

Convergence Rate Analysis of Parallel Block Coordinate Descent Method for Composite Function Minimization

Yutong Dai^{a,1}, Yang Weng^{b,1,*}

^a*Department of Mathematics, Sichuan University*

^b*Department of Mathematics, Sichuan University*

Abstract

In this paper, we propose a synchronous parallel block coordinate descent algorithm for minimizing a composite function, which consists of a smooth convex function plus a non-smooth but separable convex function. Our method can include some existing synchronous parallel algorithms as special cases. For the sake of using our method to solve high dimension problems, we further develop a randomized variant. We have proved that both proposed parallel algorithms are endowed with sub-linear convergence rate for general convex composite function. The experiments on solving the regularized logistic regression with ℓ_1 norm problem show that the proposed parallel algorithms are quite efficient to implement.

Keywords: Optimization, Block Coordinate Descent, Synchronous Parallel Computation, Logistic Regression with ℓ_1 regularization

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1. Introduction

Consider the problem of minimizing a non-smooth convex composite function $f(x)$ of the form,

*Corresponding author

Email address: wengyang@scu.edu.cn (Yang Weng)

$$\begin{aligned} \min f(x) = g(x) + h(x) &\triangleq g(x_1, \dots, x_K) + \lambda \sum_{i=1}^K h_i(x_i) \\ \text{subject to } x_i &\in X_i \end{aligned} \tag{1.1}$$

In (1.1), (x_1, \dots, x_K) denotes a pre-defined partition of the optimization variable x , where $x_i \in X_i \subset \mathbb{R}^{n_i}$ and $\sum_{i=1}^K n_i = n$; $X = X_1 \times \dots \times X_K$ is the feasible set for x . $g(x)$ is continuously differentiable and block-wise convex, i.e., $g(x_1, \dots, x_K)$ is convex with respect to $x_i, i = 1, \dots, K$ (once $g(x)$ is convex with respect to $x \in X$, $g(x)$ is block-wise convex.); $h(x)$ is convex and separable, not necessarily differentiable.

The Problem (1.1) indeed covers a variety of application problems in many areas, to name a few, machine learning and statistics, e.g. [1–3], and has been intensively studied. Among previous works, a family of algorithms, Block Coordinate Descent(BCD) and its variants([4–7] and references therein), are of special interests. They are iterative methods and at each iteration new iterate is derived from either approximately or exactly minimizing the chosen variables while keeping the rest fixed. Assumptions made on $g(x), h(x)$ and different block-selection rules give rise to different convergence results. We briefly summarize related works, to our best knowledge, as follows.

1.1. Related Work

There are two mainstreams for solving Problem (1.1). One is to do exact minimization at each iteration while the other is to do approximate minimization and both of them prefer to employ the Gauss-Seidel update strategy. To be specific, when updating each block, the Gauss-Seidel style fixes the rest blocks to be the most up- to-date solution. Also, there's a less frequently employed Jacobi update strategy. When updating each coordinate, the Jacobi style fixes the rest blocks to be the solution obtained from previous cycle. As for block selection rules, cyclic rules, maximum block improvement rule and random rules are commonly used[4, 5].

For exact minimization, the local linear convergence rate was established under some assumptions by Luo and Tseng[8]. Tseng[9] proved the convergence for non-differentiable functions under certain conditions. However, doing exact minimization at each iteration may be theoretically intractable or numerically infeasible, so much more recent works focus on inexact minimization. When $h(x) := 0$, (block) coordinate descent combined with different step-size selection rules is commonly used. For example, global linear(sub-linear) convergence rate for strongly (general) convex optimization was established by Beck and Tetruashvili[10]. When the non-differentiable part $h(x)$ exists, one popular approximation method is to use a $\tilde{g}(x)$ to upper bound the smooth part $g(x)$ while keep the non-smooth convex term $h(x)$. Minimizing $\tilde{g}(x) + h(x)$ is often referred as proximal (block) coordinate descent in literature (e.g. [11]). More general and unified treatments for inexact minimization can be found in [4, 5]. Hong et al. suggested minimizing upper bound function at each iteration for chosen blocks and sub-linear convergence rate was established for non-smooth separable convex function. Tseng and Yun instead used quadratic approximation function combined with Armijo rule and established linear convergence rate for non-smooth separable convex function[4]. Usually, under some mild assumptions and conditions, combined with either deterministic or random block selection rules, sub-linear and linear convergence rate was established for general convex functions and strongly convex functions respectively in inexact minimization strategy.

Many numerical experiments in above mentioned works reveal that BCD-type algorithms enjoy satisfactory performances in practice, for example, [7, 12–14]. The underlying reasons for popularity of BCD-type algorithms includes following reasons. On the one hand, it breaks high dimension optimization problems into lower ones and first order information is used and thus the cost for both computation and storage at each iteration can be significantly reduced. On the other hand, it scales well with problem dimension, for example, in [6], Shai Shalev-Shwartz and Ambuj Tewari have proved that the required iteration to achieve (expected) ϵ accuracy grows linearly with the number of variables as

60 well as the number of training samples.

1.2. Motivation

Current existing sequential BCD-type algorithms, which optimize one block on one processor at one time, may not fulfill our desire to get faster since Nesterov[15] has proved that you can get at most the $O(\frac{1}{k^2})$ convergence rate
65 when only using gradients and function evaluations. Meanwhile, in [13], Nesterov claimed that, when the number of variables is huge, his proposed accelerated method may not be that efficient to implement. At the same time, with the significant progress made in collecting and storing data, there are imperious demands for scalable optimization algorithms. Many efficient algorithms, e.g.
70 [4], scale poorly with the growth of the problem's dimension.

Though a better convergence rate for BCD-type algorithms can not be obtained, thankfully, the availability of high performance multi-core computing platforms seems to be our last hope to be faster. In addition, there have been tremendous improvements in software, which help researchers to utilize computers
75 with multi-cores more efficiently. With the accesses of abundant computation resources and powerful software, we can optimize multi-blocks simultaneously to save the runtime. Therefore, we turn to work on parallel versions of BCD-type algorithms.

As mentioned by Wright([7]), there are two commonly employed parallel strategies. One is synchronous parallelism([3, 14, 16]) and the other is asynchronous parallelism([17] and references therein). Here, we only focus on the synchronous parallelism.
80

1.3. Contributions

Motivated by [5, 18, 19], we propose a Parallel Successive Upper approximation Minimization (PSUM) algorithm and its randomized variant(RPSUM). The
85 contributions of this work are as follows:

1. Most sequential BCD-type algorithm combine Gauss-Seidel update strategy with either deterministic or random block selection rule, e.g.[5], few

convergence analysis regarding Jacobi update exists. However, to parallelize BCD-type algorithm in a synchronous setting, one must employ Jacobi update strategy. We have proved $O(\frac{1}{k})$ convergence rate using PSUM to solve convex problems.

2. Unlike existing algorithms in literature, e.g. [14, 16], our algorithms use general approximation function, which treat their algorithms as special cases. Meanwhile, minimizing the approximation function actually can potentially be much easier than directly tackle with the original function. Moreover, in [16], when features are highly correlated, it fails to parallelize. But, our algorithm can also deal with this problem and accomplish full parallelism, that is, we can simultaneously update up to K blocks while the variables are highly correlated.
3. We find the idea behind our propose PSUM algorithm is similar with the algorithm in [3]. But our work is different in three aspects. First, PSUM is a parameter free method¹, that is, we do not need choose step-size to guarantee the convergence of our algorithm. On the contrary, the algorithm in [3] needs fine tuning step-size . Second, assumptions made on approximation function are less strict than those made in [3]. Third, the proof technique is also different, as we fully exploit the Jacobi update strategy.
4. When confronted with high dimension problem, optimizing with respect all blocks at the same time seems to be impractical, as we usually can't have enough processors. So we devised RPSUM, which simultaneously optimizes randomly chosen p blocks, coinciding with the maximal processors you have. Also, many literatures(e.g. [7]) can account for the reason for choosing blocks at random since they have reported that randomized block selection rule can outperform deterministic ones.

