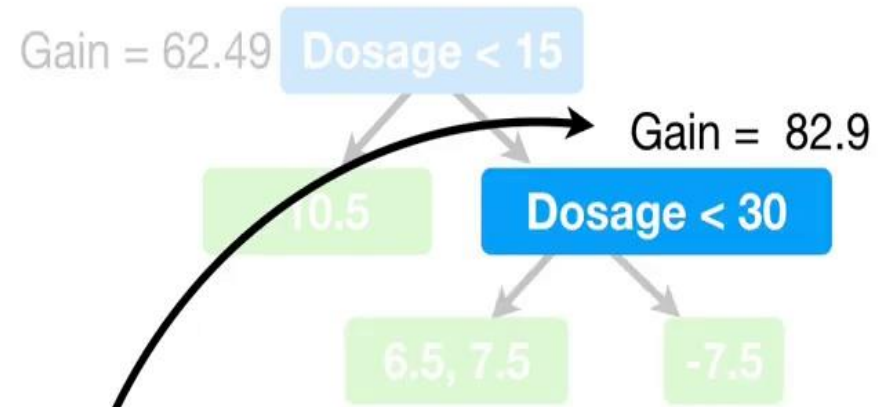
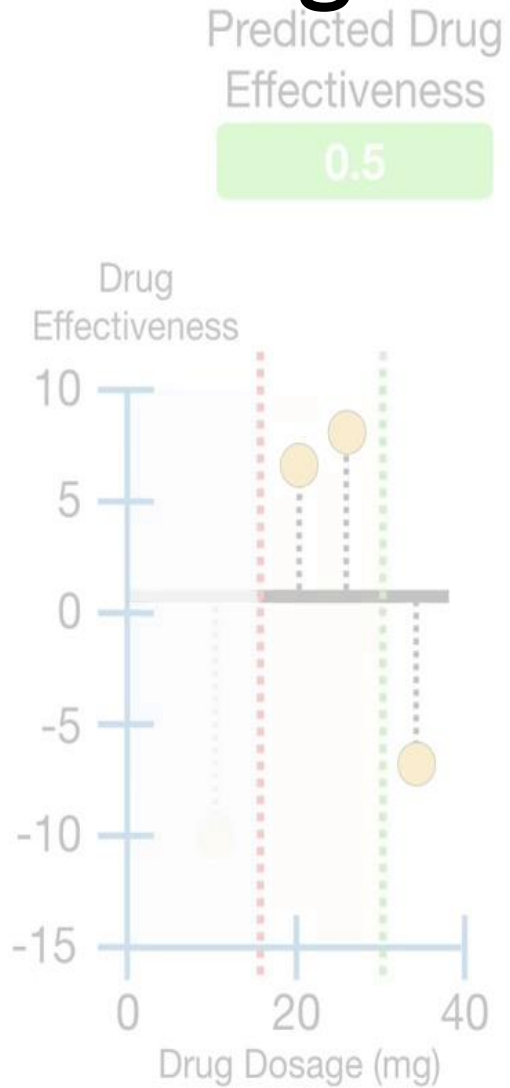
0.5



$$\text{Gain} = 55.12 + 10.56 - 3.2 = 62.48$$

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

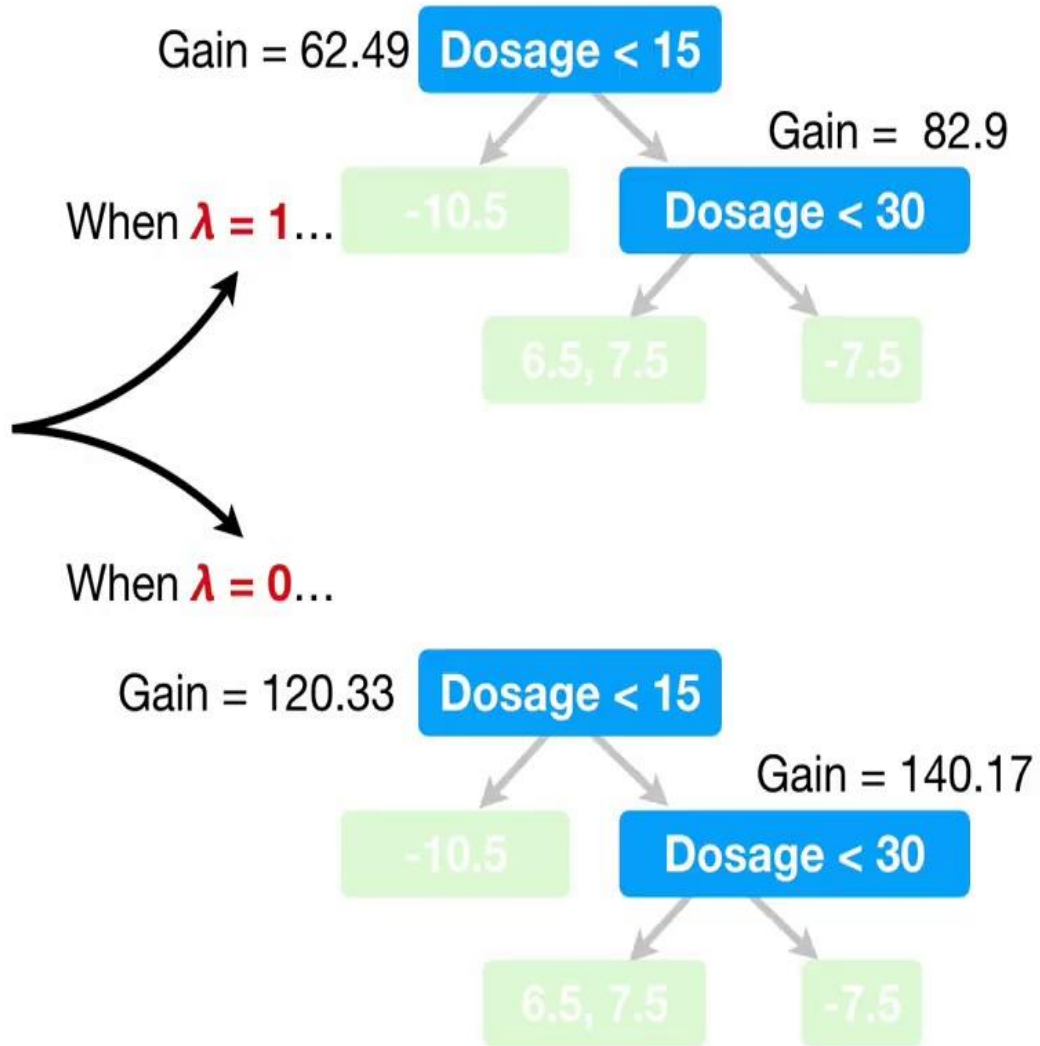
# Tree Pruning



Similarly, when  $\lambda = 1$ , the **Gain** for the next branch is smaller than before.

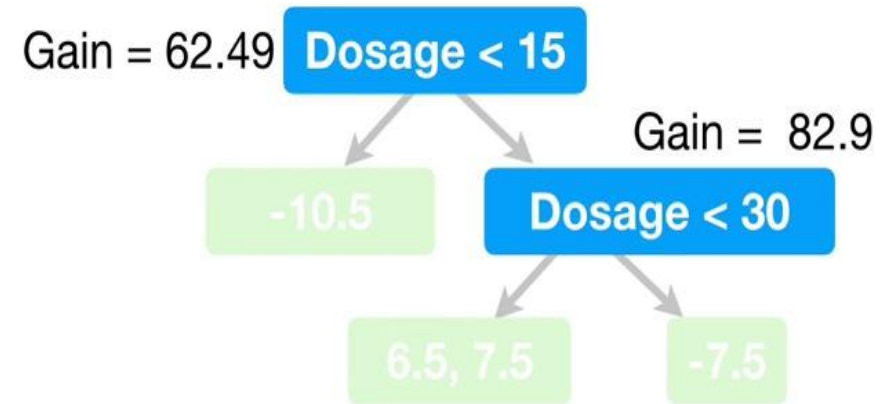
# Tree Pruning

So when  $\lambda > 0$ , it is easier to prune leaves because the values for **Gain** are smaller.



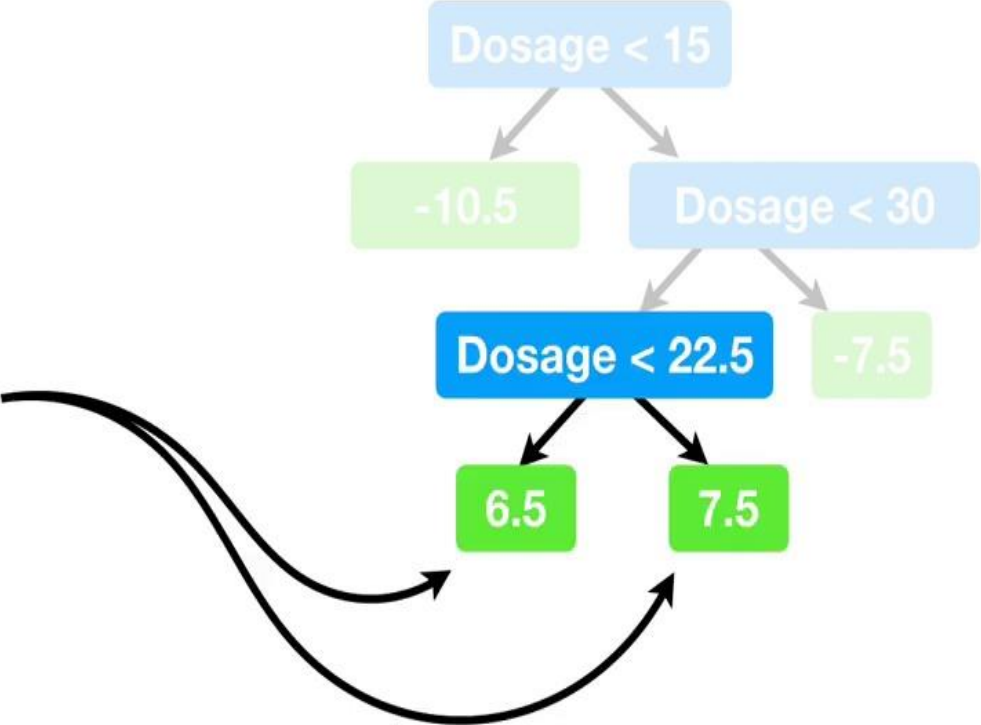
# Tree Pruning

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



# Tree Pruning

For this example, imagine we split this node into two leaves.

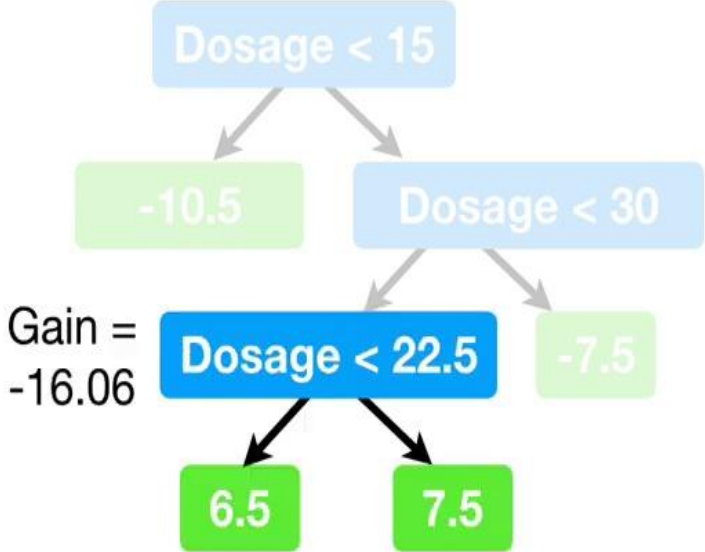




# Tree Pruning

That means the **Gain** is...  
**-16.06.**

$$\text{Gain} = 21.12 + 28.12 - 65.3 = -16.06$$

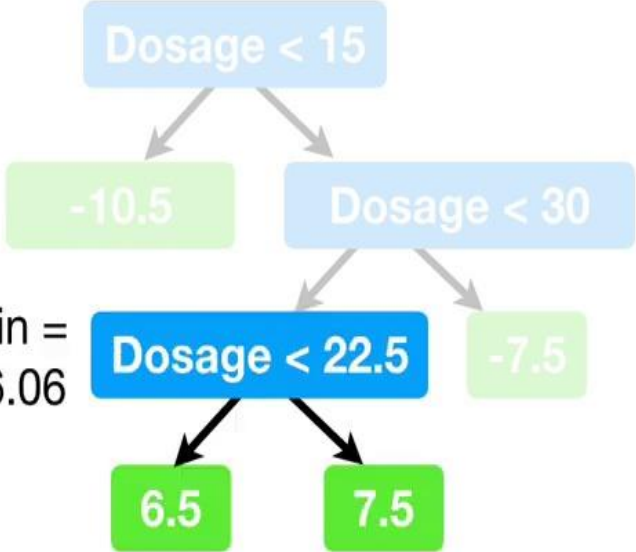


# Tree Pruning

That means the **Gain** is...  
**-16.06.**

Gain =  
-16.06

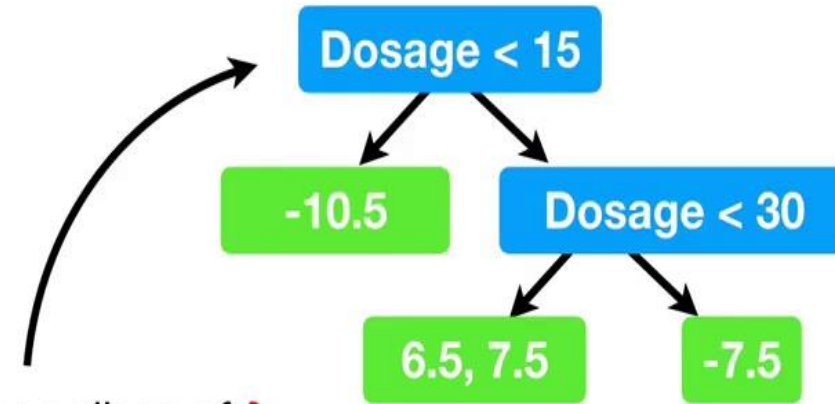
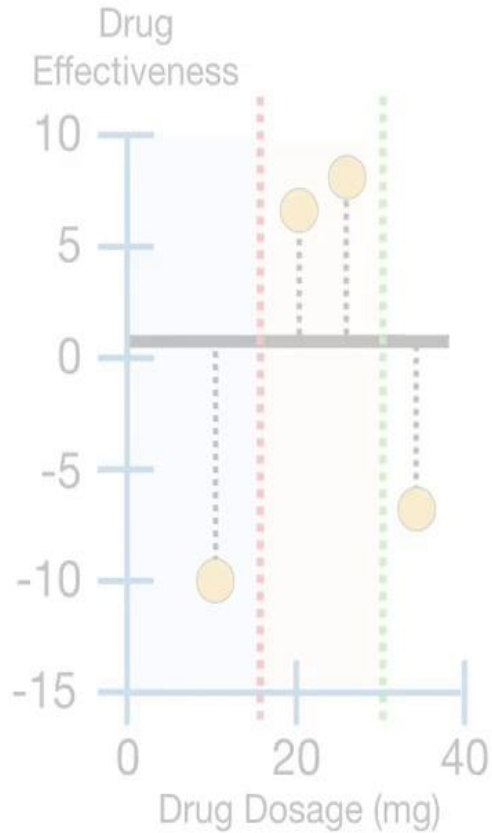
$$\text{Gain} = 21.12 + 28.12 - 65.3 = -16.06$$



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

# Leaves Update

Predicted Drug Effectiveness  
0.5

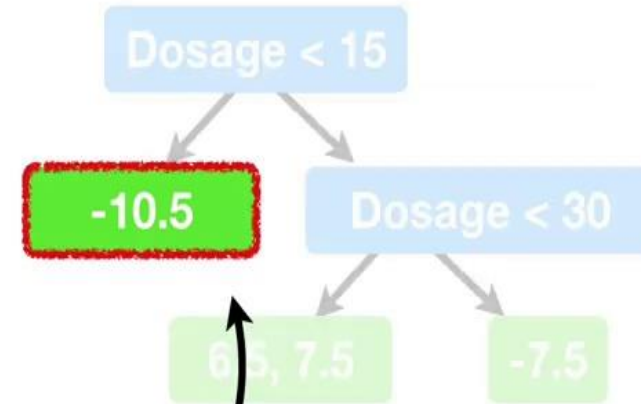
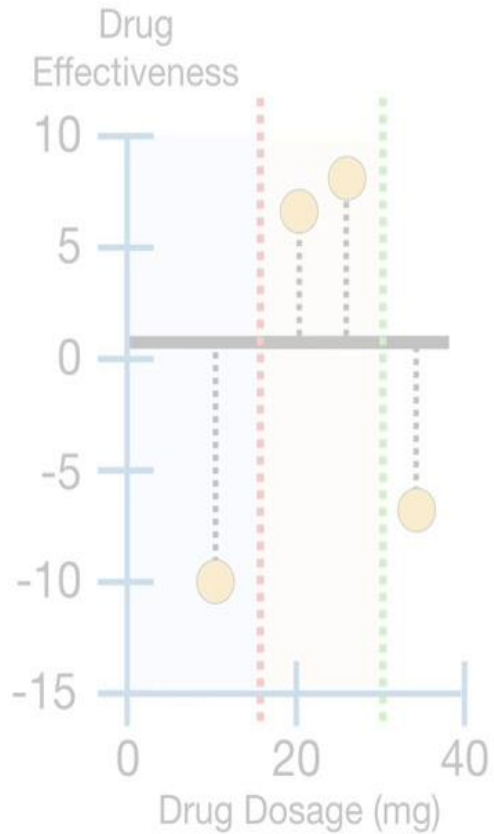


For now, regardless of  $\lambda$  (**lambda**) and  $\gamma$  (**gamma**), let's assume this is the tree we are working with...

# Leaves Update

Predicted Drug Effectiveness

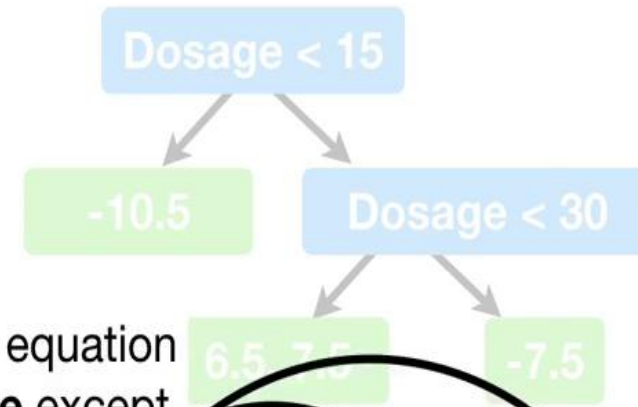
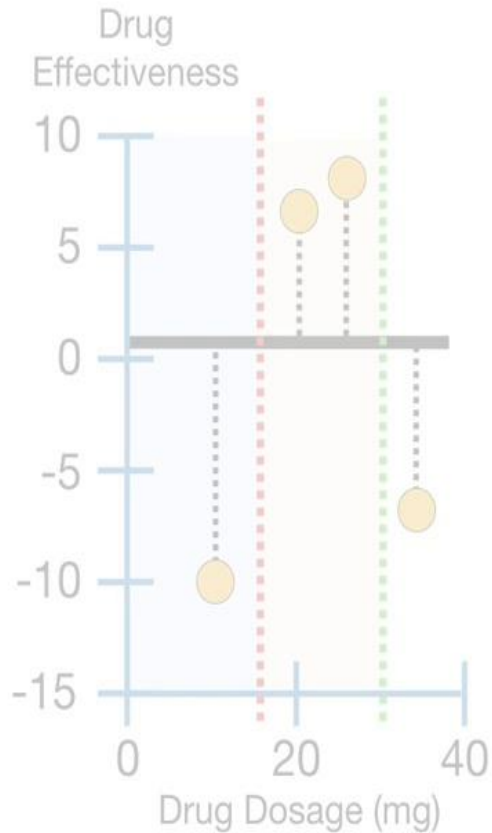
0.5



...and determine the **Output Values** for the leaves.

# Leaves Update

Predicted Drug Effectiveness  
0.5



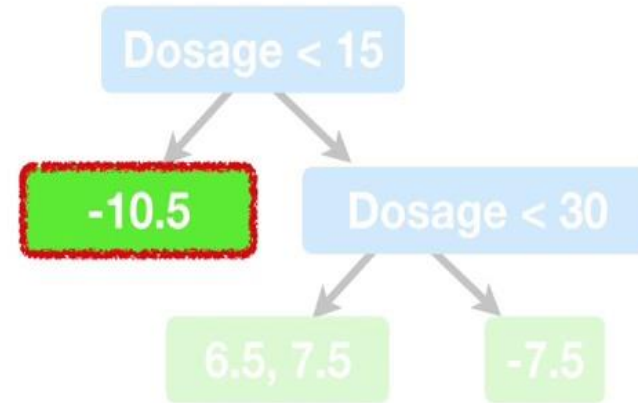
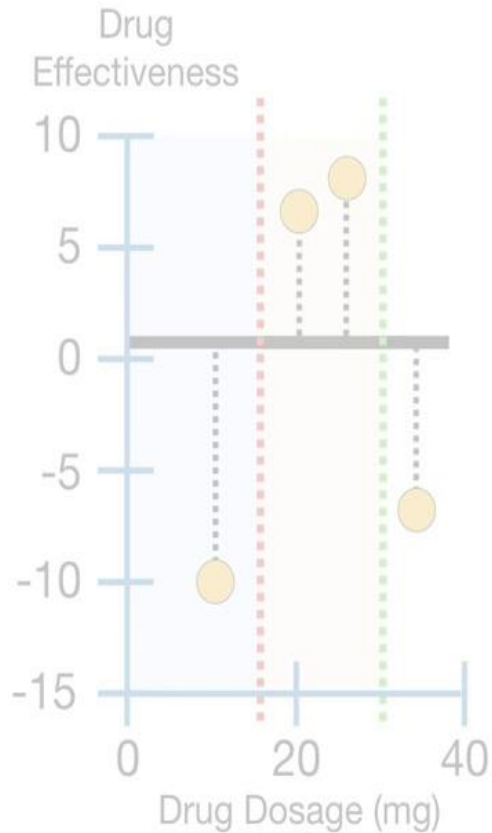
**NOTE:** The **Output Value** equation is like the **Similarity Score** except we do not square the sum of the residuals.

$$\text{Output Value} = \frac{\text{Sum of Residuals}}{\text{Number of Residuals} + \lambda}$$

$$\text{Similarity Score} = \frac{\text{Sum of Residuals, Squared}}{\text{Number of Residuals} + 1}$$

# Leaves Update

Predicted Drug Effectiveness  
0.5



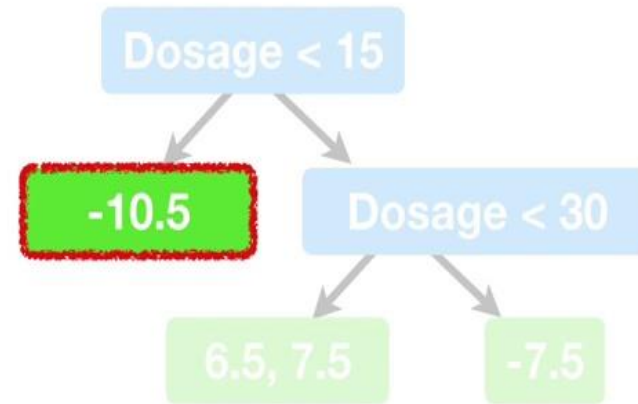
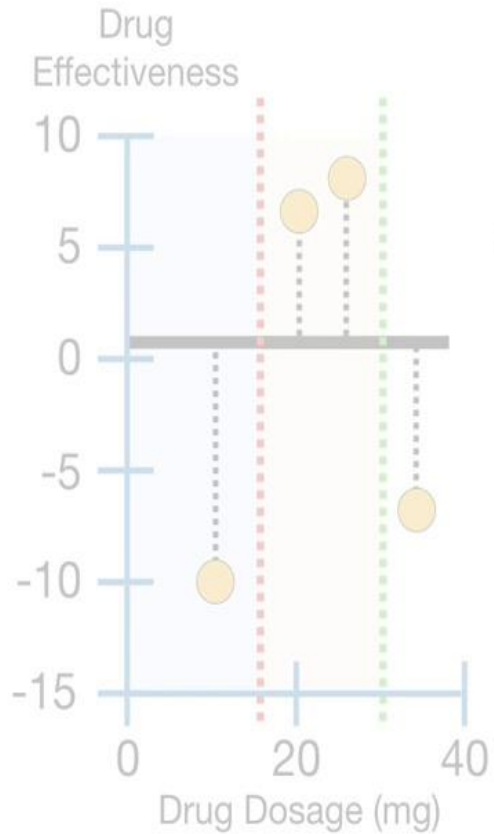
If  $\lambda = 0$ , then there is no **Regularization** and the **Output Value = -10.5**.

