

Chapter 9

Matrix completion & recommender systems

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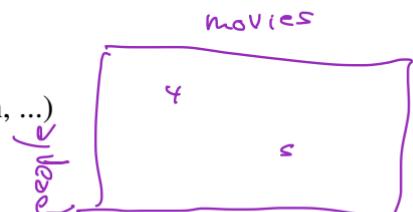
9.0 Introduction

This chapter discusses the important problem of **matrix completion**, where we know some, but not all, elements of a matrix and want to “complete” the matrix by filling in the missing entries. Obviously this problem is **ill posed** in general because one could assign arbitrary values to the missing entries, unless one assumes some **model** for the matrix elements. The most common model is that the matrix is **low rank**.

A particularly famous application of **low-rank matrix completion (LRMC)** is the **Netflix problem**; this topic is also relevant to dynamic MR image reconstruction, and numerous other applications with missing data (incomplete observations).

Successful applications of LRMC include the following.

- **Recommender systems** using **collaborative filtering** (netflix, spotify, amazon, ...)
- Imaging: denoising, reconstruction in medical, hyper-spectral imaging
- Anomaly detection in network flows
- Source localization and target tracking in radar and sonar
- Computer vision: background subtraction, object tracking, and to represent a single scene under varying illuminations
- Environmental monitoring of soil and crop conditions, water contamination, and air pollution, also sensor calibration
- Seismological activity and modal estimation in materials and human-made structures.



9.1 Measurement model

If \mathbf{X} is a **latent** $M \times N$ “low-rank” matrix with rank $r \leq \min(M, N)$, then

$$\mathbf{X} = \sum_{k=1}^r \sigma_k \mathbf{u}_k \mathbf{v}'_k.$$

Suppose we observe only *some* of the entries of \mathbf{X} in a sampling set Ω , i.e.,

$$Y_{ij} = \begin{cases} \overset{\text{noiseless}}{X_{ij}}, & (i,j) \in \Omega \\ ? , & \text{otherwise,} \end{cases} \quad \Omega \subseteq \{1, \dots, M\} \times \{1, \dots, N\},$$

where “?” denotes a value is “missing,” e.g., the i th user has not rated the j th movie.

Key questions:

- For a given (known) sampling set Ω , can we recover all of \mathbf{X} from just the observed entries \mathbf{Y} ?
The answer will depend on the rank r , where typically $r \ll \min(M, N)$, and on the sampling set.
- How do we do it?
and on $|\Omega| \ll MN$
- How well does it work (e.g., in the presence of noise or if \mathbf{X} is not exactly low rank)?

Practical implementation

This type of topic is so important to modern data analysis problems that JULIA has its own data type, `Missing`, to represent missing values. Try this:

```
Y = [2 0; missing 4; 5 missing]
```

$$\begin{bmatrix} 2 & 0 \\ ? & 4 \\ 5 & ? \end{bmatrix}$$

The type of `Y` is `3x2 Array{Union{Missing, Int64},2}`

because it is a 3×2 array of values that are either 64-bit integers or the special type `Missing`.

We cannot use any particular value like “0” to represent a missing value because 0 might be one of the observed values!

1. What is the minimum possible rank of `Y` in this example?

A: 0

B: 1

C: 2

D: 3

E: None of these.

??

$$\boxed{X} = \underbrace{\tilde{U}_r}_{\text{rank } r} \underbrace{\Sigma_r}_{\tilde{U}_r \cdot \tilde{V}_r'} \underbrace{\tilde{V}_r'}_{r \times N}$$

Sampling conditions for LRMC

The first question to ask is whether it should even be possible to recover \tilde{X} from \tilde{Y} in the noiseless case.

If X has rank r , then we can write X as the product of a $M \times r$ matrix (think $\tilde{U}_r = U_r \Sigma_r$ for example) with the transpose of a $N \times r$ matrix \tilde{V}_r . The number of **degrees of freedom (DoF)** in this representation looks to be $Mr + Nr = (M + N)r$. However, actually there are fewer DoF because once we fix \tilde{U}_r there are constraints on the values that \tilde{V}_r can take and still have $\tilde{U}_r \tilde{V}_r' = X$. So $(M + N)r$ is an upper bound on the DoF. The total number of elements of X is MN , so if the matrix is low rank, then $(M + N)r \ll MN$ so it seems plausible that having around $(M + N)r$ samples might suffice.

A more careful analysis of the DoF is

$$\text{DoF} = rM + r(N-r) = \frac{r(M+N)-r^2}{2Nr - r^2}$$

because

- the first (if we permute appropriately) r columns (of length M) are linearly independent,
- and the remaining $N - r$ columns depend linearly on those first r columns and we need r coefficients per column to describe the linear combination.

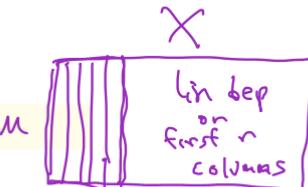
Note that if $r = N \leq M$ then $(M + N)r - r^2 = (M + N)N - N^2 = MN$, as expected.

When $M = N$ and $r \leq N/2$, a **lower bound on the number of samples** is $4Nr - 4r^2$ [1].

If $M \approx N$, then the **DoF $\approx 2Nr$** . So we need at least $O(Nr)$ samples to have any chance of recovery (unless we have additional information such as knowing that U_r and/or V_r is sparse).

