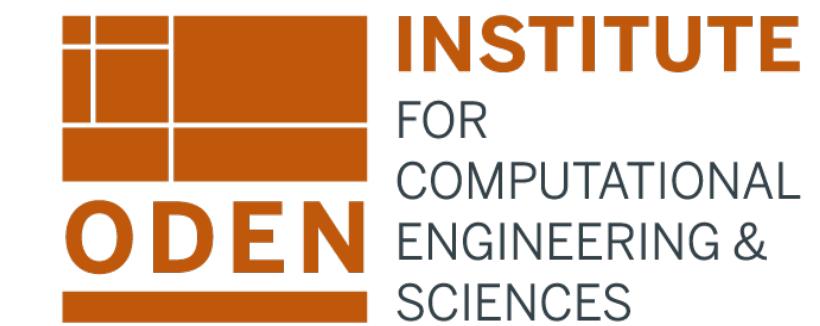


Toward Fast and Provable Data Selection under Low Intrinsic Dimension

Yijun Dong

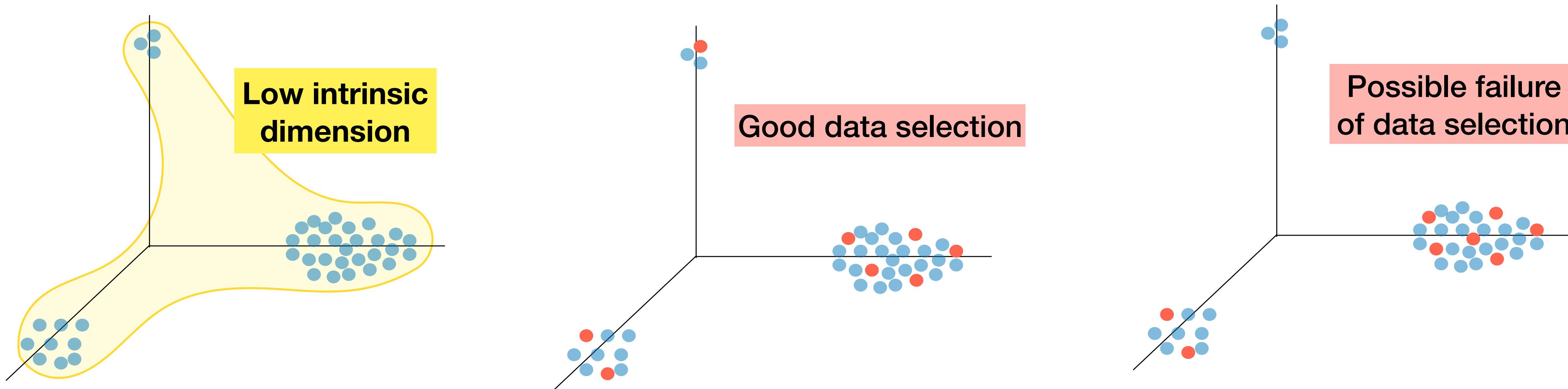
Courant Institute of Mathematical Sciences, New York University

University of Delaware Numerical Analysis and PDE Seminar
Oct 31, 2024



Low Intrinsic Dimension & Data Selection

- Low intrinsic dimension is ubiquitous in real world
 - Example: Large models can be finetuned in a lower dimension with much fewer samples than the model size Aghajanyan-Zettlemoyer-Gupta-2020
 - 341M parameters → dimension-322 subspace can be fine-tuned with less than 6K samples!
- Learning under low intrinsic dimension with limited data, data selection becomes crucial



How to **select the most informative data** for learning under **low intrinsic dimension** (e.g. finetuning)?

Outline

- Data selection for statistical models in kernel regime — **finetuning**
 - A theory on data selection for finetuning that
 - Extends the classical wisdom of **V-optimal experimental design** in low dimensions
 - To high-dimensional data selection under low intrinsic dimensions via **sketching**
 - A fast and effective data selection algorithm for finetuning: **Sketchy Moment Matching (SkMM)**
- Data selection for low-rank approximation — **interpolative decomposition (ID)**
 - Select minimum possible rows as a basis to form a good low-rank approximation
 - A fast and accurate adaptive ID algorithm: **Robust Blockwise Random Pivoting (RBRP)**

Sketchy Moment Matching: Toward Fast and Provable Data Selection for Finetuning



Hoang Phan
NYU



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Data Selection for Finetuning

- Large full dataset $X = [x_1, \dots, x_N]^\top \subset \mathcal{X}^N$, $y = [y_1, \dots, y_N] \in \mathbb{R}^N$ drawn i.i.d. from unknown distribution P
- Finetuning function class $\mathcal{F} = \{f(\cdot; \theta) : \mathcal{X} \rightarrow \mathbb{R} \mid \theta \in \Theta\}$ with parameters $\Theta \subset \mathbb{R}^r$
- Pre-trained initialization $0_r \in \mathbb{R}^r$ (without loss of generality)
- Ground truth $\theta_* \in \Theta$ such that $\mathbb{E}[y \mid x] = f(x; \theta_*)$ and $\mathbb{V}[y \mid x] \leq \sigma^2$

Select a small coresset $(X_S, y_S) \subset \mathcal{X}^n \times \mathbb{R}^n$ of size n indexed by $S \subset [N]$ such that:

$$(1) \quad \theta_S = \arg \min_{\theta \in \Theta} \frac{1}{n} \|f(X_S; \theta) - y_S\|_2^2 + \alpha \|\theta\|_2^2$$

- **Low-dimensional** data selection: $r \leq n$, (1) = linear regression ($\alpha = 0$)
- **High-dimensional** data selection: $r > n$, (1) = ridge regression ($\alpha > 0$)

Finetuning in Kernel Regime

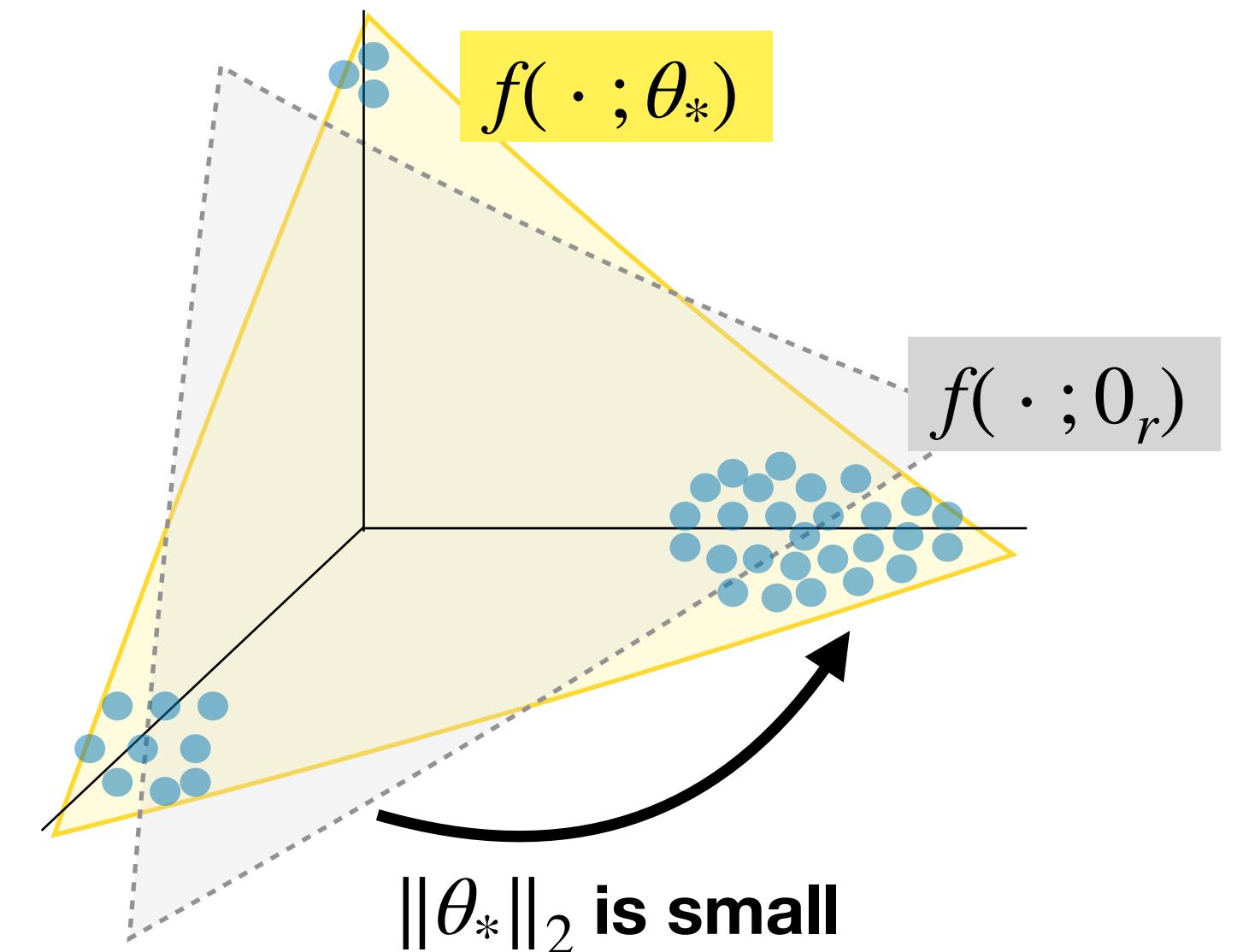
- Finetuning dynamics fall in the **kernel regime**:

$$f(x; \theta) \approx f(x; 0_r) + \nabla_{\theta} f(x; 0_r)^{\top} \theta$$

- With a **suitable pre-trained initialization** (i.e. $f(\cdot, 0_r)$ is close to $f(\cdot, \theta_*)$), $\|\theta_*\|_2$ is small
- Let $G = \nabla_{\theta} f(X; 0_r) \in \mathbb{R}^{N \times r}$ and $G_S = \nabla_{\theta} f(X_S; 0_r) \in \mathbb{R}^{n \times r}$, (1) is well approximated by:

$$(2) \quad \theta_S = \arg \min_{\theta \in \Theta} \frac{1}{n} \|G_S \theta - (y_S - f(X_S; 0_r))\|_2^2 + \alpha \|\theta\|_2^2$$

- Aim to control excess risk $\text{ER}(\theta_S) = \|\theta_S - \theta_*\|_{\Sigma}^2$ where $\Sigma = \mathbb{E}_{x \sim P} [\nabla_{\theta} f(x; 0_r) \nabla_{\theta} f(x; 0_r)^{\top}] \in \mathbb{R}^{r \times r}$



In Low Dimension: Variance Reduction

- Consider **fixed design** for simplicity: $\Sigma = \mathbb{E}_{x \sim P} [\nabla_{\theta} f(x; 0_r) \nabla_{\theta} f(x; 0_r)^{\top}] = G^{\top} G / N$
- **Low-dimensional** data selection: $\text{rank}(G_S) = r \leq n$ such that $\Sigma_S = G_S^{\top} G_S / n > 0$
- **V(ariance)-optimality** characterizes generalization: $\mathbb{E}[\text{ER}(\theta_S)] \leq \frac{\sigma^2}{n} \text{tr}(\Sigma \Sigma_S^{-1})$
- If $\Sigma \leq c_S \Sigma_S$ for some $c_S \geq \frac{n}{N}$, then $\mathbb{E}[\text{ER}(\theta_S)] \leq c_S \frac{\sigma^2 r}{n}$

Uniform sampling achieves nearly optimal sample complexity in low dimension: Assuming $\|\nabla_{\theta} f(\cdot; 0_r)\|_2 \leq B$ and $\Sigma \succeq \gamma I_r$. With probability $\geq 1 - \delta$, X_S sampled uniformly from X satisfies

$$\Sigma \leq c_S \Sigma_S \text{ for any } c_S > 1 \text{ when } n \gtrsim \frac{B^4}{\gamma^2 (1 - c_S^{-1})^2} (r + \log(1/\delta))$$

Can the **low intrinsic dimension** of fine-tuning be leveraged when $r > n$?

With Low Intrinsic Dimension: Variance + Bias

Optimal rank- t approximation
(truncated SVD)

Assumption (Low intrinsic dimension): For $\Sigma = G^\top G/N$, let $\bar{r} = \min\{t \in [r] \mid \text{tr}(\Sigma - \langle \Sigma \rangle_t) \leq \text{tr}(\Sigma)/N\}$ be the intrinsic dimension of the learning problem. Assume $\bar{r} \ll \min\{N, r\}$

Corollary (Exploitation + exploration): Given $S \subset [N]$, for $\mathcal{S} \subseteq \text{Range}(\Sigma_S)$ with $\text{rank}(P_{\mathcal{S}}) \asymp \bar{r}$, if

- **Variance** is controlled by **exploiting** information in \mathcal{S} : $P_{\mathcal{S}}(c_S \Sigma_S - \Sigma) P_{\mathcal{S}} \succeq 0$ for some $c_S \geq n/N$; and
- **Bias** is controlled by **exploring** $\text{Range}(\Sigma)$ for an informative \mathcal{S} : $\text{tr}(\Sigma P_{\mathcal{S}}^\perp) \leq \frac{N}{n} \text{tr}(\Sigma - \langle \Sigma \rangle_{\bar{r}})$. Then,

$$\mathbb{E}[\text{ER}(\theta_S)] \leq \text{variance} + \text{bias} \lesssim \frac{1}{n} (c_S \sigma^2 \bar{r} + \text{tr}(\Sigma) \|\theta_*\|_2^2)$$

- **Sample efficiency:** With suitable selection of $S \subset [N]$, the sample complexity of finetuning is **linear in the intrinsic dimension \bar{r}** , independent of the (potentially high) parameter dimension r

How to explore the intrinsic low-dimensional structure **efficiently** for data selection?

