CPSC 340:
Machine Learning and Data Mining

Multi-Dimensional Scaling

Original version of these slides by Mark Schmidt, with modifications by Mike Gelbart.

Admin

- Assignment 5:
  - Due Friday

- Assignment 6:
  - Remember to request partner

Latent-Factor Models for Visualization

- PCA for visualization:
  - We're using PCA to get the location of the z_i values.
  - We then plot the z_i values as locations in a scatterplot.
- But PCA is a parametric linear model
- PCA may not find obvious low-dimensional structure.
- We could use change of basis or kernels: but still need to pick basis.

Multi-Dimensional Scaling

- Multi-dimensional scaling (MDS) is a crazy idea:
  - Let's directly optimize the z_i values.
    - "Gradient descent on the points in a scatterplot".
  - Needs a "cost" function saying how "good" the z_i locations are.
    - Traditional MDS cost function:

$$f(Z) = \sum_{i=1}^{n} \sum_{j=i+1}^{n} (\|z_i - z_j\| - \|x_i - x_j\|)^2$$

"Try to make scatterplot distances match high-dimensional distance"

sum over pairs of examples

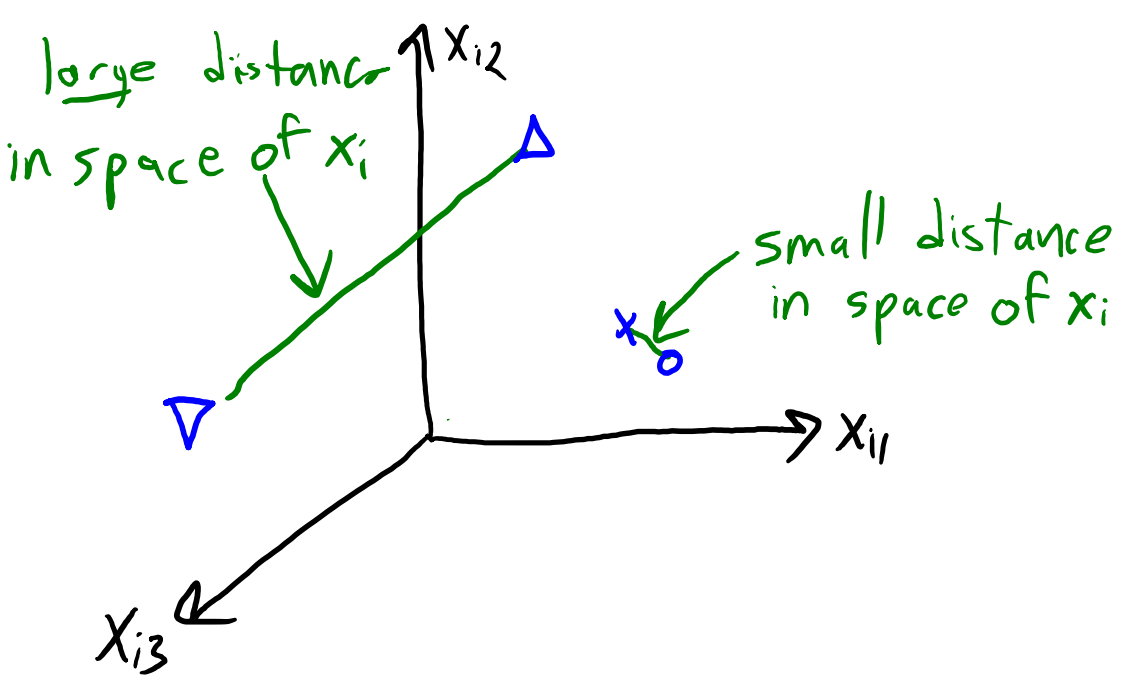
distance in scatterplot

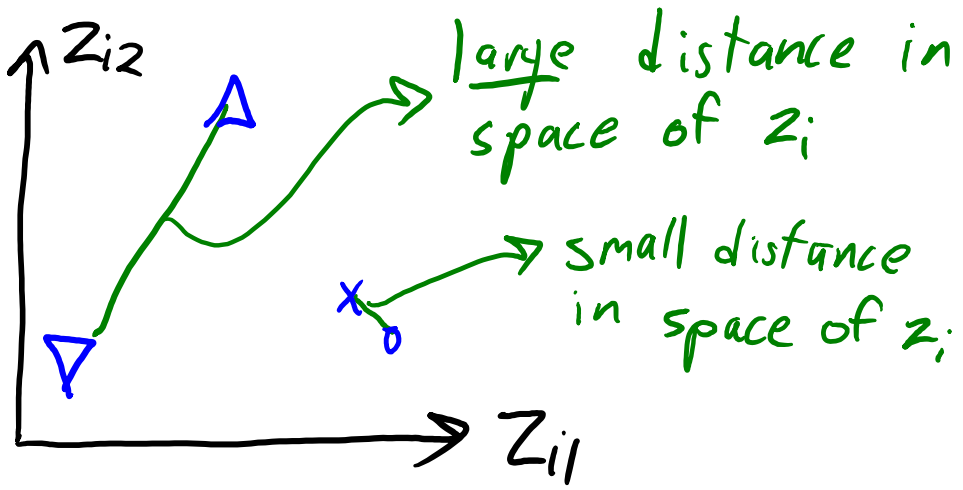
Distance between points in original 'd' dimensions

Multi-Dimensional Scaling

- Multi-dimensional scaling (MDS):
  - Directly optimize the final locations of the z_i values.

$$f(Z) = \sum_{i=1}^{n} \sum_{j=i+1}^{n} \left(\|z_i - z_j\| - \|x_i - x_j\| \right)^2$$





Multi-Dimensional Scaling

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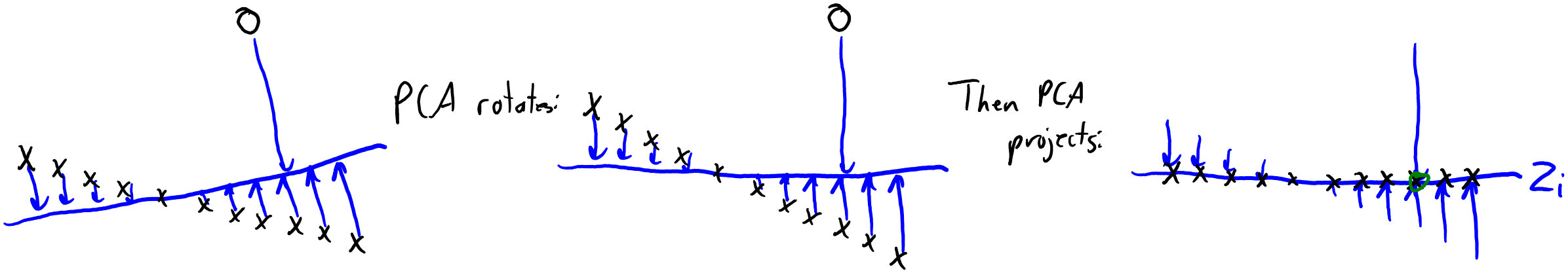
  - Non-parametric dimensionality reduction and visualization:
    - No 'W': just trying to make z_i preserve high-dimensional distances between x_i.



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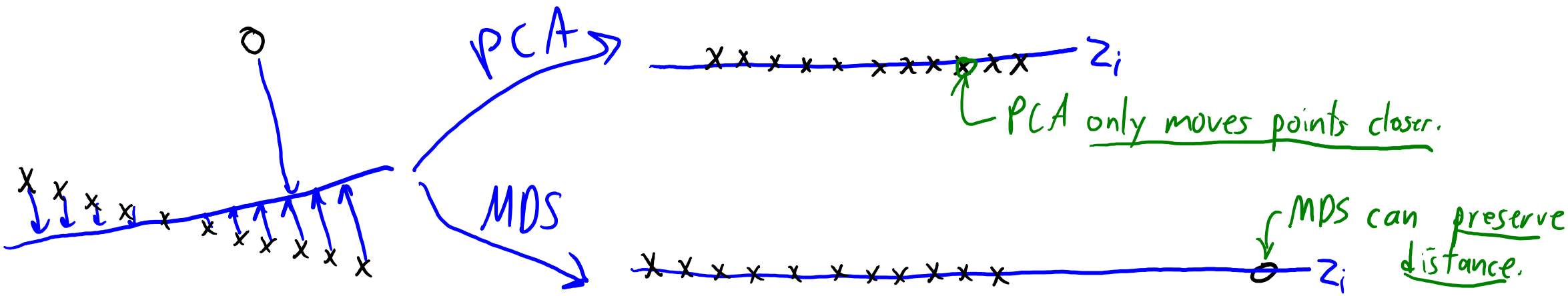

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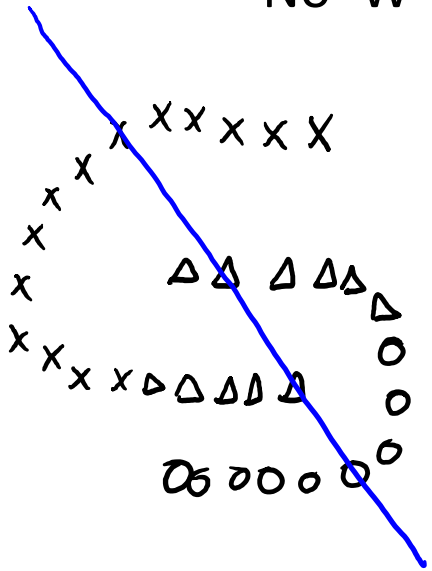

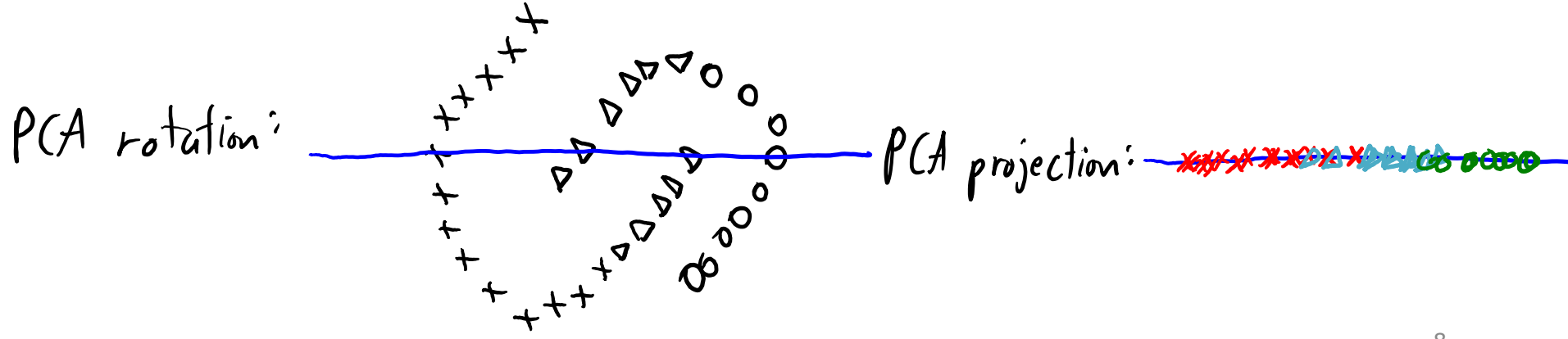

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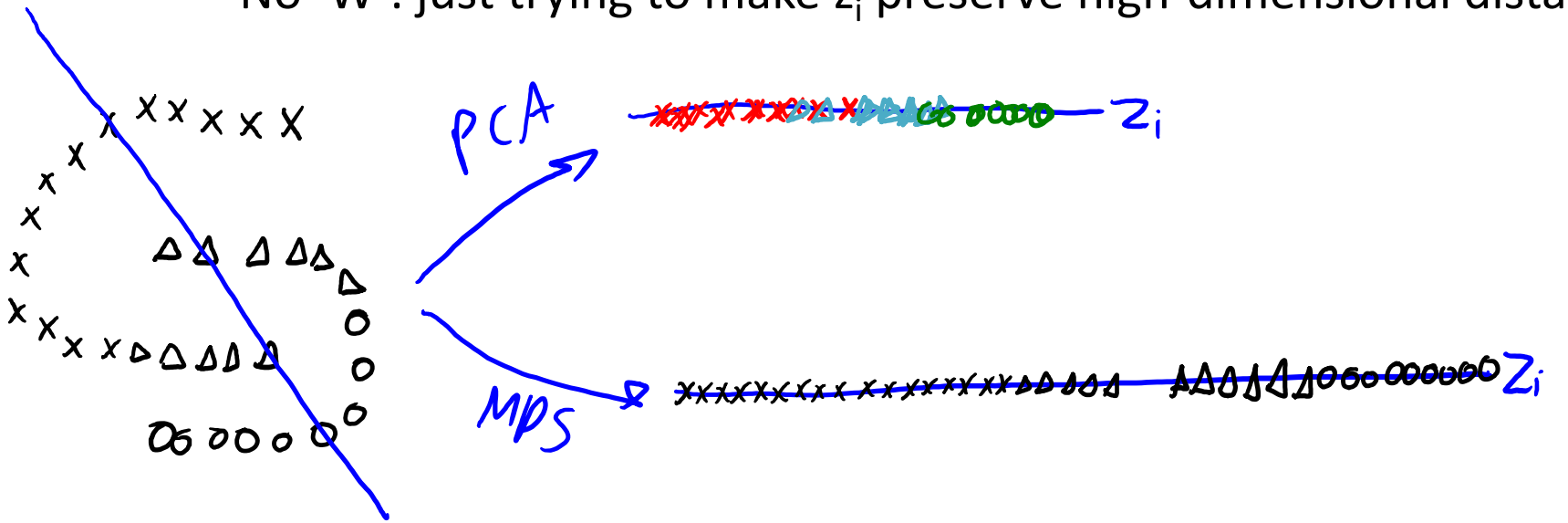

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- Cannot use SVD to compute solution:
  - Instead, do gradient descent on the z_i values.
  - You "learn" a scatterplot that tries to visualize high-dimensional data.
  - Not convex and sensitive to initialization.

Different MDS Cost Functions

- MDS default objective: squared difference of Euclidean norms:

$$f(Z) = \sum_{i=1}^{n} \sum_{j=i+1}^{n} \left(\|z_i - z_j\| - \|x_i - x_j\| \right)^2$$

- But we can make z_i match different distances/similarities:

$$f(Z) = \sum_{i=1}^{n} \sum_{j=i+1}^{n} d_3\left(d_2(z_i, z_j) - d_1(x_i, x_j)\right)$$

  - Where the functions are not necessarily the same:
    - d_1 is the high-dimensional distance we want to match.
    - d_2 is the low-dimensional distance we can control.
    - d_3 controls how we compare high-/low-dimensional distances.