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

# Leaves Update

Predicted Drug Effectiveness

0.5

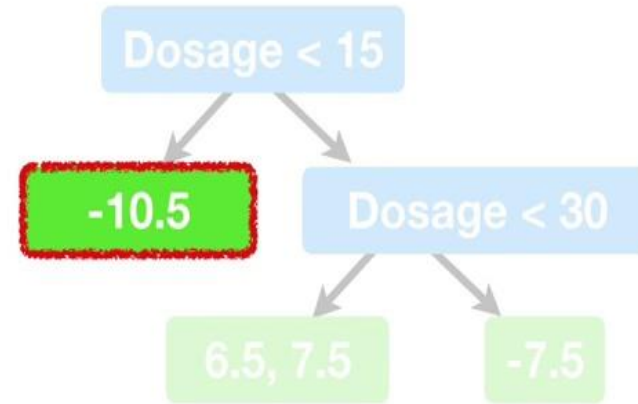
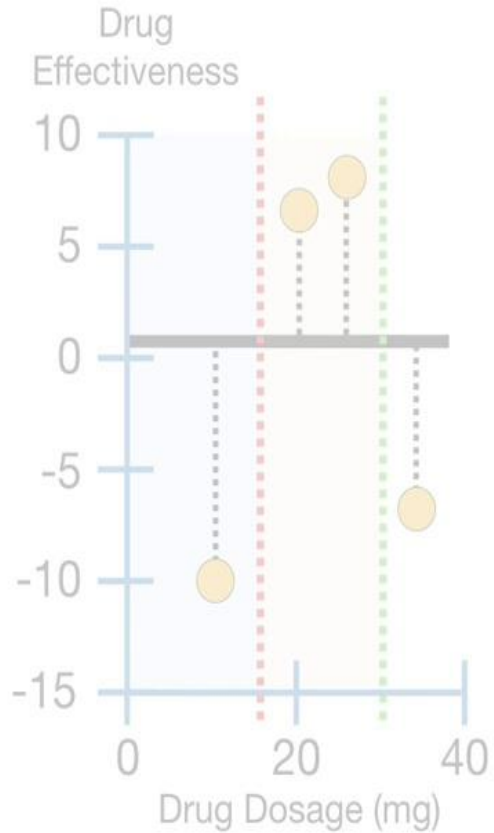


On the other hand, if  $\lambda = 1$ ...

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

# Leaves Update

Predicted Drug Effectiveness  
0.5



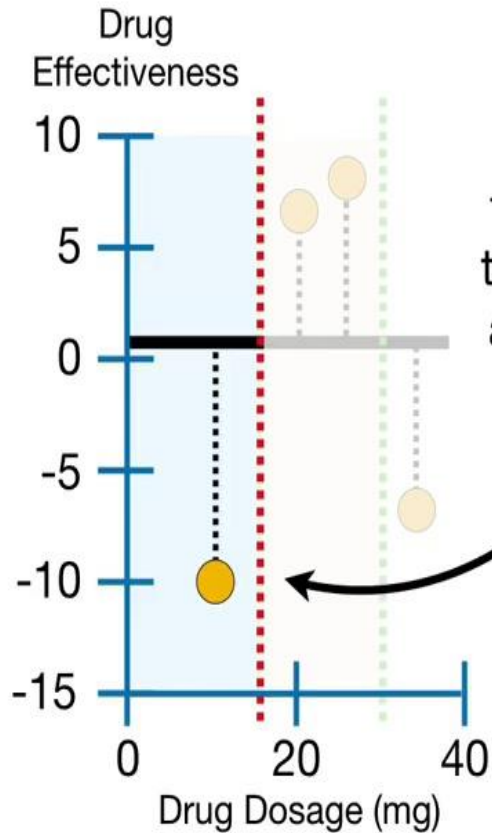
...the **Output Value** = -5.25.

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

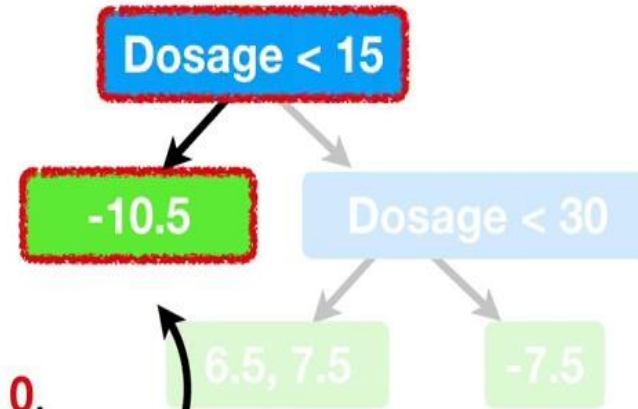


# Leaves Update

Predicted Drug Effectiveness  
0.5



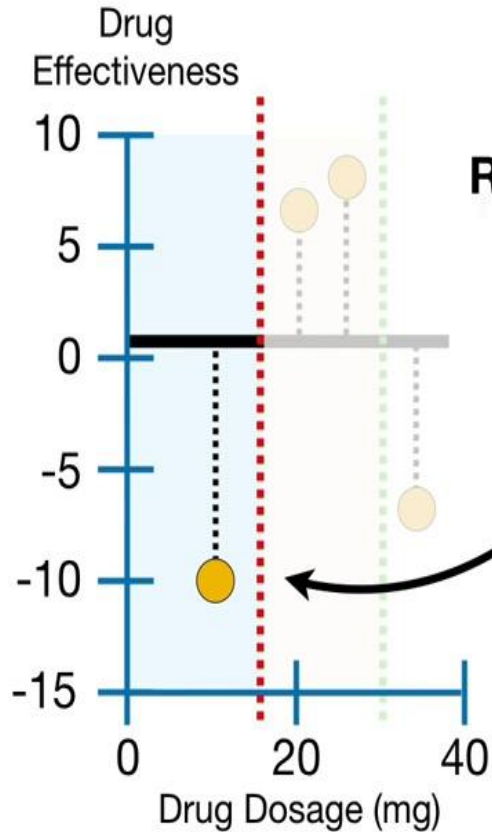
In other words, when  $\lambda > 0$ , then it will reduce the amount that this individual observation adds to the overall prediction.



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

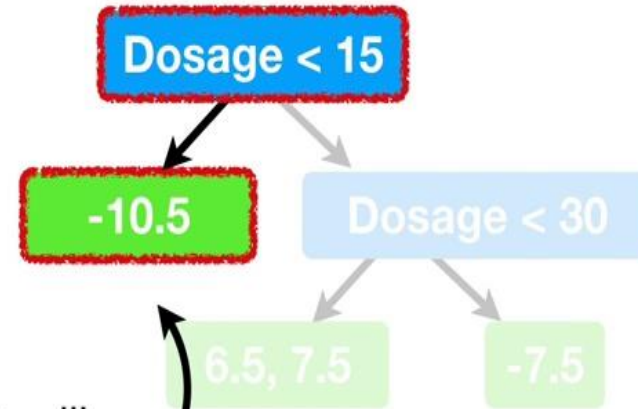
# Leaves Update

Predicted Drug Effectiveness  
0.5



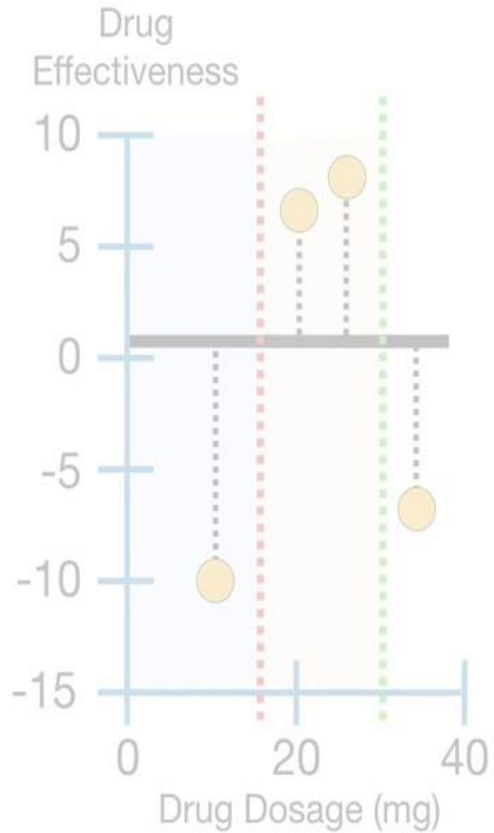
Thus,  $\lambda$  (**lambda**), the **Regularization Parameter**, will reduce the prediction's sensitivity to this individual observation.

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

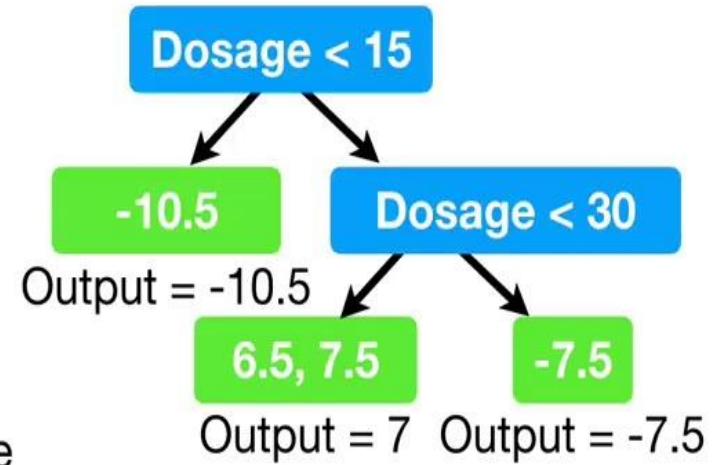


# Leaves Update

Predicted Drug Effectiveness  
0.5

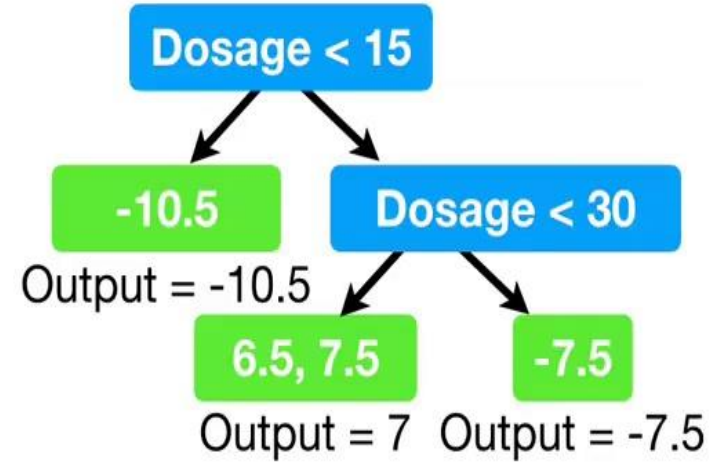
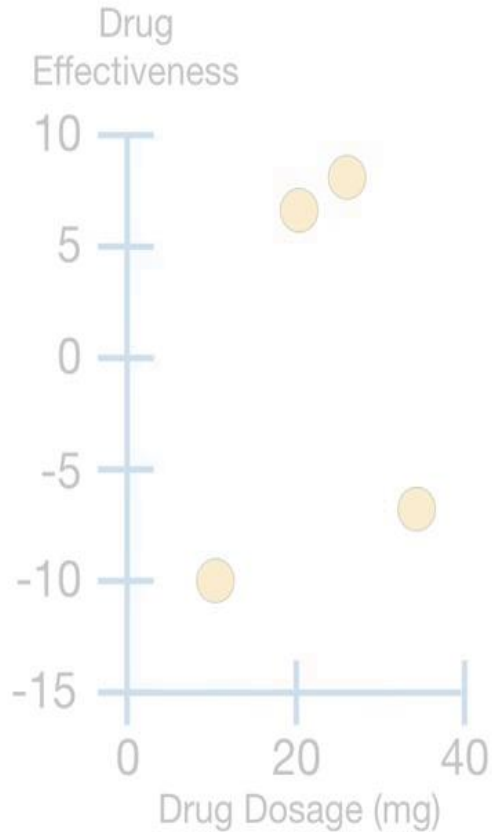


Now, at long last, the first tree is complete!



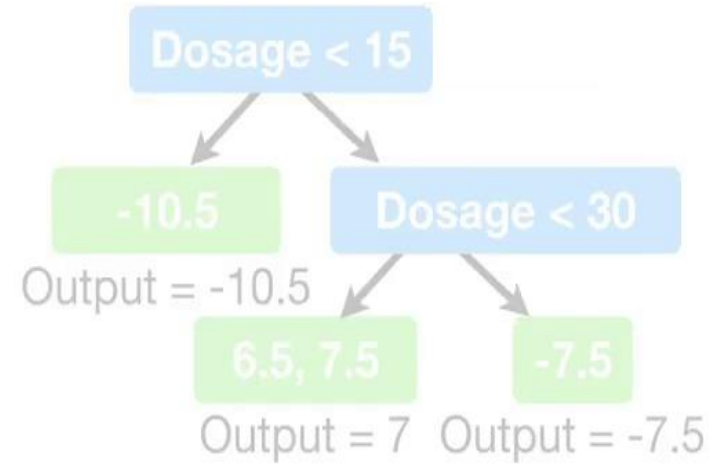
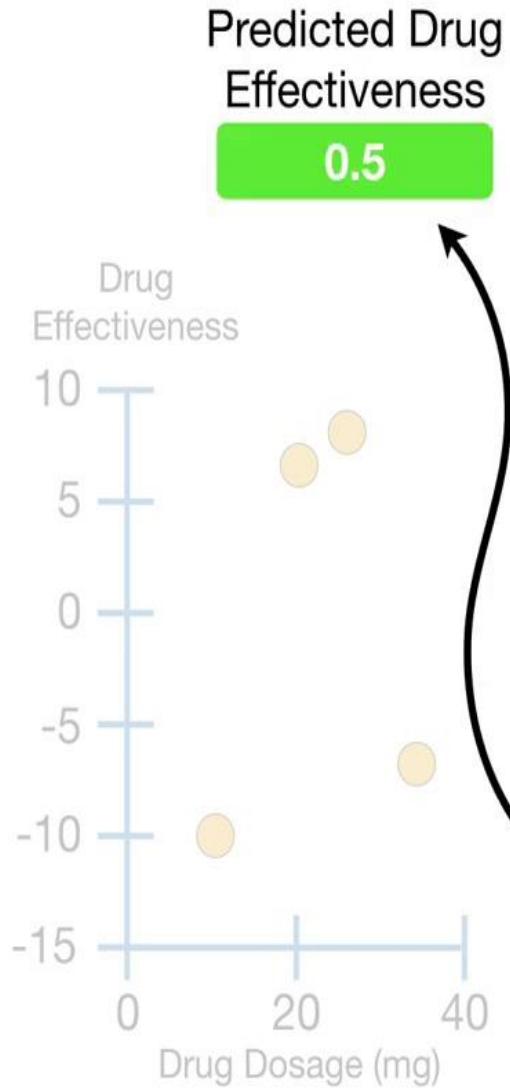
# Tree Prediction

Predicted Drug  
Effectiveness  
0.5



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

# Tree Prediction



And just like unextreme **Gradient Boost, XGBoost** makes new predictions by starting with the initial **Prediction...**

# Tree Prediction

Predicted Drug Effectiveness

0.5

+ Learning Rate X

Dosage < 15

-10.5

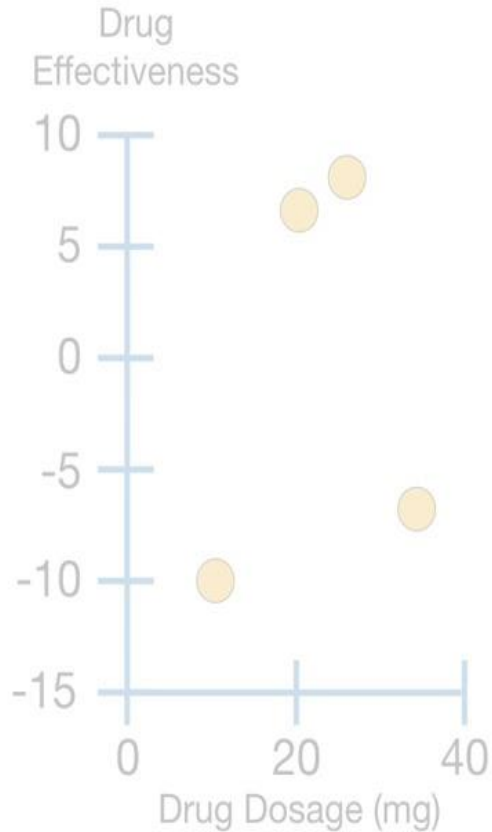
Dosage < 30

Output = -10.5

6.5, 7.5

-7.5

Output = 7    Output = -7.5



...and adding the output of the **Tree**, scaled by a **Learning Rate**.

