



Computational Methods in Statistics with Applications Singular Value Decomposition

Maya Neytcheva, Lars Eldén

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Matrix factorizations/decompositions



Matrix factorizations I

- ► LU, LDU, Cholesky LDL^T
- ► Tridiagonalization $Q^TAQ = T$, A symmetric Aasen's algorithm: $A = LTL^T$
- ▶ Bidiagonalization $Q^TAV = B$, A(m, n), B upper bi-diagonal
- **.** . . .
- ▶ QR

Golub, Van Loan, *Matrix Computations*, many editions. Note: Some of the algorithms are not numerically stable.



Matrix factorizations

• Schur decomposition A=Q*TAny real matrix A can be decomposed into a unitary matrix Utimes an upper triangular matrix T, which has the eigenvalues of Aon its diagonal. Note: Eigenvalue-revealing factorization



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- Eigenvalue decomposition
- A square. If all its eigenvectors are linearly independent, then $A = QDQ^T$, where Q is orthogonal and D is diagonal, containing the eigenvalues of A.
- Singular value decomposition *SVD*Question: Can we diagonalize a general matrix using unitary matrices?

$$Q_1 A Q_2^T = \Sigma$$



Singular value decomposition

SVD

Let A(m, n), $n \le m$ or $n \ge m$, $rank(A) = rank(A^*)k$.

Definition

If there exist $\mu \neq 0$ and vectors ${\bf u}$ and ${\bf v}$, such that

$$A\mathbf{v} = \mu \mathbf{u}$$
 and $A^* \mathbf{u} = \mu \mathbf{v}$

then μ is called a singular value of A, and \mathbf{u}, \mathbf{v} are a pair of singular vectors, corresponding to μ .



The existence of singular values and vectors is shown...

via the following construction:

$$A\mathbf{v} = \mu\mathbf{u}, \quad A^*\mathbf{u} = \mu\mathbf{v}$$

can be written as

$$\widetilde{A} \begin{bmatrix} \mathbf{v} \\ \mathbf{u} \end{bmatrix} = \begin{bmatrix} 0 & A \\ A^* & 0 \end{bmatrix} \begin{bmatrix} \mathbf{v} \\ \mathbf{u} \end{bmatrix} = \mu \begin{bmatrix} \mathbf{v} \\ \mathbf{u} \end{bmatrix}$$

The matrix \widetilde{A} is selfadjoint, has real eigenvalues and a complete eigenvector space.

Furthermore, μ^2 is an eigenvalue of A^*A with eigenvector ${\bf u}$ and of AA^* with eigenvector ${\bf v}$, because

$$\begin{array}{cccc} A\mathbf{v} = \mu\mathbf{u}, & \rightarrow & A^*A\mathbf{v} = \mu A^*\mathbf{u} = \mu^2\mathbf{v} \\ A^*\mathbf{u} = \mu\mathbf{v}, & \rightarrow & AA^*\mathbf{u} = \mu A\mathbf{v} = \mu^2\mathbf{u} \end{array}$$



Singular Value Decomposition

Theorem (SVD)

Any $m \times n$ matrix A with dimensions, say, $m \ge n$, can be factorized as

$$A = U \begin{pmatrix} \Sigma \\ 0 \end{pmatrix} V^{T},$$

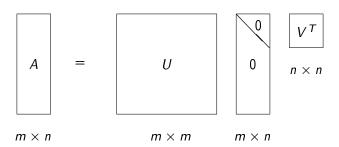
where $U \in \mathbb{R}^{m \times m}$ and $V \in \mathbb{R}^{n \times n}$ are orthogonal, and $\Sigma \in \mathbb{R}^{m \times n}$ is diagonal,

$$\Sigma = \operatorname{diag}(\sigma_1, \, \sigma_2, \, \dots, \, \sigma_n),$$

$$\sigma_1 \ge \sigma_2 \ge \dots \ge \sigma_n \ge 0.$$

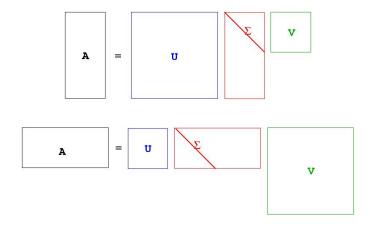


SVD





SVD

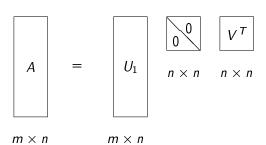




Thin SVD

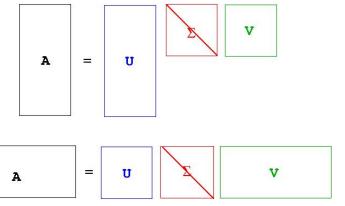
Partition
$$U = (U_1 \ U_2)$$
, where $U_1 \in \mathbb{R}^{m \times n}$,

