

XGBoost (classification)

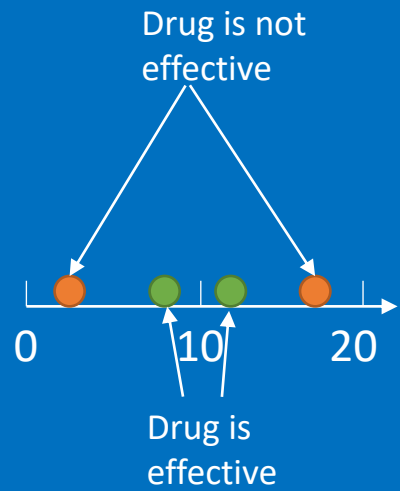
Drug dosage	Drug usefulness
5	No
8	Yes
11	Yes
18	No

Assuming that we
have this very simple
dataset

Step 1: make the initial prediction

Drug dosage	Drug usefulness
5	No
8	Yes
11	Yes
18	No

Plot it out, we have

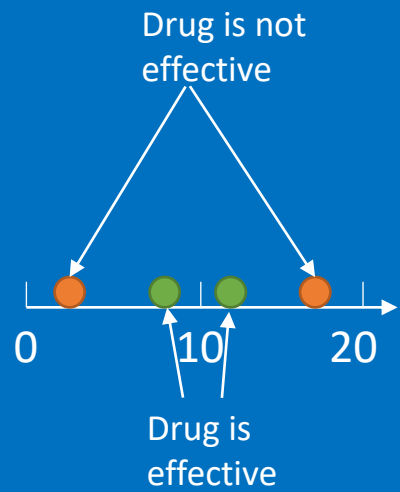


Step 1: make the initial prediction

For example, the probability of the dosage is effective: 0.5

Drug dosage	Drug usefulness
5	No
8	Yes
11	Yes
18	No

Plot it out, we have



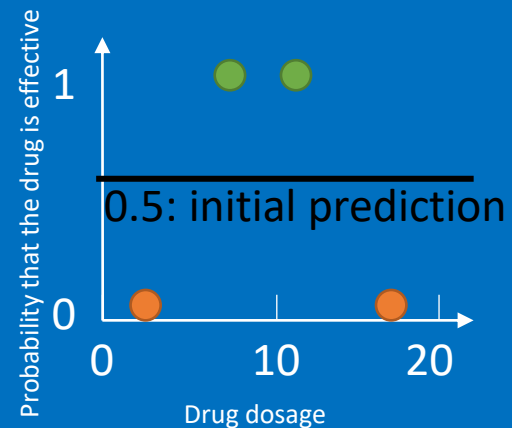
Step 1: make the initial prediction

For example, the probability of the dosage is effective: 0.5

In other words, does not matter how many dosages we take, the initial prediction for the usefulness for the drug is 50%

Drug dosage	Drug usefulness
5	No
8	Yes
11	Yes
18	No

Plot it out, we have



Step 1: make the initial prediction

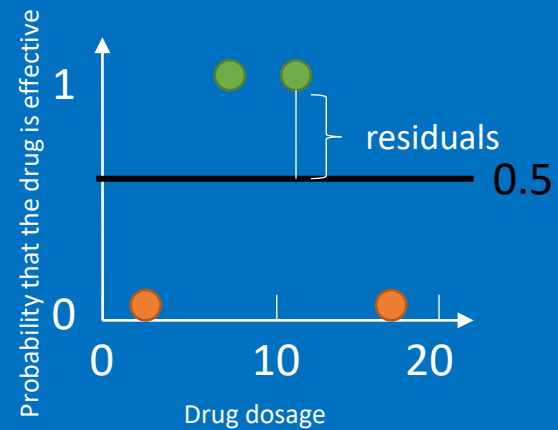
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5



Plot it out, we have



Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

For example, the probability of the dosage is effective: 0.5

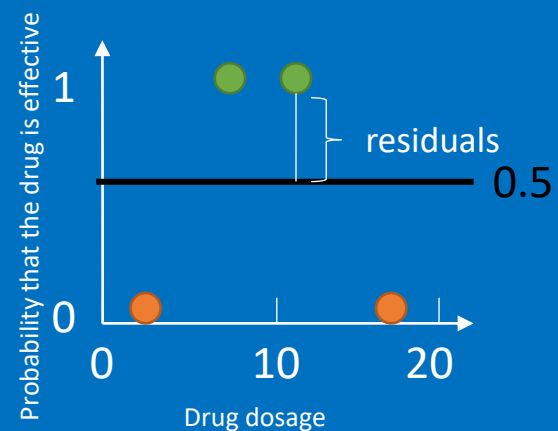
Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

3.1 All the tree starts from a leaf, with all the residuals

-0.5, 0.5, 0.5, -0.5

Plot it out, we have



Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

3.1 All the tree starts from a leaf, with all the residuals

-0.5, 0.5, 0.5, -0.5

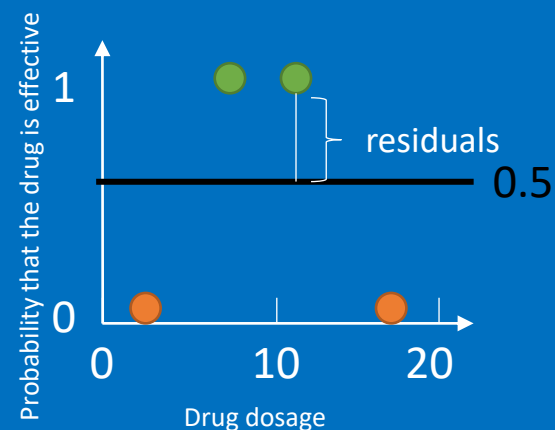
Note that for regression:

$$\text{Similarity score} = \frac{(\sum \text{residuals})^2}{\text{number of residuals} + \lambda}$$

3.2 Calculate the Similarity score using

$$\text{Similarity score} = \frac{(\sum \text{residuals})^2}{\sum \text{previous prob}_i \times (1 - \text{previous prob}_i) + \lambda}$$

Plot it out, we have



Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

3.1 All the tree starts from a leaf, with all the residuals

-0.5, 0.5, 0.5, -0.5

Note that for regression:

$$\text{Similarity score} = \frac{(\sum \text{residuals})^2}{\text{number of residuals} + \lambda}$$

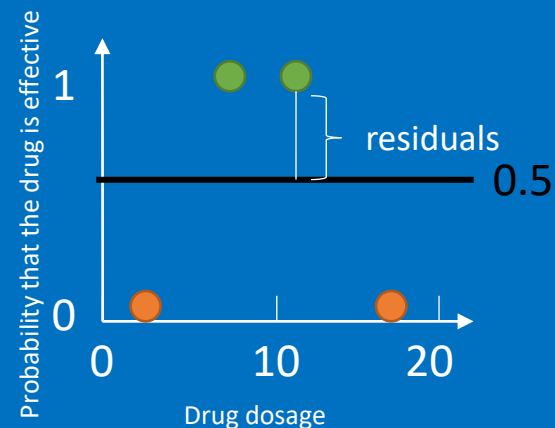
3.2 Calculate the Similarity score using

$$\text{Similarity score} = \frac{(\sum \text{residuals})^2}{\sum \text{previous prob}_i \times (1 - \text{previous prob}_i) + \lambda}$$

For this case we have

$$\text{Similarity score} = \frac{(-0.5 + 0.5 + 0.5 - 0.5)^2}{\sum \text{previous prob}_i \times (1 - \text{previous prob}_i) + \lambda} = 0$$

