Streaming Algorithms for High-Dimensional Robust Statistics

Ilias Diakonikolas ¹ Daniel M. Kane ² Ankit Pensia ¹ Thanasis Pittas ¹

Abstract

We study high-dimensional robust statistics tasks in the streaming model. A recent line of work obtained computationally efficient algorithms for a range of high-dimensional robust statistics tasks. Unfortunately, all previous algorithms require storing the entire dataset, incurring memory at least quadratic in the dimension. In this work, we develop the first efficient streaming algorithms for high-dimensional robust statistics with nearoptimal memory requirements (up to logarithmic factors). Our main result is for the task of highdimensional robust mean estimation in (a strengthening of) Huber's contamination model. We give an efficient single-pass streaming algorithm for this task with near-optimal error guarantees and space complexity nearly-linear in the dimension. As a corollary, we obtain streaming algorithms with near-optimal space complexity for several more complex tasks, including robust covariance estimation, robust regression, and more generally robust stochastic optimization.

1. Introduction

This work studies high-dimensional learning in the presence of a constant fraction of arbitrary outliers. Outlier-robust learning in high dimensions is motivated by pressing machine learning (ML) applications, including ML security (Barreno et al., 2010; Biggio et al., 2012; Steinhardt et al., 2017; Tran et al., 2018; Diakonikolas et al., 2019a) and exploratory analysis of datasets with natural outliers (Rosenberg et al., 2002; Paschou et al., 2010; Li et al., 2008). This field has its roots in robust statistics, a branch of statistics initiated in the 60s with the pioneering works of Tukey and Huber (Tukey, 1960; Huber, 1964). Early work developed minimax optimal estimators for various robust estimation tasks,

Authors are listed in alphabetical order. ¹University of Wisconsin-Madison ²University of California, San Diego. Correspondence to: Ankit Pensia <ankitp@cs.wisc.edu>, Thanasis Pittas pittas@wisc.edu>.

Proceedings of the 39th International Conference on Machine Learning, Baltimore, Maryland, USA, PMLR 162, 2022. Copyright 2022 by the author(s).

albeit with runtimes exponential in the dimension. A recent line of work in computer science, starting with (Diakonikolas et al., 2016; Lai et al., 2016), developed polynomial time robust estimators for a range of high-dimensional statistical tasks. Algorithmic high-dimensional robust statistics is by now a relatively mature field, see, e.g., (Diakonikolas & Kane, 2019; Diakonikolas et al., 2021a) for surveys.

This recent progress notwithstanding, even for the basic task of mean estimation, previous robust estimators require the entire dataset in main memory. This space requirement can be a major bottleneck in large-scale applications, where an algorithm has access to a very large stream of data. Indeed, practical machine learning methods are typically simple iterative algorithms that make a single pass over the data and require a small amount of storage — with stochastic gradient descent being the prototypical example (Bottou, 2010; Bottou et al., 2018). Concretely, in prior applications of robust statistics in data analysis (Diakonikolas et al., 2017a) and data poisoning defenses (Diakonikolas et al., 2019a), the storage requirements of the underlying algorithms were observed to significantly hinder scalability¹. This discussion motivates the following natural question:

Can we develop efficient robust estimators in the streaming model with (near-) optimal space complexity?

We emphasize that this broad question is meaningful and interesting even ignoring computational considerations. While any method requires space complexity $\Omega(d)$, where d is the dimension of the problem (to store a single sample), it is not obvious that a matching upper bound exists. We note that it is relatively simple to design O(d)-memory streaming algorithms with sample complexity exponential in d. But it is by no means clear whether there exists an estimator with near-linear space requirements and $\operatorname{poly}(d)$ sample complexity (independent of its runtime).

1.1. Our Results

In this work, we initiate a systematic investigation of highdimensional robust statistics in the streaming model. We start by focusing on the most basic task — that of robust mean estimation. Our main result is the first space-efficient streaming algorithm for robust mean estimation under nat-

¹See, for example, Section 5.1 of (Diakonikolas et al., 2017b).

ural distributional assumptions. Our computationally efficient algorithm makes a single pass over the data, uses near-optimal space, and matches the error guarantees of previous polynomial-time algorithms for the problem.

Given this result, we leverage the fact that several robust statistics tasks can be reduced to robust mean estimation to obtain near-optimal space, single-pass streaming algorithms for more complex statistical tasks.

To formally state our contributions, we require some basic definitions. We start with the standard streaming model.

Definition 1.1 (Single-Pass Streaming Model). Let S be a fixed set. In the *one-pass streaming model*, the elements of S are revealed one at a time to the algorithm, and the algorithm is allowed a single pass over these points.

Our robust estimators work in the following contamination model, where the adversary can corrupt the true distribution in total variation distance (for distributions P and Q, we use $d_{\text{TV}}(P,Q)$ to denote their total variation distance).

Definition 1.2 (TV-contamination). Given a parameter $\epsilon < 1/2$ and a distribution class \mathcal{D} , the adversary specifies a distribution D' such that there exists $D \in \mathcal{D}$ with $d_{\mathrm{TV}}(D, D') \leq \epsilon$. Then the algorithm draws i.i.d. samples from D'. We say that the distribution D' is an ϵ -corrupted version of the distribution D in total variation distance.

The distribution D' in Definition 1.2 can be adversarially selected (and can even depend on our learning algorithm). Since Huber's contamination model (Huber, 1964) only allows additive errors, TV-contamination is a stronger model.

STREAMING ALGORITHM FOR ROBUST MEAN ESTIMATION

The main result of this paper is the following (see Theorem D.2 for a more general statement):

Theorem 1.3 (Streaming Robust Mean Estimation). Let \mathcal{D} be a distribution family on \mathbb{R}^d and $0 < \epsilon < \epsilon_0$ for a sufficiently small constant $\epsilon_0 > 0$. Let P be an ϵ -corrupted version of D in total variation distance for some $D \in \mathcal{D}$ with unknown mean μ_D . There is a single-pass streaming algorithm that, given ϵ and \mathcal{D} , reads a stream of n i.i.d. samples from P, runs in sample near-linear time, uses memory d polylog (d/ϵ) , and outputs an estimate $\hat{\mu}$ that, with probability at least 9/10, satisfies the following:

- 1. If \mathcal{D} is the family of distributions with identity-bounded covariance, then $n = \tilde{O}\left(d^2/\epsilon\right)$ and $\|\hat{\mu} \mu_D\|_2 = O(\sqrt{\epsilon})$.
- 2. If \mathcal{D} is the family of identity-covariance subgaussian distributions, then $n = \tilde{O}\left(d^2/\epsilon^2\right)$ and $\|\widehat{\mu} \mu_D\|_2 = O(\epsilon\sqrt{\log(1/\epsilon)})$.

We note that the above error guarantees are information-theoretically optimal, even in absence of resource constraints. While prior work had obtained efficient robust mean estimators matching these error guarantees (Diakonikolas et al., 2016; 2017a; Steinhardt et al., 2018), all previous algorithms with dimension-independent error incurred space complexity $\Omega(d^2)$.

BEYOND ROBUST MEAN ESTIMATION

Using the algorithm of Theorem 1.3 as a black-box, we obtain the first efficient single-pass streaming algorithms with near-optimal space complexity for a range of more complex statistical tasks. These contributions are presented in detail in Appendix E. Here we highlight some of these results.

Our first application is a streaming algorithm for robust covariance estimation.

Theorem 1.4 (Robust Gaussian Covariance Estimation). Let Q be a distribution on \mathbb{R}^d with $d_{\mathrm{TV}}(Q, \mathcal{N}(0, \Sigma)) \leq \epsilon$ and assume $\frac{1}{\kappa}\mathbf{I}_d \preceq \Sigma \preceq \mathbf{I}_d$. There is a single-pass streaming algorithm that uses $n = (d^4/\epsilon^2)\mathrm{polylog}(d, \kappa, 1/\epsilon)$ samples from Q, runs in time $nd^2\mathrm{polylog}(d, \kappa, 1/\epsilon)$, uses memory $d^2\mathrm{polylog}(d, \kappa, 1/\epsilon)$, and outputs a matrix $\widehat{\Sigma}$ such that $\|\Sigma^{-1/2}\widehat{\Sigma}\Sigma^{-1/2} - \mathbf{I}_d\|_F = O(\epsilon\log(1/\epsilon))$ with probability at least 9/10.

See Theorem E.3 for a more detailed statement.

Our second application is for the general problem of robust stochastic optimization. Here we state two concrete results for robust linear and logistic regression (see Theorem E.9 and Theorem E.12 for more detailed statements). Both of these statements are special cases of a streaming algorithm for robust stochastic convex optimization (see Corollary E.6).

Theorem 1.5 (Streaming Robust Linear Regression). Let D be the distribution of (X,Y) defined by $Y=X^T\theta^*+Z$, where $X\sim \mathcal{N}(0,\mathbf{I}_d),\ Z\sim \mathcal{N}(0,1)$ independent of X, and $\|\theta^*\|_2\leq r$. Let P be an ϵ -corruption of D in total variation distance. There is a single-pass streaming algorithm that uses $n=(d^2/\epsilon)$ polylog $(d(1+r)/\epsilon)$, samples from P, runs in time nd polylog $(d(1+r)/\epsilon)$, uses memory d polylog (dr/ϵ) , and outputs an estimate $\widehat{\theta}\in\mathbb{R}^d$ such $\|\widehat{\theta}-\theta^*\|_2=O(\sqrt{\epsilon})$ with high probability.

Theorem 1.6 (Streaming Robust Logistic Regression). Consider the following model: Let $(X,Y) \sim D$, where $X \sim \mathcal{N}(0,\mathbf{I}_d)$, $Y \mid X \sim Bern(p)$, for $p=1/(1+e^{-X^T\theta^*})$, and $\|\theta^*\|_2 = O(1)$. Let P be an ϵ -corruption of D in total variation distance. There is a single-pass streaming algorithm that uses $n=(d^2/\epsilon)$ polylog (d/ϵ) samples from P, runs in time nd polylog (d/ϵ) , uses memory d polylog (d/ϵ) , and outputs an estimate $\widehat{\theta} \in \mathbb{R}^d$ such $\|\widehat{\theta} - \theta^*\|_2 = O(\sqrt{\epsilon})$ with

high probability.

Finally, in Appendix E, we include an additional application to distributed non-convex optimization in the streaming setting.

Remark 1.7 (Bit complexity). For simplicity of presentation, in the main body of the paper, we consider the model of computation where the algorithms can store and manipulate real numbers exactly. We show in Appendix L that our algorithms can tolerate errors due to finite precision. In particular, all our algorithms (including Algorithm 5) can be implemented in the word RAM model with d polylog (d/ϵ) bits.

1.2. Overview of Techniques

In this section, we provide a brief overview of our approach to establish Theorem 1.3. We start by recalling how robust mean estimation algorithms typically work without space constraints. A standard tool in the literature is the filtering technique of (Diakonikolas et al., 2016; 2017a; Diakonikolas & Kane, 2019). The idea of the filtering method is the following: Given a set S of corrupted samples, by analyzing spectral properties of the covariance of S, we can either certify that the sample mean of S is close to the true mean of the distribution, or can construct a filter. The filter is a method for selecting some elements of S to remove, with the guarantee that it will remove more outliers than inliers. If we can efficiently construct a filter, our algorithm can then remove the selected samples from S, obtaining a cleaner dataset and repeat the process. Eventually, this procedure must terminate, giving an accurate estimate of the true mean.

We proceed to explain how to implement the filtering method in a streaming model. We start with the easier case where the dataset is stored in read-only-memory, or more generally in a *multi-pass* streaming setting. At each round of the algorithm, one has a subset S' of the original dataset S that needs to be maintained (in particular, the set of samples that has survived the filters applied thus far). To do this naïvely would require n = |S| many bits of memory, which is too much for us. A more inventive strategy would be the following: instead of storing these subsets S'explicitly, store them implicitly by instead storing enough information to reconstruct the filters used to obtain S'. This seems like a productive idea, as most filters are relatively simple. For example, a commonly used filter is to remove all points $x \in S$ for which $v^T x > t$, for some vector v and scalar t. One could store enough information to apply this filter by merely storing (v,t), which would take O(d)bits of information. Unfortunately, most filtering algorithms may require $\Omega(d)$ many iterations before attaining their final answer. Consequently, the sets S' one needs to store are not just the result of applying a single filter, but instead the result of iteratively applying $\Omega(d)$ of them. In order to

store all of these extra filters, one would need $\Omega(d^2)$ bits. (For the sake of this intuitive description, we focused on "hard-thresholding" filters. Our algorithm will actually use a soft-thresholding filter, assigning weights to each point.)

To circumvent this first obstacle, one requires as a starting point a filtering algorithm that is guaranteed to terminate after a small (namely, at most poly-logarithmic) number of iterations. Recent work (Dong et al., 2019; Diakonikolas et al., 2021b) has obtained such algorithms. Here we generalize and simplify the filtering method of (Diakonikolas et al., 2021b). This allows us to obtain an algorithm with space complexity d polylog (d/ϵ) that works in the $multipass\ streaming\ model$, where $polylog(d/\epsilon)$ passes over the same dataset are allowed.

To obtain a *single-pass* streaming algorithm, new ideas are required. In the single-pass setting, we cannot implicitly store a subset of the full dataset S; once we access some points from S, we will never be able to see them again. To deal with this issue, we will need to slightly alter our way of thinking about the algorithm. Instead of being given a set S of samples, an ϵ -fraction of which have been corrupted, we instead adopt the view of having sample access to a distribution P, which is ϵ -close in total variation distance to the inlier distribution G. Given this point of view, instead of a filter defining a procedure for removing samples from S and outputting a subset S', we think of it as a rejection sampling procedure that replaces P with a cleaner distribution P'.

This shift in perspective comes with new technical challenges. In particular, when constructing the next round of filters, we will need to compute quantities pertaining to the current distribution P of the data points. In the setting of the multiple-pass model, this imposed no problem; these quantities could be calculated exactly. This is no longer possible when we merely have sample access to P. The best one can hope for is to approximate these quantities to sufficient precision for the rest of our analysis to carry over. However, the natural estimators for some required quantities (e.g., powers of the covariance matrix) would need to access the data multiple times. Circumventing this issue requires non-trivial technical work. Roughly speaking, instead of iterating over the same dataset to approximate the desired quantities, we show that it suffices to iterate over statistically identical datasets.

1.3. Prior and Related Work

Since the dissemination of (Diakonikolas et al., 2016; Lai et al., 2016), there has been an explosion of research in algorithmic aspects of robust statistics. We now have efficient robust estimators for a range of more complex problems, including covariance estimation (Diakonikolas et al., 2016; Cheng et al., 2019a), sparse estimation tasks (Balakrishnan et al., 2017; Diakonikolas et al., 2019b; Cheng et al., 2021),

learning graphical models (Cheng et al., 2018; Diakonikolas et al., 2021c), linear regression (Klivans et al., 2018; Diakonikolas et al., 2019c; Pensia et al., 2020), stochastic optimization (Prasad et al., 2020; Diakonikolas et al., 2019a), and robust clustering/learning various mixture models (Hopkins & Li, 2018; Kothari et al., 2018; Diakonikolas et al., 2018; 2020a; 2021b; Bakshi et al., 2020a; Liu & Moitra, 2020; Bakshi et al., 2020b). The reader is referred to (Diakonikolas & Kane, 2019) for a detailed overview. We reiterate that all previously developed algorithms work in the batch setting, i.e., require the entire dataset in memory.

For the problem of robust mean estimation, (Dong et al., 2019; Diakonikolas et al., 2021b) gave filtering-based algorithms with a poly-logarithmic number of iterations. The former algorithm relies on the matrix multiplicative weights framework, while the latter is based on first principles. Our starting point in Section 3 can be viewed as a generalization and further simplification of the ideas in (Diakonikolas et al., 2021b). Specifically, our algorithm works under the stability condition (Definition 2.1), which broadly generalizes the bounded covariance assumption used in (Diakonikolas et al., 2021b).

In the context of robust supervised learning (including, e.g., our robust linear regression application), low-space streaming algorithms are known in weaker contamination models that only allow *label* corruptions, see, e.g., (Pesme & Flammarion, 2020; Shah et al., 2020; Diakonikolas et al., 2020c). We emphasize that the contamination model of Definition 1.2 is significantly more challenging, and no low-space streaming algorithms were previously known in this model. In particular, naive methods of outlier removal do not lead to dimension-independent errors, see, for example, Section 3 of (Diakonikolas et al., 2016), Section 2 of (Lai et al., 2016), and Section 1 of (Diakonikolas & Kane, 2019).

We note that the focus of our work is orthogonal to the that of (Sharan et al., 2019) that studies memory-sample tradeoff for linear regression *without* any outliers.

Finally, we note that recent work (Tsai et al., 2021) studies streaming algorithms for heavy-tailed stochastic optimization. While the goal of developing low-space streaming algorithms is qualitatively similar to the goal of our work, the algorithmic results in (Tsai et al., 2021) have no implications in the corrupted setting studied in this work.

2. Preliminaries

Basic Notation We use \mathbb{Z}_+ to denote the set of positive integers. For $n \in \mathbb{Z}_+$, we denote $[n] := \{1, \ldots, n\}$. For a vector v, we use $\|v\|_2, \|v\|_\infty$ for its ℓ_2 and infinity norms respectively. We let \otimes denote the Kronecker product for vectors. We use boldface letters for matrices. We use \mathbf{I}_d to denote the $d \times d$ identity matrix. We use $\|\mathbf{A}\|_F$ for

the Frobenius norm of \mathbf{A} . For a square matrix \mathbf{A} , we use $\|\mathbf{A}\|_2$ for its spectral norm and $\operatorname{tr}(\mathbf{A})$ for its trace. We use \preceq for the Löwner ordering of matrices. Finally, \mathbf{A}^{\flat} denotes the vector obtained by concatenating the rows of \mathbf{A} . For a distribution D, we denote $\mu_D = \mathbf{E}_{X \sim D}[X]$ and $\mathbf{\Sigma}_D = \mathbf{E}_{X \sim D}[(X - \mu_D)(X - \mu_D)^T]$. Given a weight function $w: \mathbb{R}^d \to [0,1]$, we define the re-weighted distribution D_w to be $D_w(x) := D(x)w(x)/\int_{\mathbb{R}^d} w(x)D(x)\mathrm{d}x$. We use $\mu_{w,D} = \mathbf{E}_{X \sim D_w}[X]$ for its mean and $\overline{\mathbf{\Sigma}}_{w,D}^\mu = \mathbf{E}_{X \sim D_w}[(X - \mu)(X - \mu)^T]$ for the second moment centered with respect to μ (we will usually drop μ from the notation when it is clear from the context). We use $d_{\mathrm{TV}}(P,Q)$ for the total variation distance between P,Q. For a set S, we use $\mathcal{U}(S)$ for the uniform distribution on S.

2.1. Stability Condition

Our algorithms succeed for every distribution satisfying the following standard set of deterministic conditions. (See Appendix B.2 for more details.)

Definition 2.1 $((\epsilon, \delta)$ -stable distribution). Fix $0 < \epsilon < 1/2$ and $\delta \ge \epsilon$. A distribution G on \mathbb{R}^d is (ϵ, δ) -stable with respect to $\mu \in \mathbb{R}^d$ if for any weight function $w : \mathbb{R}^d \to [0, 1]$ with $\mathbf{E}_{X \sim G}[w(X)] \ge 1 - \epsilon$ we have that

$$\left\|\mu_{w,G} - \mu\right\|_2 \leq \delta \quad \text{and} \quad \left\|\overline{\mathbf{\Sigma}}_{w,G} - \mathbf{I}_d\right\|_2 \leq \delta^2/\epsilon \;.$$

We can similarly define the same notion for sets.

Definition 2.2 $((\epsilon, \delta)$ -stable set). Fix $0 < \epsilon < 1/2$ and $\delta \ge \epsilon$. A finite set $S_0 \subset \mathbb{R}^d$ is (ϵ, δ) -stable with respect to $\mu \in \mathbb{R}^d$ if the empirical distribution $\mathcal{U}(S_0)$ is (ϵ, δ) -stable with respect to μ .

Fact 2.3 ((Diakonikolas & Kane, 2019)). A set of $O(d/(\epsilon^2 \log(1/\epsilon)))$ i.i.d. samples from an identity covariance subgaussian distribution is $(\epsilon, O(\epsilon \sqrt{\log(1/\epsilon)}))$ -stable with respect to μ with high probability. Similarly, a set of $\tilde{O}(d/\epsilon)$ i.i.d. samples from a distribution X with $\mathbf{Cov}[X] \leq \mathbf{I}_d$ contains a large subset S, which is $O(\epsilon, O(\sqrt{\epsilon}))$ -stable with respect to its mean $\mathbf{E}[X]$ with high probability.

All known filtering techniques rely on the following lemma, stating that if the empirical covariance is not too large in the Löwner order, while the distribution is close to the original in total variation distance, then the sample mean is a good estimate of the true mean.

Lemma 2.4 (Certificate Lemma). Let G be an (ϵ, δ) -stable distribution with respect to $\mu \in \mathbb{R}^d$, for some $0 < \epsilon < 1/3$ and $\delta \ge \epsilon$. Let P be a distribution with $d_{\mathrm{TV}}(P,G) \le \epsilon$. Denoting by μ_P, Σ_P the mean and covariance of P, if $\lambda_{\max}(\Sigma_P) \le 1 + \lambda$, for some $\lambda \ge 0$, then $\|\mu_P - \mu\|_2 = O(\delta + \sqrt{\epsilon \lambda})$.

3. Filtering Algorithm with Small Number of Iterations

In this section, we develop a filtering algorithm (in the batch setting) that terminates in $\operatorname{polylog}(d/\epsilon)$ iterations for any stable set. This leads to an algorithm that runs in nearlinear time, i.e., $nd\operatorname{polylog}(nd/\epsilon)$, generalizing the results of (Dong et al., 2019; Diakonikolas et al., 2021b). Crucially, this algorithm will form the building block of our streaming algorithm in Section 4. We remark that the algorithm of this section works even against the $strong-contamination\ model$ (Definition C.1), where the outliers may not be i.i.d. samples from any distribution, but are allowed to be completely arbitrary. We prove the following (see also Theorem C.2).

Theorem 3.1 (informal). Let $0 < \epsilon < \epsilon_0$, for a sufficiently small constant ϵ_0 . Let S_0 be a set of n points in \mathbb{R}^d that is (ϵ, δ) -stable with respect to μ and S be an ϵ -corrupted version of S_0 . There exists an algorithm that runs in time $nd \operatorname{polylog}(nd/\epsilon)$ and with high probability finds a vector $\widehat{\mu}$ such that $\|\widehat{\mu} - \mu\|_2 = O(\delta)$.

Theorem 3.1 applies to any stable set. By Fact 2.3, we directly obtain (i) an $O(\epsilon \sqrt{\log(1/\epsilon)})$ -accurate estimator given $O(d/(\epsilon^2/\log(1/\epsilon)))$ many ϵ -corrupted samples from an identity covariance subgaussian distribution; and (ii) an $O(\sqrt{\epsilon})$ -accurate estimator for any distribution $X \sim D$ with $\mathbf{Cov}[X] \preceq \mathbf{I}_d$, given $\tilde{O}(d/\epsilon)$ many ϵ -corrupted samples.

The pseudocode of the algorithm establishing Theorem 3.1 is presented in Algorithm 1. We will define the necessary notation as needed (see the pseudocode for details). First, we assume that the distribution over the input samples is of the form $P = (1 - \epsilon)G + \epsilon B$, where G is the uniform distribution over the stable set of inliers and B is the uniform distribution on the outliers. Although this mixture may seem to suggest that the adversary only adds points, it is without loss of generality. Indeed, in the case that the adversary also removes points, we can think of G as the distribution of the remaining inliers (which continues to be stable with slightly worse parameters; see Lemma B.12).

