### Singular Value Decomposition

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Goal: Another matrix decomposition (SVD) for low-rank matrix approximation

cf. Eigen decomposition
$$A = Q [ ]Q^{T}$$
QR decomposition
$$A = Q [ ]$$

See note on "least square fit" & Numerical Recipes Sec. 2.6





#### Rank of a Matrix

•  $N \times M$  matrix A as a mapping:  $R^M \to R^N$ 

$$M \quad \begin{bmatrix} x \\ x \end{bmatrix} \quad x (\in \mathbb{R}^M) \xrightarrow{A} b = Ax (\in \mathbb{R}^N) \quad \begin{bmatrix} b \\ b \end{bmatrix} \quad N$$

- Range of A: Vector space  $\{b = Ax | \forall x\}$
- Rank of A: Number, m, of linearly-independent vectors in the range, i.e., how many linearly-independent N-element vectors are there in the range, such that

$$b = A^{\forall} x = \sum_{v=1}^{m} c_v |v\rangle$$

#### Low Rank Approximations of a Matrix

• Rank-1 approximation:  $NM \rightarrow N + M$ 

$$\mathbf{N} \left[ \begin{array}{c} \mathbf{M} \\ \psi \end{array} \right] \cong \left[ u \right[ \begin{array}{c} v \end{array} \right] \qquad |u\rangle\langle v|\forall x\rangle \propto |u\rangle$$

• Rank-2 approximation:  $NM \rightarrow 2(N + M)$ 

$$\left[ \begin{array}{c} \psi \end{array} \right] \cong \left[ u_1 \right] w_1 \left[ \begin{array}{c} v_1 \end{array} \right] + \left[ u_2 \right] w_2 \left[ \begin{array}{c} v_2 \end{array} \right]$$

• Rank- $m \ (m << N, M)$  approximation:  $NM \to m(N + M)$ 

$$\left[ \begin{array}{c} \psi \\ \end{array} \right] \cong \sum_{v=1}^{m} \left[ u_{v} \right] w_{v} \left[ \begin{array}{c} v_{v} \\ \end{array} \right]$$

### Singular Value Decomposition

- **Problem:** Optimal approximation of an  $N \times M$  matrix  $\psi$  of rank-m (m << N)?

Theorem: An 
$$N \times M$$
 matrix  $\psi$  (assume  $N \ge M$ ) can be decomposed as 
$$\psi = UDV^T = \sum_{v=1}^M U_{iv} d_v V_{jv} = \sum_{v=1}^M u_i^{(v)} d_v v_j^{(v)}$$

where  $U \in \mathbb{R}^N \times \mathbb{R}^M$  &  $V \in \mathbb{R}^M \times \mathbb{R}^M$  are column orthogonal & D is diagonal

$$\mathbf{M} \qquad U^T U = V^T V = I_M$$
 
$$\mathbf{N} \begin{bmatrix} \psi \end{bmatrix} = \begin{bmatrix} U \\ U \end{bmatrix} \begin{bmatrix} \text{See appendix on polar \& singular decompositions} \\ d_1 \\ \ddots \\ d_M \end{bmatrix} \begin{bmatrix} V^T \\ W \times \mathbf{M} \end{bmatrix}$$

• Theorem: Sort the SVD diagonal elements in descending order,  $d_1 \ge d_2 \ge ... \ge$  $d_M \ge 0$ , & retain the first m terms  $\psi^{(m)} = \sum_{i=1}^{m} u^{(v)} d_v v^{(v)T}$ 

which is optimal among  $\forall$ rank-m matrices in the 2-norm sense with the error

$$\min_{rank(A)=m} ||A - \psi||_2 = ||\psi^{(m)} - \psi||_2 = d_{m+1}$$

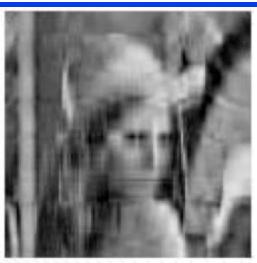
cf. singular.c & svdcmp.c cc -o singular singular.c svdcmp.c -lm

Use the program!

