

# Robust Blockwise Random Pivoting (RBRP): Fast and Accurate Adaptive Interpolative Decomposition

Yijun Dong<sup>1</sup>, Chao Chen<sup>2</sup>, Per-Gunnar Martinsson<sup>3</sup>, Katherine Pearce<sup>3</sup>

<sup>1</sup>Courant Institute, New York University

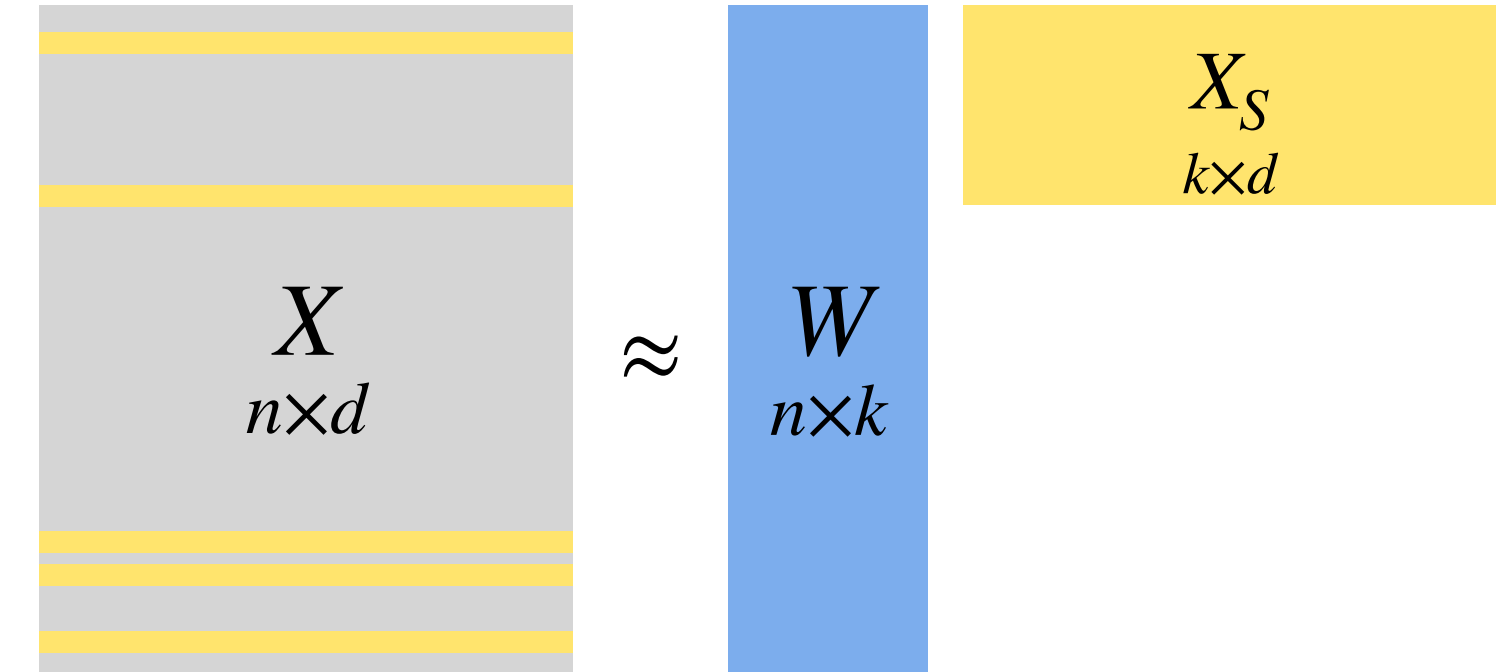
<sup>2</sup>Department of Mathematics, North Carolina State University

<sup>3</sup>Oden Institute, University of Texas at Austin

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# Interpolative Decomposition (ID)

- Given a data matrix  $X = [x_1, \dots, x_n]^\top \in \mathbb{R}^{n \times d}$
- A target rank  $1 \leq r \leq \text{rank}(X)$
- A distortion constant  $\epsilon > 0$
- Aim to construct a  **$(r, \epsilon)$ -ID** of  $X$  —  $X \approx WX_S$  such that



$$\|X - WX_S\|_F^2 \leq (1 + \epsilon) \|X - X_{\langle r \rangle}\|_F^2$$

- $S = \{s_1, \dots, s_k\} \subseteq [n]$  contains indices for a **skeleton subset** of size  $|S| = k$  (usually  $k \ll n$ )
- $X_S = [x_{s_1}, \dots, x_{s_k}]^\top \in \mathbb{R}^{k \times d}$  is the row skeleton submatrix corresponding to  $S$
- $W \in \mathbb{R}^{n \times k}$  is an interpolation matrix for the given skeleton subset  $S$
- $X_{\langle r \rangle}$  denotes the optimal rank- $r$  approximation of  $X$  (given by the truncated SVD)

# Two Stages of ID Constructions

- **Stage I: Skeleton selection**

- Find a good skeleton subset  $S$ :

$$\min_{S \subseteq [n]} \min_{W \in \mathbb{R}^{n \times |S|}} \|X - WX_S\|_F^2$$

- **Skeletonization error:**  $\mathcal{E}_X(S) := \|X - XX_S^\dagger X_S\|_F^2 = \min_{W \in \mathbb{R}^{n \times |S|}} \|X - WX_S\|_F^2$

- Naive construction of  $XX_S^\dagger$  (e.g., via QR) takes  $O(ndk)$  time (i.e.,  $k = |S|$  additional passes through  $X$ )

- **Stage II: Interpolation matrix construction**

- For some  $O(ndk)$ -time selection algorithms,  $W$  can be evaluated/approximated a posteriori in  $O(nk^2)$  time
- **Interpolation error:**  $\mathcal{E}_X(W|S) := \|X - WX_S\|_F^2$

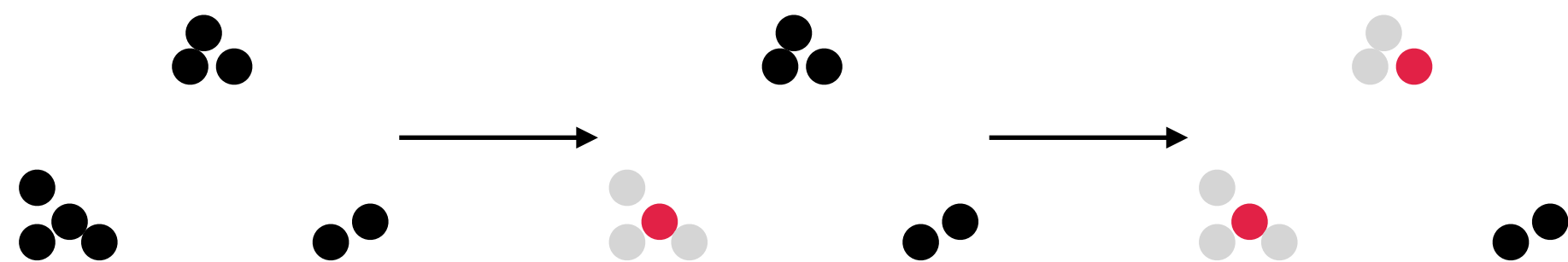
# What are Fast & Accurate ID Algorithms?