¹See the remark at the end of Section 3

1.4. Outline

The rest of this paper is organized as follows. Section 2 describes our proposed PSUM and RPSUM algorithms and necessary assumptions. In Section 3, convergence analysis and convergence rate is provided here. In Section 4, we implement our algorithms to solve the logistic regression problem with ℓ_1 regularization and also numerical results are reported here. We make extra remarks and draw conclusions in Section 5.

2. The PSUM and RPSUM Algorithms

2.1. Preliminaries

For simplicity, we need to introduce some auxiliary variables.

1. $x_{-i} = (x_1, \dots, x_{i-1}, x_{i+1}, \dots, x_K) \quad i = 1, 2, \dots, K$

$$\nabla \left(\sum_{i=1}^K g(\cdot, x_{-i}) \right) \Big|_x = \nabla g(x) \quad (2.1)$$

$$\nabla \left(\sum_{i=1}^K g(x_i, \cdot) \right) \Big|_x = (K-1) \nabla g(x) \quad (2.2)$$

2. $\xi_i^k \in \partial(h_i(\cdot)|_{x_i^k})$, where $\partial(h_i(\cdot)|_{x_i^k})$ represents a sub-gradient of $h_i(\cdot)$ at x_i^k

The notation $\nabla \left(\sum_{i=1}^K g(\cdot, x_{-i}) \right) \Big|_x$ seems to be obscure at glance. If we define a permutation matrix U partitioned as $U = [U_1, \dots, U_K]$, where $U_i \in \mathbb{R}^{n \times n_i}$ such that

$$x = \sum_{i=1}^K U_i x_i \quad x_i = U_i^T x, \quad i = 1, 2, \dots, K.$$

Then $\nabla \left(\sum_{i=1}^K g(\cdot, x_{-i}) \right) \Big|_x$ is equivalent to $\sum_{i=1}^K U_i \nabla_i g(x)$, where $\nabla_i g(x)$ is the partial gradient of g with respect to x_i .

2.2. Main Assumptions

Suppose $f(x)$ is a closed proper convex function in \mathbb{R}^n . Let $\text{dom } f$ denote the effective domain of $f(x)$ and let $\text{int}(\text{dom } f)$ denote the interior of $\text{dom } f$. We
135 assume $\text{dom } f \cap X \neq \emptyset$. Furthermore, we make assumptions on $f(x)$ regarding Problem (1.1).

Assumption A

Global minimum of Problem (1.1) can be attained on a set \mathbb{S} . There's a finite
140 R such that the level set for $f(\cdot)$ defined by x^0 is bounded, that is,

$$\max_{x \in \mathbb{S}} \max_x \{ \|x - x^*\| : f(x) < f(x^0) \} \leq R$$

Next we assume our chosen approximation function $u_i(\cdot, x)$ satisfy:

Assumption B

1. $u_i(x_i; x) = g(x)$, $\forall x \in X, \forall i \in \{1, 2, \dots, K\}$;
2. $u_i(v_i; x) \geq g(v_i, x_{-i})$, $\forall x \in X, x_i \in X_i, \forall i \in \{1, 2, \dots, K\}$;
- 145 3. $\nabla u_i(x_i; x) = \nabla_i g(x)$, $\forall x \in X, \forall i \in \{1, 2, \dots, K\}$;
4. $u_i(\cdot; x)$ has Lipschitz continuous gradient with respect to the second argument, that is, for any $x, y \in X$

$$\|\nabla u_i(\cdot; x) - \nabla u_i(\cdot; y)\| \leq G_i \|x - y\|$$

Define $G_{\max} = \max_i G_i$

We follow [5], also without causing any confusion, to call the $u_i(\cdot; x)$'s that satisfy Assumption B as a *upper-approximation* function.

2.3. The PSUM Algorithm

The PSUM algorithm is formally described as follows. We parallelize BCD algo-
150 rithms using Jacobi update strategy, which is clearly indicated in the Algorithm 1, step(6). First, we choose a feasible starting point $x^0 \in X$ and give a constant c , which is important for convergence and we shall see soon the way to set it. Then on each iteration, PSUM updates K blocks simultaneously. Until some
155 conditions are met, the algorithm is terminated. The intuition behind the key update rule is three-folds:

1. Use $u_i(z_i; x)$ to approximate the smooth part $g(x)$.
2. Keep the non-smooth regularized term $h_i(z_i)$ unchanged.
3. Use a quadratic term $c\|z_i - x_i\|^2$ to guarantee convergence.

Algorithm 1 PSUM for Problem (1.1)

- 1: **Initialization**
 - 2: $k \leftarrow 0$
 - 3: Consider $x_i^0 \in X^i$, for all $i = 1, 2, \dots, K$
 - 4: **while** not converged **do**
 - 5: **In parallel on i th processor** ($i = 1, 2, \dots, K$)
 - 6: $x_i^{k+1} \leftarrow \arg \min_{z_i \in X_i} u_i(z_i; x^k) + \lambda h_i(z_i) + c\|z_i - x_i^k\|^2$
 - 7: $k \leftarrow k + 1$
 - 8: **end while**
-

160 When n is large, but K is relative small, then this algorithm can achieve full parallelism, since you may afford K processors.

2.4. The RPSUM Algorithm

However, when K is sufficiently large, for example, if we employ coordinate descent, then the K equals to the number of variables, which is particularly
165 huge in genome domain. Under these circumstances, we may fail to achieve full parallelism, simply owing to the lack of adequate processors. Therefore, we propose the RPSUM with slight modification in the block selection rule. As the PSUM, we first choose a feasible starting point $x^0 \in X$ and give a constant c . Then on the iteration $t + 1$, Algorithm 2 chooses $p < K$ coordinates
170 independently and uniformly at random from $\{1, 2, \dots, K\}$; these form a multi-set S_{t+1} .² Blocks indexed by S_{t+1} are updated in parallel while the rest blocks are held fixed. Until some conditions are met, the algorithm is terminated. The RPSUM seems to be less efficient, as in iteration $t + 1$, $K - p$ blocks remain unchanged. But it's a trade-off since we might not be able to utilize K processors

²For the sake of intelligibility, we use t instead of k to number the iteration.

Algorithm 2 RPSUM for Problem (1.1)

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1: Initialization
2:  $t \leftarrow 0$ 
3: Consider  $x_i^0 \in X^i$ , for all  $i = 1, 2, \dots, K$ 
4: while not converged do
5:   In parallel on each of  $p$  processors
6:     Choose  $i \in \{1, 2, \dots, K\}$  uniformly at random
7:      $x_i^{t+1} \leftarrow \arg \min_{z_i \in X_i} u_i(z_i; x^t) + \lambda h_i(z_i) + c \|z_i - x_i^t\|^2$ 
8:     For those coordinates( $js$ ) that are not chosen,  $x_j^{t+1} \leftarrow x_j^t$ 
9:      $t \leftarrow t + 1$ 
10: end while

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175 when K is incredibly huge.

3. Convergence Analysis for PSUM and RPSUM

We provide detailed convergence analysis for the PSUM in this section and the proof is quite similar for RPSUM, hence only some differences will be explicitly addressed.

