

Advanced Data Visualization

CS 6965

Spring 2018

Prof. Bei Wang Phillips

University of Utah

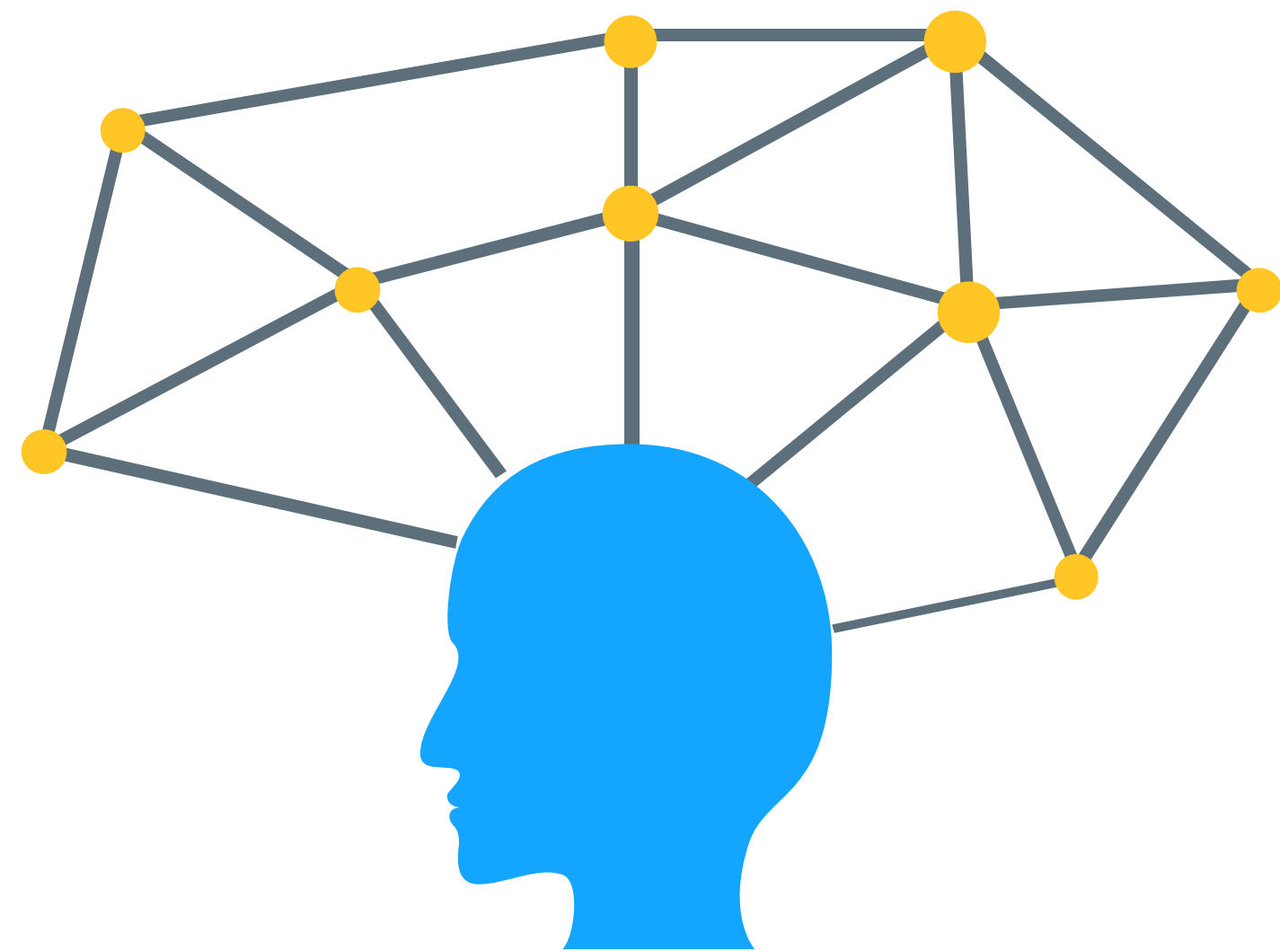


Lecture 02

Dim Reduction & Vis



HD



Visualization
is the secret weapon for
Machine learning

Roles of ML in HD data visualization

From **Black Box** to **Glass Box**:

- ML as part of data transformation in the visualization pipeline
- Visualization increase the **interpretability** of the algorithmic results (visualizing algorithm **output**)
- Visualization increases the **interpretability** of ML algorithms (visualizing algorithmic **processes**)
- (Interactive) visualization becomes part of the ML algorithm

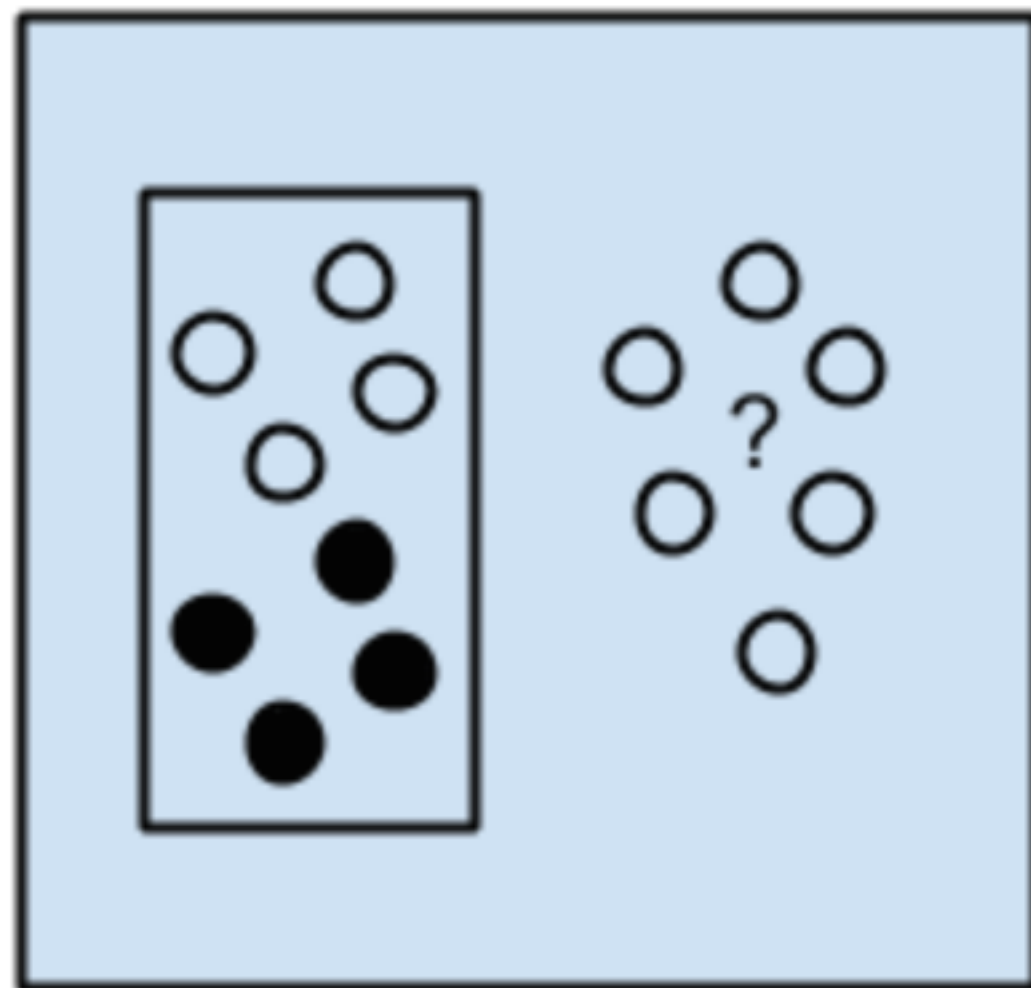
ML algorithms in a nutshell

Not a full-blown ML class, but

How to best incorporate vis into ML algorithms?

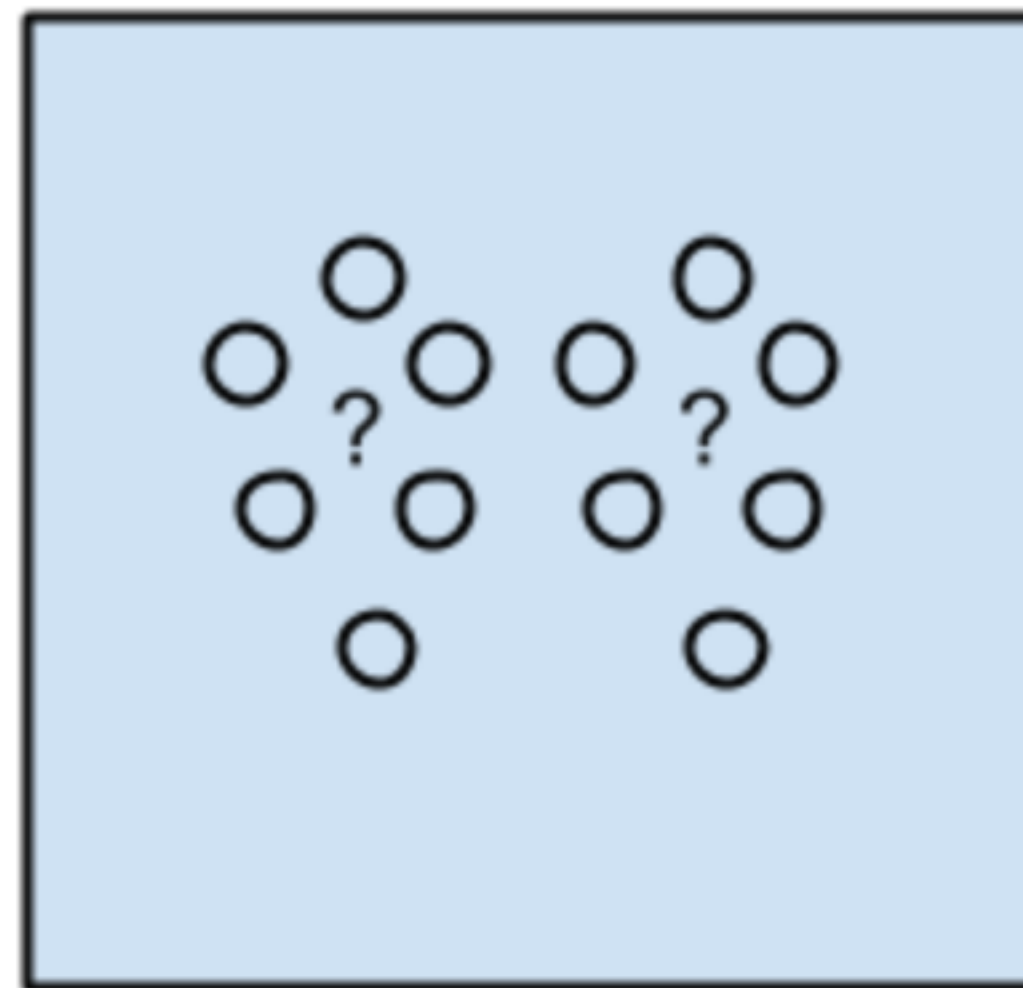
- A simple approach is to treat the ML algorithm as a black box, and build vis surrounding its input/output
- Not knowing the interworking of the algorithm (e.g. a glass box) may lead to misinterpretation of the algorithm output
- We need to have a good understanding of the **core** of some ML algorithms
- We will review **some** ML algorithms with a focus on their **inner-workings** so as to think about how visualization can be incorporated
- You are encouraged to read about ML in general (see recommended reading, and talk to the instructor)
- Keep in mind, our focus is **ML+Vis**

ML algorithm by learning styles



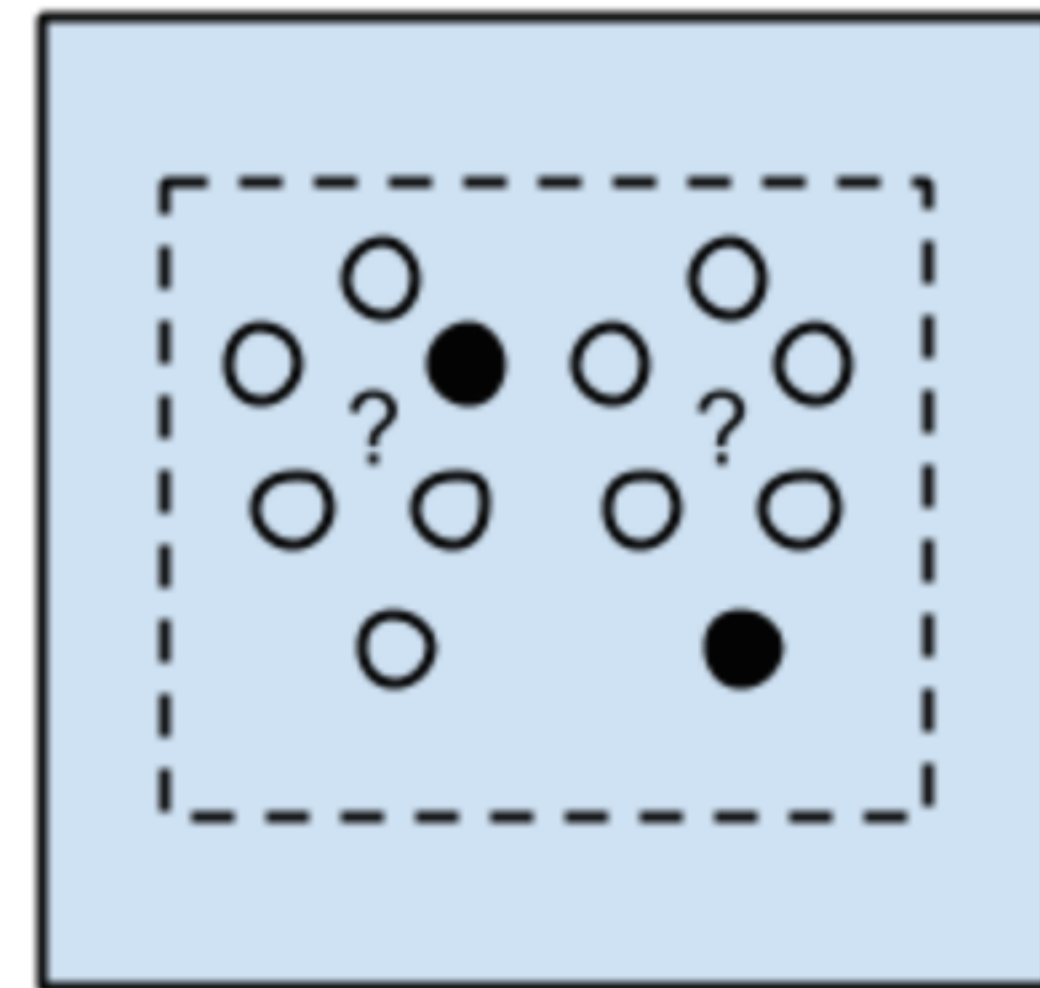
Supervised Learning

*Problems: Classification
Regression*



Unsupervised Learning

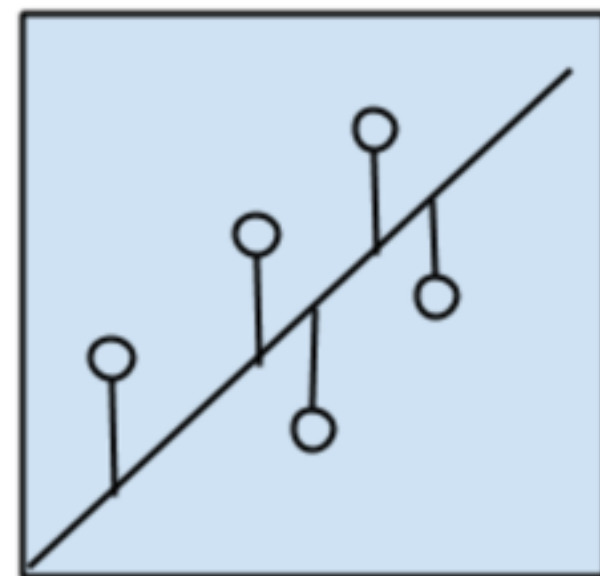
*Problems: Clustering
Dimensionality Reduction*



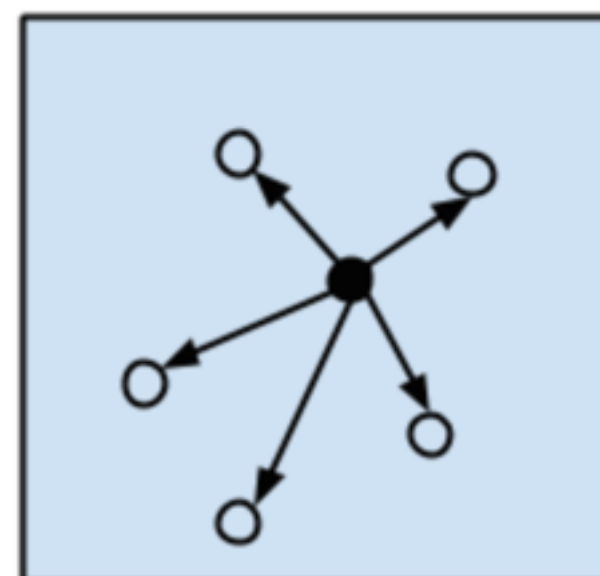
Semi-supervised Learning

*Problems: Classification
Regression*

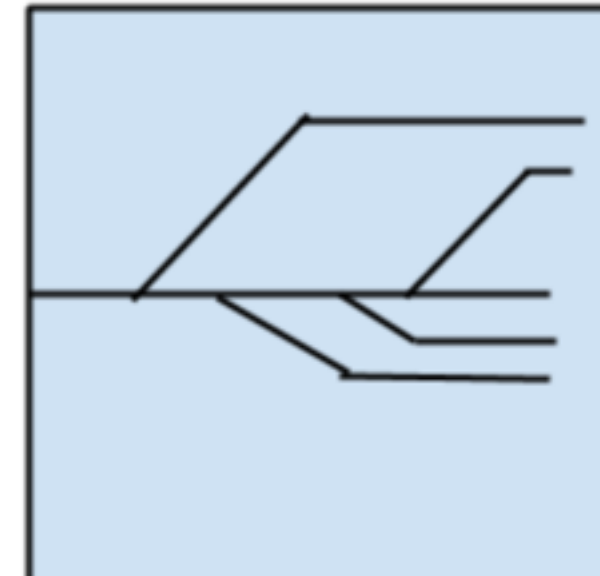
ML algorithm by similarity (how they work)