- **Skeleton complexity**: the minimum number of skeletons  $k = |S|$  that an ID algorithm needs to select in order to form a  $(r, \epsilon)$ -ID (in expectation), i.e.,  $\mathcal{E}_X(S) \leq (1 + \epsilon)\|X - X_{\langle r \rangle}\|_F^2$
- **Asymptotic complexity**: the asymptotic FLOP counts of the skeleton selection stage in an ID algorithm
- **Parallelizability**: whether the dominant cost of the skeleton selection stage in an ID algorithm can be casted as **matrix-matrix (fast)**, instead of **matrix-vector (slow)**, multiplications with  $X$  (i.e., applicability of Level 3 BLAS)
- **Error-revealing property**: the ability of an ID algorithm to evaluate  $\mathcal{E}_X(S)$  efficiently on the fly so that the target rank  $k$  does not need to be given a priori.
  - Definition: An ID algorithm is **error-revealing** if after selecting any skeleton subset  $S$ , it can evaluate the corresponding skeletonization error  $\mathcal{E}_X(S)$  efficiently in at most  $O(n)$  time.
- **ID-revealing property**: if the skeleton selection stage of an ID algorithm extracts sufficient information so that
  - **Exact/inexact-ID-revealing**:  $W = XX_S^\dagger$  can be evaluated exactly/approximated in  $O(nk^2)$  time
  - **Non-ID-revealing** otherwise

# Adaptiveness & Randomness

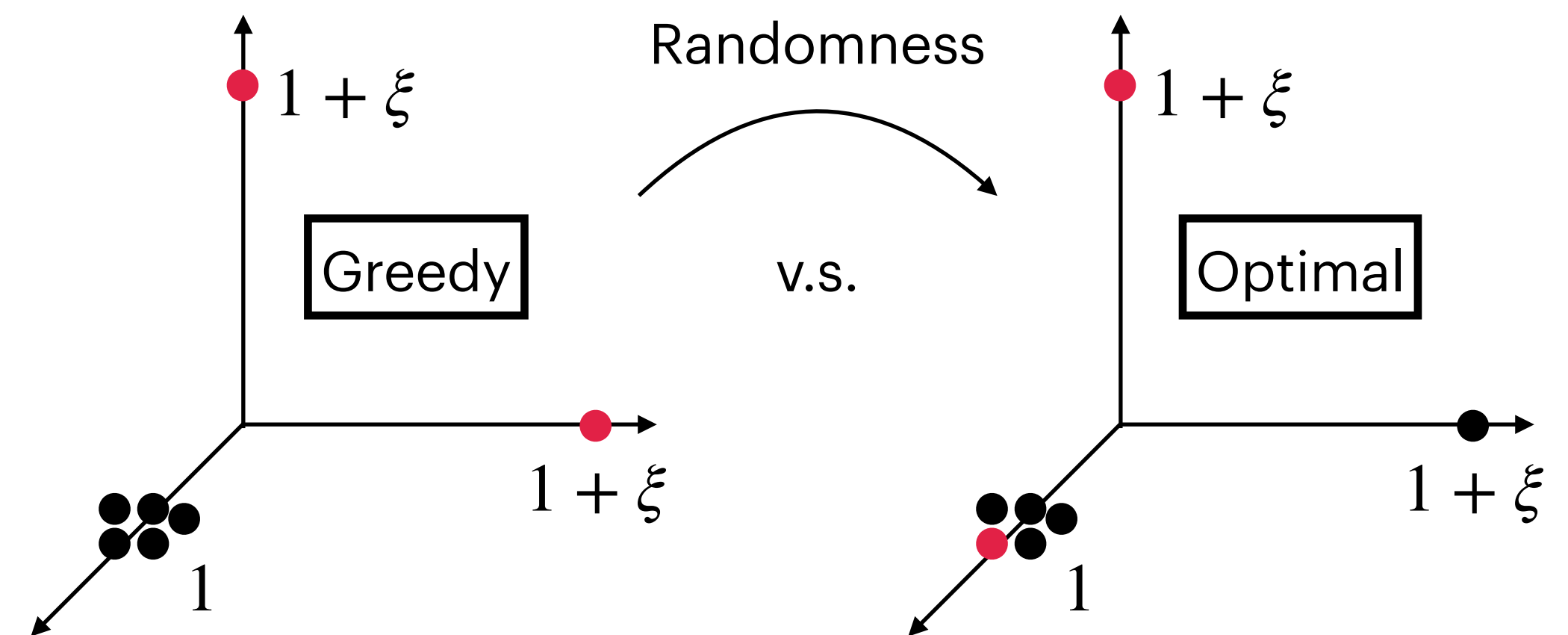
- **Adaptiveness**

- Each new skeleton selection is aware of the previously selected skeleton subset
- By selecting according to the residual
- Common adaptive residual updates:
  - Gram-Schmidt (QR)
  - Gaussian elimination (LU)



- **Randomness** (in contrast to greedy)

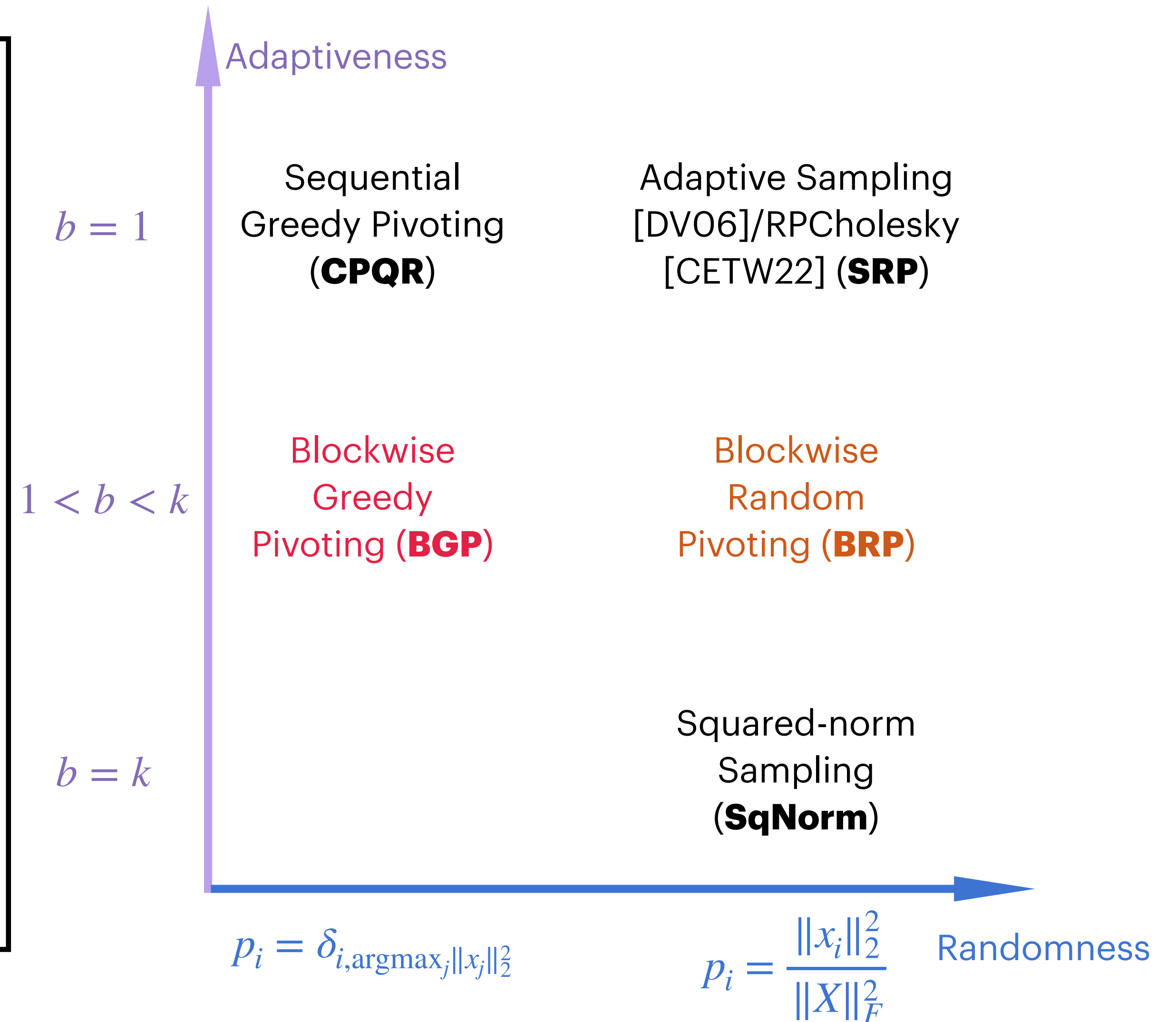
- Intuition: balance exploitation with exploration
- Effectively circumvent adversarial inputs for greedy methods
- Achieve appealing skeleton complexities in expectation
- Common randomness: sampling, sketching