Modern algorithms need $O(Nr \text{ polylog}(N))$ random samples under certain assumptions about X .

$$|\mathcal{S}| \geq O(Nr \text{ polylog}(N))$$



2. Approximately how many observations per column are the minimum needed to recover a rank- r low-rank matrix of size $N \times N$?

A: r B: $2r$ C: r^2 D: N E: $2N$

??

$$\frac{2Nr - r^2}{N} \approx 2r$$

Sampling mask

Define a $M \times N$ binary **sampling mask** matrix \mathbf{M} in terms of the sampling set Ω by:

$$M_{ij} \triangleq \begin{cases} 1, & (i, j) \in \Omega, \\ 0, & \text{otherwise.} \end{cases} \quad (9.1)$$

In words, \mathbf{M} is zero wherever we are missing a value in \mathbf{Y} .

Example. For the earlier 3×2 example we have $\mathbf{M} = \text{. !ismissing.}(\mathbf{Y})$, i.e., $\mathbf{M} = \begin{bmatrix} 1 & 1 \\ 0 & 1 \\ 1 & 0 \end{bmatrix}$.

Hereafter, let $\mathbf{Y} \in \mathbb{F}^{M \times N}$ be an matrix where

$$Y_{ij} = \begin{cases} X_{ij}, & (i, j) \in \Omega \\ 0, & \text{otherwise.} \end{cases} \quad (9.2)$$

$$\begin{bmatrix} 2 & 0 \\ 0 & 1 \\ 5 & 0 \end{bmatrix}$$

Now we *can* store a 0 in the locations of missing values because we have the separate mask \mathbf{M} to keep track of which values are sampled and which are missing.

9.2 LRMC: noiseless case

Noiseless problem statement

The (noiseless) LRMC problem is to find a matrix \hat{X} that is low-rank and that agrees with the measurements \mathbf{Y} on Ω , i.e., we want

$$\mathcal{M} \odot \hat{X} = \mathcal{M} \odot \mathbf{Y} \quad \textcircled{1} \quad \textcircled{2}$$

where \odot denotes element-wise multiplication, i.e., $\mathbf{M} \cdot \star \mathbf{X}$ in JULIA and MATLAB, also known as the **Hadamard product** or **array multiplication** (as opposed to **matrix multiplication**) and is applicable to arrays of any dimension.

(In Python, $\mathbf{M} \star \mathbf{X}$ is the Hadamard product for `array` objects in NumPy, but is ordinary matrix multiplication for `matrix` objects.) 

One “ideal” (noiseless) optimization formulation for the LRMC problem is [2]:

$$\hat{X} = \arg \min_{\mathbf{X}} \text{rank}(\mathbf{X}) \quad \text{st} \quad \mathcal{M} \odot \mathbf{X} = \mathcal{M} \odot \mathbf{Y} \quad (9.3)$$

This is a **non-convex** problem and is **NP hard** (impractical). No fast algorithm exists that is guaranteed to solve this problem formulation. Nevertheless, there are practical algorithms that often work reasonably well.



Alternating projection approach to LRMC

An alternative to (9.3) is to pick a rank $1 \leq K \ll \min(M, N)$ and seek a matrix $\hat{\mathbf{X}}$ that satisfies

$$\hat{\mathbf{X}} \in \mathcal{C} \cap \mathcal{D} \quad \mathcal{C} \triangleq \left\{ \mathbf{X} \in \mathbb{F}^{M \times N} : \underbrace{\text{rank}(\mathbf{X}) \leq K}_{\text{defn}} \right\}, \quad \mathcal{D} \triangleq \left\{ \mathbf{X} \in \mathbb{F}^{M \times N} : \mathbf{M} \odot \mathbf{X} = \mathbf{M} \odot \mathbf{Y} \right\}.$$

In words, we seek a matrix $\hat{\mathbf{X}}$ having rank at most K and that agrees with the measurements \mathbf{Y} on Ω .

The **projections onto convex sets (POCS)** approach is a classic signal processing tool for finding points in the intersection of two or more convex sets.

3. Which (if any) of the two sets \mathcal{C} and \mathcal{D} above is **convex**?

- A: neither \mathcal{C} nor \mathcal{D} B: \mathcal{C} but not \mathcal{D} C: \mathcal{D} but not \mathcal{C} D: both \mathcal{C} and \mathcal{D}

??

see Ch 6

To analyze the set \mathcal{D} :

$$\begin{aligned} \underline{\mathbf{X}, \mathbf{Z} \in \mathcal{D}} &\implies \mathbf{M} \odot \mathbf{X} = \mathbf{M} \odot \mathbf{Y} \quad \text{and} \quad \mathbf{M} \odot \mathbf{Z} = \mathbf{M} \odot \mathbf{Y} \\ &\implies \mathbf{M} \odot (\alpha \mathbf{X} + (1-\alpha)\mathbf{Z}) = \alpha \mathbf{M} \odot \mathbf{X} + (1-\alpha) \mathbf{M} \odot \mathbf{Z} \quad \text{by Hadamard distr. prop.} \\ &\implies \alpha \mathbf{M} \odot \mathbf{Y} + (1-\alpha) \mathbf{M} \odot \mathbf{Y} = \mathbf{M} \odot \mathbf{Y} \quad \forall \mathbf{X}, \mathbf{Z} \in \mathcal{D} \Rightarrow \mathcal{D} \text{ convex} \end{aligned}$$

convex combination

3'

\mathcal{D} a subspace?

