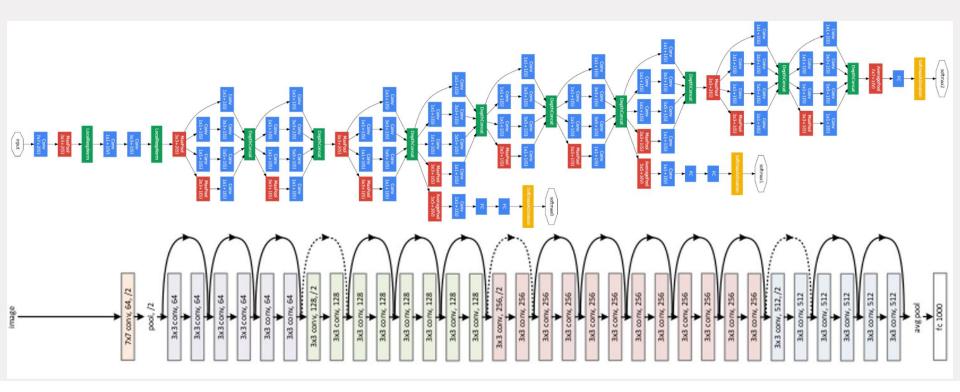


Unsupervised machine learning (8DC00)

Navchetan Awasthi

Previous lecture: deep learning models





Previous lecture: frameworks













Previous lecture: applications

Classification



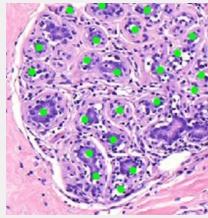
Esteva et al., Nature 2017

Regression



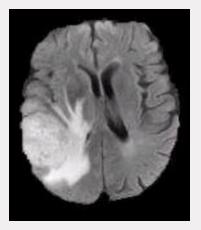
Heslinga et al., SPIE-MI 2019

Detection



Wetstein et al., SPIE-MI 2019

Segmentation



Dong et al., MIAU 2017



Previous lecture: deep learning enabled



- Hardware improvements
- Parallel computing (GPU)



- Digital data
- More (image) data
- = also true for medical imaging



Previous examples required labeled data

learning from labeled data = supervised training

Alternatives:

Unsupervised

Semi-supervised

Reinforcement Learning



Learning outcomes

- Student can describe the difference between supervised and unsupervised learning and name advantages of both methods
- Student can apply K-means to find clusters in data
- Student can explain Principal Component Analysis and motivate dimensionality reduction
- Student can explain the concept of an Autoencoder and motivate why abstract features (latent variables) can be used for a secondary task.



Lecture outline

- Supervised vs unsupervised
- K-means
- Principal component analysis
- Break (15 mins)
- Auto-encoders
- Semi-supervised



Learning strategies

Supervised

Learning from examples (=training data) that are labeled with their desired outputs. The goal is to learn general rules that maps inputs to outputs.

Unsupervised

Learning from examples without labels. The goals are:

- Learning the entire probability distribution that generated a dataset
- Finding structure in data
- Reducing dimensionality → feature learning



Another learning strategy (no exam material)

Reinforcement Learning

Learning by interacting with a dynamic environment to achieve a certain goal (such as driving a vehicle or playing a game against an opponent). The system is provided feedback in terms of rewards and punishments as it navigates its problem space.



Reinforcement learning

RESEARCH

COMPUTER SCIENCE

A general reinforcement learning algorithm that masters chess, shogi, and Go through self-play

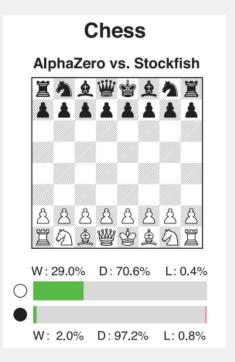
David Silver 1,2 ; Thomas Hubert 1 ; Julian Schrittwieser 1 ; Ioannis Antonoglou 1 , Matthew Lai 1 , Arthur Guez 1 , Marc Lanctot 1 , Laurent Sifre 1 , Dharshan Kumaran 1 , Thore Graepel 1 , Timothy Lillicrap 1 , Karen Simonyan 1 , Demis Hassabis 1 †