Explore Low Intrinsic Dimension: Gradient Sketching

- **Gradient sketching:** Randomly projecting the high-dimensional gradients $G = \nabla_{\theta} f(X; \theta_r) \in \mathbb{R}^{N \times r}$ with $r > n$ to a lower-dimension $m = O(\bar{r}) \ll r$ via a Johnson-Lindenstrauss transform (JLT) $\Gamma \in \mathbb{R}^{r \times m}$
- Common JLT: a Gaussian random matrix with i.i.d entries $\Gamma_{ij} \sim \mathcal{N}(0, 1/m)$

Theorem (Gradient sketching): For Gaussian embedding $\Gamma \in \mathbb{R}^{r \times m}$ with $m \geq 11\bar{r}$, let $\widetilde{\Sigma} = \Gamma^\top \Sigma \Gamma$ and $\widetilde{\Sigma}_S = \Gamma^\top \Sigma_S \Gamma$. If the coresset $S \subset [N]$ satisfies $\text{rank}(\Sigma_S) = n > m$ and the $\lceil 1.1\bar{r} \rceil$ -th largest eigenvalue $s_{\lceil 1.1\bar{r} \rceil}(\Sigma_S) \geq \gamma_S > 0$, then with probability at least 0.9 over Γ , there exists $\alpha > 0$ such that

$$\mathbb{E}[\text{ER}(\theta_S)] \lesssim \underbrace{\frac{\sigma^2}{n} \text{tr}(\widetilde{\Sigma} (\widetilde{\Sigma}_S)^\dagger)}_{\text{variance}} + \underbrace{\frac{\sigma^2}{n} \frac{1}{m\gamma_S} \|\widetilde{\Sigma} (\widetilde{\Sigma}_S)^\dagger\|_2}_{\text{sketching error}} \text{tr}(\Sigma) + \underbrace{\frac{1}{n} \|\widetilde{\Sigma} (\widetilde{\Sigma}_S)^\dagger\|_2}_{\text{bias}} \text{tr}(\Sigma) \|\theta_*\|_2^2$$

- If S further satisfies $\widetilde{\Sigma} \leq c_S \widetilde{\Sigma}_S$ for some $c_S \geq n/N$, with $m = \max\{\sqrt{\text{tr}(\Sigma)/\gamma_S}, 11\bar{r}\}$,

$$\mathbb{E}[\text{ER}(\theta_S)] \lesssim \frac{c_S}{n} (\sigma^2 m + \text{tr}(\Sigma) \|\theta_*\|_2^2)$$

Control Variance: Sketchy Moment Matching (SkMM)

Gradient sketching

- Draw a (fast) JLT (e.g. Gaussian random matrix) $\Gamma \in \mathbb{R}^{r \times m}$
- Sketch the gradients $\widetilde{G} = \nabla_{\theta} f(X; 0_r) \Gamma \in \mathbb{R}^{N \times m}$

Moment matching

- Spectral decomposition $\widetilde{\Sigma} = \widetilde{G}^T \widetilde{G} / N = V \Lambda V^T$ with $V = [v_1, \dots, v_m]$, $\Lambda = \text{diag}(\lambda_1, \dots, \lambda_m)$
- Initialize $s = [s_1, \dots, s_N]$ with $s_i = 1/n$ for $i \in [N]$ uniformly sampled and $s_i = 0$ otherwise
- Sample a size- n cores $S \subset [N]$ according to the distribution s that solves the optimization problem

$$\begin{aligned} \min_{s \in [0, 1/n]^N} \quad & \min_{\gamma = [\gamma_1, \dots, \gamma_m] \in \mathbb{R}^m} \sum_{j=1}^m (v_j^T \widetilde{G}^T \text{diag}(s) \widetilde{G} v_j - \gamma_j \lambda_j)^2 \\ \text{s.t.} \quad & \|s\|_1 = 1, \quad \gamma_j \geq 1/c_S \quad \forall j \in [m] \end{aligned}$$

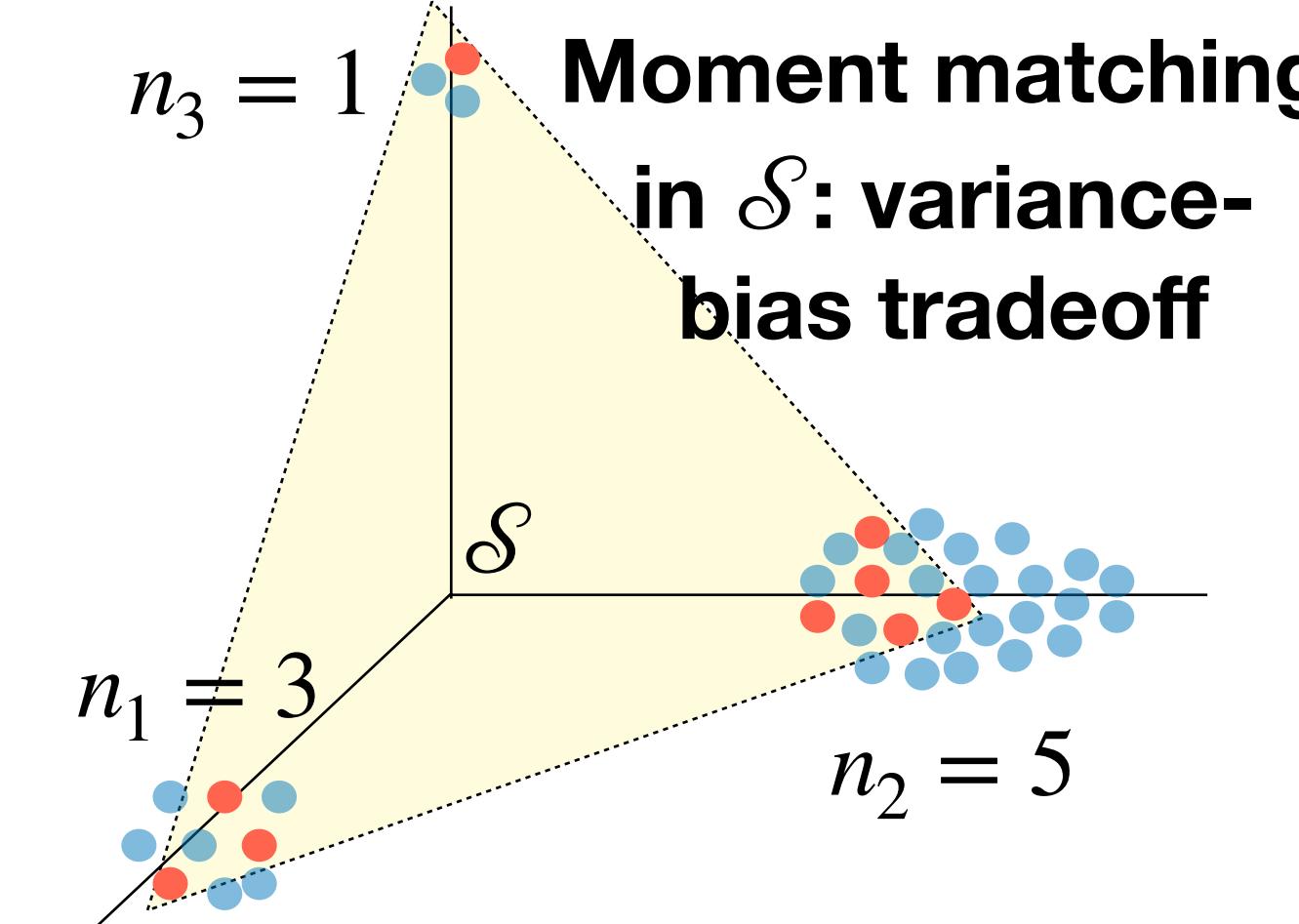
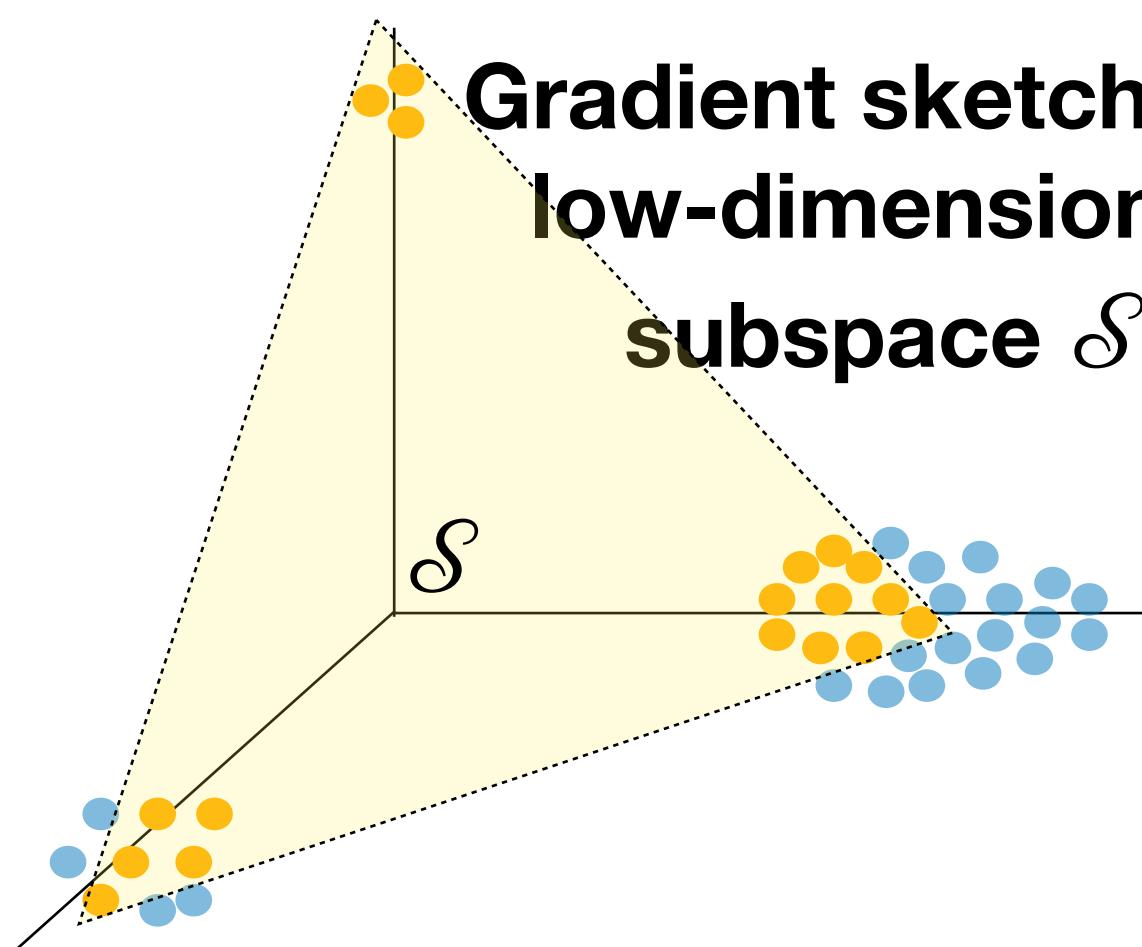
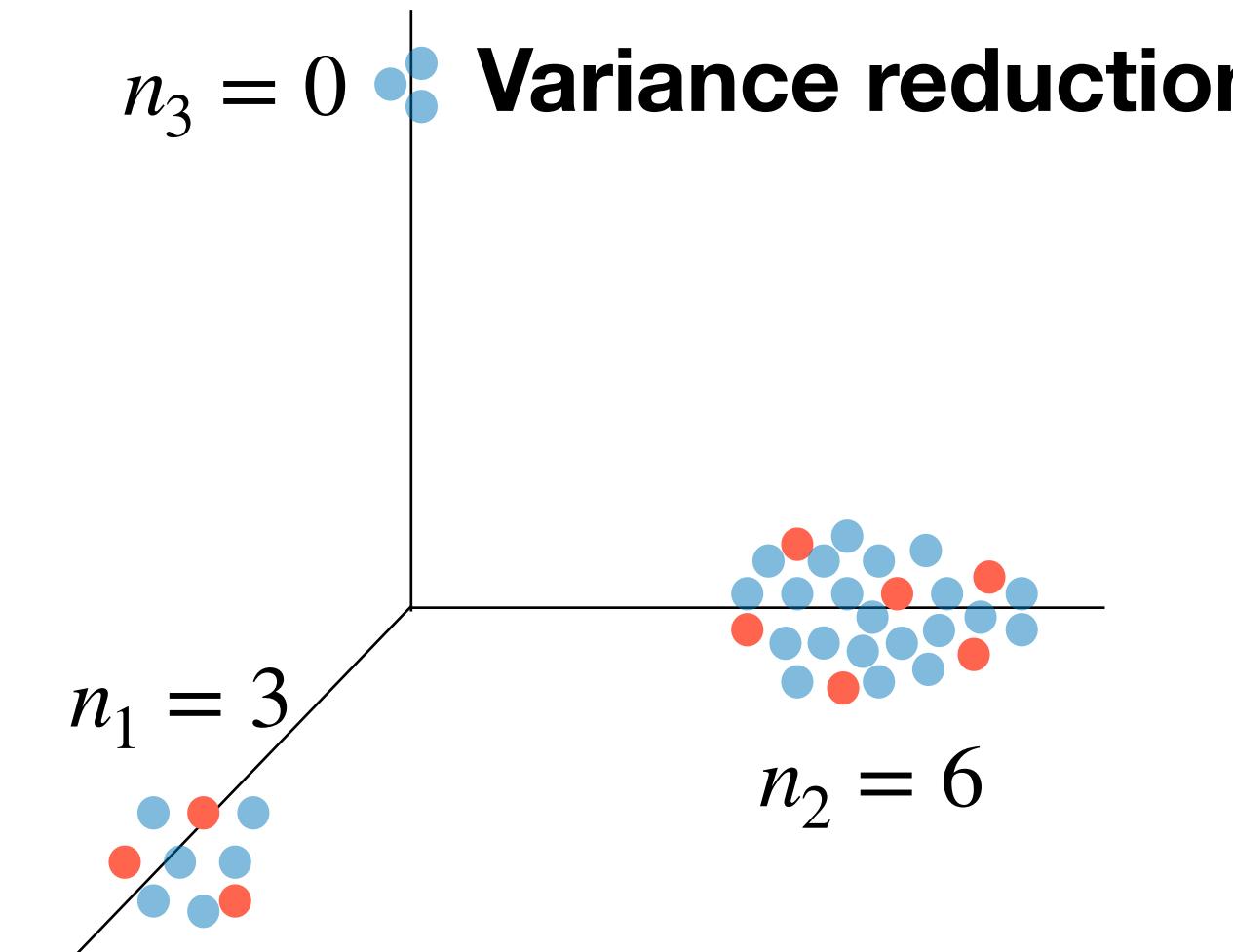
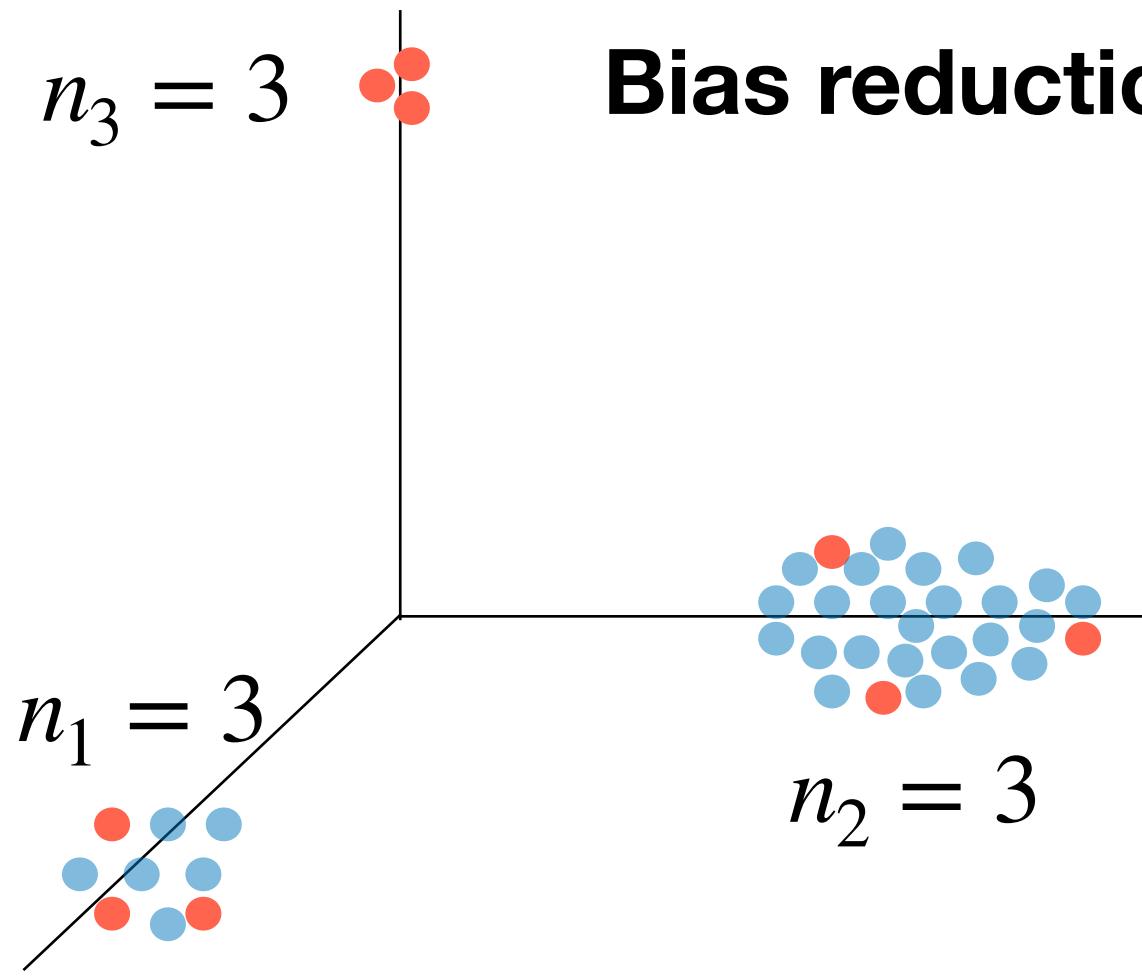
Efficiency of SkMM: (recall $m \ll \min\{N, r\}$)

