Smaller Steps for Faster Algorithms: 
A New Approach to Solving Linear Systems

by

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Abstract

In this thesis we study iterative algorithms with simple sublinear time update steps, and we show how a mix of data structures, randomization, and results from numerical analysis allow us to achieve faster algorithms for solving linear systems in a variety of different regimes.

First we present a simple combinatorial algorithm for solving symmetric diagonally dominant (SDD) systems of equations that improves upon the best previously known running time for solving such systems in the standard unit-cost RAM model. Then we provide a general method for convex optimization that improves this simple algorithm’s running time as special case. Our results include the following:

- We achieve the best known running time of $O(m \log^{3/2} n \sqrt{\log \log n} \log(\varepsilon^{-1} \log n))$ for solving Symmetric Diagonally Dominant (SDD) system of equations in the standard unit-cost RAM model.
- We obtain a faster asymptotic running time than conjugate gradient for solving a broad class of symmetric positive definite systems of equations.
- We achieve faster asymptotic convergence rates than the best known for Kaczmarz methods for solving overdetermined systems of equations, by accelerating an algorithm of Strohmer and Vershynin [55].

Beyond the independent interest of these solvers, we believe they highlight the versatility of the approach of this thesis and we hope that they will open the door for further algorithmic improvements in the future.

This work was done in collaboration with Jonathan Kelner, Yin Tat Lee, Lorenzo Orecchia, and Zeyuan Zhu, and is based on the content of [30] and [35].

Thesis Supervisor: Jonathan Kelner  
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Chapter 1

Introduction

In recent years iterative methods for convex optimization that make progress in sublinear time using only partial information about a function and its gradient have become of increased importance to both the theory and practice of computer science. From a practical perspective, the increasing volume and distributed nature of data are forcing efficient practical algorithms to be amenable to asynchronous and parallel settings where only a subset of the data is available to a single processor at any point in time. From a theoretical perspective, rapidly converging algorithms with sublinear time update steps create the hope for new faster algorithms for solving old problems.

In this thesis we present several results showing that by combining simple sublinear steps with a mix of data structures, randomization, and results from numerical analysis one can achieve faster and simpler algorithms for solving a wide array of linear systems. In the first half of this thesis we show how to use these techniques to produce a simple combinatorial algorithm that solves symmetric diagonally dominant (SDD) systems of equations in nearly linear time and improves upon the best previously known running times for solving SDD systems in the unit-cost RAM model.

In the second half of this thesis we generalize and improve upon the iterative framework applied by this SDD solver. We show to obtain a general first order method for convex optimization that can solve a broad class of linear systems faster than conjugate gradient, generically improve the convergence rate of randomized Kaczmarz [55], and obtain a faster SDD solver in the unit-cost RAM model.
1.1 SDD Systems

While the results in this thesis apply to general problems in convex optimization, they were motivated by the desire to achieve faster and simpler algorithms for solving symmetric diagonal dominant (SDD) systems of equations. A matrix \( A \in \mathbb{R}^{n \times n} \) is SDD if \( A^T = A \) and \( A_{ii} \geq \sum_{j \neq i} |A_{ij}| \) for all \( i \in [n] \). While the best known algorithm for solving a general linear system takes time \( O(n^{2.373}) \) [61], a seminal paper by Spielman and Teng [50] showed that when \( A \) is SDD one can solve \( A\vec{x} = \vec{b} \) approximately in nearly linear time. \(^1\)

Fast algorithms for solving SDD linear systems have found broad applications across both the theory and practice of computer science. They have long been central to scientific computing, where solving SDD systems is the main computational task in modeling of electrical networks of resistors and performing finite element simulations of a wide range of physical systems (see, e.g., [12]). Beyond this, SDD system solvers have been applied to foundational problems in a wide range of other fields, including machine learning, random processes, computer vision, image processing, network analysis, and computational biology (see, for example, [36, 33, 37, 60, 22]).

More recently, SDD solvers have emerged as a powerful tool in the design of graph algorithms. To every graph \( G \), one can associate a SDD matrix \( L \) called its Laplacian (see Section 2.2) such that there are deep connections between the combinatorial properties of \( G \) and the linear algebraic properties of \( L \). By exploiting these connections, researchers have used nearly linear time algorithms for solving SDD systems to break longstanding barriers and provide new algorithms for a rapidly growing list of fundamental graph problems, including maximum flow problems [16], multi-commodity flow problems [28], generating random spanning tree [27], graph sparsification [48], lossy flow problems [17], sparsest cut [46], distributed routing [26], and balanced separator [44], as well as fundamental linear algebraic problems for SDD.

---

\(^1\)Throughout this thesis we are primarily interested in approximate linear system solvers, that is algorithms that compute \( \vec{x} \in \mathbb{R}^n \) such that \( \|\vec{x} - \vec{x}_{opt}\|_A \leq \varepsilon \|\vec{x}_{opt}\|_A \) for any \( \varepsilon \in \mathbb{R} > 0 \) where \( \vec{x}_{opt} \in \mathbb{R}^n \) is a vector such that \( A\vec{x}_{opt} = \vec{b} \). By a nearly linear time SDD system solver we mean an algorithm that computes such a \( \vec{x} \) in time \( O(m \log^c n \log \varepsilon^{-1}) \) where \( m \) is the number of nonzero entries in \( A \) and \( c \geq 0 \in \mathbb{R} \) is a fixed constant.
matrices, including computing the matrix exponential [44] and the largest eigenvalue and corresponding eigenvector [53]. (See [49, 57, 59] for surveys of these solvers and their applications.)

1.1.1 Previous Nearly Linear Time SDD Solvers

The first nearly linear time algorithm for solving SDD systems was given by Spielman and Teng [50], building on a long line of previous work (e.g., [58, 20, 7, 13, 11]). Their algorithm and its analysis is a technical tour-de-force that required multiple fundamental innovations in spectral and combinatorial graph theory, graph algorithms, and computational linear algebra. Their work included the invention of spectral sparsification and ultra-sparsifiers, better and faster constructions of low-stretch spanning trees, and efficient local clustering algorithms, all of which was used to construct and analyze an intricate recursively preconditioned iterative solver. They divided this work into three papers totaling over 130 pages ([51, 53, 52]), each of which has prompted a new line of inquiry and substantial follow-up work. Their work was followed by two beautifully insightful papers by Koutis, Miller, and Peng that simplified the SDD system solver while improving its running time to \( \widetilde{O}(m \log n \log \varepsilon^{-1}) \) [31, 32].

For a more in-depth discussion of the history of this work see [53].

These algorithms all rely on the same general framework. They reduce solving general SDD systems to solving systems in graph Laplacians. Given a graph, they show how to obtain a sequence of logarithmically many successively sparser graphs that approximate it, which they construct by adding carefully chosen sets of edges to a low-stretch spanning tree. They then show that the relationship between the combinatorial properties of a graph and the spectral properties of its Laplacian enables them to use the Laplacian of each graph in this sequence as a preconditioner for the one preceding it in a recursively applied iterative solver, such as the Preconditioned Chebyshev Method or Preconditioned Conjugate Gradient.

We remark that multigrid methods (see, e.g., [15]), which are widely used in

\[ \widetilde{O}(\cdot) \] to hide lower order log terms in \( n \), e.g. \( \widetilde{O}(m) = O(m \log^c n) \) and \( \widetilde{O}(m \log^d n) = O(m \log^c n \log \log^d n) \) for some constants \( c, d \geq 0 \).
practice on graphs with sufficiently nice topologies, can be thought of as following a similar multilevel recursively preconditioned iterative framework. Indeed, Koutis et al. have an algorithm and implementation based on related techniques that they refer to as "combinatorial multigrid" [9]. Even if one does not demand provable running time bounds, we are not aware of any algorithm whose running time empirically scales nearly linearly on large classes of input graphs that does not roughly follow this general structure.\(^3\)

1.2 First Order Methods

The iterative framework applied by all known nearly linear time SDD solvers fall into the broad category of first order methods for convex optimization. That is, they all run combinatorial algorithms to set up convex optimization problems and then they solve these problems by repeatedly computing only values and gradients of the objective function being optimized. In fact, all iterative optimization algorithms in this thesis (including the new one we develop) can be similarly viewed as first order methods.

1.2.1 Previous First Order Methods

Gradient descent is one of the oldest and most fundamental first order methods in convex optimization. Given a convex differentiable function the gradient descent method is a simple greedy iterative method that computes the gradient at the current point and uses that information to perform an update and make progress. This method is central to much of scientific computing and from a theoretical perspective the standard method is well understood [41]. There are multiple more sophisticated variants of this method [25], but many of them have only estimates of local convergence rates which makes them difficult to be applied to theoretical problems and be compared in general.

\(^3\)With the exception of the new algorithms considered in this thesis.
In 1983, Nesterov [40] proposed a way to accelerate the gradient descent method by iteratively developing an approximation to the function through what he calls an *estimate sequence*. This *accelerated gradient descent method* or *fast gradient method* has the same worst case running time as conjugate gradient method and it is applicable to general convex functions. Recently, this method has been used to improve the fastest known running time of some fundamental problems in computer science, such as compressive sensing [5, 6], undirected maximum flow [16, 34, 29], linear programming [42, 8].

The accelerated gradient descent method is known to achieve an optimal (up to constants) convergence rate among all first order methods, that is algorithm that only have access to the function’s value and gradient [41]. Therefore, to further improve accelerated gradient descent one must either assume more information about the function or find a way to reduce the cost of each iteration. Using the idea of fast but crude iteration steps, Nesterov proposed a randomized coordinate descent method [43], which minimizes convex functions by updating one randomly chosen coordinate in each iteration.

Coordinate descent methods, which use gradient information about a single coordinate to update a single coordinate in each iteration, have been around for a long time [62]. Various schemes have been considered for picking the coordinate to update, such as cyclic coordinate update and the best coordinate update, however these schemes are either hard to estimate [38] or difficult to be implemented efficiently. Both the recent work of Strohmer and Vershynin [55] and Nesterov [43] (as well as the algorithms in this thesis) overcame these obstacles by showing that by performing particular randomized updates one can produce methods with provable global convergence rate and small costs per iteration.

Applying the similar ideas of accelerated gradient descent, Nesterov also proposed an accelerated variant called the *accelerated coordinate descent method (ACDM)* that achieves a faster convergence rate while still only considering a single coordinate of the gradient at a time. However, in both Nesterov’s paper [43] and later work [45], this method was considered inefficient as the computational complexity of the
naive implementation of each iteration of ACDM requires $\Theta(n)$ time to update every coordinate of the input, at which point the accelerated gradient descent method would seem preferable.

1.3 Our Results

In this thesis we present both a simple, combinatorial algorithms that solves SDD systems in nearly linear time and a general first order method for convex optimization that improves the running time of this solver as a special case.

The SDD solver uses very little of the machinery that previously appeared to be necessary for a nearly linear time algorithm. It does not require spectral sparsifiers (or variants such as ultra-sparsifiers or incremental sparsifiers), recursive preconditioning, or even the Chebyshev Method or Conjugate Gradient. To solve a SDD system all the algorithm requires is a single low-stretch spanning tree\(^4\) of $G$ (not a recursive collection of subgraphs), and a straightforward data structure. Given these, the algorithm can be described in a few lines of pseudocode, and its analysis can be made to fit on a single blackboard.

We show how to cast this solver as an instance of the coordinate descent algorithm of Nesterov [43] and we provide a a first order method for convex optimization that both strengthens and unifies this result and the randomized Kaczmarz method of Strohmer and Vershynin [55]. In particular, we present a more general version of Nesterov's ACDM and show how to implement it so that each iteration has the same asymptotic runtime as its non-accelerated variants. We show that this method is numerically stable and we show how to use this method to outperform conjugate gradient in solving a general class of symmetric positive definite systems of equations. Furthermore, we show how to cast both randomized Kaczmarz and the simple SDD solver in this framework and achieve faster running times through the use of ACDM.\(^5\)

\(^4\)This can be obtained in nearly-linear time by a simple ball-growing algorithm [3]; the constructions with the best known parameters use a more intricate, but still nearly linear time, region growing technique [18, 1, 31, 2]. See [30] for a discussion of how even this requirement can be relaxed.

\(^5\)This algorithm can also be shown to have an asymptotically tight running time in certain regimes and we refer the reader to [35] for a discussion of these lower bounds.
1.3.1 Comparison to SDD Solvers

Previous nearly-linear time SDD solvers relied on Preconditioned Chebyshev methods, whose numerical stability is quite difficult to analyze. At present, the best known results show that they can be implemented with finite-precision arithmetic, but the number of bits of precision required is $\log \kappa(L) \log^c n \log \varepsilon^{-1}$, where $\kappa(L)$ is the condition number of $L$, and $c$ is some possibly large constant [53]. Therefore, while the stated running time of the best SDD solver is $\tilde{O}(m \log n \log \varepsilon^{-1})$ [32], this is assuming arbitrary precision arithmetic. If one analyzes it in the more standard unit-cost RAM model, where one can perform operations only on $O(\log n)$-bit numbers in constant time, this introduces several additional logarithmic factors in the running time.

In contrast, we show that our algorithms are numerically stable and do not require this additional overhead in the bit precision. Our simple algorithm approximately solves both SDD systems and the dual electrical flow problem in time $O(m \log^2 n \log \log n \log (\varepsilon^{-1} n))$ \footnote{For a discussion of how to improve the running time to $O(m \log^2 n \log \log n \log (\varepsilon^{-1} n))$ using a different technique we refer the reader to [30].} and our accelerated solver solves these problems in time $O(m \log^{3/2} n \sqrt{\log \log n \log (\varepsilon^{-1} \log n)})$. As such, our algorithm gives the fastest known algorithm for solving SDD systems in the unit-cost RAM model. If one allows infinite precision arithmetic to be performed in constant time, our fastest algorithm is slower than [32] by a factor of $\tilde{O}(\log n)$ for solving SDD systems and actually faster than [32] for the dual problem of computing $\varepsilon$-approximate electric flows when $\varepsilon$ is constant.\footnote{To the best of our knowledge, to convert the SDD system solver of [32] to an $\varepsilon'$-approximate electrical flow solver, one needs to pick $\varepsilon = O(\varepsilon'^{-1} n)$.}

Due to the complexity of previous nearly linear time solvers and the intricate and delicate nature of their analyses, it was necessary to apply them as a black box. By providing a new, easy-to-understand algorithms, it is our hope that algorithms that use SDD solvers can be improved by "opening up" this black box and modifying it to take advantage of specific features of the problem, and that similar techniques can be applied to related problems (e.g., ones with additional constraints or slight deviations from linearity or diagonal dominance). Furthermore, due to the lightweight nature of
the algorithm and data structure, we hope it to be fast in practice and adaptable to work in multicore, distributed, and even asynchronous settings.

1.3.2 Comparison to General Linear System Solvers

In some sense, the principle difference between the asymptotic running time of the efficient ACDM presented in this thesis and accelerated gradient descent (or conjugate gradient in the linear system case) is that as accelerated gradient descent depends on the maximum eigenvalue of the Hessian of the function being minimized, ACDM instead depends on the trace of the Hessian and has the possibility of each iteration costing a small fraction of the cost a single iteration of accelerated gradient descent. As a result, any nontrivial bound on the trace of the Hessian and the computational complexity of performing a single coordinate update creates the opportunity for ACDM to yield improved running times.

Beyond demonstrating how ACDM yields a faster SDD solver we show that under mild assumptions ACDM solves positive definite systems with a faster asymptotic running time than conjugate gradient (and an even milder set of assumptions for Chebyshev method), and it is an asymptotically optimal algorithm for solving general systems in certain regimes.

Furthermore, consider over-constrained systems of equations where the randomized Kaczmarz method of Strohmer and Vershynin [55], which iteratively picks a random constraint and projects the current solution onto the plane corresponding to a random constraint, has been shown to have strong convergence guarantees and appealing practical performance. We show how to cast this method in the framework of coordinate descent and accelerate it using ACDM yielding improved asymptotic performance. Given the appeal of Kaczmarz methods for practical applications such as image reconstructions [21], there is hope that this could yield improved performance in practice.

We remark that while our analysis of applications of ACDM focus on solving linear systems there is hope that our ACDM algorithm will have a broader impact in both theory and practice of efficient algorithms. Just as the accelerated gradient
descent method has improved the theoretical and empirical running time of various, both linear and nonlinear, gradient descent algorithms [19, 5], we hope that ACDM will improve the running time of various algorithms for which coordinate descent based approaches have proven effective. Given the generality of our analysis and the previous difficulty in analyzing such methods, we hope that this is just the next towards a new class of provably efficient algorithms with good empirical performance.

1.4 Thesis Organization

The rest of this thesis is organized as follows. In Chapter 2 we present and analyze our simplest nearly linear time SDD solver proving that it achieves a running time of $O(m \log^2 n \log \log n \log(\varepsilon^{-1} n))$ in the unit cost RAM model. In Chapter 3 we take a more general approach and present a new first order method, the efficient accelerated coordinate descent method (ACDM) and show how ACDM yields faster algorithms for solving a variety of linear systems. In Appendix A we provide additional proof details for generalizing ACDM and showing its numerical stability.

The material in Chapter 2 is based on joint work with Jonathan Kelner, Lorenzo Orecchia, and Zeyuan Zhu [30] and the material in Chapter 3 and Appendix A is based on joint work with Yin Tat Lee [35].
Chapter 2

A Simple Nearly Linear Time Solver for SDD Systems

In this chapter we present a simple combinatorial algorithm for solving SDD systems in nearly linear time. The algorithm uses very little of the machinery that previously appeared to be necessary for such an algorithm. It does not require recursive preconditioning, spectral sparsification, or even the Chebyshev Method or Conjugate Gradient. After constructing a “nice” spanning tree of a graph associated with the linear system, the algorithms consist of the repeated application of a simple (non-recursive) update rule, which can be implemented using a lightweight data structure. The algorithms is numerically stable and achieves a faster running time in the standard unit-cost RAM model than was known for previous solvers.

The rest of this chapter is organized as follows. In Section 2.1 we provide an overview of our approach, in Section 2.2 we provide technical background that needed for our analysis, in Section 2.3 we present the algorithm, in Section 2.4 we prove the algorithm’s convergence rate of the algorithm, in Section 2.5 we present a data structure allowing us to implement the algorithm efficiently, in Section 2.6 we prove the asymptotic running time of the algorithm, and in Section 2.7 we prove that the algorithm is numerically stable and achieves the desired running time in the unit-cost RAM model.
2.1 Overview of our Approach

Here we provide a basic overview of our simple algorithm in order to motivate the definitions and analysis that follow. Using standard reductions techniques we reduce solving arbitrary SDD systems to solving $L\vec{v} = \vec{x}$, where $L \in \mathbb{R}^{n \times n}$ is the Laplacian of a weighted connected graph $G = (V, E, \vec{w})$ with $n$ vertices and $m$ edges and $\vec{x} \in \mathbb{R}^n$, (see Appendix A of [30] for details). Just to simplify the overview we suppose that $\vec{x} = \vec{e}_s - \vec{e}_t$ and $\vec{e}_s$ and $\vec{e}_t$ are the unit basis vectors corresponding to arbitrary $s, t \in V$.

Such Laplacian systems $L\vec{v} = \vec{x}$ can be viewed as electrical flow problems: each edge $e \in E$ can be viewed as a resistor of resistance $r_e = 1/w_e$, where $w_e$ is the weight of this edge, and one unit of electric current needs to be sent from $s$ to $t$. Now, if $\vec{v}$ is a valid solution to $L\vec{v} = \vec{x}$, then the entries of $\vec{v}$ can be viewed as the electrical potentials of this system, and the amount of electric current or flow $\vec{f}(e)$ on an edge $e = (i, j)$ from $i$ to $j$ is given by $(\vec{v}_i - \vec{v}_j)/r_e$. The fact that electric flows are induced by vertex potential differences is a crucial property of electrical systems that our algorithm is going to exploit.

While previous solvers worked with the potential vector $\vec{v}$, our algorithm works with the flow vector $\vec{f}$. Our algorithm begins with any arbitrary unit $s$-$t$ flow (e.g., a path from $s$ to $t$) and maintains its feasibility throughout the execution. If $\vec{f}$ were a valid electrical flow, then it would be induced by some potential vector $\vec{v} \in \mathbb{R}^V$ satisfying $\vec{f}(e) = (\vec{v}_i - \vec{v}_j)/r_e$ for all edges $e = (i, j) \in E$, and in particular, for any cycle $C$ in $G$, we would have the potential drop $\sum_{e \in C} \vec{f}(e)r_e = 0$. Exploiting this fact, our entire algorithm consists of simply repeating the following two steps.

- Randomly sample a cycle $C$ from some probability distribution.
- Compute $\sum_{e \in C} \vec{f}(e)r_e$ and add a multiple of $C$ to $\vec{f}$ to make it zero.

To turn this into a provably-correct and efficient algorithm we need to do the following:

- **Specify the cycle distribution.** The cycles are those found by adding edges to a low-stretch spanning tree. They are sampled proportional to their stretch (See Section 2.2.2 and 2.3).
• **Bound the number of iterations.** In Section 2.4, we show that repeating this process a nearly linear number of times yields an \( \varepsilon \)-approximate solution.

• **Implement the iterations efficiently.** Since a cycle may contain a large number of edges, we cannot simply update the flow edge-by-edge. In Section 2.5, we give a data structure that allows each iteration to take \( O(\log n) \) time.

While in the next chapter we show how this algorithm can be viewed as an instantiation of coordinate descent, here we note that this algorithm has a geometric interpretation and can be viewed as an application of the **randomized Kaczmarz** method of Strohmer and Vershynin [55] (see Section 3.4), which solves a linear system by randomly picking a constraint and projecting the current solution onto its feasible set. For a more detailed exposition on this fact and discussion of how the algorithm can be viewed as providing a linear approximation to \( L^t \) we refer the reader to [30].

### 2.2 Preliminaries

For the remainder of this chapter we let \( G = (V, E, w) \) be a weighted, connected, undirected graph with \( n = |V| \) vertices, \( m = |E| \) edges and edge weights \( w_e > 0 \). We think of \( w_e \) as the **conductance** of \( e \), and we define the **resistance** of \( e \) by \( r_e \overset{\text{def}}{=} 1/w_e \).

For notational convenience, we fix an orientation of the edges so that for any vertices \( a \) and \( b \) connected by an edge, exactly one of \( (a, b) \in E \) or \( (b, a) \in E \) holds.

We make extensive use of the following matrices associated with \( G \):

**Definition 2.2.1 (Incidence Matrix).** The incidence matrix \( B \in \mathbb{R}^{E \times V} \) is defined by

\[
\forall (a, b) \in E, \; \forall c \in V : \; B_{(a,b),c} = \begin{cases} 
1 & a = c \\
-1 & b = c \\
0 & \text{otherwise}
\end{cases}
\]

**Definition 2.2.2 (Resistance Matrix).** The resistance matrix, \( R \in \mathbb{R}^{E \times E} \), is the diagonal matrix where \( R_{e,e} = r_e \) for all \( e \in E \).
Definition 2.2.3 (Laplacian Matrix). The Laplacian matrix, \( L \in \mathbb{R}^{V \times V} \), is defined for all \( a, b \in V \) by

\[
L_{a,b} = \begin{cases} 
\sum_{\{a,u\} \in E} w_{a,u} & a = b \\
-w_{a,b} & \{a, b\} \in E \\
0 & \text{otherwise}
\end{cases}
\]

For a vector \( \vec{f} \in \mathbb{R}^E \) and an edge \( e = (a, b) \in E \), we write \( \vec{f}(e) = \vec{f}(a, b) \) for the coordinate of \( \vec{f} \) corresponding to \( e \), and we adopt the convention \( \vec{f}(b, a) = -\vec{f}(a, b) \). This allows us to think of \( \vec{f} \) as a flow on the graph (not necessarily obeying any conservation constraints) that sends \( \vec{f}(e) \) units of flow from \( a \) to \( b \), and thus \( -\vec{f}(e) \) units of flow from \( b \) to \( a \).