The game of chess is the longest-studied domain in the history of artificial intelligence. The strongest programs are based on a combination of sophisticated search techniques, domain-specific adaptations, and handcrafted evaluation functions that have been refined by human experts over several decades. By contrast, the AlphaGo Zero program recently achieved superhuman performance in the game of Go by reinforcement learning from self-play. In this paper, we generalize this approach into a single AlphaZero algorithm that can achieve superhuman performance in many challenging games. Starting from random play and given no domain knowledge except the game rules, AlphaZero convincingly defeated a world champion program in the games of chess and shogi (Japanese chess), as well as Go.

programmers, combined wit alpha-beta search that expar by using a large number of domain-specific adaptations these augmentations, focus Chess Engine Championsh world champion Stockfish (programs, including Deep I architectures (1, 12).

In terms of game tree o substantially harder game is played on a larger board v pieces; any captured opps sides and may subsequently on the board. The strongest as the 2017 Computer Sho world champion Elmo, ha feated human champions use an algorithm similar to puter chess programs, ago optimized alpha-beta seam domain-specific adaptatio

AlphaZero replaces the edge and domain-specific



Silver, Hubert, Schrittweiser et al., Science (2018)



Reinforcement learning in healthcare

Deep Reinforcement Learning for Sepsis Treatment

Aniruddh Raghu Cambridge University

United Kingdom

Matthieu Komorowski

Imperial College London United Kingdom m.komorowski14@imperial.ac.uk Imran Ahmed

Cambridge University United Kingdom

However, not often used in healthcare. Why?

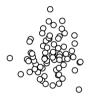
How could we employ reinforcement learning for a surgery robot?

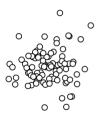


Finding clusters using K-means

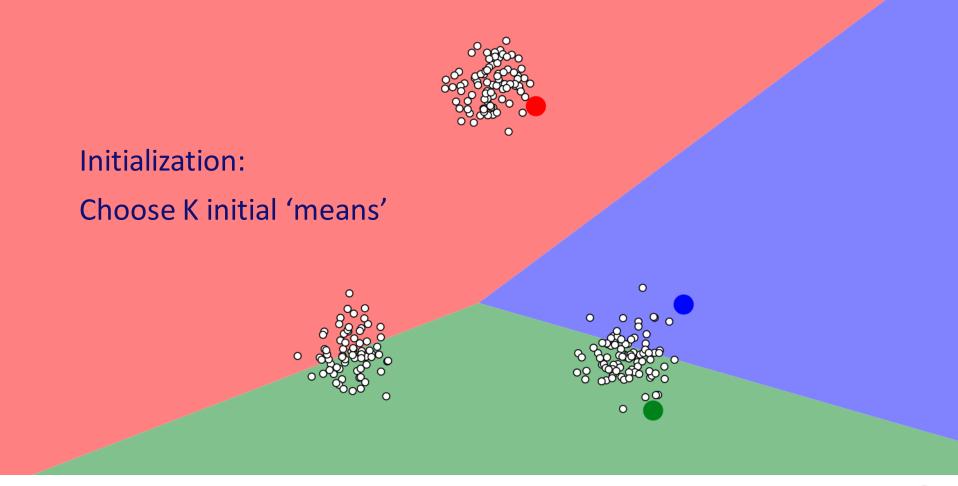
https://www.naftaliharris.com/blog/visualizing-k-means-clustering/



