

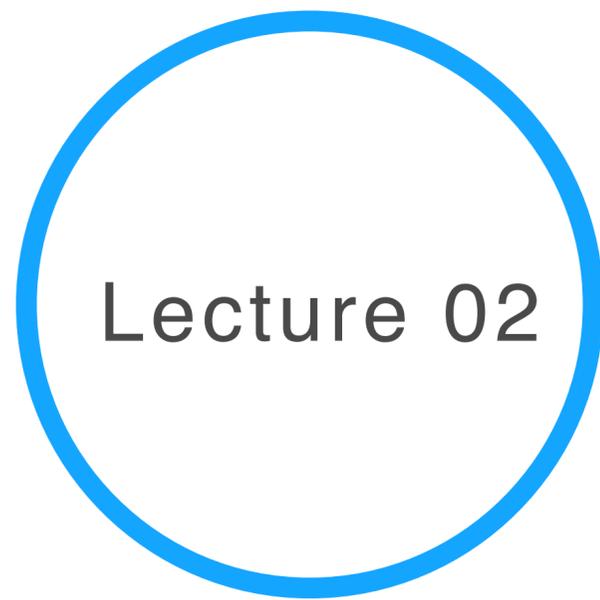
# Advanced Data Visualization

CS 6965

Fall 2019

Prof. Bei Wang Phillips

University of Utah

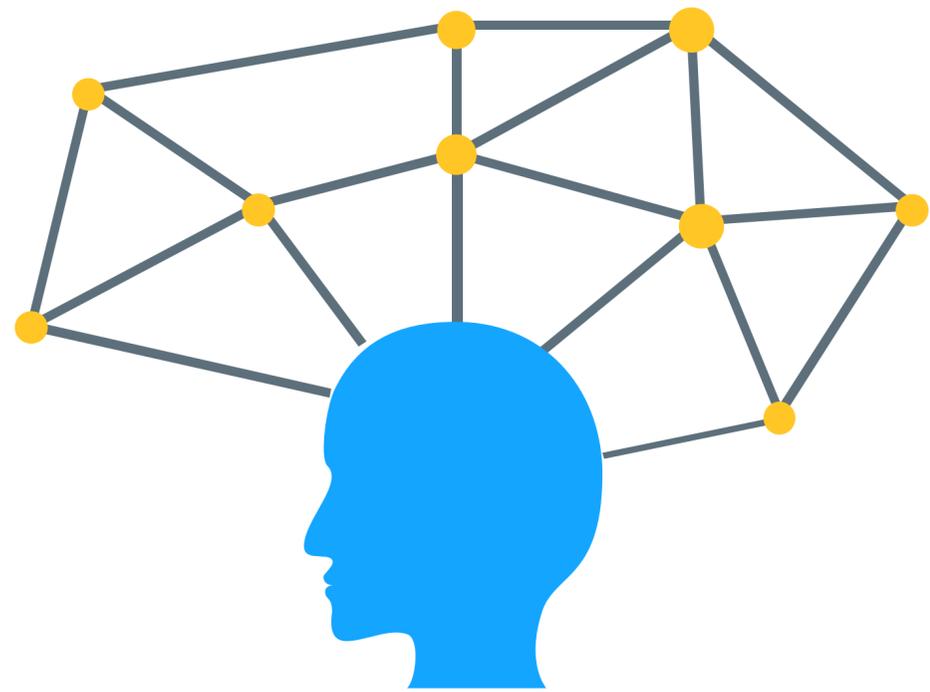


Lecture 02

# 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

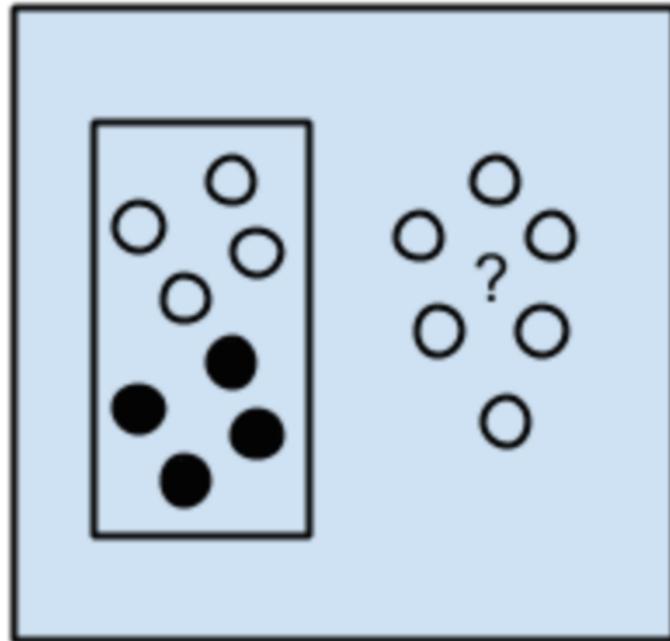
# ML algorithms in a nutshell

# Not a full-blown ML class, but

How to best incorporate vis into ML algorithms?

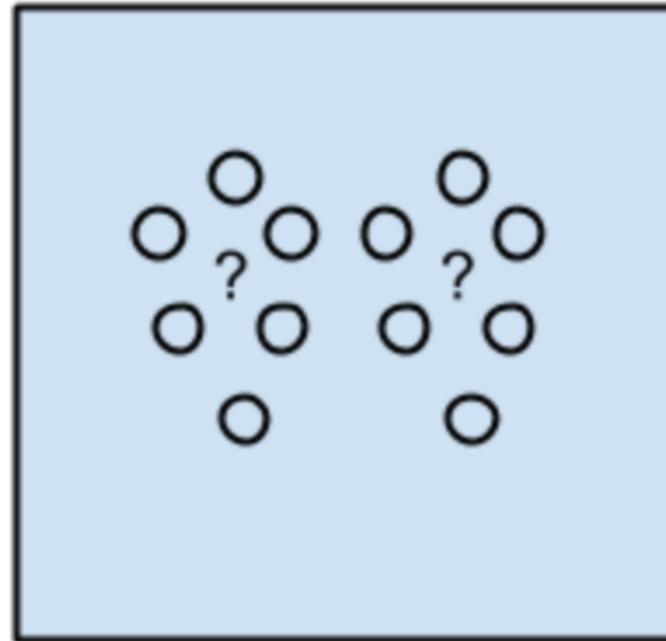
- A simple approach is to treat the ML algorithm as a black box, and build vis surrounding its input/output
- Not knowing the interworking of the algorithm (e.g. a glass box) may lead to misinterpretation of the algorithm output
- We need to have a good understanding of the **core** of some ML algorithms
- 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



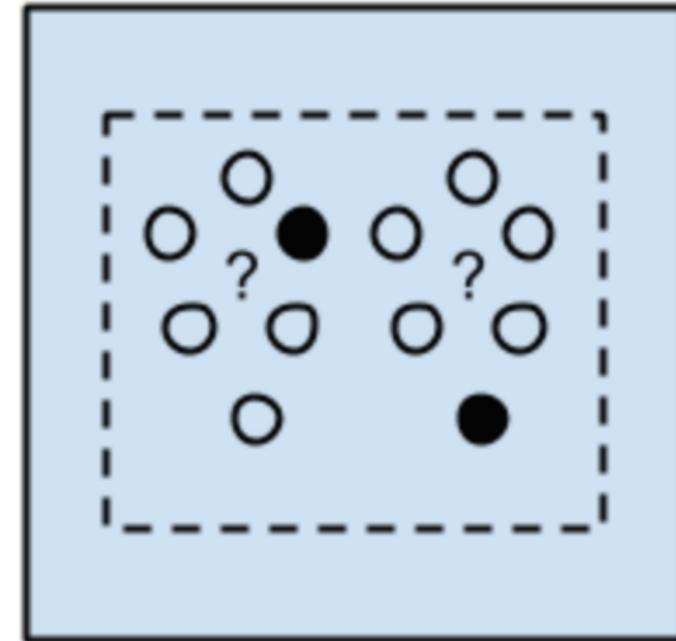
Supervised Learning

Problems: Classification  
Regression



Unsupervised Learning

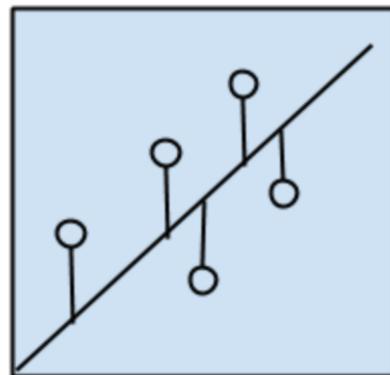
Problems: Clustering  
Dimensionality Reduction



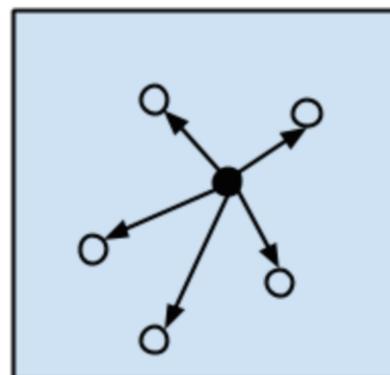
Semi-supervised Learning

Problems: Classification  
Regression

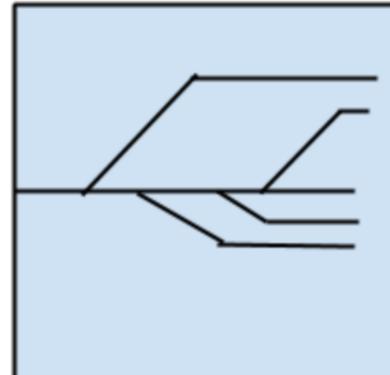
# ML algorithm by similarity (how they work)



