Machine Learning, Spring 2020

Clustering and Dimensionality Reduction

Reading Assignment: Chapter 8

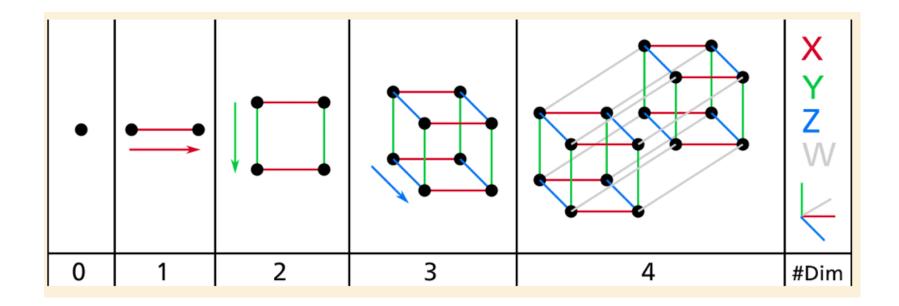
Python tutorial: http://learnpython.org/

TensorFlow tutorial: https://www.tensorflow.org/tutorials/

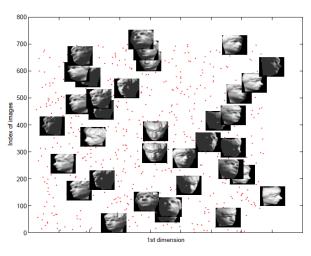
PyTorch tutorial: https://pytorch.org/tutorials/

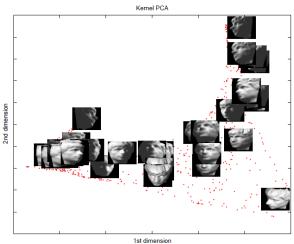
Acknowledge: The slides are partially referred to coursera online machine learning course by Prof. Andrew Ng, and NYU machine learning course. All copyrights owned by original authors.

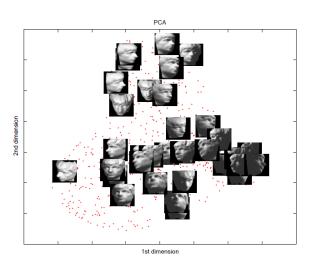
Data Examples

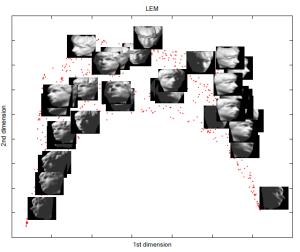


Data Examples









Curse of Dimensionality

• Cursed phenomena occur in domains such as numerical analysis, sampling, combinatorics, machine learning, data mining and databases. The common theme of these problems is that when the dimensionality increases, the volume of the space increases so fast that the available data become sparse. This sparsity is problematic for any method that requires statistical significance. In order to obtain a statistically sound and reliable result, the amount of data needed to support the result often grows exponentially with the dimensionality. Also, organizing and searching data often relies on detecting areas where objects form groups with similar properties; in high dimensional data, however, all objects appear to be sparse and dissimilar in many ways, which prevents common data organization strategies from being efficient.

Why dimensionality reduction (DR)?

- Some features may be irrelevant
- We want to visualize high dimensional data
- "Intrinsic" dimensionality may be smaller than the number of features

DR Models

- Principal Component Analysis (PCA)
- Kernel PCA
- Locally Linear Embedding (LLE)
- Laplacian Eigenmaps (LEM)
- Multidimensional Scaling (MDS)
- ISOMAP
- Semidefinite Embedding (SDE)
- Unified Framework