180 *3.1. Convergence Analysis for PSUM*

Proposition 3.1. *Suppose Assumption B holds, then for all $x, y, z \in X$, with $x = (x_1, \dots, x_K), y = (y_1, \dots, y_K), z = (z_1, \dots, z_K)$,*

$$\|\nabla\left(\sum_{i=1}^K u_i(\cdot; x)\right)\big|_z - \nabla\left(\sum_{i=1}^K u_i(\cdot; y)\right)\big|_z\| \leq \sqrt{K}G_{max}\|x - y\| \quad (3.1)$$

PROOF.

$$\begin{aligned}
\|\nabla(\sum_{i=1}^K u_i(\cdot; x))|_z - \nabla(\sum_{i=1}^K u_i(\cdot; y))|_z\| &= \|\sum_{i=1}^K (\nabla u_i(\cdot; x) - \nabla u_i(\cdot; y))|_z\| \\
&= \left\| \begin{pmatrix} \nabla u_1(z_1; x) - \nabla u_1(z_1; y) \\ \vdots \\ \nabla u_K(z_K; x) - \nabla u_K(z_K; y) \end{pmatrix} \right\| \\
&= \sqrt{\sum_{i=1}^K \|\nabla u_i(z_i; x) - \nabla u_i(z_i; y)\|^2} \\
&\leq \sqrt{K} G_{max} \|x - y\| \tag{3.2}
\end{aligned}$$

The last inequality holds due to the Assumption B.4

Proposition 3.2. *Suppose Assumption B holds then the sequence $\{x^k\}$, generated by Algorithm 1, satisfies,*

$$(x^{k+1} - x^k) \nabla g(x^{k+1}) \leq (\sqrt{K} G_{max} - 2c) \|x^{k+1} - x^k\|^2 + \lambda(h(x^k) - h(x^{k+1})) \tag{3.3}$$

PROOF. Consider the problem,

$$\tilde{x}^{k+1} \leftarrow \arg \min_{z \in X} \sum_{i=1}^K \{u_i(z_i; x^k) + \lambda h_i(z_i) + c \|z_i - x_i^k\|^2\} \tag{3.4}$$

Due to the separable structure of (3.4), we have $\tilde{x}^{k+1} = x^{k+1}$. By the optimality of x^{k+1} , we derive,

$$(x^k - x^{k+1})^T \nabla \left(\sum_{i=1}^K \{u_i(z_i; x^k) + \lambda h_i(z_i) + c \|z_i - x_i^k\|^2\} \right) \Big|_{z=x^{k+1}} \geq 0 \tag{3.5}$$

Simple operations on (3.5) gives rise to

$$\begin{aligned}
0 &\leq (x^k - x^{k+1})^T \left[\nabla \left(\sum_{i=1}^K u_i(z_i; x^k) \right) \Big|_{z=x^{k+1}} \right] + \lambda (x^k - x^{k+1})^T \xi^{k+1} - 2c \|x^{k+1} - x^k\|^2 \\
&\leq (x^k - x^{k+1})^T \left[\nabla \left(\sum_{i=1}^K u_i(z_i; x^k) \right) \Big|_{z=x^{k+1}} \right] + \lambda [h(x^k) - h(x^{k+1})] - 2c \|x^{k+1} - x^k\|^2
\end{aligned} \tag{3.6}$$

where $\xi^{k+1} \in \partial h(x^{k+1})$. The last inequality holds naturally by the definition of sub-gradient

$$h(x^k) - h(x^{k+1}) \geq (x^k - x^{k+1})^T \xi^{k+1}$$

190 Adding $(x^{k+1} - x^k)^T [\nabla(\sum_{i=1}^K u_i(z_i; x^{k+1}))|_{z=x^{k+1}}]$ on both sides of (3.6)

$$\begin{aligned} & (x^{k+1} - x^k)^T [\nabla(\sum_{i=1}^K u_i(z_i; x^{k+1}))|_{z=x^{k+1}}] \\ & \leq -2c\|x^{k+1} - x^k\|^2 + \lambda[h(x^k) - h(x^{k+1})] - \\ & (x^{k+1} - x^k)^T [\nabla(\sum_{i=1}^K u_i(z_i; x^k))|_{z=x^{k+1}} - u_i(z_i; x^{k+1})|_{z=x^{k+1}}] \end{aligned} \quad (3.7)$$

Plugging (3.1) into (3.7) and using Cauchy-Schwartz inequality, we have

$$\begin{aligned} (x^{k+1} - x^k)^T \nabla g(x^{k+1}) &= (x^{k+1} - x^k)^T [\nabla(\sum_{i=1}^K u_i(z_i; x^{k+1}))|_{z=x^{k+1}}] \\ &\leq (\sqrt{K}G_{max} - 2c)\|x^{k+1} - x^k\|^2 + \lambda[h(x^k) - h(x^{k+1})] \end{aligned} \quad (3.8)$$

Lemma 3.1 (Sufficient Descent). *Suppose Assumption A and B hold then the sequence $\{x^k\}$, generated by Algorithm 1, satisfies,*

$$f(x^{k+1}) \leq f(x^k) - \gamma\|x^{k+1} - x^k\|^2 \quad (3.9)$$

where $K\gamma = [c - (K-1)(\sqrt{K}G_{max} - 2c)] > 0$ and $c > \frac{K-1}{2K-1}\sqrt{K}G_{max}$.

PROOF. By the definition of x_i^{k+1} we derive

$$\sum_{i=1}^K u_i(x_i^{k+1}; x^k) + \lambda h(x^{k+1}) + c\|x^{k+1} - x^k\|^2 \leq \sum_{i=1}^K u_i(x_i^k; x^k) + \lambda h(x^k) \quad (3.10)$$

By the assumptions of $u_i(\cdot; x)$, we have

$$* \text{ (a) } u_i(v_i; x) \geq g(v_i, x_{-i}) \quad \forall v_i \in X_i, x \in X, i = 1, 2, \dots, K$$

195 * (b) $u_i(x_i; x) = g(x_i, x_{-i}) \quad \forall x \in X, i = 1, 2, \dots, K$

Hence, plugging (a),(b) into (3.10) and rearranging it, we obtain

$$\begin{aligned} \sum_{i=1}^K g(x_i^{k+1}, x_{-i}^k) + \lambda h(x^{k+1}) &\leq \sum_{i=1}^K g(x_i^k, x_{-i}^k) + K\lambda h(x^k) - c\|x^{k+1} - x^k\|^2 \\ &= Kg(x^k) + \lambda h(x^k) - c\|x^{k+1} - x^k\|^2 \end{aligned} \quad (3.11)$$

By convexity of $\sum_{i=1}^K g(x_i^{k+1}, \cdot)$, we have

$$\begin{aligned} \sum_{i=1}^K g(x_i^{k+1}, x_{-i}^k) &\geq \sum_{i=1}^K g(x_i^{k+1}, x_{-i}^{k+1}) + (x^k - x^{k+1})^T \nabla \left(\sum_{i=1}^K g(x_i^{k+1}, \cdot) \right) \Big|_{x^{k+1}} \\ &= Kg(x^{k+1}) + (K-1)(x^k - x^{k+1})^T \nabla g(x^{k+1}) \end{aligned} \quad (3.12)$$

Combing (3.8),(3.11),(3.12) produces

$$\begin{aligned} &Kg(x^{k+1}) + \lambda h(x^{k+1}) \\ &\leq Kg(x^k) + \lambda h(x^k) - c\|x^{k+1} - x^k\|^2 + (K-1)(x^{k+1} - x^k)^T \nabla g(x^{k+1}) \\ &\leq Kg(x^k) + \lambda h(x^k) + (K-1)\lambda[h(x^k) - h(x^{k+1})] + \\ &\quad [-c + (K-1)(\sqrt{K}G_{max} - 2c)]\|x^{k+1} - x^k\|^2 \end{aligned} \quad (3.13)$$

Rearranging the (3.13) gives

$$Kg(x^{k+1}) + K\lambda h(x^{k+1}) \leq Kg(x^k) + K\lambda h(x^k) + [-c + (K-1)(\sqrt{K}G_{max} - 2c)]\|x^{k+1} - x^k\|^2$$

200 Choose $c > \frac{K-1}{2K-1}\sqrt{K}G_{max}$, we conclude (3.9).

Lemma 3.2 (Cost-to-Go Estimate). *Suppose Assumption A and Assumption B hold, then the sequence $\{x^k\}$, generated by Algorithm 1, satisfies*

$$f(x^{k+1}) - f(x^*) \leq \theta\|x^{k+1} - x^k\| \quad (3.14)$$

where $\theta = (G_{max}K + 2c)R$

PROOF.