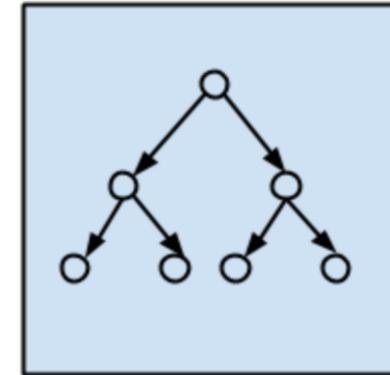
Regression Algorithms



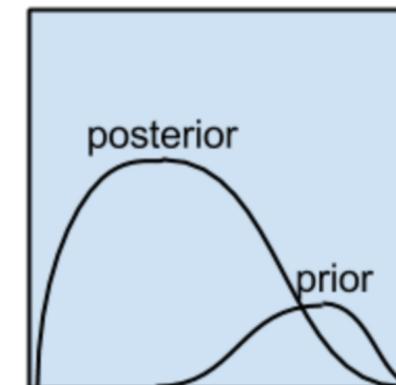
Instance-based Algorithms



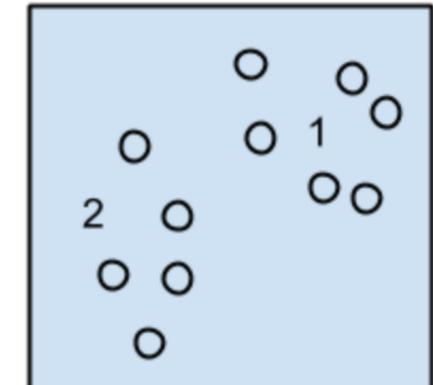
Regularization Algorithms



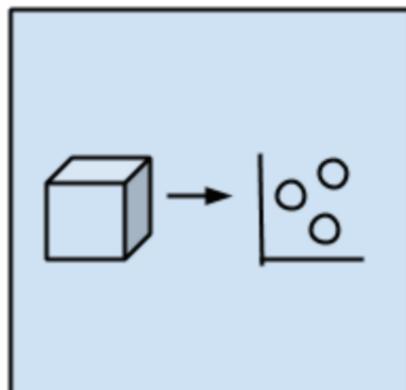
Decision Tree Algorithms



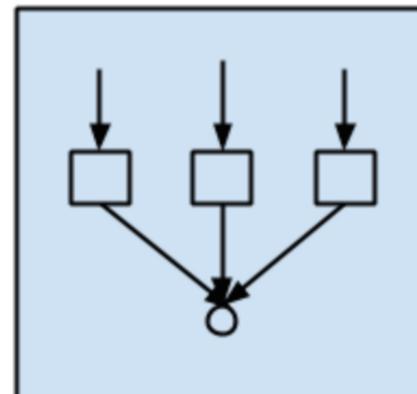
Bayesian Algorithms



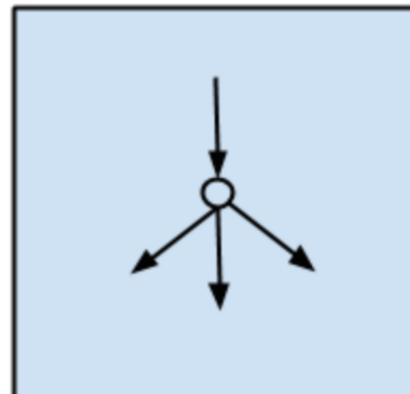
Clustering Algorithms



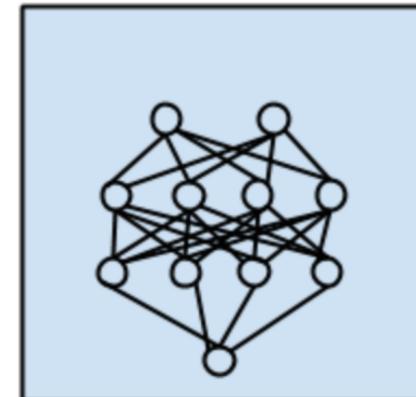
Dimensional Reduction Algorithms



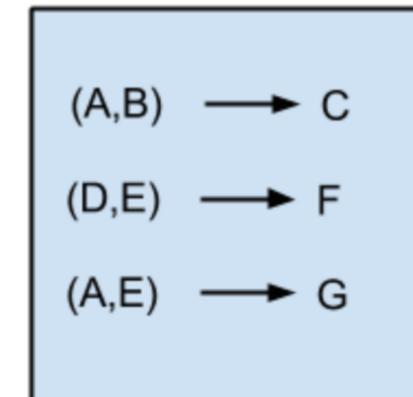
Ensemble Algorithms



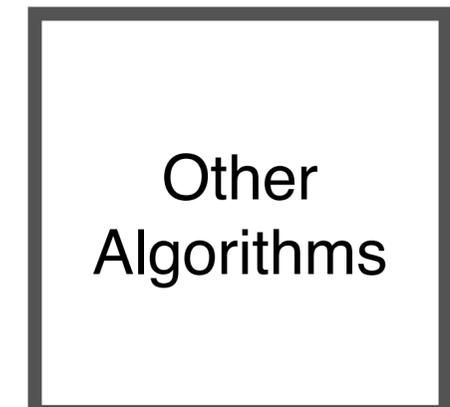
Artificial Neural Network Algorithms



Deep Learning Algorithms



Association Rule Learning Algorithms





# Advances in HD Vis

# Visualizing High-Dimensional Data: Advances in the Past Decade

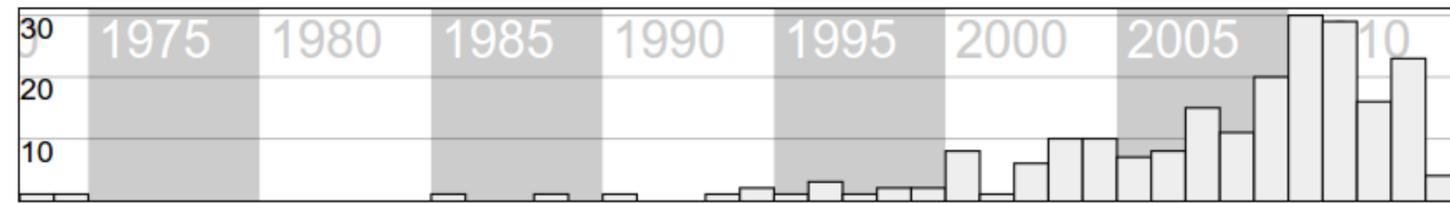
Digital library for publication **Visualizing High-Dimensional Data: Advances in the Past Decade**



Selectors

search ...

## Timeline



## Tags

**pipeline stage:** ?<sub>6</sub> data transformation<sub>137</sub> view transformation<sub>17</sub> visual mapping<sub>62</sub>  
**user involvement:** ?<sub>7</sub> computation centric<sub>61</sub> interactive exploration<sub>144</sub>  
model manipulation<sub>6</sub>  
**paper type:** ?<sub>40</sub> application<sub>7</sub> survey<sub>11</sub> system<sub>11</sub> technical<sub>147</sub> theory<sub>3</sub>  
**data type:** ?<sub>86</sub> high-dimensional function<sub>7</sub> high-dimensional point cloud<sub>1</sub>  
high-dimensional points<sub>100</sub> nominal data<sub>14</sub> spatial data<sub>6</sub> time series<sub>4</sub>  
**analysis method:** ?<sub>55</sub> clustering<sub>83</sub> data abstraction<sub>5</sub> data subset<sub>1</sub> dimension relationship<sub>9</sub>  
dimension similarity<sub>4</sub> dimensionality reduction<sub>25</sub> distance metric<sub>6</sub> feature extraction<sub>2</sub>  
histogram<sub>2</sub> optimization<sub>1</sub> precision measure<sub>5</sub> projection<sub>12</sub> quality measure<sub>1</sub> regression<sub>8</sub>  
regression?<sub>1</sub> scagnostics<sub>1</sub> segmentation<sub>1</sub> statistic<sub>2</sub> subspace<sub>14</sub> topological analysis<sub>9</sub>  
**visual method:** ?<sub>21</sub> animation<sub>6</sub> bar charts<sub>7</sub> focus+context<sub>6</sub> glyphs<sub>10</sub> heat map<sub>1</sub>  
hierarchy<sub>13</sub> isosurface<sub>4</sub> magic lens<sub>4</sub> node-link<sub>3</sub> novel visual encoding<sub>31</sub>  
parallel coordinates<sub>96</sub> pixel-based<sub>5</sub> progressive update<sub>3</sub> radviz<sub>4</sub>  
rendering enhancement<sub>4</sub> scatterplot<sub>59</sub> star coordinates<sub>2</sub> surfaces<sub>7</sub> treemap<sub>3</sub>  
volume visualization<sub>5</sub>  
**other:** <sub>5</sub> clustering<sub>1</sub> clutter reduction<sub>15</sub> comparison<sub>1</sub> high-dimensional points<sub>1</sub> data transformation<sub>1</sub>  
filtering<sub>2</sub> histogram<sub>1</sub> information<sub>1</sub> machine learning<sub>5</sub> matching<sub>1</sub> parameter exploration<sub>8</sub>  
perception<sub>4</sub> query<sub>8</sub> ranking<sub>17</sub> reordering<sub>4</sub> segmentation<sub>1</sub> sensitivity analysis<sub>4</sub> uncertainty<sub>3</sub>  
user study<sub>1</sub> view optimization<sub>1</sub> visual data mining<sub>1</sub>

