
Advanced Predictive Modeling Workshop

Tree-based Methods

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Kudakwashe Chibanda, FCAS, MAAA

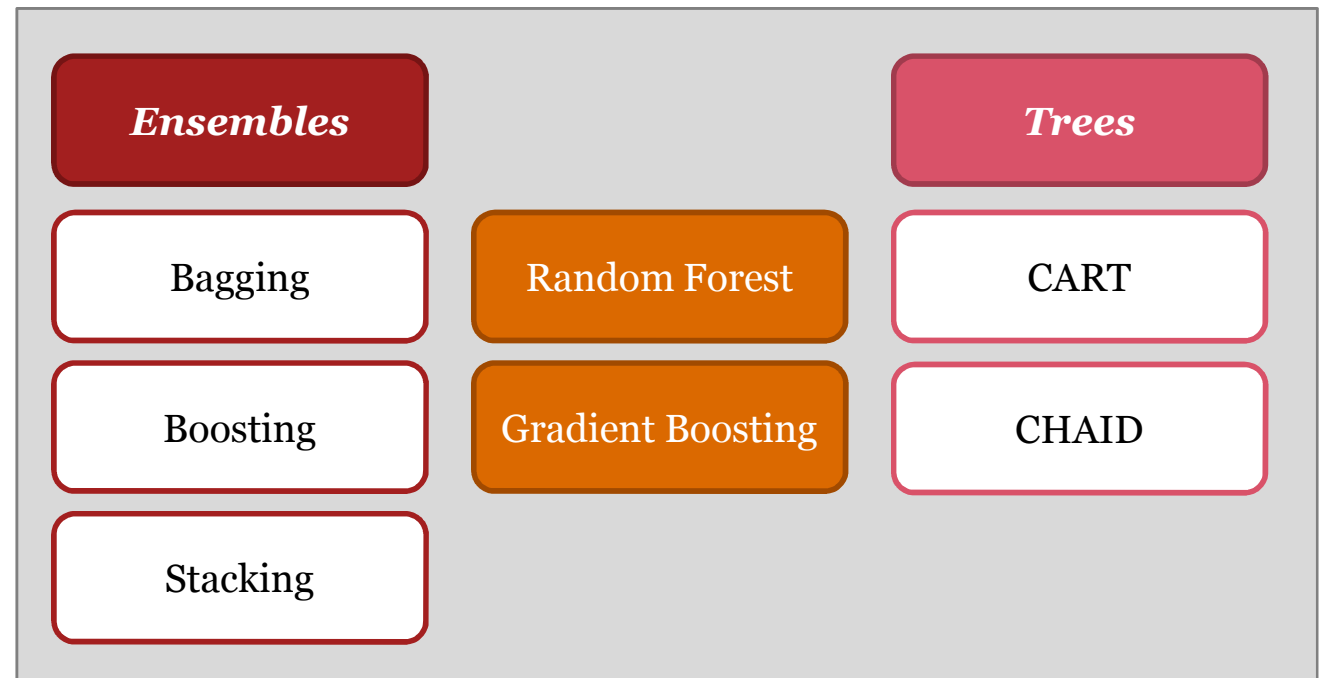
kudakwashe.chibanda@pwc.com

Jean-François Greeff, FASSA

jean.francois.greeff@pwc.com

Agenda

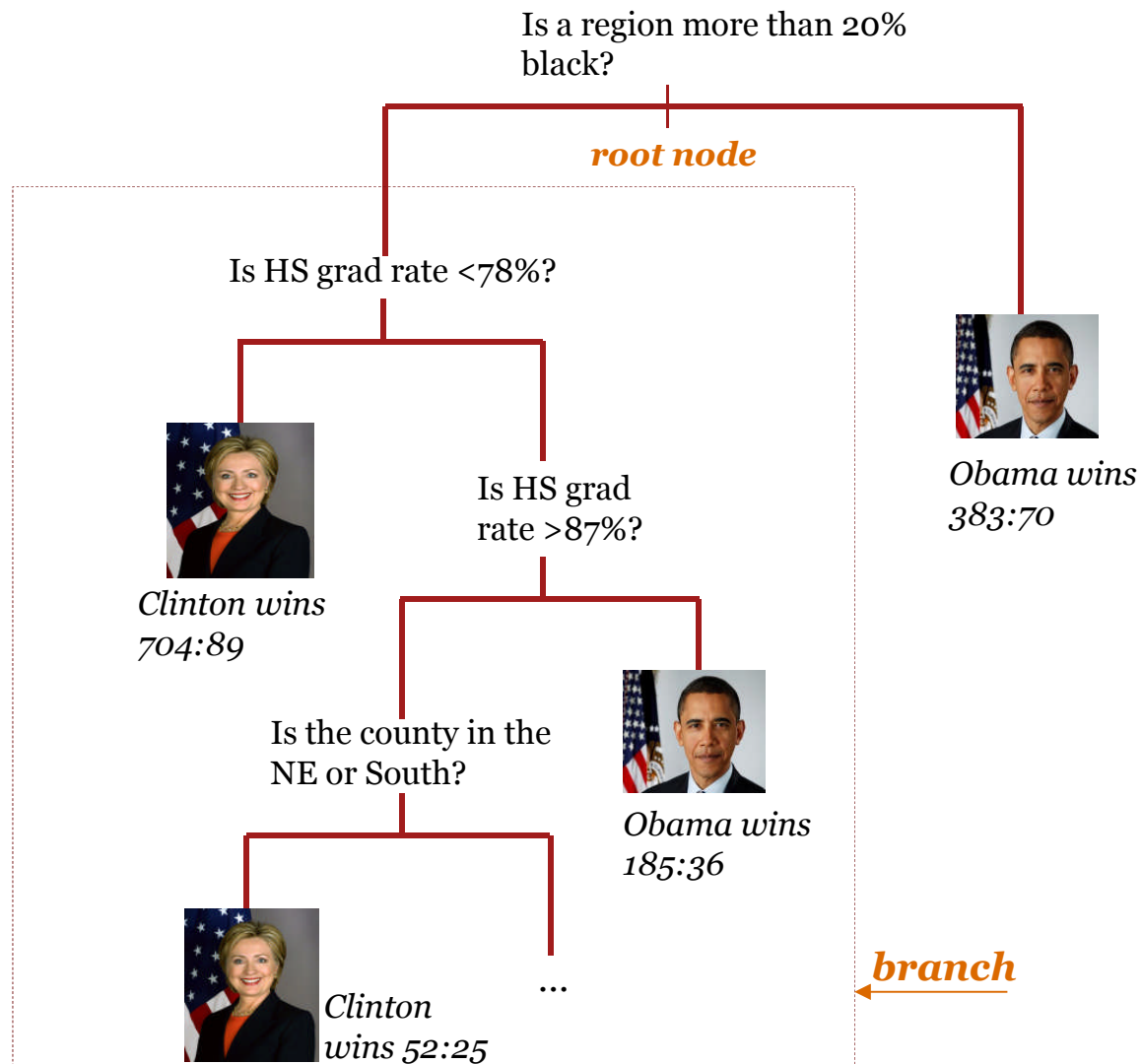
1. Decision trees
2. Ensembles
3. Classification examples
4. Regression examples



Decision trees