- A. True B. False

"Flat"

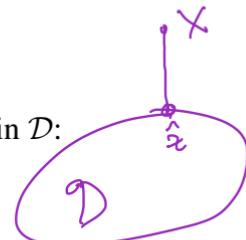
The **alternating projection** method alternates between projecting onto the sets \mathcal{C} and \mathcal{D} :

"POCS"

$$\mathbf{X}_{k+1} = \mathcal{P}_{\mathcal{C}}(\mathcal{P}_{\mathcal{D}}(\mathbf{X}_k))$$

where $\mathcal{P}_{\mathcal{D}}(\mathbf{X})$ denotes the **projection** of its argument \mathbf{X} onto the set \mathcal{D} , i.e., the closest point in \mathcal{D} :

$$\mathcal{P}_{\mathcal{D}}(\mathbf{X}) = \arg \min_{\mathbf{Z} \in \mathcal{D}} \|\mathbf{X} - \mathbf{Z}\|_F.$$



The algorithm designer gets to choose the norm for quantifying “closest.” Here we are using matrices so we work with the simplest choice, the **Frobenius norm**.

For the “data” set \mathcal{D} above, the **projection operation is very simple**:

$$[\mathcal{P}_{\mathcal{D}}(\mathbf{X})]_{i,j} = \begin{cases} Y_{i,j}, & (i,j) \in \Omega \\ X_{i,j}, & \text{otherwise.} \end{cases}$$

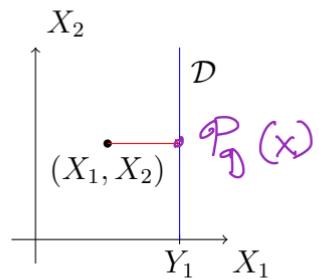
Example. Suppose $(M, N) = (2, 1)$ and $\mathbf{M} = \begin{bmatrix} 1 \\ 0 \end{bmatrix}$ so $\mathbf{Y} = \begin{bmatrix} Y_1 \\ ? \end{bmatrix}$.

$\left[\begin{array}{c} Y_1 \\ Y_2 \end{array} \right]$
Then \mathcal{D} is just $\{(X_1, X_2) \in \mathbb{R}^2 : X_1 = Y_1\}$.

For any point (X_1, X_2) , the projection onto \mathcal{D} is simply (Y_1, X_2) .

If \mathbf{M} is a logical array, then we can implement this operation “in place” in JULIA as follows:

$\mathbf{X}[\mathbf{M}] .= \mathbf{Y}[\mathbf{M}]$



Even though \mathcal{C} is not convex, finding a projection onto \mathcal{C} is fairly easy using Ch. 6:

$$\mathcal{P}_{\mathcal{C}}(\mathbf{Z}) = \underbrace{\arg \min_{\mathbf{L} : \text{rank}(\mathbf{L}) \leq K}}_{\in} \|\mathbf{Z} - \mathbf{L}\|_F = \mathbf{U}_K \boldsymbol{\Sigma}_K \mathbf{V}'_K, \quad \mathbf{Z} = \mathbf{U} \boldsymbol{\Sigma} \mathbf{V}'.$$

This is simply the low-rank approximation problem of Ch. 6. It requires an SVD of the input argument.

4. The projection $\mathcal{P}_{\mathcal{C}}(\mathbf{Z})$ is **unique** for any $M \times N$ input matrix \mathbf{Z} . (?)

A: True

B: False

??

if $\sigma_{F+}(z) < \sigma_k(z)$

5. Given $K = 1$, $\mathbf{Y} = \begin{bmatrix} 0 & 0 \\ 0 & 2 \end{bmatrix}$, $\mathbf{M} = \begin{bmatrix} 0 & 1 \\ 1 & 1 \end{bmatrix}$, $\mathbf{X}_k = \begin{bmatrix} 3 & 0 \\ 0 & 0 \end{bmatrix}$, the **POCS** update $\mathbf{X}_{k+1} = \mathcal{P}_{\mathcal{D}}(\mathcal{P}_{\mathcal{D}}(\mathbf{X}_k))$ is:

A: $\begin{bmatrix} 2 & 2 \\ 2 & 2 \end{bmatrix}$

B: $\begin{bmatrix} 0 & 0 \\ 0 & 2 \end{bmatrix}$

C: $\begin{bmatrix} 3 & 0 \\ 0 & 2 \end{bmatrix}$

D: $\begin{bmatrix} 3 & 0 \\ 0 & 0 \end{bmatrix}$

E: $\begin{bmatrix} 3 & 3 \\ 3 & 3 \end{bmatrix}$

??

alt proj.

$$\mathcal{P}_{\mathcal{D}}(\mathbf{X}_k) = \begin{bmatrix} 3 & 0 \\ 0 & 2 \end{bmatrix}$$



