

Data Selection under Low Intrinsic Dimension: from Interpolative Decomposition to Ridge Regression

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Joint Mathematics Meeting, Jan 11, 2025



Low Intrinsic Dimension & Data Selection

- Low intrinsic dimension is ubiquitous in real world
 - Example: A language model with **341M parameters** can be finetuned in a **dimension-322 subspace** with **less than 6K samples** [Aghajanyan-Zettlemoyer-Gupta-2020]
- Learning under low intrinsic dimension **with limited data, data selection becomes crucial**



Low Intrinsic Dimension & Data Selection

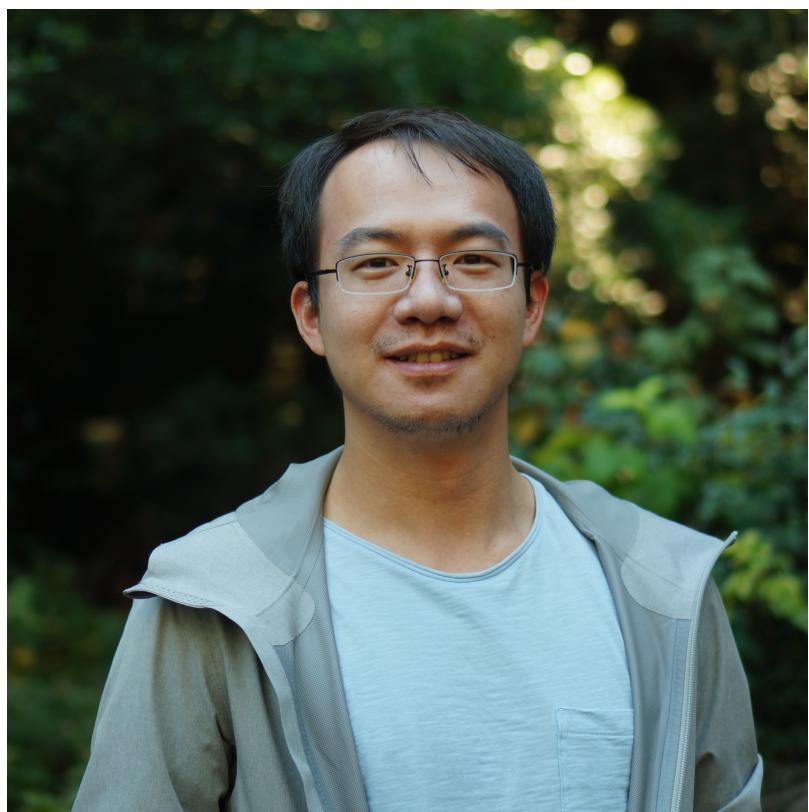
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How to **select informative data** for learning under **low intrinsic dimension**?

- Learning without noise: low-rank interpolative decomposition (ID)
- Learning with noise: low-rank approximation (bias) + variance reduction

Robust Blockwise Random Pivoting: Fast and Accurate Adaptive Interpolative Decomposition



Chao Chen
NCSU



Per-Gunnar Martinsson
UT Austin

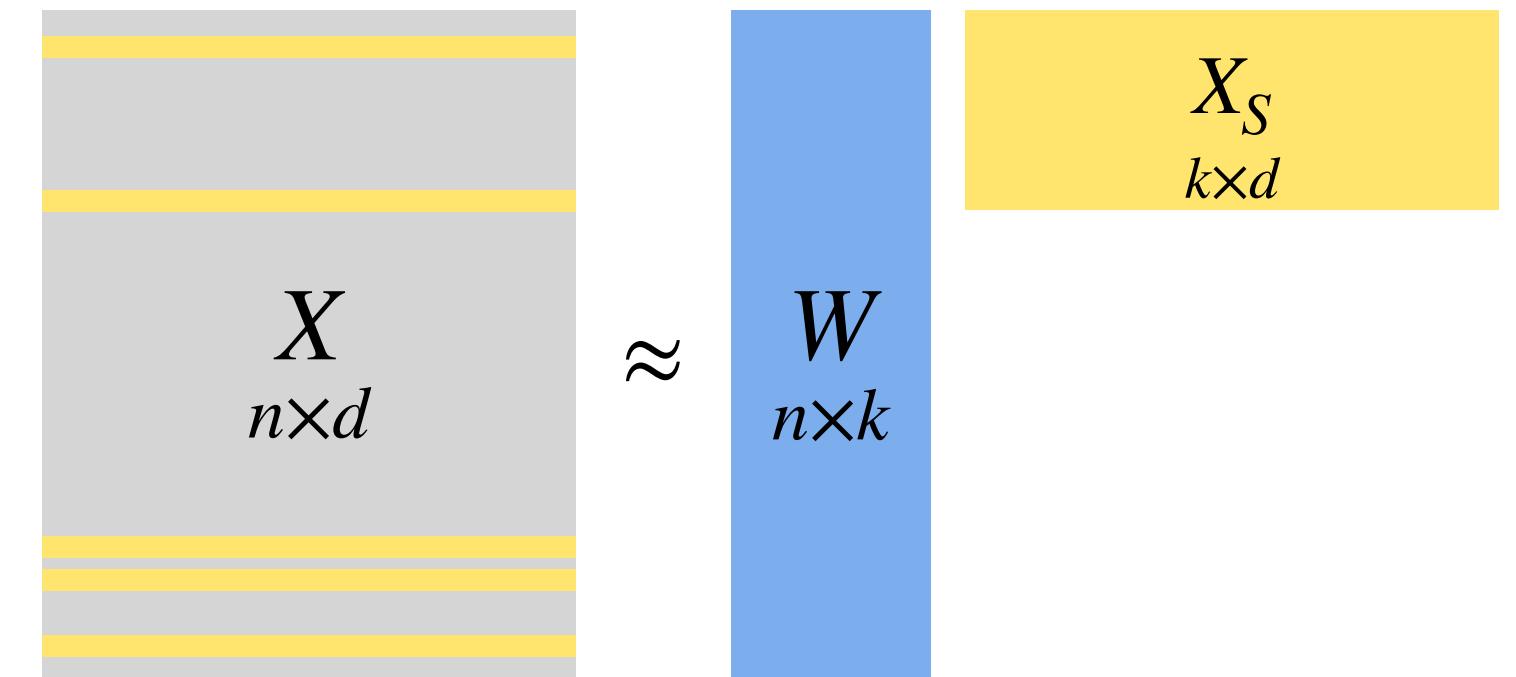


Katherine Pearce
UT Austin

Interpolative Decomposition (ID)

- Given a data matrix $X = [x_1, \dots, x_n]^\top \in \mathbb{R}^{n \times d}$
- A target rank $1 \leq r \leq \text{rank}(X)$
- An error tolerance $\tau > 0$
- Aim to construct an ID of X — $X \approx (XX_S^\dagger)X_S$ such that

$$\mathcal{E}(S) = \|X - (XX_S^\dagger)X_S\|_F^2 \leq \tau \|X\|_F^2$$

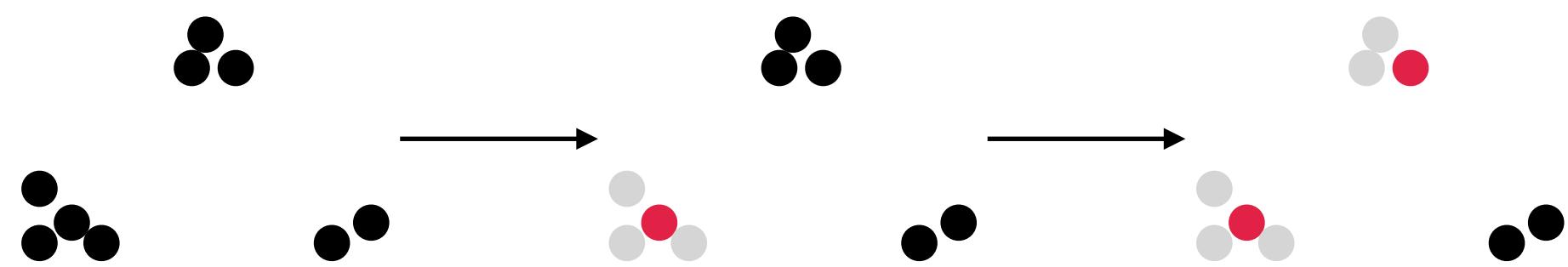


- $S = \{s_1, \dots, s_k\} \subseteq [n]$ contains indices for a **skeleton subset** of size $|S| = k$ (usually $k \ll n$)
- $X_S = [x_{s_1}, \dots, x_{s_k}]^\top \in \mathbb{R}^{k \times d}$ is the row skeleton submatrix corresponding to S
- $W = XX_S^\dagger \in \mathbb{R}^{n \times k}$ is an interpolation matrix for the given skeleton subset S

