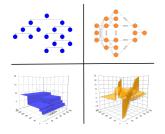
# Important Learning Algorithms in ML





## Learning goals

- General idea of important ML algorithms
- Overview of strengths and weaknesses

## CONTENTS

k-Nearest Neighbors (k-NN)

Generalized Linear Models (GLM)

Generalized Additive Models (GAM)

Classification & Regression Trees (CART)

**Random Forests** 

**Gradient Boosting** 

**Linear Support Vector Machines (SVM)** 

**Nonlinear Support Vector Machines** 

Gaussian Processes (GP)

**Neural Networks (NN)** 







## K-NN – METHOD SUMMARY

REGRESSION

CLASSIFICATION

NONPARAMETRIC

WHITE-BOX

#### General idea

- similarity in feature space (w.r.t. certain distance metric  $d(\mathbf{x}^{(i)}, \mathbf{x})$ )  $\rightarrow$  similarity in target space
- **Prediction** for **x**: construct *k*-neighborhood  $N_k(\mathbf{x})$  from *k* points closest to **x** in  $\mathcal{X}$ , then predict
  - (weighted) mean target for regression:

$$\hat{y} = \frac{1}{\sum\limits_{i:\mathbf{x}^{(i)} \in N_k(\mathbf{x})} w_i} \sum\limits_{i:\mathbf{x}^{(i)} \in N_k(\mathbf{x})} w_i y^{(i)} \text{ with } w_i = \frac{1}{d(\mathbf{x}^{(i)},\mathbf{x})}$$

- $\rightarrow$  optional: higher weights  $w_i$  for close neighbors
- most frequent class for classification:

$$\hat{y} = \underset{\ell \in \{1, \dots, g\}}{\operatorname{arg max}} \sum_{i: \mathbf{x}^{(i)} \in N_k(\mathbf{x})} \mathbb{I}(y^{(i)} = \ell)$$

 $\Rightarrow$  Estimating posterior probabilities as

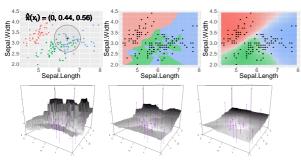
$$\hat{\pi}_{\ell}(\mathbf{x}^{(i)}) = \frac{1}{k} \sum_{i: \mathbf{x}^{(i)} \in N_k(\mathbf{x})} \mathbb{I}(\mathbf{y}^{(i)} = \ell)$$

- Nonparametric behavior: no compression of information
- Not immediately interpretable



## K-NN - METHOD SUMMARY

**Hyperparameters** Neighborhood **size** *k* (locality), **distance** metric (next page)



#### Classification

Left: Neighborhood for exemplary observation in iris, k=50 Middle: Prediction surface for k=1 Right: Prediction surface for k=50

#### Regression

Left: Prediction surface for k = 3Middle: Prediction surface for k = 7Right: Prediction surface for k = 15



- Small  $k \Rightarrow$  very local, "wiggly" decision boundaries
- Large  $k \Rightarrow$  rather global, smooth decision boundaries

## K-NN - METHOD SUMMARY

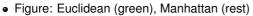
## Popular distance metrics

• Numerical feature space: Typically, Minkowski distances

$$d(\mathbf{x}, \tilde{\mathbf{x}}) = \|\mathbf{x} - \tilde{\mathbf{x}}\|_q = \left(\sum_j |x_j - \tilde{x}_j|^q\right)^{\frac{1}{q}}$$

• 
$$q=1$$
 (Manhattan):  $d(\mathbf{x}, \tilde{\mathbf{x}}) = \sum_{j} |x_j - \tilde{x}_j|$ 

• 
$$q=2$$
 (Euclidean):  $d(\mathbf{x}, \tilde{\mathbf{x}}) = \sqrt{\sum_j (x_j - \tilde{x}_j)^2}$ 







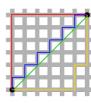
- numerical: 
$$d(x_i, x_j) = \frac{|x_i - x_j|}{\max(x) - \min(x)}$$

- categorical: 
$$d(x_i, x_j) = \begin{cases} 1, & \text{if } x_i \neq x_j \\ 0, & \text{if } x_i = x_j \end{cases}$$

- Gower distance as average over individual scores

Optional weighting for beliefs about varying feature importance

Figure Source: https://es.m.wikipedia.org/wiki/Archivo:Manhattan\_distance.svg





## **K-NN – IMPLEMENTATION & PRACTICAL HINTS**

Preprocessing Features should be standardized or normalized

### Implementation

- R: mlr3 learners (calling kknn::kknn())
  - Classification:
    - LearnerClassifKKNN
    - -fnn::knn()
  - Regression:
    - LearnerRegrKKNN
    - -fnn::knn.reg()
  - Nearest Neighbour Search in O(N log N): RANN::nn2()
- Python: From package sklearn.neighbors
  - Classification:
    - KNeighborsClassifier()
    - RadiusNeighborsClassifier() as alternative if data not uniformly sampled
  - Regression:
    - KNeighborsRegressor()
    - RadiusNeighborsRegressor() as alternative if data not uniformly sampled



## K-NN - PROS & CONS

#### **Advantages**

- + Algorithm **easy** to explain and implement
- No distributional or functional assumptions

   → able to model data of arbitrary
   complexity
- + No training or optimization required
- + local model
  - → **nonlinear** decision boundaries
- + Easy to **tune** (few hyperparameters)
   → number of neighbors k, distance metric
- Custom distance metrics can often be easily designed to incorporate domain knowledge

#### **Disadvantages**

- Sensitivity w.r.t. noisy or irrelevant features and outliers due to dependency on distance measure
- Heavily affected by curse of dimensionality
- Bad performance when feature scales are not consistent with feature relevance
- Poor handling of data **imbalances** (worse for more global model, i.e., large k)







## **GLM – METHOD SUMMARY**

REGRESSION

ASSIFICATION

PARAMETR<u>IC</u>

VHITE-BOX

FEATURE SELECTION

General idea Represent target as function of linear predictor  $\theta^{\top} \mathbf{x}$  (weighted sum of features)

 $\rightarrow$  **Interpretation:** if feature  $x_j$  increases by 1 unit, the linear predictor changes by  $\theta_i$  units