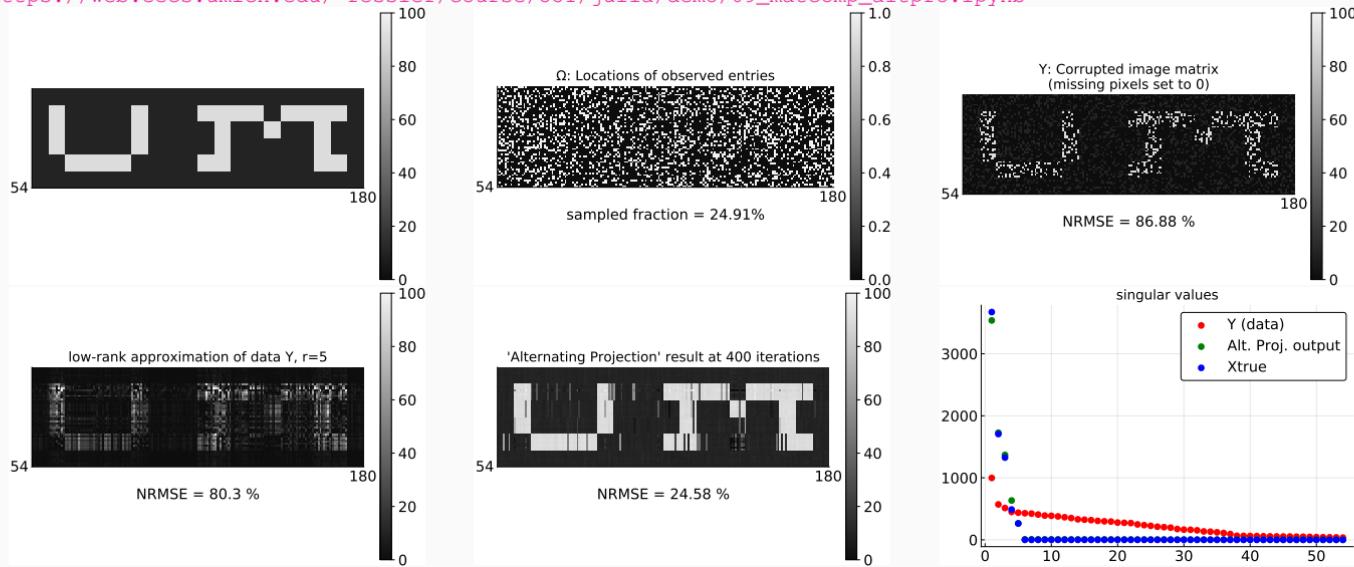
fixed point

$$\mathcal{P}_{\{X: \text{rank}(X) \leq 1\}} \left(\begin{bmatrix} 3 & 0 \\ 0 & 2 \end{bmatrix} \right) = \begin{bmatrix} 3 & 0 \\ 0 & 0 \end{bmatrix}$$

Example. This demo shows matrix completion via alternating projection with 25% sampling.

https://web.eecs.umich.edu/~fessler/course/551/julia/demo/09_matcomp_altpro.html

https://web.eecs.umich.edu/~fessler/course/551/julia/demo/09_matcomp_altpro.ipynb



The alternating projection method works better than simply making a low-rank approximation to \mathbf{Y} , but it does not account for noise in the data, so we can do better.

Convergence of alternating projection method

Because the set of rank $\leq K$ matrices is nonconvex, it is hard to say much about the convergence of the alternating projection method. Choice of initial guess \mathbf{X}_0 can affect the convergence behavior.

Example. Consider $\mathbf{Y} = \begin{bmatrix} 1 & 1 \\ ? & ? \end{bmatrix}$ with rank $K = 1$.

If $\mathbf{X}_0 = \begin{bmatrix} 1 & 1 \\ x & x \end{bmatrix}$ for any value x , then $\mathbf{X}_k = \mathbf{X}_0$ for all $k \in \mathbb{N}$.

So in this case, the set of “optimal” solutions $\mathcal{C} \cap \mathcal{D}$ is uncountably infinite!

Challenge. Find a more interesting case (if possible) where some $\mathbf{X}_0 \in \mathcal{D}$ eventually leads to a fixed point \mathbf{X}_k of the alternating projection method that is not in the set $\mathcal{C} \cap \mathcal{D}$, i.e., \mathbf{X}_k is in \mathcal{C} but not in \mathcal{D} . Assume some constraint on K , like that it has to be at least as large as the minimum possible rank of \mathbf{Y} (so the intersection of \mathcal{C} and \mathcal{D} is non-empty).

Choice of rank

In practice one must choose the rank K to perform low-rank matrix completion.

One can use **cross validation** or **minimum description length (MDL)** methods [3] among others [4].

See survey papers [5] [6].

9.3 LRMC: noisy case

Noisy problem statement

In practice, often there is noise in the data, so the model (9.2) is often unrealistic. A more realistic model is

$$Y_{ij} = \begin{cases} X_{ij} + \varepsilon_{ij}, & (i, j) \in \Omega \leftarrow \text{sampling set} \\ 0, & \text{otherwise.} \end{cases}$$

Again, it is fine to store a 0 (or any other value) in the missing locations because we have the sampling mask M available.

In this case, it would be unreasonable to insist on exact data equality $\cancel{M \odot X = M \odot Y}$.

So now we want to find \hat{X} where $\cancel{M \odot \hat{X} \approx M \odot Y}$, i.e., $\cancel{M \odot (\hat{X} - Y) \approx 0}$, and also \hat{X} is low-rank.

