

Overview and final words

DAT200

Overview

Basic Learning Algos

SVM:

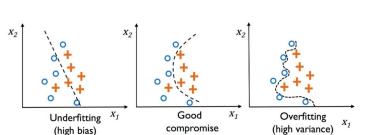
Maximize the margin

SVM

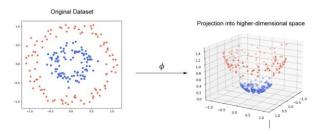
Hyperplane

for first class

Overfitting



Kernels / feature mapping

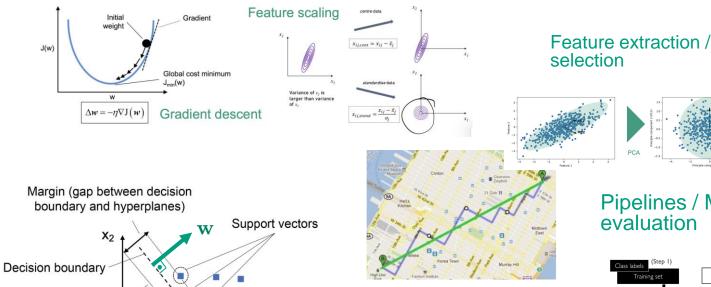




feature mapping, here:

$$\phi(x_1,x_2)=(z_1,z_2,z_3)=(x_1,x_2,x_1^2+x_2^2)$$

Clustering



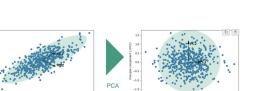
Hyperplane

for second

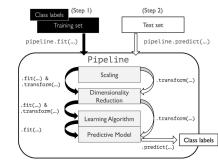
class

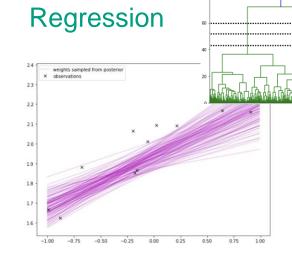
K-NN

Decision Trees

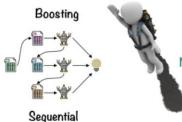


Pipelines / Model evaluation







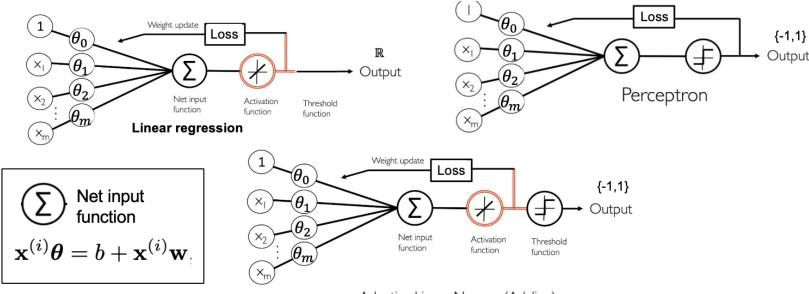


Ensembles



Basics of machine learning, classification/regression

- Simple artificial neurons
 - The Perceptron
 - Adaptive linear neuron (Adaline)
- Simple linear regression, least squares, normal equations
- Net input
- Activation function
- Threshold function
- Cost/Loss function
- Gradient descent
- Newton's method



Adaptive Linear Neuron (Adaline)



Below you see the equation for the perceptron learning rule.

$$\Delta w_j = \eta \left(y^{(i)} - \hat{y}^{(i)} \right) x_j^{(i)}$$

What does w_i represent?

- The learning rate
- A feature weight
- The prediction error
- A feature value



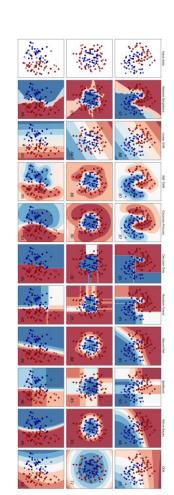
Which activation function does Adaline use?

