
Accelerated Coordinate Descent with Arbitrary Sampling and Best Rates for Minibatches

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Abstract

Accelerated coordinate descent is a widely popular optimization algorithm due to its efficiency on large-dimensional problems. It achieves state-of-the-art complexity on an important class of empirical risk minimization problems. In this paper we design and analyze an accelerated coordinate descent (ACD) method which in each iteration updates a random subset of coordinates according to an arbitrary but fixed probability law, which is a parameter of the method. While minibatch variants of ACD are more popular and relevant in practice, there is no importance sampling for ACD that outperforms the standard uniform minibatch sampling. Through insights enabled by our general analysis, we design new importance sampling for minibatch ACD which significantly outperforms previous state-of-the-art minibatch ACD in practice. We prove a rate that is at most $\mathcal{O}(\sqrt{\tau})$ times worse than the rate of minibatch ACD with uniform sampling, but can be $\mathcal{O}(n/\tau)$ times better, where τ is the minibatch size. Since in modern supervised learning training systems it is standard practice to choose $\tau \ll n$, and often $\tau = \mathcal{O}(1)$, our method can lead to dramatic speedups. Lastly, we obtain similar results for minibatch nonaccelerated CD as well, achieving improvements on previous best rates.

1 Introduction

Many key problems in machine learning and data science are routinely modeled as optimization problems

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and solved via optimization algorithms. With the increase of the volume of data used to formulate optimization models, there is a need for new efficient algorithms able to cope with the challenge. Through intensive research and algorithmic innovation during the last 10-15 years, gradient methods have become the methods of choice for large-scale optimization problems.

In this paper we consider the optimization problem

$$\min_{x \in \mathbb{R}^n} f(x), \quad (1)$$

where f a smooth and strongly convex function, and the main difficulty comes from the dimension n being very large (e.g., millions or billions). In this regime, *coordinate descent* (CD) variants of gradient methods are the state of the art.

The simplest variant of CD in each iterations updates a single variable of x by taking a one dimensional gradient step along the direction of i th unit basis vector $e_i \in \mathbb{R}^n$, which leads to the update rule

$$x^{k+1} = x^k - \alpha_i \nabla_i f(x^k) e_i \quad (2)$$

where $\nabla_i f(x^k) := e_i^\top \nabla f(x^k)$ is the i th partial derivative and α_i is a suitably chosen stepsize. The classical smoothness assumption used in the analysis of CD methods (Nesterov, 2012a) is to require the existence of constants $L_i > 0$ such that

$$f(x + te_i) \leq f(x) + t \nabla_i f(x) + \frac{L_i}{2} t^2 \quad (3)$$

holds for all $x \in \mathbb{R}^n$, $t \in \mathbb{R}$ and $i \in [n] := \{1, 2, \dots, n\}$. In this setting, one can choose the stepsizes to be $\alpha_i = 1/L_i$.

There are several rules studied in the literature for choosing the coordinate i in iteration k , including cyclic rule (Luo and Tseng, 1992; Tseng, 2001; Saha and Tewari, 2013; Wright, 2015; Gurbuzbalaban et al., 2017), Gauss-Southwell or other greedy rules (Nutini et al., 2015; You et al., 2016; Stich et al., 2017a), random (stationary) rule (Nesterov, 2012a; Richtárik and Takáč, 2014, 2016; Shalev-Shwartz and Zhang, 2014;

Lin et al., 2014; Fercoq and Richtárik, 2015) and adaptive random rules (Csiba et al., 2015; Stich et al., 2017b). In this work we focus on stationary random rules, which are popular by practitioners and well understood in theory.

Updating one coordinate at a time. The simplest randomized CD method of the form (2) chooses coordinate i in each iteration uniformly at random. If f is σ -convex¹, then this method converges in $(n \max_i L_i / \sigma) \log(1/\epsilon)$ iterations in expectation. If index i is chosen with probability $p_i \propto L_i$, then the iteration complexity improves to $(\sum_i L_i / \sigma) \log(1/\epsilon)$. The latter result is always better than the former, and can be up to n times better. These results were established in a seminal paper by Nesterov (2012a). The analysis was later generalized to arbitrary probabilities $p_i > 0$ by Richtárik and Takáč (2014), who obtained the complexity

$$\left(\max_i \frac{L_i}{p_i \sigma} \right) \log \frac{1}{\epsilon}. \quad (4)$$

Clearly, (4) includes the previous two results as special cases. Note that the importance sampling $p_i \propto L_i$ minimizes the complexity bound (4) and is therefore in this sense optimal.

Minibatching: updating more coordinates at a time. In many situations it is advantageous to update a small *subset* (*minibatch*) of coordinates in each iteration, which leads to the *minibatch CD method* which has the form

$$x_i^{k+1} = \begin{cases} x_i^k - \alpha_i \nabla_i f(x^k) & i \in S^k, \\ x_i^k & i \notin S^k. \end{cases} \quad (5)$$

For instance, it is often equally easy to fetch information about a small batch of coordinates S^k from memory at the same or comparable time as it is to fetch information about a single coordinate. If this memory access time is the bottleneck as opposed to computing the actual updates to coordinates $i \in S^k$, then it is more efficient to update all coordinates belonging to the minibatch S^k . Alternatively, in situations where parallel processing is available, one is able to compute the updates to a small batch of coordinates simultaneously, leading to speedups in wall clock time. With this application in mind, minibatch CD methods are also often called *parallel CD methods* (Richtárik and Takáč, 2016).