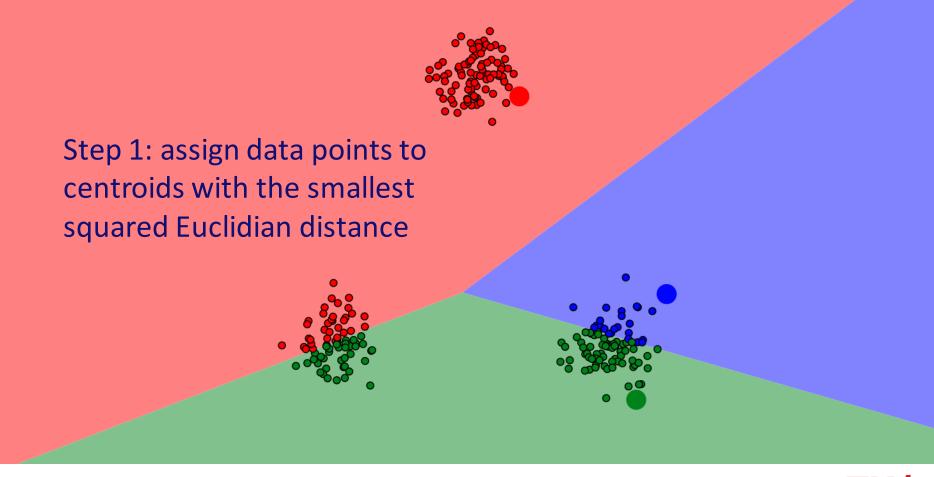














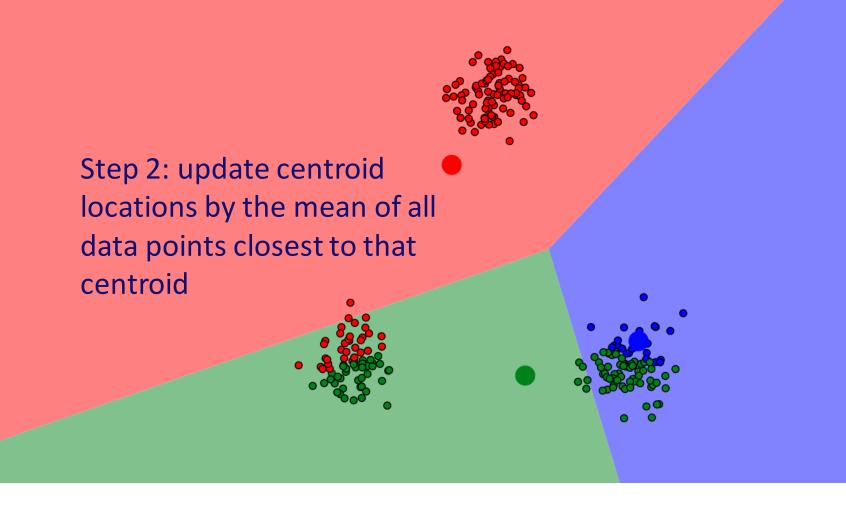
K-means – evaluate clustering performance

Average squared Euclidean distance between each point and the closest cluster:

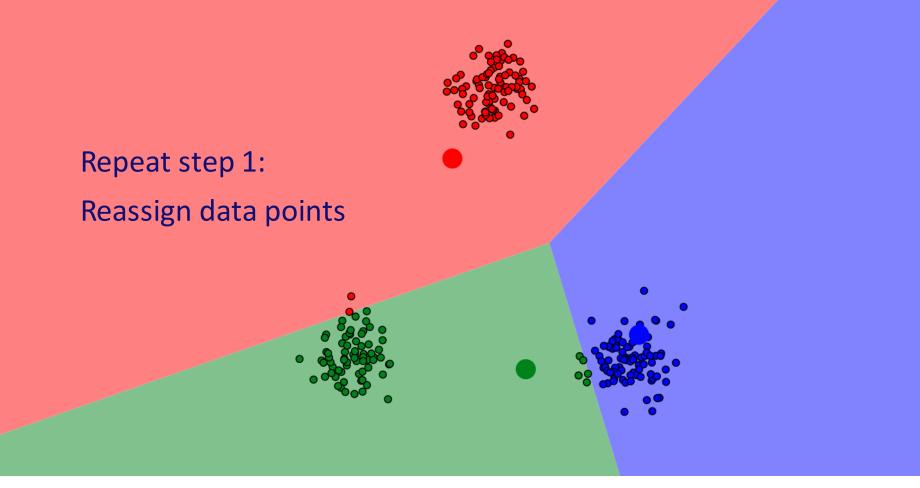
$$J(W) = \frac{1}{N} \sum_{i} \left| \left| \min_{k} (W_k - x_i) \right| \right|_2^2$$