# **SVD** for Image Compression







Original Image

5 Iterations

10 Iterations

D. Richards & A. Abrahamsen





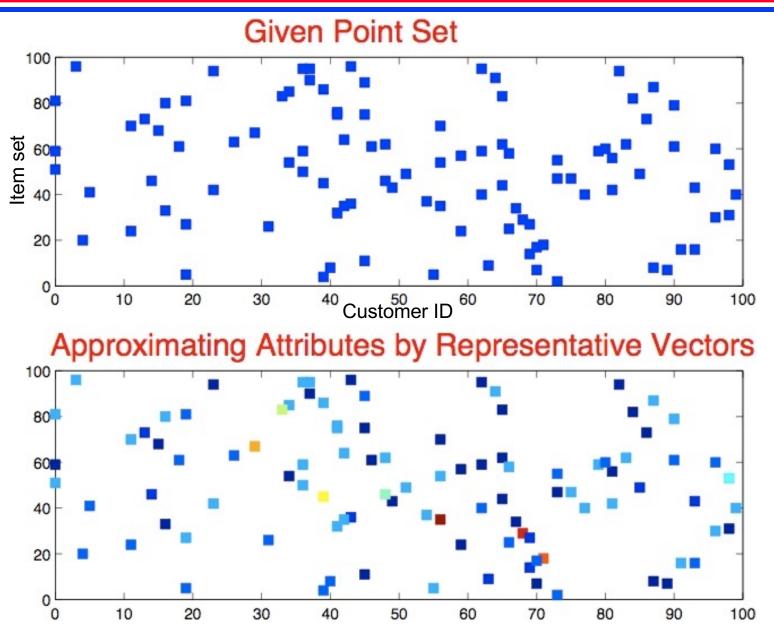


20 Iterations

60 Iterations

100 Iterations

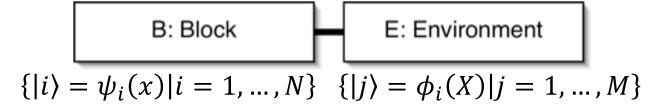
### **SVD** in Data Mining



N. Ramakrishnan & A. Y. Grama

### **Reduced Density Matrix**

Quantum system coupled to an environment



• **∀Quantum state of block + environment** 

$$|\psi\rangle = \sum_{i=1}^{N} \sum_{j=1}^{M} \psi_{ij} |i\rangle |j\rangle$$
 or  $\Psi(x,X) = \sum_{i=1}^{N} \sum_{j=1}^{M} \psi_{ij} \psi_i(x) \phi_j(X)$ 

Reduced density matrix

#### Low-Rank Approx. to Reduced Density Matrix

$$\psi \cong \psi^{(m)} = \sum_{v=1}^{m} u^{(v)} d_{v} v^{(v)T} \qquad \psi_{ij}^{(m)} = \sum_{v=1}^{m} u_{i}^{(v)} d_{v} v_{j}^{(v)}$$

$$\rho = \psi \psi^{T} \cong \psi^{(m)} \psi^{(m)T} = \sum_{v=1}^{m} \sum_{v'=1}^{m} u^{(v)} d_{v} \left( v^{(v)T} v^{(v')} \right) d_{v'} u^{(v')T}$$

$$= \sum_{v=1}^{m} \sum_{v'=1}^{m} u^{(v)} d_{v} \left( \delta_{vv'} \right) d_{v'} u^{(v')T} = \sum_{v=1}^{m} u^{(v)} d_{v}^{2} u^{(v)T} \equiv \rho^{(m)}$$

$$\rho_{ii'}^{(m)} = \sum_{v=1}^{m} u_{i}^{(v)} d_{v}^{2} u_{i'}^{(v)}$$

- Density matrix renormalization group = systematic procedure to accurately obtain a quantum ground state:
  - 1. Incrementally add environment to a block
  - 2. Solve the global (= block + environment) ground state
  - 3. Construct a low-rank approx. to represent the block with reduced d.o.f.

S. R. White, *Phys. Rev. B* **48**, 10345 ('93); G. K.-L. Chan & S. Sharma, *Annu. Rev. Phys. Chem.* **62**, 465 ('11)

#### **Entanglement Entropy**

- Entanglement entropy: A measure of the degree of quantum entanglement between two subsystems. If a state describing two subsystems A and B is a *separable* state  $|\Psi_{AB}\rangle = |\varphi_{A}\rangle|\varphi_{B}\rangle$ , then the reduced density matrix  $\rho_{A} = \text{Tr}_{B}|\Psi_{AB}\rangle\langle\Psi_{AB}| = |\varphi_{A}\rangle\langle\varphi_{A}|$  is a *pure state*. Thus, the entropy of the state is zero. A reduced density matrix having a non-zero entropy is therefore a signal of the existence of entanglement in the system.
- Area law: A quantum state satisfies an *area law* if the leading term of the entanglement entropy grows at most proportionally with the *boundary* between the two partitions. Area laws are remarkably common for ground states of local gapped quantum many-body systems. It greatly reduces the complexity of quantum many-body systems. The density matrix renormalization group and matrix product states, for example, implicitly rely on such area laws.