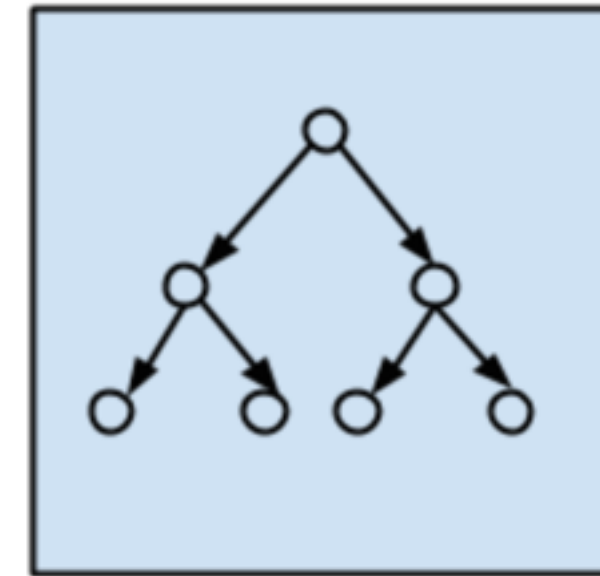
Regression Algorithms



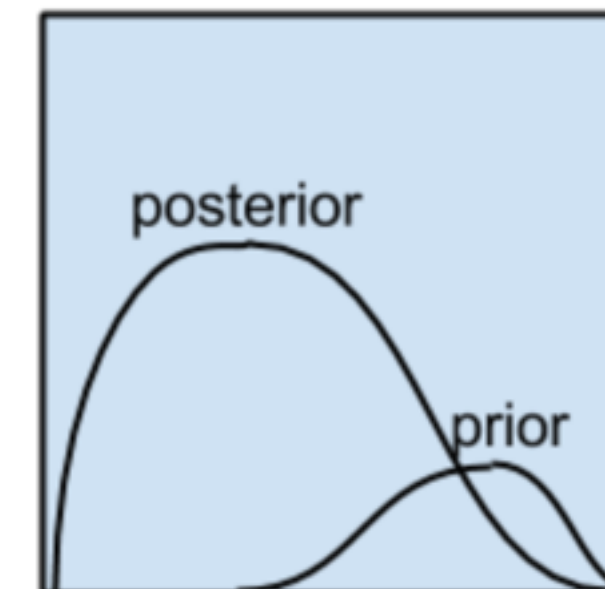
Instance-based Algorithms



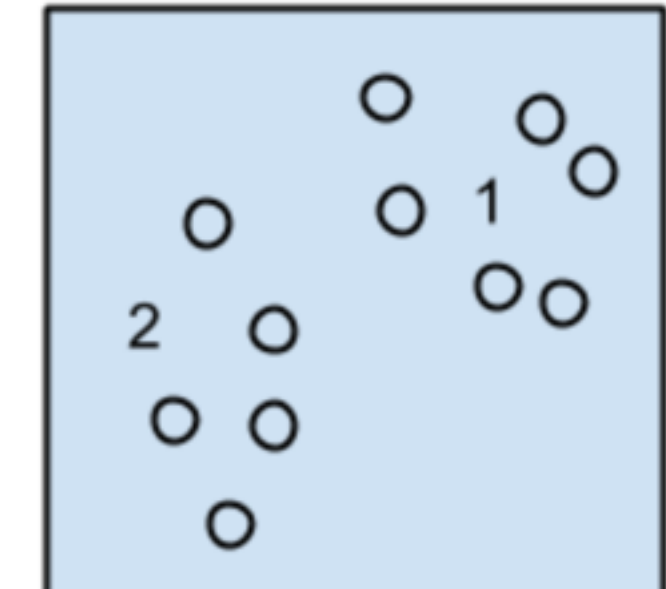
Regularization Algorithms



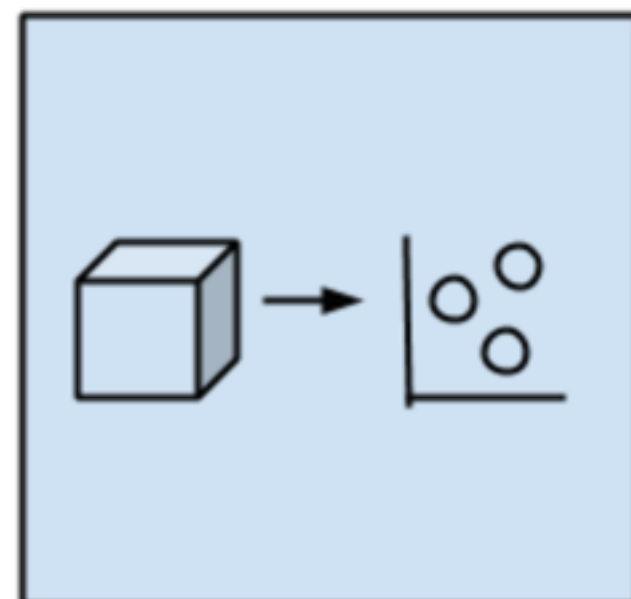
Decision Tree Algorithms



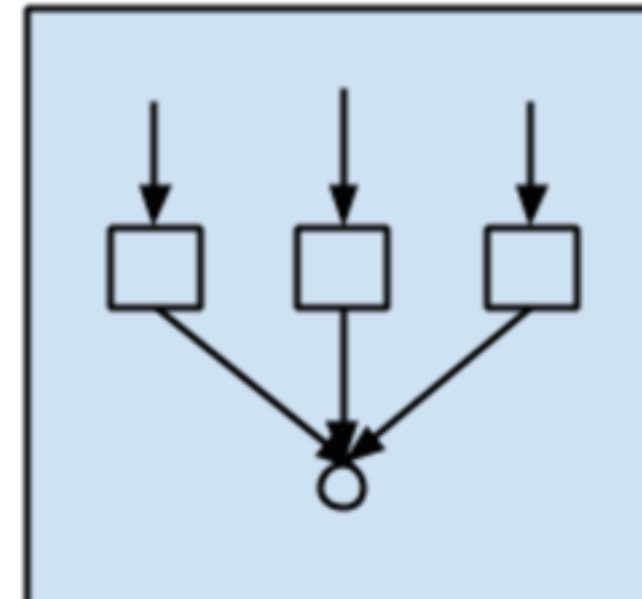
Bayesian Algorithms



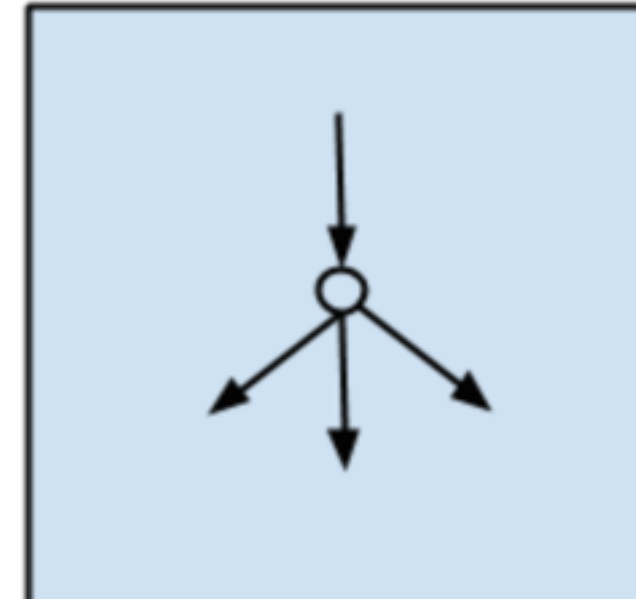
Clustering Algorithms



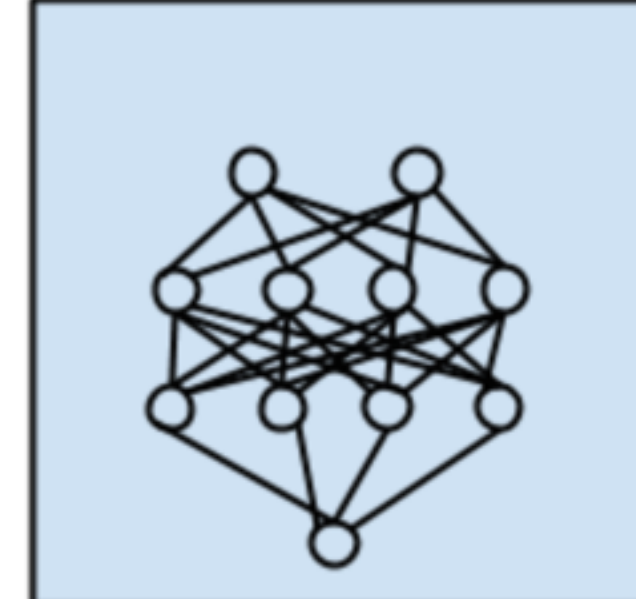
Dimensional Reduction Algorithms



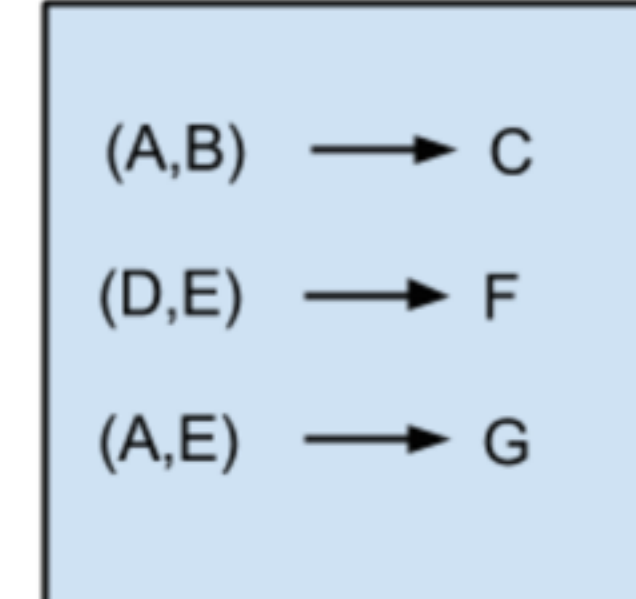
Ensemble Algorithms



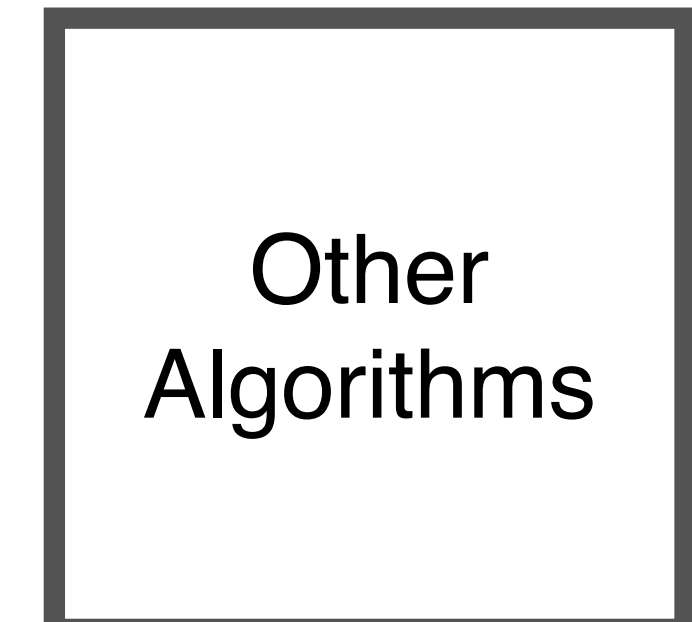
Artificial Neural Network Algorithms



Deep Learning Algorithms



Association Rule Learning Algorithms



Advances in HD Vis

Visualizing High-Dimensional Data: Advances in the Past Decade

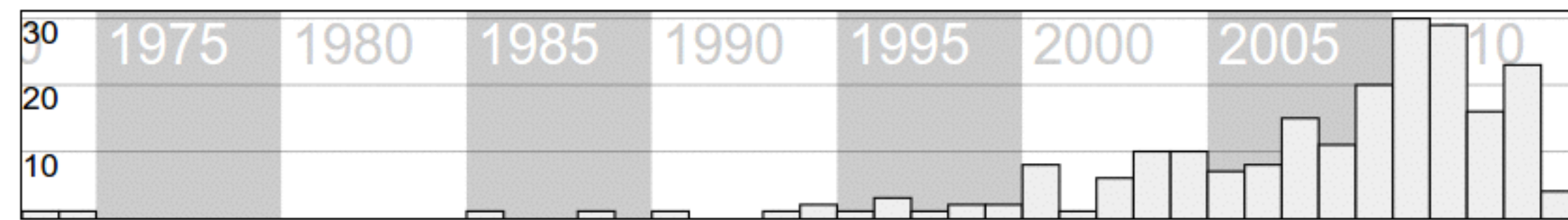
Digital library for publication **Visualizing High-Dimensional Data: Advances in the Past Decade**



Selectors

search ...

Timeline



Tags

pipeline stage: ?₆ data transformation₁₃₇ view transformation₁₇ visual mapping₆₂
user involvement: ?₇ computation centric₆₁ interactive exploration₁₄₄
model manipulation₆
paper type: ?₄₀ application₇ survey₁₁ system₁₁ technical₁₄₇ theory₃
data type: ?₈₆ high-dimensional function₇ high-dimensional point cloud₁
high-dimensional points₁₀₀ nominal data₁₄ spatial data₆ time series₄
analysis method: ?₅₅ clustering₈₃ data abstraction₅ data subset₁ dimension relationship₉
dimension similarity₄ dimensionality reduction₂₅ distance metric₆ feature extraction₂
histogram₂ optimization₁ precision measure₅ projection₁₂ quality measure₁ regression₈
regression?₁ scagnostics₁ segmentation₁ statistic₂ subspace₁₄ topological analysis₉
visual method: ?₂₁ animation₆ bar charts₇ focus+context₆ glyphs₁₀ heat map₁
hierarchy₁₃ isosurface₄ magic lens₄ node-link₃ novel visual encoding₃₁
parallel coordinates₉₆ pixel-based₅ progressive update₃ radviz₄
rendering enhancement₄ scatterplot₅₉ star coordinates₂ surfaces₇ treemap₃
volume visualization₅
other: ₅ clustering₁ clutter reduction₁₅ comparison₁ high-dimensional points₁ data transformation₁
filtering₂ histogram₁ information₁ machine learning₅ matching₁ parameter exploration₈
perception₄ query₈ ranking₁₇ reordering₄ segmentation₁ sensitivity analysis₄ uncertainty₃
user study₁ view optimization₁ visual data mining₁

216 publications

Bug Report Welcome! [LiuMaljovecWang2017] <http://www.sci.utah.edu/~shusenl/highDimSurvey/website/>