$$A = U_1 \Sigma V^T$$
,





Thin SVD





Fundamental Subspaces | |

The range of the matrix A:

$$\mathcal{R}(A) = \{ y \mid y = Ax, \text{ for arbitrary } x \}.$$

Assume that A has rank r:

$$\sigma_1 \geq \cdots \geq \sigma_r > \sigma_{r+1} = \cdots = \sigma_n = 0.$$

Outer product form:

$$y = Ax = \sum_{i=1}^{r} \sigma_{i} u_{i} v_{i}^{T} x = \sum_{i=1}^{r} (\sigma_{i} v_{i}^{T} x) u_{i} = \sum_{i=1}^{r} \alpha_{i} u_{i}.$$



Fundamental Subspaces II

The null-space of the matrix A:

$$\mathcal{N}(A) = \{x \mid Ax = 0\}.$$

$$Ax = \sum_{i=1}^{r} \sigma_i u_i v_i^T x$$

Any vector $z = \sum_{i=r+1}^{n} \beta_i v_i$ is in the null-space:

$$Az = \left(\sum_{i=1}^{r} \sigma_i u_i v_i^T\right) \left(\sum_{i=r+1}^{n} \beta_i v_i\right) = 0.$$



Fundamental Subspaces

Theorem (Fundamental subspaces)

1. The singular vectors u_1, u_2, \ldots, u_r are an orthonormal basis in $\mathcal{R}(A)$ and

$$rank(A) = dim(\mathcal{R}(A)) = r.$$

2. The singular vectors $v_{r+1}, v_{r+2}, \ldots, v_n$ are an orthonormal basis in $\mathcal{N}(A)$ and

$$\dim(\mathcal{N}(A)) = n - r.$$

- 3. The singular vectors v_1, v_2, \ldots, v_r are an orthonormal basis in $\mathcal{R}(A^T)$.
- 4. The singular vectors $u_{r+1}, u_{r+2}, \ldots, u_m$ are an orthonormal basis in $\mathcal{N}(A^T)$.



SVD matrix expansion

$$A = U\Sigma V^T$$

$$A = \sum_{i=1}^{n} \sigma_i u_i v_i^T = + + \cdots$$



SVD of a matrix with full column rank I

$$A = 1$$
 1 2 1 2 1 3 1 4



SVD of a matrix with full column rank II



Thin SVD



Rank deficient matrix I

4□ > 4団 > 4 豆 > 4 豆 > 豆 9 Q (~)



Rank deficient matrix II

SVD is rank-revealing!



Null Space

The third column of V is a basis vector in N(A):



Historical notes

SVD has many different names:

- First derivation of the SVD by Eugenio Beltrami (1873)
- ► Full proof by Camille Jordan (1874)
- James Joseph Sylvester (1889), independently discovers SVD
- ► Erhard Schmidt (1907), first to derive an optimal, low-rank approximation of a larger problem
- Hermann Weyl (1912) determination of the rank in the presence of errors
- Eckart-Young decomposition and optimality properties of SVD (1936), psychometrics
- Numerically efficient algorithms to compute the SVD works by Gene Golub 1970 (Golub-Kahan)





Best approximation / Eckart-Young Property I

Frobenius norm

$$||A||_F = \left(\sum_{i,j} a_{ij}^2\right)^{1/2}$$



Best approximation / Eckart-Young Property II

Theorem

Assume that the matrix $A \in \mathbb{R}^{m \times n}$ has rank r and choose k, such that r > k. The Frobenius norm matrix approximation problem

$$\min_{\operatorname{rank}(Z)=k} \|A-Z\|_F$$

has the solution

$$Z = A_k = U_k \Sigma_k V_k^T,$$

where
$$U_k = (u_1, \ldots, u_k)$$
, $V_k = (v_1, \ldots, v_k)$, and $\Sigma_k = \operatorname{diag}(\sigma_1, \ldots, \sigma_k)$.



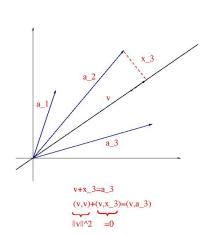
Best approximation / Eckart-Young Property III

Proof:

- (1) Observe: if $A_k = \sum_{j=1}^k \sigma_j u_j v_j^*$, then $||A A_k|| = \sigma_{k+1}$.
- (2) Observe: Consider the subspace, spanned by the first k+1 singular vectors of A, W. Then, $||Aw||_2 \ge \sigma k + 1||w||_2$, $w \in W$.
- (3) Assume that there exists a matrix B of rank k, such that $\|A-B\|_2 < \sigma_{k+1}$. Then, there exists a subspace \widehat{W} of size n-k, such that $Bw=0, w\in \widehat{W}$.
- $||Aw||_2 = ||(A B)w||_2 \le ||A B||_2 ||w||_2 \le \sigma_{k+1} ||w||_2$. From dinemsion argiments $W \cap \widehat{W} \ne \emptyset$.