¹We say that f is σ -convex if it is strongly convex with strong convexity modulus $\sigma > 0$. That is, if $f(x+h) \geq f(x) + (\nabla f(x))^T h + \frac{\sigma}{2} \|h\|^2$ for all $x, h \in \mathbb{R}^n$, where $\|h\| := (\sum_i h_i^2)^{1/2}$ is the standard Euclidean norm.

2 Arbitrary sampling and minibatching

Arbitrary sampling. Richtárik and Takáč (2016) analyzed method (5) for *uniform samplings* S^k , i.e., assuming that $\mathbb{P}(i \in S^k) = \mathbb{P}(j \in S^k)$ for all i, j . However, the ultimate generalization is captured by the notion of *arbitrary sampling* pioneered by Richtárik and Takáč (2016b). A *sampling* refers to a set-valued random mapping S with values being the subsets of $[n]$. The word *arbitrary* refers to the fact that no additional assumptions on the sampling, such as uniformity, are made. This result generalizes the results mentioned above.

M-smoothness. For minibatch CD methods it is useful to assume a more general notion of smoothness parameterized by a positive semidefinite matrix $\mathbf{M} \in \mathbb{R}^{n \times n}$. We say that f is **M-smooth** if

$$f(x+h) \leq f(x) + \nabla f(x)^T h + \frac{1}{2} h^T \mathbf{M} h \quad (6)$$

for all $x, h \in \mathbb{R}^n$. The standard L -smoothness condition is obtained in the special case when $\mathbf{M} = L\mathbf{I}$, where \mathbf{I} is the identity matrix in \mathbb{R}^n . Note that if f is **M-smooth**, then (3) holds for $L_i = \mathbf{M}_{ii}$. Conversely, it is known that if (3) holds, then (6) holds for $\mathbf{M} = n \mathbf{Diag}(L_1, L_2, \dots, L_n)$ (Nesterov, 2012a). If h has at most ω nonzero entries, then this result can be strengthened and (6) holds with $\mathbf{M} = \omega \mathbf{Diag}(L_1, L_2, \dots, L_n)$ (Richtárik and Takáč, 2016, Theorem 8). In many situations, **M-smoothness** is a very natural assumption. For instance, in the context of empirical risk minimization (ERM), which is a key problem in supervised machine learning, f is of the form $f(x) = \frac{1}{m} \sum_{i=1}^m f_i(\mathbf{A}_i x) + \frac{\sigma}{2} \|x\|^2$, where $\mathbf{A}_i \in \mathbb{R}^{q \times n}$ are data matrices, $f_i : \mathbb{R}^q \rightarrow \mathbb{R}$ are loss functions and $\sigma \geq 0$ is a regularization constant. If f_i is convex and γ_i -smooth, then f is σ -convex and **M-smooth** with $\mathbf{M} = (\frac{1}{m} \sum_i \gamma_i \mathbf{A}_i^T \mathbf{A}_i) + \sigma \mathbf{I}$ (Qu and Richtárik, 2016). In these situations it is useful to design CD algorithms making full use of the information contained in the data as captured in the smoothness matrix \mathbf{M} .

Given a sampling S and **M-smooth** function f , let $v = (v_1, \dots, v_n)$ be positive constants satisfying the ESO (expected separable overapproximation) inequality

$$\mathbf{P} \circ \mathbf{M} \preceq \mathbf{Diag}(p_1 v_1, \dots, p_n v_n), \quad (7)$$

where \mathbf{P} is the *probability matrix* associated with sampling S , defined by $\mathbf{P}_{ij} := \mathbb{P}(i \in S \ \& \ j \in S)$, $p_i := \mathbf{P}_{ii} = \mathbb{P}(i \in S)$ and \circ denotes the Hadamard (i.e., elementwise) product of matrices. From now on we define the *probability vector* as $p := (p_1, \dots, p_n) \in \mathbb{R}^n$ and let $v = (v_1, \dots, v_n) \in \mathbb{R}^n$ be the vector of ESO parameters. With this notation, (7) can be equivalently

written as $\mathbf{P} \circ \mathbf{M} \preceq \mathbf{Diag}(p \circ v)$. We say that S is *proper* if $p_i > 0$ for all i .

It can be show by combining the results of Richtárik and Takáč (2016b) and Qu and Richtárik (2016) that under the above assumptions, the minibatch CD method (5) with stepsizes $\alpha_i = 1/v_i$ enjoys the iteration complexity

$$\left(\max_i \frac{v_i}{p_i \sigma} \right) \log \frac{1}{\epsilon}. \quad (8)$$

Since in situations when $|S^k| = 1$ with probability 1 once can choose $v_i = L_i$, the complexity result (8) generalizes (4). Inequality (7) is standard in minibatch coordinate descent literature. It was studied extensively by Qu and Richtárik (2016), and has been used to analyze parallel CD methods (Richtárik and Takáč 2016; Richtárik and Takáč, 2016b; Fercoq and Richtárik, 2015), distributed CD methods (Richtárik and Takáč, 2016a; Fercoq et al., 2014), accelerated CD methods (Fercoq and Richtárik, 2015; Fercoq et al., 2014; Qu and Richtárik, 2016; Chambolle et al., 2017), and dual methods (Qu et al., 2015; Chambolle et al., 2017).