# Skeleton Selection: A General Framework

A framework for (blockwise adaptive) skeleton selection

- **Inputs:**  $X \in \mathbb{R}^{n \times d}$ ,  $\tau = (1 + \epsilon)\eta_r \in (0,1)$
- $X^{(0)} \leftarrow X$ ,  $S^{(0)} \leftarrow \emptyset$ ,  $t \leftarrow 0$
- **while**  $\mathcal{E}(S^{(t)}) > \tau \|X\|_F^2$  **do**
  - $t \leftarrow t + 1$
  - Select  $|S_t| = b$  skeletons  $S_t$  based on  $\left( p_i(X^{(t-1)}) \right)_{i \in [n]}$
  - $S^{(t)} \leftarrow S^{(t-1)} \cup S_t$
  - $X^{(t)} \leftarrow X^{(t-1)} \left( I_d - X_{S_t}^\dagger X_{S_t} \right)$
- $S \leftarrow S^{(t)}$ ,  $k = |S|$



# Skeleton Selection: Other Methods

## Sampling methods

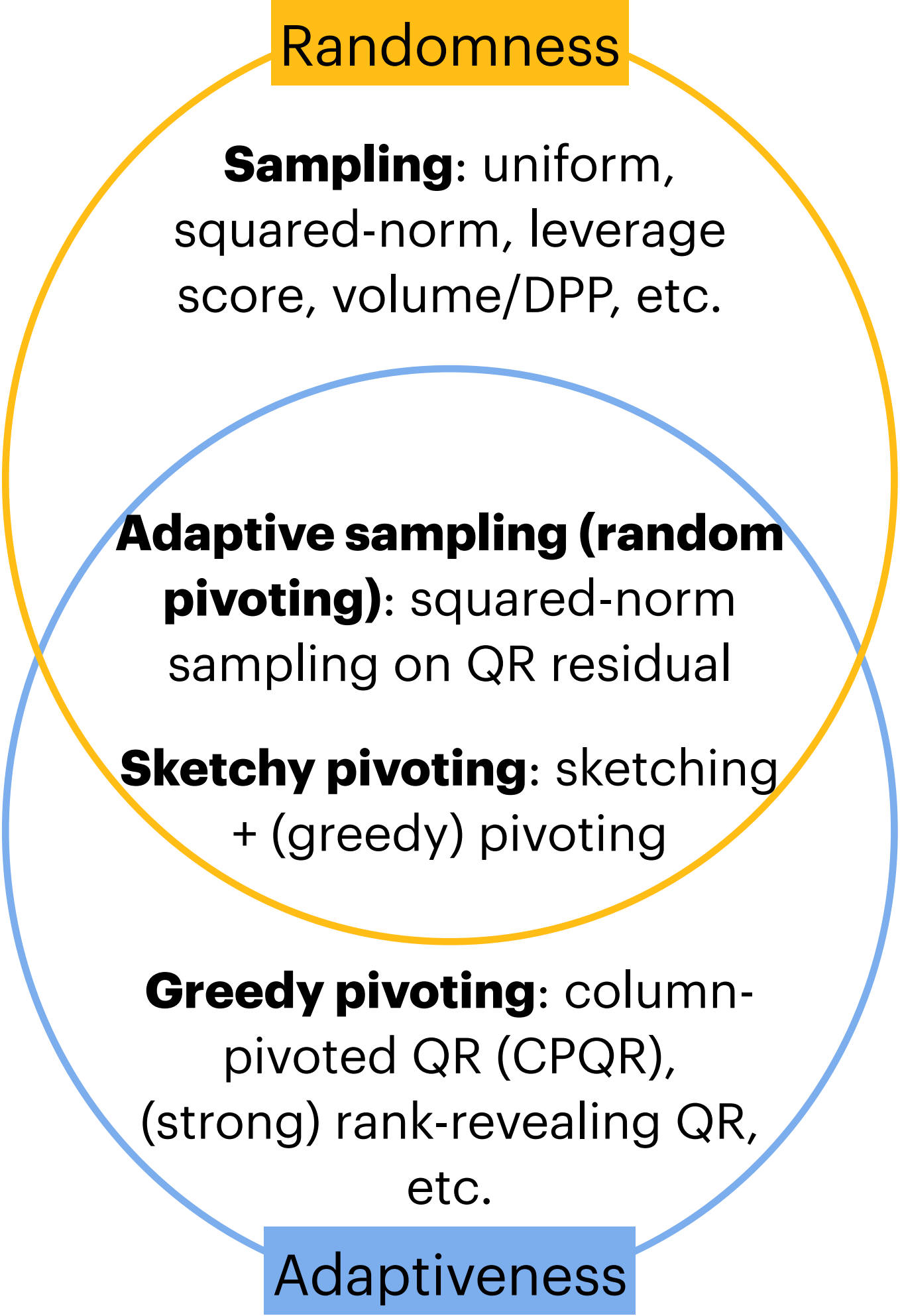
- **DPP/volume sampling** [HKPV06, BW09, DR10, KT11, GS12]
  - Pro: nearly optimal expected skeleton complexity:  
 $k \geq \frac{r}{\epsilon} + r - 1$  is sufficient for  $(r, \epsilon)$ -ID in expectation
  - Con: expensive to compute
- **Leverage score sampling** [MD09, DMMW12]
  - Pro: can be estimated efficiently for large-scale problems (e.g., tensor Khatri-Rao product)
  - Con: expensive to compute
- **Uniform sampling** [CLMMPS15]
  - Pro: linear time
  - Con: require/depend on matrix incoherence