# Tree Prediction

Predicted Drug Effectiveness

0.5

+ Learning Rate X

Dosage < 15

-10.5

Dosage < 30

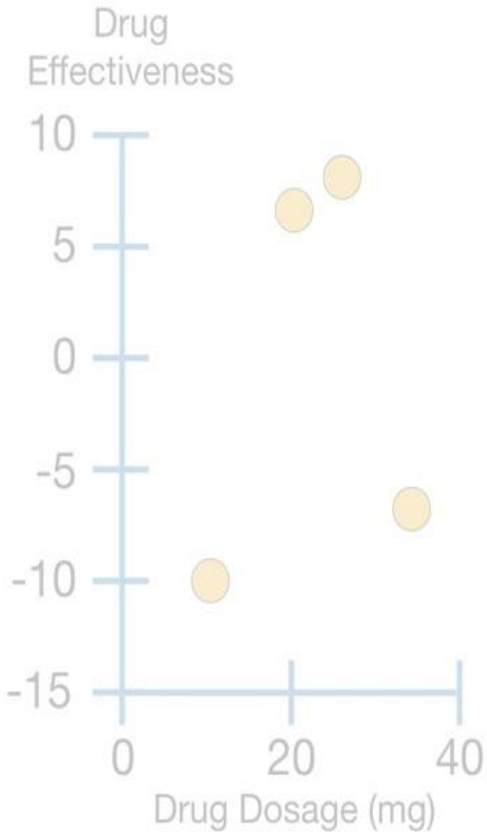
Output = -10.5

6.5, 7.5

-7.5

Output = 7

Output = -7.5



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

# Tree Prediction

Predicted Drug Effectiveness

0.5

+

0.3 X

Dosage < 15

-10.5

Output = -10.5

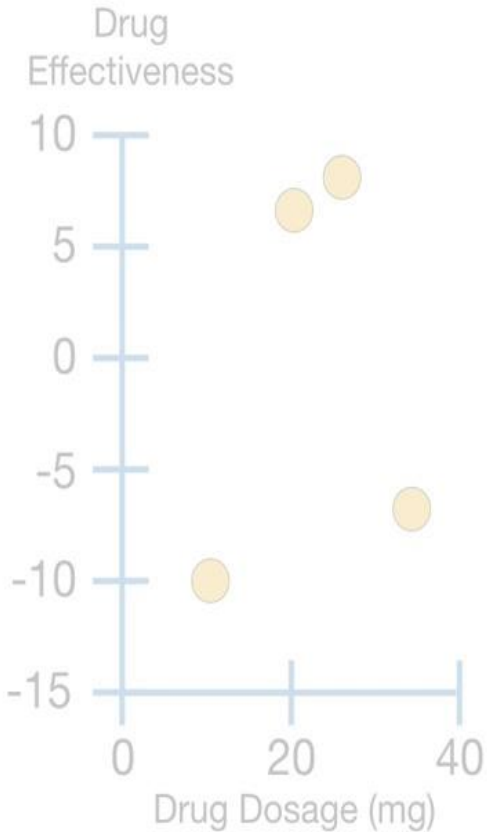
Dosage < 30

6.5, 7.5

Output = 7

-7.5

Output = -7.5



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



# Tree Prediction

Predicted Drug Effectiveness

0.5

+

0.3 X

Dosage < 15

-10.5

Output = -10.5

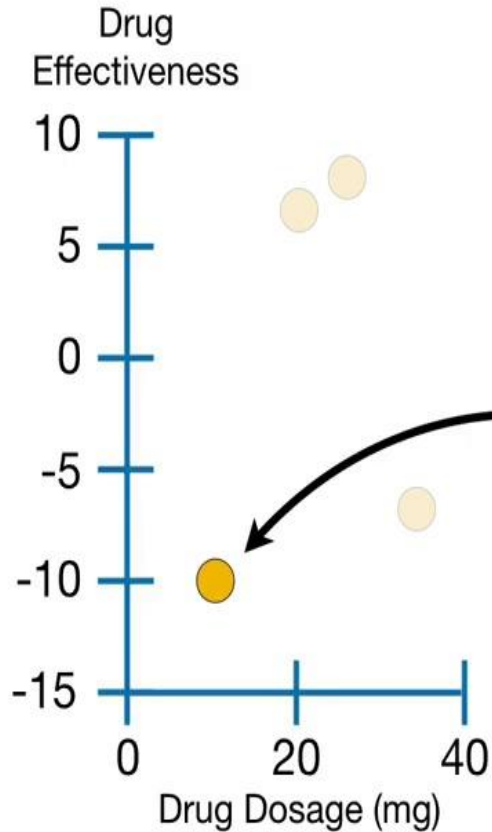
Dosage < 30

6.5, 7.5

Output = 7

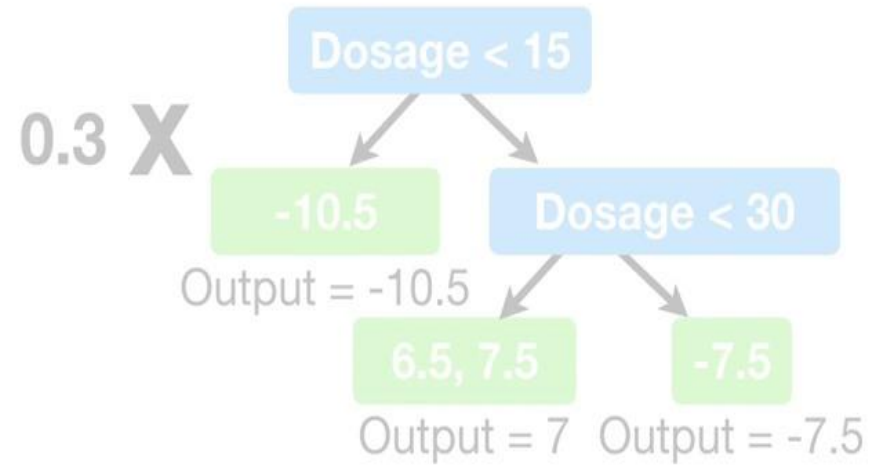
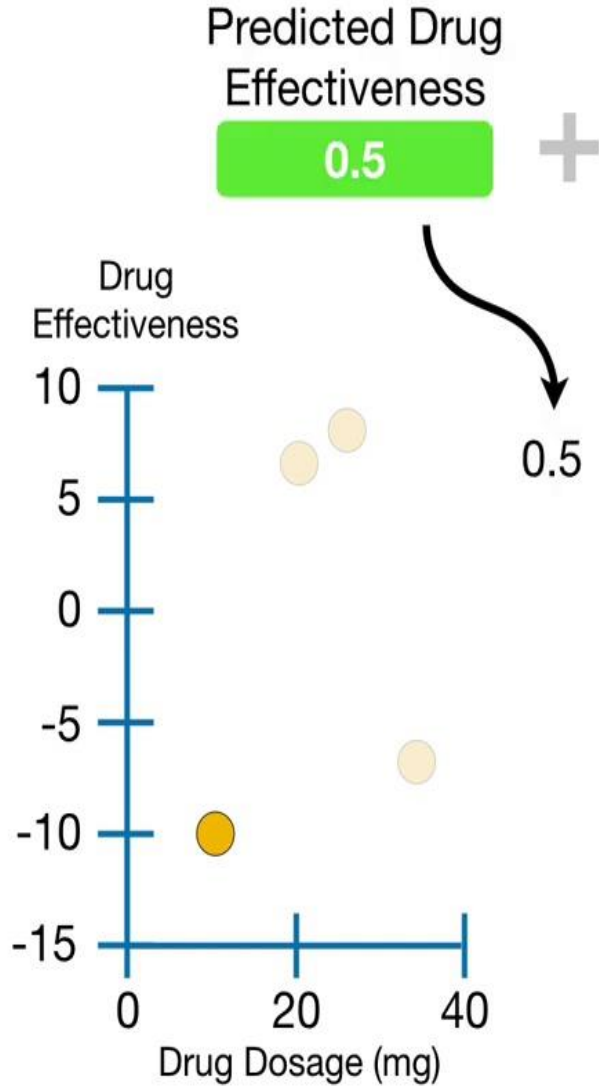
-7.5

Output = -7.5



Thus, the new **Predicted** value for this observation, with **Dosage = 10...**

# Tree Prediction



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

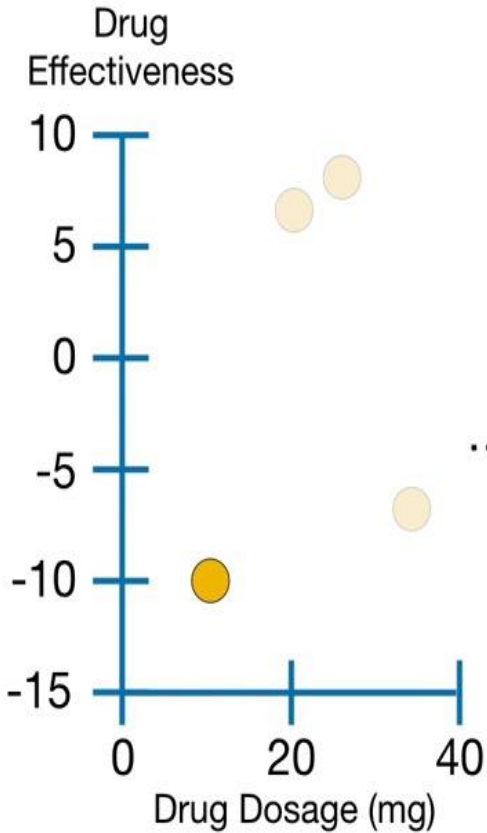
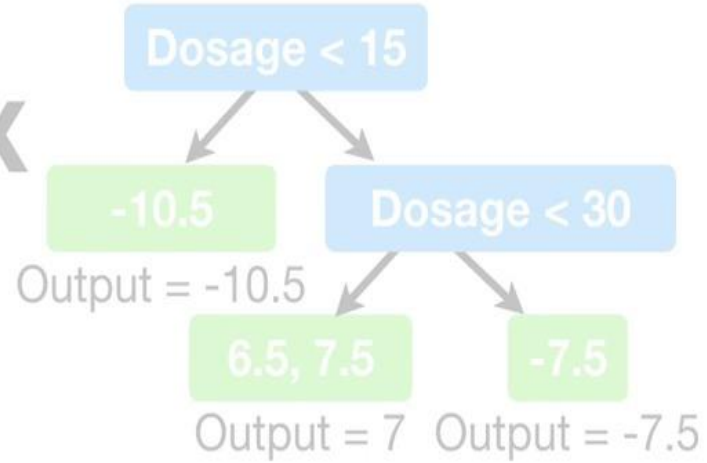
# Tree Prediction

Predicted Drug Effectiveness

0.5

+

0.3 X



0.5 + (0.3

...plus the **Learning Rate,  $\epsilon$  (eta), 0.3...**

# Tree Prediction

Predicted Drug Effectiveness

0.5

+

0.3 X

Dosage < 15

-10.5

Dosage < 30

6.5, 7.5

-7.5

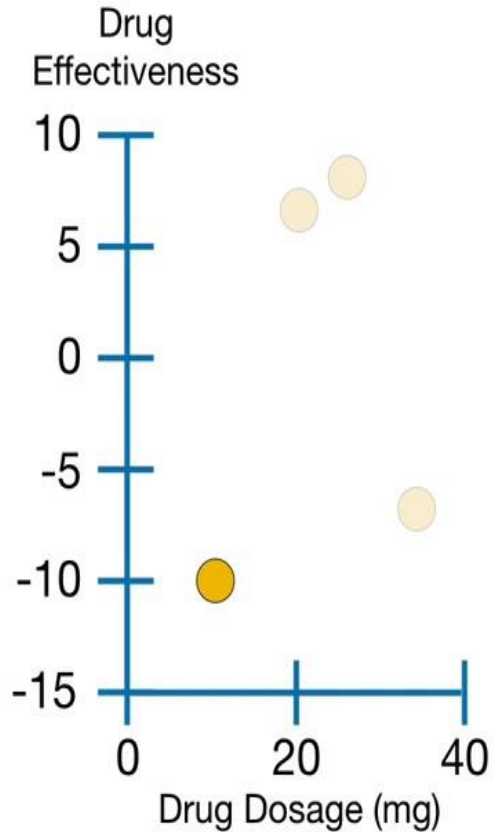
Output = -10.5

Output = 7

Output = -7.5

$0.5 + (0.3 \times -10.5)$

...times the **Output Value, -10.5...**



# Tree Prediction

Predicted Drug Effectiveness

0.5

+

0.3 X

Dosage < 15

-10.5

Output = -10.5

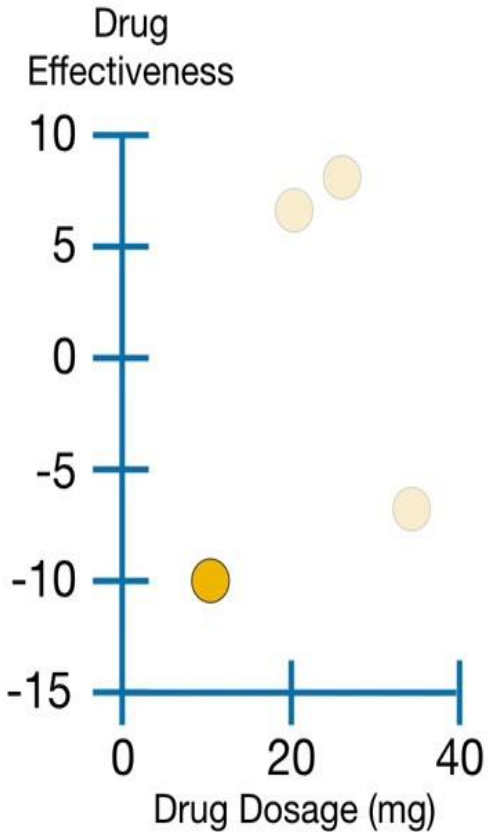
Dosage < 30

6.5, 7.5

Output = 7

-7.5

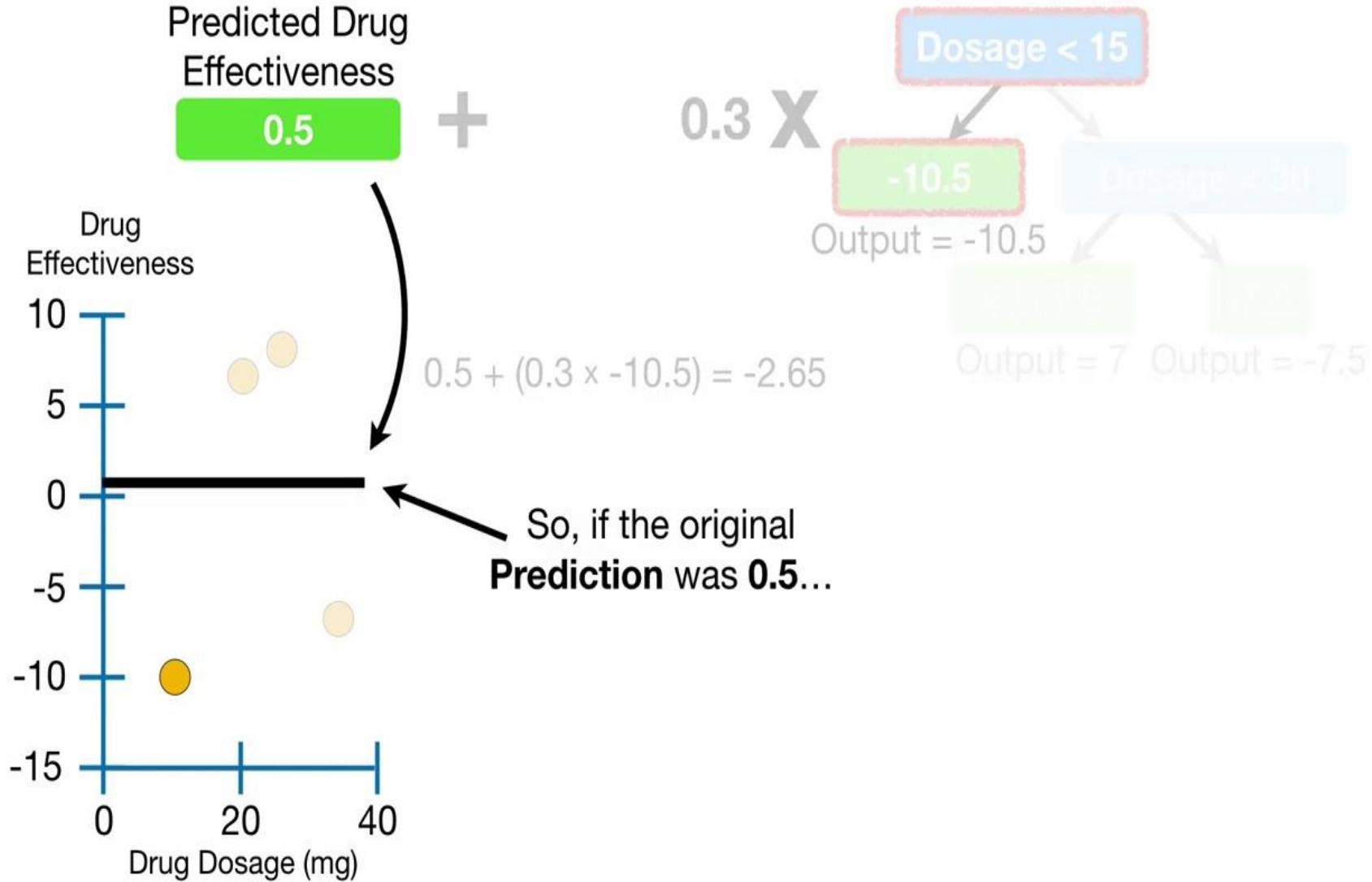
Output = -7.5



$$0.5 + (0.3 \times -10.5) = -2.65$$

...and that gives us **-2.65**.

# Tree Prediction



# Tree Prediction

Predicted Drug Effectiveness

0.5

+

0.3 X

Dosage < 15

-10.5

Output = -10.5

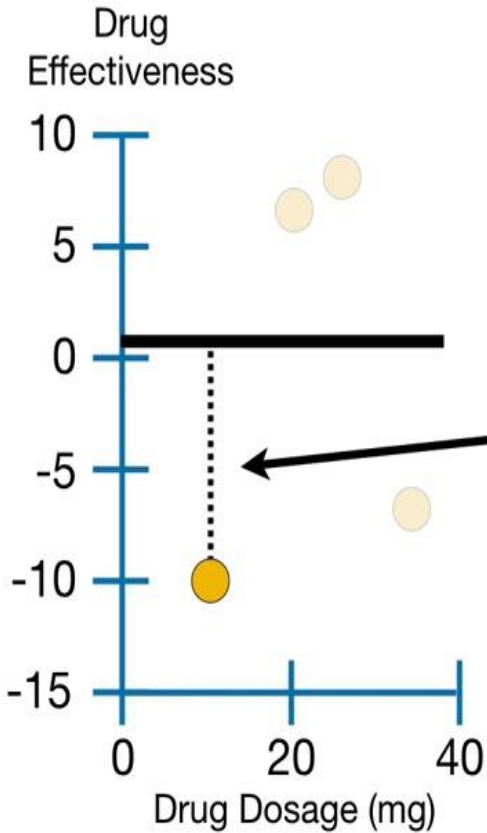
Dosage < 30

6.5, 7.5

Output = 7

-7.5

Output = -7.5



$$0.5 + (0.3 \times -10.5) = -2.65$$

...then this was the original **Residual**.

# Tree Prediction

Predicted Drug Effectiveness

0.5

+

0.3 X

Dosage < 15

-10.5

Output = -10.5

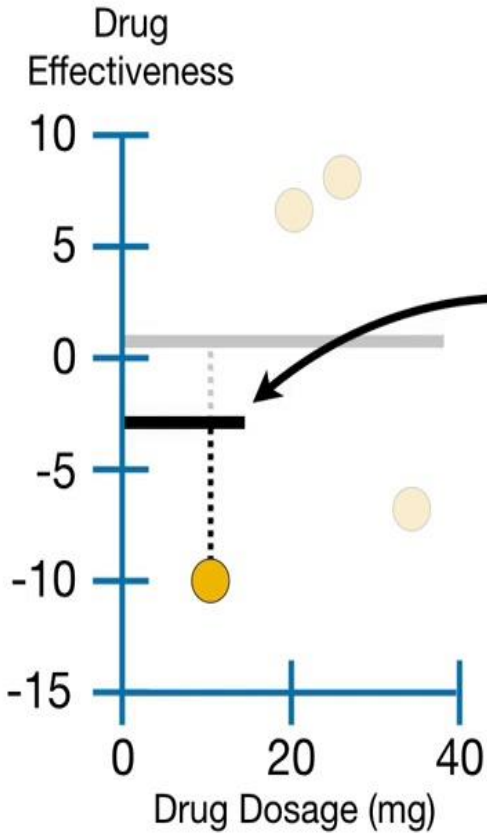
Dosage < 30

6.5, 7.5

Output = 7

-7.5

Output = -7.5



$$0.5 + (0.3 \times -10.5) = -2.65$$

The new prediction is **-2.65...**



# Tree Prediction

Predicted Drug Effectiveness

0.5

+

0.3 X

Dosage < 15

-10.5

Output = -10.5

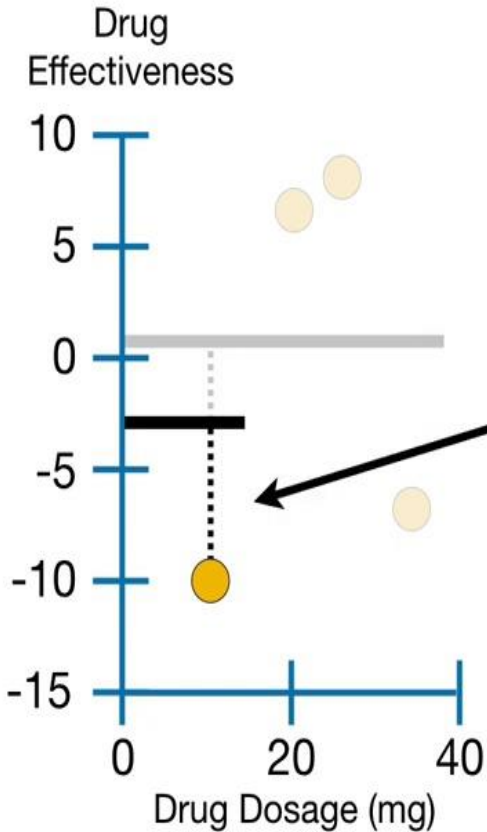
Dosage < 30

6.5, 7.5

Output = 7

-7.5

Output = -7.5



$$0.5 + (0.3 \times -10.5) = -2.65$$

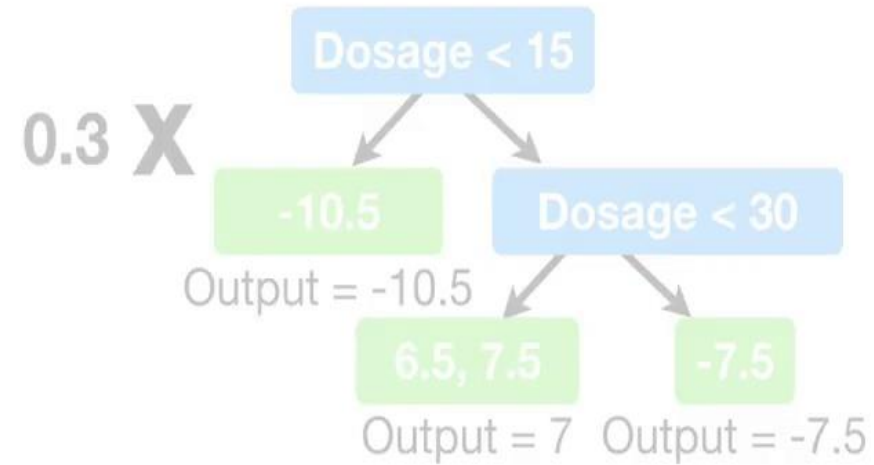
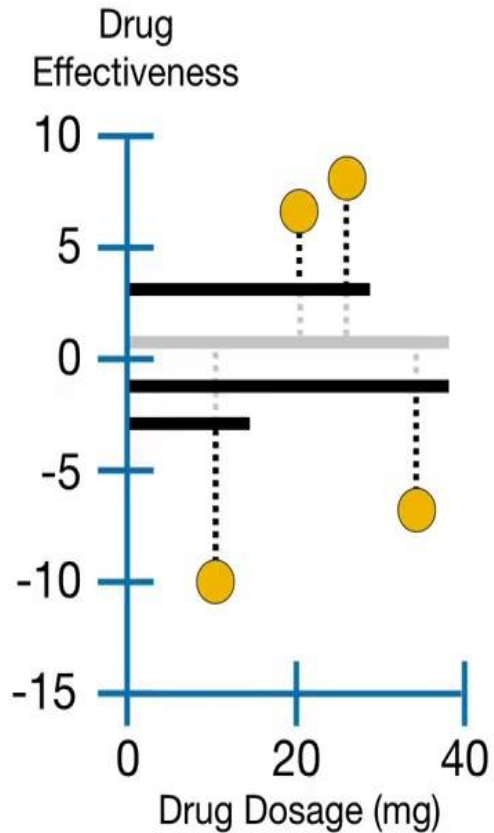
...and we see that the new **Residual** is smaller than before, so we've taken a small step in the right direction.

# Building More Trees

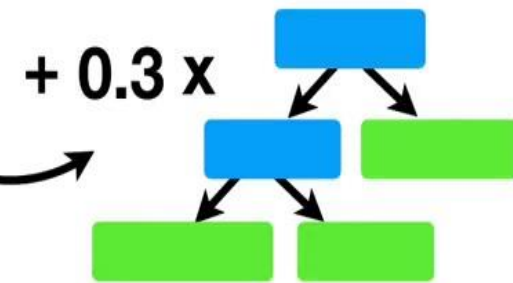
Predicted Drug Effectiveness

0.5

+



Now we build another tree based on the new Residuals...



# Building More Trees

Predicted Drug Effectiveness

0.5

+

0.3 X

Dosage < 15

-10.5

Output = -10.5

Dosage < 30

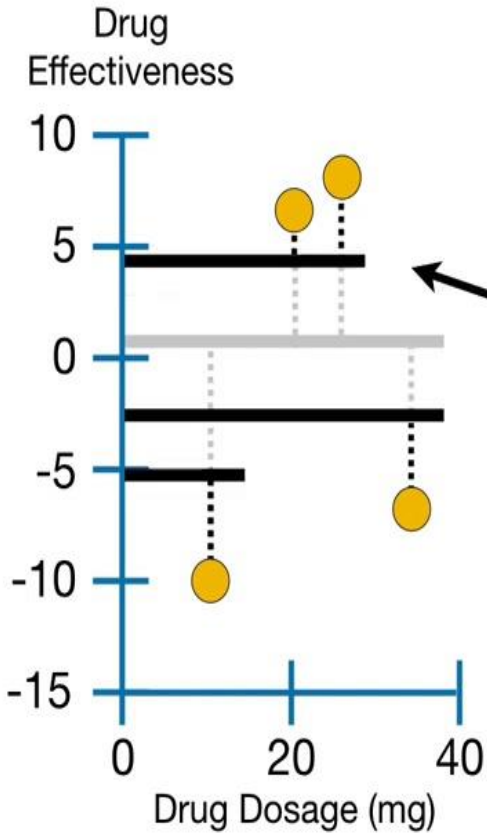
6.5, 7.5

Output = 7

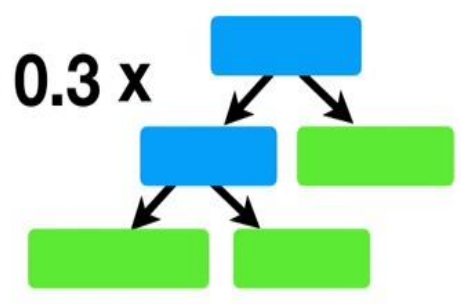
-7.5

Output = -7.5

+ 0.3 X



...and make new predictions that give us even smaller residuals...