Plot it out, we have



Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

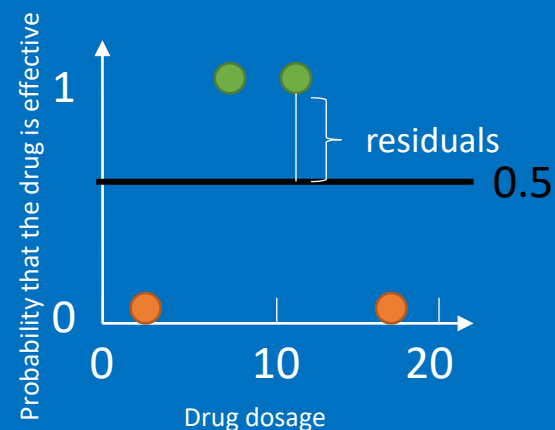
3.1 All the tree starts from a leaf, with all the residuals

-0.5, 0.5, 0.5, -0.5 0 *Similarity score*

3.2 Calculate the Similarity score

3.3 Further split the tree

Plot it out, we have



Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

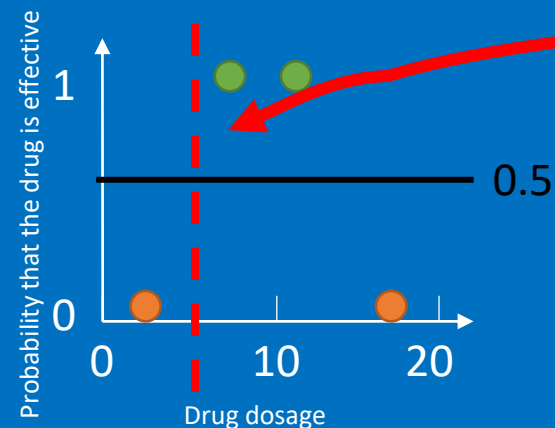
3.1 All the tree starts from a leaf, with all the residuals

-0.5, 0.5, 0.5, -0.5 0 *Similarity score*

3.2 Calculate the Similarity score

3.3 Further split the tree

Plot it out, we have



Similar to regression xgboost tree, we can start from the first data point, calculate the similarity score, and the gain. Then move to the next two points

Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

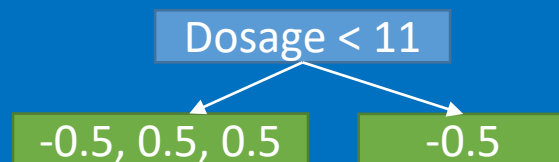
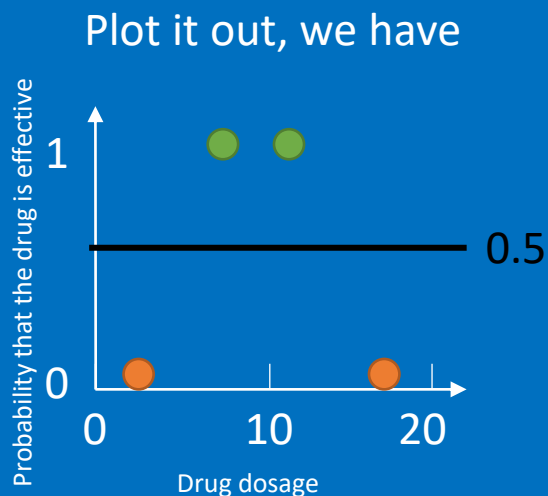
Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

3.1 All the tree starts from a leaf, with all the residuals

-0.5, 0.5, 0.5, -0.5 0 *Similarity score*

3.2 Calculate the Similarity score

3.3 Further split the tree



After gone through all the thresholds, we can tell that this tree gives the highest “Gain”

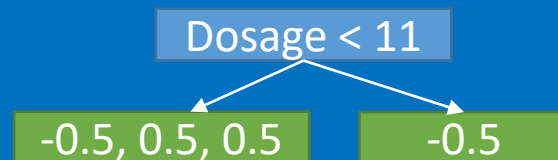
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

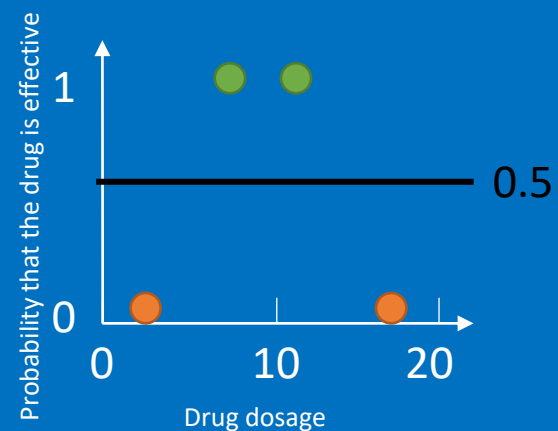
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Plot it out, we have



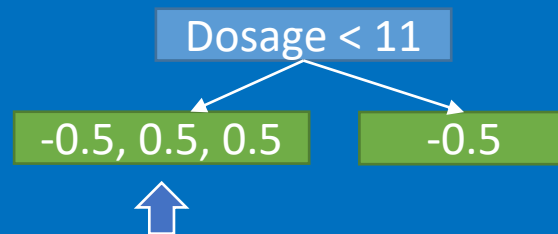
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

For example, the probability of the dosage is effective: 0.5

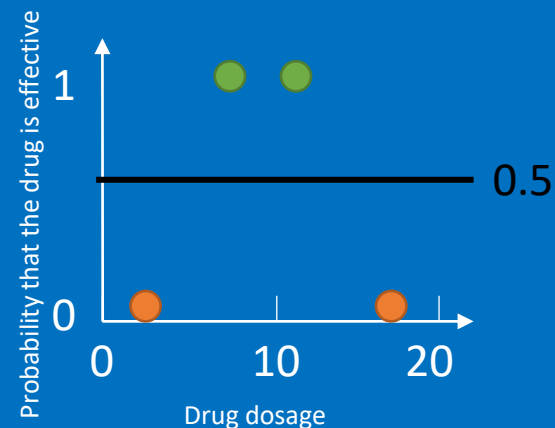
Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



We can continue split these residuals (following the same method)...

Plot it out, we have



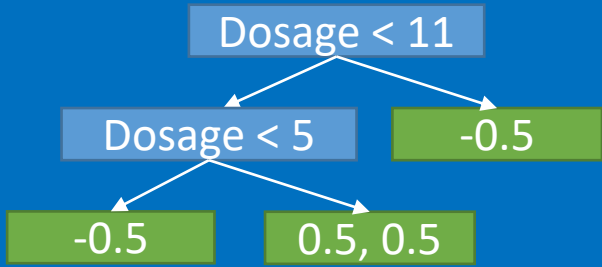
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

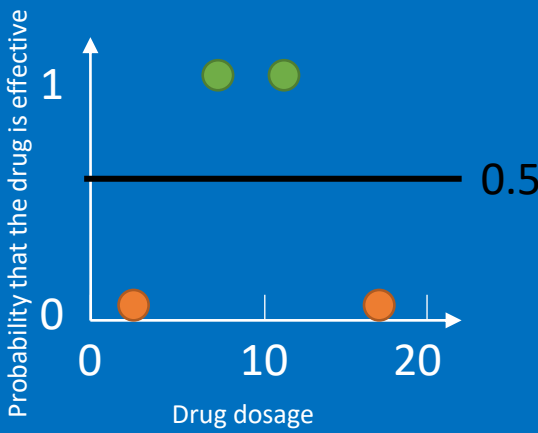
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Plot it out, we have