“If you dream of a forest, you better learn how to plant a tree”

Introduction



PwC

terminal node/leaf

Decision Tree: The Obama-Clinton Divide (from NYT April, 2008)

Features

- Non-parametric classification/regression tools
- Create splits according to measures of homogeneity

Advantages

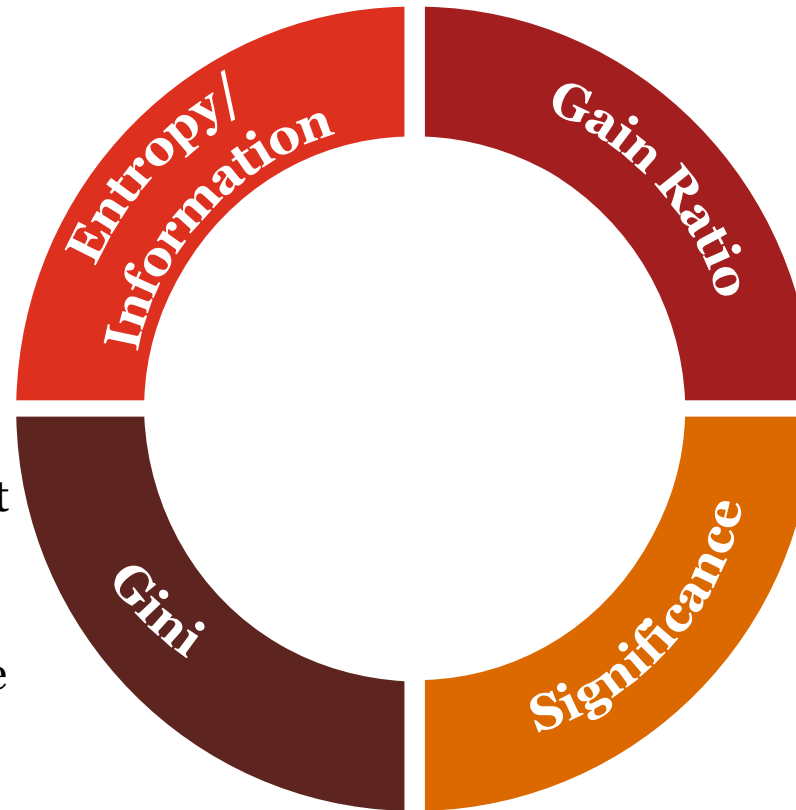
- Simple to understand and interpret
- Flexible for non-linear or complex relationships

