# A non-backtracking method for long matrix and tensor completion

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## What is tensor completion ?



#### What is tensor completion ?



Tensor T

Observed tensor  $\tilde{T}$ 

- T is an order-k tensor of size  $n \times \cdots \times n$
- The observed tensor  $\tilde{T}$  is defined as

$$ilde{T}_{i_1,...,i_k} = egin{cases} T_{i_1,...,i_k} & ext{with probability } p \ 0 & ext{with probability } 1-p \end{cases}$$

 Goal: Exactly/approximately recover T from T with very few samples (with an efficient algorithm)



Too many degrees of freedom!



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Too localized!





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• T has low CP-rank:

$$T = \sum_{i=1}^{r} \lambda_i \left( w_i^{(1)} \otimes \cdots \otimes w_i^{(k)} \right)$$

 $\Rightarrow$  *r*  $\times$  *kn* degrees of freedom





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• T is delocalized:

$$\|w_i^{(j)}\|_{\infty} \simeq n^{-1/2}$$

## Matrix completion



• Low-rank matrix completion [Candes-Recht '09, Candes-Tao '10, Keshavan-Montanari-Oh '10,...]. When r = O(1), with high probability, uniformly sampling  $O(n \log(n))$  entries with convex/ non-convex optimization is sufficient to exactly recover M.

## Matrix completion



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- Information threshold O(rn log n) [Candes-Tao '10]. Best rank dependence: O(r log r · n log n) [Ding-Chen '20].

Computational complexity problem: most tensor problems are hard [Hillar-Lim '09]

- spectral norm
- eigenvalues/singular values
- low-rank approximations

## Unfolding



$$M_{(i_1,...,i_a),(i_{a+1},...,i_k)} = T_{i_1,...,i_k}$$

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Tensor completion on  $T \leftarrow$  Matrix completion on M

## Unfolding



"Grouping" indices:

$$M_{(i_1,...,i_a),(i_{a+1},...,i_k)} = T_{i_1,...,i_k}$$

Tensor completion on  $T \leftarrow$  Matrix completion on M

If k is even: square matrix of size  $n^{k/2} \implies \tilde{O}(n^{k/2})$  samples suffice If k is odd: matrix of size  $n^{\lfloor k/2 \rfloor} \times n^{\lceil k/2 \rceil}$ 

#### Statistical-computational gap for random tensors

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- Similar gaps in the spiked tensor model  $T = \lambda v^{\otimes q} + Z$ [Montanari-Richard '14, Ben Arous-Mei-Montanari-Nica '17, Chen '18, Ben Arous-Gheissari-Jagannath '18, Wein-Alaoui-Moore '19, Perry-Wein-Bandeira '20...]



Commonly poly-time algorithms: unfolding-based [Montanari and Sun '16, Liu and Moitra '20, Cai et al. '21...]

- Unfold  $\tilde{T}$  into  $A \in \mathbb{R}^{n \times n^2}$ . If  $T = \sum_{i=1}^r x_i \otimes y_i \otimes z_i$ , unfold  $\tilde{T}$  in 3 different ways.
- Take the SVD of the hollowed matrix h(AA<sup>T</sup>) = AA<sup>T</sup> − diag(AA<sup>T</sup>) (spectral initialization) + postprocessing
- Diagonal removal improved the performance [Cai et al. '21]
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What happens if  $p \propto n^{-k/2}$  ?



**Figure:**  $T = v \otimes v \otimes v, AA^{\top} - \text{diag}(AA^{\top}), p = 20n^{-3/2}$ 



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 $A \in \mathbb{R}^{n imes n^2}$  corresponds to a (weighted) random bipartite graph with  $V_1 = [n], V_2 = [n^2].$ 



#### A random graph theory explanation

Hollowed matrix counts walks of length 2,  $V_1 \rightarrow V_2 \rightarrow V_1$ :

$$(AA^{\top})_{ij} = \sum_{k} A_{ik} A_{jk}.$$

 $h(AA^{\top})$  can be seen as the adjacency matrix of a new graph  $\tilde{G}$  (dashed edges).



Fact:  $\tilde{G}$  is still sparse (average degree  $d^2$  for  $p = dn^{-k/2}$ ).