- The sigmoid function
- The identity function
- A threshold function
- The radial basis function



Classification algorithms / classification analysis with sci-kit learn

- Binary to multi-class: One-vs-all (one classifier per class)
- Logistic regression (outputs probabilities, nonlinear loss function, regularization, ...)
- Support vector machines (margin hard/soft (slack variables), kernels)
- Decision trees (information gain, impurity, random forests, bagging, ...)
- **K-NN** (lazy learner, parametric vs. non-parametric models, distances)
- Associated main hyperparameters and their effects
- Algorithmic understanding in the detail level provided in the lectures (e.g. know the steps of PCA, majority voting, bagging/boosting, ...)

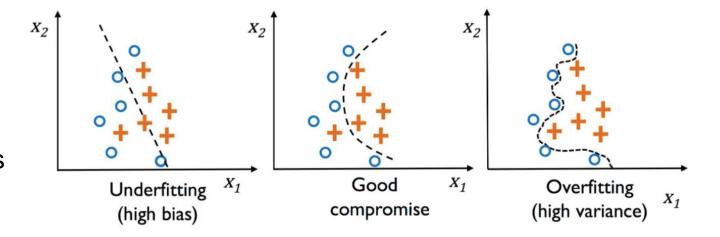




Overfitting, Bias/variance trade-off

Techniques to reduce overfitting:

- Reduce model complexity
 - Reduce the number of parameters
 - Reduce the number of input features
- Regularization (L1 / L2)
- Dimensionality reduction





A model that is overfitted has...

- High bias and low variance
- High variance and low bias
- Both high variance and high bias

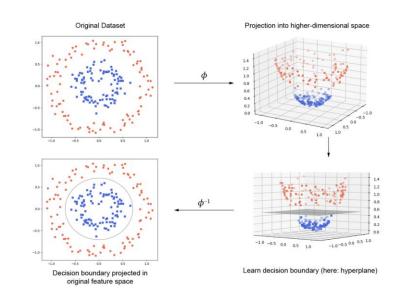


Kernels, kernel trick, feature maps

- Nonlinear decision boundaries
- Properties of the kernel function/matrix
- Feature maps
- Kernel trick
- Replace inner/dot products by kernel function calls

$$\mathbf{x}^{(i)}\mathbf{x}^{(j)}^T \longrightarrow \kappa(\mathbf{x}^{(i)},\mathbf{x}^{(j)}) = \phi(\mathbf{x}^{(i)})\phi(\mathbf{x}^{(j)})^T$$

Kernel SVM, Kernel perceptron, Kernel PCA

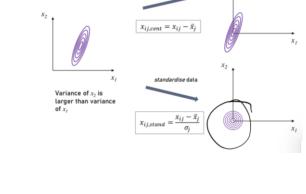


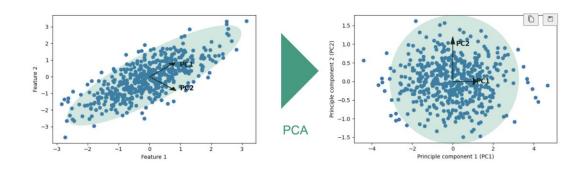
$$\kappa(\mathbf{x},\mathbf{x}'):\mathcal{X} imes\mathcal{X} o\mathbb{R}$$



Pre-processing and feature selection/extraction

- Techniques used on various data types (e.g. encoding)
- Removing or replacing / imputing missing values
- Scaling
- Feature selection (SFS/SBS, feature importances from random forests, regularization (L1/L2))
- Feature extraction / Dimension reduction
 - PCA (unsupervised), LDA (supervised)





Feature scaling



What does the following equation represent?

$$\kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \exp\left(-\frac{||\mathbf{x}^{(i)} - \mathbf{x}^{(j)}||_2^2}{2\sigma^2}\right) := \exp\left(-\gamma||\mathbf{x}^{(i)} - \mathbf{x}^{(j)}||_2^2\right)$$

- The Linear kernel
- The polynomial kernel
- The sigmoid kernel
- The radial basis function kernel



