# **Models in Machine Learning**

**An Overview** 



### **PREFACE**

### @Bernd: pls fill with wisdom

In machine learning, there's something called the "No Free Lunch" theorem. In a nutshell, it states that no one algorithm works best for every problem, and it's especially relevant for supervised learning (i.e. predictive modeling). For example, you can't say that neural networks are always better than decision trees or vice-versa. There are many factors at play, such as the size and structure of your dataset. As a result, you should try many different algorithms for your problem, while using a hold-out "test set" of data to evaluate performance and select the winner.

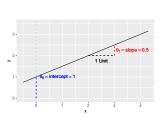
# LINEAR MODEL - FUNCTIONALITY

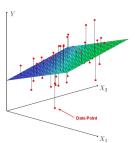
SUPERVISED PARAMETRIC LINEAR WHITE-BOX

Caro alter Tanzbesen

General idea A linear model (LM) fits a hyperplane  $\theta_0 + \boldsymbol{\theta}^T \mathbf{x}$  to minimize the distance between the data points and its closed point on the hyperplane.

Hypothesis space  $\mathcal{H} = \{\theta_0 + \boldsymbol{\theta}^T \mathbf{x} \mid (\theta_0, \boldsymbol{\theta}) \in \mathbb{R}^{p+1} \}$ 





# **LINEAR MODEL – FUNCTIONALITY**

### **Empirical risk**

- Typically, **ordinary least squares (OLS)** with a squared loss function is used for regression:  $\mathcal{R}_{emp}(\theta) = \sum_{i=1}^{n} \left( y^{(i)} \theta^{\mathsf{T}} \mathbf{x}^{(i)} \right)^2$
- Sometimes the empirical risk function is based on the absolute loss or the Huber loss
- For **logistic regression** the ERM is based on the **log loss**  $L(y, f(\mathbf{x})) = \log [1 + \exp(-yf(\mathbf{x}))].$
- In this case, the hyperplane can represent the decision boundary between two classes (classification).

### Optimization

- for **OLS**: analytically with  $\hat{\theta} = (\mathbf{X}^{\top}\mathbf{X})^{-1}\mathbf{X}^{\top}\mathbf{y}$
- for other loss functions: numerical optimization

Hyperparameters None

Runtime behavior  $\mathcal{O}(d^2 \cdot n + d^3)$  for *n* observations and *d* features

# **LINEAR MODEL - PRO'S & CON'S**

### Advantages

- simple and fast implementation; cheap computational costs
- intuitive interpretability: mean influence of features on the output and feature importance
- fits linearly separable data sets very well
- + works well independent of data size
- basis for many machine learning algorithms

### Disadvantages

- not suitable for data based on a non-linear data generating process → strong simplification of real-world problems
- strong assumptions: data is independent (multi-collinearity must be removed and normal distributed residuals ??
- tend to overfit (can be reduced by regularization)
- sensitive to outliers and noisy data

Simple method with good interpretability for linear problems, but strong assumptions and simplification of real-world problems

# **LINEAR MODEL – PRACTICAL HINTS**

#### Check assumptions???????

This model is very effective, if the following assumptions are fulfilled:

- linearity: the relationship between the mean of predicted value and the features
- homoscedasticity: The variance of residuals is equal for all features.
- independence: All observations are independent of each other.
- normality: Y is normally distributed for any fixed value of the features

### Implementation

- R: function 1m
- Python: LinearRegression from package sklearn.linear\_model, package for advanced statistical parameters statsmodels.api

### Regularization

In practice, we often use regularized models in order to **prevent overfitting** or perform feature selection. More details will follow in the subsequent chapter.

### REGULARIZED LM – FUNCTIONALITY

SUPERVISED PARAMETRIC LINEAR WHITE-BOX

#### General idea

- Linear model (LM) can overfit if we operate in high-dimensional space with not that many oberservations.
- When features are highly correlated, the least-squares estimate becomes highly sensitive to random errors in the observed response, producing a large variance in the fit.
- If we fit a linear model, we can find a compromise between generalizing the model (simple model, underfitted) and correspond closely to the data (complex model, overfitted).