We begin with a high-level explanation of Algorithm 1. At each iteration t, we assign a weight $w_t(x) \in [0,1]$ to each point x. Let P_t be the distribution on S, weighted according to w_t . Let μ_t and Σ_t be the mean and covariance of P_t , respectively. We want to assign scores to each point, using spectral properties of Σ_t and the stability of inliers, so that the scores over outliers are more than those of inliers. Essentially, if a direction v has variance larger than $1 + \Omega(\delta^2/\epsilon)$, then the stability of inliers implies that this must be due to outliers. Thus, we can assign scores based on the values $(v^T(x-\mu_t))^2$ that have provably more mass on outliers than inliers. The filters proposed in (Diakonikolas et al., 2016; 2017a) assigned scores based on a single direction, the leading eigenvector of Σ_t , and can take as many as $\Omega(d)$

iterations (see Section 1.2).

To reduce the number of iterations, we need to filter in all directions of large variance simultaneously. Letting $\mathbf{B}_t \approx \mathbf{\Sigma}_t - (1 - C_1 \delta^2 / \epsilon) \mathbf{I}_d$, we would like to filter along all directions where the eigenvalue of \mathbf{B}_t is within a constant factor from $\lambda_t := \|\mathbf{B}_t\|_2$, not necessarily the leading eigenvector of \mathbf{B}_t . As we show in Section 3.2, this can be approximately achieved by assigning scores for each point x based on $g_t(x) := \|\mathbf{M}_t(x - \mu_t)\|_2^2$, where $\mathbf{M}_t = \mathbf{B}_t^{\log d}$. At a high level, this happens because the spectrum of \mathbf{M}_t is distributed across along all large eigenvectors of \mathbf{B}_t .

```
Algorithm 1 Robust Mean Estimation in polylog iterations
```

```
1: Input: S = \{x_i\}_{i \in [n]}, \delta, \epsilon
 2: Let C_1 \geq 22, C be a sufficiently large constant, C_2 =
      100C and C_3 = 0.1.
 3: Let R = \sqrt{(d/\epsilon)(1+\delta^2/\epsilon)}.
 4: Let P = (1 - \epsilon)G + \epsilon B be the empirical distribution
      on the points from S. <sup>2</sup>
 5: Let K = \text{polylog}(d/\epsilon), L = \text{polylog}(nd/\epsilon).
 6: Obtain a naïve estimate \widehat{\mu} of \mu with \|\widehat{\mu} - \mu\|_2 \le 4R.
 7: Initialize w_1(x) \leftarrow \mathbb{1}\{\|x - \widehat{\mu}\|_2 \leq 5R\} for all x \in S.
 8: for t \in [K] do
 9:
          Let P_t be the distribution of P weighted by w_t.
10:
          Let \mu_t, \Sigma_t be the mean and covariance of P_t.
         Let \mathbf{B}_t = (\mathbf{E}_{X \sim P}[w_t(X)])^2 \mathbf{\Sigma}_t - \left(1 - C_1 \frac{\delta^2}{\epsilon}\right) \mathbf{I}_d and \mathbf{M}_t = \mathbf{B}_t^{\log d}.
          Let \lambda_t = \|\mathbf{B}_t\|_2. Compute \widehat{\lambda}_t \in [0.8\lambda_t, 1.2\lambda_t] using
12:
          power iteration.
          if \hat{\lambda}_t > C_2 \delta^2 / \epsilon then
13:
              for j \in [L] do
14:
                   z_{t,j} \sim \dot{\mathcal{U}}(\{\pm 1\}^d), v_{t,j} \leftarrow \mathbf{M}_t z_{t,j}.
15:
16:
              Denote by \mathbf{U}_t the matrix having the vectors
17:
              \frac{1}{\sqrt{L}}v_{t,j} for j \in [L] as rows.
              Let \tilde{g}_t(x) = \|\mathbf{U}_t(x - \mu_t)\|_2^2 and
18:
              \tilde{\tau}_t(x) = \tilde{g}_t(x) \mathbb{1}\{\tilde{g}_t(x) > C_3 \|\mathbf{U}_t\|_F^2 \hat{\lambda}_t/\epsilon\},
              \ell_{\text{max}} \leftarrow (dR/\epsilon)^{C \log d}, T \leftarrow 0.01 \hat{\lambda}_t \|\mathbf{U}_t\|_F^2.
19:
                                              DownweightingFilter(P, w_t,
20:
               \tilde{\tau}_t, R, T, \ell_{\max}
          end if
21:
22: end for
```

Even though assigning scores based on \mathbf{M}_t , i.e., $g_t(x)$, reduces the number of iterations, computing $g_t(x)$ for all $x \in S$ is slow. We thus use a Johnson-Lindenstrauss (JL) sketch of \mathbf{M}_t , denoted by \mathbf{U}_t . We denote by $\tilde{g}_t(x) := \|\mathbf{U}_t(x-\mu_t)\|_2$ the resulting scores. We claim that the set

23: **return** μ_t .

²Without loss of generality, outliers are within O(R) from μ in ℓ_2 -norm. This is ensured in Line 7, which removes only ϵ -fraction of inliers (Claim C.12).

Algorithm 2 Downweighting Filter

- 1: **Input**: $P, w, \tilde{\tau}, R, T, \ell_{\max}$
- 2: $r \leftarrow CdR^{2+4\log d}$.
- 3: Let $w_{\ell}(x) = w(x)(1 \tilde{\tau}(x)/r)^{\ell}$.
- 4: Find the smallest $\ell \in \{1, \dots, \ell_{\max}\}$ satisfying $\mathbf{E}_{X \sim P} \left[w_{\ell}(X) \tilde{\tau}(X) \right] \leq 2T$ using binary search.
- 5: **return** w_{ℓ} .

 $\{\tilde{g}_t(x)\}_{x\in S}$ can be calculated in time $\tilde{O}(nd)$ such that for each $x\in S$, $\tilde{g}_t(x)\approx g_t(x)$. First, we will show (see Lemma C.5) that \mathbf{U}_t can be as small as $L\times d$, where $L=\operatorname{polylog}(nd/\epsilon)$, which follows from the classical JL lemma (stating that n points can be linearly embedded into a $\log n$ -dimensional space). Also, each row of \mathbf{U}_t can be computed by repeatedly multiplying a vector $\log d$ times by \mathbf{B}_t (Line 15). Thus, by also noting that for any v, $\mathbf{\Sigma}_t v = \sum_{x\in S} w_t(x)(v^T(x-\mu_t))(x-\mu_t)/(\sum_{x\in S} w_t(x))$ can be computed in O(nd) time by calculating μ_t and $v^T(x-\mu_t)$ first, all rows of \mathbf{U}_t can be computed in time $\tilde{O}(nd)$. Thus, each iteration of Algorithm 1 runs in nearlinear time.

For the proof of correctness, we require that the JL and spectral approximations used by the algorithm are sufficiently accurate. We prove in Appendix C.2 that the following event occurs with high probability.

Condition 3.2 (Deterministic Conditions For Algorithm 1). For all $t \in [K]$, the following hold:

- 1. Spectral norm of \mathbf{B}_t : $\widehat{\lambda}_t \in [0.8\lambda_t, 1.2\lambda_t]$.
- 2. Frobenius norm: $\|\mathbf{U}_t\|_F^2 \in [0.8\|\mathbf{M}_t\|_F^2, 1.2\|\mathbf{M}_t\|_F^2]$.
- 3. *Scores*: For all $x \in S$, $\tilde{q}_t(x) \in [0.8q_t(x), 1.2q_t(x)]$.

3.1. Downweighting Filter

The final scores $\tilde{\tau}_t(x)$ used in filtering are thresholded versions of $\tilde{g}_t(x)$, as defined in Line 18. Recall that we have $P=(1-\epsilon)G+\epsilon B$, where G is the uniform distribution over inliers satisfying stability and B is the uniform distribution of the outliers. Roughly speaking, our filter guarantees two things when modifying the weights $w_t(x)$ to $w_{t+1}(x)$ (see Lemma C.6):

- 1. The weight removed from the outliers is greater than the weight removed from the inliers.
- 2. $\mathbf{E}_{X \sim P}[w_{t+1}(X)\tilde{\tau}_t(X)] \leq 2\mathbf{E}_{X \sim G}[w_t(X)\tilde{\tau}_t(X)]$, i.e., the weighted mean of scores after filtering over both inliers and outliers is at most twice the weighted mean of scores of inliers *before* filtering.

Moreover, Algorithm 2 terminates in $O(n \log(\ell_{\text{max}}))$ time. The first guarantee ensures that P_t will be never be more

than $O(\epsilon)$ -far in total variation distance from G. Regarding the second item above, using straightforward implications of stability (Appendix B.2), we show in Appendix C.2 that $\mathbf{E}_{X\sim G}[w_t(X)\tilde{\tau}_t(X)]$ is small, and thus filtering will make the corresponding expectation for both inliers and outliers comparably small. Let $\tau_t(x)$ be the thresholded scores defined using \mathbf{M}_t . Then we have the following:

Lemma 3.3 (Informal). Under the deterministic Condition 3.2, we have that $\epsilon \mathbf{E}_{X \sim B}[w_{t+1}(X)\tilde{\tau}_t(X)] < c\lambda_t \|\mathbf{M}_t\|_F^2$ for some constant $c \leq 0.01$. As a consequence, it also holds $\epsilon \mathbf{E}_{X \sim B}[w_{t+1}(X)\tau_t(X)] < c\lambda_t \|\mathbf{M}_t\|_F^2$.

The second part of the lemma holds because Item 3 of Condition 3.2 implies that $\tilde{\tau}_t(x)$ is close to $\tau_t(x)$ (Claim C.10). Lemma 3.3 is crucial towards proving our main theorem, since it essentially means that filtering reduces the variance by a multiplicative factor in the directions of \mathbf{M}_t .

3.2. Correctness of Algorithm 1: Proof of Theorem 3.1

Recall that each iteration of Algorithm 1 can be implemented in near-linear time. Thus, it remains to show that the choice $K = \operatorname{polylog}(d/\epsilon)$ suffices to guarantee correctness of our algorithm. We now sketch the proof using a potential function argument. Let Λ_t be the vector in \mathbb{R}^d containing the eigenvalues of \mathbf{B}_t . Recall that our goal is to show that $\|\mathbf{B}_t\|_2 = \|\Lambda_t\|_\infty = O(\delta^2/\epsilon)$ in polylog many iterations. Let $p{=}2\log d$. Since $\|x\|_p = \Theta(\|x\|_\infty)$ for any $x \in \mathbb{R}^d$, we are motivated to use the potential function $\phi_t := \|\Lambda_t\|_p^p$. We now focus on showing that ϕ_t decreases rapidly. Observe that for any $i \in \mathbb{Z}_+$, $\operatorname{tr}(\mathbf{B}_t^i) = \|\Lambda_t\|_i^i$. We start with the following inequalities (and explain them directly below):

$$\phi_{t+1} = \|\Lambda_{t+1}\|_{p}^{p} \le \left(d^{\frac{1}{p(p+1)}} \|\Lambda_{t+1}\|_{p+1}\right)^{p} \\
= d^{\frac{1}{p+1}} \left(\|\Lambda_{t+1}\|_{p+1}^{p+1}\right)^{\frac{p}{p+1}} \\
= d^{\frac{1}{p+1}} (\operatorname{tr}(\mathbf{B}_{t+1}^{p+1}))^{\frac{p}{p+1}} \\
= d^{\frac{1}{p+1}} (\operatorname{tr}(\mathbf{M}_{t+1}\mathbf{B}_{t+1}\mathbf{M}_{t+1}))^{\frac{p}{p+1}} \\
\le d^{\frac{1}{p}} (\operatorname{tr}(\mathbf{M}_{t}\mathbf{B}_{t+1}\mathbf{M}_{t}))^{\frac{p}{p+1}}, \tag{1}$$

where the first line uses Fact B.1, the third one uses $\operatorname{tr}(\mathbf{B}_{t+1}^i) = \|\Lambda_{t+1}\|_i^i$, the fourth one $\mathbf{M}_{t+1}\mathbf{B}_{t+1}\mathbf{M}_{t+1} = \mathbf{B}_{t+1}^{p+1}$, and the last one uses the fact $\mathbf{B}_{t+1} \preceq \mathbf{B}_t$, which holds because removing points can only make their covariance smaller; see Appendix C.4 for more details.

Then the goal becomes to bound from above the term $\operatorname{tr}(\mathbf{M}_t\mathbf{B}_{t+1}\mathbf{M}_t)$. As will be shown below, we have that $\operatorname{tr}(\mathbf{M}_t\mathbf{B}_{t+1}\mathbf{M}_t)$ is related to $\mathbf{E}_{X\sim P}[w_{t+1}(X)\tau_t(X)]$. Using the guarantees of the Downweighting filter (Lemma 3.3), we prove the following result in Appendix C.4.2 (see Lemma C.16):

Lemma 3.4 (informal). Assume Condition 3.2 holds. Then $\operatorname{tr}(\mathbf{M}_t \mathbf{B}_{t+1} \mathbf{M}_t) \leq c \lambda_t \|\mathbf{M}_t\|_F^2$ for some constant c < 0.1.

Having this at hand, we continue with Equation (3) as follows:

$$\begin{split} \phi_{t+1} &\leq d^{\frac{1}{p}} \left(\text{tr}(\mathbf{M}_{t} \mathbf{B}_{t+1} \mathbf{M}_{t}) \right)^{\frac{p}{p+1}} \\ &\leq d^{\frac{1}{p}} \left(c \| \Lambda_{t} \|_{\infty} \| \Lambda_{t} \|_{p}^{p} \right)^{\frac{p}{p+1}} \quad \text{(using Lemma 3.4)} \\ &\leq d^{\frac{1}{p}} c^{\frac{p}{p+1}} \left(\| \Lambda_{t} \|_{p} \| \Lambda_{t} \|_{p}^{p} \right)^{\frac{p}{p+1}} \\ &= d^{\frac{1}{p}} c^{\frac{p}{p+1}} \| \Lambda_{t} \|_{p}^{p} \\ &\leq 3\sqrt{c} \| \Lambda_{t} \|_{p}^{p} \leq 0.9999 \phi_{t} \;, \end{split}$$

where the third line uses that $\|\Lambda_t\|_{\infty} \leq \|\Lambda_t\|_i$ for $i \geq 1$, and the last line uses that $d^{1/p} = \exp((\log d)/(2\log d)) \leq 3$, $p/(p+1) \geq 0.5$, and c < 0.1. We thus get the desired convergence.

The final step is to bound from above the number of iterations needed for Lemma 2.4 to ensure that $\|\mu_t - \mu\|_2 = O(\delta)$. Concretely, due to our naı̈ve pruning, at the beginning of the algorithm we have the upper bound $\phi_1 \leq dR^{O(\log d)}$. In the t-th iteration, if the condition of Line 13 still holds, we have that $\phi_t \leq 0.99^t dR^{O(\log d)}$. Thus, when $t = \text{polylog}(d/\epsilon)$, we have that $\|\mathbf{B}_t\|_2 \leq (\phi_t)^{\frac{1}{2\log d}} = O(\delta^2/\epsilon)$. Otherwise, if the condition of Line 13 is violated for any t, then we directly have $\|\mathbf{B}_t\|_2 = O(\delta^2/\epsilon)$. As a consequence, we obtain that $\|\mathbf{\Sigma}_t\|_2 \leq 1 + O(\delta^2/\epsilon)$ for t = K. An application of Lemma 2.4 shows that the final weighted mean μ_t for t = K has error at most $\|\mu_t - \mu\|_2 = O(\delta)$, completing the proof of Theorem 3.1.

4. Efficient Streaming Algorithm for Robust Mean Estimation

We now turn to the main focus of our paper, and present a streaming algorithm with near-optimal space complexity for robust mean estimation. To obtain our streaming algorithm, we will appropriately modify Algorithm 1 from the previous section. This will eventually lead to a single-pass streaming algorithm with near-optimal space complexity. Due to space limitations, the final pseudocode of our streaming algorithm is Algorithm 5 in the appendix. In this section, we give an overview of the key modifications to Algorithm 1 required in tandem with a sketch of the analysis.

Moving to the streaming model requires a change in perspective: instead of having a corrupted dataset, we now have sample access to a distribution P such that $d_{\text{TV}}(P,D) \leq \epsilon$, where D is a stable distribution. We will reweight this distribution using weights, $w_t(\cdot)$, that are now functions on the whole \mathbb{R}^d instead of a fixed dataset. Thus, P_t now denotes the reweighting of the (corrupted) distribution P_t with the weights w_t . Similarly μ_t , Σ_t , \mathbf{B}_t , \mathbf{M}_t denote the quantities that pertain to the distribution P_t . The goal of our algorithm remains essentially the same: obtain P_t such that $d_{\text{TV}}(P_t, P) = O(\epsilon)$ and $\|\Sigma_t\|_2 \leq 1 + O(\delta^2/\epsilon)$; Lemma 2.4 would then imply that $\|\mu_t - \mu\|_2 = O(\delta)$.

Since weights are now functions over \mathbb{R}^d and not stored anywhere, we need an efficient way to compute $w_t(x)$ whenever required. As outlined in Section 1.2, it is sufficient to store a representation of the filters, i.e., the matrices \mathbf{U}_t and the counts ℓ_t from the downweighting filter. Then, given a point x, $w_t(x)$ can be computed in poly-logarithmically many operations. Moreover, we need a way to simulate samples from the weighted distributions P_t , which can be done using rejection sampling. We will follow the structure of Algorithm 3 and show the following: (i) it is possible to approximate the necessary quantities in Algorithm 1 using i.i.d. samples from P in the streaming model, and (ii) the size of the JL-sketch can be kept small.

We begin with the first obstacle of calculating approximate scores. Recall that the only place where M_t is used in Algorithm 1 is Line 15, where M_t is multiplied with the vectors $z_{t,j}$. Let z be an arbitrary vector. Since $\mathbf{M}_t = \mathbf{B}^{\log d}$, in the previous section we were able to compute $M_t z$ by iteratively multiplying z by \mathbf{B}_t . Since we now do not have access to \mathbf{B}_t , but only sample access to P_t , we need a sufficiently fine approximation \mathbf{B}_t of \mathbf{B}_t (obtained using i.i.d. samples). The natural approach would then be to multiply \mathbf{B}_t with z iteratively $\log d$ many times. Even though $\mathbf{B}_t z$ can be computed in a streaming fashion (as outlined in the previous section), it is not possible to compute $(\mathbf{B}_t)^{\log d}z$ without accessing the data $\log d$ times. To circumvent this issue, we use a fresh sample approximation of \mathbf{B}_t in every multiplication step. That is, we approximate $\mathbf{M}_t z$ with $\mathbf{\hat{M}}_t z$, where $\widehat{\mathbf{M}}_t := \prod_{j=1}^p \widehat{\mathbf{B}}_{t,j}$ and each $\widehat{\mathbf{B}}_{t,j}$ is computed on a different set of samples. This approach crucially leverages the fact that in the contamination model of Definition 1.2, outliers are added in a way that is oblivious to the inliers, and therefore these datasets are statistically identical and independent of each other. We show in Appendix D.3 that the resulting $\widehat{\mathbf{M}}_t$ is a sufficiently accurate approximation of

Lemma 4.1 (informal). Let the estimator $\widehat{\mathbf{M}}_t := \prod_{j=1}^p \widehat{\mathbf{B}}_{t,j}$, where $\widehat{\mathbf{B}}_{t,j}$ is the empirical version of \mathbf{B}_t on the same number of samples as in Theorem 1.3. Then it holds that $\|\widehat{\mathbf{M}}_t - \mathbf{M}_t\|_2 \le O(1/\sqrt{d}) \|\mathbf{M}_t\|_F$ with high probability.

Similarly, we need to modify the downweighting filter, since its implementation using binary search requires performing checks of the form $\mathbf{E}_{X\sim P}[w(X)\tilde{\tau}(x)]>2T$ and calculating the weighted mean exactly is no longer possible. We propose a sample-efficient estimator to approximate that expectation (see Lemma D.16 in Appendix D.3.3) and run an "approximate" variant of binary search (see Appendix D.2).

We now turn to the more technical issue of controlling the size of the JL-sketch, i.e., the number of rows, L, of the matrix $\mathbf{U}_t \in \mathbb{R}^{L \times d}$. For simplicity, assume $\widehat{\mathbf{M}}_t = \mathbf{M}_t$, and recall that $\tilde{\tau}(x)$ is the thresholded version of $\|\mathbf{U}_t(x - \mu_t)\|_2^2$,

as defined in Line 18 and $\tau(x)$ is the same score but using \mathbf{M}_t . The potential-based analysis in Section 3.2 requires that $\mathbf{E}_{X\sim P}[w_{t+1}(X)\tau_t(X)]$ is small. However, the stopping condition of the downweighting filter only implies that $\mathbf{E}_{X\sim P}[w_{t+1}(X)\tilde{\tau}_t(X)]$ is small. In the previous section, the bound on the former was obtained from the bound on the latter by using that $\|\mathbf{U}_t(x-\mu_t)\|_2 \approx \|\mathbf{M}_t(x-\mu_t)\|_2$ pointwise in the support of P (Claim C.10).

By the classical JL lemma, the size of the JL sketch, L, needs to be at most logarithmic in the size of the set Swhere we require the pointwise approximation to hold. Thus, in the previous section, L scaled as $\log |S| = \log n$. However, in the streaming model where there is no such dataset, it is far from obvious how the analysis should proceed. A näive approach would be to require the approximation to hold on a cover \tilde{S} of the support of P_t . Since $|\tilde{S}|$ scales exponentially with d, the required bound on L would be $\log |S| = \Omega(d)$, which is too large for our purposes. Luckily, we can still find a fixed set S_{cover} such that the following holds: (i) $\log |S_{\text{cover}}| = \text{polylog}(d/\epsilon)$ and (ii) the expectation of scores over $\mathcal{U}(S_{\text{cover}})$ approximates the expectation of scores over P. That is, as far as the expectation of the scores are concerned, P can be approximated by the uniform distribution over S_{cover} . Arguing as before, if $\|\mathbf{U}_t(x-\mu_t)\| \approx \|\mathbf{M}_t(x-\mu_t)\|$ for each $x \in S_{\mathrm{cover}}$, then the downweighting filter also ensures that $\mathbf{E}_{X \sim P}[w_{t+1}(X)\tilde{\tau}_t(X)]$ is small. Thus, S_{cover} can serve as a proxy dataset (only used in the analysis) to ensure that the size of the JL sketch is sufficiently bounded, i.e., that $L = \log |S_{\text{cover}}| = \text{polylog}(d/\epsilon).$

Establishing the desired upper bound on the cardinality of $S_{\rm cover}$ requires a somewhat more sophisticated argument that relies on the VC-dimension of a family of functions corresponding to the weight update rule. Due to space limitations, this proof is deferred to Appendix D.2.1.

In summary, the algorithm for the streaming setting (Algorithm 5) is obtained via a number of crucial adaptations of Algorithm 1. Specifically, we perform the following major modifications: (i) sample approximations for \mathbf{M}_t and μ_t , (ii) a slightly larger JL-sketch, (iii) an approximate binary search in the downweighting filter, and (iv) storing only the representations ℓ_t of $w_t(x)$'s and the JL-sketches \mathbf{U}_t in memory. The analysis is again based on a set of high probability events (Condition D.5) similar to Condition 3.2. Due to space limitations, the full algorithm and the proofs of the technical arguments are deferred to Appendix D.3.

Given the aforementioned differences, we again use the same potential function as in Section 3.2, and show that it decreases by a multiplicative factor in every round. This completes the analysis sketch of our streaming algorithm, thereby completing the proof sketch of Theorem 1.3.

5. Applications: Beyond Robust Mean Estimation

In this section, we develop robust streaming algorithms with near-optimal space complexity for more complex statistical tasks, specifically for robust covariance estimation and robust stochastic optimization. The main idea enabling these applications is that these tasks can be effectively reduced to robust mean estimation. (Due to space limitations, detailed statements and proofs are deferred to Appendix E.)

