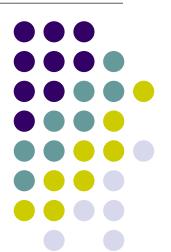
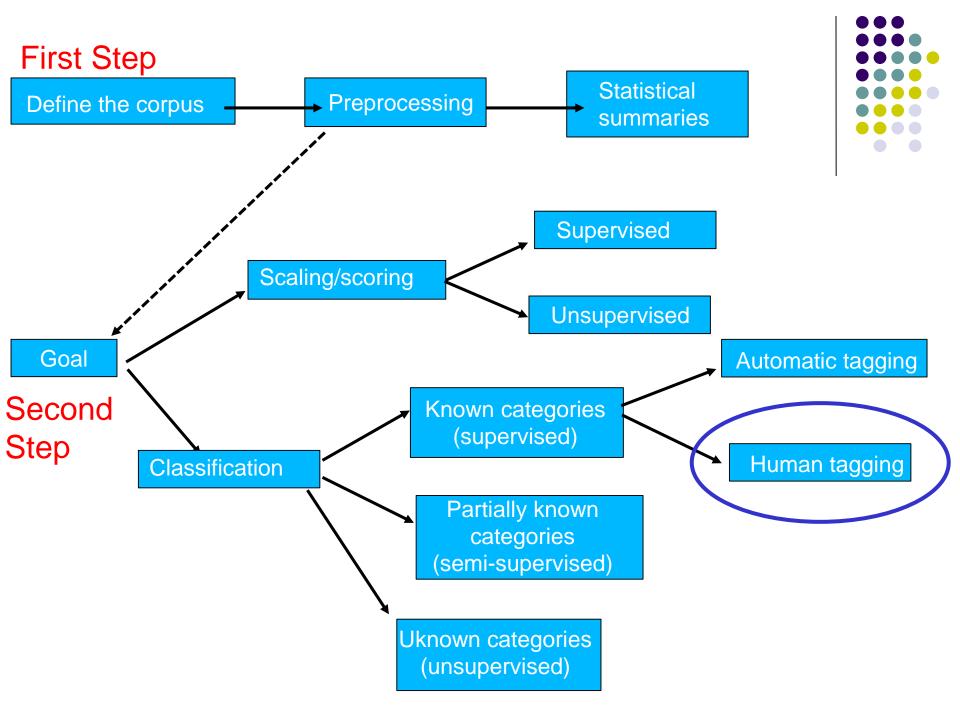
Big Data Analytics

Lecture 7

Supervised classification methods:

A review of (some) Machine Learning Algorithms (first part)









✓ Olivella, Santiago, and Shoub Kelsey (2020). Machine Learning in Political Science: Supervised Learning Models. In Luigi Curini and Robert Franzese (eds.), SAGE Handbook of Research Methods is Political Science & International Relations, London, Sage, chapter 56

ML alghoritms



- Several different possible machine learning algorithms are available out there
- We will offer an intuitive introduction to some of them (pretty standard in text analytics)
- Unfortunately we will have no time to discuss about many other of them
- However, the general logic they employ remains always the same...
- We will begin with Naïve Bayes classifiers

Naïve Bayes classifier: the algorithm allows us to predict a class, given a set of features using (Bayes) probability theorem

But what do we mean by Bayes probability theorem?

Bayesian probability incorporates the concept of *conditional probability*, the probability of event *A* given that event *B* has occurred, i.e., p(A|B)

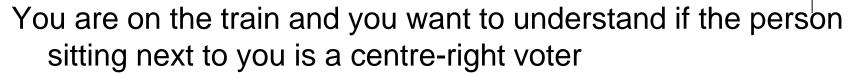
If we have just two classes, we can write:

$$p(A|B) = \frac{p(A)*p(B|A)}{p(A)*p(B|A)+p(\neg A)*p(B|\neg A)}$$

where:

p(A|B)=posterior probability p(B|A)= conditional probability or likelihood p(A)=prior probability of the outcome

An example (with no texts involved!):



You know a priori (your *priors*!) that 54% of the Italian citizens are centre-right voters (46% centre-left)

Now the person sitting next to you open a newspaper. You know that the 35% of centre-right voters read that newspaper (while it is read by 65% of centre-left voters) – your *likelihood*!

