Lecture 16: Bagging and Random Forest

CS109A Introduction to Data Science Pavlos Protopapas, Kevin Rader and Chris Tanner



ANNOUNCEMENTS

- Homework 5 (109) due tonight 11:59 pm
- Homework 5 (209) due next Wednesday 11:59 pm, Nov 6
- Homework 6 (109) to be released tomorrow morning
- No 209 for homework 6



ANNOUNCEMENTS

Homework 7 individual







HW1

HW4

HW7



Without OH or ED forum

ANNOUNCEMENTS

• After CS109B, STAT 139 ...





Outline

- Review of Decision Trees
- Bagging
- Out of Bag Error (OOB)
- Variable Importance
- Random Forests



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Question: Can you guess the equation that defines the decision boundary below?

 $-0.8x_1 + x_2 = 0 \implies x_2 = 0.8x_1 \Rightarrow Latitude = 0.8 Lon$





Complicate decision boundaries can not be explained with LogRegression.





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To learn a decision tree model, we take a greedy approach:

- 1. Start with an empty decision tree (undivided feature space)
- Choose the 'optimal' predictor on which to split and choose the 'optimal' threshold value for splitting by applying a *splitting criterion*
- 3. Recurse on each new node until **stopping condition** is met

For classification, we label each region in the model with the label of the class to which the plurality of the points within the region belong.

For regression, we predict with the average of the output values of the training points contained in the region .



The splitting criteria we've examined each minimize a loss function

- **A. For classification**, purity of the regions is a good indicator the performance of the model. Entropy as a splitting criterial minimizes the cross-entropy (greedy).
- **B. For regression**, we want to select a splitting criterion that promotes splits that improves the predictive accuracy of the model as measured by the MSE.



Stopping Conditions



Common simple stopping conditions:

- Don't split a region if all instances in the region belong to the same class.
- Don't split a region if the number of instances in the sub-region will fall below pre-defined threshold (min_samples_leaf).
- Don't split a region if the total number of leaves in the tree will exceed pre-defined threshold.

The appropriate thresholds can be determined by evaluating the model on a held-out data set or, better yet, via cross-validation.



 Compute the gain in purity, information or reduction in entropy of splitting a region R into R₁ and R₂:

$$Gain(R) = \Delta(R) = m(R) - \frac{N_1}{N}m(R_1) - \frac{N_2}{N}m(R_2)$$

where *m* is a metric like the Gini Index or entropy. Don't split if the gain is less than some pre-defined threshold (**min_impurity_decrease**).

In the place of purity gain, we can instead compute accuracy gain for splitting a region *R*

$$Gain(R) = \Delta(R) = MSE(R) - \frac{N_1}{N}MSE(R_1) - \frac{N_2}{N}MSE(R_2)$$

nd stop the tree when the gain is less than some pre-defined threshold



a





















Same issues as with classification trees. Avoid overfitting by pruning or limiting the depth of the tree and using CV.







Reduce the variance





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Hyper-parameters: Depth



Hyper-parameters: Depth

Use train/validation or cross validation to estimate the best depth.

Magic realism: Bootstrap

Magic realism: Bootstrap

Decision trees models are highly interpretable and fast to train, using our greedy learning algorithm.

However, in order to **capture a complex decision boundary** (or to approximate a complex function), we need to use a large tree (since each time we can only make axis aligned splits).

We've seen that large trees have high variance and are prone to overfitting.

For these reasons, in practice, decision tree models often underperforms when compared with other classification or regression methods.

One way to adjust for the high variance of the output of an experiment is to perform the experiment multiple times and then average the results.

The same idea can be applied to high variance models:

- 1. (Bootstrap)we generate multiple samples of training data, via bootstrapping. We train a full decision tree on each sample of data.
- 2. (Aggregate) for a given input, we output the averaged outputs of all the models for that input.

For classification, we return the class that is outputted by the plurality of the models. For regression we return the average of the outputs for each tree.