Robust Covariance Estimation Let Σ be the covariance matrix of a centered distribution D, and define $Y:=X\otimes X$ for $X\sim D$, where \otimes denotes the Kronecker product. Noting that $\mathbf{E}[Y]=\mathbf{\Sigma}^{\flat}$ (where $\mathbf{\Sigma}^{\flat}$ denotes the vector obtained by concatenating all the rows of $\mathbf{\Sigma}$), estimating $\mathbf{\Sigma}$ reduces to estimating $\mathbf{E}[Y]$. Thus, as long as $\mathbf{Cov}[Y] \preceq O(1)\mathbf{I}_{d^2}$, Theorem 1.3 yields an estimator $\hat{\mathbf{\Sigma}}$ such that $\|\hat{\mathbf{\Sigma}} - \mathbf{\Sigma}\|_F = O(\sqrt{\epsilon})$ that uses memory at most $\tilde{O}(d^2)$ (see Theorem E.1 for details). Moreover, the bounded fourth moment assumption is satisfied in the case $D=\mathcal{N}(0,\mathbf{\Sigma})$ with $\mathbf{\Sigma}\preceq\mathbf{I}_d$ (see Fact E.2). Therefore, the theorem covers the Gaussian case with bounded covariance. Using a more sophisticated approach (see (Cheng et al., 2019b) and Appendix \mathbf{E}), we can achieve the near-optimal error of $O(\epsilon \log(1/\epsilon))$ for the special case of the Gaussian distribution.

Theorem 5.1 (Streaming Robust Covariance Estimation). Let Q be such that $d_{\mathrm{TV}}(Q, \mathcal{N}(0, \Sigma)) \leq \epsilon$ and assume that $\frac{1}{\kappa}\mathbf{I}_d \preceq \Sigma \preceq \mathbf{I}_d$. There exists a single-pass streaming algorithm that draws $n = (d^4/\epsilon^2)\mathrm{polylog}(d\kappa/\epsilon)$ samples from Q, runs in time $nd^2\mathrm{polylog}(d\kappa/\epsilon)$, has space complexity $d^2\mathrm{polylog}(d\kappa/\epsilon)$, and returns a hypothesis $\widehat{\Sigma} \in \mathbb{R}^{d \times d}$ such that $\|\Sigma^{-1/2}\widehat{\Sigma}\Sigma^{-1/2} - \mathbf{I}_d\|_F = O(\epsilon\log(1/\epsilon))$ with high probability.

Robust Stochastic Convex Optimization Consider the goal of optimizing $\overline{f}(\theta) := \mathbf{E}_{z \sim D}[f(\theta;z)]$, given ϵ -corrupted samples of an unknown distribution $D \in \mathcal{D}$. A standard approach for solving the uncorrupted version of this problem is to use gradient descent, where we update $\theta_{t+1} = \theta_t - \nabla \overline{f}(\theta)$, and approximate $\nabla \overline{f}(\theta)$ using samples from D. Since we only have access to a corrupted version of D, we can use a robust mean estimation algorithm to robustly estimate $\nabla \overline{f}(\theta) = \mathbf{E}_{z \sim D}[\nabla f(\theta;z)]$ (Diakonikolas et al., 2019a; Prasad et al., 2020).

Gradient descent with an approximate gradient oracle is well-understood, see, e.g., Theorem E.5 of (Prasad et al., 2020). By using our streaming robust mean estimator as that oracle, we obtain a general streaming algorithm for robust stochastic optimization (see Corollary E.6). We highlight below a concrete application to logistic regression.

Theorem 5.2 (Streaming Robust Logistic Regression). *Consider the following model: Let* $(X,Y) \sim D$, *where* $X \sim \mathcal{N}(0,\mathbf{I}_d)$, $Y \mid X \sim Bern(p)$, for $p = 1/(1 + e^{-X^T\theta^*})$, and

 $\|\theta^*\|_2 = O(1)$. Let P be an ϵ -contamination of D in total variation. There is a single-pass streaming algorithm that uses $n = (d^2/\epsilon)$ polylog (d/ϵ) samples of P, runs in time nd polylog (d/ϵ) , uses memory d polylog (d/ϵ) , and returns $\widehat{\theta} \in \mathbb{R}^d$ such $\|\widehat{\theta} - \theta^*\|_2 = O(\sqrt{\epsilon})$ with high probability.

Distributed Non-convex Optimization Finally, we study the task of finding a second-order stationary point of a non-convex objective, i.e., a stationary point $\widehat{\theta}$ such that the Hessian at $\widehat{\theta}$ is not too negative in any direction. We consider a distributed setup where a master machine collects gradient estimates from worker machines, an ϵ -fraction of which are dishonest. Note that a vanilla robust gradient descent method may get stuck at a stationary point with a large negative Hessian direction. (Yin et al., 2019) showed that it is possible to escape such points by carefully adding spherical noise to a robust estimate of the gradient. Plugging our streaming robust mean estimator into their framework yields a memory-efficient streaming algorithm with quantitatively similar guarantees (see Appendix E.3 for details).

6. Discussion

In this work, we gave the first efficient streaming algorithm with near-optimal space complexity for outlier-robust high-dimensional mean estimation. As an application, we also obtained low-space streaming algorithms for a range of other robust estimation tasks. Our work is a first step towards understanding the space complexity of high-dimensional robust statistics in the streaming setting.

Our work suggests a number of open problems. First, the sample complexity of our mean estimation algorithm is $\tilde{O}(d^2/\epsilon^2)$, while the information-theoretic optimum (without space constraints!) is $\tilde{O}(d/\epsilon^2)$. What is the *optimal* sample-space tradeoff? A similar question can be asked for the broader tasks of covariance estimation and stochastic optimization. A more general goal is to characterize the tradeoff between space complexity, number of passes, and sample size/runtime for other robust high-dimensional statistics tasks, e.g., clustering and learning of mixture models.

Finally, another research direction concerns the considered contamination model. Throughout this paper, we focused on the TV-contamination model. One can consider an even stronger contamination model with an adaptive adversary, where the outliers can be completely arbitrary (i.e., not follow any distribution), and the adversary can additionally control the order in which the points are presented in the stream. Is it possible to obtain $\tilde{O}_{\epsilon}(d)$ -space single-pass streaming algorithms for robust mean estimation in the presence of such an adversary? While our algorithms can be shown to work in this model with a poly-logarithmic number of passes, it is not clear whether a single-pass algorithm with sub-quadratic space complexity exists in this setting.

References

- Achlioptas, D. Database-friendly random projections: Johnson-lindenstrauss with binary coins. *Journal of computer and System Sciences*, 66(4):671–687, 2003.
- Anthony, M. and Bartlett, P. L. *Neural Network Learning: Theoretical Foundations*. Cambridge University Press, first edition, November 1999.
- Bakshi, A., Diakonikolas, I., Hopkins, S. B., Kane, D., Karmalkar, S., and Kothari, P. K. Outlier-robust clustering of gaussians and other non-spherical mixtures. In 61st IEEE Annual Symposium on Foundations of Computer Science, FOCS 2020, pp. 149–159. IEEE, 2020a.
- Bakshi, A., Diakonikolas, I., Jia, H., Kane, D. M., Kothari, P. K., and Vempala, S. S. Robustly learning mixtures of k arbitrary gaussians. *CoRR*, abs/2012.02119, 2020b. URL https://arxiv.org/abs/2012.02119.
- Balakrishnan, S., Du, S. S., Li, J., and Singh, A. Computationally efficient robust sparse estimation in high dimensions. In *Proceedings of the 30th Conference on Learning Theory, COLT 2017*, pp. 169–212, 2017.
- Banks, J., Garza-Vargas, J., Kulkarni, A., and Srivastava, N. Pseudospectral shattering, the sign function, and diagonalization in nearly matrix multiplication time. In *FOCS* 2020, pp. 529–540, 2020.
- Barreno, M., Nelson, B., Joseph, A. D., and Tygar, J. D. The security of machine learning. *Machine Learning*, 81 (2):121–148, 2010.
- Biggio, B., Nelson, B., and Laskov, P. Poisoning attacks against support vector machines. In *Proceedings of the 29th International Conference on Machine Learning, ICML 2012*, 2012.
- Birgé, L. An alternative point of view on lepski's method. *Lecture Notes-Monograph Series*, pp. 113–133, 2001.
- Blum, A., Hopcroft, J., and Kannan, R. *Foundations of Data Science*. Cambridge University Press, first edition, January 2020.
- Bottou, L. Large-scale machine learning with stochastic gradient descent. In *COMPSTAT*, 2010.
- Bottou, L., Curtis, F. E., and Nocedal, J. Optimization methods for large-scale machine learning. *SIAM Review*, 60(2):223–311, 2018.
- Cheng, Y., Diakonikolas, I., Kane, D., and Stewart, A. Robust learning of fixed-structure bayesian networks. In Advances in Neural Information Processing Systems 31: Annual Conference on Neural Information Processing Systems 2018, NeurIPS 2018, pp. 10304–10316, 2018.
 Full version available at https://arxiv.org/abs/1606.07384.

- Cheng, Y., Diakonikolas, I., Ge, R., and Woodruff, D. P. Faster algorithms for high-dimensional robust covariance estimation. In *Conference on Learning Theory, COLT* 2019, pp. 727–757, 2019a.
- Cheng, Y., Diakonikolas, I., Ge, R., and Woodruff, D. P. Faster algorithms for high-dimensional robust covariance estimation. In *COLT*, pp. 727–757, 2019b.
- Cheng, Y., Diakonikolas, I., Kane, D. M., Ge, R., Gupta, S., and Soltanolkotabi, M. Outlier-robust sparse estimation via non-convex optimization. *CoRR*, abs/2109.11515, 2021. URL https://arxiv.org/abs/2109.11515.
- Diakonikolas, I. and Kane, D. M. Recent advances in algorithmic high-dimensional robust statistics. *CoRR*, abs/1911.05911, 2019.
- Diakonikolas, I., Kamath, G., Kane, D. M., Li, J., Moitra, A., and Stewart, A. Robust estimators in high dimensions without the computational intractability. In *FOCS*, pp. 655–664, 2016.
- Diakonikolas, I., Kamath, G., Kane, D. M., Li, J., Moitra, A., and Stewart, A. Being robust (in high dimensions) can be practical. In *ICML 2017*, pp. 999–1008, 2017a.
- Diakonikolas, I., Kamath, G., Kane, D. M., Li, J., Moitra, A., and Stewart, A. Being robust (in high dimensions) can be practical. *arXiv e-prints*, pp. arXiv–1703, 2017b.
- Diakonikolas, I., Kane, D. M., and Stewart, A. List-decodable robust mean estimation and learning mixtures of spherical gaussians. In *Proceedings of the 50th Annual ACM SIGACT Symposium on Theory of Computing, STOC 2018*, pp. 1047–1060, 2018. Full version available at https://arxiv.org/abs/1711.07211.
- Diakonikolas, I., Kamath, G., Kane, D., Li, J., Steinhardt, J., and Stewart, A. Sever: A robust meta-algorithm for stochastic optimization. In *ICML* 2019, pp. 1596–1606, 2019a.
- Diakonikolas, I., Karmalkar, S., Kane, D., Price, E., and Stewart, A. Outlier-robust high-dimensional sparse estimation via iterative filtering. In Advances in Neural Information Processing Systems 32: Annual Conference on Neural Information Processing Systems 2018, NeurIPS 2019, 2019b.
- Diakonikolas, I., Kong, W., and Stewart, A. Efficient algorithms and lower bounds for robust linear regression. In *Proceedings of the Thirtieth Annual ACM-SIAM Symposium on Discrete Algorithms*, SODA 2019, pp. 2745–2754, 2019c.

- Diakonikolas, I., Kane, D. M., Kongsgaard, D., J., and Tian, K. List-decodable mean estimation in nearly-pca time. *CoRR*, abs/2011.09973, 2020a. URL https://arxiv.org/abs/2011.09973.
- Diakonikolas, I., Kane, D. M., and Pensia, A. Outlier robust mean estimation with subgaussian rates via stability. In Advances in Neural Information Processing Systems 33: Annual Conference on Neural Information Processing Systems 2020, NeurIPS 2020, 2020b.
- Diakonikolas, I., Kontonis, V., Tzamos, C., and Zarifis, N. Non-convex SGD learns halfspaces with adversarial label noise. In *NeurIPS* 2020, 2020c.
- Diakonikolas, I., Kamath, G., Kane, D. M., Li, J., Moitra, A., and Stewart, A. Robustness meets algorithms. *Commun. ACM*, 64(5):107–115, 2021a.
- Diakonikolas, I., Kane, D. M., Kongsgaard, D., Li, J., and Tian, K. Clustering mixture models in almost-linear time via list-decodable mean estimation. *CoRR*, abs/2106.08537, 2021b.
- Diakonikolas, I., Kane, D. M., Stewart, A., and Sun, Y. Outlier-robust learning of ising models under dobrushin's condition. In *Conference on Learning Theory, COLT 2021*, volume 134 of *Proceedings of Machine Learning Research*, pp. 1645–1682. PMLR, 2021c.
- Dong, Y., Hopkins, S. B., and Li, J. Quantum entropy scoring for fast robust mean estimation and improved outlier detection. *NeurIPS*, 32:6067–6077, 2019.
- Goldberg, P. W. and Jerrum, M. R. Bounding the vapnik-chervonenkis dimension of concept classes parameterized by real numbers. *Machine Learning*, 18(2):131–148, 1995.
- Hopkins, S. B. and Li, J. Mixture models, robustness, and sum of squares proofs. In *Proceedings of the 50th Annual ACM SIGACT Symposium on Theory of Computing*, *STOC 2018*, pp. 1021–1034, 2018.
- Huber, P. J. Robust estimation of a location parameter. *Ann. Math. Statist.*, 35(1):73–101, 03 1964.
- Klivans, A. R., Kothari, P. K., and Meka, R. Efficient algorithms for outlier-robust regression. In *Conference On Learning Theory, COLT 2018*, pp. 1420–1430, 2018.
- Kothari, P. K., Steinhardt, J., and Steurer, D. Robust moment estimation and improved clustering via sum of squares. In *Proceedings of the 50th Annual ACM SIGACT Symposium on Theory of Computing, STOC 2018*, pp. 1035–1046, 2018.
- Lai, K. A., Rao, A. B., and Vempala, S. Agnostic estimation of mean and covariance. In *FOCS*, 2016.

- Lepskii, O. V. On a problem of adaptive estimation in gaussian white noise. *Theory of Probability & Its Applications*, 35(3):454–466, 1991.
- Li, J., Absher, D., Tang, H., Southwick, A., Casto, A., Ramachandran, S., Cann, H., Barsh, G., Feldman, M., Cavalli-Sforza, L., and Myers, R. Worldwide human relationships inferred from genome-wide patterns of variation. *Science*, 319:1100–1104, 2008.
- Liu, A. and Moitra, A. Settling the robust learnability of mixtures of gaussians. *CoRR*, abs/2011.03622, 2020.
- Paschou, P., Lewis, J., Javed, A., and Drineas, P. Ancestry informative markers for fine-scale individual assignment to worldwide populations. *Journal of Medical Genetics*, 47:835–847, 2010.
- Pensia, A., Jog, V., and Loh, P. Robust regression with covariate filtering: Heavy tails and adversarial contamination. *arXiv* preprint arXiv:2009.12976, 2020.
- Pesme, S. and Flammarion, N. Online robust regression via SGD on the 11 loss. In Advances in Neural Information Processing Systems 33: Annual Conference on Neural Information Processing Systems 2020, NeurIPS 2020, 2020.
- Prasad, A., Suggala, A. S., Balakrishnan, S., and Ravikumar, P. Robust estimation via robust gradient estimation. *Journal of the Royal Statistical Society: Series B (Statistical Methodology)*, 82(3):601–627, 2020.
- Rosenberg, N., Pritchard, J., Weber, J., Cann, H., Kidd, K., Zhivotovsky, L., and Feldman, M. Genetic structure of human populations. *Science*, 298:2381–2385, 2002.
- Shah, V., Wu, X., and Sanghavi, S. Choosing the sample with lowest loss makes SGD robust. In *AISTATS* 2020, volume 108 of *Proceedings of Machine Learning Research*, pp. 2120–2130, 2020.
- Sharan, V., Sidford, A., and Valiant, G. Memory-sample tradeoffs for linear regression with small error. In *Proceedings of the 51st Annual ACM SIGACT Symposium on Theory of Computing*, STOC 2019, pp. 890–901. ACM, 2019. doi: 10.1145/3313276.3316403. URL https://doi.org/10.1145/3313276.3316403.
- Steinhardt, J., Koh, P. W., and Liang, P. S. Certified defenses for data poisoning attacks. In *NeurIPS*, pp. 3520–3532, 2017.
- Steinhardt, J., Charikar, M., and Valiant, G. Resilience: A criterion for learning in the presence of arbitrary outliers. In *ITCS* 2018, pp. 45:1–45:21, 2018.
- Tran, B., Li, J., and Madry, A. Spectral signatures in backdoor attacks. In *NeurIPS 2018*, pp. 8011–8021, 2018.

- Tsai, C., Prasad, A., Balakrishnan, S., and Ravikumar, P. Heavy-tailed streaming statistical estimation. *CoRR*, abs/2108.11483, 2021.
- Tukey, J. W. A survey of sampling from contaminated distributions. *Contributions to probability and statistics*, 2:448–485, 1960.
- Vershynin, R. Introduction to the non-asymptotic analysis of random matrices. *arXiv preprint arXiv:1011.3027*, 2010.
- Yin, D., Chen, Y., Ramchandran, K., and Bartlett, P. Defending against saddle point attack in byzantine-robust distributed learning. In *ICML*, pp. 7074–7084, 2019.

A. Introduction

This work studies high-dimensional learning in the presence of a constant fraction of arbitrary outliers. Outlier-robust learning in high dimensions is motivated by pressing machine learning (ML) applications, including ML security (Barreno et al., 2010; Biggio et al., 2012; Steinhardt et al., 2017; Tran et al., 2018; Diakonikolas et al., 2019a) and exploratory analysis of datasets with natural outliers (Rosenberg et al., 2002; Paschou et al., 2010; Li et al., 2008). This field has its roots in robust statistics, a branch of statistics initiated in the 60s with the pioneering works of Tukey and Huber (Tukey, 1960; Huber, 1964). Early work developed minimax optimal estimators for various robust estimation tasks, albeit with runtimes exponential in the dimension. A recent line of work in computer science, starting with (Diakonikolas et al., 2016; Lai et al., 2016), developed polynomial time robust estimators for a range of high-dimensional statistical tasks. Algorithmic high-dimensional robust statistics is by now a relatively mature field, see, e.g., (Diakonikolas & Kane, 2019; Diakonikolas et al., 2021a) for surveys.

This recent progress notwithstanding, even for the basic task of mean estimation, previous robust estimators require the entire dataset in main memory. This space requirement can be a major bottleneck in large-scale applications, where an algorithm has access to a very large stream of data. Indeed, practical machine learning methods are typically simple iterative algorithms that make a single pass over the data and require a small amount of storage — with stochastic gradient descent being the prototypical example (Bottou, 2010; Bottou et al., 2018). Concretely, in prior applications of robust statistics in data analysis (Diakonikolas et al., 2017a) and data poisoning defenses (Diakonikolas et al., 2019a), the storage requirements of the underlying algorithms were observed to significantly hinder scalability³. This discussion motivates the following natural question:

Can we develop efficient robust estimators in the streaming model with (near-) optimal space complexity?

We emphasize that this broad question is meaningful and interesting even ignoring computational considerations. While any method requires space complexity $\Omega(d)$, where d is the dimension of the problem (to store a single sample), it is not obvious that a matching upper bound exists. We note that it is relatively simple to design O(d)-memory streaming algorithms with sample complexity exponential in d. But it is by no means clear whether there exists an estimator with near-linear space requirements and $\operatorname{poly}(d)$ sample complexity (independent of its runtime).

A.1. Our Results

In this work, we initiate a systematic investigation of high-dimensional robust statistics in the streaming model. We start by focusing on the most basic task — that of robust mean estimation. Our main result is the first space-efficient streaming algorithm for robust mean estimation under natural distributional assumptions. Our computationally efficient algorithm makes a single pass over the data, uses near-optimal space, and matches the error guarantees of previous polynomial-time algorithms for the problem.

Given this result, we leverage the fact that several robust statistics tasks can be reduced to robust mean estimation to obtain near-optimal space, single-pass streaming algorithms for more complex statistical tasks.

To formally state our contributions, we require some basic definitions. We start with the standard streaming model.

Definition A.1 (Single-Pass Streaming Model). Let S be a fixed set. In the *one-pass streaming model*, the elements of S are revealed one at a time to the algorithm, and the algorithm is allowed a single pass over these points.

Our robust estimators work in the following contamination model, where the adversary can corrupt the true distribution in total variation distance (for distributions P and Q, we use $d_{TV}(P,Q)$ to denote their total variation distance).

Definition A.2 (TV-contamination). Given a parameter $\epsilon < 1/2$ and a distribution class \mathcal{D} , the adversary specifies a distribution D' such that there exists $D \in \mathcal{D}$ with $d_{\mathrm{TV}}(D, D') \leq \epsilon$. Then the algorithm draws i.i.d. samples from D'. We say that the distribution D' is an ϵ -corrupted version of the distribution D in total variation distance.

The distribution D' in Definition A.2 can be adversarially selected (and can even depend on our learning algorithm). Since Huber's contamination model (Huber, 1964) only allows additive errors, TV-contamination is a stronger model.

³See, for example, Section 5.1 of (Diakonikolas et al., 2017b).

STREAMING ALGORITHM FOR ROBUST MEAN ESTIMATION

The main result of this paper is the following (see Theorem D.2 for a more general statement):

Theorem A.3 (Streaming Robust Mean Estimation). Let \mathcal{D} be a distribution family on \mathbb{R}^d and $0 < \epsilon < \epsilon_0$ for a sufficiently small constant $\epsilon_0 > 0$. Let P be an ϵ -corrupted version of D in total variation distance for some $D \in \mathcal{D}$ with unknown mean μ_D . There is a single-pass streaming algorithm that, given ϵ and \mathcal{D} , reads a stream of n i.i.d. samples from P, runs in sample near-linear time, uses memory d polylog (d/ϵ) , and outputs an estimate $\widehat{\mu}$ that, with probability at least 9/10, satisfies the following:

- 1. If \mathcal{D} is the family of distributions with identity-bounded covariance, then $n = \tilde{O}\left(d^2/\epsilon\right)$ and $\|\widehat{\mu} \mu_D\|_2 = O(\sqrt{\epsilon})$.
- 2. If \mathcal{D} is the family of identity-covariance subgaussian distributions, then $n = \tilde{O}\left(d^2/\epsilon^2\right)$ and $\|\hat{\mu} \mu_D\|_2 = O(\epsilon \sqrt{\log(1/\epsilon)})$.

We note that the above error guarantees are information-theoretically optimal, even in absence of resource constraints. While prior work had obtained efficient robust mean estimators matching these error guarantees (Diakonikolas et al., 2016; 2017a; Steinhardt et al., 2018), all previous algorithms with dimension-independent error incurred space complexity $\Omega(d^2)$.

BEYOND ROBUST MEAN ESTIMATION

Using the algorithm of Theorem A.3 as a black-box, we obtain the first efficient single-pass streaming algorithms with near-optimal space complexity for a range of more complex statistical tasks. These contributions are presented in detail in Appendix E. Here we highlight some of these results.

Our first application is a streaming algorithm for robust covariance estimation.

