# Important Learners in ML



### Learning goals

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

### **CONTENTS**

- 1 k-Nearest Neighbors (k-NN)
- 2 Generalized Linear Models (GLM)
- **3** Generalized Additive Models (GAM)
- 4 Classification & Regression Trees (CART)
- 5 Random Forests
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- 7 Linear Support Vector Machines (SVM)
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### K-NN – METHOD SUMMARY

REGRESSION

CLASSIFICATION

NONPARAMETRIC

WHITE-BOX

#### General idea

- similarity in feature space (w.r.t. certain distance metric  $d(\mathbf{x}^{(i)}, \mathbf{x})$ )  $\leadsto$  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**: 
$$\hat{y} = \frac{1}{\sum\limits_{i:\mathbf{x}^{(i)} \in N_k(\mathbf{x})} w_i} \sum\limits_{i:\mathbf{x}^{(i)} \in N_k(\mathbf{x})} w_i y^{(i)}$$
 with  $w_i = \frac{1}{d(\mathbf{x}^{(i)},\mathbf{x})}$ 

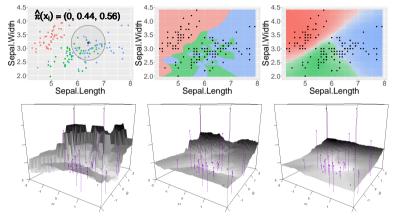
- $\rightarrow$  optional: higher weights  $w_i$  for close neighbors
- most frequent class for **classification**:  $\hat{y} = \underset{\ell \in \{1, ..., g\}}{\arg \max} \sum_{i, \mathbf{x}^{(i)} \in N_{\ell}(\mathbf{x})} \mathbb{I}(y^{(i)} = \ell)$

$$\Rightarrow$$
 Estimating posterior probabilities as  $\hat{\pi}_{\ell}(\mathbf{x}^{(i)}) = \frac{1}{k} \sum_{i:\mathbf{x}^{(i)} \in N_{\ell}(\mathbf{x})} \mathbb{I}(y^{(i)} = \ell)$ 

- Nonparametric behavior: parameters = training data; no compression of information
- Not immediately interpretable, but inspection of neighborhoods can be revealing

### 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

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### K-NN - METHOD SUMMARY

### Popular distance metrics

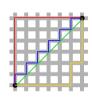
• Numerical feature space:

$$\Rightarrow$$
 Typically, **Minkowski** distances  $d(\mathbf{x}, \widetilde{\mathbf{x}}) = \|\mathbf{x} - \widetilde{\mathbf{x}}\|_q = \left(\sum_j |x_j - \widetilde{x}_j|^q\right)^{\frac{1}{q}}$ 

• 
$$q=1$$
: Manhattan distance  $o d(\mathbf{x}, \tilde{\mathbf{x}}) = \sum_{i} |x_i - \tilde{x_i}|$ 

• 
$$q=2$$
: Euclidean distance  $\to d(\mathbf{x}, \tilde{\mathbf{x}}) = \sqrt{\sum_{i} (x_i - \tilde{x}_i)^2}$ 

• Visualization: Manhattan (red, blue, yellow) vs. Euclidean (green)



- Mixed feature space:
  - Gower distance can handle numerical and categorical features, and missing data:

- 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 to account 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  $\mathcal{O}(N \log N)$ : 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

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### 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  $\rightarrow$  nonlinear decision boundaries
- + Easy to **tune** (few hyperparameters)
  - $\rightarrow$  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)

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# GENERALIZED LINEAR MODELS – METHOD SUMMARY

CLASSIFICATION

Represent target as function of linear predictor  $\theta^{\top} \mathbf{x}$  (weighted sum of features)  $\rightarrow$  **Interpretation:** if feature  $x_i$  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  $\phi(\boldsymbol{\theta}^{\top} \mathbf{x}) = \frac{1}{1 + \exp(-\boldsymbol{\theta}^{\top} \mathbf{x})} =: \pi(\mathbf{x} \mid \boldsymbol{\theta})$ ⇒ Decision rule: Linear hyperplane