Importance sampling for minibatches. It is easy to see, for instance, that if we do not restrict the class of samplings over which we optimize, then the trivial *full sampling* $S^k = [n]$ with probability 1 is optimal. For this sampling, \mathbf{P} is the matrix of all ones, $p_i = 1$ for all i , and (7) holds for $v_i = L := \lambda_{\max}(\mathbf{M})$ for all i . The minibatch CD method (5) reduces to gradient descent, and the complexity estimate (8) becomes $(L/\sigma) \log(1/\epsilon)$, which is the standard rate of gradient descent. However, typically we are interested in finding the best sampling from the class of samplings which use a minibatch of size τ , where $\tau \ll n$. While we have seen that the importance sampling $p_i = L_i / \sum_j L_j$ is optimal for $\tau = 1$, in the minibatch case $\tau > 1$ the problem of determining a sampling which minimizes the bound (8) is much more difficult. For instance, Richtárik and Takáč (2016b) consider a certain parametric family of samplings where the problem of finding the best sampling from this family reduces to a linear program.

Surprisingly, and in contrast to the situation in the $\tau = 1$ case where an optimal sampling is known and is in general non-uniform, there is no minibatch sampling that is guaranteed to outperform τ -nice sampling. We say that S is τ -nice if it samples uniformly from among all subsets of $[n]$ of cardinality τ . The probability matrix of this sampling is given by $\mathbf{P} = \frac{\tau}{n} ((1 - \beta)\mathbf{I} + \beta\mathbf{E})$, where $\beta = \frac{\tau-1}{n-1}$ (assume $n > 1$) and \mathbf{E} is the matrix of all ones, and $p_i = \frac{\tau}{n}$ (Qu and Richtárik, 2016). It follows that the ESO inequality (7) holds for $v_i = (1 - \beta)\mathbf{M}_{ii} + \beta L$. By plugging

into (8), we get the iteration complexity

$$\frac{n}{\tau} \left(\frac{(1 - \beta) \max_i \mathbf{M}_{ii} + \beta L}{\sigma} \right) \log \frac{1}{\epsilon}. \quad (9)$$

This rate interpolates between the rate of CD with uniform probabilities (for $\tau = 1$) and the rate of gradient descent (for $\tau = n$).

3 Contributions

For *accelerated coordinate descent (ACD)* without minibatching (i.e., when $\tau = 1$), the currently best known iteration complexity result, due to Allen-Zhu et al. (2016), is

$$\mathcal{O} \left(\frac{\sum_i \sqrt{L_i}}{\sqrt{\sigma}} \log \frac{1}{\epsilon} \right). \quad (10)$$

The probabilities used in the algorithm are proportional to the square roots of the coordinate-wise Lipschitz constants: $p_i \propto \sqrt{L_i}$. This is the first CD method with a complexity guarantee which does not explicitly depend on the dimension n , and is an improvement on the now-classical result of Nesterov (2012a) giving the complexity

$$\mathcal{O} \left(\sqrt{\frac{n \sum_i L_i}{\sigma}} \log \frac{1}{\epsilon} \right).$$

The rate (10) is always better than this, and can be up to \sqrt{n} times better if the distribution of L_i is extremely non-uniform. Unlike in the non-accelerated case described in the previous section, there is no complexity result for ACD with general probabilities such as (4), or with an arbitrary sampling such as (8). In fact, an ACD method was not even designed in such settings, despite a significant recent development in accelerated coordinate descent methods (Nesterov, 2012b; Lee and Sidford, 2013; Lin et al., 2014; Qu and Richtárik, 2016; Allen-Zhu et al., 2016).

Our key contributions are:

◊ **ACD with arbitrary sampling.** We design an ACD method which is able to operate with an *arbitrary sampling* of subsets of coordinates. We describe our method in Section 4.

◊ **Iteration complexity.** We prove (see Theorem 4.2) that the iteration complexity of ACD is

$$\mathcal{O} \left(\sqrt{\max_i \frac{v_i}{p_i^2 \sigma}} \log \frac{1}{\epsilon} \right), \quad (11)$$

where v_i are ESO parameters given by (7) and $p_i > 0$ is the probability that coordinate i belongs to the sampled set S^k : $p_i := \mathbb{P}(i \in S^k)$. The result of Allen-Zhu et al. (10) (NUACDM) can be recovered as a special case of (11) by focusing on samplings defined by $S^k = \{i\}$ with

Table 1: Complexity results for non-accelerated (CD) and accelerated (ACD) coordinate descent methods for σ -convex functions and arbitrary sampling S . The last row corresponds to the setup with arbitrary proper sampling S (i.e., a random subset of $[n]$ with the property that $p_i := \mathbb{P}(i \in S) > 0$). We let $\tau := \mathbf{E}[|S|]$ be the expected mini-batch size. We assume that f is \mathbf{M} -smooth (see (6)). The positive constants v_1, v_2, \dots, v_n are the ESO parameters (depending on f and S), defined in (7). The first row arises as a special of the third row in the non-minibatch (i.e., $\tau = 1$) case. Here we have $v_i = L_i := \mathbf{M}_{i,i}$. The second row is a special case of the first row for the optimal choice of the probabilities p_1, p_2, \dots, p_n .