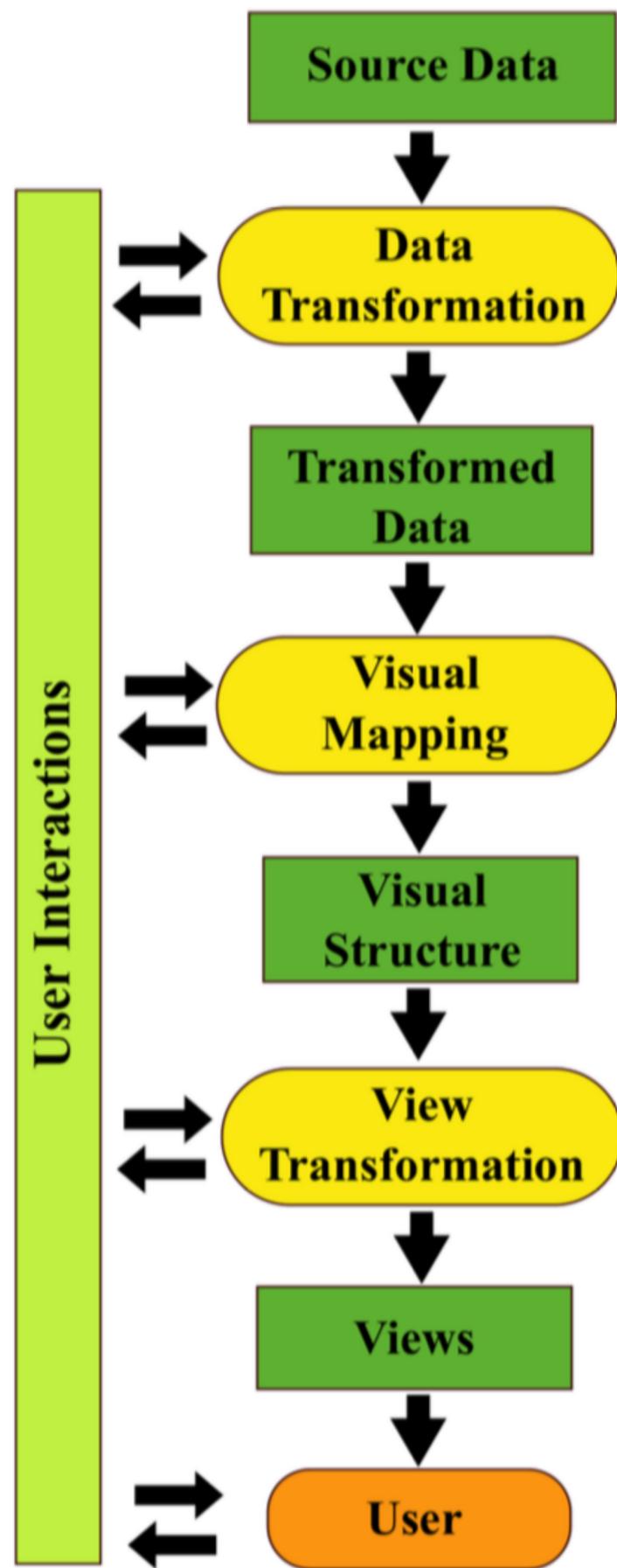
216 publications

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

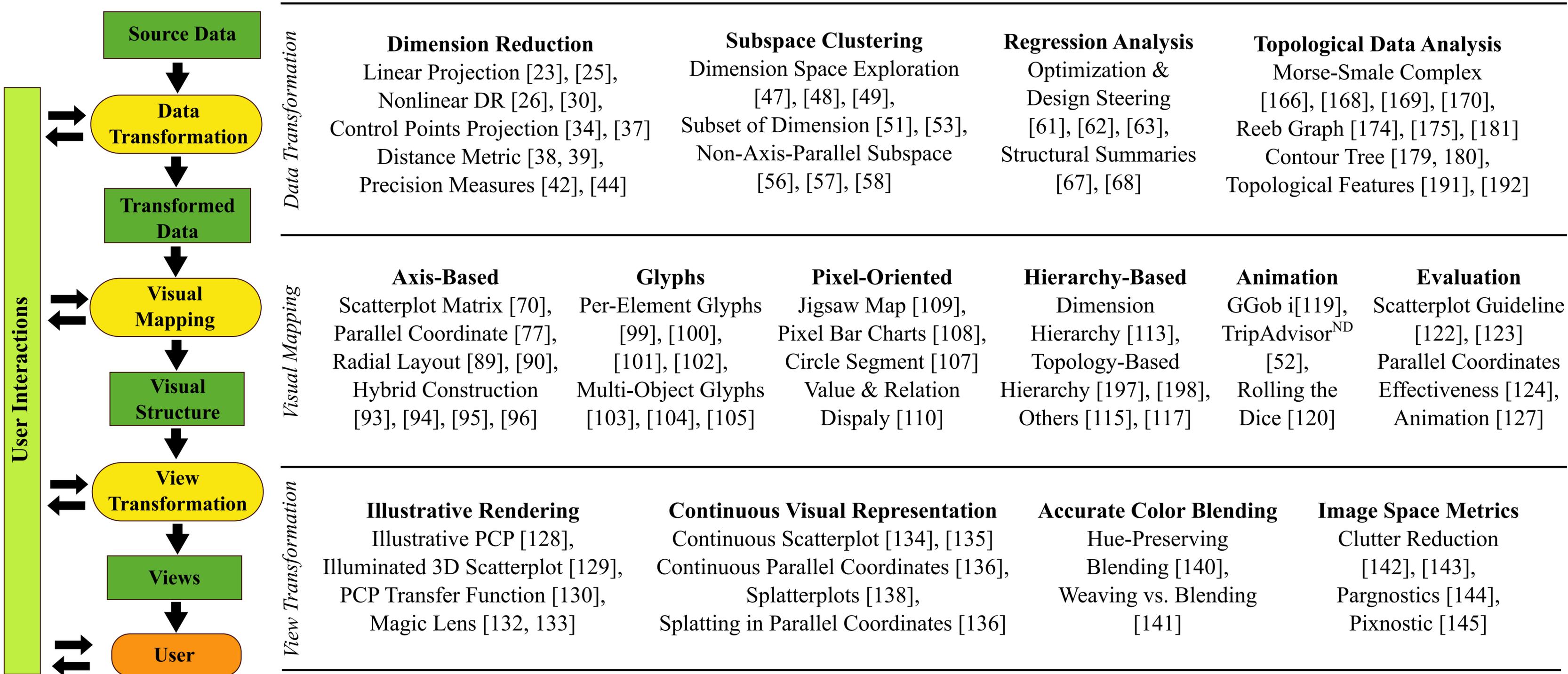
4. AnkerstBerchtoldKeim1998 [inproceedings] (1998) | PDF | DOI | Google Scholar | Google  
**Similarity clustering of dimensions for an enhanced visualization of multidimensional data**  
Ankerst, Mihael Berchtold, Stefan Keim, Daniel A  
**Abstract:** The order and arrangement of dimensions (variates) 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

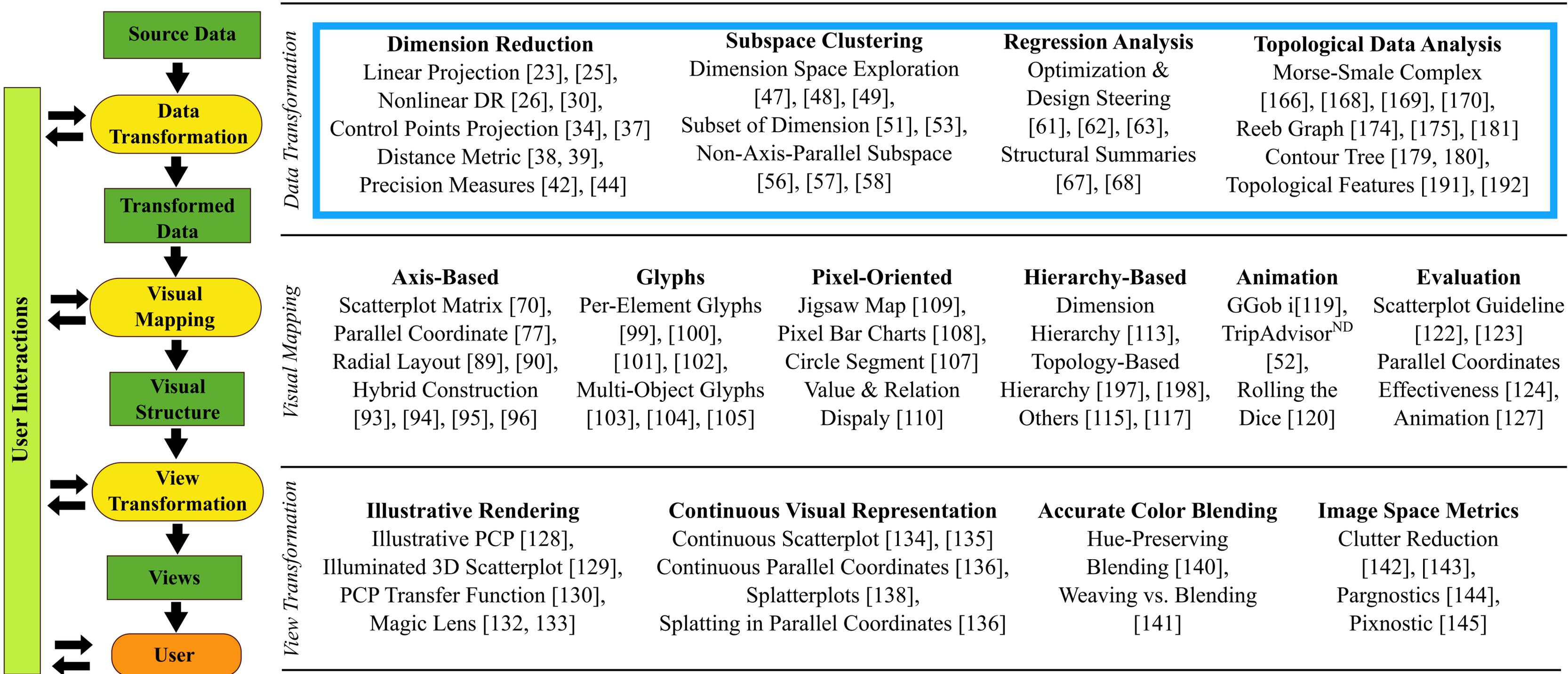
6. ArterodeOliveiraLevkowitz2004 [inproceedings] (2004) | PDF | DOI | Google Scholar | Google  
**Uncovering Clusters in Crowded Parallel Coordinates Visualizations**  
Artero, A.O. de Oliveira, M.C.F. Levkowitz, H.  
**Abstract:** The one-to-one strategy of mapping each single data item into a graphical 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**