Disadvantages

- Overfitting
- Unstable/Biased if certain classes of data dominate

Splitting Criteria

- Entropy measures the **disorderliness** for each variable level
- The **purier** the level for a given response, the more **predictable** the outcome
- The weighted average entropy across all levels of a variable gives us information
- Gini impurity is a purity measure that relies on **misclassification**
- It measures the probability that a randomly selected observation will be placed in the wrong bucket (i.e. misclassified)



- A large number of observations in a level can **bias** the information towards the entropy of the concentrated level
- To compensate, **Intrinsic Information** is calculated
- II takes size and number of levels into account i.e. penalizes large values/splits
- $Gain\ Ratio = \frac{Information\ Gain}{Intrinsic\ Information}$
- **p-values of Chi-Square** statistics can be used to split nodes
- Measure statistical significance of a variable's levels and the response (i.e. test **null hypothesis of independence**)
- Insignificant splits are merged while significant ones are tested for further splits

Purity Measures Calculations

Outlook (<i>x variable</i>)	Yes (<i>y variable</i>)	No (<i>y variable</i>)	Total (<i>by level</i>)
Sunny (node i)	3	2	5
Overcast	4	0	4
Rainy	20	30	50
Total (t branch)	27	32	59

Entropy/ Information measure purity of outcomes at each node, taking number and size of nodes into account

IG measures increase in purity from having no splits [$I(t)$] to having c splits

Information Gain and Gini also take purity at the branch (regardless of splits) into account

- **Entropy** [$H_y(i)$] = $-\sum_y p(y|i) \log(y|i)$

$$\text{Entropy(sunny)} = -\left(\frac{3}{5} \log\left(\frac{3}{5}\right) + \frac{2}{5} \log\left(\frac{2}{5}\right)\right) = 0.971$$

- **Information** [$H(t)$] = $-\sum_{i=0}^{c-1} p(i) H_y(i)$

$$H(\text{Outlook}) = \frac{5}{59} * 0.971 + \frac{4}{59} * 0 + \frac{50}{59} * 0.971 = 0.905$$

- **Information Gain** [$IG(t)$] = $I(t) - H(t)$

$$IG(t) = -\left(\frac{27}{59} \log\left(\frac{27}{59}\right) + \frac{32}{59} \log\left(\frac{32}{59}\right)\right) - 0.905 = 0.09$$

- **Intrinsic Information** [$II(t)$] =

$$II(t) = -\left(\frac{5}{59} \log\left(\frac{5}{59}\right) + \frac{4}{59} \log\left(\frac{4}{59}\right) + \frac{50}{59} \log\left(\frac{50}{59}\right)\right) = 0.767$$