The following facts follow by simple manipulations of the above definitions:

Claim 2.2.4. For all \( \vec{f} \in \mathbb{R}^E \), \( x \in \mathbb{R}^V \), \( a \in V \) and \( (a, b) \in E \):

- \( [B^T \vec{f}]_a = \sum_{(b,a) \in E} \vec{f}(b, a) - \sum_{(a,b) \in E} \vec{f}(a, b) \),
- \( L = B^T R^{-1} B \),
- \( [Bx]_{(a,b)} = x(a) - x(b) \), and
- \( x^T L x = \sum_{(a,b) \in E} r_{a,b} (x_a - x_b)^2 \).

One can interpret the first assertion in Claim 2.2.4 as saying that \( B^T \vec{f} \) is a vector in \( \mathbb{R}^V \) whose \( a \)-th coordinate indicates how much flow \( \vec{f} \) leaves (or enters, if it is negative) the graph \( G \) at vertex \( a \). We say that \( \vec{f} \) is a circulation if \( B^T \vec{f} = 0 \).

2.2.1 Electrical Flow

For any vector \( \vec{f} \in \mathbb{R}^E \), we define its energy \( \xi(\vec{f}) \) by

\[
\xi(\vec{f}) \overset{\text{def}}{=} \sum_{e \in E} r_e \vec{f}(e)^2 = \vec{f}^T R \vec{f} = \|\vec{f}\|_R^2.
\]

We fix a demand vector \( \vec{\chi} \in \mathbb{R}^V \) and we say a flow \( \vec{f} \in \mathbb{R}^E \) is feasible (with respect to \( \vec{\chi} \)), or that it meets the demands, if \( B^T \vec{f} = \vec{\chi} \). Since \( G \) is connected it is straightforward to check that there exists a feasible flow with respect to \( \vec{\chi} \) if and only if \( \sum_{v \in V} \vec{\chi}(v) = 0 \).
Definition 2.2.5 (Electrical Flow). For a demand vector $\bar{x} \in \mathbb{R}^V$ that satisfies $\sum_{a \in V} \bar{x}(a) = 0$, the electrical flow satisfying $\bar{x}$ is the unique minimizer to

$$\bar{f}_{\text{opt}} \overset{\text{def}}{=} \arg \min_{\mathbf{f} \in \mathbb{R}^E : \mathbf{B}^T \mathbf{f} = \bar{x}} \xi(\mathbf{f}) \ . \quad (2.1)$$

This quadratic program describes a natural physical problem. Given an electric circuit with nodes in $V$, for each undirected edge $e = \{a, b\} \in E$ we connect nodes $a$ and $b$ with a resistor of resistance $r_e$. Next, we fix the amount of current entering and leaving each node and denote this by demand vector $\bar{x}$. Recalling from physics that the energy of sending $i$ units of current over a resistor of resistance $r$ is $i^2 \cdot r$ the amount of electric current on each resistor is given by $\bar{f}_{\text{opt}}$. The central problem of this chapter is to efficiently compute an $\varepsilon$-approximate electrical flow.

Definition 2.2.6 ($\varepsilon$-Approximate Electrical Flow). For $\varepsilon \in \mathbb{R}^\geq 0$, we say $\bar{f} \in \mathbb{R}^E$ is an $\varepsilon$-approximate electric flow satisfying $\bar{x}$ if $\mathbf{B}^T \bar{f} = \bar{x}$, and $\xi(\bar{f}) \leq (1 + \varepsilon) \cdot \xi(\bar{f}_{\text{opt}})$.

Duality

The electric flow problem is dual to solving $\mathbf{L} \bar{x} = \bar{X}$ when $\mathbf{L}$ is the Laplacian for the same graph $G$. To see this, we note that the Lagrangian dual of (2.1) is given by

$$\max_{\bar{v} \in \mathbb{R}^V} 2\bar{v}^T \bar{X} - \bar{v}^T \mathbf{L} \bar{v} \ . \quad (2.2)$$

For symmetry we define the (dual) energy of $\bar{v} \in \mathbb{R}^V$ as $\zeta(\bar{v}) \overset{\text{def}}{=} 2\bar{v}^T \bar{X} - \bar{v}^T \mathbf{L} \bar{v}$. Setting the gradient of (2.2) to 0 we see that (2.2) is minimized by $\bar{v} \in \mathbb{R}^V$ satisfying $\mathbf{L} \bar{v} = \bar{X}$.

Let $\mathbf{L}^\dagger$ denote the Moore-Penrose psuedoinverse of $\mathbf{L}$ and let $\bar{v}_{\text{opt}} \overset{\text{def}}{=} \mathbf{L}^\dagger \bar{X}$ denote a particular set of optimal voltages. Since the primal program (2.1) satisfies Slater's condition, we have strong duality, so for all $\bar{v} \in \mathbb{R}^V$

$$\zeta(\bar{v}) \leq \zeta(\bar{v}_{\text{opt}}) = \xi(\bar{f}_{\text{opt}})$$
Therefore, for feasible $\vec{f} \in \mathbb{R}^E$ and $\vec{v} \in \mathbb{R}^V$ the duality gap, $\text{gap}(\vec{f}, \vec{v}) \overset{\text{def}}{=} \xi(\vec{f}) - \zeta(\vec{v})$ is an upper bound on both $\xi(\vec{f}) - \xi(\vec{f}_{\text{opt}})$ and $\zeta(\vec{v}_{\text{opt}}) - \zeta(\vec{v})$.

In keeping with the electrical circuit interpretation we refer to a candidate dual solution $\vec{v} \in \mathbb{R}^V$ as voltages or vertex potentials and we define $\Delta_\vec{v}(a, b) \overset{\text{def}}{=} \vec{v}(a) - \vec{v}(b)$ to be the potential drop of $\vec{v}$ across $(a, b)$. By the KKT conditions we know that

$$\vec{f}_{\text{opt}} = \mathbf{R}^{-1} \mathbf{B} \vec{v}_{\text{opt}} \quad \text{i.e.} \quad \forall e \in E : \vec{f}_{\text{opt}}(e) = \frac{\Delta_{\vec{v}_{\text{opt}}}(e)}{r_e}.$$  

For $e \in E$ we call $\frac{\Delta_{\vec{v}}(e)}{r_e}$ the flow induced by $\vec{v}$ across $e$ and we call $\vec{f}(e)r_e$ the potential drop induced by $\vec{f}$ across $e$.

The optimality conditions $\vec{f}_{\text{opt}} = \mathbf{R}^{-1} \mathbf{B} \vec{v}_{\text{opt}}$ can be restated solely in terms of flows in a well known variant of Kirchoff's Potential Law (KPL) as follows.

**Lemma 2.2.7 (KPL).** Feasible $\vec{f} \in \mathbb{R}^E$ is optimal if and only if $\vec{f}^T \mathbf{R} \vec{c} = 0$ for all circulations $\vec{c} \in \mathbb{R}^E$.

### 2.2.2 Spanning Trees and Cycle Space

Let $T \subseteq E$ be a spanning tree of $G$ and let us call the edges in $T$ the tree edges and the edges in $E \setminus T$ the off-tree edges. By the fact that $G$ is connected and $T$ spans $G$ we know that for every $a, b \in V$ there is a unique path connecting $a$ and $b$ using only tree edges.

**Definition 2.2.8 (Tree Path).** For $a, b \in V$, we define the tree path $P_{(a,b)} \subseteq V \times V$ to be the unique path from $a$ to $b$ using edges from $T$. In vector form we let $\vec{p}_{(a,b)} \in \mathbb{R}^E$ denote the unique flow sending 1 unit of flow from $a$ to $b$, that is nonzero only on $T$.

For the off-tree edges we similarly define tree cycles.

**Definition 2.2.9 (Tree Cycle).** For $(a, b) \in E \setminus T$, we define the tree cycle $C_{(a,b)} \overset{\text{def}}{=} \{(a, b)\} \cup P_{(b,a)}$ to be the unique cycle consisting of edge $(a, b)$ and $P_{(b,a)}$. In vector form we let $\vec{c}_{(a,b)}$ denote the unique circulation sending 1 unit of flow on $C_{(a,b)}$.

---

1Note that edges of $P_{(a,b)}$ are oriented with respect to the path, not the natural orientation of $G$.
Cycle Space

The tree cycles form a complete characterization of circulations in a graph. The set of all circulations \( \{ \bar{c} \in \mathbb{R}^E \mid \mathbf{B}^T \bar{c} = 0 \} \) is a well-known subspace called cycle space [10] and the tree cycles \( \{ \bar{c}_e \mid e \in E \setminus T \} \) form a basis. This yields an even more succinct description of the KPL optimality condition (Lemma 2.2.7). A feasible \( \bar{f} \in \mathbb{R}^E \) is optimal if and only if \( \bar{f}^T \mathbf{R} \bar{c}_e = 0 \) for all \( e \in E \setminus T \).

We can think of each tree cycle \( C_e \) as a long resistor consisting of its edges in series with total resistance \( \sum_{e \in C_e} r_e \) and flow induced potential drop of \( \sum_{e \in C_e} \bar{f}(e) r_e \). KPL optimality then states that \( \bar{f} \in \mathbb{R}^E \) is optimal if and only if the potential drop across each of these resistors is 0. Here we define two key quantities relevant to this view.

**Definition 2.2.10 (Cycle Quantities).** For \( e \in E \setminus T \) and \( \bar{f} \in \mathbb{R}^E \) the resistance of \( C_e \), \( R_e \), and the flow induced potential across \( C_e \), \( \Delta_{c_e}(\bar{f}) \), are given by

\[
R_e \overset{\text{def}}{=} \sum_{e' \in C_e} r_e' = \bar{c}_e^T \mathbf{R} \bar{c}_e \quad \text{and} \quad \Delta_{c_e}(\bar{f}) \overset{\text{def}}{=} \sum_{e \in C_e} r_e \bar{f}(e) = \bar{f}^T \mathbf{R} \bar{c}_e .
\]

Low-Stretch Spanning Trees

Starting with a feasible flow, our algorithm computes an approximate electrical flow by fixing violations of KPL on randomly sampled tree cycles. How well this algorithm performs is determined by how well the resistances of the off-tree edges are approximated by their corresponding cycle resistances. This quantity, which we refer to as the tree condition number, is in fact a certain condition number of a matrix whose rows are properly normalized instances of \( \bar{c}_e \) (see Section 3.4.3).

**Definition 2.2.11 (Tree Condition Number).** The tree condition number of spanning tree \( T \) is given by \( \tau(T) \overset{\text{def}}{=} \sum_{e \in E \setminus T} \frac{R_e}{r_e} \), and we abbreviate it as \( \tau \) when the underlying tree \( T \) is clear from context.

This is closely related to a common quantity associated with a spanning tree called stretch.
Definition 2.2.12 (Stretch). The stretch of \( e \in E \), \( \text{st}(e) \), and the total stretch of \( T \), \( \text{st}(T) \), are

\[
\text{st}(e) \overset{\text{def}}{=} \sum_{e' \in P_e} \frac{r_{e'}}{r_e} \quad \text{and} \quad \text{st}(T) \overset{\text{def}}{=} \sum_{e \in E} \text{st}(e).
\]

Since \( R_e = r_e \cdot (1 + \text{st}(e)) \) we see that these quantities are related by \( \tau(T) = \text{st}(T) + m - 2n + 2 \).

Efficient algorithms for computing spanning trees with low total or average stretch, i.e. low-stretch spanning trees, have found numerous applications [3, 18] and all previous nearly-linear-time SDD-system solvers [50, 53, 31, 32], including the most efficient form of the SDD solver presented in this thesis, make use of such trees. There have been multiple breakthroughs in the efficient construction of low-stretch spanning trees [3, 18, 1, 31, 2] and the latest such result is used in this thesis and stated below.

Theorem 2.2.13 ([2]). In \( O(m \log n \log \log n) \) time we can compute a spanning tree \( T \) with total stretch \( \text{st}(T) = O(m \log n \log \log n) \).

2.3 The Algorithm

Given a SDD system \( Ax = b \) we wish to efficiently compute \( x \) such that \( \| x - A^T b \|_A \leq \varepsilon \| A^T b \|_A \). Using standard reduction techniques (see Appendix A of [30]) we can reduce solving such SDD systems to solving Laplacian systems corresponding to connected graphs without a loss in asymptotic run time. Therefore it suffices to solve \( L\tilde{x} = \tilde{\chi} \) in nearly-linear time when \( L \) is the Laplacian matrix for some connected graph \( G \). Here we provide an algorithm that both solves such systems and computes the corresponding \( \varepsilon \)-approximate electric flows in \( O(m \log^2 n \log \log n \log(e^{-1} n)) \) time.

2.3.1 A Simple Iterative Approach

Our algorithm focuses on the electric flow problem. First, we compute a low stretch spanning tree, \( T \), and a crude initial feasible \( \tilde{f}_0 \in \mathbb{R}^E \) taken to be the unique \( \tilde{f}_0 \) that meets the demands and is nonzero only on tree edges. Next, for a fixed number of iterations, \( K \), we perform simple iterative steps, referred to as cycle updates, in
which we compute a new feasible \( \tilde{f}_i \in \mathbb{R}^E \) from the previous feasible \( \tilde{f}_{i-1} \in \mathbb{R}^E \) while attempting to decrease energy. Each iteration, \( i \), consists of sampling an \( e \in E \setminus T \) proportional to \( \frac{R_e}{r_e} \), checking if \( \tilde{f}_{i-1} \) violates KPL on \( C_e \) (i.e. \( \Delta_{ce}(\tilde{f}_i) \neq 0 \)) and adding a multiple of \( \tilde{c}_e \) to make KPL hold on \( C_e \) (i.e. \( \tilde{f}_i = \tilde{f}_{i-1} - \frac{\Delta_{ce}(\tilde{f}_{i-1})}{R_e} \tilde{c}_e \)). Since, \( \mathbf{B}^T \tilde{c}_e = 0 \), this operation preserves feasibility. We show that in expectation \( f_K \) is an \( \varepsilon \)-approximate electrical flow.

To solve \( \mathbf{L} \tilde{v} = \tilde{x} \) we show how to use an \( \varepsilon \)-approximate electrical flow to derive a candidate solution to \( \mathbf{L} \tilde{v} = \tilde{x} \) of comparable quality. In particular, we use the fact that a vector \( \tilde{f} \in \mathbb{R}^E \) and a spanning tree \( T \) induce a natural set of voltages, \( \tilde{v} \in \mathbb{R}^V \) which we call the tree induced voltages.

**Definition 2.3.1 (Tree Induced Voltages).** For \( \tilde{f} \in \mathbb{R}^E \) and an arbitrary (but fixed) \( s \in V,^2 \) we define the tree induced voltages \( \tilde{v} \in \mathbb{R}^V \) by \( \tilde{v}(a) \triangleq \sum_{e \in P(a,s)} \tilde{f}(e)r_e \) for \( \forall a \in V \).

Our algorithm simply returns the tree induced voltages for \( \tilde{f}_K \), denoted \( \tilde{v}_K \), as the approximate solution to \( \mathbf{L} \tilde{v} = \tilde{x} \). The full pseudocode for the algorithm is given in Algorithm 1.

### Algorithm 1: SimpleSolver

**Input**: \( G = (V, E, r) \), \( \tilde{x} \in \mathbb{R}^V \), \( \varepsilon \in \mathbb{R}^+ \)

**Output**: \( \tilde{f} \in \mathbb{R}^E \) and \( \tilde{v} \in \mathbb{R}^V \)

1. \( T := \) low-stretch spanning tree of \( G \);
2. \( \tilde{f}_0 := \) unique flow on \( T \) such that \( \mathbf{B}^T \tilde{f}_0 = \tilde{x} \);
3. \( \tilde{v}_e := \frac{1}{\tau(T)} \cdot \frac{R_e}{r_e} \) for all \( e \in E \setminus T \);
4. \( K = \lceil \tau \log \left( \frac{m(T)}{\tau(T)} \right) \rceil \);
5. for \( i = 1 \) to \( K \) do
6. \( \quad \) Pick random \( e_i \in E \setminus T \) by probability distribution \( \tilde{\nu} \);
7. \( \quad \) \( \tilde{f}_i = \tilde{f}_{i-1} - \frac{\Delta_{ce}(\tilde{f}_{i-1})}{R_e} \tilde{c}_e \);
8. \( \) end
9. return \( \tilde{f}_K \) and its tree induced voltages \( \tilde{v}_K \)

---

^2We are primarily concerned with difference between potentials which are invariant under the choice \( s \in V \).
2.3.2 Algorithm Guarantees

In the next few sections we prove that SimpleSolver both computes an ε-approximate electric flow and solves the corresponding Laplacian system in nearly linear time:

**Theorem 2.3.2 (SimpleSolver).** The output of SimpleSolver satisfies

\[
E[\xi(f)] \leq (1 + \varepsilon) \cdot \xi(f_{opt}) \quad \text{and} \quad E[\|\varepsilon - L^i \mathbf{x}\|_L] \leq \sqrt{\varepsilon} \cdot \|L^i \mathbf{x}\|_L
\]

and SimpleSolver can be implemented to run in time \(O(m \log^2 n \log \log n \log(\varepsilon^{-1}n))\).

By construction SimpleSolver outputs a feasible flow and, by choosing \(T\) to be a low stretch spanning tree with properties guaranteed by Theorem 2.2.13, we know that the number of iterations of simple solver is bounded by \(O(m \log n \log \log n \log(\varepsilon^{-1}n))\). However, in order to prove the theorem we still need to show that (1) each iteration makes significant progress, (2) each iteration can be implemented efficiently, and (3) the starting flow and final voltages are appropriately related to \(\xi(f_{opt})\). In particular we show:

1. Each iteration of SimpleSolver decreases the energy of the current flow by at least an expected \((1 - \frac{1}{n})\) fraction of the energy distance to optimality (see Section 2.4).

2. Each iteration of SimpleSolver can be implemented to take \(O(\log n)\) time (see Section 2.5).

3. \(\xi(f_0)\) is sufficiently bounded, the quality of tree voltages is sufficiently bounded, and the entire algorithm can be implemented efficiently (see Section 2.6).

---

3 Although the theorem is stated as an expected guarantee on \(f\) and \(\varepsilon\), one can easily use Markov bound and Chernoff bound to provide a probabilistic but exact guarantee.

4 Note that a naïve implementation of cycle updates does not necessarily run in sublinear time. In particular, updating \(f_i := f_{i-1} - \Delta e(f_{i-1}) e\) by walking \(C_e\) and updating flow values one by one may take more than (even amortized) sublinear time, even though \(T\) may be of low total stretch. Since \(st(T)\) is defined with respect to cycle resistances but not with respect to the number of edges in these cycles, it is possible to have a low-stretch tree where each tree cycle still has \(O(V)\) edges on it. Furthermore, even if all edges have resistances 1 and therefore the average number of edges in a tree cycle is \(\tilde{O}(\log n)\), since SimpleSolver samples off-tree edges with higher stretch with higher probabilities, the expected number of edges in a tree cycle may still be \(\Omega(V)\).
2.4 Convergence Rate Analysis

In this section we analyze the convergence rate of SimpleSolver. The central result is as follows.

**Theorem 2.4.1 (Convergence).** Each iteration \( i \) of SimpleSolver computes feasible \( \bar{f}_i \in \mathbb{R}^E \) such that

\[
\mathbb{E}[\xi(\bar{f}_i)] - \xi(\bar{f}_{\text{opt}}) \leq \left(1 - \frac{1}{r}\right)^i \left(\xi(\bar{f}_0) - \xi(\bar{f}_{\text{opt}})\right).
\]

Our proof is divided into three steps. In Section 2.4.1 we analyze the energy gain of a single algorithm iteration, in Section 2.4.2 we bound the distance to optimality in a single algorithm iteration, and in Section 2.4.3 we connect these to prove the theorem.

2.4.1 Cycle Update Progress

For feasible \( \bar{f} \in \mathbb{R}^E \), we can decrease \( \xi(\bar{f}) \) while maintaining feasibility, by adding a multiple of a circulation \( \bar{c} \in \mathbb{R}^E \) to \( \bar{f} \). In fact, we can easily optimize to pick the best multiple.

**Lemma 2.4.2 (Energy Improvement).** For \( \bar{f} \in \mathbb{R}^E \), \( \bar{c} \in \mathbb{R}^E \), and \( \alpha^* = \frac{-\bar{f}^T R \bar{c}}{\bar{c}^T R \bar{c}} \in \mathbb{R} \) we have

\[
\arg \min_{\alpha \in \mathbb{R}} \xi(\bar{f} + \alpha \bar{c}) = -\frac{\bar{f}^T R \bar{c}}{\bar{c}^T R \bar{c}} \quad \text{and} \quad \xi(\bar{f} + \alpha^* \bar{c}) - \xi(\bar{f}) = -\frac{(\bar{f}^T R \bar{c})^2}{\bar{c}^T R \bar{c}}.
\]

**Proof.** By definition of energy \( \xi(\bar{f} + \alpha \bar{c}) = (\bar{f} + \alpha \bar{c})^T R (\bar{f} + \alpha \bar{c}) = \bar{f}^T R \bar{f} + 2\alpha \bar{f}^T R \bar{c} + \alpha^2 \bar{c}^T R \bar{c} \). Setting the derivative with respect to \( \alpha \) to 0 and substituting in \( \alpha = \alpha^* \) yields the results. \( \square \)

In the special case where \( \bar{c} = \bar{c}_e \) is a tree cycle for some off-tree edge \( e \in E \setminus T \), since \( R_e = \bar{c}_e^T R \bar{c}_e \) and \( \Delta_{\alpha}(\bar{f}) = \bar{f}^T R \bar{c}_e \), this procedure is precisely the iterative step of SimpleSolver, i.e. a cycle update. The following lemma follows immediately and
states that the energy decrease of a cycle update is exactly the energy of a resistor with resistance $R_e$ and potential drop $\Delta_{\infty}(\bar{f})$.

**Lemma 2.4.3** (Cycle Update). For feasible $\tilde{f} \in \mathbb{R}^E$ and $e \in E \setminus T$ we have

$$\xi\left(\tilde{f} - \frac{\Delta_{\infty}(\bar{f})}{R_e} \bar{e}\right) - \xi(\bar{f}) = -\frac{\Delta_{\infty}(\bar{f})^2}{R_e}.$$ 

**2.4.2 Distance to Optimality**

To bound how far $\tilde{f}$ in SimpleSolver is from optimality we use that the duality gap between $\tilde{f}$ and its tree induced voltages $\bar{v}$, is an upper bound on this distance. Here we derive a simple expression for this quantity in terms of cycle potentials.

