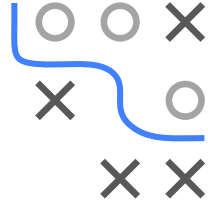
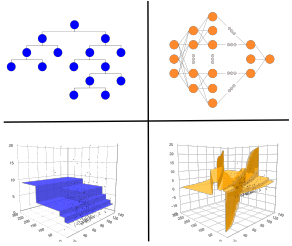
Important Learning Algorithms in ML

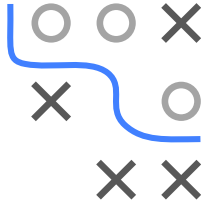




Learning goals

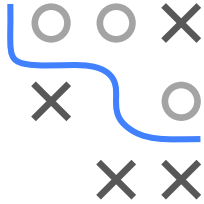
- General idea of important ML algorithms
- Overview of strengths and weaknesses

CONTENTS



K-NN – METHOD SUMMARY

REGRESSION | CLASSIFICATION | NONPARAMETRIC | WHITE-BOX



General idea

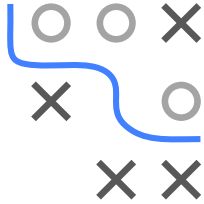
- **similarity** in feature space (w.r.t. certain **distance metric** $d(\mathbf{x}^{(i)}, \mathbf{x})$) $\rightsquigarrow$ similarity in target space
- **Prediction** for $\mathbf{x}$: construct k**-neighborhood** $N_k(\mathbf{x})$ from k points closest to $\mathbf{x}$ in $\mathcal{X}$, then predict
  - (weighted) mean target for **regression**: $\hat{y} = \frac{1}{\sum_{i:\mathbf{x}^{(i)} \in N_k(\mathbf{x})} w_i} \sum_{i:\mathbf{x}^{(i)} \in N_k(\mathbf{x})} w_i y^{(i)}$

    with $w_i = \frac{1}{d(\mathbf{x}^{(i)}, \mathbf{x})}$

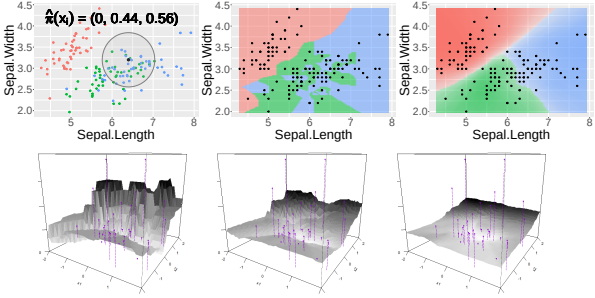
    $\rightarrow$ optional: higher weights w_i for close neighbors
  - most frequent class for **classification**: $\hat{y} = \underset{\ell \in \{1,\ldots,g\}}{\arg\max} \sum_{i:\mathbf{x}^{(i)} \in N_k(\mathbf{x})} \mathbb{I}(y^{(i)} = \ell)$

    $\Rightarrow$ Estimating posterior probabilities as $\hat{\pi}_\ell(\mathbf{x}^{(i)}) = \frac{1}{k} \sum_{i:\mathbf{x}^{(i)} \in N_k(\mathbf{x})} \mathbb{I}(y^{(i)} = \ell)$
- **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 $k = 3$
Middle: Prediction surface for $k = 7$
Right: Prediction surface for $k = 15$

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

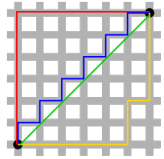
K-NN – METHOD SUMMARY

Popular distance metrics

- Numerical feature space: Typically, **Minkowski** distances

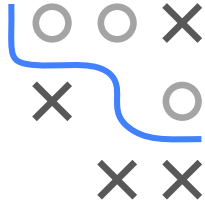
$$d(\mathbf{x}, \tilde{\mathbf{x}}) = \|\mathbf{x} - \tilde{\mathbf{x}}\|_q = \left(\sum_j |x_j - \tilde{x}_j|^q\right)^{\frac{1}{q}}$$

  - $q = 1$: **Manhattan** distance
    $\rightarrow d(\mathbf{x}, \tilde{\mathbf{x}}) = \sum_j |x_j - \tilde{x}_j|$
  - $q = 2$: **Euclidean** distance
    $\rightarrow d(\mathbf{x}, \tilde{\mathbf{x}}) = \sqrt{\sum_j (x_j - \tilde{x}_j)^2}$