Hypothesis space  $\mathcal{H} = \{\theta_0 + \boldsymbol{\theta}^T \mathbf{x} \mid (\theta_0, \boldsymbol{\theta}) \in \mathbb{R}^{p+1} \}$ 



### **REGULARIZED LM – FUNCTIONALITY**

#### **Empirical risk**

• Therefore, we minimize the empirical risk function  $\mathcal{R}_{emp}(\theta)$  plus the a complexity measure  $J(\theta)$ :

$$\mathcal{R}_{\text{reg}}(\boldsymbol{\theta}) = \mathcal{R}_{\text{emp}}(\boldsymbol{\theta}) + \lambda \cdot J(\boldsymbol{\theta}).$$

- We can use the L2-penalty for the complexity measure (**ridge regression**) with  $J(\theta) = |\theta||_2^2$ .
- Alternativly, **LASSO** (least absolute shrinkage and selection operator) uses the L1-penalty  $(J(\theta) = |\theta||_1)$ .
- Whereas both regularization methods shrink the coefficients of the model, LASSO also performs feature selection.
- Elastic net as a convex combination of Ridge and LASSO ???

### Optimization

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

Hyperparameters Shrinkage parameter  $\lambda$  [and  $\alpha$  for Elastic net??]

Runtime behavior  $\mathcal{O}(d^2 \cdot n + d^3)$  for *n* observations and *d* features

# **REGULARIZED LM – PRO'S & CON'S**

#### SAME LIKE LINEAR MODEL?

#### **Advantages**

- simple and fast implementation;
  cheap computational costs
- + intuitive interpretability: mean influence of features on the output and feature importance
- fits linearly separable data sets very well
- + works well independent of data size
- basis for many machine learning algorithms
- + prevents overfitting

### **Disadvantages**

- not suitable for data based on a non-linear data generating process
   → strong simplification of real-world problems
- strong assumptions: data is independent (multi-collinearity must be removed and normal distributed residuals 2222
- sensitive to outliers and noisy data

Simple method with good interpretability for linear problems, but strong assumptions and simplification of real-world problems.

# REGULARIZED LM – PRACTICAL HINTS

Choice of regularization parameter  $\lambda$ 

Choose  $\lambda$  with e. g. smallest sum of squared residuals through cross-validation. In the R package glmnet lambda.min is the value of  $\lambda$  that gives minimum mean cross-validated error.

### Implementation

- R: package for regularized linear model glmnet
- Python: LinearRegression from package sklearn.linear\_model, package for advanced statistical parameters statsmodels.api

# SVM - FUNCTIONALITY

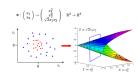
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#### General idea

- Support vector machines (SVMs) construct separating hyperplanes in a multi-dimenstional space.
- The SVM algorithm finds a decision boundary (separating hyperplane) that maximizes the distance (margin) between the closest members (support vectors) of the separate classes. (Linear SVM)
- If the data is not linearly separable, kernels transform the input space into a higher dimensional space. (Non-linear SVM)

Hypothesis space 
$$\mathcal{H} = \{ sign \sum_{i=1}^{n} \alpha_i y^{(i)} k(\mathbf{x}^{(i)}, \mathbf{x}) + \theta_0 | (\theta_0, \boldsymbol{\theta}) \in \mathbb{R}^{p+1} \}$$





# **SVM - FUNCTIONALITY**

**Empirical risk** 

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Optimization XX

Hyperparameters

- C: penalization for missclassified data points
- kernel parameters: depending if and which kernel is used (e. g. degree of the polynomial kernel or width of RBF kernel)

Runtime behavior  $\mathcal{O}(n^2 \cdot d + n^3)$  for *n* observations and *d* features

# SVM - PRO'S & CON'S

#### Advantages

- + can learn non-linear decision boundaries
- + often sparse solution
- robust against overfitting; especially in high-dimensional space
- + computational costs are low ?????