Different MDS Cost Functions

- MDS default objective function with general distances/similarities:

$$f(Z) = \sum_{i=1}^{n} \sum_{j=i+1}^{n} d_3(d_2(z_i, z_j) - d_1(x_i, x_j))$$

- PCA is a special case of MDS
  - using $d_1(x_i,x_j) = x_i^T x_j$ and $d_2(z_i,z_j) = z_i^T z_j$ and centered x_i).

Different MDS Cost Functions

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- Another possibility: $d_1(x_i,x_j) = ||x_i - x_j||_1$ and $d_2(z_i,z_j) = ||z_i - z_j||$.
  - The z_i approximate the high-dimensional L_1-norm distances.

Sammon's Mapping

- Challenge for most MDS models: they focus on large distances.
  - Leads to "crowding" effect like with PCA.
- Early attempt to address this is Sammon's mapping:
  - Weighted MDS so large/small distances are more comparable.

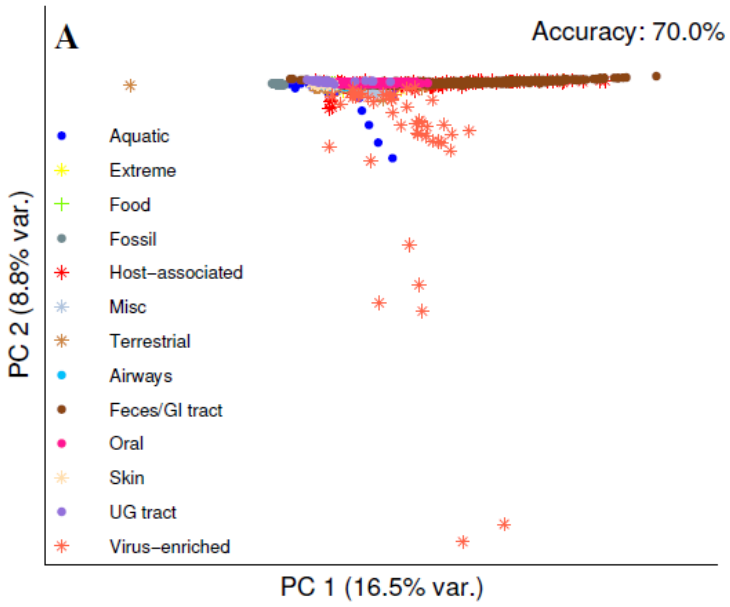
$$f(Z) = \sum_{i=1}^{n} \sum_{j=i+1}^{n} \left(\frac{d_2(z_i, z_j) - d_1(x_i, x_j)}{d_1(x_i, x_j)} \right)^2$$

  - Denominator reduces focus on large distances.

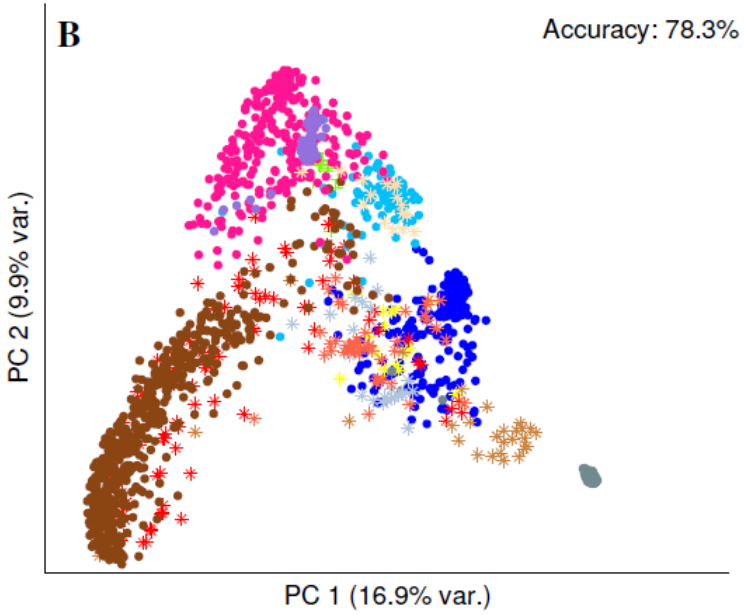
Sammon's Mapping

- Visualizing "metagenomes"

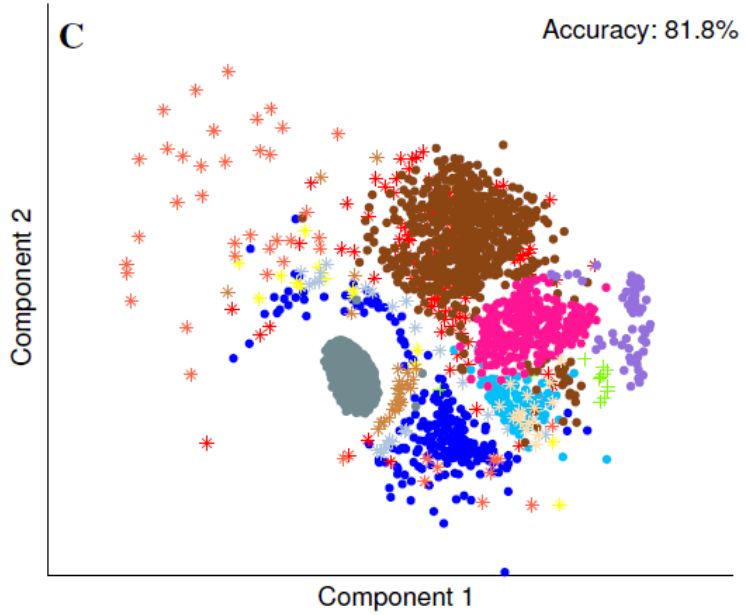
PCA



MDS



MDS + Sammon

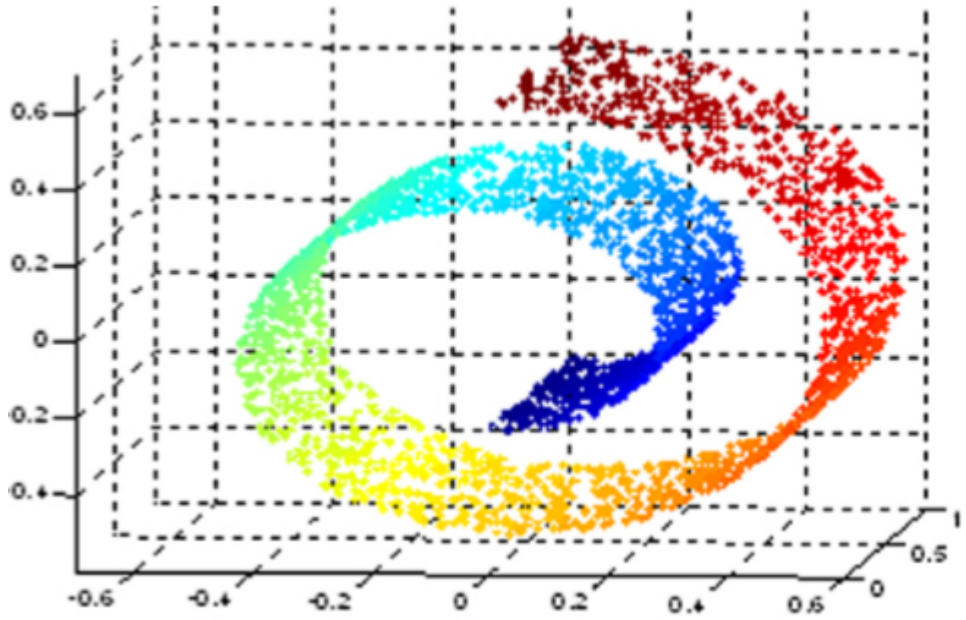


(pause)

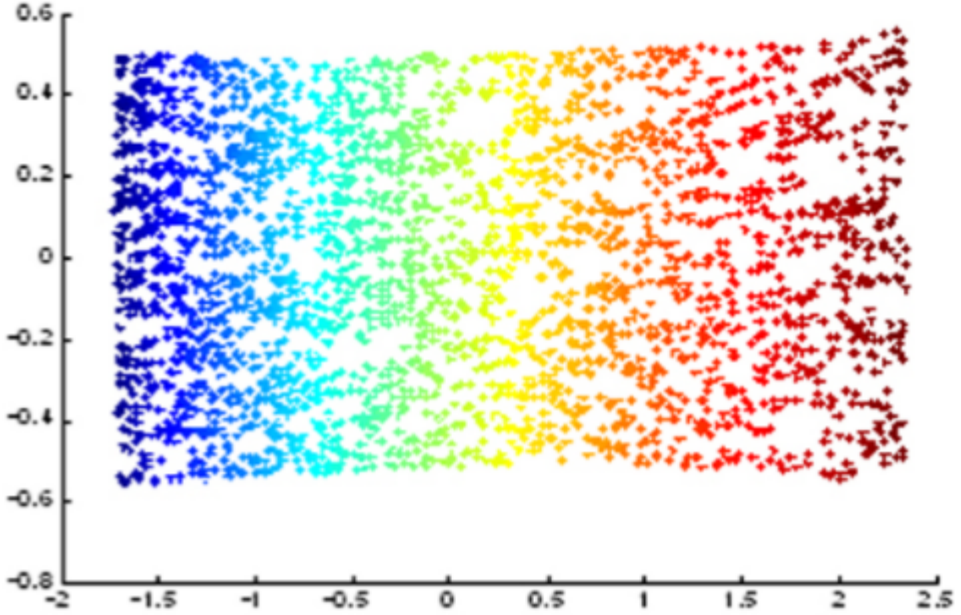
Learning Manifolds

- Consider data that lives on a low-dimensional “manifold”.
- Example is the ‘Swiss roll’:

Original data



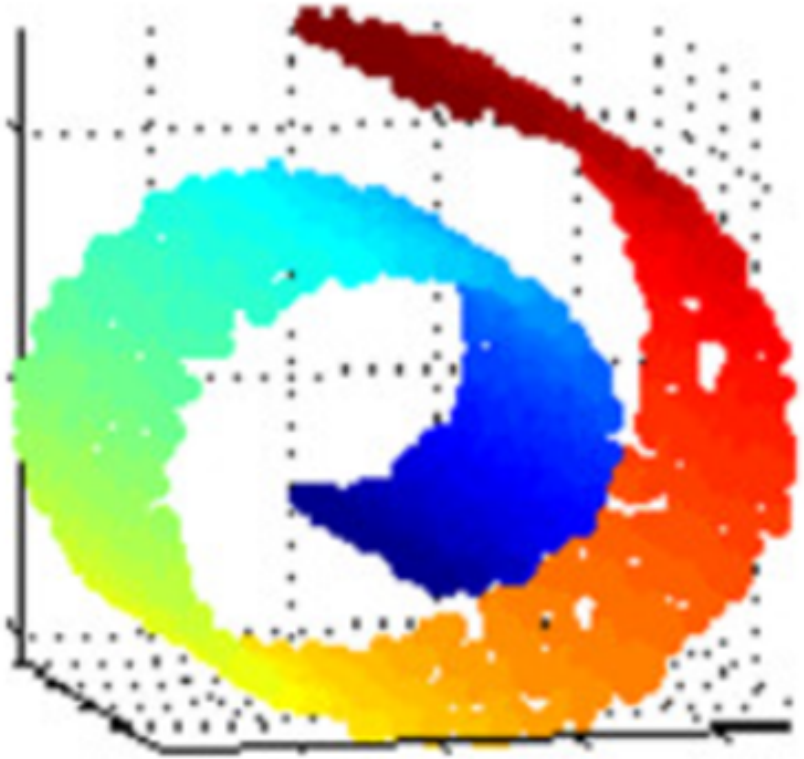
Two-dimensional manifold



http://www.biomedcentral.com/content/pdf/1471-2105-13-S7-S3.pdf

Learning Manifolds

- Consider data that lives on a low-dimensional "manifold".
  - With usual distances, PCA/MDS will not discover non-linear manifolds.



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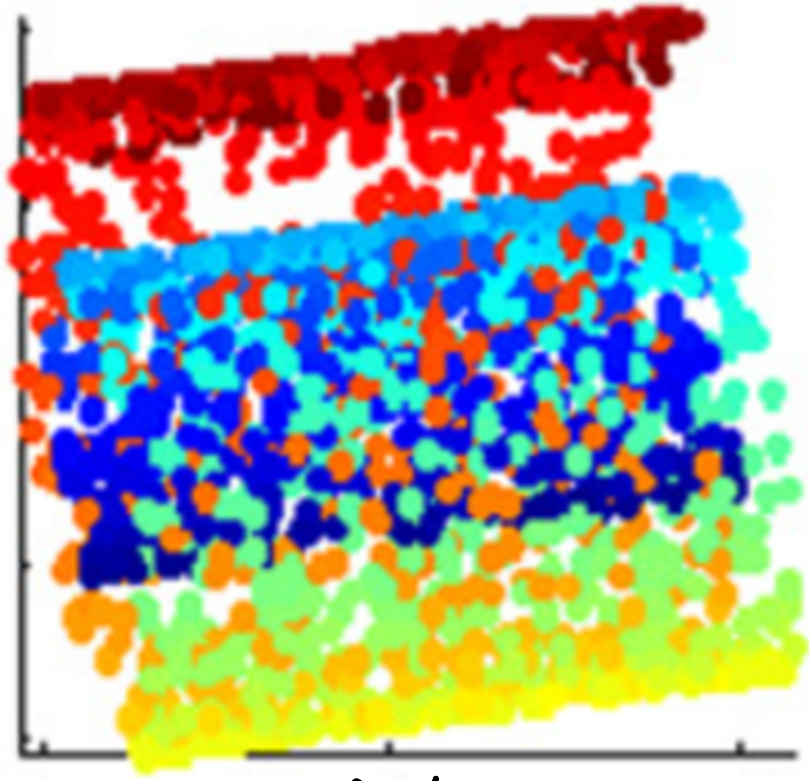
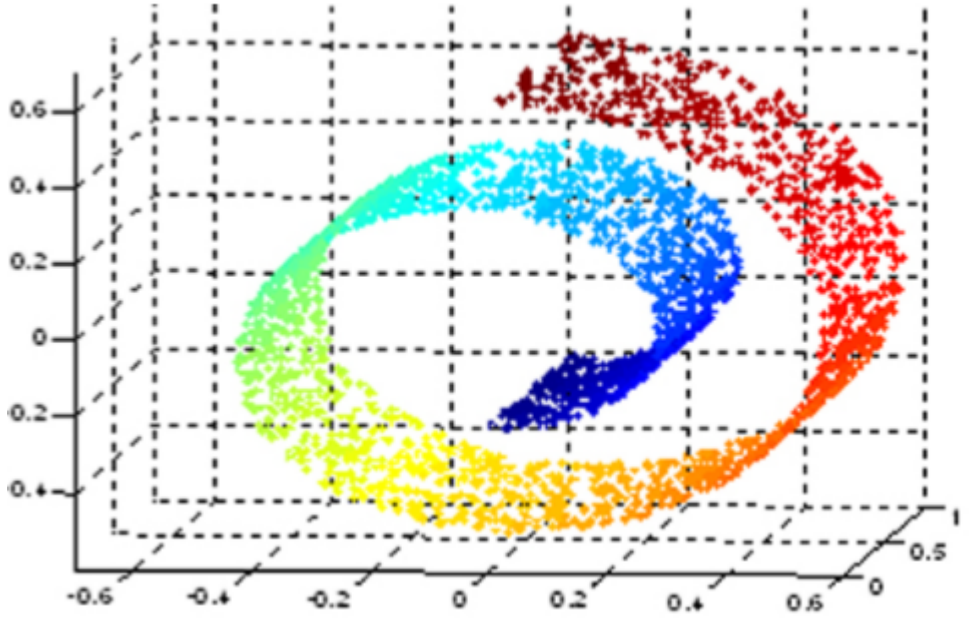