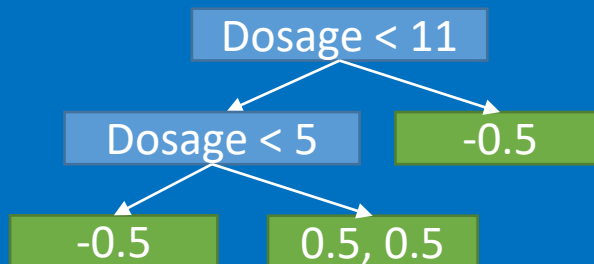
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

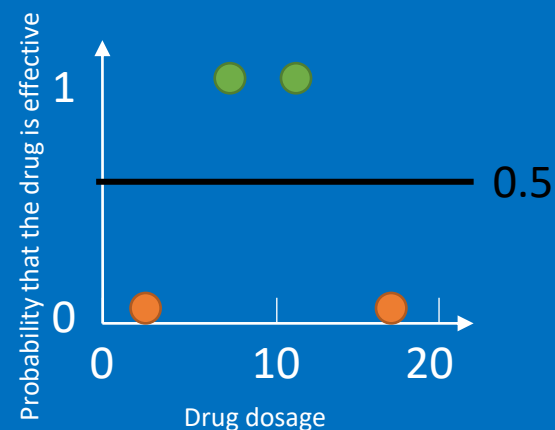
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Plot it out, we have



Note that for a xgboost tree, there are two limitations we can implement:

- How many levels of the tree can grow (like any other tree methods)
- How many residuals in each leaf: this is defined as **cover**

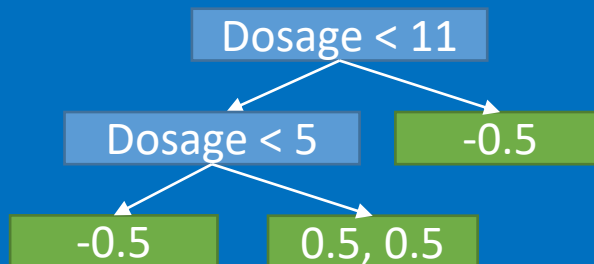
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

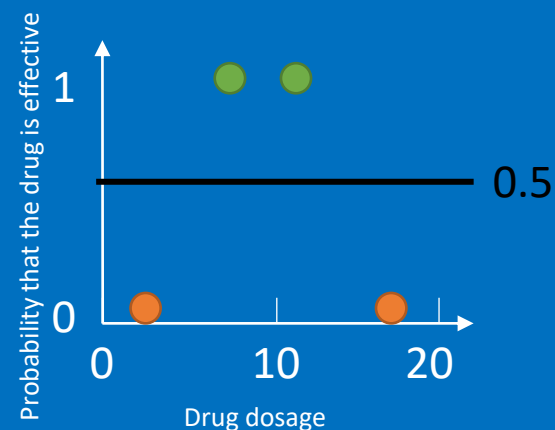
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Plot it out, we have



Note that for a xgboost tree, there are two limitations we can implement:

- How many levels of the tree can grow (like any other tree methods)
- How many residuals in each leaf: this is defined as **cover**

cover, by default, is defined as

- For regression $cover = number\ of\ residuals$
- For classification $cover = \sum [previous\ prob_i \times (1 - previous\ prob_i)]$

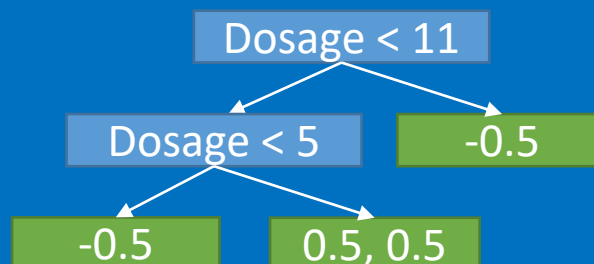
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

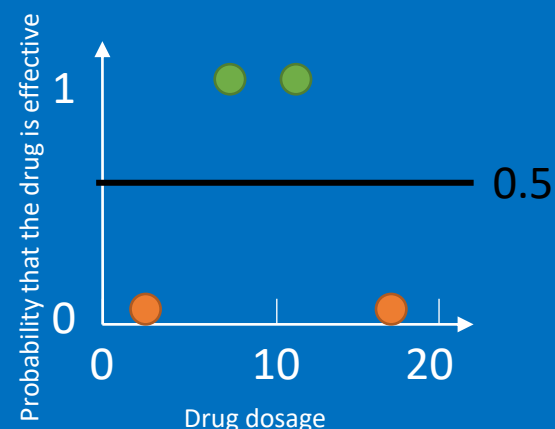
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Plot it out, we have



Note that for a xgboost tree, there are two limitations we can implement:

- How many levels of the tree can grow (like any other tree methods)
- How many residuals in each leaf: this is defined as **cover**

cover, by default, is defined as

- For regression $cover = number\ of\ residuals$
- For classification $cover = \sum [previous\ prob_i \times (1 - previous\ prob_i)]$

So for regression, if we don't set cover by ourself, we always can just have one residual in one leaf

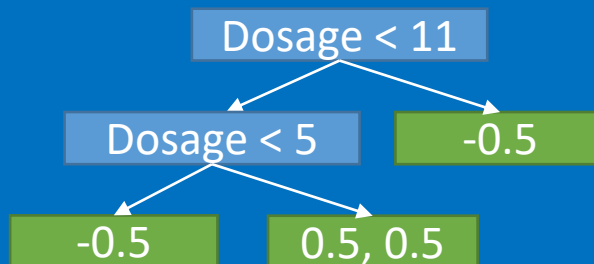
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

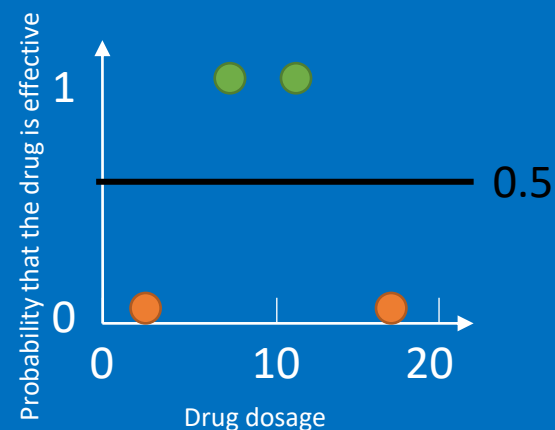
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Plot it out, we have



Note that for a xgboost tree, there are two limitations we can implement:

- How many levels of the tree can grow (like any other tree methods)
- How many residuals in each leaf: this is defined as **cover**

cover, by default, is defined as

- For regression $cover = number\ of\ residuals$
- For classification $cover = \sum [previous\ prob_i \times (1 - previous\ prob_i)]$

So for regression, if we don't set cover by ourself, we always can just have one residual in one leaf