x_i are the points, W are the cluster centroids









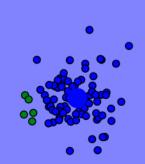




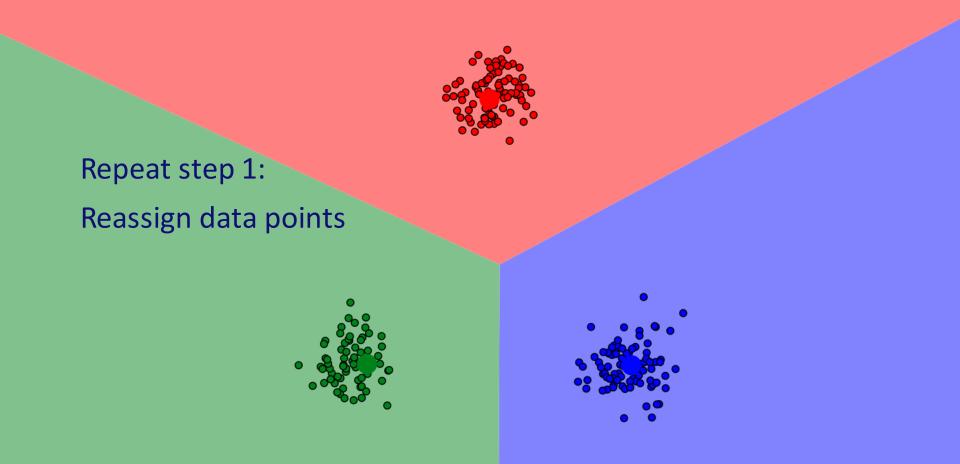
Repeat step 2:

Update centroid locations





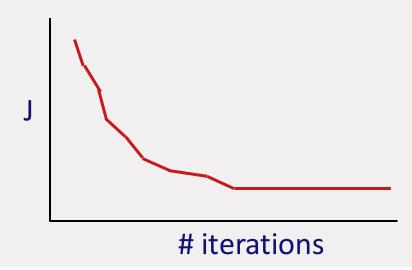






When do we stop?

- When the error J does not decrease anymore
- After n iterations





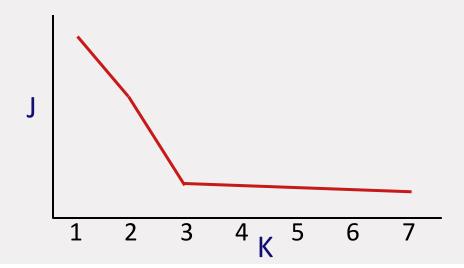
How do we choose initial centroid locations?

- Random
- Farthest points
- Manual?
 - Supervised
 - Difficult for high-dimensional data



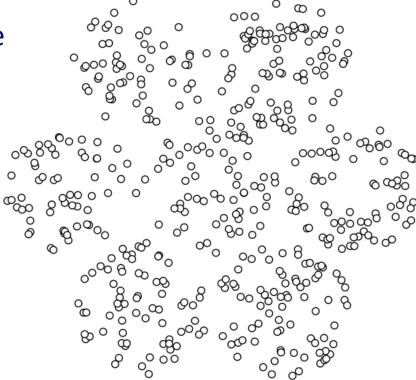
How do we choose K?