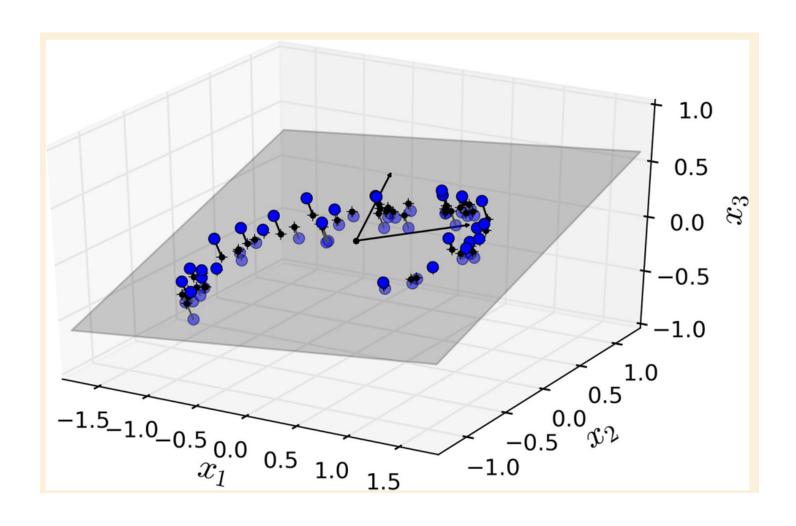
DR Models

- Linear methods
 - Principal component analysis (PCA)
 - Multidimensional scaling (MDS)
 - Independent component analysis (ICA)
- Nonlinear methods
 - Kernel PCA
 - Locally linear embedding (LLE)
 - Laplacian eigenmaps (LEM)
 - Semidefinite embedding (SDE)