Adaptiveness & Randomness

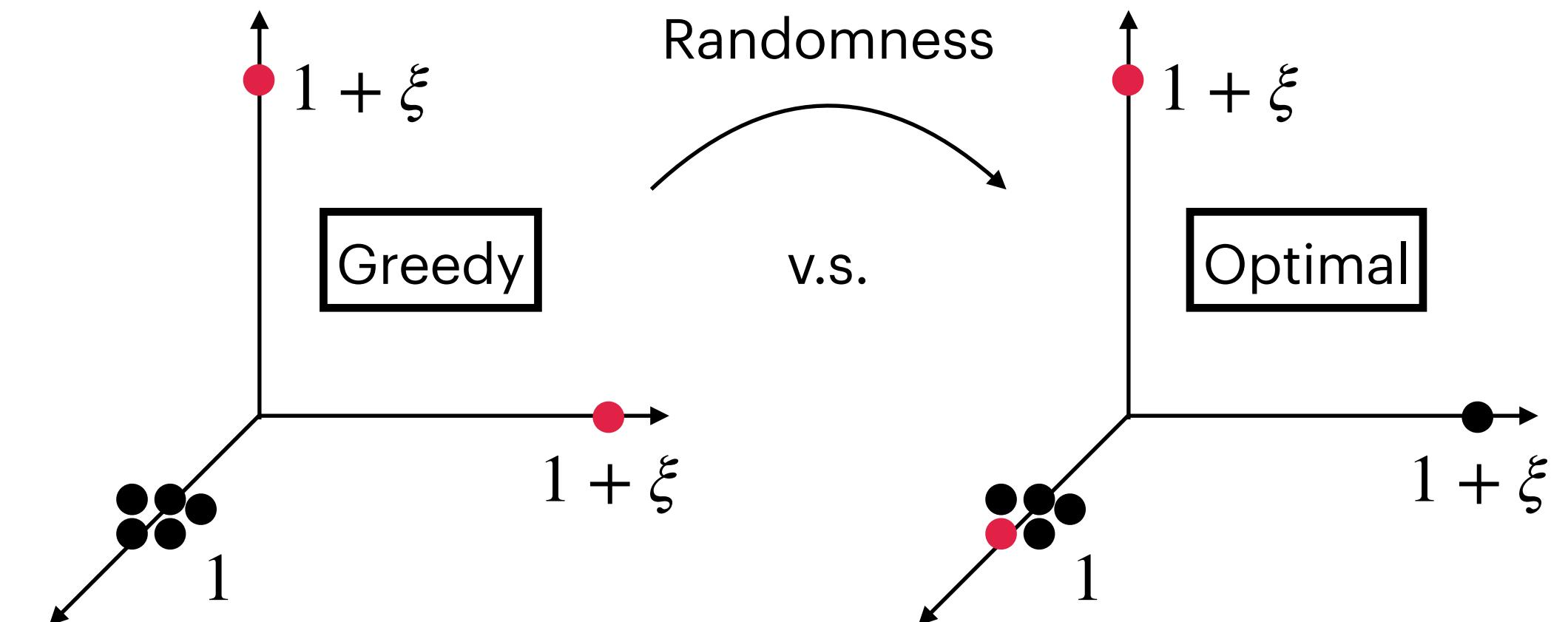
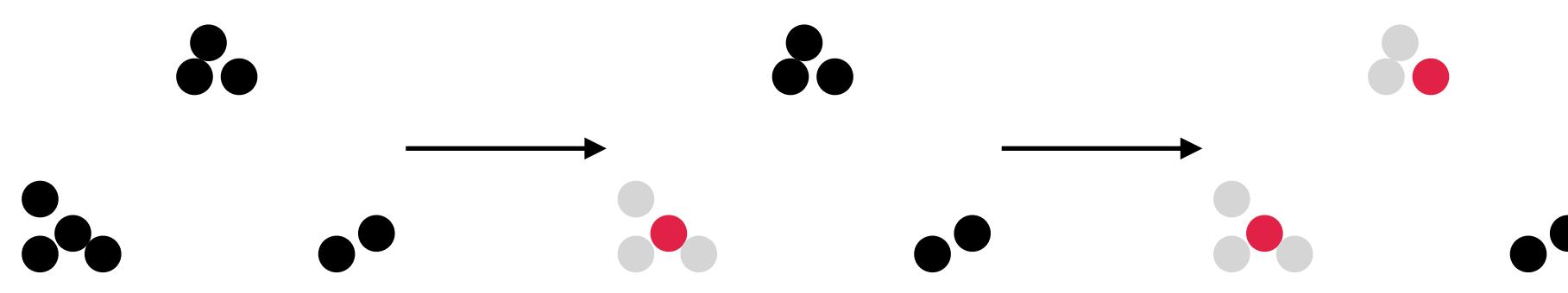
- **Adaptiveness**

- 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)



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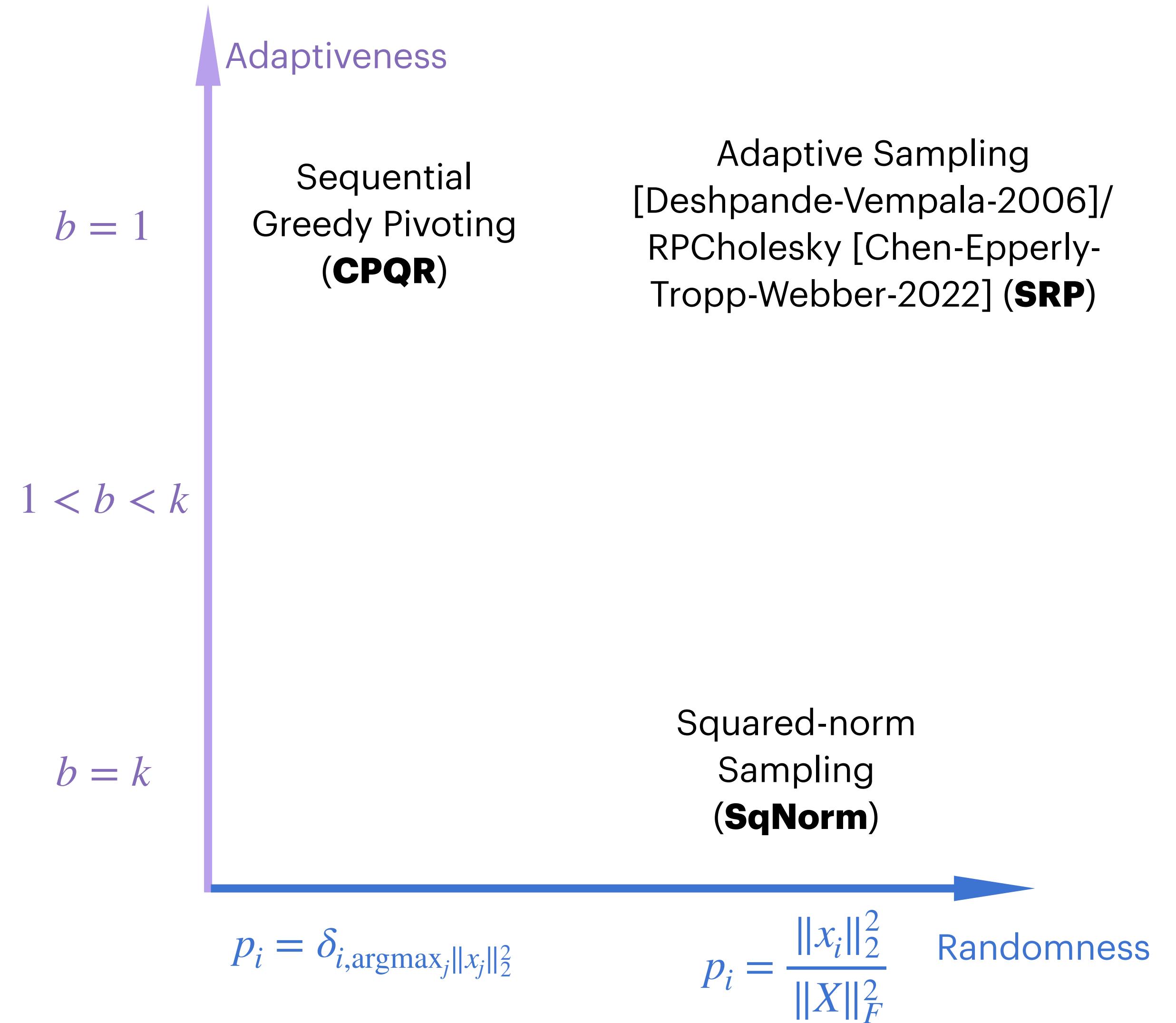
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 - Each new skeleton selection is aware of the previously selected skeleton subset
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 - Common adaptive residual updates:
 - Gram-Schmidt (QR)
 - Gaussian elimination (LU)
- **Randomness** (in contrast 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