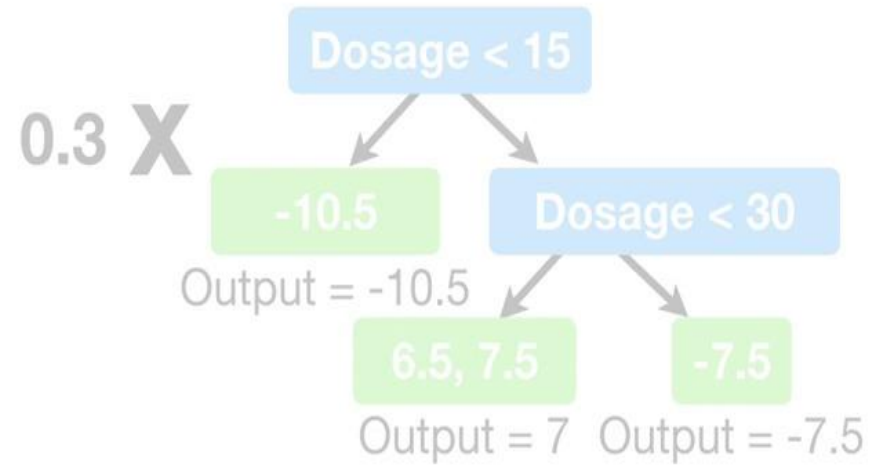
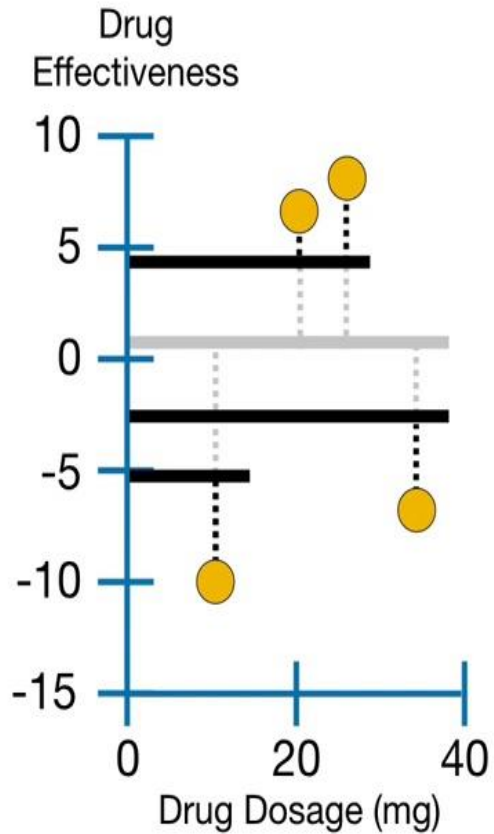


# Building More Trees

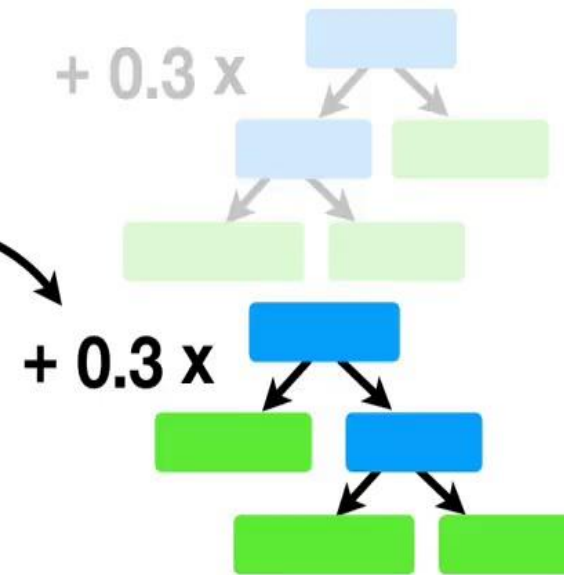
Predicted Drug Effectiveness

0.5

+



...and then build another tree based on the newest Residuals...

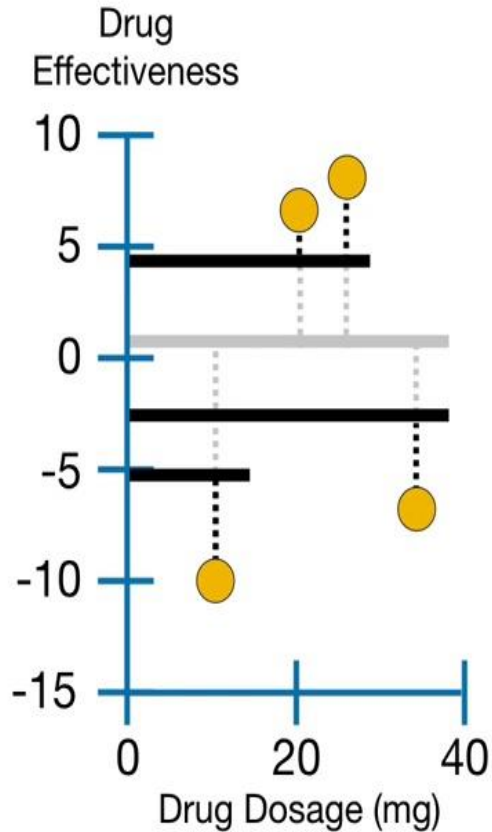


# Building More Trees

Predicted Drug Effectiveness

0.5

+



0.3 X

Dosage < 15

-10.5

Output = -10.5

Dosage < 30

6.5, 7.5

Output = 7

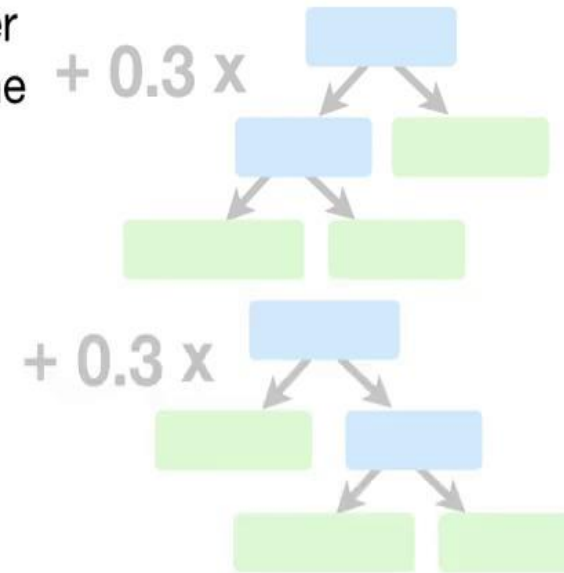
-7.5

Output = -7.5

...and we keep building trees until the **Residuals** are super small, or we have reached the maximum number.

+ 0.3 x

+ 0.3 x



# Building More Trees

Predicted Drug Effectiveness

0.5

+

0.3 X

Dosage < 15

-10.5

Output = -10.5

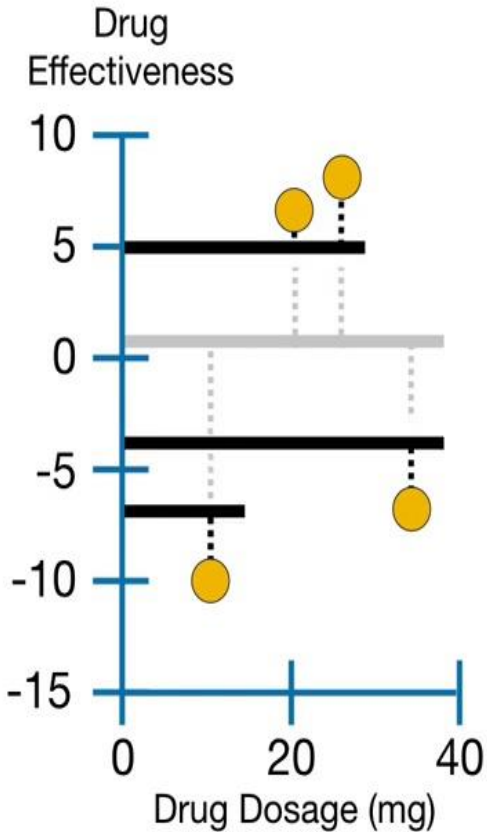
Dosage < 30

6.5, 7.5

Output = 7

-7.5

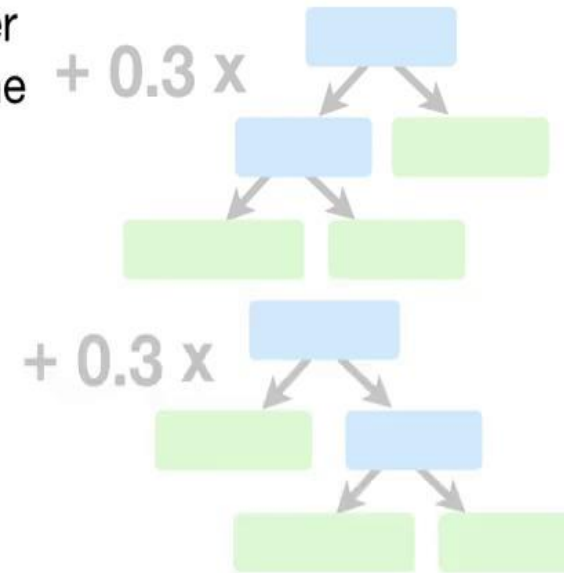
Output = -7.5



...and we keep building trees until the **Residuals** are super small, or we have reached the maximum number.

+ 0.3 x

+ 0.3 x



# Building More Trees

Predicted Drug Effectiveness

0.5

+

0.3 X

Dosage < 15

-10.5

Output = -10.5

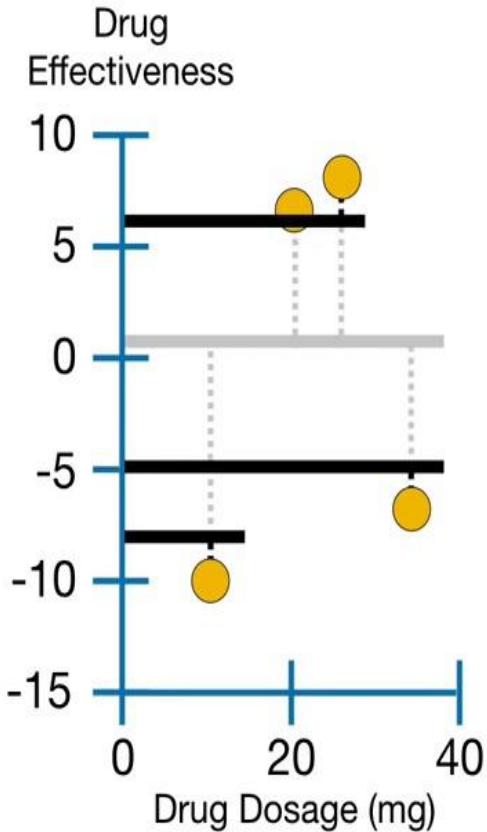
Dosage < 30

6.5, 7.5

Output = 7

-7.5

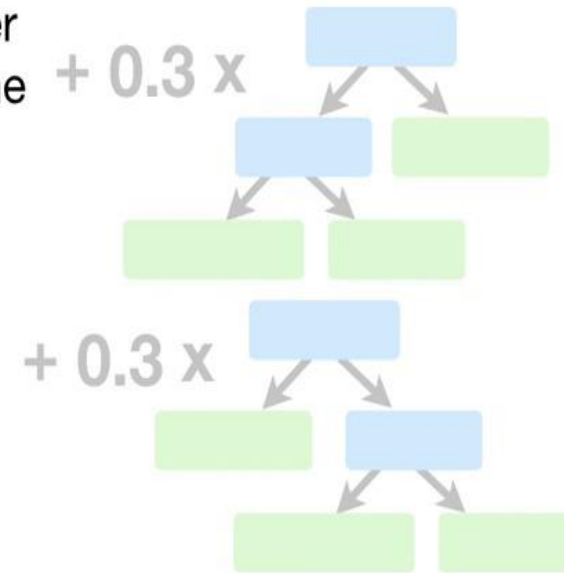
Output = -7.5



...and we keep building trees until the **Residuals** are super small, or we have reached the maximum number.

+ 0.3 x

+ 0.3 x

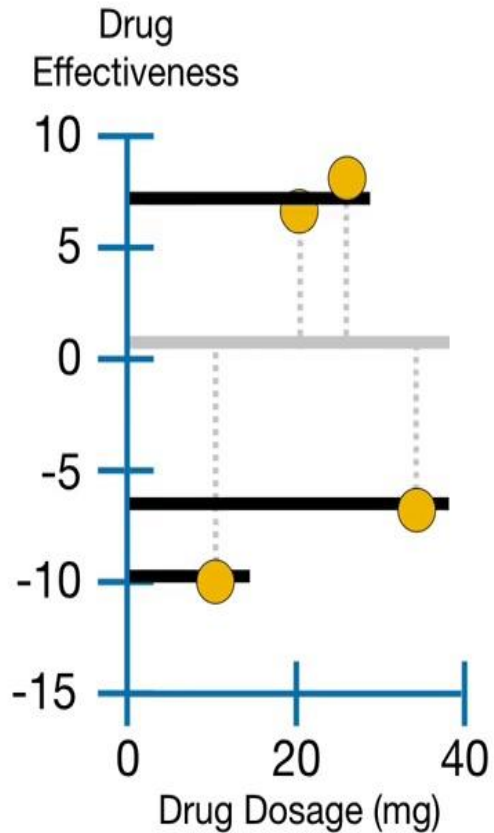


# Building More Trees

Predicted Drug Effectiveness

0.5

+



0.3 X

Dosage < 15

-10.5

Output = -10.5

Dosage < 30

6.5, 7.5

Output = 7

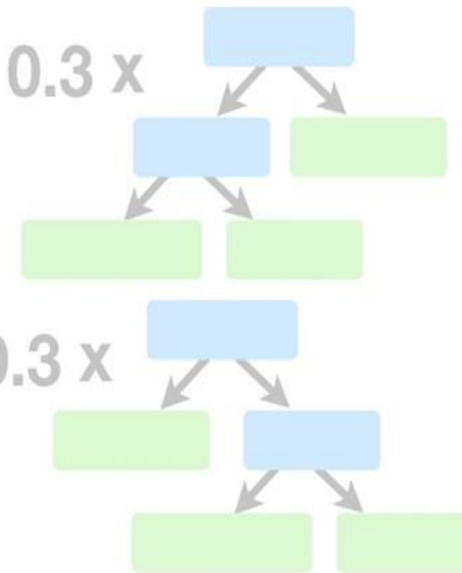
-7.5

Output = -7.5

...and we keep building trees until the **Residuals** are super small, or we have reached the maximum number.

+ 0.3 x

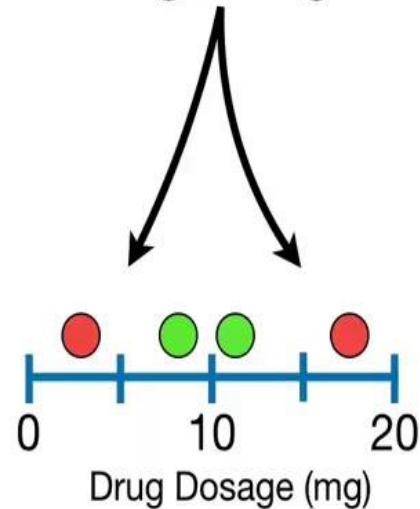
+ 0.3 x





# XGBoost for Classification

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

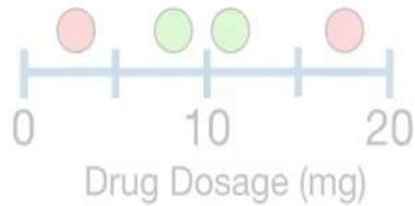


# XGBoost for Classification

Predicted Drug  
Effectiveness

0.5

The very first step in fitting  
**XGBoost** to the **Training  
Data** is to make an initial  
prediction.

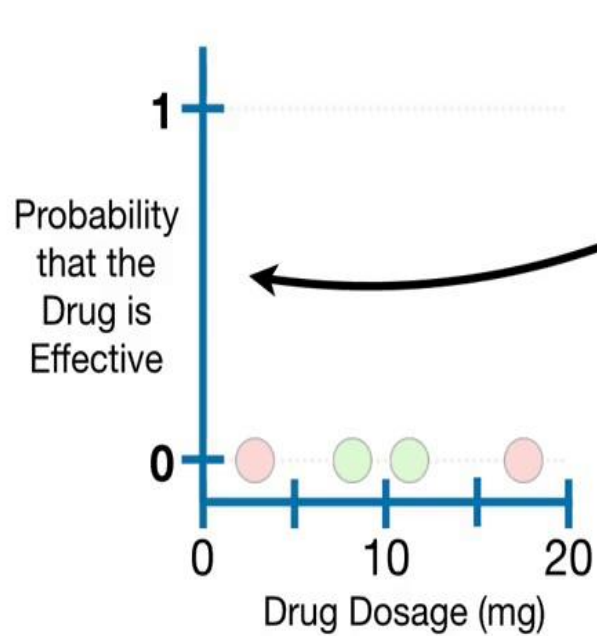


# XGBoost for Classification

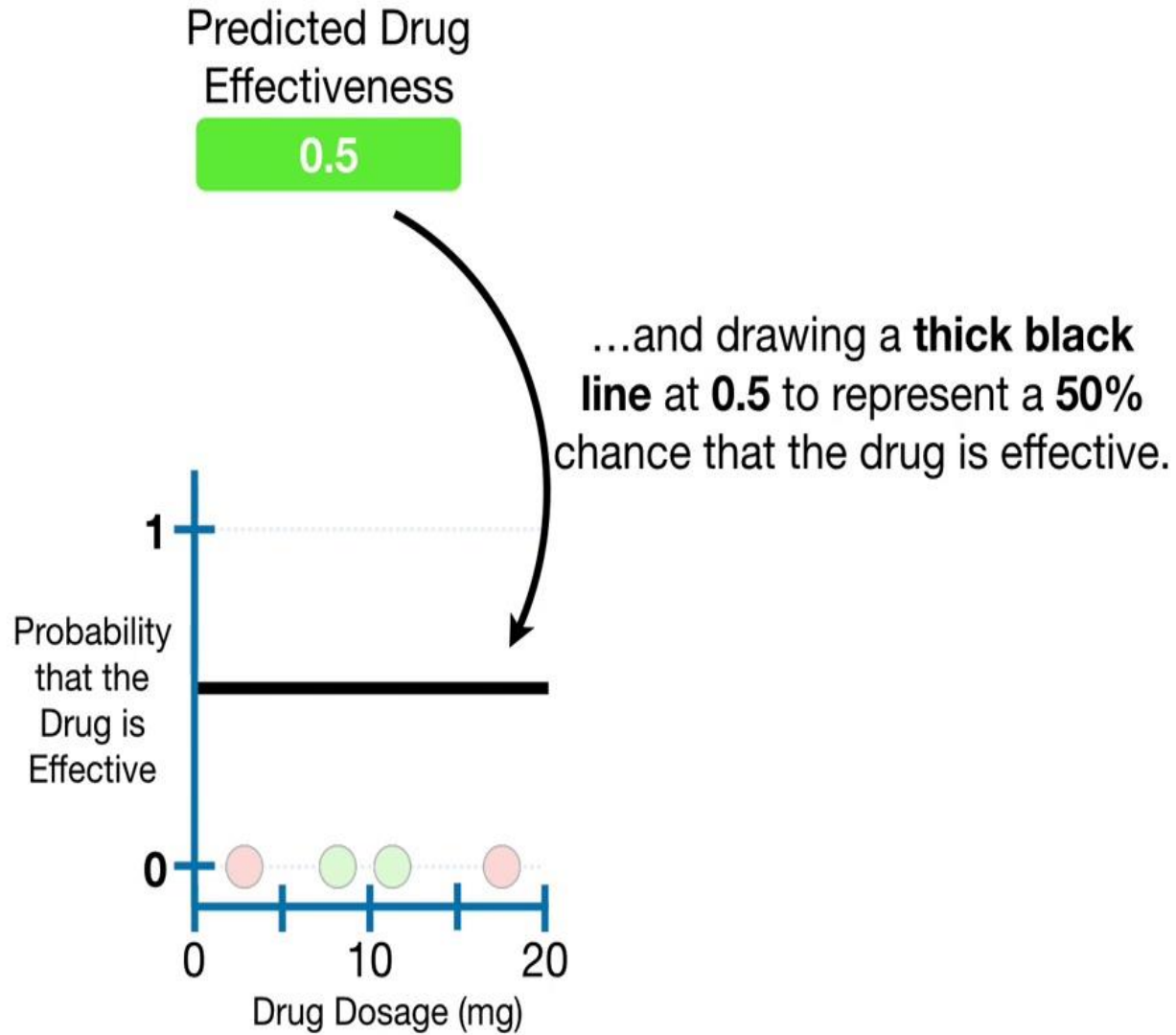
Predicted Drug  
Effectiveness

0.5

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



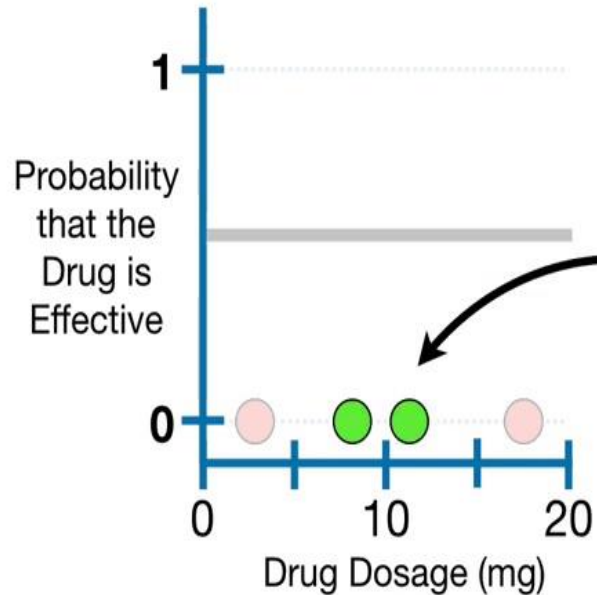
# XGBoost for Classification



# XGBoost for Classification

Predicted Drug  
Effectiveness  
0.5

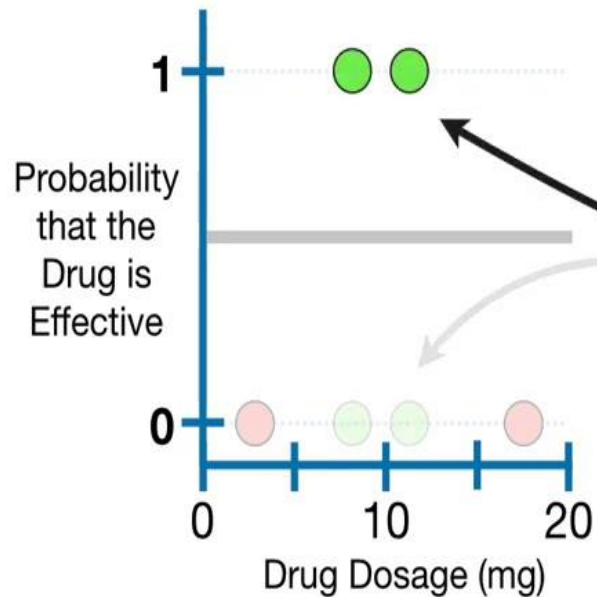
Since these two **Green Dots** represent effective dosages, we will move them to the top of the graph, where the probability that the drug is effective is **1**.