## Sketchy pivoting

- **Inputs:**  $X \in \mathbb{R}^{n \times d}$ ,  $k \leq \text{rank}(X)$ ,
- Draw JLT  $\Omega \in \mathbb{R}^{d \times k}$  (e.g.,  $\Omega_{ij} \sim \mathcal{N}(0, 1/k)$  i.i.d.)
- Sketching  $Y = X\Omega \in \mathbb{R}^{n \times k}$
- Greedy pivoting: for  $t = 1, \dots, k$ 
  - Column (row) pivoted QR (**CPQR**) [VM17]:  
 $s_t \leftarrow \underset{i}{\operatorname{argmax}} \|Y_{i,:}^{(t-1)}\|_2^2 + \text{Gram-Schmidt}$
  - LU with partial pivoting (**LUPP**) [DM23]:  
 $s_t \leftarrow \underset{i}{\operatorname{argmax}} |Y_{i,t}^{(t-1)}| + \text{Gaussian Elimination}$
- Pro: fast, accurate, robust to adversarial inputs
- Con: require prior knowledge of  $k$



# ID Algorithms with Adaptiveness & Randomness



| Algorithm                    | Skeleton Complexity  | Asymp. Cost + Parallelizability | Error-reveal | ID-reveal    |
|------------------------------|--|---------------------------------|--------------|--------------|
| <b>Greedy Pivoting</b>       | $k \geq (1 + (1 + \epsilon)\eta_r)n$   | $O(ndk)$ sequential             | ✓            | <b>Exact</b> |
| <b>Squared-norm Sampling</b> | $k \geq \frac{r-1}{\epsilon\eta_r} + \frac{1}{\epsilon}$   | $O(nd)$ parallel                | ✗            | Non          |
| <b>Random Pivoting</b>       | $k \geq k_{RP} := \frac{r}{\epsilon} + r \log \left( \min \left\{ \frac{1}{\epsilon\eta_r}, \frac{2^{r+1}}{\epsilon} \right\} \right)$ | $O(ndk)$ sequential             | ✓            | <b>Exact</b> |
| <b>Sketchy Pivoting</b>      | Conjecture: $k \gtrsim k_{RP}$   | $O(ndk)$ parallel               | ✗            | Inexact      |
| <b>RBRP</b>                  | Conjecture: $k \gtrsim k_{RP}$   | $O(ndk)$ parallel               | ✓            | <b>Exact</b> |

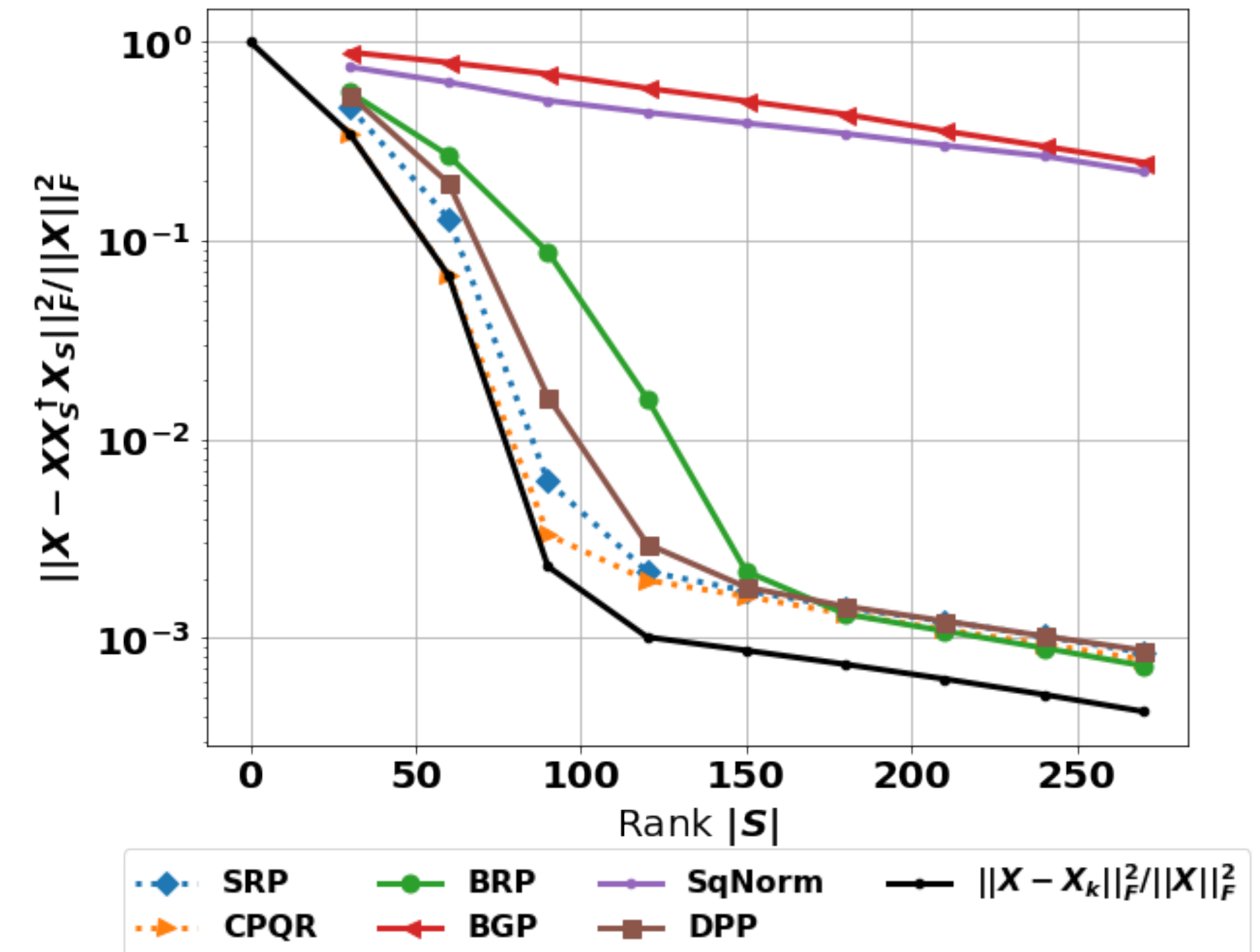
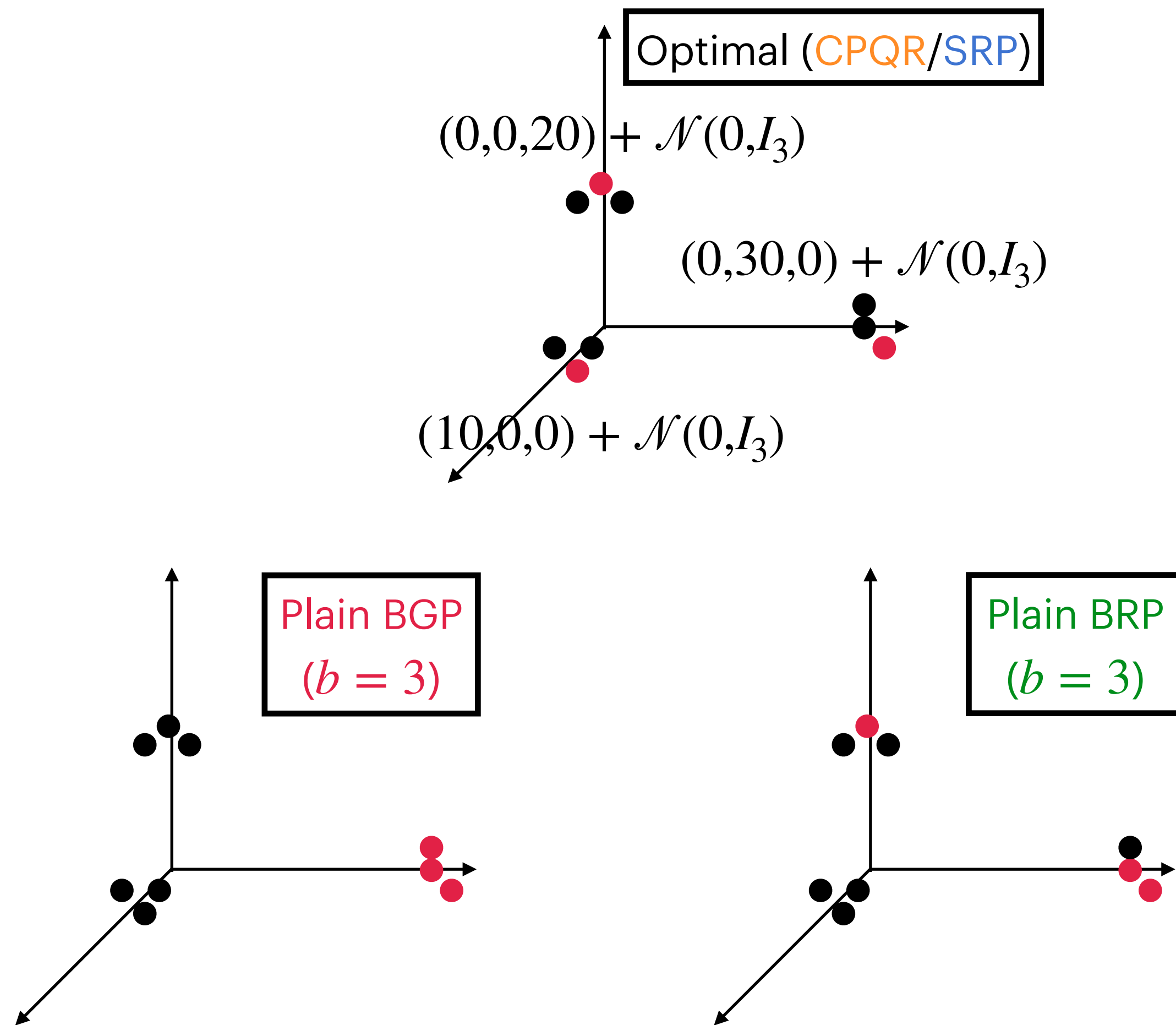
\*  $\eta_r = \|X - X_{<r>\|_F^2 / \|X\|_F^2$  quantifies the relative optimal rank- $r$  approximation error of  $X$

