Chapter 4 **Column and Column-Row Decompositions**

Column Decompositions

DMM, summer 2017 **Pauli Miettinen** [Boutsidis et al. 2010;](http://arxiv.org/pdf/0812.4293.pdf) [Strauch 2014](http://kops.uni-konstanz.de/handle/123456789/29539)

Motivation

- SVD is often hard to interpret and yields dense factorizations
	- NMF tries to address these problems with varying success
- But if original data is sparse & easy to interpret, why not use it in the decompositions?

The CX decomposition

- In the **CX decomposition** we are given a matrix *A* and a rank *k*, and we need to select *k* **columns** of *A* into matrix *C* and build matrix *X* s.t. we minimize ||*A* – *CX*||^ξ
	- ξ is either *F* or 2
	- A.k.a. **column subset selection problem** (CSSP)

Why CX?

- The columns of *C* preserve the original interpretation of columns of *A*
	- Even complex constraints are satisfied if the original data satisfied them
- Feature selection
	- Selects the columns that can be used to explain the rest
	- Compare to the dimensionality reduction

Alternative target function

- Building *C* is the hard part of CX decompositions
- Given *A* and *C*, *X* can be computed with the pseudo-inverse
	- $X = C^+A$
	- Alternative target function for CX: minimize $||\mathbf{A} - \mathbf{CC}^+\mathbf{A}||_{\mathcal{E}} = ||\mathbf{A} - \mathbf{P}_\mathbf{C}\mathbf{A}||_{\mathcal{E}}$

How to select C?

- Exhaustive: try all $\binom{n}{k}$ subsets of columns *m k* $\overline{\mathcal{L}}$
	- Not very scalable
- Try to select the columns in a clever way
	- But how?

Sample columns w.r.t. carefully selected probabilities

• Avoids deterministic worst-case scenarios

Related idea: RRQR

• The **rank-revealing QR** (RRQR) factorization of matrix *A* is that satisfies *A* **=** *QR* **=** *Q* $\begin{pmatrix} R_1^2 & R_1^2 \\ 0 & R_2^2 \end{pmatrix}$ $\sigma_k(\bm{A})$ *p*1**(***k,m***)** \leq σ_{\min} (R_{11}) $\leq \sigma_{k}'$ (A) $\sigma_{k+1}(A) \leq \sigma_{\max}(R_{22}) \leq p_2(k,m)\sigma_{k+1}(A)$ Permutation matrix *n*-by-*n* orthogonal *n*-by-*m k*th singular value of *A* Some polynomial on *k* and *m k-*by*-k* upper-triangular w/ positive diagonal *k-*by*-*(*m–k*) *(n–k)-*by*-*(*m–k*)

CX and RRQR

- Let $\bf{A}\bf{\Pi} = \bf{Q}\bf{R}$ and let $\bf{\Pi}$ _{*k*} be the first *k* columns of **Π** and *C* = *A***Π***k* some *k* columns of *A*
	- Now $||A P_C A||_{\mathcal{E}} = ||R_{22}||_{\mathcal{E}}$, $\xi = F$ or 2
	- In particular $||\mathbf{A} \mathbf{P}_{\mathbf{C}}\mathbf{A}||_2 \leq p_2(k, m)||\mathbf{A} \mathbf{A}_k||_2$
		- \cdot $A_k = U_k \Sigma_k V_k^T$ (truncated SVD)
		- $\mathbf{C} \mathbf{X}$ is $p_2(k, m)$ -approximation to SVD

Computing CX by sampling

- Let $A = U\Sigma V^T$ be the input and its SVD and V_k the truncated *V*
- Sample columns of *A* with replacement
	- Probability *pj* for selecting column *j* is

$$
p_j = ||(\mathbf{V}_k^T)_j||_2^2 / k
$$

• Sample $O(k^2 \log(1/\delta)/\epsilon^2)$ columns and repeat log(1/δ) times returning the least-error sample

Notes on sampling

• We can prove that

 $||\mathbf{A} - \mathbf{P}_c\mathbf{A}||_F \leq (1 + \epsilon) ||\mathbf{A} - \mathbf{A}_k||_F$

with probability at least $1 - \delta$

- Notice that *C* has much more than *k* columns
	- $O(k^2 \log(1/\delta)/\epsilon^2)$ with large hidden constants

Why does sampling work?

- Intuitively, if *A* is of low rank (*k* ≪ *n*), *A* should have many almost-similar columns
- If we sample many columns enough, we should get a representative for each set of similar columns
	- ⇒ We need to sample more columns than the rank
		- Or our error depends on the rank…

CX with exact *k*

- Construct larger-than-k CX decomposition as above for $c =$ O(*k* log *k*) columns (and using rank-*k* truncated SVD)
	- Let Π_1 be the *m*-by-*c* matrix that selects *c* columns s.t. $C = A\Pi_1$
	- Let D_1 be *c*-by-*c* diagonal s.t. if *j*th column is selected on round *i*, $(D_1)_{ii} = (cp_j)$ –1/2 From $A = U\Sigma V^T$
- Run RRQR algorithm for $V_k \Pi_1 D_1$ to select exactly *k* columns *T* of *V^k T* $\Pi_1\mathcal{D}_1$ with matrix $\boldsymbol{\varPi}_2$ (*c*-by-*k*)
	- return $\mathbf{C} = \mathbf{A}\mathbf{\Pi}_1\mathbf{\Pi}_2$

Notes on the exact-*k* **CX**

- Pr[$||A P_cA||_F \le \Theta(k \log^{1/2} k) ||A A_k||_F] \ge 0.8$
- The sampling phase still requires really many columns (high hidden constants)
	- But in practice something like *c* = 5*k* works
- Any RRQR algorithm can be used for the second step
	- But the analysis depends on the chosen algorithm