- **Gradient sketching** is parallelizable with input-sparsity time: for $\text{nnz}(G) = \#\text{nonzeros in } G$
 - Gaussian embedding: $O(\text{nnz}(G)m)$
 - Fast JLT (sparse sign): $O(\text{nnz}(G)\log m)$
- **Moment matching** takes $O(m^3)$ for spectral decomposition. The optimization takes $O(Nm)$ per iteration

Relaxation of $\widetilde{\Sigma} \leq c_S \widetilde{\Sigma}_S$:

- $\widetilde{\Sigma} \leq c_S \widetilde{\Sigma}_S \iff V^T ((\widetilde{G})_S^T (\widetilde{G})_S / n) V \succeq \Lambda / c_S$
- Assume Σ, Σ_S commute such that imposing m diagonal constraints is sufficient

SkMM simultaneously controls variance and bias



SkMM on Synthetic Data: Regression

Synthetic high-dimensional linear probing

- Gaussian mixture model (GMM) $G \in \mathbb{R}^{N \times r}$
- $N = 2000, r = 2400 > N$
- $\bar{r} = 8$ well separated clusters of random sizes
- Grid search for the nearly optimal $\alpha > 0$

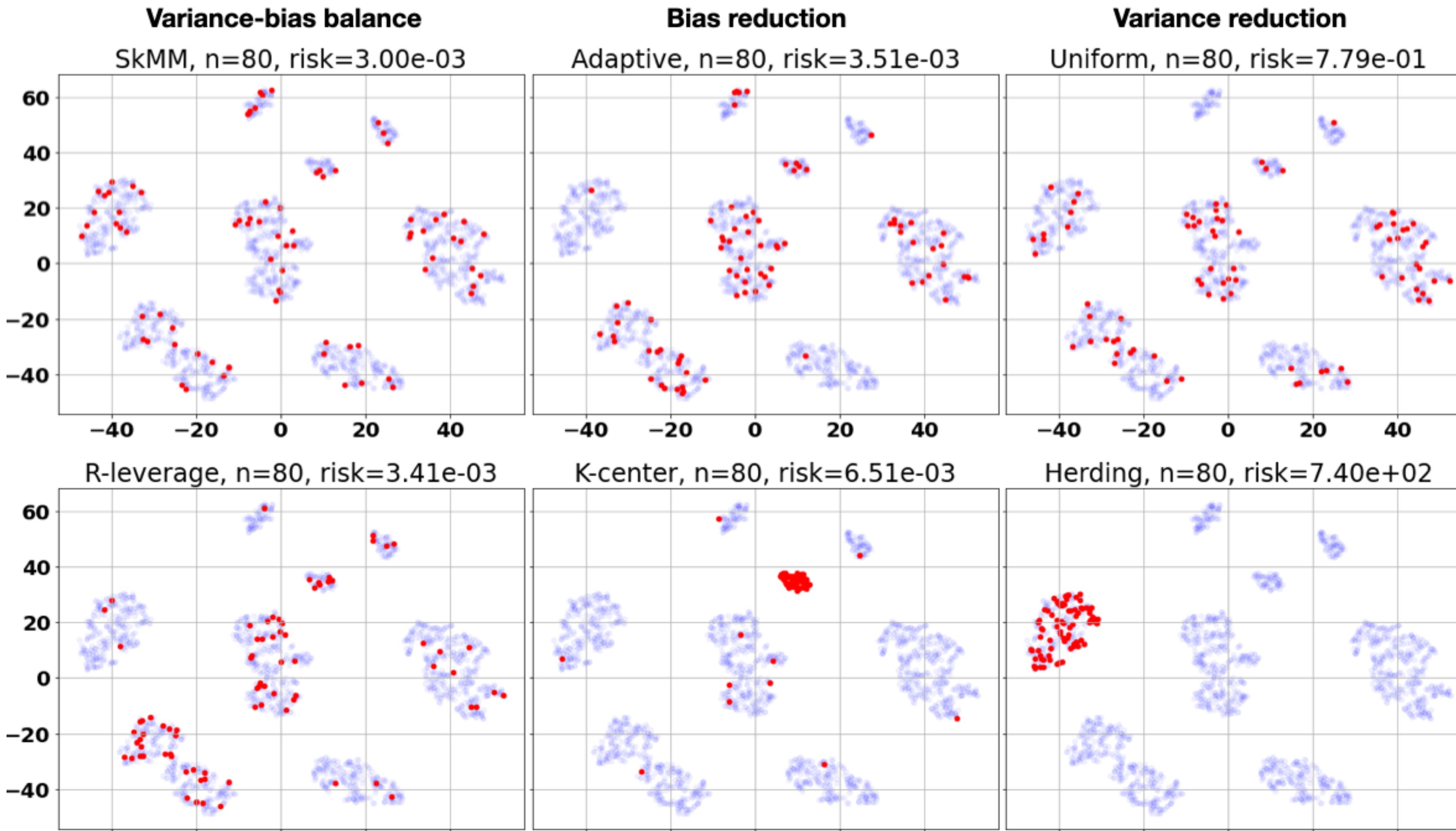
Baselines

- Herding
- Uniform sampling
- K-center greedy
- Adaptive sampling/random pivoting
- T(runcated)/R(idge) leverage score sampling

Table 1: Empirical risk $\mathcal{L}_{\mathcal{D}}(\boldsymbol{\theta}_S)$ on the GMM dataset at various n , under the same hyperparameter tuning where ridge regression over the full dataset \mathcal{D} with $N = 2000$ samples achieves $\mathcal{L}_{\mathcal{D}}(\boldsymbol{\theta}_{[N]}) = \mathbf{2.95e-3}$. For methods involving sampling, results are reported over 8 random seeds.

n	48	64	80	120	400	800	1600
Herding	7.40e+2	7.40e+2	7.40e+2	7.40e+2	7.38e+2	1.17e+2	2.95e-3
Uniform	(1.14 ± 2.71)e-1	(1.01 ± 2.75)e-1	(3.44 ± 0.29)e-3	(3.13 ± 0.14)e-3	(2.99 ± 0.03)e-3	(2.96 ± 0.01)e-3	(2.95 ± 0.00)e-3
K-center	(1.23 ± 0.40)e-2	(9.53 ± 0.60)e-2	(1.12 ± 0.45)e-2	(2.73 ± 1.81)e-2	(5.93 ± 4.80)e-2	(1.18 ± 0.64)e-1	(1.13 ± 0.70)e+0
Adaptive	(3.81 ± 0.65)e-3	(3.79 ± 1.37)e-3	(4.83 ± 1.90)e-3	(4.03 ± 1.35)e-3	(3.40 ± 0.67)e-3	(7.34 ± 3.97)e-3	(3.19 ± 0.16)e-3
T-leverage	(0.99 ± 1.65)e-2	(3.63 ± 0.49)e-3	(3.30 ± 0.30)e-3	(3.24 ± 0.14)e-3	(2.98 ± 0.01)e-3	(2.96 ± 0.01)e-3	(2.95 ± 0.00)e-3
R-leverage	(4.08 ± 1.58)e-3	(3.48 ± 0.43)e-3	(3.25 ± 0.31)e-3	(3.09 ± 0.06)e-3	(3.00 ± 0.02)e-3	(2.97 ± 0.01)e-3	(2.95 ± 0.00)e-3
SkMM	(3.54 ± 0.51)e-3	(3.31 ± 0.15)e-3	(3.12 ± 0.07)e-3	(3.07 ± 0.08)e-3	(2.98 ± 0.02)e-3	(2.96 ± 0.01)e-3	(2.95 ± 0.00)e-3