	CD	ACD
$\tau = 1, p_i > 0$	$\left(\max_i \frac{L_i}{p_i \sigma}\right) \log \frac{1}{\epsilon}$ (Richtárik and Takáč, 2014)	$\sqrt{\max_i \frac{L_i}{p_i^2 \sigma}} \log \frac{1}{\epsilon}$ (this paper)
$\tau = 1, \text{ best } p_i$	$\frac{\sum_i L_i}{\sigma} \log \frac{1}{\epsilon}; \quad p_i \propto L_i$ (Nesterov, 2012a)	$\frac{\sum_i \sqrt{L_i}}{\sqrt{\sigma}} \log \frac{1}{\epsilon}; \quad p_i \propto \sqrt{L_i}$ (Allen-Zhu et al., 2016)
arbitrary sampling S	$\left(\max_i \frac{v_i}{p_i \sigma}\right) \log \frac{1}{\epsilon}$ (Richtárik and Takáč, 2016b)	$\sqrt{\max_i \frac{v_i}{p_i^2 \sigma}} \log \frac{1}{\epsilon}$ (this paper)

probability $p_i \propto \sqrt{L_i}$ (recall that in this case $v_i = L_i$). When $S^k = [n]$ with probability 1, then our method reduces to accelerated gradient descent (AGD) (Nesterov, 1983, 2004), and since $p_i = 1$ and $v_i = L$ (the Lipschitz constant of ∇f) for all i , (11) reduces to the standard complexity of AGD: $\mathcal{O}(\sqrt{L/\sigma} \log(1/\epsilon))$.

◇ **Weighted strong convexity.** We prove a slightly more general result than (11) in which we allow the strong convexity of f to be measured in a weighted Euclidean norm with weights v_i/p_i^2 . In situations when f is naturally strongly convex with respect to a weighted norm, this more general result will typically lead to a better complexity result than (11), which is fine-tuned for standard strong convexity. There are applications when f is naturally a strongly convex with respect to some weighted norm (Allen-Zhu et al., 2016).

◇ **Minibatch methods.** We design several *new* importance samplings for minibatches, calculate the associated complexity results, and show through experiments that they significantly outperform the standard uniform samplings used in practice and constitute the state of the art. Our importance sampling leads to rates which are provably within a small factor from the best known rates, but can lead to an improvement by a factor of $\mathcal{O}(n)$. We are the first to establish such a result, both for CD (Appendix B) and ACD (Section 5).

The key complexity results obtained in this paper are summarized and compared to prior results in Table 1.

4 The algorithm

The accelerated coordinate descent method (ACD) we propose is formalized as Algorithm 1. If we removed (13) and (16) from the method, and replaced y^{k+1} in (15) by x^{k+1} , we would recover the CD method. Acceleration is obtained by the inclusion of the extrapolation

steps (13) and (16). As mentioned before, we will analyze our method under a more general strong convexity assumption.

Assumption 4.1. f is σ_w -convex with respect to the $\|\cdot\|_w$ norm. That is,

$$f(x+h) \geq f(x) + \langle \nabla f(x), h \rangle + \frac{\sigma_w}{2} \|h\|_w^2, \quad (12)$$

for all $x, h \in \mathbb{R}^n$, where $\sigma_w > 0$.

Note that if f is σ -convex in the standard sense (i.e., for $w = (1, \dots, 1)$), then f is σ_w -convex for any $w > 0$ with $\sigma_w = \min_i \frac{\sigma}{w_i}$. Considering a general σ_w -convexity allows us to get a tighter convergence rate in some cases (Allen-Zhu et al., 2016).

Algorithm 1 ACD (Accelerated coordinate descent with arbitrary sampling)

Input: i.i.d. proper samplings $S^k \sim \mathcal{D}$; ESO parameters $v \in \mathbb{R}_{++}^n$; $p_i = \mathbb{P}(i \in S^k)$ and $w_i = v_i/p_i^2$ for all $i \in [n]$; strong convexity constant $\sigma_w > 0$; stepsize parameters $\theta \approx 0.618\sqrt{\sigma_w}$ (see (20)) and $\eta = 1/\theta$

Initialize: Initial iterate $y^0 = z^0 \in \mathbb{R}^n$

for $k = 0, 1 \dots$ **do**

$$x^{k+1} = (1 - \theta)y^k + \theta z^k \quad (13)$$

$$\text{Get } S^k \sim \mathcal{D} \quad (14)$$

$$y^{k+1} = x^{k+1} - \sum_{i \in S^k} \frac{1}{v_i} \nabla_i f(x^{k+1}) e_i \quad (15)$$

$$z^{k+1} = \frac{1}{1 + \eta \sigma_w} \left(z^k + \eta \sigma_w x^{k+1} - \sum_{i \in S^k} \frac{\eta}{p_i w_i} \nabla_i f(x^{k+1}) e_i \right) \quad (16)$$

end

Using the tricks developed by Lee and Sidford (2013); Fercoq and Richtárik (2015); Lin et al. (2014), Algo-

gorithm [1](#) can be implemented so that only $|S^k|$ coordinates are updated in each iteration. We are now ready to derive a convergence rate of ACD.

Theorem 4.2 (Convergence of ACD). *Let S^k be i.i.d. proper (but otherwise arbitrary) samplings. Let \mathbf{P} be the associated probability matrix and $p_i := \mathbb{P}(i \in S^k)$. Assume f is \mathbf{M} -smooth (see [6](#)) and let v be ESO parameters satisfying [7](#). Further, assume that f is σ_w -convex (with $\sigma_w > 0$) for*

$$w_i := \frac{v_i}{p_i^2}, \quad i = 1, 2, \dots, n, \quad (17)$$

with respect to the weighted Euclidean norm $\|\cdot\|_w$ (i.e., we enforce Assumption [4.1](#)). Then

$$\sigma_w \leq \frac{\mathbf{M}_{ii} p_i^2}{v_i} \leq p_i^2 \leq 1, \quad i = 1, 2, \dots, n. \quad (18)$$