$$\begin{aligned}
& f(x^{k+1}) - f(x^*) + 2c(x^{k+1} - x^k)^T(x^{k+1} - x^*) \\
&= g(x^{k+1}) - g(x^*) + h(x^{k+1}) - h(x^*) + 2c(x^{k+1} - x^k)^T(x^{k+1} - x^*) \\
&\leq (x^{k+1} - x^*)^T \nabla g(x^{k+1}) + h(x^{k+1}) - h(x^*) + 2c(x^{k+1} - x^k)^T(x^{k+1} - x^*) \\
&= \sum_{i=1}^K (x_i^{k+1} - x_i^*) [\nabla_i g(x^{k+1}) - \nabla u_i(x_i^{k+1}; x^k)] + \sum_{i=1}^K (x_i^{k+1} - x_i^*)^T [\nabla u_i(x_i^{k+1}; x^k)] + \\
&\quad h(x^{k+1}) - h(x^*) + 2c(x^{k+1} - x^k)^T(x^{k+1} - x^*) \tag{3.15}
\end{aligned}$$

Notice that $x_i^{k+1} = \arg \min_{z_i \in X_i} u_i(z_i; x^k) + \lambda h_i(z_i) + c\|z_i - x_i^k\|^2$. Using the optimality condition of this problem, we claim that there exists some $\xi_i^{k+1} \in \partial h_i(x_i^{k+1})$ such that

$$0 \geq (x_i^{k+1} - x_i^*) [\nabla u_i(x_i^{k+1}; x^k) + \lambda \xi_i^{k+1} + 2c(x_i^{k+1} - x_i^k)] \tag{3.16}$$

$$\begin{aligned}
&\geq (x_i^{k+1} - x_i^*) [\nabla u_i(x_i^{k+1}; x^k)] + \lambda [h_i(x_i^{k+1}) - h_i(x_i^*)] + 2c(x_i^{k+1} - x_i^k)(x_i^{k+1} - x_i^*) \\
&\tag{3.17}
\end{aligned}$$

205 In view of (3.15) and (3.17), we derive

$$\begin{aligned}
& f(x^{k+1}) - f(x^*) + 2c(x^{k+1} - x^k)^T(x^{k+1} - x^*) \\
&\leq \sum_{i=1}^K (x_i^{k+1} - x_i^*) [\nabla_i g(x^{k+1}) - \nabla u_i(x_i^{k+1}; x^k)] \\
&= \sum_{i=1}^K (x_i^{k+1} - x_i^*) [\nabla u_i(x_i^{k+1}; x^{k+1}) - \nabla u_i(x_i^{k+1}; x^k)] \\
&\leq \sum_{i=1}^K \|x_i^{k+1} - x_i^*\| \|\nabla u_i(x_i^{k+1}; x^{k+1}) - \nabla u_i(x_i^{k+1}; x^k)\| \\
&\leq \sum_{i=1}^K G_i \|x^{k+1} - x^k\| \|x_i^{k+1} - x_i^*\| \\
&\leq G_{max} RK \|x^{k+1} - x^k\| \tag{3.18}
\end{aligned}$$

where the last inequality holds due to the Assumption A. Rearranging (3.18),

we obtain

$$\begin{aligned} f(x^{k+1}) - f(x^*) &\leq G_{max}RK\|x^{k+1} - x^k\| + 2c(x^k - x^{k+1})^T(x^{k+1} - x^*) \\ &\leq (G_{max}K + 2c)R\|x^{k+1} - x^k\|. \end{aligned} \quad (3.19)$$

Theorem 3.1. *Suppose Assumption A and B hold then the sequence $\{x^k\}$, generated by Algorithm 1, satisfies,*

$$\Delta_k = f(x^k) - f(x^*) \leq \frac{\alpha}{\sigma} \frac{1}{k}, \forall k \geq 1 \quad (3.20)$$

where $\sigma = \gamma/\theta^2$ and $\alpha = \max\{f(x^1) - f(x^*), 4\sigma - 2, 2\}$.

PROOF. From Lemma3.1 and Lemma3.2, we have

$$\Delta_k - \Delta_{k+1} \geq \gamma\|x^{k+1} - x^k\|^2 \geq \frac{\gamma}{\theta^2} \Delta_{k+1}^2 = \sigma \Delta_{k+1}^2$$

or equivalently,

$$\sigma \Delta_{k+1}^2 + \Delta_{k+1} \leq \Delta_k, \forall k \geq 1 \quad (3.21)$$

From (3.21) and $\Delta_1 \leq \alpha$, we derive

$$\Delta_2 = \frac{-1 + \sqrt{1 + 4\sigma\Delta_1}}{2\sigma} < \frac{-1 + \sqrt{1 + 4\sigma\alpha}}{2\sigma} = \frac{2\alpha}{1 + \sqrt{1 + 4\sigma\alpha}} \leq \frac{2\alpha}{1 + |4\sigma - 1|}$$

210 where the last inequality holds as $\alpha \geq 4\sigma - 2$. We assert that, at this point,

$\Delta_2 \leq \frac{\alpha}{2\sigma}$, since

* When $4\sigma - 1 \geq 0$, then $\Delta_2 \leq \frac{\alpha}{2\sigma}$.

* When $4\sigma - 1 < 0$, then $\Delta_2 \leq \frac{2\alpha}{2-4\sigma} \leq \frac{2\alpha}{8\sigma-4\sigma} = \frac{\alpha}{2\sigma}$.

Now, we argue that if $\Delta_k \leq \frac{\alpha}{k\sigma}$, then the following holds

$$\Delta_{k+1} \leq \frac{\alpha}{(k+1)\sigma}.$$

215 It can be seen from the fact that

$$\Delta_{k+1} \leq \frac{-1 + \sqrt{1 + \frac{4\alpha}{k}}}{2\sigma} = \frac{2\alpha}{k\sigma(1 + \sqrt{1 + \frac{4\alpha}{k}})} \leq \frac{2\alpha}{\sigma(k + \sqrt{k^2 + 4k + 1})} = \frac{\alpha}{\sigma(k + 1)} \quad (3.22)$$

where the last inequality holds due to the fact that $\alpha, k \geq 2$. Therefore, we conclude our proof.

Although techniques used to prove Theorem3.1 are quite similar to those in [5], it's noteworthy that the [5] employs the on Gauss-Seidel update strategy, so we cannot directly derive Theorem3.1 from it and need to find a new way instead. We achieve this by first establishing Lemma3.1 and Lemma3.2. The proofs of them fully exploit the Jacobi update structures, hence completely new. Based on Lemma3.1 and Lemma3.2, then Theorem3.1 can be easily derived.

3.2. Convergence Analysis for RPSUM

Before beginning our analysis, we need to introduce additional auxiliary variables for simplicity. Since the S_{t+1} is a random index set, there are actually $q = C_K^p$ possible values, and we denote them as $S_{t+1}^j, j = 1, \dots, q$. By the construction of Algorithm 2, we know x^{t+1} depends on x^t through S_{t+1} . Therefore, there are q possible values of x^{t+1} . We explicitly formulate this relation discussed above through the set function

$$\phi(S_{t+1}^j; x^t) = w_j^{t+1}, j = 1, 2, \dots, q$$

where w_j^{t+1} represents one possible value of x^{t+1} .

Lemma 3.3 (Expected Sufficient Descent). *Suppose Assumption B hold, then the sequence $\{x_t\}$, generated by Algorithm 2, satisfies,*

$$E[f(x^{t+1}) - f(x^t)] \leq -\gamma_e E\|x^{t+1} - x^t\|^2 \quad (3.23)$$

where $K\gamma_e = p(\sqrt{K}G_{max} - 2c) > 0$ and $c > \frac{1}{2}\sqrt{K}G_{max}$ and $E[\cdot]$ is taken with respect to random index sets $S_1, S_2, \dots, S_t, S_{t+1}$.

