MINIMUM-VOLUME COVERING ELLIPSOIDS: IMPROVING THE EFFICIENCY OF THE WOLFE-ATWOOD ALGORITHM FOR LARGE-SCALE INSTANCES BY POOLING AND BATCHING

Jakub Kůdela
Institute of Automation and Computer Science, Brno University of Technology, Czech Republic
Jakub.Kudela@vutbr.cz

Abstract
The Minimum-Volume Covering Ellipsoid (MVCE) problem is an important optimization problem that comes up in various areas of engineering and statistics. In this paper, we improve the state-of-the-art Wolfe-Atwood algorithm for solving the MVCE problem with pooling and batching procedures. This implementation yields significant improvements on the runtime of the algorithm for large-scale instances of the MVCE problem, which is demonstrated on quite extensive computational experiments.

Keywords: minimum-volume covering ellipsoid, Löwner-John ellipsoid, large-scale optimization, Wolfe-Atwood algorithm, pooling, batching.

1 Introduction
Let \( X = \{x_1, \ldots, x_m\} \subset \mathbb{R}^n \) be a finite set of vectors whose affine hull is \( \mathbb{R}^n \). In this paper, the problem in question will be finding the Minimum-Volume Covering Ellipsoid (MVCE) of \( X \), also known as the Löwner-John ellipsoid. The recently published book [1] gives a wonderful overview of the MVCE problem, and will serve as our primary source for the introduction and the apparatus for handling the MVCE problem. The idea of MVCE and the proofs of its existence and uniqueness come from Löwner (unfortunately in an unpublished work). The first published results about MVCE appear in the 1950’s in [2] and [3]. An earlier result by John [4] about the suitability of ellipsoids for fitting convex compact sets was also one of the pivotal starting points for the study of the MVCE problem (hence, the alternative name for the problem is deciphered!). The MVCE problem comes up in several applied and theoretical areas. It has a strong connection to the optimal design in statistics [5]. Other application in statistics include outlier identification (see [6], [7], and [8]), and clustering (see [9], [10], and [11]). Containing ellipsoids play an important role in robust optimization [12] and control theory [13]. MVCEs are also used in computational geometry and computer graphics [14], e.g. for obstacle avoidance in robotics. One of the standard methods for solving the MVCE problem, the Frank-Wolfe method, is being investigated for its suitability for solving statistical learning problems [15, 16].

The main contribution of this paper is in the efficient computation of large scale instances of the MVCE problem (in both the dimension \( n \) and the number of vectors \( m \)). Our starting point is the Wolfe-Atwood (WA) method [17, 18] with Kumar-Yıldırım initialization [19]. We describe a pooling scheme, in which we disregard a large portion of the vectors, solving the MCVE problem for only a small subset of \( X \). Once a solution of this surrogate problem is found, we look at the whole \( X \) and find the points that were the furthest away from the ellipsoid. These points are then added to the surrogate problem. This procedure is repeated until a “good enough” solution is found (the precise conditions will be defined in the following sections). A similar idea in the context of chance constrained optimization problems was investigated in [20] and [21].

2 The MVCE Problem
There are several equivalent mathematical definitions of ellipsoids. The one adopted in this paper is the following: An ellipsoid \( \mathcal{E}(H, \bar{x}) \) is a set of the form

\[
\mathcal{E}(H, \bar{x}) = \{x \in \mathbb{R}^n : (x - \bar{x})^TH(x - \bar{x}) \leq n\},
\]

where \( \bar{x} \in \mathbb{R}^n \) is the center of the ellipsoid and \( H \) is a symmetric positive definite matrix of order \( n \). Another way of thinking about ellipsoids is through affine transformations of balls in \( \mathbb{R}^n \). In particular, let \( L \) be the Cholesky factor of \( H \), i.e. \( H = LL^T \), with \( L \) being a lower triangular matrix with positive diagonal entries. Then \( x \) lies in \( \mathcal{E}(H, \bar{x}) \) iff \( \|L^T(x - \bar{x})\| \leq \sqrt{n} \), so that