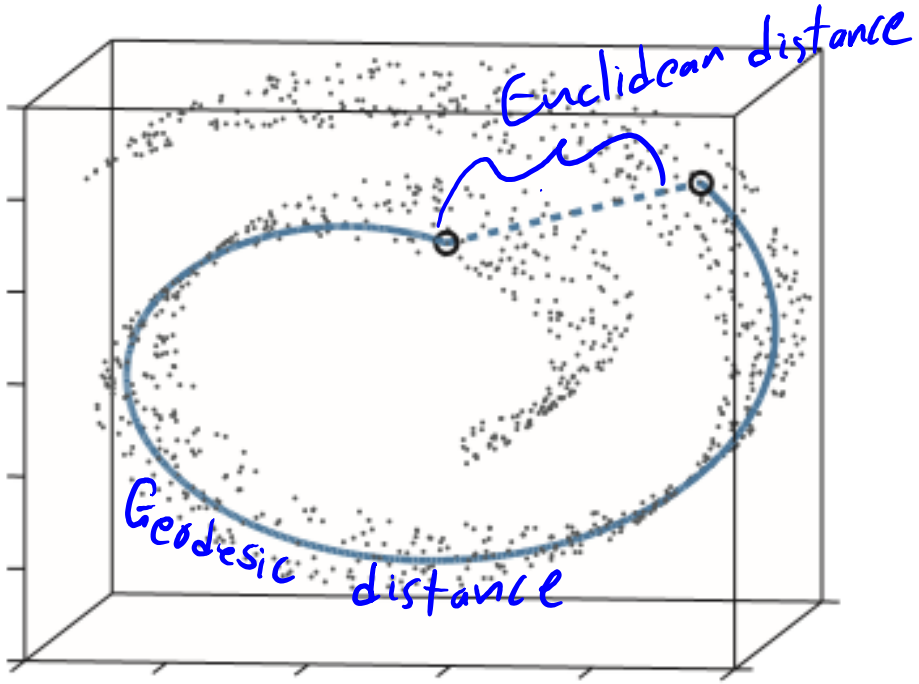
PCA

http://www.peh-med.com/content/9/1/12/figure/F1

Learning Manifolds

- Consider data that lives on a low-dimensional "manifold".
  - With usual distances, PCA/MDS will not discover non-linear manifolds.
- We need geodesic distance: the distance *through* the manifold.

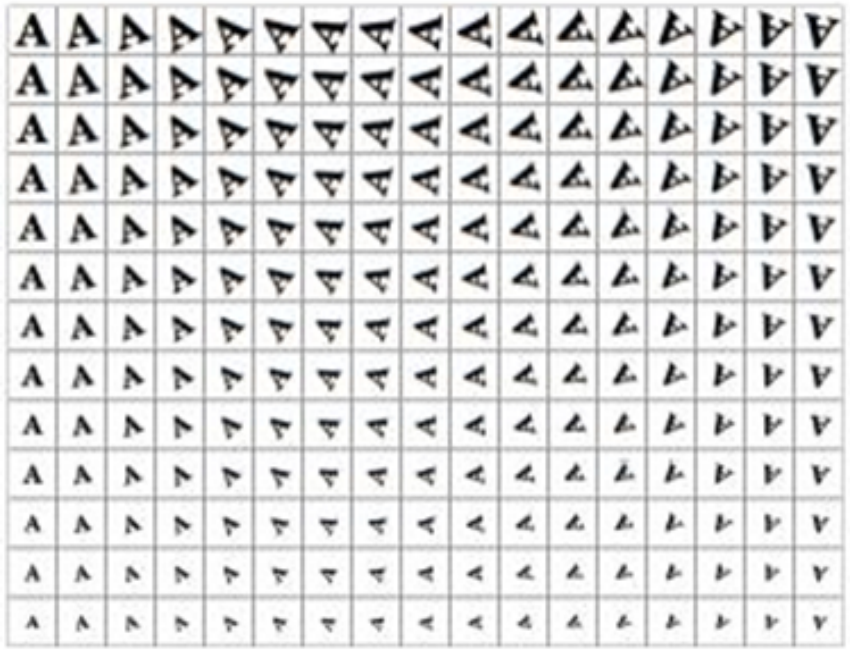




http://www.biomedcentral.com/content/pdf/1471-2105-13-S7-S3.pdf
http://wearables.cc.gatech.edu/paper_of_week/isomap.pdf

Manifolds in Image Space

- Consider slowly-varying transformation of image:

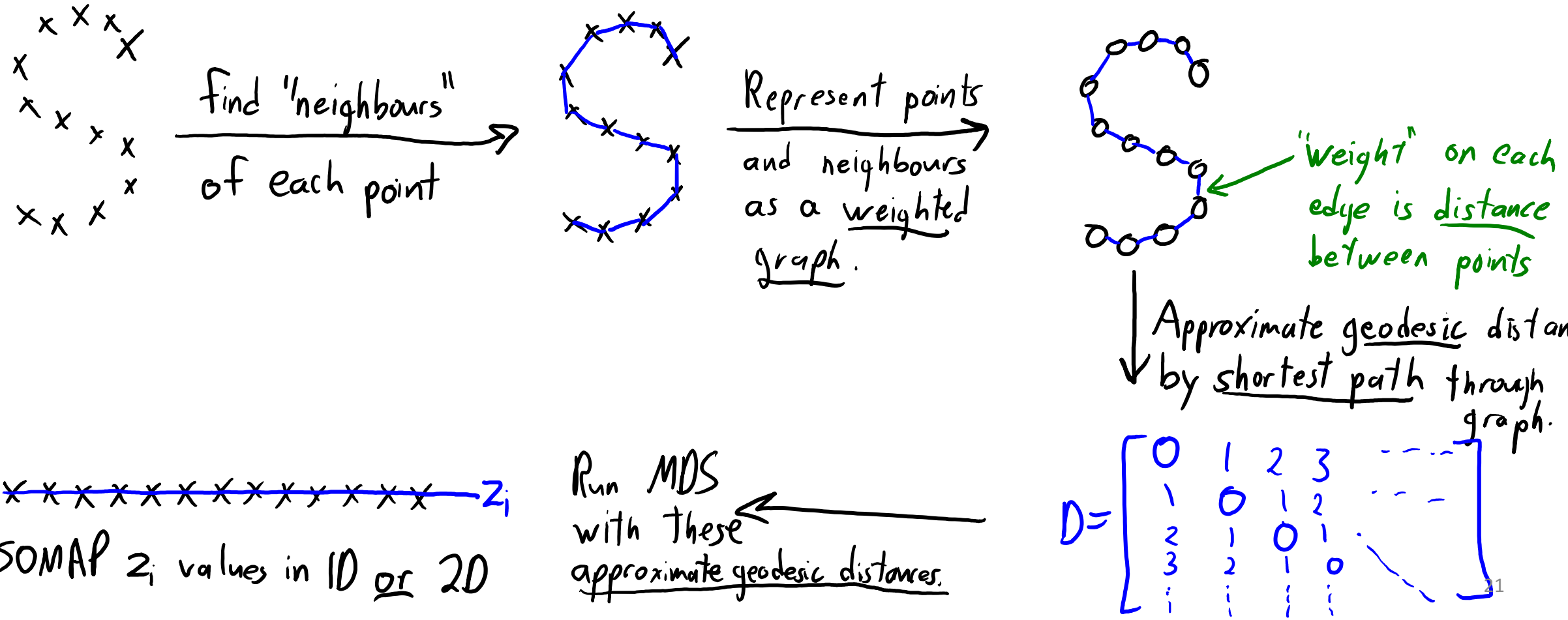


- Images are on a manifold in the high-dimensional space.
  - Euclidean distance doesn't reflect manifold structure.
  - Geodesic distance is distance through space of rotations/resizings.

https://en.wikipedia.org/wiki/Nonlinear_dimensionality_reduction

ISOMAP

- ISOMAP is latent-factor model for visualizing data on manifolds:



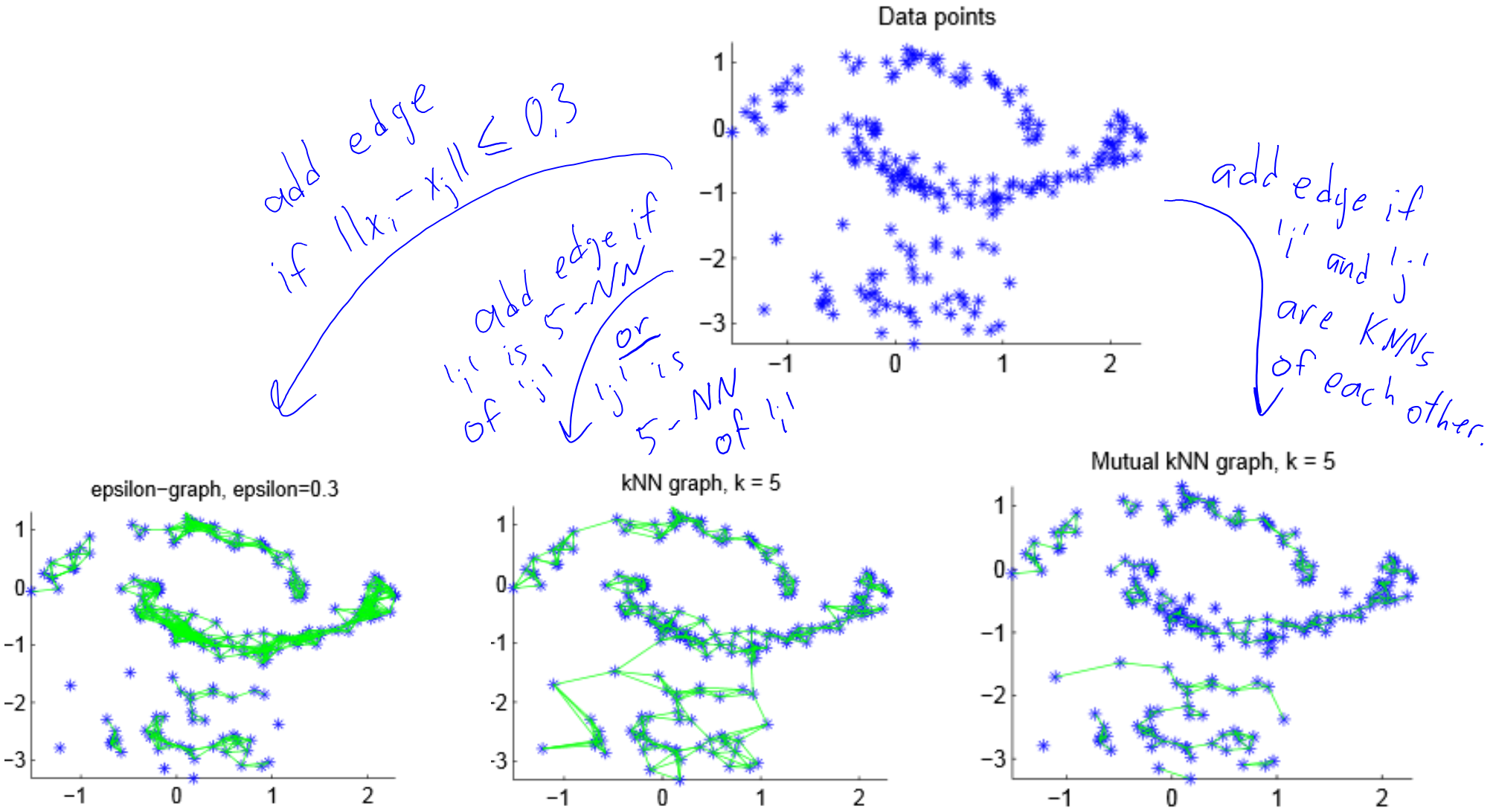
Digression: Constructing Neighbour Graphs

- Sometimes you can define the graph/distance without features:
  - Facebook friend graph.
  - Connect YouTube videos if one video tends to follow another.