In particular, if f is σ -convex with respect to the standard Euclidean norm, then we can choose

$$\sigma_w = \min_i \frac{p_i^2 \sigma}{v_i}. \quad (19)$$

Finally, if we choose

$$\begin{aligned} \theta &:= \frac{\sqrt{\sigma_w^2 + 4\sigma_w} - \sigma_w}{2} = \frac{2\sigma_w}{\sqrt{\sigma_w^2 + 4\sigma_w} + \sigma_w} \\ &\geq 0.618\sqrt{\sigma_w} \end{aligned} \quad (20)$$

and $\eta := \frac{1}{\theta}$, then the random iterates of ACD satisfy

$$\mathbb{E}[P^k] \leq (1 - \theta)^k P^0, \quad (21)$$

where $P^k := \frac{1}{\theta^2} (f(y^k) - f(x^*)) + \frac{1}{2(1-\theta)} \|z^k - x^*\|_w^2$ and x^* is the optimal solution of [1](#).

Noting that $1/0.618 \leq 1.619$, as an immediate consequence of [21](#) and [20](#) we get bound

$$k \geq \frac{1.619}{\sqrt{\sigma_w}} \log \frac{1}{\epsilon} \Rightarrow \mathbb{E}[P^k] \leq \epsilon P^0. \quad (22)$$

If f is σ -convex, then by plugging [19](#) into [22](#) we obtain the iteration complexity bound

$$1.619 \cdot \sqrt{\max_i \frac{v_i}{p_i^2 \sigma}} \log \frac{1}{\epsilon}. \quad (23)$$

Complexity [23](#) is our key result (also mentioned in [11](#) and Table [1](#)).

5 Importance sampling for minibatches

Let $\tau := \mathbb{E}[|S^k|]$ be the expected minibatch size. The next theorem provides an insightful lower bound for the complexity of ACD we established, one independent of p and v .

Theorem 5.1 (Limits of minibatch performance). *Let the assumptions of Theorem [4.2](#) be satisfied and let f be σ -convex. Then the dominant term in the rate [23](#) of ACD admits the lower bound*

$$\sqrt{\max_i \frac{v_i}{p_i^2 \sigma}} \geq \frac{\sum_i \sqrt{\mathbf{M}_{ii}}}{\tau \sqrt{\sigma}}. \quad (24)$$

Note that for $\tau = 1$ we have $\mathbf{M}_{ii} = v_i = L_i$, and the lower bound is achieved by using the importance sampling $p_i \propto \sqrt{L_i}$. Hence, this bound gives a limit on how much speedup, compared to the best known complexity in the $\tau = 1$ case, we can hope for as we increase τ . The bound says we can not hope for better than linear speedup in the minibatch size. An analogous result (obtained by removing all the squares and square roots in [24](#)) was established by [Richtárik and Takáč \(2016b\)](#) for CD.

In what follows, it will be useful to write the complexity result [23](#) in a new form by considering a specific choice of the ESO vector v .

Lemma 5.2. *Choose any proper sampling S and let \mathbf{P} be its probability matrix and p its probability vector. Let $c(S, \mathbf{M}) := \lambda_{\max}(\mathbf{P}' \circ \mathbf{M}')$, where $\mathbf{P}' := \mathbf{D}^{-1/2} \mathbf{P} \mathbf{D}^{-1/2}$, $\mathbf{M}' := \mathbf{D}^{-1} \mathbf{M} \mathbf{D}^{-1}$ and $\mathbf{D} := \text{Diag}(p)$. Then the vector v defined by $v_i := c(S, \mathbf{M}) p_i^2$ satisfies the ESO inequality [7](#) and the total complexity [23](#) becomes*

$$1.619 \cdot \frac{\sqrt{c(S, \mathbf{M})}}{\sqrt{\sigma}} \log \frac{1}{\epsilon}. \quad (25)$$

Since $\frac{1}{n} \text{Trace}(\mathbf{P}' \circ \mathbf{M}') \leq c(S, \mathbf{M}) \leq \text{Trace}(\mathbf{P}' \circ \mathbf{M}')$ and $\text{Trace}(\mathbf{P}' \circ \mathbf{M}') = \sum_i \mathbf{P}'_{ii} \mathbf{M}'_{ii} = \sum_i \mathbf{M}'_{ii} = \sum_i \mathbf{M}_{ii} / p_i^2$, we get the bounds:

$$\begin{aligned} \sqrt{\frac{c(S, \mathbf{M})}{\sigma}} \log \frac{1}{\epsilon} &\geq \sqrt{\frac{1}{n} \sum_i \frac{\mathbf{M}_{ii}}{p_i^2 \sigma}} \log \frac{1}{\epsilon} \\ \sqrt{\frac{c(S, \mathbf{M})}{\sigma}} \log \frac{1}{\epsilon} &\leq \sqrt{\sum_i \frac{\mathbf{M}_{ii}}{p_i^2 \sigma}} \log \frac{1}{\epsilon}. \end{aligned} \quad (26)$$

5.1 Sampling 1: standard uniform minibatch sampling (τ -nice sampling)

Let S_1 be the τ -nice sampling. It can be shown (see Lemma [C.3](#)) that $c(S_1, \mathbf{M}) \leq \frac{n^2}{\tau^2} ((1 - \beta) \max_i \mathbf{M}_{ii} + \beta L)$, and hence the iteration complexity [23](#) becomes

$$\mathcal{O} \left(\frac{n}{\tau} \sqrt{\frac{(1 - \beta) \max_i \mathbf{M}_{ii} + \beta L}{\sigma}} \log \frac{1}{\epsilon} \right). \quad (27)$$

This result interpolates between ACD with uniform probabilities (for $\tau = 1$) and accelerated gradient descent (for $\tau = n$). Note that the rate [27](#) is a strict improvement on the CD rate [9](#).