**Question:** How to parallelize random pivoting?  
**Answer:** Blockwise random pivoting



# Pitfall of Plain Blockwise Greedy/Random Pivoting

$k = 100$  clusters centered at  $\{10j \cdot e_j\}_{j \in [k]}$ ,  $n = 20k$ ,  $d = 500$ ,  $b = 30$

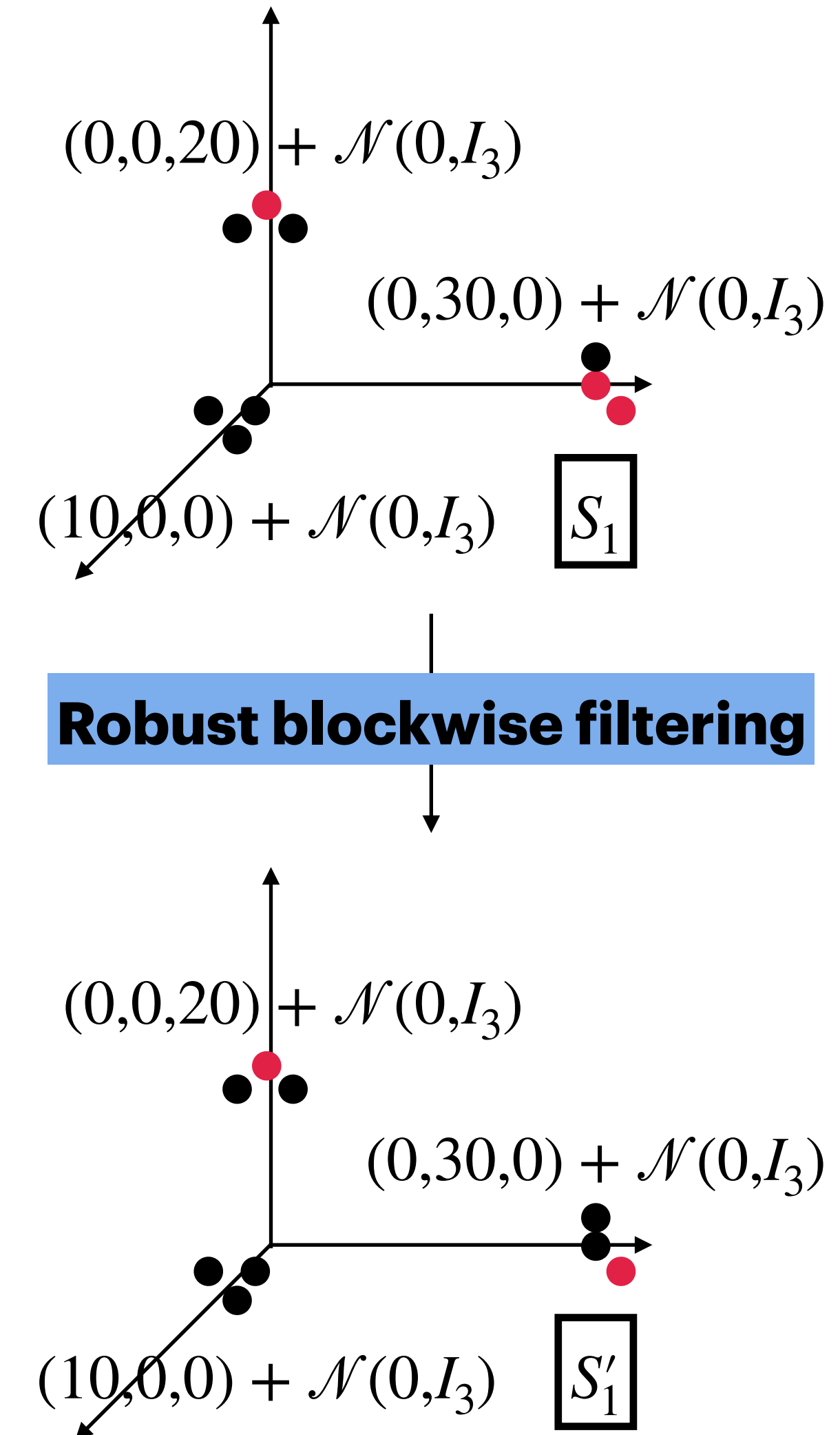


- Sequential pivoting (CPQR & SRP) is nearly optimal
- Plain blockwise pivoting (BRP/BGP, especially BGP) suffers from suboptimal skeleton complexities (up to  $b$  times)
- Squared-norm sampling (SqNorm) tends to fail