K (=number of means) is a hyperparameter

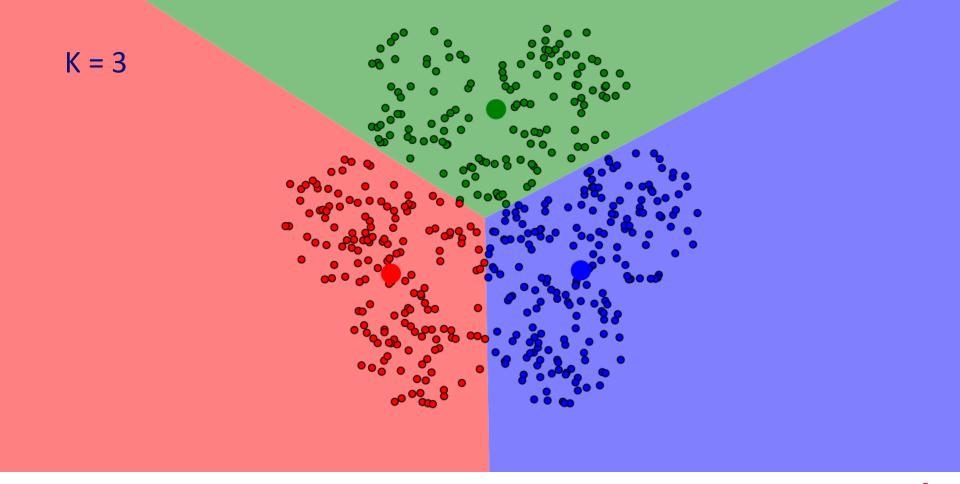




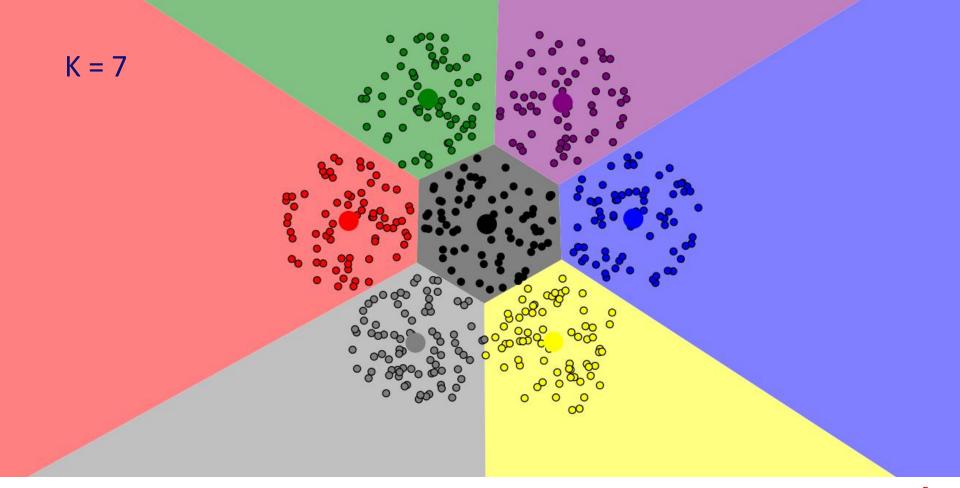
Another example



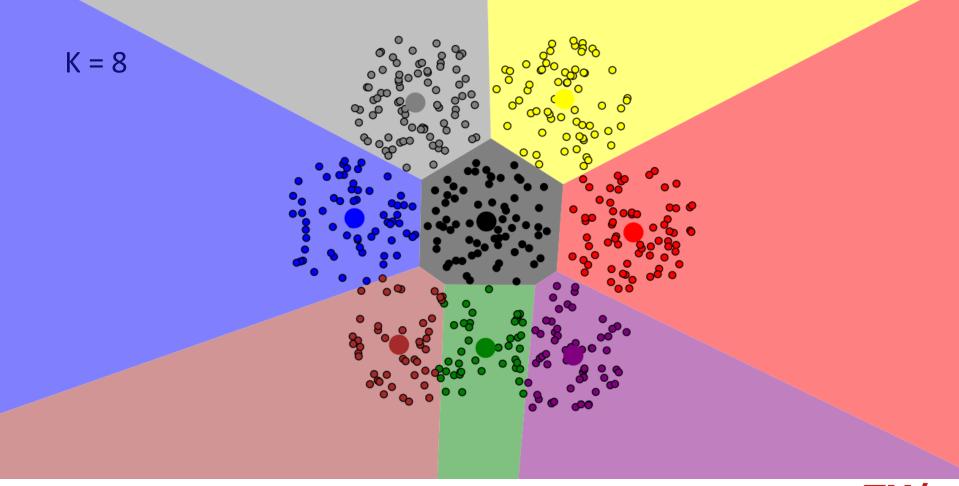












Principal Component Analysis (PCA)

Goal: Finding the principle components that describe our data.

= finding the directions in which the data shows most variation

Useful for dimensionality reduction

E.g. find low-dimensional classification boundaries

Results in better generalization!



