

Two Count Sketch Kaczmarz Algorithms for Linear Systems

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Abstract. In this paper, combining count sketch, we construct a count sketch two greedy subspace Kaczmarz (CS-2GSK) algorithm. In addition, the block count sketch Kaczmarz (BCSK) algorithm is also given, which selects multiple different rows simultaneously at each iteration. We prove that our methods converge to the unique solution of the linear systems. Finally, the numerical experiments show the high efficiency and robustness of our methods.

Keywords. Kaczmarz, count sketch, image reconstruction

1. Introduction

We only consider over-determined linear systems

$$Ax = b, \quad (1)$$

where $A \in \mathbb{R}^{m \times n}$, $b \in \mathbb{R}^m$, m is much bigger than n , and x is what we want to gain. In particular, we also require linear systems (1) to be consistent. The Kaczmarz method [1] is a common projection method for solving (1), which has been studied for many years, and its iterative scheme is

$$x_{k+1} = x_k + \frac{b_i - a_i^T x_k}{\|a_i\|^2} a_i.$$

In 2018, Bai and Wu [2] proposed the greedy randomized Kaczmarz (GRK) algorithm and Gu and Liu [3] extended the GRK algorithm to the 2GSK algorithm in 2020, and the theoretical guarantee the convergence of the 2GSK algorithm.

Numerous random sketching matrices [4–7] have been discovered by researchers over the years. The maximal weighted residual Kaczmarz [8] algorithm was used by Li and Zhang [9] in 2021, and they provided a CS-MWRK method for solving the systems (1). Numerical experiments demonstrated that the CS-MWRK method can effectively reduce computing time, but the number of iterations has significantly increased.

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In this research, we combine the count sketch and 2GSK method, and we present the count sketch two greedy subspace Kaczmarz algorithm (CS-2GSK), which is motivated by the works of Zhang and Gu. Furthermore, in order to decrease the calculation times and iteration times of the CS-2GSK method, we suggest a block count sketch Kaczmarz algorithm (BCSK), which uses a selecting criterion to choose a block.

The main content of our work is structured as follows. Section 2 begins with several lemmas and the definition of count sketch. Sections 3 and 4 detail the proposed approaches as well as the accompanying convergence. Finally, in Section 5, we show the numerical results and followed the conclusion in Section 6.

2. Preliminaries

The following is about the definition and the property of count sketch, respectively. They have been given in detail in [4, 5, 10].

Definition 2.1 A count sketch matrix $\mathbf{S} \in \mathbb{R}^{d \times m}$ is constructed in terms of ΦD . Here, $D \in \mathbb{R}^{m \times m}$ is a random diagonal matrix. Besides, each diagonal entry of D is chosen to be positive unity or minus one, and Φ is an $d \times m$ arbitrary matrix whose entries are either zeros or ones.

Lemma 2.2 If $\mathbf{S}$ is a count sketch matrix defined in **Definition 2.1** with $(n^2 + n)/(\delta\epsilon^2)$ rows, where δ is a positive constant and ϵ is a normal number less than 1, then we have

$$(1 - \epsilon)\|Ax\|_2 \leq \|\mathbf{S}Ax\|_2 \leq (1 + \epsilon)\|Ax\|_2.$$

3. The Count Sketch two greedy subspace Kaczmarz algorithm

Given a sufficiently large positive integer ℓ and initial point x_0 , then, the specific process of the CS-2GSK algorithm is in **Algorithm 2**.

Algorithm 1 The CS-2GSK algorithm

1. Input: A , b , ℓ x_0 .
2. Output: x_ℓ .
3. Initialize: Create S , compute $\tilde{A} = SA$ and $\tilde{b} = Sb$.
4. **for** $k = 0, 1, \dots, \ell - 1$ **do**
 - (a) Compute $r_k = \tilde{b} - \tilde{A}x_k$
 - (b) Select rows i_k and j_k that satisfy

$$i_k = \arg \max_{1 \leq i \leq d} \left\{ \frac{|\tilde{b}^{(i)} - \tilde{A}^{(i)}x_k|}{\|\tilde{A}^{(i)}\|_2^2} \right\}, \quad j_k = \arg \max_{i \in [m] \setminus i_k} \left\{ \frac{|\tilde{b}^{(i)} - \tilde{A}^{(i)}x_k|}{\|\tilde{A}^{(i)}\|_2^2} \right\}.$$

- (c) Set

$$x_{k+1} = x_k + \frac{r_k^{(i_k)} (\tilde{A}^{(i_k)})^T}{\|\tilde{A}^{(i_k)}\|_2^2} + \frac{r_k^{(j_k)} (\tilde{A}^{(j_k)})^T}{\|\tilde{A}^{(j_k)}\|_2^2}.$$

5. **end for.**

Regarding the convergence of the CS-2GSK method, the **Theorem 3.1** is given.