- **Gain Ratio** [$GR(t)$] = $\frac{IG(t)}{II(t)}$

$$GR(t) = \frac{0.09}{0.767} = 0.117$$

- **Gini** [$G(t)$] = $1 - \sum_y p(i|t)^2$

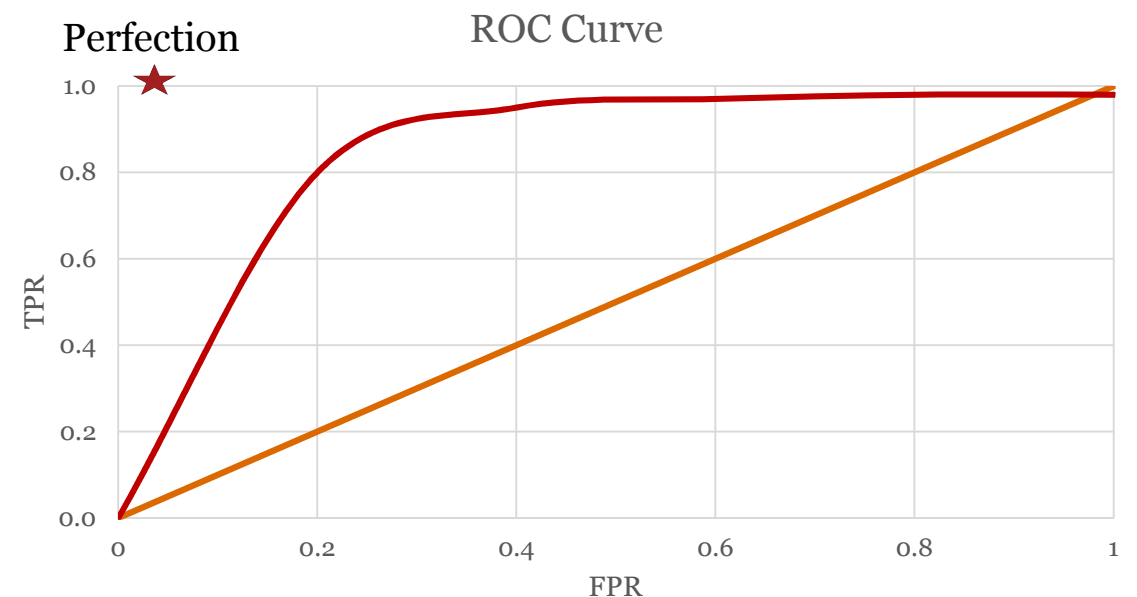
(prior to split)

$$G(t) = 1 - \left[\left(\frac{27}{59}\right)^2 + \left(\frac{32}{59}\right)^2\right] = 0.496$$

Receiver Operating Characteristic (ROC)

- To measure predictive performance in binary classifier models, we rely on **confusion matrices**
- Using a selected **threshold**, we can bucket observations into each one of the four buckets as shown in the table
- Receiver Operator Curves (ROC) are commonly used to select a threshold
 - By plotting relationship between TPR and FPR, we can determine the point that **maximizes TPR while minimizing FPR**
 - We can also summarize the information by calculating **Area Under Curve (AUC)**

		Predicted	
Actual	True Positive (TP)	False Negative (FN)	True Positive Rate (Sensitivity): $TPR = \frac{TP}{TP + FN}$
	False Positive (FP)	True Negative (TN)	False Positive Rate (Fall-out): $FPR = \frac{FP}{FP + TN}$



Types of Trees – ID3 and C4.5

ID3

- Purity measure: **Entropy**
- Methodology: at each node, calculate entropy for all variables. Select variable with minimum entropy
- Splits: can have multiple splits
- Continuous/missing data: no
- Risks: does not **prune**
 - Fix: use **stopping criteria** to avoid overfitting

C4.5

- Purity measure: **Information Gain**
- Methodology & splits: similar to ID3
- Continuous/missing data: yes
- Risks: susceptible to **outliers**
 - Fix: remove outliers

Classification And Regression Trees (CART)

Classification Trees

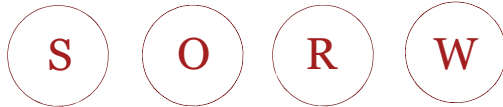
- Purity measure: **Gini impurity**
- Methodology: at each node, calculate gini for all variables. Select split with minimum gini
- Splits: **binary**
- Continuous data: requires splitting
- Risks: does not work for **multiple category** data
 - Fix: use CHAID/ID3

Regression Trees

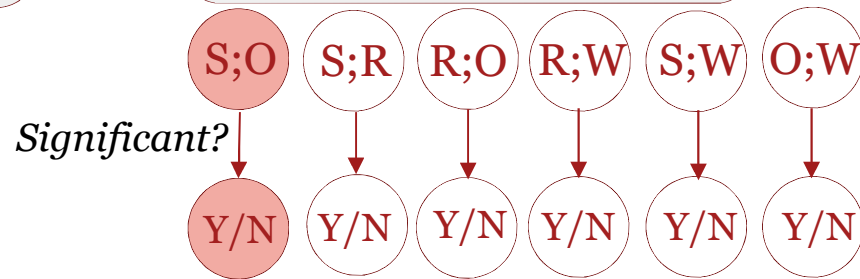
- Purity measure: **Variance reduction**
- Methodology: For each variable, the split is determined by the point that **minimizes SSE**
- Continuous/missing data: yes
- Risks: **overfitting**
 - Fix: **prune** using Sum of Square Errors (SSE)

