

Scientific Computing for Biologists

Singular Value Decomposition and Biplots

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Overview of Lecture

- Singular Value Decomposition
 - Algebra of SVD
 - Geometry of SVD
 - Relationship to Eigendecomposition
 - Applications of SVD
- Biplots
 - Simultaneous representation of rows and columns of a matrix

Hands-on Session

- SVD and Biplots in R
- SVD in Python
- Applications of SVD in R and Python
 - 'Seriation' using SVD
 - Matrix approximation and image compression using SVD

Eigendecomposition

$$\mathbf{A} = \mathbf{U}\mathbf{D}\mathbf{U}^{-1}$$

where:

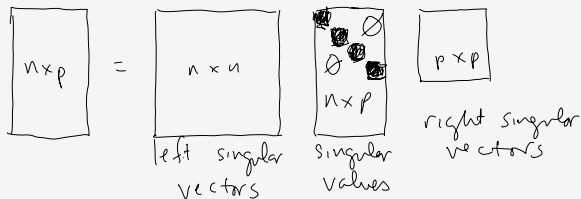
- \mathbf{U} is a matrix of eigenvectors (in columns)
- \mathbf{D} is a diagonal matrix with eigenvalues along diagonal.

when \mathbf{A} is real-valued and symmetric than \mathbf{U} is orthogonal.

Singular Value Decomposition

$$A = U S V^T \quad \text{assume } n \geq p$$

$(n \times p) \quad (n \times n) \quad (n \times p) \quad (p \times p)$



when written like this U & V are orthonormal

• sometimes written as

$$A = U S V^T = \begin{matrix} (n \times p) & (p \times p) & (p \times p) \end{matrix}$$

Diagram illustrating the compact SVD notation:

- A box labeled $n \times p$ is equal to a box labeled $n \times p$ (labeled "left singular vectors") multiplied by a box labeled $p \times p$ (labeled "singular values" and containing small circles) multiplied by a box labeled $p \times p$ (labeled "right singular vectors").

Facts about SVD

- Singular Value Decomposition is often referred to as giving the “basic structure” of a matrix
- The rank of \mathbf{A} is equivalent to the number of non-zero singular values in $\mathbf{A} = \mathbf{USV}^T$

$$\text{rank}(\mathbf{A}) \leq \min(n, p)$$

- The Euclidean norm (L_2) norm of a matrix is the relative amount it stretches a vector:

$$|\mathbf{A}|_E = \frac{|\mathbf{Ax}|}{|\mathbf{x}|}$$

The L_2 norm of \mathbf{A} is given by \mathbf{S}_{11} .

Geometric Interpretation of SVD

Any matrix, $\mathbf{A}_{n \times p}$, represents a linear transformation from $\mathbb{R}^p \mapsto \mathbb{R}^n$.

SVD can be thought of decomposing the transformation specified by \mathbf{A} into a simple form:

$$\mathbf{A} = (\text{rotation})(\text{scaling})(\text{rotation})$$

- \mathbf{U} and \mathbf{V} are orthonormal matrices \leadsto Orthonormal matrices represent rigid rotations (or rotation plus reflection)
- Diagonal matrices represent “stretching”

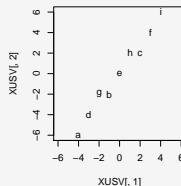
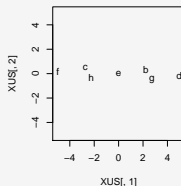
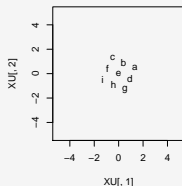
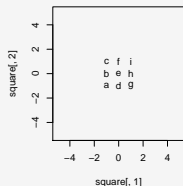
SVD Example

$$A = \begin{bmatrix} 1 & 2 \\ 3 & 4 \end{bmatrix} = U S V^T$$

$$\text{where } U = \begin{bmatrix} -0.41 & -0.91 \\ -0.91 & 0.41 \end{bmatrix} \quad S = \begin{bmatrix} 5.47 & 0 \\ 0 & 0.37 \end{bmatrix}$$

$$V^T = \begin{bmatrix} -0.58 & -0.82 \\ 0.82 & -0.58 \end{bmatrix}$$

Geometry



Relationship of SVD to Eigendecomposition

$$A = U S V^T, \text{ let } D = S S$$

$$\begin{aligned} A^T A &= (V S U^T)(U S V^T) = V S \underbrace{U^T U}_= I S V^T \\ &= V D V^T \end{aligned}$$

(since orthonormal)

so the singular values S_{ii} are $\sqrt{D_{ii}}$ where
 D_{ii} are the eigenvalues of $A^T A$

columns of V are the eigenvectors of $A^T A$

Using SVD to do PCA

let X be a mean-centered data matrix

covariance
matrix of X

$$C = \frac{1}{n} X^T X$$

By SVD we can write $X = U S V^T$

$$\begin{aligned} C &= \frac{1}{n} V S U^T U S V^T \\ &= \frac{1}{n} V S^2 V^T \end{aligned}$$

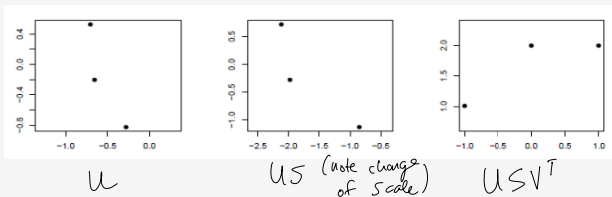
Principal Components given by columns of V

PC Scores given by $U D$

Another Way of Thinking about SVD

$$\begin{array}{lcl}
 \begin{array}{l} \text{Observations} \\ \text{in measurement} \\ \text{space} \\ A \end{array} & = & \begin{array}{l} \left(\begin{array}{l} \text{Observations} \\ \text{in PC} \\ \text{space} \end{array} \right) \\ U S \end{array} \xleftarrow{\text{rotation}} \begin{array}{l} \text{Inverse of} \\ \text{Eigenvectors} \\ V^T \end{array}
 \end{array}$$

e.g. $A = \begin{bmatrix} 1 & 2 \\ 0 & 2 \\ -1 & 1 \end{bmatrix}$



Applications of SVD

- Pseudoinverse of an arbitrary matrix
- Matrix approximation
- Motivates the Biplot and Correspondence Analysis

Pseudoinverse via SVD

The pseudoinverse of a matrix is a generalization of the concept of a matrix inverse. Only square matrices have a matrix inverse; the pseudoinverse applies to an arbitrary $n \times p$ matrix.

