

# Dimensionality reduction

<https://github.com/Maarten-vd-Sande/Teaching>

Maarten van der Sande

# Content

- ▶ Principal Component Analysis (PCA)
  - ▶ K-means clustering
- ▶ t-distributed Stochastic Neighbour Embedding (t-SNE)
  - ▶ Louvain clustering

# Disclaimers

- ▶ I have never actually *\*really\** used the methods

# Disclaimers

- ▶ I have never actually *\*really\** used the methods
- ▶ I'm going to make untrue simplifications

# Disclaimers

- ▶ I have never actually *\*really\** used the methods
- ▶ I'm going to make untrue simplifications
- ▶ Please stop me when something is unclear

# Why dimensionality reduction?

# Why dimensionality reduction?

- ▶ Visualization

# Why dimensionality reduction?

- ▶ Visualization
- ▶ Computational cost (storage, memory, processors, etc.)



# Why dimensionality reduction?

- ▶ Visualization
- ▶ Computational cost (storage, memory, processors, etc.)
- ▶ Reduce chance of over fitting

# PCA

```
run:  
python3 manual_PCA.py
```

# K-means clustering I

How to analyze this data? e.g.

- ▶ We can do regression (fit a line)
- ▶ Cluster the data

# K-means clustering I

How to analyze this data? e.g.

- ▶ We can do regression (fit a line)
- ▶ Cluster the data

Simple cluster algorithm: K-means!

# K-means clustering II

---

## Algorithm 1 K-means clustering

---

1: Initialize K random cluster centroids

2: **repeat**

    Group each point to the nearest centroid

    Move each centroid to the middle of all its points

3: **until** convergence

---

# K-means clustering III

```
run:  
python3 k-means.py
```

# t-SNE I

- ▶ t-SNE fundamentally different from PCA.

# t-SNE I

- ▶ t-SNE fundamentally different from PCA.
- ▶ t-SNE has the "*objective*" to project:
  - ▶ points that are close in high-dimensional space close in low-dimensional space
  - ▶ points that are far in high-dimensional space far in low-dimensional space



# t-SNE I

- ▶ t-SNE fundamentally different from PCA.
- ▶ t-SNE has the "*objective*" to project:
  - ▶ points that are close in high-dimensional space close in low-dimensional space
  - ▶ points that are far in high-dimensional space far in low-dimensional space
- ▶ It does so by moving points in the low-dimensional space so that its "*objective*" is maximized.

# t-SNE I

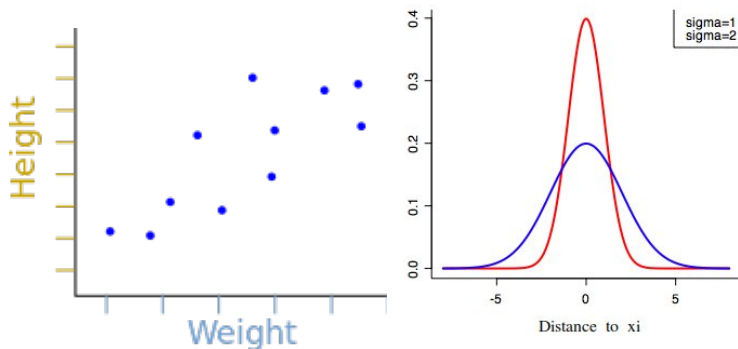
- ▶ t-SNE fundamentally different from PCA.
- ▶ t-SNE has the "*objective*" to project:
  - ▶ points that are close in high-dimensional space close in low-dimensional space
  - ▶ points that are far in high-dimensional space far in low-dimensional space
- ▶ It does so by moving points in the low-dimensional space so that its "*objective*" is maximized.
- ▶ **First we need a metric for closeness and farness!**

## t-SNE II

Closeness (measured in chance) between point  $x_i$  and point  $x_j$ :

$$p_{ji} = \exp\left(\frac{-||x_i - x_j||^2}{\sigma_i^2}\right) \quad (1)$$

this is just a normal distribution! With  $||x_i - x_j||$  the distance between  $x_i$  and  $x_j$ , and  $\sigma^2$  as the variance.



## t-SNE III

Closeness (measured in chance) between point  $x_i$  and point  $x_j$ :

$$p_{ji} = \exp\left(\frac{-||x_i - x_j||^2}{\sigma_i^2}\right) \quad (2)$$

this is just a normal distribution! With  $||x_i - x_j||$  the distance between  $x_i$  and  $x_j$ , and  $\sigma^2$  as the variance.