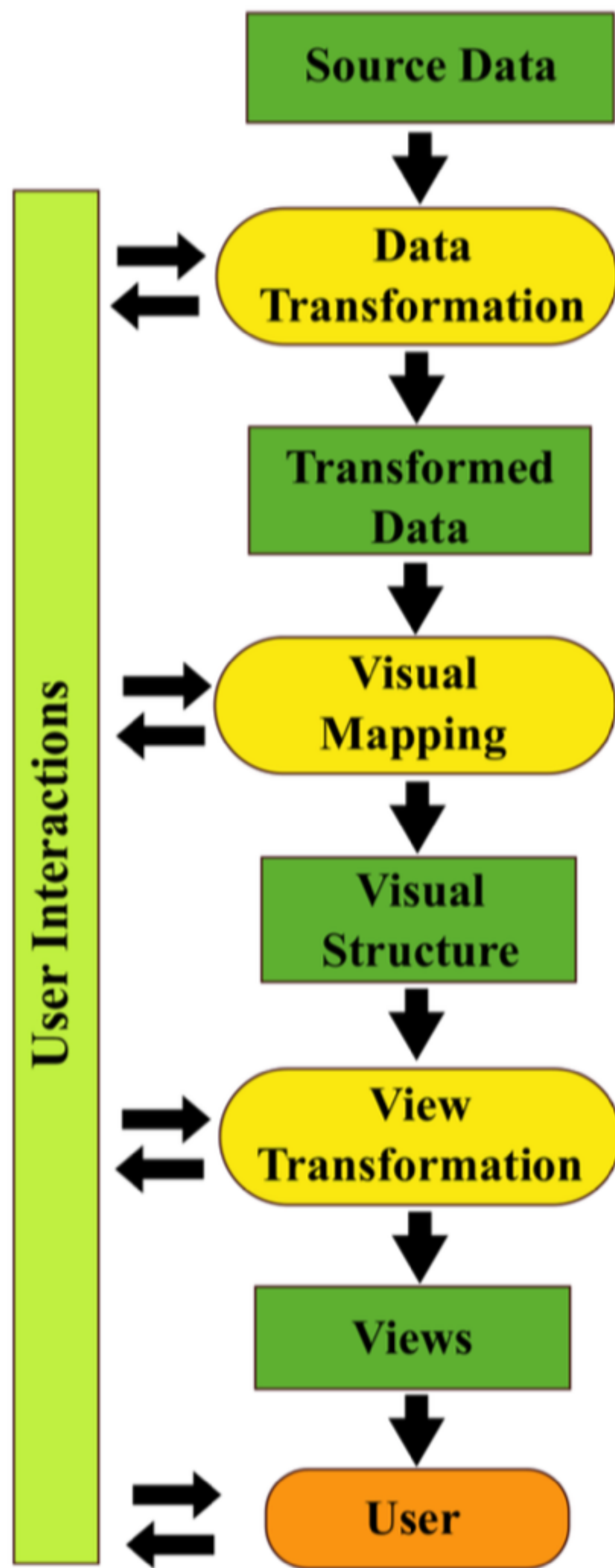
4. AnkerstBerchtoldKeim1998 [inproceedings] (1998) | PDF | DOI | Google Scholar | Google
Similarity clustering of dimensions for an enhanced visualization of multidimensional data
Ankerst, Mihael Berchtold, Stefan Keim, Daniel A
Abstract: The order and arrangement of dimensions (variates) is crucial for the effectiveness of a large number of visualization techniques such as parallel coordinates, scatterplots, recursive pattern, and many others. We describe a systematic approach to arrange the dimensions according to their similari... ▶
pipeline stage:visual mapping user involvement:computation centric paper type:technical
data type:? analysis method:dimension similarity visual method:parallel coordinates
view optimization +
select similar BibTeX



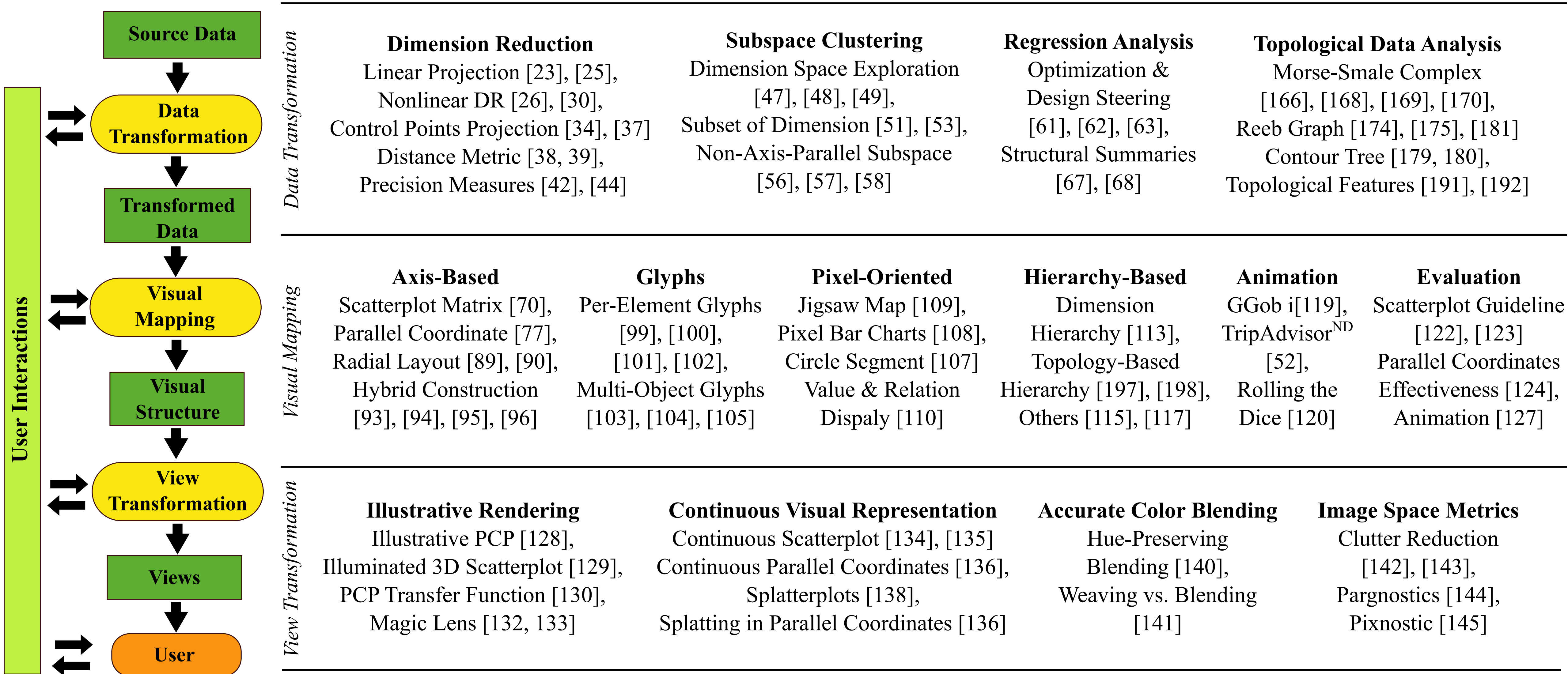
5. AnkerstKeimKriegel1996 [inproceedings] (1996) | PDF | Google Scholar | Google
Circle Segments: A Technique for Visually Exploring Large Multidimensional Data Sets
Mihael Ankerst Daniel A. Keim Hans-peter Kriegel
Abstract: In this paper, we describe a novel technique for visualizing large amounts of high-dimensional data, called 'circle segments'. The technique uses one colored pixel per data value and can therefore be classified as a pixel-per-value technique. The basic idea of the 'circle segments' visualization ... ▶
pipeline stage:visual mapping user involvement:? paper type:technical data type:?
analysis method:? visual method:pixel-based +
select similar BibTeX



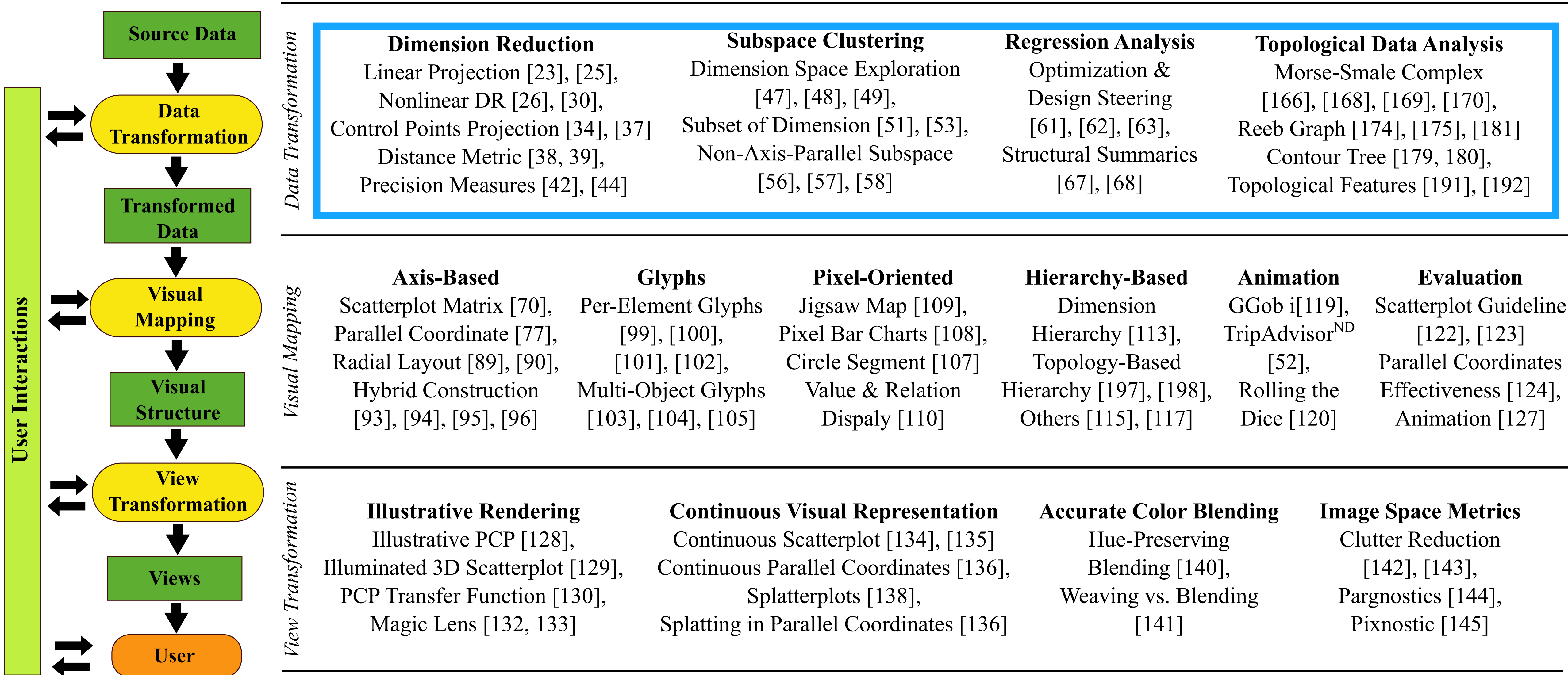
6. ArterodeOliveiraLevkowitz2004 [inproceedings] (2004) | PDF | DOI | Google Scholar | Google
Uncovering Clusters in Crowded Parallel Coordinates Visualizations
Artero, A.O. de Oliveira, M.C.F. Levkowitz, H.
Abstract: The one-to-one strategy of mapping each single data item into a graphical marker adopted in many visualization techniques has limited usefulness when the number of records and/or the dimensionality of the data set are very high. In this situation, the strong overlapping of graphical markers sever... ▶
pipeline stage:visual mapping user involvement:computation centric paper type:technical
data type:high-dimensional points analysis method:clustering visual method:parallel coordinates
view optimization +
select similar BibTeX



Visualization pipeline for high-dim data



Visualization pipeline for HD data



Visualization pipeline for HD data

ML in data transformation

Dimension Reduction

Linear Projection [23], [25],
Nonlinear DR [26], [30],
Control Points Projection [34], [37]
Distance Metric [38, 39],
Precision Measures [42], [44]

Subspace Clustering

Dimension Space Exploration
[47], [48], [49],
Subset of Dimension [51], [53],
Non-Axis-Parallel Subspace
[56], [57], [58]

Regression Analysis

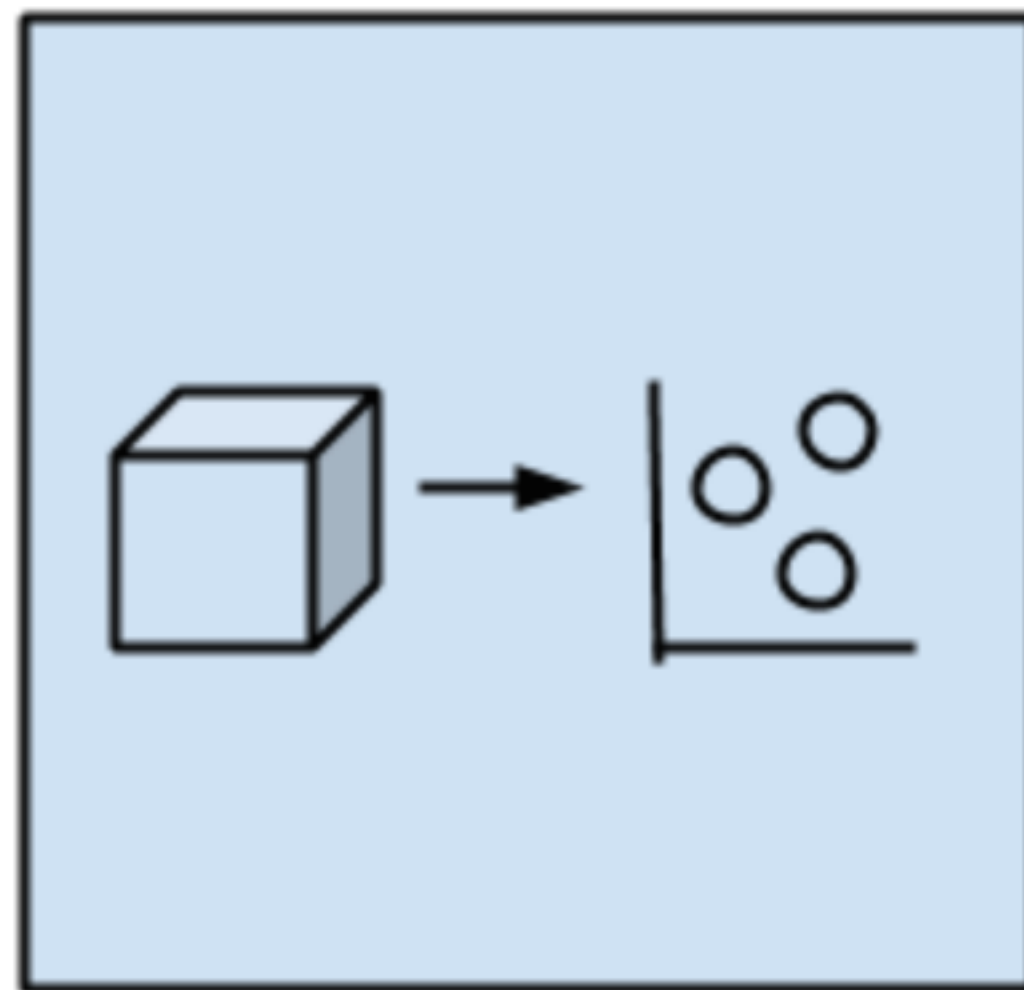
Optimization &
Design Steering
[61], [62], [63],
Structural Summaries
[67], [68]

Topological Data Analysis

Morse-Smale Complex
[166], [168], [169], [170],
Reeb Graph [174], [175], [181]
Contour Tree [179, 180],
Topological Features [191], [192]

Dimensionality Reduction (DR)

Vis+DR can be a semester worth of material...



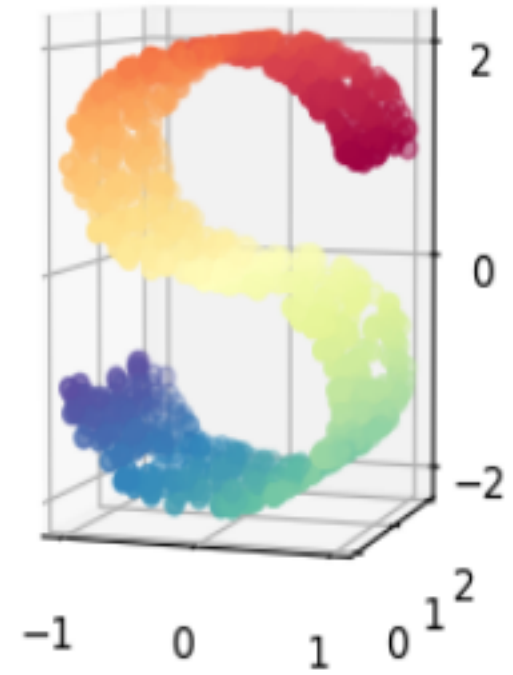
Dimensional Reduction
Algorithms

- Seek and explore the inherent structure in data
- Unsupervised
- Data compression, summarization
- Pre-processing for vis and supervised learning
- Can be adapted for classification and regression
- Well-known DR algorithms:
 - Principal Component Analysis (PCA)
 - Principal Component Regression (PCR)
 - Partial Least Squares Regression (PLSR)
 - Multidimensional Scaling (MDS)
 - Projection Pursuit
 - Linear Discriminant Analysis (LDA)
 - Mixture Discriminant Analysis (MDA)
 - ...