PROOF.

$$\begin{aligned}
E_{S_{t+1}}[f(x^{t+1}) - f(x^t)] &= E_{S_{t+1}}\left[\frac{1}{p} \sum_{i \in S_{t+1}} \{g(x_i^{t+1}, x_{-i}^{t+1}) - g(x_i^t, x_{-i}^t)\} + \lambda \sum_{i \in S_{t+1}} \{h_i(x_i^{t+1}) - h_i(x_i^t)\}\right] \\
&= \frac{1}{q} \sum_{j=1}^q [g(w_j^{t+1}) - g(x^t)] + \lambda \frac{p}{K} [h(\hat{x}^{t+1}) - h(x^t)] \\
&\stackrel{(1)}{\leq} \sum_{j=1}^q [(w_j^{t+1} - x^t)^T \nabla g(w_j^{t+1})] + \lambda \frac{p}{K} [h(\hat{x}^{t+1}) - h(x^t)] \\
&\stackrel{(2)}{=} \frac{p}{K} (\hat{x}^{t+1} - x^t)^T \nabla g(\hat{x}^{t+1}) + \lambda \frac{p}{K} [h(\hat{x}^{t+1}) - h(x^t)]
\end{aligned} \tag{3.24}$$

where \hat{x}_i^{t+1} is generated by the Algorithm 1; i.e., $\hat{x}_i^{t+1} \leftarrow \arg \min_{z_i \in X_i} u_i(z_i; x^t) + \lambda h_i(z_i) + c \|z_i - x_i^t\|^2$, $i = 1, \dots, K$ and x^t is generated by Algorithm 2. By convexity of $g(\cdot)$, (1) holds. As for (2), notice the fact that those updated coordinate i , x_i^{t+1} will be counted for C_{K-1}^{-1} times and $x_i^{t+1} = \hat{x}_i^{t+1}$.

Taking full expectation with respect to S_1, \dots, S_t on both sides of (3.24), we obtain

$$f(\tilde{x}^{t+1}) - f(\tilde{x}^t) \leq \frac{p}{K} (\tilde{x}^{t+1} - \tilde{x}^t)^T \nabla g(\tilde{x}^{t+1}) + \lambda \frac{p}{K} [h(\tilde{x}^{t+1}) - h(\tilde{x}^t)] \tag{3.25}$$

where \tilde{x}_i^{t+1} is generated by the Algorithm 1; i.e., $\tilde{x}_i^{t+1} \leftarrow \arg \min_{z_i \in X_i} u_i(z_i; \tilde{x}^t) + \lambda h_i(z_i) + c \|z_i - \tilde{x}_i^t\|^2$, $i = 1, \dots, K$.

Using the results from Lemma3.2, we have,

$$(\tilde{x}^{t+1} - \tilde{x}^t)^T \nabla g(\tilde{x}^{t+1}) \leq (\sqrt{K} G_{max} - 2c) \|\tilde{x}^{t+1} - \tilde{x}^t\|^2 + \lambda (\tilde{h}(\tilde{x}^t) - h(\tilde{x}^{t+1})) \tag{3.26}$$

Plugging (3.26) into (3.25) leads to

$$E(f(x^{t+1}) - f(x^t)) = f(\tilde{x}^{t+1}) - f(\tilde{x}^t) \leq \frac{p}{K} (\sqrt{K} G_{max} - 2c) \|\tilde{x}^{t+1} - \tilde{x}^t\|^2 = \frac{p}{K} (\sqrt{K} G_{max} - 2c) E\|x^{t+1} - x^t\|^2$$

Remark The techniques used here are more straightforward than that used in Lemma3.1, but it also requires us to choose larger c than that of Lemma3.1 to guarantee the sequence $E[f(x^t)]$ are non-increasing.

To go on our proof, we need to modify the **Assumption A** as

Assumption C

Global minimum of Problem (1.1) can be attained on a set \mathbb{E} , where all of its
 250 elements are generated by Algorithm 2. There's a finite R_e such that the level
 set for $f(\cdot)$ defined by x^0 is bounded, that is,

$$\max_{x \in \mathbb{E}} \max_x \{E\|x - x^*\|^2 : E[f(x)] < f(x^0)\} \leq R_e^2$$

where expectation is taken with respect to all of random variables. Now we give
 the counterpart of Lemma 3.2.

Lemma 3.4 (Expected Cost-to-Go Estimate). *Suppose Assumption B and
 Assumption C hold, then*

$$E[f(x^{k+1}) - f(x^*)] \leq \theta_e \sqrt{E\|x^{t+1} - x^t\|^2} \quad (3.27)$$

where $\theta_e = (pG_{max} + 2c)R_e$

255 **PROOF.** The proof is almost the same as Lemma3.2. We only address the
 different part here. It's easy to derive

$$f(x^{t+1}) - f(x^*) + 2c(x^{t+1} - x^t)^T(x^{t+1} - x^*) \leq \sum_{i \in S_{t+1}} G_i \|x^{t+1} - x^t\| \|x_i^{t+1} - x_i^*\|$$

Taking expectation with S_{t+1} on both sides and using Cauchy-Schwartz inequality,
 we derive

$$\begin{aligned} E_{S_{t+1}}[f(x^{t+1}) - f(x^*)] &\leq E_{S_{t+1}}\{\|x^{t+1} - x^t\| \sum_{i \in S_{t+1}} G_i \|x_i^{t+1} - x_i^*\|\} + E_{S_{t+1}}\{2c(x^t - x^{t+1})^T(x^{t+1} - x^*)\} \\ &\leq G_{max} \sqrt{E_{S_{t+1}}\|x^{t+1} - x^t\|^2} \sqrt{E_{S_{t+1}}\{\sum_{i \in S_{t+1}} \|x_i^{t+1} - x_i^*\|^2\}} + 2c \sqrt{E_{S_{t+1}}\|x^{t+1} - x^t\|^2} \sqrt{E_{S_{t+1}}\|x^{t+1} - x^*\|^2} \\ &\stackrel{(1)}{\leq} G_{max} \sqrt{E_{S_{t+1}}\|x^{t+1} - x^t\|^2} \sqrt{E_{S_{t+1}}\{p \sum_{i \in S_{t+1}} \|x_i^{t+1} - x_i^*\|^2\}} + 2c \sqrt{E_{S_{t+1}}\|x^{t+1} - x^t\|^2} \sqrt{E_{S_{t+1}}\|x^{t+1} - x^*\|^2} \\ &= G_{max} p \sqrt{E_{S_{t+1}}\|x^{t+1} - x^t\|^2} \sqrt{E_i\|x_i^{t+1} - x_i^*\|^2} + 2c \sqrt{E_{S_{t+1}}\|x^{t+1} - x^t\|^2} \sqrt{E_{S_{t+1}}\|x^{t+1} - x^*\|^2} \\ &\stackrel{(2)}{\leq} (pG_{max} + 2c)R_e \sqrt{E_{S_{t+1}}\|x^{t+1} - x^t\|^2} = \theta_e \sqrt{E_{S_{t+1}}\|x^{t+1} - x^t\|^2} \quad (3.28) \end{aligned}$$

The (1) inequality holds due to the fact that $(\sum_{i=1}^n a_i)^2 \leq n \sum_{i=1}^n a_i^2$, $a_i \in \mathbb{R}$;
 260 the (2) holds as the result of $E[f(x^{t+1})] \leq f(x^0)$ (derived from Lemma 3.3).
 Taking full expectation on both sides of (3.28) leads to

$$E[f(x^{t+1})] - f(x^*) \leq E[\theta_e \sqrt{E_{S_{t+1}} \|x^{t+1} - x^t\|^2}] \leq \theta_e \sqrt{E \|x^{t+1} - x^t\|^2}$$

where in the last inequality, we use the fact that $E(l(x)) \leq l(E(x))$, if $l(x)$ is concave. Hence, we conclude our claim.