# XGBoost for Classification

Predicted Drug  
Effectiveness  
0.5

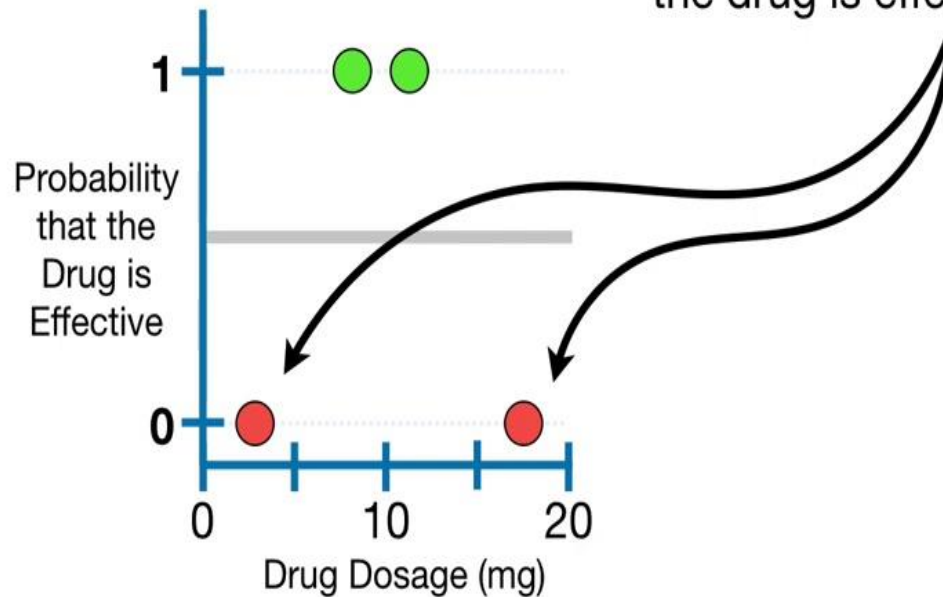
Since these two **Green Dots** represent effective dosages, we will move them to the top of the graph, where the probability that the drug is effective is 1.



# XGBoost for Classification

Predicted Drug  
Effectiveness  
0.5

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**.

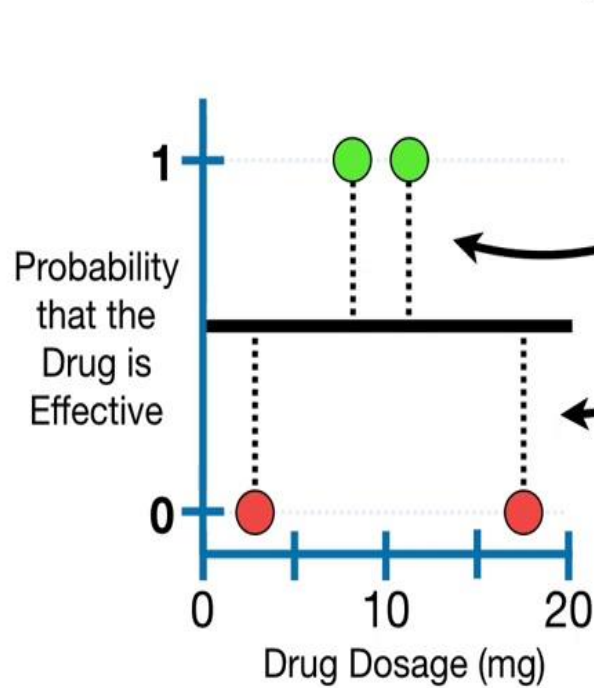


# XGBoost for Classification

Predicted Drug  
Effectiveness

0.5

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

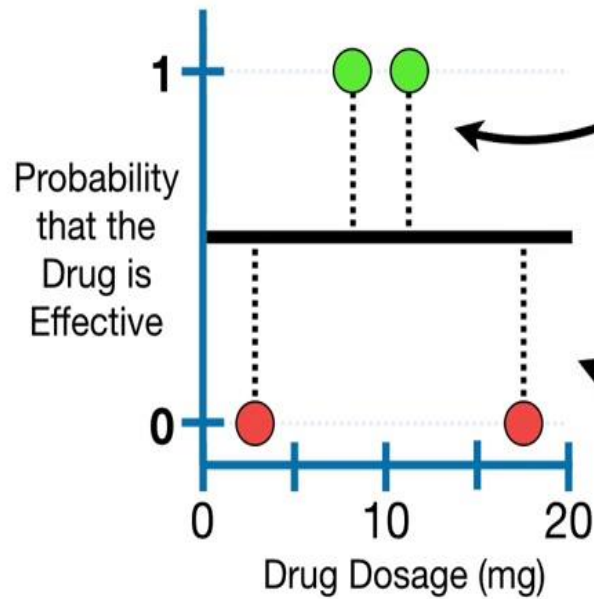
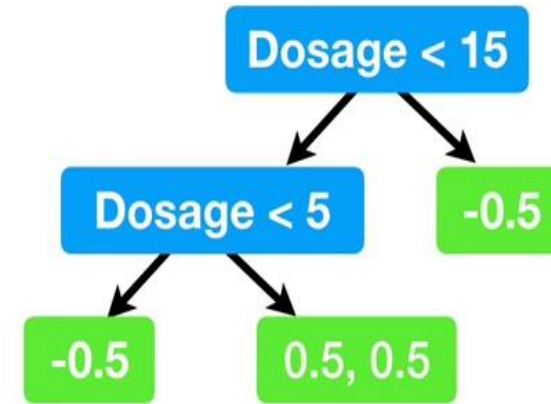




# XGBoost for Classification

Predicted Drug Effectiveness  
0.5

Now, just like we did for Regression, we fit an XGBoost Tree to the Residuals...

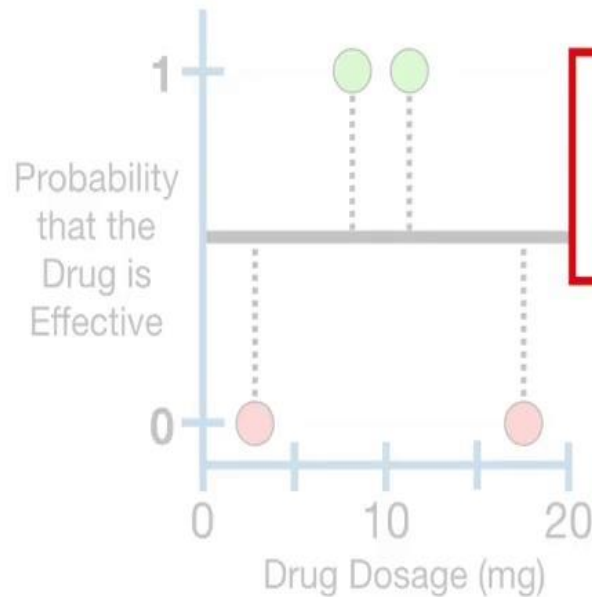
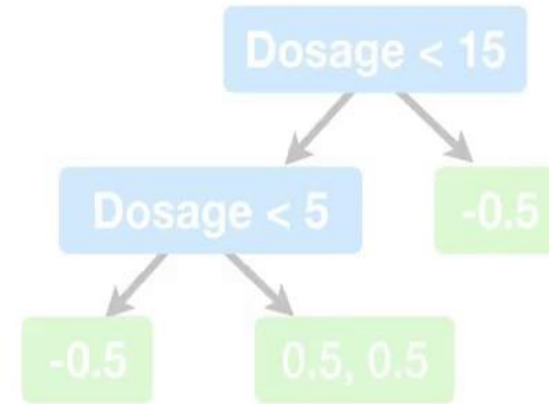


# XGBoost for Classification

Predicted Drug Effectiveness

0.5

...however, since we are using **XGBoost for Classification**, we have a new formula for the **Similarity Scores**.

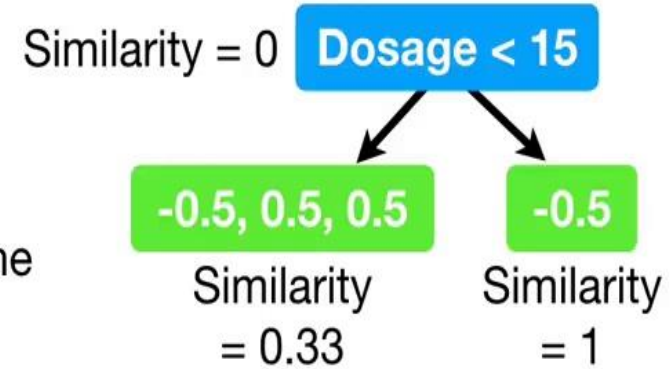
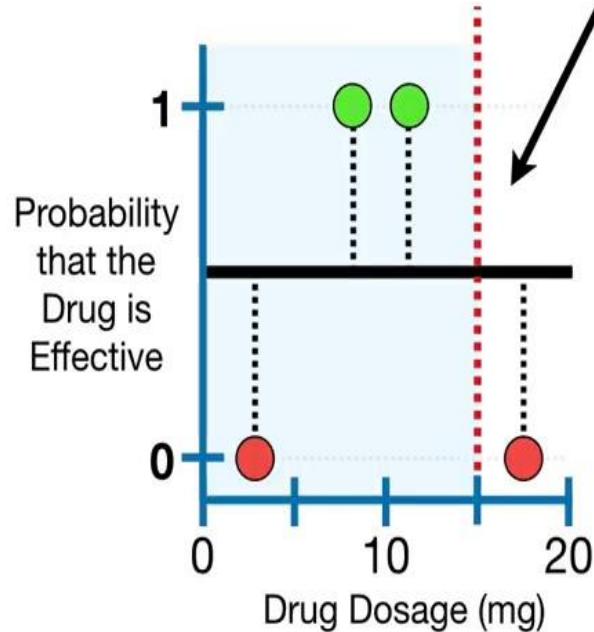


$$\frac{(\sum \text{Residual}_i)^2}{\sum [\text{Previous Probability}_i \times (1 - \text{Previous Probability}_i)] + \lambda}$$

# XGBoost for Classification

Predicted Drug Effectiveness  
0.5

So when we split the **Observations** based on the threshold **Dosage < 15**, **Gain = 1.33**.



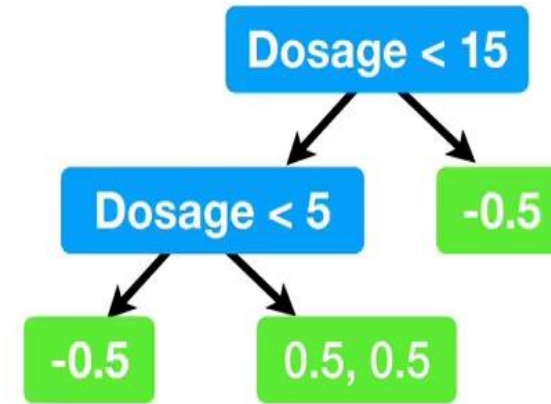
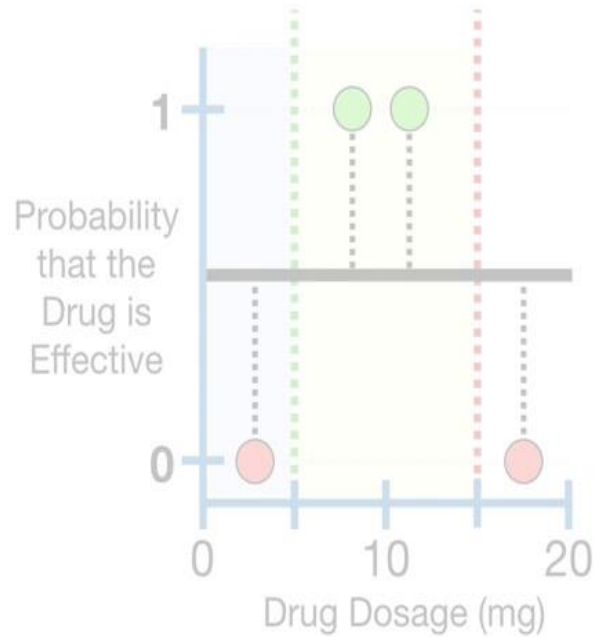
$$\text{Gain} = 0.33 + 1 - 0 = 1.33$$

# XGBoost for Classification

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.

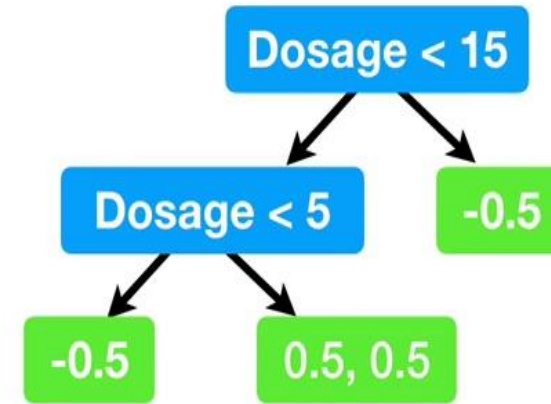
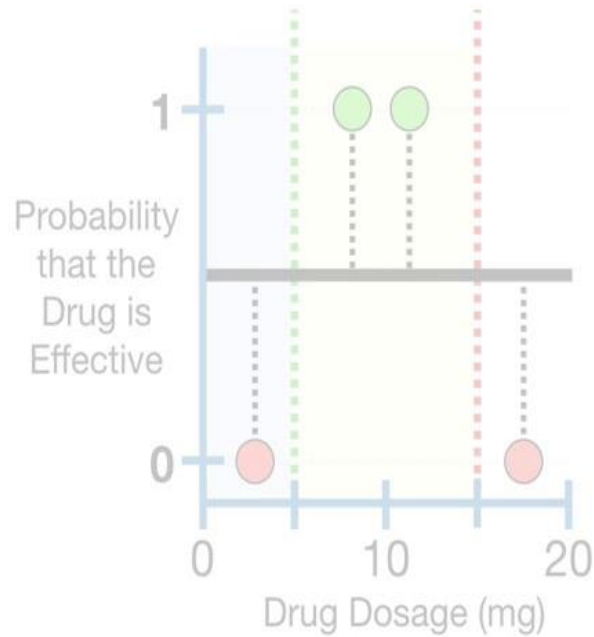


# XGBoost for Classification

Predicted Drug  
Effectiveness

0.5

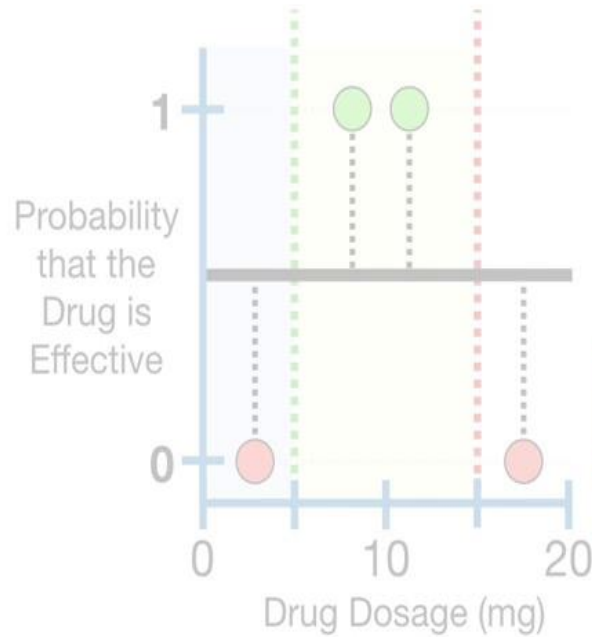
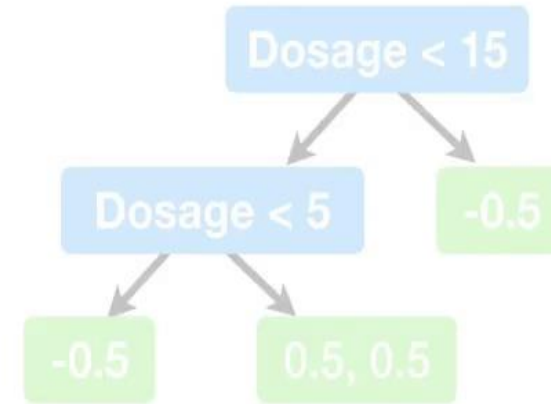
The minimum number of **Residuals**  
in each leaf is determined by  
calculating something called **Cover**.



# XGBoost for Classification

Predicted Drug Effectiveness  
0.5

**Cover** is defined as the denominator of the **Similarity Score** minus  $\lambda$  (**lambda**).



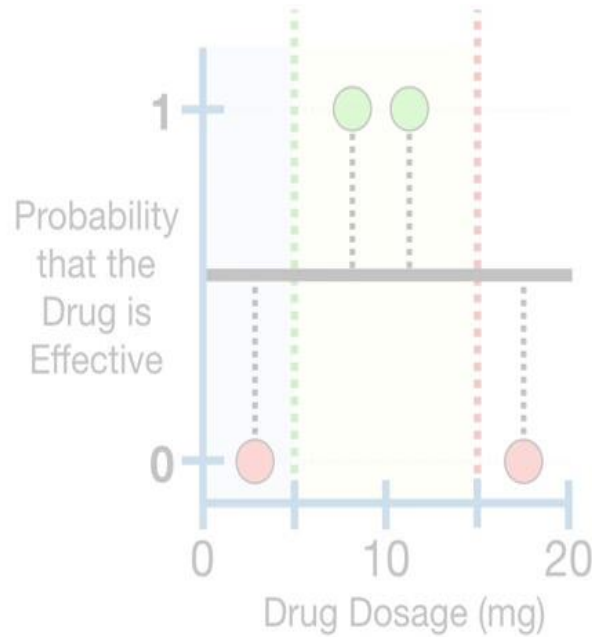
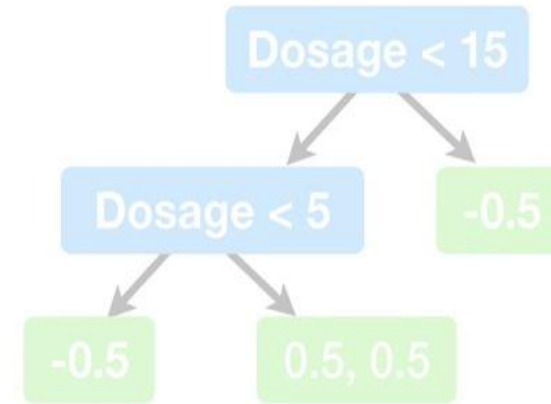
$$\text{Similarity} = \frac{(\sum \text{Residual}_i)^2}{\sum [\text{Previous Probability}_i \times (1 - \text{Previous Probability}_i)] - \lambda}$$

$$\sum [\text{Previous Probability}_i \times (1 - \text{Previous Probability}_i)] - \lambda$$

# XGBoost for Classification

Predicted Drug Effectiveness  
0.5

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

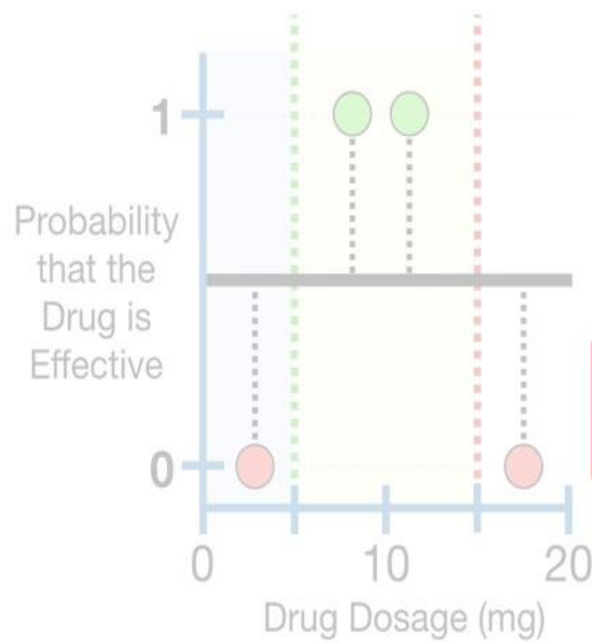
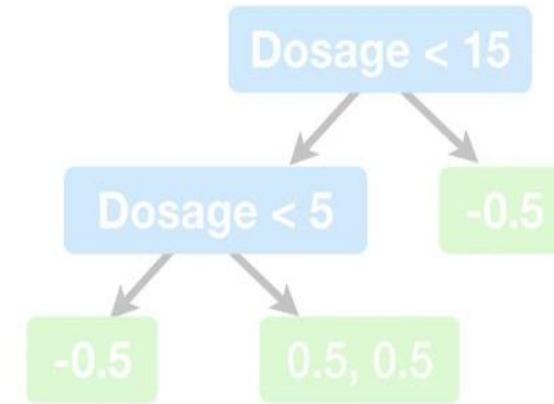


$$\text{Similarity} = \frac{(\sum \text{Residual}_i)^2}{\sum [\text{Previous Probability}_i \times (1 - \text{Previous Probability}_i)] + \lambda}$$

# XGBoost for Classification

Predicted Drug Effectiveness  
0.5

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



Similarity =

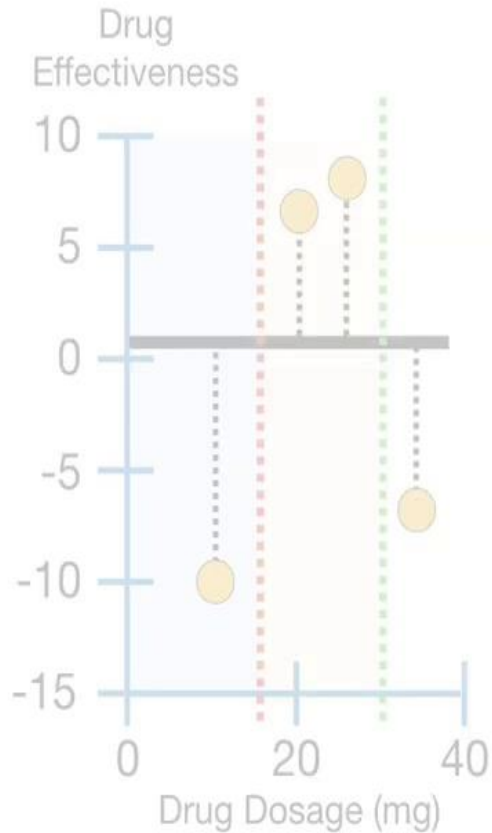
$$\left( \sum \text{Residual}_i \right)^2$$

$$\sum \left[ \text{Previous Probability}_i \times (1 - \text{Previous Probability}_i) \right] + \lambda$$

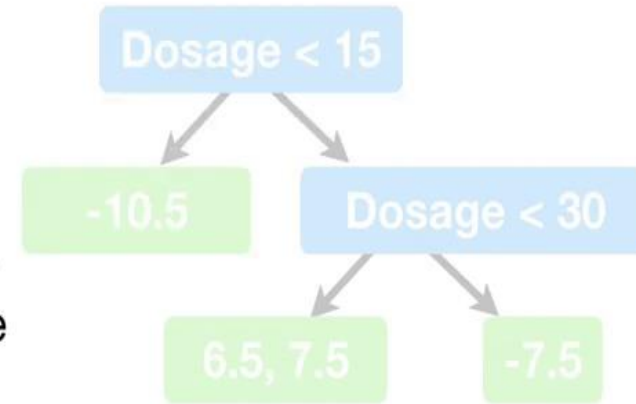


# XGBoost for Classification

Predicted Drug Effectiveness  
0.5



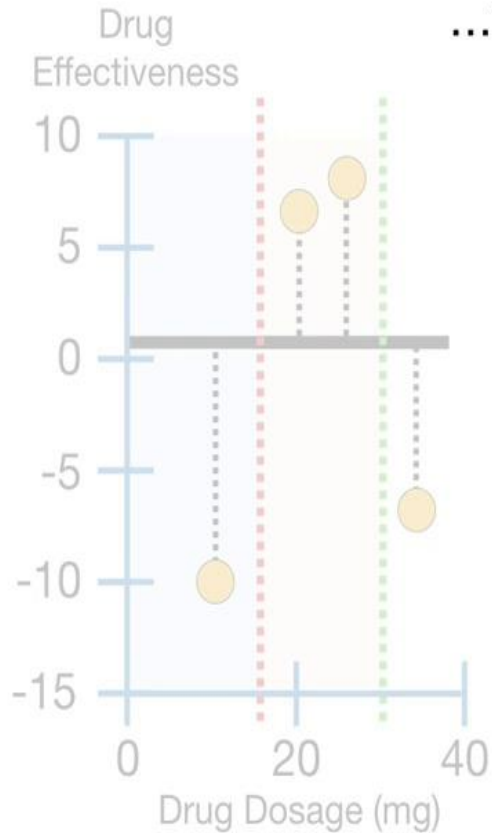
In contrast, when **XGBoost** is used for **Regression** and we are using this formula for the **Similarity Score**...