For classification, it's more complicated

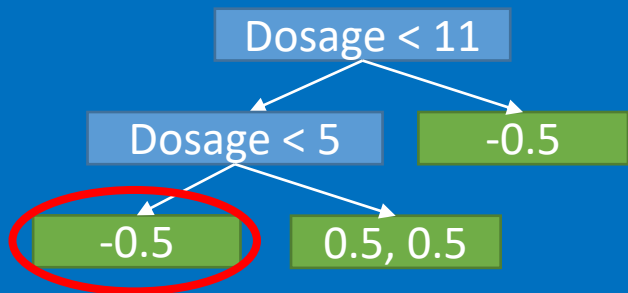
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

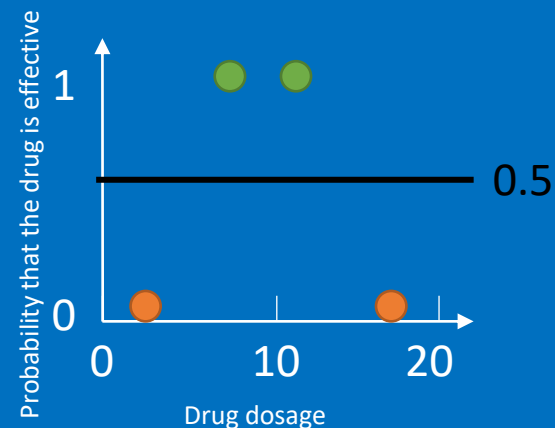
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Plot it out, we have



Note that for a xgboost tree, there are two limitations we can implement:

- How many levels of the tree can grow (like any other tree methods)
- How many residuals in each leaf: this is defined as **cover**

cover, by default, is defined as

- For regression $cover = number\ of\ residuals$
- For classification $cover = \sum [previous\ prob_i \times (1 - previous\ prob_i)]$

So for regression, if we don't set cover by ourself, we always can just have one residual in one leaf

For classification, it's more complicated

For example, the cover for this leaf is

$$cover = 0.5 \times (1 - 0.5) = 0.25$$

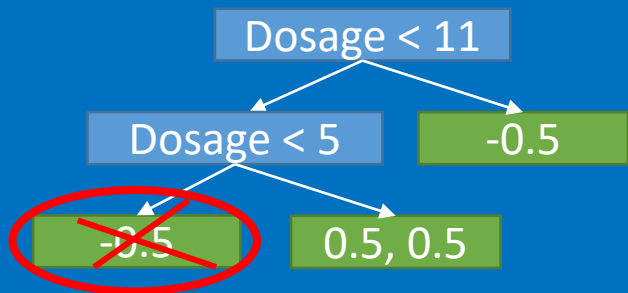
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

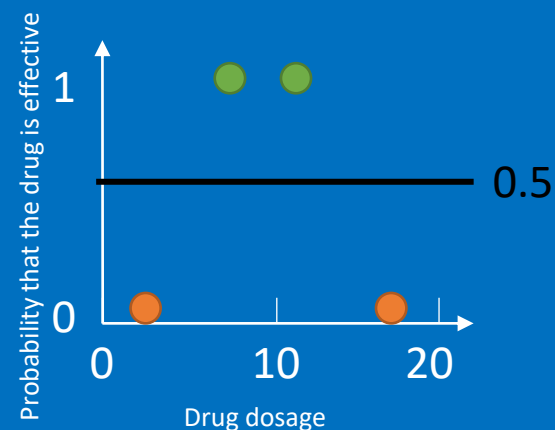
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Plot it out, we have



Note that for a xgboost tree, there are two limitations we can implement:

- How many levels of the tree can grow (like any other tree methods)
- How many residuals in each leaf: this is defined as **cover**

cover, by default, is defined as

- For regression $cover = number\ of\ residuals$
- For classification $cover = \sum [previous\ prob_i \times (1 - previous\ prob_i)]$

So for regression, if we don't set cover by ourself, we always can just have one residual in one leaf

For classification, it's more complicated

For example, the cover for this leaf is

$$cover = 0.5 \times (1 - 0.5) = 0.25$$

So this leaf will be removed

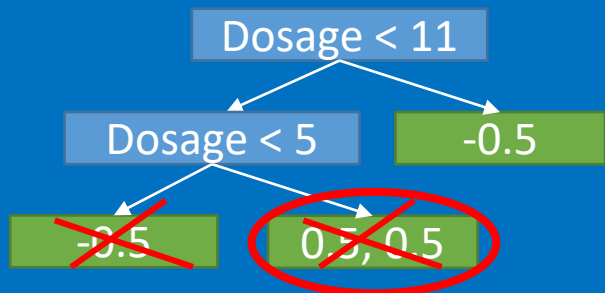
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

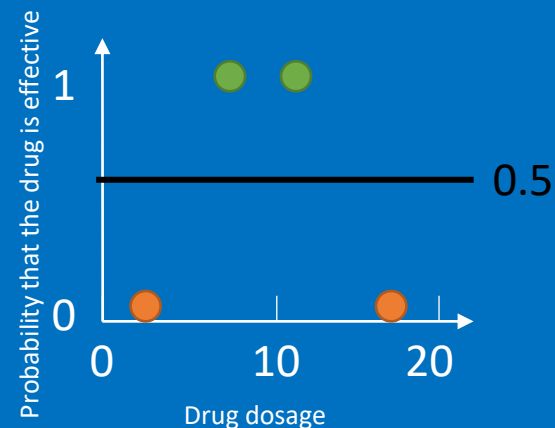
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Plot it out, we have



Note that for a xgboost tree, there are two limitations we can implement:

- How many levels of the tree can grow (like any other tree methods)
- How many residuals in each leaf: this is defined as **cover**

cover, by default, is defined as

- For regression $cover = number\ of\ residuals$
- For classification $cover = \sum [previous\ prob_i \times (1 - previous\ prob_i)]$

So for regression, if we don't set cover by ourself, we always can just have one residual in one leaf

For classification, it's more complicated

For example, the cover for this leaf is

$$cover = 0.5 \times (1 - 0.5) + 0.5 \times (1 - 0.5) = 0.5$$

So this leaf will be removed, too

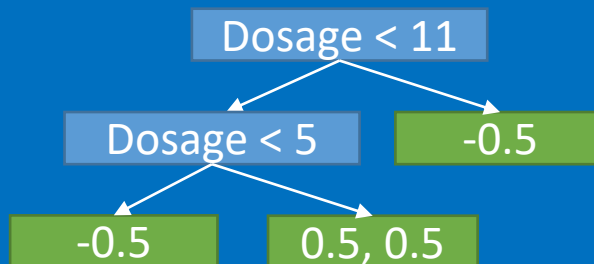
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

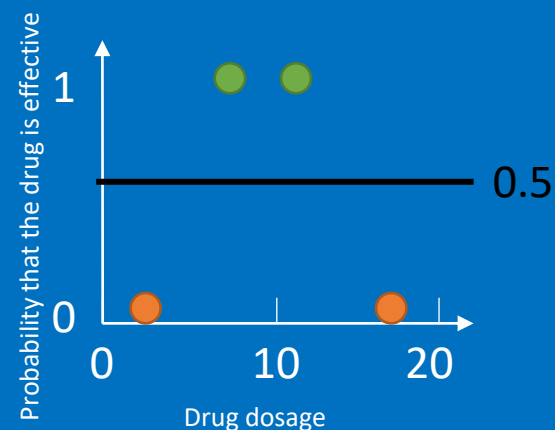
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Plot it out, we have



Note that for a xgboost tree, there are two limitations we can implement:

- How many levels of the tree can grow (like any other tree methods)
- How many residuals in each leaf: this is defined as **cover**

cover, by default, is defined as

- For regression $cover = number\ of\ residuals$
- For classification $cover = \sum [previous\ prob_i \times (1 - previous\ prob_i)]$

So for regression, if we don't set cover by ourself, we always can just have one residual in one leaf

So as you can see, by default, too many leaf will be removed in this example, if we use the default cover. In order to make this example working, let's just set **cover=0**

In some packages, cover is called min_child_weight

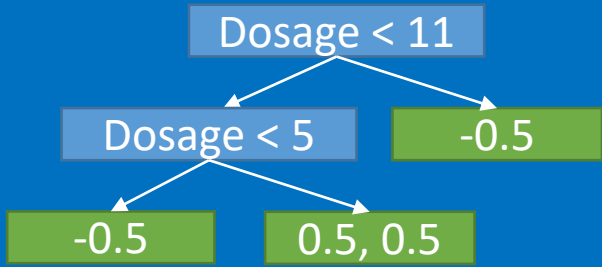
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

For example, the probability of the dosage is effective: 0.5

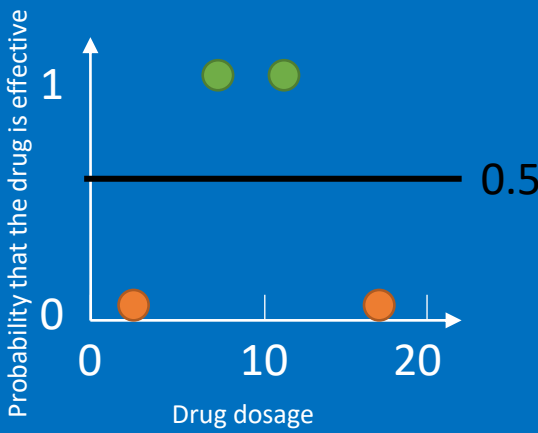
Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Step 4: Prune the tree (like the xgboost for regression)

Plot it out, we have



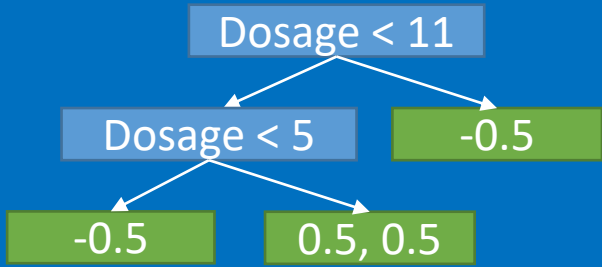
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

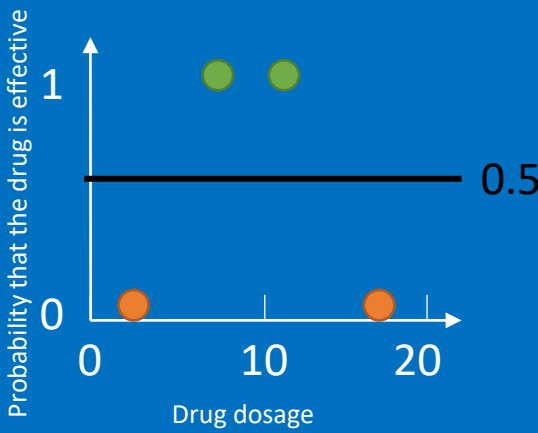
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Plot it out, we have



Step 4: Prune the tree (like the xgboost for regression)

Step 5: Determine the output value

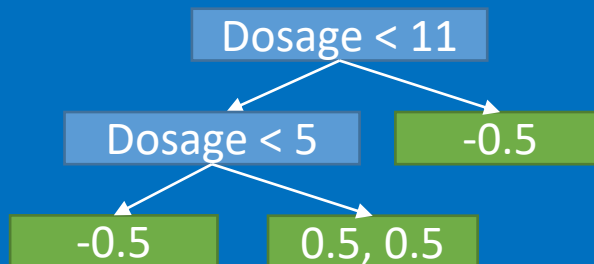
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

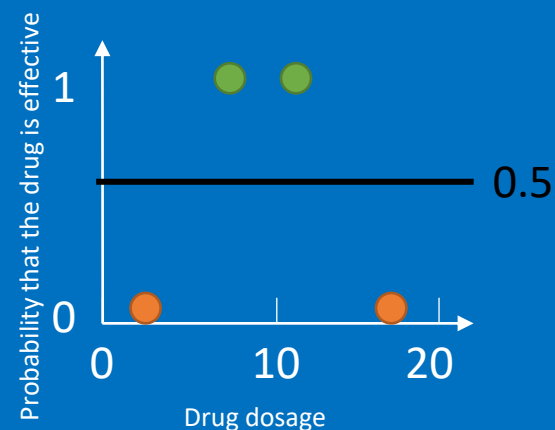
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Plot it out, we have



Step 4: Prune the tree (like the xgboost for regression)

Step 5: Determine the output value

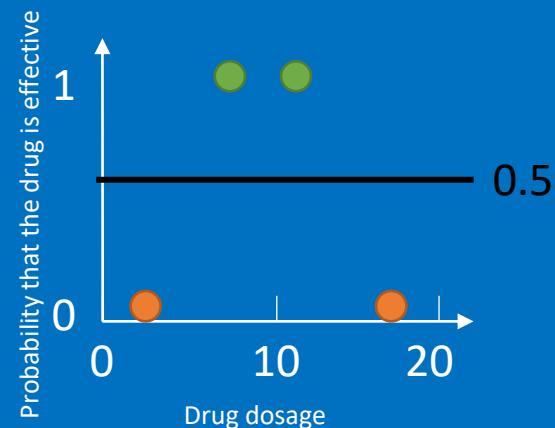
Note that for regression xgboost:

$$output = \frac{sum\ of\ residuals}{number\ of\ residuals + \lambda}$$

$$output = \frac{\sum residuals}{\sum previous\ prob_i \times (1 - previous\ prob_i) + \lambda}$$

Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Plot it out, we have

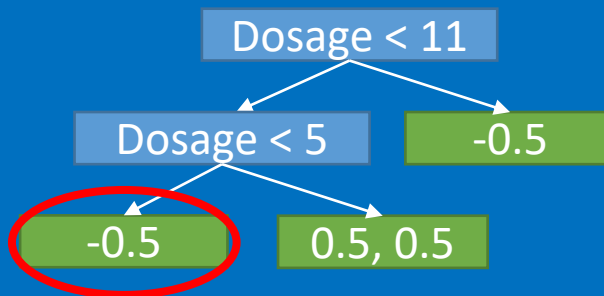


Step 1: make the initial prediction

For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Step 4: Prune the tree (like the xgboost for regression)

Step 5: Determine the output value

$$output = \frac{\sum residuals}{\sum previous prob_i \times (1 - previous prob_i) + \lambda}$$

So for this leaf, we have (assuming lambda=0):

$$output = \frac{-0.5}{0.5 \times (1 - 0.5) + 0} = -2$$

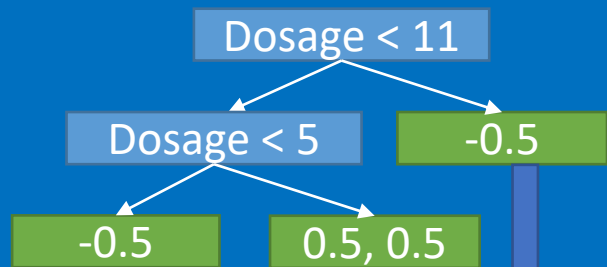
Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