**Lemma 2.4.4** (Tree Gap). For feasible $\tilde{f} \in \mathbb{R}^E$ and tree induced voltages $\bar{v} \in \mathbb{R}^V$ we have

$$\text{gap}(\tilde{f}, \bar{v}) = \sum_{e \in E \setminus T} \frac{\Delta_{\infty}(\bar{f})^2}{r_e}.$$ 

**Proof.** By definition of primal and dual energy we have $\text{gap}(\tilde{f}, \bar{v}) = \tilde{f}^T R \tilde{f} - (\bar{v}^T \bar{x} - \bar{v}^T L \bar{v})$. Using that $B^T \tilde{f} = \bar{x}$ and $L = B^T R^{-1} B$ we get

$$\text{gap}(\tilde{f}, \bar{v}) = \tilde{f}^T R \tilde{f} - 2\bar{v}^T B^T \tilde{f} - \bar{v}^T B^T R^{-1} B \bar{v} = \left(R f - B v\right)^T R^{-1} \left(R f - B v\right).$$

Therefore $\text{gap}(\tilde{f}, \bar{v}) = \sum_{e \in E} \frac{1}{r_e} \left(\bar{f}(e) r_e - \Delta_\infty(e)\right)^2$. However, by the uniqueness of tree paths, the antisymmetry of $\tilde{f} \in \mathbb{R}^E$, and the definition of tree voltages we have

$$\forall a, b \in V : \Delta_\infty(a, b) = \bar{v}(a) - \bar{v}(b) = \sum_{e \in P_{ab}} \bar{f}(e) r_e + \sum_{e \in P_{ba}} \bar{f}(e) r_e = \sum_{e \in P_{ab}} \bar{f}(e) r_e.$$ 

Therefore, $e \in T \Rightarrow \bar{f}(e) r_e - \Delta_\infty(e) = 0$ and $e \in E \setminus T \Rightarrow \bar{f}(e) r_e - \Delta_\infty(e) = \Delta_{\infty}(\bar{f})$. \qed

**2.4.3 Convergence Proof**

Here we connect the energy decrease of a cycle update (Lemma 2.4.3) and the duality gap formula (Lemma 2.4.4) to bound the the convergence of SimpleSolver.
Throughout this section we use $\tilde{v}_i$ to denote the tree induced voltages for flow $\tilde{f}_i$.

First, we show that in expectation each iteration $i$ of \texttt{SimpleSolver} decreases $\xi(\tilde{f}_{i-1})$ by a $\frac{1}{\tau}$ fraction of the duality gap.

**Lemma 2.4.5 (Expected Progress).** For iteration $i$ of \texttt{SimpleSolver} we have

$$ \mathbb{E}\left[\xi(\tilde{f}_i) - \xi(\tilde{f}_{i-1}) \left| \text{gap}(\tilde{f}_{i-1}, \tilde{v}_{i-1})\right.\right] = -\frac{\text{gap}(\tilde{f}_{i-1}, \tilde{v}_{i-1})}{\tau}. $$

**Proof.** In each iteration $i$, \texttt{SimpleSolver} picks a random $e_i \in E \setminus T$ with probability $p_{e_i}$ and adds a multiple of $\tilde{c}_{e_i}$ which by Lemma 2.4.3 decreases the energy by $\frac{\Delta_{e_i}(\tilde{f}_{i-1})^2}{R_{e_i}}$. Therefore

$$ \mathbb{E}\left[\xi(\tilde{f}_i) - \xi(\tilde{f}_{i-1}) \left| \text{gap}(\tilde{f}_{i-1}, \tilde{v}_{i-1})\right.\right] = \mathbb{E}\left[\sum_{e \in E \setminus T} p_e \left( -\frac{\Delta_{e_i}(\tilde{f}_{i-1})^2}{R_{e_i}} \right) \left| \text{gap}(\tilde{f}_{i-1}, \tilde{v}_{i-1})\right.\right] $$

Using that $p_e = \frac{1}{\tau} \cdot \frac{R_{e_i}}{R_{e}}$ and applying Lemma 2.4.4 yields the result. \qed

Next, we show that this implies that each iteration decreases the expected energy difference between the current flow and the optimal flow by a multiplicative $(1 - \frac{1}{\tau})$.

**Lemma 2.4.6 (Convergence Rate).** For all $i \geq 0$ let random variable $D_i \overset{\text{def}}{=} \xi(\tilde{f}_i) - \xi(\tilde{f}_{\text{opt}})$. Then for all iterations $i \geq 1$ we have $\mathbb{E}[D_i] \leq \left(1 - \frac{1}{\tau}\right) \mathbb{E}[D_{i-1}]$.

**Proof.** Since in each iteration of \texttt{SimpleSolver} one of a finite number of edges is chosen, clearly $D_i$ is a discrete random variable and by law of total expectation we have

$$ \mathbb{E}[D_i] = \sum_c \mathbb{E}[D_i | D_{i-1} = c] \Pr[D_{i-1} = c] $$

However, we know $D_{i-1} \leq \text{gap}(\tilde{f}_{i-1}, \tilde{v}_{i-1})$ so by Lemma 2.4.5, $\mathbb{E}[D_i | D_{i-1} = c] \leq c - \frac{\xi}{\tau}$. Therefore:

$$ \mathbb{E}[D_i] \leq \sum_c \left[\left(1 - \frac{1}{\tau}\right)c \cdot \Pr[D_{i-1} = c]\right] = \left(1 - \frac{1}{\tau}\right) \mathbb{E}[D_{i-1}]. $$

\qed
Finally, by induction on Lemma 2.4.6 and the definition of $D_i$ we prove Theorem 2.4.1.

2.5 Cycle Update Data Structure

In this section we show how to implement each iteration (i.e. cycle update) of SimpleSolver in $O(\log n)$ time. In Section 2.5.1 we formulate this as a data structure problem, in Section 2.5.2 we present a recursive solution, and in Section 2.5.3 we provide a linear algebraic interpretation that may be useful in both theory and practice.

2.5.1 The Data Structure Problem

In each iteration $i$ of SimpleSolver we pick a random $(a, b) \in E \setminus T$ and for feasible $f \in \mathbb{R}^E$ compute

$$\alpha^* = \frac{\Delta_{C(a,b)}(\bar{f})}{R(a,b)} = \bar{f}(a,b) r_{a,b} - (\bar{v}(a) - \bar{v}(b))$$

where $\bar{v}(a) = \sum_{e \in P(s,a)} \bar{f}(e) r_e$

and $s$ is an arbitrary fixed vertex in $V$ which we refer to as the root. Then $\alpha^*$ is added to the flow on every edge in the tree cycle $C(a,b)$. By the antisymmetry of flows and the uniqueness of tree paths, this update is equivalent to (1) adding $\alpha^*$ to the flow on edge $(a, b)$, (2) adding $-\alpha^*$ to the flow on every edge in $P(s,b)$ and then (3) adding $\alpha^*$ to the flow on every edge in $P(s,a)$.

Therefore, to implement cycle updates it suffices to store the flow on off-tree edges in an array and implement a data structure that supports the following operations.

* **init**($T, s \in V$): initialize a data structure $D$ given tree $T$ and root $s \in V$.
* **query**$_D(a \in V)$: return $\bar{v}(a) = \sum_{e \in P(s,a)} \bar{f}(e) r_e$.
* **update**$_D(a \in V, \alpha \in \mathbb{R})$: set $\bar{f}(e) := \bar{f}(e) + \alpha$ for all $e \in P(s,a)$.
2.5.2 Recursive Solution

While one could solve this data structure problem with a slight modification of link-cut trees [47], that dynamic data structure is overqualified for our purpose. In contrast to the problems for which link-cut trees were originally designed, in our setting the tree is static, so that much of the sophistication of link-cut trees may be unnecessary. Here we develop a very simple separator decomposition tree based structure that provides worst-case $O(\log n)$ operations that we believe sheds more light on the electric flow problem and may be useful in practice.

Our solution is based on the well known fact that every tree has a good vertex separator, tracing back to Jordan in 1869 [23].

**Lemma 2.5.1 (Tree Vertex Separator).** For a spanning tree $T$ rooted at $s$ with $n \geq 2$ vertices, in $O(n)$ time we can compute

$$(d, T_0, \ldots, T_k) = \text{tree-decompose}(T),$$

such that the removal of $d \in V$ (which might equal to $s$) disconnects $T$ into subtrees $T_0, \ldots, T_k$, where $T_0$ is rooted at $s$ and contains $d$ as a leaf, while other $T_i$'s are rooted at $d$. Furthermore, each $T_i$ has at most $n/2 + 1$ vertices.

**Proof.** We start at root $s$, and at each step follow an edge to a child whose corresponding subtree is the largest among all children. We continue this procedure until at some vertex $d$ (which might be $s$ itself) the sizes of all its subtrees have no more than $n/2$ vertices. This vertex $d$ is the desired vertex separator. Now if $s \neq d$ we let $T_0$ be the subtree above $d$ rooted at $s$ with $d$ being the leaf, and let each $T_i$ be a sub-tree below $d$ and rooted at $d$; or when $s = d$ we let $T_0, \ldots, T_k$ each denote a subtree below $d$ and rooted at $d$. By pre-computing subtree sizes $\text{tree-decompose}(T)$ runs in $O(n)$ time. \[\square\]

\[5\] For technical purposes, when $V$ has only two vertices with $s$ being the root, we always define $d \in V$ such that $d \neq s$ and define $\text{tree-decompose}(T)$ to return no sub-trees, i.e., $k = -1$. By orienting the tree away from $s$ and precomputing the number of descendents of each node we can easily implement this in $O(n)$ time.
Applying this lemma recursively induces a separator decomposition tree from which we build our data structure. To execute init\((T,s)\) we compute a vertex separator \((d,T_0,\ldots,T_k)\) and recurse separately on each subtree until we are left with a tree of only two vertices. In addition, we precompute the total weight of the intersection of the root to \(d\) path \(P_{(s,d)}\) and the root to \(a\) path \(P_{(s,a)}\), denoted by \(\text{height}(a)\) for every \(a \in V\), by taking a walk over the tree. See Algorithm 2 for details.

With this decomposition we can support \texttt{query} and \texttt{update} by maintaining just 2 values about the paths \(P_{(s,d)}\). For each subtree we maintain the following variables:

- \texttt{d_drop}, the total potential drop induced on the path \(P_{(s,d)}\), and
- \texttt{d_ext}, the total amount of flow that has been updated using the entire \(P_{(s,d)}\) path, i.e. the contribution to \(P_{(s,d)}\) from vertices beyond \(d\).

It is easy to see that these invariants can be maintained recursively and that they suffice to answer queries and updates efficiently (see Algorithm 3 and Algorithm 4). Furthermore, because the sizes of trees at least half at each recursion, init takes \(O(n \log n)\) while \texttt{query} and \texttt{update} each takes \(O(\log n)\) time.

### 2.5.3 Linear Algebra Solution

Here we unpack the recursion in the previous section to provide a linear algebraic view of our data structure that may be useful in both theory and practice. We show that the init procedure in the previous section computes for all \(a \in V\) a query vector \(\hat{q}(a)\) and an update vector \(\hat{u}(a)\) each of support size \(O(\log |V|)\), such that the entire state of the data structure is an \(O(|V|)\) dimensional vector \(\vec{x}\) initialized to \(\vec{0}\) allowing \texttt{query} and \texttt{update} to be as simple as the following,

\[
\text{query}(a) : \text{return } \hat{q}(a) \cdot \vec{x} \\
\text{update}(a, \alpha) : \vec{x} := \vec{x} + \alpha \hat{u}(a) .
\] (2.3)

To see this, first note that init recursively creates a total of \(N = O(|V|)\) subtrees \(T_1, \ldots, T_N\). Letting \texttt{d_ext}_i, \texttt{d_drop}_i, \texttt{height}_i denote \texttt{d_ext}, \texttt{d_drop}, and \texttt{height} as-
Algorithm 2: Recursive init\((T, s \in V)\)

\[
\begin{align*}
d_{\text{ext}} &:= 0, \quad d_{\text{drop}} := 0; \\
(d, T_0, \ldots, T_k) &:= \text{tree-decompose}(T); \\
\forall a \in V: \text{height}(a) &:= \sum_{e \in P(a, s) \cap P(s, d)} f_e; \\
\text{if } |V| > 2 &\text{ then} \\
\quad &\forall i \in \{0, 1, \ldots, k\}: D_i := \text{init}(T_i, d); \\
\text{end}
\end{align*}
\]

Algorithm 3: Recursive query\((a \in V)\)

\[
\begin{align*}
\text{if } a &= d & \text{then return } d_{\text{drop}}; \\
\text{else if } |V| &= 2 & \text{then return } 0; \\
\text{else if } a &\in T_0 & \text{then return } d_{\text{ext}} \cdot \text{height}(a) + \text{query}_{D_0}(a); \\
\text{else} &\quad \text{let } T_i \text{ be unique tree containing } a; \\
&\quad \text{return } d_{\text{drop}} + \text{query}_{D_i}(a); \\
\text{end}
\end{align*}
\]

Algorithm 4: Recursive update\((a \in V, \alpha \in \mathbb{R})\)

\[
\begin{align*}
d_{\text{drop}} &:= d_{\text{drop}} + \alpha \cdot \text{height}(a); \\
\text{if } |V| &= 2 & \text{then return}; \\
\text{if } a &\not\in T_0 & \text{then } d_{\text{ext}} := d_{\text{ext}} + \alpha; \\
\text{if } a &\neq d & \text{then} \\
&\quad \text{let } T_i \text{ be unique tree containing } a; \\
&\quad \text{update}_{D_i}(a, \alpha); \\
\text{end}
\end{align*}
\]

Associated with tree \(T_i\), the state of the data structure is completely represented by a vector \(\vec{x} \in \mathbb{R}^{\{e,d\} \times N}\) defined as follows

\[
\vec{x}_{c,i} \overset{\text{def}}{=} \begin{cases} 
  d_{\text{ext}}, & \text{if } c = e; \\
  d_{\text{drop}}, & \text{if } c = d.
\end{cases}
\]

It is easy to see that given vertex \(a \in V\), in order for \text{query}(a) or \text{update}(a, \alpha) to affect tree \(T_i\) in the decomposition, it is necessary that \(a\) is a vertex in \(T_i\) and \(a\) is not the root \(s_i\) of \(T_i\). Accordingly, we let \(T(a) \overset{\text{def}}{=} \{i \in [N] \mid a \in T_i \text{ and } a \neq s_i\}\) be the indices of such trees that \(a\) may affect.
Next, in order to fully specify the query vector $\vec{q}(a)$ and the update vector $\vec{u}(a)$, we further refine $T(a)$ by defining $T_0(a) \overset{\text{def}}{=} \{ i \in T(a) \mid a \text{ is in } T_0 \text{ of } \text{tree-decompose}(T) \}$ and $T_+(a) \overset{\text{def}}{=} T(a) \setminus T_0(a)$. Then, one can carefully check that the following definitions of $\vec{q}(a), \vec{u}(a) \in \mathbb{R}^{[e,d] \times N}$ along with their query and update in (2.3), exactly coincide with our recursive definitions in Algorithm 3 and Algorithm 4 respectively.

$$
\vec{q}(a)(c,i) \overset{\text{def}}{=} \begin{cases} 
\text{height}_i(a) & c = e \text{ and } i \in T_0(a) \\
1 & c = d \text{ and } i \in T_+(a) \\
0 & \text{otherwise}
\end{cases}
$$

and

$$
\vec{u}(a)(c,i) \overset{\text{def}}{=} \begin{cases} 
1 & c = e \text{ and } i \in T_+(a) \\
\text{height}_i(a) & c = d \text{ and } i \in T(a) \\
0 & \text{otherwise}
\end{cases}
$$

Furthermore, the properties of init and tree-decompose immediately implies that each vertex $a$ can appear in at most $O(\log n)$ trees where it is not the root, so $|T(a)| = O(\log n)$. As a consequence, $\vec{q}(a)$ and $\vec{u}(a)$ each has at most $O(\log n)$ non-zero entries. It is also not hard to see that we can pre-compute these vectors in $O(n \log n)$ time using our recursive init.

Applying these insights to the definition of a cycle update we see that this data structure allows each cycle update of SimpleSolver to be implemented as a single dot product and a single vector addition where one vector in each operation has $O(\log n)$ non-zero entries. Since these vectors can be precomputed each cycle update can be implemented by iterating over a fixed array of $O(\log n)$ pointers and performing simple arithmetic operations. We hope this may be fast in practice.

### 2.6 Asymptotic Running-Time

Here we give the remaining analysis needed to prove the correctness of SimpleSolver and prove Theorem 2.3.2. First we bound the energy of $f_0$. 36
Lemma 2.6.1 (Initial Energy). \( \xi(\vec{f}_0) \leq \text{st}(T) \cdot \xi(\vec{f}_{\text{opt}}) \).

Proof. Recall that \( \vec{f}_0 \in \mathbb{R}^E \) is the unique vector that meets the demands and is nonzero only on \( T \). Now, if for every \( e \in E \) we sent \( \vec{f}_{\text{opt}}(e) \) units of flow along \( P_e \) we achieve a vector with the same properties. Therefore, \( \vec{f}_0 = \sum_{e \in E} \vec{f}_{\text{opt}}(e)\vec{p}_e \) and by applying the Cauchy-Schwarz inequality we get

\[
\xi(\vec{f}_0) = \sum_{e \in T} r_e \left( \sum_{e' \in E \mid e \in T_e} \vec{f}_{\text{opt}}(e') \right)^2 \leq \sum_{e \in T} \left[ \left( \sum_{e' \in E \mid e \in T_e} \frac{r_e}{r_{e'}} \right) \left( \sum_{e' \in E \mid e \in T_e} \frac{r_{e'}}{r_e} \right) \left( \vec{f}_{\text{opt}}(e') \right)^2 \right] .
\]

Applying the crude bound that

\[
\forall e \in E : \sum_{e' \in E \mid e \in T_e} r_{e'} \vec{f}_{\text{opt}}(e')^2 \leq \xi(\vec{f}_{\text{opt}})
\]

and noting that by the definition of stretch

\[
\sum_{e \in T} \sum_{e' \in E \mid e \in T_e} \frac{r_e}{r_{e'}} = \sum_{e' \in E \mid e \in T_e} \sum_{e \in T_e} \frac{r_e}{r_{e'}} = \text{st}(T)
\]

the result follows immediately. \( \square \)

Next, we show how approximate optimality is preserved within polynomial factors when rounding from an \( \varepsilon \)-approximate electric flow to tree induced voltages.

Lemma 2.6.2 (Tree Voltage Rounding). Let \( \vec{f} \in \mathbb{R}^E \) be a primal feasible flow with \( \xi(\vec{f}) \leq (1 + \varepsilon) \cdot \xi(\vec{f}_{\text{opt}}) \) for \( \varepsilon > 0 \) and let \( \vec{v} \in \mathbb{R}^V \) be the tree induced voltages. Then the following holds

\[
\|\vec{v} - \mathbf{L}^\dagger \vec{\chi}\|_L \leq \sqrt{\varepsilon \cdot \tau} \cdot \|\mathbf{L}^\dagger \vec{\chi}\|_L .
\]

Proof. By Lemma 2.4.5 we know that one random cycle update from SimpleSolver is expected to decrease \( \xi(\vec{f}) \) by \( \text{gap}(\vec{f}, \vec{v})/\tau \). Therefore by the optimality of \( \vec{f}_{\text{opt}} \) we have

\[
\xi(\vec{f}) - \frac{\text{gap}(\vec{f}, \vec{v})}{\tau} \geq \xi(\vec{f}_{\text{opt}})
\]

\[6\text{This lemma follows immediately from the well known fact that } T \text{ is a } \text{st}(T)\text{-spectral sparsifier of } G, \text{ cf. } [53, \text{Lemma 9.2}], \text{ but a direct proof is included here for completeness.}\]
and since $\xi(\tilde{f}) \leq (1 + \varepsilon) \cdot \xi(\tilde{f}_{\text{opt}})$ and $\xi(\tilde{f}) \leq \xi(\tilde{f}_{\text{opt}})$ we have $\text{gap}(\tilde{f}_{\text{opt}}, \tilde{v}) \leq \varepsilon \cdot \tau \cdot \xi(\tilde{f}_{\text{opt}})$. Finally, using that $\tilde{v}_{\text{opt}} = L^T\tilde{x}$, $\tilde{f}_{\text{opt}} = R^{-1}B\tilde{v}_{\text{opt}}$, and $L = B^TR^{-1}B$ it is straightforward to check

$$\text{gap}(\tilde{f}_{\text{opt}}, \tilde{v}) = \|\tilde{v} - L^T\tilde{x}\|_L^2 \quad \text{and} \quad \xi(\tilde{f}_{\text{opt}}) = \|L^T\tilde{x}\|_L^2.$$ 

Finally, we have everything we need to prove the correctness of SimpleSolver.

**Proof of Theorem 2.3.2.** Applying Theorem 2.4.1, Lemma 2.6.1, and the definition of $K$ immediately yields the following.

$$\mathbb{E}[\xi(\tilde{f}_K)] - \xi(\tilde{f}_{\text{opt}}) \leq \left(1 - \frac{1}{\tau}\right)\tau \log(\varepsilon^{-1}\text{st}(T)\tau) \left(\text{st}(T) \xi(\tilde{f}_{\text{opt}}) - \xi(\tilde{f}_{\text{opt}})\right).$$

Using several crude bounds and applying Lemma 2.6.2 we get

$$\mathbb{E}[\xi(\tilde{f}_K)] \leq \left(1 + \frac{\varepsilon}{\tau}\right)\xi(\tilde{f}_{\text{opt}}) \quad \text{and} \quad \mathbb{E}\|\tilde{v}_K - L^T\tilde{x}\|_L \leq \sqrt{\varepsilon} \cdot \|L^T\tilde{x}\|_L$$

which is stronger than what we need to prove for the theorem.

All that remains is to bound the running time. To do this, we implement the algorithm SimpleSolver using the latest low-stretch spanning tree construction (Theorem 2.2.13) and construct a spanning tree $T$ of stretch $\text{st}(T) = O(m \log n \log \log n)$ in time $O(m \log n \log \log n)$ yielding $\tau = O(m \log n \log \log n)$.

Next, to compute $\tilde{f}_0$ we note that given any demand vector $\tilde{x} \in \mathbb{R}^V$ with $\sum_i \tilde{x}_i = 0$, the quantity $\tilde{f}_0(e)$ on a tree edge $e \in T$ is uniquely determined by the summation of $\tilde{x}_v$ where $v$ is over the vertices on one side of $e$. Therefore, $\tilde{f}_0$ can be computed via a DFS in $O(n)$ time.

To compute $R_e$ for each off-tree edge $e \in E \setminus T$ we could either use Tarjan's off-line LCA algorithm [56] that runs in a total of $O(m)$ time, or simply use our own data structure in $O(m \log n)$ time. In fact, one can initiate a different instance of our data structure on $T$, and for each off-tree edge $(a, b) \in E \setminus T$, one can call update($b, 1$)
and update \( (a, -1) \), so that \( R_e = \text{query}(b) - \text{query}(a) + r_e \).

Finally, we can initialize our data structure in \( O(n \log n) \) time, set the initial flow values in \( O(n \log n) \) time, perform each cycle update in \( O(\log n) \) time, and compute all the tree voltages in \( O(n \log n) \) time using the work in section Section 2.5. Since the total number of iterations of our algorithm is easily seen to be \( O(m \log n \log(n \varepsilon^{-1}) \log \log n) \) we get the desired total running time.

2.7 Numerical Stability

Up until this point our analysis has assumed that all arithmetic operations are exact. In this section, we show that SimpleSolver is numerically stable and achieves the same convergence guarantees when implemented with finite-precision arithmetic.

For simplicity of exposition, we assume that the resistances \( r_e \) and the coordinates of the demand vector \( \bar{X} \) are all represented as \( b \)-bit integers with absolute values bounded by \( N = 2^b \). We show that SimpleSolver works when arithmetic operations are implemented with \( O(\max(b, \log n, \log 1/\varepsilon)) \) bits of precision (which is necessary simply to guarantee we can represent an answer meeting the required error bound).