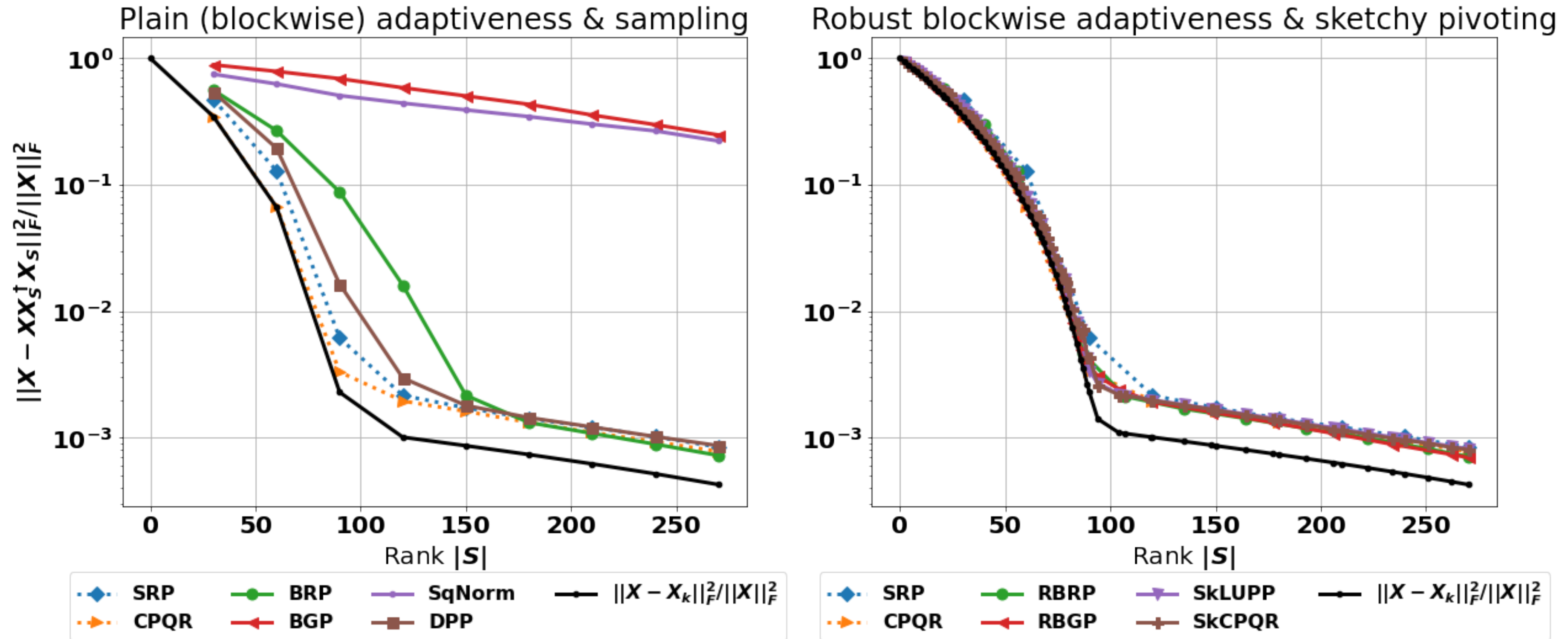
# Robust Blockwise Random Pivoting

## Robust Blockwise Random Pivoting (RBRP)

- **Inputs:**  $X \in \mathbb{R}^{n \times d}$ ,  $\tau = (1 + \epsilon)\eta_r \in (0,1)$
- $X^{(0)} \leftarrow X$ ,  $S^{(0)} \leftarrow \emptyset$ ,  $t \leftarrow 0$
- **while**  $\mathcal{E}(S^{(t)}) > \tau \|X\|_F^2$  ( $t \leftarrow t + 1$ ) **do**
  - Select  $|S_t| = b$  skeletons  $S_t$  based on  $\left(p_i(X^{(t-1)})\right)_{i \in [n]}$
  - **Robust blockwise filtering (RBF)**
    - $\pi \leftarrow \text{CPQR} \left(X_{S_t}^{(t-1)}\right) \in S_b$  (SRP and CPQR both work)
    - $\min_{S'_t = S_t(\pi(1:b'))} b'$  s.t.  $\|X_{S_t} - X_{S'_t}\|_F^2 < \tau_b \|X_{S_t}\|_F^2$  (e.g.,  $\tau_b = \frac{1}{b}$ )
  - $S^{(t)} \leftarrow S^{(t-1)} \cup S'_t$  and  $X^{(t)} \leftarrow X^{(t-1)} \left(I_d - X_{S'_t}^\dagger X_{S_t}\right)$
- $S \leftarrow S^{(t)}$ ,  $k = |S|$

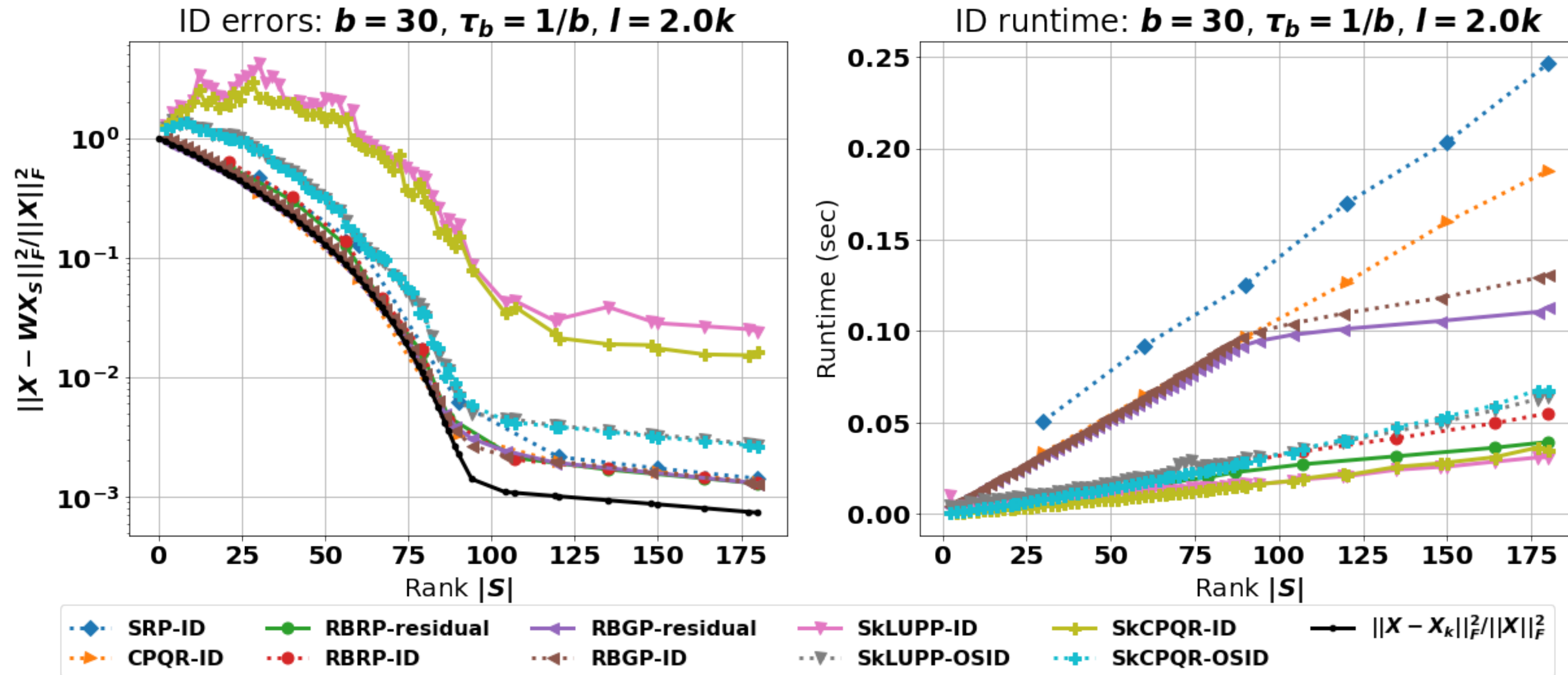


# Robust Blockwise Random Pivoting: Robustness



- GMM with  $k = 100$  clusters centered at  $\{10j \cdot e_j\}_{j \in [k]}$ ,  $\Sigma = I_d$ ,  $n = 20k$ ,  $d = 500$ ,  $b = 30$
- Robust blockwise filtering (RBRP and RBGP) brings nearly optimal skeleton complexities