\[
\mathcal{E}(H, \bar{x}) = \{x = \bar{x} + (\sqrt{n}L^{-T})z : z \in \mathbb{R}^n, \|z\| \leq 1\}.
\]
The volume of the ellipsoid can therefore be computed as the volume of the unit ball times $|\det(\sqrt{n}L^{-T})|$. Using the fact that $H = LL^T$, and $\det H = (\det L)^2$, we get

$$\text{vol}(E(H, \bar{x})) = \frac{n^{n/2} \Omega_n}{\sqrt{\det H}},$$

where $\Omega_n$ is the volume of a unit ball in $\mathbb{R}^n$. This means that if we seek to minimize the volume of the ellipsoid, we can frame it as a minimization of the negative of the determinant of its shape matrix $H$. The problem of finding the MVCE of $\mathcal{X}$ then attains the following form:

$$\begin{aligned}
\minimize_{H \in S_n^+, \bar{x} \in \mathbb{R}^n} & - \ln \det(H) \\
\text{subject to} & (x_i - \bar{x})^T H (x_i - \bar{x}) \leq n, \quad i = 1, \ldots, m,
\end{aligned} \tag{1}$$

where $S_n^+$ is the space of symmetric positive definite matrices of order $n$. In the problem (1) the “centering” variable $\bar{x}$ causes computational difficulties [1]. These difficulties can be circumvented by “lifting” the MVCE problem to a higher dimension and solving an equivalent problem, that has $\bar{x} = 0$, see [22]. This equivalent problem is then convex [23] and has the following form:

$$\begin{aligned}
\minimize_{H \in S_n^+, u \in \mathbb{R}^m} & - \ln \det(H) \\
\text{subject to} & x_i^T H x_i \leq n, \quad i = 1, \ldots, m.
\end{aligned} \tag{2}$$

We say that $H$ is feasible for (2) if it satisfies all $m$ constraints and is positive definite. Denoting $u_i \geq 0$ the Lagrange multiplier for the $i$th constraint, the Lagrangian $L(H, u)$ for (2) is

$$L(H, u) = - \ln \det(H) + \sum_{i=1}^{m} u_i (x_i^T H x_i).$$

If we denote by $e$ the vector of ones in $\mathbb{R}^m$, by $U$ the diagonal matrix from the values of $u$:

$$U = \text{Diag}(u) \in S_m^+,$$

and by $X$ the matrix $X = [x_1, \ldots, x_m] \in \mathbb{R}^{n \times m}$, we can rewrite the Lagrangian as

$$L(H, u) = - \ln \det(H) + H X U X^T - u e^T u.$$ 

After a series of manipulations (see [1]) we obtain a dual of (2):

$$\begin{aligned}
\maximize_{u \in \mathbb{R}^m} & \ln \det(X U X^T) \\
\text{subject to} & e^T u = 1, \\
& u \geq 0.
\end{aligned} \tag{3}$$

If the affine hull of $\mathcal{X}$ is $\mathbb{R}^n$ (or, equivalently, if $X$ has full row rank), the optimal solutions to (2) and (3) are unique and attain the same optimal objective value [1] (i.e., strong duality holds).

Another important result that concerns our investigation is the following: there is a finite subset of $\mathcal{X}$ of cardinality at most $n(n + 1)/2$, such that the MVCE containing this subset is also the MVCE for the whole set $\mathcal{X}$ [1]. This small subset is called a core set. The implication being that the optimal solution to (3) has only $n(n + 1)/2$ positive components. Notice that this number does not depend on the number of points $m$.