Theorem 3.1 Let each row of $A \in \mathbb{R}^{m \times n}$ not be orthogonal to each other. Solving linear system (1) by CS-2GSK method can produce a series of sequence $\{x_k\}_{k=0}^\infty$. Then the sequence $\{x_k\}_{k=0}^\infty$ converges to the unique solution of $Ax = b$. Moreover, the solution error obeys

$$\|x_{k+1} - x_\star\|_2^2 \leq \prod_{q=0}^k (1 - \lambda_{\min}(P_q P_q^T)) \|x_0 - x_\star\|_2^2, \quad k = 0, 1, 2, \dots,$$

where $P_k = P_{i_k} - P_{j_k}$ with

$$P_{i_k} = \frac{(\tilde{A}^{(i_k)})^T}{\|\tilde{A}^{(i_k)}\|_2} \cdot \frac{\tilde{A}^{(i_k)}}{\|\tilde{A}^{(i_k)}\|_2} \quad \text{and} \quad P_{j_k} = \frac{(\tilde{A}^{(j_k)})^T}{\|\tilde{A}^{(j_k)}\|_2} \cdot \frac{\tilde{A}^{(j_k)}}{\|\tilde{A}^{(j_k)}\|_2},$$

where $\lambda_{\min}(P_q P_q^T)$ denotes the minimum eigenvalue of $P_q P_q^T$.

Proof. In fact, we only let

$$P_{i_k} = \frac{(\tilde{A}^{(i_k)})^T}{\|\tilde{A}^{(i_k)}\|_2} \cdot \frac{\tilde{A}^{(i_k)}}{\|\tilde{A}^{(i_k)}\|_2} \quad \text{and} \quad P_{j_k} = \frac{(\tilde{A}^{(j_k)})^T}{\|\tilde{A}^{(j_k)}\|_2} \cdot \frac{\tilde{A}^{(j_k)}}{\|\tilde{A}^{(j_k)}\|_2}.$$

The rest of the proof can be completed along the same lines as in [3, Theorem 1], we omit the details. $\square$

4. The block count sketch Kaczmarz algorithm

The basic goal of the block count sketch Kaczmarz (BCSK) algorithm is to gather all of the indices that are closest to the greatest entry as the i_k and j_k in the CS-2GSK

method. Specifically, we select the index set by

$$|r_k^{(j_k)}|^2 \geq \alpha \max_{1 \leq j \leq n} |r_k^{(j)}|^2, \quad j_k \in \{1, 2, \dots, n\}. \quad \alpha \in [0, 1)$$

It's clear that BCSK algorithm is retrieving more rows as α increases.

Below, we give the specific process of the BCSK algorithm in **Algorithm 2**.

Algorithm 2 The BCSK algorithm

1. Input: A, b, α, ℓ, x_0 .
2. Output: x_ℓ .
3. Initialize: Create $\mathbf{S}$, compute $\tilde{A} = \mathbf{S}A$ and $\tilde{b} = \mathbf{S}b$.
4. For $k = 0, 1, \dots, \ell - 1$ **do**
 - (a) Compute $r_k = \tilde{b} - \tilde{A}x_k$
 - (b) Compute target block $\tau_k = \{j_k \mid |r_k^{(j_k)}|^2 \geq \alpha \max_{1 \leq j \leq n} |r_k^{(j)}|^2\}$
 - (c) Set

$$x_{k+1} = x_k + \tilde{A}_{\tau_k}^\dagger r_k.$$

5. **end for**.

Remark 4.1 In practice, the CGLS [11–13] can be used to calculate the approximation of $\tilde{A}_{\tau_k}^\dagger$ in the fourth step of **Algorithm 2**.

Next, we will show the convergence of the BCSK method by **Theorem 4.2**.

Theorem 4.2 Let the condition in **Theorem 3.1** be satisfied. Solving linear system (1) by BCSK method can produce a series of sequence $\{x_k\}_{k=0}^\infty$. Then the sequence $\{x_k\}_{k=0}^\infty$ will converge to the unique solution of (1). Moreover, the solution error obeys

$$\|x_{k+1} - x_\star\|_2^2 \leq \prod_{q=0}^k \left(1 - \frac{\alpha |\tau_q|}{m - |\tau_{q-1}|} \cdot \frac{(1 - \varepsilon)^2 \lambda_{\min}(A^T A)}{(1 + \varepsilon)^2 \sigma_{\max}^2(A)}\right) \|x_0 - x_\star\|_2^2. \quad (2)$$

where $|\tau_q|$ represents the cardinality of τ_q and $\alpha \in (0, 1]$.