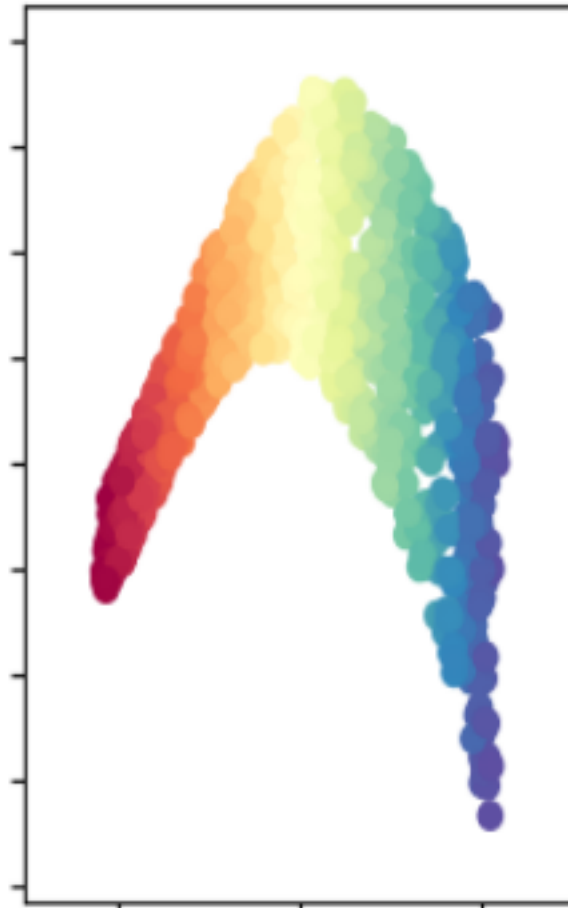
Linear vs nonlinear DR

- Linear: Principal Component Analysis (PCA)
- Nonlinear DR, Manifold learning:
 - Isomap
 - Locally Linear Embedding (LLE)
 - Hessian Eigenmapping
 - Spectral Embedding
 - Multi-dimensional Scaling (MDS)
 - t-distributed Stochastic Neighbor Embedding (t-SNE)

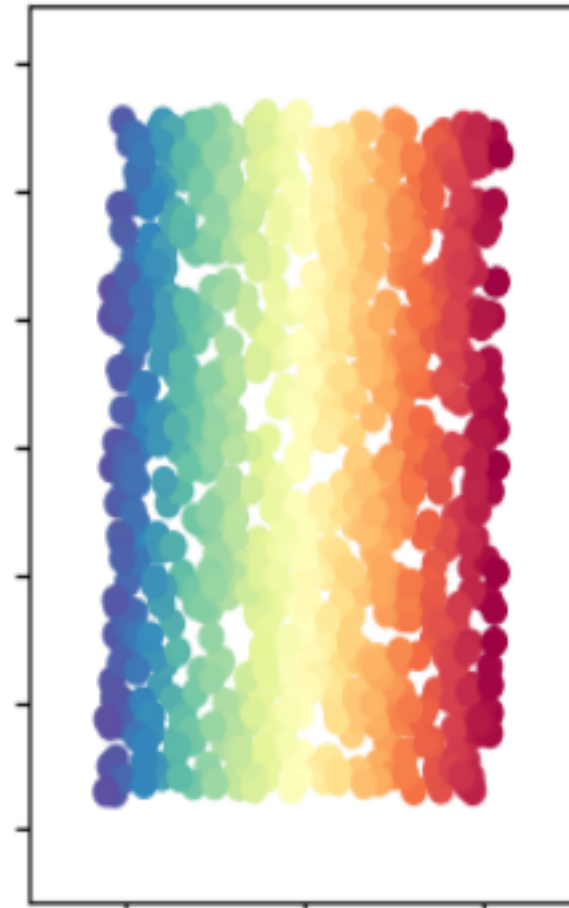
Manifold Learning with 1000 points, 10 neighbors



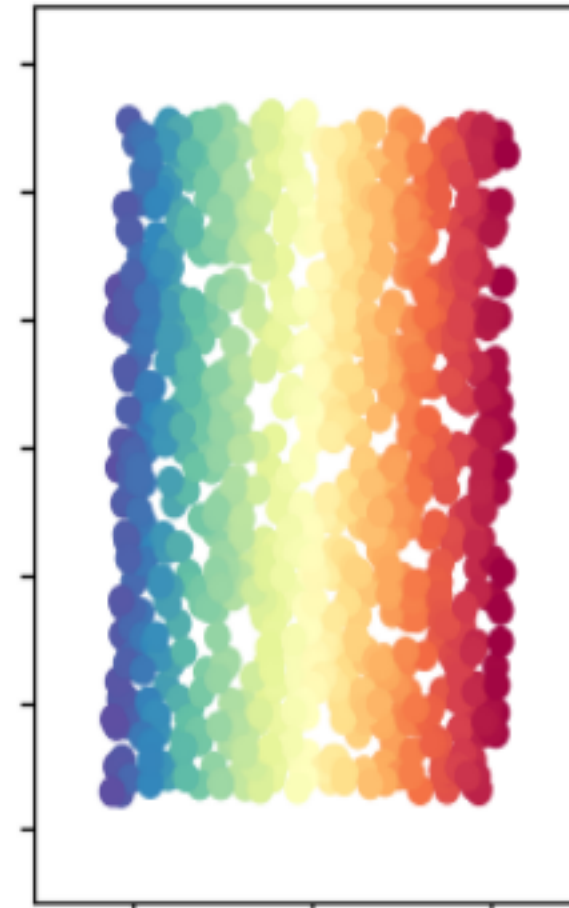
LLE (0.23 sec)



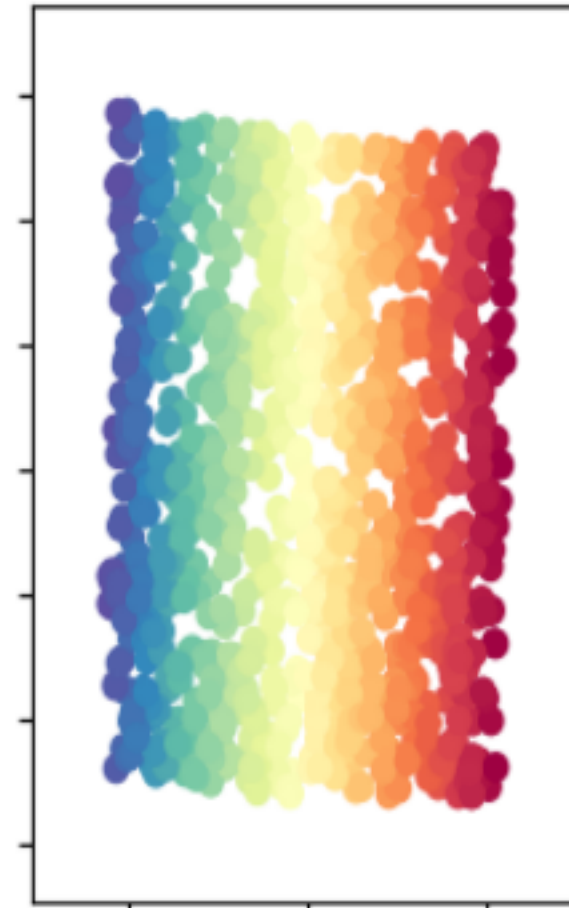
LTSA (0.37 sec)



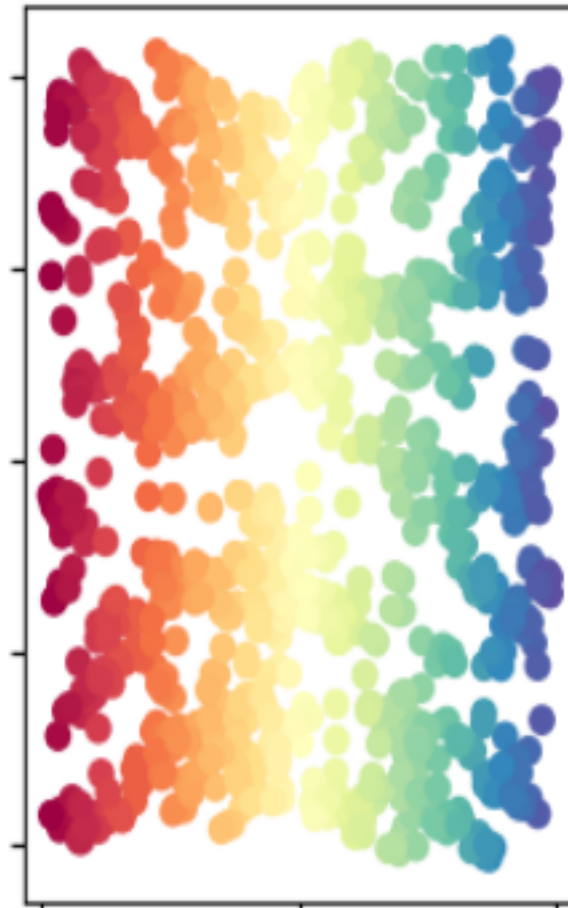
Hessian LLE (0.52 sec)



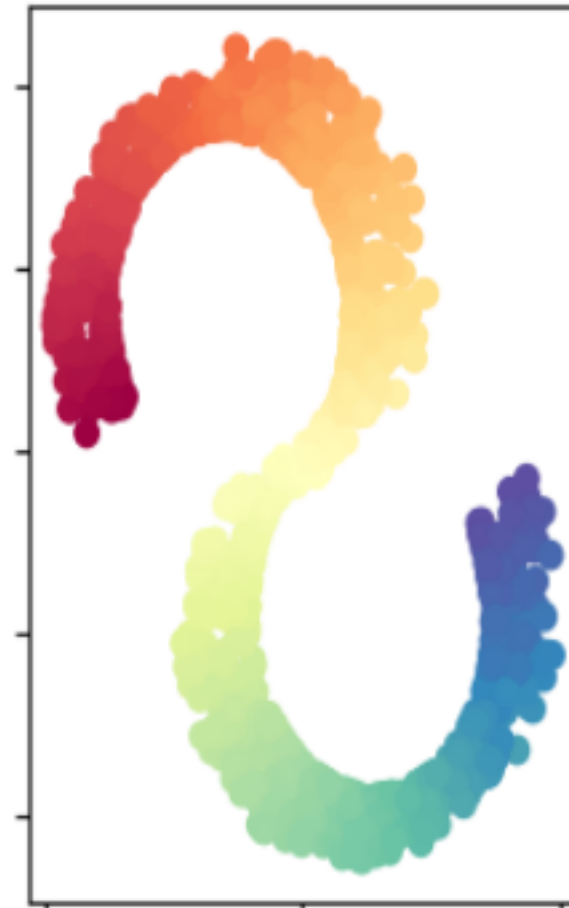
Modified LLE (0.43 sec)



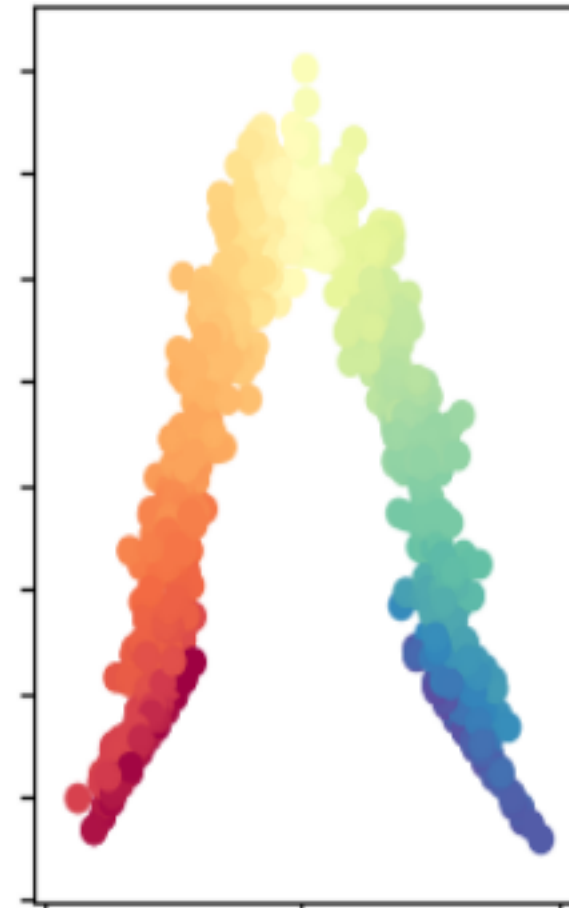
Isomap (0.46 sec)



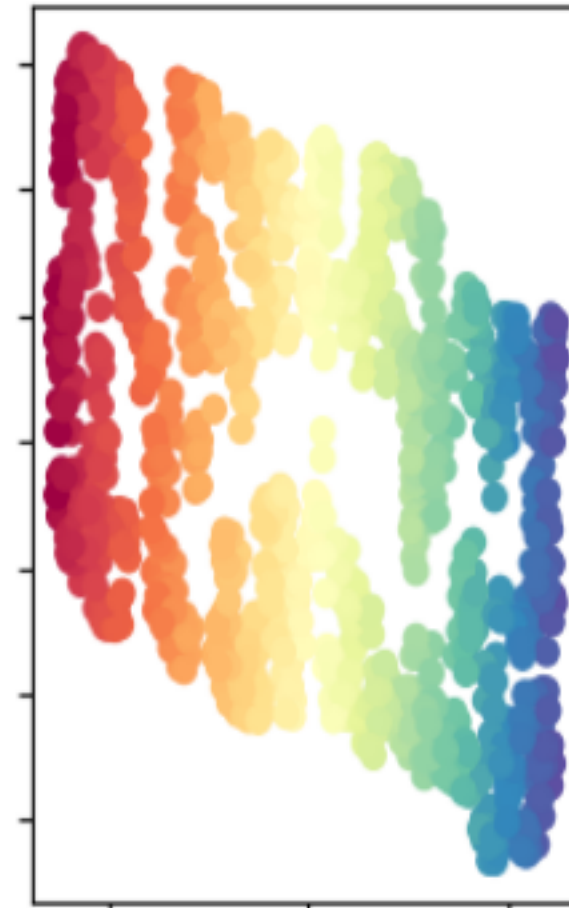
MDS (2.1 sec)



SpectralEmbedding (0.22 sec)

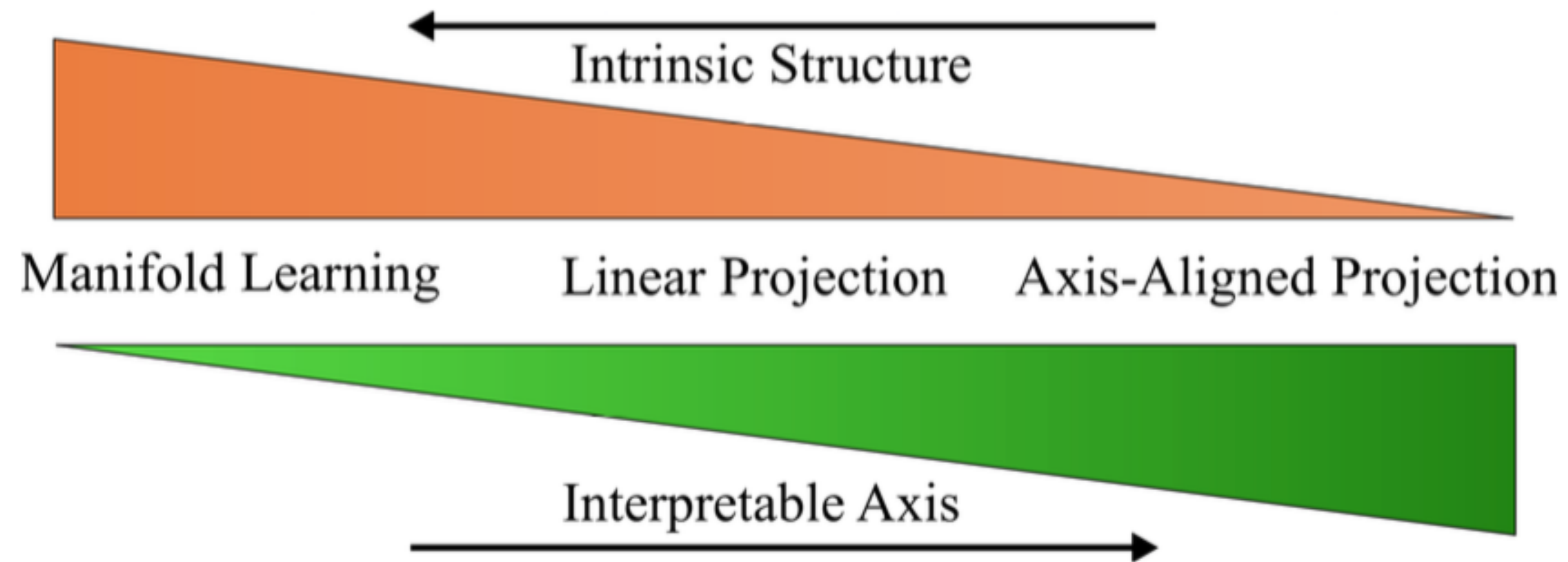


t-SNE (17 sec)



Manifold learning

Interpretability trade off



DR and Vis Overview

How do we proceed from here

- Give two case studies involving DR + Vis
 - Case 1: PCA + Vis (simple)
 - Case 2: SNE and t-SNE + Vis (more involved)
- We do not go through all (but some of) the mathematical details of these algorithms, but instead give a high-level overview of what the algorithm is trying to do
- You are encouraged to follow references and recommended readings to obtain in-depth understanding of these algorithms
- You can use these case studies to think about what might be a good final project