Chi-square Automatic Interaction Detector (CHAID)

Step 1: Discretize continuous variables. For categorical variables, pair levels

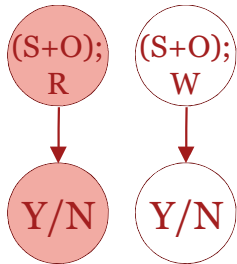


Step 2: Perform Chi-Square test for each pair's significance with response



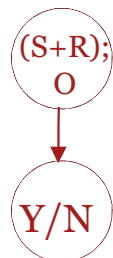
Apply **Bonferroni Adjustment** to penalize for multiple testing

Step 3: Merge pair with least significance & repeat test until **stopping criteria**



PwC

Step 4: Test whether merged categories should be further split



Only test combinations not previously tested

Step 5: Repeat step 1-4 for every variable to determine optimal split

Signature characteristic of CHAID is its ability to handle multiple categories

Step 6: Select root node based on variable with smallest χ^2 with response

Classification Example

Ensembles

Weak Learners and Strong Classifiers

Weak learners

Performs well only on a subset of the domain

May be unstable with small perturbations in data

May be biased in its predictions

Typically what we **have**

Strong classifiers

Performs well over the whole domain

Stable across small changes in the data

Unbiased in its predictions

What we **want**



Illustration of Ensembling (1)

Situation

- Transmit binary signal from A to B
- Ensure that signal uncorrupted

Ensemble approach

- Use 3 independent signal carriers
- Majority vote (Choose bits where 2+ of three carriers agree)

Consequence

- Reconstruct signal with reduced error

	Signal	Accuracy
Original signal	0100101001000110	
Signal 1	0100001001000110	93.75%
Signal 2	0100101001000111	93.75%
Signal 3	0100100101000110	87.5%
Combined Signal	0100101001000110	100%

	Signal	Accuracy
Original signal	0100101001000110	
Signal 1	0100000101010101	60.00%
Signal 2	0000111000111110	60.00%
Signal 3	0100000000010010	66.67%
Combined Signal	0100000000010110	73.33%

Illustration of Ensembling (2)

- 3 signals with probability of corruption 30% per bit

- $P(\text{All correct}) = 0.7^3 = 34.29\%$
- $P(2 \text{ correct}) = 3 \times (0.7^2 \times 0.3) = 44.09\%$
- $P(1 \text{ correct}) = 3 \times (0.7^1 \times 0.3^2) = 18.90\%$
- $P(\text{None correct}) = 0.3^3 = 2.70\%$

