Gradient boosting: Distance to target

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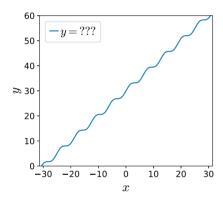
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Contents

An introduction to additive modeling An introduction to boosted regression The intuition behind gradient boosting Gradient boosting regression by example Measuring model performance Choosing hyper-parameters GBM algorithm to minimize L2 loss

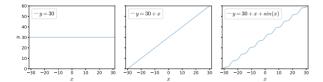
An introduction to additive modeling

Before we get into boosting, let's look at an example of what mathematicians call *additive modeling* because it is the foundation of boosting. The idea is quite simple: we are going to add a bunch of simple terms together to create a more complicated expression. In the machine learning world, that expression (function) represents a model mapping some observation's feature, x, to a scalar target value, y. It's a useful technique because we can often conjure up the simple terms more easily than cracking the overall function in one go. Consider the following curve that shows y as some unknown but nontrivial function of x.



Let's assume that the function is composed of several simple terms and try to guess what they are. After adding each term, we can reassess the situation to help us figure out the next term to add by considering the difference between the current combined function and the desired target function.

Our first approximation might be the horizontal line y=30 because we can see that the y-intercept (at x=0) is 30. Check out the first plot in the following graph sequence. Naturally, the slope is wrong so we should add in the 45 degree line y=x (with slope 60/60=1) that runs through the squiggly target function, which gives us the second graph. It turns out that the squiggly bit comes from our friend the sine function so we can add that term, which leads us to the final plot matching our target function:



Decomposing a complicated function into simpler subfunctions is nothing more than the divide and conquer strategy that we programmers use all the time. In this case, we are dividing a potentially very complicated function into smaller, more manageable bits. For example, let's call our target function F(x) then we have y = F(x) = 30 + x + sin(x) and can abstract away the individual terms, also as functions, giving us the addition of three subfunctions:

$$F(x) = f_1(x) + f_2(x) + f_3(x)$$

where:

$$f_1(x) = 30$$

$$f_2(x) = x$$

$$f_3(x) = \sin(x)$$

More generally, mathematicians describe the decomposition of a function into the addition of M subfunctions like this:

$$F_M(\mathbf{x}) = f_1(\mathbf{x}) + ... + f_M(\mathbf{x}) = \sum_{m=1}^{M} f_m(\mathbf{x})$$

The sigma \sum notation is a for-loop that iterates m from 1 to M, accumulating the sum of the subfunction, f_m , results.

In the machine learning world, we're given a set of (x, y) data points rather than a continuous function, as we have here. The goal is to create a function that draws a nice curve through the data points. We call that function a *model* and it maps x to y, thus, making predictions given some unknown x. Adding up a bunch of subfunctions to create a composite function that models some data points is then called *additive modeling*. Gradient boosting machines use additive modeling to gradually nudge an approximate model towards a really good model, by adding simple submodels to a composite model.

An introduction to boosted regression

<u>Boosting</u> is a loosely-defined strategy that combines multiple simple models into a single composite model. The idea is that, as we introduce more simple models, the overall model becomes a stronger and stronger predictor. In boosting terminology, the simple models are called *weak models* or *weak learners*.

In the context of regression, we make numerical predictions, such as rent prices, based upon information, such as square footage, about an entity (an observation). To keep things simple in this article, we'll work with a single feature per entity but, in general, each observation has a vector of features; let's call it \mathbf{x} . Given \mathbf{x} , we'd like to learn scalar target value y for a bunch of (\mathbf{x}_i, y_i) pairs. Training a regression model is a matter of fitting a function through the data points (\mathbf{x}_i, y_i) as best we can. Given a single feature vector \mathbf{x} and scalar target value y for a single observation, we can express a composite model that predicts (approximates) \hat{y} as the addition of M weak models $f_m(\mathbf{x})$:

$$\hat{\mathbf{y}} = \sum_{m=1}^{M} f_m(\mathbf{x})$$

Mathematicians represent both the weak and composite models as functions, but in practice the models can be anything including knearest-neighbors or regression trees. Since everyone uses trees for boosting, we'll focus on implementations that use regression trees for weak models, which also happens to greatly simplify the mathematics. Later, we'll stack N feature vectors as rows in a matrix, $X = [\mathbf{x}_1, \mathbf{x}_2, ..., \mathbf{x}_N]$, and N targets into a vector, $\mathbf{y} = [y_1, y_2, ..., y_N]$ for N observations.

It's often the case that an additive model can build the individual $f_m(\mathbf{x})$ terms independently and in parallel, but that's not the case for boosting. Boosting constructs and adds weak models in a stage-wise fashion, one after the other, each one chosen to improve the overall model performance. The boosting strategy is greedy in the sense that choosing $f_m(\mathbf{x})$ never alters previous functions. We could choose to stop adding weak models when $\hat{y} = F_M(\mathbf{x})$'s performance is good enough or when $f_m(\mathbf{x})$ doesn't add anything. In practice, we choose the number of stages, M, as a hyper-parameter of the overall model. Allowing M to grow arbitrarily increases the risk of overfitting.

Because greedy strategies choose one weak model at a time, you will often see boosting models expressed using this equivalent, recursive formulation:

$$F_m(\mathbf{x}) = F_{m-1}(\mathbf{x}) + f_m(\mathbf{x})$$

That says we should tweak the previous model with $f_m(\mathbf{x})$ to get the next model.

Boosting itself does not specify how to choose the weak learners. Boosting does not even specify the form of the $f_m(\mathbf{x})$ models, but the form of the weak model dictates the form of the meta-model. For example, if all weak models are linear models, then the resulting meta-model is a simple linear model. If we use tiny regression trees as the weak models, the result is a forest of trees whose predictions are added together.

Let's see if we can design a strategy for picking weak models to create our own boosting algorithm for a single observation. Then, we can extend it to work on the multiple observations we'd encounter in practice.

The intuition behind gradient boosting

To construct a boosted regression model, let's start by creating a crappy model, $f_0(\mathbf{x})$, that predicts an initial approximation of y given feature vector \mathbf{x} . Then, let's gradually nudge the overall $F_m(\mathbf{x})$ model towards the known target value y by adding one or more tweaks, $\Delta_m(\mathbf{x})$:

$$\hat{y} = f_0(\mathbf{x}) + \Delta_1(\mathbf{x}) + \Delta_2(\mathbf{x}) + \dots + \Delta_M(\mathbf{x})$$

$$= f_0(\mathbf{x}) + \sum_{m=1}^{M} \Delta_m(\mathbf{x})$$

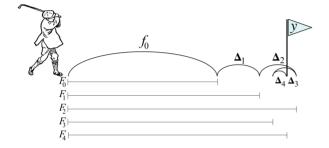
$$= F_M(\mathbf{x})$$

Or, using a recurrence relation, let:

$$F_0(\mathbf{x}) = f_0(\mathbf{x})$$

 $F_m(\mathbf{x}) = F_{m-1}(\mathbf{x}) + \Delta_m(\mathbf{x})$

It might be helpful to think of this boosting approach as a golfer initially whacking a golf ball towards the hole at y but only getting as far as $f_0(\mathbf{x})$. The golfer then repeatedly taps the ball more softly, working the ball towards the hole, after reassessing direction and distance to the hole at each stage. The following diagram illustrates 5 strokes getting to the hole, y, including two strokes, Δ_2 and Δ_3 , that overshoot the hole. (Golfer clipart from http://etc.usf.edu/clipart/)