Principal Component Analysis

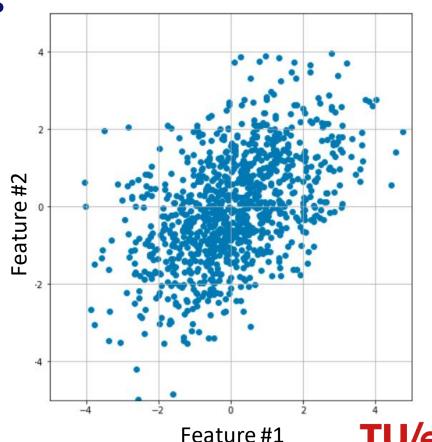
Example data set

M-by-2 matrix X containing M points Sampled from 2D Gaussian distribution

$$\mu_1 = 0$$

$$\mu_2 = 0$$

$$\sum_{i=1}^{n} = \begin{pmatrix} 2 & 1 \\ 1 & 2 \end{pmatrix}$$



Principal Component Analysis

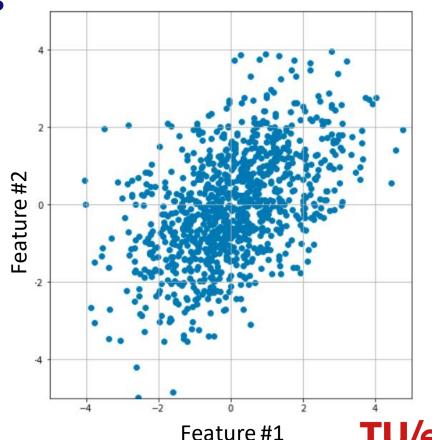
Example data set

M-by-2 matrix X containing M points Sampled from 2D Gaussian distribution

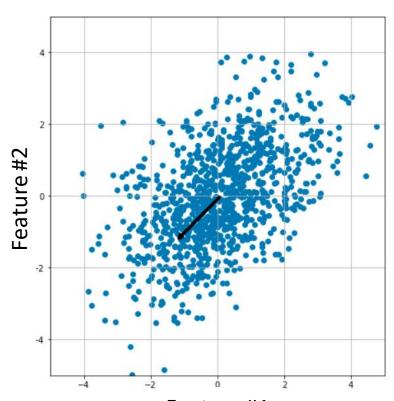
$$\mu_1 = 0$$

$$\mu_2 = 0$$

$$\sum_{i=1}^{n} = \begin{pmatrix} 2 & 1 \\ 1 & 2 \end{pmatrix}$$

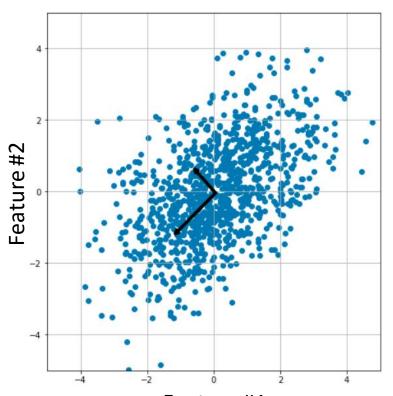


Principal Component 1





Principal component are orthogonal to one another





Finding the principal components

Center data by subtracting mean of each variable

$$\widehat{X} = X - \overline{X}$$

Calculate covariance matrix

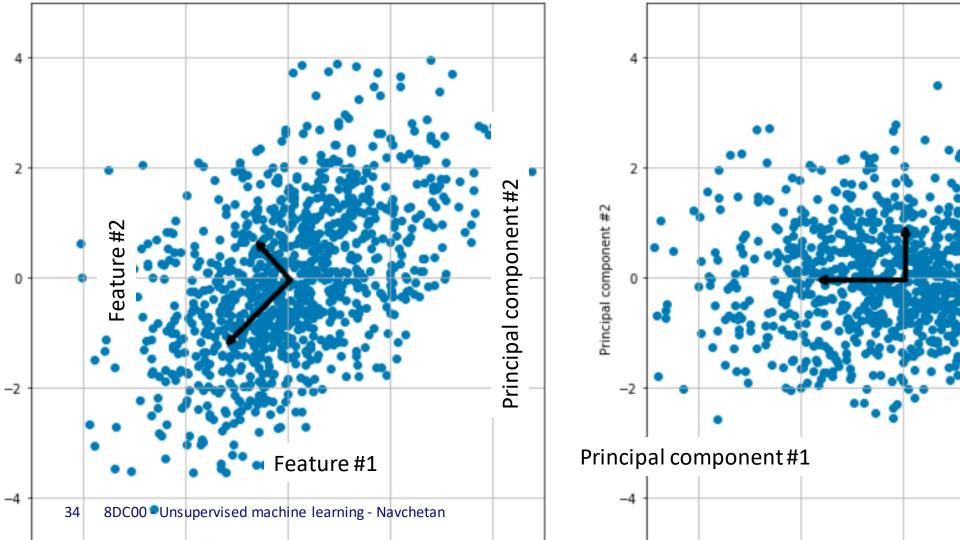
$$\sum = \frac{1}{M-1} X^T X$$

• Singular value decomposition (SVD) to find a matrix \boldsymbol{U} that contains eigenvectors, ordered by largest to smallest variance

→ principal components

lacksquare Multiply \widehat{X} with $oldsymbol{U}$ to obtain $oldsymbol{X_{pca}}$





Dimensionality reduction

Instead of using all eigenvectors from \boldsymbol{U} we can select a set of n principal components.