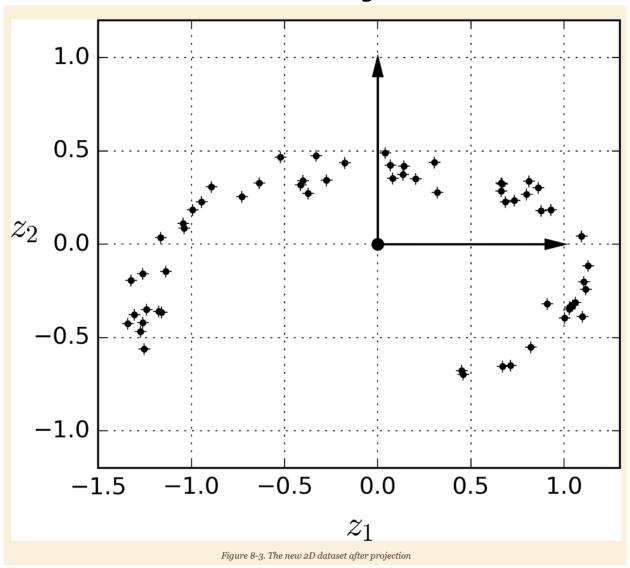
DR Approaches

- Projection
- Manifold Learning

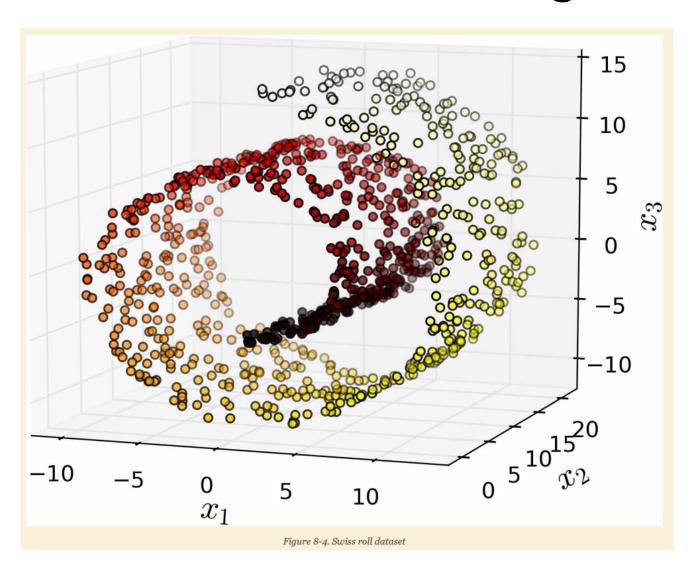
Projection



After Projection



Manifold Learning



Projection Vs Unrolling

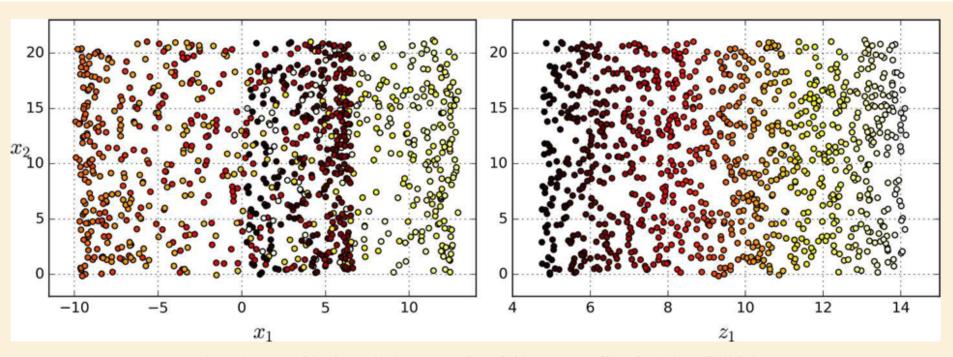


Figure 8-5. Squashing by projecting onto a plane (left) versus unrolling the Swiss roll (right)

Projection Algorithm: PCA

 Principal Component Analysis (PCA) is by far the most popular dimensionality reduction algorithm. First it identifies the hyperplane that lies closest to the data, and then it projects the data onto it.

Key Mechanism: Preserving the Variance of data

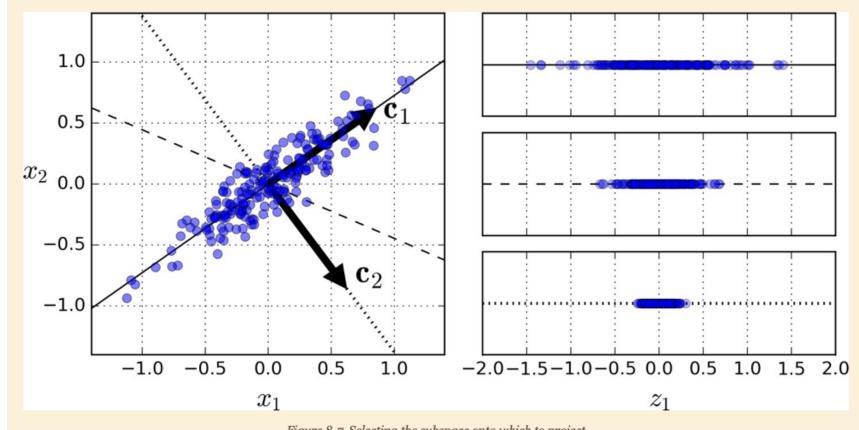


Figure 8-7. Selecting the subspace onto which to project

Preserving the Variance

- Before you can project the training set onto a lower-dimensional hyperplane, you first need to choose the right hyperplane. For example, a simple 2D dataset is represented on the left of Figure above, along with three different axes (i.e., one-dimensional hyperplanes). On the right is the result of the projection of the dataset onto each of these axes. As you can see, the projection onto the solid line preserves the maximum variance, while the projection onto the dotted line preserves very little variance, and the projection onto the dashed line preserves an intermediate amount of variance.
- It seems reasonable to select the axis that preserves the maximum amount of variance, as it will most likely lose less information than the other projections. Another way to justify this choice is that it is the axis that minimizes the mean squared distance between the original dataset and its projection onto that axis.