Theorem A.4 (Robust Gaussian Covariance Estimation). Let Q be a distribution on \mathbb{R}^d with $d_{\mathrm{TV}}(Q, \mathcal{N}(0, \Sigma)) \leq \epsilon$ and assume $\frac{1}{\kappa}\mathbf{I}_d \leq \Sigma \leq \mathbf{I}_d$. There is a single-pass streaming algorithm that uses $n = (d^4/\epsilon^2)\mathrm{polylog}(d, \kappa, 1/\epsilon)$ samples from Q, runs in time $nd^2\mathrm{polylog}(d, \kappa, 1/\epsilon)$, uses memory $d^2\mathrm{polylog}(d, \kappa, 1/\epsilon)$, and outputs a matrix $\widehat{\Sigma}$ such that $\|\Sigma^{-1/2}\widehat{\Sigma}\Sigma^{-1/2} - \mathbf{I}_d\|_F = O(\epsilon \log(1/\epsilon))$ with probability at least 9/10.

See Theorem E.3 for a more detailed statement.

Our second application is for the general problem of robust stochastic optimization. Here we state two concrete results for robust linear and logistic regression (see Theorem E.9 and Theorem E.12 for more detailed statements). Both of these statements are special cases of a streaming algorithm for robust stochastic convex optimization (see Corollary E.6).

Theorem A.5 (Streaming Robust Linear Regression). Let D be the distribution of (X,Y) defined by $Y = X^T \theta^* + Z$, where $X \sim \mathcal{N}(0,\mathbf{I}_d)$, $Z \sim \mathcal{N}(0,1)$ independent of X, and $\|\theta^*\|_2 \leq r$. Let P be an ϵ -corruption of D in total variation distance. There is a single-pass streaming algorithm that uses $n = (d^2/\epsilon)$ polylog $(d(1+r)/\epsilon)$ samples from P, runs in time nd polylog $(d(1+r)/\epsilon)$, uses memory d polylog (dr/ϵ) , and outputs an estimate $\hat{\theta} \in \mathbb{R}^d$ such $\|\hat{\theta} - \theta^*\|_2 = O(\sqrt{\epsilon})$ with high probability.

Theorem A.6 (Streaming Robust Logistic Regression). Consider the following model: Let $(X,Y) \sim D$, where $X \sim \mathcal{N}(0,\mathbf{I}_d)$, $Y \mid X \sim Bern(p)$, for $p = 1/(1+e^{-X^T\theta^*})$, and $\|\theta^*\|_2 = O(1)$. Let P be an ϵ -corruption of D in total variation distance. There is a single-pass streaming algorithm that uses $n = (d^2/\epsilon)$ polylog (d/ϵ) samples from P, runs in time nd polylog (d/ϵ) , uses memory d polylog (d/ϵ) , and outputs an estimate $\widehat{\theta} \in \mathbb{R}^d$ such $\|\widehat{\theta} - \theta^*\|_2 = O(\sqrt{\epsilon})$ with high probability.

Finally, in Appendix E, we include an additional application to distributed non-convex optimization in the streaming setting. *Remark* A.7 (Bit complexity). For simplicity of presentation, in the main body of the paper, we consider the model of computation where the algorithms can store and manipulate real numbers exactly. We show in Appendix L that our algorithms can tolerate errors due to finite precision. In particular, all our algorithms (including Algorithm 5) can be implemented in the word RAM model with $d \operatorname{polylog}(d/\epsilon)$ bits.

A.2. Overview of Techniques

In this section, we provide a brief overview of our approach to establish Theorem A.3. We start by recalling how robust mean estimation algorithms typically work without space constraints. A standard tool in the literature is the filtering technique of

(Diakonikolas et al., 2016; 2017a; Diakonikolas & Kane, 2019). The idea of the filtering method is the following: Given a set S of corrupted samples, by analyzing spectral properties of the covariance of S, we can either certify that the sample mean of S is close to the true mean of the distribution, or can construct a filter. The filter is a method for selecting some elements of S to remove, with the guarantee that it will remove more outliers than inliers. If we can efficiently construct a filter, our algorithm can then remove the selected samples from S, obtaining a cleaner dataset and repeat the process. Eventually, this procedure must terminate, giving an accurate estimate of the true mean.

We proceed to explain how to implement the filtering method in a streaming model. We start with the easier case where the dataset is stored in read-only-memory, or more generally in a *multi-pass* streaming setting. At each round of the algorithm, one has a subset S' of the original dataset S that needs to be maintained (in particular, the set of samples that has survived the filters applied thus far). To do this naïvely would require n = |S| many bits of memory, which is too much for us. A more inventive strategy would be the following: instead of storing these subsets S' explicitly, store them implicitly by instead storing enough information to reconstruct the filters used to obtain S'. This seems like a productive idea, as most filters are relatively simple. For example, a commonly used filter is to remove all points $x \in S$ for which $v^T x > t$, for some vector v and scalar t. One could store enough information to apply this filter by merely storing (v,t), which would take O(d) bits of information. Unfortunately, most filtering algorithms may require $\Omega(d)$ many iterations before attaining their final answer. Consequently, the sets S' one needs to store are not just the result of applying a single filter, but instead the result of iteratively applying $\Omega(d)$ of them. In order to store all of these extra filters, one would need $\Omega(d^2)$ bits. (For the sake of this intuitive description, we focused on "hard-thresholding" filters. Our algorithm will actually use a soft-thresholding filter, assigning weights to each point.)

To circumvent this first obstacle, one requires as a starting point a filtering algorithm that is guaranteed to terminate after a small (namely, at most poly-logarithmic) number of iterations. Recent work (Dong et al., 2019; Diakonikolas et al., 2021b) has obtained such algorithms. Here we generalize and simplify the filtering method of (Diakonikolas et al., 2021b). This allows us to obtain an algorithm with space complexity $d \operatorname{polylog}(d/\epsilon)$ that works in the *multi-pass streaming model*, where $\operatorname{polylog}(d/\epsilon)$ passes over the same dataset are allowed.

To obtain a *single-pass* streaming algorithm, new ideas are required. In the single-pass setting, we cannot implicitly store a subset of the full dataset S; once we access some points from S, we will never be able to see them again. To deal with this issue, we will need to slightly alter our way of thinking about the algorithm. Instead of being given a set S of samples, an ϵ -fraction of which have been corrupted, we instead adopt the view of having sample access to a distribution P, which is ϵ -close in total variation distance to the inlier distribution S. Given this point of view, instead of a filter defining a procedure for removing samples from S and outputting a subset S', we think of it as a rejection sampling procedure that replaces S with a cleaner distribution S.

This shift in perspective comes with new technical challenges. In particular, when constructing the next round of filters, we will need to compute quantities pertaining to the current distribution P of the data points. In the setting of the multiple-pass model, this imposed no problem; these quantities could be calculated exactly. This is no longer possible when we merely have sample access to P. The best one can hope for is to approximate these quantities to sufficient precision for the rest of our analysis to carry over. However, the natural estimators for some required quantities (e.g., powers of the covariance matrix) would need to access the data multiple times. Circumventing this issue requires non-trivial technical work. Roughly speaking, instead of iterating over the same dataset to approximate the desired quantities, we show that it suffices to iterate over statistically identical datasets.

A.3. Prior and Related Work

Since the dissemination of (Diakonikolas et al., 2016; Lai et al., 2016), there has been an explosion of research in algorithmic aspects of robust statistics. We now have efficient robust estimators for a range of more complex problems, including covariance estimation (Diakonikolas et al., 2016; Cheng et al., 2019a), sparse estimation tasks (Balakrishnan et al., 2017; Diakonikolas et al., 2019b; Cheng et al., 2021), learning graphical models (Cheng et al., 2018; Diakonikolas et al., 2021c), linear regression (Klivans et al., 2018; Diakonikolas et al., 2019c; Pensia et al., 2020), stochastic optimization (Prasad et al., 2020; Diakonikolas et al., 2019a), and robust clustering/learning various mixture models (Hopkins & Li, 2018; Kothari et al., 2018; Diakonikolas et al., 2018; 2020a; 2021b; Bakshi et al., 2020a; Liu & Moitra, 2020; Bakshi et al., 2020b). The reader is referred to (Diakonikolas & Kane, 2019) for a detailed overview. We reiterate that all previously developed algorithms work in the batch setting, i.e., require the entire dataset in memory.

For the problem of robust mean estimation, (Dong et al., 2019; Diakonikolas et al., 2021b) gave filtering-based algorithms

with a poly-logarithmic number of iterations. The former algorithm relies on the matrix multiplicative weights framework, while the latter is based on first principles. Our starting point in Appendix C can be viewed as a generalization and further simplification of the ideas in (Diakonikolas et al., 2021b). Specifically, our algorithm works under the stability condition (Definition B.8), which broadly generalizes the bounded covariance assumption used in (Diakonikolas et al., 2021b).

In the context of robust supervised learning (including, e.g., our robust linear regression application), low-space streaming algorithms are known in weaker contamination models that only allow *label* corruptions, see, e.g., (Pesme & Flammarion, 2020; Shah et al., 2020; Diakonikolas et al., 2020c). We emphasize that the contamination model of Definition A.2 is significantly more challenging, and no low-space streaming algorithms were previously known in this model.

Finally, we note that recent work (Tsai et al., 2021) studies streaming algorithms for heavy-tailed stochastic optimization. While the goal of developing low-space streaming algorithms is qualitatively similar to the goal of our work, the algorithmic results in (Tsai et al., 2021) have no implications in the corrupted setting studied in this work.

A.4. Organization

The structure of this paper is as follows: In Appendix B, we record the notation and technical background that will be used throughout the paper. In Appendix C, we design a filter-based algorithm for robust mean estimation under the stability condition with a poly-logarithmic number of iterations. In Appendix D, we build on the algorithm from Appendix C, to obtain our single-pass streaming algorithm for robust mean estimation under the stability condition. Finally, in Appendix E, we obtain our streaming algorithms for more complex robust estimation tasks. To facilitate the flow of the presentation, some proofs of intermediate lemmas are deferred to Appendices G to L.

B. Preliminaries

B.1. Notation and Basic Facts

Basic Notation We use \mathbb{Z}_+ to denote the set of positive integers. For $n \in \mathbb{Z}_+$, we denote $[n] := \{1, \dots, n\}$ and use \mathcal{S}^{d-1} for the d-dimensional unit sphere. For a vector v, we let $\|v\|_2$ denote its ℓ_2 -norm. We use boldface letters for matrices. We use \mathbf{I}_d to denote the $d \times d$ identity matrix. For a matrix \mathbf{A} , we use $\|\mathbf{A}\|_F$ and $\|\mathbf{A}\|_2$ to denote the Frobenius and spectral norms respectively. For $\mathbf{A} \in \mathbb{R}^{m \times n}$, we use \mathbf{A}^b to denote the nm-dimensional vector obtained by concatenating the rows of \mathbf{A} . We say that a symmetric $d \times d$ matrix \mathbf{A} is PSD (positive semidefinite), and write $\mathbf{A} \succeq 0$, if for all vectors $x \in \mathbb{R}^d$ we have that $x^T \mathbf{A} x \ge 0$. We denote $\lambda_{\max}(\mathbf{A}) := \max_{u \in \mathcal{S}^{d-1}} x^T \mathbf{A} x$. We write $\mathbf{A} \preceq \mathbf{B}$ when $\mathbf{B} - \mathbf{A}$ is PSD. For a matrix $\mathbf{A} \in \mathbb{R}^{d \times d}$, $\operatorname{tr}(\mathbf{A})$ denotes the trace of the matrix \mathbf{A} . We use \otimes to denote the Kronecker product. For the sake of conciseness, we sometimes use $x = a \pm b$ as a shorthand for $a - b \le x \le a + b$. We use $a \le b$, to denote that there exists an absolute universal constant C > 0 (independent of the variables or parameters on which a and b depend) such that $a \le Cb$. Similarly, we use the notation $a \ge b$ to denote that $b \le a$. We use c, c', C, C' to denote absolute constants that may change from line to line, whereas we use constants C_1, C_2, C_3, \ldots to denote fixed absolute constants that are important for our algorithms. We use $\widetilde{O}(\cdot)$ to ignore poly-logarithmic factors in all variables appearing inside the parentheses. For the sake of simplicity, we sometimes omit rounding non-integer quantities to integer ones. For example, we treat logarithmic factors as integers when they appear in the sample complexity or number of iterations of an algorithm. We use poly(·) to indicate a quantity that is polynomial in its arguments. Similarly, polylog(·) denotes a quantity that is polynomial in the logarithm of its arguments.

Probability Notation For a random variable X, we use $\mathbf{E}[X]$ for its expectation. For a set S, we use $\mathcal{U}(S)$ to denote the uniform distribution on S. We use $\mathcal{N}(\mu, \Sigma)$ to denote the Gaussian distribution with mean μ and covariance matrix Σ . For a distribution D on \mathbb{R}^d , we denote $\mu_D = \mathbf{E}_{X \sim D}[X]$ and $\Sigma_D = \mathbf{E}_{X \sim D}[(X - \mu_D)(X - \mu_D)^T]$. Moreover, given a weight function $w : \mathbb{R}^d \to [0,1]$, we define the re-weighted distribution D_w to be $D_w(x) := D(x)w(x)/\int_{\mathbb{R}^d} w(x)D(x)\mathrm{d}x$. We use $\mu_{w,D} = \mathbf{E}_{X \sim D_w}[X]$ for its mean and $\overline{\Sigma}_{w,D}^\mu = \mathbf{E}_{X \sim D_w}[(X - \mu)(X - \mu)^T]$ for the second moment that is centered with respect to μ (we will often drop μ from the notation when it is clear from the context). We use $\mathbb{1}\{x \in E\}$ to denote the indicator function of the set E.

Basic Facts We will use the following two basic facts.

Fact B.1. Let $x \in \mathbb{R}^d$ and $p \ge 1$. Then $||x||_{p+1} \le ||x||_p \le ||x||_{p+1} d^{\frac{1}{p(p+1)}}$.

Fact B.2. If A, B, C are symmetric $d \times d$ matrices satisfying $A \succeq 0$ and $B \preceq C$, we have that $tr(AB) \leq tr(AC)$.

Proof. Since **A** is PSD, we can consider its spectral decomposition $\mathbf{A} = \sum_{i=1}^{d} \lambda_i v_i v_i^T$, where $\lambda_i \geq 0$. Using the linearity of trace operator, we have that

$$\operatorname{tr}(\mathbf{A}\mathbf{B}) = \sum_{i=1}^{d} \lambda_{i} \operatorname{tr}(v_{i} v_{i}^{T} \mathbf{B}) = \sum_{i=1}^{d} \lambda_{i} \operatorname{tr}(v_{i}^{T} \mathbf{B} v_{i}) \leq \sum_{i=1}^{d} \lambda_{i} \operatorname{tr}(v_{i}^{T} \mathbf{C} v_{i}) = \sum_{i=1}^{d} \lambda_{i} \operatorname{tr}(v_{i} v_{i}^{T} \mathbf{C}) = \operatorname{tr}(\mathbf{A}\mathbf{C}) ,$$

where the inequality uses that $\mathbf{B} \leq \mathbf{C}$ and $\lambda_i \geq 0$.

We will use the notion of total variation distance, defined below.

Definition B.3. Let P,Q be two probability distributions on \mathbb{R}^d . The total variation distance between P and Q, denoted by $d_{\mathrm{TV}}(P,Q)$, is defined as $d_{\mathrm{TV}}(P,Q) = \sup_{A \subseteq \mathbb{R}^d} |P(A) - Q(A)|$. For continuous distributions P,Q with densities p,q, we have that $d_{\mathrm{TV}}(P,Q) = \frac{1}{2} \int_{\mathbb{R}^d} |p(x) - q(x)| \mathrm{d}x$.

Whenever $d_{\text{TV}}(P,Q) = \epsilon$, it is sometimes helpful to consider the decomposition below.

Fact B.4. Let a domain \mathcal{X} . For any $\epsilon \in [0,1]$ and for any two distributions D_1, D_2 on \mathcal{X} with $d_{\mathrm{TV}}(D_1, D_2) = \epsilon$, there exist distributions D, Q_1, Q_2 such that $D_1 = (1 - \epsilon)D + \epsilon Q_1$ and $D_2 = (1 - \epsilon)D + \epsilon Q_2$.

This decomposition can be achieved by the following choice of Q_1 , Q_2 , and D:

$$Q_1(x) = \begin{cases} \frac{D_1(x) - D_2(x)}{\epsilon} \;, & \text{if } D_1(x) > D_2(x) \\ 0 \;, & \text{otherwise} \end{cases}, \quad Q_2(x) = \begin{cases} \frac{D_2(x) - D_1(x)}{\epsilon} \;, & \text{if } D_2(x) > D_1(x) \\ 0 \;, & \text{otherwise} \end{cases},$$

and $D(x) = \min\{D_1(x), D_2(x)\}/(1 - \epsilon)$. In light of Fact B.4, the adversary that performs corruption in total variation distance can be thought of as "both additive and subtractive" adversary.

Concentration Inequalities We will also require following standard results regarding concentration of random variables: **Fact B.5** ((Vershynin, 2010)). Consider a distribution D on \mathbb{R}^d that has zero mean and is supported in an ℓ_2 -ball of radius R from the origin. Denote by Σ its covariance matrix and denote by $\Sigma_N = (1/n) \sum_{i=1}^N X_i X_i^T$ the empirical covariance matrix using N samples $X_i \sim D$. There is a constant C such that for any $0 < \epsilon' < 1$ and $0 < \tau < 1$, if $N > C\epsilon'^{-2} \|\Sigma\|_2^{-1} R^2 \log(d/\tau)$, we have that $\|\Sigma - \Sigma_N\|_2 \le \epsilon' \|\Sigma\|_2$, with probability at least $1 - \tau$.

Fact B.6 (Gaussian Norm Concentration). For every $\beta > 0$, $\Pr_{X \sim \mathcal{N}(0, \mathbf{I}_d)}[|||X||^2 - d| > \beta] \le 2e^{-c\beta^2/d}$, where c > 0 is a universal constant.

Fact B.7 ((Achlioptas, 2003)). Let $0 < \gamma < 1$ and $u_1, \ldots, u_N \in \mathbb{R}^d$. Let z_j for $j \in [L]$ drawn from the uniform distribution on $\{\pm 1\}^d$. There exists a constant C > 0 such that, if $L > C \log(N/\gamma)$, then, with probability at least $1 - \gamma$, we have that $0.8||u_i||^2 \le \frac{1}{L} \sum_{j=1}^L (z_j^T u_i)^2 \le 1.2||u_i||^2$ for all $i \in [N]$.

B.2. Stability Condition and Its Properties

Our results will hold for every distribution satisfying the following key property (Diakonikolas & Kane, 2019).

Definition B.8 ((ϵ, δ) -stable distribution). Fix $0 < \epsilon < 1/2$ and $\delta \ge \epsilon$. A distribution G on \mathbb{R}^d is (ϵ, δ) -stable with respect to $\mu \in \mathbb{R}^d$ if for any weight function $w : \mathbb{R}^d \to [0, 1]$ with $\mathbf{E}_{X \sim G}[w(X)] \ge 1 - \epsilon$ we have that

$$\left\|\mu_{w,G} - \mu\right\|_2 \leq \delta$$
 and $\left\|\overline{\mathbf{\Sigma}}_{w,G} - \mathbf{I}_d\right\|_2 \leq \delta^2/\epsilon$.

We call a set of points S (ϵ, δ) -stable when the uniform distribution on S is stable :

Definition B.9 $((\epsilon, \delta)$ -stable set). Fix $0 < \epsilon < 1/2$ and $\delta \ge \epsilon$. A finite set $S_0 \subset \mathbb{R}^d$ is (ϵ, δ) -stable with respect to $\mu \in \mathbb{R}^d$ if the empirical distribution $\mathcal{U}(S_0)$ is (ϵ, δ) -stable with respect to μ .

We begin by stating some examples of stable distributions (see (Diakonikolas & Kane, 2019) for more details). If G is a subgaussian distribution with identity covariance, then G is (ϵ, δ) -stable with $\delta = O(\epsilon \sqrt{\log(1/\epsilon)})$. If G is a distribution with covariance at most identity, i.e., $\Sigma_G \leq \mathbf{I}_d$, then G is (ϵ, δ) -stable with $\delta = O(\sqrt{\epsilon})$. Interpolating these two results, we have that if G is a distribution with identity covariance and bounded k-th moment for $k \geq 4$, i.e., $(\mathbf{E}_{X \sim G}[|v^T(X - \mu)|^k])^{1/k} = O(1)$, then G is (ϵ, δ) -stable with $\delta = O(\epsilon^{1-1/k})$. Furthermore, it is known that $\operatorname{poly}(d/\epsilon)$ i.i.d. samples from these distributions also yields a set that contains a large stable subset (see, for example, (Diakonikolas & Kane, 2019; Diakonikolas et al., 2020b; Dong et al., 2019; Diakonikolas et al., 2016)):

Fact B.10 ((Diakonikolas & Kane, 2019)). A set of $O(d/(\epsilon^2 \log(1/\epsilon)))$ i.i.d. samples from an identity covariance subgaussian distribution is $(\epsilon, O(\epsilon \sqrt{\log(1/\epsilon)}))$ -stable with respect to μ with high probability. Similarly, a set of $\tilde{O}(d/\epsilon)$ i.i.d. samples from a distribution X with $\mathbf{Cov}[X] \leq \mathbf{I}_d$ contains a large subset S, which is $O(\epsilon, O(\sqrt{\epsilon}))$ -stable with respect to its mean $\mathbf{E}[X]$ with high probability.

The basic fact regarding stability, which is the starting point of many robust estimation algorithms, is that any slight modification of a stable distribution can not perturb the mean by a large amount, unless it significantly changes its covariance (see, for example, (Diakonikolas et al., 2016; Lai et al., 2016; Diakonikolas & Kane, 2019)). Here we require a slightly different statement than that of (Diakonikolas & Kane, 2019), and hence provide a proof in Appendix G for completeness.

Lemma B.11 (Certificate Lemma). Let G be an (ϵ, δ) -stable distribution with respect to $\mu \in \mathbb{R}^d$, for some $0 < \epsilon < 1/3$ and $\delta \geq \epsilon$. Let P be a distribution with $d_{\text{TV}}(P, G) \leq \epsilon$. Denoting by μ_P, Σ_P the mean and covariance of P, if $\lambda_{\text{max}}(\Sigma_P) \leq 1 + \lambda$, for some $\lambda \geq 0$, then $\|\mu_P - \mu\|_2 = O(\delta + \sqrt{\epsilon \lambda})$.

Given Fact B.4, we can essentially think of an ϵ -corrupted version of a stable distribution as a mixture of a stable distribution with a noise distribution, as shown below (see Appendix G for a proof).

Lemma B.12. For any $0 < \epsilon < 1/2$ and $\delta \ge \epsilon$, if a distribution G is $(2\epsilon, \delta)$ -stable with respect to $\mu \in \mathbb{R}^d$, and P is an ϵ -corrupted version of G in total variation distance, there exist distributions G_0 and B such that $P = (1 - \epsilon)G_0 + \epsilon B$ and G_0 is (ϵ, δ) -stable with respect to μ .