In particular, if the entries of the input and \( \varepsilon \) can be stored in \( \log n \)-bit words, our running time guarantees hold in the standard unit cost RAM model, which only allows arithmetic operations on \( O(\log n) \)-bit numbers to be done in constant time.

We start our algorithm by multiplying the demand vector \( \bar{X} \) by \( \lceil 4mN^2/\varepsilon \rceil \), to ensure that in \( \tilde{f}_{\text{opt}} \) there exist at least \( \lceil 4mN^2/\varepsilon \rceil \) total units of flow on the edges. Since the resistances are at least \( 1 \), this guarantees that \( \xi(\tilde{f}_{\text{opt}}) \geq \lceil 4mN^2/\varepsilon \rceil \). Next, we ensure that throughout our algorithm all flow vectors \( \tilde{f}_i \) are integer vectors. At each iteration of SimpleSolver, when we are given a primal feasible flow \( \tilde{f} \) and want to compute the optimal cycle update \( \alpha^* \) of \( f/ \text{opt} R e \), we round this fraction to the nearest integer and suppose we pick some \( \tilde{d} \in Z \) such that \( |\alpha^* - \tilde{d}| \leq \frac{1}{2} \).

Based on the modifications above, it is easy to see that our algorithm can be run on a RAM machine with word size \( O(\max(b, \log n, \log 1/\varepsilon)) \), since all flow values are integers bounded above by \( O(\text{poly}(N,n,1/\varepsilon)) \). All we have left to show is the
convergence analysis for such integral updates.

We do so by strengthening our cycle update lemma, Lemma 2.4.3. For each cycle update, if we add \( \bar{\alpha} = \alpha^*(1 + \delta) \) units of flow on the cycle instead of \( \alpha^* \), the corresponding energy decrease is

\[
\xi(\tilde{f} - \alpha^*(1 + \delta)\tilde{c}_e) - \xi(\tilde{f}) = (\tilde{f} - \alpha^*(1 + \delta)\tilde{c}_e)^T R(\tilde{f} - \alpha^*(1 + \delta)\tilde{c}_e) - \tilde{f}^T R \tilde{f}
\]

\[
= -2\alpha^*(1 + \delta)\tilde{f}^T R \tilde{c}_e + (\alpha^*(1 + \delta))^2 \tilde{c}_e^T R \tilde{c}_e
\]

\[
= -\frac{\Delta_{c_e}(\tilde{f})^2}{R_e}(1 - \delta^2),
\]

where the last equality has used the fact that \( R_e = \tilde{c}_e^T R \tilde{c}_e \). We have \( |\delta| \leq \frac{1}{2\alpha^*} \), so, as long as \( \alpha^* \geq 1 \), the decrease in the energy decrease is at least \( \frac{3}{4} \) of what it would be for \( \delta = 0 \). We call an off-tree edge "good" if its \( \alpha^* > 1 \), and "bad" otherwise. We can rewrite the duality gap as

\[
gap(\tilde{f}, \overline{c}) = \sum_{e \in E \setminus T} \frac{\Delta_{c_e}(\tilde{f})^2}{r_e} \leq \sum_{e \in E \setminus T} \frac{\Delta_{c_e}(\tilde{f})^2}{R_e} \frac{R_e}{r_e}
\]

\[
= \sum_{e \in E \setminus T} \frac{\Delta_{c_e}(\tilde{f})^2}{R_e^2} \frac{R_e}{r_e} + \sum_{e \in E \setminus T} \frac{\Delta_{c_e}(\tilde{f})^2}{R_e} \frac{R_e}{r_e}
\]

\[
\leq mN^2 + \sum_{e \in E \setminus T} \frac{\Delta_{c_e}(\tilde{f})^2}{R_e} \frac{R_e}{r_e}.
\]

As a consequence, if one samples\(^7\) each tree cycle \( \tilde{c}_e \) with probability to \( \frac{R_e}{r_e} \), then the expected energy decrease is at least:

\[
\mathbb{E} \left[ \xi(\tilde{f}_i) - \xi(\tilde{f}_{i-1}) \big| \text{gap}(\tilde{f}_{i-1}, \overline{v}_{i-1}) \right] \leq \sum_{e \in E \setminus T} \frac{R_e}{r_e} \left( -\frac{\Delta_{c_e}(\tilde{f})^2}{4R_e^2} \right)
\]

\[
\leq -\frac{\mathbb{E} \left[ \text{gap}(\tilde{f}_{i-1}, \overline{v}_{i-1}) - mN^2 \right]}{4\tau/3} \leq -\frac{\xi(\tilde{f}_{i-1}) - \xi(\tilde{f}_{\text{opt}}) - mN^2}{4\tau/3}.
\]

\(^7\)Recall that exact sampling takes expected \( O(1) \) time on a machine with \( O(\log n) \)-sized words.
If one defines a random variable $D_i \overset{\text{def}}{=} \xi(f_i) - \xi(f_{\text{opt}}) - mN^2$, then

$$
\mathbb{E}[D_i|D_{i-1}] \leq \left( 1 - \frac{1}{4\pi/3} \right) D_{i-1} .
$$

Using the same analysis as before, after $K = \frac{4\pi}{3} \log \frac{2\text{att}(T)}{\varepsilon p}$, we have with probability at least $1 - p$ that $\xi(f_K) \leq (1 + \frac{\varepsilon}{2}) (\xi(f_{\text{opt}}) + mN^2)$ $\leq (1 + \varepsilon) \xi(f_{\text{opt}})$ as desired.
Chapter 3

Efficient Accelerated Coordinate Descent

In the previous chapter we saw how repeated application of a simple computational efficient update rule can be used to obtain a nearly linear time SDD solver. This idea of using simple sublinear-time iterative steps to solve a convex optimization problem is in fact an old one [24, 62, 4]. It is an algorithmic design principle that has seen great practical success [39, 21, 4] but has been notoriously difficult to analyze. In the past few years great strides have been taken towards developing a theoretical understanding of randomized variants of these approaches. Beyond the results in the previous chapter, in 2006 Strohmer and Vershynin [55] showed that a particular sublinear update algorithm for solving overconstrained linear systems called randomized Kaczmarz converges exponentially and in 2010 Nesterov [43] analyzed randomized analog of gradient descent that updates only a single coordinate in each iteration, called coordinate gradient descent method and provided a computationally inefficient but theoretically interesting accelerated variant, called accelerated coordinate gradient descent method (ACDM).

In this chapter we provide a framework that both strengthens and unifies these results. We present a more general version of Nesterov’s ACDM and show how to implement it so that each iteration has the same asymptotic runtime as its non-accelerated variants. We show that this method is numerically stable and optimal
under certain assumptions. Then we show how to use this method to outperform conjugate gradient in solving a general class of symmetric positive definite systems of equations. Furthermore, we show how to cast both randomized Kaczmarz and SimpleSolver in this framework and achieve faster running times using ACDM.

The rest of this chapter is organized as follows. In Section 3.1, we introduce the problem and function properties that we will use for optimization. In Section 3.2, we briefly review the mathematics behind gradient descent, accelerated gradient descent, and coordinate descent. In Section 3.3, we present our general ACDM implementation, prove correctness, and show how to implement the update steps efficiently. In Section 3.4, we show how to apply ACDM to achieve faster runtimes for various linear system solving scenarios and in Appendix A we provide missing details of the proof of ACDM convergence and provide numerical stability proofs.

3.1 Preliminaries

For the remainder of this chapter we change notation slightly and focus on the general unconstrained minimization problem $^1$

$$\min_{\bar{x} \in \mathbb{R}^n} f(\bar{x})$$

where the objective function $f : \mathbb{R}^n \rightarrow \mathbb{R}$ is continuously differentiable and convex, meaning that

$$\forall \bar{x}, \bar{y} \in \mathbb{R}^n : f(\bar{y}) \geq f(\bar{x}) + \langle \nabla f(x), \bar{y} - \bar{x} \rangle.$$  

We let $f^* \overset{\text{def}}{=} \min_{\bar{x} \in \mathbb{R}^n} f(\bar{x})$ denote the minimum value of this optimization problem and we let $\bar{x}^* \overset{\text{def}}{=} \arg\min_{\bar{x} \in \mathbb{R}^n} f(\bar{x})$ denote an arbitrary point that achieves this value.

To minimize $f$, we restrict our attention to first-order iterative methods, that is algorithms that generate a sequence of points $\bar{x}_1, \ldots, \bar{x}_k$ such that $\lim_{k \rightarrow \infty} f(\bar{x}_k) = f^*$.

---

$^1$Many of the results in this chapter can be generalized to constrained minimization problems [43], problems where $f$ is strongly convex with respect to different norms [43], and problems where each coordinate is a higher dimension variable (i.e. the block setting) [45]. However, for simplicity we focus on the basic coordinate unconstrained problem in the Euclidian norm.
while only evaluating the objective function, \( f \), and its gradient \( \nabla f \), at points. In other words, other than the global function parameters related to \( f \) that we define in this section, we assume that the algorithms under consideration have no further knowledge regarding the structure of \( f \). To compare such algorithms, we say that an iterative method has convergence rate \( r \) if \( f(x_k) - f^* \leq O((1 - r)^k) \) for this method.

Now, we say that \( f \) has convexity parameter \( \sigma \) with respect to some norm \( \| \cdot \| \) if the following holds

\[
\forall \bar{x}, \bar{y} \in \mathbb{R}^n : \quad f(\bar{y}) \geq f(\bar{x}) + \langle \nabla f(\bar{x}), \bar{y} - \bar{x} \rangle + \frac{\sigma}{2} \| \bar{y} - \bar{x} \|^2 \quad (3.1)
\]

and we say \( f \) strongly convex if \( \sigma > 0 \). We refer to the right hand side of (3.1) as the lower envelope of \( f \) at \( \bar{x} \) and for notational convenience when the norm \( \| \cdot \| \) is not specified explicitly we assume it to be the standard Euclidean norm \( \| x \| \overset{\text{def}}{=} \sqrt{\sum_i x_i^2} \).

Furthermore, we say \( f \) has \( L \)-Lipschitz gradient if

\[
\forall \bar{x}, \bar{y} \in \mathbb{R}^n : \quad \| \nabla f(\bar{y}) - \nabla f(\bar{x}) \| \leq L \| \bar{y} - \bar{x} \|
\]

The definition is related to an upper bound on \( f \) as follows:

**Lemma 3.1.1.** \([41, \text{Thm 2.1.5}]\) For continuously differentiable \( f : \mathbb{R}^n \to \mathbb{R} \) and \( L > 0 \), it has \( L \)-Lipschitz gradient if and only if

\[
\forall \bar{x}, \bar{y} \in \mathbb{R}^n : \quad f(\bar{y}) \leq f(\bar{x}) + \langle \nabla f(\bar{x}), \bar{y} - \bar{x} \rangle + \frac{L}{2} \| \bar{x} - \bar{y} \|^2. \quad (3.2)
\]

We call the right hand side of (3.2) the upper envelope of \( f \) at \( \bar{x} \).

The convexity parameter \( \mu \) and the Lipschitz constant of the gradient \( L \) provide lower and upper bounds on \( f \). They serve as the essential characterization of \( f \) for first-order methods and they are typically the only information about \( f \) provided to both gradient and accelerated gradient descent methods (besides the oracle access to \( f \) and \( \nabla f \)). For twice differentiable \( f \), these values can also be computed by properties of the Hessian of \( f \) by the following well known lemma:
Lemma 3.1.2 ([43]). Twice differentiable $f : \mathbb{R}^n \rightarrow \mathbb{R}$ has convexity parameter $\mu$ and $L$-Lipschitz gradient with respect to norm $\|\cdot\|$ if and only if $\forall \vec{x} \in \mathbb{R}^n$ the Hessian of $f$ at $\vec{x}$, $\nabla^2 f(\vec{x}) \in \mathbb{R}^{n \times n}$ satisfies

$$\forall \vec{y} \in \mathbb{R}^n : \mu \|\vec{y}\|^2 \leq \vec{y}^T (\nabla^2 f(\vec{x})) \vec{y} \leq L \|\vec{y}\|^2$$

To analyze coordinate-based iterative methods, that is iterative methods that only consider one component of the current point or current gradient in each iteration, we need to define several additional parameters characterizing $f$. For all $i \in [n]$, let $\vec{e}_i \in \mathbb{R}^n$ denote the standard basis vector for coordinate $i$, let $f_i(\vec{x}) \in \mathbb{R}^n$ denote the partial derivative of $f$ at $\vec{x}$ along $\vec{e}_i$, i.e. $f_i(\vec{x}) \overset{\text{def}}{=} \vec{e}_i^T \nabla f(\vec{x})$, and let $\vec{f}_i(\vec{x})$ denote the corresponding vector, i.e $\vec{f}_i(\vec{x}) \overset{\text{def}}{=} f_i \cdot \vec{e}_i$. We say that $f$ has component-wise Lipschitz continuous gradient with Lipschitz constants $\{L_i\}$ if

$$\forall \vec{x} \in \mathbb{R}^n, \forall t \in \mathbb{R}, \forall i \in [n] : |f_i(\vec{x} + t \cdot \vec{e}_i) - f_i(\vec{x})| \leq L_i \cdot |t|$$

and for all $\alpha \geq 0$ we let $S_\alpha \overset{\text{def}}{=} \sum_{i=1}^{n} L_i^\alpha$ denote the total component-wise Lipschitz constant of $\nabla f$. Later we will see that $S_\alpha$ has a similar role for coordinate descent as $L$ has for gradient descent.

We give two examples for convex functions induced by linear systems and calculate their parameters. Note that even though one example can be deduced from the other, we provide both as the analysis allows us to introduce more notation.

**Example 3.1.3.** Let $f(\vec{x}) \overset{\text{def}}{=} \frac{1}{2} \langle \mathbf{A}\vec{x}, \vec{x} \rangle - \langle \vec{b}, \vec{x} \rangle$ for symmetric positive definite matrix $\mathbf{A} \in \mathbb{R}^{n \times n}$. Since $\mathbf{A} = \mathbf{A}^T$ clearly $\nabla f(\vec{x}) = \mathbf{A}\vec{x} - \vec{b}$ and $\nabla^2 f(\vec{x}) = \mathbf{A}$. Therefore, by Lemma 3.1.2, $L$ and $\sigma$ satisfy

$$\sigma \|\vec{x}\|^2 \leq \vec{x}^T \mathbf{A}\vec{x} \leq L \|\vec{x}\|^2 .$$

Consequently, $\sigma$ is the the smallest eigenvalue $\lambda_{\min}$ of $\mathbf{A}$ and $L$ is the largest eigenvalue $\lambda_{\max}$ of $\mathbf{A}$. Furthermore, $\forall i \in [n]$ we see that $f_i(\vec{x}) = \vec{e}_i^T (\mathbf{A}\vec{x} - \vec{b})$ and therefore $L_i$
satisfies
\[ \forall t \in \mathbb{R} : |t| \cdot |A_{ii}| = |e_i^T A (te_i)| \leq L_i |t| . \]

Since the positive definiteness of \( A \) implies that \( A \) is positive on diagonal, we have \( L_i = A_{ii} \), and consequently \( S_1 = \text{tr}(A) = \sum_{i=1}^{n} A_{ii} = \sum_{i=1}^{n} \lambda_i \) where \( \lambda_i \) are eigenvalues of \( A \).

**Example 3.1.4.** Let \( f(\bar{x}) = \frac{1}{2} \|A\bar{x} - b\|^2 \) for any matrix \( A \). Then the gradient \( \nabla f(\bar{x}) = A^T (A\bar{x} - \bar{b}) \) and the hessian \( \nabla^2 f(\bar{x}) = A^T A \). Hence, \( \sigma \) and \( L \) satisfy
\[ \sigma \|\bar{x}\|^2 \leq \bar{x}^T A^T A \bar{x} \leq L \|\bar{x}\|^2 \]
and we see that \( \sigma \) is the the smallest eigenvalue \( \lambda_{\min} \) of \( A^T A \) and \( L \) is the largest eigenvalue \( \lambda_{\max} \) of \( A^T A \). As in the previous example, we therefore have \( L_i = \|a_i\|^2 \) where \( a_i \) is the \( i \)-th column of \( A \) and \( S_1 = \sum \|a_i\|^2 = \|A\|_F^2 \), the Frobenius norm of \( A \).

### 3.2 Review of Previous Iterative Methods

In this section, we briefly review several standard iterative first-order method for smooth convex minimization. This overview is by no means all-inclusive, our goal is simply to familiarize the reader with numerical techniques we will make heavy use of later and motivate our presentation of the accelerated coordinate descent method. For a more comprehensive review, there are multiple good references, e.g. [41], [14].

In Section 3.2.1, we briefly review the standard gradient descent method. In Section 3.2.2, we show how to improve gradient descent by motivating and reviewing Nesterov's *accelerated gradient descent method* [40] through a more recent presentation of his via *estimate sequences* [41]. In Section 3.2.3, we review Nesterov's *coordinate gradient descent method* [43]. In the next section we combine these concepts to present a general and efficient accelerated coordinate descent scheme.
3.2.1 Gradient Descent

Given an initial point $x_0 \in \mathbb{R}^n$ and step sizes $h_k \in \mathbb{R}$, the *gradient descent method* applies the following simple iterative update rule:

\[
\forall k \geq 0 : \ x_{k+1} := x_k - h_k \nabla f(x_k).
\]

For $h_k = \frac{1}{k}$, this method simply chooses the minimum point of the upper envelope of $f$ at $x_k$:

\[
\tilde{x}_{k+1} = \arg\min_{\tilde{y}} \left\{ f(\tilde{x}_k) + \langle \nabla f(\tilde{x}_k), \tilde{y} - \tilde{x}_k \rangle + \frac{L}{2} \| \tilde{y} - \tilde{x}_k \|^2 \right\}.
\]

Thus, we see that the gradient descent method is a greedy method that chooses the minimum point based on the worst case estimate of the function based on the value of $f(x_k)$ and $\nabla f(x_k)$. It is well known that it provides the following guarantee [41, Cor 2.1.2, Thm 2.1.15]

\[
f(\tilde{x}_k) - f^* \leq \frac{L}{2} \cdot \min \left\{ \left( 1 - \frac{\sigma}{L} \right)^k, \frac{4}{k+4} \right\} \| \tilde{x}_0 - \bar{x} \|^2.
\] (3.3)

3.2.2 Accelerated Gradient Descent

To speed up the greedy and memory-less gradient descent method, one could make better use of the history and pick the next step to be the smallest point in upper envelope of all points computed. Formally, one could try the following update rule

\[
\tilde{x}_{k+1} := \arg\min_{\tilde{y} \in \mathbb{R}^n} \min_{\sum_{k=1}^n t_k \tilde{x}_k = \tilde{y}} \left\{ \sum_{k=1}^n t_k \left( f(\tilde{x}_k) + \langle \nabla f(\tilde{x}_k), \tilde{y} - \tilde{x}_k \rangle + \frac{L}{2} \| \tilde{y} - \tilde{x}_k \|^2 \right) \right\}.
\]

However, this problem is difficult to solve efficiently and requires storing all previous points. To overcome this problem, Nesterov [40, 41] suggested to use a quadratic function to estimate the function. Formally, we define an *estimate sequence* as follows:

\[^2\text{Note that our definition deviates slightly from Nesterov's [41, Def 2.2.1] in that we include condition 3.5.}\]
Definition 3.2.1 (Estimate Sequence). A triple of sequences \( \{\phi_k(x), \eta_k, \bar{x}_k\}_{k=0}^{\infty} \) is called an estimate sequence of \( f \) if \( \lim_{k \to \infty} \eta_k = 0 \) and for any \( \bar{x} \in \mathbb{R}^n \) and \( k \geq 0 \) we have
\[
\phi_k(\bar{x}) \leq (1 - \eta_k)f(\bar{x}) + \eta_k \phi_0(\bar{x})
\]
(3.4)
and
\[
f(\bar{x}_k) \leq \min_{\bar{x} \in \mathbb{R}^n} \phi_k(\bar{x}).
\]
(3.5)

An estimate sequence of \( f \) is an approximate lower bound of \( f \) which is slightly above \( f^* \). This relaxed definition allows us to find a better computable approximation of \( f \) instead of relying on the worst case upper envelope at each step.

A good estimate sequence gives an efficient algorithm [41, Lem 2.2.1] by the following
\[
\lim_{k \to \infty} f(\bar{x}_k) - f^* \leq \lim_{k \to \infty} \eta_k (\phi_0(\bar{x}^*) - f^*) = 0.
\]
Since an estimate sequence is an approximate lower bound, a natural computable candidate is to use the convex combination of lower envelopes of \( f \) at some points.

Since it can be shown that any convex combinations of lower envelopes at evaluation points \( \{y_k\} \) satisfies (3.4) under some mild condition, additional points \( \{y_k\} \) other than \( \{x_k\} \) can be used to tune the algorithm. Nesterov’s accelerated gradient descent method can be obtained by tuning the free parameters \( \{y_k\} \) and \( \{\eta_k\} \) to satisfy (3.5). Among all first order methods, this method is optimal up to constants in terms of number of queries made to \( f \) and \( \nabla f \). The performance of the accelerated gradient descent method can be characterized as follows: [41]
\[
f(\bar{x}_k) - f^* \leq L \cdot \min \left\{ \left(1 - \sqrt{\frac{\sigma}{L}}\right)^k, \frac{4}{(k+2)^2} \right\} \|\bar{x}_0 - \bar{x}^*\|^2.
\]
(3.6)

3.2.3 Coordinate Descent

The coordinate descent method of Nesterov [43] is a variant of gradient descent in which only one coordinate of the current iterate is updated at a time. For a fixed \( \alpha \in \mathbb{R} \), each iteration \( k \) of coordinate descent consists of picking a random a random
coordinate $i_k \in [n]$ where

$$Pr[i_k = j] = P_\alpha(j) \quad \text{where} \quad P_\alpha(j) \overset{\text{def}}{=} \frac{S_\alpha}{L_{ik}}$$

and then performing a gradient descent step on that coordinate:

$$\bar{x}_{k+1} := \bar{x}_k - \frac{1}{L_{ik}} \tilde{f}_i(\bar{x}_k).$$

To analyze this algorithm's convergence rate, we define the norm $\| \cdot \|_{1-\alpha}$, its dual $\| \cdot \|_{1-\alpha}^*$, and the inner product $\langle \cdot , \cdot \rangle_{1-\alpha}$ which induces this norm as follows:

$$\| \bar{x} \|_{1-\alpha} \overset{\text{def}}{=} \sqrt{\sum_{i=1}^{n} L_i^{1-\alpha} \bar{x}_i^2} \quad \text{and} \quad \| \bar{x} \|_{1-\alpha}^* \overset{\text{def}}{=} \sqrt{\sum_{i=1}^{n} L_i^{-(1-\alpha)} \bar{x}_i^2}$$

and

$$\langle \bar{x}, \bar{y} \rangle_{1-\alpha} \overset{\text{def}}{=} \sum_{i=1}^{n} L_i^{1-\alpha} \bar{x}_i \bar{y}_i$$

and we let $\sigma_{1-\alpha}$ denote the convexity parameter of $f$ with respect to $\| \cdot \|_{1-\alpha}$.