Hypothesis space  $\mathcal{H} = \{f : \mathcal{X} \to \mathbb{R} \mid f(\mathbf{x}) = \phi(\boldsymbol{\theta}^{\top}\mathbf{x})\}$ , with suitable transformation  $\phi(\cdot)$ , e.g.,

- Linear Regression:  $\mathcal{Y} = \mathbb{R}$ ,  $\phi$  identity
- Logistic Regression:  $\mathcal{Y} = \{0, 1\}$ , logistic sigmoid

$$\phi(\boldsymbol{\theta}^{\top}\mathbf{x}) = \frac{1}{1 + \exp(-\boldsymbol{\theta}^{\top}\mathbf{x})} =: \pi(\mathbf{x} \mid \boldsymbol{\theta})$$

$$\Rightarrow \text{Decision rule: Linear hyperplane}$$







Logistic function for bivariate input and loss-minimal  $\boldsymbol{\theta}$ 



Corresponding separating hyperplane



## **GLM – METHOD SUMMARY**

#### Loss functions

- Lin. Regr.:
  - Typically, based on quadratic loss (OLS estimation):

$$L(y,f)=(y-f)^2$$

- Log. Regr.: Based on bernoulli / log / cross-entropy loss
  - Loss based on scores

$$L(y, f) = \ln(1 + \exp(-y \cdot f)) \text{ for } y \in \{-1, +1\}$$
  
 $L(y, f) = -y \cdot f + \log(1 + \exp(f)) \text{ for } y \in \{0, 1\}$ 

Loss based on probabilities:

$$L(y,\pi) = \ln(1 + \exp(-y \cdot \log(\pi))) \text{ for } y \in \{-1, +1\}$$
  

$$L(y,\pi) = -y \log(\pi) - (1-y) \log(1-\pi) \text{ for } y \in \{0, 1\}$$



# **GLM – METHOD SUMMARY**

# × 0 0 × × ×

#### Optimization

- Minimization of the empirical risk
- ullet For **OLS**: analytical solution  $\hat{oldsymbol{ heta}} = (\mathbf{X}^{ op}\mathbf{X})^{-1}\mathbf{X}^{ op}\mathbf{y}$
- For other loss functions:
  - Log. Regr.: Convex problem, solvable via second-order optimization methods (e.g. BFGS)
  - Else: Numerical optimization

## Multi-class extension of logistic regression

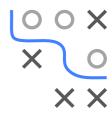
Estimate class-wise scoring functions:

$$\Rightarrow \pi: \mathcal{X} \rightarrow [0,1]^g, \ \pi(\mathbf{x}) = (\pi_1(\mathbf{x}), \dots, \pi_g(\mathbf{x})), \ \sum_{k=1}^g \pi_k(\mathbf{x}) = 1$$

• Achieved through **softmax** transformation:

$$\pi_k(\mathbf{x} \mid \boldsymbol{\theta}) = \exp(\boldsymbol{\theta}_k^{\top} \mathbf{x}) / \sum_{j=1}^g \exp(\boldsymbol{\theta}_j^{\top} \mathbf{x})$$

- Multi-class log-loss:  $L(y, \pi(\mathbf{x})) = -\sum_{k=1}^{g} \mathbb{I}_{\{y=k\}} \log(\pi_k(\mathbf{x}))$
- Predict class with maximum score (or use thresholding variant)



#### General idea

- Unregularized LM: risk of overfitting in high-dimensional space with only few observations
- Goal: avoidance of overfitting by adding penalty term

## Regularized empirical risk

- Empirical risk function **plus complexity penalty**  $J(\theta)$ , controlled by shrinkage parameter  $\lambda > 0$ :  $\mathcal{R}_{reg}(\theta) := \mathcal{R}_{emp}(\theta) + \lambda \cdot J(\theta)$
- ullet Ridge regression: L2 penalty  $J(oldsymbol{ heta}) = \|oldsymbol{ heta}\|_2^2$
- LASSO regression: L1 penalty  $J(\theta) = \|\theta\|_1$

### Optimization under regularization

- Ridge: analytically with  $\hat{\theta}_{Ridge} = (\mathbf{X}^{\top}\mathbf{X} + \lambda \mathbf{I})^{-1}\mathbf{X}^{\top}\mathbf{y}$
- LASSO: numerically with, e.g., (sub-)gradient descent

### Choice of regularization parameter

- Standard hyperparameter optimization problem
- ullet E.g., choose  $\lambda$  with minimum mean cross-validated error



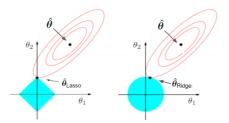
### Ridge vs. LASSO

## Ridge

- ullet Global shrinkage  $\Rightarrow$  overall smaller but still dense heta
- Applicable with large number of influential features, correlated variables' coefficients are shrinked by equal amount

#### LASSO

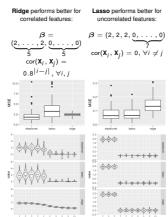
- Actual variable selection by shrinking coefficients to zero
- Suitable for sparse problems, ineffective with correlated features (randomly selecting one)





- Neither overall better ⇒ elastic net
- Weighted combination of Ridge and LASSO
- Introducing additional penalization coefficient:

$$\mathcal{R}_{\text{reg}}(\theta) = \mathcal{R}_{\text{emp}}(\theta) + \lambda \cdot P_{\alpha}(\theta), \text{ with } P_{\alpha}(\theta) = [\alpha \cdot \|\theta\|_{1} + (1 - \alpha) \cdot \frac{1}{2} \cdot \|\theta\|_{2}^{2}]$$





## **GLM – IMPLEMENTATION**

### Implementation

- R:
  - Unregularized: mlr3 learner LearnerRegrLM, calling stats::lm()/mlr3 learner LearnerClassifLogReg, calling stats::glm()
  - Regularized / ElasticNet: mlr3 learners LearnerClassifGlmnet
     / LearnerRegrGlmnet, calling glmnet::glmnet()
  - For large classification data: mlr3 learner
     LearnerClassifLiblineaR, calling LiblineaR::LiblineaR()
     uses fast coordinate descent
- Python: From package sklearn.linear\_model
  - Unregularized:
    - LinearRegression()
    - -LogisticRegression(penalty = None)
  - Regularized:
    - Linear regression: Lasso(), Ridge(), ElasticNet()
    - Logistic regression: LogisticRegression(penalty = {'11',
      '12', 'elasticnet'})
  - Package for advanced statistical models: statsmodels.api