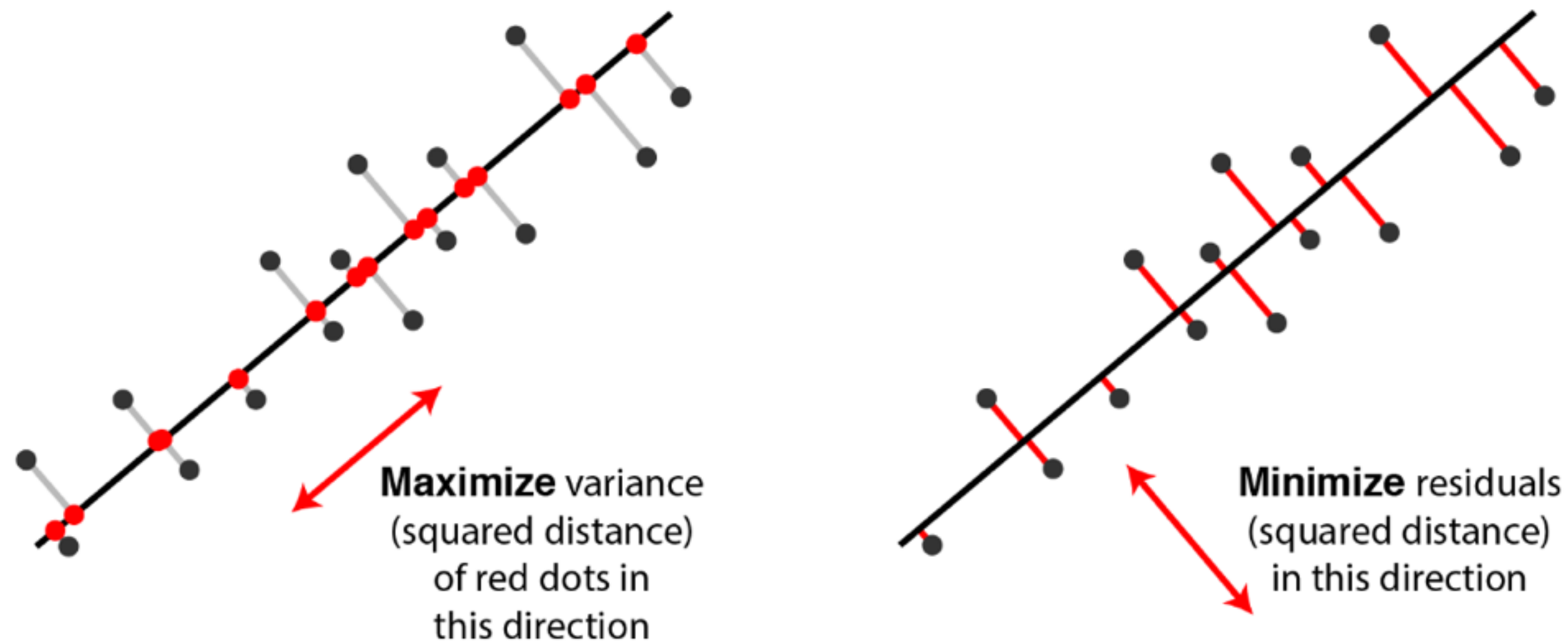
Vis + DR: PCA

A case study with a linear DR method

Three interpretation of PCA

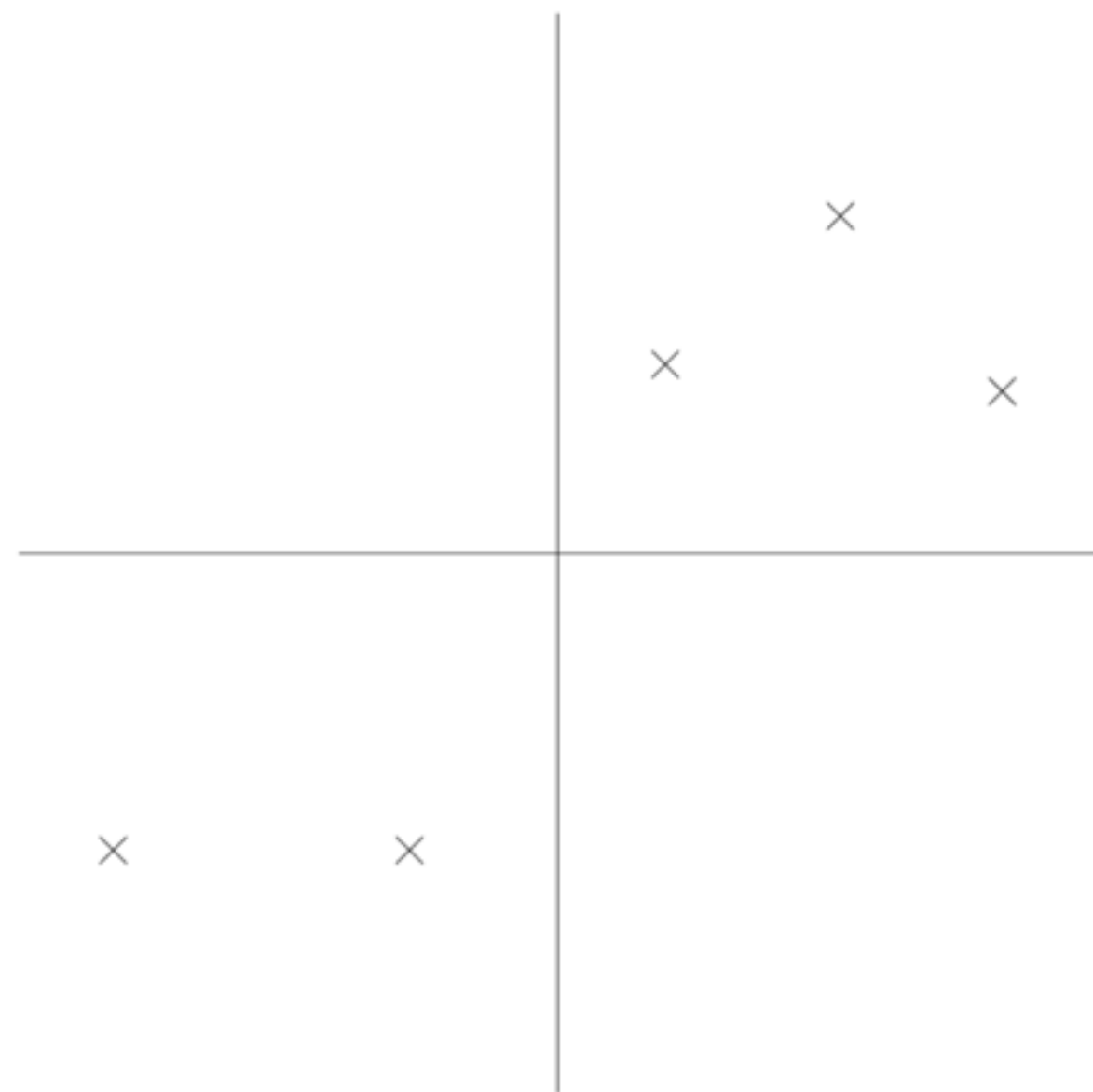
PCA can be interpreted in 2 different ways:

- Maximize the variance of projection along each component (dimension).
- Minimize the reconstruction error, that is, the squared distance between the original data and its projected coordinates.

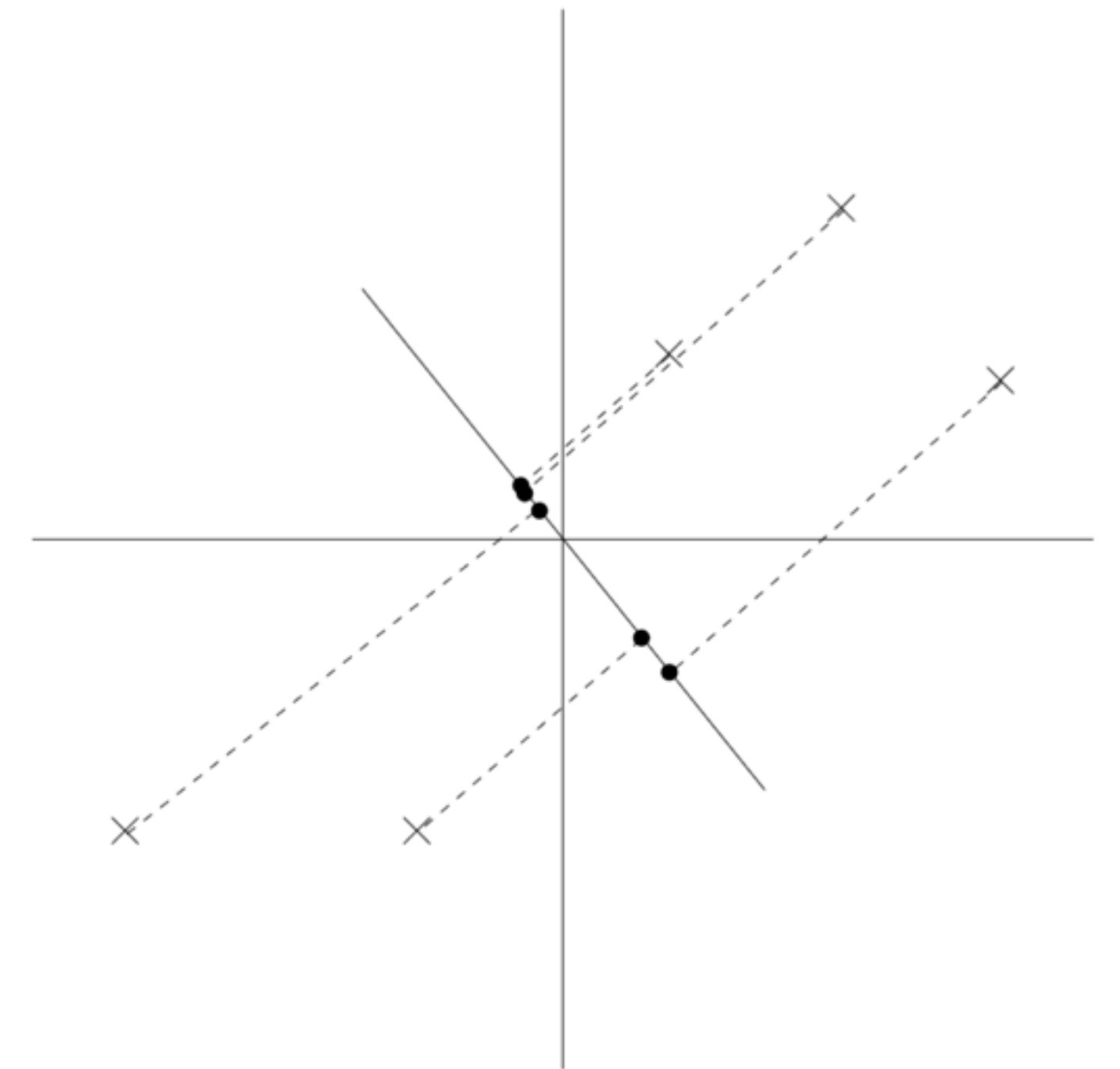


Two equivalent views of principal component analysis.

PCA at a glance

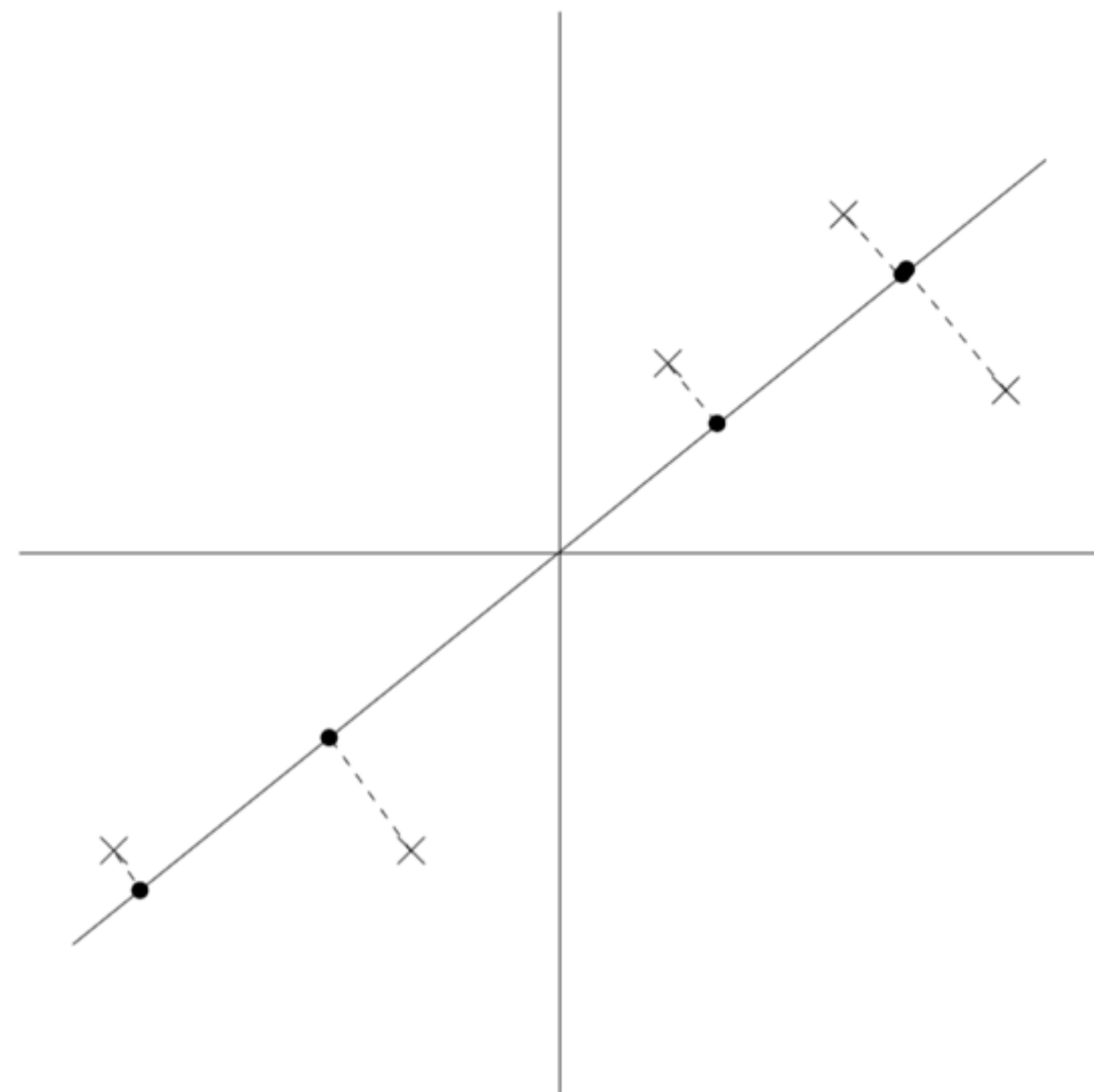


Data after normalization



A projection with small variance

PCA at a glance



A projection with large variance

- PCA automatically choose project direction that maximizes the variance
- The direction of maximum variance in the input space happens to be the same as the principal eigenvector of the covariance matrix of the data
- PCA algorithm: finding the **eigenvalues and eigenvectors** of the covariance matrix.
- The eigenvectors with the largest eigenvalues correspond to the dimensions that have the strongest correlation in the dataset; this is the principle component.

Eigenvalues and eigenvectors

For a given matrix \mathbf{A} , what are the vectors \mathbf{x} for which the product \mathbf{Ax} is a scalar multiple of \mathbf{x} ? That is, what vectors \mathbf{x} satisfy the equation

$$\mathbf{Ax} = \lambda\mathbf{x}$$

for some scalar λ ?

Eigen decomposition theorem

Let \mathbf{P} be a **matrix** of **eigenvectors** of a given **square matrix** \mathbf{A} and \mathbf{D} be a **diagonal matrix** with the corresponding eigenvalues on the diagonal. Then, as long as \mathbf{P} is a **square matrix**, \mathbf{A} can be written as an **eigen decomposition**

$$\mathbf{A} = \mathbf{P} \mathbf{D} \mathbf{P}^{-1},$$

where \mathbf{D} is a **diagonal matrix**. Furthermore, if \mathbf{A} is **symmetric**, then the columns of \mathbf{P} are **orthogonal vectors**.

Covariance matrix

$$Q = XX^T = \begin{bmatrix} \mathbf{x}_1 - \bar{\mathbf{x}} & \mathbf{x}_2 - \bar{\mathbf{x}} & \cdots & \mathbf{x}_n - \bar{\mathbf{x}} \end{bmatrix} \begin{bmatrix} (\mathbf{x}_1 - \bar{\mathbf{x}})^T \\ (\mathbf{x}_2 - \bar{\mathbf{x}})^T \\ \vdots \\ (\mathbf{x}_n - \bar{\mathbf{x}})^T \end{bmatrix}$$

X: data; each col is a data point; each row is a dim.
Don't want to explicitly compute Q: can be huge!
Instead, using SVD, singular value decomposition.

Singular value decomposition (SVD)

Any $m \times n$ matrix X can be decomposed into three matrices:

$$X = U\Sigma V^T$$

- U is $m \times m$ and its columns are orthonormal vectors (i.e. perpendicular)
- Σ is $n \times n$ and its columns are orthonormal vectors
- D is $m \times n$ diagonal and its diagonal elements are called the singular values of X

Relation between PCA and SVD

Simply put, the PCA viewpoint requires that one compute the eigenvalues and eigenvectors of the covariance matrix, which is the product $\mathbf{X}\mathbf{X}^\top$, where \mathbf{X} is the data matrix. Since the covariance matrix is symmetric, the matrix is diagonalizable, and the eigenvectors can be normalized such that they are orthonormal:

$$\mathbf{X}\mathbf{X}^\top = \mathbf{W}\mathbf{D}\mathbf{W}^\top$$

On the other hand, applying SVD to the data matrix \mathbf{X} as follows:

$$\mathbf{X} = \mathbf{U}\mathbf{\Sigma}\mathbf{V}^\top$$

and attempting to construct the covariance matrix from this decomposition gives

$$\mathbf{X}\mathbf{X}^\top = (\mathbf{U}\mathbf{\Sigma}\mathbf{V}^\top)(\mathbf{U}\mathbf{\Sigma}\mathbf{V}^\top)^\top$$

$$\mathbf{X}\mathbf{X}^\top = (\mathbf{U}\mathbf{\Sigma}\mathbf{V}^\top)(\mathbf{V}\mathbf{\Sigma}\mathbf{U}^\top)$$

and since \mathbf{V} is an orthogonal matrix ($\mathbf{V}^\top\mathbf{V} = \mathbf{I}$),

$$\mathbf{X}\mathbf{X}^\top = \mathbf{U}\mathbf{\Sigma}^2\mathbf{U}^\top$$

and the correspondence is easily seen (the square roots of the eigenvalues of $\mathbf{X}\mathbf{X}^\top$ are the singular values of \mathbf{X} , etc.)