## **SVD** for Rapid Genome Sequencing

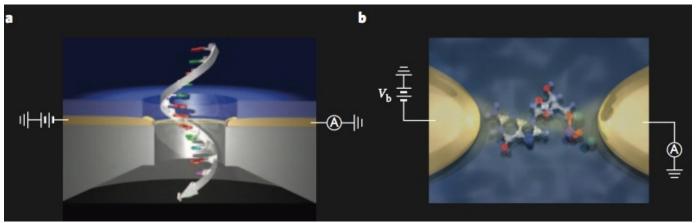
• \$10M Archon X prize for decoding 100 human genomes in 10 days & \$10K per genome (http://genomics.xprize.org): Preemptive attack on diseases



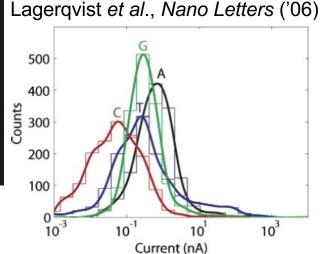




Quantum tunneling current for rapid DNA sequencing?



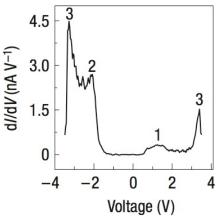
Tsutsui et al., Nature Nanotechnology ('10)



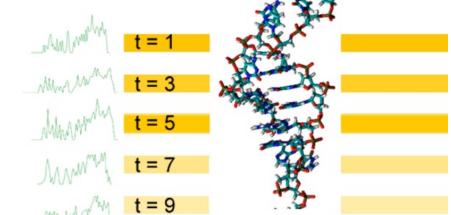
• Tunneling current alone cannot distinguish the 4 nucleotides (A, C, G, T)

### Rapid DNA Sequencing via Data Mining

Use tunneling current (I)-voltage (V) characteristic (or electronic density-ofstates) as the 'fingerprints' of the 4 nucleotides

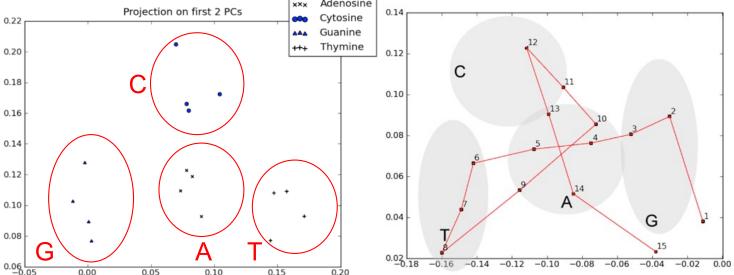


Shapir et al., Nature Materials ('08)



Principal component analysis (PCA) & fuzzy c-means clustering clearly distinguish the 4 nucleotides

H. Yuen et al., IJCS 4, 352 ('10)





http://www.henryyuen.net/

Viterbi algorithm for even higher-accuracy sequencing

### SVD vs. PCA (in Economics)

SVD of N (number of companies)  $\times$  T (number of time points) of stock-price time series

$$\Xi^{T} = U \sum_{T \times N} V^{T}$$

$$T \times N \times N \times N \times N$$

**Stock correlation matrix** 

$$\mathbf{C}_{N\times N} = \mathbf{\Xi} \quad \mathbf{\Xi}^{T}$$

$$N\times N = T \times N$$

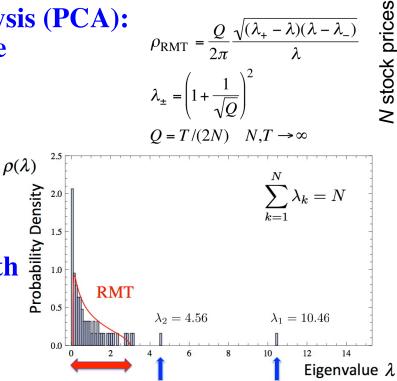
**Principal component analysis (PCA):** Eigen decomposition of the correlation matrix

$$C = \Xi \Xi^{T}$$

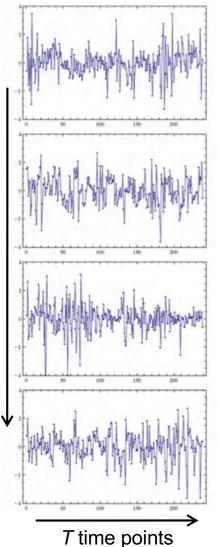
$$= V \Sigma \widetilde{U^{T} U} \Sigma V^{T}$$

$$= V \Sigma^{2} V^{T}$$

**Compare the spectrum with** that of random matrix theory (RMT) for judging statistical significance



 $\rho_{\rm RMT} = \frac{Q}{2\pi} \frac{\sqrt{(\lambda_+ - \lambda)(\lambda - \lambda_-)}}{\lambda}$ 



Y. Kichikawa et al., Proc. Comp. Sci. **60**, 1836 ('15)