Which is your update belief that the person sitting next to you is a centre-right voter?

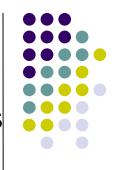
p(CR|N) = p(CR) p(N|CR)/[(p(CR) p(N|CR)+p(CL) p(N|CL))] ==.54*.35/(.54*.35+.46*.65) = .387



Within a text-analytics framework, the goal (as already discussed!) is to infer the probability that document i belongs to category k given word profile \mathbf{W}_i (i.e., the probability of a text belonging to category k given that its predictors – features – values are $\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_p$

This can be written as $p(C_k|x_1, x_2,...,x_p)$)

More formally, the Bayesian formula for calculating this probability is:



$$p(C_k|W_i) = \frac{p(C_k) * p(W_i|C_k)}{p(W_i)}$$

In plain English:

 $p(C_k|W_i)$ =posterior probability: by combining our observed information, we are updating our *a priori* information on probabilities to compute a posterior probability that an observation has class C_k

 $p(W_i|C_k)$ = conditional probability or likelihood

 $p(C_k)$ =prior probability of the outcome (i.e., the average probabily of that outcome in the training-set)

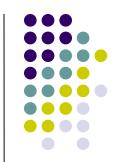
 $p(\mathbf{W}_i)$ =evidence (the word profile we observe)

In other words, the Bayesian formula is simply:



$$Posterior = \frac{prior * likelihood}{evidence}$$

From above, we can drop $p(\mathbf{W}_i)$ - i.e., the evidence - from the denominator since it is a constant across the different categories for each document



An example: let's say we have a training-set on 1000 pieces of fruit. The fruit being a Banana, Orange or some Other fruit. We know **3 variables of each fruit**, whether it's long or not, sweet or not and yellow or not:

Fruit	Long	Sweet	Yellow	Total
Banana	400	350	450	500
Orange	0	150	300	300
Other	100	150	50	200
Total	500	650	800	1000

)

Fruit	Long	Sweet	Yellow	Total
Banana	400	350	450	500
Orange	0	150	300	300
Other	100	150	50	200
Total	500	650	800	1000

From the table we know that, in our training-set: 50% of the fruits are bananas; 30% are oranges; 20% are other fruits (our *priors*!)

Based on our table, we can also say the following:

Out of 500 bananas 400 (0.8) are Long, 350 (0.7) are Sweet and 450 (0.9) are Yellow; Out of 300 oranges 0 are Long (0.0), 150 (0.5) are Sweet and 300 (1.0) are Yellow; From the remaining 200 fruits, 100 (0.5) are Long, 150 (0.75) are Sweet and 50 (0.25) are Yellow. All these values refer to conditional probabilities or likelihood!



Now let's say we're given the variables of a piece of fruit and we need to predict the class (out-of-sample prediction!)

If we're told that the additional fruit is Long, Sweet and Yellow, we can classify it using the Bayes formula and subbing in the values for each outcome, whether it's a Banana, an Orange or Other Fruit

The one with the highest probability (score) being the **winner** class!



p(Banana|Long, Sweet, Yellow)=p(Banana=0.5)*p(Long|Banana=0.8)*p(Sweet|Ban

ana=0.7)*p(Yellow|Banana=0.9)=**0.252**

p(Orange|Long, Sweet, Yellow)=p(Orange=0.3)*p(Long|Orange=0)*p(Sweet|Orange=0.5)*p(Yellow|Orange=1)=**0**

p(Other|Long, Sweet, Yellow)=p(Other=0.2)*p(Long|Other=0.5)*p(Sweet|Other=0.75)*p(Yellow|Other=0.25)=**0.01875**

In this case, based on the **highest score**, we can classify this Long, Sweet and Yellow fruit as a Banana

- In the case of text-classification, instead of Banana, Orange and Other you will have some specific categories (say, Positive, Negative and Neutral if we are applying a sentiment analysis) and instead of Long, Sweet and Yellow we will have the features (aka: words) included in our DfM
- The logic however followed in predicting the category of each text included in the test-set remains exactly the same as the one we just discussed!
- Given the features included in that (unread) text, we will estimate its posterior probability to belong to each given category...
- ...then we will assign that (unread) text to the category with the largest posterior probability among the one we estimated!