Feature extraction: Principal Component Analysis (PCA)

Summary of steps (centering but not standardization is included in PCA from sklearn)

- 1. Center (covariance matrix) or standardize (correlation matrix) the data. [→ Always at least center (subtract the mean), but don't standardize if the scale carries significant meaning, e.g. several features measuring similar things (e.g. signal intensity at different locations)]
- 2. Compute the covariance/correlation matrix ($\Sigma = \frac{1}{n-1} X^T X$)
- 3. Compute the **eigendecomposition** of Σ [\rightarrow 2./3. may be replaced by one step by performing a singular value decomposition directly on the centered data]
- 4. Rank the **eigenvectors** according to the values of the corresponding **eigenvalues** of Σ
- 5. Keep $k \leq m$ features, and construct a projection matrix W from the "top" k eigenvectors
- 6. Transform the d -dimensional input data set with m features to the new k features (principal components), using the projection matrix W to obtain the new k -dimensional feature subspace

$$T = XW$$



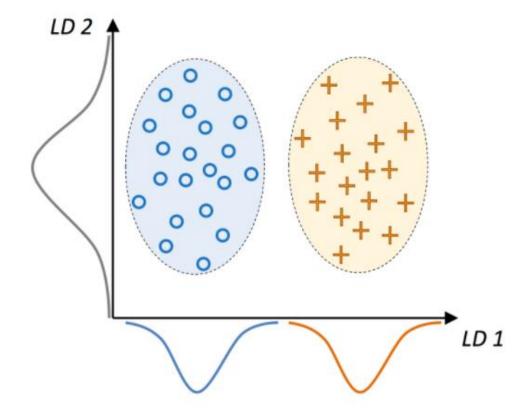
PCA is often used for... (multiple correct answers)

- Dimensionality reduction
- Classification
- Feature extraction
- Data visualisation
- Prediction
- Interpretation of patterns in the data
- Clustering

Feature extraction using Linear Discriminant Analysis



- The concept of LDA
- In the binary classification example to the right a dataset has been projected onto two *linear* discriminants.
- Which of the LD's would you choose to classify the dataset?





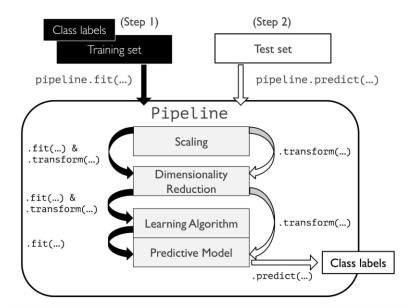
Pipelines, cross-validation and model selection

- Pipelines
- Holdout cross-validation
- K-fold cross-validation
- Learning & validation curves
- Grid search
- Randomized search
- Nested cross-validation

- Confusion matrix
- Important metrics (precision, recall, F1)
- ROC-curve and ROC-AUC

Multi-class classification (micro- vs macro-

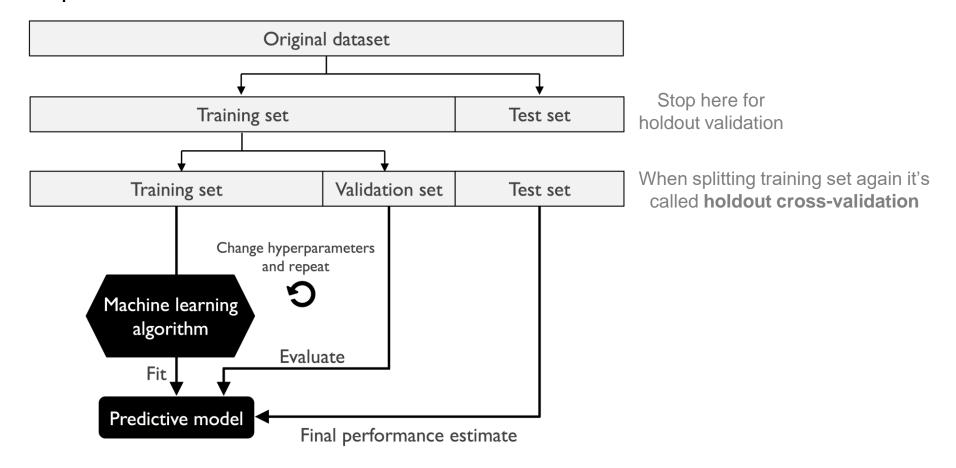
averaging)