## **GLM - PROS & CONS**

#### **Advantages**

- + **Simple, fast** implementation
- + **Analytical** solution for L2 loss
- Applicable for any dataset size, where observations
   number of features
- + Flexibility **beyond linearity** with polynomials, trigonometric transformations, interaction terms etc.
- Intuitive interpretability via feature effects
- + Statistical hypothesis **tests** for effects available

## **Disadvantages**

- Nonlinearity of many real-world problems
- Further restrictive assumptions: linearly independent features, homoskedastic residuals, normality of conditional response
- Sensitivity: outliers, noise
- LM can **overfit** (e.g., many features and few observations)
- Feature interactions must be handcrafted
  - $\rightarrow$  practically infeasible for higher orders





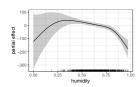


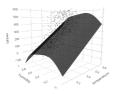
# GAM – METHOD SUMMARY CLASSIFICATION

General idea

- Same as GLM, but introduce flexibility through nonlinear (smooth) effects  $f_i(x_i)$
- Typically, combination of linear & smooth effects
- Smooth effects also conceivable for feature interactions

Hypothesis space 
$$\mathcal{H} = \left\{ f : \mathcal{X} \to \mathbb{R} \mid f(\mathbf{x}) = \phi \left( \theta_0 + \sum_{j=1}^p f_j(x_j) \right) \right\}$$
, suitable transformation  $\phi(\cdot)$ , intercept  $\theta_0$ , smooth functions  $f_i(\cdot)$ 





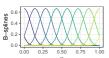
Prediction of bike rentals from smooth term of humidity (left: partial effect) and linear term of temperature (right: bivariate prediction).

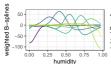


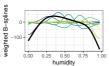
## **GAM – METHOD SUMMARY**

#### Smooth functions

- Non-/semi-/parametric approaches conceivable
- Frequently: express f<sub>j</sub> as weighted sum of basis functions → model linear in weight coefficients again
  - Use fixed basis of functions b<sub>1</sub>,..., b<sub>K</sub> and estimate associated coefficients γ<sub>1</sub>,..., γ<sub>K</sub>
     → f<sub>i</sub>(x<sub>i</sub>) = ∑<sup>K<sub>i</sub></sup><sub>k=1</sub> γ<sub>i,k</sub>b<sub>k</sub>(x<sub>i</sub>)
  - Popular types of basis functions
    - Polynomial → smoothing/TP-/B-splines
    - Radial → Kriging
    - Trigonometric → Fourier/wavelet forms
- Alternatives: local regression (LOESS), kernel-smoothing, . . .







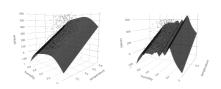
Left: B-spline basis with 9 basis functions. Middle: BFs weighted with coefficients estimated for humidity. Right: sum of weighted BFs in black (= partial effect).



## **GAM – METHOD SUMMARY**

#### Regularization

- Smooth functions possibly very flexible → regularization vital to prevent overfitting
- Control smoothness
  - Basis-function approaches: control number; impose penalty on coefficients (e.g., magnitude or differences between coefficients of neighboring components) & control associated hyperparameter



Prediction surfaces for bike rentals with 9 (left) and 500 (right) basis functions in smooth humidity term. Higher number of basis functions yields more local, less smooth model.



# **GAM - METHOD SUMMARY**



#### Loss functions

- Same as in GLM
  - → essentially: use negative log-likelihood

#### Optimization

- Coefficients (of smooth + linear terms): penalized MLE, Bayesian inference
- Smoothing hyperparameters: typically, generalized cross-validation

## **GAM – IMPLEMENTATION**

# × 0 0 × × ×

#### Implementation

- R: mlr3 learner LearnerRegrGam, calling mgcv::gam()
  - Smooth terms: s(..., bs="<basis>") or te(...) for multivariate (tensorproduct) effects
  - Link functions: family={Gamma, Binomial, ...}
- Python: GLMGam from package statsmodels; package pygam

## **GAM - PROS & CONS**

#### **Advantages**

- + Simple and fast
- + Applicable for any dataset size, where number of observations
   ≫ number of features
- + High **flexibility** via smooth effects
- + Easy to **combine** linear & nonlinear effects
- + Rather intuitive **interpretability** via feature effects
- + Statistical hypothesis **tests** for effects available

### **Disadvantages**

- Sensitivity w.r.t. outliers and noisy data
- Feature interactions must be handcrafted
  - → practically infeasible for higher orders
- Harder to optimize than GLM
- Additional hyperparameters (type of smooth functions, smoothness degree, ...)







## **CART – METHOD SUMMARY**

REGRESSION

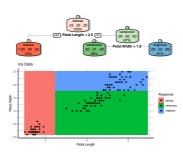
CLASSIFICATION

NONPARAMETRIC

WHITE-BC

General idea

- Start at root node containing all data
- Perform repeated axis-parallel binary splits in feature space to obtain rectangular partitions at terminal nodes Q<sub>1</sub>,..., Q<sub>M</sub>
- Splits based on reduction of node impurity
  - → empirical risk minimization (ERM)
- In each step:
  - Find optimal split (feature-threshold combination)
    - $\rightarrow$  greedy search
  - Assign constant prediction  $c_m$  to all obs. in  $Q_m$ 
    - $\rightarrow$  Regression:  $c_m$  is average of y
    - $\rightarrow$  Classif.:  $c_m$  is majority class (or class proportions)
  - Stop when a pre-defined criterion is reached
    - → See Complexity control



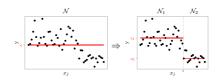


## **CART – METHOD SUMMARY**

Hypothesis space 
$$\mathcal{H} = \left\{ f(\mathbf{x}) : f(\mathbf{x}) = \sum_{m=1}^{M} c_m \mathbb{I}(\mathbf{x} \in Q_m) \right\}$$
  