# Robust Blockwise Random Pivoting: Efficiency



- Robust blockwise filtering (RBRP and RBGP) brings nearly optimal skeleton complexities
- RBGP tends to be slowed down much more significantly than RBRP by robust blockwise filtering
- For ID: RBRP-ID is almost as fast as sketchy pivoting (SkLUPP-ID/SkCPQR-ID), while enjoying much better interpolation error  $\mathcal{E}_X(W|S) = \mathcal{E}_X(S)$  thanks to its exact-ID-revealing property.



# Exact- v.s. Inexact- ID-revealing Algorithms

- **Exact-ID-revealing algorithms**

- Sequential/blockwise random/greedy pivoting algorithms (SRP, CPQR, BRP, BGP, RBRP, RBGP)
- The skeleton selection process generates sufficient information for solving the least square problem

$$\min_{W \in \mathbb{R}^{n \times k}} \|X - WX_S\|_F^2 \text{ in } O(nk^2) \text{ time}$$

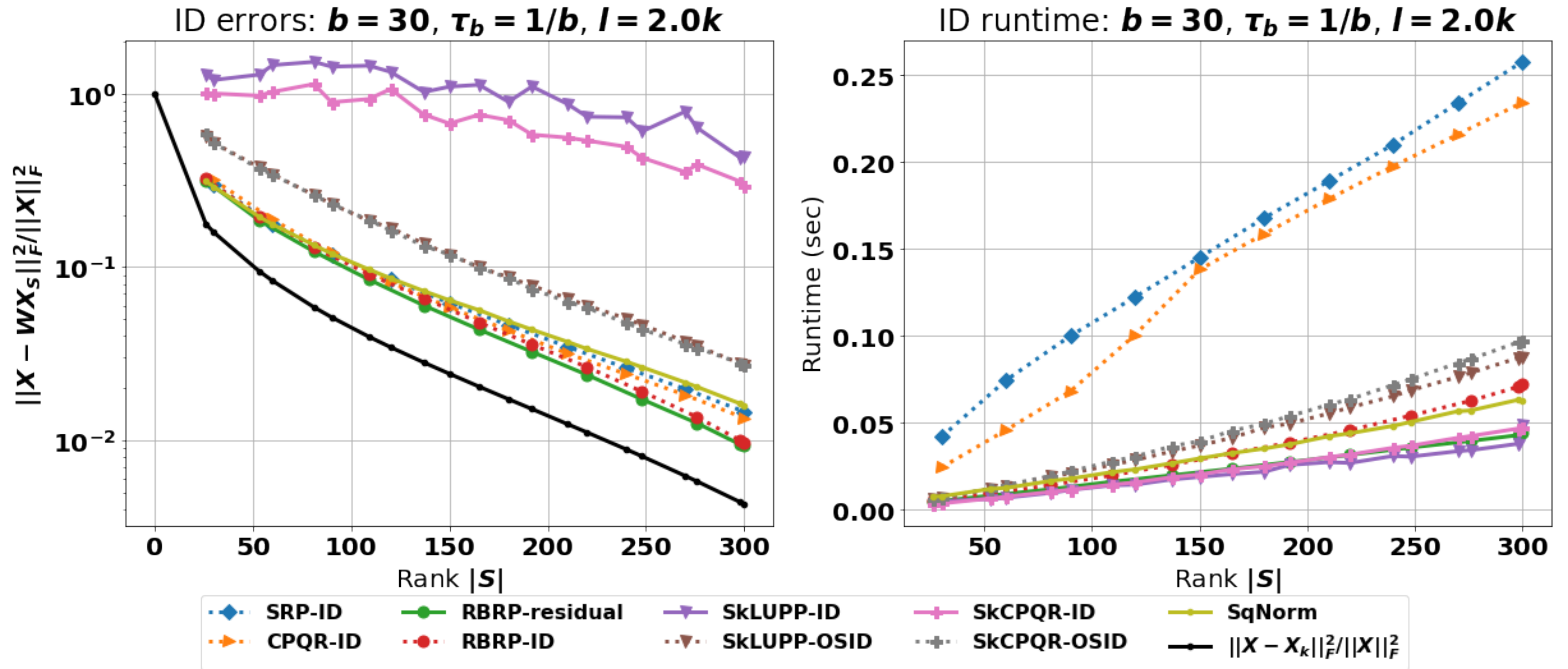
- **Inexact-ID-revealing algorithms**

- Sketchy pivoting algorithms (SkLUPP, SkCPQR)
- The skeleton selection process generates sufficient information for solving the **sketched** least square problem