The holdout method ("validation" partition)

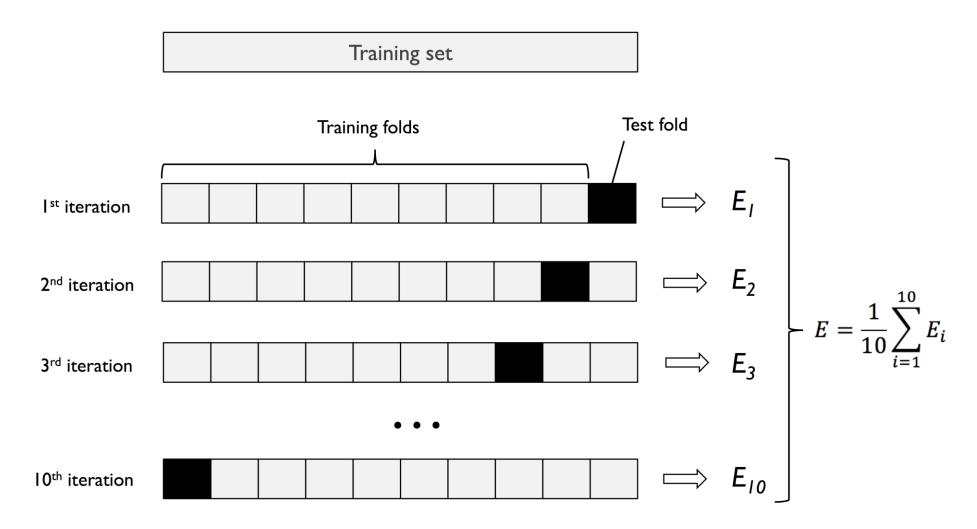


 The method of splitting the full dataset into training partition and an evaluation partition is referred to as the hold out method





K-fold cross-validation (CV)





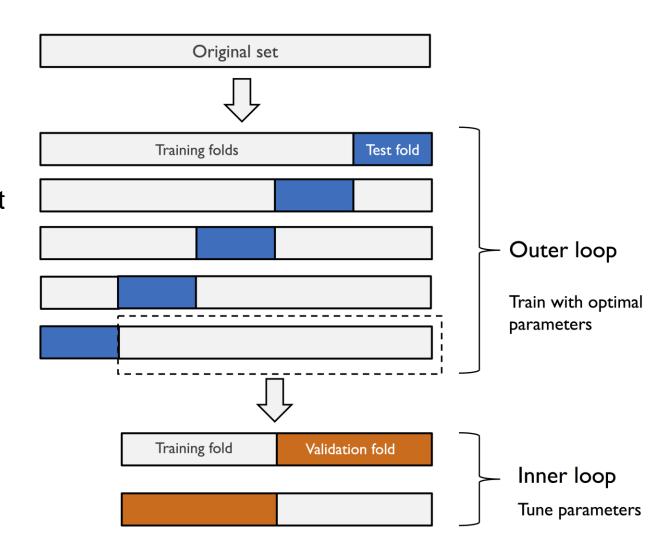
Grid search & randomized search

- Grid search:
 - Brute-force exhaustive search through grid of specified set of hyperparameters
- Randomized search:
 - We don't specify a grid of hyperparameter combinations to search exhaustively
 - Instead we specify
 - A range of possible hyperparameter values (could be continous)
 - Parameter specific probability distributions
 - Max set of iterations



Nested cross-validation

- Cross-validation loop within a crossvalidation loop
- Addresses the fact that the initial split between the training/val set and test set is also sensitive to how the split is done
- Becomes very computationally expensive
- Is rarely done when you are working with datasets of over a certain size