Empirical risk

 Splitting feature x<sub>j</sub> at split point t divides a parent node N<sub>p</sub> into two child nodes:

$$N_{I} = \{(\mathbf{x}, y) \in N_{p} : x_{j} \leq t\}$$
 and  $N_{r} = \{(\mathbf{x}, y) \in N_{p} : x_{j} > t\}$ 



• Compute empirical risks in child nodes and minimize their sum to find best split (impurity reduction):

$$\underset{j,t}{\arg\min} \mathcal{R}(N_p, j, t) = \underset{j,t}{\arg\min} \mathcal{R}(N_t) + \mathcal{R}(N_r)$$
 (1)

Note: If  $\mathcal{R}$  is the average instead of the sum of loss functions, we need to reweight:  $\frac{|N_{pl}|}{|N|}\mathcal{R}(N_{pl})$ 



## **CART – METHOD SUMMARY**

 $\times$ 

- In general, compatible with arbitrary losses typical choices:
  - g-way classification:

$\textbf{Brier score} \rightarrow \textbf{Gini} \text{ impurity}$	<b>Bernoulli</b> loss $ o$ <b>entropy</b> impurity
$\mathcal{R}(N_p) = \sum_{(\mathbf{x}, y) \in N_p} \sum_{k=1}^g \hat{\pi_k}^{(N_p)} (1 - \hat{\pi_k}^{(N_p)})$	$\mathcal{R}(N_p) = -\sum_{(\mathbf{x}, \mathbf{y}) \in N_p} \sum_{k=1}^g \hat{\pi_k}^{(N_p)} \log \hat{\pi_k}^{(N_p)}$

Regression (quadratic loss):

$$\mathcal{R}(N_p) = \sum_{(\mathbf{x}, y) \in N_p} (y - c)^2$$
 with  $c = \frac{1}{|N_p|} \sum_{(\mathbf{x}, y) \in N_p} y$ 

#### Optimization

- Exhaustive search over all split candidates, choice of risk-minimal split
- In practice: reduce number of split candidates (e.g., using quantiles instead of all observed values)

# **CART – IMPLEMENTATION & PRACTICAL HINTS**

## Hyperparameters and complexity control

- Unless interrupted, splitting continues until we have pure leaf nodes (costly + overfitting)
- Hyperparameters: Complexity (i.e., number of terminal nodes) controlled via tree depth, minimum number of observations per node, maximum number of leaves, minimum risk reduction per split, ...
- Limit tree growth / complexity via
  - Early stopping: stop growth prematurely
    - $\rightarrow$  hard to determine good stopping point without many guesses
  - Pruning: grow deep trees; prune in risk-optimal manner afterwards



# **CART – IMPLEMENTATION & PRACTICAL HINTS**

## **Implementations**

- R:
  - CART: mlr3 learners LearnerClassifRpart / LearnerRegrRpart, calling rpart::rpart()
  - Conditional inference trees: partykit::ctree()
    mitigates overfitting by controlling tree size via p-value-based
    splitting
  - Model-based recursive partitioning: partykit::mob()
     fits a linear model within each terminal node of the decision tree
  - Rule-based models: Cubist::cubist() for regression and
     C50::C5.0() for classification; more flexible frameworks for fitting various types of models (e.g., GLMs) within a tree's terminal nodes
- Python: DecisionTreeClassifier / DecisionTreeRegressor from package scikit-learn



## **CART – DUAL PURPOSE**



### **Dual purpose of CART**

- Exploration purpose to obtain interpretable decision rules (here: performance/tuning is secondary)
- Prediction model: CART as base learner in ensembles (bagging, random forest, boosting) can improve stability and performance (if tuned properly), but becomes less interpretable

## **CART - PROS & CONS**

### **Advantages**

- + **Easy** to understand & visualize (interpretable)
- + Built-in feature selection
   → e.g., when features are not used for splitting
- Applicable to categorical features
   → e.g., 2<sup>m</sup> possible binary splits for m categories
  - $\rightarrow$  trick for regr. with L2-loss and binary classif.: categories can be sorted  $\Rightarrow m-1$  binary splits
- Handling of missings possible via surrogate splits
- + Models interactions.
- + Fast well scalable
- + High **flexibility** with custom split criteria or leaf-node prediction rules

## **Disadvantages**

- Rather poor generalization
- High variance/instability: model can change a lot when training data is minimally changed
- Can **overfit** if tree is grown too deep
- Not well-suited to model linear relationships
- Bias toward features with many unique values or categories





# **Random Forests**

## RANDOM FORESTS – METHOD SUMMARY

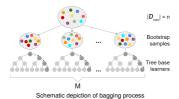
CLASSIFICATION

NONPARAMETRIC

General idea

FEATURE SELECTION

- Bagging ensemble of M tree base learners fitted on bootstrap data samples
  - ⇒ Reduce **variance** by ensembling while slightly increasing **bias** by bootstrapping
    - Use unstable, **high-variance** base learners by letting trees grow to full size
    - Promoting decorrelation by random subset of candidate features for each split
- Predict via averaging (regression) or majority vote (classification) of base learners



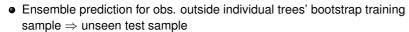


Prediction surface for iris data with 500-tree ensemble



## RANDOM FORESTS – METHOD SUMMARY

Hypothesis space 
$$\mathcal{H} = \left\{ f(\mathbf{x}) : f(\mathbf{x}) = \frac{1}{M} \sum_{m=1}^{M} \sum_{t=1}^{T^{[m]}} \mathbf{c}_t^{[m]} \mathbb{I}(\mathbf{x} \in Q_t^{[m]}) \right\}$$
 Empirical risk & Optimization Just like tree base learners Out-of-bag (OOB) error



- Use resulting loss as unbiased estimate of generalization error
- Mainly useful for tuning and less for model comparison as we usually compare all models uniformly by CV



## RANDOM FORESTS – METHOD SUMMARY

#### Feature importance

- Based on improvement in split criterion: aggregate improvements by all splits using j-th feature
- Based on **permutation:** permute *j*-th feature in OOB observations and compute impact on OOB error