- Correction made for 44.09% of the bits
- Expected accuracy of 78.38% per bit

Only if signals ***uncorrelated***

Bagging

Bootstrap Aggregation

Algorithm

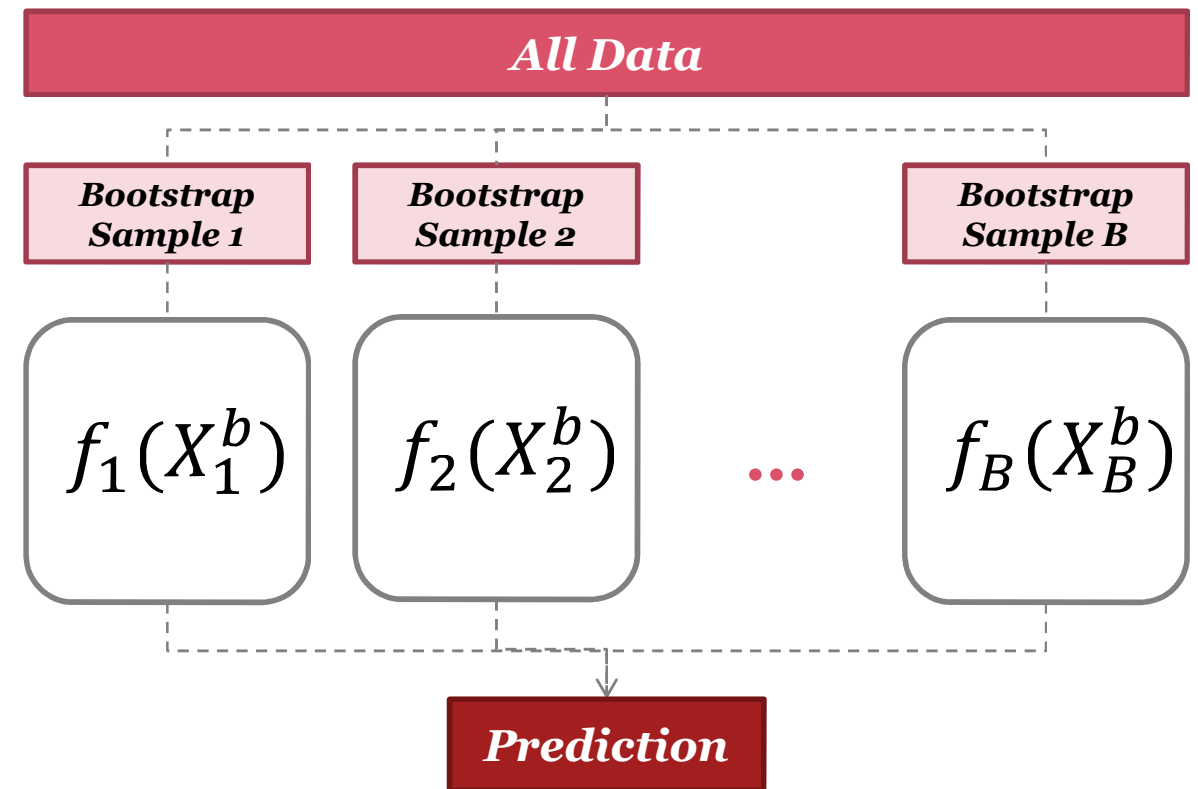
1. Create bootstrap resample of data
2. Fit model on each resample
3. Scoring:
 - Classification: Majority vote
 - Regression: Mean/Median score

Advantages

- Produces more stable predictions – i.e. reduces variance
- Less likely to over-fit data

Disadvantages

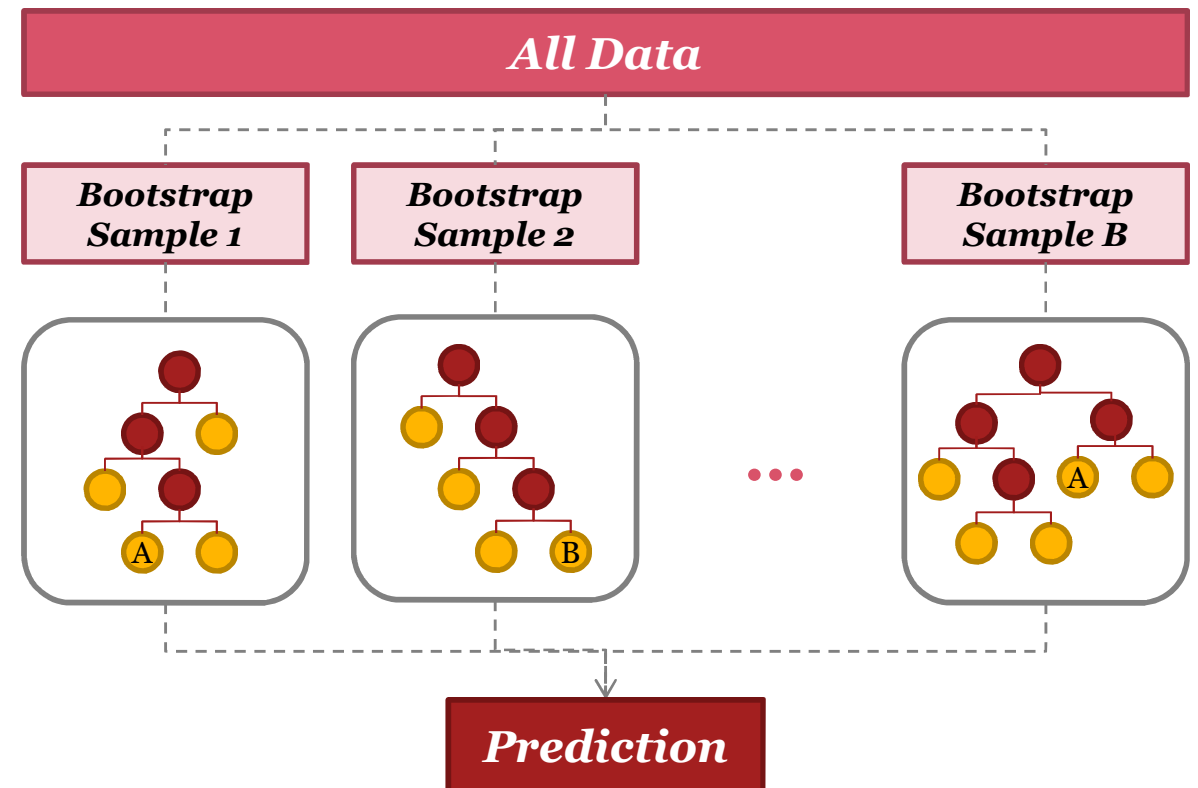
- Generates a “black box”