We continue with some technical claims related to stability that we prove in Appendix G. Let G be an (ϵ, δ) -stable distribution with respect to μ and w a weight function with $\mathbf{E}_{X \sim G}[w(X)] \geq 1 - \epsilon$. Denoting by G_w the re-weighted distribution $G_w(x) := G(x)w(x)/\int_{\mathbb{R}^d} w(x)G(x)\mathrm{d}x$, the stability of G directly implies that $1 - \delta^2/\epsilon \leq \mathbf{E}_{X \sim G_w}[(v^T(X - \mu))^2] \leq 1 + \delta^2/\epsilon$. We require a generalization of this fact for a matrix \mathbf{U} in place of v and an arbitrary vector v in place of v:

Lemma B.13. Fix $0 < \epsilon < 1/2$ and $\delta \ge \epsilon$. Let $w : \mathbb{R}^d \to [0,1]$ such that $\mathbf{E}_{X \sim G}[w(X)] \ge 1 - \epsilon$ and let G be an (ϵ, δ) -stable distribution with respect to $\mu \in \mathbb{R}^d$. For any matrix $\mathbf{U} \in \mathbb{R}^{d \times d}$ and any vector $b \in \mathbb{R}^d$, we have that

$$\underset{X \sim G_m}{\mathbf{E}} \left[\| \mathbf{U}(X-b) \|_2^2 \right] = \| \mathbf{U} \|_F^2 (1 \pm \delta^2/\epsilon) + \| \mathbf{U}(\mu-b) \|_2^2 \pm 2\delta \, \| \mathbf{U} \|_F^2 \| \mu - b \|_2 \; .$$

We use this to show Corollary B.14, which will be required when proving correctness of our algorithm. Although its exact role will become clearer later on, the corollary will be relevant to our analysis because we will filter out outliers using scores of the form $\|\mathbf{U}(x-b)\|_2^2$ for each point x.

Corollary B.14. Fix $0 < \epsilon < 1/2$ and $\delta \ge \epsilon$. Let G be an (ϵ, δ) -stable distribution with respect to $\mu \in \mathbb{R}^d$. Let a matrix $\mathbf{U} \in \mathbb{R}^{d \times d}$ and a function $w : \mathbb{R}^d \to [0, 1]$ with $\mathbf{E}_{X \sim G}[w(X)] \ge 1 - \epsilon$. For the function $\tilde{g}(x) = \|\mathbf{U}(x - b)\|_2^2$, we have that

$$(1-\epsilon)\|\mathbf{U}\|_F^2(1-\delta^2/\epsilon-2\delta\|b-\mu\|_2) \leq \underset{X \sim G}{\mathbf{E}}[w(X)\tilde{g}(X)] \leq \|\mathbf{U}\|_F^2\left(1+\delta^2/\epsilon+\|b-\mu\|_2^2+2\delta\|b-\mu\|_2\right) \; .$$

C. Filtering Algorithm with Small Number of Iterations

In this section, we develop a filtering algorithm (in the batch setting) that terminates in $polylog(d/\epsilon)$ iterations for any stable set. This leads to an algorithm that runs in near-linear time, i.e., $nd polylog(nd/\epsilon)$, generalizing the results of (Dong et al., 2019; Diakonikolas et al., 2021b). Crucially, this algorithm will form the building block of our streaming algorithm in Appendix D. We remark that the algorithm of this section works even against the *strong-contamination model* (Definition C.1 below), where the outliers may not be i.i.d. samples from any distribution, but are allowed to be completely arbitrary.

Definition C.1 (Strong Contamination Model). Given a parameter $0 < \epsilon < 1/2$ and a class of distributions \mathcal{D} , the strong adversary operates as follows: The algorithm specifies a number of samples n, then the adversary draws a set of n i.i.d. samples from some $D \in \mathcal{D}$ and after inspecting them, removes up to ϵn of them and replaces them with arbitrary points. The resulting set is given as input to the learning algorithm. We call a set ϵ -corrupted if it has been generated by the above process.

The main result of this section is the following.

Theorem C.2. Let $d \in \mathbb{Z}_+$, $0 < \tau < 1$, $0 < \epsilon < \epsilon_0$ for a sufficiently small constant ϵ_0 , and $\delta \ge \epsilon$. Let S_0 be a set of n points that is $(C\epsilon, \delta)$ -stable with respect to the (unknown) vector $\mu \in \mathbb{R}^d$, for a sufficiently large constant C > 0. Let S be an ϵ -corrupted version of S_0 in the strong contamination model. There exists an algorithm that given ϵ, δ, τ , and S, runs in time nd polylog $(d, n, 1/\epsilon, 1/\tau)$, and outputs a vector $\hat{\mu}$ such that, with probability at least $1 - \tau$, it holds $\|\mu - \hat{\mu}\|_2 = O(\delta)$.

We note that Theorem C.2 applies to any stable set. By Fact B.10, we directly obtain (i) an $O(\epsilon \sqrt{\log(1/\epsilon)})$ -accurate estimator given $O(d/(\epsilon^2/\log(1/\epsilon)))$ many ϵ -corrupted samples from an identity covariance subgaussian distribution; and (ii) an $O(\sqrt{\epsilon})$ -accurate estimator for any distribution $X \sim D$ with $\mathbf{Cov}[X] \preceq \mathbf{I}_d$, given $\tilde{O}(d/\epsilon)$ many ϵ -corrupted samples.

C.1. Setup and Algorithm Description

The pseudocode of the algorithm establishing Theorem C.2 is presented in Algorithm 3. We will define the necessary notation as needed (see the pseudocode for details). First, we assume that the distribution over the input samples is of the form $P = (1 - \epsilon)G + \epsilon B$, where G is the uniform distribution over the stable set of inliers and B is the uniform distribution on the outliers. Although this mixture may seem to suggest that the adversary only adds points, it is without loss of generality. Indeed, in the case that the adversary also removes points, we can think of G as the distribution of the remaining inliers (which continues to be stable with slightly worse parameters; see Lemma B.12).

We begin with a high-level explanation of Algorithm 3. At each iteration t, we assign a weight $w_t(x) \in [0,1]$ to each point x. Let P_t be the distribution on S, weighted according to w_t . Let μ_t and Σ_t be the mean and covariance of P_t , respectively. We want to assign scores to each point, using spectral properties of Σ_t and the stability of inliers, so that the scores over outliers are more than those of inliers. Essentially, if a direction v has variance larger than $1 + \Omega(\delta^2/\epsilon)$, then the stability of inliers implies that this must be due to outliers. Thus, we can assign scores based on the values $(v^T(x-\mu_t))^2$ that have provably more mass on outliers than inliers. The filters proposed in (Diakonikolas et al., 2016; 2017a) assigned scores based on a single direction, the leading eigenvector of Σ_t , and can take as many as $\Omega(d)$ iterations (see Appendix A.2).

To reduce the number of iterations, we need to filter in all directions of large variance simultaneously. Letting $\mathbf{B}_t \approx \mathbf{\Sigma}_t - (1 - C_1 \delta^2 / \epsilon) \mathbf{I}_d$, we would like to filter along all directions where the eigenvalue of \mathbf{B}_t is within a constant factor from $\lambda_t := \|\mathbf{B}_t\|_2$, not necessarily the leading eigenvector of \mathbf{B}_t . As we show in Appendix C.4, this can be approximately achieved by assigning scores for each point x based on $g_t(x) := \|\mathbf{M}_t(x - \mu_t)\|_2^2$, where $\mathbf{M}_t = \mathbf{B}_t^{\log d}$. At a high level, this happens because the spectrum of \mathbf{M}_t is distributed across along *all* large eigenvectors of \mathbf{B}_t .

Algorithm 3 Robust Mean Estimation in polylog iterations

```
1: Input: S = \{x_i\}_{i \in [n]}, \delta, \epsilon
 2: Let C_1 \ge 22, C be a sufficiently large constant, C_2 = 100C and C_3 = 0.1.
 3: Let R = \sqrt{(d/\epsilon)(1 + \delta^2/\epsilon)}.
 4: Let P = (1 - \epsilon)G + \epsilon B be the empirical distribution on the points from S. <sup>4</sup>
 5: Let K = \text{polylog}(d/\epsilon), L = \text{polylog}(nd/\epsilon).
 6: Obtain a naïve estimate \widehat{\mu} of \mu with \|\widehat{\mu} - \mu\|_2 \le 4R.
 7: Initialize w_1(x) \leftarrow \mathbb{1}\{\|x - \widehat{\mu}\|_2 \leq 5R\} for all x \in S.
 8: for t \in [K] do
          Let P_t be the distribution of P weighted by w_t.
 9:
          Let \mu_t, \Sigma_t be the mean and covariance of P_t.

Let \mathbf{B}_t = (\mathbf{E}_{X \sim P}[w_t(X)])^2 \mathbf{\Sigma}_t - \left(1 - C_1 \frac{\delta^2}{\epsilon}\right) \mathbf{I}_d and \mathbf{M}_t = \mathbf{B}_t^{\log d}.
10:
11:
           Let \lambda_t = \|\mathbf{B}_t\|_2. Compute \hat{\lambda}_t \in [0.8\lambda_t, 1.2\lambda_t] using power iteration.
12:
           if \hat{\lambda}_t > C_2 \delta^2 / \epsilon then
13:
               for j \in [L] do
14:
               z_{t,j} \sim \mathcal{U}(\{\pm 1\}^d), v_{t,j} \leftarrow \mathbf{M}_t z_{t,j}. end for
15:
16:
               Denote by \mathbf{U}_t the matrix having the vectors \frac{1}{\sqrt{L}}v_{t,j} for j \in [L] as rows.
17:
               Let \tilde{g}_t(x) = \|\mathbf{U}_t(x - \mu_t)\|_2^2 and
18:
               \tilde{\tau}_t(x) = \tilde{g}_t(x) \mathbb{1}\{\tilde{g}_t(x) > C_3 \|\mathbf{U}_t\|_F^2 \hat{\lambda}_t/\epsilon\},
               \ell_{\text{max}} \leftarrow (dR/\epsilon)^{C \log d}, T \leftarrow 0.01 \hat{\lambda}_t \|\mathbf{U}_t\|_F^2.
19:
               w_{t+1} \leftarrow \text{DownweightingFilter}(P, w_t, \tilde{\tau}_t, R, T, \ell_{\text{max}})
20:
           end if
21:
22: end for
23: return \mu_t.
```

Algorithm 4 Downweighting Filter

- 1: Input: $P, w, \tilde{\tau}, R, T, \ell_{\max}$
- 2: $r \leftarrow CdR^{2+4\log d}$.
- 3: Let $w_{\ell}(x) = w(x)(1 \tilde{\tau}(x)/r)^{\ell}$.
- 4: Find the smallest $\ell \in \{1, \dots, \ell_{\max}\}$ satisfying $\mathbf{E}_{X \sim P} [w_{\ell}(X)\tilde{\tau}(X)] \leq 2T$ using binary search.
- 5: **return** w_{ℓ} .

Even though assigning scores based on \mathbf{M}_t , i.e., $g_t(x)$, reduces the number of iterations, computing $g_t(x)$ for all $x \in S$ is slow. We thus use a Johnson-Lindenstrauss (JL) sketch of \mathbf{M}_t , denoted by \mathbf{U}_t . We denote by $\tilde{g}_t(x) := \|\mathbf{U}_t(x - \mu_t)\|_2$ the resulting scores. We claim that the set $\{\tilde{g}_t(x)\}_{x \in S}$ can be calculated in time $\tilde{O}(nd)$ such that for each $x \in S$, $\tilde{g}_t(x) \approx g_t(x)$. First, we will show (see Lemma C.5) that \mathbf{U}_t can be as small as $L \times d$, where L is some polylog $(nd/(\epsilon\tau))$, which follows from the classical JL lemma (stating that n points can be linearly embedded into a $\log n$ -dimensional space). Also, each row of \mathbf{U}_t can be computed by repeatedly multiplying a vector $\log d$ times by \mathbf{B}_t (Line 15). By the following remark, all rows of \mathbf{U}_t can be computed in time $\tilde{O}(nd)$ and thus, each iteration of Algorithm 3 runs in near-linear time.

Remark C.3. (Efficient Implementation) Note that for any $v \in \mathbb{R}^d$, the vector $\mathbf{B}_t v$ can be calculated in O(nd) time. This is because $\mathbf{\Sigma}_t v = \sum_{x \in S} w_t(x) (v^T(x - \mu_t)) (x - \mu_t) / (\sum_{x \in S} w_t(x))$ which means that the result can be computed in O(nd) time by calculating μ_t and $v^T(x - \mu_t)$ first. Regarding Line 7, an approximate large eigenvector can be computed via power iteration, i.e., starting from a random Gaussian vector and multiplying by \mathbf{B}_t iteratively $\log d$ many times (see, for example, (Blum et al., 2020)). As mentioned above, each of these multiplications can be done in O(nd) time.

For the proof of correctness, we require that the JL and spectral approximations used by the algorithm are sufficiently accurate. We prove that the following event occurs with high probability.

Condition C.4 (Deterministic Conditions For Algorithm 3). For all $t \in [K]$, the following hold:

- 1. Spectral norm of \mathbf{B}_t : $\widehat{\lambda}_t \in [0.8\lambda_t, 1.2\lambda_t]$.
- 2. Frobenius norm: $\|\mathbf{U}_t\|_F^2 \in [0.8\|\mathbf{M}_t\|_F^2, 1.2\|\mathbf{M}_t\|_F^2]$.
- 3. *Scores*: For all $x \in S$, $\tilde{g}_t(x) \in [0.8g_t(x), 1.2g_t(x)]$.

C.2. Establishing the Deterministic Conditions

In this section, we establish that Condition C.4 holds with high probability. Regarding Item 1 of this condition, an approximate large eigenvector can be computed via power iteration as described in Remark C.3. This gives us an algorithm that runs in time $O(nd \log d \log(K/\tau))$ and satisfies Item 1 with probability $1-\tau$.

We now move to the other two conditions. The claim is that instead of using the matrix \mathbf{M}_t to calculate the scores, it suffices to store and use only a small set of random projections $\{\mathbf{M}_t z_{t,j}\}_{j \in [L]}$. This is exactly the Johnson-Lindenstrauss sketch that is computed in Line 14 of Algorithm 3. Using Fact B.7, we get the following guarantees (see Appendix H.1 for the proof).

Lemma C.5. Fix a set of n points $x_1, \ldots, x_n \in \mathbb{R}^d$. For $t \in [K]$, define $g_t(x) := \|\mathbf{M}_t(x - \mu_t)\|_2^2$ and let $\tilde{g}_t(x), v_{t,j}$ as in Algorithm 3. If C is a sufficiently large constant and $L = C \log((n+d)K/\tau)$, with probability at least $1 - \tau$, for every $t \in [K]$ we have the following:

- 1. $0.8g_t(x_i) \leq \tilde{g}_t(x_i) \leq 1.2g_t(x_i)$ for every $i \in [n]$,
- $2. \ \ 0.8 \|\mathbf{M}_t\|_F^2 \leq \left(\tfrac{1}{L} \sum_{j=1}^L \|v_{t,j}\|_2^2 \right) \leq 1.2 \|\mathbf{M}_t\|_F^2.$

This concludes the proof that Condition C.4 is satisfied with high probability.

C.3. Downweighting Filter

We use the following re-weighting procedure also used in (Dong et al., 2019). Recall that P denotes the empirical distribution on the samples, which we write as $P = (1 - \epsilon)G + \epsilon B$, where G and B are the contributions from the good and bad samples respectively. Roughly speaking, our filter guarantees two things when going from the weights w(x) to w'(x):

⁴Without loss of generality, outliers are within O(R) from μ in ℓ_2 -norm. This is ensured in Line 7, which removes only ϵ -fraction of inliers (Claim C.12).

- 1. The weight removed from the outliers is greater than the weight removed from the inliers.
- 2. $\mathbf{E}_{X \sim P}[w'(X)\tilde{\tau}(X)] \leq 2\mathbf{E}_{X \sim G}[w(X)\tilde{\tau}(X)]$, i.e., the weighted mean of scores after filtering over both inliers and outliers is at most twice the weighted mean of scores of inliers before filtering.

Regarding the first guarantee, since the fraction of outliers is at most ϵ , this ensures that the filtered distribution P_t will never be more than $O(\epsilon)$ -far in total variation distance from the initial (corrupted) distribution, and thus the condition of the certificate lemma that $d_{\text{TV}}(P, P_t) \leq O(\epsilon)$ will always be satisfied. The second guarantee ensures that the filtering step reduces the average score significantly. We prove the following in Appendix H.2.

Lemma C.6. Let $P=(1-\epsilon)G+\epsilon B$ be the empirical distribution on n samples, as in Algorithm 3. If $(1-\epsilon)\mathbf{E}_{X\sim G}[w(X)\tilde{\tau}(X)] \leq T$, $\|\tilde{\tau}\|_{\infty} \leq r$, and $\ell_{\max} > r/T$, then Algorithm 4 modifies the weight function w to w' such that

1.
$$(1 - \epsilon) \mathbf{E}_{X \sim G}[w(X) - w'(X)] < \epsilon \mathbf{E}_{X \sim B}[w(X) - w'(X)],$$

2.
$$\mathbf{E}_{X \sim P} \left[w'(X) \tilde{\tau}(X) \right] \leq 2T$$
,

and the algorithm terminates after $O(\log(\ell_{\max}))$ iterations, each of which takes O(n) time.

We note that the two conditions $\|\tilde{\tau}\|_{\infty} \leq r$, $\ell_{\max} > r/T$ of Lemma C.6 hold by our choice of ℓ_{\max} and r inside Algorithm 3 and Algorithm 4 as follows. For $\|\tilde{\tau}\|_{\infty}$, we have the following upper bound

$$\tilde{\tau}_t(x) \le \tilde{g}_t(x) \le \|\mathbf{U}_t(x - \mu_t)\|_2^2 \lesssim R^2 \|\mathbf{U}_t\|_2^2 \lesssim R^2 \|\mathbf{M}_t\|_F^2 \lesssim R^2 \|\mathbf{\Sigma}_t\|_2^{2\log d} = O(dR^{2+4\log d}),$$
(2)

where we used the guarantee of our JL sketch that $\|\mathbf{U}_t\|_2^2 \le 1.2 \|\mathbf{M}_t\|_F^2$ (Lemma C.5). A crude upper bound on r/T follows from the following inequalities:

$$\frac{r}{T} \lesssim \frac{dR^{2+4\log d}}{\widehat{\lambda}_t \|\mathbf{M}_t\|_T^2} \lesssim \frac{dR^{2+4\log d}}{\lambda_t \|\mathbf{M}_t\|_F^2} \lesssim \left(\frac{dR}{\delta^2/\epsilon}\right)^{O(\log d)},$$

where the first inequality uses the values of r and T as set in the algorithm, the second inequality uses Items 1 and 2 of the deterministic conditions of Condition C.4, and the last inequality uses the fact that $\|\mathbf{M}_t\|_F^2 \ge \|\mathbf{M}_t\|_2^2$ and $\|\mathbf{M}_t\|_2^2$ cannot be smaller than $(C_2\delta^2/\epsilon)^{O(\log d)}$ (otherwise Line 20 terminates the algorithm).

We now use the guarantees of Algorithm 4 as follows. We first show that the weighted mean of the inliers' scores is small.

Lemma C.7. Under the setting of Algorithm 3 and the deterministic Condition C.4, we have that $\mathbf{E}_{X \sim G}[w_t(X)\tau_t(X)]$ and $\mathbf{E}_{X \sim G}[w_t(X)\tilde{\tau}_t(X)]$ are bounded from above by $c\lambda_t \|\mathbf{M}_t\|_F^2$ for some constant c of the form $c = C/C_2$, where C_2 is the constant used in Line 13 and C is some absolute constant.

The proof is based on stability arguments from Appendix B.2 and can be found in Appendix H.3.

Remark C.8. In our analysis, it will be important that the constant c in Lemma C.7 can be made sufficiently smaller than 1, for example, c < 0.01. This can be achieved by choosing C_2 to be a large enough constant.

Using Algorithm 4, we get that the weighted sum of scores after filtering is also small.

Lemma C.9. Under the setting of Algorithm 3 and the deterministic Condition C.4, we have that $\epsilon \mathbf{E}_{X \sim B}[w_{t+1}(X)\tilde{\tau}_t(X)] < c\lambda_t \|\mathbf{M}_t\|_F^2$, with c being of the form $c = C/C_2$, where C_2 is the constant used in Line 13 and C is some absolute constant. Furthermore, $\epsilon \mathbf{E}_{X \sim B}[w_{t+1}(X)\tau_t(X)] < c\lambda_t \|\mathbf{M}_t\|_F^2$.

Proof. The first claim follows by the stopping condition of the algorithm, Lemma C.7, and the fact that $w_{t+1} \leq w_t$. We now prove the second conclusion by relating τ to $\tilde{\tau}$: Recall that we denote by S the ϵ -corrupted version of the original set of samples S_0 . Since $\tilde{g}_t(x)$ is within a constant factor of $g_t(x)$ (Condition C.4) for all $x \in S$, the scores $\tilde{\tau}_t(x)$ and $\tau_t(x)$ are comparable (up to an additive term) as shown below.

Claim C.10. In the setting of Algorithm 3 and under the Condition C.4, if $x \in S$, we have that $\tau_t(x) \leq 1.25\tilde{\tau}_t(x) + 3C_3(\lambda_t/\epsilon)\operatorname{tr}(\mathbf{M}_t^2)$, where C_3 is the constant used in Algorithm 3.

We prove Claim C.10 in Appendix H.5. Using Claim C.10, we have the following set of inequalities:

$$\begin{split} \epsilon \cdot \mathop{\mathbf{E}}_{X \sim B}[w_{t+1}(X)\tau_t(X)] &= \frac{1}{n} \sum_{x_i \in S \backslash S_0} w_{t+1}(x_i)\tau_t(x_i) \\ &\leq 3C_3\lambda_t \ \|\mathbf{M}_t\|_F^2 + \frac{1}{n} 1.25 \sum_{i \in S \backslash S_0} w_{t+1}(x_i)\tilde{\tau}_t(x_i) \qquad \text{(using Claim C.10)} \\ &= 3C_3\lambda_t \|\mathbf{M}_t\|_F^2 + 1.25\epsilon \mathop{\mathbf{E}}_{X \sim B}[w_{t+1}(X)\tilde{\tau}_t(X)] \\ &\leq 3C_3\lambda_t \|\mathbf{M}_t\|_F^2 + 1.25c\lambda_t \|\mathbf{M}_t\|_F^2 \qquad \text{(using the first part of Lemma C.9)} \\ &< 10(C/C_2)\lambda_t \|\mathbf{M}_t\|_F^2 \ . \qquad \text{(using the value of C_3)} \end{split}$$

The last inequality above uses the fact that the constant C_3 is chosen to be $C_3 = C/C_2$ in Algorithm 3, where C is a sufficiently large constant.

C.4. Correctness of Algorithm 3: Proof of Theorem C.2

The rest of this section is dedicated to proving Theorem C.2. We first state the correctness of the naïve approximation step of Line 6, then record the invariants of the algorithm in Appendix C.4.1, and finally show in Appendix C.4.2 that it suffices for the number of iterations K to be bounded by some $\operatorname{polylog}(d, R, 1/\epsilon)$.

The naïve approximation step of Line 6 is based on the following folklore fact (see Appendix H.4 for more details).

Claim C.11. Let the fraction of outliers be $\epsilon < 1/10$ and a parameter $0 < \tau < 1$. Let the distribution $P = (1 - \epsilon)G + \epsilon B$. Let R > 0, $\mu \in \mathbb{R}^d$ be such that $\Pr_{X \sim G}[\|X - \mu\|_2 > R] \le \epsilon$. There is an estimator $\widehat{\mu}$ on $k = O(\log(1/\tau))$ samples from P such that $\|\widehat{\mu} - \mu\|_2 \le 4R$ with probability at least $1 - \tau$. Furthermore, $\widehat{\mu}$ can be computed in time $O(k^2d)$ and memory O(kd).

Claim C.12 below gives a valid upper bound on R using the (ϵ, δ) -stability of the good distribution.

Claim C.12. If
$$R = \sqrt{\frac{d}{\epsilon}(1 + \delta^2/\epsilon)}$$
, then $\Pr_{X \sim G}[\|X - \mu\|_2 > R] \le \epsilon$.