Linear regression hyperplane



Logistic sigmoid function



Logistic function for bivariate input and loss-minimal A



Corresponding separating hyperplane

# GENERALIZED LINEAR MODELS – 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:

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

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

### GENERALIZED LINEAR MODELS – METHOD SUMMARY

#### 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:  $\pi_k(\mathbf{x} \mid \boldsymbol{\theta}) = \exp(\boldsymbol{\theta}_k^{\top} \mathbf{x}) / \sum_{i=1}^g \exp(\boldsymbol{\theta}_i^{\top} \mathbf{x})$
- ullet Multi-class log-loss:  $L(y,\pi(\mathbf{x})) = -\sum\limits_{k=1}^g \mathbb{I}_{\{y=k\}} \log(\pi_k(\mathbf{x}))$
- 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

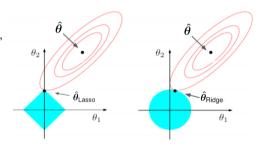
- Empirical risk function **plus complexity penalty**  $J(\theta)$ , controlled by shrinkage parameter  $\lambda > 0$ :  $\mathcal{R}_{\text{reg}}(\theta) := \mathcal{R}_{\text{emp}}(\theta) + \lambda \cdot J(\theta)$
- Ridge regression: L2 penalty  $J(\theta) = \|\theta\|_2^2$
- ullet LASSO regression: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

#### Optimization under regularization

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

#### Choice of regularization parameter

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



### GENERALIZED LINEAR MODELS – REGULARIZATION

Ridge vs. LASSO

### Ridge

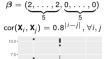
- ullet Global shrinkage  $\Rightarrow$  overall smaller but still dense  $oldsymbol{ heta}$
- Applicable with large number of influential features, handling correlated variables by shrinking their coefficients by equal amount

#### LASSO

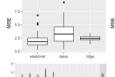
- Actual variable selection by shrinking coefficients for irrelevant features all the way to zero
- Suitable for sparse problems, ineffective with correlated features (randomly selecting one)
- Neither overall better ⇒ compromise: elastic net
  - Weighted combination of Ridge and LASSO
  - Introducing additional penalization coefficient:

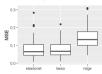
$$\mathcal{R}_{\mathsf{reg}}(oldsymbol{ heta}) = \mathcal{R}_{\mathsf{emp}}(oldsymbol{ heta}) + \lambda \cdot P_{lpha}(oldsymbol{ heta}), ext{ with } P_{lpha}(oldsymbol{ heta}) = [lpha \cdot \|oldsymbol{ heta}\|_1 + (1-lpha) \cdot rac{1}{2} \cdot \|oldsymbol{ heta}\|_2^2]$$

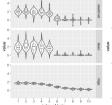
Ridge performs better for correlated features: **Lasso** performs better for uncorrelated features:

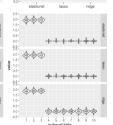












### GENERALIZED LINEAR MODELS – IMPLEMENTATION

### 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
- Python: From package sklearn.linear\_model
  - Unregularized:
    - LinearRegression()
    - LogisticRegression(penalty = None)
  - Regularized:
    - Linear regression: Lasso().Ridge().ElasticNet()
    - Logistic regression: Logistic Regression (penalty = {'11', '12', '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 ≫ 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 w.r.t. outliers and noisy data (especially with L2 loss)
- Also a LM can **overfit** (e.g., many features and few observations)
- Feature interactions must be handcrafted
  - → practically infeasible for higher orders

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### GENERALIZED ADDITIVE MODELS – METHOD SUMMARY

REGRESSION

CLASSIFICATION

NON)PARAMETRIC

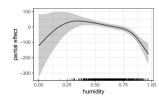
WHITE-BOX

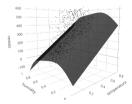
**FEATURE SELECTION** 

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



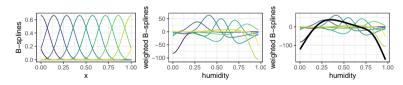


Prediction of bike rentals from smooth term of humidity (left: partial effect) and linear term of temperature (right: bivariate prediction).