Note a possible problem of the logic just explained: given that naïve Bayes uses the product of variable probabilities conditioned on each class, we run into a serious problem when new data includes a variable value that **never occurs** for one or more levels of a response class

This is what happens with p(Long|Orange) = 0. This "0" will ripple through the entire multiplication of all variable and will always force the posterior probability to be zero for that class

This is clear a HUGE PROBLEM when dealing with a sparse DfM (as it is usually the case)!

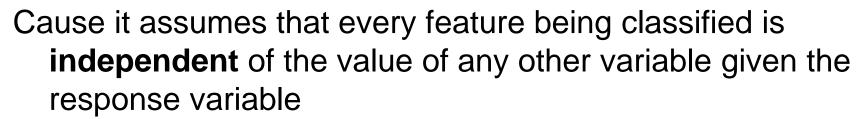


A solution to this problem involves using the *Laplace* smoother

The Laplace smoother adds a small number to each of the counts in the frequencies for each feature, which ensures that each feature has a nonzero probability of occurring for each class

Typically, a value of 1 for the Laplace smoother is employed, but this is a *tuning parameter* to incorporate and optimize with cross validation (more on this later on!)

Why Naïve?



In the previous example each of the three variables (Long, Sweet, Yellow) are considered to contribute independently to the probability that the fruit is a Banana, regardless of any correlations between features

By making this assumption we can simplify our calculation such that the posterior probability is simply the product of the probability distribution for each individual variable conditioned on the response category





Variables, however, aren't always independent!

Although the model is *clearly wrong* – quite often features are not conditionally independent - it has proven to be a useful classifier for a diverse set of tasks

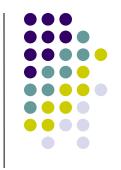
The same is true for text analysis: assuming that features (i.e., words) are generated indipendently is wrong given that the use of words is highly correlated in any data set. However, Naïve Bayes classifier can still be useful (remember the First Principle of text-analysis!)

- Naïve Bayes classifier algorithm has a simple, but powerful (and fast!), approach to learning the relationship between words and categories
- Moreover, it has been shown to perform surprisingly well with very small amounts of training data that most other classifiers, would find significantly insufficient
- As a result, if you find yourself with a small amount of training data Naïve Bayes would be a good bet!
- Likewise, Naïve Bayes' simplicity prevents it from fitting its training data too closely and therefore does not tend toward **overfitting** especially on smaller datasets like other approaches do

However, Naïve Bayes classifier also behaves differently on the other end of the spectrum, when provided with large amounts of training data

As it is fed increasing quantities of training data, the performance of the Naïve Bayes classifier plateaus above a certain threshold

Its simplicity prevents it from benefiting incrementally from training data past a certain point

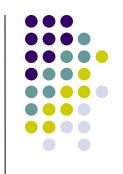




Jordan Goldmeier @Option_Explicit

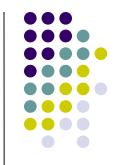
Where do data scientists go camping?

In random forests



To understand random forest, let's start with what we mean by a *Decision tree model*

A decision tree is basically a set of rules used to classify data into categories. In particular it is the set of rules which best partitions the data



More in details...

...For a single-tree model, the goal is to partition the space of predictor features (i.e. the set of *all unique combinations of predictor values*) into *B* non-overlapping and exhaustive regions, R₁, R₂, ..., R_B, that are **relatively homogeneous** with respect to the outcome *y*, thus improving overall predictive accuracy by sorting observations into their respective bins

An example: Given only the gender and weight of a person, can we predict whether they are Japanese or American (our 2 classes/categories)?