Since these problems cannot usually be solved exactly, we will be interested in finding $\epsilon$-primal feasible or $\epsilon$-approximately optimal solutions: A feasible $u$ for the problem (3) is said to be $\epsilon$-primal feasible if $H(u) = (X U X^T)^{-1}$ satisfies

$$x_i^T H(u) x_i \leq (1 + \epsilon)n, \quad i = 1, \ldots, m.$$ 

Additionally, if it satisfies

$$x_i^T H(u) x_i \geq (1 - \epsilon)n \text{ if } u_i > 0, \quad i = 1, \ldots, m,$$

we say that $u$ is $\epsilon$-approximately optimal.
Algorithm 1 WA Algorithm

1: procedure Step 0:
2:   Choose $u$ feasible for (3) and $\epsilon > 0$.
3:   Compute $\omega = \omega(u)$ and a (scaled) Cholesky factorization of $XUX^T$.
4:   Go to Step 1.
5: procedure Step 1:
6:   Given the current iterate $u$ and its associated $\omega = \omega(u)$:
7:   Compute $\epsilon_+ = \max_{k} \{\omega_k - n\}/n$, with $h = i$ attaining the maximum.
8:   Compute $\epsilon_- = \max_{k} \{(n - \omega_h)/n : u_h > 0\}$, with $h = j$ attaining the maximum.
9:   if $\max \epsilon_+, \epsilon_- \leq \epsilon$ then STOP: $u$ is $\epsilon$-approximately optimal.
10:  else
11:      if $\epsilon_+ > \epsilon_-$ then go to Step 2;
12:      else go to Step 3.
13: procedure Step 2:
14:   Compute $\lambda^* = \frac{\omega_i - n}{(n+1)\omega_i}$ and update $u \leftarrow (1 + \lambda^*)^{-1}(u + \lambda^* e_i)$.
15:   Go to Step 4.
16: procedure Step 3:
17:   Compute $\lambda^* = \frac{\omega_i - n}{(n+1)\omega_i}$ and set $\lambda = \max\{-u_j, \lambda^*\}$.
18:   Update $u \leftarrow (1 + \lambda)^{-1}(u + \lambda e_j)$.
19:   Go to Step 4.
20: procedure Step 4:
21:   Update $\omega$ and a scaled Cholesky factorization of $XUX^T$.
22:   Go to Step 1.

3 Solving the MVCE Problem

The primary approaches for solving the MVCE problem have been first-order methods (a modified Frank-Wolfe method [24]) applied to the dual problem (3), denoted here as the WA method [17, 18]. There exists a similar method for solving the MVCE problem attributed to Fedorov [25] and Wynn [26] that was proposed at around the same time as the WA method. However, this method was found computationally inferior to the WA method in several numerical studies [27]. A parallel line of research was conducted in second-order methods for solving the primal problem (2). The resulting dual reduced Newton method with an active-set strategy [28] was successful in being faster than the WA method [27], but could not be used for truly large-scale problems because of its memory requirements (for problems in higher dimensions $n$).

In our baseline implementation, we will use the WA method, with its form is taken from [1]. The individual steps are described in Algorithm 1. An important factor in the efficiency of the WA algorithm is the starting value of $u$. Notable initialization schemes are the due to Khachiyan [29], and Kumar and Yıldırım [19]. The Kachiyan initialization is quite straightforward, choosing $u = e/m$. The Kumar and Yıldırım (KY) strategy is a little bit more involved (and also work a bit better), and is described in Algorithm 2. The Steps 0 through 2 of the algorithm are due to [30]. Another increase in the effectiveness of the implementation of the WA method can be made by eliminating points that do not belong to the core set. The method for carrying out the elimination was developed in [31] and the positive impact on the runtime of the algorithm was investigated in [32].