### GENERALIZED ADDITIVE MODELS – METHOD SUMMARY

#### Smooth functions

- Nonparametric/semiparametric/parametric approaches conceivable
- Frequently: express f<sub>i</sub> as weighted sum of basis functions ~> model linear in weight coefficients again
  - Use fixed basis of functions  $b_1, \ldots, b_K$  and estimate associated coefficients  $\gamma_1, \ldots, \gamma_K \rightarrow f_l(x_l) = \sum_{k=1}^{K_l} \gamma_{l,k} b_k(x_l)$
  - Popular types of basis functions
    - Polynomial → smoothing/TP-/B-splines
    - Radial ~ Kriging
    - Trigonometric → Fourier/wavelet forms
- Alternatives: local regression (LOESS), other kernel-smoothing approaches, . . .

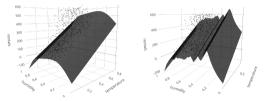


Left: B-spline basis with 9 basis functions. Middle: BFs weighted with coefficients estimated for humidity. Right: sum of weighted BFs in black (= partial effect).

### GENERALIZED ADDITIVE MODELS – METHOD SUMMARY

### Regularization

- Smooth functions possibly very flexible → 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 → 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.

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

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# **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

### **Advantages**

- + Simple and fast implementation
- Applicable for any dataset size, as long as number of observations ≫ number of features
- + High **flexibility** via smooth effects
- + Easy to **combine** linear & nonlinear effects
- Intuitive interpretability via feature effects (though not quite as straightforward as in GLM)
- $+\;$  Statistical hypothesis **tests** for effects available

#### **Disadvantages**

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

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### **CART – METHOD SUMMARY**

REGRESSION

CLASSIFICATION

NONPARAMETRIC

WHITE-BOX

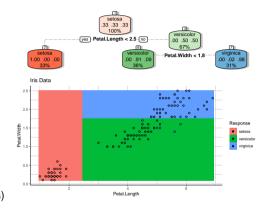
FEATURE SELECTION

### 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

   → empirical risk minimization (ERM)
- In each step:
  - Find optimal split (feature-threshold combination)
     → greedy search
  - Assign constant prediction  $c_m$  to all obs. in  $Q_m$ 
    - $\rightarrow$  Regression:  $c_m$  is average of y
    - ightarrow Classif.:  $c_m$  is majority class (or class proportions)
  - Stop when a pre-defined criterion is reached
    - → See Complexity control

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

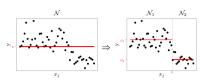


### **CART – METHOD SUMMARY**

#### **Empirical risk**

• Splitting **feature**  $x_j$  **at split point** t divides a parent node  $\mathcal N$  into two child nodes:

$$\mathcal{N}_1 = \{(\mathbf{x}, y) \in \mathcal{N} : x_j \leq t\} \text{ and } \mathcal{N}_2 = \{(\mathbf{x}, y) \in \mathcal{N} : x_j > t\}$$



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

$$\mathop{\mathsf{arg\,min}}_{j,t} \mathcal{R}(\mathcal{N},j,t) = \mathop{\mathsf{arg\,min}}_{j,t} \mathcal{R}(\mathcal{N}_1) + \mathcal{R}(\mathcal{N}_2)$$

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

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

#### 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
    - ightarrow hard to determine good stopping point before actually trying all combinations
  - **Pruning:** grow deep trees and cut back in risk-optimal manner afterwards

### 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

#### **Advantages**

- + **Easy** to understand & visualize (**interpretable**)
- + Built-in **feature selection** 
  - ightarrow 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**, even of higher order
- + Fast computation and good scalability
- 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

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### RANDOM FORESTS – METHOD SUMMARY