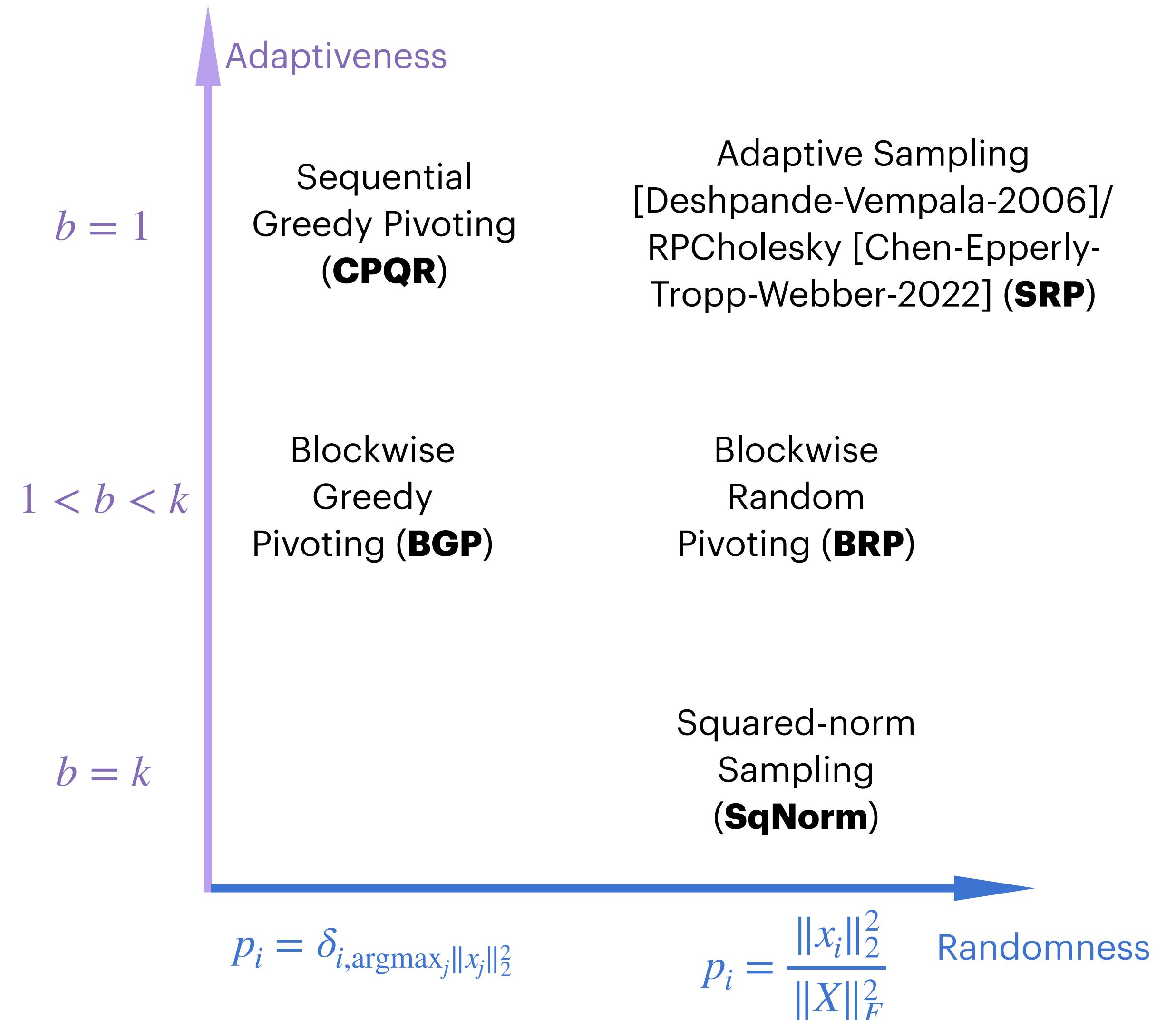
- **Inputs:** $X \in \mathbb{R}^{n \times d}$, $\tau \in (0,1)$
- $X^{(0)} \leftarrow X$, $S^{(0)} \leftarrow \emptyset$, $t \leftarrow 0$
- **while** $\mathcal{E}(S^{(t)}) > \tau \|X\|_F^2$ **do**
 - $t \leftarrow t + 1$
 - Select $|S_t| = b$ skeletons S_t based on $\left(p_i(X^{(t-1)})\right)_{i \in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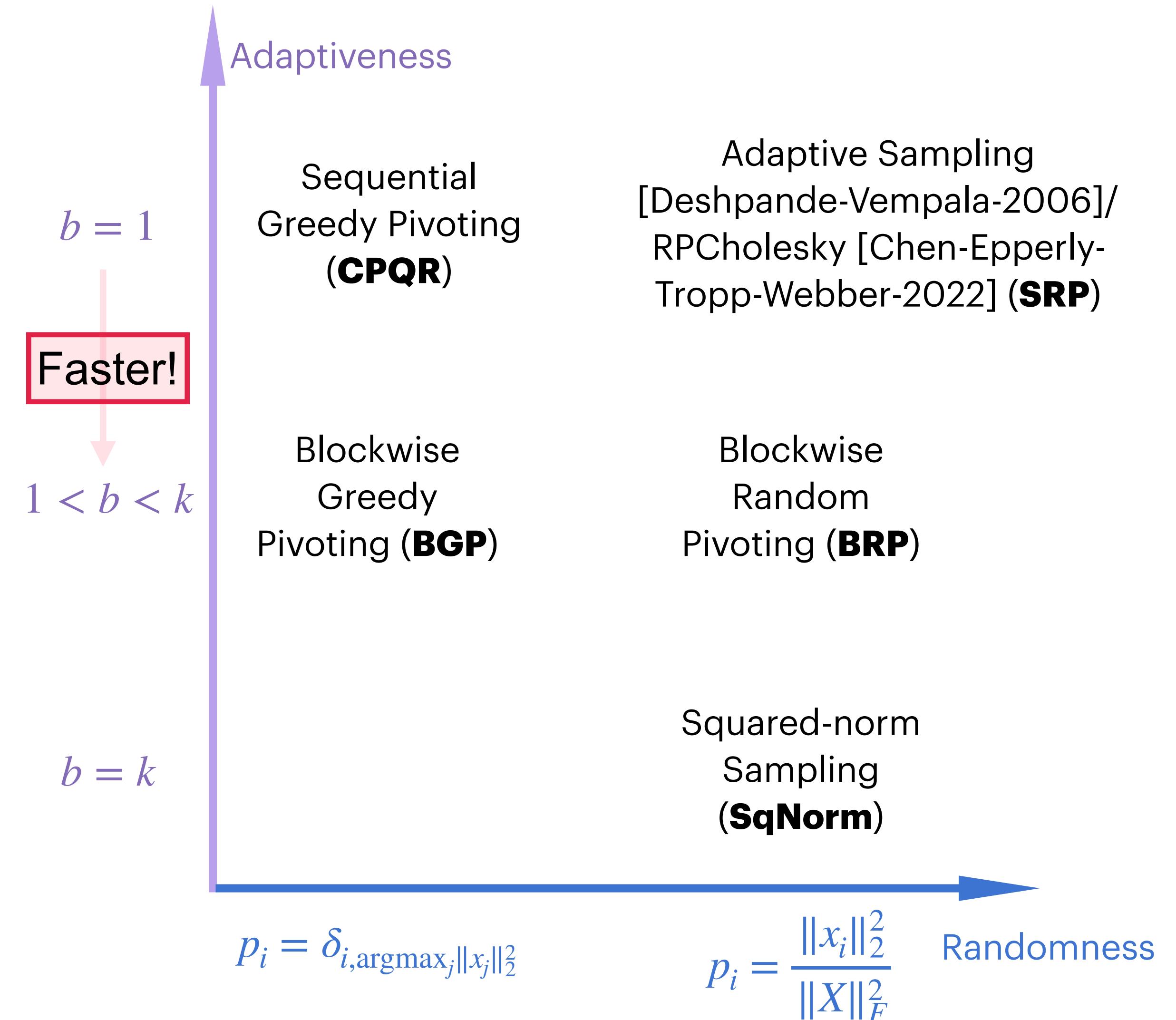
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Skeleton Selection: Other Methods

Sampling methods

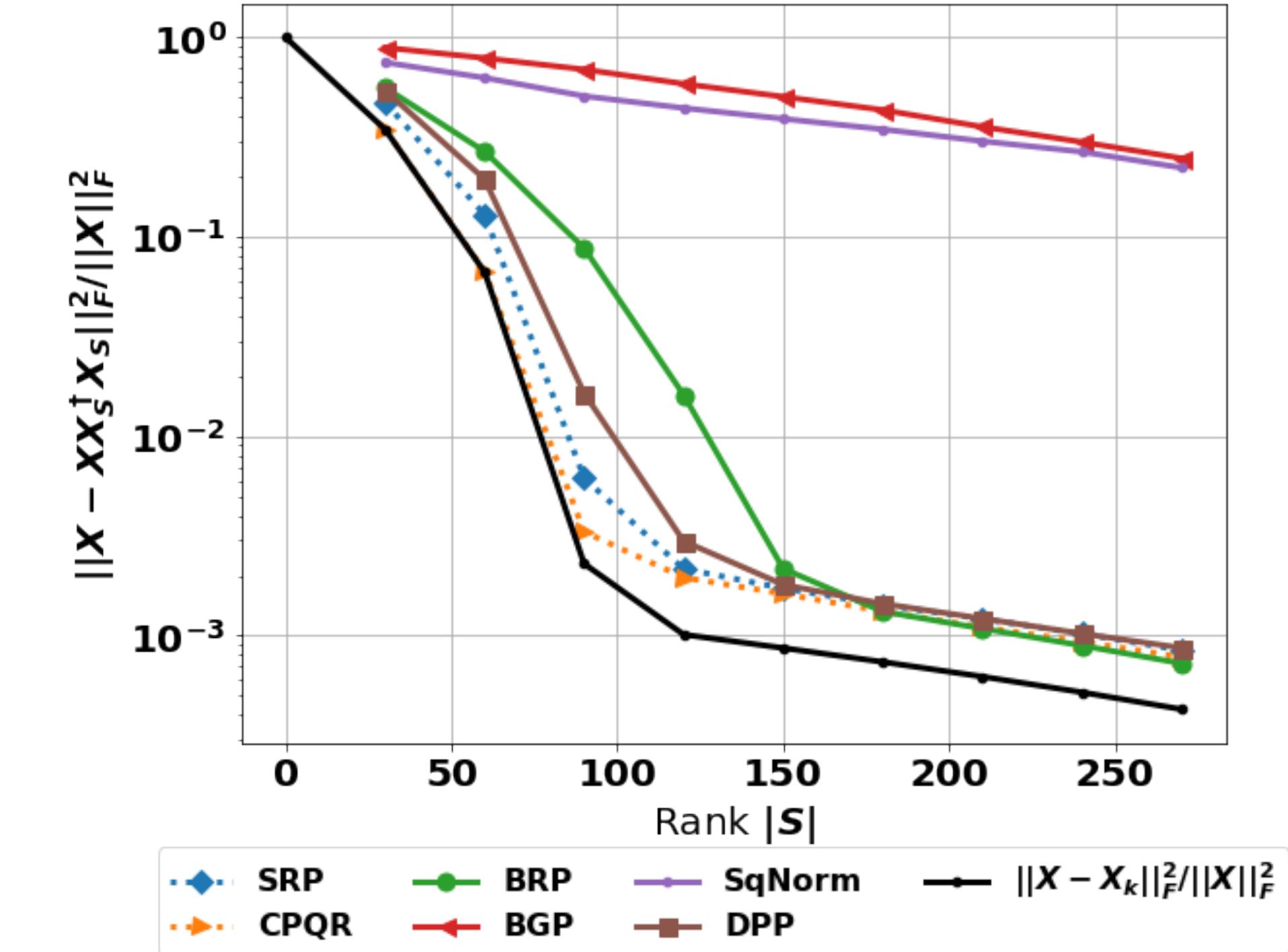
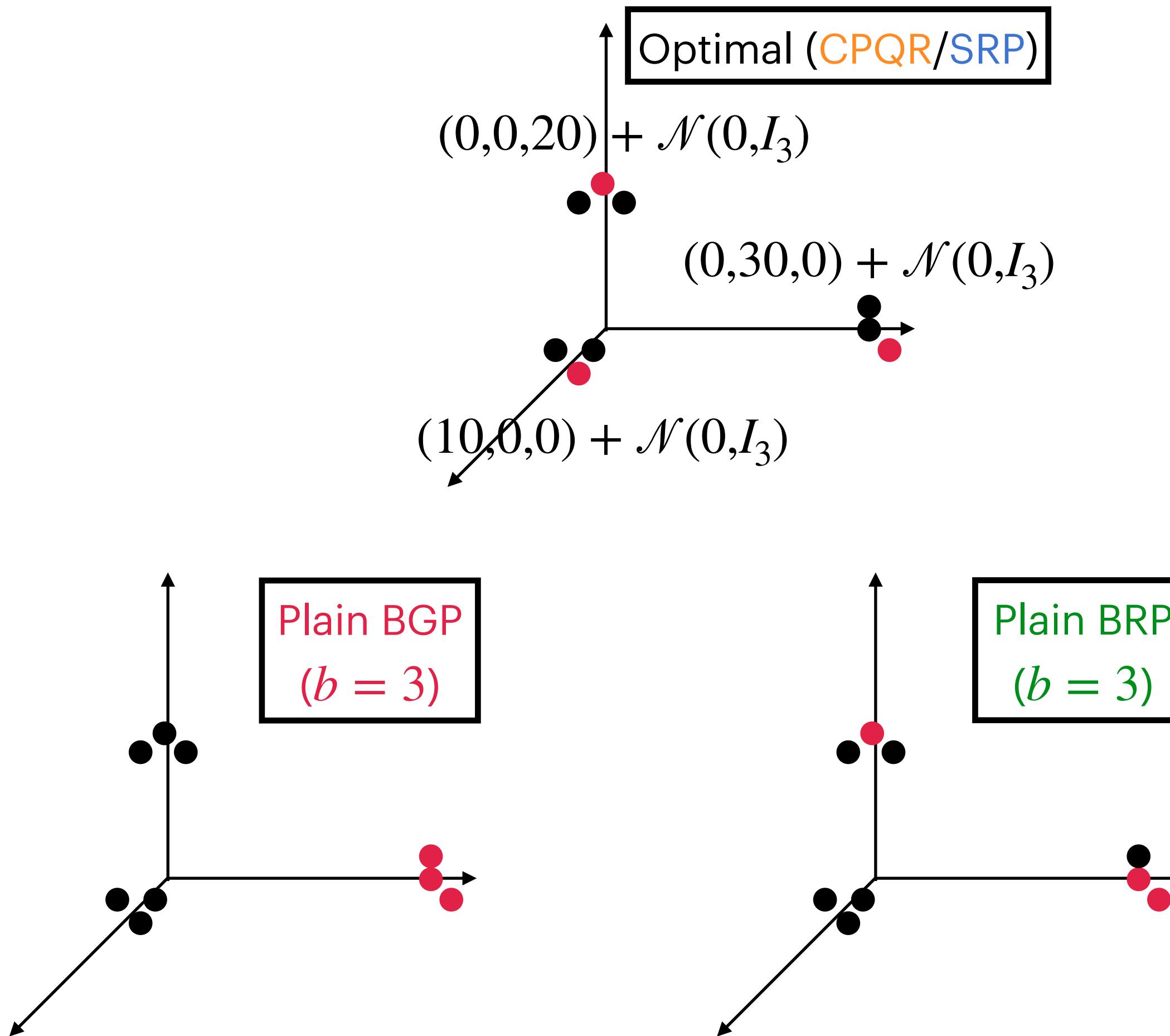
- **DPP/volume sampling** [Deshpande-Rademacher-Vempala-Wang-2006, Belabbas-Wolfe-2009, etc.]
 - Pro: nearly optimal expected skeleton complexity
 - Con: expensive to compute
- **Leverage score sampling** [Mahoney-Drineas-2009, Cohen-Musco-Musco-2017, etc.]
 - Pro: can be estimated efficiently for large-scale 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