- But we can also convert from features x_i to a "neighbour" graph:
  - Approach 1 ("epsilon graph"): connect x_i to all x_j within some threshold ε.
    - Like we did with density-based clustering.
  - Approach 2 ("KNN graph"): connect x_i to x_j if:
    - x_j is a KNN of x_i **OR** x_i is a KNN of x_j.
  - Approach 2 ("mutual KNN graph"): connect x_i to x_j if:
    - x_j is a KNN of x_i **AND** x_i is a KNN of x_j.

http://ai.stanford.edu/~ang/papers/nips01-spectral.pdf

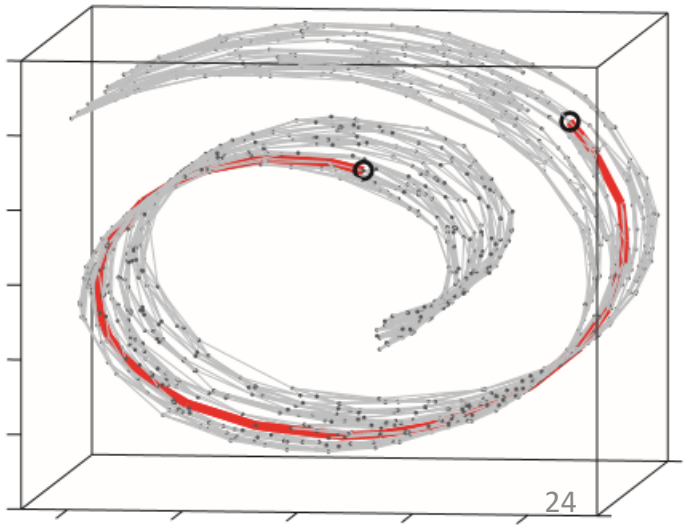
Converting from Features to Graph



http://www.kyb.mpg.de/fileadmin/user_upload/files/publications/attachments/Luxburg07_tutorial_4488%5B0%5D.pdf

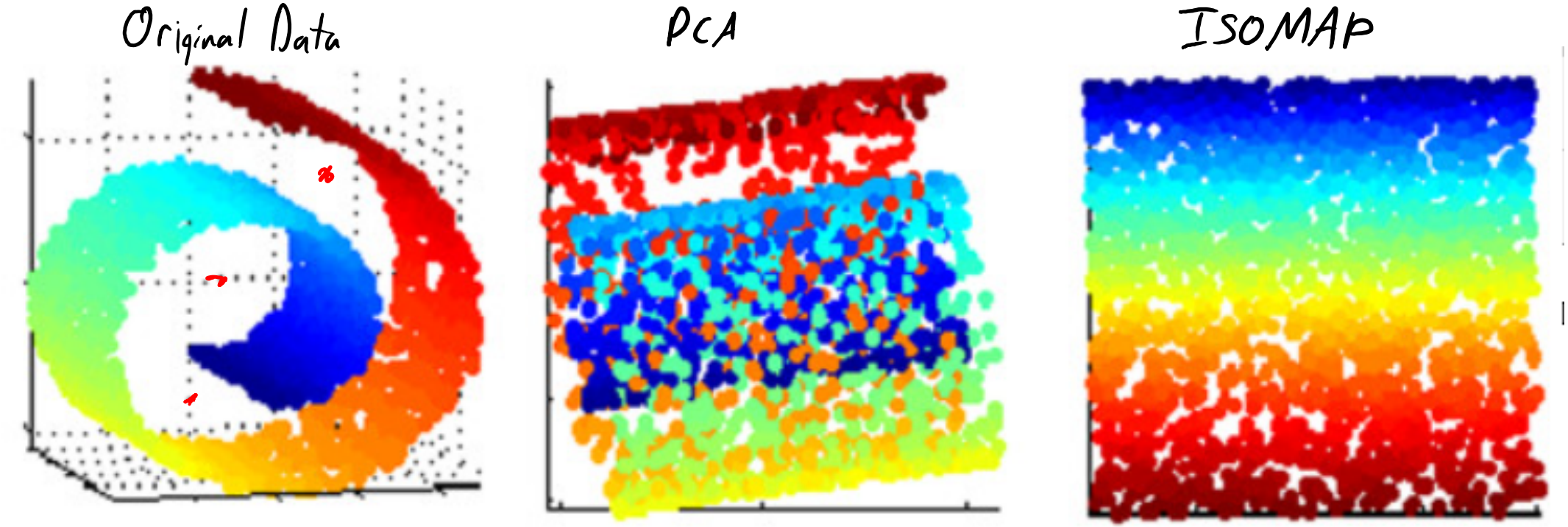
ISOMAP

- ISOMAP is latent-factor model for visualizing data on manifolds:
  1. Find the neighbours of each point.
     - Usually “k-nearest neighbours graph”, or “epsilon graph”.
  2. Compute edge weights:
     - Usually distance between neighbours.
  3. Compute weighted shortest path between all points.
     - Dijkstra or other shortest path algorithm.
  4. Run MDS using these distances.



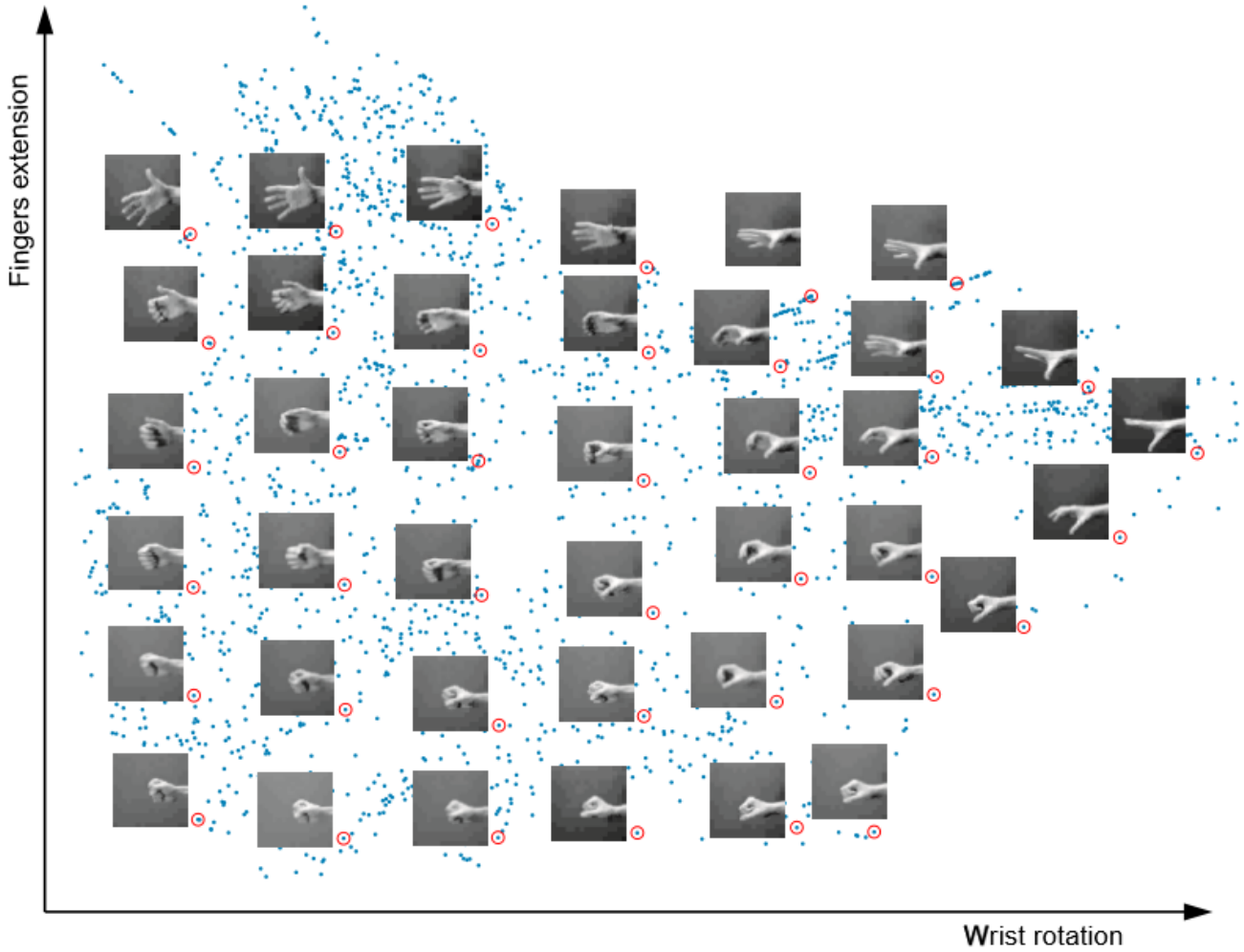
ISOMAP

- ISOMAP can "unwrap" the roll:
  - Shortest paths are approximations to geodesic distances.



- Sensitive to having the right graph:
  - Points off of manifold and gaps in manifold cause problems.

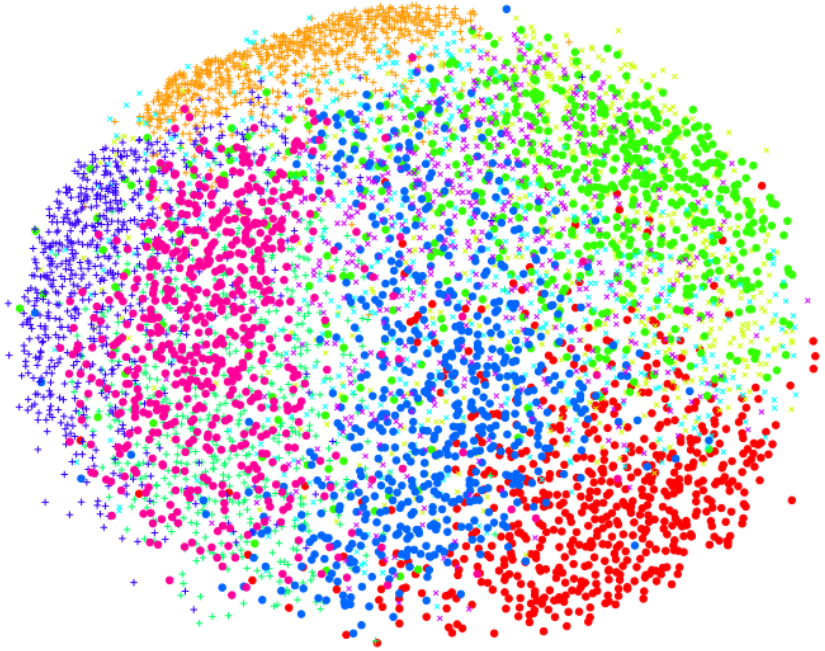
ISOMAP on Hand Images



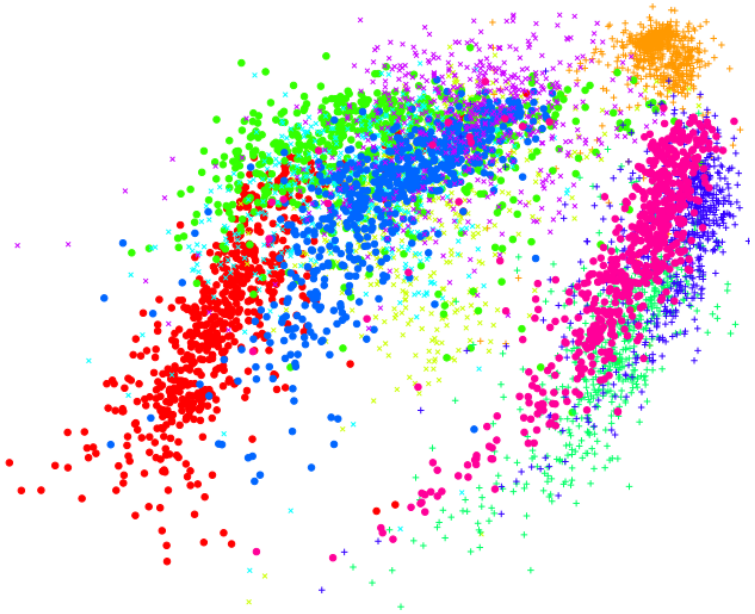
http://wearables.cc.gatech.edu/paper_of_week/isomap.pdf

MNIST digits: Sammon's Map vs. ISOMAP vs. PCA

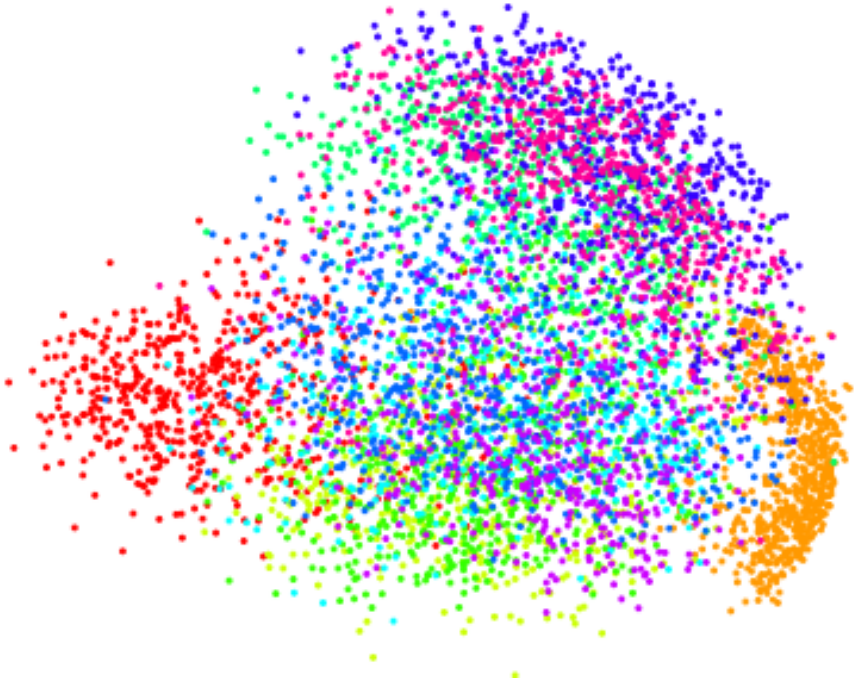
Sammon Map



ISOMAP



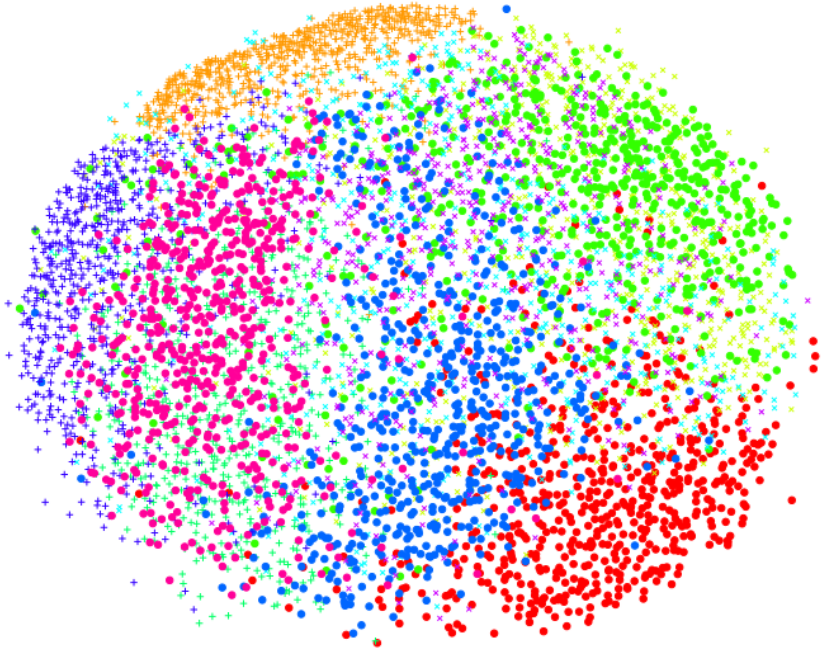
PCA



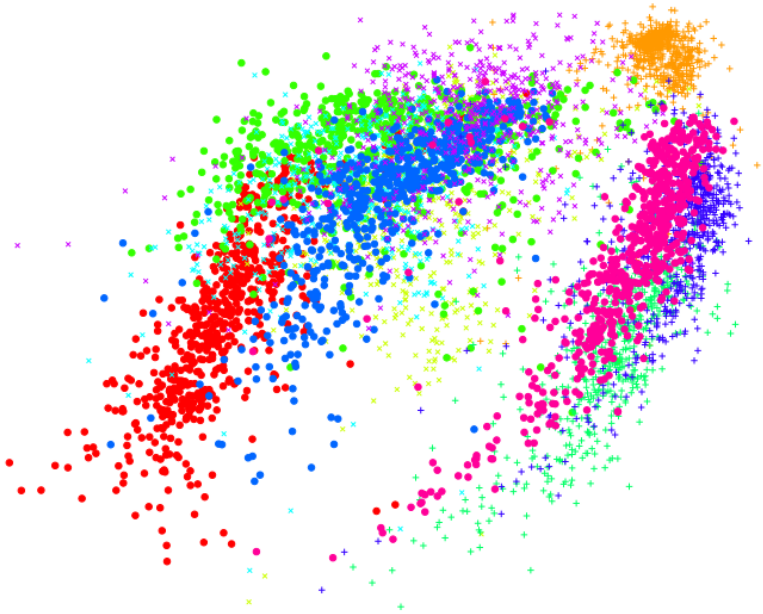
http://lvdmaaten.github.io/publications/papers/JMLR_2008.pdf