This method is called **Bagging** (Breiman, 1996), short for, of course, Bootstrap Aggregating.

Note that bagging enjoys the benefits of:

- 1. High expressiveness by using full trees each model is able to approximate complex functions and decision boundaries.
- 2. Low variance averaging the prediction of all the models reduces the variance in the final prediction, assuming that we choose a sufficiently large number of trees.

Bagging (regression)

Bagging (classification)

Question: Do you see any problems?

- Still some overfitting if the trees are too large.
- If trees are too shallow it can still underfits.
- Interpretability:

The **major drawback** of bagging (and other **ensemble methods** that we will study) is that the averaged model is no longer easily interpretable - i.e. one can no longer trace the 'logic' of an output through a series of decisions based on predictor values!

Case of underfitting

Case of underfitting

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Cross Validations

Out-of-Bag Error

Original Data		Bo	otstrap	Sample 1		Decision Tree	e 1	Used and unused d		ata
X	V		X	V				X	V	
X X ₁	y_1		X X ₄	y_4				X X ₁	<i>y</i> ₁	
X ₂	y ₂		X ₁₄	y ₁₄				X ₂	y ₂	
X ₃	У 3		X ₁₁	y ₁₁			_	X ₃	У 3	
X_4	y ₄		<i>X</i> ₂	У ₂			0	X_4	y ₄	
X_5	У ₅		<i>X</i> ₃₅	y ₃₅	V		-	X_5	У ₅	
:	:		÷	:			:	÷		
X _n	Уn		X _k	y _k				X _n	Уn	

					_		_			
Original Data		Bo	Bootstrap Sample 2		2	Decision Tree 2		Used and unused data		
X	Y		X	Y				X	Υ	
X ₁	y_1		X ₅	${\mathcal Y}_5$				X ₁	y_1	
X ₂	y ₂		X ₃	У 3				X ₂	У ₂	
X ₃	У 3		X ₁₂	y ₁₂				X ₃	У 3	
X_4	y ₄		<i>X</i> ₄₃	y ₄₃				X_4	y ₄	
<i>X</i> ₅	У ₅		<i>X</i> ₁	y ₁				X_5	У ₅	
:	:		:	:				:		
						0 1	0 1			
X _n	Уn		X _k	y _k				X _n	Уn	

X	Υ
X ₁	${\mathcal Y}_1$
X ₂	y ₂
X ₃	У 3
:	:
X _i	Уi
:	:
X _n	Уn

Point-wise out-of-bag error

We average the point-wise out-of-bag error over the full training set.

Classification

$$Error_{OOB} = \sum_{i}^{n} e_{i} = \sum_{i}^{n} \mathbb{I}(\hat{y}_{i,pw} \neq y_{i})$$

Regression

$$Error_{OOB} = \sum_{i}^{n} e_{i} = \sum_{i}^{n} (y_{i} - \hat{y}_{i,pw})^{2}$$

Bagging is an example of an **ensemble method**, a method of building a single model by training and aggregating multiple models.

With ensemble methods, we get a new metric for assessing the predictive performance of the model, the **out-of-bag error**.

Given a training set and an ensemble of models, each trained on a bootstrap sample, we compute the **out-of-bag error** of the averaged model by

- 1. For each point in the training set, we average the predicted output for this point over the models whose bootstrap training set excludes this point. We compute the error or squared error of this averaged prediction. Call this the point-wise out-of-bag error.
- 2. We average the point-wise out-of-bag error over the full training set.

Question: Do you see any problems?

- Still some overfitting if the trees are too large.
- If trees are too shallow it can still underfits.
- Interpretability:

The **major drawback** of bagging (and other **ensemble methods** that we will study) is that the averaged model is no longer easily interpretable - i.e. one can no longer trace the 'logic' of an output through a series of decisions based on predictor values!

Variable Importance for Bagging

Bagging improves prediction accuracy at the expense of interpretability.

Calculate the total amount that the MSE (for regression) or Gini index (for classification) is decreased due to splits over a given predictor, averaged over all *B* trees.