Performing SVD on data matrix

X is the (normalized) data matrix, perform SVD on X :

$$X = UDV^T$$

- The columns of U are the eigenvectors of covariance matrix: XX^T
- The columns of V are the eigenvectors of $X^T X$
- The squares of the diagonal elements of D are the eigenvalues of XX^T and $X^T X$

PCA related readings

- Many PCA lectures are available on the web
- Reading materials
 - <http://www.cse.psu.edu/~rtc12/CSE586Spring2010/lectures/pcaLectureShort.pdf>
 - <http://cs229.stanford.edu/notes/cs229-notes10.pdf>
- Things you should pay attention when using PCA
 - Make sure the data is centered: normalize mean and variance

Using PCA with scikit-learn

```
import numpy as np
from sklearn.decomposition import PCA
X = np.array([[ -1, -1], [-2, -1], [-3, -2], [ 1,  1], [ 2,  1], [ 3,  2]])
pca = PCA(n_components=2)
pca.fit(X)

print(pca.explained_variance_ratio_)

print(pca.singular_values_)
```

iPCA: interactive PCA

iPCA: An Interactive System for PCA-based Visual Analytics

UNC Charlotte

Dong Hyun Jeong Caroline Ziemkiewicz
William Ribarsky Remco Chang













Simon Fraser University
Brian Fisher

Source: <http://www.knowledgeviz.com/iPCA/> [JeongZiemkiewiczFisher2009]

Video also available at: <http://www.cs.tufts.edu/~remco/publication.html>

iPCA extension: collaborative sys



Button	Meaning	Button	Meaning
	Go back to the initial state		Delete the selected item(s)
	Individual item selection		Partition the selected item(s) into a new workspace
	Range item(s) selection		Close the application
	Manipulation		Create a new application
	Trail enable – on/ off		Rotate the application
	Cancel the selected item(s)		Make the sliderbar panel appear / disappear

Vis + DR: t-SNE

A case study with a nonlinear DR method

The material from this section is heavily drawn from Jaakko Peltonen
http://www.uta.fi/sis/mtt/mtts1-dimensionality_reduction/drv_lecture10.pdf

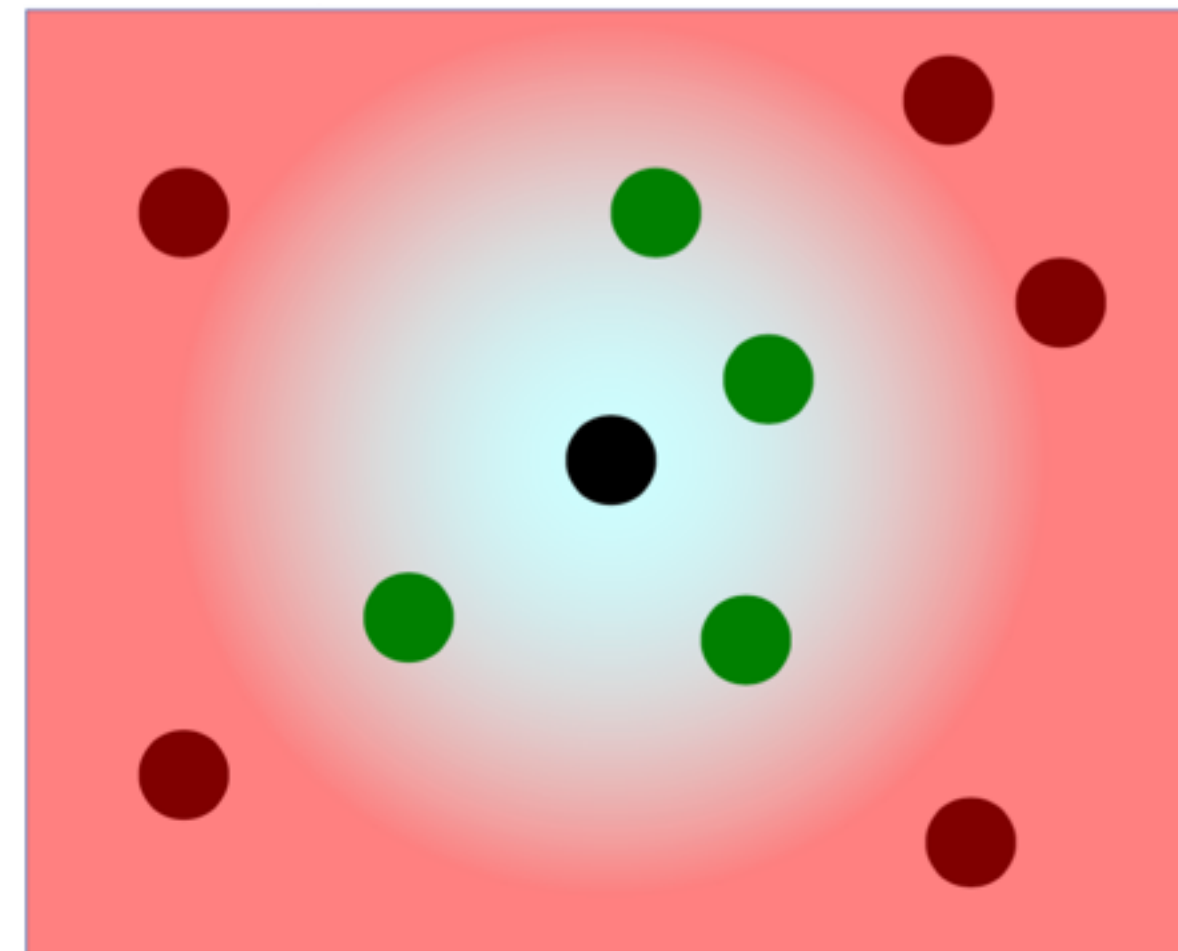
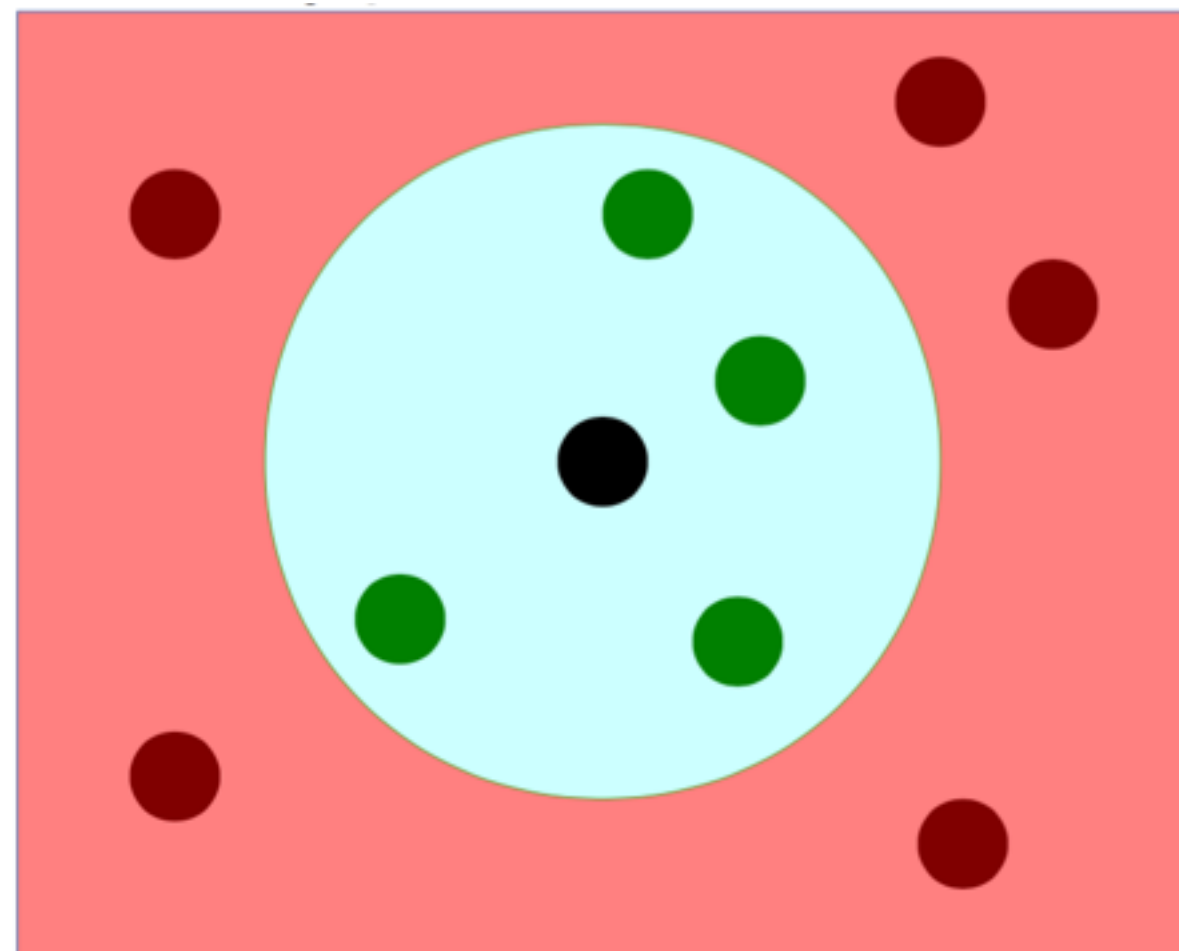
DR: preserving distances

$$C = \frac{1}{a} \sum_{ij} w_{ij} (d_X(x_i, x_j) - d_Y(y_i, y_j))^2$$

- Many DR methods focus on **preserving distances**, e.g. the above is the cost function for a particular DR method called **metric MDS**
- An alternative idea is **preserving neighborhoods**.

DR: preserving neighborhoods

- Neighbors are an important notion in data analysis, e.g. social networks, friends, twitter followers...
- Object nearby (in a metric space) are considered neighbors
- Consider **hard neighborhood** and **soft neighborhood**
- Hard: each point is a neighbor (green) or a non-neighbor (red)
- Soft: each point is a neighbor (green) or a non-neighbor (red) with some weight

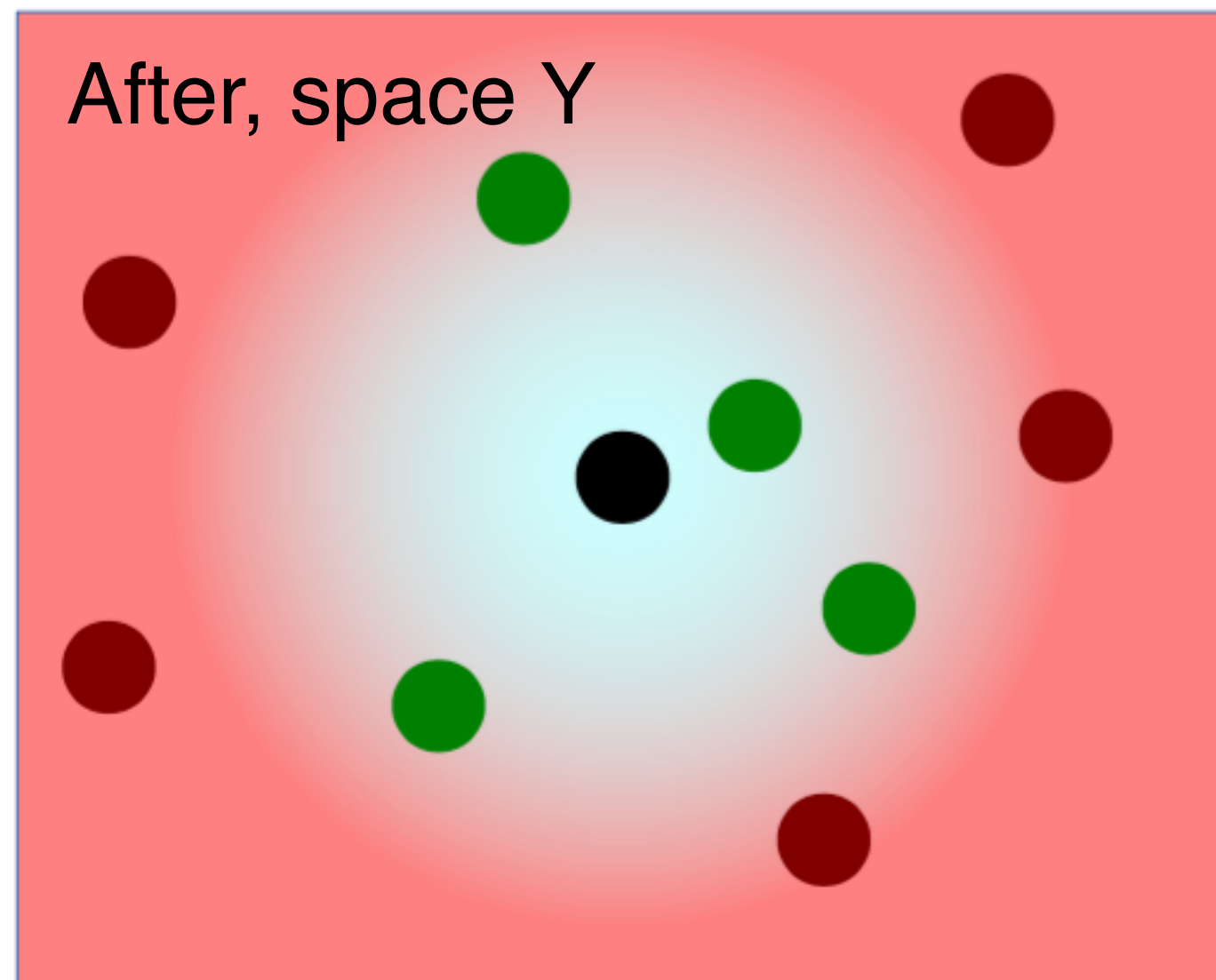
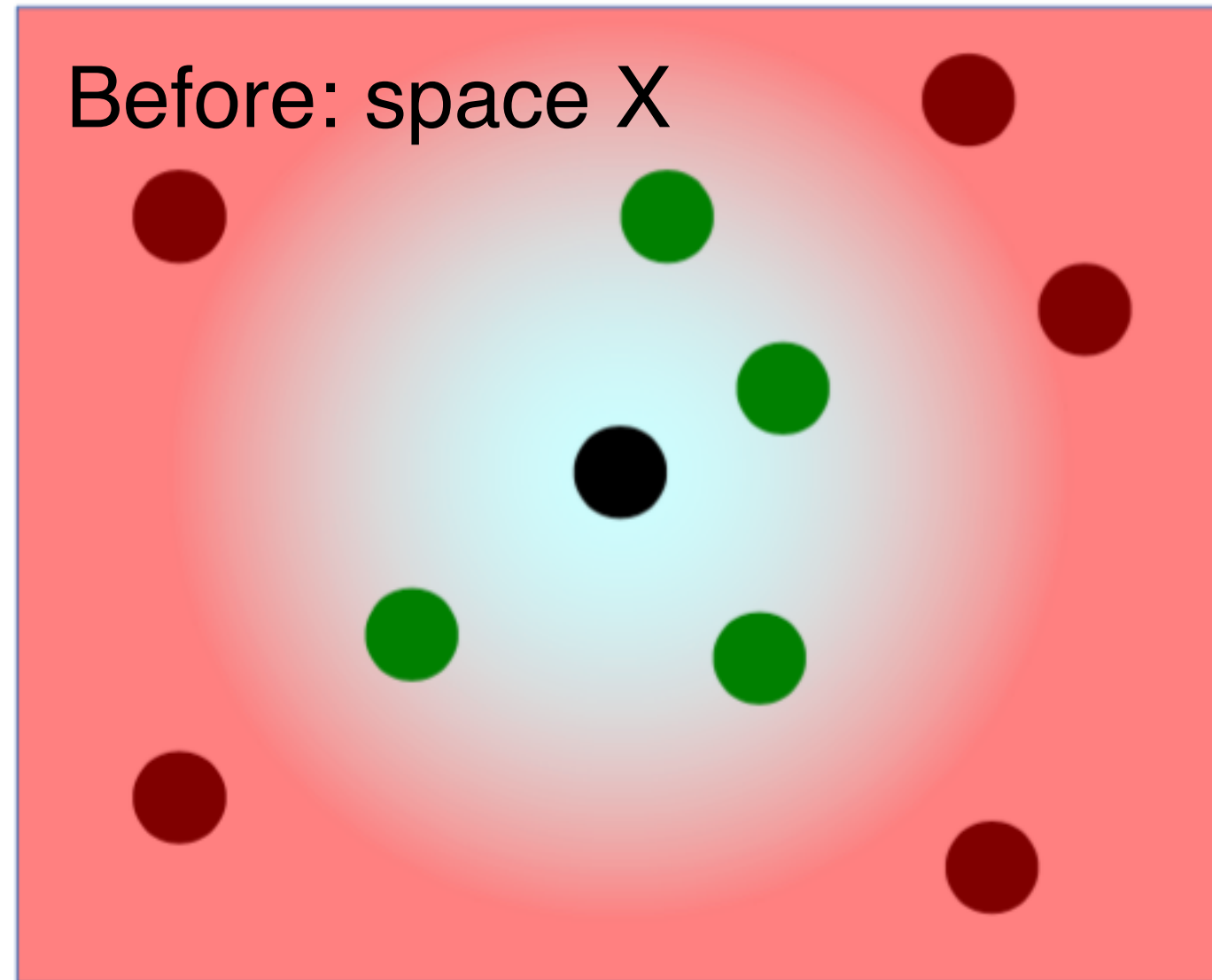