Manhattan vs. Euclidean (green)

https://es.m.wikipedia.org/wiki/Archivo:Manhattan_distance.svg



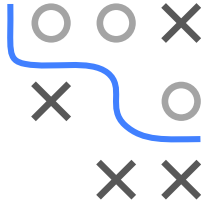
- Mixed feature space:
  - **Gower distance** for numerical, categorical and missing data:

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

    - 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

K-NN – IMPLEMENTATION & PRACTICAL HINTS

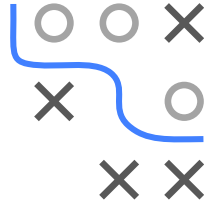


Preprocessing Features should be standardized or normalized

Implementation

- **R:** `mlr3` learners (calling `kknn::kknn()`)
  - **Classification:**
    - `LearnerClassifKKNN`
    - `fnn::knn()`
  - **Regression:**
    - `LearnerRegrKKNN`
    - `fnn::knn.reg()`
  - Nearest Neighbour Search in $\mathcal{O}(N \log N)$: `RANN::nn2()`

K-NN – IMPLEMENTATION & PRACTICAL HINTS



- **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

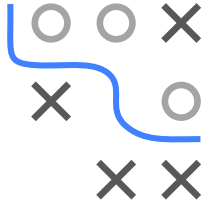
K-NN – PROS & CONS

Advantages

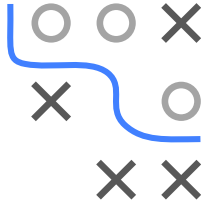
- \+ Algorithm **easy** to explain and implement
- \+ No distributional or functional **assumptions** → able to model data of **arbitrary complexity**
- \+ No **training** or **optimization** required
- \+ **local model** → **nonlinear** decision boundaries
- \+ Easy to **tune** (few hyperparameters) → 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)



GENERALIZED LINEAR MODELS – METHOD SUMMARY



REGRESSION CLASSIFICATION PARAMETRIC WHITE-BOX

FEATURE SELECTION

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

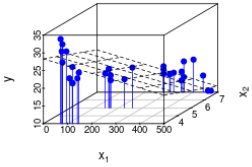
$\rightarrow$ **Interpretation:** if feature x_j increases by 1 unit, the linear predictor changes by θ_j units

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

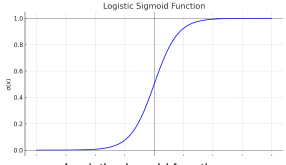
- **Linear Regression**: $\mathcal{Y} = \mathbb{R}$, ϕ identity
- **Logistic Regression**: $\mathcal{Y} = \{0, 1\}$, logistic sigmoid
  $\phi(\boldsymbol{\theta}^\top \mathbf{x}) = \frac{1}{1+\exp(-\boldsymbol{\theta}^\top \mathbf{x})} =: \pi(\mathbf{x} \mid \boldsymbol{\theta})$
  $\Rightarrow$ Decision rule: Linear hyperplane

GENERALIZED LINEAR MODELS – METHOD SUMMARY

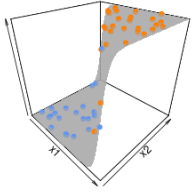
Loss functions



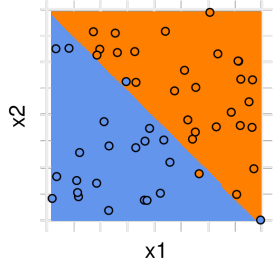
Linear regression hyperplane



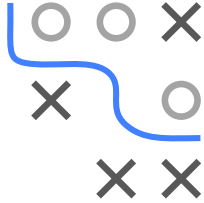
Logistic sigmoid function



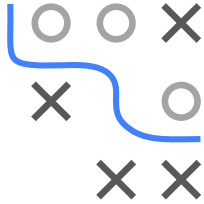
Logistic function for bivariate input and loss-minimal $\boldsymbol{\theta}$



Corresponding separating hyperplane



GENERALIZED LINEAR MODELS – METHOD SUMMARY



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

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

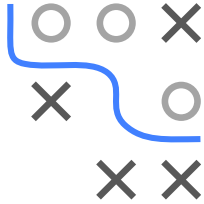
- **Log. Regr.**: Based on **bernoulli** / **log** / **cross-entropy** loss
  - Loss based on scores