Table 2: New complexity results for ACD with minibatch size $\tau = \mathbb{E}[|S^k|]$ and various samplings (we suppress $\log(1/\epsilon)$ factors in all expressions). Constants: σ = strong convexity constant of f , $L = \lambda_{\max}(\mathbf{M})$, $\beta = (\tau - 1)/(n - 1)$, $1 \leq \gamma \leq \sqrt{n}$, and $\omega \leq \mathcal{O}(\sqrt{\tau})$ (ω can be as small as $\mathcal{O}(\tau/n)$).

Lower bound	$S_1 : p_i = \frac{\tau}{n}$	$S_2 : \frac{p_i^2}{\mathbf{M}_{ii}} \propto 1$	$S_3 : \frac{p_i^2}{\mathbf{M}_{ii}} \propto 1 - p_i$
$\frac{\sum_i \sqrt{\mathbf{M}_{ii}}}{\tau \sqrt{\sigma}}$	$\frac{n\sqrt{(1-\beta)} \max_i \mathbf{M}_{ii} + \beta L}{\tau \sqrt{\sigma}}$	$\frac{\gamma \sum_i \sqrt{\mathbf{M}_{ii}}}{\tau \sqrt{\sigma}}$	$\omega \frac{n\sqrt{(1-\beta)} \max_i \mathbf{M}_{ii} + \beta L}{\tau \sqrt{\sigma}}$
(24)	= uniform ACD for $\tau = 1$ = AGD for $\tau = n$	$\leq \sqrt{n} \times$ lower bound $\bullet \tau \leq \frac{\sum_j \sqrt{\mathbf{M}_{jj}}}{\max_i \mathbf{M}_{ii}}$	\bullet fastest in practice \bullet any τ allowed

5.2 Sampling 2: importance sampling for minibatches

Consider now the sampling S_2 which includes every $i \in [n]$ in S_2 , independently, with probability $p_i = \tau \frac{\sqrt{\mathbf{M}_{ii}}}{\sum_j \sqrt{\mathbf{M}_{jj}}}$. This sampling was not considered in the literature before. Note that $\mathbb{E}[|S_2|] = \sum_i p_i = \tau$. For this sampling, bounds (26) become:

$$\begin{aligned} \sqrt{\frac{c(S, \mathbf{M})}{\sigma}} \log \frac{1}{\epsilon} &\geq \frac{\sum_i \sqrt{\mathbf{M}_{ii}}}{\tau \sqrt{\sigma}} \log \frac{1}{\epsilon} \\ \sqrt{\frac{c(S, \mathbf{M})}{\sigma}} \log \frac{1}{\epsilon} &\leq \frac{\sqrt{n} \sum_i \sqrt{\mathbf{M}_{ii}}}{\tau \sqrt{\sigma}} \log \frac{1}{\epsilon}. \end{aligned} \quad (28)$$

Clearly, with this sampling we obtain an ACD method with complexity within a \sqrt{n} factor from the lower bound established in Theorem 5.1. For $\tau = 1$ we have $\mathbf{P}' = \mathbf{I}$ and hence

$$\begin{aligned} c(S, \mathbf{M}) &= \lambda_{\max}(\mathbf{I} \circ \mathbf{M}') = \lambda_{\max}(\mathbf{Diag}(\mathbf{M}')) \\ &= \max_i \mathbf{M}_{ii}/p_i^2 = \left(\sum_j \sqrt{\mathbf{M}_{jj}}\right)^2. \end{aligned}$$

Thus, the rate of ACD achieves the lower bound in (28) (see also (10)) and we recover the best current rate of ACD in the $\tau = 1$ case, established by Allen-Zhu et al. (2016). However, the sampling has an important limitation: it can be used for $\tau \leq \sum_j \sqrt{\mathbf{M}_{jj}} / \max_i \mathbf{M}_{ii}$ only as otherwise the probabilities p_i exceed 1.

5.3 Sampling 3: another importance sampling for minibatches

Now consider sampling S_3 which includes each coordinate i within S_3 independently, with probability p_i satisfying the relation $p_i^2/\mathbf{M}_{ii} \propto 1 - p_i$. This is equivalent to setting

$$p_i := \frac{2\mathbf{M}_{ii}}{\sqrt{\mathbf{M}_{ii}^2 + 2\mathbf{M}_{ii}\delta} + \mathbf{M}_{ii}}, \quad (29)$$

where δ is a scalar for which $\sum_i p_i = \tau$. This sampling was not considered in the literature before. Probability

vector p was chosen as (29) for two reasons: i) $p_i \leq 1$ for all i , and therefore the sampling can be used for all τ in contrast to S_1 , and ii) we can prove Theorem 5.3.

Let $c_1 := c(S_1, \mathbf{M})$ and $c_3 := c(S_3, \mathbf{M})$. In light of (25), Theorem 5.3 compares S_1 and S_3 and says that ACD with S_3 has at most $\mathcal{O}(\sqrt{\tau})$ times worse rate compared to ACD with S_1 , but has the capacity to be $\mathcal{O}(n/\tau)$ times better. We prove in Appendix B a similar theorem for CD. We stress that, despite some advances in the development of importance samplings for minibatch methods (Richtárik and Takáč, 2016b; Csiba and Richtárik, 2018), S_1 was until now the state-of-the-art in theory for CD. We are the first to give a provably better rate in the sense of Theorem B.3. The numerical experiments show that S_3 consistently outperforms S_1 , and often dramatically so.

