CS456: Machine Learning Unsupervised Dimensionality Reduction

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- To understand issues found in high-dimensional space
- To understand what principal component in PCA is
- Be able to apply PCA for realworld problem

- Curse of Dimensionality
- Principal Component Analysis (PCA)

- The working of machine learning algorithms depends in some way on the geometry of data
 - lengths of vectors, distances, angles
- High dimensional geometry is different from low dimensional geometry

What happens in HD? [1/2]

Concentration of norms: generate points in m-dimensional space and measure their lengths



Figure: Credit: A.Kaban, CS-Bham

What happens in HD? [2/2]

Near-orthogonality: generate points in *m*-dimensional space and measure their angles



d × 10 ~

Figure: Credit: A.Kaban, CS-Bham

- We can see from the plots that
 - As *m* increases, any two random vectors end up being orthogonal to each other
 - As *m* increases, any random vectors ends up having about the same length
- We also need a lot (exponentially) more data to cover the space as *m* increases
- Training time is also increased significantly as *m* grows

- Surprisingly, high dimensionality makes data more linearly separable
- Think of kernel method, or feature learnt by convolutional neural networks
- It is easier for algorithm to find separating hyperplane
 - providing that there's no noise in data

Dimensionality reduction approaches

Feature selection

- Find subset of features
- Feature projection (feature extraction)
 - learn a function $\phi(\cdot)$ which transforms data from HD to lower dimensional space
 - In general, we aim at minimising reconstruction error

$$\epsilon_{recon} = ||X - \phi^{-1}(\phi(X))||^2$$

Principal Component Analysis (PCA)

- An unsupervised algorithm for dimensionality reduction
- Reduce dimensionality of the data while trying to preserve data structure

Find low-dimensional projection with largest spread



Figure: Applied Multivariate Statistics: ETZ

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CS456: Machine Learning

Toy data

Assuming data is in 2D space



Figure: Applied Multivariate Statistics: ETZ

• Normally, data lives In the standard basis defined by two basis vectors {[0 1], [1 0]} 1



 $^1\mathsf{A}$ basis is a set of linearly independent vectors that can represent any vector in a given vector space

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Other basis systems

• Imagine using another valid basis system



• Observe: data is more spread along one of the new basis directions (orange vector)

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- Find new basis system that maximises variance in all directions
- These directions are known as Principal Components (PC)
- Hopefully we can select a few of these PCs and project the data onto this new basis

Projection onto one direction

• Projection of a vector
$$\mathbf{x} = \begin{bmatrix} x^1 \\ x^2 \\ \cdots \\ x^m \end{bmatrix}$$
 on to a vector $\mathbf{a} = \begin{bmatrix} a^1 \\ a^2 \\ \cdots \\ a^m \end{bmatrix}$ is the linear combination

$$\mathbf{a}^T \mathbf{x} = \sum_{i=1}^m a^i x^i$$

• Usually, **a** is kept as a unit vector

• To generalise one direction projection, the projection of **x** onto a set of linearly independent vectors (*basis*) A is

$$A^{T}\mathbf{x} = \begin{bmatrix} a_{1}^{1} & a_{2}^{1} & a_{3}^{1} \\ a_{1}^{2} & a_{2}^{2} & a_{3}^{2} \\ \vdots & \vdots & \vdots \\ a_{1}^{m} & a_{2}^{m} & a_{3}^{m} \end{bmatrix}^{T} \begin{bmatrix} x^{1} \\ \vdots \\ x^{m} \end{bmatrix} = \begin{bmatrix} p^{1} \\ p^{2} \\ p^{3} \end{bmatrix}$$

• **p** is a vector in 3D space , compared to the original **x** in *m*D space

- Assumed data matrix X and its projection $\mathbf{a}^T X$ are mean-centred
- The spread of the projection is

$$\sigma_{\mathbf{a}}^{2} = (\mathbf{a}^{T} X) (\mathbf{a}^{T} X)^{T}$$
$$= \mathbf{a}^{T} X X^{T} \mathbf{a}$$
$$= \mathbf{a}^{T} V \mathbf{a}$$