## Hyperparameters

- Ensemble size, i.e., number of trees
- Complexity of base learners, e.g., tree depth, min-split, min-leaf-size
- Number of split candidates
  - i.e., number of features to be considered at each split
  - $\Rightarrow$  frequently used heuristics with total of p features:  $\left\lfloor \sqrt{p} \right\rfloor$  for classification,  $\left\lfloor p/3 \right\rfloor$  for regression



## **RANDOM FORESTS – PRACTICAL HINTS**

## **Extremely Randomized Trees**

- Variance of trees can be further increased by randomizing split points instead of using the optimal one
- Alternatively consider k random splits and pick the best one according to impurity

#### **Tuning**

- Ensemble size should not be tuned as it only decreases variance —
   choose sufficiently large ensemble
- While default values for number of split points is often good, tuning it can still improve performance
- Tuning the minimum samples in leafs and minimum samples for splitting can be beneficial but usually no huge performance increases



## **RANDOM FORESTS – IMPLEMENTATION**



#### Implementation

- R:mlr3 learners LearnerClassifRanger / LearnerRegrRanger, calling ranger::ranger() as a highly efficient and flexible implementation
- **Python:** RandomForestClassifier / RandomForestRegressor from package scikit-learn

## **RANDOM FORESTS – PROS & CONS**

#### **Advantages**

- + Retains most of **trees**' advantages (e.g., feature selection, feature interactions)
- + Fairly **good predictor**: mitigating base learners' variance through bagging
- + Quite **robust** w.r.t. small changes in data
- Good with high-dimensional data, even in presence of noisy features
- + Easy to parallelize
- Robust to its hyperparameter configuration
- + Intuitive measures of **feature importance**

## **Disadvantages**

- Loss of individual trees' interpretability
- Can be suboptimal for regression when extrapolation is needed
- Bias toward selecting features with many categories (same as CART)
- Rather large model size and slow inference time for large ensembles
- Typically inferior in performance to tuned gradient tree boosting.





# **Gradient Boosting**

## **GRADIENT BOOSTING – METHOD SUMMARY**

REGRESSION

CLASSIFICATION

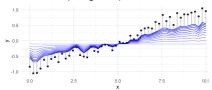
(NON)PARAMETRIC

BLACK-BOX

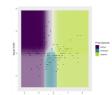
FEATURE SELECTION

General idea

- Sequential ensemble of M base learners by greedy forward stagewise additive modeling
  - In each iteration a base learner is fitted to current pseudo residuals
     one boosting iteration is one approximate gradient step in function space
  - Base learners are typically trees, linear regressions or splines
- Predict via (weighted) sum of base learners



Boosting prediction function with GAM base learners for univariate regression problem after 10 iterations



Boosting prediction surface with tree base learners for iris data after 100 iterations (right: contour lines of discriminant functions)



## **GRADIENT BOOSTING – METHOD SUMMARY**

Hypothesis space 
$$\mathcal{H} = \left\{ f(\mathbf{x}) : f(\mathbf{x}) = \sum_{m=1}^{M} \beta m b(\mathbf{x}, \theta m) \right\}$$
  
Empirical risk

- In general, compatible with any differentiable loss
- Base learner in iteration m is fitted on **Pseudo residuals**:

$$\tilde{r}^{(i)} = -\frac{\partial L(\mathbf{y}^{(i)}, \mathbf{f}(\mathbf{x}^{(i)}))}{\partial f(\mathbf{x}^{(i)})}$$
 by minimizing the **L2-loss**:  $\sum_{i=1}^{n} (\tilde{r}^{(i)} - b(\mathbf{x}^{(i)}, \boldsymbol{\theta}))^2$ 

#### Optimization

- Same optimization procedure as base learner, while keeping the current ensemble fmdh fixed
  - ⇒ Efficient and generally applicable since inner loss is always L2
- $\beta m$  is found via **line search** or fixed to a **small constant value** and combined with the leaf values ctm for tree base learners:  $ctmt = \beta m \cdot ctm$



## **GRADIENT BOOSTING – METHOD SUMMARY**

## **Hyperparameters**

- Ensemble size, i.e., number of base learners
- Complexity of base learners (depending on type used)
- Learning rate  $\beta$ , i.e., impact of next base learner

#### **Scalable Gradient Boosting**

- Feature and data subsampling for each base learner fit
- Parallelization and approximate split finding for tree base learners
- GPU accelaration



## **GRADIENT BOOSTING – PRACTICAL HINTS**

#### **Explainable / Componentwise Gradient Boosting**

- Base learners of simple linear regression models or splines, selecting a single feature in each iteration
- Allows feature selection and creates an interpretable model since uniand bivariate effects can be visualized directly.
- Feature interactions can be learned via ranking techniques (e.g., GA<sup>2</sup>M, FAST)

#### Tuning

- Use early-stopping to determine ensemble size
- Various regularization parameters, e.g., L1/L2, number of leaves, ... that need to be carefully tuned
- Tune learning rate and base learner complexity hyperparameters on log-scale



## **GRADIENT BOOSTING – IMPLEMENTATION**

#### **Gradient Tree Boosting**

- R:mlr3 learners LearnerClassifXgboost / LearnerRegrXgboost, LearnerClassifLightGBM / LearnerRegrLightGBM
- Python: GradientBoostingClassifier / GradientBoostingRegressor from package scikit-learn, XGBClassifier / XGBRegressor from package xgboost, lgb.train from package lightgbm
- $\Rightarrow \mathtt{LightGBM}$  current state-of-the-art but slightly more complicated to use than  $\mathtt{xgboost}$

#### **Componentwise Gradient Boosting**

- R: mboost from package mboost, boostLinear / boostSplines from package compboost
- Python: /
- $\Rightarrow$  mboost very flexible but slow while compboost is much faster with limited features



## **GRADIENT BOOSTING - PROS & CONS**

## **Advantages**

- Retains most of base learners' advantages
- Very good predictor due to aggressive loss minimization, typically only outperformed by heterogenous stacking ensembles
- High flexibility via custom loss functions and choice of base learner
- Highly efficient implementations exist (lightgbm / xgboost) that work well on large (distributed) data sets
- + Componentwise boosting: Good combination of
  - (a) high performance
  - (b) interpretable model
  - (c) feature selection