MNIST digits: Sammon's Map vs. ISOMAP vs. t-SNE

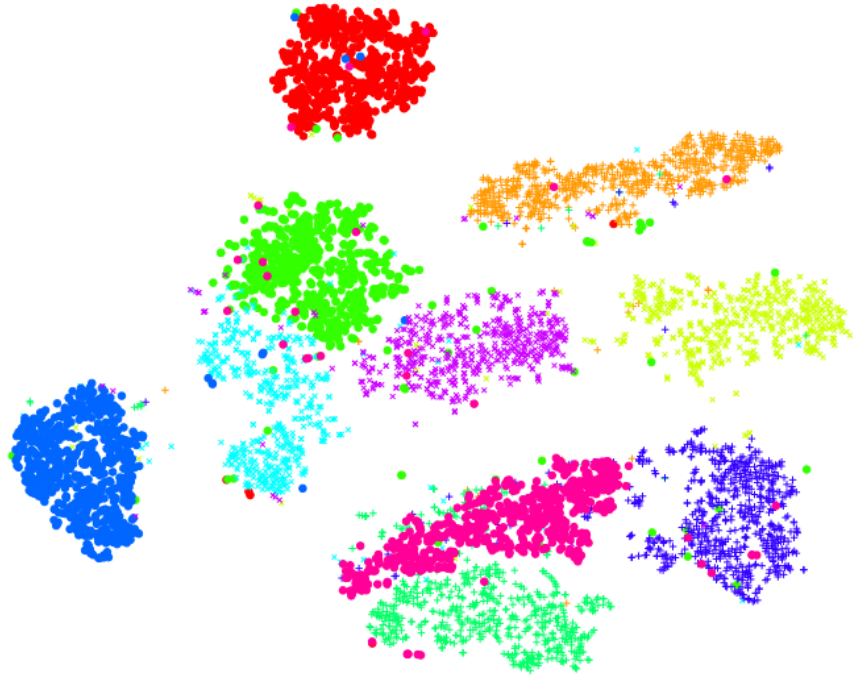
Sammon Map



ISOMAP



t-SNE



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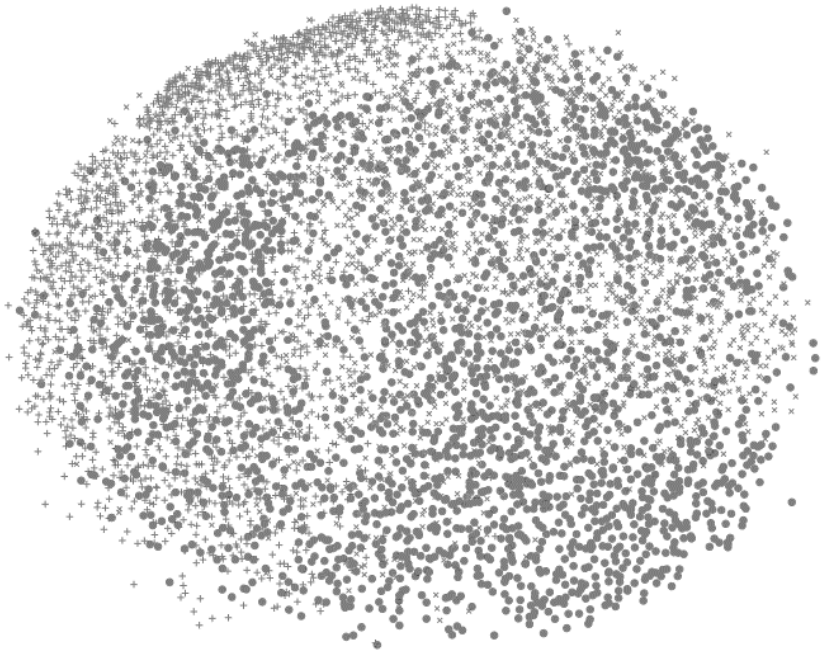
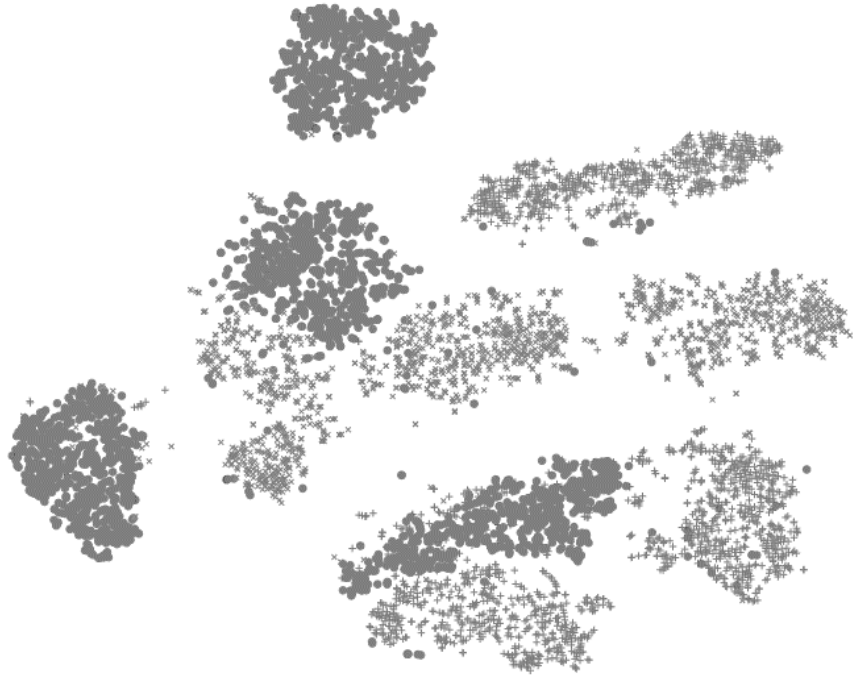
MNIST digits: Sammon's Map vs. ISOMAP vs. t-SNE

Sammon Map



ISOMAP

t-SNE



Remember this is unsupervised, algorithms do not know the labels.

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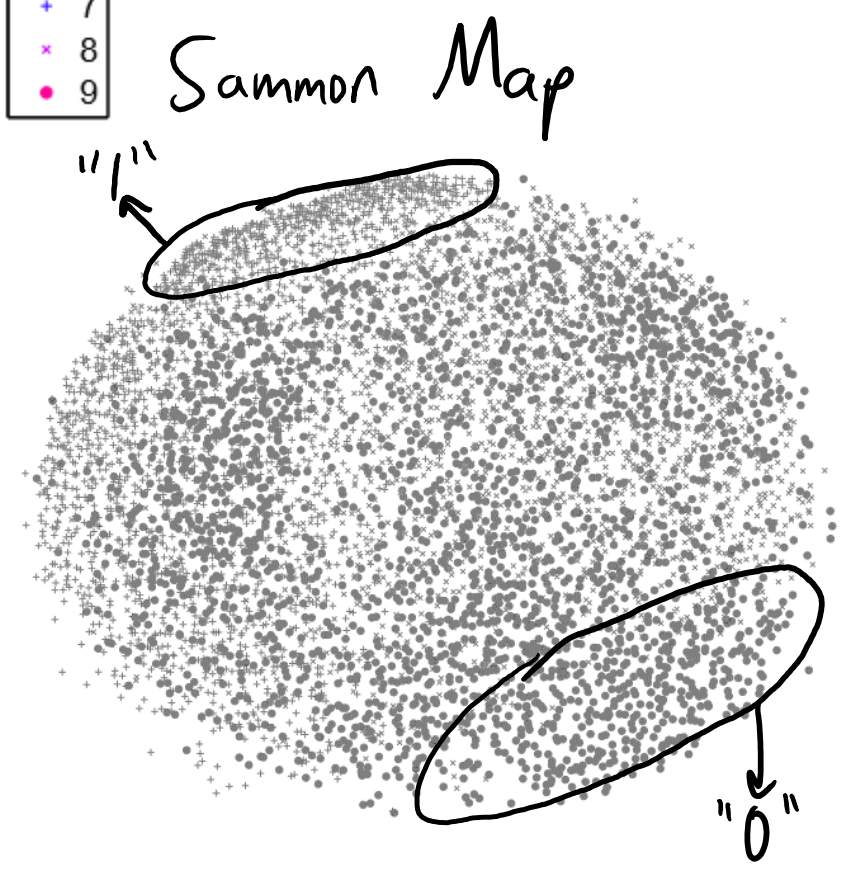