Random Forests

Bagging Decision Trees

- Introduced by Leo Breiman (2001)
- Uses bagging to improve decision trees
- De-correlates trees by sampling
 - Data with replacement
 - Columns/features at each node
- Produces out-of-bag error rates
- Produces variable importance measure
- Parameters to tune*:
 1. Number of trees
 2. Number of features to select at each node



Boosting

Algorithm

- Rather than fitting models to bootstrap samples of the data – boosting fits sequential models focusing on areas of poor performance
- Subsequent models correct errors of previous models

Advantages

- Decrease bias in predictions

Disadvantages

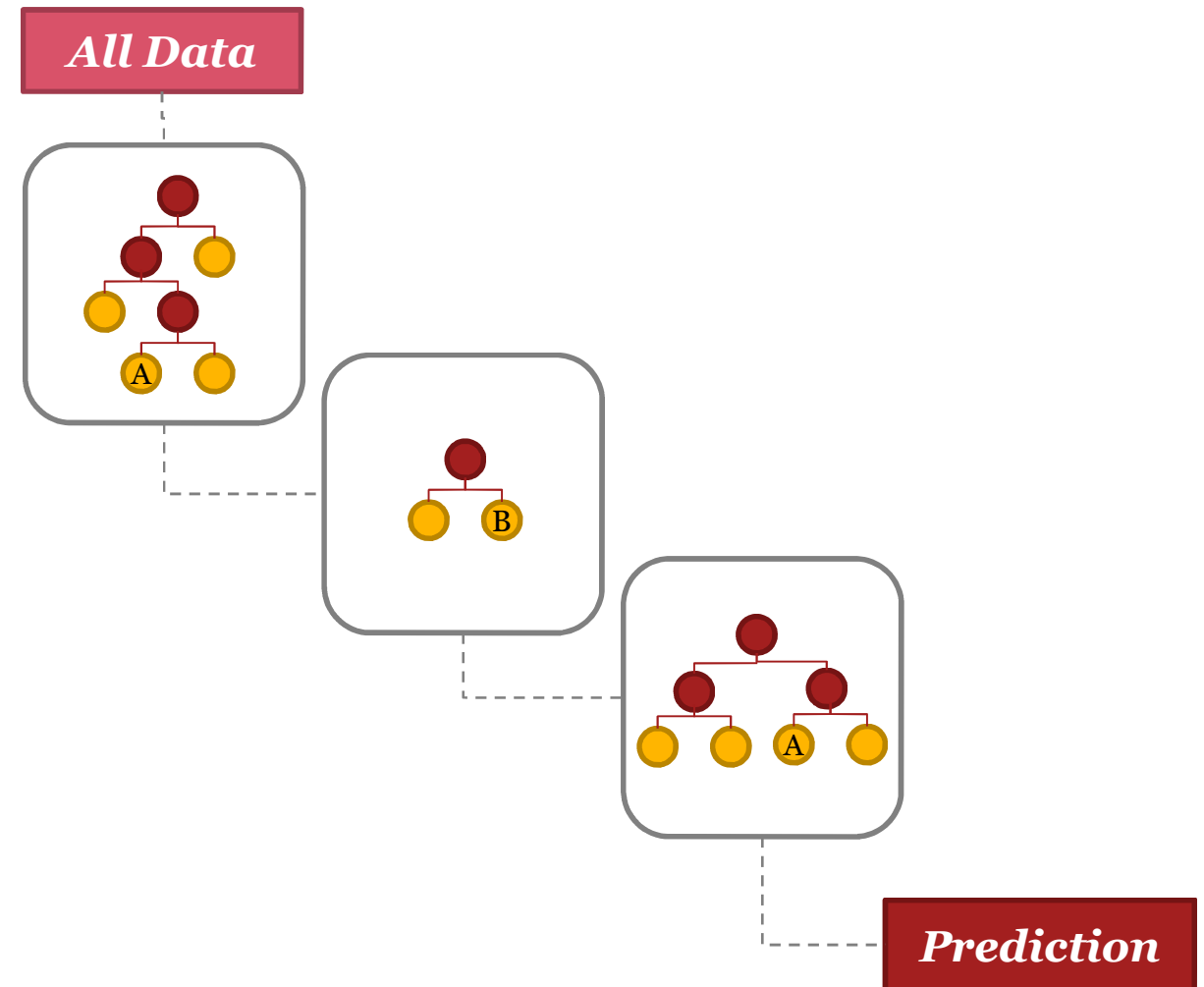
- Generates a “black box”
- May be sensitive to outliers and noise