After the initial stroke, the golfer determines the appropriate nudge by computing the difference between y and the first approximation, $y-f_0(\mathbf{x})$. (We can let \mathbf{x} be the hole number 1-18, but it doesn't really matter since we're only working with one observation for illustration purposes.) This difference is usually called the *residual* or *residual vector*, but it's helpful for gradient boosting to think of this as the vector pointing from the current prediction, $\hat{y}=F_m(\mathbf{x})$, to the true y. (In the <u>second article</u>, we will look at just the sign of the direction, not magnitude; we'll call that the *sign vector* to distinguish from the residual vector.) Using the residual vector as our nudge, means training $\Delta_m(\mathbf{x})$ on value $y-F_{m-1}(\mathbf{x})$ for our base weak models. As with any machine learning model, our Δ_m models will not have perfect recall and precision, so we should expect Δ_m to give a noisy prediction instead of exactly $y-F_{m-1}(\mathbf{x})$.

As an example, let's say that the hole is at y=100 yards, $f_0(\mathbf{x})=70$. Manually boosting, we might see a sequence like the following, depending on the imprecise Δ_m strokes made by the golfer:

$_{m}^{\mathbf{Stage}}$	$egin{array}{l} { m Boosted} \\ { m Model} \end{array}$	$egin{aligned} \operatorname{Model} \ \operatorname{Output} \hat{y} \end{aligned}$	Train Δ_m on $y - F_{m-1}$	$egin{aligned} ext{Noisy} \ ext{Prediction } \Delta_m \end{aligned}$
0	F_0	70		
1	$F_1 = F_0 + \Delta_1$	70+15=85	100 - 70 = 30	$\Delta_1 = 15$
2	$F_2 = F_1 + \Delta_2$	85+20=105	100-85=15	$\Delta_2 = 20$
3	$F_3 = F_2 + \Delta_3$	105 - 10 = 95	100 - 105 = -5	$\Delta_3 =$ -10
4	$F_4 = F_3 + \Delta_4$	95+5=100	100-95=5	$\Delta_4 = 5$

A GBM implementation would also support a so-called learning rate, η , that speeds up or slows down the overall approach of \hat{y} to y, which helps to reduce the likelihood of overfitting.

To show how flexible this technique is, consider training the weak models on just the direction of y, rather than the magnitude and direction of y. In other words, we would train the $\Delta_m(\mathbf{x})$ on $sign(y-F_{m-1}(\mathbf{x}))$, not $y-F_{m-1}(\mathbf{x})$. The sign(z) (or sgn(z)) function expresses the direction as one of $\{-1,0,+1\}$, but both $sign(y-\hat{y})$ and $y-\hat{y}$ point us in suitable directions. We'll see in <u>Gradient boosting performs gradient descent</u> that training on the residual vector optimizes the overall model for the squared error loss function and training on the sign vector optimizes the absolute error loss function.

If you understand this golfer example, then you understand the key intuition behind boosting for regression, at least for a single observation. Yup, that's it, but there are several things to reinforce before moving on:

- $\bullet \ \ \, \text{The weak models learn direction } \textbf{vectors} \text{ with direction information, not just magnitudes}.$
- The initial model $f_0(\mathbf{x})$ is trying to learn target y given \mathbf{x} , but the $\Delta_m(\mathbf{x})$ tweaks are trying to learn direction vectors given \mathbf{x} .
- All weak models, $f_0(\mathbf{x})$ and $\Delta_m(\mathbf{x})$, train on direction vectors that are some function of the original feature vectors, \mathbf{x} .
- Two common direction vector choices are the residual, $y F_{m-1}(\mathbf{x})$, and the sign, $sign(y F_{m-1}(\mathbf{x}))$.

Let's walk through a concrete example to see what gradient boosting looks like on more than one observation.