#### **Disadvantages**

- Loss of base learners' potential interpretability
- Many hyperparameters to be carefully tuned
- Hard to **parallelize** (→ solved by efficient implementation)







## **LINEAR SVM - METHOD SUMMARY**

CLASSIFICATION

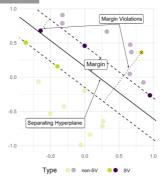
EGRESSION

PARAMETRIC

WHITE-BOX

## General idea (Soft-margin SVM)

- Find linear decision boundary (separating hyperplane) that
  - maximizes distance (margin γ) to closest points (support vectors, SVs) on each side of decision boundary
  - while minimizing margin violations (points either on wrong side of hyperplane or between dashed margin line and hyperplane)





Soft-margin SVM with margin violations

- 3 types of training points
  - non-SVs with no impact on decision boundary
  - SVs that are margin violators and affect decision boundary
  - SVs located exactly on dashed margin lines and affect decision boundary

## **LINEAR SVM – METHOD SUMMARY**

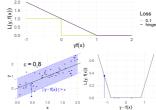
Hypothesis space (primal)  $\mathcal{H} = \{ f(\mathbf{x}) : f(\mathbf{x}) = \boldsymbol{\theta}^{\top} \mathbf{x} + \theta_0 \}$ 

#### **Empirical risk**

Soft-margin SVM as **L2-regularized ERM**:

$$\frac{1}{2} \|\boldsymbol{\theta}\|_{2}^{2} + C \sum_{i=1}^{n} L(y^{(i)}, f(\mathbf{x}^{(i)}))$$

- $\|\theta\| = 1/\gamma$  ( $\hat{=}$  maximizing margin)
- C > 0: penalization for margin violations
- Loss aims at minimizing margin violations
  - $\rightarrow$  Classif. (**hinge** loss):  $L(y, f) = \max(1 yf, 0)$
  - $\rightarrow$  Regr. ( $\epsilon$ -insensitive loss):  $L(y, f) = \max(|y f| \epsilon, 0)$





## **LINEAR SVM – METHOD SUMMARY**

**Dual problem** SVMs as a constraint optimization (primal) problem (maximize margin s.t. constraints on obs. to limit margin violations) can be formulated as a Lagrangian dual problem with Lagrange multipliers  $\alpha_i \geq 0$ :

$$\max_{\alpha v \in \mathbb{R}^n} \Leftrightarrow \quad \text{s.t.} \quad 0 \leq \alpha_i \leq \textit{C} \ \, \forall i \in \{1, \dots, n\} \text{ and } \sum_{i=1}^n \alpha_i \textit{y}^{(i)} = 0$$

**Solution** Non-SVs have  $\alpha_i = 0$  as they do not affect the hyperplane

$$f(\mathbf{x}) = \sum_{i=1}^{n} \alpha_i \mathbf{y}^{(i)} \langle \mathbf{x}^{(i)}, \mathbf{x} \rangle + \theta_0$$

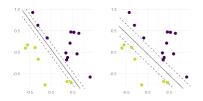


## **LINEAR SVM - METHOD SUMMARY**

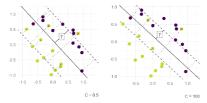
#### Optimization

- Typically, tackling dual problem (though feasible in corresponding primal)
   via quadratic programming
- Popular: sequential minimal optimization ⇒ iterative algorithm based on breaking down objective into bivariate quadratic problems with analytical solutions

**Hyperparameters** Cost parameter *C* to control maximization of the margin vs. minimizing margin violations



Hard-margin SVM: margin is maximized by boundary on the right



Soft-margin SVM: large margin and few margin violations on the right (best trade-off)



## **LINEAR SVM - PRACTICAL HINTS**

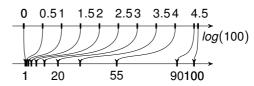


#### Preprocessing

Features should be scaled before applying SVMs (applies generally to regularized models)

## **Tuning**

- Tuning of cost parameter C advisable
   ⇒ strong influence on resulting
   hyperplane
- C it is often tuned on a log-scale grid for optimal and space-filling search space



## **LINEAR SVM - IMPLEMENTATION**

## Implementation

- R: mlr3 learners LearnerClassifSVM / LearnerRegrSVM, calling e1071::svm() with linear kernel (libSVM interface). Further implementations in mlr3extralearners based on
  - kernlab::ksvm() allowing custom kernels
  - LiblineaR::LiblineaR() for a fast implementation with linear kernel
- Python: sklearn.svm.SVC from package scikit-learn/package libSVM





# **Nonlinear Support Vector Machines**

## **NONLINEAR SVM – METHOD SUMMARY**

NONPARAMETRIC

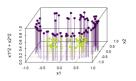
**BLACK-BOX** 

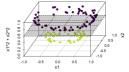
#### General idea

- Move beyond linearity by mapping data to transformed space where they are linearly separable
- Kernel trick
  - No need for explicit construction of feature maps
  - Replace inner product of feature map  $\phi: \mathcal{X} \to \Phi$  by kernel:  $\langle \phi x, \phi xt \rangle = kxxt$









Mapping to 3D space and subsequent linear separation - implicitly handled by kernel in nonlinear



## **NONLINEAR SVM - METHOD SUMMARY**

Hypothesis space

#### Primal:

$$\mathcal{H} = \{ f(\mathbf{x}) : f(\mathbf{x}) = \operatorname{sign} (\boldsymbol{\theta}^{\top} \phi \mathbf{x} + \theta_0) \}$$