100 trees, max_depth=10

In practice, the ensembles of trees in Bagging tend to be highly correlated.

Suppose we have an extremely strong predictor, x_j , in the training set amongst moderate predictors. Then the greedy learning algorithm ensures that most of the models in the ensemble will choose to split on x_j in early iterations.

That is, each tree in the ensemble is identically distributed, with the expected output of the averaged model the same as the expected output of any one of the trees.

Improving on Bagging

Recall, for *B* number of identically and independently distributed variable, *X*, with variance σ^2 , the variance of the estimate of the mean is :

$$\operatorname{var}(\hat{\mu}_x) = \frac{\sigma^2}{B}$$

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Improving on Bagging

For B number of identically but not independently distributed variables with pairwise correlation ρ and variance σ^2 , the variance of their mean is

 $\operatorname{var}(\hat{\mu}_x) \propto \sigma^2 (1 + \rho^2)/B$

Estimates of means for correlated xs, $\rho = 0.5$, for 100 Xs. Here we show the results for 3000 simulations

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Question: Do you see any problems?

- Still some overfitting if the trees are too large
- If trees are too shallow it can still underfits.
- interpretability
- The major drawback of bagging (and other ensemble methods that we will study) is that the averaged model is no longer easily interpretable i.e. one can no longer trace the 'logic' of an output through a series of decisions based on predictor values!

You will be UNAWARE OF WHAT I'M SAYING *4 OUT OF *8 MINUTES

Zeenat Potia

Random Forests

Random Forest is a modified form of bagging that creates ensembles of independent decision trees.

To de-correlate the trees, we:

- train each tree on a separate bootstrap sample of the full training set (same as in bagging)
- 2. for each tree, at each split, we **randomly** select a set of J' predictors from the full set of predictors.

From amongst the J' predictors, we select the optimal predictor and the optimal corresponding threshold for the split.

Random forest models have multiple hyper-parameters to tune:

- 1. the number of predictors to randomly select at each split
- 2. the total number of trees in the ensemble
- 3. the minimum leaf node size

In theory, each tree in the random forest is full, but in practice this can be computationally expensive (and added redundancies in the model), thus, imposing a minimum node size is not unusual.

There are standard (default) values for each of random forest hyperparameters recommended by long time practitioners, but generally these parameters should be tuned through **OOB** (making them data and problem dependent).

e.g. number of predictors to randomly select at each split:

- $\sqrt{N_j}$ for classification
- $-\frac{N}{3}$ for regression

Using out-of-bag errors, training and cross validation can be done in a single sequence - we cease training once the out-of-bag error stabilizes

Same as with Bagging:

Calculate the total amount that the RSS (for regression) or Gini index (for classification) is decreased due to splits over a given predictor, averaged over all *B* trees.

<u>Alternative:</u>

- Record the prediction accuracy on the oob samples for each tree.
- Randomly permute the data for column *j* in the *oob* samples the record the accuracy again.
- The decrease in accuracy as a result of this permuting is averaged over all trees, and is used as a measure of the importance of variable *j* in the random forest.

Variable Importance for RF

100 trees, max_depth=10

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100 trees, max_depth=10

When the number of predictors is large, but the number of relevant predictors is small, random forests can perform poorly.

Question: Why?

In each split, the chances of selected a relevant predictor will be low and hence most trees in the ensemble will be weak models.

Final Thoughts on Random Forests (cont.)

Increasing the number of trees in the ensemble generally does not increase the risk of overfitting.

Again, by decomposing the generalization error in terms of bias and variance, we see that increasing the number of trees produces a model that is at least as robust as a single tree.

However, if the number of trees is too large, then the trees in the ensemble may become more correlated, increase the variance.

Final Thoughts on Random Forests (cont.)

Probabilities:

- Random Forrest Classifier (and bagging) can return probabilities.
- Question: How?

- Unbalance dataset
- Weighted samples
- Categorical data
- Missing data
- Different implementations

AND BOOSTING