Let's train a decision-tree algorithm by using the following training set:

Weight (lbs.)/Sex/Nationality

195 M American

190 M American

160 F American

165 F American

165 M Japanese

160 M Japanese

130 F Japanese

140 F Japanese

For

Key idea: the procedure to create decision trees is **recursive**. For a set (**S**) of observations, the following algorithm is applied:

Step 1: If every observation in **S** is the **same class** or if **S** is very small, the tree becomes an **endpoint**, labeled with the *most* frequent class

American

.50 .50

100%

Clearly the initial group with all our 8 observations does not satisfy Step 1! Therefore, we need to move to Step 2

Step 2: If **S** is too large and it contains more than one class (as in our case!), find the **best rule** based **on one feature** to split it into subsets, one for each class (different rules can be used in this respect

✓ The aim is always the same: the "best" branching rule is
the one that results in the most information gain)



So given our initial **S**, on which feature should we focus?

- ✓ Clearly not the gender feature! If we divide our initial S according to gender (Male/Female) we would in fact remain with two sub-groups that will have the same problems of the initial S (50% of the two groups). No information gain at all!
- ✓ Therefore, let's focus on the weight feature. But weight is a continuous feature! Therefore which rule should we apply with respect to weight? For example we could apply the following one: if weight is larger or lower than 150

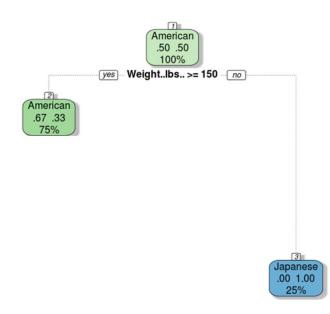


The decision tree: first feature selection rule

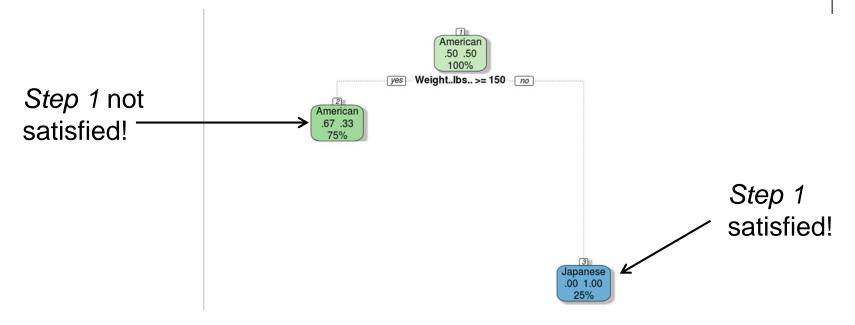




What's the result of this first rule?



If you had to go to step 2, apply step 1 to each new subset

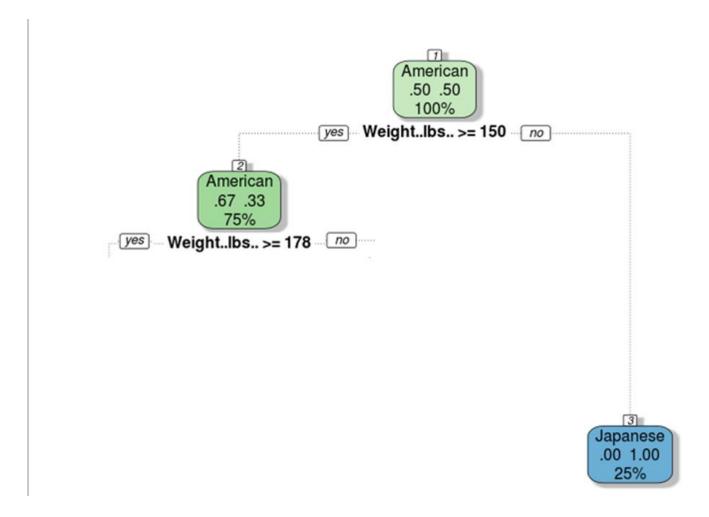


If your subsets need to go to step 2, apply step 1 to the subsubsets, etc.