Follow the results derived in Theorem 3.1, we can immediately assert

Theorem 3.2. *Suppose Assumption B and Assumption C hold, then the sequence $\{x^t\}$, generated by Algorithm 2, satisfies,*

$$\Delta_t = E[f(x^t) - f(x^*)] \leq \frac{\alpha_e}{\sigma_e} \frac{1}{t}, \forall t \geq 1 \quad (3.29)$$

265 where $\sigma_e = \gamma_e / \theta_e^2$ and $\alpha_e = \max\{E[f(x^1)] - f(x^*), 4\sigma_e - 2, 2\}$.

Remark PSUM and RPSUM are claimed to be parameter-free, due to the fact that no specific c is required in the convergence analysis once *Assumption B* is satisfied. The convergence results hold for sufficiently large c . However, in numerical experiments, the Lipschitz constant G_{max} is required. In some cases,
 270 the Lipschitz constant can be derived from computing the maximum eigenvalue of the Hessian, like our experiments given below. In other cases, there are some numerical methods can be applied to find an appropriate estimation of Lipschitz constant, for example the method in [13].

4. Application

275 In this section, we use the proposed RPSUM to solve regularized logistic regression problem. The setup for logistic regression are as usual. Suppose we have N pairs of observation (x_i, y_i) , where $x_i \in \mathbb{R}^p$ are predict variables and $y_i \in \mathbb{R}$ are binary response variables. Without loss of generality, we assume that x_{ij} are

standardized: $\sum_{i=1}^N x_{ij} = 0$, $\frac{1}{N} \sum_{i=1}^N x_{ij}^2 = 1$, for $j = 1, 2, \dots, p$. and response
 280 variable y_i take value in $\{0, 1\}$.

The logistic regression model represents the class-conditional probabilities for all pairs of (x_i, y_i) through a linear function of the predictors

$$\begin{aligned} P(y_i = 0|x_i) &= \frac{1}{1 + e^{-(\beta_0 + x_i^T \beta)}} \\ P(y_i = 1|x_i) &= \frac{1}{1 + e^{(\beta_0 + x_i^T \beta)}} \end{aligned} \quad (4.1)$$

where $\beta \in \mathbb{R}^p$, $\beta_0 \in \mathbb{R}$. Equivalently, (4.1) implies that

$$\frac{P(y_i = 0|x_i)}{P(y_i = 1|x_i)} = \beta_0 + x_i^T \beta \quad (4.2)$$

In order to find the proper β_0, β , a widely employed strategy is to maximize the
 285 log likelihood

$$\ell(\beta_0, \beta) = \sum_{i=1}^N [y_i(\beta_0 + x_i^T \beta) - \ln(1 + e^{\beta_0 + x_i^T \beta})] \quad (4.3)$$

which is a concave function of parameter β_0, β .

The ℓ_1 -regularized logistic regression is to maximize log-likelihood function with ℓ_1 penalty

$$\max_{(\beta_0, \beta) \in \mathbb{R}^{p+1}} \ell(\beta_0, \beta) - \lambda |\beta|_1 \quad (4.4)$$

Solving Problem (4.4) is equivalent to tackle with the following minimization
 290 problem

$$\min_{(\beta_0, \beta) \in \mathbb{R}^{p+1}} f(\beta_0, \beta) = -\frac{1}{2N} \ell(\beta_0, \beta) + \lambda |\beta|_1 \triangleq g(\beta_0, \beta) + h(\beta_0, \beta) \quad (4.5)$$

Since $\ell(\beta_0, \beta)$ is concave, $f(\beta_0, \beta)$ is convex, hence Algorithms1 and Algorithm2 can be employed.

4.1. RPSUM for Logistic Regression with ℓ_1 regularization

To solve Problem (4.5), we need first to partition the argument $(\beta_0, \beta) \in \mathbb{R}^{p+1}$ into $p+1$ coordinates(blocks), that is $(\beta_0, \beta) = (\beta_0, \beta_1, \dots, \beta_p)$ and then specify the form of $u_j(\cdot; \beta_0, \beta)$ as

$$u_j(z_j; \beta_0, \beta) = g(\beta_0, \beta) + (z_j - \beta_j) \nabla_j g(\beta_0, \beta) + \frac{L}{2} (z_j - \beta_j)^2, j = 0, 1, 2, \dots, p \quad (4.6)$$

Since the largest eigenvalue of $\nabla^2 g(\beta_0, \beta_p)$, denoted as $\lambda_{max}(\nabla^2 g(\beta_0, \beta_p))$, is no bigger than $\frac{1}{4}$, therefore we set L in (4.6) to $\frac{1}{4}$. Next, we dive into the core of both Algorithms1 and Algorithm2. Due to the explicit form of Problem (4.5), we can solve the low dimension problem

$$\min_{z_j \in \mathbb{R}} u_j(z_j; \beta_0, \beta) + \lambda |z_j| + c \|z_i - \beta_j\|^2 = (z_j - \beta_j) \nabla_j g(\beta_0, \beta) + (\frac{1}{8} + c)(z_j - \beta_j)^2 + \lambda |z_j| \quad j = 1, 2, \dots, p \quad (4.7)$$

precisely in a closed form since it's a lasso type problem. For $j = 0$ case, the above (4.7) simply becomes quadratic optimization problem, which takes form

$$\min_{z_0 \in \mathbb{R}} (z_0 - \beta_0) \nabla_0 g(\beta_0, \beta) + (\frac{1}{8} + c)(z_0 - \beta_0)^2 \quad (4.8)$$

where $\nabla_0 g(\beta_0, \beta)$ is the partial gradient with respect to β_0 . Solving (4.7) is equivalent to solve

$$\min_{z_j \in \mathbb{R}} (\frac{1}{8} + c) z_j^2 + m_j z_j + \lambda |z_j| \quad j = 1, 2, \dots, p$$

and thus

$$z_j = \begin{cases} \frac{-m_j - \lambda}{2(\frac{1}{8} + c)} & m_j < 0, |m_j| > \lambda \\ \frac{-m_j + \lambda}{2(\frac{1}{8} + c)} & m_j > 0, |m_j| > \lambda \\ 0 & \text{otherwise} \end{cases} \quad j = 1, 2, \dots, p$$

where $m_j = \nabla_j f(\beta_0, \beta) - 2(\frac{1}{8} + c)\beta_j$. As for (4.8), simple calculations imply that

$$z_0 = \frac{-m_0}{2(\frac{1}{8} + c)}.$$

Now we give a pseudo-code for solving for Logistic Regression with ℓ_1 regularization to conclude this subsection.(See Algorithm3.)

Algorithm 3 RPSUMfor Logistic Regression with ℓ_1 regularization

```

1: Initialization
2:  $t \leftarrow 0$ 
3: Set  $\beta_0 = 0, \beta = 0_p$ , where  $0_p$  represent a  $p$  dimension vector whose elements
   are all 0.
4: while not converged do
5:   In parallel on  $q$  processors
6:     Choose  $j \in \{0, 1, 2, \dots, p\}$  uniformly at random.
7:      $\beta_j^{t+1} \leftarrow z_j^t$ , where  $z_j^t$  are defined as above but use  $\beta_0^t, \beta^t$  where
       appropriate.
8:     For those coordinates( $js$ ) that are not chosen,  $\beta_j^{t+1} \leftarrow \beta_j^t$ 
9:      $t \leftarrow t + 1$ 
10: end while

```

310 4.2. Experiments

In this section, we perform our proposed algorithms for Logistic Regression with ℓ_1 regularization on both real and simulated datasets. We make a few but important statements on the way our numerical experiments conducted.