Pitfall of Plain Blockwise Greedy/Random Pivoting

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

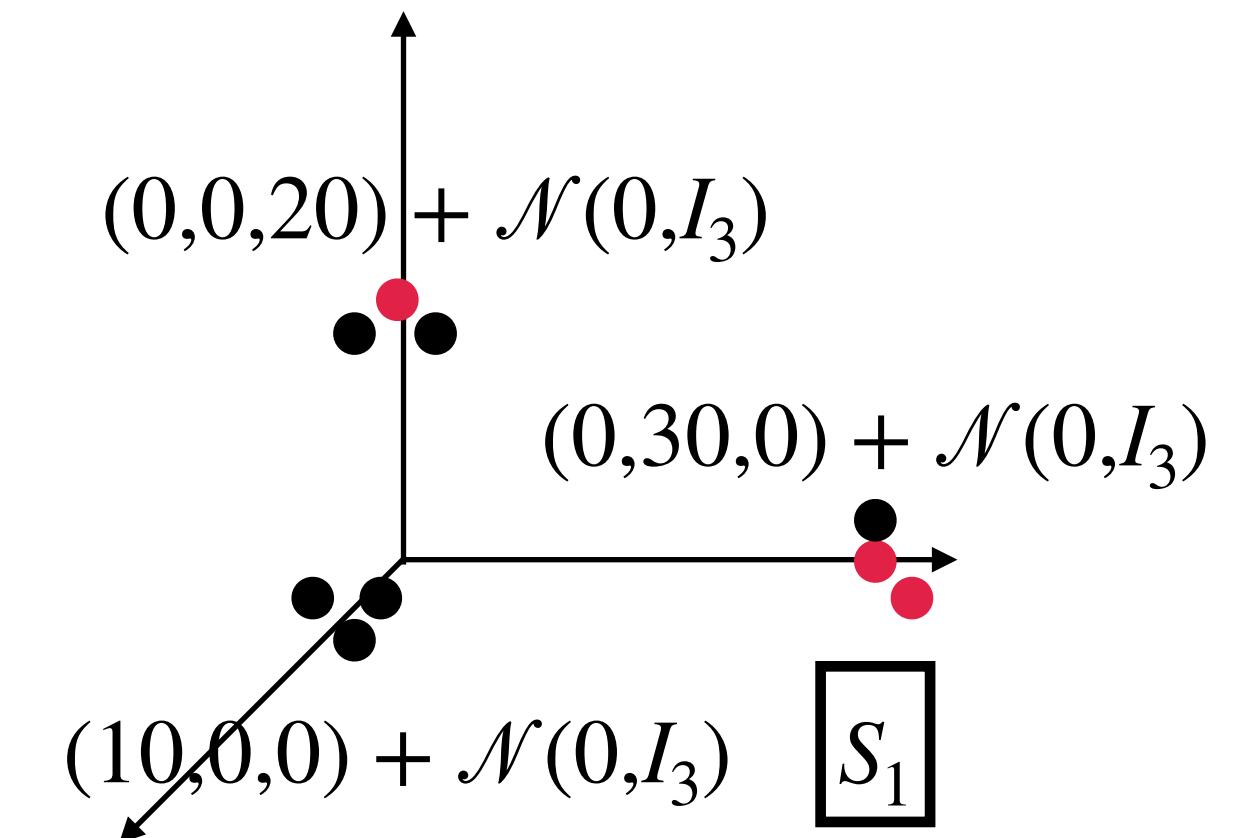


- Sequential pivoting (CPQR & SRP) is nearly optimal
- Plain blockwise pivoting (BRP/BGP, especially BGP) suffers from suboptimal skeleton complexities (up to b times)
- Squared-norm sampling (SqNorm) tends to fail

Robust Blockwise Random Pivoting

Robust Blockwise Random Pivoting (RBRP)

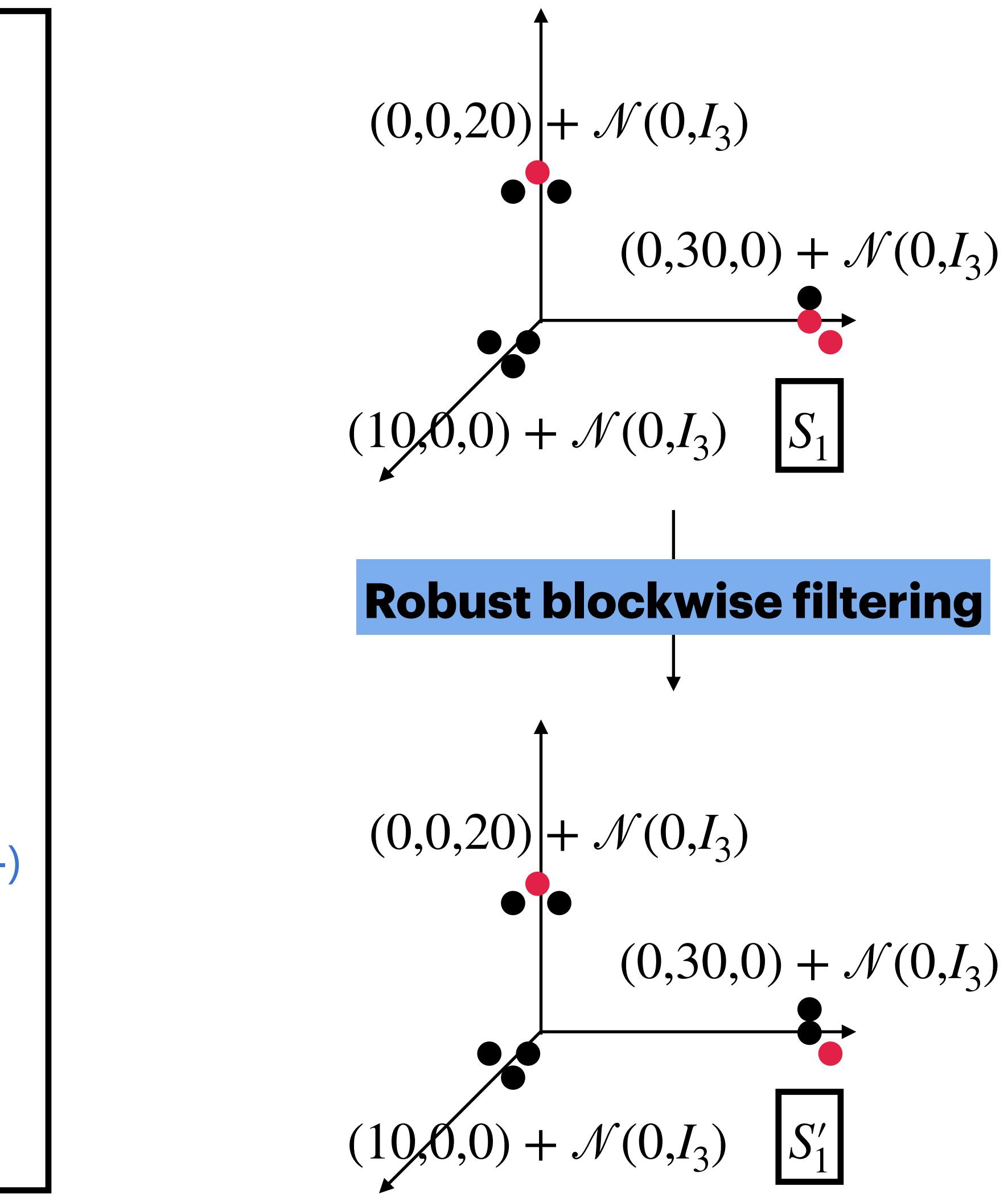
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 - Select $|S_t| = b$ skeletons S_t based on $\left(p_i(X^{(t-1)})\right)_{i \in [n]}$
 - **Robust blockwise filtering (RBF)**
 - $\pi \leftarrow \text{CPQR}\left(X_{S_t}^{(t-1)}\right) \in S_b$ (SRP and CPQR both work)
 - $\min_{S'_t=S_t(\pi(1:b'))} b' \text{ s.t. } \|X_{S_t} - X_{S'_t}\|_F^2 < \tau_b \|X_{S_t}\|_F^2$ (e.g., $\tau_b = \frac{1}{b}$)
 - $S^{(t)} \leftarrow S^{(t-1)} \cup S'_t$ and $X^{(t)} \leftarrow X^{(t-1)} \left(I_d - X_{S'_t}^\dagger X_{S'_t} \right)$
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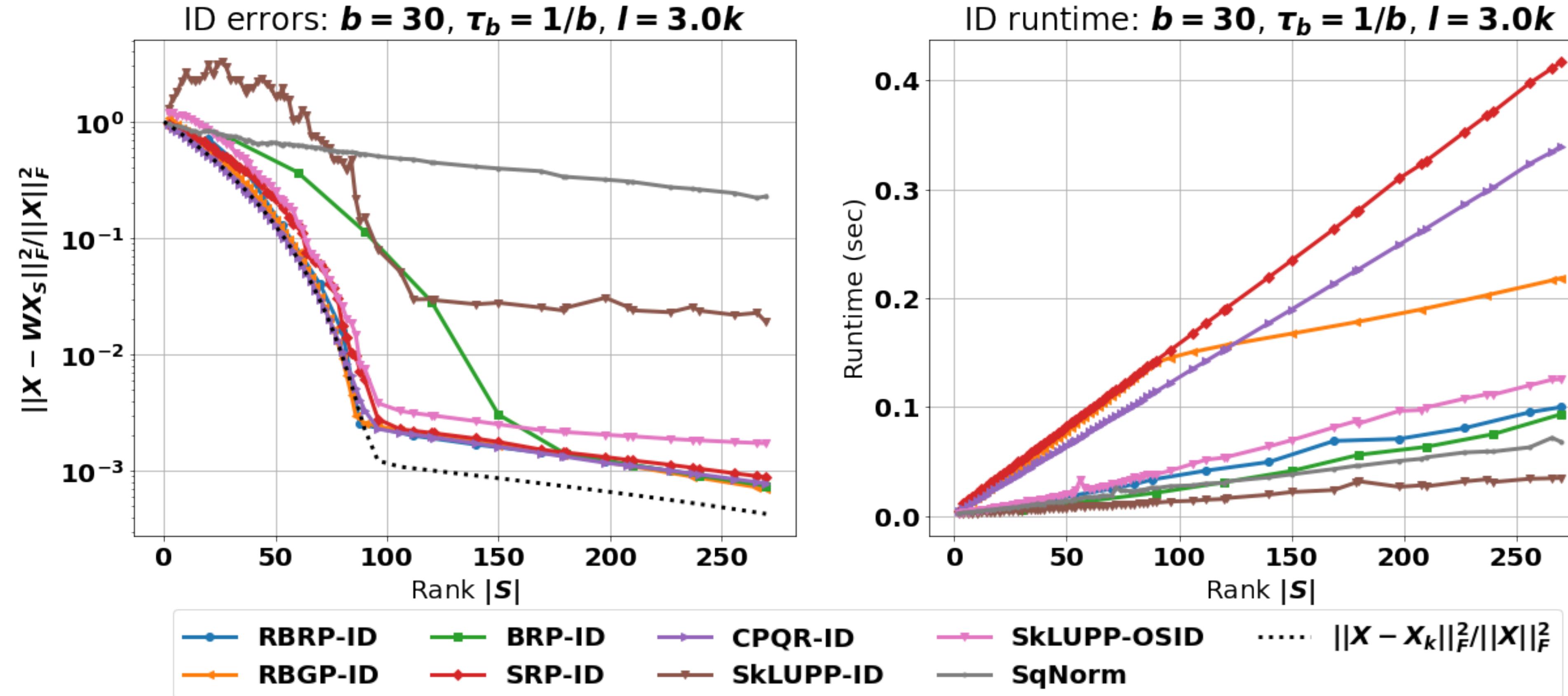
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Robust Blockwise Random Pivoting: Efficiency