Given an $n \times p$ matrix \mathbf{A} find matrix \mathbf{A}^+ such that:

$$\begin{aligned}\mathbf{A}\mathbf{A}^+\mathbf{A} &= \mathbf{A} \\ \mathbf{A}^+\mathbf{A}\mathbf{A}^+ &= \mathbf{A}^+ \\ (\mathbf{A}\mathbf{A}^+)^T &= \mathbf{A}\mathbf{A}^+ \\ (\mathbf{A}^+\mathbf{A})^T &= \mathbf{A}^+\mathbf{A}\end{aligned}$$

Moore-Penrose Inverse via SVD:

$$\begin{aligned}\text{if } \mathbf{A} &= \mathbf{U}\mathbf{S}\mathbf{V}^T \\ \mathbf{A}^+ &= \mathbf{V}\mathbf{S}^+\mathbf{U}^T\end{aligned}$$

where \mathbf{S}^+ has the reciprocal of non-zero elements of \mathbf{S} .

SVD for Matrix Approximation

If $\mathbf{A} = \mathbf{USV}^T$ then the optimal (least-squares) k -dimensional approximation of \mathbf{A} (where $k < \text{rank}(\mathbf{A})$) is given by:

$$\tilde{\mathbf{A}} = \mathbf{US}^*\mathbf{V}^T$$

where:

$$\begin{aligned}\mathbf{S}_{ii}^* &= \mathbf{S}_{ii} \text{ for } i \leq k \\ \mathbf{S}_{ii}^* &= 0 \text{ for } i > k\end{aligned}$$

Biplots

- Technique for simultaneously displaying row and column data
- Invented by K. Gabriel (see also papers by Gower)

Given data matrix X , write

$$X = U S V^T$$

$(n \times p)$ $(n \times p)$ $(p \times p)$ $(p \times p)$

$$\tilde{X}_k = U S^* T \quad \begin{matrix} k\text{-dimensional} \\ \text{(approximation to } X\text{)} \end{matrix}$$

reduce \tilde{X} to a product

$$\tilde{X} = G H^T$$

$$\text{where } G = U(S^*)^\alpha \quad H^T = (S^*)^{1-\alpha} V^T$$

(row effects) (column effects)

if $\alpha = 1$, PCs are "spherical"

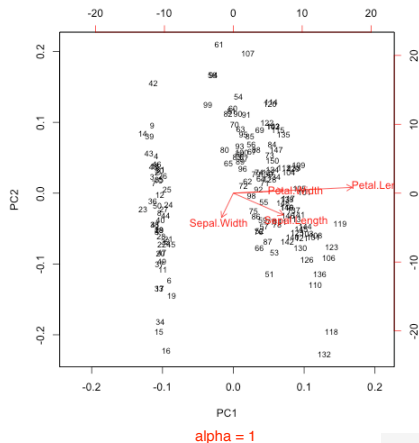
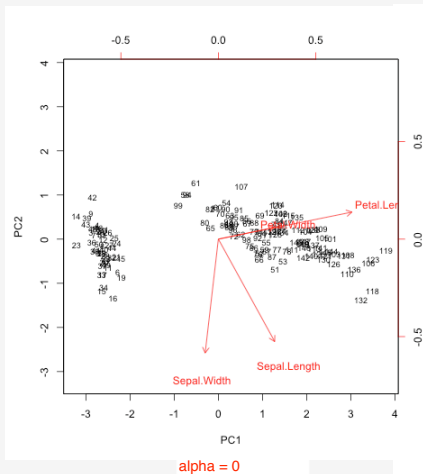
Biplots

$$\begin{aligned}\mathbf{G} &= \mathbf{U}(\mathbf{S}^*)^\alpha \text{ (row effects)} \\ \mathbf{H}^T &= (\mathbf{S}^*)^{1-\alpha} \mathbf{V}^T \text{ (columns effects)}\end{aligned}$$

Different choices of α emphasize different relationships in the data.

- $\alpha = 0$, column-metric preserving biplot; optimally approximates variance-covariance structure. Cosine of angles between vectors approximate correlations; distances between points approximate Mahalanobis distance (“correlation biplot”)
- $\alpha = 1$, row-metric preserving biplot; optimally approximates Euclidean distances among observations. Coordinates of observations correspond to PC scores; coordinates of variables correspond to eigenvector coefficients (“distance biplot”)
- $\alpha = 0.5$, optimally approximates observational values (“symmetric biplot”)

Biplots, Example



Anderson's famous iris data set.

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- PCA plot showing the first two principal components (Comp.1 and Comp.2) for 200 samples. The x-axis (Comp.1) ranges from -0.15 to 0.15, and the y-axis (Comp.2) ranges from -0.15 to 0.15. Red arrows indicate the direction of increasing 'CLCW' (top right) and 'RW' (bottom right). The plot shows a clear separation of samples along the x-axis, with higher values of CLCW and RW associated with higher Comp.1 values.