Proof. By Markov's inequality, we have that

$$\Pr_{X \sim G} \left[\|X - \mu\|_2^2 \ge \frac{d}{\epsilon} \left(1 + \delta^2 / \epsilon \right) \right] \le \epsilon \frac{\mathbf{E}_{X \sim G} [\|X - \mu\|_2^2]}{d(1 + \delta^2 / \epsilon)} \le \epsilon.$$

C.4.1. INVARIANTS OF ALGORITHM 3

Recall that the end goal is to obtain a filtered version, P_t , of P that is not too far from P in total variation distance $d_{\text{TV}}(P, P_t) = O(\epsilon)$, and satisfies that $\|\mathbf{B}_t\|_2 = O(\delta^2/\epsilon)$. For the first condition to be satisfied, we ensure that the Downweighting filter removes more weight from G than B (Lemma C.6). Using this, we show that $\mathbf{E}_{X \sim G}[w_t(X)] \ge 1 - O(\epsilon)$, which implies the bound on the total variation distance (Claim C.13). The proofs are deferred to Appendix H.5.

Claim C.13. Under Condition C.4, Algorithm 3 maintains the following invariant: $\mathbf{E}_{X \sim G}[w_t(X)] \ge 1 - 3\epsilon$. In particular, if $\epsilon \le 1/8$, then $d_{\mathrm{TV}}(P_t, P) \le 9\epsilon$.

The following properties of \mathbf{B}_t as PSD operator will also be useful later on.

Claim C.14. Under Condition C.4, if
$$C_1 \ge 22$$
, $\mathbf{B}_t \succeq (0.5C_1\delta^2/\epsilon)\mathbf{I}_d$ for every $t \in [K]$.

The proof of Claim C.14 is provided in Appendix H.5. Although just showing that $\mathbf{B}_t \succeq 0$ would suffice for this section, the slightly stronger bound of the above claim will be useful in Appendix D. Claim C.14 follows from Claim C.13 and the stability of G. In particular, the stability of G implies that $\overline{\Sigma}_G \succeq (1 - \delta^2/\epsilon)\mathbf{I}_d$. We now prove the following claim, which is the reason for having the multiplicative factor of $\mathbf{E}_{X\sim P}[w_t(X)]^2$ in the definition of \mathbf{B}_t .

Claim C.15. We have that $\mathbf{B}_{t+1} \leq \mathbf{B}_t$ for every $t \in [K]$.

Proof. We use the alternative definition of the covariance matrix: Let X, Y be i.i.d. from P, then

$$\Sigma_t = \frac{1}{2(\mathbf{E}_{X \sim P}[w_t(X)])^2} \sum_{X,Y \sim P} [w_t(X)w_t(Y)(X - Y)(X - Y)^T] .$$

Since $w_{t+1}(x) \leq w_t(x)$ for all x, this completes the proof.

C.4.2. REDUCING THE POTENTIAL FUNCTION

Recall that each iteration of Algorithm 3 can be implemented in near-linear time. Thus, it remains to show that the choice $K = C \log d \log(dR/\epsilon)$ suffices to guarantee correctness of our algorithm. We now sketch the proof using a potential function argument. Let Λ_t be the vector in \mathbb{R}^d containing the eigenvalues of \mathbf{B}_t . Recall that our goal is to show that $\|\mathbf{B}_t\|_2 = \|\Lambda_t\|_\infty = O(\delta^2/\epsilon)$ in polylog many iterations. Let $p := 2 \log d$. Since $\|x\|_p = \Theta(\|x\|_\infty)$ for any $x \in \mathbb{R}^d$, we are motivated to use the potential function $\phi_t := \|\Lambda_t\|_p^p$. We now focus on showing that ϕ_t decreases rapidly. Observe that for any $i \in \mathbb{Z}_+$, $\operatorname{tr}(\mathbf{B}_t^i) = \|\Lambda_t\|_i^i$. We start with the following inequalities (and explain them directly below):

$$\phi_{t+1} = \|\Lambda_{t+1}\|_{p}^{p} \le \left(d^{\frac{1}{p(p+1)}} \|\Lambda_{t+1}\|_{p+1}\right)^{p}$$

$$= d^{\frac{1}{p+1}} \left(\|\Lambda_{t+1}\|_{p+1}^{p+1}\right)^{\frac{p}{p+1}}$$

$$= d^{\frac{1}{p+1}} \left(\operatorname{tr}(\mathbf{B}_{t+1}^{p+1})\right)^{\frac{p}{p+1}}$$

$$= d^{\frac{1}{p+1}} \left(\operatorname{tr}(\mathbf{M}_{t+1}\mathbf{B}_{t+1}\mathbf{M}_{t+1})\right)^{\frac{p}{p+1}}$$

$$\le d^{\frac{1}{p}} \left(\operatorname{tr}(\mathbf{M}_{t}\mathbf{B}_{t+1}\mathbf{M}_{t})\right)^{\frac{p}{p+1}}, \tag{3}$$

where the first line uses Fact B.1, the third one uses $\operatorname{tr}(\mathbf{B}_{t+1}^i) = \|\Lambda_{t+1}\|_i^i$, the fourth line uses that $\mathbf{M}_{t+1}\mathbf{B}_{t+1}\mathbf{M}_{t+1} = \mathbf{B}_{t+1}^{p+1}$, and the last one uses the fact $\mathbf{B}_{t+1} \leq \mathbf{B}_t$, which holds because removing points can only make their covariance smaller; see Appendix C.4 for more details.

Then the goal becomes to bound from above the term $\operatorname{tr}(\mathbf{M}_t\mathbf{B}_{t+1}\mathbf{M}_t)$. The claim is that $\operatorname{tr}(\mathbf{M}_t\mathbf{B}_{t+1}\mathbf{M}_t)$ is related to $\mathbf{E}_{X\sim P}[w_{t+1}(X)\tau_t(X)]$, and thus can be bounded by $c\lambda_t\|\mathbf{M}_t\|_F^2$. Using the guarantees of the Downweighting filter (Lemma C.9), we prove the following result:

Lemma C.16. Consider the setting of Algorithm 3 and assume that Condition C.4 holds. Then $\operatorname{tr}(\mathbf{M}_t\mathbf{B}_{t+1}\mathbf{M}_t) \leq c\lambda_t \|\mathbf{M}_t\|_F^2$ for some c of the form C/C_2 , where C_2 is the constant used in Line 13 and C is some absolute constant.

Before giving the details regarding Lemma C.16, we first show that it suffices to prove that the potential function decreases by a multiplicative factor. In the rest of the proof, we will assume that c < 0.1, which can be guaranteed by taking C_2 to be a sufficiently large constant (cf. Remark C.8). We continue with Equation (3) as follows:

$$\begin{split} \phi_{t+1} & \leq d^{\frac{1}{p}} \left(\text{tr}(\mathbf{M}_{t} \mathbf{B}_{t+1} \mathbf{M}_{t}) \right)^{\frac{p}{p+1}} \\ & \leq d^{\frac{1}{p}} \left(c \| \Lambda_{t} \|_{\infty} \| \Lambda_{t} \|_{p}^{p} \right)^{\frac{p}{p+1}} \\ & \leq d^{\frac{1}{p}} c^{\frac{p}{p+1}} \left(\| \Lambda_{t} \|_{p} \| \Lambda_{t} \|_{p}^{p} \right)^{\frac{p}{p+1}} \\ & = d^{\frac{1}{p}} c^{\frac{p}{p+1}} \| \Lambda_{t} \|_{p}^{p} \\ & \leq 3 \sqrt{c} \| \Lambda_{t} \|_{p}^{p} \leq 0.9999 \phi_{t} \;, \end{split}$$
 (using $\| \Lambda_{t} \|_{\infty} \leq \| \Lambda_{t} \|_{i}$ for $i \geq 1$)

where the last line uses that $d^{1/p} = \exp(\frac{\log d}{2\log d}) \le 3$, $p/(p+1) \ge 0.5$, and c < 1. We thus get the desired convergence.

The final step is to bound the number of iterations needed for Lemma B.11 to ensure that $\|\mu_t - \mu\|_2 = O(\delta)$. Concretely, due to our naïve pruning, at the beginning of the algorithm we have the upper bound $\phi_1 \leq dR^{O(\log d)}$. After K iterations, we have that $\phi_K \leq 0.99^K dR^{O(\log d)}$. Setting K as

$$K = C \log d \log \left(\frac{dR}{\delta^2 / \epsilon} \right) \tag{4}$$

suffices to have that $\|\mathbf{B}_K\|_2 \leq (dO(\delta^2/\epsilon)^{2\log d})^{\frac{1}{2\log d}} = O(\delta^2/\epsilon)$. This implies that

$$\|\mathbf{\Sigma}_{K}\|_{2} \leq \frac{1}{\mathbf{E}_{X \sim P}[w_{K}(X)]^{2}} (\|\mathbf{B}_{K}\|_{2} + 1) \leq 1 + \left(\frac{1}{(1 - 3\epsilon)^{2}} - 1\right) + O\left(\frac{\delta^{2}}{\epsilon}\right) \leq 1 + O\left(\frac{\delta^{2}}{\epsilon}\right) , \tag{5}$$

where we used that $\mathbf{E}_{X\sim P}[w_K(X)])^2 \geq 1-3\epsilon$, $\delta \geq \epsilon$, and $\epsilon \leq \epsilon_0$. An application of Lemma B.11 shows that the estimate has error at most $\|\mu_t - \mu\|_2 = O(\delta)$. This completes the proof of Theorem C.2. The rest of the section focuses on proving Lemma C.16.

Proof Sketch of Lemma C.16 Before giving the full proof of Lemma C.16, we provide a brief proof sketch. By the definition of \mathbf{B}_{t+1} , we have the following (the full proof is deferred to the end of this section)

$$\operatorname{tr}(\mathbf{M}_{t}\mathbf{B}_{t+1}\mathbf{M}_{t}) \leq \underset{X \sim P}{\mathbf{E}}[w_{t+1}(X)] \underset{X \sim P}{\mathbf{E}}[w_{t+1}(X)g_{t}(X)] - \left(1 - C_{1}\frac{\delta^{2}}{\epsilon}\right) \|\mathbf{M}_{t}\|_{F}^{2}.$$

In order to bound from above $\mathbf{E}_{X\sim P}[w_{t+1}(X)g_t(X)]$, one can consider the contribution due to inliers (distribution G) and contribution due to outliers (distribution B). Using the stability of inliers and Corollary B.14, we have that $\mathbf{E}_{X\sim G}[w(X)g_t(X)] \leq (1+c\lambda_t)\|\mathbf{M}_t\|_F^2$, for any weight function w satisfying the conditions of Corollary B.14. We know that w_{t+1} satisfies them because of our invariant in Claim C.13. Turning to the contribution of outliers, we want to bound $\epsilon \cdot \mathbf{E}_{X\sim B}[w_{t+1}(X)g_t(X)]$. By definition, we have that $g_t(x) \leq \tau_t(x) + C_3\lambda_t\|\mathbf{M}_t\|_F^2/\epsilon$, and thus we get that the desired expression is bounded from above by $\epsilon \mathbf{E}_{X\sim B}[w_{t+1}(X)\tau_t(X)] + C_3\lambda_t\|\mathbf{M}_t\|_F^2$. The first expression was bounded from above in Lemma C.9 by using the downweighting filter, and the second is small because of how C_3 is set in our algorithm.

This completes the proof sketch of Lemma C.16. We now provide the complete proof.

Proof of Lemma C.16. We will bound the contribution of inliers and outliers to the quantity $\mathbf{E}_{X\sim P}[w_{t+1}(X)g_t(X)]$ from above. Recall from our notation that the decomposition into inliers and outliers is $P=(1-\epsilon)G+\epsilon B$. For the inliers, we use Corollary B.14 with $\mathbf{U}=\mathbf{M}_t$ and $b=\mu_t$ to obtain the following:

$$\mathbf{E}_{X \sim G}[w_{t+1}(X)g_t(X)] \le \|\mathbf{M}_t\|_F^2 \left(1 + \frac{\delta^2}{\epsilon} + \|\mu_t - \mu\|_2^2 + 2\delta\|\mu_t - \mu\|_2\right) \le \|\mathbf{M}_t\|_F^2 \left(1 + c\lambda_t\right) , \tag{6}$$

where the last inequality uses that, by the certificate lemma (Lemma B.11), every term except the first in the previous expression is less than a sufficiently small fraction of λ_t .

Regarding the outliers, we decompose their contribution to $\mathbf{E}_{X\sim P}[w_{t+1}(X)g_t(X)]$ into two sets: (i) the set of points with projection greater than the threshold $C_3\|\mathbf{M}_t\|_F^2\lambda_t/\epsilon$ used in Line 18 of the algorithm, and (ii) the set of points with smaller projection. Concretely, letting $L_t:=\{x\in\mathbb{R}^d:g_t(x)>C_3\|\mathbf{M}_t\|_F^2\lambda_t/\epsilon\}$, we have that

$$\epsilon \underset{X \sim B}{\mathbf{E}}[w_{t+1}(X)g_t(X)] = \epsilon \underset{X \sim B}{\mathbf{E}}[w_{t+1}(X)\tau_t(X)] + \epsilon \underset{X \sim B}{\mathbf{E}}[w_{t+1}(X)g_t(X)\mathbb{1}\{x \notin L_t\}]
\leq c\lambda_t \|\mathbf{M}_t\|_F^2 + \epsilon C_3 \|\mathbf{M}_t\|_F^2 \lambda_t / \epsilon \leq c' \|\mathbf{M}_t\|_F^2 \lambda_t ,$$
(7)

where the first inequality follows from Lemma C.9 and the second inequality follows from the choice of C_3 in Algorithm 3.

We also use the following relation on Σ_{t+1} :

$$\Sigma_{t+1} = \underset{X \sim P_{t+1}}{\mathbf{E}} \left[(X - \mu_{t+1})(X - \mu_{t+1})^T \right]
\leq \underset{X \sim P_{t+1}}{\mathbf{E}} \left[(X - \mu_t)(X - \mu_t)^T \right]
= \frac{1}{\mathbf{E}_{X \sim P}[w_{t+1}(X)]} \underset{X \sim P}{\mathbf{E}} \left[w_{t+1}(X)(X - \mu_t)(X - \mu_t)^T \right].$$
(8)

Recalling the definition $\mathbf{B}_{t+1} = (\mathbf{E}_{X \sim P}[w_{t+1}(X)])^2 \mathbf{\Sigma}_{t+1} - \left(1 - C_1 \frac{\delta^2}{\epsilon}\right) \mathbf{I}_d$, Equation (8) implies that $\mathbf{B}_{t+1} \preceq \mathbf{F}_{t+1}$, where $\mathbf{F}_{t+1} := (\mathbf{E}_{X \sim P}[w_{t+1}(X)]) \mathbf{E}_{X \sim P}[w_{t+1}(X)(X - \mu_t)(X - \mu_t)^T] - \left(1 - C_1 \frac{\delta^2}{\epsilon}\right) \mathbf{I}_d$. Using Fact B.2 along with the fact that $\mathbf{B}_{t+1} \succeq 0$ (Claim C.14), we get the following:

$$\operatorname{tr}(\mathbf{M}_{t}\mathbf{B}_{t+1}\mathbf{M}_{t}) = \operatorname{tr}(\mathbf{M}_{t}^{2}\mathbf{B}_{t+1}) < \operatorname{tr}(\mathbf{M}_{t}^{2}\mathbf{F}_{t+1})$$
(using $\operatorname{tr}(\mathbf{ABC}) = \operatorname{tr}(\mathbf{CAB})$)

$$= \operatorname{tr}\left(\mathbf{M}_{t}\left(\sum_{X\sim P}[w_{t+1}(X)]\sum_{X\sim P}[w_{t+1}(X)(X-\mu_{t})(X-\mu_{t})^{T}] - \left(1-C_{1}\frac{\delta^{2}}{\epsilon}\right)\mathbf{I}_{d}\right)\mathbf{M}_{t}\right)$$

$$= \sum_{X\sim P}[w_{t+1}(X)]\sum_{X\sim P}[w_{t+1}(X)\operatorname{tr}((X-\mu_{t})^{T}\mathbf{M}_{t}^{2}(X-\mu_{t}))] - \left(1-C_{1}\frac{\delta^{2}}{\epsilon}\right)\|\mathbf{M}_{t}\|_{F}^{2}$$

$$= \sum_{X\sim P}[w_{t+1}(X)]\sum_{X\sim P}[w_{t+1}(X)g_{t}(X)] - \left(1-C_{1}\frac{\delta^{2}}{\epsilon}\right)\|\mathbf{M}_{t}\|_{F}^{2}$$

$$\leq \left(1+c\lambda_{t}+c'\lambda_{t}-(1-C_{1}\delta^{2}/\epsilon)\right)\|\mathbf{M}_{t}\|_{F}^{2}$$

$$\leq c''\lambda_{t}\|\mathbf{M}_{t}\|_{F}^{2}.$$
(using Equations (6) and (7))
$$\leq c''\lambda_{t}\|\mathbf{M}_{t}\|_{F}^{2}.$$

This concludes the proof.

D. Efficient Streaming Algorithm for Robust Mean Estimation

We now turn to the main focus of this paper and present a low-memory algorithm for robust mean estimation. Our algorithm works in two setups: (i) the single-pass streaming setting, where a set of i.i.d. samples from an ϵ -corrupted distribution in total variation distance (Definition A.2) arrive *one at a time* (Definition A.1), and (ii) the strong-contamination model (Definition C.1), where the algorithm is allowed poly-logarithmically many passes over the input stream (defined below).

Definition D.1 (Streaming Model in k Passes). For a fixed set S, the elements of S are revealed to the algorithm one at a time. This process is repeated k times. The sequence of elements in S within each pass can be arbitrary.

Our main result is the following theorem for the single-pass streaming model, which is a generalized version of Theorem A.3:

Theorem D.2 (Robust Mean Estimation in Single-Pass Streaming Model). Let $d \in \mathbb{Z}_+$, $0 < \tau < 1$, $0 < \epsilon < \epsilon_0$ for a sufficiently small constant ϵ_0 , and $\delta \ge \epsilon$. Let D be a distribution which is $(C\epsilon, \delta)$ -stable with respect to the (unknown) vector $\mu \in \mathbb{R}^d$, for a sufficiently large constant C > 0. Let R be any radius such that $\Pr_{X \sim D}[\|X - \mu\|_2 > R] \le \epsilon$. Let P be a distribution with $d_{TV}(P, D) \le \epsilon$. There exists an algorithm that given ϵ, δ, τ and

$$n = O\left(R^2 \max\left(d, \frac{\epsilon}{\delta^2}, \frac{(1 + \delta^2/\epsilon)d}{\delta^2 R^2}, \frac{\epsilon^2 d}{\delta^4}, \frac{R^2 \epsilon^2}{\delta^2}, \frac{R^2 \epsilon^4}{\delta^6}\right) \operatorname{polylog}\left(d, \frac{1}{\epsilon}, \frac{1}{\tau}, R\right)\right)$$
(9)

i.i.d. samples from P in a stream according to the model of Definition A.1, runs in time $nd \operatorname{polylog}(d, 1/\epsilon, 1/\tau, R)$, uses memory $d \operatorname{polylog}(d, 1/\epsilon, 1/\tau, R)$, and returns a vector $\widehat{\mu}$ such that, with probability at least $1 - \tau$, it holds that $\|\mu - \widehat{\mu}\|_2 = O(\delta)$.

Note that Theorem A.3 in Appendix A.1 is a special case of Theorem D.2 for the two important families of distributions: (i) subgaussian distributions with identity covariance, and (ii) distributions with bounded covariance.

- 1. For subgaussian distributions with identity covariance, we have that $R = \Theta(\sqrt{d \log(1/\epsilon)})$, $\delta = O(\epsilon \sqrt{\log(1/\epsilon)})$, and thus $n = \tilde{O}\left(d^2/\epsilon^2\right)$.
- 2. For distributions with covariance at most identity, we have that $R = \Theta(\sqrt{d/\epsilon})$, $\delta = O(\sqrt{\epsilon})$, and thus $n = \tilde{O}\left(d^2/\epsilon\right)$.

In order to obtain a low-memory algorithm for the robust mean estimation problem, we start with an obstacle that one faces when trying to modify the existing Algorithm 3 to that setting. The issue is that, since n can be much larger than d, we cannot even store the weight function w_t . Fortunately, this can be handled by freshly computing the scores $w_t(x)$ for any given x, whenever we need them. This requires us to store only $\{(\mathbf{U}_t, \ell_t) : t \in [K]\}$, where \mathbf{U}_t is the Johnson-Lindenstrauss sketch at the iteration t, and ℓ_t is the corresponding count from the downweighting filter. This can be achieved with additional poly-logarithmic memory. Thus, Algorithm 3 can be readily extended to setting (ii), giving us Corollary D.3.

Corollary D.3 (Robust Mean Estimation in Multiple Passes Streaming Model). Let $d \in \mathbb{Z}_+$, $0 < \tau < 1$ and $0 < \epsilon < \epsilon_0$ for a sufficiently small constant ϵ_0 , and $\delta \geq \epsilon$. Let S be an ϵ -corrupted version of a set that is $(C\epsilon, \delta)$ -stable with respect to the (unknown) vector $\mu \in \mathbb{R}^d$, for a sufficiently large constant C. Denote by n the cardinality of S. There exists an algorithm that operates in the streaming model of Definition D.1 with $k = \text{polylog}(d, 1/\epsilon, 1/\tau)$ and, given ϵ, δ, τ and T, runs in time $nd \text{ polylog}(d, 1/\epsilon, 1/\tau)$, uses additional memory $d \text{ polylog}(d, 1/\epsilon, 1/\tau)$, and finds a vector $\hat{\mu}$ such that, with probability at least $1 - \tau$, it holds $\|\mu - \hat{\mu}\|_2 = O(\delta)$.

In the main body of this section, we prove Theorem D.2.

D.1. Setup and Algorithm Description

Moving to the single-pass streaming model and Theorem D.2 requires a change in perspective: instead of having a corrupted dataset, we now have sample access to a distribution P such that $d_{\text{TV}}(P,D) \leq \epsilon$, where D is a stable distribution. We will reweight this distribution using weights, $w_t(\cdot)$, that are now functions on the whole \mathbb{R}^d instead of a fixed dataset. Thus, P_t now denotes the reweighting of the (corrupted) distribution P with the weights w_t . Similarly μ_t , Σ_t , \mathbf{B}_t , \mathbf{M}_t denote the quantities that pertain to the distribution P_t . The goal of our algorithm remains essentially the same: obtain P_t such that $d_{\text{TV}}(P_t, P) = O(\epsilon)$ and $\|\Sigma_t\|_2 \leq 1 + O(\delta^2/\epsilon)$; Lemma B.11 would then imply that $\|\mu_t - \mu\|_2 = O(\delta)$. Before presenting the pseudocode of Algorithm 5, we identify two problems that arise in generalizing our results from Appendix C and provide an overview of their solutions:

Calculating Scores Recall that the only place where \mathbf{M}_t is used in Algorithm 3 is Line 15, where \mathbf{M}_t is multiplied with the vectors $z_{t,j}$. Let z be an arbitrary vector. Since $\mathbf{M}_t = \mathbf{B}^{\log d}$, in the previous section we were able to compute $\mathbf{M}_t z$ by iteratively multiplying z by \mathbf{B}_t . Since we now do not have access to \mathbf{B}_t , but only sample access to P_t , we need a sufficiently fine approximation $\hat{\mathbf{B}}_t$ of \mathbf{B}_t (obtained using i.i.d. samples). The natural approach would then be to multiply $\hat{\mathbf{B}}_t$ with z iteratively $\log d$ many times. Even though $\hat{\mathbf{B}}_t z$ can be computed in a streaming fashion (as outlined in the previous section), it is not possible to compute $(\hat{\mathbf{B}}_t)^{\log d} z$ without accessing the data $\log d$ times. To circumvent this issue, we use a fresh sample approximation of \mathbf{B}_t in every multiplication step. That is, we approximate $\mathbf{M}_t z$ by $\widehat{\mathbf{M}}_t z$, where $\widehat{\mathbf{M}}_t := \prod_{j=1}^p \widehat{\mathbf{B}}_{t,j}$ and each $\widehat{\mathbf{B}}_{t,j}$ is computed on a different set of samples. This approach crucially leverages the fact that in the contamination model of Definition A.2, outliers are added in a way that is oblivious to the inliers, and therefore these datasets are statistically identical and independent of each other. We show in Appendix D.3 that the resulting $\widehat{\mathbf{M}}_t$ is a sufficiently accurate approximation of \mathbf{M}_t . Similarly, we need to modify the Downweighting filter, since its implementation using binary search requires performing checks of the form $\mathbf{E}_{X\sim P}[w(X)\tilde{\tau}(x)] > 2T$ and calculating the weighted mean exactly is no longer possible. We propose a sample-efficient estimator to approximate that expectation (see Lemma D.16 in Appendix D.3.3) and run an "approximate" variant of binary search (see Appendix D.2).