- 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
- RBGP can be slowed down more significantly than RBRP by robust blockwise filtering

Summary and Questions

- **Blockwise pivoting** exploits the efficiency of Level-3 BLAS, bringing much **better hardware efficiency** than sequential pivoting
- For adversarial inputs, **plain blockwise pivoting can pick up redundant points**
- **Robust Blockwise Random Pivoting (RBRP)** leverages **robust blockwise filtering (RBF)**, a local greedy filtering step with negligible additional cost, as an effective remedy for such vulnerability
- Alternative to RBF, Epperly-Tropp-Webber-2024 showed that **rejective sampling** can also serve as a remedy for a closely related problem of Cholesky decomposition

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With the shared virtue of **low intrinsic dimension**, are there connections between ID and finetuning?

Beyond low-rank approximation, are “redundant” points necessarily bad?

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 $\text{Var}[y \mid x] \leq \sigma^2$

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 falls in the Kernel Regime

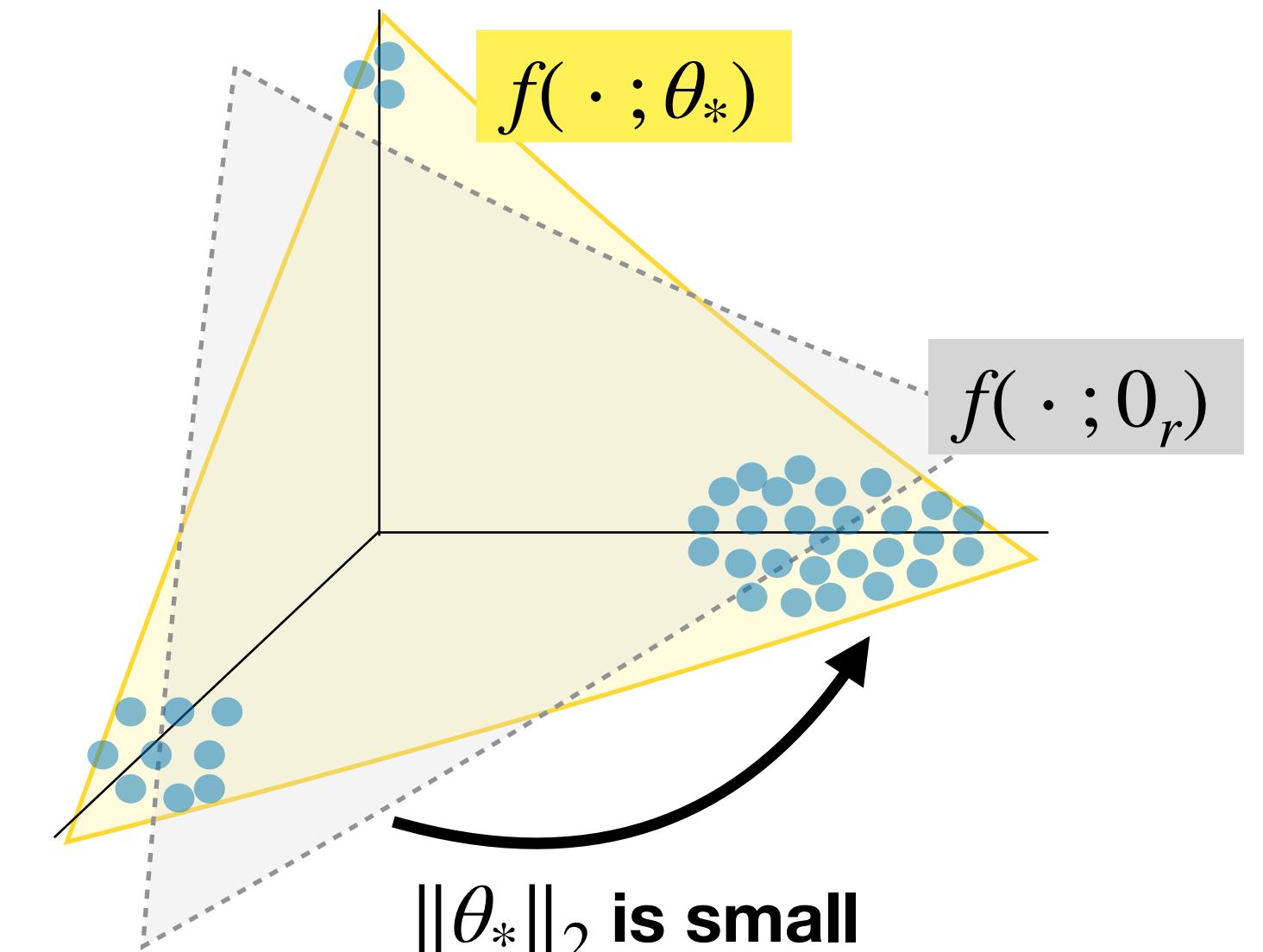
- Finetuning dynamics fall in the **kernel regime**:

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

- 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 the excess risk $\text{ER}(\theta_S) = \|\theta_S - \theta_*\|_{\Sigma}^2$ where $\Sigma = \mathbb{E}_{x \sim P} [\nabla_{\theta} f(x; 0_r) \nabla_{\theta} f(x; 0_r)^{\top}] \in \mathbb{R}^{r \times r}$
- Let $\Sigma_S = G_S^{\top} G_S / n \geq 0$



Finetuning falls in the Kernel Regime

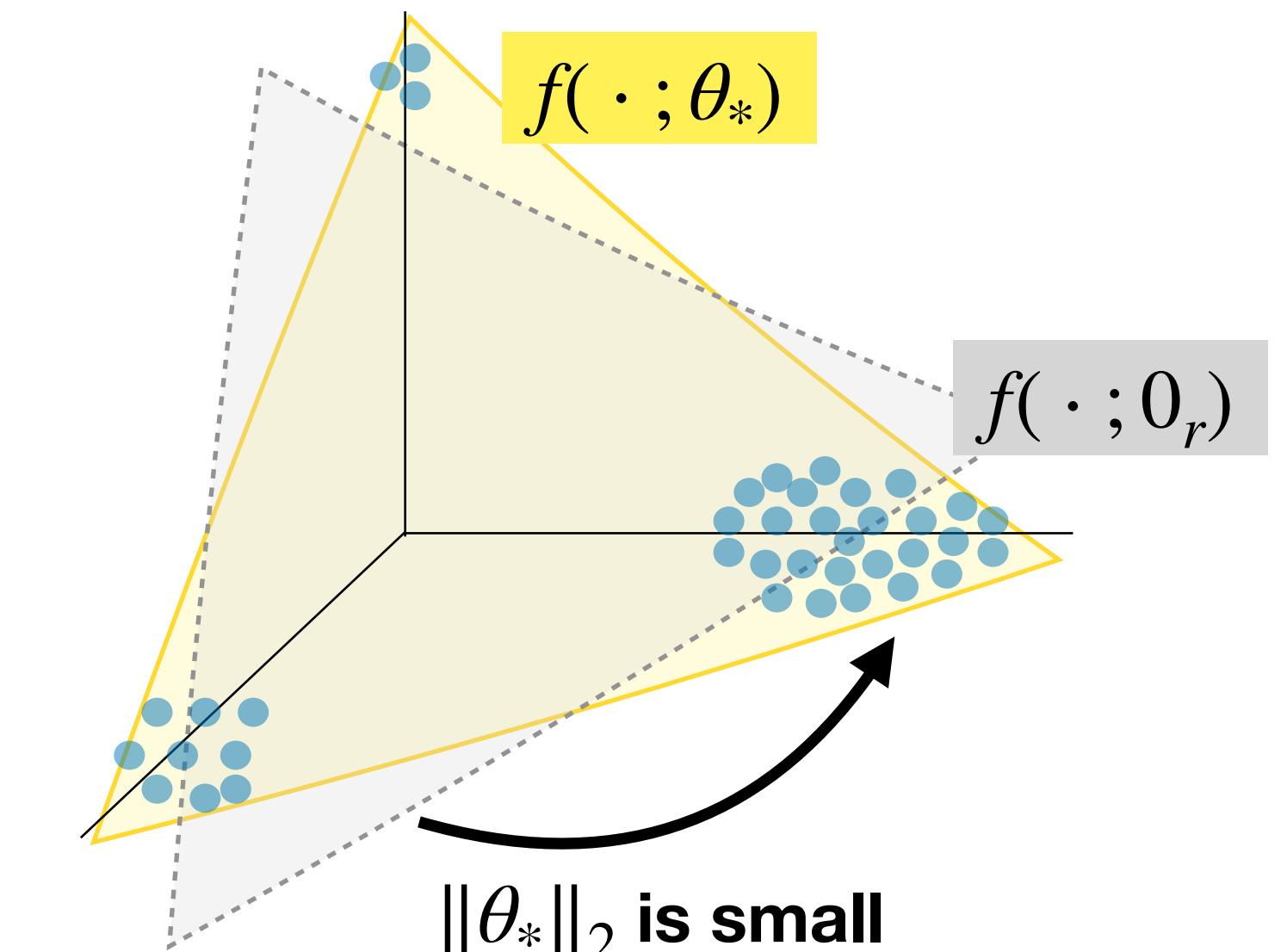
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- Let $\Sigma_S = G_S^{\top} G_S / n \geq 0$