### **Disadvantages**

- not easy tunable as it is highly important to choose the right kernel
- does not scale well to larger data sets

XX

# **SVM - PRACTICAL HINTS**

#### Kernels

For projecting the input data into another onto a higher dimension, mainly these three types of kernels are used:

- Linear kernel: the dot product of the given observations
- Polynomial kernel: curved lines in the input space
- Radial basis function (RBF) kernel: complex regions in the input space (e. g. spirals)

### Implementation

- R: package for regularized linear model XXX
- Python: XX from package XXX, package for XXXX statsmodels.api

### **CART – FUNCTIONALITY**

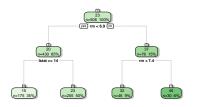
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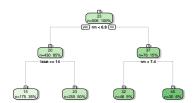
FEATURE SELECTION

#### General idea

- Starting from a root node, classification & regression trees (CART) perform repeated **binary splits** of the data according to feature values, thereby subsequently dividing the input space  $\mathcal{X}$  into T rectangular partitions  $Q_t$ .
- Observations are passed along until each ends up in exactly one leaf node (unless **stopped early** or **pruned**).
- In each step, CART find the optimal feature-threshold combination to split by.
- Leaf node t is assigned response ct.

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





# CART – FUNCTIONALITY

#### **Empirical risk**

- Empirical risk is calculated for each potential terminal node  $\mathcal{N}_t$  of a split.
- In general, trees can handle any type of loss function. Typical choices are:
  - Classification (for g classes):

• Using **Brier score** 
$$\mathcal{R}(\mathcal{N}_t) = \sum_{(\mathbf{x}, y) \in \mathcal{N}_t} \sum_{k=1}^g (\mathbb{I}(y=k) - \pi_k(\mathbf{x}))^2$$

• Using **Bernoulli loss** 
$$\mathcal{R}(\mathcal{N}_t) = \sum_{(\mathbf{x}, y) \in \mathcal{N}_t} \sum_{k=1}^g \mathbb{I}(y = k) \cdot \log(\pi_k(\mathbf{x}))$$

• Regression: Using *quadratic loss* 
$$\mathcal{R}(\mathcal{N}_t) = \sum\limits_{(\mathbf{x},y) \in \mathcal{N}_t} (y-c_t)^2$$

#### Optimization

**Exhaustive** search over all (randomly selected) split candidates in each node to minimize empirical risk in the child nodes (greedy optimization)

Hyperparameters Complexity, i.e., number of leaves T

Runtime behavior  $\mathcal{O}(n^2 \cdot d)$  for *n* observations and *d* features

# CART - PRO'S & CON'S

### **Advantages**

- Easy to understand, interpret & visualize
- Automatic handling of non-numerical features
- + Built-in feature selection
- + Automatic handling of missings
- Interaction effects between features easily possible, even of higher orders
- Fast computation and good scalability
- High flexibility (custom split criteria or leaf-node prediction rules)

### Disadvantages

- Rather low accuracy (at least, without bagging or boosting)
- High variance/instability: strong dependence on training data
- Therefore, poor generalization & risk of overfitting
- Several steps required for modeling linear relationships
- In presence of categorical features, bias towards features with many categories

Simple and good with feature selection, but not the best predictor

# CART – PRACTICAL HINTS

### Pruning / early stopping

Unless interrupted, splitting will go on until each leaf node contains a single observation (expensive + overfitting!)

ightarrow Use **pruning** and **stopping criteria** to limit complexity

### Implementation

R: package rpart

Python: DecisionTreeClassifier / DecisionTreeRegressor from package scikit-learn

Complexity controlled via tree depth, minimum number of observations per node, maximum number of leaves, minimum risk reduction per split, ...

### Bagging / boosting

Since CART are instable predictors on their own, they are typically ensembled to form a *random forest* (*bagging*) or used in combination with *boosting*.