Cover Argument We now turn to the more technical issue of controlling the size of the JL-sketch, i.e., the number of rows, L, of the matrix $\mathbf{U}_t \in \mathbb{R}^{L \times d}$. For simplicity, assume $\widehat{\mathbf{M}}_t = \mathbf{M}_t$, and recall that $\tilde{\tau}(x)$ is the thresholded version of $\|\mathbf{U}_t(x - \mu_t)\|_2^2$, as defined in Line 18 and $\tau(x)$ is the same score but using \mathbf{M}_t . The potential-based analysis in Appendix C requires that $\mathbf{E}_{X \sim P}[w_{t+1}(X)\tau_t(X)]$ is small. However, the stopping condition of the Downweighting filter implies only that $\mathbf{E}_{X \sim P}[w_{t+1}(X)\tilde{\tau}_t(X)]$ is small. In Appendix C, the bound on the former was obtained from the bound on the latter by using that $\|\mathbf{U}_t(x - \mu_t)\|_2 \approx \|\mathbf{M}_t(x - \mu_t)\|_2$ pointwise in the support of P (Claim C.10).

By the classical JL lemma, the size of the JL sketch, L, needs to be at most logarithmic in the size of the set S where we require the pointwise approximation to hold. Thus, in the previous section, L scaled as $\log |S| = \log n$. However, in the streaming model where there is no such dataset, it is far from obvious how the analysis should proceed. A näive approach would be to require the approximation to hold on a cover \tilde{S} of the support of P_t . Since $|\tilde{S}|$ scales exponentially with d, the required bound on L would be $\log |\tilde{S}| = \Omega(d)$, which is too large for our purposes. Luckily, we can still find a fixed set S_{cover} such that the following holds: (i) $\log |S_{\text{cover}}| = \text{polylog}(d/\epsilon)$, and (ii) the expectation of scores over $\mathcal{U}(S_{\text{cover}})$ approximates the expectation of scores over P. That is, as far as the expectations of the scores are concerned, P can be approximated by the uniform distribution over S_{cover} . Arguing as before, if $\|\mathbf{U}_t(x-\mu_t)\| \approx \|\mathbf{M}_t(x-\mu_t)\|$ for each $x \in S_{\text{cover}}$, then the downweighting filter also ensures that $\mathbf{E}_{X \sim P}[w_{t+1}(X)\tilde{\tau}_t(X)]$ is small. Thus, S_{cover} can serve as a proxy dataset (used only in the analysis) to ensure that the size of the JL sketch is sufficiently bounded, i.e., that $\log |S_{\text{cover}}| \leq C \text{polylog}(d/\epsilon)$.

Establishing the desired upper bound on the cardinality of S_{cover} requires a somewhat more sophisticated argument that relies on the VC-dimension of a family of functions corresponding to the weight update rule. This result is stated in Appendix D.2.1.

We now present the algorithm more formally. We start by clarifying the notation used.

Notation regarding Algorithm 5: The quantities involved in the algorithm and its analysis now are based on the underlying data distribution P as well as its approximations. We note that P_t , μ_t , Σ_t , \mathbf{B}_t , \mathbf{M}_t , λ_t are functionals of the distribution P and are primarily used in the analysis. The parameters $\widehat{\lambda}_t$, $\widehat{\mathbf{M}}_t$ are approximations for $\|\mathbf{B}_t\|_2$ and \mathbf{M}_t

respectively that the algorithm forms using samples from P_t . Regarding score functions, $g_t(x) = \|\mathbf{M}_t(x - \mu_t)\|_2^2$ is as before. The computations however use only the Johnson-Lindenstrauss versions $\tilde{g}_t(x) := \frac{1}{L} \sum_{i=1}^L (v_{t,i}^T(x - \widehat{\mu}_t))^2$, which can also be written as $\|\mathbf{U}_t(x - \widehat{\mu}_t)\|_2^2$ in matrix form, by defining \mathbf{U}_t to have the vectors $\frac{1}{\sqrt{L}}v_{t,i}$ as its rows. Note that $\tilde{g}_t(x)$ is defined using $\hat{\mu}_t$ instead of μ_t . Finally, we denote by $\tau_t(x) = g_t(x)\mathbb{1}\{g_t(x) > C_3\|\mathbf{M}_t\|_F^2\lambda_t/\epsilon\}$ and $\tilde{\tau}_t(x) = \tilde{g}_t(x)\mathbb{1}\{\tilde{g}_t(x) > C_3\|\mathbf{U}_t\|_F^2\hat{\lambda}_t/\epsilon\}$.

Remark D.4. Recalling Lemma B.12, we may again treat the input distribution as a mixture $P = (1 - \epsilon)G + \epsilon B$, where G is a distribution that is $(C'\epsilon, \delta)$ -stable with respect to μ .

Algorithm 5 Robust Mean Estimation In Single-Pass Streaming Model

```
1: function RobustMeanStreaming(P, \delta, \epsilon, \tau)
          Let R be such that \Pr_{X \sim G}[\|X - \mu\|_2 > R] \le \epsilon.
          Let P = (1 - \epsilon)G + \epsilon B. Without loss of generality, we assume that the points added by the adversary are within
 3:
          O(R) from \mu in Euclidean norm (see Appendix C.1).
          Let K = \Theta(\log d \log(dR/\epsilon).
 4:
          Let L = \text{polylog}(d, R, 1/\epsilon, 1/\tau).
 5:
          Let r = CdR^{2+4\log d}.
 6:
          Obtain a naïve estimation \widehat{\mu} of \mu such that \|\widehat{\mu} - \mu\|_2 \leq 4R.
 7:
          Let w: \mathbb{R}^d \to [0,1] be the weight function. Initialize w_0(x) \leftarrow \mathbb{1}\{\|x-\widehat{\mu}\|_2 \leq 5R\} for all x \in T and \ell_1 \leftarrow 0.
 8:
 9:
          for t \in [K] do
              Define w_t(x) = w_{t-1}(x)(1 - \tilde{\tau}_t(x)/r)^{\ell_t}.
10:
              Let P_t be the distribution of P weighted by w_t, i.e., P_t(x) = P(x)w_t(x)/\mathbf{E}_{X\sim P}[w_t(X)].
11:
              Let \mu_t be the mean of P_t.
12:
13:
              Let \Sigma_t be the covariance matrix of P_t.
              Let \mathbf{B}_t = (\mathbf{E}_{X \sim P}[w_t(X)])^2 \mathbf{\Sigma}_t - \left(1 - C_1 \frac{\delta^2}{\epsilon}\right) \mathbf{I}_d and \mathbf{M}_t = \mathbf{B}_t^{\log d}.
14:
              Compute an O(\delta)-accurate estimator \widehat{\mu}_t of \mu_t (see Lemma D.11).
15:
              Let \tilde{n} = C'' R^2 (\log d)^2 \max\left(d, \frac{\epsilon^2 d}{\delta^4}, \frac{R^2 \epsilon^2}{\delta^2}, \frac{R^2 \epsilon^4}{\delta^6}\right) \log\left(\frac{dK \log d}{\tau}\right).
16:
              For k \in [\log d], let \hat{\mathbf{B}}_{t,k} be an empirical version of \mathbf{B}_t over \tilde{n} independent samples (see Appendix D.3 for more
17:
              details).
              Define \widehat{\mathbf{M}}_t := \prod_{k=1}^{\log d} \widehat{\mathbf{B}}_{t,k}
18:
              Let \lambda_t = \|\mathbf{B}_t\|_2 and an approximation \widehat{\lambda}_t such that \widehat{\lambda}_t/\lambda_t \in [0.8, 1.2].
19:
              if \widehat{\lambda}_t > C_2 \delta^2 / \epsilon then
20:
21:
                  for j \in [L] do
                      z_{t,j} \sim \mathcal{U}(\{\pm 1\}^d).
22:
                      v_{t,j} \leftarrow \widehat{\mathbf{M}}_t z_{t,j}.
23:
                  end for
24:
                  Denote by \mathbf{U}_t the matrix with rows \frac{1}{\sqrt{L}}v_{t,j} for j\in[L].
25:
                 Let \tilde{g}_t(x) = \|\mathbf{U}_t(x - \hat{\mu}_t)\|_2^2 and \tilde{\tau}_t(x) = \tilde{g}_t(x) \mathbb{1}\{\tilde{g}_t(x) > C_3 \|\mathbf{U}_t\|_F^2 \hat{\lambda}_t/\epsilon\}.

\ell_{\max} \leftarrow \left(\frac{dR}{\delta^2/\epsilon}\right)^{\Theta(\log d)}.
26:
27:
                   \ell_t \leftarrow \text{DownweightingFilter}(P, w_t, \tilde{\tau}_t, R, c\hat{\lambda}_t ||\mathbf{U}_t||_F^2, \ell_{\text{max}}).
28:
29:
              end if
          end for
30:
          return an O(\delta) approximation \hat{\mu}_t of the mean \mu_t of the distribution P_t (see Lemma D.11).
31:
32: end function
```

As already mentioned, Algorithm 5 uses two levels of approximation: the first level is approximating the true distributional quantities by taking samples, and the second is preserving the latter quantities using the JL sketch. If both of these approximations are sufficiently accurate, the correctness of Algorithm 5 would follow similarly to Algorithm 3. Of course, the challenge is to ensure that these approximations hold over the entire distribution, while controlling the sample and memory complexity of the algorithm. As we show in Appendix D.2.1, this can be achieved by restricting our attention to a finite set (cover) of sufficiently large cardinality. Thus, the deterministic conditions that we require now also involve

the cover set, which we denote by S_{cover} . The reader may think of S_{cover} as a fixed set, which will be specified later on (Lemma D.9).

Condition D.5 (Deterministic Conditions for Algorithm 5). Let S_{cover} denote the cover of Lemma D.9 for $\epsilon' = \text{poly}(d, R, 1/\epsilon)^{\log d}$. Our condition consists of the following event:

- 1. Estimator $\widehat{\mu}_t$: For all $t \in [K]$, we have that $\|\widehat{\mu}_t \mu_t\|_2 \leq \delta/100$.
- 2. For every $t \in [K]$, if $\|\mathbf{B}_t\|_2 \ge (C_1/2)\delta^2/\epsilon$ and $\mathbf{E}_{X \sim P}[w_t(X)] \ge 1 O(\epsilon)$, we have that:
 - (a) Spectral norm of \mathbf{B}_t : $\widehat{\lambda}_t \in [0.8\lambda_t, 3\lambda_t]$.
 - (b) Frobenius norm: $\|\mathbf{U}_t\|_F^2 \in [0.8\|\mathbf{M}_t\|_F^2, 1.2\|\mathbf{M}_t\|_F^2]$.
 - (c) Scores: $\tilde{g}_t(x) \geq 0.2g_t(x) 0.8(\delta^2/\epsilon^2) \|\mathbf{M}_t\|_F^2$, for all $x \in S_{\text{cover}}$.
- 3. Stopping condition: Let $T_t := c \widehat{\lambda}_t \|\mathbf{U}_t\|_F^2$. For every $w : \mathbb{R}^d \to [0,1]$, the algorithm has access to an estimator f(w) for the quantity $\mathbf{E}_{X \sim P}[w(X)\tilde{\tau}_t(X)]$, such that $\widehat{F}(P) > T_t/2$ whenever $\mathbf{E}_{X \sim P}[w(X)\tilde{\tau}_t(X)] > T_t$. This estimator is accurate when called $O(\log(d)\log(dR/\epsilon))$ times in every iteration $t \in [K]$.

We note that the Item 3 above is needed to evaluate the stopping condition in the downweighting filter. For every $t \in [K]$, the stopping condition is evaluated at most $O(\log(\ell_{\max}))$ times, with $\ell_{\max} = O(dR^{2+\log d}/(\widehat{\lambda}_t \|\mathbf{U}_t\|^2)$ (using Lemma C.6 with $r = C_4 dR^{2+4\log d}$ and $T := O(\widehat{\lambda}_t \|\mathbf{U}_t\|_F^2)$). This means that we require the estimator in Item 3 to be accurate on $O(K\log(d)\log(dR/\epsilon))$ calls.

D.2. Correctness of Algorithm 5

The analysis in this section is along the same lines as that of Appendix C.4. The naïve estimation step of Line 7 is the same as that used in Algorithm 3 (see Appendix H.4).

Given Condition D.5, we first show the correctness of Algorithm 5 and leave the task of establishing Condition D.5 for Appendix D.3. The proof of correctness would largely follow by our work done in Appendix C.4. There are two adjustments needed in these arguments.

The first concerns Lemma C.6, since the algorithm cannot perform exact binary search. Instead, it can use the approximate oracle of Item 3 of Condition D.5, resulting in a multiplicative constant in the final guarantee. For completeness, we prove correctness for this case in Appendix I.1.

Lemma D.6. In the context of Algorithm 5, if $(1 - \epsilon) \mathbf{E}_{X \sim G}[w(X)\tilde{\tau}(X)] \leq T$, $\|\tilde{\tau}\|_{\infty} \leq r$, and $\ell_{\max} > r/T$, then Algorithm 6 modifies the weight function w to w' such that $(i) (1 - \epsilon) \mathbf{E}_{X \sim G}[w(X) - w'(X)] < \epsilon \mathbf{E}_{X \sim B}[w(X) - w'(X)]$, and (ii) upon termination we have $\mathbf{E}_{X \sim P}[w'(X)\tilde{\tau}(X)] \leq 54T$. Furthermore, if the estimator of Line 3 is set to be that of Lemma D.16, the algorithm terminates after $O(\log(\ell_{\max}))$ iterations, each of which uses $O((R^2\epsilon/\delta^2)\log(1/\tau))$ samples, takes O(nd) time and memory $O(\log(1/\tau))$.

The second adjustment is regarding the analog of Lemma C.9, i.e., $\epsilon \mathbf{E}_{X \sim B}[w_{t+1}\tau_t(X)]$ is small (the bound on $\epsilon \mathbf{E}_{X \sim B}[w_{t+1}\tilde{\tau}_t(X)]$ follows from the stopping condition as before). Since the support is unbounded, we use an argument based on a fixed cover to show that the downweighting filter succeeds with the JL-sketch of size L. The statement is given below.

Lemma D.7. Under the deterministic Condition D.5 and the context of Algorithm 5, we have that $\mathbf{E}_{X \sim B}[w_{t+1}(X)\tau_t(X)] \leq 5 \mathbf{E}_{X \sim B}[w_{t+1}(X)\tilde{\tau}_t(X)] + c(\lambda_t/\epsilon)\|\mathbf{M}_t\|_F^2$, where c is of the form C/C_2 with C being a sufficiently large constant and C_2 being the constant used in Algorithm 5.

The next section is dedicated to proving Lemma D.7. Here we just show that Lemma D.7 suffices to prove the analog of Lemma C.9 below.

Lemma D.8. Consider the context of Algorithm 5 and assume that Condition D.5 holds. Then $\epsilon \cdot \mathbf{E}_{X \sim B}[w_{t+1}(X)\tau_t(X)] \leq c'\lambda_t \|\mathbf{M}_t\|_F^2$, for some constant c'' of the form C''/C_2 , where C_2 is the constant used in Line 20 and C' is some absolute constant.

Algorithm 6 Downweighting Filter using Approximate Oracle

```
1: function DownweightingFilter(P, w, \tilde{\tau}, R, T, \ell_{\text{max}})
        r \leftarrow CdR^{2+4\log d}.
 2:
        Denote by f(\ell) be an estimator close to \mathbf{E}_{X \sim P}[w(X)(1 - \tilde{\tau}(X)/r)^{\ell}\tilde{\tau}(X)] (see Lemma D.16 for details).
 3:
        L \leftarrow \{1, 2 \dots, \ell_{\text{max}}\}
 4:
 5:
        while |L| > 2 do
           Let \ell be the element in the middle of L.
 6:
 7:
           if f(\ell) > 9T then
 8:
              Discard all elements smaller than \ell from L.
 9:
10:
              Discard all elements greater than \ell from L.
11:
           end if
        end while
12:
        return any \ell of L satisfying 4T \leq f(\ell) \leq 36T.
13:
    end function
```

Proof. Denoting by c, c', c'' > 0 constants that are all multiples of $1/C_2$, we have the following:

$$\epsilon \mathop{\mathbf{E}}_{X \sim B}[w_{t+1}(X)\tau_t(x)] \le 5\epsilon \mathop{\mathbf{E}}_{X \sim B}[w_{t+1}(X)\tilde{\tau}_t(x)] + c\lambda_t \|\mathbf{M}_t\|_F^2 \le 5c'\lambda_t \|\mathbf{M}_t\|_F^2 + c\lambda_t \|\mathbf{M}_t\|_F^2 \le c''\lambda_t \|\mathbf{M}_t\|_F^2 ,$$

where the first inequality uses Lemma D.7 and the second inequality uses Lemma D.6.

Letting $\phi_t := \operatorname{tr}(\mathbf{M}_t^2)$ denote the potential function, the above result allows us to follow the same steps as in Appendix C.4.2 to prove that $\phi_{t+1} \leq 0.9999\phi_t$ exactly as in Appendix C.4.2. Thus, we get that after K iterations, we have that $\|\mathbf{\Sigma}_t\|_2 \lesssim \delta^2/\epsilon$. Under Item 1 of Condition D.5, we have that the final estimate $\hat{\mu}_t$ satisfies that $\|\hat{\mu}_t - \mu\|_2 = O(\delta)$. This completes the proof of correctness of Algorithm 5.

D.2.1. PROOF OF LEMMA D.7 VIA A COVER ARGUMENT

To outline the idea of proving Lemma D.7, recall the proof in the setting of Appendix C. There, we just required that $\tilde{g}_t(x)/g_t(x) \in [0.8, 1.2]$ for all samples x in our dataset, which can be translated to some relation between $\tilde{\tau}(x)$ and $\tau(x)$. Then, since B was the empirical distribution on ϵn of these points, the desired condition followed. However, in our case we cannot use pointwise relationships, since the distribution B may be continuous and the Johnson-Lindenstrauss argument might not work for the entire \mathbb{R}^d with $\operatorname{polylog}(d)$ vectors. The idea is first to relate $\mathbf{E}_{X \sim B}[w_{t+1}(X)\tau_t(X)]$ to a discrete expectation over N (not too many) points from a fixed set, then use the relationship between $\tilde{\tau}$ and τ for these points, and finally relate that discrete expectation back to $\mathbf{E}_{X \sim B}[w_{t+1}(X)\tilde{\tau}_t(X)]$. The existence of a cover of a small size is stated in the following.

Lemma D.9. Consider the setting of Algorithm 5, where B is the distribution of outliers supported in a ball of radius R around μ . Let $r' := \left(CdR^2 + 1 + C_1\delta^2/\epsilon\right)^{C\log d}$ for sufficiently large constant C. Denote by ϵ the contamination rate and let an arbitrary $\epsilon' \in (0,1)$. There exists a set S_{cover} of $N = \frac{1}{\epsilon'^3}d^4K^2L^2(dR\epsilon/\delta^2)^{O(\log d)}$ points x_1, \ldots, x_N lying in the ball of radius R around μ , such that for all $t \in [K]$, for all choices of the vectors $z_{t,j}$ of Line 22 of Algorithm 5 it holds

$$\left| \sum_{X \sim B} \left[\frac{1}{r'} w_{t+1}(X) \tilde{\tau}_t(X) \right] - \frac{1}{N} \sum_{i=1}^N \frac{1}{r'} w_{t+1}(x_i) \tilde{\tau}_t(x_i) \right| \le \epsilon'$$
and
$$\left| \sum_{X \sim B} \left[\frac{1}{r'} w_{t+1}(X) \tau_t(X) \right] - \frac{1}{N} \sum_{i=1}^N \frac{1}{r'} w_{t+1}(x_i) \tau_t(x_i) \right| \le \epsilon'.$$

We prove this result in Appendix I.2. Here we show how it implies the desired condition, following the proof sketch from the start of this section.

Proof of Lemma D.7. Let $r' := \left(CdR^2 + 1 + C_1\delta^2/\epsilon\right)^{C\log d}$ and $\epsilon' \in (0,1)$. Applying Lemma D.9, let S_{cover} be the corresponding cover of cardinality N. From the guarantee of approximation of \tilde{g}_t for every $x \in S_{\text{cover}}$, we get the following approximation for $\tilde{\tau}_t(x)$ for $x \in S_{\text{cover}}$ (proved in Appendix I.2).

Claim D.10. Let S be the cover of Lemma D.9 with r' and ϵ' as defined above. Suppose that the deterministic condition Condition D.5 holds. If $x \in S_{cover}$, then $\tau_t(x) \leq 5\tilde{\tau}_t(x) + (18C_3 + 12/C_2)(\lambda_t/\epsilon) \|\mathbf{M}_t\|_F^2$, where C_3 and C_2 are the constants used in Algorithm 5.

Using Claim D.10 and Lemma D.9, we obtain the following series of inequalities:

$$\begin{split} & \underset{X \sim B}{\mathbf{E}}[w_{t+1}(X)\tau_{t}(X)] = r' \underset{X \sim B}{\mathbf{E}} \left[\frac{1}{r'} w_{t+1}(X)\tau_{t}(X) \right] \\ & \leq \epsilon' r' + r' \frac{1}{N} \sum_{i=1}^{N} \frac{1}{r'} w_{t+1}(x_{i})\tau_{t}(x_{i}) \\ & = \epsilon' r' + \frac{1}{N} \sum_{i=1}^{N} w_{t+1}(x_{i})\tau_{t}(x_{i}) \\ & \leq \epsilon' r' + (18C_{3} + 12/C_{2})(\lambda_{t}/\epsilon) \|\mathbf{M}_{t}\|_{F}^{2} + 5 \frac{1}{N} \sum_{i=1}^{N} w_{t+1}(x_{i})\tilde{\tau}_{t}(x_{i}) \\ & \leq 6\epsilon' r' + (18C_{3} + 12/C_{2})(\lambda_{t}/\epsilon) \|\mathbf{M}_{t}\|_{F}^{2} + 5 \underset{X \sim B}{\mathbf{E}}[w_{t+1}(X)\tilde{\tau}_{t}(X)] \\ & \leq 5 \underset{X \sim B}{\mathbf{E}}[w_{t+1}(X)\tilde{\tau}_{t}(X)] + (19C_{3} + 12/C_{2})(\lambda_{t}/\epsilon) \|\mathbf{M}_{t}\|_{F}^{2}. \end{split} \tag{using Lemma D.9 for $\tilde{\tau}_{t}$)} \\ & = 5 \underset{X \sim B}{\mathbf{E}}[w_{t+1}(X)\tilde{\tau}_{t}(X)] + (19C_{3} + 12/C_{2})(\lambda_{t}/\epsilon) \|\mathbf{M}_{t}\|_{F}^{2}. \end{aligned} \tag{using the definition of ϵ')}$$

For the last line above, we want to choose ϵ' such that $\epsilon' \leq \frac{C_3 \lambda_t}{\epsilon r'} \|\mathbf{M}_t\|_F^2$. Since $\|\mathbf{M}_t\|_F^2 \geq (C_2 \delta^2/\epsilon)^{2\log d}$ (otherwise the algorithm has already terminated), it suffices to choose an ϵ' that satisfies $\epsilon' \gtrsim \frac{(C_2 \delta^2/\epsilon)^{2\log d}}{\epsilon (CdR^2 + 1 + C_1 \delta^2/\epsilon)^{C\log d}}$. This gives an upper bound on the cardinality of the set S_{cover} , which gives the upper bound on the size of the JL-sketch, i.e., L. We provide explicit calculations in Remark I.3.