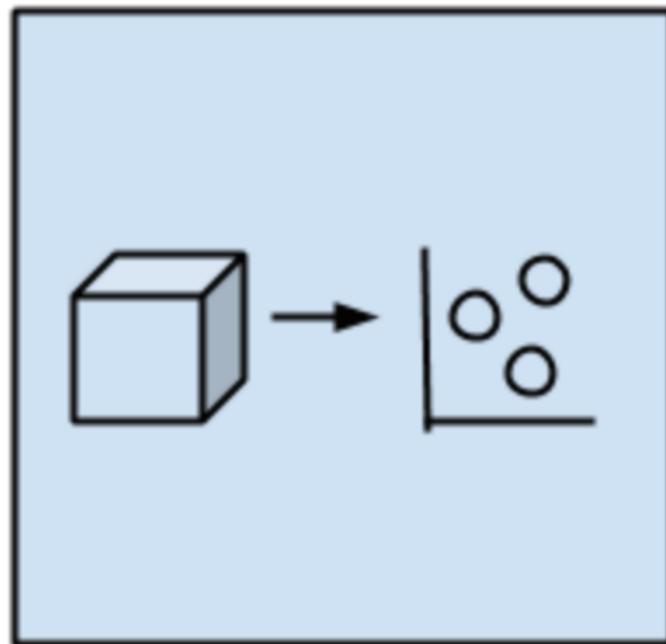
Optimization &  
Design Steering  
[61], [62], [63],  
Structural Summaries  
[67], [68]

## **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...



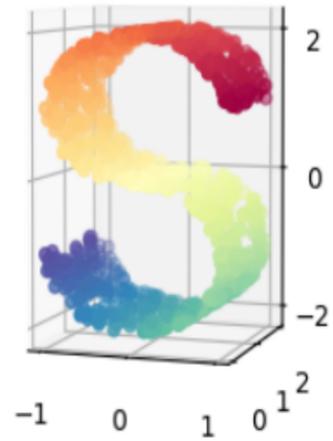
Dimensional Reduction  
Algorithms

- 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)
  - ...

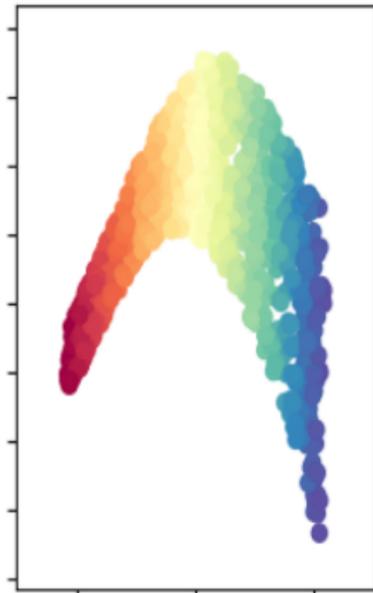
# Linear vs nonlinear DR

- 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)

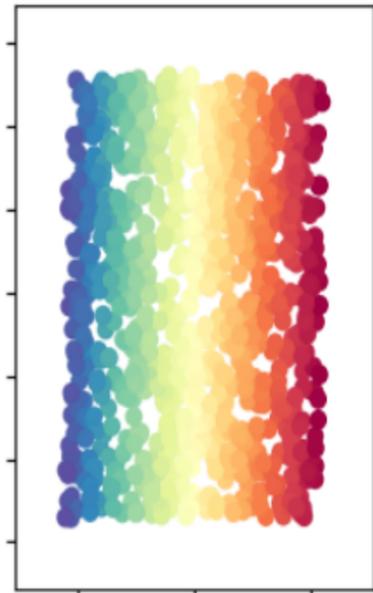
Manifold Learning with 1000 points, 10 neighbors



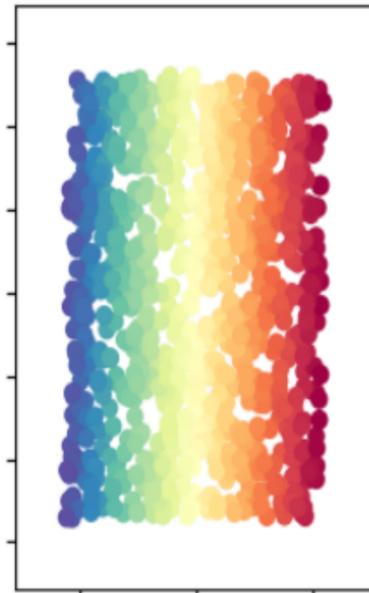
LLE (0.23 sec)



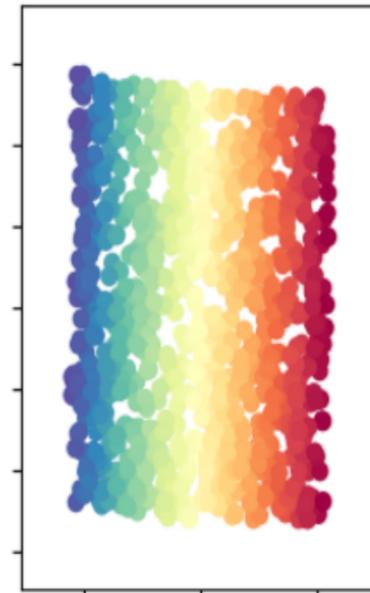
LTSA (0.37 sec)



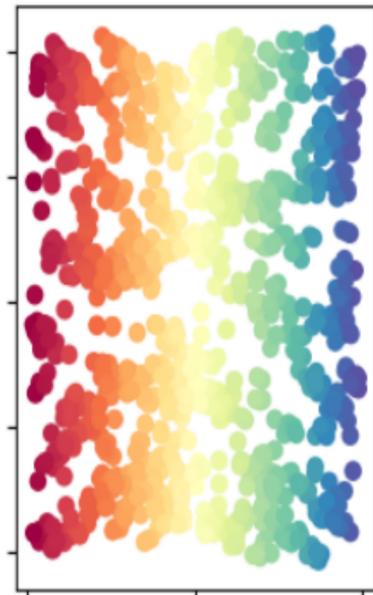
Hessian LLE (0.52 sec)



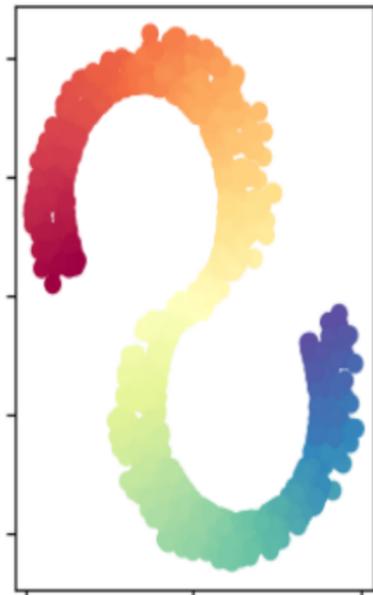
Modified LLE (0.43 sec)



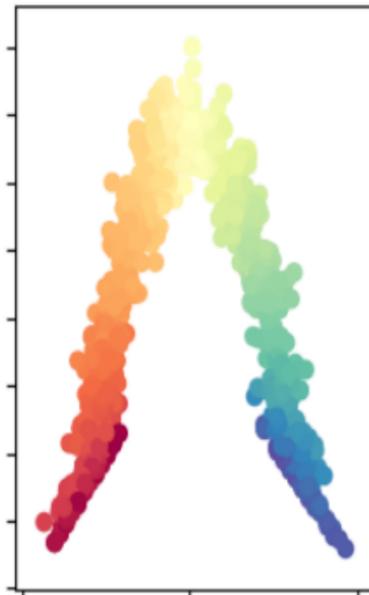
Isomap (0.46 sec)



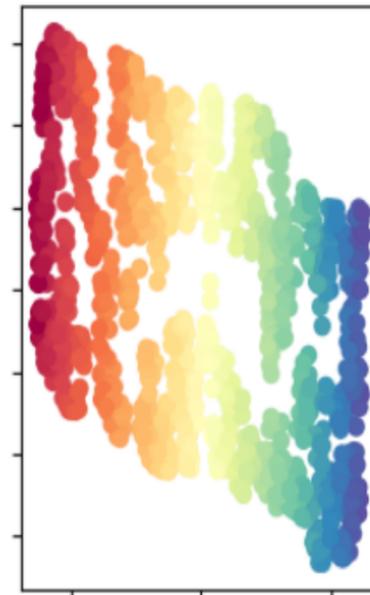
MDS (2.1 sec)



SpectralEmbedding (0.22 sec)