- An Intel Xeon machine is used for experiments. All codes are written in Python and the multiprocessing library is used for parallel computation.
- We denote the lower bound $\frac{1}{2}\sqrt{K}G_{max}$ as \bar{c} and to guarantee the theoretic convergence, we set $c = \bar{c}(1 + 0.01)$.

- We make slight twist to our algorithms to address the unavoidable communication overhead[20]. In one iteration, we update 40 scalar variables on each processor sequentially. We still refer it as a method that updates q (number of processors available) variables in parallel. To be more specific, if we have 4 threads available , then in each iteration, we will update 160 scalar variables, with 40 scalar variables processed by per processor. As for how to assign the 40 scalar variables for each processor, we employ two rules. One is cyclic rule that corresponds to PSUM while the other is random rules that represents the RPSUM.
- To make our results comparable, we set $q = 1$ in our program to represent the serial counterparts.

4.2.1. Real Data

We use Leukemia [21], which is gene-expression data with a binary response indicating type of leukemia—AML vs ALL, to conduct the numerical experiments. We used the preprocessed data of [12]. The results are given in Figure 1.

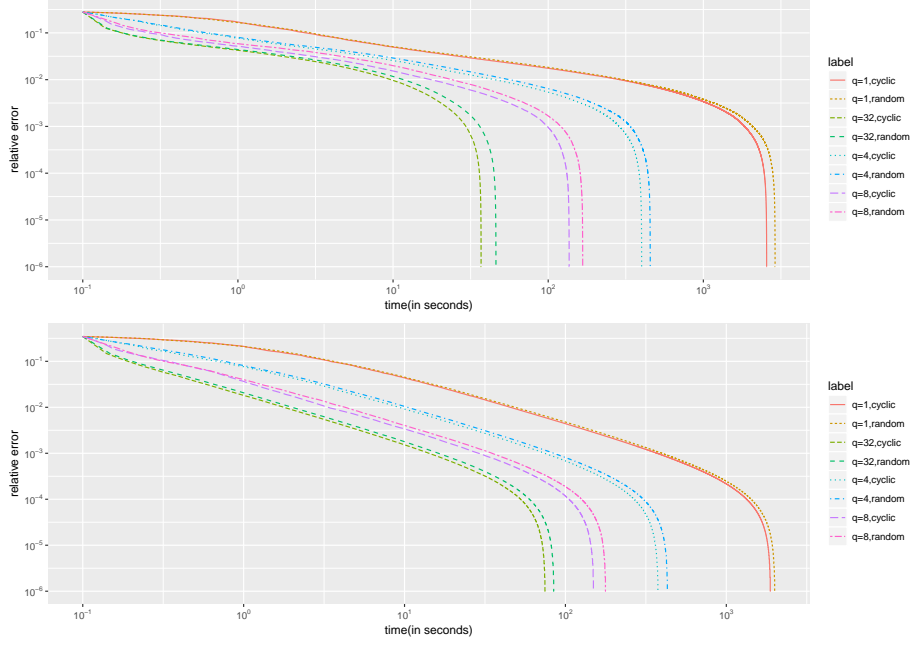


Figure 1: Performance comparison for RPSUM and PSUM with different degrees of regularization. The top figure represents the strong regularization situation with $\lambda = 10^{-2}$ while the bottom one correspond to the $\lambda = 10^{-6}$.

The y axis represents the relative error, which is defined by $f(x^{t+1}) - f^*$. It's quite clear that, with the increase with of available processors, we gain the near-linear speed up. Also, the number of data access for each method is near the same (see Table 1), so the comparison is really fair. It's also worth noting that the unpleasant phenomenon in [3] does not occur in this groups of experiments, which means we may assert that the more processors you have, more time you can save.

Methods	$\lambda = 10^{-2}$	$\lambda = 10^{-6}$
q=1,cyclic	2410880	1760880
q=1,random	2502640	1761320
q=4,cyclic	2401600	2254080
q=4,random	2399520	2253440
q=8,cyclic	2387200	2486720
q=8,random	2390400	2489920
q=32,cyclic	22627840	2728960
q=32,random	2659840	2731520

Table 1: Number of data access of Leukemia dataset with different q.

340 4.2.2. Simulated Data

We follow the simulation setup described in [12].

- First, generate the coefficient

$$\beta_j = (-1)^j e^{\left(\frac{2(j-1)}{20}\right)}, j = 1, 2, \dots, p.$$

- Second, generate the design matrix X of size $n \times p$. Then each row of X represents one observation, denoted as x_i .
- 345 • Third, generate the error vector ε of size $n \times 1$, with each element drawn from the Normal Distribution $N(0, 1)$. Together with X and β , we produce the working response Z ,

$$Z = X\beta + \varepsilon.$$

- Finally, define $p_i = 1/(1 + \exp(-Z[i])), i = 1, 2, \dots, N$, where $Z[i]$ is the i -th element of Z . And use $Z[i]$ to generate the y_i with $\Pr(y_i = 1) = p_i$,
350 $\Pr(y_i = 0) = 1 - p_i$.

We use the set up mentioned above to generate two datasets, one is of size $N = 1000, p = 10000$ the other is of size $N = 10000, p = 1000$.

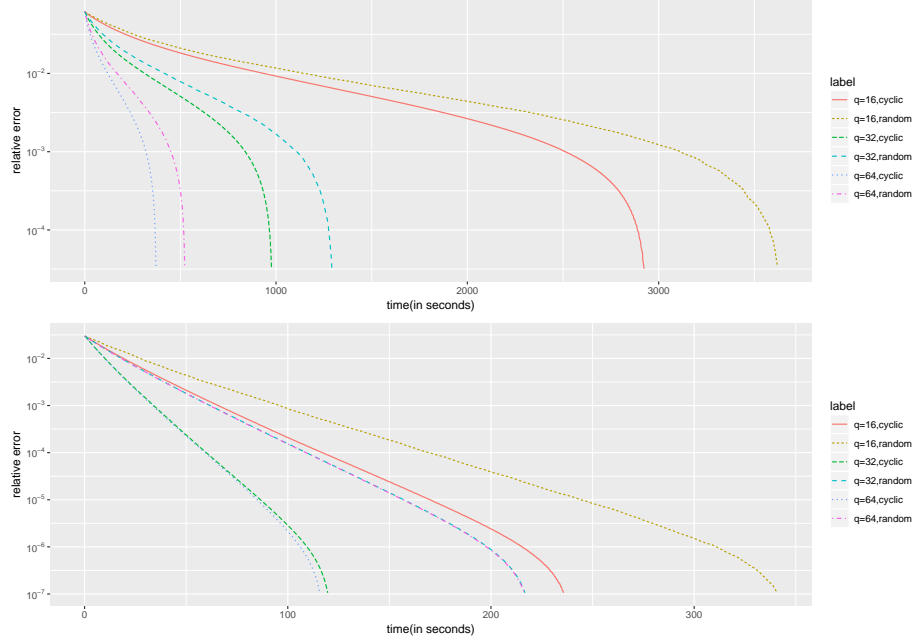


Figure 2: The above figure corresponds to the experiment results on the dataset $N=1000, p=10000$, while the bottom one represents the results of the dataset $N=10000, p=1000$.

We observe two interesting things here.

- Solving the $N > p$ problem is much easier than solving the $p > N$ problem.
- The increase in the number of processors does not guarantee the decrease of time to reach the same global optimal. In fact, after some points, the increase of processors consumes more time to reach the same global optimal. This is as the result of the communication overhead overwhelm the computation time. This observation coincides with results in [3]. It means that though in theory you can use as many as processor as you like, in fact, it does not guarantee the linear computation gain.

4.3. Discussions

In this subsection, we make some remarks about the technique details in designing and implementing our proposed approach.

365 *Jacobi versus Gauss-Seidel.* Many literatures indicate that methods with Jacobi update strategy is usually slower than those with Gauss-Seidel update strategy, e.g. [22]. Here, we also give a simple verification example below, see Figure 3 .