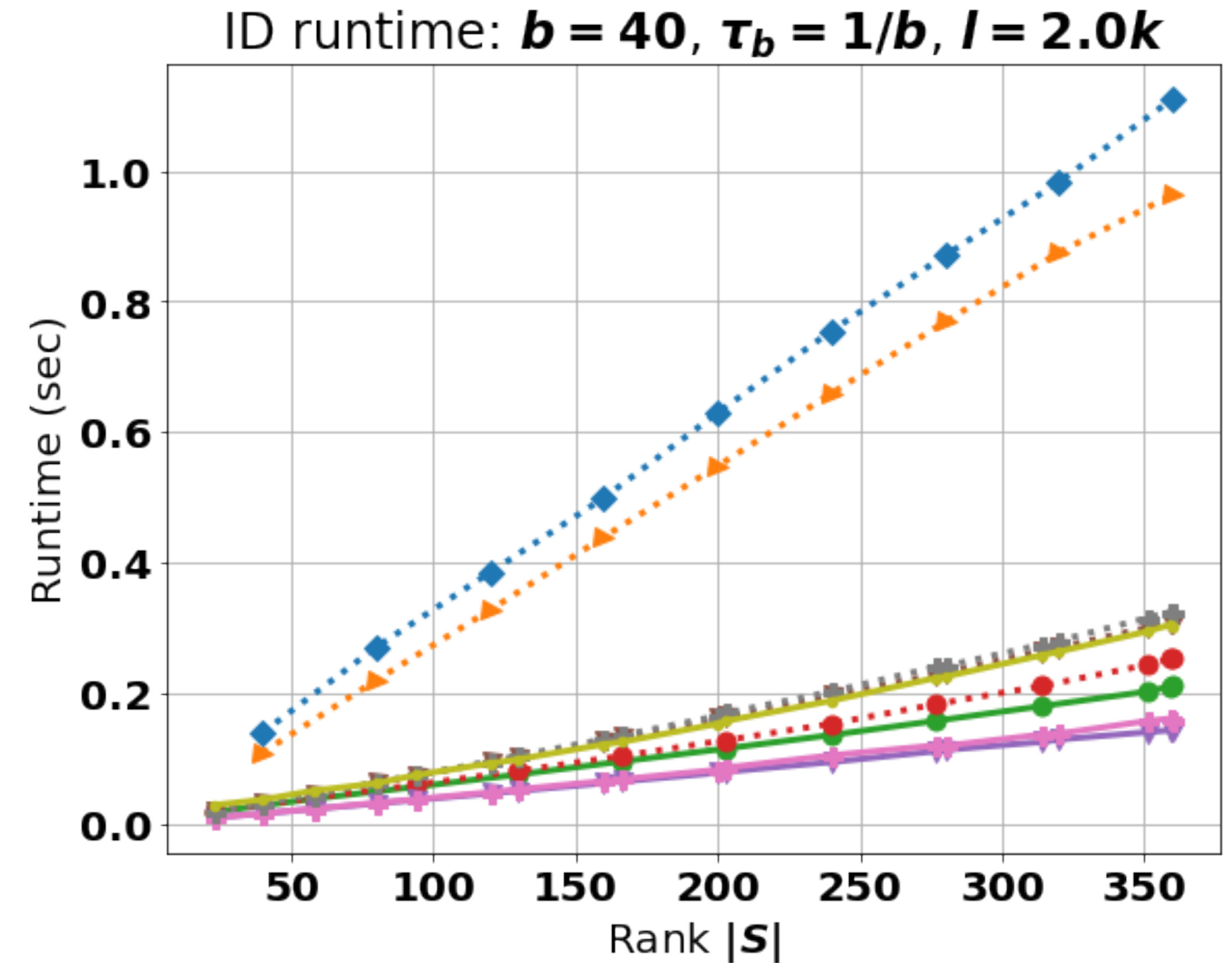
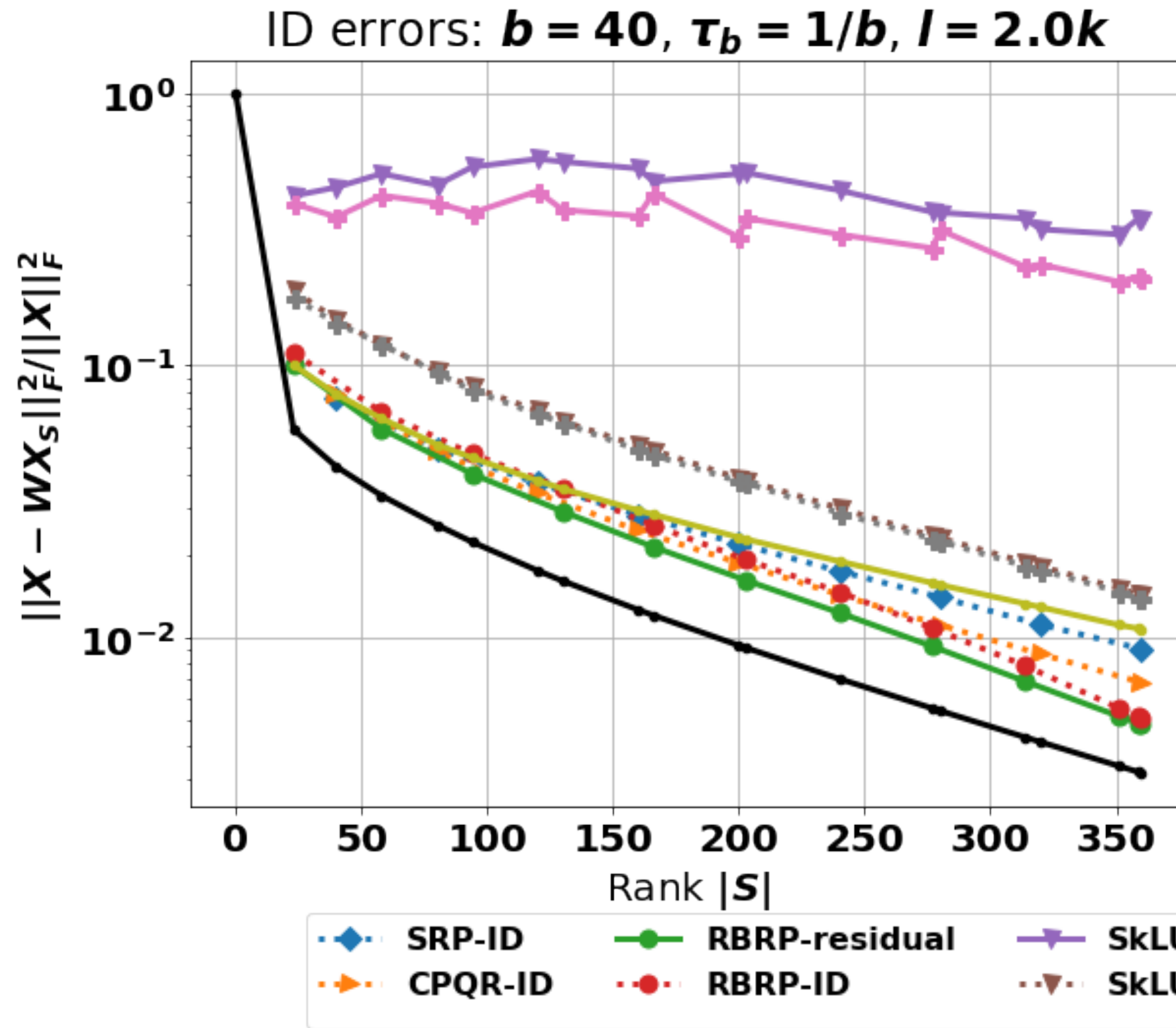
$$\min_{W \in \mathbb{R}^{n \times k}} \|X\Omega - WX_S\Omega\|_F^2 \text{ in } O(nk^2) \text{ time}$$

- Oversampled sketchy ID (**OSID**): for  $|S| = k$ 
  - Sketching with oversampling  $Y = X\Omega \in \mathbb{R}^{n \times l}$  such that  $l = O(k)$
  - $W = YY_S^\dagger$  can be computed in  $O(nlk) = O(nk^2)$  time
- Suboptimal interpolation error:  $\mathcal{E}_X(W|S) - \mathcal{E}_X(S) = O(k/l)$

# More Numerical Comparisons: MNIST



# More Numerical Comparisons: CIFAR-10





# Summary

- A fast & accurate ID algorithm that finds  $\|X - WX_S\|_F^2 \leq (1 + \epsilon)\|X - X_{\langle r \rangle}\|_F^2$ 
  - With **nearly optimal skeleton complexity** in practice
  - Computationally efficient in terms of both **asymptotic complexity** and **parallelizability**
  - **Error-revealing** without requiring prior knowledge of the target skeleton subset size
  - **Exact-ID-revealing** where the optimal interpolation matrix can be computed efficiently
- **Combining adaptiveness and randomness** is a key for designing robust skeleton selection algorithms with competitive skeleton complexity
- A critical challenge is to **relax the sequential nature of adaptive selection**
- We introduced **Robust Blockwise Random Pivoting (RBRP)**, a parallelizable blockwise adaptive selection scheme that achieves comparable skeleton complexity as its sequential counterpart

# Thank You! Happy to take any questions



arXiv: <https://arxiv.org/abs/2309.16002>



GitHub: [https://github.com/dyjdongyijun/Robust\\_Blockwise\\_Random\\_Pivoting](https://github.com/dyjdongyijun/Robust_Blockwise_Random_Pivoting)