For example, the probability of the dosage is effective: **0.5**

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)



Step 4: Prune the tree (like the xgboost for regression)

Step 5: Determine the output value

$$output = \frac{\sum residuals}{\sum previous prob_i \times (1 - previous prob_i) + \lambda}$$

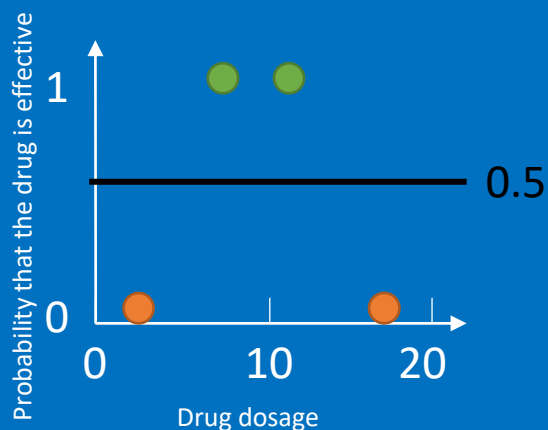
Output

-2

2

-2

Plot it out, we have



Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

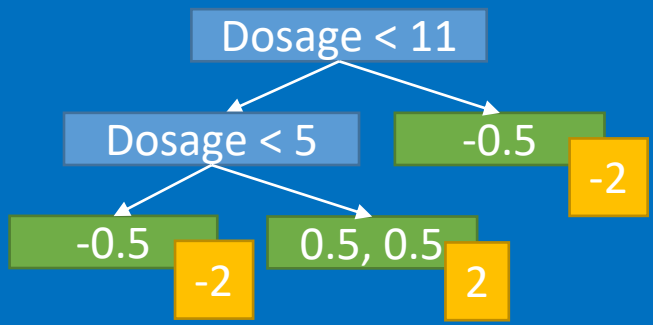
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

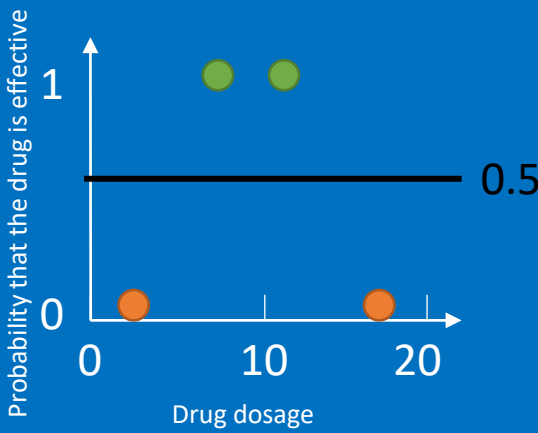
Step 4: Prune the tree (like the xgboost for regression)

Step 5: Determine the output value



So the first tree is completed

Plot it out, we have



Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

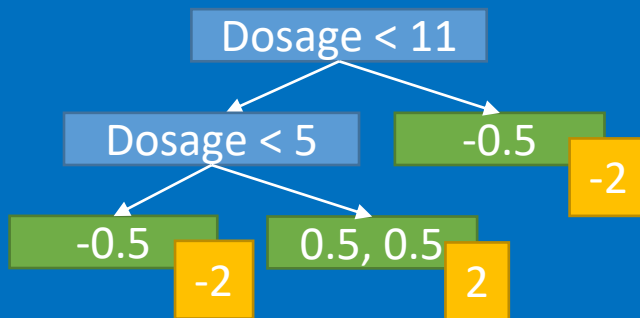
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

Step 4: Prune the tree (like the xgboost for regression)

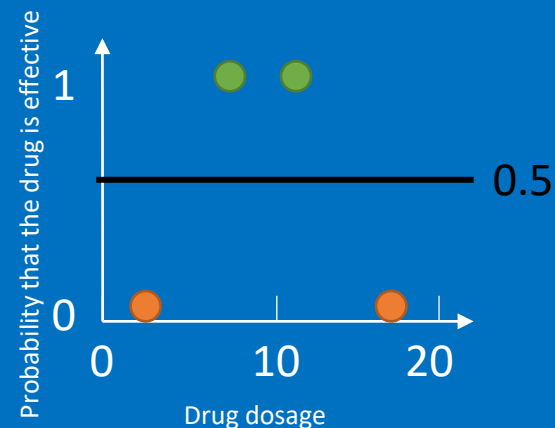
Step 5: Determine the output value



So the first tree is completed

Step 6: Start making predictions

Plot it out, we have



Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

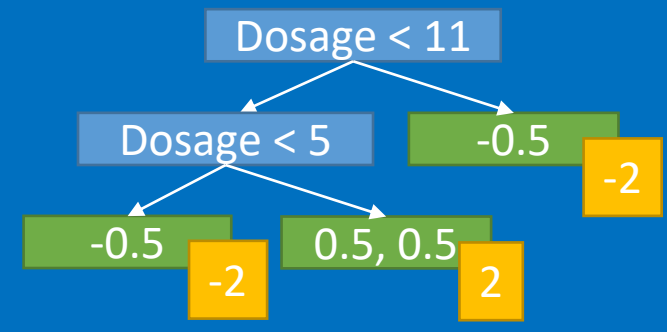
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

Step 4: Prune the tree (like the xgboost for regression)

Step 5: Determine the output value

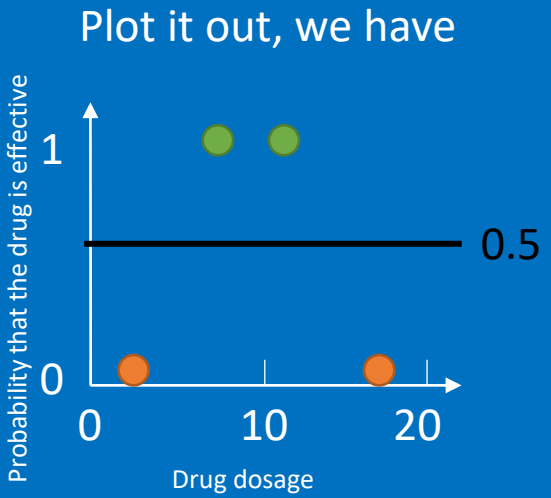


So the first tree is completed

Step 6: Start making predictions

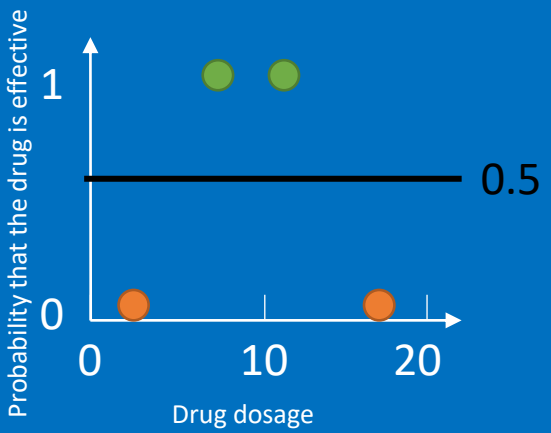
6.1: first we need to convert the previous prob to log(odds)