One possible LRMC formulation uses a (non-convex) **rank constraint**:

$$\hat{X} = \underset{\substack{\text{set} \\ X: \text{rank}(X) \leq K}}{\operatorname{argmin}} \|M \odot (X - Y)\|_F \quad (9.4)$$

Another possible LRMC formulation uses a (non-convex) **rank regularizer**:

$$\hat{X} = \underset{\substack{\text{non-convex function} \\ X \in \mathbb{R}^{m \times n}}}{\operatorname{argmin}} \|M \odot (X - Y)\|_F^2 + \beta \text{rank}(X) \quad (9.5)$$

The following LRMC optimization formulation is the **convex relaxation** of the rank regularizer, using instead a **nuclear norm** regularizer:

$$\hat{\mathbf{X}} = \arg \min_{\mathbf{X}} \frac{1}{2} \|\mathbf{M} \odot (\mathbf{X} - \mathbf{Y})\|_F^2 + \beta \|\mathbf{X}\|_* \quad (9.6)$$

There are fast and practical algorithms for all three of these formulations, and there are convergence guarantees for the convex formulation.

Unfortunately, the optimization tools presented in Ch. 8 (GD, PGD, PSD) are inapplicable to the above problems either because the cost function is not differentiable or it involves a nonconvex set.

Even the cost function (9.6) that uses the (convex) nuclear norm $\|\mathbf{X}\|_*$ is not differentiable.

So we need an approach that is more sophisticated than gradient-based methods.

6. What is nuclear norm $\|\mathbf{x}\|_*$ of a 1×1 "matrix" $[x]$?

A: x

B: $|x|$

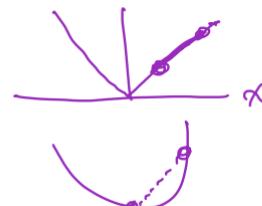
C: $\mathbb{I}_{\{x \geq 0\}}$

D: $\max(0, x)$

E: x^2

??

$$x = \begin{cases} \text{sign}(x) |x| & \text{if } x \neq 0 \\ 0 & \text{if } x = 0 \end{cases}$$



- not differentiable
- convex
- not strictly convex

Majorize-minimize (MM) iterations

To solve optimization problems like (9.6), we first consider **majorize-minimize (MM)** algorithms.

To solve an optimization problem like

$$\arg \min_{x \in \mathcal{X}} \Psi(x)$$

we first design a **majorizer** or **surrogate function** $\phi_k(x)$ that satisfies the following two conditions:

$$\begin{aligned} \Psi(x_k) &= \phi_k(x_k) && \text{matched value} \\ \Psi(x) &\leq \phi_k(x) && \forall x \text{ lies above} \end{aligned}$$

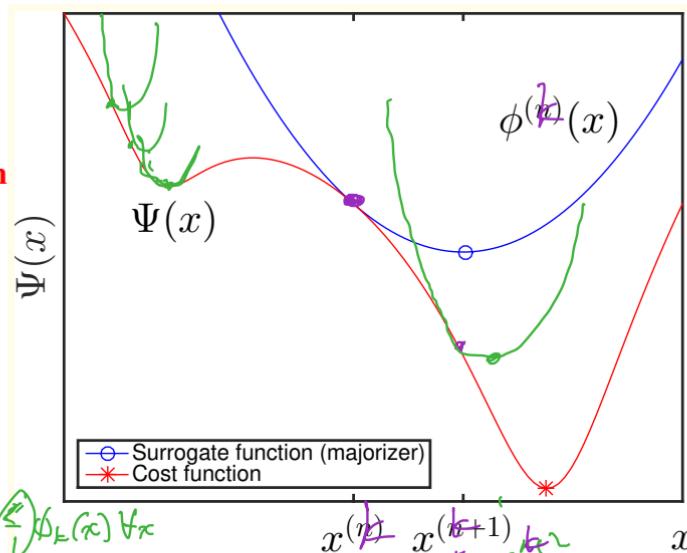
Then the MM algorithm update is simply:

$$x_{k+1} = \arg \min_{x \in \mathcal{X}} \phi_k(x).$$

$$\phi_k(x_{k+1}) \leq \phi_k(x) \quad \forall x$$

Any MM algorithm will monotonically decrease the cost function because

$$\Psi(x_{k+1}) \leq \phi_k(x_{k+1}) \leq \phi_k(x_k) = \Psi(x_k).$$



Sandwich
inequality

Next we design a MM algorithm for LRMC problems like (9.6).

MM methods for LRM

First define the complement of the mask matrix as follows:

$$\widetilde{\mathbf{M}} \triangleq \mathbf{1}_M \mathbf{1}'_N - \mathbf{M}, \quad \widetilde{M}_{i,j} = \begin{cases} 0, & (i,j) \in \Omega \\ 1, & (i,j) \notin \Omega. \end{cases}$$

$$\widetilde{\mathbf{M}} = \mathbf{1}_M \mathbf{1}'_N - \mathbf{M}$$

We focus now on the quadratic data-fit term in (9.4) (9.5) (9.6):

$$q(\mathbf{X}) \triangleq \|\mathbf{M} \odot (\mathbf{X} - \mathbf{Y})\|_F^2.$$

Now define the following quadratic function:

$$Q(\mathbf{X}; \mathbf{Z}) \triangleq \|\mathbf{X} - \mathbf{Z} + \mathbf{M} \odot (\mathbf{Z} - \mathbf{Y})\|_F^2 = \|\mathbf{X} - (\widetilde{\mathbf{M}} \odot \mathbf{Z} + \mathbf{M} \odot \mathbf{Y})\|_F^2 \quad (9.7)$$

no \mathbf{M} here!!