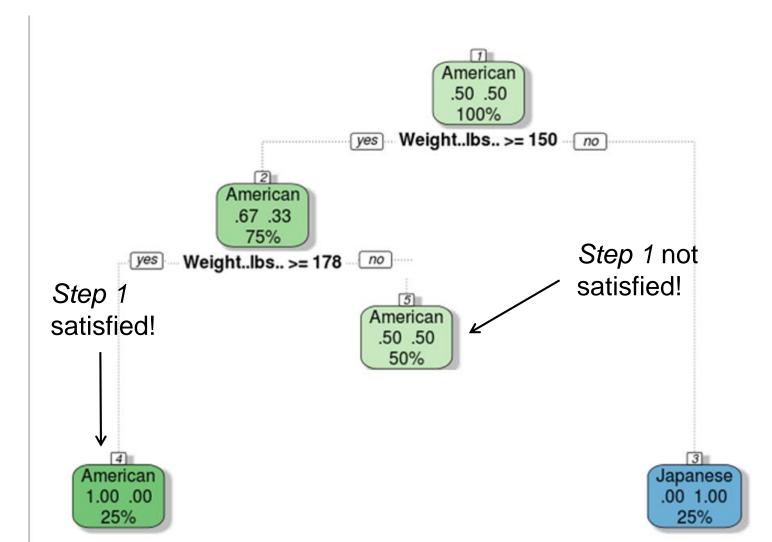
When everything is split up appropriately (into buckets that are very small or entirely one class), you have a set of rules that look like a tree!

The decision tree: second feature selection rule



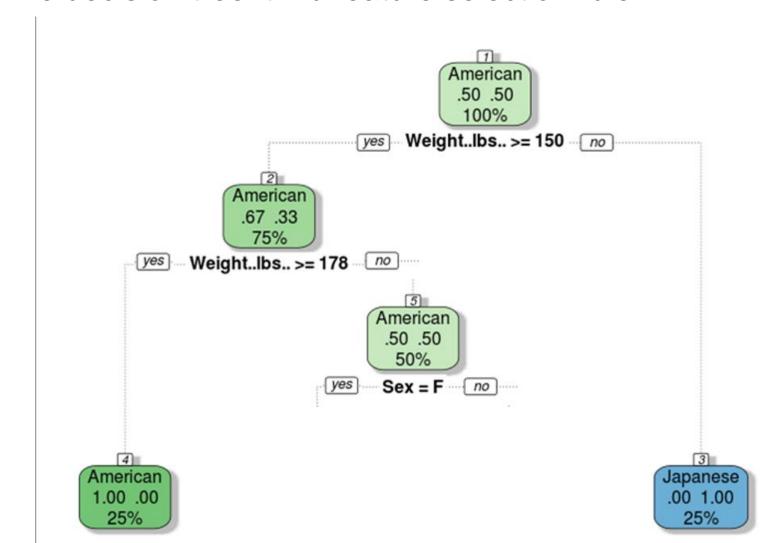


The decision tree: second feature selection rule



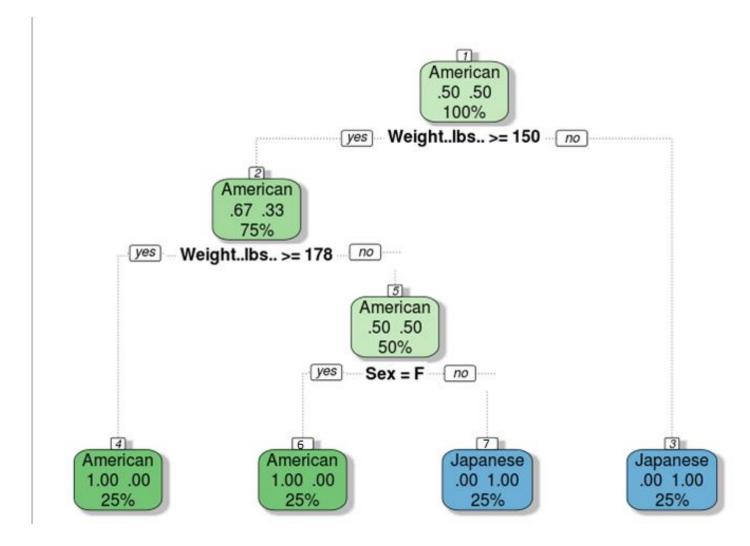
The decision tree: third feature selection rule











Wanna replicate it? Two lines of command!