AdaBoost	GBM
Adaptive Boosting	Gradient Boosted Machines / Models
Fits model to weighted distribution of the data. More weight is given to observations that have the highest error rate.	Fits model to the residual of the prior models.

Gradient Boosted Trees

Boosting Decision Trees

- Introduced by Jerome Friedman (1999)
- Uses boosting to improve decision trees
- XGBoost algorithm most common
 - Stochastic gradient descent
 - Feature sub-sampling
- Parameters to tune*:
 1. Number of trees
 2. Depth of trees
 3. Learning rate



Stacking

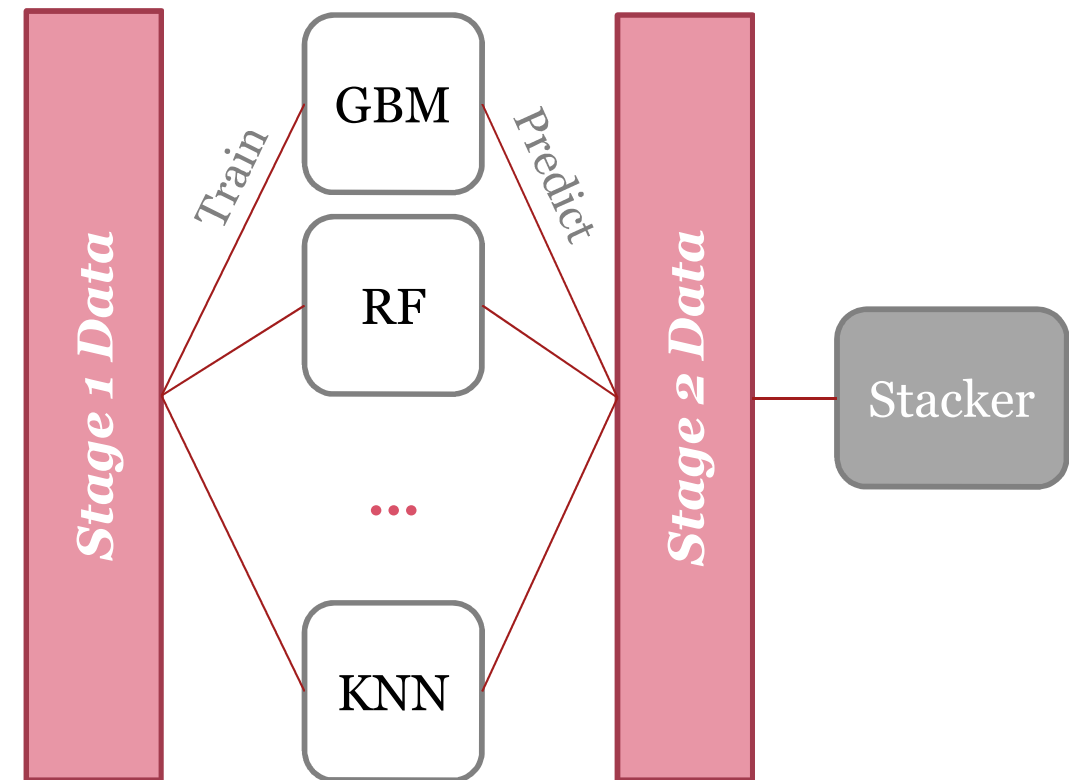
Stacked generalization & Blending

Algorithm

- Two stages of model fitting
 - First Stage: Fit base learners to data
 - Second Stage: Fit meta-learner to predictions of base learners

Considerations

- Different approaches to how the stacking is performed
- Careful consideration needs to be given to what data is used at what stages
- Need diverse models



Classification Example

Predictive Modeling Applications

Advanced Predictive Modeling Workshop

Tree-based Methods

Q&A



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