SkMM on Synthetic Data: Regression



SkMM for Classification: Linear Probing (LP)

Table 2: Accuracy and F1 score (%) of LP over CLIP on StanfordCars

	n	2000	2500	3000	3500	4000
Uniform Sampling	Acc	67.63 \pm 0.17	70.59 \pm 0.19	72.49 \pm 0.19	74.16 \pm 0.22	75.40 \pm 0.16
	F1	64.54 \pm 0.18	67.79 \pm 0.23	70.00 \pm 0.20	71.77 \pm 0.23	73.14 \pm 0.12
Herding [90]	Acc	67.22 \pm 0.16	71.02 \pm 0.13	73.17 \pm 0.22	74.64 \pm 0.18	75.71 \pm 0.29
	F1	64.07 \pm 0.23	68.28 \pm 0.15	70.64 \pm 0.28	72.22 \pm 0.26	73.26 \pm 0.39
Contextual Diversity [1]	Acc	67.64 \pm 0.13	70.82 \pm 0.23	72.66 \pm 0.12	74.46 \pm 0.17	75.77 \pm 0.12
	F1	64.51 \pm 0.17	68.18 \pm 0.25	70.05 \pm 0.11	72.13 \pm 0.15	73.35 \pm 0.07
Glister [43]	Acc	67.60 \pm 0.24	70.85 \pm 0.27	73.07 \pm 0.26	74.63 \pm 0.21	76.00 \pm 0.20
	F1	64.50 \pm 0.34	68.07 \pm 0.38	70.47 \pm 0.35	72.18 \pm 0.25	73.69 \pm 0.24
GraNd [63]	Acc	67.27 \pm 0.07	70.38 \pm 0.07	72.56 \pm 0.05	74.67 \pm 0.06	75.77 \pm 0.12
	F1	64.04 \pm 0.09	67.48 \pm 0.09	69.81 \pm 0.08	72.13 \pm 0.05	73.44 \pm 0.13
Forgetting [79]	Acc	67.59 \pm 0.10	70.99 \pm 0.05	72.54 \pm 0.07	74.81 \pm 0.05	75.74 \pm 0.01
	F1	64.85 \pm 0.13	68.53 \pm 0.07	70.30 \pm 0.05	72.59 \pm 0.04	73.74 \pm 0.02
DeepFool [59]	Acc	67.77 \pm 0.29	70.73 \pm 0.22	73.24 \pm 0.22	74.57 \pm 0.23	75.71 \pm 0.15
	F1	64.16 \pm 0.68	68.49 \pm 0.53	70.93 \pm 0.32	72.44 \pm 0.27	73.79 \pm 0.15
Entropy [19]	Acc	67.95 \pm 0.11	71.00 \pm 0.10	73.28 \pm 0.10	75.02 \pm 0.08	75.82 \pm 0.06
	F1	64.55 \pm 0.10	67.95 \pm 0.12	70.68 \pm 0.12	72.46 \pm 0.12	73.29 \pm 0.04
Margin [19]	Acc	67.53 \pm 0.14	71.19 \pm 0.09	73.09 \pm 0.14	74.66 \pm 0.11	75.57 \pm 0.13
	F1	64.16 \pm 0.15	68.33 \pm 0.14	70.37 \pm 0.17	72.03 \pm 0.11	73.14 \pm 0.20
Least Confidence [19]	Acc	67.68 \pm 0.11	70.99 \pm 0.14	73.04 \pm 0.05	74.65 \pm 0.09	75.58 \pm 0.08
	F1	64.09 \pm 0.20	68.03 \pm 0.20	70.30 \pm 0.07	72.02 \pm 0.10	73.15 \pm 0.12
SkMM-LP	Acc	68.27 \pm 0.03	71.53 \pm 0.05	73.61 \pm 0.02	75.12 \pm 0.01	76.34 \pm 0.02
	F1	65.29 \pm 0.03	68.75 \pm 0.06	71.14 \pm 0.03	72.64 \pm 0.02	74.02 \pm 0.10

StanfordCar dataset

- 196 imbalanced classes
- $N = 16,185$ images

Linear probing (LP)

- CLIP-pre-trained ViT
- $r = 100,548$

Last-two-layer finetuning (FT)

- ImageNet-pre-trained ResNet18
- $r = 2,459,844$

SkMM for Classification: Last-two-layer Finetuning (FT)

Table 3: Accuracy and F1 score (%) of FT over (the last two layers of) ResNet18 on StanfordCars

	n	2000	2500	3000	3500	4000
Uniform Sampling	Acc	29.19 \pm 0.37	32.83 \pm 0.19	35.69 \pm 0.35	38.31 \pm 0.16	40.35 \pm 0.26
	F1	26.14 \pm 0.39	29.91 \pm 0.16	32.80 \pm 0.37	35.38 \pm 0.19	37.51 \pm 0.23
Herding [90]	Acc	29.19 \pm 0.21	32.42 \pm 0.16	35.83 \pm 0.24	38.30 \pm 0.19	40.51 \pm 0.19
	F1	25.90 \pm 0.24	29.48 \pm 0.23	32.89 \pm 0.27	35.50 \pm 0.22	37.56 \pm 0.21
Contextual Diversity [1]	Acc	28.50 \pm 0.34	32.66 \pm 0.27	35.67 \pm 0.32	38.31 \pm 0.15	40.53 \pm 0.18
	F1	25.65 \pm 0.40	29.79 \pm 0.29	32.86 \pm 0.31	35.55 \pm 0.14	37.81 \pm 0.23
Glister [43]	Acc	29.16 \pm 0.26	32.91 \pm 0.19	36.03 \pm 0.20	38.16 \pm 0.12	40.47 \pm 0.16
	F1	26.33 \pm 0.19	30.05 \pm 0.28	33.26 \pm 0.18	35.41 \pm 0.14	37.63 \pm 0.17
GraNd [63]	Acc	28.59 \pm 0.17	32.67 \pm 0.20	35.83 \pm 0.16	38.58 \pm 0.15	40.70 \pm 0.11
	F1	25.66 \pm 0.15	29.70 \pm 0.22	32.76 \pm 0.16	35.72 \pm 0.15	37.83 \pm 0.11
Forgetting [79]	Acc	28.61 \pm 0.31	32.48 \pm 0.28	35.18 \pm 0.24	37.78 \pm 0.22	40.24 \pm 0.13
	F1	25.64 \pm 0.25	29.58 \pm 0.30	32.38 \pm 0.20	35.16 \pm 0.18	37.41 \pm 0.14
DeepFool [59]	Acc	24.97 \pm 0.20	29.02 \pm 0.17	32.60 \pm 0.18	35.59 \pm 0.24	38.20 \pm 0.22
	F1	22.11 \pm 0.11	26.08 \pm 0.29	29.83 \pm 0.27	32.92 \pm 0.33	35.47 \pm 0.22
Entropy [19]	Acc	28.87 \pm 0.13	32.84 \pm 0.20	35.64 \pm 0.20	37.96 \pm 0.11	40.29 \pm 0.27
	F1	25.95 \pm 0.17	30.03 \pm 0.17	32.85 \pm 0.23	35.19 \pm 0.12	37.33 \pm 0.34
Margin [19]	Acc	29.18 \pm 0.12	32.73 \pm 0.15	35.67 \pm 0.30	38.27 \pm 0.20	40.58 \pm 0.06
	F1	26.15 \pm 0.12	29.66 \pm 0.05	32.86 \pm 0.30	35.61 \pm 0.17	37.77 \pm 0.07
Least Confidence [19]	Acc	29.05 \pm 0.07	32.88 \pm 0.13	35.66 \pm 0.18	38.25 \pm 0.20	39.91 \pm 0.09
	F1	26.18 \pm 0.04	30.03 \pm 0.14	32.79 \pm 0.15	35.42 \pm 0.16	37.14 \pm 0.12
SkMM-FT	Acc	29.44 \pm 0.09	33.48 \pm 0.04	36.11 \pm 0.12	39.18 \pm 0.03	41.77 \pm 0.07
	F1	26.71 \pm 0.10	30.75 \pm 0.05	33.24 \pm 0.05	36.38 \pm 0.05	39.07 \pm 0.10

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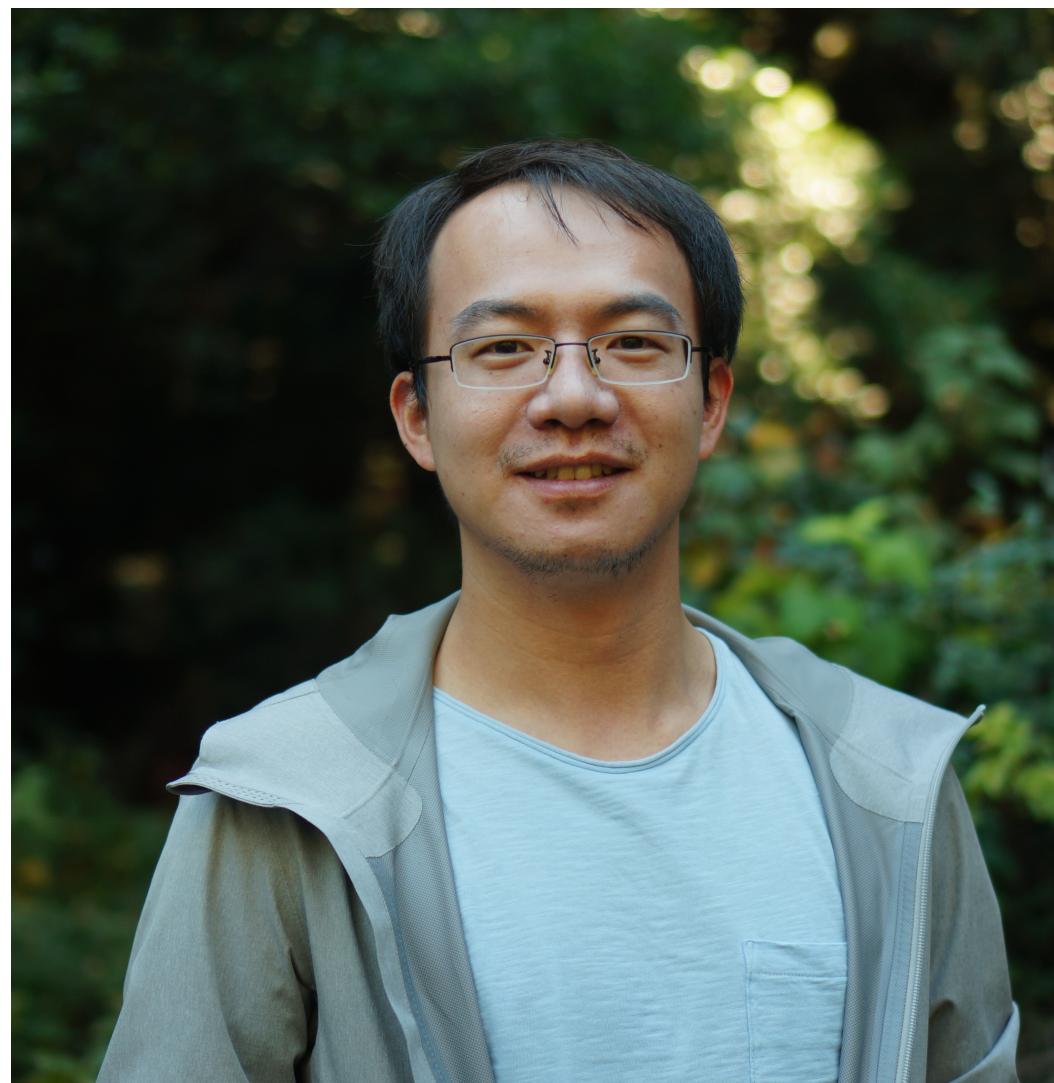
Takeaways

- A rigorous generalization analysis on data selection for finetuning
 - Low-dimensional data selection: variance reduction (V -optimality)
 - **High-dimensional data selection:** variance-bias tradeoff
- **Gradient sketching** provably finds a low-dimensional parameter subspace \mathcal{S} with small bias
 - Reducing variance over \mathcal{S} preserves the fast-rate generalization $O(\dim(\mathcal{S})/n)$
- **SkMM** — a scalable two-stage data selection method for finetuning that simultaneously
 - **Explores** the high-dimensional parameter space via **gradient sketching** and
 - **Exploits** the information in the low-dimensional subspace via **moment matching**