• We see that spread is a function of projection direction ${\bf a}$ and $m \times m$ covariance matrix V

Objective function of projection direction

- Maximising a^TVa makes no sense, because we can increase the spread by multiplying a by some large number
- We have to impose size constraint on \mathbf{a} , e.g., $\mathbf{a}^T \mathbf{a} = 1$
- We then arrive at an objective function

$$u = \mathbf{a}^T V \mathbf{a} - \lambda (\mathbf{a}^T \mathbf{a} - 1)$$

- $\lambda > 0$ is a parameter imposing the size constraint ²
 - Think of λ like the C parameter in SVM

²Its name is Lagrange multiplier often employed in constrained optimisation

- Objective function is convex, calculus helps finding stationary point
- Differentiating the objective function w.r.t **a** and equating to zero

$$\frac{\partial u}{\partial \mathbf{a}} = 2V\mathbf{a} - 2\lambda\mathbf{a} = 0 \tag{1}$$
$$V\mathbf{a} = \lambda\mathbf{a} \tag{2}$$

- Eq.(2) is one type of Linear system of equations called the Characteristic Equations
- If we can solve the system for **a** we will have maximum spread direction

$$V\mathbf{a} = \lambda \mathbf{a}$$

- For an $m \times m$, real and symmetric matrix V, there are m possible solution vectors
- For symmetric matrices, eigenvectors for distinct eigenvalues are orthogonal
- Each of the solutions \mathbf{a}_i is known as eigenvector of V
- Each eigenvector is associated with an eigenvalue λ_i

Eigenvector problem refresher

Definition

A nonzero vector ${\bf x}$ is an eigenvector of a square matrix A if there exists a scalar λ such that

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

- λ is an eigenvalue of A associated with eigenvector ${\bf x}$
- The zero vector can not be an eigenvector even though

$$A0 = \lambda 0$$

• But $\lambda = 0$ can be an eigenvalue

- Given $A = \begin{bmatrix} 2 & 1 \\ 1 & 2 \end{bmatrix}$, find its eigenvectors and eigenvalues
- Solution:
 - 1 from definition $Ax = \lambda x$

$$2 (A - \lambda I)x = 0$$

- 3 Since x is nonzero, we know that $(A \lambda I)$ is not invertible
- 4 So determinant of $(A \lambda I)$ must be zero

$$5 |A - \lambda I| = 0$$

- The *m* eigenvectors form to a new basis system
- The eigenvector **a** with largest eigenvalue is the projection direction with maximum spread (most important)
- PCA selects k most important from m eigenvectors where k < m
- PCA projects dataset X onto the new basis formed by k eigenvectors

•
$$X_r = \begin{bmatrix} a_1^1 & a_2^1 & \dots & a_k^1 \\ a_1^2 & a_2^2 & \dots & a_k^2 \\ \vdots & \vdots & \vdots & \vdots \\ a_1^m & a_2^m & \dots & a_k^m \end{bmatrix}^T X$$

- Standardising X
- Calculate a covariance matrix $V = XX^T$
- Find all the eigenvectors of V
- Select k most important principal components according to eigenvalues and put it in a matrix A
- Project X onto A by calculating $X_r = A^T X$
- The reducted data is in X_r

- The structure of data can be defined as sum of spreads in all direction
- From Va_i = λ_ia_i, we see that λ_i quantifies the spread of data after projecting on a_i
- The loss in structure information by selecting only k PCs is

$$\frac{\sum_{i=k+1}^{m} \lambda_i}{\sum_{i=1}^{m} \lambda_i}$$

• Usually, we stop throwing PCs away when the loss exceeds the predefined threshold

- Computing V takes $O(nm^2)$
- PCA complexity is then $O(nm^2)$ + Complexity of solving eigenvector
- PCA can be applied to large dataset (scale well with *n*) but it does not scale well with dimensionality *m*
 - Slow for high-dimensional data

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