

# Machine Learning – winter term 2016/17 –

# Chapter 06: Dimensionality Reduction

(and Anomaly Detection)

Prof. Adrian Ulges Masters "Computer Science" DCSM Department University of Applied Sciences RheinMain

### Outline



### 1. Dimensionality Reduction: Motivation

- 2. Feature Selection
- 3. Principal Component Analysis
- 4. PCA Example: "Eigenfaces"
- 5. Anomaly Detection

# Dimensionality Reduction: Motivation



### The Challenge

- ► Reminder: In machine learning, we are usually given d-dimensional samples x<sub>1</sub>,..., x<sub>n</sub> ∈ ℝ<sup>d</sup>
- The number of features d can be high!



 Goal: Reduce d while preserving (or even improving) discriminativity

### Why?

- ▶ Efficiency: faster training, faster application, less storage
- Avoid overfitting: overcoming the curse of dimensionality
- Better interpretability (and maybe visualization) of data

# Dimensionality Reduction: Overview

### Approaches

- feature selection
- feature derivation, typically, by applying transformations

### Example: K-Means for Feature Derivation

- ► Approach: map samples x to clusters (vector quantization)
- ► Alternative 1: store only the cluster number (1 dimension!)
- Alternative 2: store the distances to all centers
  (K dimensions) (see sklearn > KMeans > transform())



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### Feature Selection: Strategies

- ► Goal: Find a subset of features *F* ⊆ {1, ..., *d*} for which we can solve our machine learning problem 'best' (for example: minimizing classification error)
- ▶ This is a **search problem** (brute-force effort: O(2<sup>d</sup>))
- There are three common approaches: wrappers, filters, and embedded methods

### 1. Wrappers

- Wrappers use an explicit evaluation of feature subsets (training and validating classifiers)
- Search can be done in a greedy fashion (adding or removing the 'best' features to the feature set), or by backtracking
- Benefit: It takes the **underlying classifier** into account!
- Usually the most reliable way, but very slow

### Feature Selection: Strategies



### 2. Filters

Assess feature quality by a proxy measure

example: mutual information between feature X and class labels C

$$I(X,C) = \sum_{x \in X} \sum_{c \in C} p(x,c) \cdot \log_2 \left( \frac{P(x,c)}{P(x) \cdot P(c)} \right)$$

- Search strategy: rank features by their quality, pick the K top ones (K determined via cross-validation)
- Filters are cheaper than wrappers, but not as accurate

# Feature Selection: Strategies

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- 3. Embedded Methods
  - ... treat feature selection as a part of model construction (i.e., we find classifier and features in the same process)
  - Optimization is driven towards models with few features
  - Example: logistic regression with regularization
    - the classifier penalizes feature weights, shrinking them to zero
    - features with weight zero are "filtered"!





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# Principal Component Analysis (PCA)

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### Idea (unsupervised!)

- The dataset exhibits a large stretch along some directions. These are the *important* directions.
- Along other directions, there is only little variation.
   These directions are merely *noise* and can be discarded.
- PCA is about finding the *important* directions (or principal components) of the data

### Principal Components: Formalization

- In the following, we assume our data points to be centered around the origin (otherwise, we simply shift the data beforehand)
- What is the most important direction? What is the second most important direction, etc.?

### **PCA: Illustration**

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### What are the Principal Components here?



# Principal Components: Derivation



Obviously, the principal components seem to be related to the data's covariance matrix. Let's find out  $how^1$ 

<sup>1</sup>cmp. Marsland, page 228f

### Principal Components: Derivation



### Principal Components: Derivation



### Principal Components: Illustration





# PCA: Training



function PCA\_TRAIN( $\mathbf{x}_1, ..., \mathbf{x}_n, K$ ) // given: samples  $\mathbf{x}_1, ..., \mathbf{x}_n \in \mathbb{R}^d$  $\mu := \frac{1}{\pi} \sum_{i} \mathbf{x}_{i}$ 2 for i = 1, ..., n:  $\mathbf{x}_i := \mathbf{x}_i - \mu$  // shift the samples to mean zero 4 stack the samples into an  $n \times d$  data matrix X 5  $\Sigma := \frac{1}{n} \left( X^T \cdot X \right)$ // covariance matrix 6 compute  $\Sigma$ 's eigenvalues  $\lambda_1, ..., \lambda_d$  and eigenvectors  $\mathbf{p}_1, ..., \mathbf{p}_d$ (sorted in descending order of the eigenvalues) 8 stack  $\mathbf{p}_1, \dots, \mathbf{p}_K$  as rows into a  $K \times d$ -Matrix  $P_K$ 9 return  $P_{K}, \mu$ 

#### Remarks

- **Training**: Compute the top *K* eigenvectors of the data's covariance matrix
- ▶ **Application**: reduces the samples from *d* to *K* dimensions

1function PCA\_APPLY(x)// given: a new sample x2return  $P_K \cdot (\mathbf{x} - \mu)$ // dimension of return value: K

### PCA: Remarks

Heuristic for choosing K: Preserve α (e.g., 90%) of the eigenvector's total energy

$$\mathcal{K} := \min_{\mathcal{K} \in \{1,...,d\}} \; \; ext{such that} \; \; \sum_{k=1} \lambda_k \geq lpha \cdot \sum_{k=1} \lambda_k$$

Κ

d

► Given a reduced K-dimensional feature y = (y<sub>1</sub>,..., y<sub>K</sub>), we can reconstruct x (to some extent) from x':

$$\mathbf{x}' = \mu + y_1 \cdot \mathbf{p}_1 + y_2 \cdot \mathbf{p}_2 + \dots + y_K \cdot \mathbf{p}_K$$

• We call  $||\mathbf{x} - \mathbf{x}'||$  the reconstruction error.