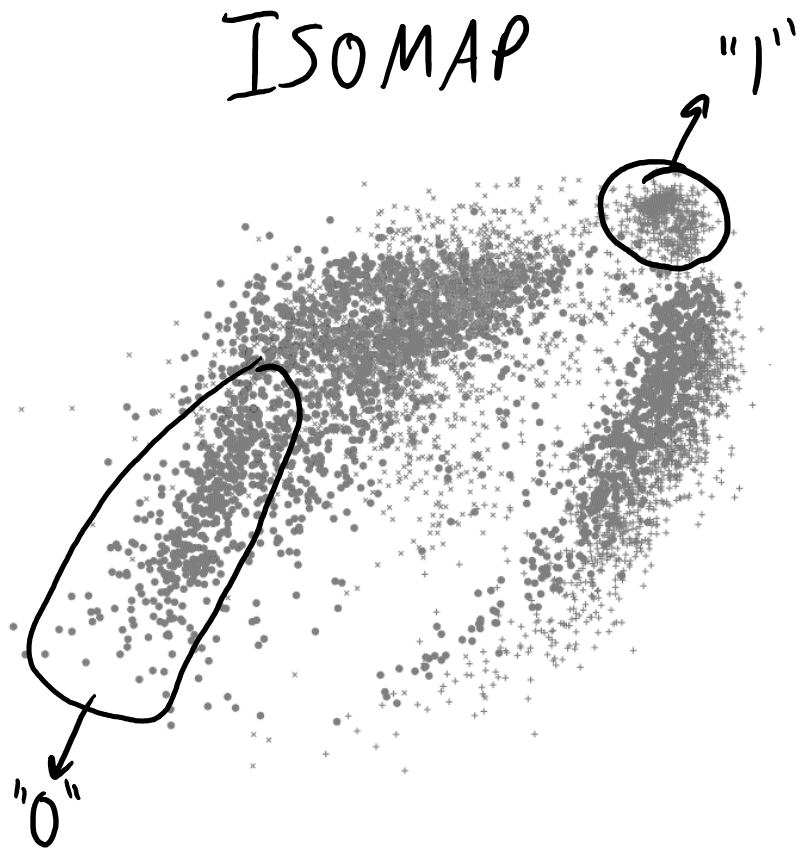

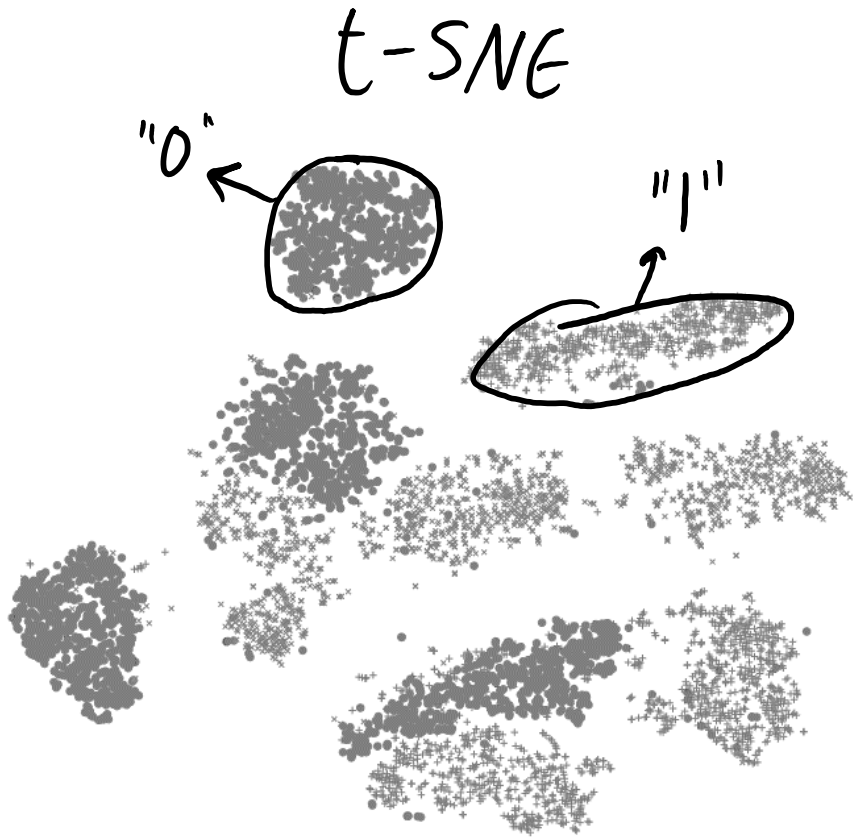

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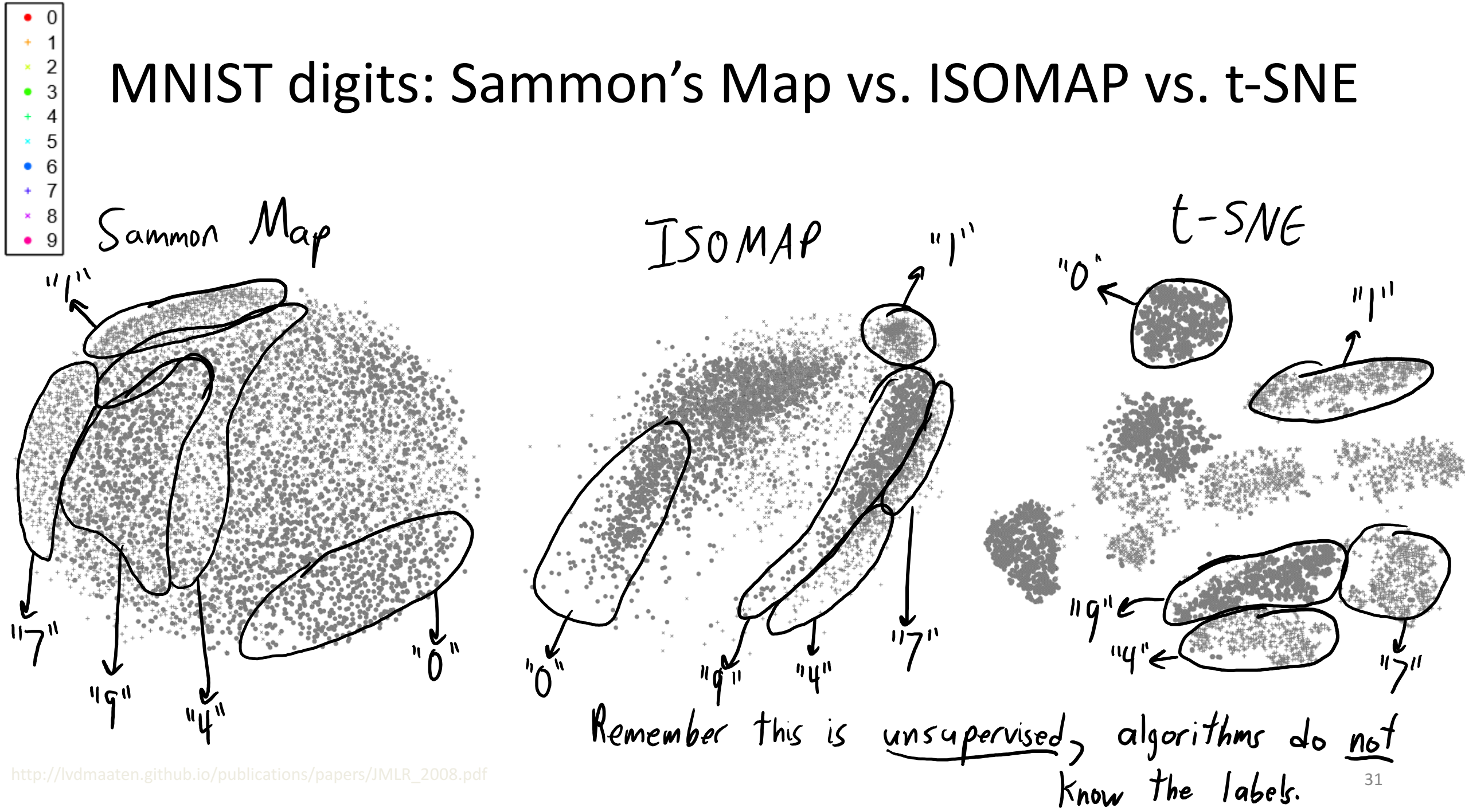

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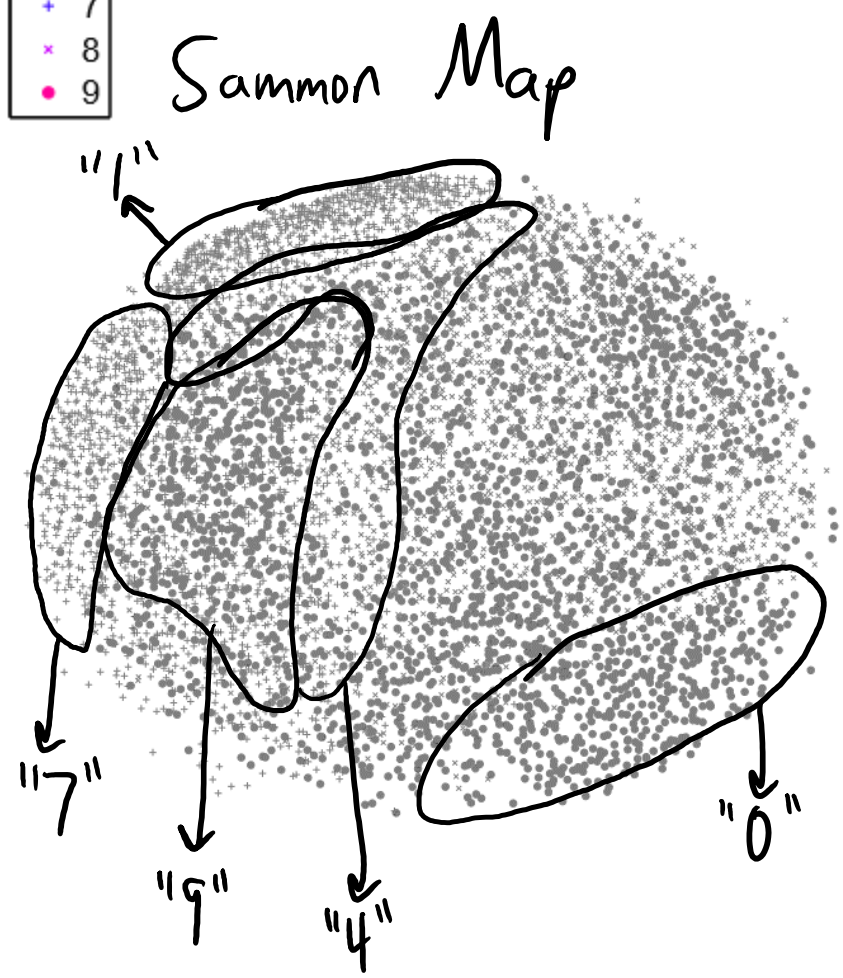
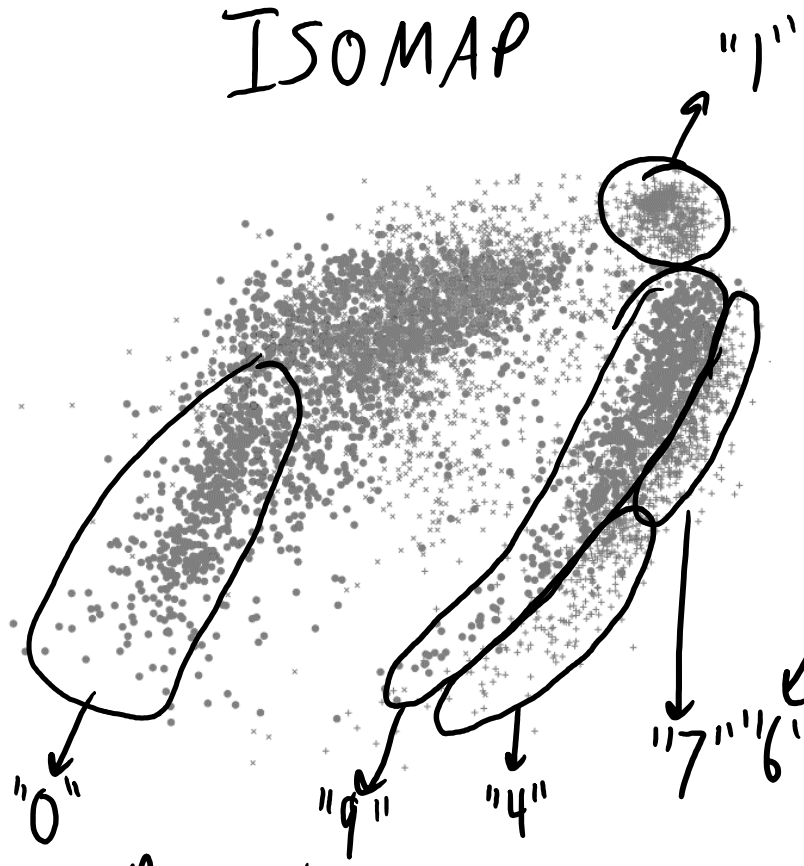
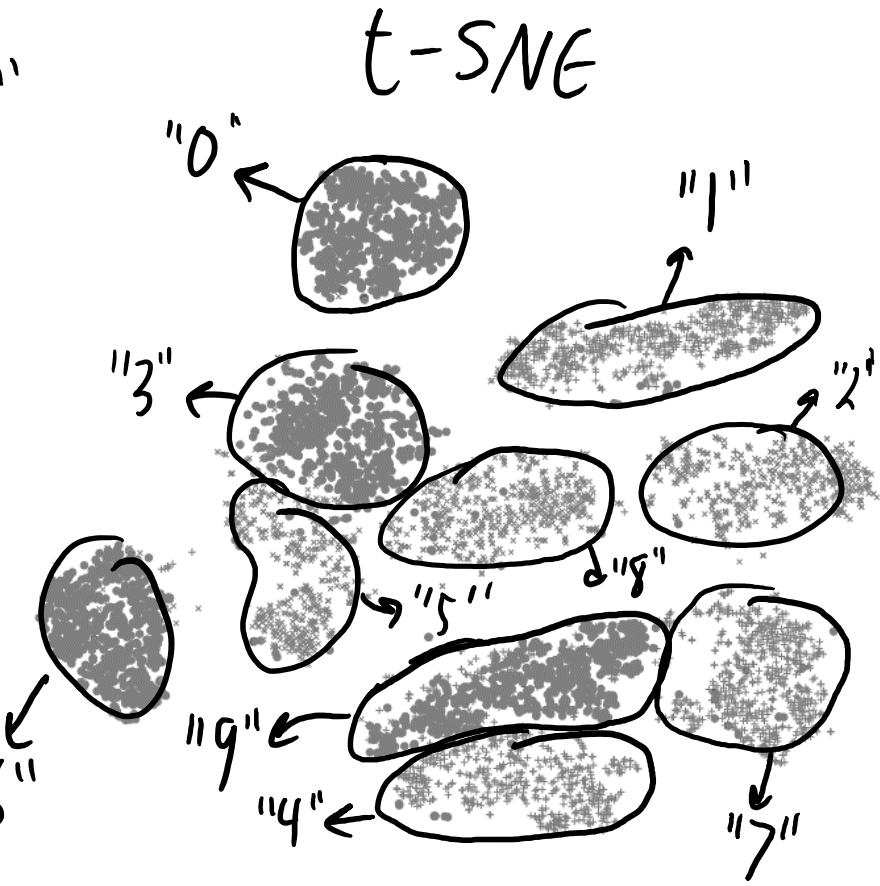
http://lvdmaaten.github.io/publications/papers/JMLR_2008.pdf

MNIST digits: Sammon's Map vs. ISOMAP vs. t-SNE





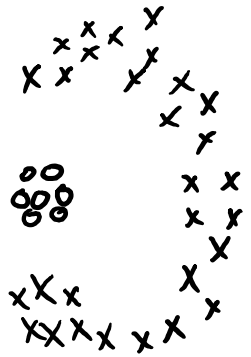


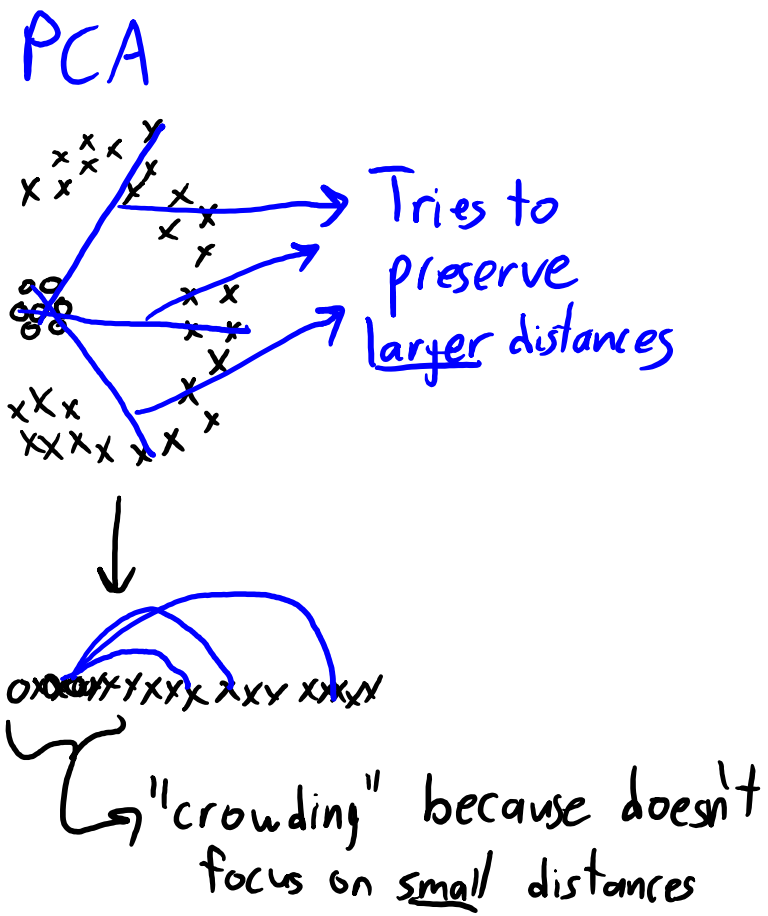
Remember this is unsupervised, algorithms do not know the labels.

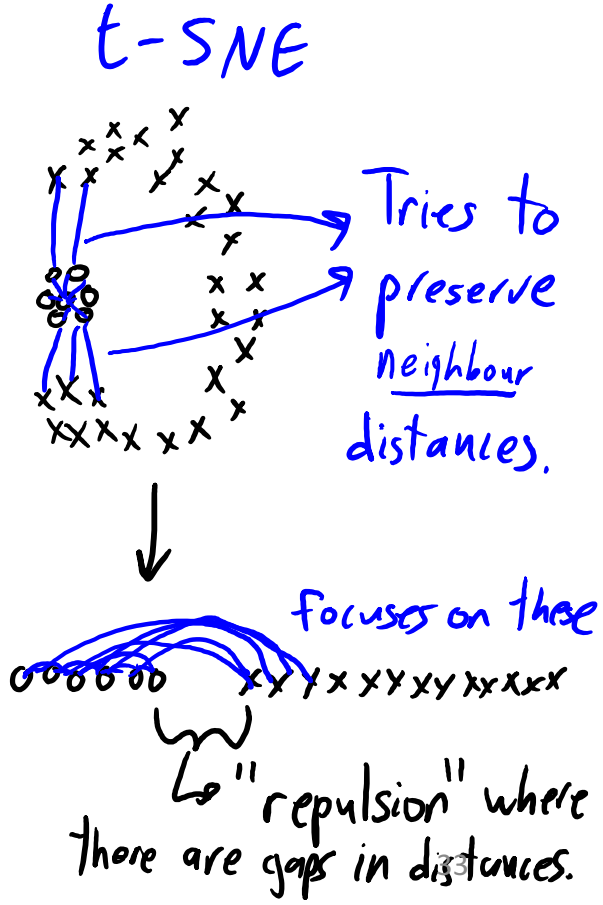
http://lvdmaaten.github.io/publications/papers/JMLR_2008.pdf

t-Distributed Stochastic Neighbour Embedding

- One key idea in t-SNE:
  - Focus on neighbour distances by allowing large variance in large distances.