Proof. From **Algorithm 2**, we have

$$\begin{aligned} x_{k+1} - x_\star &= x_k - x_\star + \tilde{A}_{\tau_k}^\dagger (\tilde{b}_{\tau_k} - \tilde{A}_{\tau_k} x_k) \\ &= x_k - x_\star - \tilde{A}_{\tau_k}^\dagger \tilde{A}_{\tau_k} (x_k - x_\star) \\ &= (I - \tilde{A}_{\tau_k}^\dagger \tilde{A}_{\tau_k}) (x_k - x_\star). \end{aligned}$$

Since $\tilde{A}_{\tau_k}^\dagger \tilde{A}_{\tau_k}$ is an orthogonal projector, we get

$$\begin{aligned} \|x_{k+1} - x_\star\|_2^2 &= \|(I - \tilde{A}_{\tau_k}^\dagger \tilde{A}_{\tau_k})(x_k - x_\star)\|_2^2 \\ &= \|x_k - x_\star\|_2^2 - \|\tilde{A}_{\tau_k}^\dagger \tilde{A}_{\tau_k} (x_k - x_\star)\|_2^2 \end{aligned} \quad (3)$$

As explained in [2], we have

$$\|A(x_k - x_*)\|_2^2 \geq \sigma_r^2(A) \|x_k - x_*\|_2^2,$$

which leads to

$$\begin{aligned} \|\tilde{A}_{\tau_k}^\dagger \tilde{A}_{\tau_k}(x_k - x_*)\|_2^2 &\geq \frac{1}{\sigma_{\max}^2(\tilde{A}_{\tau_k})} \|\tilde{A}_{\tau_k}(x_k - x_*)\|_2^2 \\ &\geq \frac{1}{(1+\varepsilon)^2 \sigma_{\max}^2(A)} \|\tilde{A}_{\tau_k}(x_k - x_*)\|_2^2 \\ &= \frac{1}{(1+\varepsilon)^2 \sigma_{\max}^2(A)} \sum_{i_k \in \tau_k} |\tilde{A}^{(i_k)}(x_k - x_*)|^2 \\ &\geq \frac{1}{(1+\varepsilon)^2 \sigma_{\max}^2(A)} \sum_{i_k \in \tau_k} \alpha \max_{1 \leq i \leq m} |\tilde{A}^{(i)}(x_k - x_*)|^2 \\ &\geq \frac{|\tau_k|}{(1+\varepsilon)^2 \sigma_{\max}^2(A)} \alpha \max_{1 \leq i \leq m} |\tilde{A}^{(i)}(x_k - x_*)|^2. \end{aligned} \quad (4)$$

From the iteration of the BCSK, we have

$$\begin{aligned} (\tilde{b} - \tilde{A}x_{k+1})_{\tau_k} &= \tilde{b}_{\tau_k} - \tilde{A}_{\tau_k}x_k - \tilde{A}_{\tau_k}\tilde{A}_{\tau_k}^\dagger(\tilde{b}_{\tau_k} - \tilde{A}_{\tau_k}x_k) \\ &= \tilde{b}_{\tau_k} - \tilde{A}_{\tau_k}x_k - \tilde{A}_{\tau_k}\tilde{A}_{\tau_k}^\dagger\tilde{b}_{\tau_k} + \tilde{A}_{\tau_k}\tilde{A}_{\tau_k}^\dagger\tilde{A}_{\tau_k}x_k \\ &= \tilde{b}_{\tau_k} - \tilde{A}_{\tau_k}x_k - \tilde{A}_{\tau_k}\tilde{A}_{\tau_k}^\dagger\tilde{b}_{\tau_k} + \tilde{A}_{\tau_k}x_k \\ &= \tilde{b}_{\tau_k} - \tilde{A}_{\tau_k}\tilde{A}_{\tau_k}^\dagger\tilde{b}_{\tau_k} \\ &= \tilde{A}_{\tau_k}x_* - \tilde{A}_{\tau_k}\tilde{A}_{\tau_k}^\dagger\tilde{A}_{\tau_k}x_* \\ &= \tilde{A}_{\tau_k}x_* - \tilde{A}_{\tau_k}x_* \\ &= 0, \end{aligned}$$

which implies the inequality

$$\|\tilde{b} - \tilde{A}x_k\|_2^2 = \sum_{i \in [m] \setminus \tau_{k-1}} |\tilde{b}^{(i)} - \tilde{A}^{(i)}x_k|^2 \leq (m - |\tau_{k-1}|) \max_{1 \leq i \leq m} |\tilde{b}^{(i)} - \tilde{A}^{(i)}x_k|^2.$$