Using the definition of coordinate-wise Lipschitz constant, each step can be shown to have the following guarantee on expected improvement [43]

$$f(\bar{x}_k) - \mathbb{E} [f(\bar{x}_{k+1})] \geq \frac{1}{2S_\alpha} (\| \nabla f(\bar{x}_k) \|_{1-\alpha}^*)^2.$$

and further analysis shows the following convergence guarantee coordinate descent [43]

$$\mathbb{E} [f(\bar{x}_k)] - f^* \leq \min \left\{ \frac{2S_\alpha}{k+4} \left( \max_{f(\bar{y}) \leq f(\bar{x}_0)} \| \bar{y} - \bar{x}^* \|_{1-\alpha}^2 \right), \left( 1 - \frac{\sigma_{1-\alpha}}{S_\alpha} \right)^k (f(\bar{x}_0) - f^*) \right\}.$$

### 3.3 General Accelerated Coordinate Descent

In this section, we present our general and iteration-efficient *accelerated coordinate descent method (ACDM)*. In particular, we show how to improve the asymptotic con-
vergence rate of any coordinate descent based algorithm without paying asymptotic cost in the running time due to increased cost in querying and updating a coordinate. We remark that the bulk of the credit for conceiving of such a method belongs to Nesterov [43] who provided a different proof of convergence for such a method for the $\alpha = 0$ case, however we note that changes to the algorithm were necessary to deal with the $\alpha = 1$ case used in all of our applications.

The rest of this section is structured as follows. In Section 3.3.1, we introduce and prove the correctness of general ACDM through what we call (probabilistic) estimation sequences, in Section 3.3.2, we present the numerical stability results we achieve for this method, and in Section 3.3.3, we show how to implement this method efficiently. In Appendix A.2, we include some additional details for the correctness proof and in Appendix A.3, we provide the details of the numerical stability proof.

3.3.1 ACDM by Probabilistic Estimate Sequences

Following the spirit of the estimate sequence proof of accelerated gradient descent [41], here we present a proof of ACDM convergence through what we call a (probabilistic) estimation sequence.

**Definition 3.3.1 ((Probabilistic) Estimate Sequence).** A triple of sequences denoted, 

\[ \{\phi_k(\mathbf{x}), \eta_k, \mathbb{E}_k\}_{k=0}^{\infty}, \]  

where $\phi_k : \mathbb{R}^n \rightarrow \mathbb{R}$ and $\mathbb{E}_k \in \mathbb{R}^n$ are chosen according to some probability distribution is called a (probabilistic) estimate sequence of $f$ if $\lim_{k \rightarrow 0} \eta_k = 0$ and for all $k \geq 0$ we have

\[ \forall \mathbf{x} \in \mathbb{R}^n : \mathbb{E}[\phi_k(\mathbf{x})] \leq (1 - \eta_k)f(\mathbf{x}) + \eta_k\mathbb{E}[\phi_0(\mathbf{x})] \tag{3.7} \]

and

\[ \mathbb{E}[f(\mathbb{E}_k)] \leq \min_{\mathbf{x} \in \mathbb{R}^n} \mathbb{E}[\phi_k(\mathbf{x})] \tag{3.8} \]

A probabilistic estimation sequence gives a randomized minimization method due to the following

\[ \lim_{k \rightarrow \infty} \mathbb{E}[f(\mathbb{E}_k)] - f^* \leq \lim_{k \rightarrow \infty} \eta_k (\mathbb{E}[\phi_0(x^*)] - f^*) = 0. \]
Since a probabilistic estimation sequence can be constructed using random partial derivatives, rather than a full gradient computations, there is hope that in certain cases probabilistic estimation sequences require less information for fast convergence and therefore outperform their deterministic counterparts.

Similar to the accelerated gradient descent method, in the following lemma we first show how to combine a sequence of lower envelopes to satisfy condition (3.7) and prove that it preserves a particular structure on the current lower bound.

**Lemma 3.3.2** (Probabilistic) Estimate Sequence Construction. Let \( \phi_0(\vec{x}) \in \mathbb{R}^n \to \mathbb{R} \) and \( \{\vec{y}_k, \theta_k, i_k\}_{k=0}^\infty \) be such that

- Each \( \vec{y}_k \in \mathbb{R}^n \)
- Each \( \theta_k \in (0,1) \) and \( \sum_{k=0}^\infty \theta_k = \infty \)
- Each \( i_k \in [n] \) is chosen randomly such that \( \forall i \in [n] \) we have \( \Pr[i_k = i] = \frac{L_i^2}{8n} \).

Then the pair of sequences \( \{\phi_k(\vec{x}), \eta_k\}_{k=0}^\infty \) defined by

- \( \eta_0 = 1 \) and \( \eta_{k+1} = (1 - \theta_k)\eta_k \)
- \( \phi_{k+1}(\vec{x}) = (1 - \theta_k)\phi_k(\vec{x}) + \theta_k \left[ f(\vec{y}_k) + \frac{S_{i_k}}{L_{i_k}} (\vec{f}_{i_k}(\vec{y}_k), \vec{x} - \vec{y}_k) + \frac{\sigma_1 - \alpha}{2} \|\vec{x} - \vec{y}_k\|_{1-\alpha}^2 \right] \)

satisfies condition (3.7). Furthermore, if \( \phi_0(\vec{x}) = \phi_0^* + \frac{\zeta_0}{2} \|\vec{e}_0\|_{1-\alpha}^2 \), then this process produces a sequence of quadratic functions of the form \( \phi_k(\vec{x}) = \phi_k^* + \frac{\zeta_k}{2} \|\vec{x} - \vec{e}_k\|_{1-\alpha}^2 \) where

\[
\zeta_{k+1} = (1 - \theta_k)\zeta_k + \theta_k \sigma_{1-\alpha} \tag{3.9}
\]

\[
v_{k+1} = \frac{1}{\zeta_{k+1}} \left[ (1 - \theta_k)\zeta_k v_k + \theta_k \sigma_{1-\alpha} \vec{y}_k - \frac{S_{i_k}}{L_{i_k}} \vec{f}_{i_k}(\vec{y}_k) \right] \tag{3.10}
\]

\[
\phi_{k+1}^* = (1 - \theta_k)\phi_k^* + \theta_k f(\vec{y}_k) - \frac{\theta_k^2 S_{i_k}^2 (\vec{f}_{i_k}(\vec{y}_k))^2}{2\zeta_{k+1} L_{i_k}^{1+\alpha}}
+ \frac{\theta_k (1 - \theta_k)\zeta_k}{\zeta_{k+1}} \left( \frac{\sigma_{1-\alpha}}{2} \|\vec{y}_k - \vec{v}_k\|_{1-\alpha}^2 + \frac{S_{i_k} (\vec{f}_{i_k}(\vec{y}_k), \vec{v}_k - \vec{y}_k)_{1-\alpha}}{L_{i_k}} \right).
\]

**Proof.** The proof follows from direct calculations however for completeness and for later use we provide a proof of an even more general statement in Appendix A.1. \( \square \)
In the following theorem, we show how to choose $\bar{x}$, $\bar{y}$ and $\theta$ to satisfy the condition (3.8) and thereby derive a simple form of the general accelerated coordinate descent method. We also show that the number of iterations required for ACDM is $\tilde{O}\left(\frac{S_{\alpha}}{\sigma_{1-\alpha}}\right)$ which is strictly better than the number of iterations required for coordinate descent method, $\tilde{O}\left(\frac{S_{\alpha}}{\sigma_{1-\alpha}}\right)$. Note that while several of the definitions specifications in the following theorem statement may at first glance seem unnatural our proof will show that they are nearly forced in order to achieve certain algorithm design goals.

**Theorem 3.3.3** (Simple ACDM). For all $i \in [n]$ let $\bar{L}_i \overset{\text{def}}{=} \max(L_i^x, S_{\alpha}/n)$ \(^3\) and let $\bar{S}_{\alpha} \overset{\text{def}}{=} \sum_{i=1}^{n} \bar{L}_i$. Furthermore, for any $\bar{x}_0 \in \mathbb{R}^n$ and for all $k \geq 0$, let

$$
\bar{y}_0 = \bar{x}_0 \quad \phi_0 = f(\bar{x}_0) \quad \phi_0(\bar{x}) = \phi_0^* + \frac{\sigma_{1-\alpha}}{2} \| \bar{x} - \bar{x}_0 \|^2_{1-\alpha} \quad \theta_k = \sqrt{\frac{\sigma_{1-\alpha}}{2 \bar{S}_{\alpha} n}}
$$

Then applying Lemma 3.3.2 with these parameters, $\bar{L}_i$ as the coordinate-wise gradient Lipshitz constants, and choosing $\bar{y}_k$ and $\bar{x}_k$ such that

$$
\forall k \geq 1 : \frac{\theta_k \zeta_k}{\zeta_{k+1}} (\bar{v}_k - \bar{y}_k) + \bar{v}_k - \bar{y}_k = 0 \quad \text{and} \quad \bar{x}_k = \bar{y}_{k-1} - \frac{1}{\bar{L}_{ik-1}} \bar{f}_{ik}(\bar{y}_{k-1})
$$

yields a probabilistic estimate sequence. This accelerated coordinate descent method satisfies

$$
\forall k \geq 0 : \mathbb{E} [f(\bar{x}_k)] - f^* \leq \left(1 - \frac{1}{2} \sqrt{\frac{\sigma_{1-\alpha}}{S_{\alpha} n}}\right)^k \left(f(\bar{x}_0) - f^* + \sigma_{1-\alpha} \| \bar{x}_0 - \bar{x}_0 \|^2_{1-\alpha}\right).
$$

(3.11)

**Proof.** By construction we know that condition (3.7) of probabilistic estimation sequences holds. It remains to show that $\bar{x}_k$ satisfies condition (3.8) and analyze the convergence rate. To prove (3.8), we proceed by induction to prove the following

\(^3\)We introduce this to avoid small sampling probabilities as they cause an issue in achieving the optimal $\tilde{O}\left(\sqrt{\frac{S_{\alpha n}}{\sigma_{1-\alpha}}}\right)$ convergence rate. This modification can be avoided by choosing a different distribution over coordinate updates in compute $x_{k+1}$ which makes the algorithm more complicated and potentially more expensive.

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\[ \forall k \geq 0 : \mathbb{E}_k [f(\bar{x}_k)] \leq \mathbb{E}_k \left[ \min_{\bar{x} \in \mathbb{R}^n} \phi_k(\bar{x}) \right] = \mathbb{E}_k [\phi_k^*] . \]

where \( \mathbb{E}_k \) indicates the expectation up to iteration \( k \). The base case \( f(\bar{x}_0) \leq \phi_0(\bar{x}_0) \) is trivial and we proceed by induction assuming that \( \mathbb{E}_k [f(\bar{x}_k)] \leq \mathbb{E}_k [\phi_k^*] \). By Lemma (3.3.2) and the inductive hypothesis we get:

\[
\mathbb{E}_k [\phi_{k+1}^*] \geq \mathbb{E}_k \left[ (1 - \theta_k) f(\bar{x}_k) + \theta_k f(\bar{y}_k) - \frac{\theta_k^2 \tilde{S}_0^2 (f_{i_k}(\bar{y}_k))^2}{2 \zeta_{k+1}^\frac{1}{1+\alpha}} \right.
\]

\[
+ \frac{\theta_k (1 - \theta_k) \zeta_{k+1}}{\xi_{k+1}} \left( \frac{\sigma_1 - \alpha}{2} \| \bar{y}_k - \bar{y}_k \|_1^2 + \frac{\tilde{S}_0}{L_{i_k}} \right) f(\bar{y}_k), \frac{\theta_k \zeta_k}{\xi_{k+1}} (\bar{y}_k - \bar{y}_k) \right] + \langle \nabla f(\bar{y}_k), \bar{x}_k - \bar{y}_k \rangle \right] \]

where for notational convenience we drop the expectation in each of the variables.

By convexity \( f(\bar{x}_k) \geq f(\bar{y}_k) + \langle \nabla f(\bar{y}_k), \bar{x}_k - \bar{y}_k \rangle \) so applying this and the definitions of \( \| \cdot \|_{1-\alpha} \) and \( \langle \cdot, \cdot \rangle_{1-\alpha} \) we get that \( \phi_{k+1}^* \) is lower bounded by

\[
f(\bar{y}_k) - \frac{\theta_k^2 \tilde{S}_0^2 f_{i_k}(\bar{y}_k)^2}{2 \zeta_{k+1}^\frac{1}{1+\alpha}} + (1 - \theta_k) \left( \frac{\tilde{S}_0}{L_{i_k}} \left( \frac{\theta_k \zeta_k}{\xi_{k+1}} (\bar{y}_k - \bar{y}_k) \right) + \langle \nabla f(\bar{y}_k), \bar{x}_k - \bar{y}_k \rangle \right) \]

Using that \( \forall i \in [n], \Pr[i_k = i] = \frac{L_i}{\tilde{S}_0} \) we get that \( \mathbb{E}_{k+1} [\phi_{k+1}^*] \) is lower bounded by

\[
\mathbb{E}_{k+1} \left[ f(\bar{y}_k) - \frac{\theta_k^2 \tilde{S}_0^2 (f_{i_k}(\bar{y}_k))^2}{2 \zeta_{k+1}^\frac{1}{1+\alpha}} \right] + (1 - \theta_k) \left( \nabla f(\bar{y}_k), \frac{\theta_k \zeta_k}{\xi_{k+1}} (\bar{y}_k - \bar{y}_k) + \bar{x}_k - \bar{y}_k \right) \right] \]

From this formula we see that \( \bar{y}_k \) was chosen specifically to cancel the second term so

\[
\mathbb{E}_{k+1} [\phi_{k+1}^*] \geq \mathbb{E}_{k+1} \left[ f(\bar{y}_k) - \frac{\theta_k^2 \tilde{S}_0^2 (f_{i_k}(\bar{y}_k))^2}{2 \zeta_{k+1}^\frac{1}{1+\alpha}} \right] = \sum_{i=1}^n \left[ f(\bar{y}_k) - \frac{\theta_k^2 \tilde{S}_0^2 f_i(\bar{y}_k)^2}{2 \zeta_{k+1}^\frac{1}{1+\alpha}} \right] \]

and it simply remains to choose \( \theta_k \) and \( \bar{x}_{k+1} \) so that \( \mathbb{E}_{k+1} [f(\bar{x}_{k+1})] \) is smaller than this quantity.

To meet condition (3.8), we simply need to choose \( \{\theta_k\}_{k=0}^\infty \) so \( \frac{\theta_k \zeta_k}{\xi_{k+1}} = \frac{1}{2\eta} \). Using
\( \bar{L}_k \geq S_{\alpha} \frac{n}{\sqrt{n}} \geq \frac{\bar{S}_\alpha}{2 \sqrt{n}} \), we have

\[
\mathbb{E}_{k+1} [\phi_{k+1}^*] \geq \sum_{i=1}^{n} \left[ f(\bar{y}_k) - \frac{1}{2 \bar{S}_\alpha} \frac{f_i(\bar{y}_k)^2}{\bar{L}_i^{1-\alpha}} \right]
\]

To compute \( \bar{x}_{k+1} \), we use the fact that applying Lemma 3.1.1 to the formula \( f(\bar{y}_k - t \bar{f}_i(\bar{y}_k)) \) yields

\[
f\left( \bar{y}_k - \frac{1}{\bar{L}_i} \bar{f}_i(\bar{y}_k) \right) \leq f(\bar{x}) + \left( \bar{f}_i(\bar{y}_k), -\frac{1}{\bar{L}_i} \bar{f}_i(\bar{y}_k) \right) + \frac{\bar{L}_i}{2} \left( \frac{1}{\bar{L}_i} \bar{f}_i(\bar{y}_k) \right)^2 \leq f(\bar{y}_k) - \frac{f_i(\bar{y}_k)^2}{2 \bar{L}_i}
\]

and therefore for \( \bar{x}_{k+1} \) as defined we have

\[
\mathbb{E}_{k+1} [f(\bar{x}_{k+1})] \leq \sum_{i=1}^{n} \left[ f(\bar{y}_k) - \frac{f_i(\bar{x})^2}{2 \bar{S}_\alpha \bar{L}_i^{1-\alpha}} \right] \leq \mathbb{E}_{k+1} [\phi_{k+1}^*].
\]

Recalling that \( \zeta_{k+1} = (1 - \theta_k) \zeta_k + \theta_k \sigma_{1-\alpha} \) we see that choosing \( \zeta_0 = \sigma_{1-\alpha} \) implies \( \zeta_k = \sigma_{1-\alpha} \) for all \( k \) and therefore choosing \( \theta_k = \sqrt{\frac{\sigma_{1-\alpha}}{2 \bar{S}_\alpha n}} \) completes the proof that the chosen parameters produce a probabilistic estimate sequence. Furthermore, we see that this choice implies that \( \eta_k = \left( 1 - \sqrt{\frac{\sigma_{1-\alpha}}{2 \bar{S}_\alpha n}} \right)^k \). Therefore, by the definition of a probabilistic estimate sequence and the fact that \( \bar{S}_\alpha \geq 2 \bar{S}_\alpha \), equation (3.11) follows.

\[ \Box \]

### 3.3.2 Numerical Stability

In the previous section, we provided a simple proof how to achieve an ACDM with a convergence rate of \( \tilde{O}\left(\sqrt{\frac{\bar{S}_\alpha}{\sigma_{1-\alpha}}}\right) \). While sufficient for many purposes, the algorithm does not achieve the ideal dependence on initial error for certain regimes. For consistency with [43] we perform the change of variables

\[
\forall k \geq 0 : \alpha_k \overset{\text{def}}{=} \frac{\theta_k \zeta_k}{\zeta_k + \theta_k \sigma_{1-\alpha}}, \quad \beta_k \overset{\text{def}}{=} \frac{(1 - \theta_k) \zeta_k}{\zeta_k + \theta_k \sigma_{1-\alpha}}, \quad \gamma_k \overset{\text{def}}{=} \frac{\bar{S}_\alpha \theta_k}{\zeta_k + \theta_k \sigma_{1-\alpha}}
\]

and by better tuning \( \theta_k \) we derive the following algorithm.
**Accelerated Coordinate Descent Method**

1. Define $L_i = \max(L_i, (S_i/n)^{1/\alpha})$ and $\tilde{S}_i = \sum L_i$

2. Define $\tilde{x}_0 = x_0, a_0 = \frac{1}{2n}, b_0 = 2$

3. For $k \geq 0$ iterate:
   
   3a. Find $\alpha_k, \beta_k, \gamma_k \geq \frac{1}{2n}$ such that $\gamma_k^2 - \frac{\gamma_k}{2n} = \left(1 - \frac{\gamma_k a_k - a_k}{\tilde{S}_i}\right) \frac{a_k^2}{b_k^2} = \beta_k b_k^2 = \beta_k \gamma_k \frac{1 - a_k}{a_k}$.

   3b. $\tilde{y}_k = \alpha_k \tilde{u}_k + (1 - \alpha_k)\tilde{x}_k$.

   3c. Choose $i_k$ according to $P_{\alpha}(i) = \frac{L_i}{\tilde{S}_i}$.

   3d. $\tilde{x}_{k+1} = \tilde{y}_k - \frac{1}{L_{\tilde{i}_k}} \tilde{f}_{i_k}(\tilde{y}_k), \tilde{v}_{k+1} = \beta_k \tilde{u}_k + (1 - \beta_k)\tilde{y}_k - \frac{\gamma_k}{L_{\tilde{i}_k}} \tilde{f}_{i_k}(\tilde{y}_k)$.

   3e. $b_{k+1} = \frac{b_k}{\sqrt{\gamma_k}}$ and $a_{k+1} = \gamma_k b_{k+1}$.

In Appendix A.2 and A.3, we give a proof of convergence of this method in the unit-cost RAM model by studying the following potential function [43] for suitable constants $a_k, b_k \in \mathbb{R}$,

$$a_k \left( E[f(\tilde{x}_k)] - f^* \right) + b_k E \left[ \| \tilde{u}_k - x_v \|_{1-\alpha}^2 \right].$$

The following theorem states that error in vector updates does not grow too fast and Lemma A.2.1 further states that errors in the coefficients $\alpha, \beta,$ and $\gamma$ also does not grow too fast. Hence, $O(\log n)$ bits of precision is sufficient to implement ACDM with the following convergence guarantees. Below we state this result formally and we refer the reader to Appendix A.3 for the proof.

**Theorem 3.3.4** (Numerical Stability of ACDM). Suppose that in each iteration of ACDM step 3d has additive error $\varepsilon$, i.e. there exists $\tilde{\varepsilon}_{1,k}, \tilde{\varepsilon}_{2,k} \in \mathbb{R}^n$ with $\| \tilde{\varepsilon}_{1,k} \|_{1-\alpha} \leq \varepsilon$ and $\| \tilde{\varepsilon}_{2,k} \|_{1-\alpha} \leq \varepsilon$ such that step 3 is

$$\tilde{x}_{k+1} := \tilde{y}_k - \frac{1}{L_{\tilde{i}_k}} \tilde{f}_{i_k}(\tilde{y}_k) + \tilde{\varepsilon}_{1,k} \quad \text{and} \quad \tilde{v}_{k+1} := \beta_k \tilde{u}_k + (1 - \beta_k)\tilde{y}_k - \frac{\gamma_k}{L_{\tilde{i}_k}} \tilde{f}_{i_k}(\tilde{y}_k) + \tilde{\varepsilon}_{2,k}.$$

If $\varepsilon < \frac{\sigma_{1-\alpha}^2}{8\tilde{S}_i}$ and $k \geq \sqrt{\frac{2\tilde{S}_i n}{\sigma_{1-\alpha}}}$, then we have the convergence guarantee

$$\sigma_{1-\alpha} E \left[ \| \tilde{v}_{k+1} - \tilde{x}_v \|_{1-\alpha}^2 \right] + (E[f(\tilde{x}_{k+1})] - f^*) \leq \delta_k$$

(3.12)
\[ \delta_k \overset{\text{def}}{=} 24kS_\alpha \epsilon^2 + 32\sigma_{1-\alpha} \left( 1 - \frac{1}{5} \sqrt{\frac{\sigma_{1-\alpha}}{S_\alpha n}} \right)^k \left( \|x_0 - x^*\|_{1-\alpha} + \frac{1}{S_\alpha^2} (f(x_0) - f^*) \right) \]

where

and the additional convergence guarantee that \( \frac{1}{k} \sum_{j=k}^{2k-1} \|\nabla f(\bar{y}_k)\|_{1-\alpha}^2 \leq 2000 \frac{S_\alpha}{n} \delta_k. \)

This theorem provides useful estimates for the error \( f(\bar{x}_{k+1}) - f^* \), the residual \( \|\vec{v}_{k+1} - \vec{x}^*\|_{1-\alpha}^2 \), and the norm of gradient \( \|\nabla f(\bar{y}_k)\|_{1-\alpha}^* \). Note how the estimate depends on the initial error \( f(\bar{x}_0) - f^* \) mildly as compared to Theorem 3.3.3. We use this fact in creating an efficient SDD solver in Section 3.4.3.

### 3.3.3 Efficient Iteration

In both Nesterov's paper [43] and later work [45] the original ACDM proposed by Nesterov was not recommend since a naive implementation takes \( O(n) \) time to update the vector \( \vec{v}_k \), and therefore is likely slower than accelerated gradient descent. This implementation issue is a potential serious problem under the assumption that the oracle to compute gradient can only accept a single vector as input. However, if we make the mild assumption that we can compute \( \nabla f(t\vec{x} + s\vec{y}) \) for \( s, t \in \mathbb{R} \) and \( \vec{y}, \vec{y} \in \mathbb{R} \) in the same asymptotic runtime as it takes to compute \( \nabla f(\bar{x}) \) (i.e. we do not need to compute the sum explicitly), then we can implement ACDM without additional asymptotic computational costs per iteration as compared to the cost of the coordinate descent method performing an update on the given coordinate. In the following lemma, we prove this fact and show that the technique can be executed in the unit-cost RAM model without additional asymptotic costs. Note that the method we propose does introduce potential numerical problems which we argue are minimal.