Theorem 5.3. *The leading complexity terms c_1 and c_3 of ACD (Algorithm (5)) with samplings S_1 , and S_3 , respectively, defined in Lemma 5.2, compare as follows:*

$$c_3 \leq 2 \frac{(2n - \tau)(n\tau + n - \tau)}{(n - \tau)^2} c_1 = \mathcal{O}(\tau) c_1. \quad (30)$$

Moreover, there exists \mathbf{M} where $c_3 \leq \mathcal{O}(\frac{\tau^2}{n^2}) c_1$.

In real world applications, minibatch size τ is limited by hardware and in typical situations, one has $\tau \ll n$, oftentimes $\tau = \mathcal{O}(1)$. The importance of Theorem 5.3 is best understood from this perspective.

6 Experiments

We perform extensive numerical experiments to justify that minibatch ACD with importance sampling works well in practice. Here we present a few selected experiment only; more can be found in Appendix D.

In most of plots we compare of both accelerated and non-accelerated CD with all samplings S_1, S_2, S_3 introduced in Sections 5.1, 5.2 and 5.3 respectively. We refer to ACD with sampling S_3 as AN (Accelerated Nonuniform), ACD with sampling S_1 as AU, ACD with sampling S_2 as

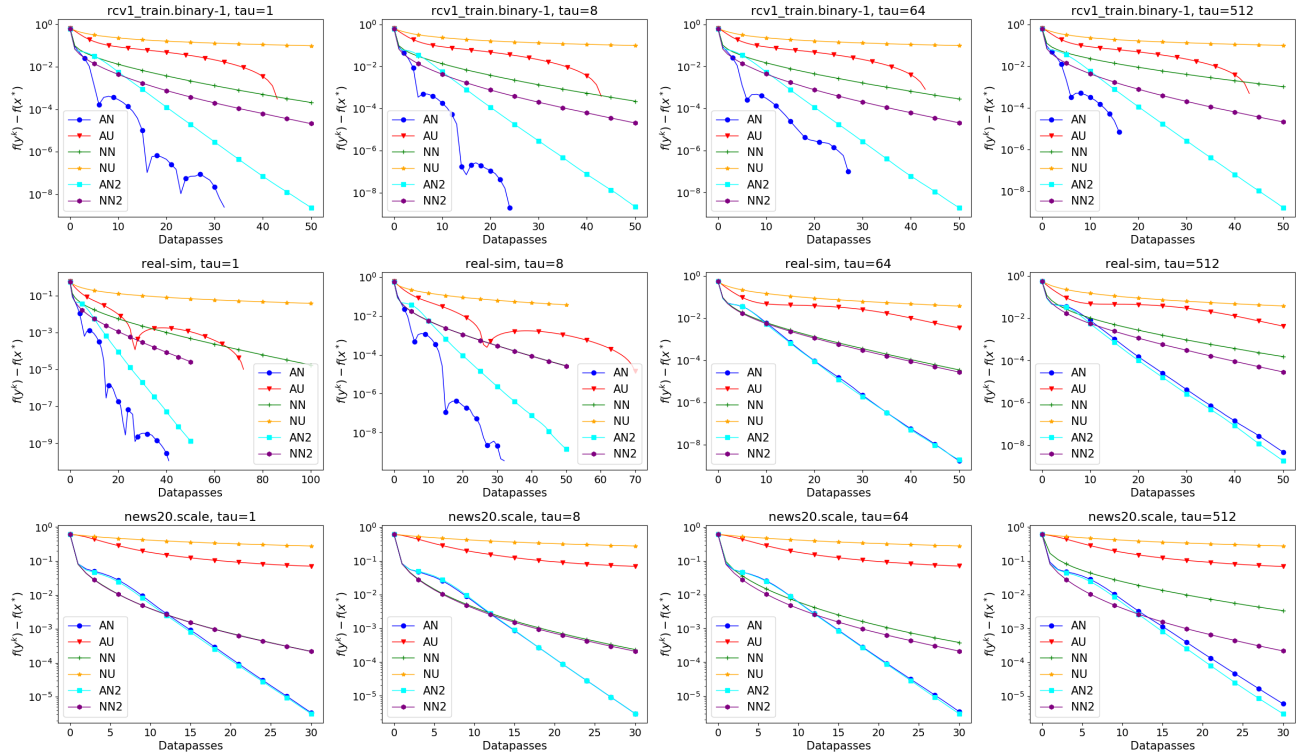


Figure 1: Six variants of coordinate descent (AN, AU, NN, NU, AN2 and AU2) applied to a logistic regression problem, with minibatch sizes $\tau = 1, 8, 64$ and 512 .

AN2, CD with sampling S_3 as NN, CD with sampling S_1 as NU and CD with sampling S_2 as NN2. We compare the methods for various choices of the expected minibatch sizes τ and on several problems.