Hence, for any $\alpha \in (0, 1]$, combining **Lemma 2.2**, we have

$$\max_{1 \leq i \leq m} |\tilde{b}^{(i)} - \tilde{A}^{(i)}x_k|^2 \geq \frac{\alpha}{m - |\tau_{k-1}|} \|\tilde{A}(x_k - x_*)\|_2^2 \geq \frac{(1-\varepsilon)^2 \alpha}{m - |\tau_{k-1}|} \|A(x_k - x_*)\|_2^2.$$

As a result, we see that

$$\begin{aligned} \|\tilde{A}_{\tau_k}^\dagger \tilde{A}_{\tau_k}(x_k - x_*)\|_2^2 &\geq \frac{|\tau_k|}{m - |\tau_{k-1}|} \cdot \frac{(1-\varepsilon)^2 \alpha}{(1+\varepsilon)^2 \sigma_{\max}^2(A)} \|A(x_k - x_*)\|_2^2 \\ &\geq \frac{\alpha |\tau_k|}{m - |\tau_{k-1}|} \cdot \frac{(1-\varepsilon)^2 \lambda_{\min}(A^T A)}{(1+\varepsilon)^2 \sigma_{\max}^2(A)} \|x_k - x_*\|_2^2. \end{aligned} \quad (5)$$

Thus, substituting Eqs.(5) and (4) into Eq.(3), we obtain

$$\|x_{k+1} - x_\star\|_2^2 \leq \left(1 - \frac{\alpha|\tau_k|}{m - |\tau_{k-1}|} \cdot \frac{(1 - \varepsilon)^2 \lambda_{\min}(A^T A)}{(1 + \varepsilon)^2 \sigma_{\max}^2(A)}\right) \|x_k - x_\star\|_2^2.$$

Then,

$$\|x_{k+1} - x_\star\|_2^2 \leq \prod_{q=0}^k \left(1 - \frac{\alpha|\tau_q|}{m - |\tau_{q-1}|} \cdot \frac{(1 - \varepsilon)^2 \lambda_{\min}(A^T A)}{(1 + \varepsilon)^2 \sigma_{\max}^2(A)}\right) \|x_0 - x_\star\|_2^2.$$

Thus, we complete the proof. $\square$

5. Numerical experiments

In this part, we will use BCSK and CS-2GSK methods to solve (1) and compare with the 2GSK [3] and CS-MWRK [9] methods in terms of the number of iteration steps (denoted as IT) and the computing time (denoted as CPU). All experiments were on a personal computer with 2.00 GHz central processing unit (Intel(R) Core(TM) i5-1038NG7 CPU), 16.00GB memory, and Windows 10 operating system. In addition, the version of MATLAB we use is R2018b.

Example 5.1 We consider the case of stochastic systems in this experiment. The matrix $A \in \mathbb{R}^{m \times n}$ and the exact solution $x_\star$ are gained randomly by using the MATLAB function *randn*(m, n). All the experiments start from zero vector and we repeat the experiment fifty times and average it. Otherwise, the experiment stops if

$$RES = \frac{\|x_k - x_\star\|_2^2}{\|x_\star\|_2^2} \leq 10^{-6}$$

or IT exceeds 200,000.

The numerical results are listed in **Table 1** illustrates that the CS-2GSK method has its pros and cons. More particular, CS-2GSK requires fewer IT than CS-MWRK and significantly less CPU time than 2GSK when $m \gg n$. Additionally, CS-2GSK needs more IT than the 2GSK method. This is due to the fact that the CS-2GSK approach converges a little more slowly due to its bigger convergence factor. In addition, we observe that IT of CS-2GSK approach lowers as d increases, suggesting that the CS-2GSK method converges more quickly. The computation of $i_k = \arg \max_{1 \leq i \leq d} \left\{ \frac{|\tilde{b}^{(i)} - \tilde{A}^{(i)} x_k|}{\|\tilde{A}^{(i)}\|_2} \right\}$

and $j_k = \arg \max_{i \in [m] \setminus i_k} \left\{ \frac{|\tilde{b}^{(i)} - \tilde{A}^{(i)} x_k|}{\|\tilde{A}^{(i)}\|_2} \right\}$ in each iteration will take longer as d rises. As a result, the CS-2GSK method's overall processing time grows. Then, we can see that, for all test values of d , BCSK method uses substantially less CPU and has smaller IT than the other three algorithms.