End of Part 4: Key Concepts

- We discussed linear latent-factor models:

$$f(W,Z) = \sum_{i=1}^{n}\sum_{j=1}^{d}((w^j)^T z_i - x_{ij})^2$$

$$= \sum_{i=1}^{n} ||W^T z_i - x_i||^2$$

$$= ||ZW - X||_F^2$$

- Represent 'X' as linear combination of latent factors 'w_c'.
  - Latent features 'z_i' give a lower-dimensional version of each 'x_i'.
  - When k=1, finds direction that minimizes squared orthogonal distance.
- Applications:
  - Outlier detection, dimensionality reduction, data compression, features for linear models, visualization, factor discovery, filling in missing entries.

End of Part 4: Key Concepts

- We discussed linear latent-factor models:

$$f(W,Z) \approx \sum_{i=1}^{n} \sum_{j=1}^{d} ((w^j)^T z_i - x_{ij})^2$$

- Principal component analysis (PCA):
  - Often uses orthogonal factors and fits them sequentially (via SVD).
- Non-negative matrix factorization:
  - Uses non-negative factors giving sparsity.
  - Can be minimized with projected gradient.
- Many variations are possible:
  - Different regularizers (sparse coding) or loss functions (robust/binary PCA).
  - Missing values (recommender systems) or change of basis (kernel PCA).

End of Part 4: Key Concepts

- We discussed multi-dimensional scaling (MDS):
  - Non-parametric method for high-dimensional data visualization.
  - Tries to match distance/similarity in high-/low-dimensions.
    - "Gradient descent on scatterplot points".
- Main challenge in MDS methods is "crowding" effect:
  - Methods focus on large distances and lose local structure.
- Common solutions:
  - Sammon mapping: use weighted cost function.
  - ISOMAP: approximate geodesic distance using via shortest paths in graph.
  - t-SNE: give up on large distances and focus on neighbour distances.

Summary

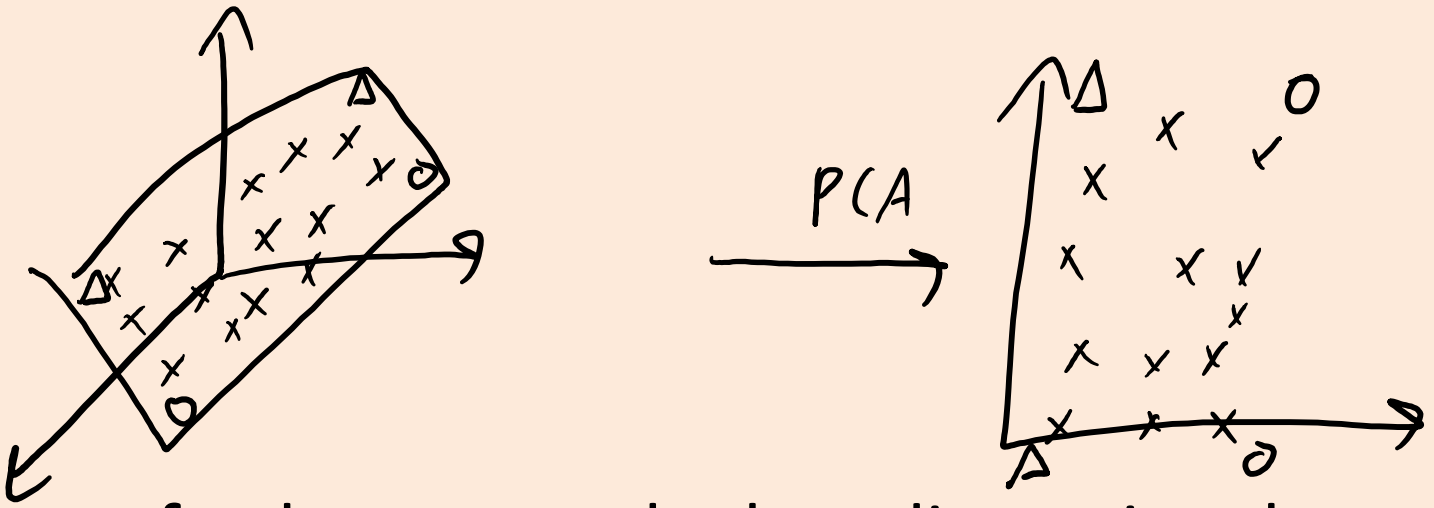
- Multi-dimensional scaling is a non-parametric latent-factor model.
- Different MDS distances/losses/weights usually gives better results.
- Manifold learning focuses on low-dimensional curved structures.
- ISOMAP is most common approach:
  - Approximates geodesic distance by shortest path in weighted graph.
- t-SNE is a promising recent MDS method.

Related method to ISOMAP

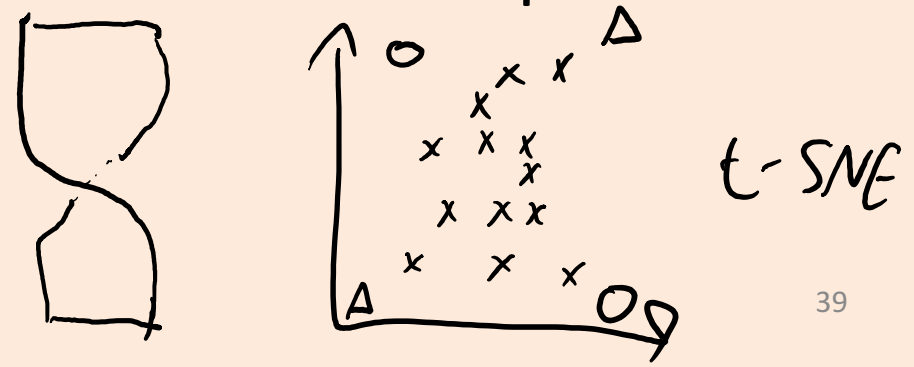
- “local linear embedding”.

Does t-SNE always outperform PCA?

- Consider 3D data living on a 2D hyper-plane:



- PCA can perfectly capture the low-dimensional structure.
- T-SNE can capture the local structure, but can “twist” the plane.
  - It doesn’t try to get long distances correct.



Latent-Factor Representation of Words

- For natural language, we often represent words by an index.
  - E.g., “cat” is word 124056.
- But this may be inefficient:
  - Should “cat” and “kitten” share parameters in some way?
- We want a latent-factor representation of individual words:
  - Closeness in latent space should indicate similarity.
  - Distances could represent meaning?
- Recent alternative to PCA/NMF is word2vec…

Word2Vec

- Two variations on objective in word2vec:
  - Try to predict word from surrounding words (continuous bag of words).
  - Try to predict surrounding words from word (skip-gram).

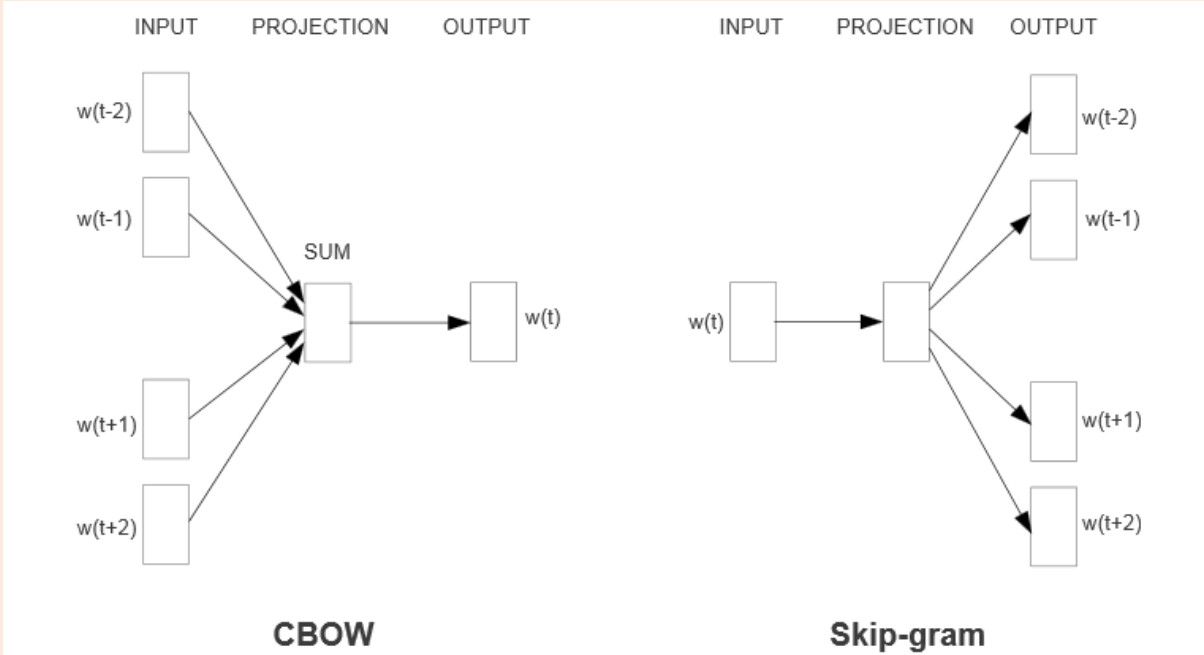


Figure 1: New model architectures. The CBOW architecture predicts the current word based on the context, and the Skip-gram predicts surrounding words given the current word.

https://arxiv.org/pdf/1301.3781.pdf

Word2Vec

- In both cases, each word ‘i’ is represented by a vector z_i.
- In continuous bag of words, we optimize the likelihood:

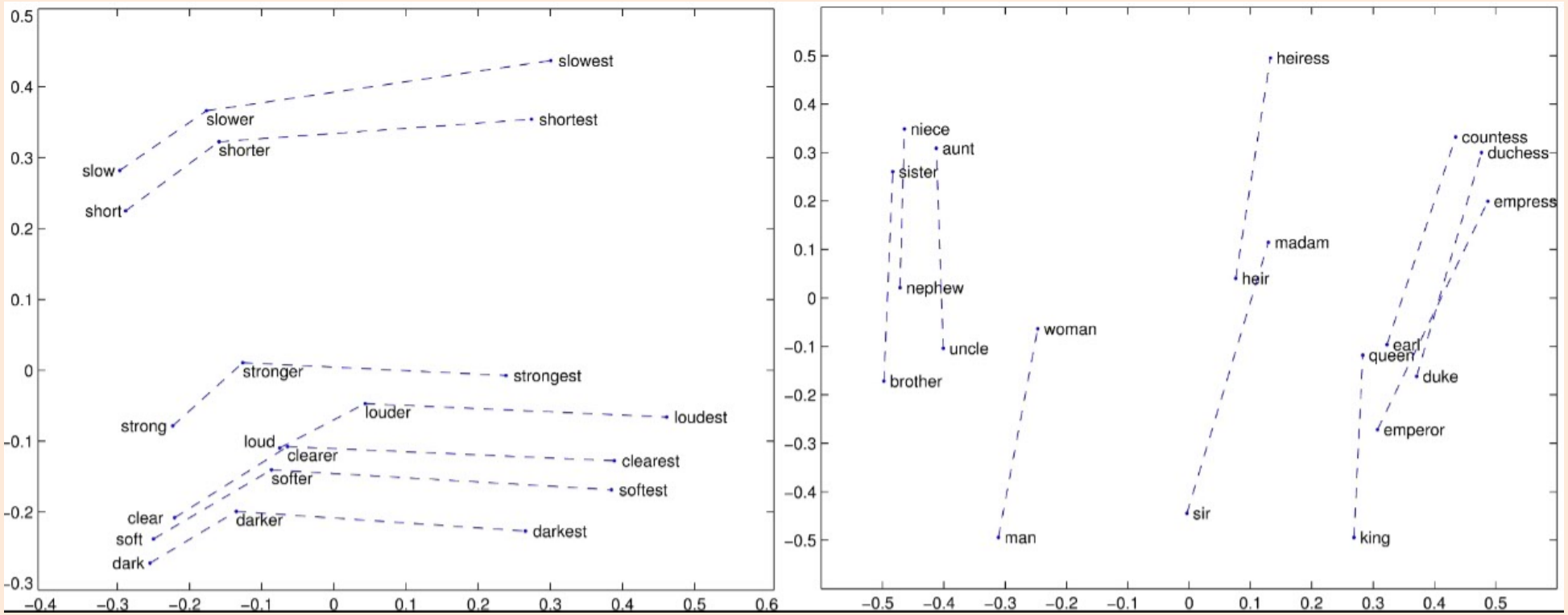
$$p(x_i \mid x_{surround}) = \prod_{j \in surround} p(x_i \mid x_j) \qquad \text{(independence assumption)}$$

$$= \prod_{j \in surround} \frac{\exp(z_i^T z_j)}{\sum_{c=1}^{K} \exp(z_c^T z_j)} \qquad \text{(softmax over all words)}$$

- Denominator sums over all words.
- For skip-gram it will be over all possible surrounding words.
  - Common trick to speed things up: samples terms in denominator.
    - “Negative sampling”.

Word2Vec Example

- MDS visualization of a set of related words:



- Distances between vectors might represent semantics.

http://ruder.io/secret-word2vec/

Word2Vec

- Subtracting word vectors to find related vectors.