Gradient boosting regression by example

Imagine that we have square footage data on five apartments and their rent prices in dollars per month as our training data:

sqfeet	rent
750	1160
800	1200
850	1280
900	1450
950	2000

where row i is an observation with one-dimensional feature vector \mathbf{x}_i (bold \mathbf{x}) and target scalar value y_i . Matrix $X = [\mathbf{x}_1, \mathbf{x}_2, ..., \mathbf{x}_n]$ holds all feature vectors and \mathbf{y} (bold \mathbf{y}) is the entire rent vector $\mathbf{y} = [y_1, y_2, ..., y_n]$. $F_m(\mathbf{x}_i)$ yields a predicted value but $F_m(X)$ yields a predicted target vector, one value for each \mathbf{x}_i .

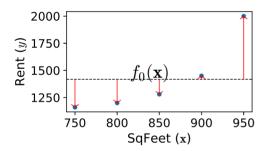
From this data, we'd like to build a GBM to predict rent price given square footage. To move towards \mathbf{y} from any $\hat{\mathbf{y}}$, we need a direction vector. Let's start with $\mathbf{y} - \hat{\mathbf{y}}$ and then, in <u>Heading in the right direction</u>, we'll see how GBM works for $sign(\mathbf{y} - \hat{\mathbf{y}})$.

Let's use the mean (average) of the rent prices as our initial model: $F_0(\mathbf{x}_i) = f_0(\mathbf{x}_i) = 1418$ for all i: We use the mean because that is the single value that minimizes the mean squared error between it and the y_i values. (We'll seen shortly that GBMs whose weak models ar trained on residual vectors optimize the mean squared error.) Once we have F_0 , we compute $\mathbf{y} - F_0$, the residual between the target and the previous estimate:

\mathbf{sqfeet}	rent	F_0	$\mathbf{y} - F_0$
750	1160	1418	-258
800	1200	1418	-218
850	1280	1418	-138
900	1450	1418	32
950	2000	1418	582

(Many articles and Friedman's original paper call the $\mathbf{y}-F_m$ values pseudo-responses and use notation $\tilde{\mathbf{y}}$.)

The last column shows not only the direction but the magnitude of the difference between where we are, $F_0(X)$, and where we want to go, y. The red vectors in the following diagram are a visualization of the residual vectors from our initial model to the rent target values.



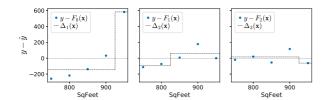
Next, we train a weak model, Δ_1 , to predict that residual vector given \mathbf{x}_i for all i observations. A perfect model, Δ_1 , would yield exactly $\mathbf{y}-F_0(X)$, meaning that we'd be done after one step since $F_1(X)$ would be $F_1(X)=F_0(X)+(\mathbf{y}-F_0(X))$, or just \mathbf{y} . Because it imperfectly captures that difference, $F_1(X)$ is still not quite \mathbf{y} , so we need to keep going for a few stages. Let's add learning rate, η , to our recurrence relation:

$$F_m(X) = F_{m-1}(X) + \eta \Delta_m(X)$$

We'll discuss the learning rate below, but for now, please assume that our learning rate is $\eta=1.0$, so $F_1=F_0+\Delta_1$, $F_2=F_1+\Delta_2$, and so on. The following table summarizes the intermediate values of the various key "players":

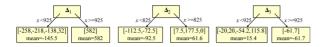
Δ_1	F_1	\mathbf{y} - F_1	Δ_2	F_2	$y - F_2$	Δ_3	F_3
-145.5	1272.5	-112.5	- 92.5	1180	-20	15.4	1195.4
-145.5	1272.5	- 72.5	- 92.5	1180	20	15.4	1195.4
-145.5	1272.5	7.5	61.7	1334.2	- 54.2	15.4	1349.6
-145.5	1272.5	177.5	61.7	1334.2	115.8	15.4	1349.6
582	2000	0	61.7	2061.7	-61.7	-61.7	2000

It helps to keep in mind that we are always training on the residual vector $\mathbf{y}-F_{m-1}$ but get imperfect model Δ_m . The best way to visualize the learning of $\mathbf{y}-F_{m-1}$ residual vectors by weak models, Δ_m , is by looking at the residual vectors and model predictions horizontally on the same scale Y-axis:



The blue dots are the residual vector elements used to train Δ_m weak models, the dashed lines are the predictions made by Δ_m , and the dotted line is the origin at o. Notice how the residual vector elements (blue dots) get smaller as we add more weak models.