Singular vectors, another view



Consider the rows of A(m, n) as points in an n-dimensional space and find the best linear fit through the origin.

$$egin{aligned} \mathbf{v}_1 &= rg\max_{\|\mathbf{v}\|=1} \|A\mathbf{v}\|_2^2, \ \sigma_1 &= \|A\mathbf{v}_1\|_2 \end{aligned}$$
 $egin{aligned} \mathbf{v}_2 &= rg\max_{\|\mathbf{v}\|=1,\mathbf{v}\perp\mathbf{v}_1} \|A\mathbf{v}\|_2^2 \end{aligned}$



Principal Component Analysis (PCA) I

Data matrix $\mathbb{R}^{m \times n} \ni X = U \Sigma V^T$

Each column of X is an observation of a real-valued random vector with mean zero.

The right singular vectors v_i are called *principal components* directions of X. The vector

$$z_1 = Xv_1 = \sigma_1 u_1$$

has the largest sample variance amongst all normalized linear combinations of the columns of X:

$$\operatorname{Var}(z_1) = \operatorname{Var}(Xv_1) = \frac{\sigma_1^2}{m}.$$



Principal Component Analysis (PCA) II

The normalized variable u_1 is called the *normalized first principal* component of X.

The second principal component is the vector of largest sample variance of the deflated data matrix $X - \sigma_1 u_1 v_1^T$, and so on.



Test example borrowed from Computational Statistics with Application to Bioinformatics Prof. William H. Press Spring Term, 2008, The University of Texas at Austin



Example

Consider some gene expression data, represented by the so-called 'design matrix' $X = \{X_{ij}\}$

Each column of X corresponds to a separate observation, in this case, a separate micro array experiment under a different condition. N rows are genes (1:500) and M columns are the corresponding responses.

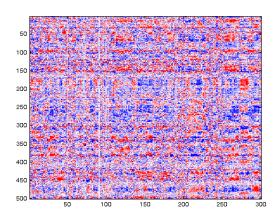
Assumptions:

- the individual experiments (columns of X) have zero mean.
- scale data to unit standard deviation.



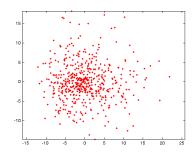
```
load yeastarray_t2.txt;
size(yeastarray_t2)
ans = 500 300
vclip = prctile(yeastarray_t2(:),[1,99])
vclip = -204 244
data = max(yclip(1), min(yclip(2), yeastarray_t2));
dmean= mean(data,1);
dstd = std(data, 1);
data = (data - repmat(dmean, [size(data, 1), 1]))./...
               repmat (dstd, [size(data, 1), 1]);
genecolormap = [\min(1, (1:64)/32); 1-abs(1-(1:64)/32);
                 min(1.(64-(1:64))/32)1':
figure (1), clf, colormap (genecolormap);
image(20*data+32)
```







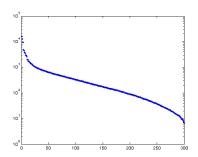
```
[U S V] = svd(data, 0);
PCAcoords = U*S;
plot(PCAcoords(:,1),PCScoords(:,2),'r.')
axis equal
```





The squares of the singular values are proportional to the portion of the total variance (L_2 norm of X) that each accounts for.

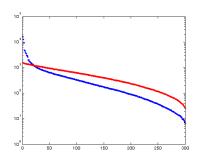
```
ssq = diag(S).^2;
semilogy(ssq,'.b')
```





We can produce fake data and compare:

```
fakedata = randn(500,300);
[Uf Sf Vf] = svd(fakedata,0);
sfsq = diag(Sf).^2;
semilogy(sfsq,'.r')
```

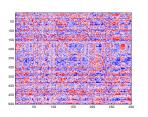


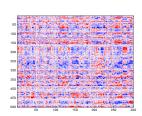


For the data in this example, a sensible use of PCA (i.e., SVD) would be to project the data into the subspace of the first 20 SVs, where we can be sure that it is not noise.

```
% Truncate the first 20 singular values/vectors
strunc = diag(S);
strunc(21:end) = 0;
filtdata20 = U*diag(strunc)*V';
figure (2), clf, colormap (genecolormap);
image(20*filtdata20+32)
% Truncate the first 5 singular values/vectors
strunc(6:end) = 0;
filtdata5 = U*diag(strunc)*V';
figure (3), clf, colormap (genecolormap);
image (20*filtdata5+32)
```

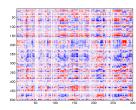






(a) The original

(b) truncate to 20

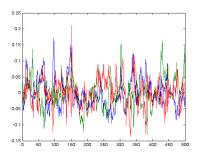


(c) Truncate to 5



How to interpret the singular vectors? The first three vectors u are 'eigengenes', the linear combination of genes that explain the most data.