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

  - Loss based on probabilities:

$$\begin{aligned} L(y, \pi) &= \ln(1 + \exp(-y \cdot \log(\pi))) \quad \text{for } y \in \{-1, +1\} \\ L(y, \pi) &= -y \log(\pi) - (1 - y) \log(1 - \pi) \quad \text{for } y \in \{0, 1\} \end{aligned}$$

GENERALIZED LINEAR MODELS – METHOD SUMMARY



Optimization

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

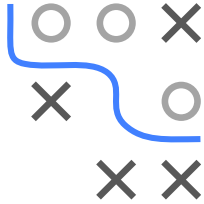
Multi-class extension of logistic regression

- Estimate **class-wise** scoring functions:

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

- 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)

GENERALIZED LINEAR MODELS – REGULARIZATION

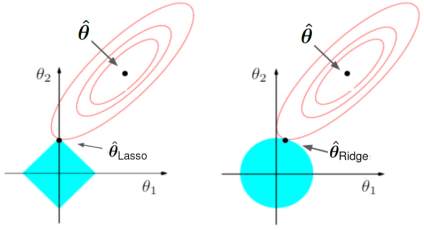


General idea

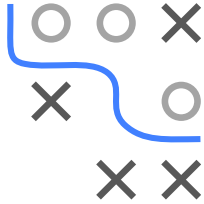
- Unregularized LM: risk of **overfitting** in high-dimensional space with only few observations
- **Goal**: avoidance of overfitting by adding **penalty term**

Regularized empirical risk

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



GENERALIZED LINEAR MODELS – REGULARIZATION



Optimization under regularization

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

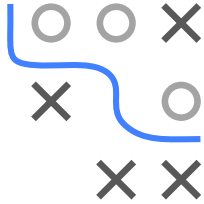
Choice of regularization parameter

- Standard hyperparameter optimization problem
- E.g., choose λ with minimum mean cross-validated error

Ridge vs. LASSO

- **Ridge**
  - Global shrinkage $\Rightarrow$ overall smaller but still dense $\boldsymbol{\theta}$
  - 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)

GENERALIZED LINEAR MODELS – REGULARIZATION



- Neither overall better $\Rightarrow$ **elastic net**
- Weighted combination of Ridge and LASSO
- Introducing additional penalization coefficient:
  $\mathcal{R}_{\text{reg}}(\boldsymbol{\theta}) = \mathcal{R}_{\text{emp}}(\boldsymbol{\theta}) + \lambda \cdot P_\alpha(\boldsymbol{\theta})$, with
  $P_\alpha(\boldsymbol{\theta}) = [\alpha \cdot \|\boldsymbol{\theta}\|_1 + (1-\alpha) \cdot \frac{1}{2} \cdot \|\boldsymbol{\theta}\|_2^2]$

Ridge performs better for correlated features:

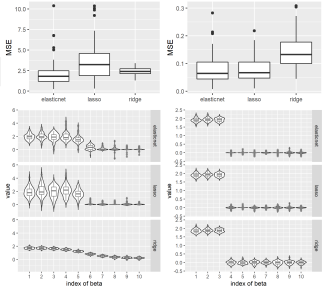
$$\boldsymbol{\beta} = (\underbrace{2, \ldots, 2}_{5}, \underbrace{0, \ldots, 0}_{5})$$

$$\text{cor}(\mathbf{X}_i, \mathbf{X}_j) = 0.8^{|i-j|}, \forall i, j$$

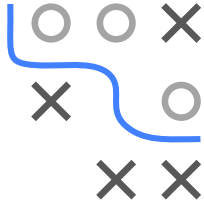
Lasso performs better for uncorrelated features:

$$\boldsymbol{\beta} = (2, 2, 2, \underbrace{0, \ldots, 0}_{7})$$

$$\text{cor}(\mathbf{X}_i, \mathbf{X}_j) = 0, \forall i \neq j$$



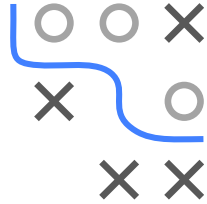
GENERALIZED LINEAR MODELS – IMPLEMENTATION