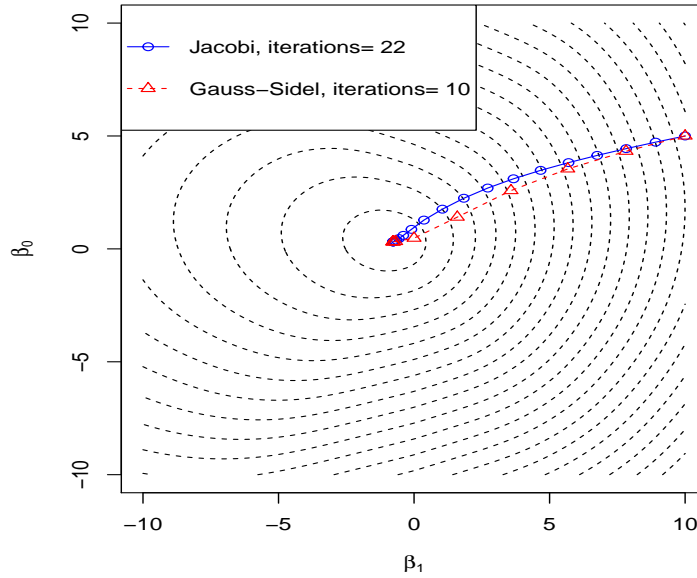


Figure 3: A simple comparison between Jacobi update strategy and Gauss-Seidel update strategy.

Clearly, the iterations of Jacobi style are twice as many as those of Gauss-Seidel's. And it seems that Jacobi strategy is much more conservative than the
 370 Gauss-Seidel strategy. Unfortunately, in the synchronous parallelism setting, the Gauss-Seidel strategy is not a qualified candidate. However, in the asynchronous parallelism setting, it's worth considering the Gauss-Seidel strategy. In [17], Liu et al. propose the "inconsistent" read model, which is really close

to the real asynchronous computing. The only defect in their algorithm is that
 375 it has to first wait for all processors to finish updating assigned coordinate and
 make current iterate information available for all processors, then the individual
 coordinate can go on its work. For detailed discussion, see [17]. Hence, to over-
 come this shortcoming, we wonder if it is possible to achieve true asynchronous
 parallelism by canceling the "synchronization" step in [17]. We shall dive into
 380 it and provide some theoretical analysis.

Smaller c. In Problem (1.1), x_i is updated by solving the following problem

$$\arg \min_{z_i \in X_i} u_i(z_i; x^t) + \lambda h_i(z_i) + c \|z_i - x_i^t\|^2.$$

As indicated in the Theorem3.1 and Theorem3.2, there's a lower bound \bar{c} on
 the parameter c to guarantee the theoretical convergence of our proposed PSUM
 and RPSUM algorithms. However, we observed in our numerical experiments
 385 that we can choose a c' , which is a little bit smaller than the \bar{c} , to reach the
 same optimal solution with fewer iterations and less time.

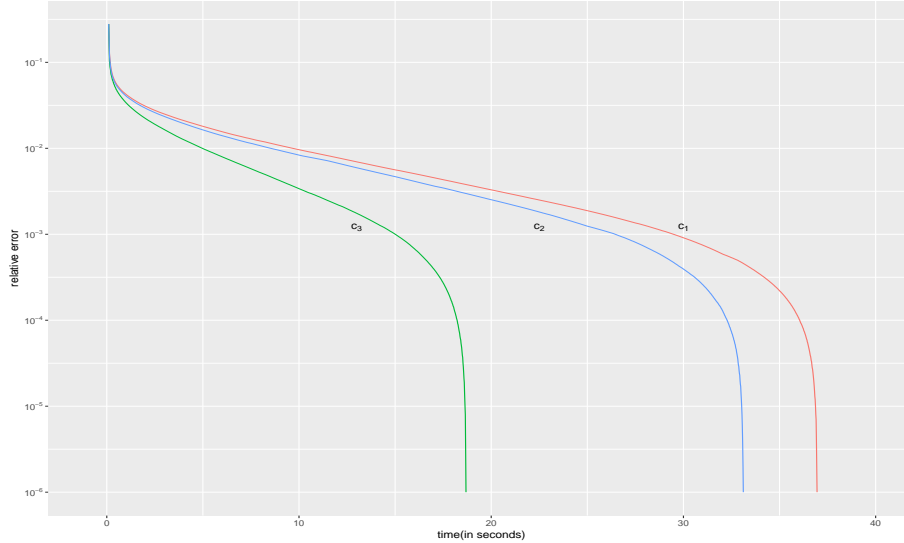


Figure 4: Different c results in different number of iterations to reach the same global optimal solution.

This phenomenon can be explained by the fact that the instructed function $u_i(z_i; x^t) + \lambda h_i(z_i) + c||z_i - x_i^t||^2$ is not a locally tight upper bound function of the $f(z)$. The smaller c' is, the upper approximating function approximates $f(z)$ better. To visualize this assertion, please see Figure 5.

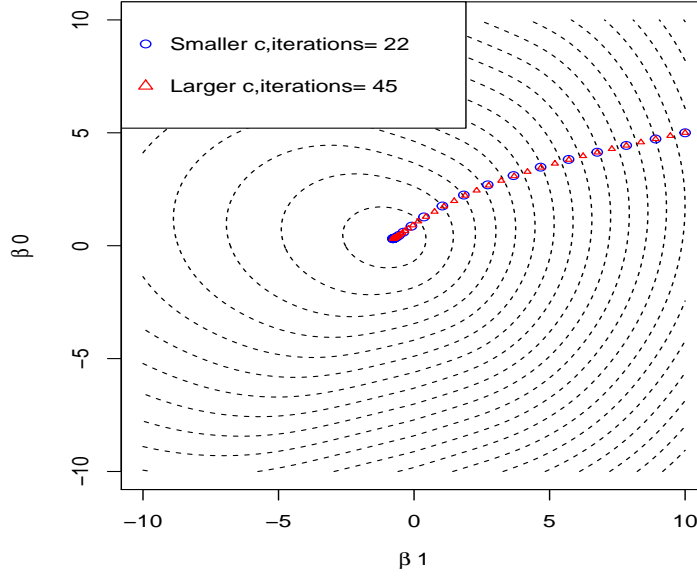


Figure 5: An illustration for the role c plays in our proposed algorithms.

It can be seen from this figure that, the smaller c' results an aggressive stride, hence fewer iterations while the larger c will produce a conservative step-size . It's also worth pointing out that c can not be too small, otherwise the PSUM and RPSUM algorithms will diverge.

Comparisons with other algorithms. Admittedly, we do not compare our algorithms with current existing algorithms because it is somewhat unfair to make comparisons when the numerical implementation details and source code of other algorithms are unknown, like algorithms in [3]. Since we cannot reproduce some results, we do not think it is appropriate to put the results here. Indeed,

400 authors in [3] only report their own algorithms' results for the similar reasons.
 Meanwhile, our algorithms do not address sparse cases in particular, hence some
 greedy methods can outperform our general methods in well-designed problems.

5. Conclusions

The fundamental problem in many application scenarios is how to minimize a
 405 non-smooth convex composite function in an efficient manner. A family of pop-
 ular solvers for this problem are serial block coordinate descent (BCD) but they
 are not efficient in many practical applications. Thanks to the development of
 multi-core parallel processing technology, it makes the speed boosting feasible
 by parallelizing these BCD-like algorithms. Towards this end, we propose par-
 410 allel block coordinate descent algorithms called PSUM and RPSUM, which are
 endowed with sub-linear convergence rate for general convex composite function.
 Different from existing algorithms, the proposed PSUM and RPSUM algorithms
 employ general approximation function to optimize at each iteration. Therefore,
 some existing synchronous parallel algorithms can be treated as special cases of
 415 the PSUM. The experiments on solving the regularized logistic regression with
 ℓ_1 norm problem show that the proposed parallel algorithms are quite efficient
 to implement.

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