For example, we can select the eigenvectors that contain 95% of the variance.

More info → PCA demo!





Questions so far?

PCA Example:

https://setosa.io/ev/principal-component-analysis/



Break!



https://www.independent.co.uk/news/science/





Lecture outline

- Supervised vs unsupervised
- K-means
- Principal component analysis
- Break (15 mins)
- Auto-encoders
- Semi-supervised
- Self-supervised

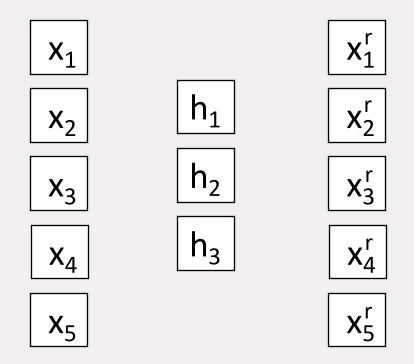


Awesome explanation

https://stats.stackexchange.com/questions/2691/making-sense-of-principal-component-analysis-eigenvectors-eigenvalues



Autoencoder



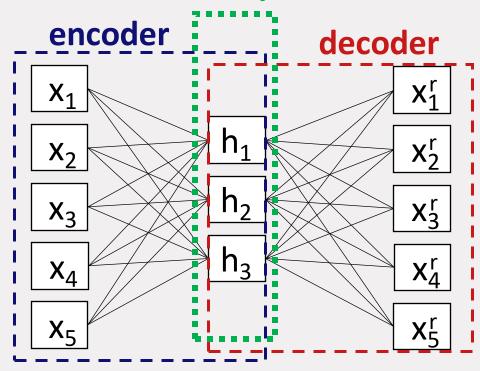
Goal = reconstruct input x_i , using a restricted number of latent variables h_i



latent space encoder decoder X_3



latent space



Encoder:

$$h = f(x)$$

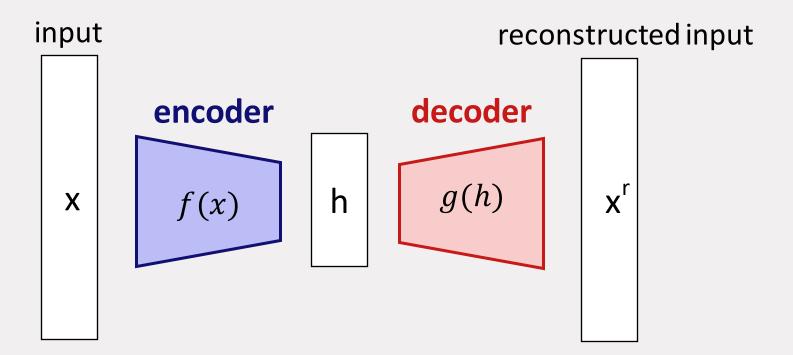
Decoder:

$$x^r = g(\mathbf{h})$$

Penalize dissimmilarity



Autoencoder – a more general representation





Autoencoder

- Encoder/decoder can me simple or complex
- For example: a deep convolutional neural network

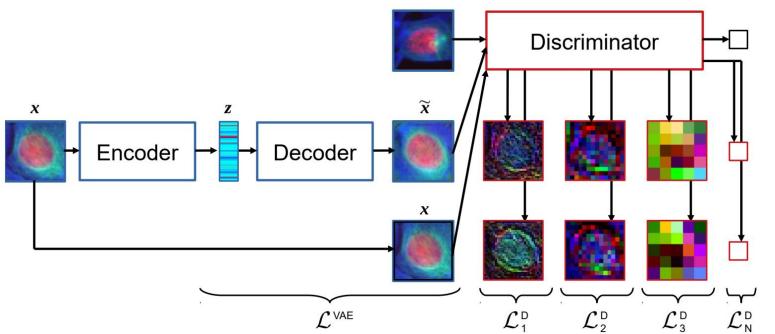
Applications

- Dimension reduction!
- Latent variables can be used for secondary objective, e.g. classification
- Denoising (by adding noise to the input and reconstructing the original)
- Generative models generating new (image) data