Confusion matrix

- It is a square matrix showing the classes that a model predicts versus the classes of the ground truth
- Let 1 be positive and 0 be negative
- **TP**: Positive sample model predicts positive
- **FP**: Negative sample model predicts positive
- **FN**: Positive sample model predicts negative
- TN: Negative sample model predicts negative

Predicted class PNTrue False positives negatives (TP) (FN) Actual class False True positives negatives (FP) (TN)



Different evaluation metrics

- Precision (PRE), Recall (REC) and F1-score are central metrics in this course
- Precision seeks to measure the amount of TP's in relation to FP's
- Recall (also known as sensitivity in medicine) is equivilant to TPR

$$\mathbf{Recall} = \frac{TP}{TP + \mathbf{FN}} \ \mathbf{Precision} = \frac{TP}{TP + \mathbf{FP}}$$

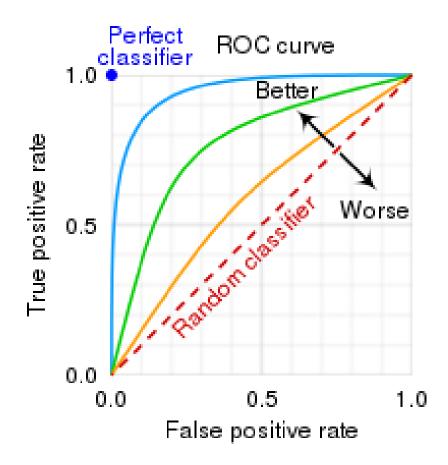
- Often, optimizing for recall might come at the cost of lowering precision
- F1-score is a metric that seeks to combine precision and recall

$$\mathbf{F1} = 2 \times \frac{\text{Precision} \times \text{Recall}}{\text{Precision} + \text{Recall}}$$



Receiver Operator Curve and Area Under Curve

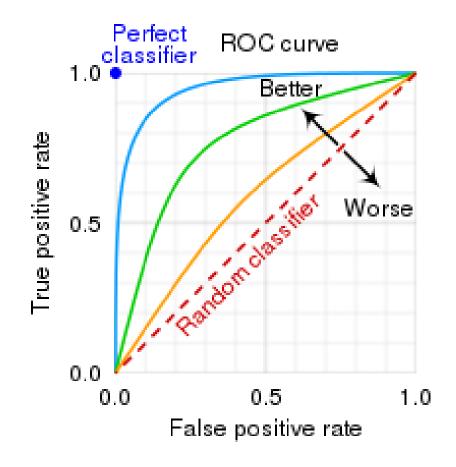
- The Receiver Operator Curve (ROC) is a graphical representation of a classifier performance.
- Plots TPR versus FPR for a binary classifier at different decision thresholds
- Can be understood as trying to visualize how much better a model is than random guessing
- Can be used with any classifier that applies a decision boundary (the majority of classifiers)
- Online illustration





Receiver Operator Curve and Area Under Curve

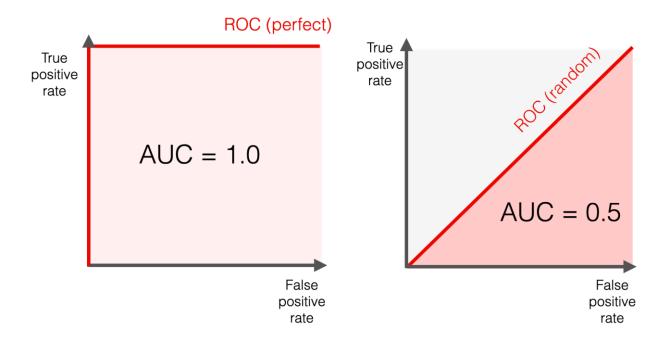
- Illustrates the trade-off between FPR and TPR
- To compute the ROC classifiers must output a probabilistic output/softmax output which can be thresholded
- Diagonal curve is the performance of random guessing (worst possible score)
- Blue curve is better





Receiver Operator Curve and Area Under Curve

- ROC-AUC: Area Under the Receiver Operating Characteristic Curve
- Quantitative measure of overall classifier performance at all possible thresholds





Ensemble Learning

Majority voting

Combinatorial argument

Weight and posterior probabilities

Bagging (parallel)

Repeatedly sample with replacement from the training data.