$r = \text{number of finetunable parameters}$
 $(n < r \text{ in overparametrized regime})$

Qs: Are there connections between ID and finetuning?

- In the **noiseless setting** $\sigma = 0$, the generalization error is controlled by the bias:

$$\mathbb{E}[\text{ER}(\theta_S)] \leq \text{tr}(\Sigma - \Sigma G_S^\dagger G_S) \|\theta_*\|_2^2$$

Low-rank approximation error of ID!

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Low-rank approximation error of ID!

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

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

- For a noiseless finetuning problem, accurate ID brings good data selection
- In high-dimensional data selection, **bias** is controlled by the **low-rank approximation error**
- Will see: learning with noise $\sigma > 0$, “redundant” points are critical for **variance reduction!**

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



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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: $\mathbb{E}[\text{ER}(\theta_S)] \leq \frac{\sigma^2}{n} \text{tr}(\Sigma \Sigma_S^{-1})$

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

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

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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 $\bar{r} \ll \min\{N, r\}$

Can the **low intrinsic dimension** of fine-tuning be leveraged when $r > n$ (Σ_S is low-rank)?

With Low Intrinsic Dimension: Variance + Bias

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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,

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

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- **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

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- **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 $\Gamma_{ij} \sim \mathcal{N}(0, 1/m)$

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Theorem (Gradient sketching): For Gaussian embedding $\Gamma \in \mathbb{R}^{r \times m}$ with $m \geq 11\bar{r}$, let $\widetilde{\Sigma} = \Gamma^T \Sigma \Gamma$ and $\widetilde{\Sigma}_S = \Gamma^T \Sigma_S \Gamma$. 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

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

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

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

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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

Control Variance: Sketchy Moment Matching (SkMM)

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- Initialize $s = [s_1, \dots, s_N]$ with $s_i = 1$ if $i \in S$ and $s_i = 0$ otherwise
- Sample a size- n cores $S \subset [N]$ that solves the optimization problem

$$\min_{s \in [0, 1/n]^N} \min_{\gamma = [\gamma_1, \dots, \gamma_m] \in \mathbb{R}^m} \sum_{j=1}^m (v_j^T \widetilde{G}^T \text{diag}(s) \widetilde{G} v_j - \gamma_j \lambda_j)^2$$

s.t. $\|s\|_1 = 1, \quad \gamma_j \geq 1/c_S \quad \forall j \in [m]$

Select $S \subset [N]$ of size $|S| = n$ that reduces the sketched V-optimality:

$$\text{tr}(\widetilde{\Sigma} (\widetilde{\Sigma}_S)^\dagger)$$

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: $O(\text{nnz}(G)m)$
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Sampling takes $O(m^3)$ for spectral

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

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- Assume Σ, Σ_S commute such that imposing m diagonal constraints is sufficient

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 $\alpha > 0$

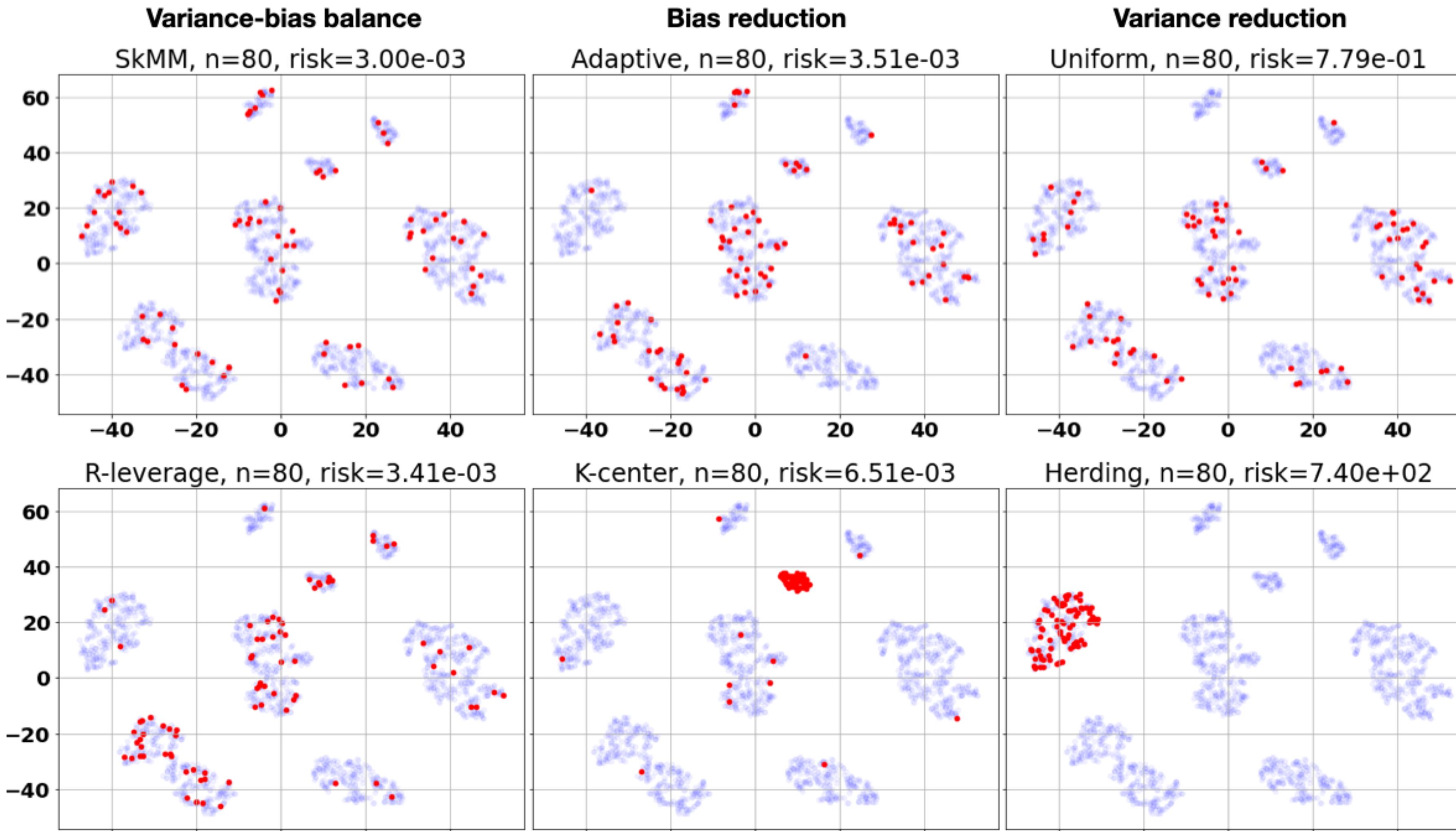
Baselines

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

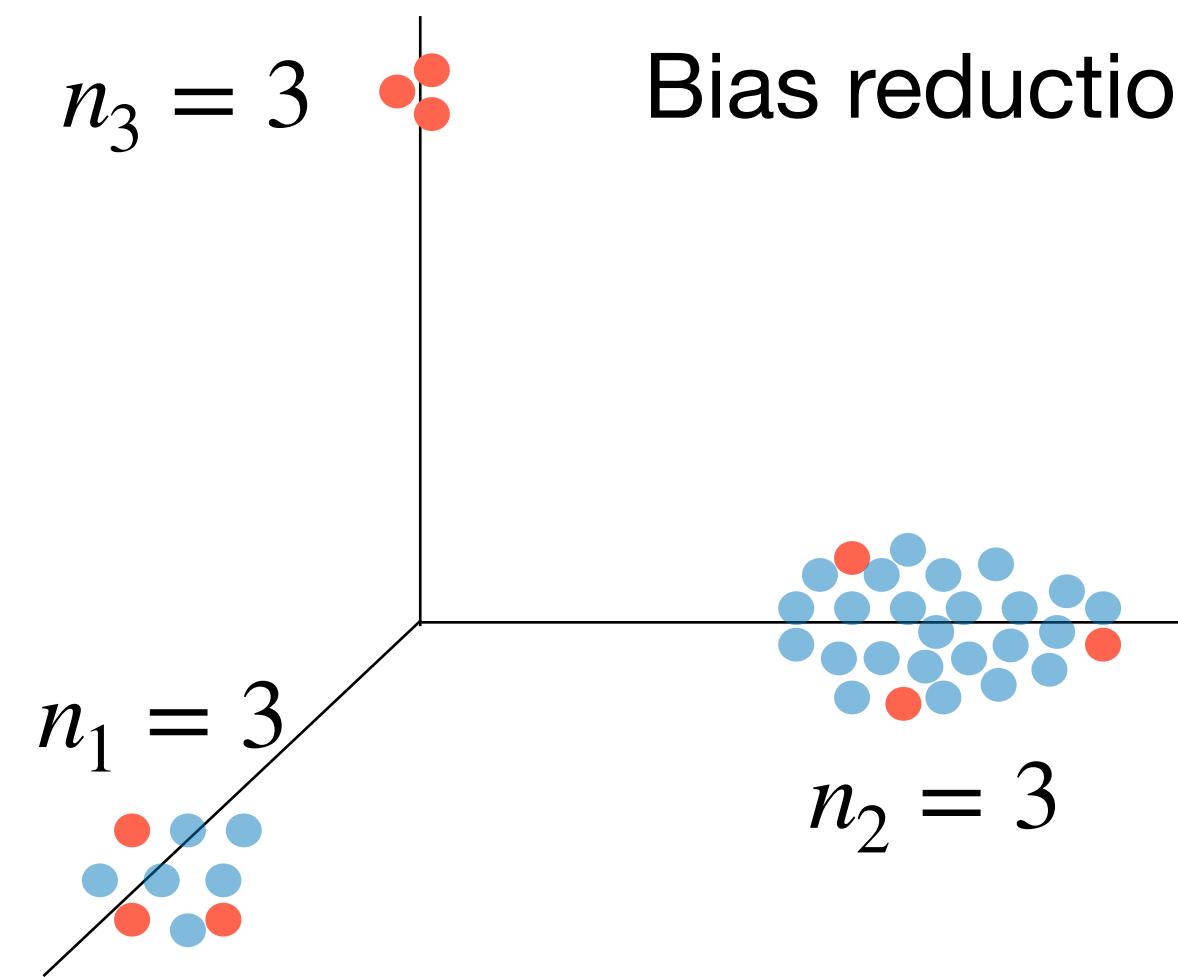
Table 1: Empirical risk $\mathcal{L}_{\mathcal{D}}(\boldsymbol{\theta}_S)$ on the GMM dataset at various n , under the same hyperparameter tuning where ridge regression over the full dataset \mathcal{D} with $N = 2000$ samples achieves $\mathcal{L}_{\mathcal{D}}(\boldsymbol{\theta}_{[N]}) = \mathbf{2.95e-3}$.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