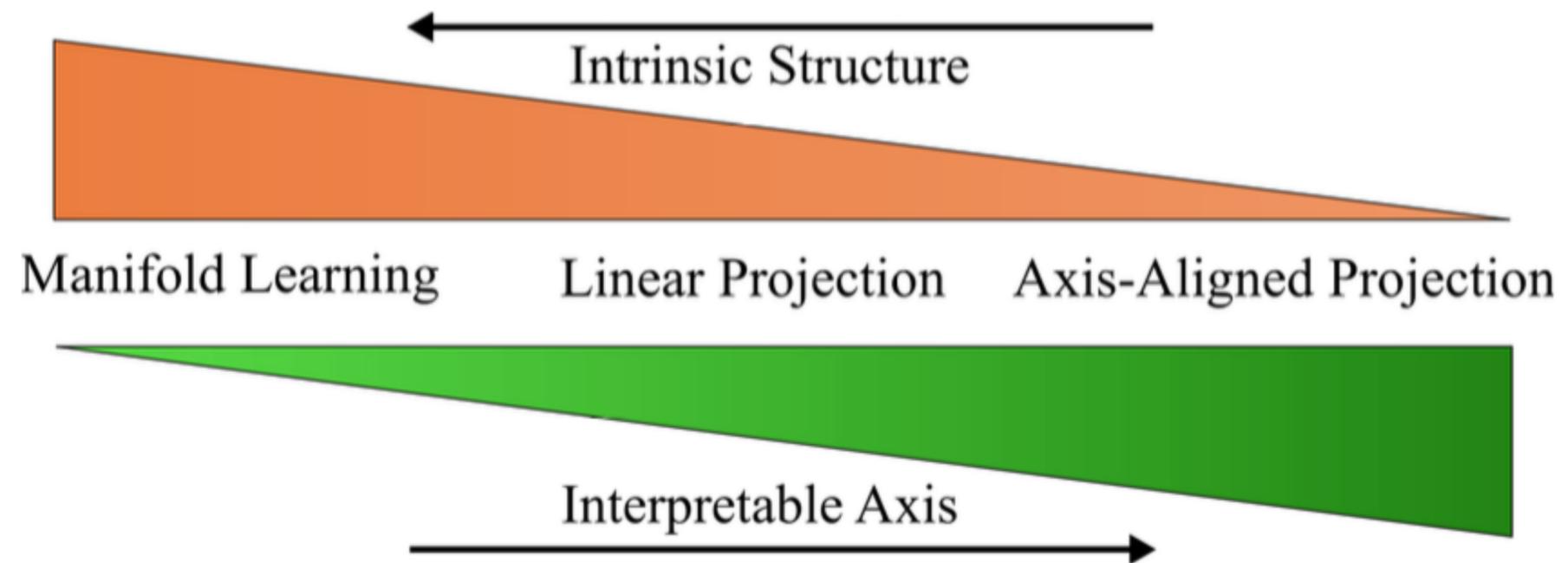


t-SNE (17 sec)



# 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

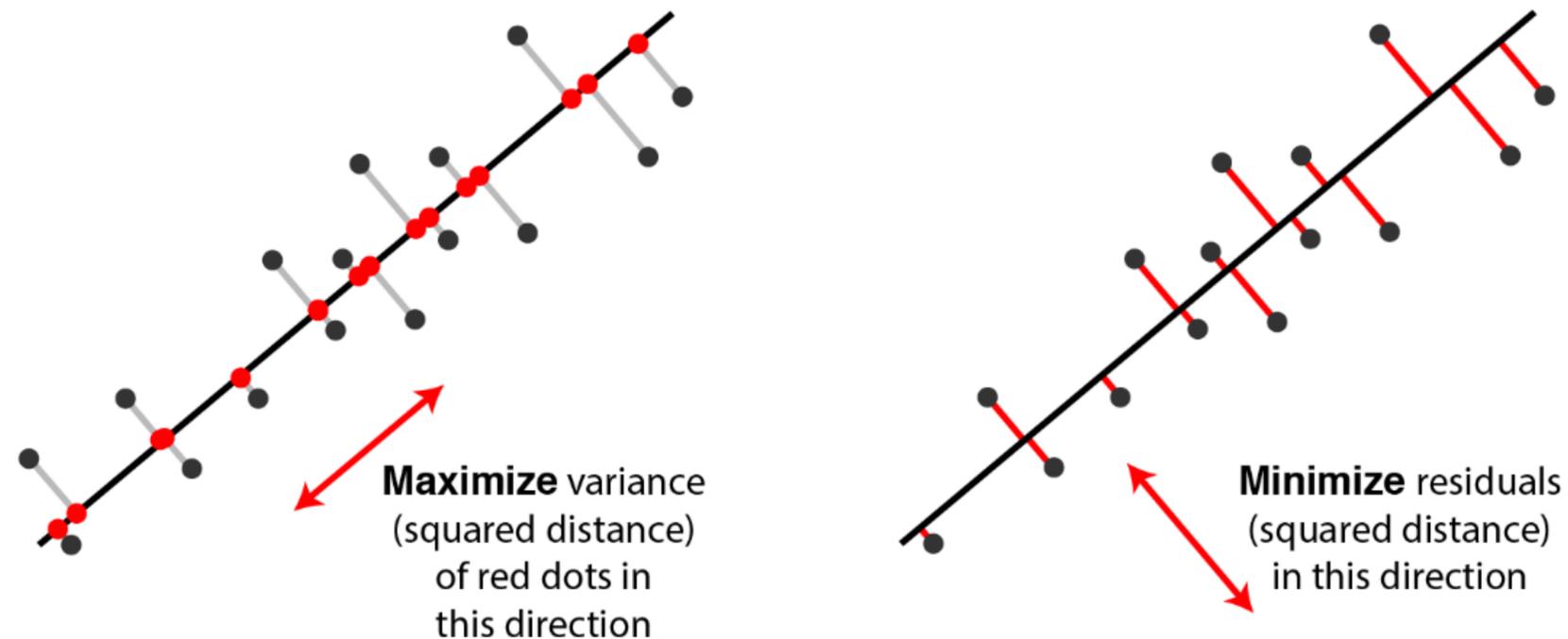
# Vis + DR: PCA

A case study with a linear DR method

# Three interpretation of PCA

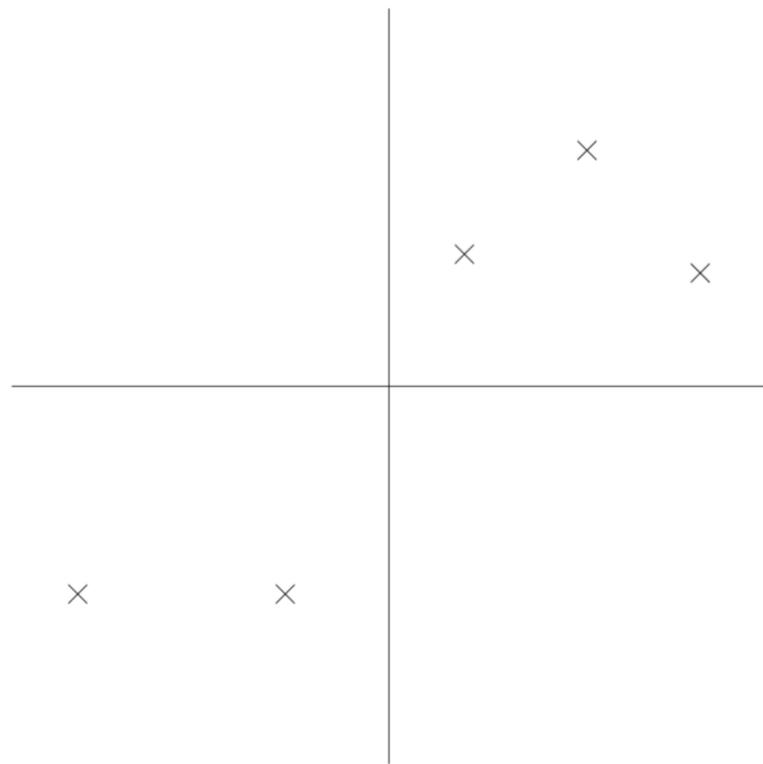
PCA can be interpreted in 2 different ways:

- 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.

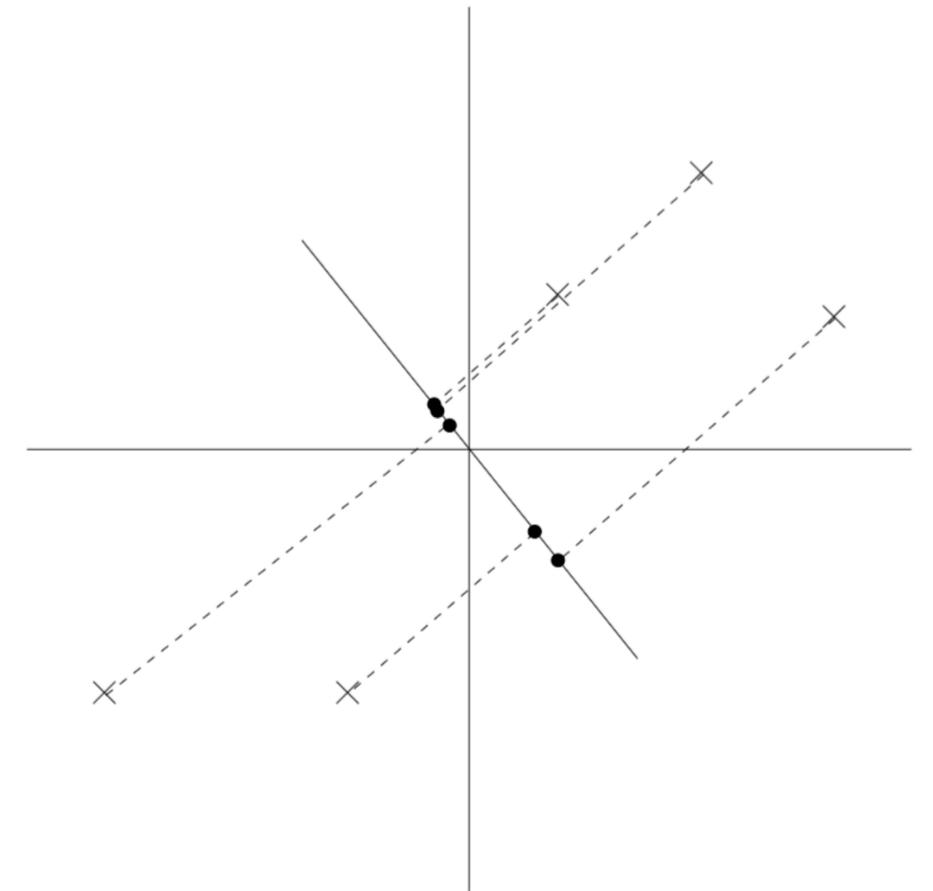


Two equivalent views of principal component analysis.