### **PCA: Illustration**





### From PCA to Whitening

We learned above: Given a feature vector  $\mathbf{x}$  (and the  $d \times d$  matrix P with  $\Sigma$ 's eigenvectors as rows), by applying the transformation  $P \cdot \mathbf{x}$ , the **covariance matrix** of the data becomes

$$\begin{pmatrix} \lambda_1 & 0 & \dots & 0 \\ 0 & \lambda_2 & 0 & \dots & 0 \\ \dots & \dots & \dots & \dots & \dots \\ 0 & \dots & \dots & 0 & \lambda_d \end{pmatrix}$$

Each value  $\lambda_i$  is a variance within one dimension. We turn these variances into 1, simply by dividing by the standard deviation  $\sqrt{\lambda_i}$ . This leads to the **whitening** transform (see Chapter 'Features'):

$$Diag(\frac{1}{\sqrt{\lambda_1}}, \frac{1}{\sqrt{\lambda_2}}, ..., \frac{1}{\sqrt{\lambda_d}}) \cdot P \cdot \mathbf{x}.$$

Applying this transformation turns the data's covariance matrix into the identity I and decorrelates the features.



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- We can apply PCA to images, simply by scaling images to a standard resolution (say, N × M) and stacking all pixel values into sample vectors of dimension d = N × M
- Note: The principal components are (N · M)-dimensional, too! (i.e., we can visualize them as images)
- ► Example: PCA for face recognition → "eigenfaces" (apply PCA to lots of face images)





Mean face (top left) and the first 7 principal components (=eigenfaces)

Some eigenfaces from a different face dataset



\*



Reconstruction of a face using  $0, 8, 16, \dots$  eigenfaces





- Which of these images does not show a human face?
- Approach: Compare the original image with its PCA reconstruction



faces are <u>well</u> reconstructed



non-faces are poorly reconstructed

### **Eigenfaces: Discussion**

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The Eigenfaces approach for **face detection** is **simple and powerful**, but it has some shortcomings:

- The results depend strongly on *illumination*, *shadows*, and *local changes*, e.g. glasses or beards
  - include those effects in the training set
  - learn smaller components (eye, mouth, ...)
  - (illumination): normalize all images
- Only faces of *fixed size* are detected
  - create scaled versions of the input image before searching
- Already small rotations of the head change the result
  - include rotated faces into the training set.
  - try to make the image upright before the applying PCA

Since eigenfaces [8], there have been at least two generations of more elaborate face detection methods [9, 7].

### PCA: Discussion



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# Anomaly Detection: Mission Statement images from [1] [3] [2]

**Goal**: identify samples that do not conform to an expected pattern, or to other samples in the dataset [6].

### Applications

- credit card fraud detection
- detecting tumours in imagery
- detecting technical failures
- finding errors in text
- network intrusion detection





# Anomaly Detection: Types of Anomalies image from [6]

- point anomalies: individual samples that are unusual (typically, in high distance) to the flock
- contextual anomalies: Samples are described by contextual features (e.g., location) and behavioral features (e.g., the temperature). An anomaly occurs if the behavioral features are unusual given the contextual ones (e.g., the temperature is unusually high given the location).
- collective anomalies: a combination of samples that is unusual (whilst the individual samples are not necessarily).
   Example: ... http-web smtp-mail buffer-overflow ssh ftp ...

Our focus here will be on **point anomalies**.



# Anomaly Detection: Characteristics



### Learning Setups

- ► Usually, there are two labels: *normal* vs. *abnormal*
- Labeled training data can be really difficult to find!
- supervised techniques: Training data from both classes given (but often highly imbalanced)
- semi-supervised techniques: Training data only for the normal class
- unsupervised techniques: training data without labels (there may be anomalies, but we do not know when/where)

Absolute Distance as an Anomaly Criterion? image from [6]



### Anomaly Detection: Methods

There is a plethora of anomaly detection methods [6]

- ... some using regular classifiers
- ... some using density-based modeling
- ... some using rule mining

▶ ...

We will only look at two of the most prominent ones:

- a density-based method ("local outlier factor")
- a classification-based method ( "one-class SVMs" ) ightarrow later



# Local Outlier Factor (LOF) [5] $_{image from [4])}$

- Idea: A sample x is a point anomaly if the point density in its surrounding is lower than in its nearest neighbors' surroundings.
- Anomaly Measure: Measure the distance to x's neighbors, measure the same distance for each neighbor, and compare.





### Derivation



# LOF: Derivation



### LOF: Example 2 image from [4]



### LOF: Discussion



### References

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