# **RANDOM FOREST – FUNCTIONALITY**

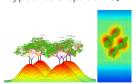
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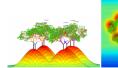
FEATURE SELECTION

#### General idea

- Random forests (RF) are bagging ensembles: they combine M CART (base learners) to form a strong learner.
- They use **complex** trees with low bias and compensate for single trees' high variance by aggregating them in a **decorrelated** manner.
- Each tree is trained a on a bootstrap sample of the data and only on a random subset of features to incur variability.
- Aggregation takes place via averaging (regression) or majority voting (classification).

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 FOREST – FUNCTIONALITY

#### **Empirical risk**

- Applicable with any kind of loss function (just like tree base learners)
- Computation of empirical risk for all potential child nodes in all trees

#### Optimization

**Exhaustive** search over all (randomly selected) split candidates in each node of each tree to minimize empirical risk in the child nodes (greedy optimization)

### **Hyperparameters**

- Ensemble size, i.e., number of trees
- Complexity, i.e., number of leaves T of each base learner
- Number of split candidates, i.e., number of features to be considered as splitting variables at each split
  - ightarrow Frequently used heuristics:  $\lfloor \sqrt{p} \rfloor$  for classification and  $\lfloor p/3 \rfloor$  for regression

Runtime behavior  $\mathcal{O}(M \cdot n^2 \cdot d)$  for M trees, n observations and d features

# **RANDOM FOREST – PRO'S & CON'S**

#### Advantages

- Translation of most of base learners' advantages (e.g., inherent variable selection, handling of missing data)
- + Fairly good at **prediction**: improved accuracy through bagging
- Inherent computation of feature importance
- + Quite stable wrt changes in the data
- Good with high-dimensional data, even in presence of noisy covariates
- + Applicable to unbalanced data
- Easy to parallelize
- + Rather easy to tune

#### Disadvantages

- Translation of some of base learners' disadvantages (e.g., trouble to model linear relations, bias towards features with many categories)
- Loss of single trees' interpretability
  black-box method
- Hard to visualize
- Often suboptimal for regression
- Often still inferior in performance to other methods (e.g., boosting)

### Fairly good predictor, but black-box method

# **RANDOM FOREST – PRACTICAL HINTS**

#### **Pre-processing**

Inherent feature selection of random forests, but high **computational costs** for large number of features

→ Upstream feature selection (e.g., via PCA) might be advisable

### Implementation

R: package ranger

Python: RandomForestClassifier / RandomForestRegressor from package scikit-learn

### Tuning

### Overall limited tunability

Number of split candidates often more impactful than number of trees

# GRADIENT BOOSTING – FUNCTIONALITY

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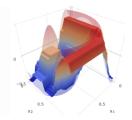
FEATURE SELECTION

General idea Gradient boosting (GB) is an **ensemble** method that constructs a strong learner from weak base learners (frequently, CART).

As opposed to **bagging**, however, base learners are assembled in a **sequential**, stage-wise manner: in each iteration, GB improves the current model by adding a new component that minimizes empirical risk. The final model is a weighted sum of base learners  $b(\mathbf{x}, \theta^{[m]})$  with weights  $\beta^{[m]}$ .

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

- base learner to current point-wise residuals
- gradient step in function space
- Fitting of each base learner using information from previously added ones



# GRADIENT BOOSTING – FUNCTIONALITY

#### **Empirical risk**

- Outer loss: Loss used to compute pseudo-residuals how large is the error of the current model fit?
  - → Arbitrary differentiable loss function
- Inner loss: Loss used to fit next base learner component to current pseudo-residuals
  - → Typically, *quadratic loss* (desirable optimization properties)

Optimization Functional gradient descent for outer optimization loop, procedure for inner one depending on inner loss

### **Hyperparameters**

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

Runtime behavior  $\mathcal{O}(M \cdot n \cdot d)$  for M base learners, n observations and d features