**Lemma 3.3.5 (Efficient Iterations).** For \( S_\alpha = O(\text{poly}(n)) \) and \( \sigma_{1-\alpha} = \Omega(\text{poly}(\frac{1}{n})) \) each iteration of ACDM can be implemented in \( O(1) \) time plus the time to make one oracle call of the form \( \bar{f}_i(t\bar{x} + s\bar{y}) \) for \( s, t \in \mathbb{R} \) and \( \bar{y}, \bar{y} \in \mathbb{R}^n \), using at most an additional \( O(\log n) \) bits of precision so long as the number of iterations of ACDM is \( O\left( \sqrt{\frac{S_\alpha n}{\sigma_{1-\alpha}}} \log(n) \right) \).
Proof. By the specification of the ACDM algorithm, we have

\[
\begin{align*}
\tilde{y}_{k+1} &= \alpha_{k+1} \tilde{v}_{k+1} + (1 - \alpha_{k+1}) \tilde{x}_{k+1} \\
&= \alpha_{k+1} \left( \beta_k \tilde{v}_k + (1 - \beta_k) \tilde{y}_k + \frac{\gamma_k}{L_{ik}} \tilde{f}_i(\tilde{y}) \right) + (1 - \alpha_{k+1}) \left( \tilde{y}_k - \frac{1}{L_{ik}} \tilde{f}_i(\tilde{y}) \right) \\
&= \alpha_{k+1} \beta_k \tilde{v}_k + (\alpha_{k+1} (1 - \beta_k) + (1 - \alpha_{k+1})) \tilde{y}_k - \frac{1 - \alpha_{k+1} - \alpha_{k+1} \gamma_k}{L_{ik}} \tilde{f}_i(\tilde{y}) \\
\end{align*}
\]

we see that each update step of ACDM can be rewritten as

\[
\begin{pmatrix}
\tilde{v}_{k+1}^T \\
\tilde{y}_{k+1}^T
\end{pmatrix}
= A_k
\begin{pmatrix}
\tilde{v}_k^T \\
\tilde{y}_k^T
\end{pmatrix}
- s_k
\]

where

\[
A_k = \begin{pmatrix}
\beta_k & 1 - \beta_k \\
\alpha_{k+1} \beta_k & 1 - \alpha_{k+1} \beta_k
\end{pmatrix}
\quad\text{and}\quad
s_k = \begin{pmatrix}
\frac{2k}{L_{ik}} \tilde{f}_i(\tilde{y})^T \\
\frac{1 - \alpha_{k+1} - \alpha_{k+1} \gamma_k}{L_{ik}} \tilde{f}_i(\tilde{y})^T
\end{pmatrix}
\]

Therefore to implement ACDM in each iteration we can just maintain vectors \( \tilde{v}, \tilde{y} \in \mathbb{R}^n \) and matrix \( B_k \in \mathbb{R}^{2 \times 2} \) such that

\[
\begin{pmatrix}
\tilde{v}_k^T \\
\tilde{y}_k^T
\end{pmatrix}
= B_k
\begin{pmatrix}
\tilde{v}^T \\
\tilde{y}^T
\end{pmatrix}
\]

With this representation each update step is

\[
B_{k+1} \begin{pmatrix}
\tilde{v}^T \\
\tilde{y}^T
\end{pmatrix}
= A_k B_k \begin{pmatrix}
\tilde{v}^T \\
\tilde{y}^T
\end{pmatrix}
- s_k
\]

and by our oracle assumption, we can implement the following equivalent update step

\[
B_{k+1} := A_k B_k \quad\text{and}\quad
\begin{pmatrix}
\tilde{v}^T \\
\tilde{y}^T
\end{pmatrix}
:= \begin{pmatrix}
\tilde{v}^T \\
\tilde{y}^T
\end{pmatrix}
- B_{k+1}^{-1} s_k
\]

in time \( O(1) \) plus the time needed for one oracle call.

To complete the lemma, we simply need to show that this scheme is numerically stable. Note that

\[
\det(A_k) = \beta_k (1 - \alpha_{k+1} \beta_k) - \alpha_{k+1} \beta_k (1 - \beta_k) = \beta_k (1 - \alpha_{k+1})
\]

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Now by Lemma A.2.1, we know that $\alpha_k \leq 32 \max \left\{ \frac{\sigma_1 - \alpha}{k}, \sqrt{\frac{\sigma_1 - \alpha}{25n}} \right\}$ for $k > 1$ and we know that $\beta_k \geq 1 - \frac{1}{2} \sqrt{\frac{\sigma_1 - \alpha}{25n}}$ for all $k$. Therefore, letting $\kappa \overset{\text{def}}{=} \sqrt{\frac{\sigma_1 - \alpha}{25n}}$ and recalling that we assumed that the total number of iterations is $O(\kappa^{-1} \log n)$ we get

$$
\det(B_k) \geq (1 - \kappa)^{O(\kappa^{-1} \log n)} (1 - 32\kappa)^{O(\kappa^{-1} \log n)} \prod_{k=2}^{\kappa-1} \left(1 - \frac{32}{k}\right)
$$

$$
= \Omega \left( \text{poly} \left( \frac{1}{n} \right) e^{-32 \log \kappa} \right) = \Omega \left( \text{poly} \left( \frac{1}{n} \right) \right).
$$

Hence, $O(\log n)$ bits of precision suffice to implement this method. 

\[\square\]

### 3.4 Faster Linear System Solvers

In this section, we show how ACDM can be used to achieve asymptotic runtimes that outperform various state-of-the-art methods for solving linear systems in a variety of settings. In Section 3.4.1, we show how to use coordinate descent to outperform conjugate gradient under minor assumptions regarding the eigenvalues of the matrix under consideration, in Section 3.4.2, we show how to improve the convergence rate of randomized Kaczmarz type methods, and in Section 3.4.3, we show how to use the ideas to achieve the current fastest known numerically stable solver for symmetric diagonally dominant (SDD) systems of equations.

#### 3.4.1 Comparison to Conjugate Gradient Method

Here we compare the performance of ACDM to conjugate gradient (CG) and show that under mild assumptions about the linear system being solved ACDM achieves a better asymptotic running time.

For symmetric positive definite (SPD) matrix $A \in \mathbb{R}^{n \times n}$ and vector $\bar{b} \in \mathbb{R}^n$, we solve the linear system of equations $A\bar{x} = \bar{b}$ via the following equivalence unconstrained quadratic minimization problem:

$$
\min_{\bar{x} \in \mathbb{R}^n} f(\bar{x}) \overset{\text{def}}{=} \frac{1}{2} \langle A\bar{x}, \bar{x} \rangle - \langle \bar{b}, \bar{x} \rangle.
$$

(3.13)
Let \( m \) denote the number of nonzero entries in \( A \), let \( \text{nnz}_i \) denote the number of nonzero entries in the \( i \)th row of \( A \). To make the analysis simpler, we assume that the nonzero entries is somewhat uniform, namely \( \forall i \in [n] \) we have \( \text{nnz}_i = O\left(\frac{m}{n}\right) \). This assumption is trivially met for dense matrices, finite difference matrices, etc. Letting \( 0 \leq \lambda_1 \leq \ldots \leq \lambda_n \) denote the eigenvalues of \( A \), we get the following running time guarantee for ACDM.

**Theorem 3.4.1 (ACDM on SPD Systems).** Assume \( A \) is a SPD matrix with the property that \( \text{nnz}_i = O\left(\frac{m}{n}\right) \) for all \( i \in [n] \). Let the numerical rank \( r(A) = \sum_{i=1}^{n} \lambda_i / \lambda_n \).

ACDM applied to (3.13) with \( \alpha = 1^4 \) produces an approximate solution in

\[
\tilde{O} \left( m \sqrt{\frac{r(A)}{n}} \sqrt{\frac{\lambda_n}{\lambda_1}} \log \frac{1}{\epsilon} \right)
\]

time with \( \epsilon \) error in \( A \) norm in expectation.

**Proof.** The running time of ACDM with \( \alpha = 1 \) depends on \( \sigma_0 \) and \( S_1 \). From Example 3.1.3, the total component-wise Lipschitz constant \( S_1 \) is the trace of \( A \), which is \( \sum \lambda_i \) and the convexity parameter \( \sigma_0 \) is \( \lambda_1 \), therefore by Theorem 3.3.4 the convergence rate of ACDM is \( \sqrt{\frac{\lambda_1}{n \sum_{i=1}^{n} \lambda_i}} \) as desired. Furthermore, the running time of each step depends on the running time of the oracle, i.e. computing \( f_i(x) = (Ax)_i \), which by our assumption on \( \text{nnz}_i \) takes time \( O\left(\frac{m}{n}\right) \).

To compare with conjugate gradient, we know that one crude bound for the rate of convergence of conjugate gradient is \( O\left(\sqrt{\frac{\lambda_1}{\lambda_n}}\right) \). Hence the total running time of CG to produce an epsilon approximate solution is \( \tilde{O} \left( m \sqrt{\frac{\lambda_n}{\lambda_1}} \log \frac{1}{\epsilon} \right) \). Therefore, with this bound ACDM is always faster or matches the running time since the numerical rank of \( A \) is always less than or equals to \( n \). Thus, we see that when the numerical rank of \( A \) is \( o(n) \), ACDM will likely \(^5\) have a faster asymptotic running time.

To be more fair in our comparison to conjugate gradient, we note that in [54] tighter bound on the performance of CG was derived and they showed that in fact

\(^4\)Here we only consider the \( \alpha = 1 \) case for easier comparison with conjugate gradient.

\(^5\)The running time of CG may be asymptotic faster when the eigenvalues form clusters.

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CG has a running time of $\tilde{O}\left(\sum \text{nnz}_i \left(\frac{\sum_{j=1}^{\text{nnz}_i} \lambda_j}{\lambda_1}\right)^{1/3}\right)$ implying that ACDM is faster than CG when $\sum_{i=1}^{n} \lambda_i \leq n^3 \lambda_1$ and it is usually satisfied. In the extreme cases that the condition is false, CG will need to run for $O(n)$ iterations at which point an exact answer could be computed.

3.4.2 Accelerating Randomized Kaczmarz

The Kaczmarz method [24] is an iterative algorithm to solve $A\vec{x} = b$ for any full row rank matrix $A \in \mathbb{R}^{m \times n}$. Letting $a_i \in \mathbb{R}^n$ denote the $i$-th row of the matrix $A$, we know that the solution of $A\vec{x} = \vec{b}$ is the intersection of the hyperplanes $H_i \overset{\text{def}}{=} \{x : \langle a_i, \vec{x} \rangle = b_i\}$. The Kaczmarz method simply iteratively picks one of these hyperplanes and projects onto it by the following formula:

$$x_{k+1} := \text{proj}_{H_{i_k}}(\vec{x}_k) \text{ where } \text{proj}_{H_{i_k}}(\vec{x}_k) + \frac{\vec{b}_{i_k} - \langle a_{i_k}, \vec{x}_k \rangle}{\|a_{i_k}\|^2}a_{i_k}.$$

There are many schemes that can be chosen to pick the hyperplane $i_k$, many of which are difficult to analyze and compare, but in a breakthrough result, Strohmer and Vershynin in 2008 analyzed the randomized schemes which sample the hyperplane with probability proportional to $\|a_i\|^2$. They proved the following

**Theorem 3.4.2** (Strohmer and Vershynin [55]). *The Kaczmarz method samples row $i$ with probability proportionally to $\|a_i\|^2$ at each iteration and yields the following

$$\forall k \geq 0 : \mathbb{E}\left[\|\vec{x}_k - \vec{x}^*\|_2^2\right] \leq (1 - \kappa(A)^{-2})^k \|\vec{x}_0 - \vec{x}^*\|^2$$

where $\vec{x}^* \in \mathbb{R}^n$ is such that $A\vec{x}^* = \vec{b}$, $\kappa(A) \overset{\text{def}}{=} \|A^{-1}\|_2 \cdot \|A\|_F$ is the relative condition number of $A$, $A^{-1}$ is the left inverse of $A$, $\|A^{-1}\|_2$ is the smallest non-zero spectral value of $A$ and $\|A\|_F^2 \overset{\text{def}}{=} \sum a_{ij}^2$ is the Frobenius norm of $A$.*

Here we show show to cast this algorithm as an instance of coordinate descent and obtain an improved convergence rate by applying ACDM. We remark that accelerated Kaczmarz will sample rows with a slightly different probability distribution. As long
as this does not increase the expected computational cost of an iteration, it will yield an algorithm with a faster asymptotic running time.

**Theorem 3.4.3** (Accelerated Kaczmarz). The ACDM method samples row \(i\) with probability proportionally to \(\max\left\{\|a_i\|_2^2, \frac{\|A\|^2}{m}\right\}\) and performs extra \(O(1)\) work at each iteration. It yields the following

\[
\forall k \geq 0 : \mathbb{E}\|\tilde{x}_k - \tilde{x}^*\|_2^2 \leq 3 \left(1 - \frac{\kappa(A)^{-1}}{2\sqrt{m}}\right)^k \|\tilde{x}_0 - \tilde{x}^*\|_2^2 .
\]

**Proof.** To cast Strohmer and Vershynin's randomized Kaczmarz algorithm in the framework of coordinate descent, we consider minimizing the objective function of theorem directly, i.e.

\[
\min_{y \in \mathbb{R}^m} 2\frac{1}{2} \|\mathbf{A}y - \mathbf{b}\|_2^2.
\]

Since \(\mathbf{A}\) has full row rank, we write \(\mathbf{x} = \mathbf{A}^T \tilde{y}\) and consider the equivalent problem \(\min_{y \in \mathbb{R}^m} 2\frac{1}{2} \|\mathbf{A}^T \tilde{y} - \mathbf{x}^*\|_2^2\). Expanding the objective function and using \(\mathbf{A}\tilde{x}^* = \tilde{b}\), we get

\[
\|\mathbf{A}^T \tilde{y} - \mathbf{x}^*\|_2^2 = \|\mathbf{A}^T \tilde{y}\|_2^2 - 2\langle \tilde{b}, \tilde{y} \rangle + \|\mathbf{x}^*\|_2^2 .
\]

Therefore, we attempt solve the following equivalent problem using accelerated coordinate descent.

\[
\min_{\tilde{y} \in \mathbb{R}^m} f(\tilde{y}) \quad \text{where} \quad f(\tilde{y}) \overset{\text{def}}{=} \frac{1}{2} \|\mathbf{A}^T \tilde{y}\|_2^2 - \langle \tilde{b}, \tilde{y} \rangle .
\]

From Example 3.1.4, we know that the \(i\)-th direction component-wise Lipschitz constant is \(L_i = \|a_i\|^2\) where \(a_i\) is the \(i\)-th row of \(\mathbf{A}\) and we know that \(\nabla f(\tilde{y}) = \mathbf{A}\mathbf{A}^T \tilde{y} - \mathbf{b}\). Therefore, each step of the coordinate descent method consists of the following

\[
\tilde{y}_{k+1} := \tilde{y}_k - \frac{1}{L_{i_k}} f_{i_k}(\tilde{y}_k) = \tilde{y}_k - \frac{1}{\|a_{i_k}\|^2} \left(\mathbf{A}\mathbf{A}^T \tilde{y}_k - \tilde{b}\right)_{i_k} .
\]

Recalling that we had performed the transformation \(\mathbf{x} = \mathbf{A}^T \tilde{y}\) we see that the corre-

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\(^6\)We ignore the thresholding here for illustration purpose.
sponding step in $\tilde{x}$ is

$$
\tilde{x}_{k+1} = \tilde{x}_k - \frac{1}{\|a_i\|^2} (A\tilde{x}_k - \tilde{b})_k = \tilde{x}_k + \frac{\tilde{b}_{ik} - \langle \tilde{a}_{ik}, \tilde{x}_k \rangle}{\|a_i\|_2^2} \tilde{a}_{ik}.
$$

Therefore, the randomized coordinate descent method applied this way yields precisely the randomized Kaczmarz method of Strohmer and Vershynin.

However, to apply the ACDM method and provide complete theoretical guarantees we need to address the problem that $f$ as we have constructed it is not strongly convex. This is clear by the fact that the null space of $A^T$ may be non trivial.

To remedy this problem we let $Z \subseteq \mathbb{R}^m$ denote the null space of $A^T$, i.e. $Z \overset{df}{=} \{ \tilde{x} \in \mathbb{R}^m \mid A^T \tilde{x} = 0 \}$, and we define the semi-norm $\| \cdot \|_Z$ on $\mathbb{R}^m$ by $\| y \|_Z \overset{df}{=} \inf_{z \in Z} \| y + z \|$. Now it is not hard to see that $f$ is strongly convex under this seminorm with convexity parameter $\sigma_Z = \| A^{-1} \|_2^{-2}$. Furthermore, one can prove similarly to the proof of Lemma 3.3.2 and Theorem 3.3.3 that the algorithm in this theorem achieves the desired convergence rate.

To see this, we let $P \in \mathbb{R}^{mxm}$ denote the orthogonal projection onto the image of $A$ we note that $\| \tilde{y} \|_Z = \sqrt{\tilde{y}_T P \tilde{y}}$. Therefore we can apply the general form of Lemma 3.3.2 proved in Appendix A.1 to derive a similar estimate sequence result for this norm. Unfortunately, the resulting estimate sequence requires working with $P^T f_i(k \tilde{y}_k)$. However, the proof of Theorem 3.3.3 still works using the same algorithm because of the facts that $P^T \nabla f(\tilde{x}) = \nabla f(\tilde{x})$,

$$
\forall \tilde{y} \in \mathbb{R}^n, \forall \tilde{z} \in Z : f(\tilde{y} + \tilde{z}) = f(\tilde{y}) \quad \text{and} \quad \nabla f(\tilde{y} + \tilde{z}) = \nabla(\tilde{y}) ,
$$

and $\| \tilde{x} \|_2^2 \geq \| \tilde{x} \|_Z^2$. This gives the same convergence guarantee with $\sigma_{1-\alpha}$ replaced with $\| A^{-1} \|_2^{-2}$, $S_{\alpha} = \| A \|_F^2$ and $\| \cdot \|_1-\alpha = \| \cdot \|_Z$. The desired convergence rate follows from the strong convexity of $f$.

To justify the cost of each iteration, note that although the iterations work on the $\tilde{y}$ variable, we can apply $A^T$ on all iterations and just do the operations on $\tilde{x}$ variables which is more efficient. \[ \square \]
3.4.3 Faster SDD Solvers in the Unit-cost RAM Model

Here we show how to cast the SimpleSolver algorithm of Chapter 2 as an instance of coordinate descent, and by applying ACDM we obtain an algorithm for solving SDD systems that has an asymptotic running time of \(O(m \log^{3/2} n \sqrt{\log \log n \log(e^{-1} \log n)})\) in the unit-cost RAM model. More precisely we prove the following.

**Theorem 3.4.4.** By applying ACDM to SimpleSolver, we can produce \(\vec{v} \in \mathbb{R}^V\) such that \(\|\vec{v} - L^T \vec{x}\|_L \leq \varepsilon \|L^T \vec{x}\|_L\) in time \(O(m \log^{3/2} n \sqrt{\log \log n \log(e^{-1} \log n)})\).\(^7\)

**Proof.** Recall that instead of computing \(\vec{v}\) directly, we can instead focus on the following electric flow problem \(^8\)

\[
\min_{B^T \vec{f} = \vec{x}} \frac{1}{2} \xi(\vec{f}) \quad \text{where} \quad \xi(\vec{f}) \overset{\text{def}}{=} \|\vec{f}\|_R^2 \overset{\text{def}}{=} \sqrt{\sum_{e \in E} r_e \vec{f}_e^2}
\]

Now to turn this into an unconstrained minimization problem we first let \(\vec{f}_0\) be feasible flow, i.e. a vector such that \(B^T \vec{f}_0 = \vec{x}\). With this, the problem can be simplified to

\[
\min_{B^T \vec{f} = \vec{x}} \frac{1}{2} \|\vec{f} + \vec{c}\|_R^2.
\]

However, from Section 2.2 we know that \(\{\vec{c} \in \mathbb{R}^E \mid B^T \vec{c} = 0\}\) is simply the set of circulations of the graph, i.e. *cycle space*, and for spanning tree \(T\) the set of tree cycles for the off tree edges, i.e. \(\{\vec{c}_e \mid e \in E \setminus T\}\) form a basis. Therefore, using this basis we see that we can simplify the problem further to

\[
\min_{\vec{g} \in \mathbb{R}^E \setminus T} \frac{1}{2} \|\vec{f} + C\vec{g}\|_R^2 \quad \text{where} \quad C \overset{\text{def}}{=} [c_{e_1}, c_{e_2}, \cdots] \in \mathbb{R}^{E \times E \setminus T}.
\]

Now, for every *off-tree edge*, i.e. \(e \in E \setminus T\), there is only one cycle \(c_e\) that passes through it. So, if we let \(R_{E \setminus T} \in \mathbb{R}^{E \setminus T \times E \setminus T}\) be the diagonal matrix for the resistances

\(^7\)Although not the focus of this section, we remark that this procedure produces an \(\varepsilon\)-approximate electric flow in time \(O(m \log^{3/2} n \sqrt{\log \log n \log(e^{-1} \log n)})\) in the unit-cost RAM model. Furthermore, by the definition of ACDM, we see that the operators produced by this algorithm are linear and by similar analysis as in \([30]\) the algorithm can be used to produce a linear approximation to \(L^T\).

\(^8\)To make the analysis cleaner we scale the objective function by \(\frac{1}{2}\). This has no substantial effect on our ability to compute approximate voltages from approximate solutions.
of the off tree edges we have that for any $\tilde{y} \in E \setminus T$

$$\tilde{y}^T \mathbf{C}^T \mathbf{R} \mathbf{C} \tilde{y} \geq \tilde{y}^T \mathbf{C}^T \mathbf{R}_{E \setminus T} \mathbf{C} \tilde{y} = \sum_{e \in E \setminus T} r_e \tilde{y}(e)^2 \geq \left( \min_{e \in E \setminus T} r_e \right) \|\tilde{y}\|_2^2.$$  

Therefore, the convexity parameter of $\|z_0 + \mathbf{C} \tilde{y}\|_R^2$ is at least $\min_{e \in E \setminus T} r_e$. However, this could be wasteful if the resistances vary, so rescale the space, $\tilde{y} = \mathbf{R}^{1/2} \tilde{y}$, to get the following problem:

$$\min_{\tilde{y} \in \mathbb{R}^E \setminus T} g(\tilde{y}) \text{ where } g(\tilde{y}) = \frac{1}{2} \left\| \mathbf{f}_0 + \mathbf{C} \mathbf{R}^{-1/2} \tilde{y} \right\|_R^2.$$  

By the reasoning above, the convexity parameter of $g$ with respect to the Euclidian norm is 1 and we bound the $e$-th direction component-wise Lipschitz constant by

$$\forall e \in E \setminus T : L_e = \frac{1}{r_e} \left\| \mathbf{R}^{-1/2} \mathbf{C}^T \mathbf{R} \mathbf{C} \mathbf{R}^{-1/2} \right\|_e = \frac{\mathbf{C} \mathbf{R}^{-1/2} \mathbf{e}_e}{r_e} = \frac{R_e}{r_e} = \text{st}(e) + 1$$

Therefore we see that

$$\frac{S_1}{\sigma_0} = S_1 = \sum_{e \in E \setminus T} \frac{R_e}{r_e} = \tau(T) = O(m + \text{st}(T))$$

Thereby justifying our naming of $\tau$ as "tree condition number."