This function is a **majorizer** of $q(\mathbf{X})$ because

$$\begin{aligned} q(\mathbf{X}) &= Q(\mathbf{X}; \mathbf{X}) \\ q(\mathbf{X}) &\leq Q(\mathbf{X}; \mathbf{z}) \quad \forall \mathbf{z} \in \mathbb{R}^{m \times n} \end{aligned}$$

Proof of majorization inequality:

(Read)

$$\begin{aligned} Q(\mathbf{X}; \mathbf{Z}) &= \|\mathbf{X} - \mathbf{Z} + \mathbf{M} \odot (\mathbf{Z} - \mathbf{Y})\|_F^2 = \|(\widetilde{\mathbf{M}} + \mathbf{M}) \odot (\mathbf{X} - \mathbf{Z}) + \mathbf{M} \odot (\mathbf{Z} - \mathbf{Y})\|_F^2 \\ &= \|\widetilde{\mathbf{M}} \odot (\mathbf{X} - \mathbf{Z}) + \mathbf{M} \odot (\mathbf{X} - \mathbf{Y})\|_F^2 = \|\widetilde{\mathbf{M}} \odot (\mathbf{X} - \mathbf{Z})\|_F^2 + \|\mathbf{M} \odot (\mathbf{X} - \mathbf{Y})\|_F^2 \\ &\geq \|\mathbf{M} \odot (\mathbf{X} - \mathbf{Y})\|_F^2 = q(\mathbf{X}), \end{aligned}$$

where the 4th equality holds because $\widetilde{\mathbf{M}} \odot \mathbf{M} = \mathbf{0}$.

The design of the function (9.7) uses a 2nd-order Taylor expansion of $q(\mathbf{X})$ about \mathbf{Z} and then uses the fact that the elements of \mathbf{M} are all 0 or 1. The above proof verifies that Q is a majorizer, which is all that is needed here. ♦♦

Now we use the majorizer (9.7) to develop a few different LRMC algorithms.

LRMC by iterative low-rank approximation

First consider LRMC using the rank constraint (9.4):

$$\hat{\mathbf{X}} = \arg \min_{\mathbf{X} : \text{rank}(\mathbf{X}) \leq K} \| \mathbf{M} \odot (\mathbf{X} - \mathbf{Y}) \|_{\text{F}}^2.$$

The MM algorithm update for this formulation is simply

$$\begin{aligned}\mathbf{X}_{k+1} &= \arg \min_{\mathbf{X} : \text{rank}(\mathbf{X}) \leq K} Q(\mathbf{X}; \mathbf{X}_k) \\ &= \arg \min_{\mathbf{X} : \text{rank}(\mathbf{X}) \leq K} \underbrace{\| \mathbf{X} - \tilde{\mathbf{X}}_k \|_{\text{F}}^2}_{\text{yellow box}} \\ &= \arg \min_{\mathbf{X} : \text{rank}(\mathbf{X}) \leq K} \| \mathbf{X} - \tilde{\mathbf{X}}_k \|_{\text{F}}^2, \quad \tilde{\mathbf{X}}_k \triangleq \widetilde{\mathbf{M}} \odot \mathbf{X}_k + \mathbf{M} \odot \mathbf{Y} = \begin{cases} Y_{i,j}, & (i, j) \in \Omega \\ [\mathbf{X}_k]_{i,j}, & (i, j) \notin \Omega. \end{cases} \Rightarrow \mathcal{P}_{\mathcal{D}}(\mathbf{X}_k)\end{aligned}$$

This algorithm alternates between the following two steps.

- Take the current guess \mathbf{X}_k and replace all the values at sampled locations with the measurements from \mathbf{Y} to get $\tilde{\mathbf{X}}_k$. (As mentioned earlier, this can be done “in place” using $\mathbf{X}[\mathbf{M}] . = \mathbf{Y}[\mathbf{M}]$)
- Perform low-rank (rank at most K) approximation (using **SVD**) to $\tilde{\mathbf{X}}_k$ to get the next iterate \mathbf{X}_{k+1} .

This process is exactly the same as the alternating projection method described earlier. So now we know that it is a MM method that decreases the Frobenius norm cost function monotonically.

(There is still no guarantee of convergence of $\{\mathbf{X}_k\}$ to a global minimizer because the rank constraint set is nonconvex.)

LRMC by iterative singular value hard thresholding

Now consider LRMC using the rank regularizer (9.5):

$$\hat{\mathbf{X}} = \arg \min_{\mathbf{X} \in \mathbb{F}^{M \times N}} \| \mathbf{M} \odot (\mathbf{X} - \mathbf{Y}) \|_{\text{F}}^2 + \beta \operatorname{rank}(\mathbf{X}).$$

The MM algorithm update for this formulation is simply

$$\begin{aligned}\mathbf{X}_{k+1} &= \arg \min_{\mathbf{X} \in \mathbb{F}^{M \times N}} Q(\mathbf{X}; \mathbf{X}_k) + \\ &\quad \underbrace{\phi_k(\mathbf{x})}_{\phi_k(\mathbf{x})} \\ &= \arg \min_{\mathbf{X} \in \mathbb{F}^{M \times N}} \| \mathbf{X} - (\tilde{\mathbf{M}} \odot \mathbf{X}_k + \mathbf{M} \odot \mathbf{Y}) \|_{\text{F}}^2 + \beta \operatorname{rank}(\mathbf{X}) \\ &= \arg \min_{\mathbf{X} \in \mathbb{F}^{M \times N}} \| \mathbf{X} - \tilde{\mathbf{X}}_k \|_{\text{F}}^2 + \beta \operatorname{rank}(\mathbf{X}).\end{aligned}$$

This algorithm alternates between the following two steps.