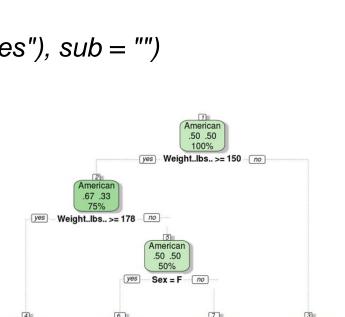
library(rpart)

library(rattle)

nation <- read.csv("Nationality.csv", stringsAsFactors=FALSE)

fit <- rpart(Nationality ~ Sex + Weight, method="class", data=nation, minsplit=2, minbucket=1)

fancyRpartPlot(fit, palettes = c("Greens", "Blues"), sub = "")





We can now use the decision tree (and its rules) just obtained to estimate the nationality of any individual not original included in our training-set

For example, if I am telling you that we have a female that weights more than 178 pounds...

...by applying the decision tree just fitted, we would predict that individual as being an American one!

In this example the tree can **perfectly explain** the data

This is a **serious limitations**! In the real world, there is overlap: there are fat (not many, still...) Japanese people and skinny (not many, still...) Americans

In other words, growing a single, deep tree using binary recursive splitting can result in a grossly overt model. In turn, this high level of in-sample predictive accuracy usually comes at the expense of high estimator variance, as single trees grown recursively can often times yield wildly different predictions as a result of small changes in the training set...overfitting!!!

So, what to do?

Trees are usually "pruned" to avoid **overfitting**. The pruning algorithm removes final nodes so that the model is a little more general and will tend to generalize better to new, unseen data

"Pruning" however is not always an advisable solution to the problem of overfitting!

The sequential nature of the recursive splitting algorithm means in fact that the structure of the tree is often highly sensitive to small changes in the observations included in the training-set

So what to do to avoid overfitting (part 2)?

Bootstrap aggregating (bagging)!

We already discuss about bootstrapping! Do you remember?

In essence bootstrapping **repeatedly draws independent samples** from our data set to create bootstrap data sets.

This sample is usually performed with *replacement*, which means that the same observation can be sampled more than once

Crucially given the problem discussed with respect to decision-trees, this computation is robust to (i.e., less affected from) sample specific characteristics

Bootstrap aggregating (bagging)!

Now back to Bagging!

Bagging follows three simple steps:

- 1. Create *m bootstrap samples* from the training data. Bootstrapped samples allow us to create many slightly different data sets but with the same distribution as the overall training set. Note: the bootstrap samples *must not* (and usually do not!) necessary have the same size of the original training-set. Why? Give me a minute!
- 2. For each bootstrap sample train a single, unpruned classification tree
- Average individual predictions from each tree to create an overall average predicted value

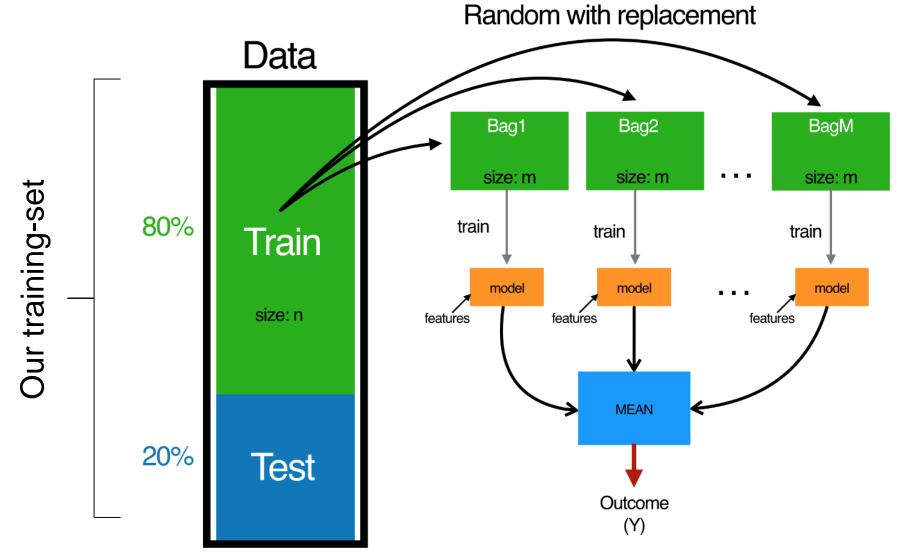


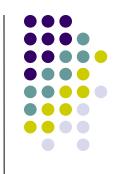
Bootstrap aggregating (bagging)!

