Introduction to Statistical Machine Learning

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Outline

- Bayesian additive regression trees (BART)
- Boruta
- Neural networks
- Generalized additive models for location, scale and shape (GAMLSS)

BART •00000

- Bayesian Additive Regression Trees (BART; Chipman et al., 2010) is a Bayesian model based on a sum of trees
- Very good prediction properties
- Deals with interactions and non-linear relationships
- Based on a probability distribution so 'easily' extendable
- A Bayesian model, so have probabilistic uncertainty intervals for any required quantities
- A focus of research, with many extended versions already proposed in the literature

$$y = \sum_{j=1}^{M} g(X, T_j, \Theta_j, \mu_j) + \varepsilon; \quad \varepsilon \sim \mathsf{N}(0, \sigma^2)$$

- y is the response and X are the covariates
- *T* is the tree structure (parameter)
- ullet Θ is a set of split variables and values (parameters)
- μ is the set of terminal node values (parameters)
- The algorithm works by guessing initial values of all these (usually a stump), then proposing and accepting/rejecting new trees
- The big problem that occurs here is that the number of parameters changes when the tree structure changes
- The main reason that BART works is that we can collapse over the tree structure when we use a normally distributed prior on the μ terminal node parameters

- Standard BART MCMC uses a back-fitting approach where each tree is fixed and the others updated in turn
- Requires us to propose and accept/reject a new tree at each iteration for M trees
- We put extra prior distributions on the size and shape of the trees to keep them small, and to keep the μ values from dominating the predictions
- Default BART has four moves to generate new trees, all of which are reversible.
 - Grow: picks a random terminal node and splits it in two by choosing a random split variable and split value
 - Prune: picks a pair of adjacent terminal nodes and merges them
 - Swap: picks two random internal nodes and swaps their split variables and values
 - Change: picks a random internal node and changes the split variable and value

RART

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■ Tree prior can be used to favour shallow/deep trees:

$$\mathsf{P}(\mathsf{node}\ d\ \mathsf{is}\ \mathsf{non\text{-}terminal}) = \frac{\alpha}{(1+d)^\beta}$$

- Default settings of $\alpha=2$ and $\beta=0.95$ give
 - depth of 1: 0.05
 - depth of 2: 0.55
 - depth of 3: 0.28
 - depth of 4: 0.09
 - depth of 5: 0.03
- but trees will grow deeper if the data demands it!

RART

- Extensions to BART.
 - BART-BMA (Hernandez et al., 2018, Statistics and Computing)
 - Soft-BART (Linero and Yang, 2018, JRSSB)
 - MOTR-BART (Prado et al., 2021, Statistics and Computing)
 - HEBART (Wundervald et al., 2022, ArXiv)
 - AMBARTI (Prado, Sarti et al., 2023, AOAS)
 - GP-BART (Maia et al., 2024, CSDA)
 - CSP-BART (Prado et al., under review)
 - ZaNI-Multinomial BART (Menezes et al., in progress)
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Boruta

Boruta

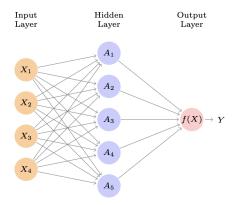
- Boruta is an algorithm used for feature selection in machine learning, helping to identify the most important variables in a dataset.
- It is a wrapper method, which means it relies on a machine learning model to evaluate the importance of features.
- Boruta specifically uses the random forest algorithm for assessing feature importance.
- It creates shadow features by shuffling the original features, maintaining their distribution but breaking any relationship with the target variable.
- The importance of original features is compared to the maximum importance of shadow features.
- This process is repeated multiple times to ensure robust selection. Each feature is given a score in each iteration.