Now normalize (divide by the sum of all chances):

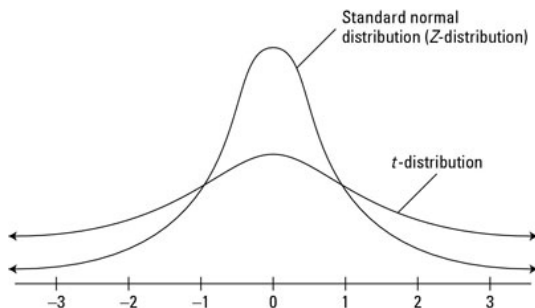
$$p_{j|i} = \frac{p_{ji}}{\sum_k p_{ki}} \quad (3)$$

## t-SNE IV

Our measure for distance in low dimensional space:

$$q_{ji} = 1 + ||x_i - x_j||^{-1} \quad (4)$$

this is a t-distribution! With  $||x_i - x_j||$  the distance between  $x_i$  and  $x_j$ . Luckily we do not have to do any difficult tricks with perplexity here.



## t-SNE IV

Our measure for distance in low dimensional space:

$$q_{ji} = 1 + ||x_i - x_j||^{-1} \quad (5)$$

this is a t-distribution! With  $||x_i - x_j||$  the distance between  $x_i$  and  $x_j$ . Luckily we do not have to do any difficult tricks with perplexity here.

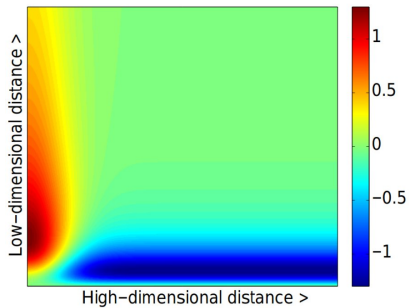
Now normalize (divide by the sum of all chances):

$$q_{j|i} = \frac{q_{ji}}{\sum_k q_{ki}} \quad (6)$$

The "*objective*" is to maximize is the cost function:

$$C = \sum_i \sum_j p_{i|j} \log \frac{p_{i|j}}{q_{i|j}} \quad (7)$$

## t-SNE VI



(c) Gradient of t-SNE.



## t-SNE VII

Now all we have to do is set the variance  $\sigma_i$  for each dot. This is done with the perplexity (hyper)parameter:

## t-SNE VII

Now all we have to do is set the variance  $\sigma_i$  for each dot. This is done with the perplexity (hyper)parameter:

$$Perp(P_i) = 2^{-\sum_j p_{j|i} \log_2 p_{j|i}} \quad (8)$$

Now all we have to do is set the variance  $\sigma_i$  for each dot. This is done with the perplexity (hyper)parameter:

$$\text{Perp}(P_i) = 2^{-\sum_j p_{j|i} \log_2 p_{j|i}} \quad (8)$$

All you need to remember:

**A higher perplexity value results in a higher variance**

Now all we have to do is set the variance  $\sigma_i$  for each dot. This is done with the perplexity (hyper)parameter:

$$\text{Perp}(P_i) = 2^{-\sum_j p_{j|i} \log_2 p_{j|i}} \quad (8)$$

All you need to remember:

**A higher perplexity value results in a higher variance**

```
python3 perplexity.py
```

## t-SNE VIII

```
python3 tSNE.py
```

# Louvain clustering I

Let's imagine a magical function ( $Q$ ) exists that tells us how well our clusters fit on the  $p_j|i$  matrix.

---

## Algorithm 2 Louvain clustering

---

```
1: Each datapoint gets its own cluster.  $c=\{0, 1, \dots, n\}$ 
2: repeat
3:   for each datapoint  $i$  do
4:     for each cluster  $j$  do
5:       if  $Q(c[j]) > Q(c[i])$  then
6:          $c[i] = c[j]$ 
7:       end if
8:     end for
9:   end for
10: until convergence
```

---

## Louvain clustering II

Luckily for us, this magical function exists!

## Louvain clustering II

Luckily for us, this magical function exists!

$$Q = \frac{1}{2m} \sum_{ij} (A_{ij} - \frac{k_i k_j}{2m}) \delta(c_i, c_j) \quad (9)$$

where  $A_{ij}$  represents edge weight between nodes  $i$  and  $j$ ,  
 $k_i$  the sum of the weights attached to node  $i$ ,  
 $2m$  is the sum of all the weights in the graph,  
 $c_i$  is the cluster  $i$  belongs to, and  
 $\delta()$  is one if  $c_i == c_j$  and zero otherwise.



## Louvain clustering II

Luckily for us, this magical function exists!

$$Q = \frac{1}{2m} \sum_{ij} (A_{ij} - \frac{k_i k_j}{2m}) \delta(c_i, c_j) \quad (9)$$

where  $A_{ij}$  represents edge weight between nodes  $i$  and  $j$ ,  
 $k_i$  the sum of the weights attached to node  $i$ ,  
 $2m$  is the sum of all the weights in the graph,  
 $c_i$  is the cluster  $i$  belongs to, and  
 $\delta()$  is one if  $c_i == c_j$  and zero otherwise.

```
python3 louvain.py
```