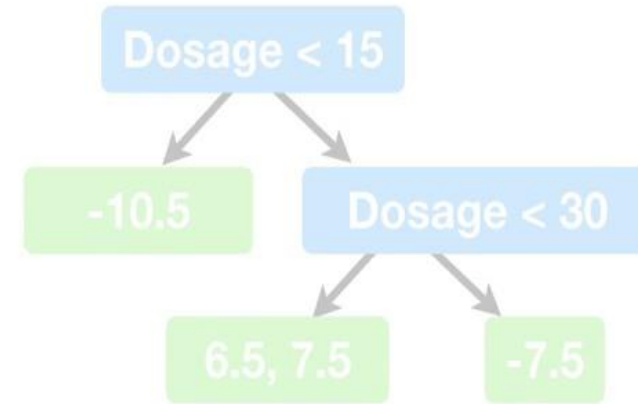
$$\text{Similarity Score} = \frac{\text{Sum of Residuals, Squared}}{\text{Number of Residuals} + \lambda}$$

# XGBoost for Classification

Predicted Drug Effectiveness  
0.5



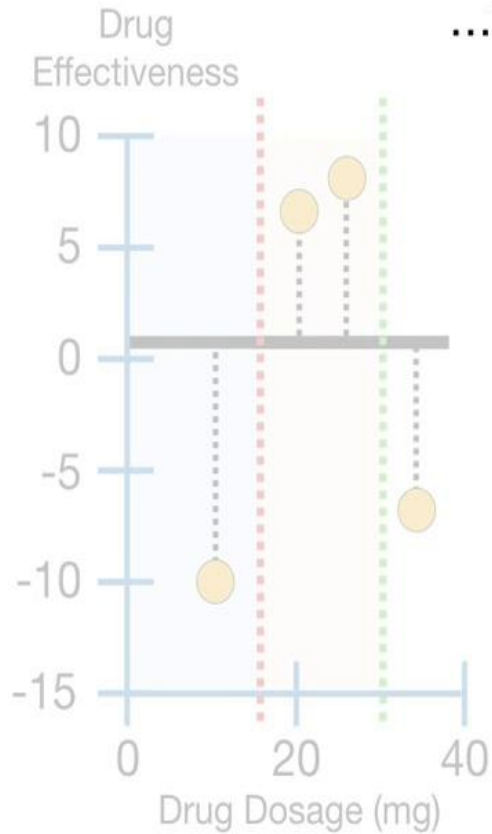
...then **Cover** is equal to...



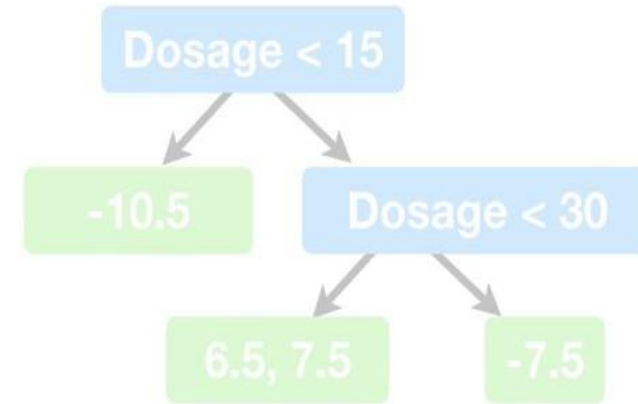
$$\text{Similarity Score} = \frac{\text{Sum of Residuals, Squared}}{\text{Number of Residuals} + \lambda}$$

# XGBoost for Classification

Predicted Drug Effectiveness  
0.5



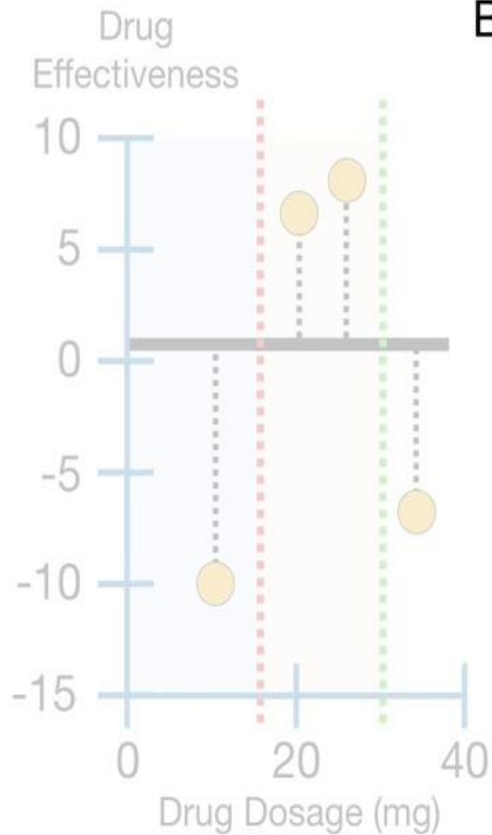
...then **Cover** is equal to...



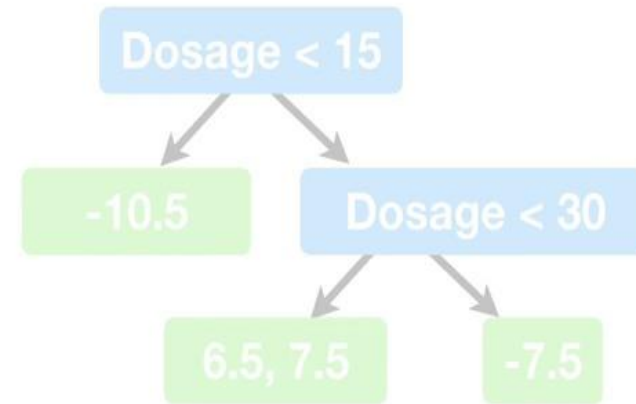
$$\text{Similarity Score} = \frac{\text{Sum of Residuals Squared}}{\text{Number of Residuals} + \lambda}$$

# XGBoost for Classification

Predicted Drug Effectiveness  
0.5



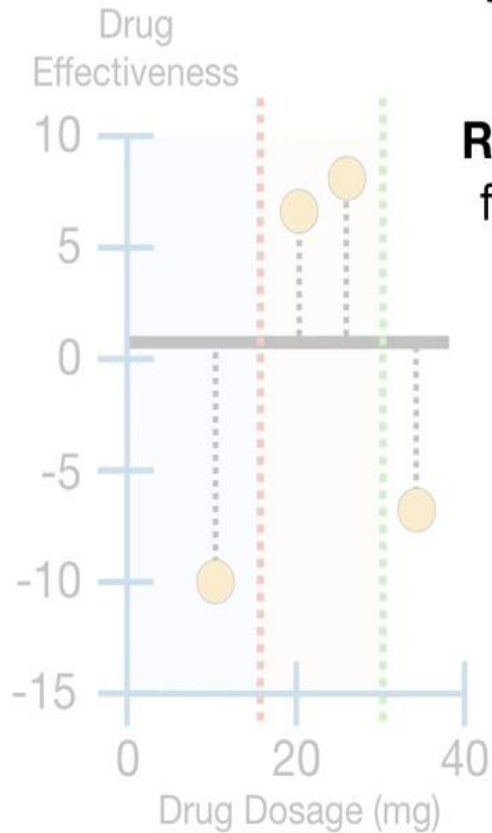
By default, the minimum value for **Cover** is 1.



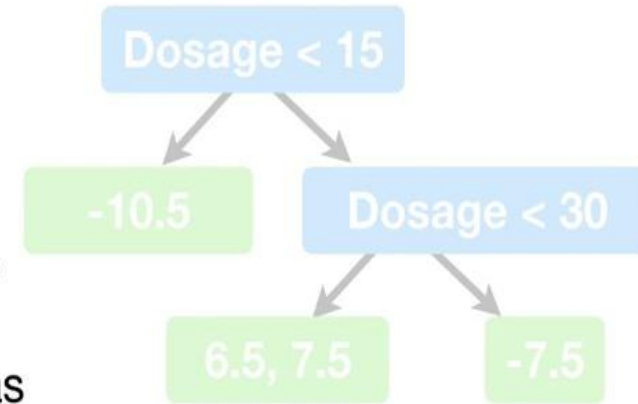
$$\text{Similarity Score} = \frac{\text{Sum of Residuals Squared}}{\text{Number of Residuals} + \lambda}$$

# XGBoost for Classification

Predicted Drug Effectiveness  
0.5



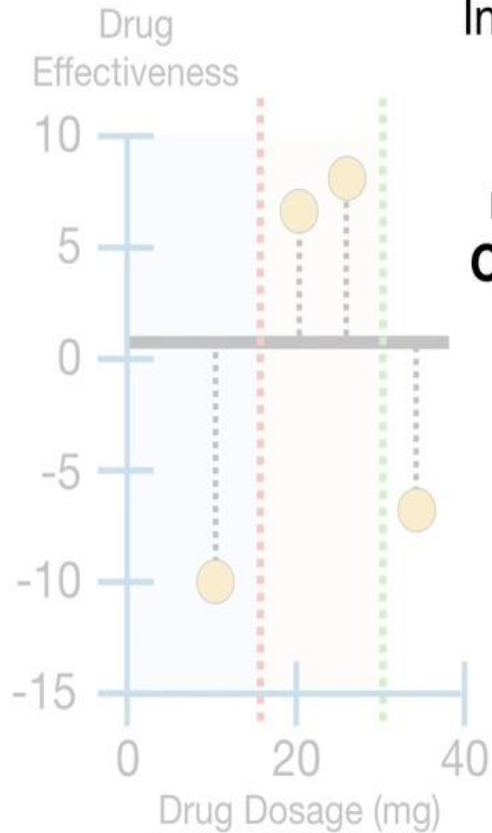
Thus, by default, when we use **XGBoost** for **Regression**, we can have as few as **1 Residual** per leaf.



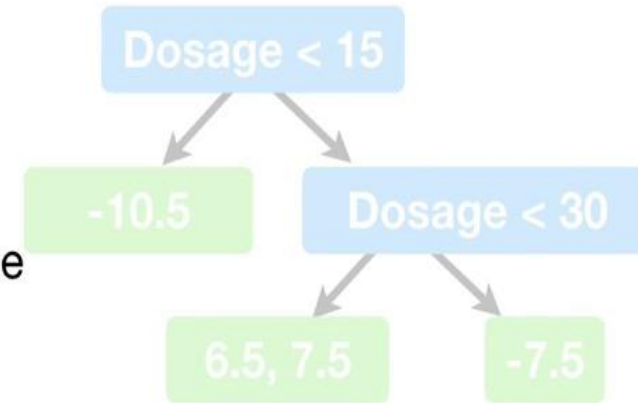
$$\text{Similarity Score} = \frac{\text{Sum of Residuals Squared}}{\text{Number of Residuals} + \lambda}$$

# XGBoost for Classification

Predicted Drug Effectiveness  
0.5



In other words, when we use **XGBoost** for **Regression** and use the default minimum value for **Cover**, **Cover** has no effect on how we grow the tree.

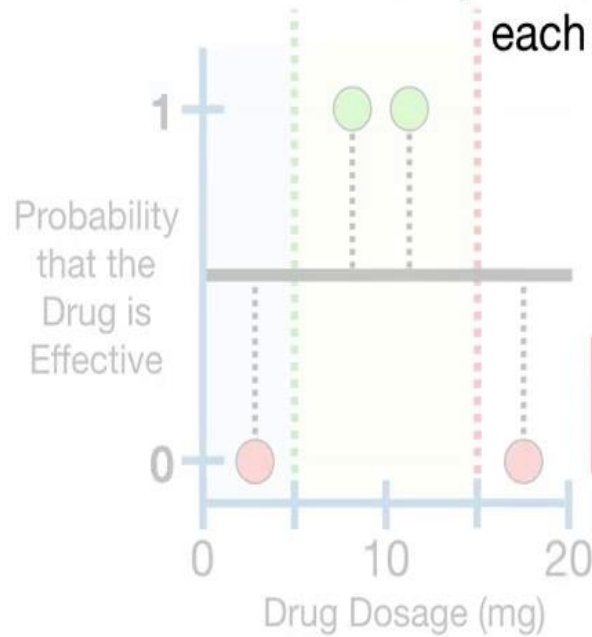
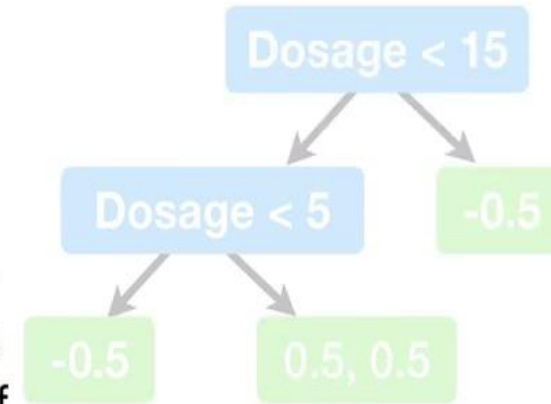


$$\text{Similarity Score} = \frac{\text{Sum of Residuals Squared}}{\text{Number of Residuals} + \lambda}$$

# XGBoost for Classification

Predicted Drug Effectiveness  
0.5

In contrast, things are way more complicated when we use **XGBoost** for **Classification** because **Cover** depends on the previously predicted probability of each **Residual** in a leaf.



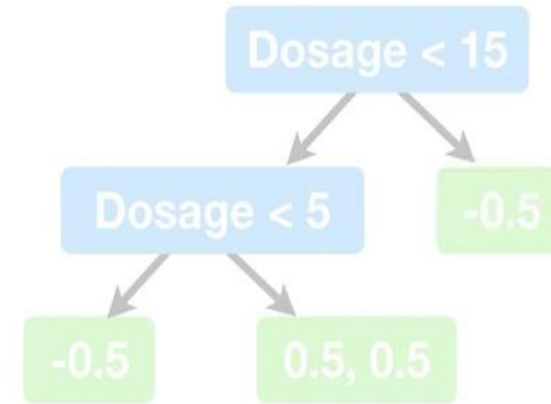
Similarity =

$$\left( \sum \text{Residual}_i \right)^2$$

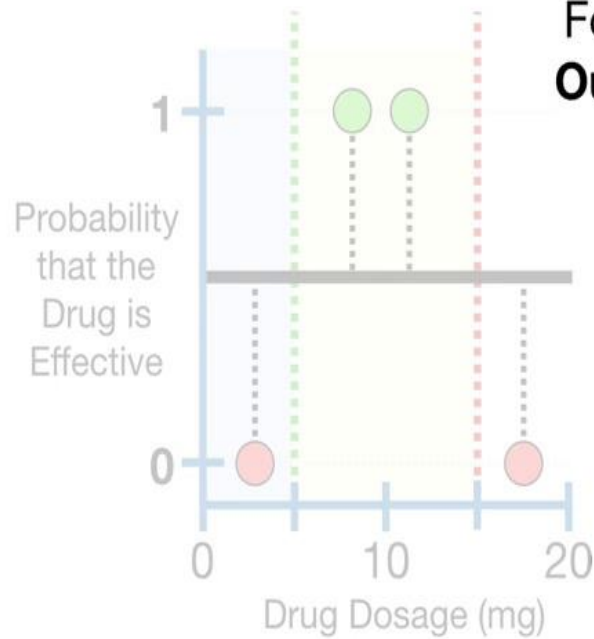
$$\sum \left[ \text{Previous Probability}_i \times (1 - \text{Previous Probability}_i) \right] + \lambda$$

# XGBoost for Classification

Predicted Drug Effectiveness  
0.5



For **Classification**, the **Output Value** for a leaf is...

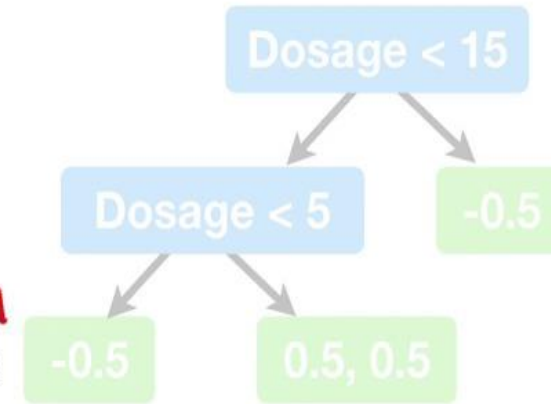


$$\frac{(\sum \text{Residual}_i)}{\sum [\text{Previous Probability}_i \times (1 - \text{Previous Probability}_i)] + \lambda}$$

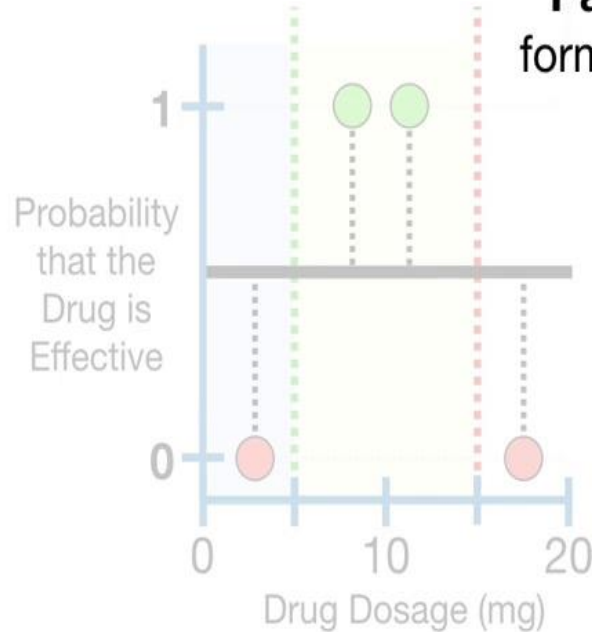


# XGBoost for Classification

Predicted Drug Effectiveness  
0.5



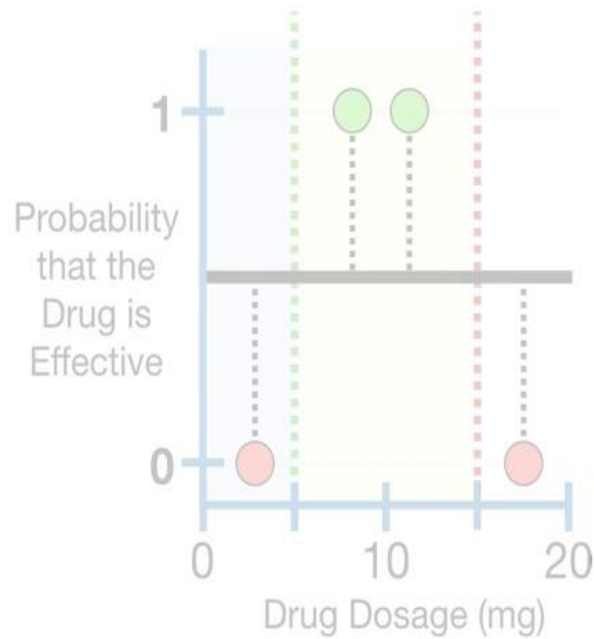
**NOTE:** With the exception of  $\lambda$  (**lambda**), the **Regularization Parameter**, this is the same formula we used for unextreme **Gradient Boost**.



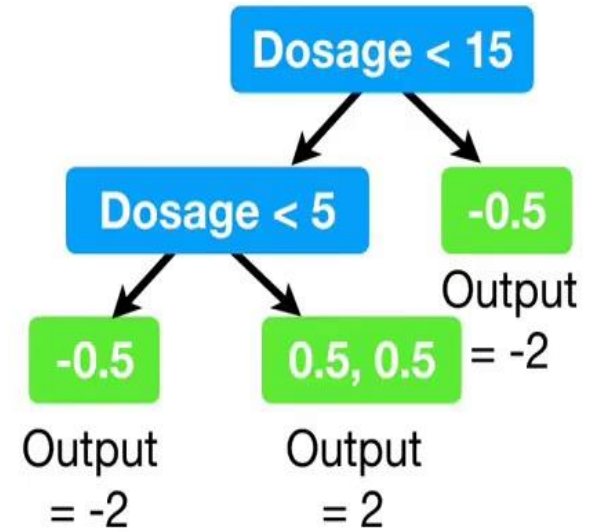
$$\frac{(\sum \text{Residual}_i)}{\sum [\text{Previous Probability}_i \times (1 - \text{Previous Probability}_i)] + \lambda}$$

# XGBoost for Classification

Predicted Drug Effectiveness  
0.5



The first tree is complete!