3.1 WA with Pooling and Batching

In a similar fashion to the point elimination [31], our idea of improving the efficiency of WA centers around the core set. However, instead of reducing the size of the solved problem by eliminating points that do not belong to the core set, we take what might be conceived as the direct opposite approach. We devise a surrogate problem that is considerably smaller – a subset $\hat{X}$ of $X$ – find the MVCE of $\hat{X}$ by Algorithm 1, and check, whether the MVCE of $\hat{X}$ contains all the points of $X$. If it does, we are done – $\hat{X}$ had to contain the core set of $X$ and the MVCE of the two sets is identical. If, however, some points of $X$ lie outside the MCVE of $\hat{X}$, the core sets must differ and we need to update $\hat{X}$. In the update, we find the points in $\hat{X}$ that are the furthers away from the MVCE of $\hat{X}$. The maximum number of points that are added during the update is called a batch size and is denoted by $k$. Effectiveness of batching was previously investigated, in the context of stochastic programming, in [33]. Moreover, we found it computationally advantageous to use the resulting $u$‘s from the computations of MVCE of $\hat{X}$ as starting points for the updated iterations. The individual steps are summarized in Algorithm 3.

The selection of points that will be added to the surrogate problem $\hat{X}$ requires the computation of the
Selection strategy of experiments, we will vary only the batch size parameter $k$.

Compared to the WA (Algorithm 1), our Algorithm 3 uses two additional “hyperparameters” – the size of the batch $k$, and the size (an selection strategy) of the starting subset $\tilde{X}$. In the following computational experiments, we will vary only the batch size parameter $k$, leaving the investigation into the impact of the selection strategy of $\tilde{X}$ as our future work. We will use two different setting for the computational experiments, both involving random generation of the set $X$ of different places (not simply appended).

Following quantity

$$||L^{-T}x_i||^2 - n$$

for each $x_i \in X$. Since $L$ comes from a Cholesky factorization it is an upper-triangular matrix, and the computation of the inverse does not pose significant computational burden. When implementing Step 2 of Algorithm 3, one should be careful about properly handling the optimal multipliers $\tilde{u}$. We suppose that the newly added points to $\tilde{X}$ will be appended to the corresponding matrix $\tilde{X}$, hence the instruction to append the corresponding number of zeros to $\tilde{u}_S$. If, for whatever reason, one decided to first somehow sort the individual vectors $x_i \in \tilde{X}$ before constructing the matrix $\tilde{X}$, the added zeros to the starting values $\tilde{u}_S$ would appear on different places (not simply appended).

## 4 Computational Experiments

Compared to the WA (Algorithm 1), our Algorithm 3 uses two additional “hyperparameters” – the size of the batch $k$, and the size (an selection strategy) of the starting subset $\tilde{X}$. In the following computational experiments, we will vary only the batch size parameter $k$, leaving the investigation into the impact of the selection strategy of $\tilde{X}$ as our future work. We will use two different setting for the computational experiments, both involving random generation of the set $X$ and centered around the origin.

In the first setting each of the $m$ vectors $x \in X \subset \mathbb{R}^n$ follows a multivariate normal distribution $x \sim N(0, \Sigma)$, where the individual values $\sigma_{i,j}$ of the covariance matrix $\Sigma$ follow a standard normal distribution $\sigma_{i,j} \sim N(0, 1)$, $i, j = 1, \ldots, n$. This configuration, however, results in the majority of points lying close to the surface of an ellipsoid (whose shape is guided by $\Sigma$), which is quite a special situation [1].
In the second setting, we will generate instances that should not have the majority of points close to an ellipsoidal boundary (this procedure is, yet again, taken from [1]). We generate seconds over 10 independent runs. The experiments with "-" were deemed not computationally interesting.

Table 1: The results from the computational experiments, the first setting. Geometric average of runtimes in seconds over 10 independent runs. The experiments with "-" were deemed not computationally interesting.