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

GENERALIZED LINEAR MODELS – IMPLEMENTATION



- **Python:** From package `sklearn.linear_model`
  - **Unregularized:**
    - `LinearRegression()`
    - `LogisticRegression(penalty = None)`
  - **Regularized:**
    - *Linear regression:* `Lasso()`,`Ridge()`,`ElasticNet()`
    - *Logistic regression:* `LogisticRegression(penalty = {'l1', 'l2', 'elasticnet'})`
  - Package for advanced **statistical** models: `statsmodels.api`

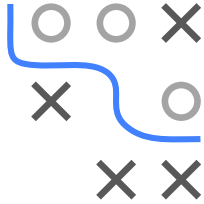
GENERALIZED LINEAR MODELS – PROS & CONS

Advantages

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

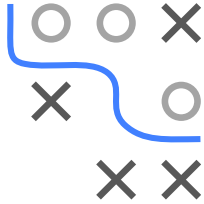
Disadvantages

- — **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



GENERALIZED ADDITIVE MODELS – METHOD SUMMARY

REGRESSION | CLASSIFICATION | (NON)PARAMETRIC | WHITE-BOX | FEATURE SELECTION

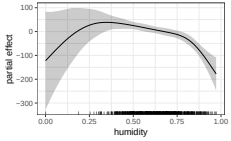


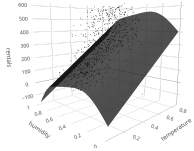
General idea

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

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

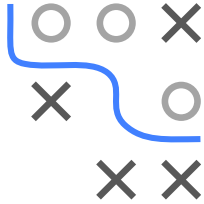
suitable transformation $\phi(\cdot)$, intercept θ_0, smooth functions $f_j(\cdot)$





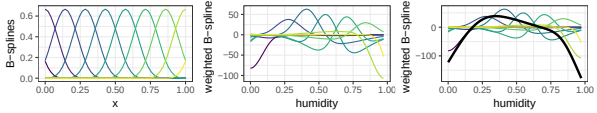
Prediction of `bike rentals` from smooth term of `humidity` (left: partial effect) and linear term of `temperature` (right: bivariate prediction).

GENERALIZED ADDITIVE MODELS – METHOD SUMMARY



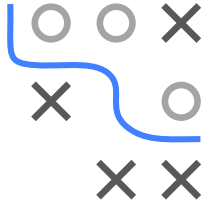
Smooth functions

- Non-/semi-/parametric approaches conceivable
- Frequently: express f_j as weighted sum of **basis functions** $\rightsquigarrow$ model **linear** in weight coefficients again
  - Use fixed basis of functions $b_1, \ldots, b_K$ and estimate associated coefficients $\gamma_1, \ldots, \gamma_K$
    $\rightsquigarrow f_j(x_j) = \sum_{k=1}^{K_j} \gamma_{j,k} b_k(x_j)$
  - Popular types of basis functions
    - Polynomial $\rightsquigarrow$ smoothing/TP-/B-**splines**
    - Radial $\rightsquigarrow$ **Kriging**
    - Trigonometric $\rightsquigarrow$ **Fourier/wavelet** forms
- Alternatives: **local regression (LOESS)**, kernel-smoothing, ...



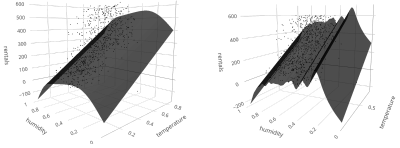
Left: B-spline basis with 9 basis functions. Middle: BFs weighted with coefficients estimated for `humidity`. Right: sum of weighted BFs in black (= partial effect).

GENERALIZED ADDITIVE MODELS – METHOD SUMMARY



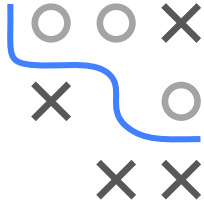
Regularization

- Smooth functions possibly very flexible $\rightsquigarrow$ 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
  - **Local smoothers**: control width of smoothing window (larger $\rightsquigarrow$ smoother)



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.