Non-Negative CX

Motivation

- If data is non-negative, so is *C*
	- But *X* can contain negative values in standard CX
	- Non-negative *X* yields "parts-of-whole" interpretation similar to NMF
		- Selected columns are "pure" while others are mixtures of the pure columns
	- Non-negativity also improves sparsity

The non-negative CX decomposition

• In the **non-negative CX decomposition** (NNCX) we are given a non-negative matrix *A* and a rank *k*, and we need to select *k* **columns** of *A* into matrix *C* and build a nonnegative matrix *X* s.t. we minimize ||*A* – *CX*||*^F*

Geometry of NNCX

Columns in *C* Columns not in *C* Convex cone Projections

Cones and columns

- Consider the cone spanned by columns of *A*, cone(*A*)
	- If removing column *j* of *A* changes the cone, that column is **extremal**
		- Otherwise it is **internal**
	- Selecting all extremal columns to *C* gives us *A* = *CX* with nonnegative *X*

Algorithm for NNCX

- When we cannot select all extremal columns, we must choose which of them to select
	- Our goal is to maximize the volume of the convex cone
	- Finding the extremal columns is not easy
	- Given the columns, we must compute the non-negative projection

The convex_cone algorithm

- Set *R* ← *A*
- **• repeat**
	- **•** Select column *c* with highest norm in the residual *R*
		- **•** Normalize *c* to unit norm
	- **•** Solve nonnegative *x* that minimizes ||*R cx T* ||
	- **•** Set *R* ← *R cx T*
- **• until** *k* columns are selected
- **•** Set *C* to the columns of *A* corresponding to the selected *c* and solve nonnegative *X* minimizing $||A - CX||_F$

Solving for non-negative *X*

- Given *C*, finding non-negative *X* is the same as with NMF
	- Convex optimization with linear constraints
	- Or truncated-to-zero pseudo-inverse

Application: Neuroimaging

- Record brain cell activity over time
	- Every row is one frame
- Assume some columns contain the pure glomerulus signal
	- *C* identifies these signals
	- *X* explains how the signals are mixed in the brain images

found locations marked Movie frame with real and

Algorithm

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Application cont'd in space the signals from *T* are located. Successful selection of pure signals into *T* leads to sparse images in *S* (cp. Section 7.4.5) that allow for interpretation of the time series and images as the signal and shape of glomeruli.

Top-10 rows of X from NNCX decomposition shows performed on a calcium imaging movie of the honeybee AL (as in Section 7.5.1). the shape and location of glomeruli

Column-Row Decompositions

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The CUR decomposition

- In the **CUR decomposition** we are given matrix *A* and integers *c* and *r*, and our task is to select *c* columns of *A* to matrix *C* and *r* rows to matrix *R*, and build *c*-by-*r* matrix *U* minimizing ||*A* – *CUR*||*^F*
	- Often $c = r = k$

Why CUR?

- If selecting the actual columns in CX is good, selecting the actual columns and rows must be even better
	- We find prototypical columns *and rows*
	- *U* is usually small, so if *C* and *R* are sparse, storing CUR takes little space

Solving CUR: general idea

- CUR is two-sided CX
- Simple algorithm idea:
	- Solve CX for **A** and **A**^{*T*} and solve for *U* given **C** and *R*
		- \cdot $U = C^+AR^+$
- Better algorithms take into account the columns selected to *C* when computing *R*

Simple CUR algorithm

- Sample columns proportional to their L₂-norm
- Sample rows proportional to their *L*₂-norm
- Build $W = A[R, C]$ (the sub-matrix of columns in *R* and rows in *C*)
	- Let $W = X\Sigma Y^T$ be an SVD of W, and set $U \leftarrow Y(\mathbf{\Sigma}^+)^2 \mathbf{X}^T$

Fancier CUR algorithm

- Find *C* similar to exact-*k* CX earlier
	- Sample *O*(*k*/ε) additional columns
- Find $\mathbf{Z} \in \text{span}(\mathbf{C})$, $\mathbf{Z}^T \mathbf{Z} = \mathbf{I}$, such that $||A^T - A^TZZ^T||_F \leq (1 + O(\epsilon)) ||A - CX^*||_F$ Optimal X
	- Use *Z* to get the probabilities for sampling O(*k* log *k*) rows of *A* and reduce that to O(*k*) rows
	- Sample *O*(*k*/ε) additional rows
- Set $\boldsymbol{U} = \boldsymbol{X}^* \boldsymbol{Z}^T \boldsymbol{A} \boldsymbol{R}^+$

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Comments on the Boutsidis & Woodruff algorithm

- Slight variations of the above algorithm achieve:
	- selects the smallest number of rows and columns for $(1+\epsilon)$ approximation
	- matrix *U* has the smallest possible rank

CX and CUR summary

- Rows and columns of the original data should be interpretable
	- Also admit local constraints in the data
- CX and CUR decompositions are forms of feature selection
	- Applications when we need "prototypical" rows and columns

Literature

- Drineas, Mahoney & Muthukrishnan (2006): *Subspace sampling and relative-error matrix approximation: Column-based methods*. In APPROX/RANDOM '06
- Boutsidis, Mahoney & Drineas (2010): *An improved approximation algorithm for the column subset selection problem*. arXiv:0812.4293v2
- Boutsidis & Woodruff (2014): *Optimal CUR matrix decompositions*. arXiv:1405.7910v2
- Strauch (2014): *Column subset selection with applications to neuroimaging data.* PhD thesis, U. Konstanz