Probabilistic neighborhood

- Derive a probability of point j to be picked as a neighbor of i in the input space

$$P_{ij} = \frac{\exp(-d_{ij}^2)}{\sum_{k \neq i} \exp(-d_{ik}^2)}$$

Preserving nbhds before & after DR



$$P_{ij} = \frac{\exp(-||x_i - x_j||^2)}{\sum_{k \neq i} \exp(-||x_i - x_k||^2)}$$

Probabilistic **input** neighborhood:
Probability to be picked as a neighbor in space X (input coordinates)

$$Q_{ij} = \frac{\exp(-||y_i - y_j||^2)}{\sum_{k \neq i} \exp(-||y_i - y_k||^2)}$$

Probabilistic **output** neighborhood:
Probability to be picked as a neighbor in space Y (display coordinates)

Stochastic neighbor embedding

- Compare neighborhoods between the input and output!
- Using Kullback-Leibler (KL) divergence
- KL divergence: relative entropy (amount of surprise when encounter items from 1st distribution when they are expected to come from the 2nd)
- KL divergence is nonnegative and 0 iff the distributions are equal
- **SNE: minimizes the KL divergence** using gradient descent

$$C = \sum_i \sum_j p_{ij} \log \frac{p_{ij}}{q_{ij}}$$

SNE: choose the size of a nbhd

- How to set the size of a neighborhood? Using a scale parameter: σ_i

$$d_{ij}^2 = \frac{\|x_i - x_j\|^2}{2\sigma_i^2}$$

- The scale parameter can be chosen without knowing much about the data, but...
- It is better to choose the parameter based on local neighborhood properties, and for each point
- E.g., in sparse region, distance drops more gradually

SNE: choose a scale parameter

Choose an **effective** number of neighbors:

- In a uniform distribution over k neighbors, the entropy is $\log(k)$
- Find the scale parameter using binary search so that the entropy of \mathcal{P}_{ij} becomes $\log(k)$ for a desired value of k .

SNE: gradient descent

- Adjusting the output coordinates using [gradient descent](#)
- [Gradient descent](#): iterative process to find the minimal of a function
- Start from a random initial output configuration, then iteratively take steps along the gradient
- Intuition: using forces to pull and push pairs of points to make input and output probabilities more similar

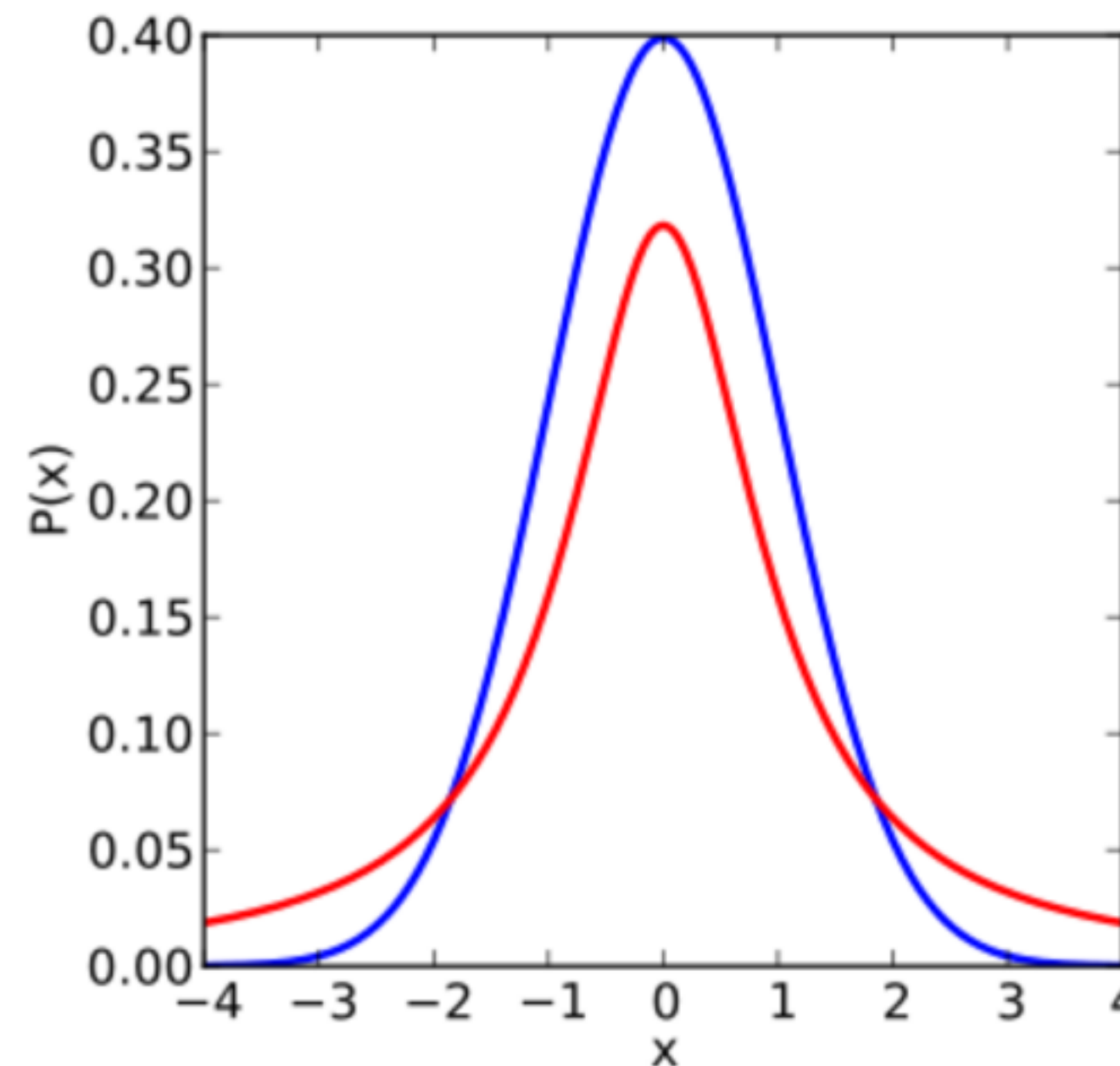
$$\frac{\partial C}{\partial y_i} = 2 \sum_j (y_i - y_j) (p_{ij} - q_{ij} + p_{ji} - q_{ji})$$

SNE: the crowding problem

- When embedding neighbors from a high-dim space into a low- dim space, there is too little space near a point for all of its close-by neighbors.
- Some points end up too far-away from each other
- Some points that are neighbors of many far-away points end up crowded near the center of the display.
- In other words, these points end up **crowded in the center** to stay close to all of the far-away points.
- t-SNE: using heavy-tailed distributions (i.e., t-distributions) to define neighbors on the display, to resolve the crowding problem

t-distributed SNE

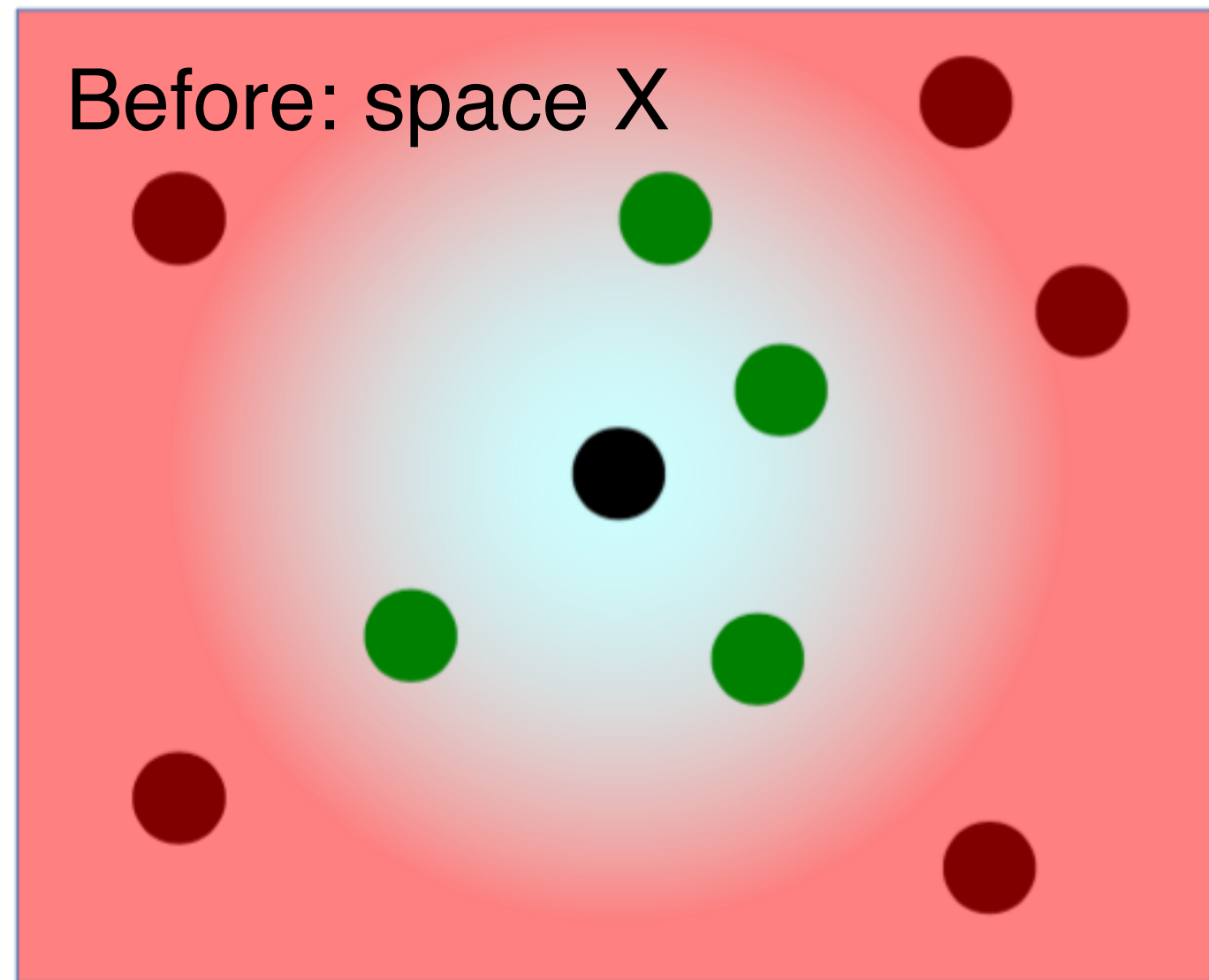
- Avoids crowding problem by using a more heavy-tailed neighborhood distribution in the low-dim output space than in the input space.
- Neighborhood probability falls off less rapidly; less need to push some points far off and crowd remaining points close together in the center.
- Use student-t distribution with 1 degree of freedom in the output space
- t-SNE (joint prob.); SNE (conditional prob.)



Blue: normal dist.

Red: student-t dist. with 1 deg. of freedom

t-SNE: preserving nbhds

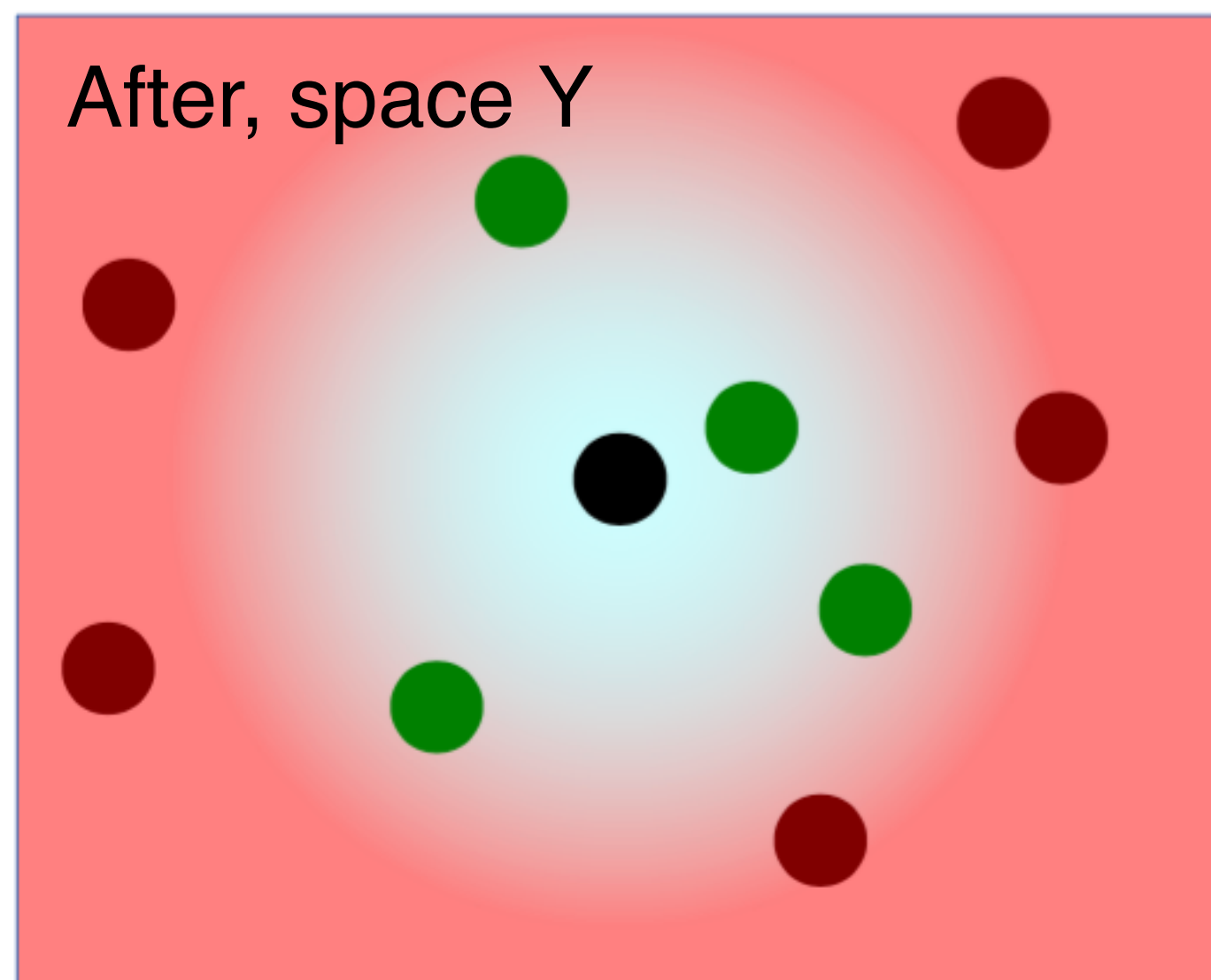


$$p_{j|i} = \frac{\exp(-||x_i - x_j||^2 / 2\sigma_i^2)}{\sum_{k \neq i} \exp(-||x_i - x_k||^2 / 2\sigma_i^2)}$$

$$p_{ij} = \frac{p_{j|i} + p_{i|j}}{2n}$$

Probabilistic **input** neighborhood:

Probability to be picked as a neighbor in space X (input coordinates)