Bagging combines and averages therefore multiple models

Averaging across multiple trees reduces the variability of any one tree and reduces overfitting, which improves predictive performance





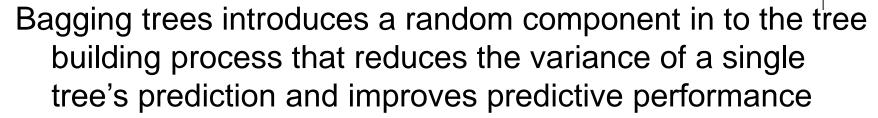


- As written earlier, the bootstrap samples have not usually the same size of the original training-set
- On average, a bootstrap sample will contain as a default 63% of the training data (a parameter you can change!)
- This leaves therefore about 37% of the data out of the bootstrapped sample. This is the *out-of-bag (OOB)* sample. What's the advantage of that?



- Imagine that we have 500 documents in the training-set, and imagine that our bootstrapped sample is based on 400 documents
- We can then train the ML algorithm on such 400 documents, and then using such model to classify the remaining 100 documents (not used to train the ML algorithm) in the OOB sample
- Then we can contrast the prediction of our model with the "true" value of the OOB sample (that we know about, given that such sample is included in the training-set after all!) to evaluate the accuracy of our prediction!
- We can use the **OOB observations** to produce therefore a *natural cross-validation* (?!?) process: more on this later on!

Still, bagging for itself cannot be enough...



However, the trees in bagging are not completely independent of each other since all the original predictors are considered at every split of every tree

Therefore, trees from different bootstrap samples typically have similar structure to each other (especially at the top of the tree) due to underlying relationships

And so? How to reduce the correlation among trees?



The Random Forest (RF) idea!

Let's inject more randomness into the tree-growing process. RF achieve this in two ways:

Bootstrap: similar to bagging, each tree is grown to a bootstrap resampled data set

Split-variable randomization (this is new!): each time a split is to be performed, the search for the split variable is limited to a random subset of *m* of the *p* variables (features). This is a tuning parameter. When m=p, the randomization amounts to using only step 1 and is the same as *bagging*



The basic algorithm for a random forest can be therefore generalized to the following:

- 1. Given training data set
- 2. Select the number of trees to build (ntrees)
- 3. for i = 1 to ntrees do
- 4. | Generate a bootstrap sample of the original data
- 5. | Grow a tree to the bootstrapped data
- 6. | for each split do
- 7. | | Select m variables at random from all p variables
- 8. | Pick the best variable/split-point among the m
- 9. | | Split the node into two child nodes
- 10. | end
- 11. | Use typical tree model stopping criteria to determine when a tree is complete (but do not prune)
- 12. end

By fitting a tree (with no pruning) to each bootstrapped sample <u>and</u> by restricting the choice of each splitting variable to a random subset of predictors, we are sure that each boostrapped tree provides a truly different "perspective" on the prediction problem. All this, of course, minimizes the risk of overfitting!

The final prediction is going to be a function of each prediction in each random sample, for example it can be the average of each prediction

Often, ML models are considered "black boxes" due to their complex inner-workings

After all, it is precisely due to their complexity that they are typically more accurate for predicting **nonlinear or rare phenomena**

Indeed, a ML algorithm would allow you to discover quite easily non-linear relationships difficult to detect ex-ante...