#### Dual:

$$\mathcal{H} = \left\{ f(\mathbf{x}) : f(\mathbf{x}) = \text{sign}\left(\sum_{i=1}^{n} \alpha_i y^{(i)} k(\mathbf{x}^{(i)}, \mathbf{x}) + \theta_0\right) \mid \alpha_i \geq 0, \sum_{i=1}^{n} \alpha_i y^{(i)} = 0 \right\}$$
Dual problem Kernelize dual (soft-margin) SVM problem, replacing all inner

products by kernels:

$$\max_{\alpha v} \sum_{i=1}^{n} \alpha_{i} - \frac{1}{2} \sum_{i=1}^{n} \sum_{j=1}^{n} \alpha_{i} \alpha_{j} y^{(i)} y^{(j)} k(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}), \text{ s.t. } 0 \leq \alpha_{i} \leq C, \sum_{i=1}^{n} \alpha_{i} y^{(i)} = 0.$$

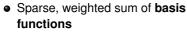
**Hyperparameters** Cost *C* of margin violations, kernel hyperparameters (e.g., width of RBF kernel)



## NONLINEAR SVM – METHOD SUMMARY

Interpretation as basis function approach

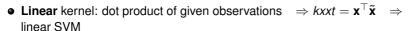
 Representer theorem: solution of dual soft-margin SVM problem is  $\theta = \sum_{i=1}^n \beta_i \phi(\mathbf{x}^{(i)})$ 



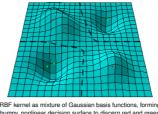
$$\rightarrow \beta_j = 0$$
 for non-SVs

 Result: local model with smoothness. depending on kernel

Common kernels



- **Polynomial** kernel of degree  $d \in \mathbb{N}$ : monomials (i.e., feature interactions) up to d-th order  $\Rightarrow kxxt = (\mathbf{x}^{\top}\tilde{\mathbf{x}} + b)^d$ , b > 0
- Radial basis function (RBF) kernel: infinite-dimensional feature space, allowing for perfect separation of all finite datasets  $\Rightarrow kxxt = \exp(-\gamma \|\mathbf{x} - \tilde{\mathbf{x}}\|_2^2)$  with bandwidth parameter  $\gamma > 0$



RBF kernel as mixture of Gaussian basis functions, forming bumpy, nonlinear decision surface to discern red and green



## **NONLINEAR SVM - IMPLEMENTATION & HINTS**

#### **Tuning**

- High sensitivity w.r.t. hyperparameters, especially those of kernel
   tuning very important
- For RBF kernels, use RBF sigma heuristic to determine bandwidth

#### Implementation

- R: mlr3 learners LearnerClassifSVM / LearnerRegrSVM, calling e1071::svm() with nonlinear kernel (libSVM interface), kernlab::ksvm() allowing custom kernels
- Python: sklearn.svm.SVC from package scikit-learn / package libSVM



## SVM - PRO'S & CON'S

## **Advantages**

- + Often **sparse** solution (w.r.t. observations)
- Robust against overfitting (regularized); especially in high-dimensional space
- Stable solutions (w.r.t. changes in train data)
  - ightarrow Non-SV do not affect decision boundary
- - minimum

#### **Disadvantages**

- **Long** training times  $\rightarrow O(n^2p + n^3)$
- Confined to linear model
- Restricted to continuous features
- Optimization can also fail or get stuck



## **NONLINEAR SVM - PRO'S & CON'S**



## Advantages (nonlinear SVM)

- + Can learn nonlinear decision boundaries
- + **Very flexible** due to custom kernels
  - ightarrow RBF kernel yields local model
  - $\rightarrow$  kernel for time series, strings etc.

#### **Disadvantages (nonlinear SVM)**

- Poor interpretability due to complex kernel
- Not easy tunable as it is highly important to choose the right kernel (which also introduces further hyperparameters)





## **GAUSSIAN PROCESSES (GP) – SUMMARY**

REGRESSION

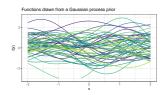
CLASSIFICATION

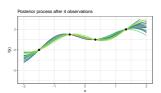
NONPARAMETRIC

**PROBABILISTIC** 

#### General idea

- GPs model a distribution over potential functions f that fit the observed data
- Assumptions:
  - *n*-observations follow a *n*-dimensional Normal distribution
  - The closer observations are, the higher they are correlated
- A **kernel** function  $k(\mathbf{x}^{(i)}, \mathbf{x}^{(j)})$  quantifies the similarity between two observations and induces the covariance matrix of the distribution.
- **Predict** via the maximum a-posteriori (MAP) estimate.







## GAUSSIAN PROCESSES (GP) – SUMMARY

Hypothesis space 
$$\mathcal{H} = \left\{ \boldsymbol{f} = \left[ f\left(\boldsymbol{x}^{(1)}\right), \ldots, f\left(\boldsymbol{x}^{(n)}\right) \right] \sim \mathcal{N}\left(\boldsymbol{m}, \boldsymbol{K}\right) \mid \boldsymbol{m} \in \mathbb{R}^{n}, \boldsymbol{K} \in \mathbb{R}^{n \times n} \right\}$$

## **Empirical risk**

- The risk is estimated by using the posterior of a conditional Normal distribution
- Most kernels have length scale parameters that need to be estimated

#### Optimization

- The kernel parameters can be learned using maximum likelihood estimation
- This requires inverting the  $n \times n$  -covariance matrix



## **GP – IMPLEMENTATION & PRACTICAL HINTS**

## Hyperparameters

- The most important hyperparameter is the choice of the kernel function  $k(\mathbf{x}^{(i)},\mathbf{x}^{(j)})$
- Common kernel choices for "standard" data are:
  - Linear or polyomial
  - Squared-exponential (infinitely differentiable)
  - Matérn (further generalization of the Squared-exponential kernel)
- Special kernels for all kind of data situation exist, e.g., a Exp-Sine-Squared kernel for periodic data
- Kernels can be composed by multiplying or addition to create more expressive structures



## **GP – IMPLEMENTATION & PRACTICAL HINTS**

## Scalable GPs for larger data

- Low-rank approximations of the covariance by using only a representative subset of inducing points
- Using a kernel that creates a sparse coviariance matrix

## **Noisy GPs**

- Having an interpolator might not be suitable if the data is noisy
- A noisy GP adds a **nugget** effect to the kernel  $k(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) + \sigma \delta_{ij}$ , creating a Gaussian process regression model