GENERALIZED ADDITIVE MODELS – METHOD SUMMARY

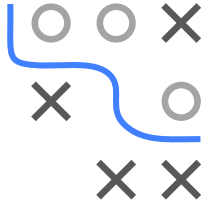


Loss functions Same as in GLM $\rightsquigarrow$ essentially: use **negative log-likelihood**

Optimization

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

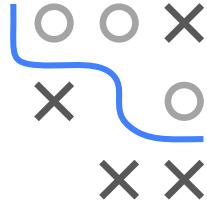
GENERALIZED ADDITIVE MODELS – IMPLEMENTATION



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`

GENERALIZED ADDITIVE MODELS – PROS & CONS



Advantages

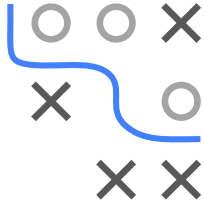
- \+ **Simple and fast**
- \+ Applicable for any **dataset size**, as long as number of observations $\gg$ number of features
- \+ High **flexibility** via smooth effects
- \+ Easy to **combine** linear & nonlinear effects
- \+ Rather intuitive **interpretability** via feature effects
- \+ Statistical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

Disadvantages

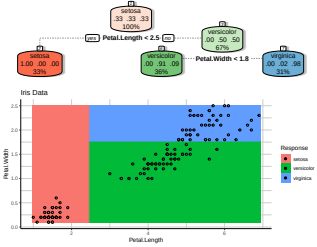
- — **Sensitivity** w.r.t. outliers and noisy data
- — Feature **interactions** must be handcrafted
  $\rightarrow$ practically infeasible for higher orders
- — Harder to **optimize** than GLM
- — Additional **hyperparameters** (type of smooth functions, smoothness degree, . . .)

CART – METHOD SUMMARY

REGRESSION | CLASSIFICATION | NONPARAMETRIC | WHITE-BOX | FEATURE SELECTION

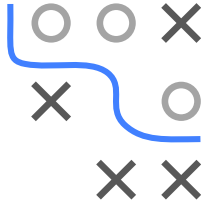


General idea (CART – Classification and Regression Trees)



- Start at root node containing all data
- Perform repeated **axis-parallel binary splits** in feature space to obtain **rectangular partitions** at terminal nodes $Q_1, \ldots, Q_M$
- Splits based on reduction of node **impurity** $\rightarrow$ 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
    $\rightarrow$ See **Complexity control**

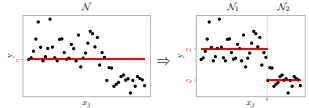
CART – METHOD SUMMARY



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_j **at split point** t divides a parent node N_p into two child nodes:



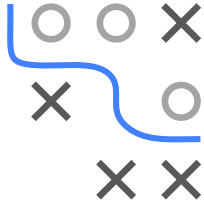
  $N_l = \{(\mathbf{x}, y) \in N_p : x_j \leq t\}$ and $N_r = \{(\mathbf{x}, y) \in N_p : x_j > t\}$

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

$$\arg\min_{j,t} \mathcal{R}(N_p, j, t) = \arg\min_{j,t} \mathcal{R}(N_l) + \mathcal{R}(N_r) \tag{1}$$

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

CART – METHOD SUMMARY



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

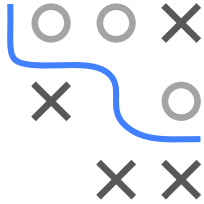
Brier score $\rightarrow$ **Gini** impurity	**Bernoulli** loss $\rightarrow$ **entropy** impurity
$\mathcal{R}(N_p) = \sum_{(\mathbf{x},y)\in N_p} \sum_{k=1}^{g} \hat{\pi}_k^{(N_p)} \left(1 - \hat{\pi}_k^{(N_p)}\right)$	$\mathcal{R}(N_p) = - \sum_{(\mathbf{x},y)\in N_p} \sum_{k=1}^{g} \hat{\pi}_k^{(N_p)} \log \hat{\pi}_k^{(N_p)}$

  - Regression (**quadratic** loss): $\mathcal{R}(N_p) = \sum_{(\mathbf{x},y)\in N_p} (y - c)^2$ with $c = \frac{1}{\lvert N_p \rvert} \sum_{(\mathbf{x},y)\in N_p} y$

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

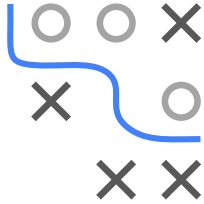


Hyperparameters and complexity control

- 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 before actually trying all combinations
  - **Pruning:** grow deep trees and cut back in risk-optimal manner afterwards