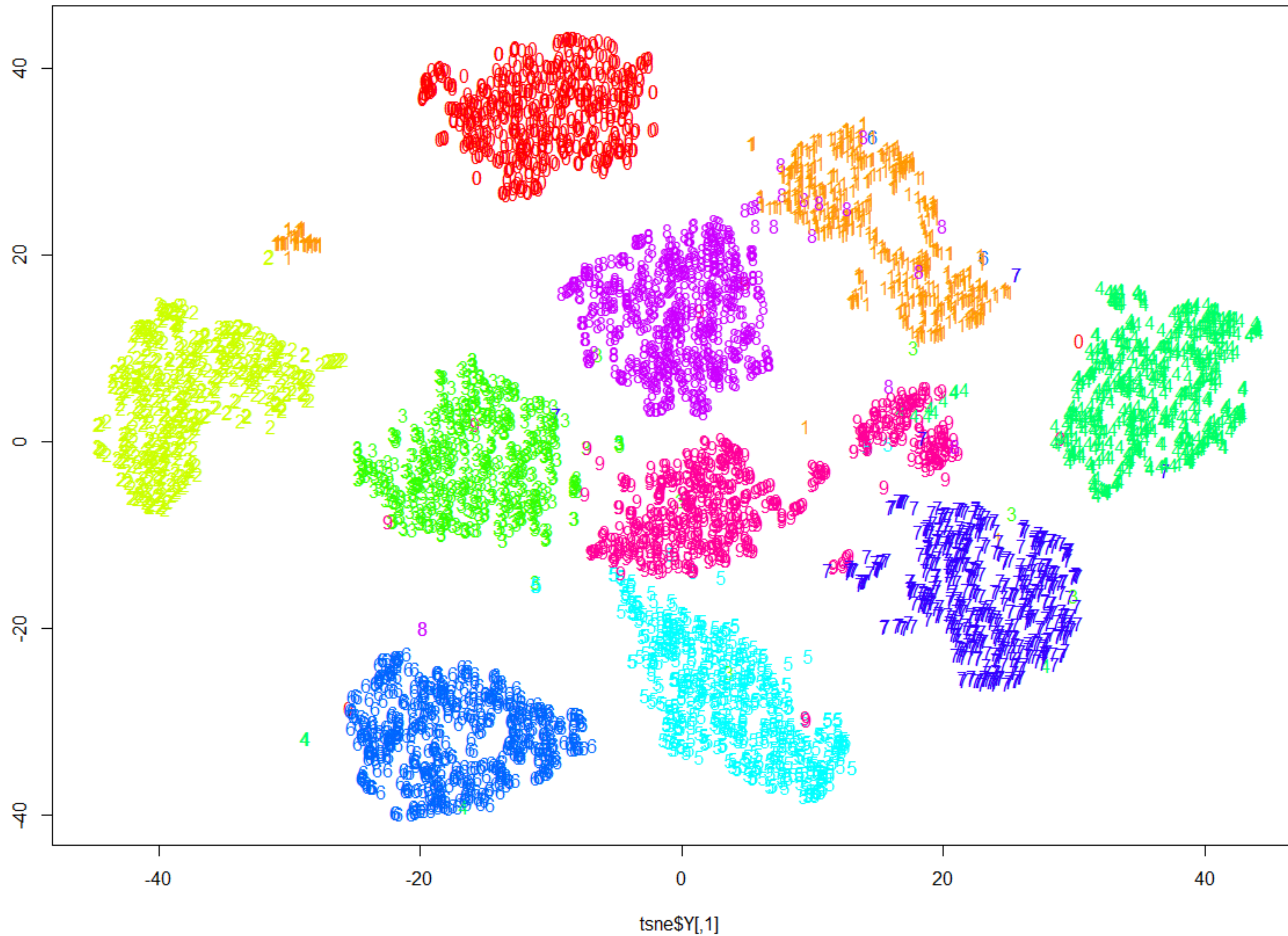


$$q_{ij} = \frac{(1 + ||y_i - y_j||^2)^{-1}}{\sum_{k \neq l} (1 + ||y_k - y_l||^2)^{-1}}$$

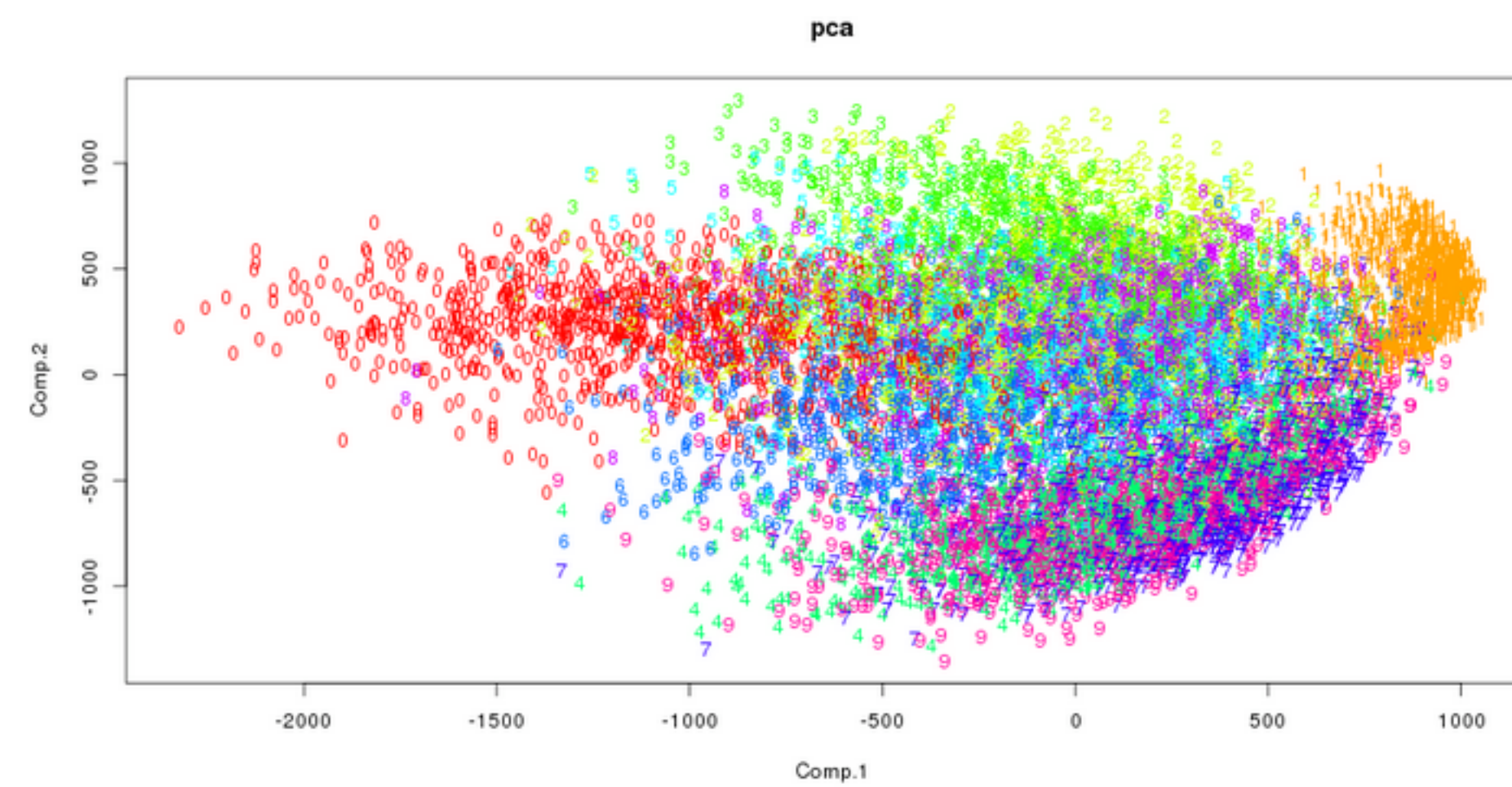
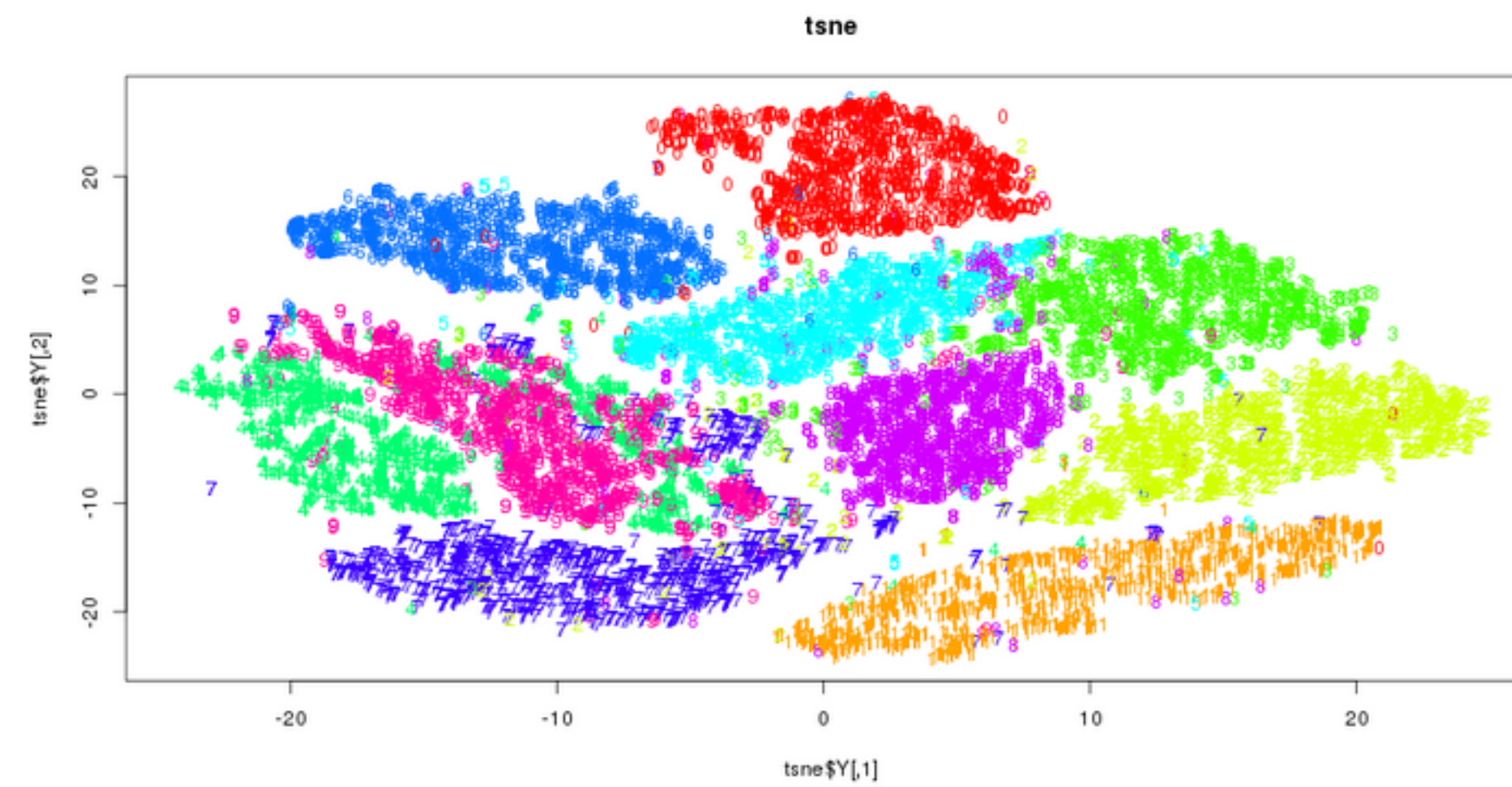
Probabilistic **output** neighborhood:

Probability to be picked as a neighbor in space Y (display coordinates)

Classic t-SNE result



t-SNE vs PCA



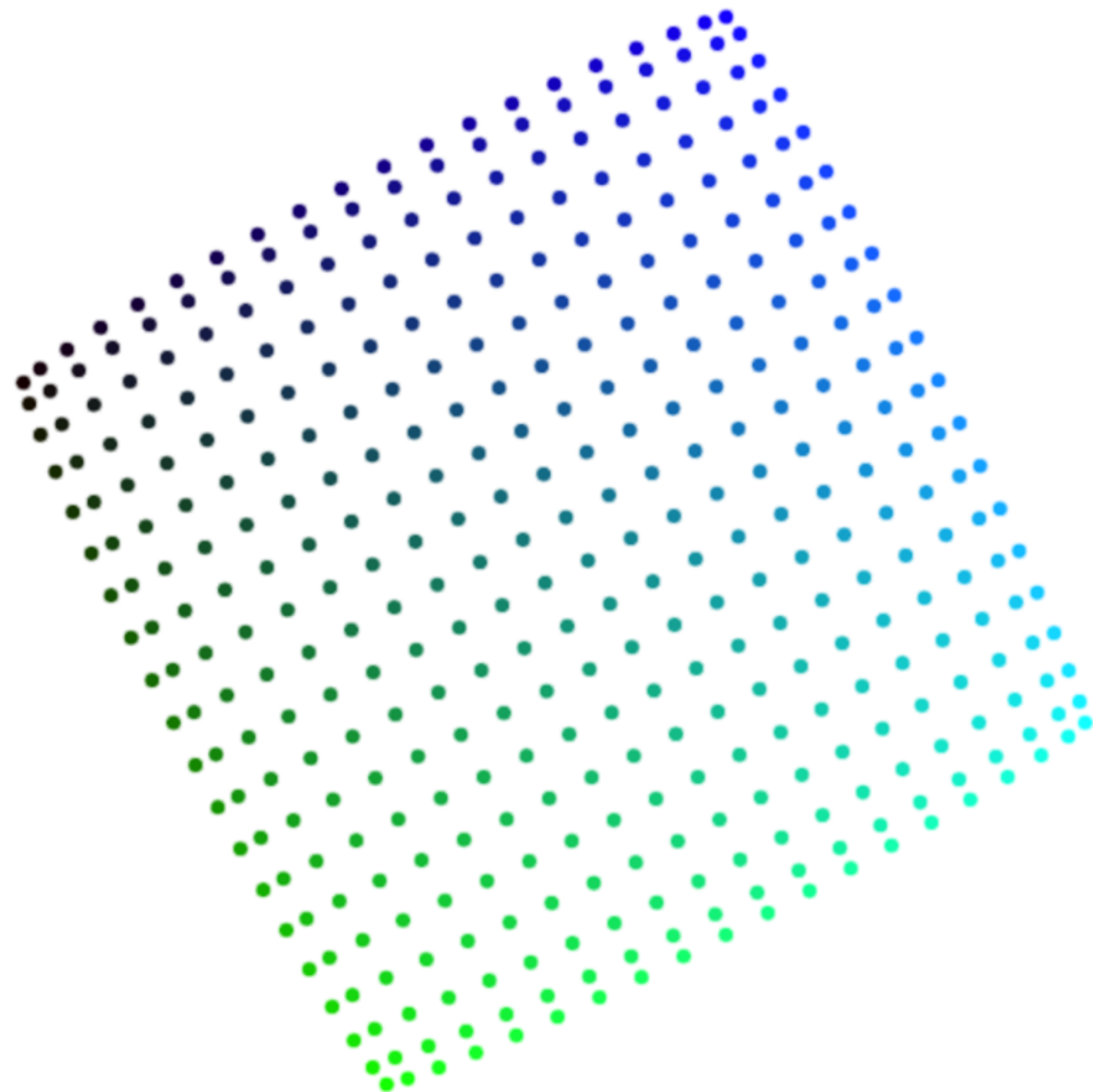
t-SNE

- t-SNE: minimize KL divergence.
- Nonlinear DR.
- Perform diff. transformation on diff. regions: main source of confusing.
- Parameter: **perplexity**, how to balance attention between local and global aspects of your data; guess the # of close neighbor each point has.
- “The performance of t-SNE is fairly robust under different settings of the perplexity. The most appropriate value depends on the density of your data. Loosely speaking, one could say that a larger / denser dataset requires a larger perplexity. Typical values for the perplexity range between 5 and 50.” (Laurens van der Maaten)

What is perplexity anyway?

- “Perplexity is a measure for information that is defined as 2 to the power of the Shannon entropy. The perplexity of a fair die with k sides is equal to k . In t-SNE, the perplexity may be viewed as a knob that sets the number of effective nearest neighbors. It is comparable with the number of nearest neighbors k that is employed in many manifold learners.”

How not to misread t-SNE



Step 420

Points Per Side 20

Perplexity 10

Epsilon 5

A square grid with equal spacing between points. Try convergence at different sizes.

Playing with t-SNE

- http://scikit-learn.org/stable/auto_examples/manifold/plot_t_sne_perplexity.html
- <https://lvdmaaten.github.io/tsne/>

Weakness of t-SNE

- Not clear how it performs on general DR tasks
- Local nature of t-SNE makes it sensitive to intrinsic dim of the data
- Not guaranteed to converge to global minimum

Take home message

- Even a simple DR method like PCA can have interesting visualization aspects to it
- Using visualization to manipulate the input to the ML algorithm, and at the same time understanding the interworking of the algorithm
- Cooperative analysis, mobile devices, virtual reality?

- t-SNE is useful, but only when you know how to **interpret** it
- Those hyper-parameters, such as perplexity, really matter
- Use visualization to interpret the ML algorithm
- Educational purposes to distill algorithms as glass boxes

Getting ready for Project 1

- Scikit-learn tutorial:
 - <http://scikit-learn.org/stable/tutorial/basic/tutorial.html>
- Install and read the documentation of kepler-mapper:
 - <https://github.com/MLWave/kepler-mapper>
- Interactive Data Visualization for the Web, 2nd Ed.
 - <http://alignedleft.com/work/d3-book-2e>

Potential Final Projects

- Inspired by:
 - <http://setosa.io/ev/principal-component-analysis/>
 - <https://distill.pub/2016/misread-tsne/>
- Extending Embedding Projector: Interactive Visualization and Interpretation of Embeddings
 - <https://opensource.googleblog.com/2016/12/open-sourcing-embedding-projector-tool.html>
 - <http://projector.tensorflow.org/>
 - https://www.tensorflow.org/versions/r1.2/get_started/embedding_viz

Can you create a web-based tools that give good visual interpretation of two linear DR and two nonlinear DR techniques?



Thanks!

Any questions?

You can find me at: beiwang@sci.utah.edu

CREDITS

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- ☐ Vector Icons by [Matthew Skiles](#)

Presentation Design

This presentation uses the following typographies and colors:

Free Fonts used:

<http://www.1001fonts.com/oswald-font.html>

<https://www.fontsquirrel.com/fonts/open-sans>

Colors used