Finally, applying the coordinate descent method to $g$ and letting $e_k \in E \setminus T$ denote the off-tree edge i.e., coordinate, picked in iteration $k$, we get

$$\tilde{y}_{k+1} := \tilde{y}_k - \frac{1}{L_{e_k}} \tilde{g}_{e_k}(\tilde{y}_k) \quad \text{(Definition of Update Step)}$$

$$= \tilde{y}_k - \frac{1}{L_{e_k}} \left( \mathbf{R}^{-1/2} \mathbf{C}^T \mathbf{R} (\mathbf{f}_0 + \mathbf{C} \mathbf{R}^{-1/2} \tilde{y}) \right)_{e_k} \cdot \mathbf{1}_{e_k} \quad \text{(Computation of } \nabla g(\tilde{y})\text{)}$$

$$= \tilde{y}_k - \frac{1}{L_{e_k} r_{e_k}^{1/2}} \sum_e \tilde{e}_{e_k}(e) r_{e_k}(\mathbf{f}_0 + \mathbf{C} \mathbf{R}^{-1/2} \tilde{y})(e) \cdot \mathbf{1}_{e_k} \quad \text{(Definition of } \mathbf{C} \text{ and } \mathbf{R})$$

$$= \tilde{y}_k - \frac{1}{L_{e_k} r_{e_k}^{1/2}} \sum_{e \in \tilde{e}_{e_k}} r_{e_k}(\mathbf{f}_0 + \mathbf{C} \mathbf{R}^{-1/2} \tilde{y})(e) \cdot \mathbf{1}_{e_k} \quad \text{(Definition of } C_{e_k})$$

---

9To avoid confusion between a flow vector and the objective function in just this section we use $g$ instead of $f$ to denote the objective function.
Recalling $\bar{y} = R^{-1/2}\bar{y}$ and the derivation of $L_e$, we can write this equivalently as

$$
\bar{y}_{k+1} := \bar{y}_k - \left[ \frac{1}{R_e} \sum_{e \in \mathcal{E}_k} r_e (\bar{f}_0 + C\bar{y})(e) \right] \cdot \bar{e}_k.
$$

Noting that the $\bar{f}$ with $B^T \bar{f} = \bar{\chi}$ corresponding to $\bar{y}$ is $\bar{f} = \bar{f}_0 + C\bar{y}$, we write the update equivalently as

$$
\bar{f}_{k+1} := \bar{f}_k - \left[ \frac{1}{R_e} \sum_{e \in \mathcal{E}_k} r_e \bar{f}_k(e) \right] \cdot \bar{e}_k,
$$

which is precisely the cycle update of SimpleSolver. Thus we see that SimpleSolver is an instantiation of coordinate descent where the data structure is used to make sure that calls to $g_{e_k}$ and updates to $\bar{y}_{e_k}$ can be implemented in $O(\log n)$. Therefore, by applying ACM we can obtain a faster algorithm. Note that to apply ACM efficiently computations of $g_{e_k}$ need to be performed on the sum of two vectors without explicitly summing. However, since here $\nabla g$ is linear, we can just call the oracle on the two vectors separately and use two separate data structure for updating coordinates.

While the above insight suffices to compute electric flows a little more work needs to be done to compute the desired voltages. However, by Lemma 2.6.2 we know that it suffices to show that $\|\nabla g(\bar{y})\|_2^2 \leq \varepsilon f^*$. Note that

$$
\|\nabla g(\bar{y})\|_2^2 = \|R^{-1/2}C^T R(\bar{f}_0 + C R^{-1/2}\bar{y})\|_2^2 = \sum_{e \in \mathcal{E}\setminus T} \frac{1}{r_e} \left( \sum_{e' \in \mathcal{E}_e} g(e') r_e' \right)^2
$$

and therefore if we choose $\bar{y}_0 = \bar{0}$. Then, we see that

$$
\|\bar{y}_0 - \bar{y}^*\|_2^2 = \|\bar{y}^*\|_2^2 = \|R^{1/2}_{E\setminus T} \bar{y}\|_2^2 \leq 2f^*,
$$

and using Lemma 2.6.1 we have that

$$
\frac{1}{s^2} (g(\bar{y}_0) - g^*) \leq \frac{g^*}{s} \leq g^*.
$$
Therefore so long as we choose our spanning tree using Theorem 2.2.13 we have \( S_{n} / |E(T)| = O(\log n \log \log n) \) and after \( k = O(m \sqrt{\log n \log \log n \log \log n}) \) iterations of ACDM, by Theorem 3.3.4, we have that

\[
\frac{1}{k} \sum_{j=k}^{2k-1} \left\| \nabla g(\tilde{y}_k) \right\|_2^2 \leq \varepsilon f^*.
\]

Therefore, if we stop ACDM at random iteration between \( k \) and \( 2k - 1 \), we have that \( \mathbb{E} \left\| \nabla g(\tilde{y}_k) \right\|_2^2 \leq \varepsilon g^* \) as desired. Therefore, in time \( O\left( m \log 3/2 n \sqrt{\log n \log (\varepsilon^{-1} \log n)} \right) \) we can compute the desired \( \tilde{u} \). \( \square \)
Appendix A

Additional ACDM Proofs

Here we present additional proofs needed to complete the analysis Chapter 3.

A.1 Probabilistic Estimate Sequence Form

Here we prove a stronger variant of Lemma 3.3.2 that is useful for Section 3.4.2. Throughout this section we let $A \in \mathbb{R}^{n \times n}$ denote a symmetric positive semidefinite matrix, we let $\|x\|_A \overset{\text{def}}{=} \sqrt{x^T A x}$ be the induced norm, we let $\langle x, y \rangle_A \overset{\text{def}}{=} x^T A y$ be the inner product which induces this norm, and we let $A^\dagger$ denote the Moore-penrose pseudoinverse of $A$. Taking $A$ to be the diagonal matrix with $A_{ii} = L_1 - \alpha$ and applying the following lemma yields Lemma 3.3.2 as an immediate corollary.

Lemma A.1.1 (General (Probabilistic) Estimate Sequence Construction). Let $A$ be a positive semidefinite matrix such that $f$ is strongly convex with respect to the norm $\|\cdot\|_A$ with convexity parameter $\sigma_A$. Furthermore let $\phi_0(x), \{y_k, \theta_k, i_k\}_{k=0}^\infty$ be such that

- $\phi_0 : \mathbb{R}^n \rightarrow \mathbb{R}$ is an arbitrary function
- Each $y_k \in \mathbb{R}^n$
- Each $\theta_k \in (0, 1)$ and $\sum_{k=0}^\infty \theta_k = \infty$
- Each $i_k \in [n]$ is chosen randomly with probability $p_{ik}$.

Then the pair of sequences $\{\phi_k(x), \eta_k\}_{k=0}^\infty$ defined by

- $\eta_0 = 1$ and $\eta_{k+1} = (1 - \theta_k)\eta_k$,
\[\phi_{k+1}(\bar{x}) := (1 - \theta_k)\phi_k(\bar{x}) + \theta_k \left[f(\bar{y}_k) + \frac{1}{p_{ik}}(\tilde{f}_u(\bar{y}_k), \bar{x} - \bar{y}_k) + \sigma_A ||\bar{x} - \bar{y}_k||^2_A\right]\]

satisfies condition (3.7). Furthermore, if \(\phi_0(\bar{x}) = \phi_0^* + \frac{\zeta_0}{2} ||\bar{x} - \bar{v}_0||^2_A\), then this process produces a sequence of quadratic functions of the form \(\phi_k(\bar{x}) = \phi_k^* + \frac{\zeta_k}{2} ||\bar{x} - \bar{v}_k||^2_A\) where

\[
\begin{align*}
\zeta_{k+1} &= (1 - \theta_k)\zeta_k + \theta_k\sigma_A, \\
\bar{v}_{k+1} &= \frac{1}{\zeta_{k+1}} \left[(1 - \theta_k)\zeta_k \bar{v}_k + \theta_k \sigma_A \bar{y}_k - \frac{\theta_k}{p_{ik}} A^T \tilde{f}_u(\bar{y}_k)\right], \\
\phi_{k+1}^* &= (1 - \theta_k)\phi_k^* + \theta_k f(\bar{y}_k) - \frac{\sigma^2_A}{2\zeta_{k+1}p_{ik}^2} ||\tilde{f}_u(\bar{y}_k)||^2_A, \\
&\quad + \frac{\theta_k(1 - \theta_k)\zeta_k}{\zeta_{k+1}} \left(\frac{\sigma_A}{2} ||\bar{y}_k - \bar{v}_k||^2_A + \frac{1}{p_{ik}} \langle A^T \tilde{f}_u(\bar{y}_k), \bar{v}_k - \bar{y}_k \rangle_A\right).
\end{align*}
\]

Proof. First prove by induction on \(k\) that \(\mathbb{E}[\phi_k(\bar{x})] \leq (1 - \eta_k)f(\bar{x}) + \eta_k\mathbb{E}[\phi_0(\bar{x})]\).

The base case, \(k = 0\), follows trivially from \(\eta_0 = 1\). Assuming it holds for some \(k\), explicitly writing out the expectation over \(i_k\) yields for all \(\bar{x} \in \mathbb{R}^n\), \(\mathbb{E}[\phi_{k+1}(\bar{x})]\) is given by the following

\[
\mathbb{E} \left[ \sum_{i_k} p_{ik} \left( (1 - \theta_k)\phi_k(\bar{x}) + \theta_k \left[f(\bar{y}_k) + p_{ik}^{-1}(\tilde{f}_u(\bar{y}_k), \bar{x} - \bar{y}_k) + \frac{\sigma_A}{2} ||\bar{x} - \bar{y}_k||^2_A\right)\right]\right].
\]

Applying the inductive hypothesis, we get that \(\mathbb{E}[\phi_{k+1}(\bar{x})]\) is equal to the following.

\[
(1 - \theta_k)(((1 - \eta_k)f(\bar{x}) + \eta_k\mathbb{E}[\phi_0(\bar{x})]) + \theta_k \mathbb{E}\left[\left[f(\bar{y}_k) + (\nabla f(\bar{y}_k), \bar{x} - \bar{y}_k) + \frac{\sigma_A}{2} ||\bar{x} - \bar{y}_k||^2_A\right]\right].
\]

By strong convexity and the definition of \(\eta_{k+1}\), we then have that

\[
\mathbb{E}[\phi_{k+1}(\bar{x})] \leq (1 - \theta_k)(1 - \eta_k)f(\bar{x}) + \theta_k f(\bar{x}) + (1 - \theta_k)\eta_k \mathbb{E}[\phi_0(\bar{x})] \\
= (1 - \eta_{k+1})f(\bar{x}) + \eta_{k+1} \mathbb{E}[\phi_0(\bar{x})].
\]

Next, we prove the form of \(\phi_k\) again by induction on \(k\). Suppose that \(\phi_k = \phi_k^* + \frac{\zeta_k}{2} ||\bar{x} - \bar{v}_k||^2_A\). Now, we prove the form of \(\phi_{k+1}\). By the update rule and the inductive

\[\text{Note that our proof was heavily influenced by the proof in [41] for estimate sequences.}\]
hypothesis, we see that

$$\forall \bar{x} \in \mathbb{R}^n : \nabla^2 \phi_{k+1}(\bar{x}) = (1 - \theta_k) \nabla^2 \phi_k(\bar{x}) + \theta_k \sigma_\mathbf{A} \mathbf{A} = [(1 - \theta_k) \zeta_k + \theta_k \sigma_\mathbf{A}] \mathbf{A}.$$

Consequently, $$\phi_{k+1}(\bar{x}) = \phi_k^* + \frac{\zeta_{k+1}}{2} ||\bar{x} - \bar{v}_{k+1}||_\mathbf{A}^2$$ for $$\zeta_{k+1}$$ as desired and $$\phi^*_{k+1}$$ and $$\bar{v}_{k+1}$$ yet to be determined.

To compute $$\bar{v}_{k+1}$$, we note that $$\nabla \phi_k(\bar{x}) = \zeta_k \mathbf{A}(\bar{x} - \bar{v}_k)$$. Therefore by update rule

$$\nabla \phi_{k+1}(\bar{x}) = (1 - \theta_k) \zeta_k \mathbf{A}(\bar{x} - \bar{v}_k) + \frac{\theta_k}{p_{i_k}} \bar{f}_{i_k} + \theta_k \sigma_\mathbf{A} \mathbf{A}(\bar{x} - \bar{y}_k)$$

$$= \zeta_{k+1} \mathbf{A} \bar{x} - (1 - \theta_k) \zeta_k \mathbf{A} \bar{v}_k + \frac{\theta_k}{p_{i_k}} \bar{f}_{i_k} - \theta_k \sigma_\mathbf{A} \mathbf{A} \bar{y}_k$$

Therefore, we have that $$\bar{v}_{k+1}$$ must satisfy

$$\zeta_{k+1} \mathbf{A} \bar{x} - \zeta_{k+1} \mathbf{A} \bar{v}_{k+1} = \zeta_{k+1} \mathbf{A} \bar{x} - (1 - \theta_k) \zeta_k \mathbf{A} \bar{v}_k + \frac{\theta_k}{p_{i_k}} \bar{f}_{i_k} \bar{y}_k - \theta \sigma_\mathbf{A} \mathbf{A} \bar{y}_k.$$

So, applying $$\mathbf{A}^\dagger$$ to each side yields the desired formula for $$\bar{v}_{k+1}$$.

Finally, to compute $$\phi^*_{k+1}$$, we note that, by the form of $$\phi_{k+1}(\bar{x})$$ and the update rule for creating $$\phi_{k+1}(\bar{y}_k)$$, looking at $$\phi_{k+1}(\bar{y}_k)$$ yields

$$\phi^*_{k+1} + \frac{\zeta_{k+1}}{2} ||\bar{y}_k - \bar{v}_{k+1}||_\mathbf{A}^2 = (1 - \theta_k) \phi_k(\bar{y}_k) + \theta_k f(\bar{y}_k)$$

$$= (1 - \theta_k) \left[ \phi^*_k + \frac{\zeta_k}{2} ||\bar{y}_k - \bar{v}_k||_\mathbf{A}^2 \right] + \theta_k f(\bar{y}_k).$$

Now, by value of $$\bar{v}_{k+1}$$, we see that

$$\zeta_{k+1}^2 ||\bar{v}_{k+1} - \bar{y}_k||_\mathbf{A}^2 = \left\| (1 - \theta_k) \zeta_k \bar{v}_k + (\theta_k \sigma_\mathbf{A} - \zeta_{k+1}) \bar{y}_k - \frac{\theta_k}{p_{i_k}} \mathbf{A}^\dagger \bar{f}_{i_k}(\bar{y}_k) \right\|_\mathbf{A}^2$$

$$= \left\| (1 - \theta_k) \zeta_k (\bar{v}_k - \bar{y}_k) - \frac{\theta_k}{p_{i_k}} \mathbf{A}^\dagger \bar{f}_{i_k}(\bar{y}_k) \right\|_\mathbf{A}^2$$

$$= (1 - \theta_k)^2 \zeta_k^2 ||\bar{v}_k - \bar{y}_k||_\mathbf{A}^2 + \frac{2(1 - \theta_k) \theta_k \zeta_k}{p_{i_k}} (\mathbf{A}^\dagger \bar{f}_{i_k}(\bar{y}_k), \bar{v}_k - \bar{y}_k)_\mathbf{A}$$

$$+ \frac{\theta_k^2}{p_{i_k}^2} \left\| \mathbf{A}^\dagger \bar{f}_{i_k}(\bar{y}_k) \right\|_\mathbf{A}^2.$$
Combining these two formulas and noting that

\[(1 - \theta_k) \frac{\zeta_k}{2} - \frac{\zeta_{k+1}}{2} \cdot \frac{1}{\zeta_{k+1}^2} (1 - \theta_k)^2 \zeta_k^2 = \frac{(1 - \theta_k) \zeta_k}{2 \zeta_{k+1}} [\zeta_{k+1} - (1 - \theta_k) \zeta_k] = \frac{\theta_k (1 - \theta_k) \zeta_k}{\zeta_{k+1}} \cdot \frac{\sigma_\Lambda}{2}
\]

yields the desired form of \( \phi_{k+1}^* \) and completes the proof. \( \Box \)

## A.2 Bounding ACDM Coefficients

In the following lemma we prove bounds on \( \alpha_k, \beta_k, \gamma_k, a_k \) and \( b_k \) required for Theorem 3.3.4 and Lemma 3.3.5.

**Lemma A.2.1 (ACDM Coefficients).** In ACDM \( \gamma_k \) is increasing to \( \sqrt{\frac{\delta_\alpha}{2n\sigma_1 - \alpha}} \) and \( \beta_k \) is decreasing to \( 1 - \sqrt{\frac{\sigma_1 - \alpha}{2\delta_\alpha n}} \). Furthermore, for all \( k \geq 1 \) we have \( \alpha_k \leq 32 \max \left( \frac{1}{k^2}, \sqrt{\frac{\sigma_1 - \alpha}{2\delta_\alpha n}} \right) \) and for all \( k \geq 0 \) we have

\[
\begin{align*}
\alpha_k & \geq \frac{\delta_\alpha}{2n\sigma_1 - \alpha} \left( \left( 1 + \frac{1}{2} \sqrt{\frac{\sigma_1 - \alpha}{2\delta_\alpha n}} \right)^{k+1} - \left( 1 - \frac{1}{2} \sqrt{\frac{\sigma_1 - \alpha}{2\delta_\alpha n}} \right)^{k+1} \right), \quad (A.1) \\
b_k & \geq \left( \left( 1 + \frac{1}{2} \sqrt{\frac{\sigma_1 - \alpha}{2\delta_\alpha n}} \right)^{k+1} + \left( 1 - \frac{1}{2} \sqrt{\frac{\sigma_1 - \alpha}{2\delta_\alpha n}} \right)^{k+1} \right). \quad (A.2)
\end{align*}
\]

**Proof.** We begin by estimating \( \gamma_k \). Using that \( \gamma_k = \alpha_{k+1} b_{k+1} \) and defining \( \gamma_{-1} \overset{\text{def}}{=} a_0/b_0 = 1/(4n) \) we know that \( \gamma_{k+1}^2 - \gamma_{k+1} = \left( 1 - \frac{\gamma_k + 1}{2n} \right) \gamma_k^2 \) and therefore \( \frac{\delta_\alpha}{2n\sigma_1 - \alpha} - \frac{\gamma_k^2}{\delta_\alpha} = \left( 1 - \frac{\gamma_k + 1}{2n} \right) \frac{\gamma_k^2}{\delta_\alpha} \) for all \( k \geq 0 \). Consequently, \( \gamma_k \) is increasing to \( \sqrt{\frac{\delta_\alpha}{2n\sigma_1 - \alpha}} \) and since \( \beta_k = 1 - \frac{\gamma_k}{\delta_\alpha} \), we have that \( \beta_k \) is decreasing to \( 1 - \sqrt{\frac{\sigma_1 - \alpha}{2\delta_\alpha n}} \). Furthermore since \( 2\delta_\alpha n \geq \sigma_1 - \alpha \) we have that all \( \beta_k \in (0, 1), b_k \leq b_{k+1}, \) and \( a_k \leq a_{k+1} \).

Using these insights we can now estimate the growth of coefficients \( a_k \) and \( b_k \). As in [43] the growth can be estimated by a recursive relation between \( a_k \) and \( b_k \). For \( b_k \), our definitions imply

\[
b_k^2 = \beta_k b_{k+1}^2 = \left( 1 - \frac{\sigma_1 - \alpha}{\delta_\alpha} \gamma_k \right) b_{k+1}^2 = \left( 1 - \frac{\sigma_1 - \alpha}{\delta_\alpha} \frac{a_{k+1}}{b_{k+1}} \right) b_{k+1}^2.
\]

Therefore since \( b_k \leq b_{k+1} \) we have \( \frac{\sigma_1 - \alpha}{\delta_\alpha} a_{k+1} b_{k+1} \leq b_{k+1}^2 - b_k^2 \leq 2b_{k+1}(b_{k+1} - b_k) \) and
consequently
\[ b_{k+1} \geq b_k + \frac{\sigma_{1-a}}{2S_a} a_{k+1}. \]  

(A.3)

To bound \( a_k \), the definitions imply
\[ \frac{a_{k+1}^2}{b_{k+1}^2} - \frac{a_k}{2nb_{k+1}} = \frac{\gamma_k^2}{2n} = \beta_k \frac{a_k}{b_k} = \frac{a_k^2}{b_{k+1}^2}. \]

Therefore, since \( a_k \leq a_{k+1} \) we have \( \frac{1}{2n} a_{k+1} b_{k+1} = a_k^2 - a_k^2 \leq 2a_{k+1}(a_{k+1} - a_k) \) and consequently
\[ a_{k+1} \geq a_k + \frac{1}{4n} b_{k+1} \geq a_k + \frac{1}{4n} b_k. \]  

(A.4)

Using (A.4) and (A.3) we can prove the growth rates (A.1) and (A.2) by induction.

All that remains is to prove the upper bound on \( a_k \). Note that \( \frac{1-a_k}{\alpha_k} = \frac{2n\gamma_k-1}{\beta_k} \) and hence \( a_k \sim \frac{\beta_k}{2n\gamma_k-1} \sim \frac{1}{2n\gamma_k} \). Therefore, we wish to find a lower bound of \( \gamma_k \), which in turn can be found by a upper bound of \( \beta_k \). Recalling that \( b_{k+1} = \frac{b_k}{\sqrt{\beta_{k-1}}} \) we see that
\[ b_k^2 = \frac{b_k}{\sqrt{\beta_{k-1}}} \leq \frac{b_{k-1}^2}{\beta_{k-1}}. \]

Therefore, by our lower bound on \( b_k \) and the fact that \( b_0 = 2 \), we see that when \( k \leq 2 \sqrt{\frac{2S_a\pi}{\sigma_{1-a}}} \), we have
\[
\beta_k \leq 2^{2/k} \left( \left( 1 + \frac{1}{2} \sqrt{\frac{\sigma_{1-a}}{2S_a n}} \right)^k + \left( 1 - \frac{1}{2} \sqrt{\frac{\sigma_{1-a}}{2S_a n}} \right)^k \right)^{-2/k}
\leq 2^{2/k} \left( 2 + \frac{\sigma_{1-a}}{8S_a n} k(k-1) \right)^{-2/k} \leq \left( 1 + \frac{\sigma_{1-a}}{16S_a n} k(k-1) \right)^{-2/k}
\leq \exp \left( -\frac{\sigma_{1-a}}{8S_a n} (k-1) \right) \leq 1 - \frac{\sigma_{1-a}}{16S_a n} (k-1). 
\]

Using \( \beta_k = 1 - \frac{\gamma_k \sigma_{1-a}}{S_a} \), we get \( \gamma_k \geq \frac{1}{16} \min \left( \frac{k-1}{n}, \sqrt{\frac{2S_a}{\sigma_{1-a} n}} \right) \). Therefore since \( \frac{1}{\alpha_k} - 1 = \frac{1-a_k}{\alpha_k} = \frac{2n\gamma_k-1}{\beta_k} \geq n\gamma_k - 1 \) for all \( k \) we have that \( \alpha_k \leq \frac{1}{n\gamma_k} \leq 16 \max \left\{ \frac{1}{k-1}, \sqrt{\frac{\sigma_{1-a}}{2S_a n}} \right\} \) and the result follows immediately. \( \Box \)
A.3 Numerical Stability of ACDM

Proof of Theorem 3.3.4. Following the structure in [43] we begin by defining $T_i(\bar{x}) \overset{\text{def}}{=} \bar{x} - \frac{1}{L_i} \tilde{f}_i(\bar{x})$ for all $\bar{x} \in \mathbb{R}^n$ and $r_{k}^{2} \overset{\text{def}}{=} \|\tilde{v}_k - \bar{x}^*\|_{1-\alpha}^2$ for all $k \geq 0$. Expanding $\tilde{v}_k$ yields

$$r_{k+1}^2 = \|\tilde{\varepsilon}_{2,k} + \beta_k \tilde{v}_k + (1 - \beta_k)\bar{y}_k - \frac{\gamma_k}{L_{i_k}} \tilde{f}_k(\bar{y}_k) - \bar{x}^*\|_{1-\alpha}^2$$

$$\leq \left(1 + \frac{1}{t}\right) \|\tilde{\varepsilon}_{2,k}\|_{1-\alpha}^2 + (1 + t)\|\beta_k \tilde{v}_k + (1 - \beta_k)\bar{y}_k - \frac{\gamma_k}{L_{i_k}} \tilde{f}_k(\bar{y}_k) - \bar{x}^*\|_{1-\alpha}^2$$

$$\leq \left(1 + \frac{1}{t}\right) \|\tilde{\varepsilon}_{2,k}\|_{1-\alpha}^2 + (1 + t)\|\beta_k \tilde{v}_k + (1 - \beta_k)\bar{y}_k - \bar{x}^*\|_{1-\alpha}^2 + (1 + t)\frac{\gamma_k^2}{L_{i_k}^2} \tilde{f}_k(\bar{y}_k)^2$$

$$+ 2\frac{\gamma_k}{L_{i_k}} (1 + t) \left(\tilde{f}_k(\bar{y}_k), \bar{x}^* - \beta_k \tilde{v}_k - (1 - \beta_k)\bar{y}_k\right)$$

$$\leq \left(1 + \frac{1}{t}\right) \|\tilde{\varepsilon}_{2,k}\|_{1-\alpha}^2 + (1 + t)\|\beta_k \tilde{v}_k + (1 - \beta_k)\bar{y}_k - \bar{x}^*\|_{1-\alpha}^2$$

$$+ 2\frac{\gamma_k^2}{L_{i_k}^2} (1 + t) \left(f(\bar{y}_k) - f(T_i(\bar{y}_k))\right)$$

$$+ 2\frac{\gamma_k}{L_{i_k}} (1 + t) \left(\tilde{f}_k(\bar{y}_k), \bar{x}^* - \bar{y}_k + \frac{\beta_k(1 - \alpha_k)\bar{x}_k - \bar{y}_k}{\alpha_k}\right),$$

(A.5)

where $t$ is yet to be determined.