In Figure 1, we report on a logistic regression problem with a few selected LibSVM [Chang and Lin (2011)] datasets. For larger datasets, pre-computing both strong convexity parameter σ and v may be expensive (however, recall that for v we need to tune only one scalar). Therefore, we choose ESO parameters v from Lemma 5.2, while estimating the smoothness matrix as $10 \times$ its diagonal. An estimate of the strong convexity σ for acceleration was chosen to be the minimal diagonal element of the smoothness matrix. We provide a formal formulation of the logistic regression problem, along with more experiments applied to further datasets in Appendix D.2, where we choose v and σ in full accord with the theory.

Coordinate descent methods which allow for separable proximal operator were proven to be efficient to solve ERM problem, when applied on dual (Shalev-Shwartz and Tewari, 2011; Shalev-Shwartz and Zhang, 2013, 2014; Zhao and Zhang, 2015). Although we do not develop proximal methods in this paper, we empirically demonstrate that ACD allows for this extension as well. As a specific problem to solve, we choose dual of

SVM with hinge loss. Figure 2 presents the results. A detailed description of the experiment is presented in Appendix D.3. The results are indeed in favour of ACD with importance sampling. Therefore, ACD is not only suitable for *big dimensional* problems, it can handle the *big data* setting as well.

Finally, in Appendix D.1 we present several synthetic examples in order to shed more light on acceleration and importance sampling, and to see how its performance depends on the data. We also study how minibatch size influences the convergence rate. All the experimental results clearly show that acceleration, importance sampling and minibatching have a significant impact on practical performance of CD methods. Moreover, the difference in the performance of samplings S_2 and S_3 is negligible, and therefore we recommend using S_3 as it is not limited by the bound on expected minibatch size τ .

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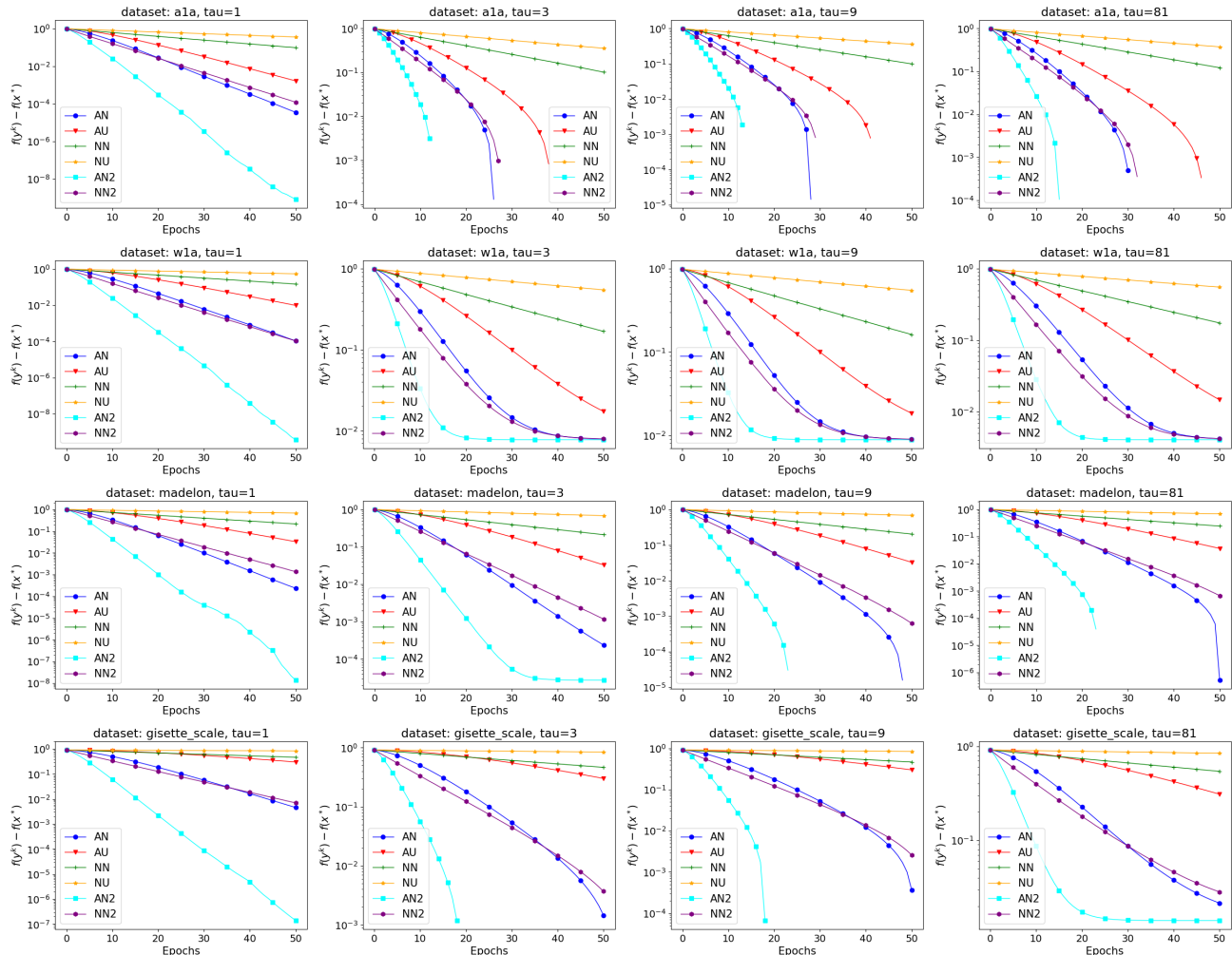


Figure 2: Six variants of coordinate descent (AN, AU, NN, NU, AN2 and AU2) applied to a dual SVM problem, with minibatch sizes $\tau = 1, 2, 8$ and 81 .

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