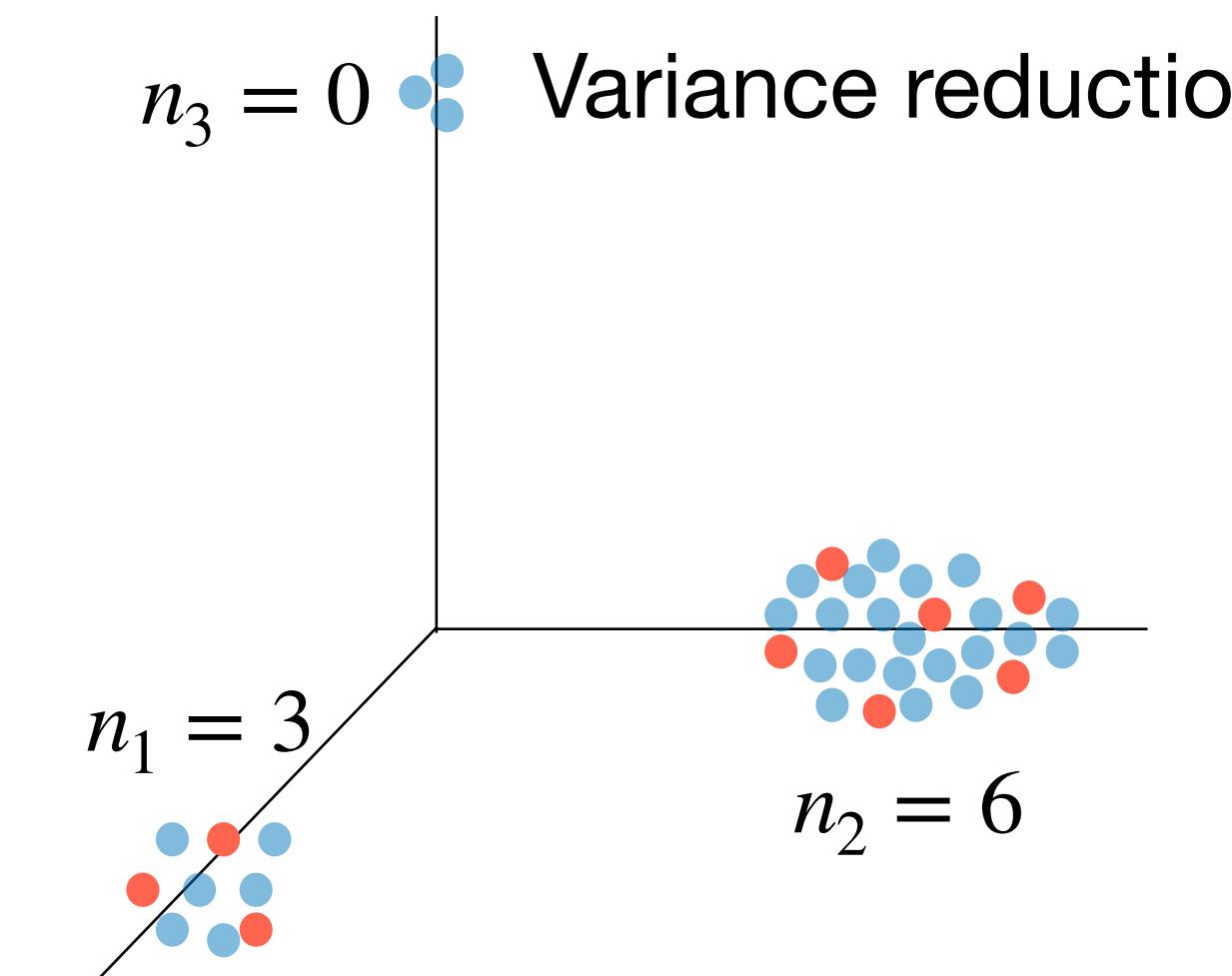
SkMM on Synthetic Data: Regression



SkMM simultaneously controls variance and bias

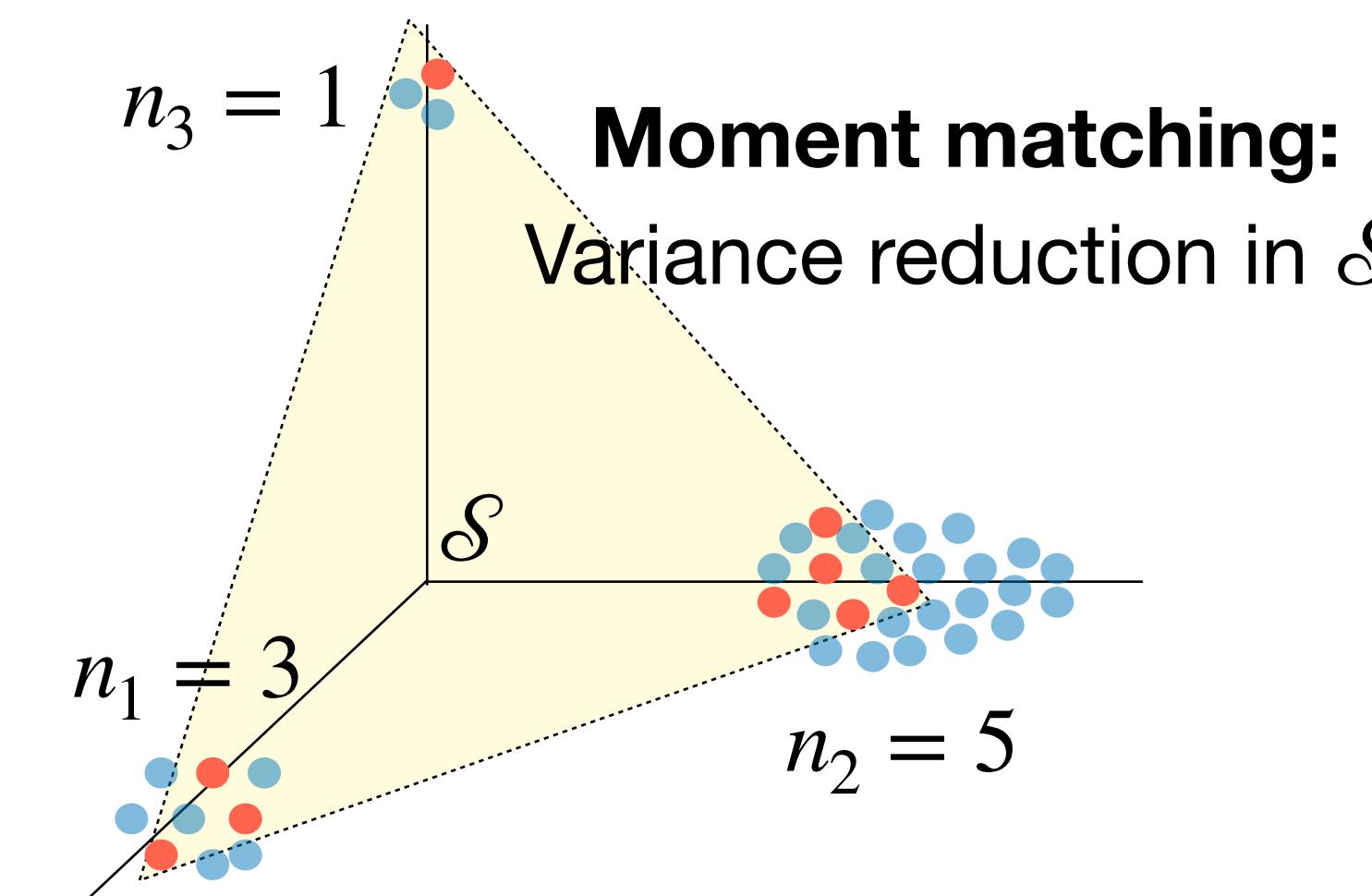
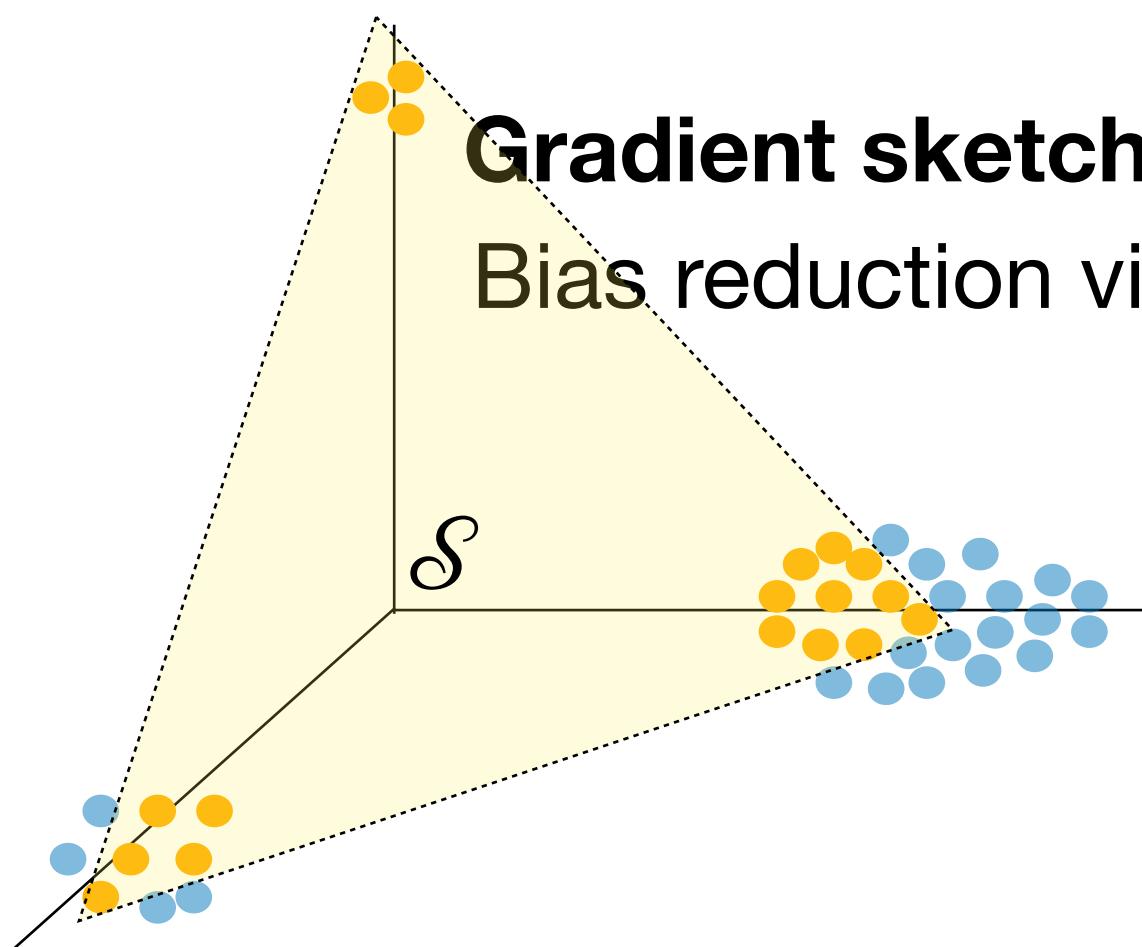
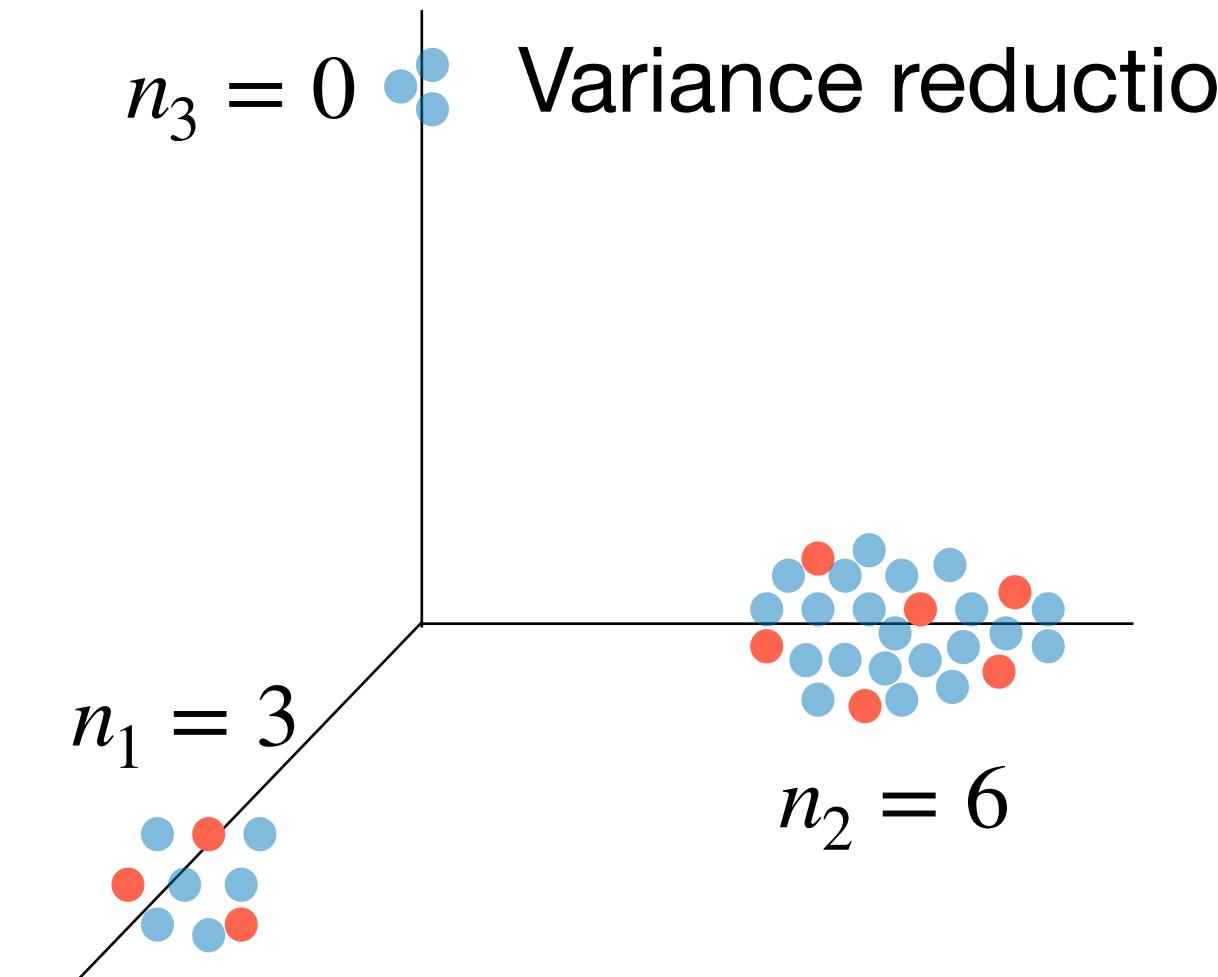
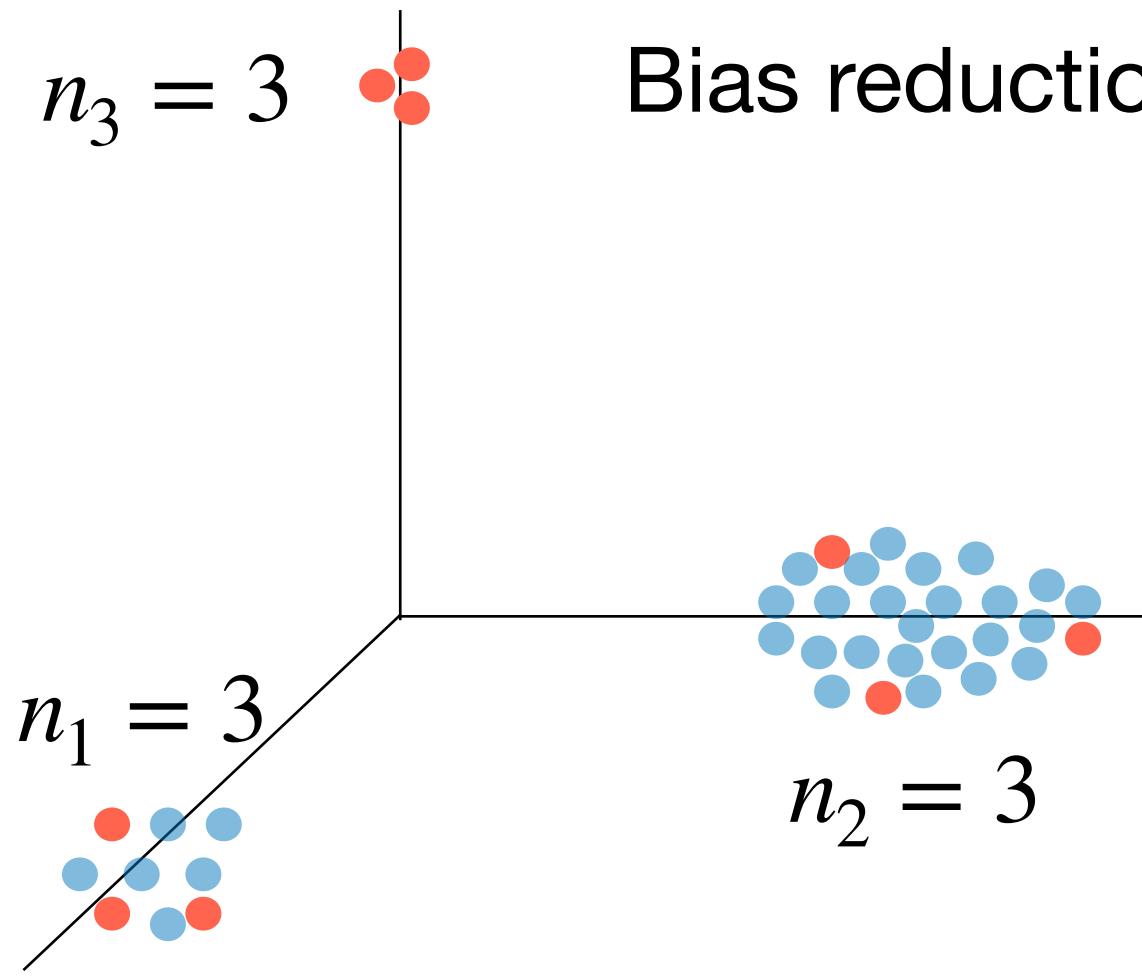


Bias reduction



Variance reduction

SkMM simultaneously controls variance and bias



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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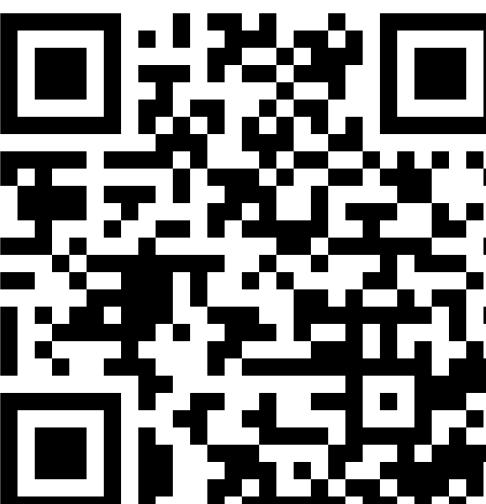
Takeaways

- 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**

Thank You!



Dong, Y., Phan, H., Pan, X., & Lei, Q. Sketchy Moment Matching: Toward Fast and Provable Data Selection for Finetuning. In *The Thirty-eighth Annual Conference on Neural Information Processing Systems*.



Dong, Y., Chen, C., Martinsson, P. G., & Pearce, K. (2023). Robust blockwise random pivoting: Fast and accurate adaptive interpolative decomposition. *arXiv preprint arXiv:2309.16002*.