# **GRADIENT BOOSTING – PRO'S & CON'S**

#### **Advantages**

- Powerful off-the-shelf method for supercharging weak learners' performance
- Translation of most of base learners' advantages (e.g., for tree boosting: inherent variable selection, handling of missing data)
- High predictive accuracy that is hard to outperform
- High flexibility (custom loss functions, many tuning options)
- + Applicable to unbalanced data

#### **Disadvantages**

- Hardly interpretable black-box method
- Hard to visualize
- Prone to overfitting
- Sensitive to outliers
- Hard to tune (high sensitivity to variations in hyperparameter values)
- Rather slow in training
- Hard to parallelize

High-performing predictor, but rather delicate to handle

# **GRADIENT BOOSTING – PRACTICAL HINTS**

XGBoost (extreme gradient boosting)

Fast, efficient implementation of gradient-boosted decision trees that has become **state of the art** for many machine learning problems

ightarrow Clever modeling techniques + computational speed

Stochastic gradient boosting (SGB)

Faster, approximate version of GB that performs each iteration only on random subset of the data

ightarrow Potentially preferrable for high-dimensional data sets

#### Implementation

R: packages gbm, xgboost

Python: GradientBoostingClassifier / GradientBoostingRegressor from package scikit-learn, XGBClassifier / XGBRegressor from package xgboost

#### **Tuning**

### Overall limited tunability

Number of split candidates often more impactful than number of trees

# **NEURAL NETWORK – FUNCTIONALITY**

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#### General idea

- A *neural network (NN)* is a model architecture loosely inspired by the human brain. It consists of various **neurons**, organized in **layers**. These layers are assembled through functional connections.
- Data enter the network in the **input layer** and sequentially pass through h **hidden layers**, each of which performs a linear transformation  $\phi^{(j)}$  and a non-linear activation  $\sigma^{(j)}$ .
- The **output layer** receives the repeatedly transformed inputs, performs a final transformation  $\phi$  and outputs predictions (suitably scaled by  $\tau$ ).

### Hypothesis space

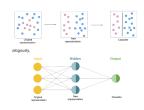
$$\mathcal{H} = \left\{ f(\mathbf{x}) : f(\mathbf{x}) = \tau \circ \phi \circ \sigma^{(h)} \circ \phi^{(h)} \circ \sigma^{(h-1)} \circ \phi^{(h-1)} \circ \dots \circ \sigma^{(1)} \circ \phi^{(1)}(\mathbf{x}) \right\}$$

- Pass observations along until each ends up in exactly one leaf node
- In each step, find the optimal feature-threshold combination to split by



### **NEURAL NETWORK – FUNCTIONALITY**

 NNs use "hidden layers" between inputs and outputs in order to model intermediary representations of the data:



### Empirical risk XX

Optimization NNs are optimised by **backpropagation** which consists of two steps:

- Forward pass: Predict result witch current weights and calculate empirical loss.
- Backward pass: Calculate error contribution of each weight and update the weights by the negative gradient descent.

Hyperparameters Number of hidden neurons, Dropout, initial weights, activation function, learning rate, number of epochs, batch size

Runtime behavior ???

# **NEURAL NETWORK – PRO'S & CON'S**

#### **Advantages**

- can solve complex, non-linear regression or classification problems
- also suitable for unstructed data (e. g. image, audio and text data)
- + very accuarate
- + can easily be updated
- reduces the need for feature engeneering

### Disadvantages

- requires a huge amount of data
- black-model → hard to interpret or explain
  - computationally expensive  $\rightarrow$  slow to train and forecast
- tend to overfit
- requires much expertise for tuning

Computationally expensive models which can learn complex functions and are suitable for unstructred data like text or pictures.

# **NEURAL NETWORK – PRACTICAL HINTS**

**RNNS** 

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**CNNS** 

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#### Implementation

R: package for regularized linear model XXX

Python: XX from package XXX, package for XXXX statsmodels.api