#### Implementation

- R:mlr3 learners LearnerClassifGausspr / LearnerRegrGausspr, calling kernlab::gausspr()
- Python: GaussianProcessClassifier /
  GaussianProcessRegressor from package scikit-learn, gpytorch
  for a modular, scalable, efficient and GPU accelerated implementation
  built on torch



## **GP - PROS & CONS**

#### **Advantages**

- GPs allow to quantify prediction uncertainty induced by both intrinsic noise in the problem and errors in the parameter estimation process
- A GP is a function interpolator and will predict the exact value of a training point
- The choice of kernel function allows considerable flexibility for problem specific characteristics
- + Automatic relevance determination (ARD) determines the importance of features

## **Disadvantages**

- GPs are **not sparse**, i.e., they require the full training data for prediction
- GP training requires  $\mathcal{O}(n^3)$ , i.e., it scales cubically in the number of observations
- GPs cannot handle categorical features.
- GPs are not particularly easy to understand conceptually





## **Neural Networks (NN)**

REGRESSION

CLASSIFICATION

(NON)PARAMETRIC

BLACK-BOX

General idea

- Learn composite function through series of nonlinear feature transformations, represented as neurons, organized hierarchically in layers
  - Basic neuron operation: 1) affine **transformation**  $\phi$  (weighted sum of inputs), 2) nonlinear **activation**  $\sigma$
  - Combinations of simple building blocks to create a complex model
- Optimize via mini-batch stochastic gradient descent (SGD) variants:
  - Gradient of each weight can be inferred from the computational graph of the network
    - → **Automatic Differentiation** (AutoDiff)
  - Updating weights based on the loss is done via Backpropagation

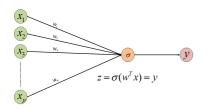
Hypothesis space

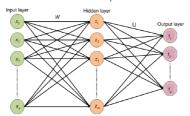
$$\mathcal{H} = \left\{ f(\mathbf{x}) : f(\mathbf{x}) = \tau \circ \phi \circ \sigma^{(h)} \circ \phi^{(h)} \circ \sigma^{(h-1)} \circ \phi^{(h-1)} \circ \dots \circ \sigma^{(1)} \circ \phi^{(1)}(\mathbf{x}) \right\}$$



#### **Architecture**

- Input layer: original features x
- Hidden layers: nonlinear transformation of previous layer  $\phi^{(h)} = \sigma^{(h-1)}(\phi^{(h-1)})$
- ullet Output layer: number of output neurons and activation depends on problem  $au(\phi)$ 
  - Regression: one output neuron,  $\tau = \text{identity}$
  - Binary classification: one output neuron,  $\tau = \frac{1}{1 + \exp(-\theta^\top \mathbf{x})}$  (logistic sigmoid)
  - Multiclass Classification: g output neurons,  $\tau_j = \frac{\exp(f_j)}{\sum_{j=1}^g \exp(f_j)}$  (softmax)







Empirical risk In general, compatible with any differentiable loss

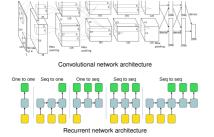
#### Optimization

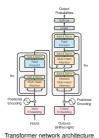
- Variety of different optimizers, mostly based on some form of stochastic gradient descent (SGD)
- Improvements:
  - (1) Accumulation of previous gradients  $\rightarrow$  Momentum
  - (2) Weight specific scaling based on previous squared gradients  $\rightarrow$  RMSProb
    - $\Rightarrow$  **ADAM** combines (1) and (2)
  - (3) Learning rate schedules, e.g., decaying or cyclical learning rates
- Training progress is measured in full passes over the full training data, called epochs
- Batch size is a hyperparameter and limited by input data dimension



Network types Large variety of architectures for different data modelities

- Feedforward NNs / multi-layer perceptrons (MLPs): sequence of fully-connected layers ⇒ tabular data
- Convolutional NNs (CNNs): sequence of feature map extractors with spatial awareness ⇒ images, time series
- Recurrent NNs (RNNs): handling of sequential, variable-length information ⇒ times series, text, audio
- Transformers: Learning invariances from data, handling multiple/any data modalities







#### Hyperparameters

#### Architecture:

- Lots of design choices ⇒ tuning problem of its own.
- Typically: hierachical optimization of components (cells) and macro structure of network
  - → Neural Architecture Search (NAS)
- Many predifined (well working) architectures exist for standard tasks

#### • Training:

- Initial learning rate and various regularization parameters
- Number of epochs is determined by early-stopping
- Data-augmentation, e.g., applying random rotations to input images





#### Foundation models

- Enormous models trained on vast amounts of (general) data, e.g., all of wikipedia, in self-supervised fashion
- Used as starting point (pre-trained) and fine-tuned via transfer or few-shot learning for other tasks requiring little data
- Examples: GPT-3 for language, CLIP for vision-language, . . .

## **NEURAL NETWORKS – PRACTICAL HINTS**

#### **General hints**

- Instead of NAS, use a standard architecture and tune training hyperparameters
- Training pipeline (data-augmentation, training schedules, ...) is more crucial than the specific architecture
- While NNets are state-of-the-art for computer vision (CV) and natural language processing (NLP), we recommend not to use them for tabular data because alternatives perform better
- Computational efforts for training (and inference) can be very high, requiring specific hardware.
  - ightarrow Using a service (esp. for foundation models) can be more cost efficient



## **NEURAL NETWORKS – IMPLEMENTATION**

# × co × x

#### Implementation

- R: Use python libraries (below) via reticulate, but not really recommended except for toy applications.
- Python libraries:
  - keras for simple high level API
  - PyTorch for flexible design with a focus on research
  - TensorFlow for flexible design with a focus on deployment / industry
  - huggingface for pre-trained / foundation models

## **NEURAL NETWORKS - PROS & CONS**

#### **Advantages**

- Applicable to complex, nonlinear problems
- + Very **versatile** w.r.t. architectures
- + State-of-the-art for CV and NLP
- + Strong **performance** if done right
- Built-in **feature extraction**, obtained by intermediate representations
- + Easy handling of **high-dimensional** data
- + **Parallelizable** training

#### **Disadvantages**

- Typically, high computational cost
- High demand for training data
- Strong tendency to overfit
- Requiring lots of tuning expertise
- Black-box model hard to interpret or explain