# PCA at a glance

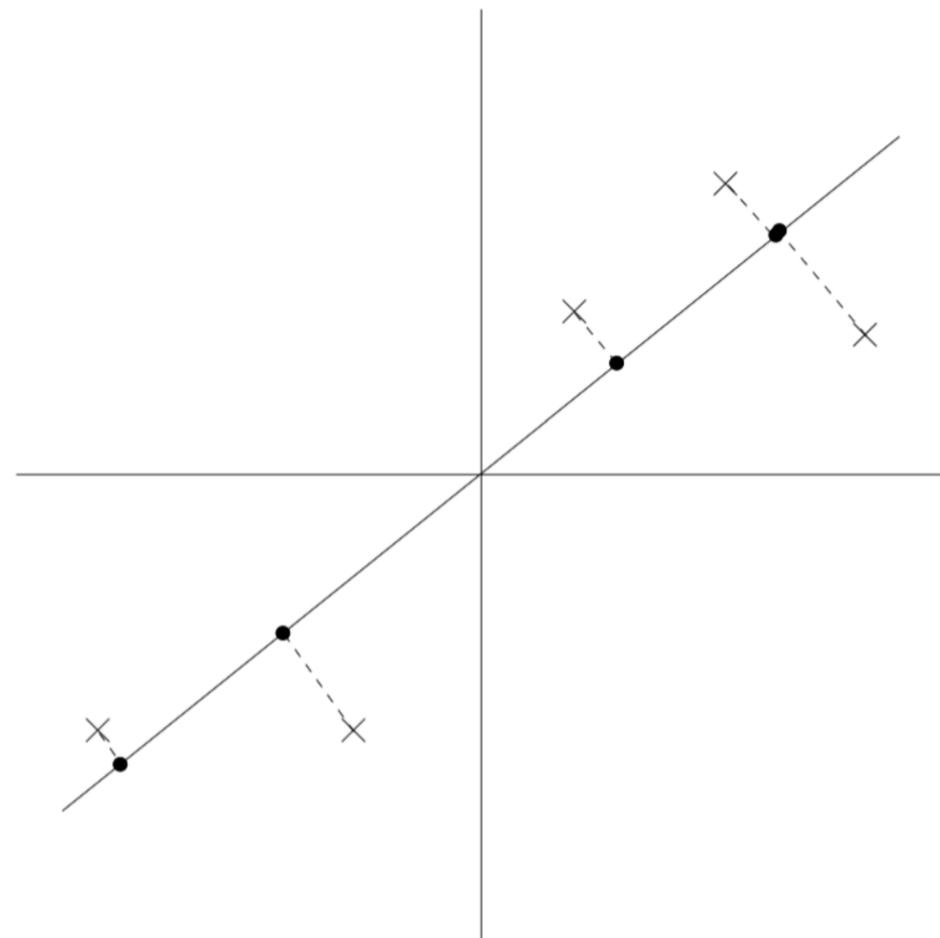


Data after normalization



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  $\lambda$ ?

# 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)
- $\Sigma$  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](http://www.uta.fi/sis/mtt/mtts1-dimensionality_reduction/drv_lecture10.pdf)

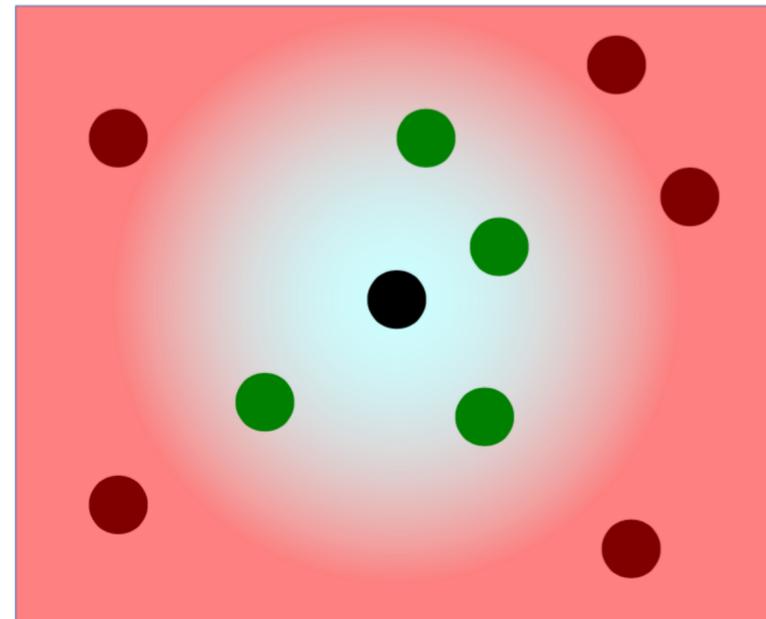
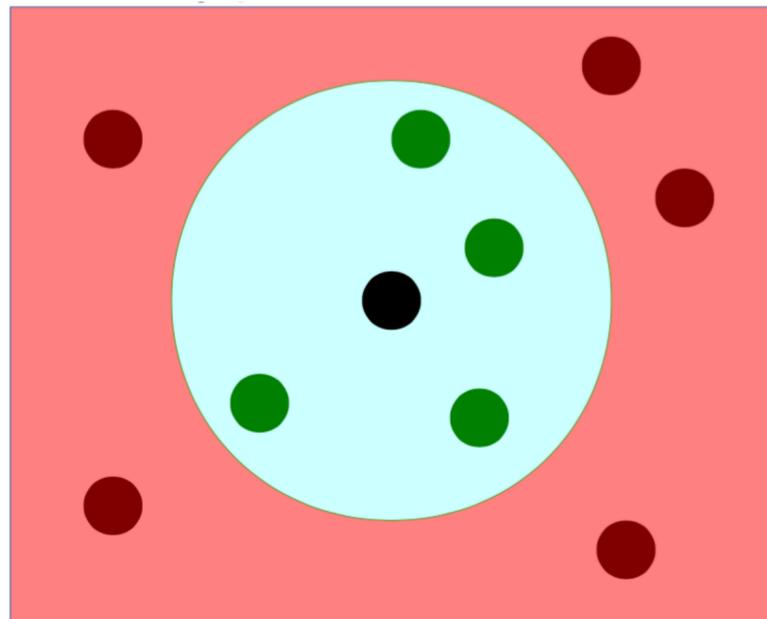
# DR: preserving distances

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

- 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

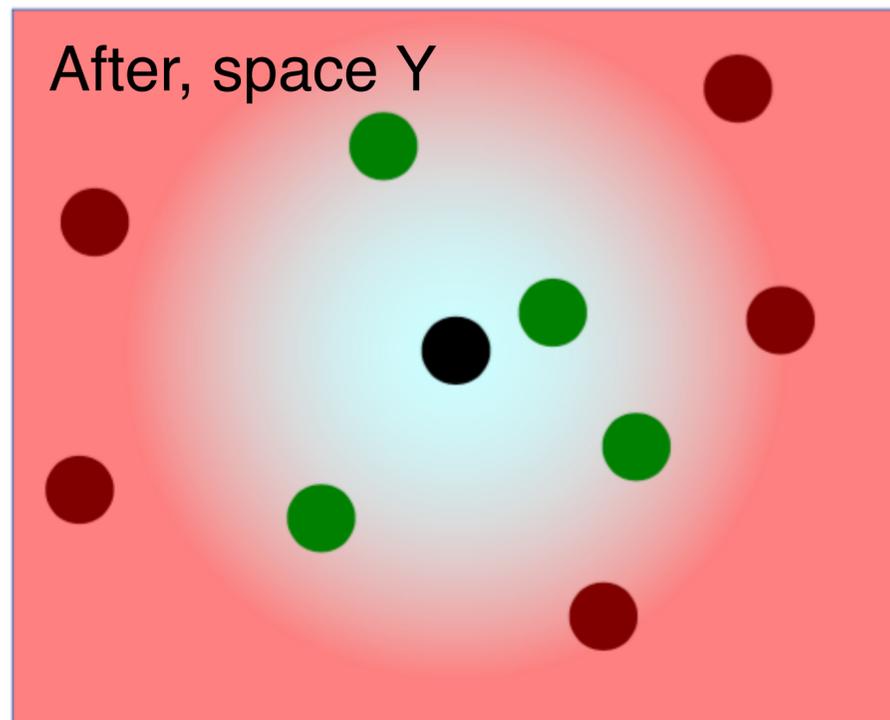
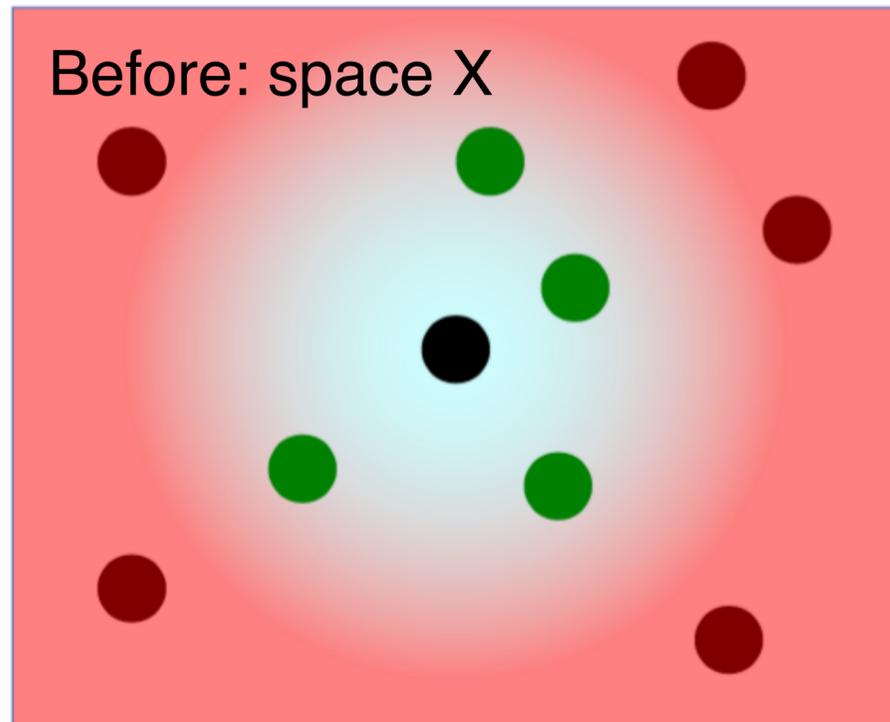