For example, in a recent paper (Jordan et al. 2022), it has been suggested the following procedure:

- 1. Estimate a parametric model that tests theoretically grounded hypotheses
- 2. Use a machine learning approach on the same set of theoretical predictors to evaluate the robustness of the initial parametric tests
- 3. Adjust the initial parametric model to account for any nuances revealed in the machine learning approach

- Unfortunately, more accuracy often comes at the expense of interpretability, and interpretability is crucial
- It is not enough to identify a ML model that optimizes predictive performance (via for example cross-validation: more on this later on)
- Understanding and trusting model results is a hallmark of good (social and political) science!
- Luckily, several advancements have been made to aid in interpreting ML models over the years.
- Interpreting ML models is an emerging field that has become known as "Interpretable Machine Learning" (IML)

- Approaches to model interpretability can be broadly categorized as providing *global or local explanations*
- Global interpretations help us understand the inputs and their entire modeled relationship with the prediction target
- Global interpretability in particular is about understanding how the model makes predictions, based on a holistic view of its features and how they influence the underlying model structure

It answers questions regarding which features are relatively influential as well as how these features influence the response variable (i.e., in a positive or negative way?)



Global interpretations is therefore crucial to assess the content validity (does the measure capture the theoretical concept?) of a ML algorithm

This is highly relevant when one is not only interested in high predictive power but also in accurately measuring latent concepts of interest

For example, the most influential features found by our ML algorithm in affecting a given class-labels should make sense from a theoretical point of view rather than being based on "accidental, meaningless regularities and confounding patterns"

"In their effort to match human annotations or given ground truths, algorithmic classifiers show little interest in separating valid variation in the material from accidental, meaningless regularities and confounding patterns. Relying on salient, correlated patterns identified in the data, these tools may still frequently guess correctly, while potentially introducing systematic biases into the analysis"

Baden, C., C. Pipal, M. Schoonvelde, and M. A. C. G. van der Velden. 2021. "Three Gaps in Computational Text Analysis Methods for Social Sciences: A Research Agenda." Communication Methods and Measures (December 27, 2021): 1–18. ISSN: 1931-2458, 1931-2466

For an interesting paper discussing how the results of a seemingly "good ML algorithm" in terms of accuracy, could present in reality a poor "content validity":

Michael Jankowski and Robert A. Huber (2022). When Correlation Is Not Enough: Validating Populism Scores from Supervised Machine-Learning Models, *Political Analysis*, DOI: 10.1017/pan.xxxx.xx

- Local interpretations help us on the other hand to understand model predictions for a single row of data or a group of similar rows (i.e., the marginal impact of a feature across its values on our DV, holding all the other features at their actual value for each observation i)
- In text analytics (given that we are dealing with huge DfM wherein the features per-se are not the main focus of our interest) we are mainly interested in global interpretations
- However if you use ML for other aims, local interpretations become VERY important!
- For an interesting paper illustrating the power of local interpretations: Soren Jordan, Hannah L. Paul, Andrew Q. Philips, How to Cautiously Uncover the "Black Box" of Machine Learning Models for Legislative Scholars, *Legislative Studies Quarterly*, 2022, https://onlinelibrary.wiley.com/doi/abs/10.1111/lsq.12378

R pakcages to install

```
install.packages("e1071", repos='http://cran.us.r-project.org')
install.packages("randomForest", repos='http://cran.us.r-project.org')
install.packages('caret', repos='http://cran.us.r-project.org',
   dependencies = TRUE)
install.packages("naivebayes", repos='http://cran.us.r-project.org',
   dependencies = TRUE)
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install.packages("iml", repos='http://cran.us.r-project.org')
install.packages("future", repos='http://cran.us.r-project.org')
install.packages("future.callr", repos='http://cran.us.r-project.org')
install.packages("gridExtra", repos='http://cran.us.r-project.org')
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"... operationalization [of a concept] is replaced by powerful algorithms trained to identify any patterns and indicators that correlate with provided annotations, effectively supplanting validity with predictive performance [...] In their effort to match human annotations or given ground truths, algorithmic classifiers show little interest in separating valid variation in the material from accidental, meaningless regularities and confounding patterns. Relying on salient, correlated patterns identified in the data, these tools may still frequently guess correctly, while potentially introducing systematic biases into the analysis"

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https://www.tandfonline.com/doi/full/10.1080/19312458.2021.201557