Outline

- Data selection for statistical models in kernel regime — **finetuning**
 - A theory on data selection for finetuning that
 - Extends the classical wisdom of **V-optimal experimental design** in low dimensions
 - To high-dimensional data selection under low intrinsic dimensions via **sketching**
 - A fast and effective data selection algorithm for finetuning: **Sketchy Moment Matching (SkMM)**
- Data selection for low-rank approximation — **interpolative decomposition (ID)**
 - Select minimum possible rows as a basis to form a good low-rank approximation
 - A fast and accurate adaptive ID algorithm: **Robust Blockwise Random Pivoting (RBRP)**

Robust Blockwise Random Pivoting: Fast and Accurate Adaptive Interpolative Decomposition



Chao Chen
UT Austin → NCSU



Per-Gunnar Martinsson
UT Austin

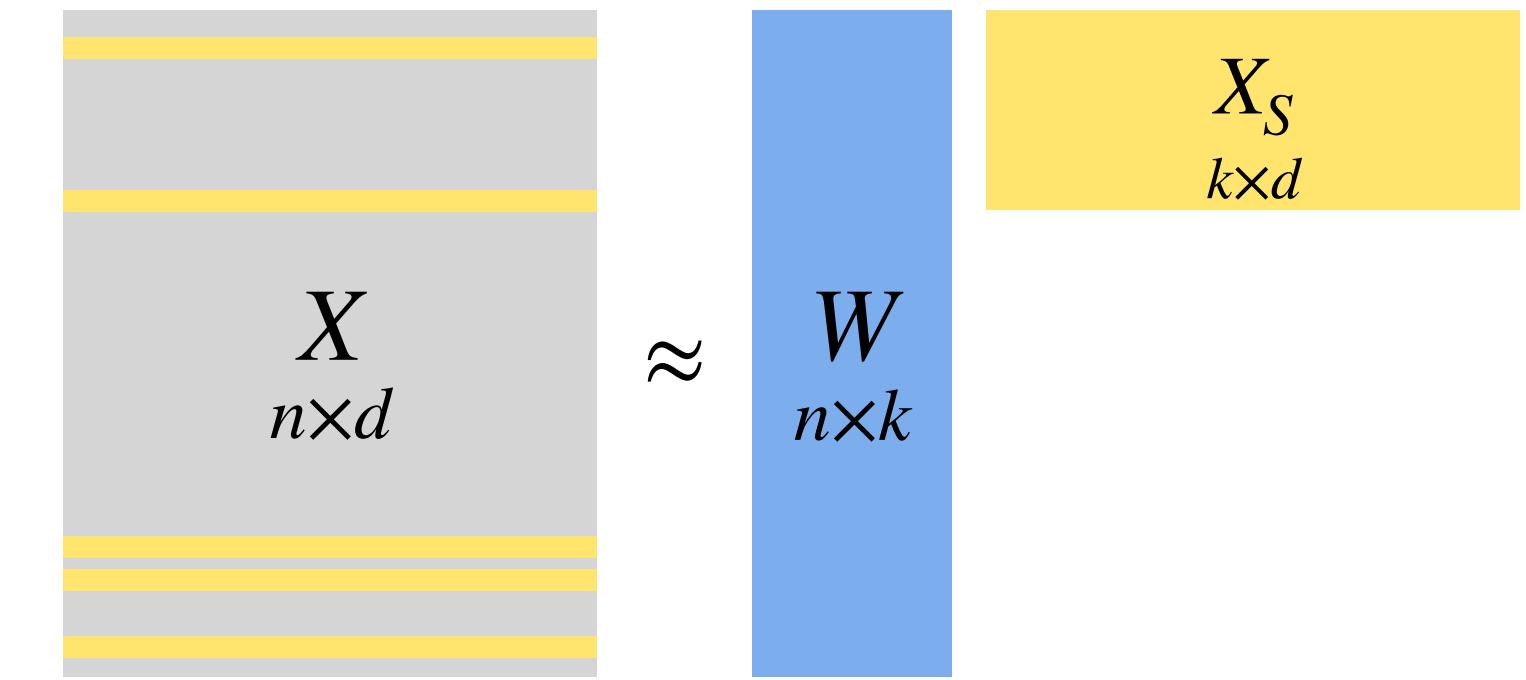


Katherine Pearce
UT Austin

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, ϵ) -ID of X — $X \approx WX_S$ such that

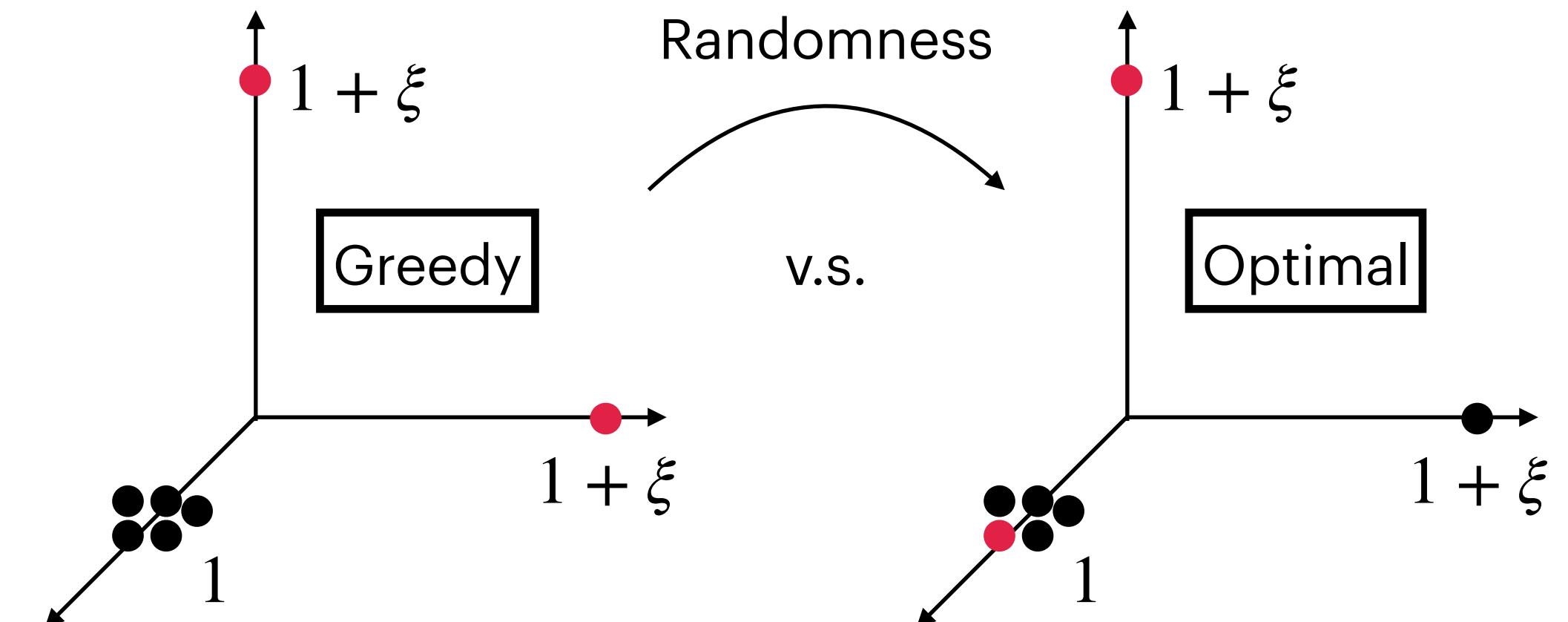
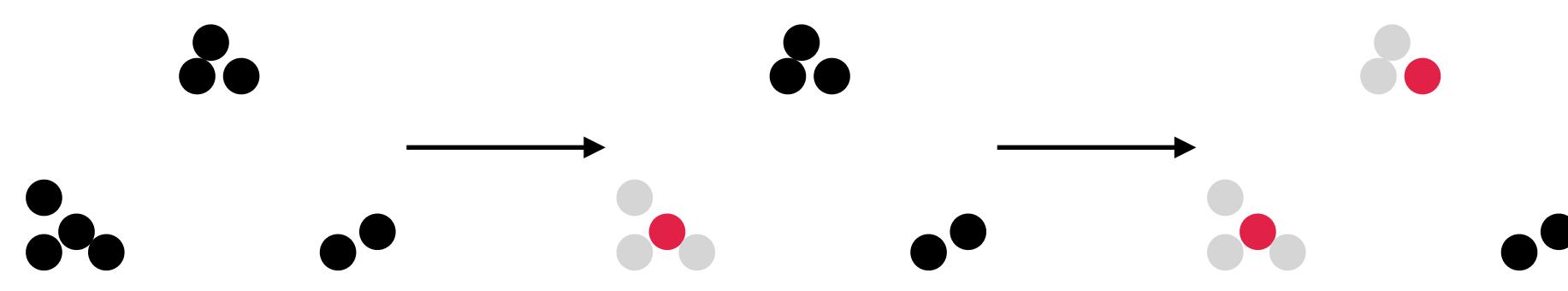
$$\|X - WX_S\|_F^2 \leq (1 + \epsilon)\|X - X_{\langle r \rangle}\|_F^2$$



- $S = \{s_1, \dots, s_k\} \subseteq [n]$ 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)

Adaptiveness & Randomness

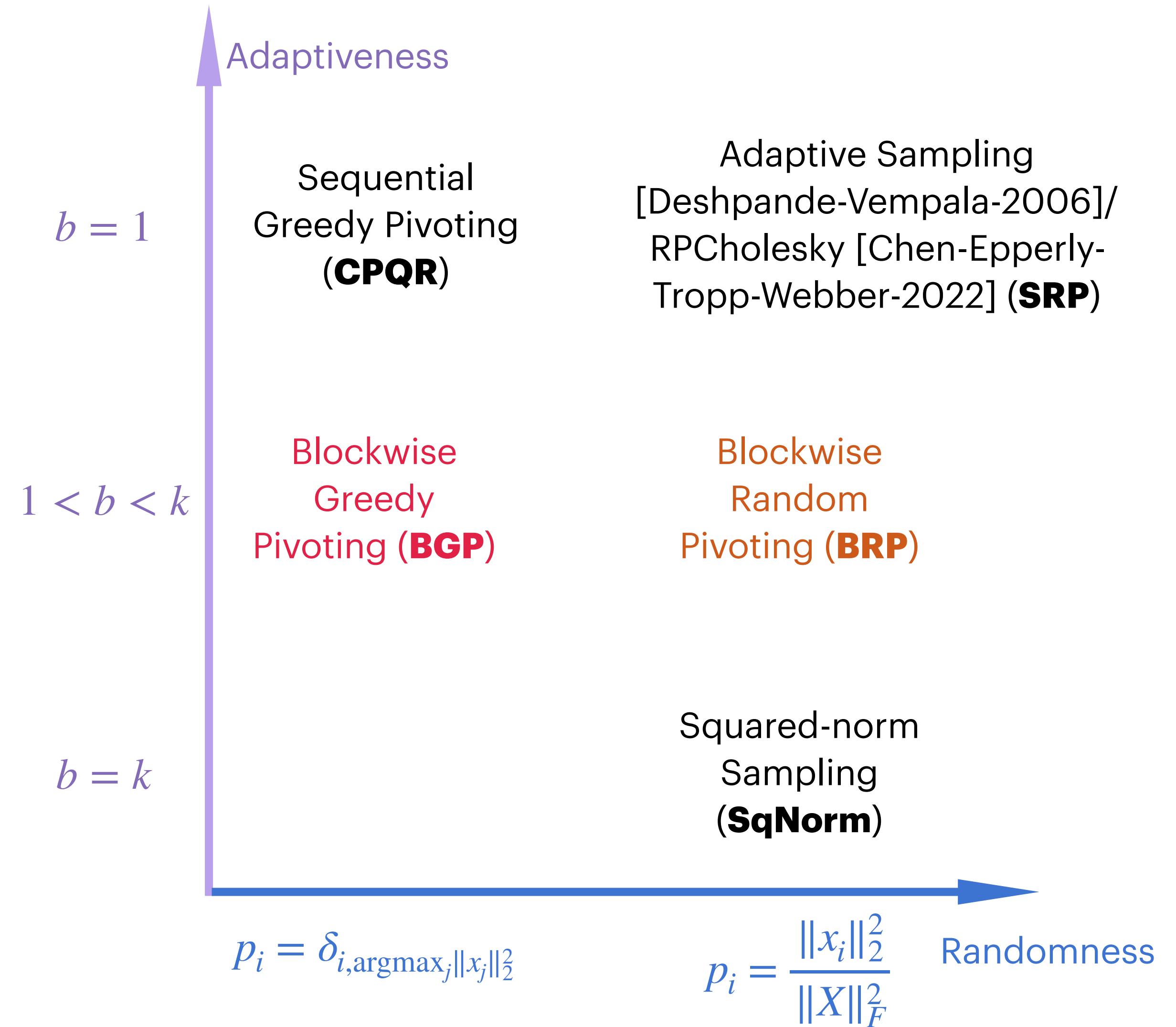
- **Adaptiveness**
 - Each new skeleton selection is aware of the previously selected skeleton subset
 - By selecting according to the residual
 - Common adaptive residual updates:
 - Gram-Schmidt (QR)
 - Gaussian elimination (LU)
- **Randomness** (in contrast to greedy)
 - Intuition: balance exploitation 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** [Deshpande-Rademacher-Vempala-Wang-2006, Belabbas-Wolfe-2009, etc.]
 - Pro: nearly optimal expected skeleton complexity
 - Con: expensive to compute
- **Leverage score sampling** [Mahoney-Drineas-2009, Cohen-Musco-Musco-2017, etc.]
 - Pro: can be estimated efficiently for large-scale problems (e.g., tensor Khatri-Rao product)
 - Con: expensive to compute
- **Uniform sampling** [Cohen-Lee-Musco-Musco-Peng-Sidford-2015]
 - 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 $t = 1, \dots, k$
 - Row pivoted QR (**CPQR**) [Voronin-Martinsson-2017]:
$$s_t \leftarrow \underset{i}{\operatorname{argmax}} \|Y_{i,:}^{(t-1)}\|_2^2 + \text{Gram-Schmidt}$$
 - LU with partial pivoting (**LUPP**) [D-Martinsson-2023]:
$$s_t \leftarrow \underset{i}{\operatorname{argmax}} |Y_{i,t}^{(t-1)}| + \text{Gaussian Elimination}$$
- Pro: fast, accurate, robust to adversarial inputs
- Con: require prior knowledge of k