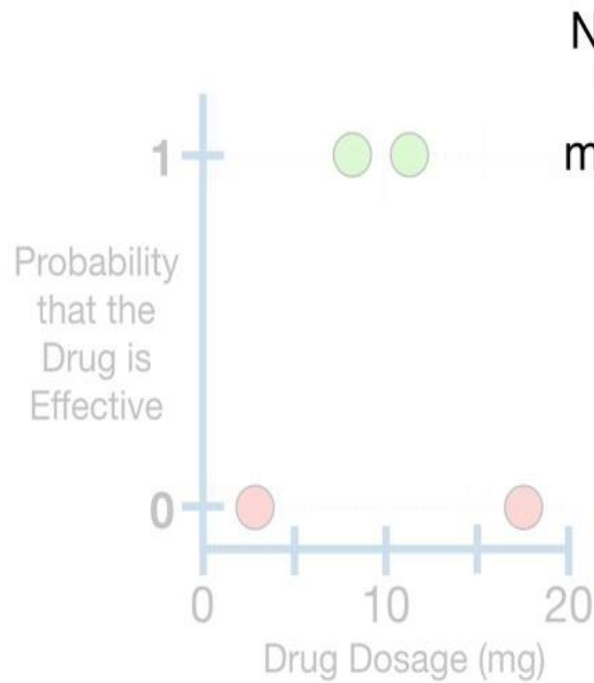


$$-0.5 / (0.5 \times (1 - 0.5) + 0) = -2$$

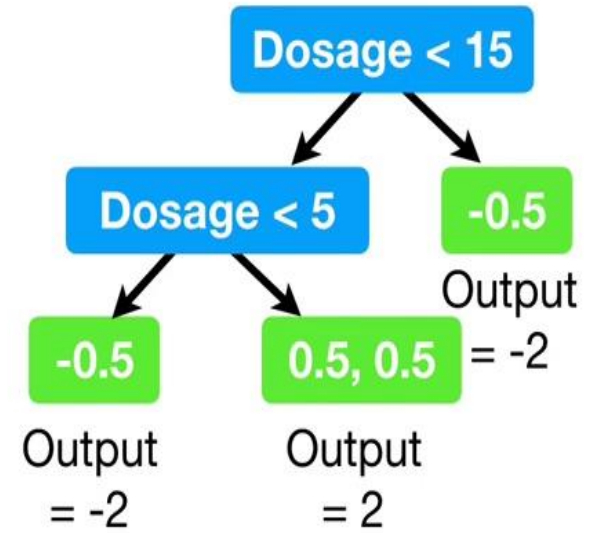
$$\frac{(\sum \text{Residual}_i)}{\sum [\text{Previous Probability}_i \times (1 - \text{Previous Probability}_i)] + \lambda}$$

# XGBoost for Classification

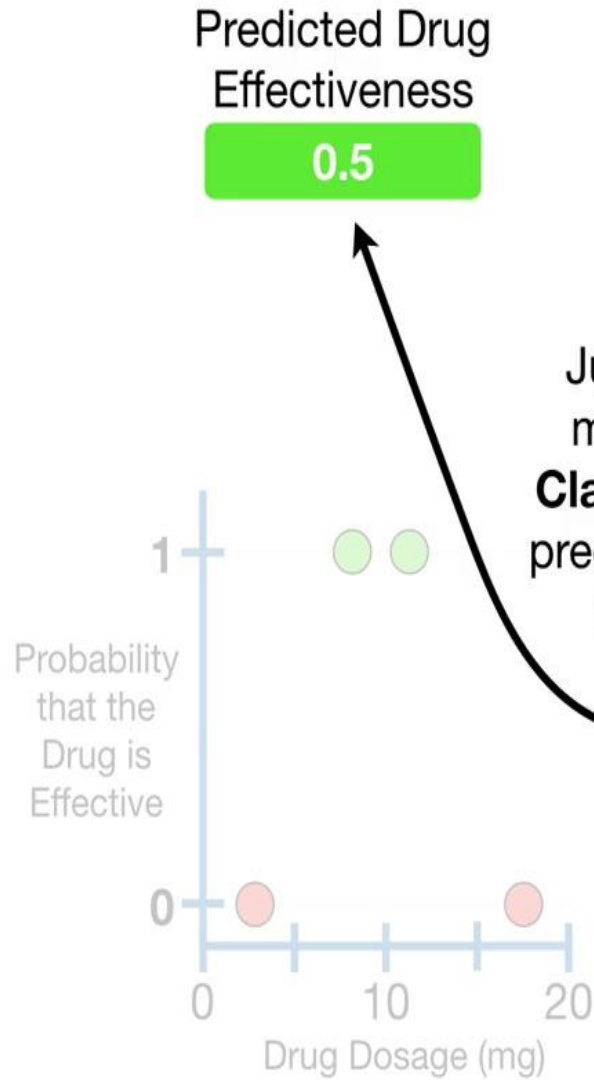
Predicted Drug Effectiveness  
0.5



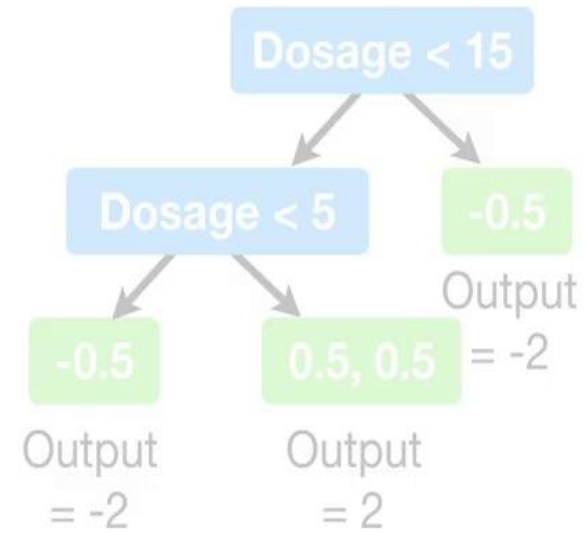
Now that we have built the first tree, we can make new **Predictions**.



# XGBoost for Classification



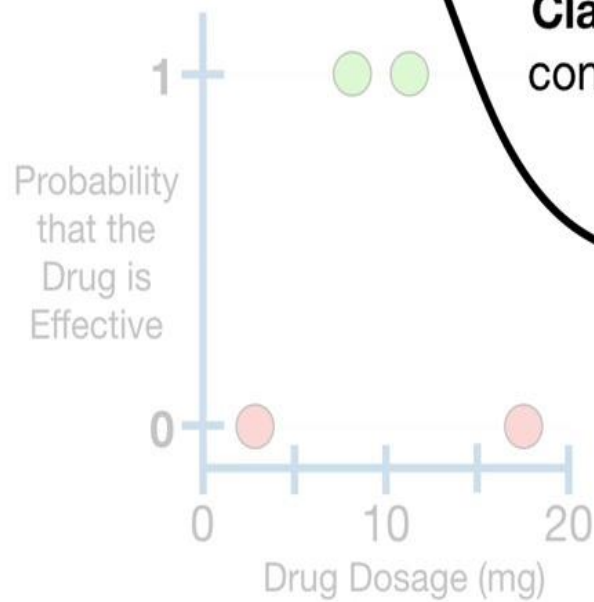
Just like other boosting methods, **XGBoost for Classification** makes new predictions by starting with the initial prediction.



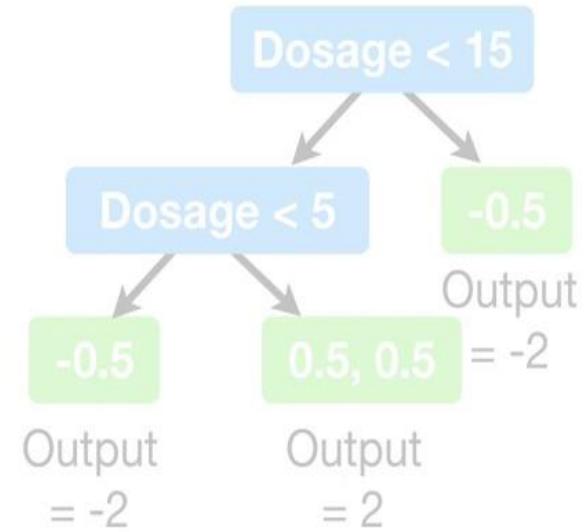
# XGBoost for Classification

Predicted Drug Effectiveness

0.5



However, just like with unextreme **Gradient Boost for Classification**, we need to convert this probability to a **log(odds)** value.

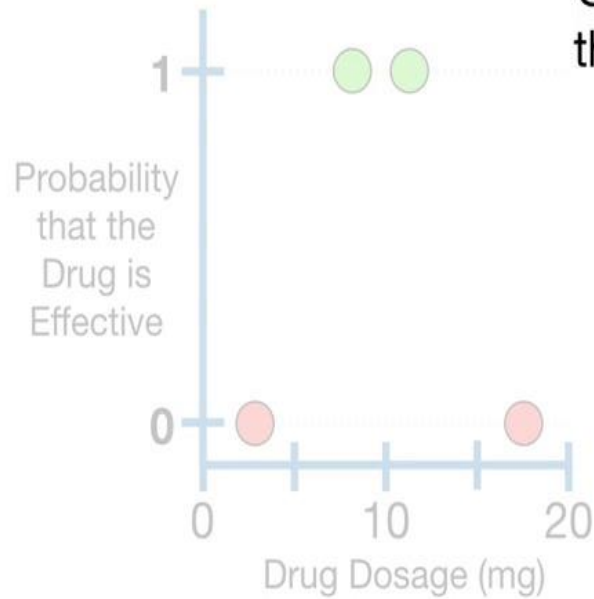


# XGBoost for Classification

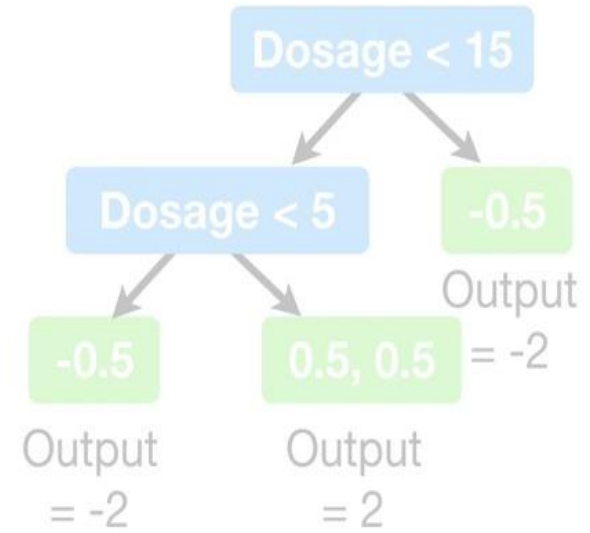
Predicted Drug  
Effectiveness

0.5

Output =  $\log(\text{odds}) = 0$



So let's put that under the initial prediction so we don't forget.

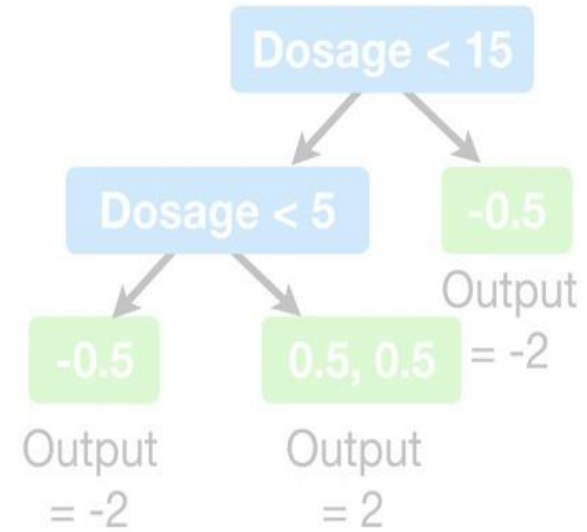
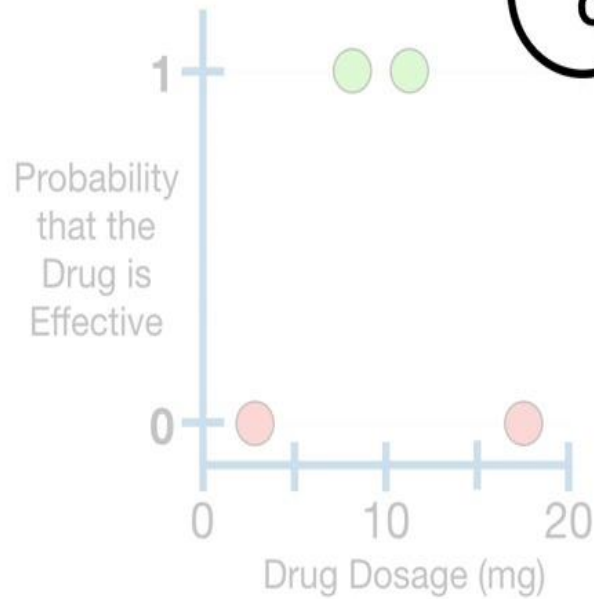


# XGBoost for Classification

Predicted Drug Effectiveness  
**0.5**  
Output =  $\log(\text{odds}) = 0$

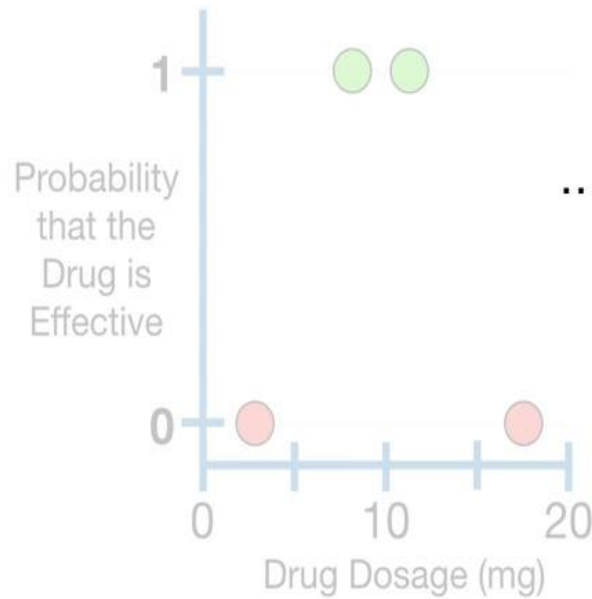
+

Now, just like unextreme **Gradient Boost for Classification**, we add the  **$\log(\text{odds})$**  of the initial prediction...

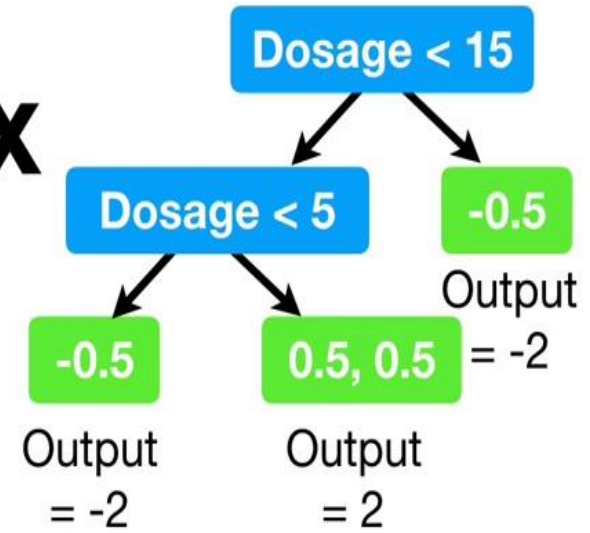


# XGBoost for Classification

Predicted Drug Effectiveness  
**0.5** +  
Output =  $\log(\text{odds}) = 0$



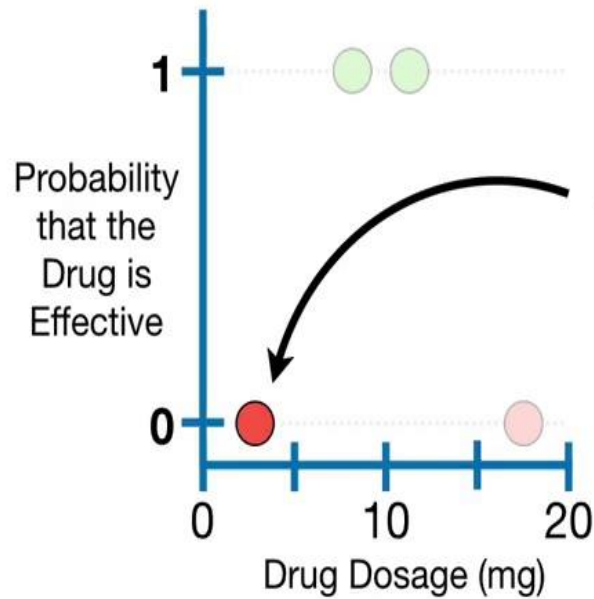
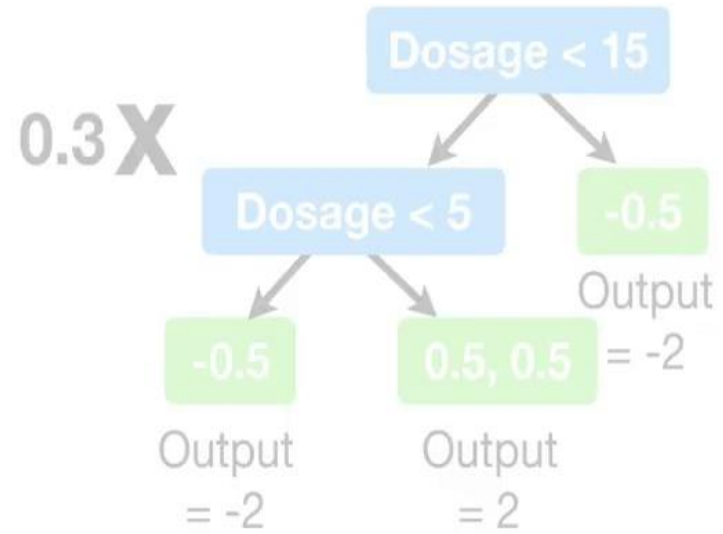
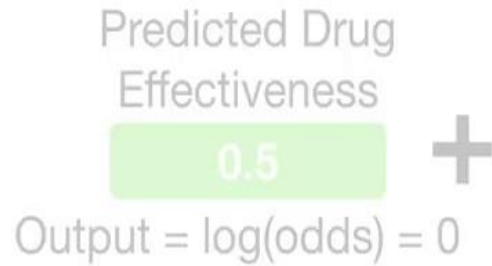
Learning Rate **X**



...to the output of the **Tree**, scaled by a **Learning Rate**.



# XGBoost for Classification

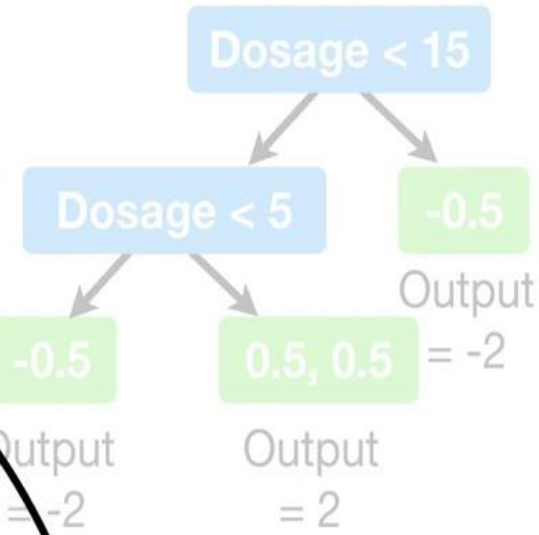


Thus, the new **Predicted** value for this observation, with **Dosage = 2...**

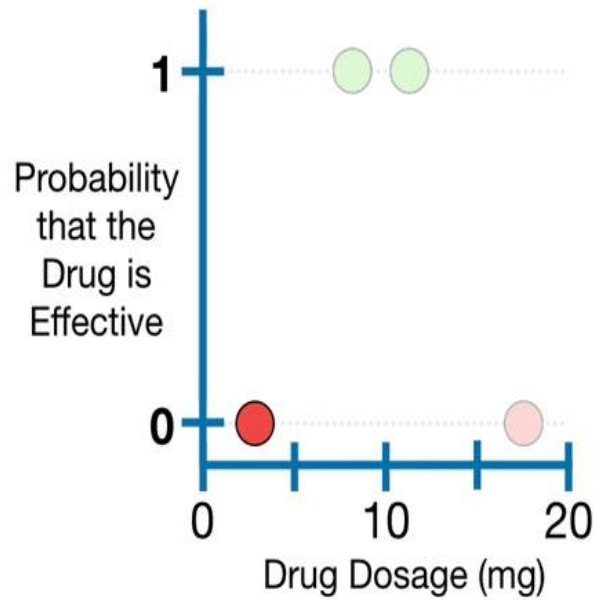
# XGBoost for Classification

Predicted Drug Effectiveness  
0.5  
Output =  $\log(\text{odds}) = 0$

0.3 X



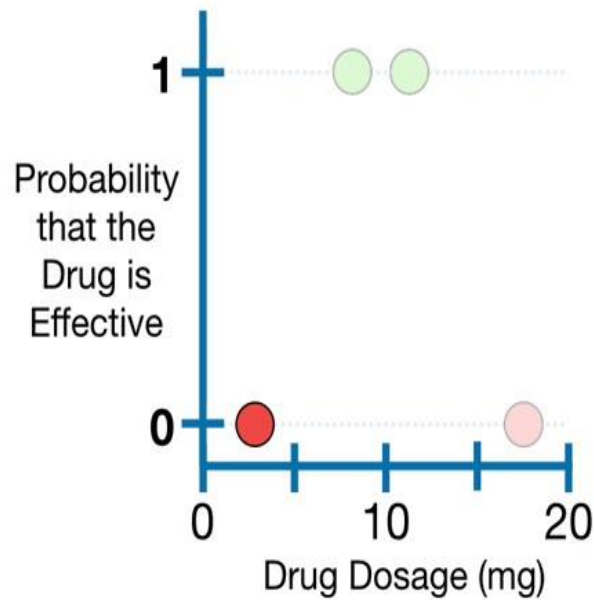
...is the **log(odds)** of original prediction, 0...