Principal Components

• PCA identifies the axis that accounts for the largest amount of variance in the training set. In Figure 8-7, it is the solid line. It also finds a second axis, orthogonal to the first one, that accounts for the largest amount of remaining variance. In this 2D example there is no choice: it is the dotted line. If it were a higher-dimensional dataset, PCA would also find a third axis, orthogonal to both previous axes, and a fourth, a fifth, and so on as many axes as the number of dimensions in the dataset. The unit vector that defines the i th axis is called the i th principal component (PC). In Figure 8-7, the 1 st PC is c 1 and the 2 nd PC is c 2. In Figure 8-2 the first two PCs are represented by the orthogonal arrows in the plane, and the third PC would be orthogonal to the plane (pointing up or down).

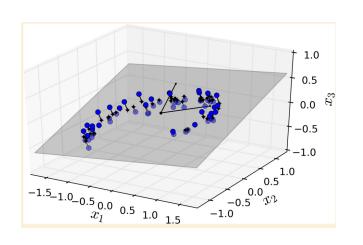
PCA Algorithm

- PCA algorithm:
 - 1. X ← Create N x d data matrix, with one row vector
 x_n per data point
 - -2. X subtract mean x from each row vector x_n in X
 - 3. Σ ← covariance matrix of X
 - Find eigenvectors and eigenvalues of Σ (through a standard matrix factorization Singular Value Decoposition (SVD))
 - PC's ← the M eigenvectors with largest eigenvalues

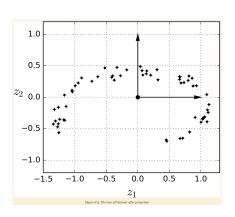
Projecting Down to d Dimensions

 Once we have identified all the principal components, you can reduce the dimensionality of the dataset down to d dimensions by projecting it onto the hyperplane defined by the first d principal components.

Projection:
$$\mathbf{X}_{d\text{-proj}} = \mathbf{X}\mathbf{W}_{d}$$

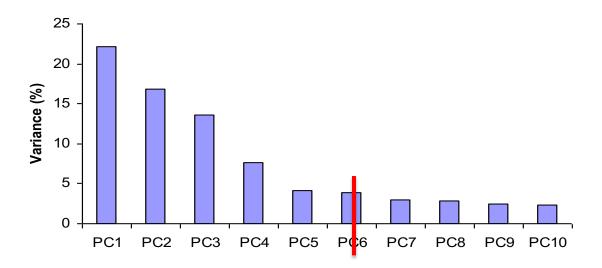






How many components?

- Distribution of Eigen-values (energy percentage)
- Choose the first few eigen-vectors to cover 80-90% of the variance of the data



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Kernel PCA

- KPCA algorithm:
 - 0: Kernel Tricks: Nonlinear mapping

$$\Phi: x \to \mathcal{H} \qquad x \mapsto \Phi(x)$$

- 1. Update X with new mapped X \leftarrow $\Phi(x)$
- 2. X ← Create N x d data matrix, with one row vector
 x_n per data point
- 3. X subtract mean x from each row vector x_n in X
- -4. Σ ← covariance matrix of X
- Find eigenvectors and eigenvalues of Σ (through a standard matrix factorization Singular Value Decoposition (SVD))
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Kernel PCA

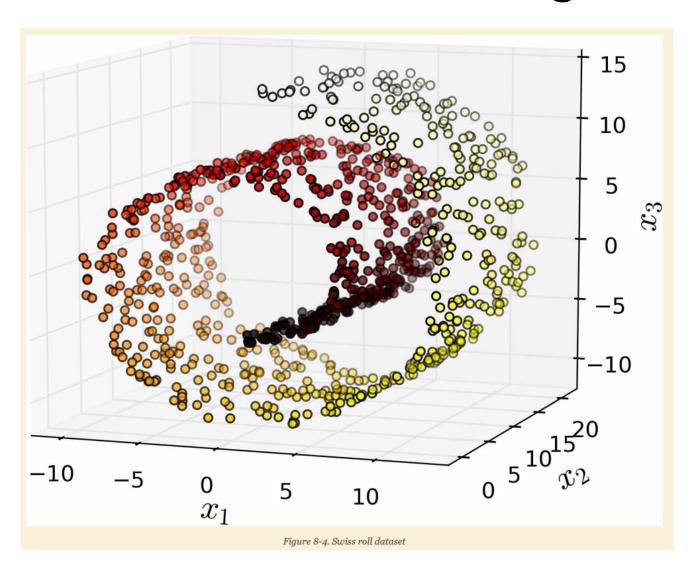
- History: S. Mika et al, NIPS, 1999
- Data may lie on or near a nonlinear manifold, not a linear subspace
- Find principal components that are nonlinearly to the input space via nonlinear mapping
- Objective
- Solution found by SVD:
 U contains the eigenvectors of