Two Stages of ID Constructions

- Stage I: Skeleton selection
 - Find a good skeleton subset S :

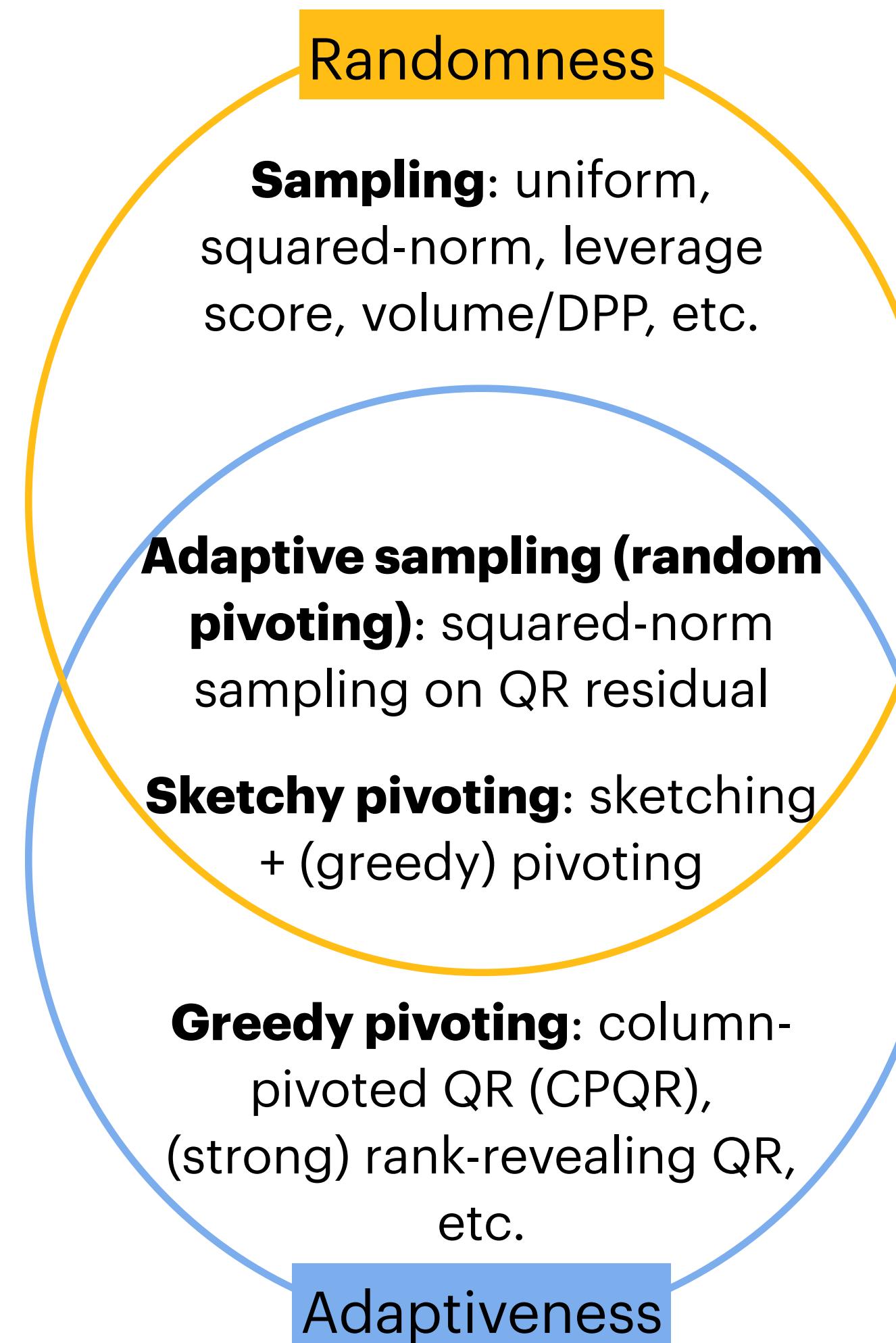
$$\min_{S \subset [n]} \min_{W \in \mathbb{R}^{n \times |S|}} \|X - WX_S\|_F^2$$

- **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 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, ϵ) -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

ID Algorithms with Adaptiveness & Randomness



Epperly-Tropp-Webber-2024 proved this conjecture for an analogous algorithm based on rejective sampling!

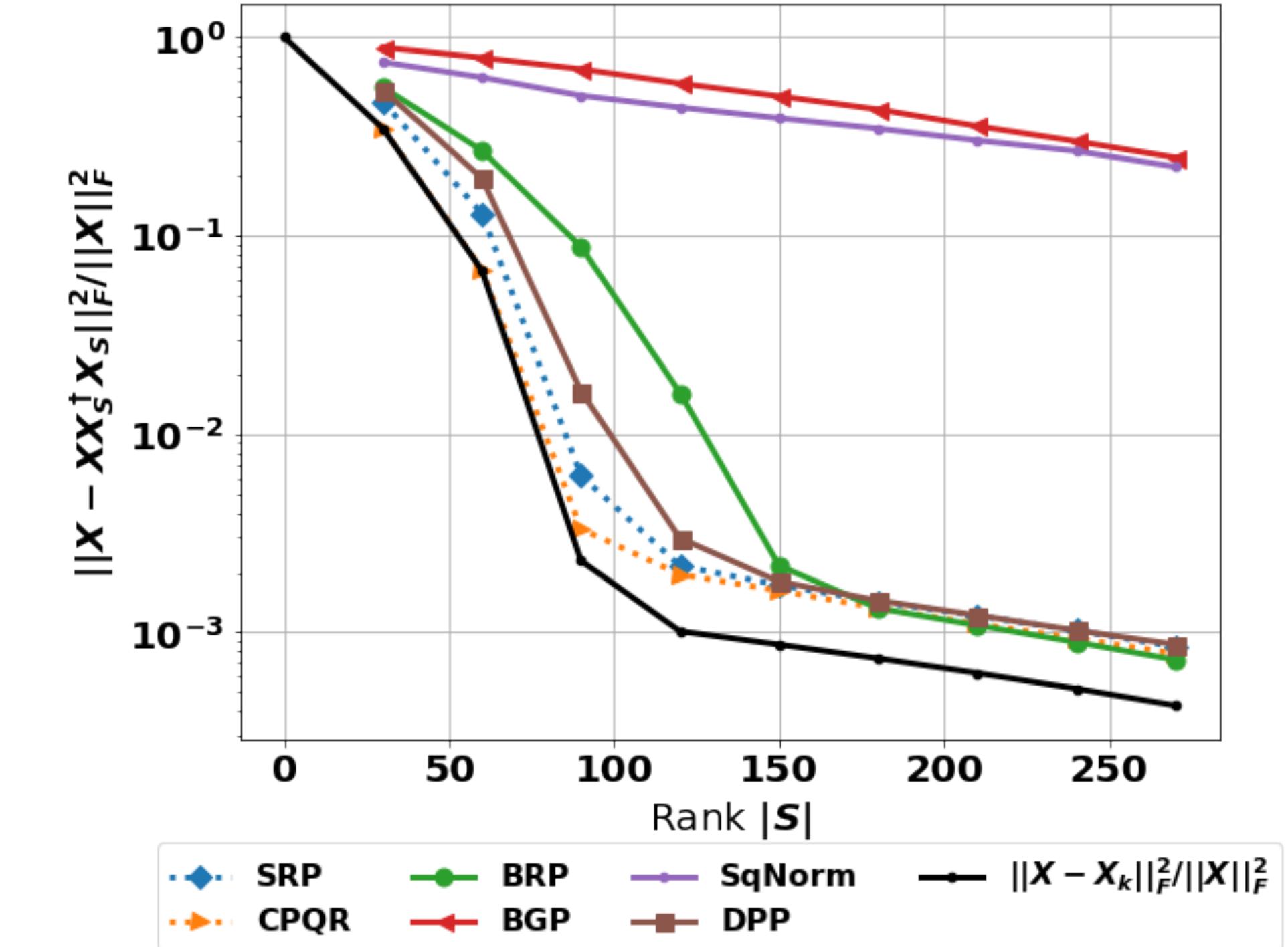
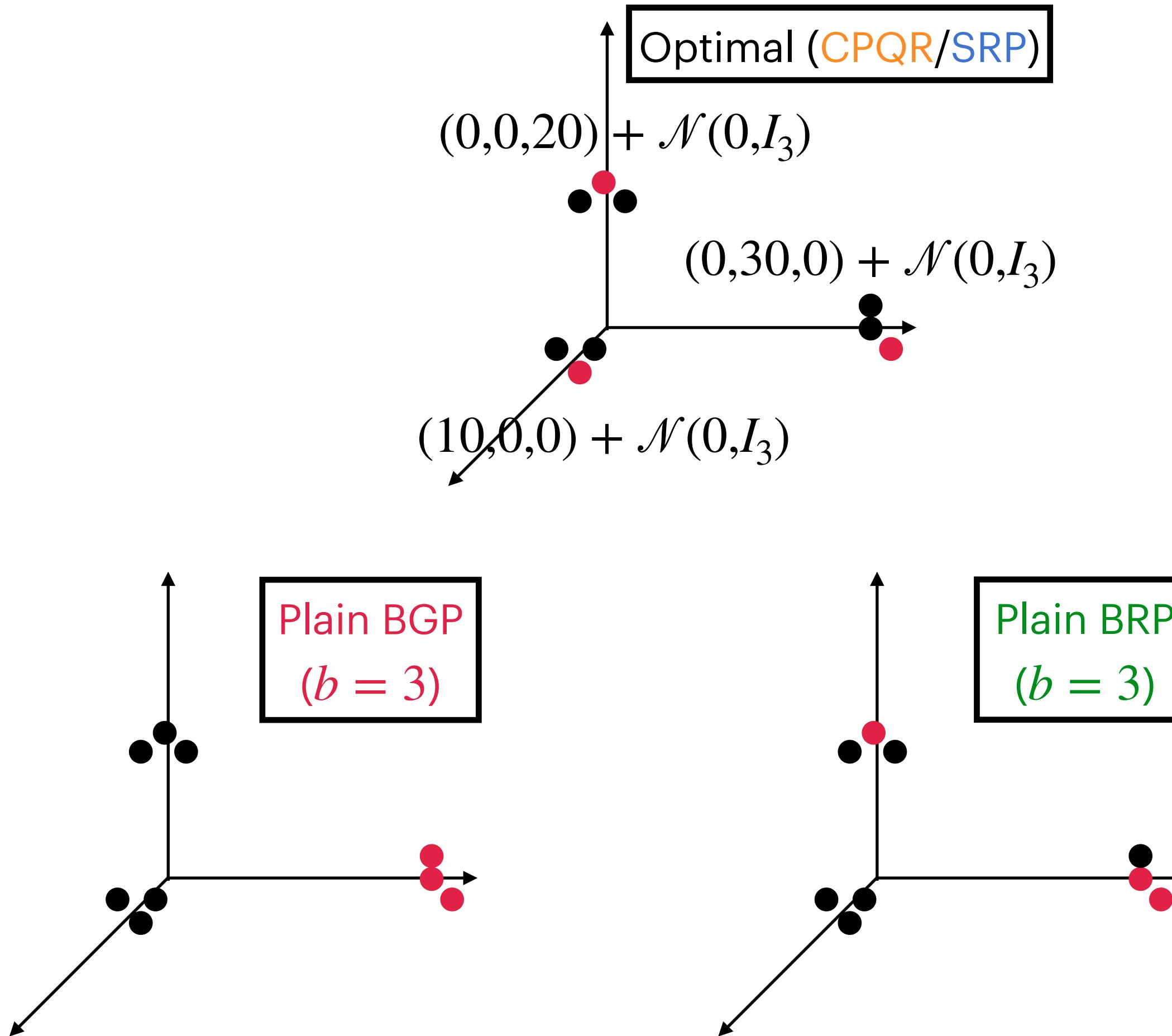
Algorithm	Skeleton Complexity	Asymp. Cost + Parallelizability	Error-reveal	ID-reveal
Greedy Pivoting	$k \geq (1 + (1 + \epsilon)\eta_r)n$	$O(ndk)$ sequential	✓	Exact
Squared-norm Sampling	$k \geq \frac{r - 1}{\epsilon\eta_r} + \frac{1}{\epsilon}$	$O(nd)$ parallel	✗	Non
Random Pivoting	$k \geq k_{RP}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