The equation is $\log(odds) = \log\left(\frac{p}{1-p}\right)$



Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Plot it out, we have



Step 1: make the initial prediction

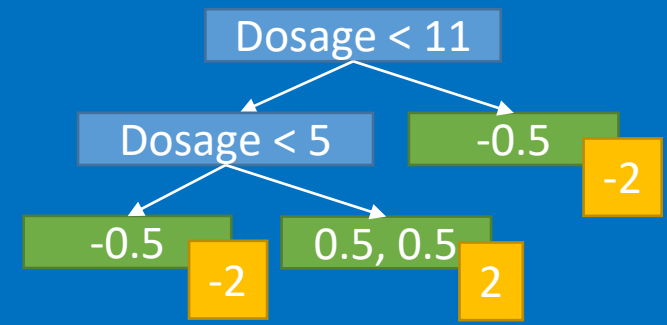
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

Step 4: Prune the tree (like the xgboost for regression)

Step 5: Determine the output value



So the first tree is completed

Step 6: Start making predictions

6.1: first we need to convert the previous prob to log(odds)

The equation is $\log(odds) = \log\left(\frac{p}{1-p}\right)$

The previous probability is 0.5, so log(odds) = 0

Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

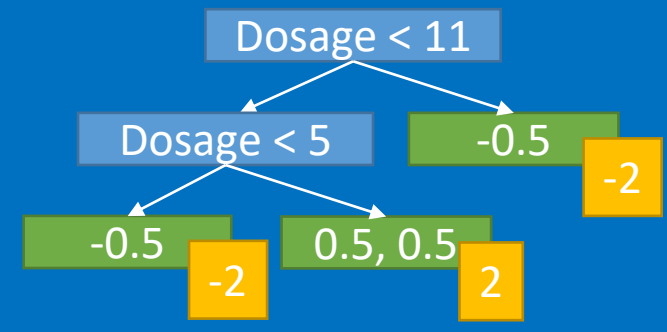
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

Step 4: Prune the tree (like the xgboost for regression)

Step 5: Determine the output value



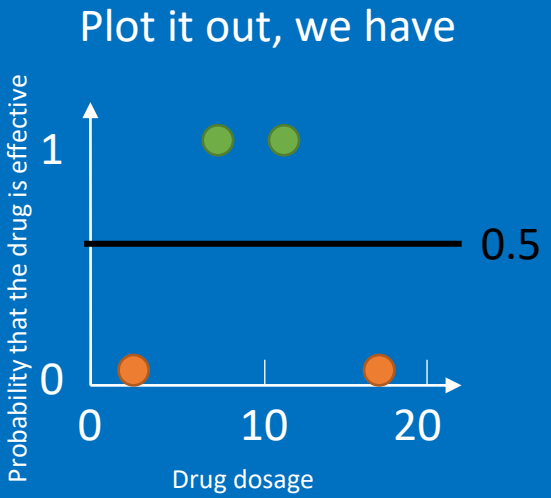
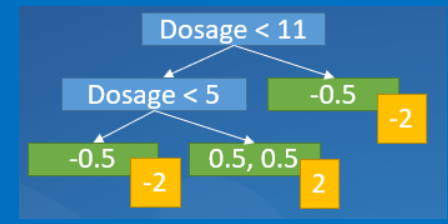
So the first tree is completed

Step 6: Start making predictions

6.1: first we need to convert the previous prob to $\log(\text{odds})=0$

6.2: making prediction with

$$0.5 + 0.3 \times \log(\text{odds})=0$$



Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

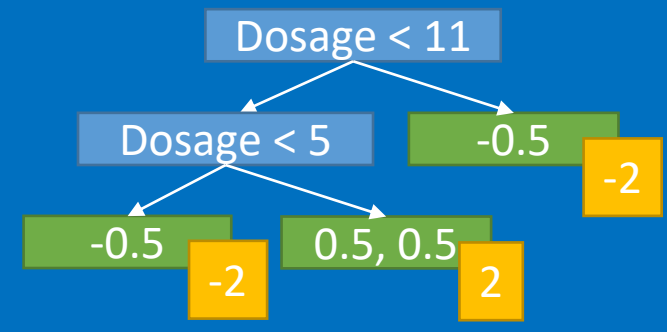
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

Step 4: Prune the tree (like the xgboost for regression)

Step 5: Determine the output value



So the first tree is completed

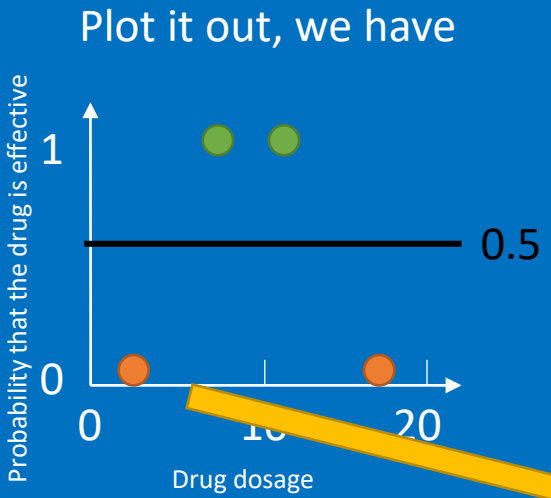
Step 6: Start making predictions

6.1: first we need to convert the previous prob to $\log(\text{odds})=0$

6.2: making prediction with



For example, for the first sample, we can have the $\log(\text{odds})$ prediction as $0+0.3 \times (-2)=-0.6$



Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

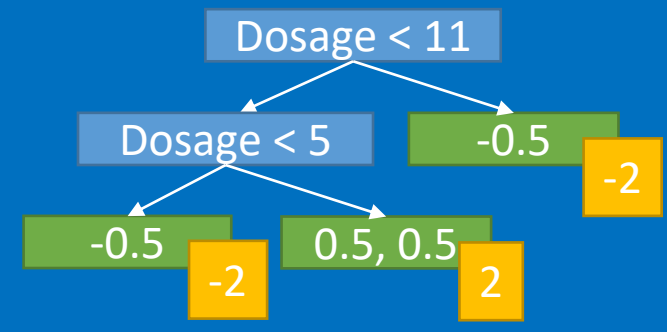
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

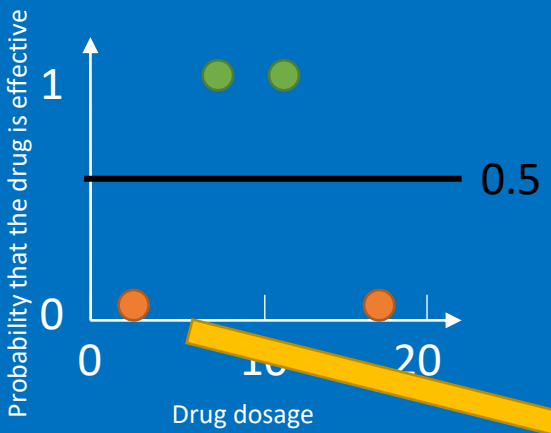
Step 4: Prune the tree (like the xgboost for regression)

Step 5: Determine the output value



So the first tree is completed

Plot it out, we have



Step 6: Start making predictions

6.1: first we need to convert the previous prob to $\log(\text{odds})=0$

6.2: making prediction with



Then we need to convert $\log(\text{odds})$ back to probability as:

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

For example, for the first sample, we can have the $\log(\text{odds})$ prediction as $0 + 0.3 \times (-2) = -0.6$

Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

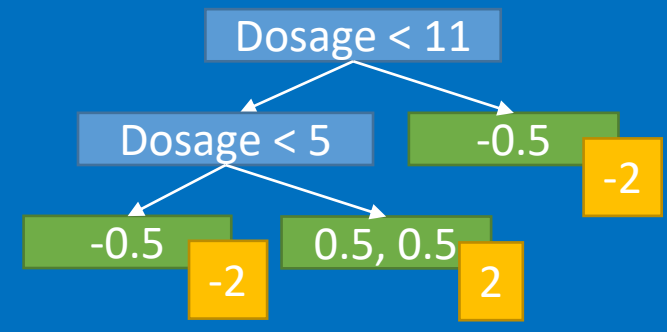
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

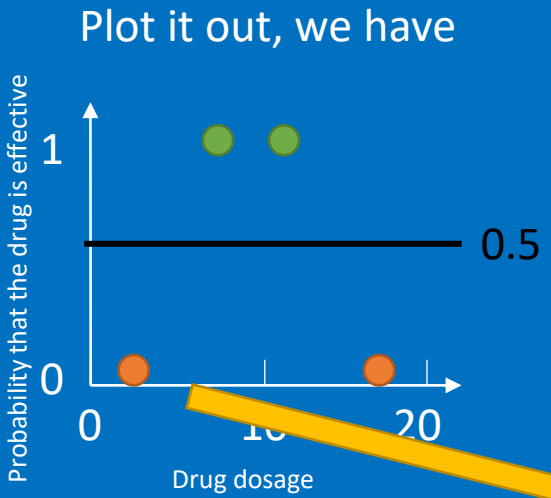
Step 4: Prune the tree (like the xgboost for regression)

Step 5: Determine the output value



So the first tree is completed

So for the first sample, the prediction of probability is

$$p = \frac{e^{\log(odds)}}{1 + e^{\log(odds)}} = \frac{e^{-0.6}}{1 + e^{-0.6}} = 0.35$$


Step 6: Start making predictions

6.1: first we need to convert the previous prob to log(odds)=0

6.2: making prediction with



Then we need to convert log(odds) back to probability as:

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

For example, for the first sample, we can have the log(odds) prediction as 0+0.3x(-2)=-0.6

Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

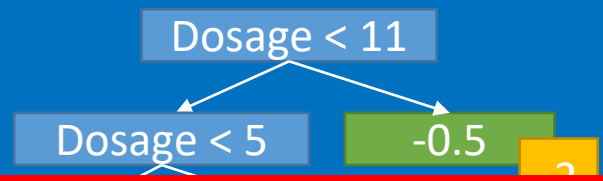
For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

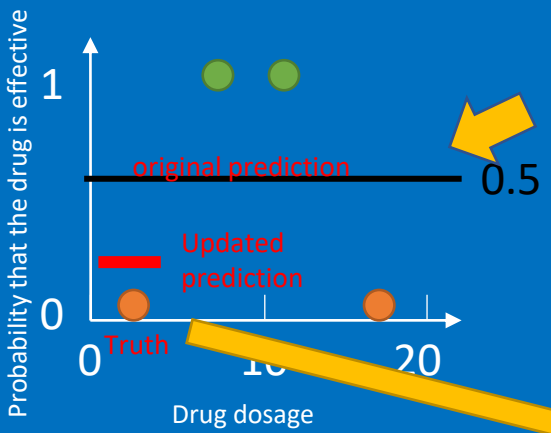
Step 4: Prune the tree (like the xgboost for regression)

Step 5: Determine the output value



So the first tree is completed

Plot it out, we have



The original prediction is 0.5, and the new prediction is 0.35 (get reduced from the original), and it is getting closer to the truth

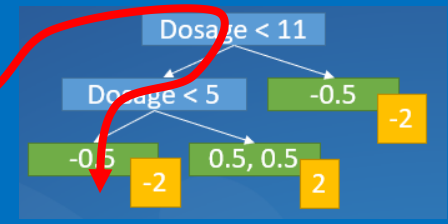
So for the first sample, the prediction of probability is

$$p = \frac{e^{\log(odds)}}{1 + e^{\log(odds)}} = \frac{e^{-0.6}}{1 + e^{-0.6}} = 0.35$$

6.1: first we need to convert the previous prob to $\log(odds)=0$

6.2: making prediction with

$$0.5 + 0.3 \times \log(odds)=0$$



Then we need to convert $\log(odds)$ back to probability as:

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

For example, for the first sample, we can have the $\log(odds)$ prediction as $0+0.3 \times (-2)=-0.6$

Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

For example, the probability of the dosage is effective: 0.5

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

Step 4: Prune the tree (like the xgboost for regression)

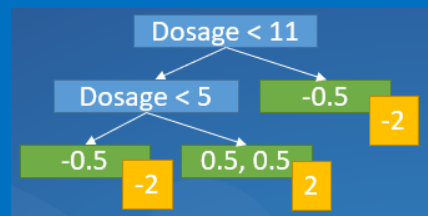
Step 5: Determine the output value

Step 6: Start making predictions

6.1: first we need to convert the previous prob to $\log(\text{odds})=0$

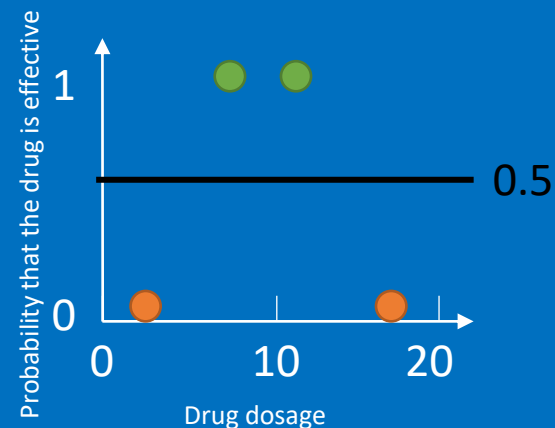
6.2: making prediction with

$$0.5 + 0.3 \times \log(\text{odds})$$



We repeat the process from Step 1 to Step 6 but with updated prediction, every time we get reduced residuals, until the residuals do not change much anymore ...

Plot it out, we have



Drug dosage	Drug usefulness	Residuals
5	No	-0.5
8	Yes	0.5
11	Yes	0.5
18	No	-0.5

Step 1: make the initial prediction

For example, the probability of the dosage is effective: **0.5**

Step 2: we get the residuals for all the samples

Step 3: grow a xgboost tree to the residuals (Similar method used for regression)

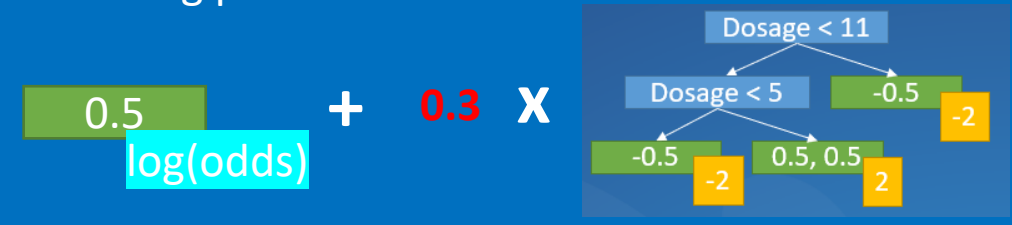
Step 4: Prune the tree (like the xgboost for regression)

Step 5: Determine the output value

Step 6: Start making predictions

6.1: first we need to convert the previous prob to $\log(\text{odds})=0$

6.2: making prediction with



We repeat the process from Step 1 to Step 6 but with updated prediction, every time we get reduced residuals, until the residuals do not change much anymore ...

So the final xgboost tree for classification will be something like this:

