# Computational Methods in Statistics with Applications <br> Singular Value Decomposition 

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## Matrix factorizations/decompositions

## Matrix factorizations I

- LU, LDU, Cholesky $L D L^{T}$
- Tridiagonalization $Q^{T} A Q=T, A$ - symmetric Aasen's algorithm: $A=L T L^{T}$
- Bidiagonalization $Q^{T} A V=B, A(m, n), B$ - upper bi-diagonal
- $Q R$

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

## Matrix factorizations

- Schur decomposition $A=Q * T$

Any real matrix $A$ can be decomposed into a unitary matrix $U$ times an upper triangular matrix $T$, which has the eigenvalues of $A$ on its diagonal. Note: Eigenvalue-revealing factorization

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- 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 $A(m, n), n \leq m$ or $n \geq m, \operatorname{rank}(A)=\operatorname{rank}\left(A^{*}\right) k$.

## Definition

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

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

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

## 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

$$
\tilde{A}\left[\begin{array}{l}
\mathbf{v} \\
\mathbf{u}
\end{array}\right]=\left[\begin{array}{cc}
0 & A \\
A^{*} & 0
\end{array}\right]\left[\begin{array}{l}
\mathbf{v} \\
\mathbf{u}
\end{array}\right]=\mu\left[\begin{array}{l}
\mathbf{v} \\
\mathbf{u}
\end{array}\right]
$$

The matrix $\widetilde{A}$ is selfadjoint, has real eigenvalues and a complete eigenvector space.
Furthermore, $\mu^{2}$ is an eigenvalue of $A^{*} A$ with eigenvector $\mathbf{u}$ and of $A A^{*}$ with eigenvector v, because

$$
\begin{aligned}
A \mathbf{v}=\mu \mathbf{u}, & \rightarrow \quad A^{*} A \mathbf{v}=\mu A^{*} \mathbf{u}=\mu^{2} \mathbf{v} \\
A^{*} \mathbf{u}=\mu \mathbf{v}, & \rightarrow \quad A A^{*} \mathbf{u}=\mu A \mathbf{v}=\mu^{2} \mathbf{u}
\end{aligned}
$$

Singular Value Decomposition

## Theorem (SVD)

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

$$
A=U\binom{\Sigma}{0} V^{T}
$$

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,

$$
\begin{aligned}
& \Sigma=\operatorname{diag}\left(\sigma_{1}, \sigma_{2}, \ldots, \sigma_{n}\right) \\
& \sigma_{1} \geq \sigma_{2} \geq \cdots \geq \sigma_{n} \geq 0
\end{aligned}
$$

SVD


SVD


## Thin SVD

Partition $U=\left(U_{1} U_{2}\right)$, where $U_{1} \in \mathbb{R}^{m \times n}$,

$$
A=U_{1} \Sigma V^{T}
$$


$m \times n$
$m \times n$

## Thin SVD



## Fundamental Subspaces I

The range of the matrix $A$ :

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

Assume that $A$ has rank $r$ :

$$
\sigma_{1} \geq \cdots \geq \sigma_{r}>\sigma_{r+1}=\cdots=\sigma_{n}=0
$$

Outer product form:

$$
y=A x=\sum_{i=1}^{r} \sigma_{i} u_{i} v_{i}^{\top} x=\sum_{i=1}^{r}\left(\sigma_{i} v_{i}^{\top} x\right) u_{i}=\sum_{i=1}^{r} \alpha_{i} u_{i}
$$

## Fundamental Subspaces II

The null-space of the matrix $A$ :

$$
\begin{gathered}
\mathcal{N}(A)=\{x \mid A x=0\} . \\
A x=\sum_{i=1}^{r} \sigma_{i} u_{i} v_{i}^{\top} x
\end{gathered}
$$

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

$$
A z=\left(\sum_{i=1}^{r} \sigma_{i} u_{i} v_{i}^{T}\right)\left(\sum_{i=r+1}^{n} \beta_{i} v_{i}\right)=0
$$

## 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

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

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

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

3. The singular vectors $v_{1}, v_{2}, \ldots, v_{r}$ are an orthonormal basis in $\mathcal{R}\left(A^{T}\right)$.
4. The singular vectors $u_{r+1}, u_{r+2}, \ldots, u_{m}$ are an orthonormal basis in $\mathcal{N}\left(A^{T}\right)$.

## SVD matrix expansion

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

$$
A=\sum_{i=1}^{n} \sigma_{i} u_{i} v_{i}^{T}=\mid \overline{ }+\sqrt{ }+\cdots
$$

SVD of a matrix with full column rank I

$$
\begin{aligned}
& A=\begin{array}{ll}
1 & 1 \\
1 & 2
\end{array} \\
& 1 \\
& 3 \\
& 14 \\
& \text { >> }[\mathrm{U}, \mathrm{~S}, \mathrm{~V}]=\operatorname{svd}(\mathrm{A})
\end{aligned}
$$

## SVD of a matrix with full column rank II

$$
\begin{aligned}
& \begin{array}{rrrr}
\mathrm{U}= & 0.2195 & -0.8073 & 0.0236 \\
& 0.3833 & -0.3912 & -0.4393 \\
& 0.5472 & 0.0249 & 0.8472 \\
& 0.7110 & 0.4410 & -0.3921
\end{array} \\
& S=5.7794 \\
& 0 \\
& 0.7738 \\
& \begin{array}{rrr}
\mathrm{V}= & 0.3220 & -0.9467 \\
& 0.9467 & 0.3220
\end{array}
\end{aligned}
$$

## Thin SVD

$$
\begin{aligned}
& \text { >> }[\mathrm{U}, \mathrm{~S}, \mathrm{~V}]=\operatorname{svd}(\mathrm{A}, 0) \\
& \mathrm{U}=0.2195-0.8073 \\
& 0.3833-0.3912 \\
& 0.5472 \quad 0.0249 \\
& 0.7110 \quad 0.4410 \\
& S=5.7794 \\
& 0 \quad 0.7738 \\
& \begin{array}{rlr}
\mathrm{V}= & 0.3220 & -0.9467 \\
& 0.9467 & 0.3220
\end{array}
\end{aligned}
$$

## Rank deficient matrix I

$$
\begin{array}{rlrl}
\gg & A(:, 3)=A(:, 1)+0.5 * A(:, 2) \\
A= & 1.0000 & 1.0000 & 1.5000 \\
& 1.0000 & 2.0000 & 2.0000 \\
& 1.0000 & 3.0000 & 2.5000 \\
& 1.0000 & 4.0000 & 3.0000 \\
\\
\gg & {[U, S, V]=\operatorname{svd}(A, 0)} \\
U= & \\
& \\
0.2612 & -0.7948 & -0.5000 \\
& 0.4032 & -0.3708 & 0.8333 \\
& 0.6871 & 0.0533 & -0.1667
\end{array}
$$

## Rank deficient matrix II

$$
\begin{array}{rrr}
S=7.3944 & 0 & 0 \\
0 & 0.9072 & 0 \\
0 & 0 & 0 \\
& & \\
V=0.2565 & -0.6998 & 0.6667 \\
0.7372 & 0.5877 & 0.3333 \\
& 0.6251 & -0.4060
\end{array}-0.6667 \text { rer }
$$

SVD is rank-revealing!

## Null Space

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

$$
\begin{aligned}
& \text { >> } A * V(:, 3) \\
& \text { ans }= \\
& 1.0 \mathrm{e}-15 * \\
& 0 \\
& -0.2220 \\
& -0.2220 \\
& 0
\end{aligned}
$$

## 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

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

## 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 $U_{k}=\left(u_{1}, \ldots, u_{k}\right), V_{k}=\left(v_{1}, \ldots, v_{k}\right)$, and
$\Sigma_{k}=\operatorname{diag}\left(\sigma_{1}, \ldots, \sigma_{k}\right)$.

## Best approximation / Eckart-Young Property III

## Proof:

(1) Observe: if $A_{k}=\sum_{j=1}^{k} \sigma_{j} u_{j} v_{j}^{*}$, then $\left\|A-A_{k}\right\|=\sigma_{k+1}$.
(2) Observe: Consider the subspace, spanned by the first $k+1$ singular vectors of $A, W$. Then, $\|A w\|_{2} \geq \sigma k+1\|w\|_{2}, w \in W$. (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 $B w=0, w \in \widehat{W}$.
$\|A w\|_{2}=\|(A-B) w\|_{2} \leq\|A-B\|_{2}\|w\|_{2} \leq \sigma_{k+1}\|w\|_{2}$. From dinemsion argiments $W \cap \widehat{W} \neq \emptyset$.

## 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.

$$
\begin{gathered}
\mathbf{v}_{1}=\arg \max _{\|\mathbf{v}\|=1}\|A \mathbf{v}\|_{2}^{2}, \sigma_{1}=\left\|A \mathbf{v}_{1}\right\|_{2} \\
\mathbf{v}_{2}=\arg \max _{\|\mathbf{v}\|=1, \mathbf{v} \perp \mathbf{v}_{\mathbf{1}}}\|A \mathbf{v}\|_{2}^{2}
\end{gathered}
$$

$$
\left.\begin{array}{l}
\mathrm{v}+\mathrm{x} \_3=\mathrm{a} \_3 \\
\underbrace{}_{\|\mathrm{v}\| \wedge} \mathrm{v}_{2} \\
(\mathrm{v}, \mathrm{v}) \\
+(\underbrace{\mathrm{v}, \mathrm{x}-3}_{=0})
\end{array}\right)=(\mathrm{v}, \mathrm{a}-3)
$$

## Principal Component Analysis (PCA) ।

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

$$
z_{1}=X v_{1}=\sigma_{1} u_{1}
$$

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

$$
\operatorname{Var}\left(z_{1}\right)=\operatorname{Var}\left(X v_{1}\right)=\frac{\sigma_{1}^{2}}{m}
$$

## 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' $X=\left\{X_{i j}\right\}$
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
yclip = prctile(yeastarray_t2(:), [1,99])
yclip = -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)]';
figure(1),clf,colormap(genecolormap);
image(20*data+32)
```


[U S V] = svd(data, 0);
PCAcoords $=\mathrm{U} * S$;
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

```
pdata = randn(500,300);
pdata(101:200,51:100) = pdata(101:200,51:100) + 1;
pdata(301:400,201:250) = pdata(301:400,201:250) - 1;
pmean = mean(pdata,1);
pstd = std(pdata,1);
pdata = (pdata - repmat(pmean,[size(pdata,1),1]))./...
    repmat(pstd,[size(pdata,1),1]);
colormap (genecolormap)
image (20*pdata+32)
```



## 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))
```


(d)

(e)

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


(f)

(g)

## Solving Least Squares problems by SVD

$$
\begin{aligned}
& A \mathbf{x}=\mathbf{b}, A(m, n) \\
& A=U \Sigma V \\
& U \Sigma V \mathbf{x}=\mathbf{b} \rightarrow \mathbf{x}=V\left(\Sigma^{-1}\left(U^{T} \mathbf{b}\right)\right)
\end{aligned}
$$

## Least Squares by SVD ।

$$
\begin{aligned}
& \mathrm{A}=1 \quad 1 \\
& 12 \\
& 13 \\
& 14 \\
& 15 \\
& \text { >> }[\mathrm{U} 1, \mathrm{~S}, \mathrm{~V}]=\operatorname{svd}(\mathrm{A}, 0) \\
& \mathrm{U} 1=0.1600 \quad-0.7579 \\
& 0.2853-0.4675 \\
& 0.4106-0.1772 \\
& 0.5359 \quad 0.1131 \\
& 0.66120 .4035
\end{aligned}
$$

## Least Squares by SVD II

$$
\begin{aligned}
& \begin{array}{ccc}
S=7.6912 & 0 & V= \\
0 & 0.9194 & 0.2669 \\
\hline
\end{array} \\
& \text { >> } \mathrm{x}=\mathrm{V} *\left(\mathrm{~S} \backslash\left(\mathrm{U} 1^{\prime} * \mathrm{~b}\right)\right) \\
& \begin{aligned}
\mathrm{x}= & 4.2360 \\
& 3.2260
\end{aligned}
\end{aligned}
$$

## Least Squares by SVD, in $R$ I

```
> 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

\$v

|  | $[, 1]$ | $[, 2]$ |
| ---: | ---: | ---: |
| $[1]$, | 0.2669336 | 0.9637149 |
| $[2]$, | 0.9637149 | -0.2669336 |

> $x=A . s v d \$ v \% * \%$ diag (1/A.svd\$d) $\% * \% ~ t(A . s v d \$ u) ~ \% * \%$
$>\mathrm{X}$
$\begin{array}{cc} & {[, 1]} \\ {[1,]} & 4.236 \\ {[2,]} & 3.226\end{array}$

## 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 $\mathrm{A} 1=[\mathrm{A} \mathrm{A}(:, 1)+0.5 * \mathrm{~A}(:, 2)]$
singvall=svd(A1)

## Linear dependence II

## Result:

$$
\begin{array}{cccc}
A=\begin{array}{lll}
1 & 1 & \\
1 & 2 & \\
1 & 3 & \\
1 & 4 & \\
& & \\
& & \\
& & \\
& 1.0000 & 1.0000
\end{array} & 1.5000 \\
& 1.0000 & 2.0000 & 2.0000 \\
& 1.0000 & 3.0000 & 2.5000 \\
& 1.0000 & 4.0000 & 3.0000
\end{array}
$$

$$
\text { singvall }=7.3944
$$

$$
0.9072
$$

## Almost linear dependence I

$A 2=[A A(:, 1)+0.5 * A(:, 2)+0.0001 * r a n d n(4,1)]$ singval2=svd(A2)

$$
\begin{array}{rlrl}
\mathrm{A} 2= & 1.0000 & 1.0000 & 1.4999 \\
& 1.0000 & 2.0000 & 2.0001 \\
& 1.0000 & 3.0000 & 2.5000 \\
& 1.0000 & 4.0000 & 3.0001
\end{array}
$$

singval2 $=7.3944$
0.9072
0.0001

## Almost linear dependence? |

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

$$
Q_{L} A Q_{R}=B=\left[\begin{array}{ccccc}
* & * & & & \\
& * & * & & \\
& \ddots & \ddots & * & \\
& & & & *
\end{array}\right]
$$

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 $4 m n^{2}-4 / 3 n^{3}$. The cost for SVD: $4 m^{2} n+8 m n^{2}+9 n^{3}$.