Example: Unsupervised representation learning to capture single-cell phenotypic variation

Lafarge et al., MIDL 2019

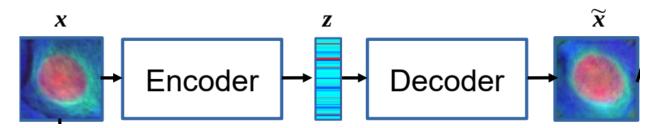




Example: Unsupervised representation learning to capture single-cell phenotypic variation

Lafarge et al., MIDL 2019

- Human MCF7 cells
- Treated with different compounds
- Latent space representation (here called z) used to predict compounds with 1-nearest-neighbors





we will train the autoencoder to map noisy digits images to clean digits images.



```
from keras.datasets import mnist
import numpy as np
(x train, ), (x test, ) = mnist.load data()
x train = x train.astype('float32') / 255.
x test = x test.astype('float32') / 255.
x train = np.reshape(x train, (len(x train), 28, 28, 1))
x \text{ test} = \text{np.reshape}(x \text{ test}, (\text{len}(x \text{ test}), 28, 28, 1))
noise factor = 0.5
x train noisy = x train + noise factor * np.random.normal(loc=0.0, scale=1.0, size=x train.shape)
x test noisy = x test + noise factor * np.random.normal(loc=0.0, scale=1.0, size=x test.shape)
x train noisy = np.clip(x train noisy, \theta., 1.)
x test noisy = np.clip(x test noisy, 0., 1.)
```



```
n = 10
plt.figure(figsize=(20, 2))
for i in range(1, n + 1):
    ax = plt.subplot(1, n, i)
    plt.imshow(x_test_noisy[i].reshape(28, 28))
    plt.gray()
    ax.get_xaxis().set_visible(False)
    ax.get_yaxis().set_visible(False)
plt.show()
```



Can autoencoder learn to recover the original digits?



```
input img = keras.Input(shape=(28, 28, 1))
x = layers.Conv2D(32, (3, 3), activation='relu', padding='same')(input img)
x = layers.MaxPooling2D((2, 2), padding='same')(x)
x = layers.Conv2D(32, (3, 3), activation='relu', padding='same')(x)
encoded = layers.MaxPooling2D((2, 2), padding='same')(x)
# At this point the representation is (7, 7, 32)
x = layers.Conv2D(32, (3, 3), activation='relu', padding='same')(encoded)
x = layers.UpSampling2D((2, 2))(x)
x = layers.Conv2D(32, (3, 3), activation='relu', padding='same')(x)
x = layers.UpSampling2D((2, 2))(x)
decoded = layers.Conv2D(1, (3, 3), activation='sigmoid', padding='same')(x)
autoencoder = keras.Model(input img, decoded)
autoencoder.compile(optimizer='adam', loss='binary crossentropy')
```









'Supervised' learning terminology

Supervised methods

Weakly Supervised

Semisupervised Unsupervised methods

Selfsupervised



Semi-supervised

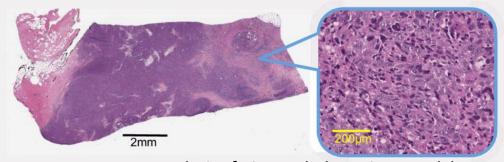
Global labels

E.g. A single diagnosis is based on a series of retinal fundus images

NIH National Eye Institute

Partially labeled data

E.g. Only cells in part of a whole slide histopathology image are segmented



Analysis of Histopathology, Jimenez-del-Toro



Some remarks on semi-supervised learning

- Fewer labeled data needed
- For many medical applications data is still limited,
 e.g. because disease is rare
- Use knowledge from a related task

Humans also learn in a semi-supervised fashion



Summary

- Supervised versus unsupervised
- Finding structures (e.g. K-means)
- Dimension reduction (e.g. PCA, autoencoders)
- Semi-supervised learning