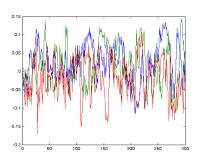
plot(U(:,1:3))





The first three vectors v are 'eigenarrays', the linear combination of experiments that explain the most data.

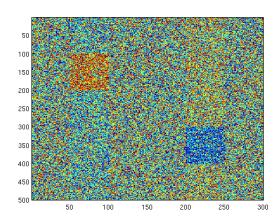
plot(V(:,1:3))





Consider a toy example





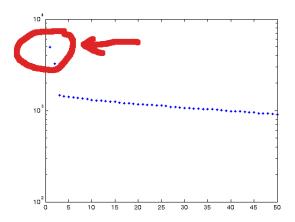


Consider a toy example

```
[Up Sp Vp] = svd(pdata,0);
spsq = diag(Sp).^2;
semilogy(spsq(1:50),'.b')
```

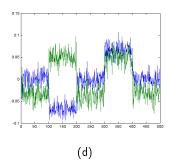
Should we expect the eigengenes/eigenarrays to show the separate main effects?

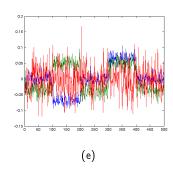






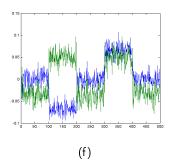
```
plot(Up(:,1:2)),
plot(Up(:,1:3))
```

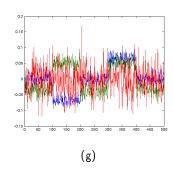






```
plot(Vp(:,1:2)),
plot(Vp(:,1:3))
```







Solving Least Squares problems by SVD

$$A\mathbf{x} = \mathbf{b}, A(m, n)$$

$$A = U\Sigma V$$

$$U\Sigma V\mathbf{x} = \mathbf{b} \rightarrow \mathbf{x} = V(\Sigma^{-1}(U^T\mathbf{b}))$$



Least Squares by SVD |

$$A = 1$$
 1 $b = 7.9700$
1 2 10.2000
1 3 14.2000
1 4 16.0000
1 5 21.2000



Least Squares by SVD II

$$S = 7.6912$$
 0 $V = 0.2669$ -0.9637 0 0.9194 0.9637 0.2669 >> $x=V*(S\setminus(U1'*b))$ $x = 4.2360$ 3.2260



Least Squares by SVD, in R | 1

```
> A.svd<-svd(A)
> A.svd
$d
[1] 7.6912131 0.9193696
$u
          [,1]
                    [,2]
[1,] 0.1600071 0.7578903
[2,] 0.2853078 0.4675462
[3,] 0.4106086 0.1772020
[4,] 0.5359094 -0.1131421
[5,] 0.6612102 -0.4034862
```



Least Squares by SVD, in R II

Linear dependence - SVD

Theorem

Let the singular values of A satisfy

$$\sigma_1 \geq \cdots \geq \sigma_r > \sigma_{r+1} = \cdots = \sigma_n = 0.$$

Then the rank of A is equal to r.

Rank = the number of linearly independent columns of A.



Linear dependence I

```
A=[1 1; 1 2; 1 3; 1 4]
singval=svd(A)
% Third col=linear combination of first two
A1=[A A(:,1)+0.5*A(:,2)]
singval1=svd(A1)
```



Linear dependence II

Result:

```
singval = 5.7794
                      0.7738
A1 = 1.0000 1.0000 1.5000
    1.0000 2.0000 2.0000
    1.0000 3.0000 2.5000
    1.0000 4.0000 3.0000
singval1 = 7.3944
         0.9072
```



Almost linear dependence I

```
A2=[A A(:,1)+0.5*A(:,2)+0.0001*randn(4,1)]
singval2=svd(A2)
```



Almost linear dependence? I

Run Matlab demo

~/.../STAT/Labs/Lab_QR_SVD/Small_singular_values.m



Computing the SVD in a numerically efficient way



Computing the SVD

1. Transform A to bidiagonal form by unitary transformations

$$Q_L A Q_R = B = \begin{bmatrix} * & * & & & & \\ & * & * & & & \\ & \ddots & \ddots & * & \\ & & & * \end{bmatrix}$$

2. Diagonalize B by two orthogonal transformations

$$\widetilde{Q}_L B \widetilde{Q}_R = \widetilde{Q}_L Q_L A Q_R \widetilde{Q}_R = \Sigma$$

The cost for the bidiagonalization is $4mn^2 - 4/3n^3$. The cost for SVD: $4m^2n + 8mn^2 + 9n^3$.