REGRESSION

CLASSIFICATION

NONPARAMETRIC

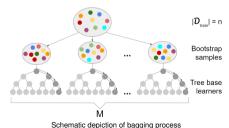
BLACK-BOX

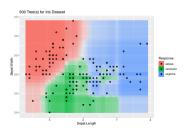
FEATURE SELECTION

#### General idea

- Bagging ensemble of *M* tree base learners fitted on bootstrap data samples
  - ⇒ 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 
$$\mathcal{H} = \left\{ f(\mathbf{x}) : f(\mathbf{x}) = \frac{1}{M} \sum_{m=1}^{M} \sum_{t=1}^{T^{[m]}} c_t^{[m]} \mathbb{I}(\mathbf{x} \in Q_t^{[m]}) \right\}$$





Prediction surface for iris data with 500-tree ensemble

### RANDOM FORESTS – METHOD SUMMARY

Empirical risk & Optimization 

Just like tree base learners

### Out-of-bag (OOB) error

- Ensemble prediction for obs. outside individual trees' bootstrap training sample ⇒ 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

#### 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

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# **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

- 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

#### Implementation

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

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### **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.

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### 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

10.0

- 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

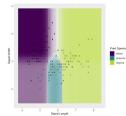
7.5

• Predict via (weighted) sum of base learners

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

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

5.0



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

## **GRADIENT BOOSTING – METHOD SUMMARY**

#### **Empirical risk**

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

$$\tilde{r}^{(i)} = -\frac{\partial L(y^{(i)}, t(\mathbf{x}^{(i)}))}{\partial t(\mathbf{x}^{(i)})}$$
 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  $\hat{i}^{[m-1]}$  fixed  $\Rightarrow$  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  $c_i^{[m]}$  for tree base learners:  $\tilde{c}_i^{[m]} = \beta^{[m]} \cdot c_i^{[m]}$

#### 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

### **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<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

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### 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
- ⇒ 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: /
- ⇒ mboost very flexible but slow while compboost is much faster with limited features

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## **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 (→ solved by efficient implementation)

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# **LINEAR SVM – METHOD SUMMARY**

CLASSIFICATION

REGRESSION

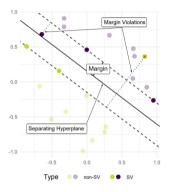
PARAMETRIC

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



Soft-margin SVM with margin violations

# LINEAR SVM - METHOD SUMMARY

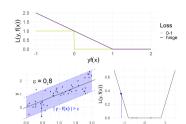
Empirical risk Soft-margin SVM as L2-regularized ERM:

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

- $\|\boldsymbol{\theta}\| = 1/\gamma$  ( $\hat{=}$  maximizing margin)
- C > 0: penalization for margin violations
- Loss aims at minimizing margin violations

$$\rightarrow$$
 Classif. (**hinge** loss):  $L(y, f) = \max(1 - yf, 0)$ 

$$ightarrow$$
 Regr. ( $\epsilon$ -insensitive loss):  $L\left(y,f
ight)=\max(|y-f|-\epsilon,0)$ 



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  $\alpha_i \ge 0$ :

$$\max_{\boldsymbol{\alpha} \in \mathbb{R}^n} \sum_{i=1}^n \alpha_i - \frac{1}{2} \sum_{i=1}^n \sum_{i=1}^n \alpha_i \alpha_i y^{(i)} y^{(j)} \left\langle \mathbf{x}^{(i)}, \mathbf{x}^{(j)} \right\rangle \quad \text{s.t.} \quad 0 \le \alpha_i \le C \quad \forall i \in \{1, \dots, n\} \text{ and } \sum_{i=1}^n \alpha_i y^{(i)} = 0$$

Hypothesis space (dual) Non-SVs have  $\alpha_i = 0$  as they do not affect the hyperplane

$$\mathcal{H} = \left\{ f(\mathbf{x}) \ : \ f(\mathbf{x}) = \operatorname{sign}\left(\sum_{i=1}^n \alpha_i y^{(i)} \langle \mathbf{x}^{(i)}, \mathbf{x} \rangle + \theta_0\right) \mid \alpha_i \geq 0, \sum_{i=1}^n \alpha_i y^{(i)} = 0 \right\}$$