Approach: Manifold Learning

DR Models

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Manifold Learning



Projection Vs Unrolling

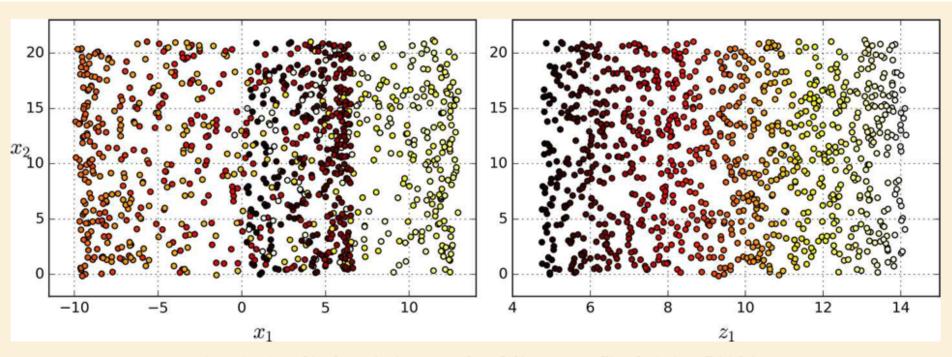


Figure 8-5. Squashing by projecting onto a plane (left) versus unrolling the Swiss roll (right)

Local Linearly Embedding

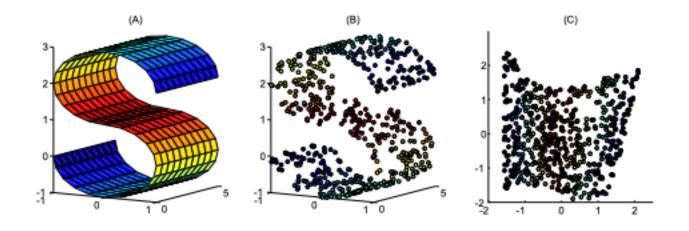
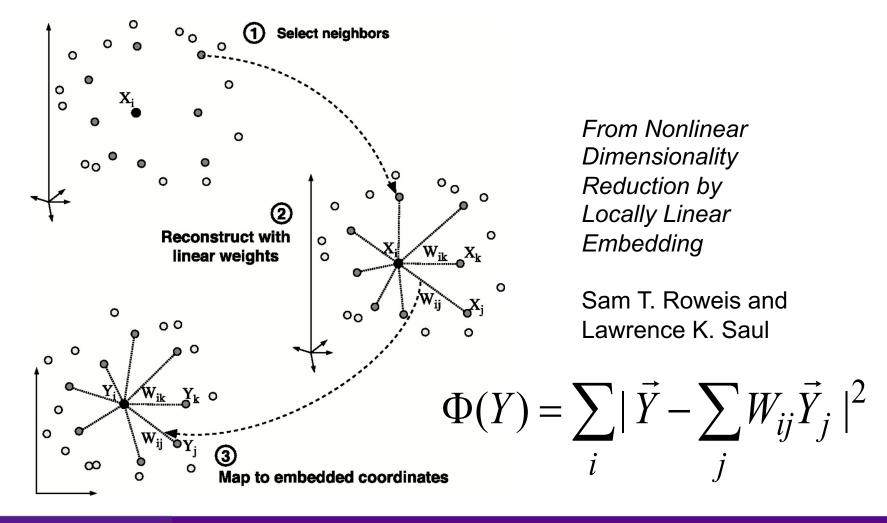


Figure 1: The problem of nonlinear dimensionality reduction, as illustrated for three dimensional data (B) sampled from a two dimensional manifold (A). An unsupervised learning algorithm must discover the global internal coordinates of the manifold without signals that explicitly indicate how the data should be embedded in two dimensions. The shading in (C) illustrates the neighborhood-preserving mapping discovered by LLE.