Table 8: *Examples of the word pair relationships, using the best word vectors from Table 4 (Skip-gram model trained on 783M words with 300 dimensionality).*

Relationship	Example 1	Example 2	Example 3
France - Paris	Italy: Rome	Japan: Tokyo	Florida: Tallahassee
big - bigger	small: larger	cold: colder	quick: quicker
Miami - Florida	Baltimore: Maryland	Dallas: Texas	Kona: Hawaii
Einstein - scientist	Messi: midfielder	Mozart: violinist	Picasso: painter
Sarkozy - France	Berlusconi: Italy	Merkel: Germany	Koizumi: Japan
copper - Cu	zinc: Zn	gold: Au	uranium: plutonium
Berlusconi - Silvio	Sarkozy: Nicolas	Putin: Medvedev	Obama: Barack
Microsoft - Windows	Google: Android	IBM: Linux	Apple: iPhone
Microsoft - Ballmer	Google: Yahoo	IBM: McNealy	Apple: Jobs
Japan - sushi	Germany: bratwurst	France: tapas	USA: pizza

Table 8 shows words that follow various relationships. We follow the approach described above: the relationship is defined by subtracting two word vectors, and the result is added to another word. Thus for example, *Paris* - *France* + *Italy* = *Rome*. As it can be seen, accuracy is quite good, although

https://arxiv.org/pdf/1301.3781.pdf

Graph Drawing

- A closely-related topic to MDS is graph drawing:
  - Given a graph, how should we display it?
  - Lots of interesting methods: https://en.wikipedia.org/wiki/Graph_drawing

Bonus Slide: Multivariate Chain Rule

- Recall the univariate chain rule:

$$\frac{d}{dw}\left[f(g(w))\right] = f'(g(w))\,g'(w)$$

- The multivariate chain rule:

$$\underbrace{\nabla\left[f(g(w))\right]}_{d\times 1} = \underbrace{f'(g(w))}_{1\times 1}\underbrace{\nabla g(w)}_{d\times 1}$$

- Example:

$$\nabla\left[\tfrac{1}{2}(w^T x_i - y_i)^2\right]$$

$$= \nabla\left[f(g(w))\right]$$

with $g(w) = w^T x_i - y_i$ → $\nabla g(w) = x_i$

and $f(r_i) = \frac{1}{2} r_i^2$ → $f'(r_i) = r_i$

$$\nabla\left[f(g(w))\right] = r_i\, x_i$$

$$= (w^T x_i - y_i) x_i$$

Bonus Slide: Multivariate Chain Rule for MDS

- General MDS formulation:

$$\underset{Z \in \mathbb{R}^{n \times k}}{\operatorname{argmin}} \sum_{i=1}^{n} \sum_{j=i+1}^{n} g\left(d_1(x_i, x_j), d_2(z_i, z_j)\right)$$

- Using multivariate chain rule we have:

$$\nabla_{z_i} g\left(d_1(x_i, x_j), d_2(z_i, z_j)\right) = g'\left(d_1(x_i, x_j), d_2(z_i, z_j)\right) \nabla_{z_i} d_2(z_i, z_j)$$

- Example: If $d_1(x_i, x_j) = \|x_i - x_j\|$ and $d_2(z_i, z_j) = \|z_i - z_j\|$ and $g(d_1, d_2) = \frac{1}{2}(d_1 - d_2)^2$

$$\nabla_{z_i} g\left(d_1(x_i, x_j), d_2(z_i, z_j)\right) = -\underbrace{\left(d_1(x_i, x_j) - d_2(z_i, z_j)\right)}_{g'(d_1, d_2)} \underbrace{\left[-\frac{(z_i - z_j)}{2\|z_i - z_j\|}\right]}_{\nabla_{z_i} d_2(z_i, z_j)}$$

↳ Assuming $z_i \neq z_j$

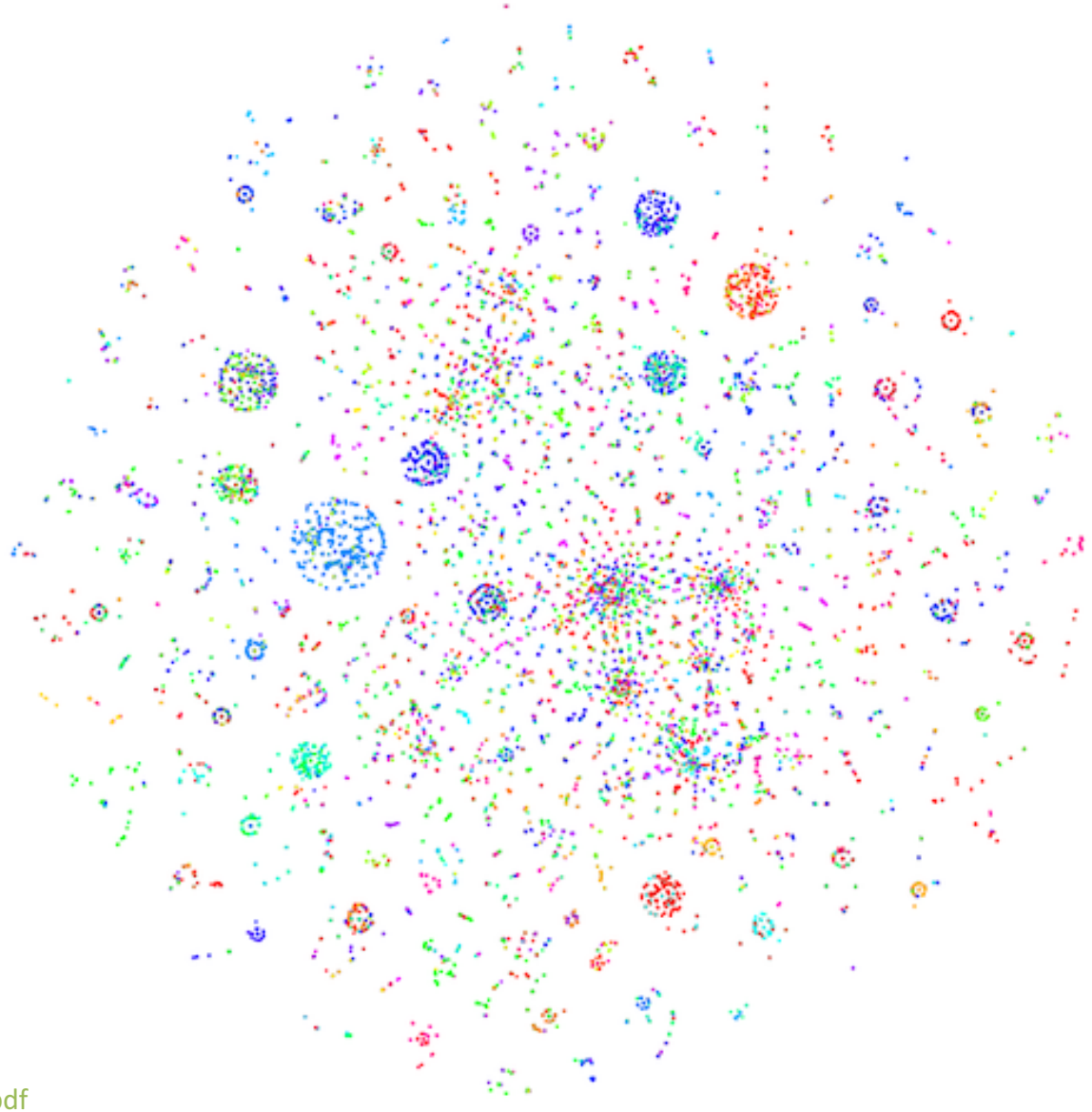
($g'(d_1, d_2)$: move distances closer)

($\nabla_{z_i} d_2(z_i, z_j)$: how distance changes in z-space)

t-Distributed Stochastic Neighbour Embedding

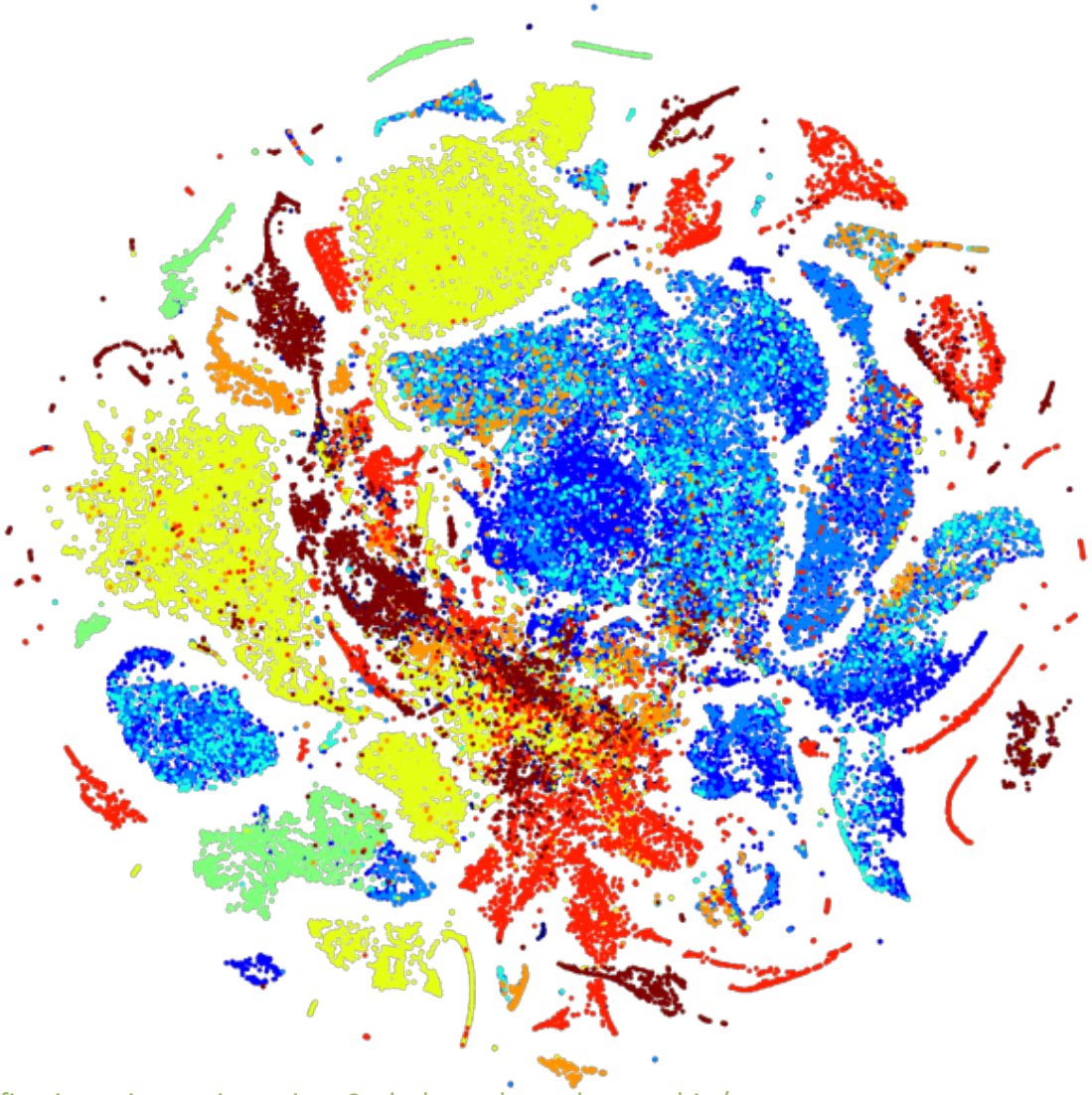
- t-SNE is a special case of MDS (specific d_1, d_2, and d_3 choices):
  - d_1: for each x_i, compute probability that each x_j is a 'neighbour'.
    - Computation is similar to k-means++, but most weight to close points (Gaussian).
    - Doesn't require explicit graph.
  - d_2: for each z_i, compute probability that each z_j is a 'neighbour'.
    - Similar to above, but uses student's t (grows really slowly with distance).
    - Avoids 'crowding', because you have a huge range that large distances can fill.
  - d_3: Compare x_i and z_i using an entropy-like measure:
    - How much 'randomness' is in probabilities of x_i if you know the z_i (and vice versa)?
- Interactive demo: https://distill.pub/2016/misread-tsne

t-SNE on Wikipedia Articles



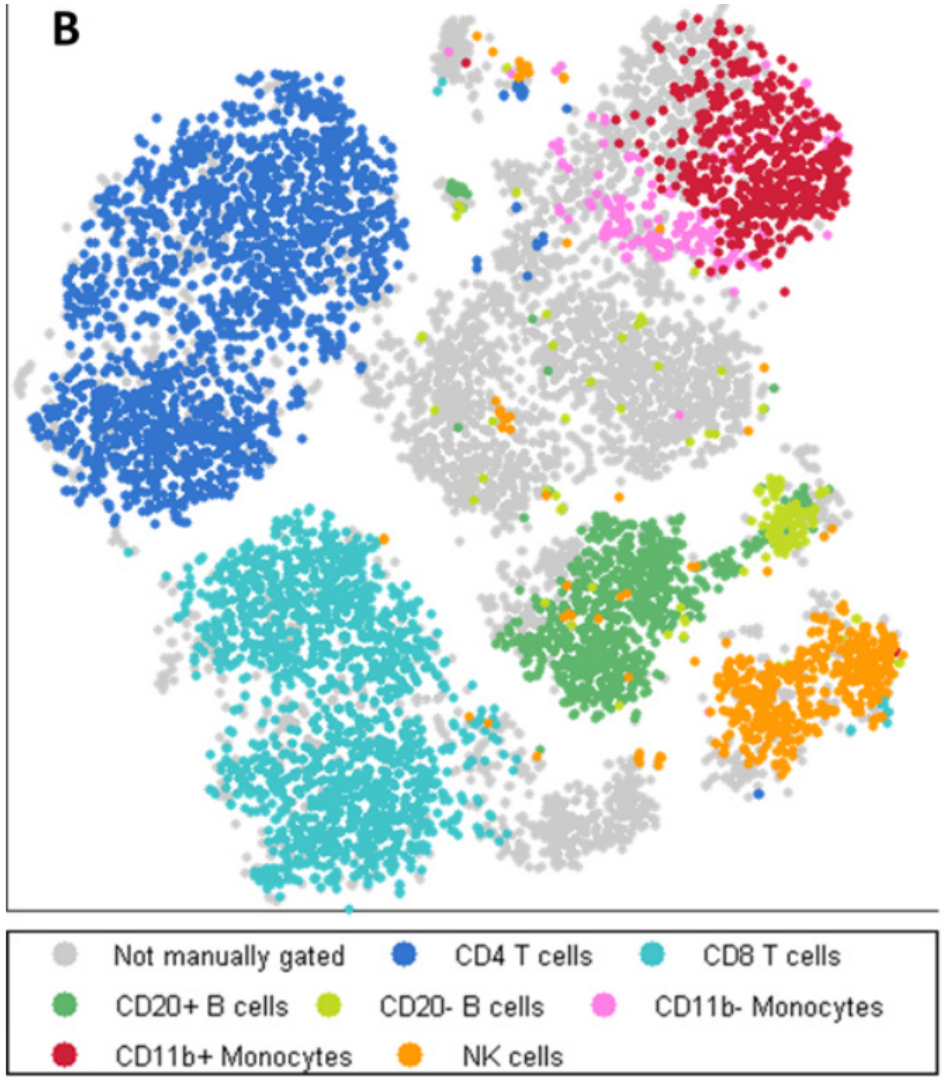
http://jasneet.me/assets/files/wiki_tsne_report.pdf

t-SNE on Product Features



http://blog.kaggle.com/2015/06/09/otto-product-classification-winners-interview-2nd-place-alexander-guschin/

t-SNE on Leukemia Heterogeneity



http://www.ncbi.nlm.nih.gov/pmc/articles/PMC4076922/