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

## Matrix Decomposition

A matrix decomposition is a factoring of a matrix into simpler parts. Some familiar factorizations:

$$12 = 3 \times 4 = 2 \times 6$$
$$x^2 - 4 = (x - 2)(x + 2)$$

Matrix decomposition is the same idea!

### Eigendecomposition

You've already been introduced to one way to decompose a square matrix, A:

$$A = VDV^{-1}$$

where:

- V is a matrix of eigenvectors (in columns)
- D is a diagonal matrix with eigenvalues along diagonal.

when A is real-valued and symmetric than V is orthgonal.

## Singular Value Decomposition

$$A = U \leq V^{T} \qquad \text{assume } n \geq p$$

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

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#### Facts about SVD

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

$$rank(A) \leq min(n, p)$$

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

$$|\mathsf{A}|_E = \frac{|\mathsf{A}\mathsf{x}|}{|\mathsf{x}|}$$

The  $L_2$  norm of A is given by  $S_{11}$ .

#### Geometric Interpretation of SVD

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

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

- U and V are orthonormal matrices ~ Orthonormal matrices represent rigid rotations (or rotation plus reflection)
- Diagonal matrices represent "stretching"

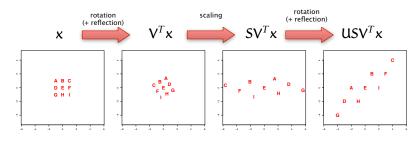
#### SVD Example

$$A = \begin{bmatrix} 3 & 1 \\ 2 & 2 \end{bmatrix} = USV^T$$

where

$$U = \begin{bmatrix} -0.75 & -0.66 \\ -0.66 & 0.75 \end{bmatrix}, S = \begin{bmatrix} 4.13 & 0 \\ 0 & 0.97 \end{bmatrix}, V^{T} = \begin{bmatrix} -0.86 & -0.50 \\ 0.50 & -0.86 \end{bmatrix}$$

#### Geometry



## Relationship of SVD to Eigendecomposition

$$A = USV^{T}$$
 (1)

$$A^{T}A = (VSU^{T})(USV^{T})$$
 (2)

$$= VSU^TUSV^T$$
 (3)

$$= VSSV^{T}$$
 (4)

Equation 4 follows from the fact that U is orthonormal ( $U^TU = I$ ) If we let D = SS, we can rewrite equation 4 as:

$$A^{T}A = VDV$$
 Eigendecomposition! (5)

- The singlular values  $S_{ii}$  are  $\sqrt{D_{ii}}$  where  $d_{ii}$  are the eigenvalues of  $A^TA$ .
- The columns of V are the eigenvectors of  $A^TA$ .

### Using SVD to do PCA

Let X be a mean-centered  $n \times p$  data matrix. The covariance matrix is given by:

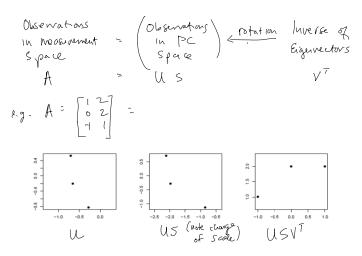
$$C = \frac{1}{n-1} X^T X$$

By SVD we can write  $X = USV^T$ , therefore:

$$C = \frac{1}{n-1} VSU^{T}USV^{T}$$
$$= \frac{1}{n-1} VSSV^{T}$$

- The PC vectors are given by the columns of V
- The PC scores are given by UD, where D = SS

### Another Way of Thinking about SVD



### Applications of SVD

- Pseudoinverse of an arbirary 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 A find matrix A<sup>+</sup> such that:

$$AA^{+}A = A$$

$$A^{+}AA^{+} = A^{+}$$

$$(AA^{+})^{T} = AA^{+}$$

$$(A^{+}A)^{T} = A^{+}A$$

Moore-Penrose Inverse via SVD:

if 
$$A = USV^T$$
  
 $A^+ = VS^+U^T$ 

where  $S^+$  has the reciprocal of non-zero elements of S.

## **SVD** for Matrix Approximation

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

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

where:

$$S_{ii}^{\star} = S_{ii} \text{ for } i \leq k$$
  
 $S_{ii}^{\star} = 0 \text{ for } i > k$ 

## **Biplots**

```
· Technique for simultaneously displaying row and column data
 . Invented by K. Gabriel ( see also papers by
Given dota matrix X, unk
          X = U \leq V^{T}
         (nxp) (nxp) (pxp) (pxp)
         \widetilde{X}_{k} = V S^{*} T  (approximation to X)
reduce & to a product
      \widetilde{Y}_{i} = GH^{T}
         where G= U(S*) +1= (S*) -2 VT
               (row effects) (column effects)
    if L= |, PCs are "sphered"
```

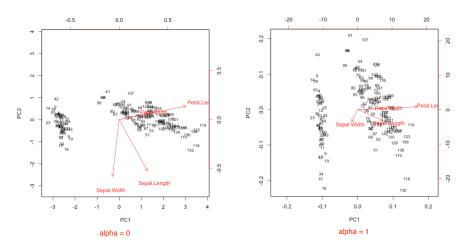
#### **Biplots**

$$G = U(S^*)^{\alpha}$$
 (row effects)  
 $H^T = (S^*)^{1-\alpha}V^T$  (columns effects)

Different choices of  $\alpha$  emphasize different relationships in the data.

- **a**  $\alpha = 0$ , 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")

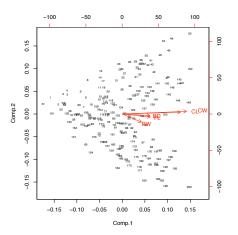
#### Biplots, Example



Anderson's famous iris data set.

#### Biplots, Example II

- Observations drawn as points in space of PCs
- Variables drawn as vectors in PC space



Biplot ( $\alpha = 1$ ) for dataset of five morphological measurements on crabs.