Using standard properties of $f$, we bound the error induced by $\tilde{\varepsilon}_{1,k}$ as follows:

$$f(T_i(\bar{y}_k)) = f(\bar{x}_{k+1} - \tilde{\varepsilon}_{1,k})$$

(Assumption of Step 3d)

$$\geq f(\bar{x}_{k+1}) - \langle \tilde{\varepsilon}_{1,k}, \nabla f(\bar{x}_{k+1}) \rangle$$

(Convexity of $f$)

$$= f(\bar{x}_{k+1}) - \langle \tilde{\varepsilon}_{1,k}, \nabla f(\bar{x}_{k+1}) - \nabla f(\bar{x}^*) \rangle$$

($\nabla f(\bar{x}^*) = 0$)

$$\geq f(\bar{x}_{k+1}) - \|\tilde{\varepsilon}_{1,k}\|_{1-\alpha} \|\nabla f(\bar{x}_{k+1}) - \nabla f(\bar{x}^*)\|_{1-\alpha}^*$$

(Cauchy Schwarz)

$$\geq f(\bar{x}_{k+1}) - \tilde{\varepsilon}_{S_\alpha} \|\bar{x}_{k+1} - \bar{x}^*\|_{1-\alpha}$$

(Lemma 2 of [43])

$$\geq f(\bar{x}_{k+1}) - \frac{\tilde{\varepsilon}_{S_\alpha}}{\sigma_{1-\alpha}} (f(\bar{x}_{k+1}) - f^*)$$

(A.6)
Taking the expectation of both sides of (A.5) in $i_k$ and using (A.6) and $L_k \geq \frac{\delta}{2\alpha}$ yield

$$
\mathbb{E}_{i_k} \left[ (r_{k+1}^2) \right] \leq \left( 1 + \frac{1}{t} \right) \tau_k^2 + (1 + t) \beta_k r_k^2 + (1 + t) (1 - \beta_k) \| \tilde{y}_k - \tilde{x} \|^2_{1-\alpha} \nabla f(\tilde{y}_k) \cdot (\tilde{x} - \tilde{y}_k) + \frac{\beta_k (1 - \alpha_k)}{\alpha_k} \nabla f(\tilde{y}_k) \cdot (\tilde{x} - \tilde{y}_k)
$$

Now since we defined coefficients so the following hold

$$
1 - \beta_k = \frac{\gamma_k}{\bar{S}_\alpha} \sigma_{1-\alpha} \quad \text{and} \quad \gamma_k^2 = \frac{\gamma_k}{2n} = \frac{\gamma_k \beta_k (1 - \alpha_k)}{2n \alpha_k},
$$

we see that the terms for $\| \tilde{y}_k - \tilde{x} \|^2_{1-\alpha}$ and $f(\tilde{y}_k)$ cancel and we obtain

$$
\mathbb{E}_{i_k} \left[ r_{k+1}^2 \right] \leq \left( 1 + \frac{1}{t} \right) \tau_k^2 + (1 + t) \beta_k r_k^2 - 4(1 + t) \frac{\gamma_k^2}{\bar{S}_\alpha} \left( \mathbb{E}_k \left[ f(\mathbb{E}_k+1) \right] - \frac{\epsilon_\alpha}{\sigma_{1-\alpha}} (\mathbb{E}_k \left[ f(\mathbb{E}_k+1) \right] - f^*) \right) + 2(1 + t) \frac{\gamma_k}{\bar{S}_\alpha} \left( f^* + \frac{\beta_k (1 - \alpha_k)}{\alpha_k} f(\mathbb{E}_k) \right).
$$

Multiplying both sides by $\frac{\delta_k b_{k+1}^2}{n}$ and recalling the following

$$
b_{k+1}^2 = \frac{1}{\beta_k} b_k^2, \quad \alpha_{k+1}^2 = \gamma_k b_{k+1}^2, \quad \beta_k^2 = \frac{\gamma_k \beta_k (1 - \alpha_k)}{2n \alpha_k},
$$

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we get
\[
\frac{\tilde{S}_a b_{k+1}^2}{n} E_{ik} (r_{k+1}^2) \leq \left( 1 + \frac{1}{t} \right) \frac{\tilde{S}_a b_{k+1}^2}{n} e^2 + (1 + t) \frac{\tilde{S}_a b_{k+1}^2}{n} r_k^2 - 4(1 + t) \cdot \\
\left( a_{k+1}^2 \left( 1 - \frac{\varepsilon \tilde{S}_a}{\sigma_{1-\alpha}} \right) \left( E_{ik} f(\bar{x}_{k+1}) - f^* \right) - a_k^2 (f(\bar{x}_k) - f^*) \right).
\]

Dropping the expectation in each variable we have that in expectation to iteration $k$
\[
\frac{\tilde{S}_a b_{k+1}^2 (r_{k+1}^2)}{n} + (1 + t) \left( 1 - \frac{\varepsilon \tilde{S}_a}{\sigma_{1-\alpha}} \right) 4a_{k+1}^2 (f(\bar{x}_{k+1}) - f^*) \\
\leq \left( 1 + \frac{1}{t} \right) \frac{\tilde{S}_a b_{k+1}^2}{n} e^2 + (1 + t) \left( \frac{\tilde{S}_a b_{k+1}^2}{n} r_k^2 + 4a_k^2 (f(\bar{x}_k) - f^*) \right).
\]

Letting $t \overset{\text{def}}{=} \frac{2 \varepsilon \tilde{S}_a}{\sigma_{1-\alpha}}$ and writing $r_k^2 = \frac{\tilde{S}_a}{n} E_{ik} [r_k^2], \phi_k = E_{ik} [f(\bar{x}_{k+1})] - f^*$, we have
\[
b_{k+1}^2 r_{k+1}^2 + 4a_{k+1}^2 \phi_{k+1} \leq \frac{\tilde{S}_a}{n} \left( 1 + \frac{1}{t} \right) b_{k+1}^2 e^2 + (1 + t) \left( b_{k+1}^2 r_k^2 + 4a_k^2 \phi_k \right) \\
\leq \frac{\tilde{S}_a}{n} e^2 \left( 1 + \frac{1}{t} \right) \sum_{j=1}^{k+1} (1 + t)^{k+1-j} b_j^2 \\
+4(1 + t)^{k+1} \left( r_0^2 + \frac{1}{2n \tilde{S}_a} \phi_0 \right).
\] (A.7)

Now, we claim the following inequalities:
\[
\begin{align*}
(1 + t)^{k+1-j} b_j^2 & \text{ is increasing} \quad (1) \\
\sqrt{\frac{\tilde{S}_a}{2n \sigma_{1-\alpha}}} & \geq \gamma_k \geq \frac{1}{2n} \quad (2) \\
\frac{1}{2} \sqrt{\frac{\tilde{S}_a}{2n \sigma_{1-\alpha}}} \left( 1 + \frac{1}{2} \sqrt{\frac{\sigma_{1-\alpha}}{25n}} \right)^{k+1} & \geq a_k \quad (3)
\end{align*}
\]
Using the claims above and (A.7), we get

\[
E_{ik} \left(2\sigma_{1-\alpha}||\vec{e}_{k+1} - \vec{x}^*||^2_{1-\alpha} + 4(f(\vec{x}_{k+1}) - f^*)\right)
= \frac{2n\sigma_{1-\alpha}r^2}{S_k} + 4\phi_{k+1}
\leq \frac{b^2_{k+1}}{a^2_{k+1}}\gamma^2_{k+1} + 4\phi_{k+1}
\leq \frac{\tilde{S}_\alpha e^2}{n} \left(1 + \frac{1}{t}\right)(k + 1)\frac{b^2_{k+1}}{a^2_{k+1}} + \frac{4(1 + t)^{k+1}}{a^2_{k+1}} \left(\tilde{\gamma}_0^2 + \frac{1}{2n}\phi_0\right)
\leq 6\tilde{S}_\alpha e^2(k + 1) + \frac{32n\sigma_{1-\alpha}}{\tilde{S}_\alpha} \left(1 + \frac{2\tilde{S}_\alpha}{\sigma_{1-\alpha}}\right)^{k+1} \left(1 + \frac{1}{2\sqrt{\sigma_{1-\alpha}}}\right)^{-(2(k+1))} \left(\tilde{\gamma}_0^2 + \frac{\phi_0}{2n}\right)
\]

By the assumption that \(\frac{2\tilde{S}_\alpha}{\sigma_{1-\alpha}} < \frac{1}{2\sqrt{2\tilde{S}_\alpha}}\), inequality (3.12) follows from the following

\[
E_{ik} \left(2\sigma_{1-\alpha}||\vec{e}_{k+1} - \vec{x}^*||^2_{1-\alpha} + 4(f(\vec{x}_{k+1}) - f^*)\right)
\leq 12k\tilde{S}_\alpha e^2 + 32\sigma_{1-\alpha} \left(1 + \frac{1}{2\sqrt{\sigma_{1-\alpha}}}\right)^{-k} \left(||\vec{x}_0 - \vec{x}^*||^2_{1-\alpha} + \frac{1}{2\tilde{S}_\alpha^2}\right)\left(f(x_0) - f^*\right)
\leq 24k\tilde{S}_\alpha e^2 + 32\sigma_{1-\alpha} \left(1 - \frac{1}{5}\right) \left(||\vec{x}_0 - \vec{x}^*||^2_{1-\alpha} + \frac{1}{\tilde{S}_\alpha^2}\right)\left(f(x_0) - f^*\right).
\]

Now, we prove the claims.

Claim (1): Since \(a_{k+1} = \gamma_kb_{k+1} \geq \frac{1}{2n}b_{k+1}\) and (A.3), we have \(b_{k+1} \geq (1 + \frac{\sigma_{1-\alpha}}{4\tilde{S}_\alpha n})b_k\).

By the assumption \(t = \frac{2\tilde{S}_\alpha}{\sigma_{1-\alpha}} < \frac{1}{4\tilde{S}_\alpha n}\), we see that \((1 + t)^{k+1}b_{k+1}^2\) is increasing.

Claim (2): Follows from Lemma A.2.1.

Claim (3): Using Lemma A.2.1 and \(k \geq \sqrt{\frac{\sigma_{1-\alpha}}{2\tilde{S}_\alpha n}}\), we have

\[
a_k \geq \sqrt{\frac{\tilde{S}_\alpha}{2n\sigma_{1-\alpha}}} \left(1 + \frac{1}{2\sqrt{\sigma_{1-\alpha}}}\right)^{k+1} = \left(1 + \frac{1}{2\sqrt{\sigma_{1-\alpha}}}\right)\left(1 + \frac{1}{2\sqrt{\sigma_{1-\alpha}}}\right)^{k+1}
\geq \sqrt{\frac{\tilde{S}_\alpha}{2n\sigma_{1-\alpha}}} \left(1 + \frac{1}{2\sqrt{\sigma_{1-\alpha}}}\right)^{k+1} - 1
\geq \frac{1}{2\sqrt{2n\sigma_{1-\alpha}}} \left(1 + \frac{1}{2\sqrt{\sigma_{1-\alpha}}}\right)^{k+1}.
\]

To bound the norm of the gradient, we show that if \(||\nabla f(\vec{y}_k)||^2_{1-\alpha}\) is large for
many steps then \( f(\bar{x}_k) \) decreases substantially. For notational simplicity we omit the expectation symbol and using (A.6), we have

\[
f(\bar{x}_{k+1}) \leq f(T_k(\bar{y}_k)) + \frac{\varepsilon \tilde{S}_\alpha}{\sigma_{1-\alpha}} (f(\bar{x}_{k+1}) - f^*) \]

\[
\leq f(\bar{y}_k) - \frac{1}{\tilde{S}_\alpha} (\|\nabla f(\bar{y}_k)\|_{1-\alpha}^*)^2 + \frac{\varepsilon \tilde{S}_\alpha}{\sigma_{1-\alpha}} (f(\bar{x}_{k+1}) - f^*).
\]

To bound \( f(\bar{y}_k) \), we consider the lower envelopes at \( \bar{y}_k \) and obtain

\[
f(\bar{y}_k) \leq f(\bar{x}_k) - \langle \nabla f(\bar{y}_k), \alpha_k(\bar{x}_k - \bar{y}_k) \rangle\]

\[
\leq f(\bar{x}_k) + \alpha_k \|\nabla f(\bar{y}_k)\|_{1-\alpha}^* \|\bar{x}_k - \bar{y}_k\|_{1-\alpha}^*
\]

\[
\leq f(\bar{x}_k) + \alpha_k \|\nabla f(\bar{y}_k)\|_{1-\alpha}^* \left( \frac{1}{\sqrt{\sigma_{1-\alpha}}} \sqrt{f(\bar{x}_k) - f^*} + \|\bar{y}_k - \bar{x}^*\|_{1-\alpha} \right).
\]

Combining with Lemma A.2.1, we have

\[
f(\bar{x}_{k+1}) \leq f(\bar{x}_k) - \frac{1}{\tilde{S}_\alpha} (\|\nabla f(\bar{y}_k)\|_{1-\alpha}^*)^2 + \frac{\varepsilon \tilde{S}_\alpha}{\sigma_{1-\alpha}} \delta_{k+1} + 3 \sqrt{\frac{\sigma_{1-\alpha}}{n \tilde{S}_\alpha}} \|\nabla f(\bar{y}_k)\|_{1-\alpha}^* \sqrt{\delta_k}.
\]

Summing up from \( k \) to \( 2k \) then yields

\[
f(\bar{x}_{2k}) \leq f(\bar{x}_k) - \sum_{j=k}^{2k-1} \left( \frac{\|\nabla f(\bar{y}_j)\|_{1-\alpha}^*}{\tilde{S}_\alpha} \right)^2 + \frac{\varepsilon \tilde{S}_\alpha}{\sigma_{1-\alpha}} \sum_{j=k}^{2k} \delta_j + 32 \sqrt{\frac{1}{n \tilde{S}_\alpha}} \sum_{j=k}^{2k-1} \sqrt{\delta_j} \|\nabla f(\bar{y}_j)\|_{1-\alpha}^*
\]

Since \( f(\bar{x}_k) - f(\bar{x}_{2k}) \leq f(\bar{x}_k) - f^* \leq \delta_k \), we have

\[
\frac{1}{\tilde{S}_\alpha} \sum_{j=k}^{2k-1} \left( \|\nabla f(\bar{y}_j)\|_{1-\alpha}^* \right)^2 \leq \delta_k + \frac{\varepsilon k \tilde{S}_\alpha}{\sigma_{1-\alpha}} \sum_{j=k+1}^{2k} \delta_j + 32 \sqrt{\frac{1}{n \tilde{S}_\alpha}} \sum_{j=k}^{2k-1} \|\nabla f(\bar{y}_j)\|_{1-\alpha}^* \sqrt{\delta_j}
\]

\[
\leq \delta_k \left( 1 + \frac{\varepsilon k \tilde{S}_\alpha}{\sigma_{1-\alpha}} \right) + 32 \sqrt{\frac{1}{n \tilde{S}_\alpha}} \sum_{j=k}^{2k-1} \|\nabla f(\bar{y}_j)\|_{1-\alpha}^* \sqrt{\delta_j}
\]

\[
\leq \delta_k \left( 1 + \frac{\varepsilon k \tilde{S}_\alpha}{\sigma_{1-\alpha}} \right) + 32 \sqrt{\frac{k \delta_k}{n}} \sqrt{\frac{1}{\tilde{S}_\alpha}} \sum_{j=k}^{2k-1} \left( \|\nabla f(\bar{y}_j)\|_{1-\alpha}^* \right)^2.
\]
Solving this quadratic equation about \( \sqrt{\frac{1}{\tilde{S}_\alpha} \sum_{j=k}^{2k-1} (\| \nabla f(\tilde{y}_k)\|_{1-\alpha})^2} \), we get
\[
\sqrt{\frac{1}{\tilde{S}_\alpha} \sum_{j=k}^{2k-1} (\| \nabla f(\tilde{y}_k)\|_{1-\alpha})^2} \leq 16\sqrt{\frac{k\delta_k}{n}} + \frac{1}{2} \sqrt{\left(32\sqrt{\frac{k\delta_k}{n}}\right)^2 + 4\delta_k \left(1 + \frac{\varepsilon k\tilde{S}_\alpha}{\sigma_{1-\alpha}}\right)}
\]

Since \( 1 + \frac{\varepsilon k\tilde{S}_\alpha}{\sigma_{1-\alpha}} \leq 2 \) and \( k \geq n \) by assumptions we have
\[
\sqrt{\frac{1}{\tilde{S}_\alpha} \sum_{j=k}^{2k-1} (\| \nabla f(\tilde{y}_k)\|_{1-\alpha})^2} \leq 16\sqrt{\frac{k\delta_k}{n}} + \frac{1}{2} \sqrt{\left(32^2 + 8\right)\frac{k\delta_k}{n}} \leq 33\sqrt{\frac{k\delta_k}{n}}.
\]

Therefore \( \frac{1}{\tilde{S}_\alpha} \sum_{j=k}^{2k-1} (\| \nabla f(\tilde{y}_k)\|_{1-\alpha})^2 \leq 2000\frac{k\delta_k}{n} \), as desired.

Lemma A.3.1 (Coefficient Stability in ACDM). Step 3a of ACDM can be computed by the following formulas
\[
\gamma_{k+1} = \min\left\{ \frac{1}{2} \left[ \left(1 - \frac{1}{2n} - \frac{\gamma_k^2 \sigma_{1-\alpha}}{\tilde{S}_\alpha}\right) + \sqrt{\left(\frac{1}{2n} - \frac{\gamma_k^2 \sigma_{1-\alpha}}{\tilde{S}_\alpha}\right)^2 + 4\gamma_k^2} \right], \sqrt{\frac{\tilde{S}_\alpha}{2n\sigma_{1-\alpha}}} \right\},
\]
\[
\beta_{k+1} = 1 - \frac{\gamma_{k+1} \sigma_{1-\alpha}}{\tilde{S}_\alpha},
\]
\[
\alpha_{k+1} = \frac{\beta_{k+1}}{\beta_{k+1} + 2n\gamma_{k+1} - 1} = \frac{\frac{2n}{\gamma_k} - \frac{\sigma_{1-\alpha}}{2n\tilde{S}_\alpha}}{1 - \frac{\sigma_{1-\alpha}}{2n\tilde{S}_\alpha}}.
\]

and can be implemented using \( O(\log n) \) bits precision to obtain any \( \text{poly}(\frac{1}{n}) \) additive accuracy.

Proof. Since \( \gamma_k = \frac{a_{k+1}}{b_{k+1}} \) we know that \( \gamma_k^2 = \frac{1}{2n} - \frac{\gamma_{k+1}^2 \sigma_{1-\alpha}}{\tilde{S}_\alpha} = \left(1 - \frac{\gamma_{k+1} \sigma_{1-\alpha}}{\tilde{S}_\alpha}\right) \gamma_k^2 \), and by re-arranging terms we get that \( \gamma_{k+1}^2 - \left(\frac{1}{2n} - \frac{\gamma_k^2 \sigma_{1-\alpha}}{\tilde{S}_\alpha}\right) \gamma_{k+1} - \gamma_k^2 = 0 \). Applying the quadratic formula and Lemma A.2.1 then yields the formula for \( \gamma_{k+1} \) and the other formulas follow by direct computation.

Now, expanding the formula for \( \gamma_{k+1} \) and using \( 2\tilde{S}_\alpha n \geq \sigma_{1-\alpha} \) implies that \( \gamma_k \geq \frac{1}{2n} \) and therefore the denominator of \( \alpha_{k+1} \) is larger than \( \beta_{k+1} \). Consequently the equations of \( \beta \) and \( \alpha \) depends continuously on \( \gamma \). Thus, it is suffices to show that \( O(\log n) \) bits precision suffice to compute \( \gamma_k \). To prove this, we define \( f : \mathbb{R}^n \rightarrow \mathbb{R} \) denote the
quadratic formula so that with full precision \( \gamma_{k+1} = f(\gamma_k) \). Direct calculation yields

\[
f'(\gamma) = \gamma \sigma_{1-\alpha} S_\alpha + \frac{\left( \gamma^2 \sigma_{1-\alpha} S_\alpha - \frac{1}{2n} \right) \gamma \sigma_{1-\alpha} S_\alpha + 2\gamma \sqrt{\left( \frac{1}{2n} - \gamma^2 \sigma_{1-\alpha} S_\alpha \right)^2 + 4\gamma^2}}{\sqrt{\left( \frac{1}{2n} - \gamma^2 \sigma_{1-\alpha} S_\alpha \right)^2 + 4\gamma^2}}
\]

Since \( 0 \leq \gamma \leq \sqrt{\frac{S_\alpha}{2n\sigma_{1-\alpha}}} \) we have that \( 0 \leq f'(\gamma) \leq 1 - \frac{\gamma \sigma_{1-\alpha} S_\alpha}{1 - \gamma^2 \sigma_{1-\alpha} S_\alpha} < 1 \) and therefore, \( f \) is a contraction mapping. Consequently, if the calculation of \( \gamma \) has up to \( \varepsilon \) additive error, \( k \) steps of calculation can accumulate up to \( k\varepsilon/(1 - \max f'(\gamma)) \) additive error. Hence, \( O(\log n) \) bits of precision suffice. \( \square \)
Bibliography