In the unweighted (Erdős-Rényi) case:

- if d<sup>2</sup> ≳ √ log(n) / log log(n): spectrum of G̃ concentrates [Feige-Ofek '05, Benaych-Georges-Bordenave-Knowles '20]
- if d<sup>2</sup> ≪ √ log(n)/log log(n): no concentration, spectrum dominated by high-degree vertices [Benaych-Georges-Bordenave-Knowles '19]

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In the unweighted (Erdős-Rényi) case:

- if  $d^2 \gtrsim \sqrt{\frac{\log(n)}{\log\log(n)}}$ : spectrum of  $\tilde{G}$  concentrates [Feige-Ofek '05, Benaych-Georges-Bordenave-Knowles '20]
- if d<sup>2</sup> ≪ √ log(n)/log log(n): no concentration, spectrum dominated by high-degree vertices [Benaych-Georges-Bordenave-Knowles '19]

## $\Rightarrow$ Naive unfolding (probably) doesn't work

Recap:

- existing methods do not reach the exact conjectured threshold for tensor completion (no results for "weak recovery").
- It is not a technical but a conceptual issue
- it suffices to solve matrix completion for a rank-r long matrix

Our solution: a new non-backtracking matrix for sparse long matrices Community detection in stochastic block models  $\mathcal{G}(n, \frac{a}{n}, \frac{b}{n})$ .

- Unknown partition  $\sigma \in \{-1,1\}^n$ . Generate a random graph G = ([n], E). *i*, *j* is connected with probability  $p = \frac{a}{n}$  if  $\sigma_i = \sigma_j$  and with probability  $q = \frac{b}{n}$  otherwise.
- goal: recover  $\sigma$  from G



## A detour through community detection

 $\mathbb{E}[A]$  is low-rank, and  $v_2(\mathbb{E}[A]) = \sigma \Rightarrow$  spectral method on A?

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 $\mathbb{E}[A]$  is low-rank, and  $v_2(\mathbb{E}[A]) = \sigma \Rightarrow$  spectral method on A? No!



 $p = \frac{a}{n}, q = \frac{b}{n}$ . High-degree vertices dominate the spectrum.  $v_2$  localized around high-degree vertices.

[Krivelevich-Sudakov '01, Benaych-Georges, Bordenave, Knowles '19, Alt-Ducatez-Knowles '23]

#### Non-backtracking matrix for graphs

Proposed in [Krzakala et al. '13]

Defined on the oriented edges of G:

$$\vec{E} = \{u \to v : \{u, v\} \in E\}, |\vec{E}| = 2|E|.$$

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The non-backtracking matrix *B* is defined: for  $u \rightarrow v, x \rightarrow y \in \vec{E}$ ,

$$B_{u\to v,x\to y}=\mathbf{1}_{v=x}\mathbf{1}_{u\neq y}.$$



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#### Non-backtracking spectral method



 If (a - b)<sup>2</sup> > 2(a + b), then the second eigenvector of B can be used to detect the community structure. [Bordenave, Lelarge, Massoulié '18]

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- If (a b)<sup>2</sup> > 2(a + b), then the second eigenvector of B can be used to detect the community structure. [Bordenave, Lelarge, Massoulié '18]
- *B* is non-Hermitian: avoid the localization effect from high degree vertices when *G* is very sparse.
- Can be generalized for very sparse matrix completion: estimate a low-rank structure from sparse observations with O(n) many samples.
   [Bordenave-Coste-Nadakuditi '23]

#### Long matrix reconstruction

• Rectangular matrix M of size  $n \times m$  ( $m \gg n$ ), with SVD

$$M = \sum_{i=1}^{r} \nu_i \phi_i \psi_i^{\top}, \quad MM^{\top} = \sum_{i=1}^{r} \nu_i^2 \phi_i \phi_i^{\top}$$

- Masking matrix X with  $X_{ij} \sim \text{Ber}(p)$ ,  $p = \frac{d}{\sqrt{mn}}$ .
- Observed matrix:

$$A = rac{X \circ M}{p}$$
 so that  $\mathbb{E}[A] = M$ 

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Assumptions:

$$r, \sqrt{n} \|\phi_i\|_{\infty} = O(\mathsf{polylog}(n))$$

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Estimating the full SVD of M needs O(m) [Bordenave-Coste-Nadakuditi '23]!