# Probabilistic neighborhood

- 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:  $\sigma_i$

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

- 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

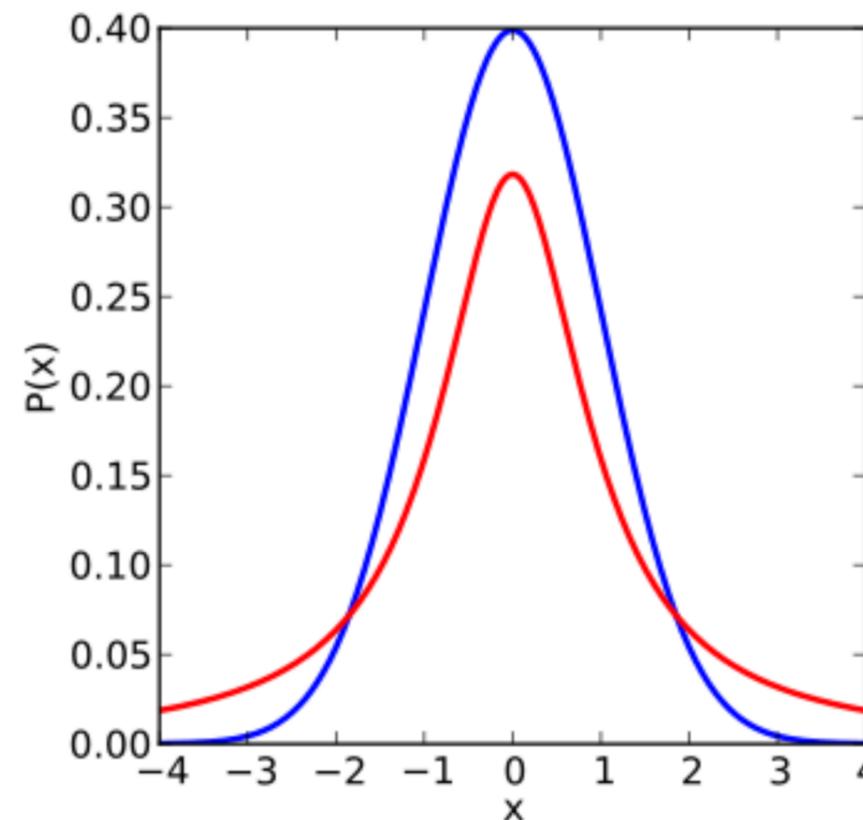
$$\frac{\partial C}{\partial y_i} = 2 \sum_j (y_i - y_j) (p_{ij} - q_{ij} + p_{ji} - q_{ji})$$

# 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

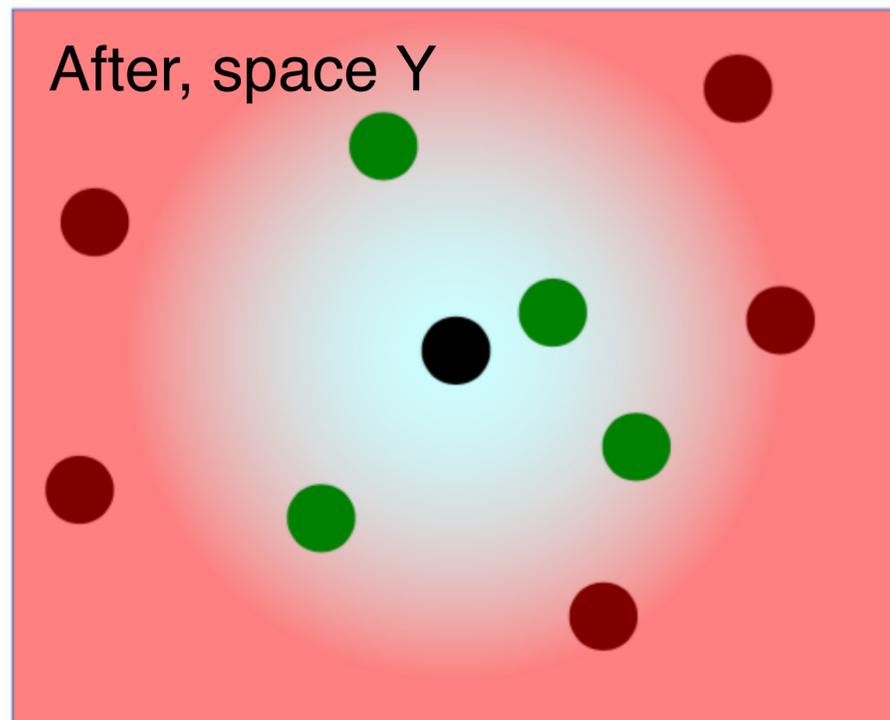
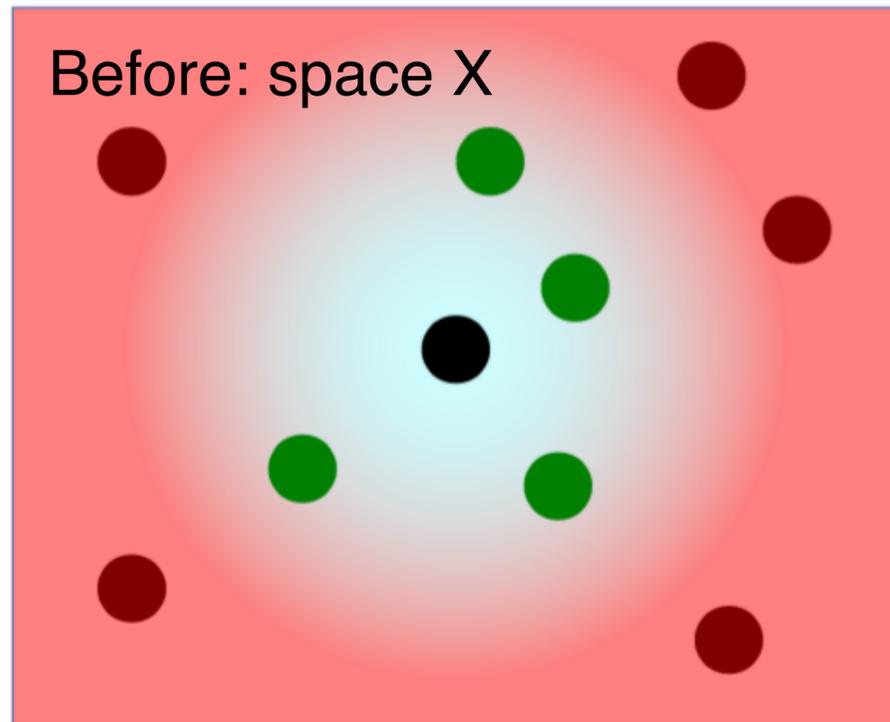
- Avoids crowding problem by using a more heavy-tailed neighborhood distribution in the low-dim output space than in the input space.
- Neighborhood probability falls off less rapidly; less need to push some points far off and crowd remaining points close together in the center.
- Use student-t distribution 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:

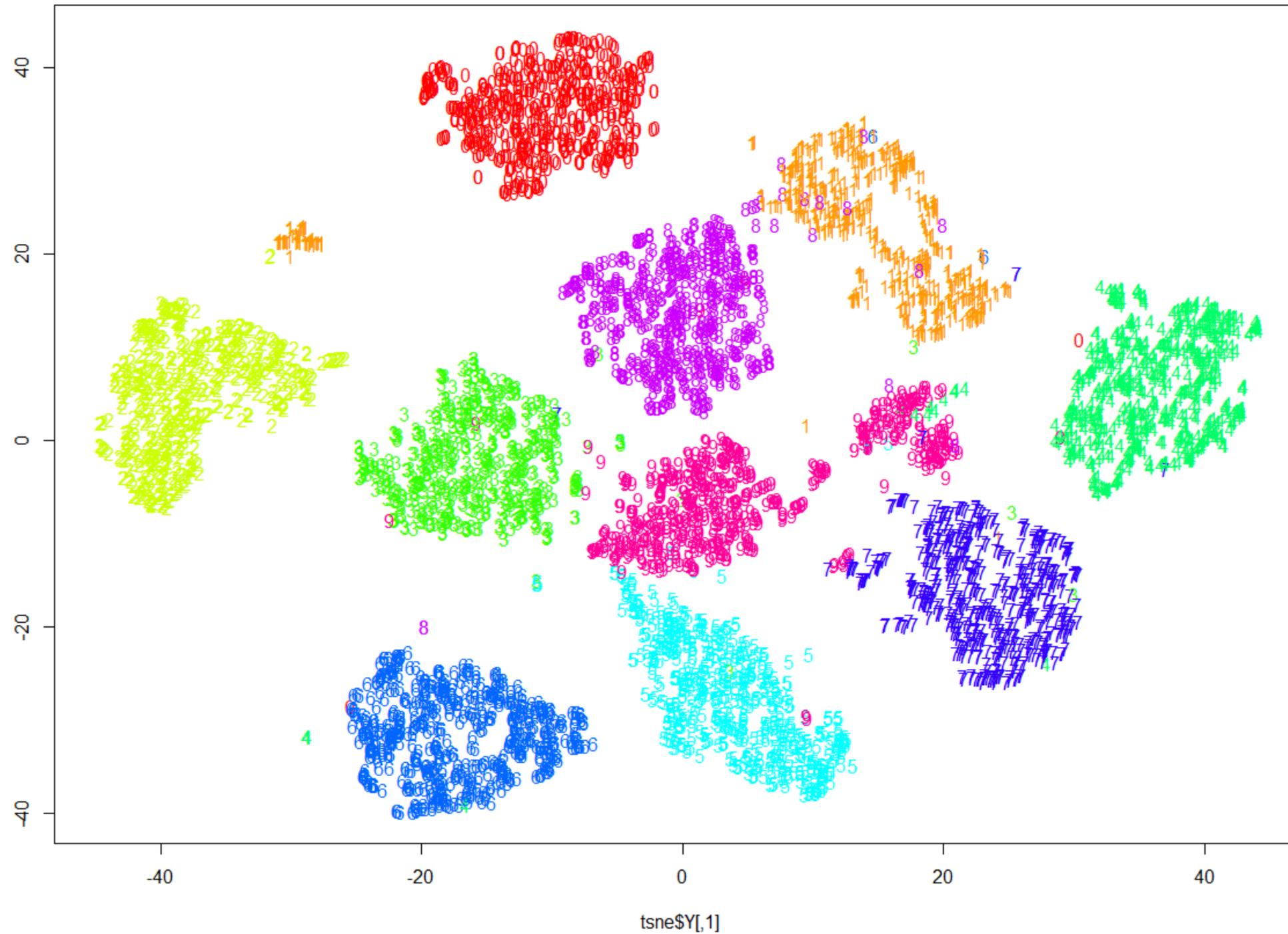
Probability to be picked as a neighbor in space X (input coordinates)