Example 5.2 In this example, we use the function *paralleltomo*(N, θ, p) in the MATLAB package *AIR TOOLS* [14]. We set $N = 30$, $\theta = 0 : 2 : 178^\circ$ and $p = 120$, then A is a 10800×1600 matrix. What's more, we run these methods a thousand times and then observe how clear the image recovered.

Table 1. Numerical results for the 2GSK, BCSK (with $\alpha = 0.16$), CS-MWRK and CS-2GSK methods with random matrix.

$m \times n$	d	IT				CPU			
		2GSK	CS-MWRK	BCSK	CS-2GSK	2GSK	CS-MWRK	BCSK	CS-2GSK
5000×50	10n	34.0000	85.2000	1.1200	56.0000	0.0069	0.0109	0.0013	0.0100
	20n	34.0000	67.4800	1.0400	49.4400	0.0075	0.0100	0.0044	0.0091
	30n	32.0000	61.5400	1.0600	48.9800	0.0084	0.0128	0.0034	0.0106
	40n	35.0000	58.5000	1.0000	47.6000	0.0075	0.0138	0.0028	0.0103
	50n	31.0000	56.5600	1.0200	47.2800	0.0075	0.0125	0.0044	0.0134
5000×100	10n	60.0000	172.2600	1.3800	119.7000	0.0219	0.0159	0.0075	0.0150
	20n	64.0000	137.6000	1.1200	103.1400	0.0222	0.0175	0.0166	0.0166
	30n	60.0000	126.6200	1.1000	99.0800	0.0200	0.0297	0.0097	0.0306
	40n	60.0000	122.0600	1.0800	97.2200	0.0213	0.0437	0.0091	0.0362
	50n	61.0000	118.6600	1.0800	95.5800	0.0213	0.0750	0.0088	0.0316
5000×150	10n	92.0000	260.8600	2.1200	189.7400	0.0547	0.0431	0.0128	0.0316
	20n	95.0000	213.4600	1.4400	164.1200	0.0603	0.0587	0.0219	0.0547
	30n	97.0000	198.6200	1.4000	157.5800	0.0597	0.1744	0.0184	0.0881
	40n	90.0000	192.2600	1.3200	154.0800	0.0512	0.2334	0.0244	0.1441
	50n	95.0000	191.0000	1.1400	151.1000	0.0591	0.2972	0.0262	0.1844
5000×200	10n	131.0000	354.3600	4.1600	267.0400	0.1244	0.0847	0.0338	0.0666
	20n	131.0000	295.5000	2.0800	233.1800	0.1247	0.3050	0.0259	0.1672
	30n	131.0000	281.5600	1.9000	222.7800	0.1259	0.4469	0.0344	0.2822
	40n	135.0000	271.9200	1.6800	212.9000	0.1256	0.5791	0.0384	0.3781
	50n	132.0000	265.7800	1.3000	204.1200	0.1278	0.7116	0.0397	0.5525

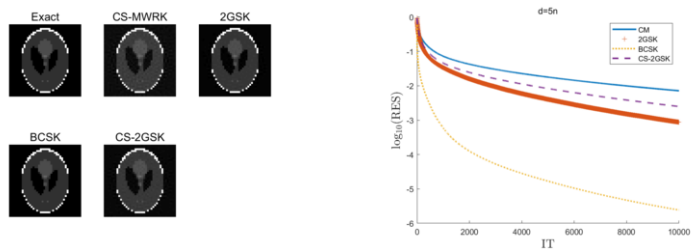


Figure 1. Numerical results for the 2GSK, BCSK (with $\alpha = 0.16$), CS-MWRK and CS-2GSK methods with paralleltomo test problem.

From the left of **Figure 1**, we see again that, under the same IT, BCSK algorithm can achieve better reconstruction results than CS-2GSK, CS-MWRK and 2GSK algorithms with generally smaller working blocks. The right side of **Figure 1** shows that, as IT increases, the RES of the BCSK method decays considerably more fast than those of CS-2GSK, CS-MWRK, and 2GSK algorithms. In fact, we can see that at the thousandth iteration, the RES of BCSK is less than 10^{-6} .

6. Conclusion

In order to solve a sizable consistent linear system (1), we modified the GRK algorithm in this study to the count sketch two greedy subspace Kaczmarz (CS-2GSK) method and the block count sketch Kaczmarz (BCSK) method. We offer theoretical assurances that the two algorithms would converge. We demonstrate certain situations in the experiments section, where the BCSK algorithm performs better than the 2GSK al-

gorithm and the CS-MWRK algorithm in terms of IT and CPU times. In light of this, the BCSK algorithm could be a helpful tool when compared with the CS-2GSK algorithm and the CS-MWRK algorithm.

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