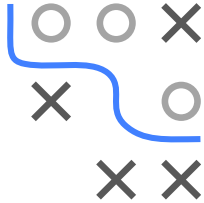
CART – IMPLEMENTATION & PRACTICAL HINTS

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 – PROS & CONS



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

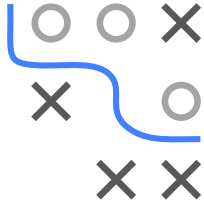
CART – PROS & CONS

Advantages

+ **Easy** to understand & visualize (**interpretable**)
+ Built-in **feature selection**
  $\rightarrow$ e.g., when features are not used for splitting
+ Applicable to **categorical** features
  $\rightarrow$ e.g., 2^m 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

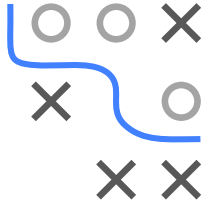
Disadvantages

− 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 – METHOD SUMMARY

REGRESSION | CLASSIFICATION | NONPARAMETRIC | BLACK-BOX | FEATURE SELECTION

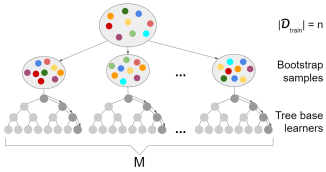


General idea

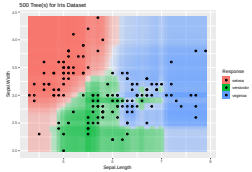
- **Bagging ensemble** of M tree **base learners** fitted on **bootstrap** data samples
  $\Rightarrow$ Reduce **variance** by ensembling while slightly increasing **bias** by bootstrapping
  - Use unstable, **high-variance** base learners by letting trees grow to full size
  - Promoting **decorrelation**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

Hypothesis space $\mathcal{H} = \left\{ f(\mathbf{x}) : f(\mathbf{x}) = \frac{1}{M} \sum_{m=1}^{M} \sum_{t=1}^{T^{[m]}} c_t^{[m]} \mathbb{I}(\mathbf{x} \in Q_t^{[m]}) \right\}$

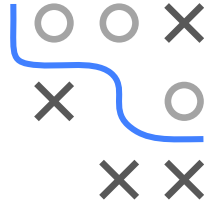
RANDOM FORESTS – METHOD SUMMARY



Schematic depiction of bagging process



Prediction surface for `iris` data with 500-tree ensemble

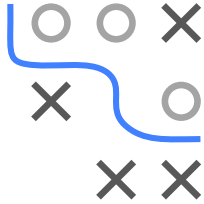


Empirical risk & Optimization Just like tree base learners

Out-of-bag (OOB) error

- Ensemble prediction for obs. outside individual trees' bootstrap training sample $\Rightarrow$ unseen test sample
- 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



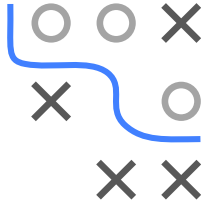
Feature importance

- 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: $\lfloor \sqrt{p} \rfloor$ for classification, $\lfloor p/3 \rfloor$ for regression

RANDOM FORESTS – IMPLEMENTATION & PRACTICAL HINTS



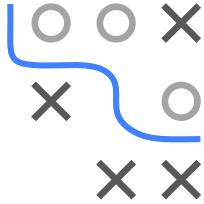
Extremely Randomized Trees

- 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 $\longrightarrow$ 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 benificial but no huge performance increases are to be expected

RANDOM FORESTS – IMPLEMENTATION & PRACTICAL HINTS



Implementation

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

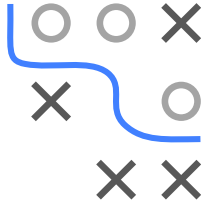
RANDOM FORESTS – PROS & CONS

Advantages

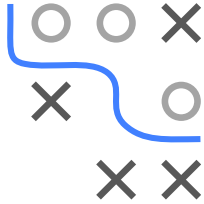
+ Retains most of **trees'** advantages (e.g., feature selection, feature interactions)
+ Fairly **good predictor**: mitigating base learners' variance through bagging
+ Quite **robust** w.r.t. small changes in data
+ Good with **high-dimensional** data, even in presence of noisy features
+ Easy to **parallelize**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