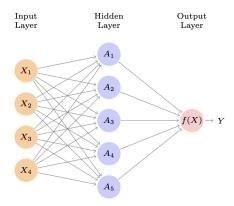
Boruta

- Decision Making:
 - Important/Confirmed: A feature is considered important if its importance is significantly higher than the shadow features across many iterations.
 - Unimportant/Rejected: A feature is deemed unimportant if its importance is consistently lower.
 - Tentative: Features that do not clearly stand out as important or unimportant are marked as tentative.
- Computationally Intensive: While robust, Boruta can be computationally intensive due to the multiple iterations and comparisons.

- Neural networks are computational models inspired by the human brain's architecture
- They consist of layers of interconnected nodes (neurons), where each connection has a weight that adjusts during training
- The network learns by adjusting these weights based on the error of its predictions compared to the actual outcomes
- The training involves iterative algorithms (e.g. backpropagation, stochastic gradient descent)
- Neural networks are widely used in fields such as computer vision, natural language processing, and predictive analytics
- Current research focusses on improving their interpretability, robustness, and efficiency, as well as exploring novel architectures like convolutional neural networks (CNNs) and recurrent neural networks (RNNs) for specialized applications

A simple neural network:





$$Y = f(X) = \beta_0 + \beta_1 A_1 + \dots + \beta_5 A_5$$

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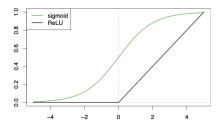
$$= \beta_0 + \sum_{k=1}^K \beta_k A_k$$

$$= \beta_0 + \sum_{k=1}^K \beta_k h_k(X)$$

$$= \beta_0 + \sum_{k=1}^K \beta_k g\left(w_{k0} + \sum_{j=1}^p w_{kj} X_j\right)$$

■ All parameters $\beta_0, \ldots, \beta_K, w_{10}, \ldots, w_{Kp}$ are estimated from the data

- $\mathbf{g}(\cdot)$ are called *activation functions*
- Different options can be used, such as the sigmoid $g(x) = \frac{1}{1 + e^{-x}}$ and ReLU (Rectified Linear Unit) $q(x) = x, x \ge 0$ and q(x) = 0, x < 0.



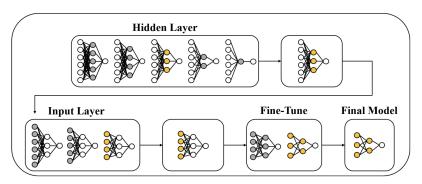
- \blacksquare The nonlinearity in q is essential to capture complex nonlinearities and interaction effects between the predictors
- **Example:** take $X = (X_1, X_2)$ and $q(x) = x^2$, with

$$egin{aligned} eta_0 &= 0, & eta_1 &= rac{1}{4}, & eta_2 &= -rac{1}{4}, \ w_{10} &= 0, & w_{11} &= 1, & w_{12} &= 1, \ w_{20} &= 0, & w_{21} &= 1, & w_{22} &= -1. \end{aligned}$$

Statistical perspective: thinking of the neural network as a statistical model

$$y_i = \mathsf{NN}(x_i) + \varepsilon_i$$

- Allows for computation of likelihood and information criteria
- R package selectnn (McInerney and Burke, 2024)



GAMLSS

GAMLSS

 Generalized Additive Models for Location, Scale, and Shape (GAMLSS: Rigby and Stasinopoulos, 2005) are a very flexible semi-parametric modelling framework

$$Y \sim f(\mu, \sigma, \nu, \tau)$$

- Includes many distributions with up to 4 parameters
- Allows for distributional regression, i.e. all parameters can be modelled with covariates
- Spline and loess smoothing, as well as random effects allowed
- Unified framework for model diagnostics

GAMLSS

- The gamlss package is the main implementation
- Extra features available through companion packages
- The package gamlss.add includes extra features, such as the inclusion of regression trees and neural networks in the linear predictor for any parameter of interest
- \blacksquare e.g. $\mu = \mathsf{NN}(x)$ or $\mu = \mathsf{CART}(x)$