The predictions are step functions because we've used a *regression tree stump* as our base weak model with split points 925, 825, and 925. Here are the three stumps implementing our Δ_m weak models:

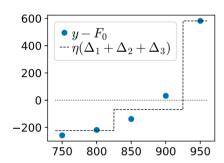


Regression tree stumps

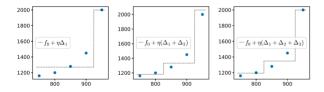
A regression tree stump is a regression tree with a single root and two children that splits on a single (feature) variable, which is what we have here, at a single threshold. (If we had more than a single value in our feature vectors, we'd have to build a taller tree that tested more variables; to avoid overfitting, we don't want very tall trees, however.) If a test feature value is less than the threshold, the model yields the average of the training target samples in the left leaf. If the test feature value is greater than or equal to the threshold, the model yields the average of the training target examples in the right leaf.

The feature value split location is chosen to minimize the variance of the target elements in each group. For example, imagine single-valued features for 5 observations [0,1,4,4,3] and target values [3.1,3,20,22,25]. A good place to split the feature values is between 1 and 4. That means separating target values in the two leaves into very similar groups: [3.1,3] and [20,22,25].

The composite model sums together all of the weak models so let's visualize the sum of the weak models:



If we add all of those weak models to the initial f_0 average model, we see that the full composite model is a very good predictor of the actual rent values:



It's worth pointing out something subtle with the learning rate and the notation used in the graphs: $f_0 + \eta(\Delta_1 + \Delta_2 + \Delta_3)$. That makes it look like the learning rate could be applied all the way at the end as a global learning rate. Mathematically, the formula is correct but it hides the fact that each weak model, Δ_m , is trained on $\mathbf{y} - F_{m-1}(X)$ and $F_{m-1}(X)$ is a function of the learning rate: $F_{m-1}(X) = F_{m-2}(X) + \eta \Delta_{m-1}(X)$. Friedman calls this incremental shrinkage.

Measuring model performance

How good is our model? To answer that, we need a loss or cost function, $L(\mathbf{y}, \hat{\mathbf{y}})$ or $L(y_i, \hat{y}_i)$, that computes the cost of predicting $\hat{\mathbf{y}}$ instead of \mathbf{y} . The loss across all N observations is just the average (or the sum if you want since N is a constant once we start training) of all the individual observation losses:

$$L(\mathbf{y}, F_M(X)) = \frac{1}{N} \sum_{i=1}^{N} L(y_i, F_M(\mathbf{x}_i))$$

The mean squared error (MSE) is the most common, and what we are optimizing in this article:

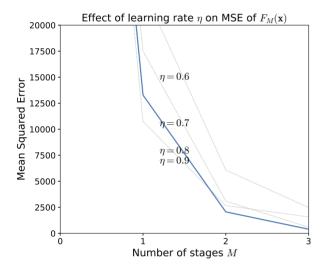
$$L(\mathbf{y}, F_M(X)) = \frac{1}{N} \sum_{i=1}^{N} (y_i - F_M(\mathbf{x}_i))^2$$

In the final article, <u>Gradient boosting performs gradient descent</u> we show that training our Δ_m on the residual vector leads to a minimization of the mean squared error loss function.