* $\eta_r = \|X - X_{\leq r}\|_F^2 / \|X\|_F^2$ 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 $\{10j \cdot e_j\}_{j \in [k]}$, $n = 20k$, $d = 500$, $b = 30$

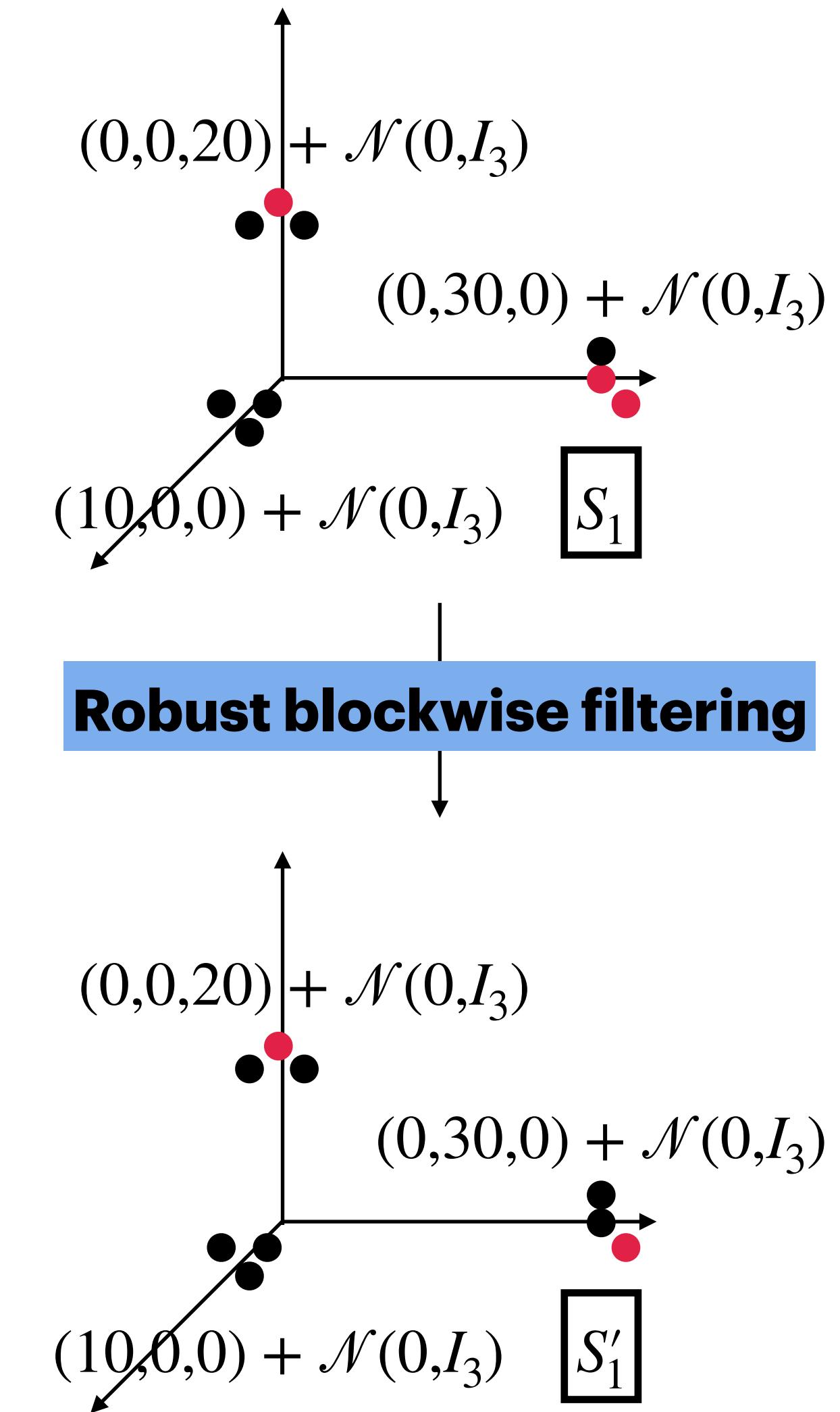


- Sequential pivoting (CPQR & SRP) is nearly optimal
- Plain blockwise pivoting (BRP/BGP, especially BGP) suffers from suboptimal skeleton complexities (up to b times)
- Squared-norm sampling (SqNorm) tends to fail

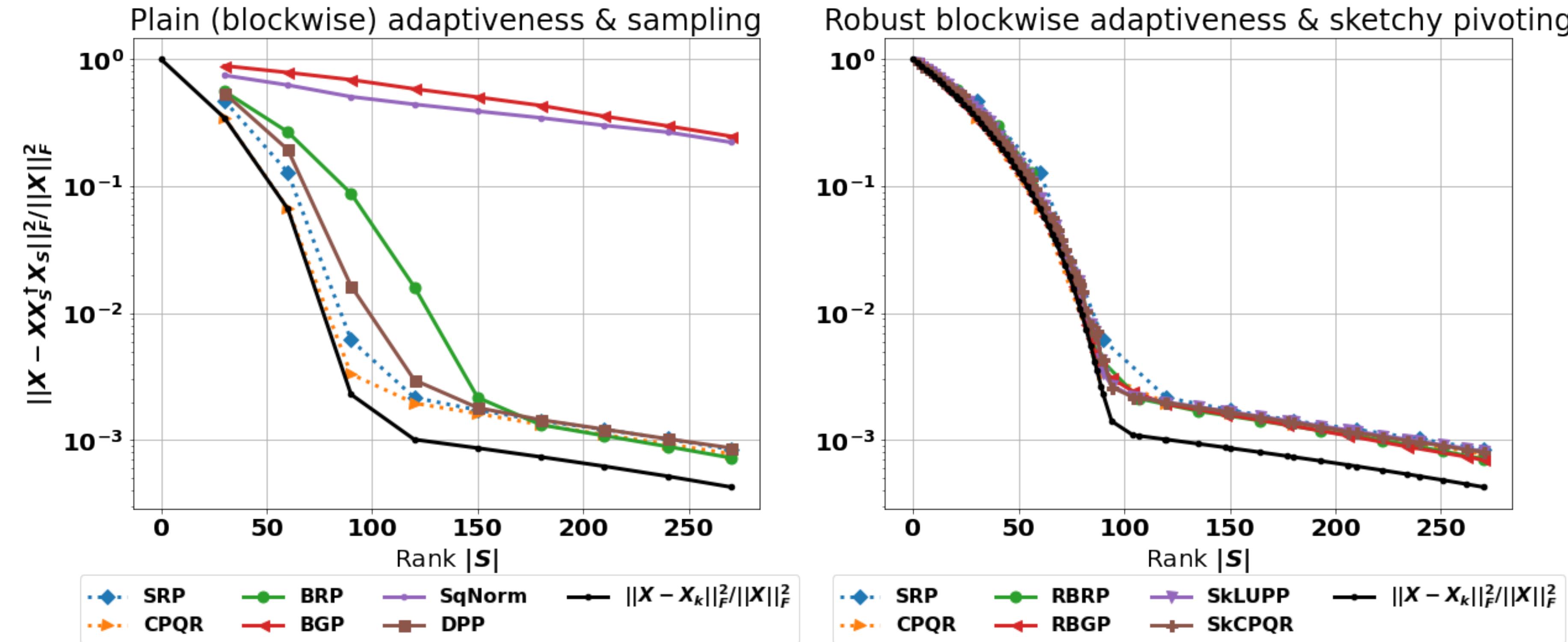
Robust Blockwise Random Pivoting

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)
 - $\min_{S'_t=S_t(\pi(1:b'))} b' \text{ s.t. } \|X_{S_t} - X_{S'_t}\|_F^2 < \tau_b \|X_{S_t}\|_F^2$ (e.g., $\tau_b = \frac{1}{b}$)
 - $S^{(t)} \leftarrow S^{(t-1)} \cup S'_t$ and $X^{(t)} \leftarrow X^{(t-1)} \left(I_d - X_{S'_t}^\dagger X_{S'_t} \right)$
 - $S \leftarrow S^{(t)}$, $k = |S|$