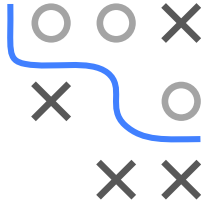


REGRESSION | CLASSIFICATION | (NON)PARAMETRIC | BLACK-BOX | FEATURE SELECTION

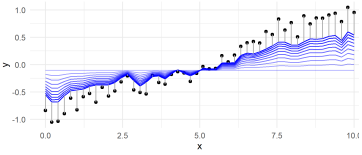
General idea

- **Sequential ensemble** of M **base learners** by greedy forward stagewise additive modeling
  - In each iteration a base learner is fitted to current **pseudo residuals** $\Rightarrow$ 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

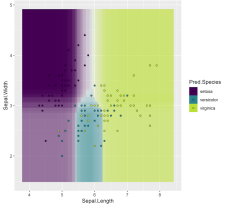
GRADIENT BOOSTING – METHOD SUMMARY



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



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

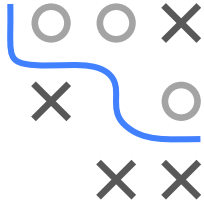


Boosting prediction surface with tree base learners for `iris` data after 100 iterations (*right:* contour lines of discriminant functions)

Empirical risk

- In general, compatible with any **differentiable** loss
- Base learner in iteration m is fitted on **Pseudo residuals**: $\tilde{r}^{(i)} = -\frac{\partial L\left(y^{(i)}, f\left(\mathbf{x}^{(i)}\right)\right)}{\partial f\left(\mathbf{x}^{(i)}\right)}$ by minimizing the **L2-loss**: $\sum_{i=1}^{n} (\tilde{r}^{(i)} - b(\mathbf{x}^{(i)}, \boldsymbol{\theta}))^2$

GRADIENT BOOSTING – METHOD SUMMARY



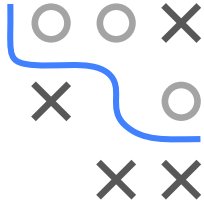
Optimization

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

Hyperparameters

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

GRADIENT BOOSTING – PRACTICAL HINTS



Scalable Gradient Boosting

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

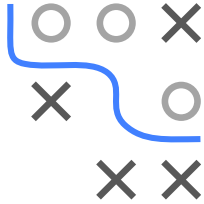
Explainable / Componentwise Gradient Boosting

- 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 uni- and bivariate effects can be visualized directly.
- Feature interactions can be learned via ranking techniques (e.g., GA^{2}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`

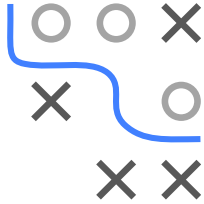
$\Rightarrow$ `LightGBM` current state-of-the-art but slightly more complicated to use than `xgboost`

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

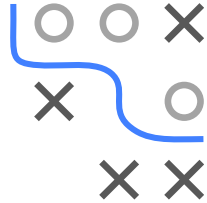
- \+ Retains of most of **base learners'** advantages
- \+ Very **good predictor** due to aggressive loss minimization, typically only outperformed by heterogenous **stacking ensembles**
- \+ High **flexibility** via custom loss functions and choice of base learner
- \+ Highly efficient implementations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

Disadvantages

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

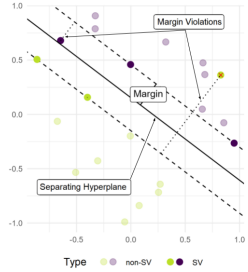
LINEAR SVM – METHOD SUMMARY

CLASSIFICATION | REGRESSION | 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**)

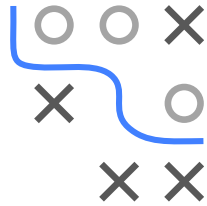


Soft-margin SVM with margin violations

- 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

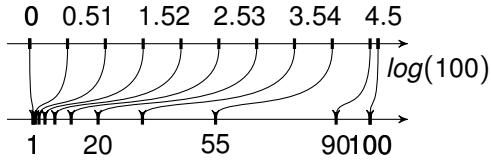
Hypothesis space (primal) $\mathcal{H} = \{f(\mathbf{x}) \,:\, f(\mathbf{x}) = \boldsymbol{\theta}^\top \mathbf{x} + \theta_0\}$