Reference: An Introduction to Locally Linear Embedding, https://cs.nyu.edu/~roweis/lle/papers/lleintro.pdf https://statweb.stanford.edu/~tibs/sta306bfiles/isomap.pdf

Fit locally, Think Globally



Locally Linear Embedding (LLE)

- History: S. Roweis and L. Saul, Science, 2000
- Procedure
 - 1. Identify the neighbors of each data point
 - 2. Compute weights that best linearly reconstruct the point from its neighbors

$$\min_{\mathbf{w}} \sum_{i=1}^{N} \|\mathbf{x}_{i} - \sum_{j=1}^{k} w_{ij} \mathbf{x}_{N_{i}(j)}\|^{2}$$

3. Find the low-dimensional embedding vector which is best reconstructed by the weights determined in Step 2

$$\min_{Y} \sum_{i=1}^{N} \|\mathbf{y}_{i} - \sum_{j=1}^{k} w_{ij} \mathbf{y}_{N_{i}(j)}\|^{2} \iff \min_{Y} \operatorname{tr}(Y^{\top}YL) \quad \text{Centering Y with unit variance}$$

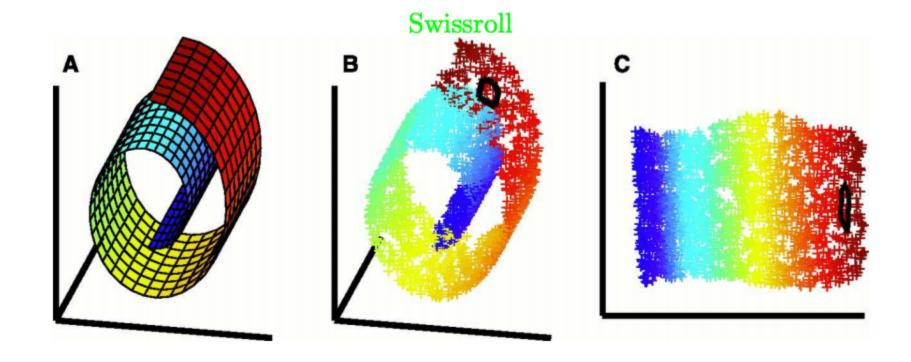
Solution:

• Minimize

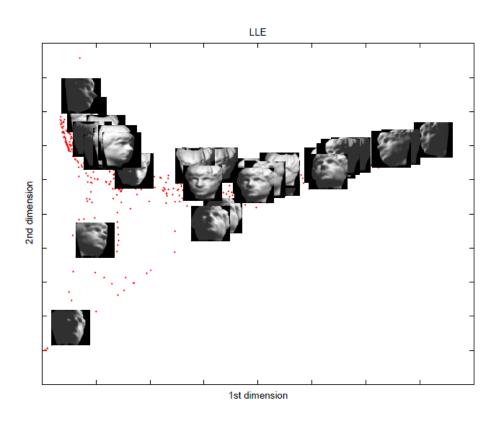
$$tr[(Y - WY)^T(Y - WY)] = tr[Y(I - W)^T(I - W)Y]$$

- W is $N \times N$; Y is $N \times d$, for some small d < p.
- solutions Y are the bottom eigenvectors of $M = (I W)^T (I W)$.
- They also assume 1^TY = 0 (i.e. they are centered at the origin), and (1/N)Y^TY = I, the identity matrix in d dimensions. This means they discard the smallest eigenvector of M and keep the next d.

After Embedding



LLE Example



ISOMAP Idea?