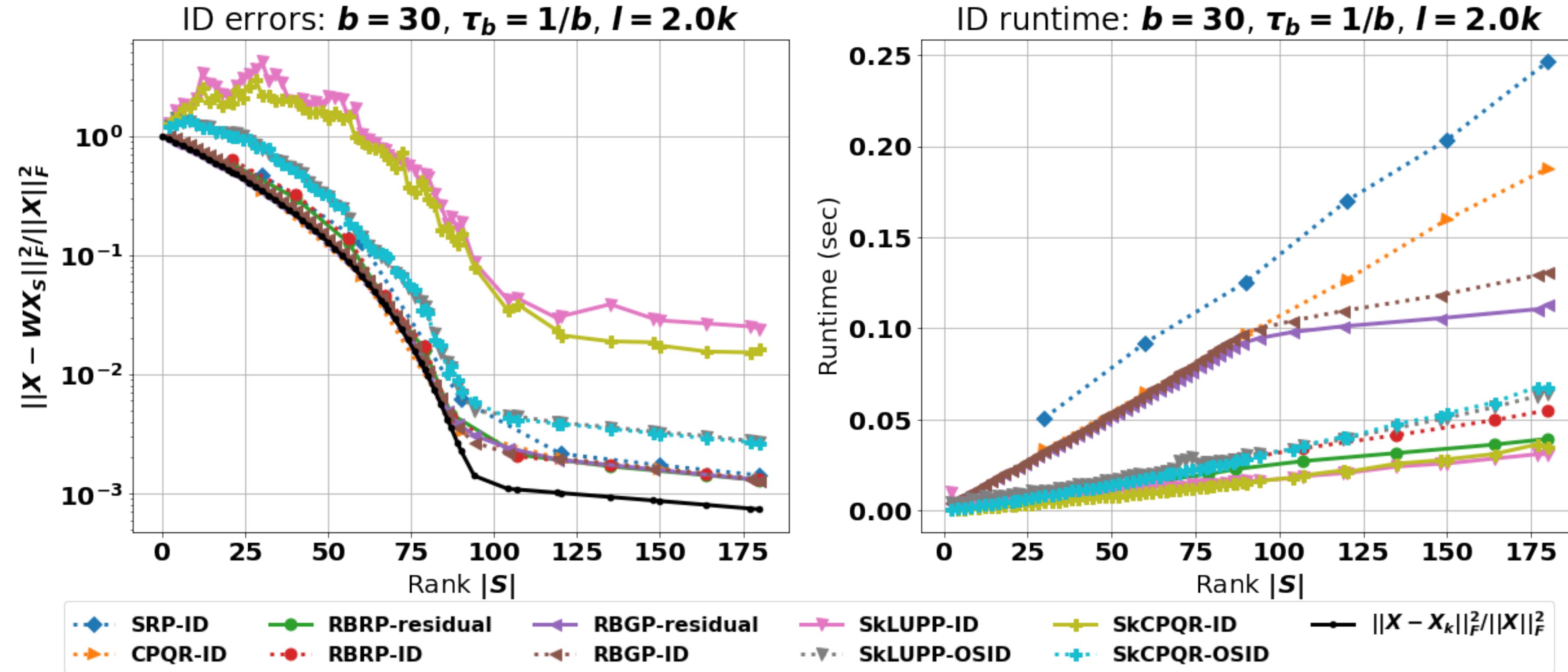


Robust Blockwise Random Pivoting: Robustness



- 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

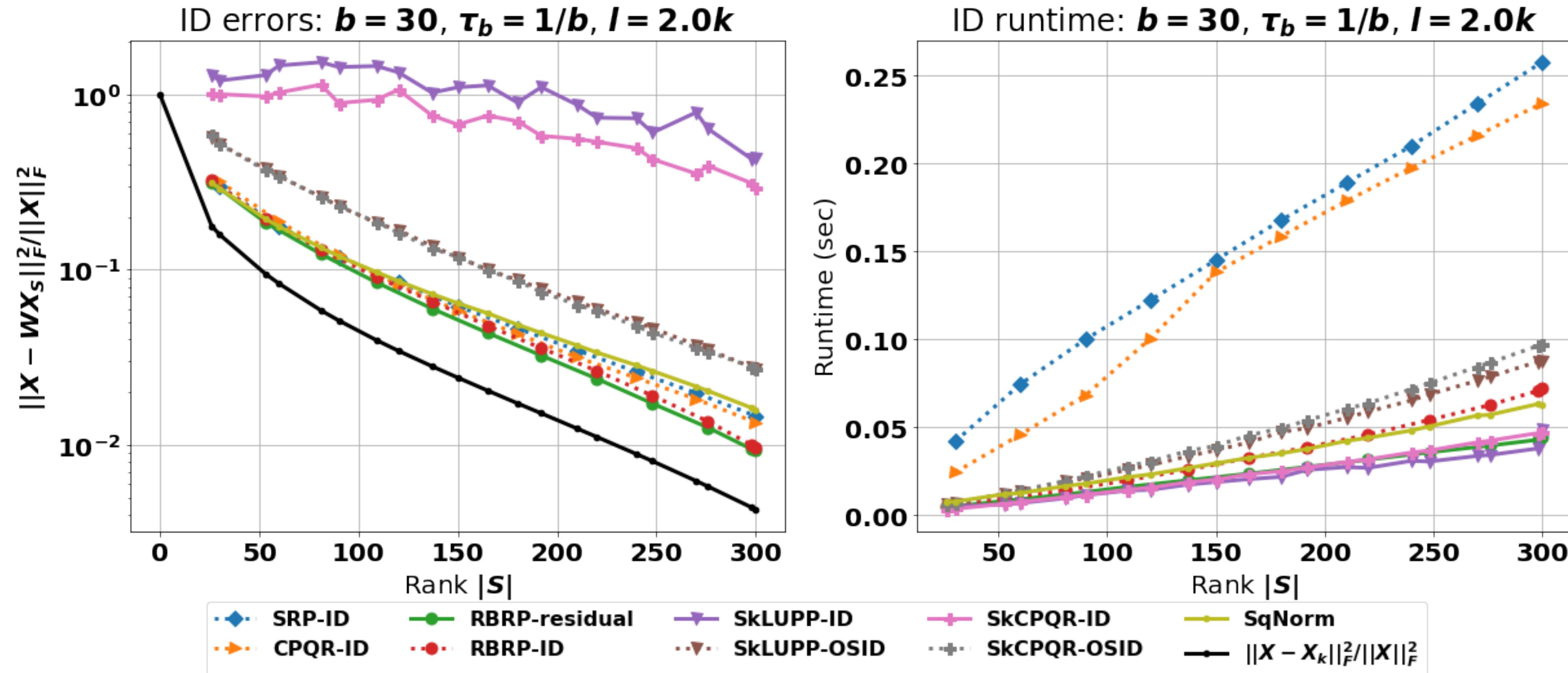


- Robust blockwise filtering (**RBRP** and **RBGP**) brings nearly optimal skeleton complexities
- **RBGP** tends to be slowed down much more significantly than **RBRP** by robust blockwise filtering
- For ID: **RBRP-ID** is almost as fast as sketchy pivoting (**SkLUPP-ID/SkCPQR-ID**), while enjoying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

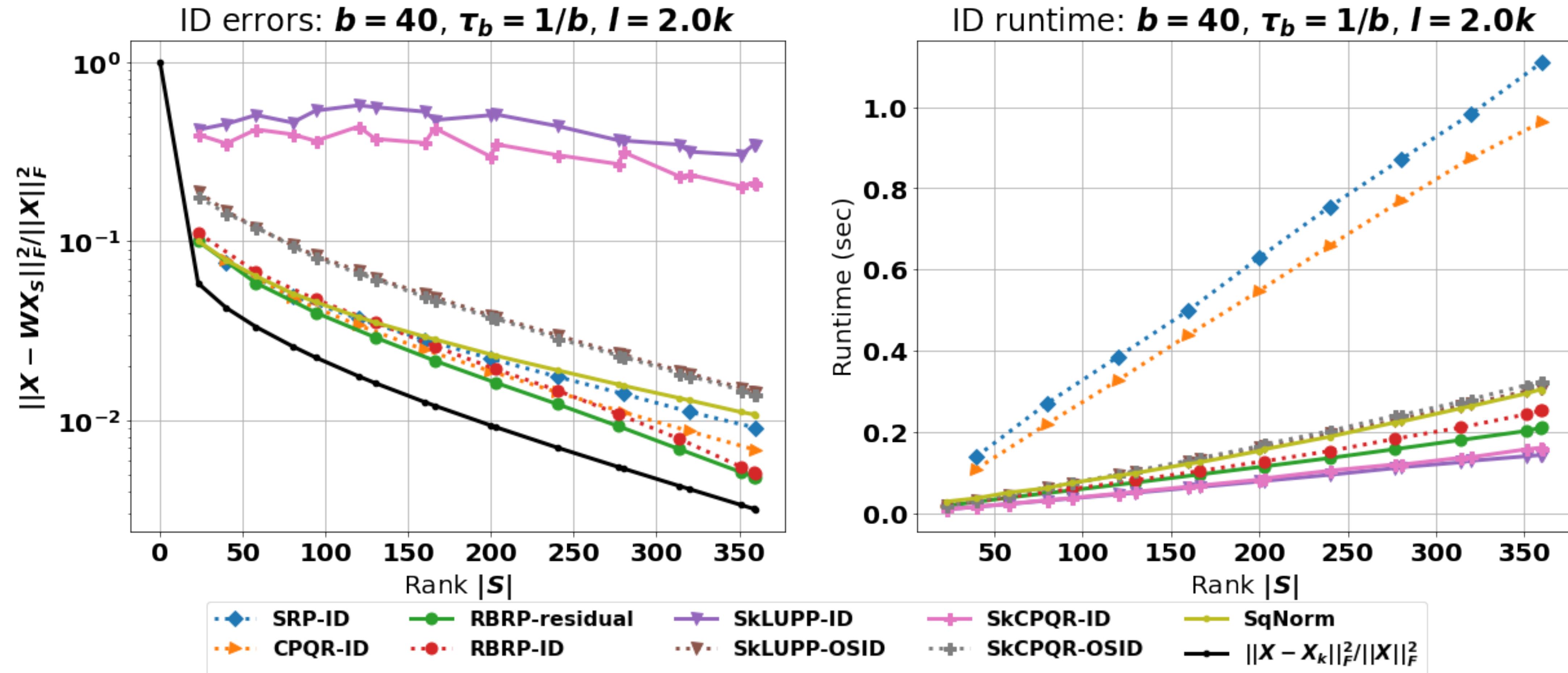
Exact- v.s. Inexact-ID-revealing Algorithms

- **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



Takeaways

- 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



SkMM: <https://arxiv.org/pdf/2407.06120>



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

Qs: Are there connections between ID and finetuning?

Theorem (Variance-bias tradeoff): Given a coresset S of size n , let $P_{\mathcal{S}} \in \mathbb{R}^{r \times r}$ be the orthogonal projector onto any subspace $\mathcal{S} \subset \text{Range}(\Sigma_S)$, and $P_{\mathcal{S}}^\perp = I_r - P_{\mathcal{S}}$. There exists $\alpha > 0$ such that (2) satisfies

$$\mathbb{E}[\text{ER}(\theta_S)] \leq \min_{\mathcal{S} \subset \text{Range}(\Sigma_S)} \underbrace{\frac{2\sigma^2}{n} \text{tr}(\Sigma(P_{\mathcal{S}} \Sigma_S P_{\mathcal{S}})^\dagger)}_{\text{variance}} + \underbrace{2\text{tr}(\Sigma P_{\mathcal{S}}^\perp) \|\theta_*\|_2^2}_{\text{bias}}$$

Bias reduction \asymp learning in noiseless settings \asymp minimizing skeletonization error of ID

- In the noiseless setting $\sigma = 0$, the expected excess risk is controlled by the variance only:

$$\mathbb{E}[\text{ER}(\theta_S)] \leq \min_{\mathcal{S} \subset \text{Range}(\Sigma_S)} 2\text{tr}(\Sigma P_{\mathcal{S}}^\perp) \|\theta_*\|_2^2$$

- $\text{tr}(\Sigma P_{\mathcal{S}}^\perp) = \|GP_{\mathcal{S}}^\perp\|_F^2/N$ can be viewed as a low-rank approximation error
- Taking $\mathcal{S} = \text{Row}(G_S)$, $\text{tr}(\Sigma P_{\mathcal{S}}^\perp)$ is effectively the skeletonization error of the ID associated with S

Qs: What's wrong with adaptive sampling for finetuning?

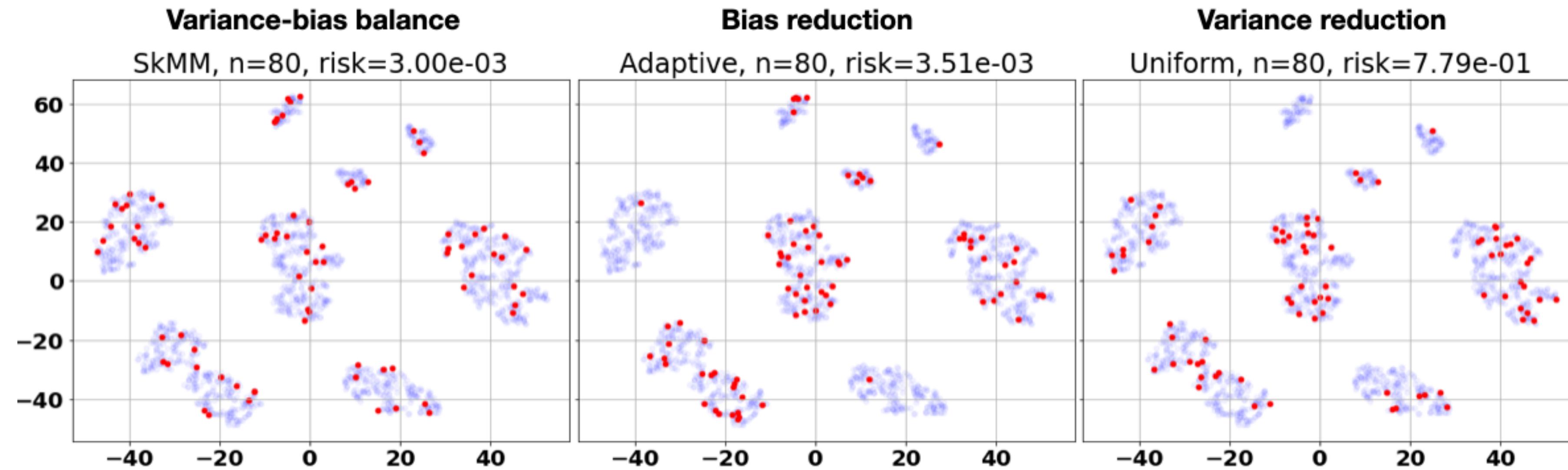


Table 1: Empirical risk $\mathcal{L}_{\mathcal{D}}(\boldsymbol{\theta}_S)$ on the GMM dataset at various n , under the same hyperparameter tuning where ridge regression over the full dataset \mathcal{D} with $N = 2000$ samples achieves $\mathcal{L}_{\mathcal{D}}(\boldsymbol{\theta}_{[N]}) = \mathbf{2.95e-3}$. For methods involving sampling, results are reported over 8 random seeds.

n	48	64	80	120	400	800	1600
Hherding	7.40e+2	7.40e+2	7.40e+2	7.40e+2	7.38e+2	1.17e+2	2.95e-3
Uniform	$(1.14 \pm 2.71)e-1$	$(1.01 \pm 2.75)e-1$	$(3.44 \pm 0.29)e-3$	$(3.13 \pm 0.14)e-3$	$(2.99 \pm 0.03)e-3$	$(2.96 \pm 0.01)e-3$	$(2.95 \pm 0.00)e-3$
K-center	$(1.23 \pm 0.40)e-2$	$(9.53 \pm 0.60)e-2$	$(1.12 \pm 0.45)e-2$	$(2.73 \pm 1.81)e-2$	$(5.93 \pm 4.80)e-2$	$(1.18 \pm 0.64)e-1$	$(1.13 \pm 0.70)e+0$
Adaptive	$(3.81 \pm 0.65)e-3$	$(3.79 \pm 1.37)e-3$	$(4.83 \pm 1.90)e-3$	$(4.03 \pm 1.35)e-3$	$(3.40 \pm 0.67)e-3$	$(7.34 \pm 3.97)e-3$	$(3.19 \pm 0.16)e-3$
T-leverage	$(0.99 \pm 1.65)e-2$	$(3.63 \pm 0.49)e-3$	$(3.30 \pm 0.30)e-3$	$(3.24 \pm 0.14)e-3$	$(2.98 \pm 0.01)e-3$	$(2.96 \pm 0.01)e-3$	$(2.95 \pm 0.00)e-3$
R-leverage	$(4.08 \pm 1.58)e-3$	$(3.48 \pm 0.43)e-3$	$(3.25 \pm 0.31)e-3$	$(3.09 \pm 0.06)e-3$	$(3.00 \pm 0.02)e-3$	$(2.97 \pm 0.01)e-3$	$(2.95 \pm 0.00)e-3$
SkMM	$(3.54 \pm 0.51)e-3$	$(3.31 \pm 0.15)e-3$	$(3.12 \pm 0.07)e-3$	$(3.07 \pm 0.08)e-3$	$(2.98 \pm 0.02)e-3$	$(2.96 \pm 0.01)e-3$	$(2.95 \pm 0.00)e-3$