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



Tuning

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

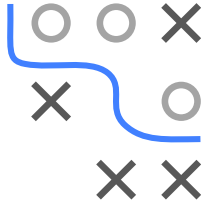


Implementation

- **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 SVM – METHOD SUMMARY

CLASSIFICATION | REGRESSION | NONPARAMETRIC | BLACK-BOX



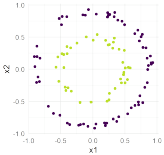
General idea

- 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**: $\langle \phi x, \phi xt \rangle = kxxt$

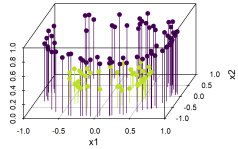
Hypothesis space

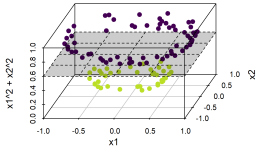
$$\mathcal{H} = \left\{ f(\mathbf{x}) \ : \ f(\mathbf{x}) = \text{sign}\left(\boldsymbol{\theta}^\top \phi x + \theta_0\right) \right\} \text{ (primal)}$$

$$\mathcal{H} = \left\{ f(\mathbf{x}) \ : \ f(\mathbf{x}) = \text{sign}\left(\sum_{i=1}^{n} \alpha_i y^{(i)} k(\mathbf{x}^{(i)}, \mathbf{x}) + \theta_0\right) \ | \ \alpha_i \geq 0, \sum_{i=1}^{n} \alpha_i y^{(i)} = 0 \right\} \text{ (dual)}$$



Nonlinear problem in original space

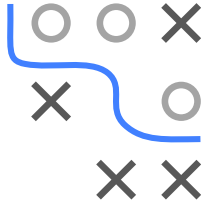




Mapping to 3D space and subsequent linear separation – implicitly handled by kernel in nonlinear SVM

Dual problem **Kernelize** dual (soft-margin) SVM problem, replacing all inner products by kernels:

NONLINEAR SVM – IMPLEMENTATION & PRACTICAL HINTS



Common kernels

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

Tuning

- High sensitivity w.r.t. hyperparameters, especially those of kernel $\Rightarrow$ **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`

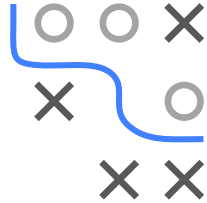
SVM – PRO'S & CON'S

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)
  → Non-SV do not affect decision boundary
+ Convex optimization problem
  → local minimum $\hat{=}$ global minimum

Disadvantages

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



Advantages (nonlinear SVM)

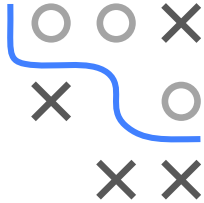
+ Can learn **nonlinear decision boundaries**
+ **Very flexible** due to custom kernels
  → RBF kernel yields local model

Disadvantages (nonlinear SVM)

− Poor **interpretability** due to complex kernel
− **Not easy tunable** as it is highly important to choose the right kernel (which also

NEURAL NETWORKS – METHOD SUMMARY

REGRESSION | CLASSIFICATION | (NON)PARAMETRIC | BLACK-BOX



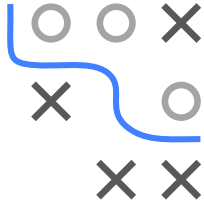
General idea

- Learn **composite function** through series of nonlinear feature transformations, represented as **neurons**, organized hierarchically in **layers**
  - Basic neuron operation: 1) affine **transformation** ϕ (weighted sum of inputs), 2) nonlinear **activation** σ
  - 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 infered from the **computational graph** of the network
    $\rightarrow$ **Automatic Differentiation** (AutoDiff)
  - Algorithm to compute weight updates based on the loss is called **Backpropagation**

Hypothesis space

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

NEURAL NETWORKS – IMPLEMENTATION & PRACTICAL HINTS



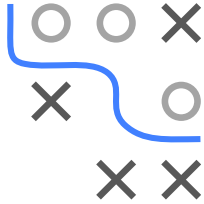
General hints

- 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.
  $\rightarrow$ Using a service (esp. for foundation models) can be more cost efficient

Implementation

- **R:** Use python libraries (below) via `reticulate`, but not really recommended except for toy applications.
- **Python libraries:**
  - `keras` for simple high level API

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