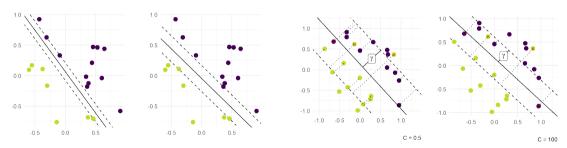
### **LINEAR SVM – METHOD SUMMARY**

#### 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)

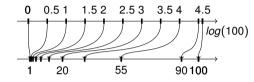
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### LINEAR SVM – IMPLEMENTATION & PRACTICAL HINTS

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



#### 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

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# **NONLINEAR SVM – METHOD SUMMARY**

CLASSIFICATION

REGRESSION

NONPARAMETRIC

BLACK-BOX

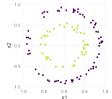
#### General idea

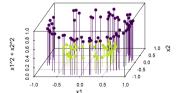
- Move beyond linearity by mapping data to transformed space where they are linearly separable
- Kernel trick
  - No need for explicit construction of feature maps
  - Replace inner product of feature map  $\phi: \mathcal{X} \to \Phi$  by **kernel**:  $\langle \phi(\mathbf{x}), \phi(\tilde{\mathbf{x}})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

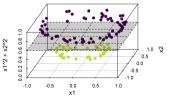
### Hypothesis space

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

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







Nonlinear problem in original space

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

### NONLINEAR SVM – METHOD SUMMARY

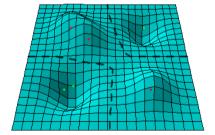
Dual problem Kernelize dual (soft-margin) SVM problem, replacing all inner products by kernels:

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

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

Interpretation as basis function approach

- Representer theorem: solution of dual soft-margin SVM problem is  $\theta = \sum_{i=1}^{n} \beta_{i} \phi(\mathbf{x}^{(i)})$
- Sparse, weighted sum of **basis functions**  $\rightarrow \beta_i = 0$  for non-SVs
- Result: local model with smoothness depending on kernel



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

## **NONLINEAR SVM – IMPLEMENTATION & PRACTICAL HINTS**

#### Common kernels

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

### Tuning

- ◆ High sensitivity w.r.t. hyperparameters, especially those of kernel ⇒ 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
- + Convex optimization problem
  - → local minimum  $\hat{=}$  global minimum

#### Advantages (nonlinear SVM)

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

#### **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

#### 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)

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# GAUSSIAN PROCESSES (GP) – METHOD SUMMARY

REGRESSION

CLASSIFICATION

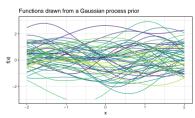
NONPARAMETRIC

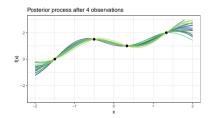
**PROBABILISTIC** 

#### 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 induced the coviariance matrix of the distribution.
- Predict via the maximum a-posteriori (MAP) estimate.

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





# GAUSSIAN PROCESSES (GP) – METHOD SUMMARY

### **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

#### 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

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## **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

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# **GAUSSIAN PROCESSES (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

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REGRESSION

CLASSIFICATION

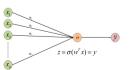
(NON)PARAMETRIC

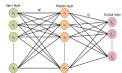
BLACK-BOX

#### 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 infered from the computational graph of the network
     → Automatic Differentiation (AutoDiff)
  - Algorithm to compute weight updates based on the loss is called **Backpropagation**

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





#### **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  $au(\phi)$ 
  - ullet Regression: one output neuron,  $au={
    m identity}$
  - 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)

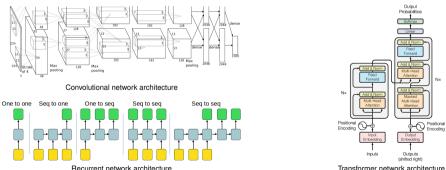
Empirical risk In general, compatible with any differentiable loss

#### 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  $\to \mathbf{RMSProb}$ 
    - $\Rightarrow$  **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 – 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.
  - ightarrow 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
  - 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

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## **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

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