Train the classifier on each sample and predict new data.

Vote for final prediction.

Random forest (decision trees)

Boosting (sequential)

Very simple individual classifiers (weak learners), usually a decision tree stump (a tree with only one split).

Focus on the samples that are hard to classify

Re-learning based on up-weighting of previous misclassifications.

Ada-Boost

Train m weak learners; compute error rate; compute weight coeff. (inverse error), update sample weights; repeat

Gradient-Boost/XG-Boost

Initial prediction (mean); train m trees on the loss from the previous tree; combine models with learning rate for final prediction

XG: 'unique' trees; gain & similarity score



Regression

Linear regression

Analytical solution vs OLS-regression (gradient descent)

Evaluation metrics (MSE; RMSE; R²)

Simple, Multiple

Residuals vs fitted values

(4) Model assumptions (i. independent errors, ii. normal dist. error, iii. equal variance, iv, linearity

None-linear patterns

Outliers

RANSAC

Outliers; iterate through sub sample, fit model, determent consensus set (inliners/outliners)

Nonlinearity

Transformation; polynomial regression

Regularization

L1 vs L2 vs Elastic

Decision tree regression



Clustering

K-means clustering

Pick k random centroid; cluster based on distance; move centroids to centre; repeat

Groups represented by (Euclidean) distance to single point/object per cluster

Need to set k, simple, tends towards spherical clusters

Suitable for large data sets

K-mean ++

Initiate each centroid using probability weights according to their distance to existing centroids

Soft/fuzzy clustering

Pick k random centroid; cluster based on membership probabilities (m); move centroids to centre; repeat

Also needs k, a bit more complicated, cluster result are similar

Optimal number of clusters - determine k?

Elbow; iterate through a range of k; pick k where distortion (SSE) stops to improve

Silhouette analysis; iterate through range of k; estimate cohesion (intra cluster) and separation (closest cluster) scores; calculate silhouette values, pick k where all clusters above average

Density based clustering

Radius (ε) and minimum number of points

Identify core points; for each core points; assign cluster to density connected points; remaining points are noise

Non-spherical clusters, no need to pick a *k*-value

Hierarchical clustering

Agglomerative; top-down vs Divisive; bottom-up

Distance metric and linkage (merging criterion for clusters)

Single, complete, Wards, Average; Centroid, Median

Can be cut at desired level or number of clusters afterwards



What is the elbow method in clustering?

A technique to determine the optimal number of clusters by plotting the sum of _____ against the number of _____ and selecting the elbow point.

Squared Errors
Absolute Differences
Variances

Classes
Neighbors
Clusters



What is the difference between LDA and PCA?

• LDA is a _____ method that considers the _____,

| Supervised | Cluster centers |
| Cluster centers |
| Cluster centers |
| Class labels |
| Feature weights

while PCA is an _____ method that does not.

Supervised
Unsupervised



What is an ensemble in machine learning?

- A. A model trained on one subset
- B. Converts unsupervised to supervised
- C. Combines multiple models for better performance



$$J_m = \sum_{i=1}^n \sum_{j=1}^k w_{ij}^m \|x_i - \mu_j\|^2$$

What does the parameter m in the Fuzzy C-Means objective function control?

- A. Number of clusters
- B. Distance metric
- C. Fuzziness of the clustering
- D. Learning rate



Rearrange the following steps into the correct order for the K-Means algorithm:

- A. Repeat until centroids no longer change significantly.
- B. Assign each data point to the nearest centroid.
- C. Update each centroid based on the mean of its assigned points.
- D. Randomly initialize k centroids.

Answer:

$$D \rightarrow B \rightarrow C \rightarrow A$$