| Batch size | m | 5  | 10  | 20  | 50  | 100  | 200  | 500  | 1000  | 5000  | 10000  | speedup |
|------------|---|----|-----|-----|-----|------|------|------|-------|-------|--------|---------|
| 1000       | 0.005 | 0.015 | 0.017 | 0.014 | 0.016 | 0.018 | 0.014 | -    | -     | -     | -      | 0.382   |
| 5000       | 0.012 | 0.025 | 0.032 | 0.024 | 0.021 | 0.026 | 0.028 | -    | -     | -     | -      | 0.594   |
| 10000      | 0.017 | 0.034 | 0.043 | 0.032 | 0.025 | 0.028 | 0.030 | -    | -     | -     | -      | 0.681   |
| 50000      | 0.067 | 0.058 | 0.115 | 0.063 | 0.044 | 0.037 | 0.043 | -    | -     | -     | -      | 2.134   |
| 100000     | 0.189 | 0.179 | 0.144 | 0.147 | 0.134 | 0.119 | 0.105 | 0.085 | 0.096 | 0.131 | 0.145  | 2.242   |
| 500000     | 1.360 | 0.260 | 0.255 | 0.235 | 0.203 | 0.198 | 0.199 | 0.185 | 0.150 | 0.168 | 0.191  | 9.145   |
| 1000000    | 1.585 | 1.194 | 1.025 | 0.851 | 0.806 | 0.749 | 0.658 | 0.622 | 0.549 | 0.537 | 0.578  | 2.965   |

In the second setting, we will generate instances that should not have the majority of points close to an ellipsoidal boundary (this procedure is, yet again, taken from [1]). We generate $m$ independent standard Cauchy random variable and an $n \times m$ matrix $A$ of independent standard normal variables. Then we set each point $x_i$ to be the normalized ith column of $A$ times the ith Cauchy sample. The points generated by this procedure satisfy rotational symmetry, and their distances from the origin are Cauchy.

In the computational experiments, we compare the WA method with the KY initialization and point elimination (the implementation is taken directly from [1]) to our method described in Algorithm 3. We set the optimality parameter $\varepsilon = 10^{-7}$ for both methods. The selection of $\tilde{X}$ is quite straightforward — we simply pick the first $n$ vectors of $\tilde{X}$. We implemented the algorithms in MATLAB 2019a and used a machine with a six-core AMD Ryzen 5 2600X 3.6 GHz processor and 32 GB RAM.

The magnitude of computational experiments was quite extensive. We varied the number of vectors $m$ between 1,000 and 5,000,000, the number of dimension $n$ between 5 and 200, and the size of the batch $k$ between 10 and 10,000. Each of the experiments (for different values of $m, n$, and $k$) was carried out 10 times (in both settings). The results are summarized in Table 1 (for the first setting) and Table 2 (for the second one).

It is apparent from the results, that the pooling and batching procedures described in Algorithm 3 can substantially decrease the computational efforts for solving the MVCE problem for the truly large-scale instances.
Table 2: The results from the computational experiments, the second setting. Geometric average of runtimes "optimal" batch size (the  \( k \)) of the Algorithm 3 with "optimally" chosen batch size). It also showed, that at least for the first setting, the speedup, although still significant, was much smaller when compared to the first setting. Moreover, the "optimal" batch size \( k \) quite clearly depends on the dimension of the MVCE problem. An important observation is that there is no clear similarity to the second setting seems to be between \( k \) and \( n \) depend on the dimension of the MVCE problem. A reasonable rule of thumb for the choice of \( k \) is \( k \approx \frac{n}{5} \).
5 Conclusion

The MVCE problem comes up in various real-world problems in engineering and statistics. In this paper, we described an improving pooling and batching scheme for the Wolfe-Atwood method for large-scale MVCE instances. We conducted quite extensive computational experiments and showed that the “optimal” batch size $k$ depends on the structure of the problem – there is a qualitative difference in its dependence on the dimension $n$ of the problem between first and second experimental setting. The computational experiments showed great promise, regarding the implementability and the usefulness of the proposed algorithm, but there is still much work to be done. The proposed method should be implemented on real-life problems, to get a clearer picture of the improvements it provides. The importance of the selection of the starting set $\tilde{X}$ and different selection criteria for updating said set should be further looked into. Another possible line of research constitutes the inclusion of the discarding methodology developed in [21] for “Ellipsoidal peeling” in experiment design [34].

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