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

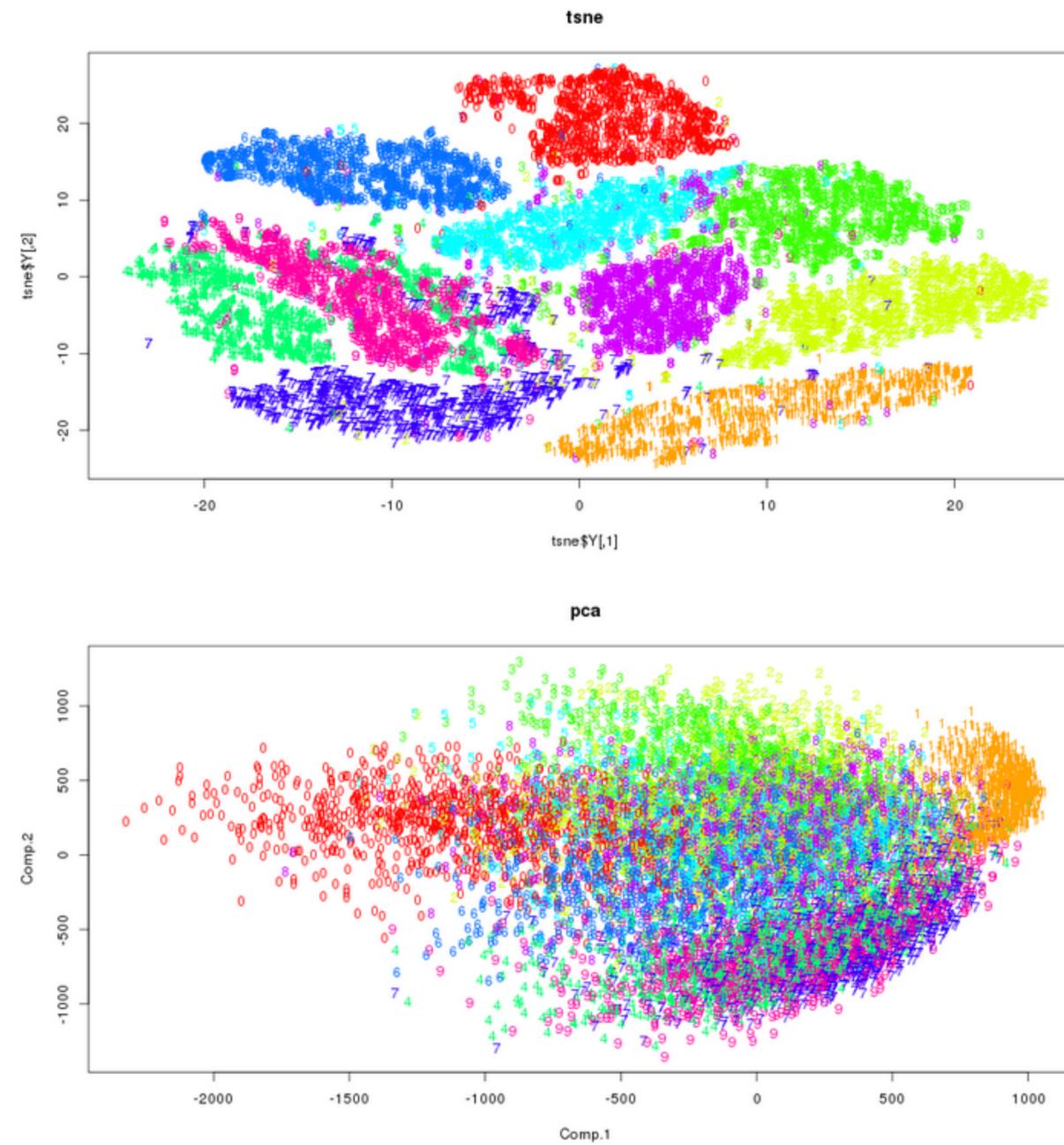
Probabilistic **output** neighborhood:

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

# Classic t-SNE result



# t-SNE vs PCA



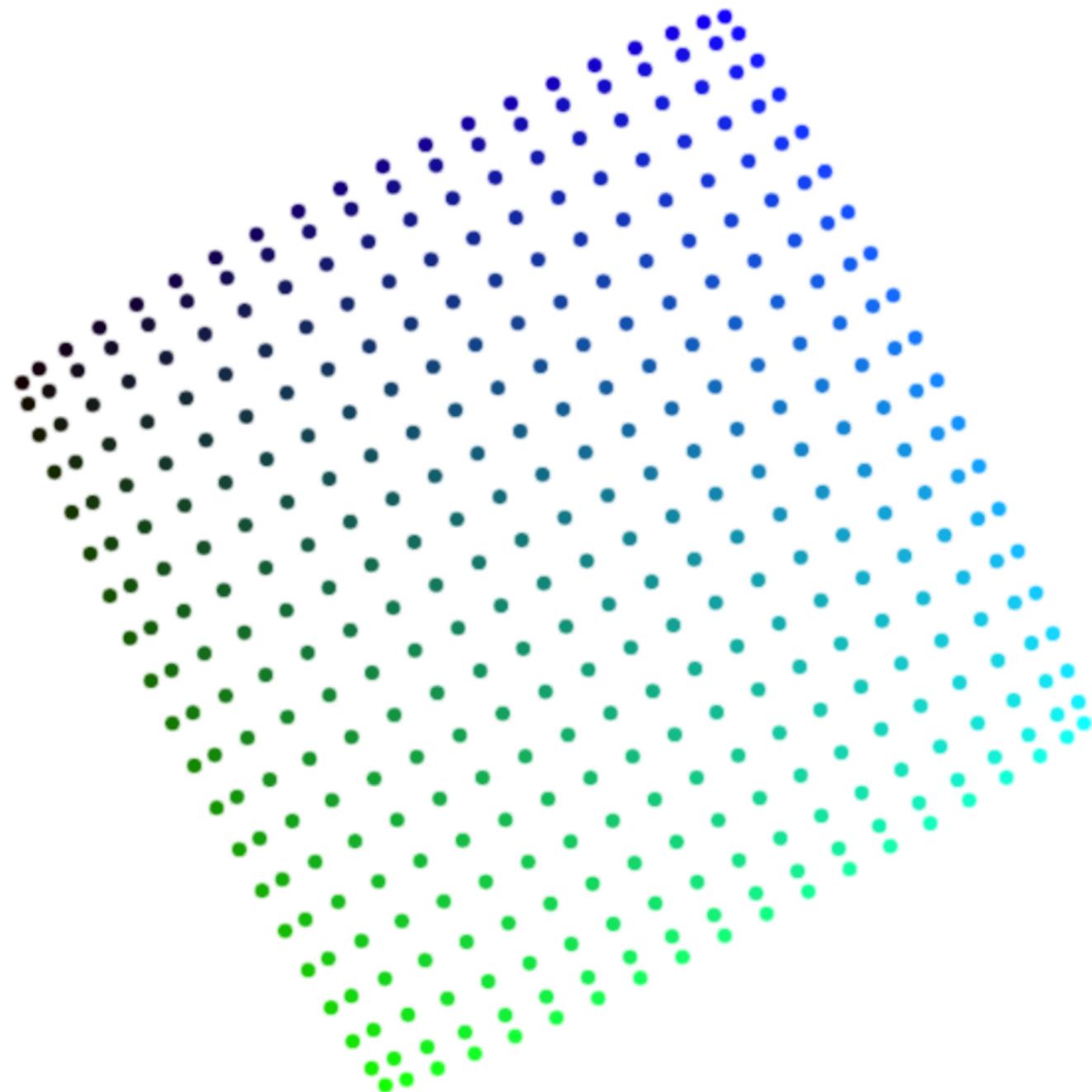
# t-SNE

- 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](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>
- UMAP:
  - <https://umap-learn.readthedocs.io/en/latest/>
- 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](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](mailto:beiwang@sci.utah.edu)

# CREDITS

Special thanks to all people who made and share these awesome resources for free:

- ☐ Presentation template designed by [Slidesmash](#)
- ☐ Photographs by [unsplash.com](#) and [pexels.com](#)
- ☐ 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