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 $\Rightarrow$  B has size  $\sim n^2 m p^2 = d^2 n$ : independent from m

Defined B index by  $\vec{E}$  as

$$B_{ef} = \begin{cases} A_{f_1f_2}A_{f_3f_2} & \text{if } e_3 = f_1 \text{ and } e_2 \neq f_2 \\ 0 & \text{otherwise} \end{cases}$$

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Two important thresholds:

$$\vartheta_1 = \sqrt{\rho/d}$$

• decreases as  $d^{-1/2}$ 

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 $\vartheta_2 = L/d$ 

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Total threshold (Signal-to-noise ratio):

 $\vartheta = \max(\vartheta_1, \vartheta_2)$ 

#### Theorem (Stephan-Z. '24)

(Outliers) For any ν<sub>i</sub> satisfying ν<sub>i</sub> > ϑ, there exists an eigenvalue λ<sub>i</sub> of B with

$$|\lambda_i - \nu_i^2| = O(n^{-c})$$

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Similar to the Kesten-Stigum threshold in community detection [Bordenave-Lelarge-Massoulié '18, Mossel-Neeman-Sly '18]

## Results: eigenvalues



**Figure:** *M* is of rank-2, spectrum of B, d = 3

## Need an embedding procedure from $\mathbb{R}^{\vec{E_2}}$ to $\mathbb{R}^n$

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• For a right eigenvector  $\xi^R$  of *B*:

$$\zeta^R(x) = \sum_{e:e_1=x} A_{e_1e_2} A_{e_3e_2} \xi^R(e), \quad \forall x \in [n].$$

• For a left eigenvector  $\xi^L$ :

$$\zeta^{L}(x) = \sum_{e:e_{1}=x} \xi^{L}(e)$$

#### Theorem (Stephan-Z. '24)

Assume that  $\nu_i > \vartheta$ , and let  $\xi_i^{L/R}$  the left/right eigenvectors associated to  $\lambda_i$ . Then, there exists a  $\gamma_i$  such that

$$\gamma_i = 1 - O(d^{-1})$$

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Weak recovery when  $d \rightarrow \infty$ . Explicit  $\gamma_i$  when d is fixed.

#### Our results: eigenvectors



**Figure:** B, d = 3

- $T = x_1 \otimes \cdots \otimes x_k$ ,  $x_i \in \frac{1}{\sqrt{n}} \{\pm 1\}$ . Sample with probability  $p = \frac{d}{n^{k/2}}$ .
- Unfold T
   in k different ways ( the most unbalanced unfolding) [Ben Arous, Huang, Huang '23]. Apply the non-backtracking method to Unfold(T

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- When sample size is  $\alpha n^{k/2}$  with  $\alpha > 1$ , one can find unit eigenvectors such that

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•  $T = \sum_{i=1}^{r} \lambda_i \left( w_i^{(1)} \otimes \cdots \otimes w_i^{(k)} \right)$ , under the orthonormal condition on  $w_1^{(j)}, \ldots, w_r^{(j)}$ , the same analysis apply.  $O(n^{k/2})$  samples for nontrivial approximation.



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- "independent" wedges
- Associated weight:

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   W<sub>e</sub> = A<sub>xY</sub>A<sub>zY</sub>, Y ~ Unif([m])
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- Bulk eigenvalues: high moment methods on random bipartite graphs.
- Top eigenvalues and eigenvectors: local tree approximation. Galton-Watson tree with random weights [Stephan and Massoulié '20]

## Conclusions

- Very sparse and long matrices are not standard in random matrix theory.
   We define a new non-backtracking matrix tailored for it (random bipartite graphs with more attention to V<sub>1</sub>).
- The corresponding spectral method for tensor completion reaches the conjectured threshold in [Barak-Moitra '15].

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- The corresponding spectral method for tensor completion reaches the conjectured threshold in [Barak-Moitra '15].
- Does not work for finite aspect ratio (m = O(n)) considered in [Bordenave-Coste-Nadakuditi '23]. Is there a unified spectral algorithm for all aspect ratios?
- Statistical-computational gap between O(n) and  $O(n^{k/2})$  samples:
  - Possible with O(n) samples and polynomial-time algorithms with non-uniform/ adaptive sampling [Haselby-Iwen-Karnik-Wang '24]
  - Rank-1 case is different, can be estimated with O(n) samples
     [Stephan-Z. '24, Gomez-Leos, López '24] by solving linear systems.
  - can we justify this gap with a hardness proxy ?

# Thank you!

L. Stephan, Y. Zhu, A non-backtracking method for long matrix and tensor completion, COLT 2024.