- Take the current guess \mathbf{X}_k and replace all the values at sampled locations with the measurements from \mathbf{Y} to get $\tilde{\mathbf{X}}_k$. (This can be done “in place.”)
- Apply **singular value hard thresholding** to $\tilde{\mathbf{X}}_k$ to get the next iterate \mathbf{X}_{k+1} .

This MM method decreases the rank-regularized Frobenius norm cost function monotonically.

LRMC by iterative singular value soft thresholding (ISTA)

Now consider LRMC using the **convex nuclear norm** regularizer (9.6):

$$\hat{\mathbf{X}} = \arg \min_{\mathbf{X} \in \mathbb{F}^{M \times N}} \frac{1}{2} \|\mathbf{M} \odot (\mathbf{X} - \mathbf{Y})\|_F^2 + \beta \|\mathbf{X}\|_*.$$

7. This cost function (also in the Demo) is (convex, strictly convex)?

A: F,F

B: F,T

C: T,F

D: T,T

??

The MM algorithm update for this formulation is simply

original $\underset{\mathbf{X}}{\arg \min} q(\mathbf{X}) + \beta \|\mathbf{X}\|_*$

$$\mathbf{X}_{k+1} = \arg \min_{\mathbf{X} \in \mathbb{F}^{M \times N}} \frac{1}{2} Q(\mathbf{X}; \mathbf{X}_k) + \beta \|\mathbf{X}\|_*$$

majorize

$$= \arg \min_{\mathbf{X} \in \mathbb{F}^{M \times N}} \frac{1}{2} \|\mathbf{X} - (\tilde{\mathbf{M}} \odot \mathbf{X}_k + \mathbf{M} \odot \mathbf{Y})\|_F^2 + \beta \|\mathbf{X}\|_* = \arg \min_{\mathbf{X} \in \mathbb{F}^{M \times N}} \frac{1}{2} \|\mathbf{X} - \tilde{\mathbf{X}}_k\|_F^2 + \beta \|\mathbf{X}\|_*$$

solved in Ch-6

This algorithm alternates between the following two steps.

- o Take the current guess \mathbf{X}_k and replace all the values at sampled locations with the measurements from \mathbf{Y} to get $\tilde{\mathbf{X}}_k$. (This can be done “in place.”)
- o Apply **singular value soft thresholding** to $\tilde{\mathbf{X}}_k$ to get the next iterate \mathbf{X}_{k+1} .

SVST

This MM method (aka **ISTA**) decreases its cost function monotonically. Because the cost function is convex, with some additional work one can show that the sequence $\{\mathbf{X}_k\}$ converges to a global minimizer.



Iterative soft-thresholding algorithm (ISTA)

In many modern applications, including LRMC, we have a **composite cost function** of the form:

$$\arg \min_{\mathbf{x}} f(\mathbf{x}) + g(\mathbf{x}) \quad (9.8)$$

where $f(\mathbf{x})$ is convex and has a Lipschitz gradient (smooth) but $g(\mathbf{x})$ is convex but *not necessarily* smooth.

To develop an algorithm for such problems, we first reinterpret the GD step for minimizing $f(\mathbf{x})$ as follows:

$$\begin{aligned} \mathbf{x}_{k+1} &= \arg \min_{\mathbf{x}} q(\mathbf{x}; \mathbf{x}_k) = \mathbf{x}_k - \alpha \nabla f(\mathbf{x}_k) \\ q(\mathbf{x}; \mathbf{x}_k) &\triangleq f(\mathbf{x}_k) + \langle \nabla f(\mathbf{x}_k), \mathbf{x} - \mathbf{x}_k \rangle + \frac{1}{2\alpha} \|\mathbf{x} - \mathbf{x}_k\|_2^2. \end{aligned}$$

Proof (by completing the square):

$$\begin{aligned} q(\mathbf{x}; \mathbf{x}_k) &= f(\mathbf{x}_k) + \langle \nabla f(\mathbf{x}_k), \mathbf{x} - \mathbf{x}_k \rangle + \frac{1}{2\alpha} \|\mathbf{x} - \mathbf{x}_k\|_2^2 \\ &= f(\mathbf{x}_k) - \frac{\alpha}{2} \|\nabla f(\mathbf{x}_k)\|_2^2 + \frac{1}{2\alpha} (\|\alpha \nabla f(\mathbf{x}_k)\|_2^2 + 2 \langle \alpha \nabla f(\mathbf{x}_k), \mathbf{x} - \mathbf{x}_k \rangle + \|\mathbf{x} - \mathbf{x}_k\|_2^2) \\ &= \left(f(\mathbf{x}_k) - \frac{\alpha}{2} \|\nabla f(\mathbf{x}_k)\|_2^2 \right) + \frac{1}{2\alpha} \|\mathbf{x} - (\mathbf{x}_k - \alpha \nabla f(\mathbf{x}_k))\|_2^2 \\ \implies \mathbf{x}_{k+1} &= \mathbf{x}_k - \alpha \nabla f(\mathbf{x}_k). \end{aligned}$$

This alternate perspective on GD is the key to extending the GD approach to composite cost functions like (9.8). Consider the following iteration:

$$\begin{aligned}\mathbf{x}_{k+1} &= \arg \min_{\mathbf{x}} \left\{ f(\mathbf{x}_k) + \langle \nabla f(\mathbf{x}_k), \mathbf{x} - \mathbf{x}_k \rangle + \frac{1}{2\alpha} \|\mathbf{x} - \mathbf{x}_k\|_2^2 + g(\mathbf{x}) \right\} \\ &= \arg \min_{\mathbf{x}} \left\{ \frac{1}{2\alpha} \|\mathbf{x} - (\mathbf{x}_k - \alpha \nabla f(\mathbf{x}_k))\|_2^2 + g(\mathbf{x}) \right\}.\end{aligned}$$

This algorithm is called **iterative soft-thresholding algorithm (ISTA)**, aka the **iterative shrinkage thresholding algorithm**, or more generally, the **proximal gradient method (PGM)**.

More concisely, ISTA uses the following two steps per iteration:

$$\begin{aligned}\tilde{\mathbf{x}}_k &\triangleq \mathbf{x}_k - \alpha \nabla f(\mathbf{x}_k) && \text{(usual GD step)} \\ \mathbf{x}_{k+1} &= \arg \min_{\mathbf{x}} \frac{1}{2\alpha} \|\mathbf{x} - \tilde{\mathbf{x}}_k\|_2^2 + g(\mathbf{x}) = \text{prox}_{\alpha g}(\tilde{\mathbf{x}}_k) && \text{(proximal step)}\end{aligned}\tag{9.9}$$

The first step is a conventional GD step. The second step is called a **proximity operation**; see Ch. 6.

The method is convergent [7] (in the sense of p. 5.41 in Ch. 5) when the convex part $f(\mathbf{x})$ is smooth, *i.e.*, its gradient $\nabla f(\mathbf{x})$ has Lipschitz constant L , and we choose $0 < \alpha < 2/L$. We do not need $f(\mathbf{x})$ to be quadratic! Convex and smooth (Lipschitz gradient) is sufficient.

Demo

Recall that Nesterov's accelerated gradient method converges faster than GD for problems with smooth cost functions. There are also accelerated versions of ISTA including (**FISTA**) [8] and (**POGM**) [9].

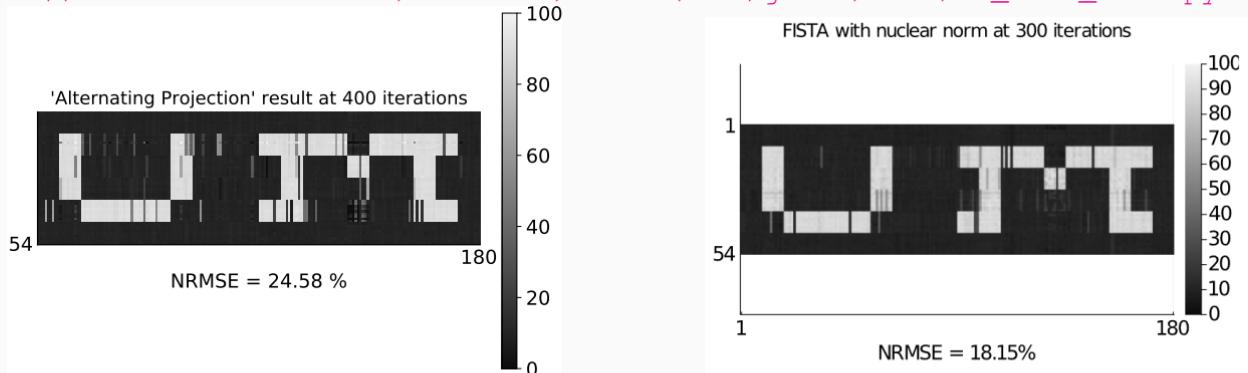
An alternative approach to the LRMC optimization problem is the ADMM method [10]. ADMM requires a tuning parameter for fast convergence. When well tuned, ADMM can be faster than FISTA.

Historically, the original work on this problem used an **EM algorithm** perspective [11].

Example. See the demo notebook created by Dr. Greg Ongie:

https://web.eecs.umich.edu/~fessler/course/551/julia/demo/09_lrmc_nuc.html

https://web.eecs.umich.edu/~fessler/course/551/julia/demo/09_lrmc_nuc.ipynb



9.4 Summary

Matrix completion using low-rank models is a rich area with numerous applications.

Low-rank matrix *approximation* is an easier problem and Ch. 6 provided nice SVD-based solutions.

LRMC is a more challenging problem (due to the missing data) so algorithms for LRMC typically are iterative.

Interestingly, LRMC algorithms usually use ingredients from low-rank matrix approximation. This pattern of using methods for simpler problems as part of some iterative approach for solving more complicate problem is quite common.

For image processing applications, often low-rank models are applied to collections of image patches, rather than to the entire image, *e.g.*, [12–16].

This topic is a very active research area, with a growing body of work related to convergence and performance guarantees, *e.g.*, [17] [18]. and heteroscedastic data [19].

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