Choosing hyper-parameters

We've discussed two GBM hyper-parameters in this article, the number of stages M and the learning rate η . Both affect model accuracy. The more stages we use, the more accurate the model, but the more likely we are to be overfitting. The primary value of the learning rate, or "shrinkage" as some papers call it, is to reduce overfitting of the overall model. As Chen and Guestrin say in $\underline{XGBoost: A Scalable Tree}$ $\underline{Boosting System}$, "shrinkage reduces the influence of each individual tree and leaves space for future trees to improve the model." Friedman recommends a low learning rate like 0.1 and a larger number of stages. In practice, people do a grid search over the hyper-parameter space looking for the best model performance. (Grid search can be very expensive given all of the model construction involved.) For example, see the article by Aarshay Jain: $\underline{Complete Guide to Parameter Tuning in XGBoost}$ or the article by Jason Brownlee called \underline{Tune} $\underline{Learning Rate for Gradient Boosting with XGBoost in Python}$.

The following graph shows how the mean squared error changes as we add more weak models, illustrated with a few different learning rates.



A value of $\eta=0.8$ looks like it reaches the minimum error at the last stage, M=3, so that might be a good starting point for the learning rate.

We stopped at M=3 for purposes of a simple explanation of how boosting works. As we said, practitioners often use a grid search to optimize hyper-parameters, such as M, but one could also keep adding stages until performance stops improving. The risk in that case would be overfitting the model.

As a side note, the idea of using a learning rate to reduce overfitting in models that optimize cost functions to learn, such as deep learning neural networks, is very common. Rather than using a constant learning rate, though, we could start the learning rate out energetically and gradually slow it down as the model approached optimality; this proves very effective in practice.

Ok, let's tie all of this together. A gradient boosting regression model, $F_M(X)$, adds together an initial weak model, $f_0(X)$, that predicts the average ${\bf y}$ value, and the predictions of ${\it M}$ weak models, $\Delta_m(X)$, that nudge $\hat{{\bf y}}$ towards ${\bf y}$. Each $\Delta_m(X)$ is trained on a residual vector that measures the direction and magnitude of the true target ${\bf y}$ from the previous model, ${\bf y}-F_{m-1}(X)$. The new prediction $F_m(X)$ is the addition of the previous model and a nudge, $\Delta_m(X)$, multiplied by a learning rate: $F_m(X)=F_{m-1}(X)+\eta\Delta_m(X)$. Hyper-parameters η and ${\it M}$ are determined by grid search.

If you more-or-less followed this discussion, then congratulations! You understand the key elements of gradient boosting for regression. That's all there is to it. Really. As we'll see in the next article, <u>Gradient boosting: Heading in the right direction</u>, we can use a different direction vector than the residual, but the basic mechanism is the same. Using the sign of the residual rather than the residual vector itself, will have the effect of minimizing a different loss function than mean squared error (it'll minimize mean absolute value).

You might've heard that gradient boosting is very complex mathematically, but that's only if we care about generalizing gradient boostin to work with any loss function (with associated direction vector), rather than the two we discuss in the first two articles of this series (residual and sign vectors). If you want to get funky with the math and see the cool relationship of gradient boosting with gradient descent, check out our last article in the series, <u>Gradient boosting performs gradient descent</u>. Also check out the next article, <u>Gradient</u>

boosting: Heading in the right direction that goes through this example again but this time training weak models on the sign of the residual not the residual vector.

GBM algorithm to minimize L2 loss

For completeness, here is the boosting algorithm, adapted from Friedman's LS Boost and LAD TreeBoost, that optimizes the L_2 loss function using regression tree stumps:

Algorithm: $l2boost(X, \mathbf{y}, M, \eta)$ returns model F_M Let $F_0(X) = \frac{1}{N} \sum_{i=1}^N y_i$, mean of target \mathbf{y} across all observations for m=1 to M do

Let $\mathbf{r}_{m-1} = \mathbf{y} - F_{m-1}(X)$ be the residual direction vector Train regression tree Δ_m on \mathbf{r}_{m-1} , minimizing squared error $F_m(X) = F_{m-1}(X) + \eta \Delta_m(X)$ end
return F_M