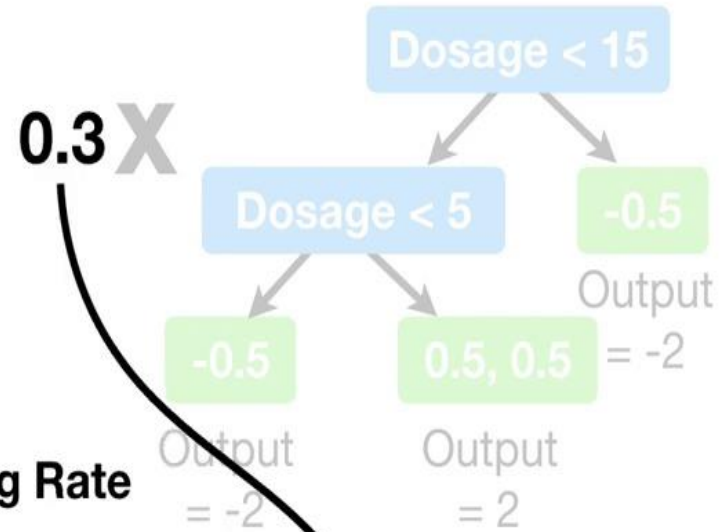
log(odds) Prediction = 0

# XGBoost for Classification

Predicted Drug Effectiveness  
**0.5** +  
Output =  $\log(\text{odds}) = 0$



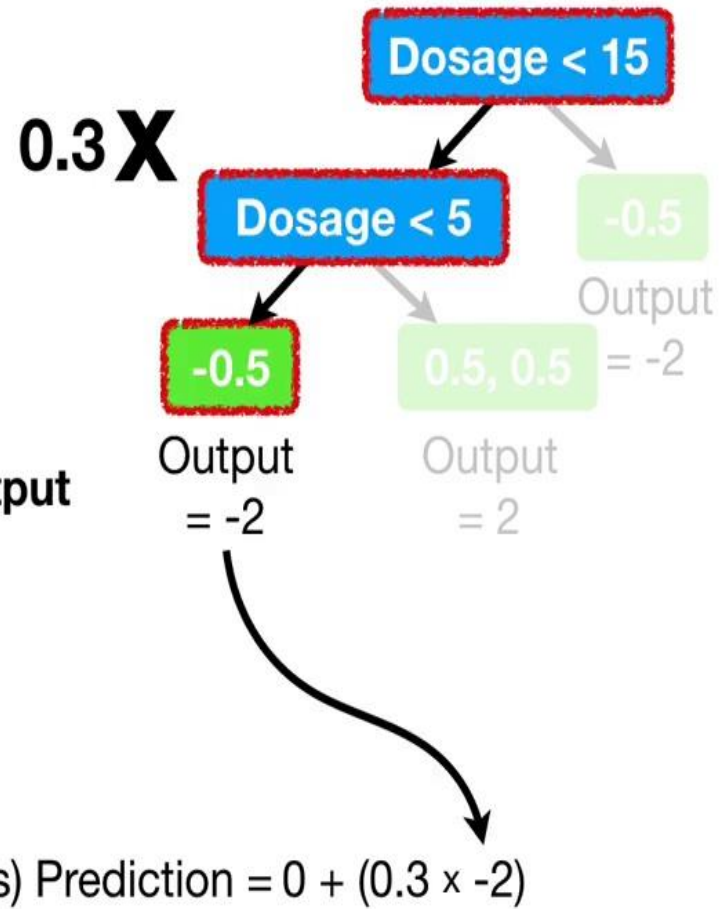
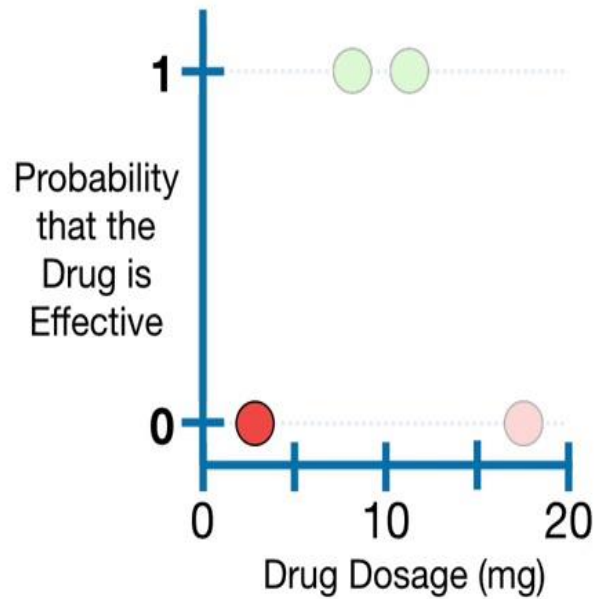
...plus the **Learning Rate** ( $\epsilon$ , eta), **0.3**...



$$\log(\text{odds}) \text{ Prediction} = 0 + (0.3)$$

# XGBoost for Classification

Predicted Drug Effectiveness  
**0.5** +  
Output =  $\log(\text{odds}) = 0$

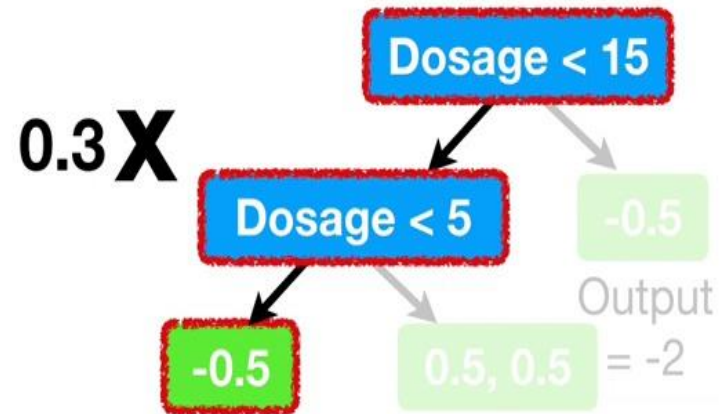
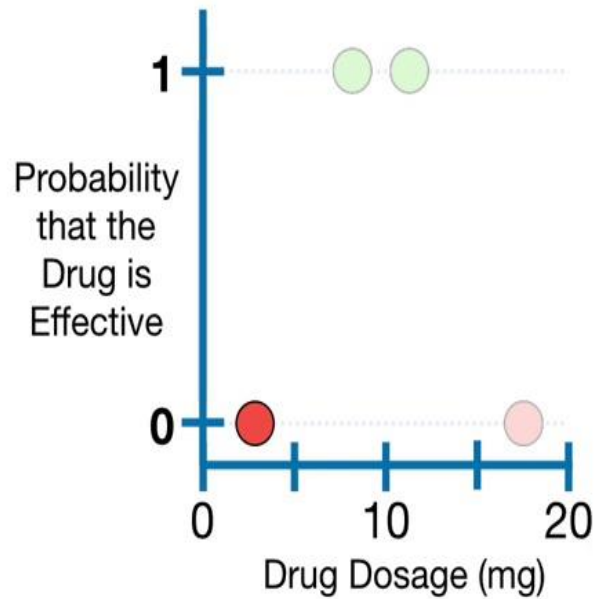


...times the **Output Value, -2...**

$$\log(\text{odds}) \text{ Prediction} = 0 + (0.3 \times -2)$$

# XGBoost for Classification

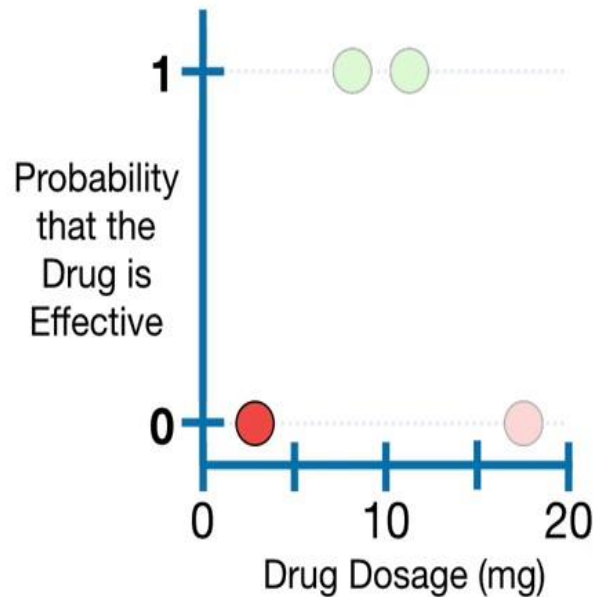
Predicted Drug Effectiveness  
**0.5** +  
Output =  $\log(\text{odds}) = 0$



...and that gives us a **log(odds)** value = **-0.6**.

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

# XGBoost for Classification

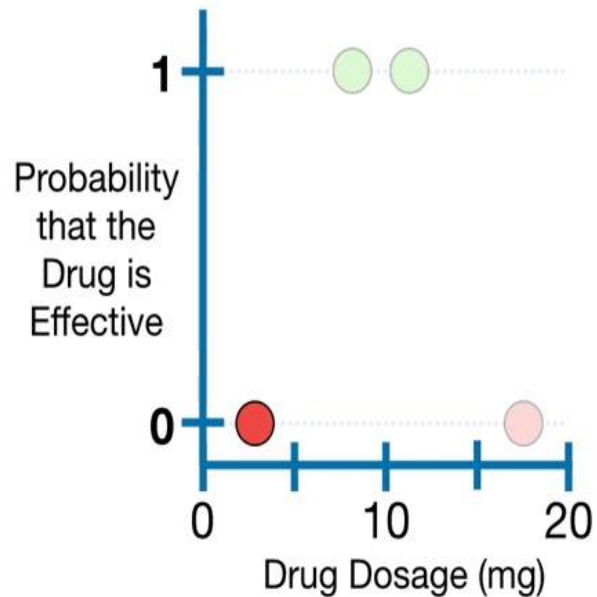


To convert a **log(odds)** value into a probability, we plug it into the **Logistic Function**.

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

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

# XGBoost for Classification



...and the the new  
predicted probability is  
**0.35.**

$$\text{Probability} = \frac{e^{-0.6}}{1 + e^{-0.6}} = 0.35$$

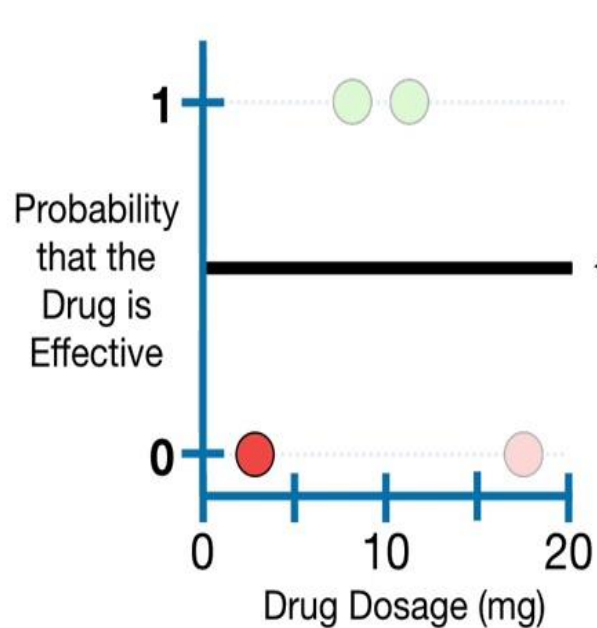
$$\text{log(odds) Prediction} = 0 + (0.3 \times -2) = -0.6$$

# XGBoost for Classification

Predicted Drug  
Effectiveness

0.5

Output =  $\log(\text{odds}) = 0$



Remember, the original  
**Prediction** was **0.5**...

$$\text{Probability} = \frac{e^{-0.6}}{1 + e^{-0.6}} = 0.35$$

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

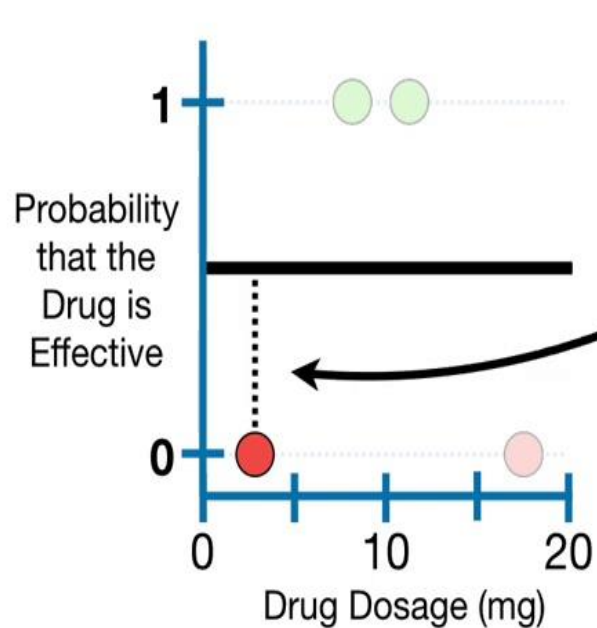


# XGBoost for Classification

Predicted Drug  
Effectiveness

0.5

Output =  $\log(\text{odds}) = 0$



...and this was the original **Residual**.

$$\text{Probability} = \frac{e^{-0.6}}{1 + e^{-0.6}} = 0.35$$

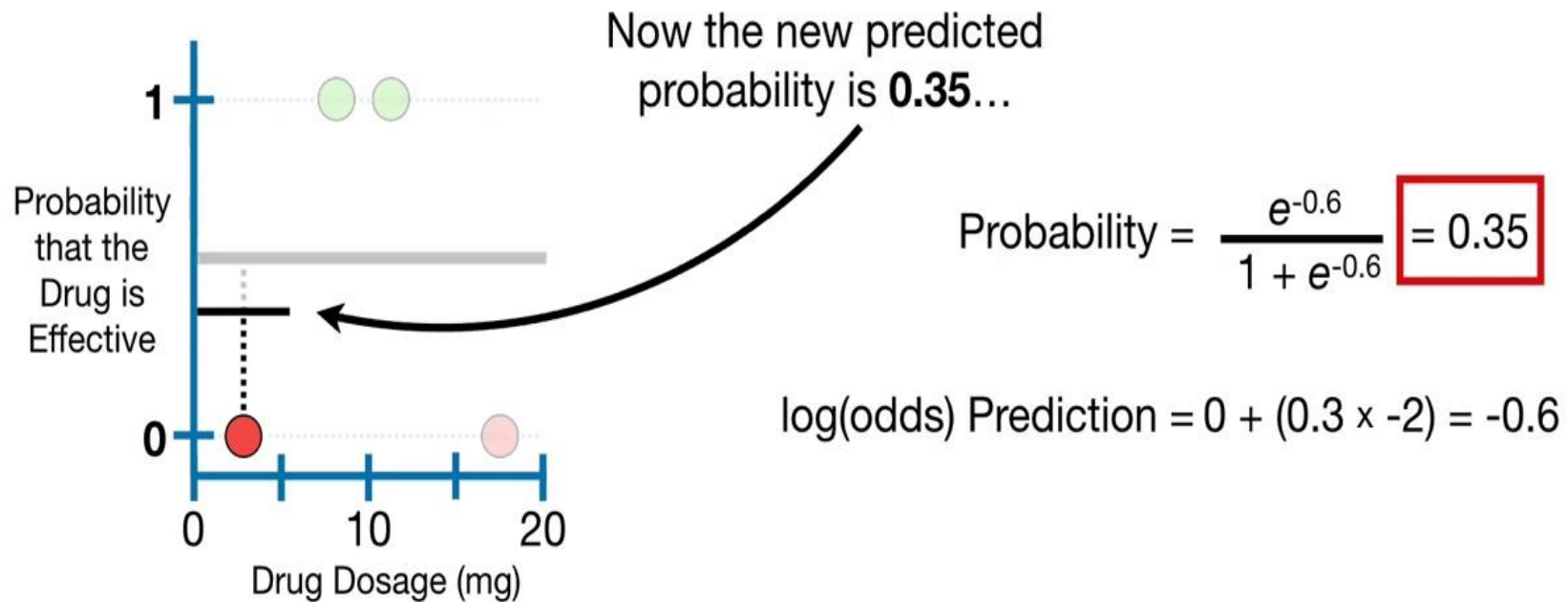
$$\log(\text{odds}) \text{ Prediction} = 0 + (0.3 \times -2) = -0.6$$

# XGBoost for Classification

Predicted Drug  
Effectiveness

0.5

Output =  $\log(\text{odds}) = 0$

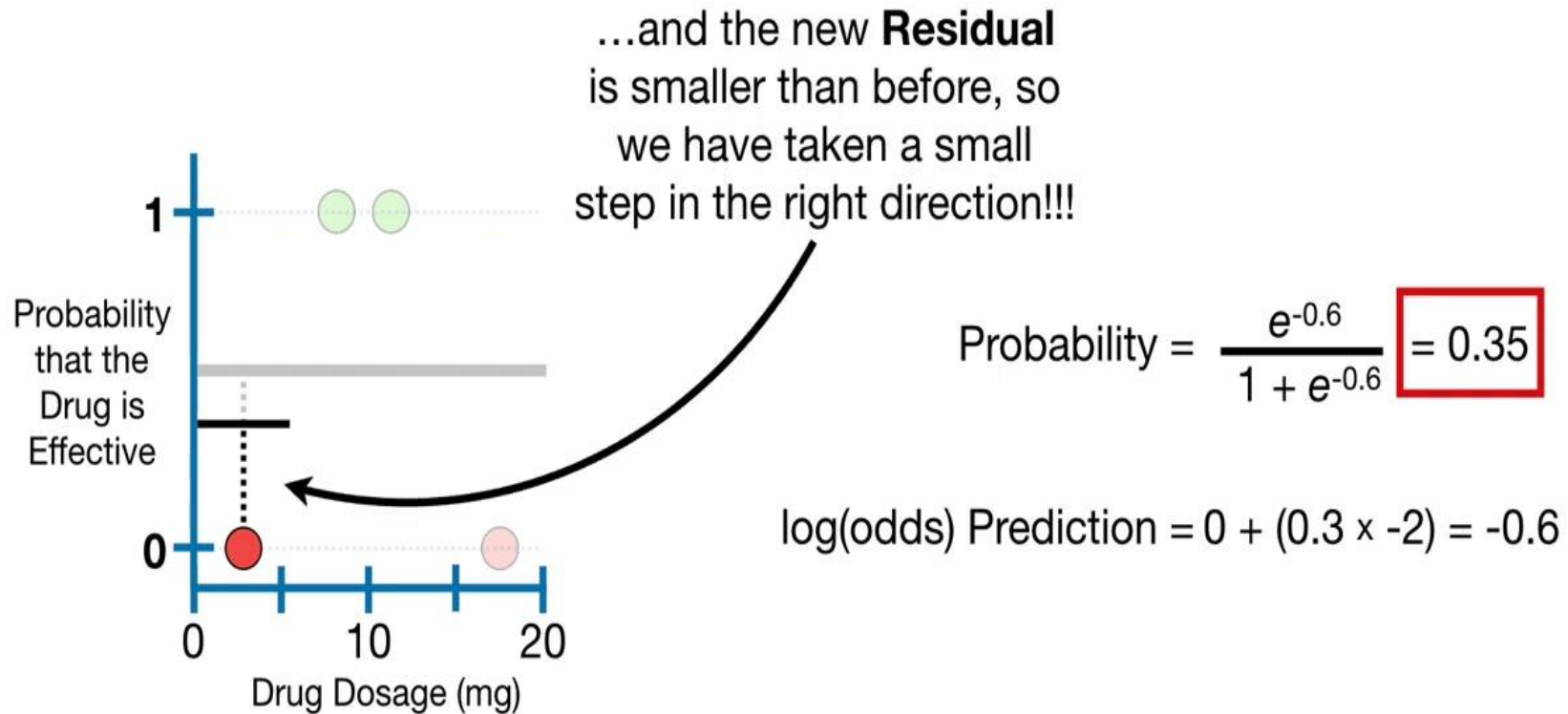


# XGBoost for Classification

Predicted Drug  
Effectiveness

0.5

Output =  $\log(\text{odds}) = 0$



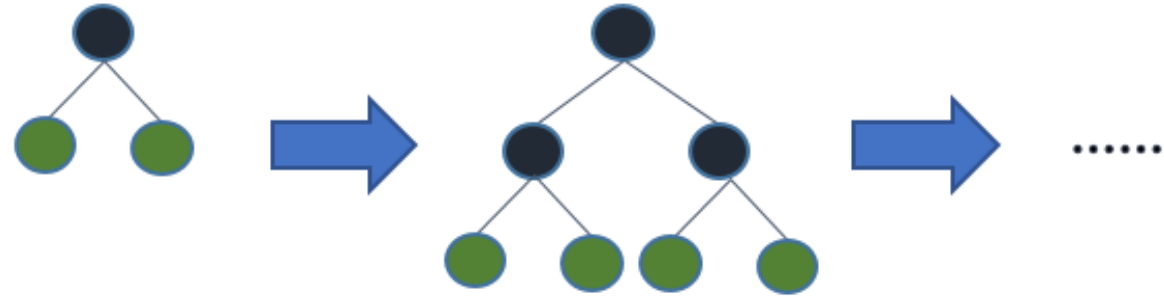
# LightGBM

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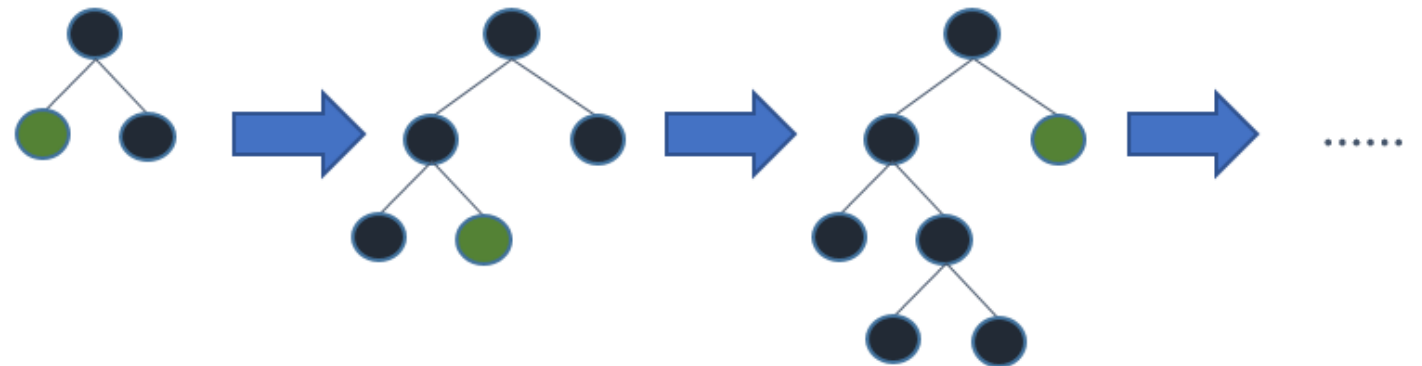
- LightGBM (by Microsoft) is a distributed high-performance framework that uses decision trees for classification, and regression tasks.
- Advantage w.r.t. XGBoost
  - Faster training speed and accuracy resulting from LightGBM being a histogram-based algorithm that performs bucketing of values (also 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)

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- 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)

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- 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

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- Handling Categorical Features
- Handling Missing Values



# References

- Mason, L., Baxter, J., Bartlett, P., & Frean, M. (1999). Boosting algorithms as gradient descent. *Advances in neural information processing systems*, 12.
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- 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).
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