D.3. Establishing Condition D.5

Throughout this section, we assume sample access to the distribution P_t . As mentioned earlier, Algorithm 4 can simulate this by drawing a sample x from P, calculating $w_t(x)$ (with poly-logarithmic cost in terms of running time and memory), and rejecting the sample with probability $1 - w_t(x)$. With high probability, rejection sampling can increase the sample complexity by at most a constant factor because $\mathbf{E}[w_t(X)] \geq 1 - O(\epsilon)$ (cf. Claim C.13).

D.3.1. ITEM 1

We establish Item 1 in the following, which is proved in Appendix I.4.

Lemma D.11. In the setting of Algorithm 5, there exist estimators $\widehat{\mu}_t$ such that, with probability at least $1 - \tau$, for all $t \in [K]$ we have that $\|\widehat{\mu}_t - \mu_t\|_2 \le \delta/100$. Furthermore, each $\widehat{\mu}_t$ can be computed on a stream of $n = O\left(\frac{R^2}{\delta^2/\epsilon}\log(K/\tau) + \frac{d(1+\delta^2/\epsilon)}{\delta^2}\log(K/\tau)\right)$ independent samples from P_t , in time $O(nd\log(K/\tau))$ and using memory $O(d\log(K/\tau))$.

D.3.2. ITEMS 2A TO 2C

Given that Item 1 holds, in this section, we show that Items 2a to 2c of Condition D.5 hold with high probability if sufficiently many samples from the underlying corrupted distribution P are drawn. Let the scores $\widehat{g}_t(x) := \|\widehat{\mathbf{M}}_t(x - \mu_t)\|_2^2$, where $\widehat{\mathbf{M}}_t$ is the following sample-based estimator of \mathbf{M}_t :

- 1. Draw a batch S_0 of \tilde{n} samples from P and let the estimate $\widehat{W}_t = \mathbf{E}_{X \sim \mathcal{U}(S_0)}[w_t(X)]$.
- 2. Let P'_t be the distribution of the differences $(X X')/\sqrt{2}$ for two independent $X, X' \sim P_t$.
- 3. Draw $\log d$ batches $S_1, \ldots, S_{\log d}$ of \tilde{n} samples, each from P'_t .
- 4. For $k \in [\log d]$,

(a) Let
$$\widehat{\Sigma}_{t,k} = \frac{1}{\tilde{n}} \sum_{x \in S_k} xx^T$$
.

(b) Let
$$\widehat{\mathbf{B}}_{t,k} = \widehat{W}_t^2 \widehat{\boldsymbol{\Sigma}}_{t,k} - (1 - C_1 \delta^2 / \epsilon) \mathbf{I}_d$$
.

5. Return
$$\widehat{\mathbf{M}}_t = \prod_{k=1}^{\log d} \widehat{\mathbf{B}}_{t,k}$$
.

Remark D.12. Algorithm 5 does not need to calculate or store $\widehat{\mathbf{M}}_t$ because it requires only that we can calculate products of $\widehat{\mathbf{M}}_t$ with vectors z as in Line 22. This operation can be implemented in linear runtime and memory. Given the description of the estimator above, it suffices to show how to multiply $\widehat{\mathbf{\Sigma}}_{t,k}$ by a vector z in linear time and memory. To this end, we observe that $\widehat{\mathbf{\Sigma}}_{t,k}z=\frac{1}{\bar{n}}\sum_{x\in S_k}x(x^Tz)$, thus by calculating the inner product (x^Tz) first, the result can be found in O(nd) time in a streaming fashion.

We will show that Items 2a to 2c of Condition D.5 follow if $\|\widehat{\mathbf{M}}_t - \mathbf{M}_t\|_2 \le 0.01 \min\left(\frac{\delta/\epsilon}{R}, \frac{1}{\sqrt{d}}\right) \|\mathbf{M}_t\|_F$ (cf. Lemma D.15) and $\|\widehat{\mu}_t - \mu_t\|_2 = O(\delta)$ (cf. Lemma D.11).

Lemma D.13. Suppose that the estimators $\widehat{\mathbf{M}}_t$ in Algorithm 5 are defined by the procedure as in Lines 1 to 5 above. Let C be a sufficiently large constant and assume that the dimension is $d = \Omega(1)^5$. Further assume that $\|\widehat{\mu}_t - \mu\| \leq 0.01\delta$. If $\widetilde{n} \geq CR^2(\log d)^2 \max\left(d, \frac{\epsilon^2 d}{\delta^4}, \frac{R^2 \epsilon^2}{\delta^2}, \frac{R^2 \epsilon^4}{\delta^6}\right) \log\left(\frac{K d \log d}{\tau}\right)$, then Items 2a to 2c hold with probability at least $1 - \tau$.

Proof. For now, we will assume that for all $t \in [K]$, $\|\widehat{\mathbf{M}}_t - \mathbf{M}_t\|_2 \le 0.01 \min\left(\frac{\delta/\epsilon}{R}, \frac{1}{\sqrt{d}}\right) \|\mathbf{M}_t\|_F$ with the claimed sample complexity. This follows from Lemma D.15 and a union bound. We will prove each of these conditions separately.

Proof of Item 2c: Denote $T := 0.1\delta/\epsilon$. Fix an iteration $t \in [K]$. We will prove that the conditions hold in the t-th iteration with probability at least $1 - \tau/K$ and then a union bound will conclude the proof.

Define $\widehat{g}_t(x) := \|\mathbf{M}_t(x-\widehat{\mu}_t)\|_2^2$. Since $\|\widehat{\mu}_t-\mu_t\|_2 \le 0.01\delta$, we have the following relation between \widehat{g}_t and $g_t: \|\mathbf{M}_t(x-\widehat{\mu}_t)\|_2^2 \ge 0.5 \|\mathbf{M}_t(x-\mu_t)\|_2^2 - T^2 \|\mathbf{M}_t\|_F^2$, i.e., $\widehat{g}_t(x) \ge 0.5 g_t(x) - T^2 \|\mathbf{M}_t\|_F^2$. We have that $\|\widehat{\mathbf{M}}_t - \mathbf{M}_t\|_2 \le 0.1 (T/R) \|\mathbf{M}_t\|_F$ with probability at least $1 - \tau/K$ (see Lemma D.15). This means that for any point x with $\|x-\widehat{\mu}_t\|_2 \le 2R$, we have $\|\mathbf{M}_t(x-\widehat{\mu}_t)\|_2 \le \|\widehat{\mathbf{M}}_t(x-\widehat{\mu}_t)\|_2 + 0.2T \|\mathbf{M}_t\|_F$, which implies that $\|\mathbf{M}_t(x-\widehat{\mu}_t)\|_2^2 \le 2\|\widehat{\mathbf{M}}_t(x-\widehat{\mu}_t)\|_2^2 + 0.08T^2 \|\mathbf{M}_t\|_F^2$, or equivalently

$$\|\widehat{\mathbf{M}}_t(x - \widehat{\mu}_t)\|_2^2 \ge 0.5\widehat{g}_t(x) - 0.04T^2 \text{tr}(\mathbf{M}_t^2)$$
 (10)

The final step is taking the Johnson-Lindenstrauss sketch of $\widehat{\mathbf{M}}_t$, which gives the matrix \mathbf{U}_t used in the definition of \widetilde{g}_t . By repeating the proof of Lemma C.5 with $\widehat{\mathbf{M}}_t$ in place of \mathbf{M}_t , we get that if $L = C \log((|S_{\text{cover}}| + d)K/\tau)$, then $\widetilde{g}_t(x) \geq 0.8 \|\widehat{\mathbf{M}}_t(x-\widehat{\mu}_t)\|_2$ for all the points in the set S_{cover} (the cover from Lemma D.9). The value used for L in Algorithm 5 satisfies this condition (c.f. Remark I.3). Combining this with Equation (10) and the relation between \widehat{g}_t and g_t , we get that $\widetilde{g}_t(x) \geq 0.4\widehat{g}_t(x) - 0.04T^2 \|\mathbf{M}_t\|_F^2 \geq 0.2g_t(x) - 0.44T^2 \|\mathbf{M}_t\|_F^2$.

Proof of Item 2b: Again, fix a $t \in [K]$. We have that $\|\widehat{\mathbf{M}}_t - \mathbf{M}_t\|_2 \le \frac{0.01}{\sqrt{d}} \|\mathbf{M}_t\|_F$ with probability $1 - \tau/K$ using Lemma D.15. We thus have that $\|\mathbf{M}_t - \widehat{\mathbf{M}}_t\|_F \le \sqrt{d} \|\mathbf{M}_t - \widehat{\mathbf{M}}_t\|_2 \le 0.01 \|\mathbf{M}_t\|_F$, which implies

$$\left| \| \mathbf{M}_t \|_F - \| \widehat{\mathbf{M}}_t \|_F \right| \le 0.01 \| \mathbf{M}_t \|_F$$
 (11)

It is easy to check that this is stronger than what we initially wanted. Indeed, squaring Equation (11) and using $\|\widehat{\mathbf{M}}_t\|_F \le 1.01 \|\mathbf{M}_t\|_F$ gives

$$\|\widehat{\mathbf{M}}_t\|_F^2 \le 2\|\mathbf{M}_t\|_F \|\widehat{\mathbf{M}}_t\|_F - \|\mathbf{M}_t\|_F^2 + (0.01)^2 \|\mathbf{M}_t\|_F^2 < 1.1 \|\mathbf{M}_t\|_F^2$$

which means that $\|\widehat{\mathbf{M}}_t\|_F^2 - \|\mathbf{M}_t\|_F^2 \le 0.1 \|\mathbf{M}_t\|_F^2$. For the other bound, Equation (11) implies

$$\begin{aligned} \|\mathbf{M}_{t}\|_{F}^{2} &\leq 2\|\mathbf{M}_{t}\|_{F}\|\widehat{\mathbf{M}}_{t}\|_{F} - \|\widehat{\mathbf{M}}_{t}\|_{F}^{2} + (0.01)^{2}\|\mathbf{M}_{t}\|_{F}^{2} \\ &\leq 2\|\widehat{\mathbf{M}}_{t}\|_{F}^{2} + 2(0.01)\|\mathbf{M}_{t}\|_{F}\|\widehat{\mathbf{M}}_{t}\|_{F} - \|\widehat{\mathbf{M}}_{t}\|_{F}^{2} + (0.01)^{2}\|\mathbf{M}_{t}\|_{F}^{2} \end{aligned}$$

⁵This is without loss of generality as we could avoid the JL-sketch when d = O(1).

$$< 1.1 \|\mathbf{M}_t\|_F^2$$
,

which means that $\|\mathbf{M}_t\|_F^2 - \|\widehat{\mathbf{M}}_t\|_F^2 \le 0.1 \|\mathbf{M}_t\|_F^2$. Therefore we obtain the following

$$|\|\mathbf{M}_t\|_F^2 - \|\widehat{\mathbf{M}}_t\|_F^2| \le 0.1 \|\mathbf{M}_t\|_F^2.$$
(12)

Finally, the Johnson-Lindenstrauss step is exactly as described in the proof of Item 2c.

Proof of Item 2a: Since we cannot access the same samples twice, the power-iteration algorithm now uses a different dataset in every step. Let the matrix $\widehat{\mathbf{M}}_t$ as in the beginning of Appendix D.3. We have already shown in the previous paragraph that, with probability $1-\tau$, for all $t\in[K]$, $\widehat{\mathbf{M}}_t$ has Frobenius norm close to that of \mathbf{M}_t (Equation (12)). For the rest of the proof, we condition on this event. Consider the algorithm that calculates $v=\widehat{\mathbf{M}}_t w$, where $w\sim\mathcal{N}(0,\mathbf{I}_d)$ (this can be done in the streaming model by multiplying with $\widehat{\mathbf{B}}_{t,k}$ iteratively; moreover this multiplication can be implemented in time $O(\tilde{n}d)$). We claim that the value $\widehat{\lambda}_t=\|v\|_2^{1/\log d}$ satisfies the desired relation. First, we note that with probability 0.9 we have that

$$0.8\operatorname{tr}(\widehat{\mathbf{M}}_{t}^{2}) \le \|v\|_{2}^{2} \le 1.2\operatorname{tr}(\widehat{\mathbf{M}}_{t}^{2}). \tag{13}$$

This is because $||v||_2^2 = \operatorname{tr}(w^T \widehat{\mathbf{M}}_t^T \widehat{\mathbf{M}}_t w) = \operatorname{tr}(ww^T \widehat{\mathbf{M}}_t^T \widehat{\mathbf{M}}_t)$ and $\Pr[0.8d \le ||w||_2^2 \le 1.2d] \le e^{-cd} \le 0.1$ (by Gaussian norm concentration, assuming that the dimension is sufficiently large). Equations (12) and (13) imply that $0.7\operatorname{tr}(\mathbf{M}_t^2) \le ||v||_2^2 \le 1.4\operatorname{tr}(\mathbf{M}_t^2)$. Furthermore, we have that

$$||v||_{2}^{\frac{1}{\log d}} \le (1.4\operatorname{tr}(\mathbf{M}_{t}^{2}))^{\frac{1}{2\log d}} \le \left(1.4d||\mathbf{B}_{t}||_{2}^{2\log d}\right)^{\frac{1}{2\log d}} \le 2||\mathbf{B}_{t}||_{2}.$$
(14)

Similarly, for the lower bound, we have that

$$||v||_{\frac{1}{\log d}}^{\frac{1}{\log d}} \ge (0.7 \operatorname{tr}(\mathbf{M}_t^2))^{\frac{1}{2\log d}} \ge (0.7)^{\frac{1}{2\log d}} ||\mathbf{B}_t||_2 \ge (0.7) ||\mathbf{B}_t||_2,$$
(15)

where in the last inequality we assumed that the dimension is sufficiently large. Putting Equations (12) to (14) together, with probability at least 0.9, we have that $0.7\|\mathbf{B}_t\|_2 \le \|v\|_2^{1/\log d} \le 3\|\mathbf{B}_t\|_2$. By repeating the procedure $O(\log(1/\tau))$ times and taking the median, we boost the probability of failure to τ . By union bound, choosing $\tau = \tau/K$ makes the event hold for all iterations $t \in [K]$.

The remainder of this section is dedicated to showing that $\|\widehat{\mathbf{M}}_t - \mathbf{M}_t\|_2 \leq \min\left(\frac{\delta/\epsilon}{R}, \frac{0.01}{\sqrt{d}}\right) \|\mathbf{M}_t\|_F$. We require the following lemma, which we prove in Appendix I.3. We use $\prod_{i=1}^p \mathbf{B}_i$ is to denote the matrix product $\mathbf{B}_1 \mathbf{B}_2 \cdots \mathbf{B}_p$.

Lemma D.14. Let $\mathbf{A}, \mathbf{B}, \mathbf{B}_1, \dots, \mathbf{B}_p$ be symmetric $d \times d$ matrices and define $\mathbf{M} = \mathbf{B}^p, \mathbf{M}_S = \prod_{i=1}^p \mathbf{B}_i$. If $\|\mathbf{B}_i - \mathbf{B}\|_2 \le \delta \|\mathbf{B}\|_2$, then $\|\mathbf{M}_S - \mathbf{B}^p\|_2 \le p\delta(1+\delta)^p \|\mathbf{B}\|_2^p$.

We are now ready to prove our main technical result.

Lemma D.15. Assume that $\|\mathbf{B}_t\|_2 \ge (C_1/2)\delta^2/\epsilon$ and $\mathbf{E}_{X\sim P}[w_t(X)] \ge 1 - O(\epsilon)$ hold in the t-th iteration of Algorithm 5. If \widehat{W} and every $\widehat{\mathbf{B}}_{t,k}$ in the product $\widehat{\mathbf{M}}_t = \prod_{k=1}^{\log d} \widehat{\mathbf{B}}_{t,k}$ is calculated using

$$\tilde{n} \geq CR^2 (\log d)^2 \max\left(d, \frac{\epsilon^2 d}{\delta^4}, \frac{R^2 \epsilon^2}{\delta^2}, \frac{R^2 \epsilon^4}{\delta^6}\right) \log\left(\frac{d \log d}{\tau}\right)$$

samples, where C is a sufficiently large constant, we have that

$$\|\widehat{\mathbf{M}}_t - \mathbf{M}_t\|_2 \le 0.01 \min\left(\frac{\delta/\epsilon}{R}, \frac{1}{\sqrt{d}}\right) \|\mathbf{M}_t\|_F$$

with probability at least $1-\tau$.

Proof. Let $T := \delta/\epsilon$ and $p := \log d$ for brevity. Using Lemma D.14 we have that $\|\widehat{\mathbf{M}}_t - \mathbf{M}_t\|_2 \le p\gamma e^{\gamma p} \|\mathbf{M}_t\|_2$, where $\gamma > 0$ is such that

$$\|\widehat{\mathbf{B}}_{t,k} - \mathbf{B}_t\|_2 \le \gamma \|\mathbf{B}_t\|_2 \tag{16}$$

for all $k \in [p]$. Therefore, for the lemma to hold, it suffices that $p\gamma e^{\gamma p} \leq 0.01 \min\left(\frac{T\|\mathbf{M}_t\|_F}{R\|\mathbf{M}_t\|_2}, \frac{1}{\sqrt{d}}\right)$. For that, it suffices to choose $\gamma = \frac{0.01}{3p} \min\left(\frac{1}{\sqrt{d}}, \frac{T\|\mathbf{M}_t\|_F}{R\|\mathbf{M}_t\|_2}\right)$. At this point, we also assume two things: First, that the estimate $\widehat{\boldsymbol{\Sigma}}_{t,k}$ (defined in step 4a) is such that $\|\widehat{\boldsymbol{\Sigma}}_{t,k} - \boldsymbol{\Sigma}_t\|_2 \leq \epsilon' \|\boldsymbol{\Sigma}_t\|_2$ for some ϵ' to be specified later on. Second, that we have an estimate \widehat{W}_t for $\mathbf{E}_{X\sim P}[w_t(X)]$ such that

$$\widehat{W}_t = \underset{X \sim P}{\mathbf{E}}[w_t(X)] + \eta , \qquad (17)$$

with $|\eta| \le \xi$ for some $\xi \le 1$ to be decided later. By Hoeffding's inequality, if we compute \widehat{W} as shown in Step 1, then $\log(2/\tau)/\xi^2$ samples suffice to guarantee that Equation (17) holds with probability $1-\tau/2$. We now focus on Equation (16). We note that

$$\|\widehat{\mathbf{B}}_{t,k} - \mathbf{B}_{t}\|_{2} = \left\| \left(\sum_{X \sim P_{t}} [w_{t}(X)] + \eta \right)^{2} \widehat{\mathbf{\Sigma}}_{t,k} - \sum_{X \sim P_{t}} [w_{t}(X)]^{2} \mathbf{\Sigma}_{t} \right\|_{2}$$

$$\leq \sum_{X \sim P_{t}} [w_{t}(X)]^{2} \|\widehat{\mathbf{\Sigma}}_{t,k} - \mathbf{\Sigma}_{t}\|_{2} + (\eta^{2} + 2\eta \sum_{X \sim P_{t}} [w_{t}(X)]) \|\widehat{\mathbf{\Sigma}}_{t,k}\|_{2}$$

$$\leq \|\widehat{\mathbf{\Sigma}}_{t,k} - \mathbf{\Sigma}_{t}\|_{2} + 3\xi (\|\widehat{\mathbf{\Sigma}}_{t,k} - \mathbf{\Sigma}\|_{2} + \|\mathbf{\Sigma}_{t}\|_{2})$$

$$= (1 + 3\xi) \|\widehat{\mathbf{\Sigma}}_{t,k} - \mathbf{\Sigma}_{t}\|_{2} + 3\xi \|\mathbf{\Sigma}_{t}\|_{2}.$$

By choosing $\xi = \min(1, \epsilon'/3)$ and $\epsilon' = \frac{1}{5}\gamma \|\mathbf{B}_t\|_2/\|\mathbf{\Sigma}_t\|_2$, the above implies that Equation (16) holds. Thus, it suffices to show that $\|\widehat{\mathbf{\Sigma}}_{t,k} - \mathbf{\Sigma}_t\|_2 \le \epsilon' \|\mathbf{\Sigma}_t\|_2$ for our choice of ϵ' . Note that Fact B.5 is not directly applicable to the distribution P_t since it does not have zero mean. This is why we are working with samples of the form $(X - X')/\sqrt{2}$. By Fact B.5 with ϵ' set as above and $\tau = \tau/(2p)$, we have the following upper bound on the sufficient number of samples:

$$\tilde{n} = C \frac{R^{2}}{\epsilon^{\prime 2} \| \mathbf{\Sigma}_{t} \|_{2}} \log \left(\frac{2pd}{\tau} \right)$$

$$\lesssim \frac{R^{2}}{\gamma^{2}} \frac{\| \mathbf{\Sigma}_{t} \|_{2}}{\| \mathbf{B}_{t} \|_{2}^{2}} \log \left(\frac{pd}{\tau} \right)$$

$$\lesssim \frac{R^{2} \epsilon}{\delta^{2} \gamma^{2}} \frac{\| \mathbf{\Sigma}_{t} \|_{2}}{\| \mathbf{B}_{t} \|_{2}} \log \left(\frac{pd}{\tau} \right)$$

$$\lesssim \frac{R^{2} \epsilon}{\delta^{2} \gamma^{2}} \max \left(1, \frac{\epsilon}{\delta^{2}} \right) \log \left(\frac{pd}{\tau} \right)$$

$$\lesssim \frac{R^{2} \epsilon}{\delta^{2}} \max \left(d, \frac{R^{2} \epsilon^{2}}{\delta^{2}} \frac{\| \mathbf{M}_{t} \|_{2}^{2}}{\| \mathbf{M}_{t} \|_{F}^{2}} \right) \max \left(1, \frac{\epsilon}{\delta^{2}} \right) \log \left(\frac{pd}{\tau} \right)$$

$$\leq \frac{R^{2} p^{2} \epsilon}{\delta^{2}} \max \left(d, \frac{R^{2} \epsilon^{2}}{\delta^{2}} \frac{\| \mathbf{M}_{t} \|_{F}^{2}}{\| \mathbf{M}_{t} \|_{F}^{2}} \right) \max \left(1, \frac{\epsilon}{\delta^{2}} \right) \log \left(\frac{pd}{\tau} \right)$$

$$\leq \frac{R^{2} p^{2} \epsilon}{\delta^{2}} \max \left(d, \frac{R^{2} \epsilon^{2}}{\delta^{2}} \right) \max \left(1, \frac{\epsilon}{\delta^{2}} \right) \log \left(\frac{pd}{\tau} \right)$$

$$\leq \frac{R^{2} p^{2} \epsilon}{\delta^{2}} \max \left(d, \frac{\epsilon d}{\delta^{2}}, \frac{R^{2} \epsilon^{2}}{\delta^{2}}, \frac{R^{2} \epsilon^{3}}{\delta^{4}} \right) \log \left(\frac{pd}{\tau} \right),$$
(19)

Equation (18) is derived as follows: First we note that $\|\mathbf{\Sigma}_t\|_2 \leq \frac{\|\mathbf{B}_t\|_2 + 1}{\mathbf{E}_{X \sim P}[w_t(X)]^2} \lesssim \|\mathbf{B}_t\|_2 + 1$, where the last inequality uses our assumption that $\mathbf{E}_{X \sim P}[w_t(X)] \geq 1 - O(\epsilon)$. We combine this with $\|\mathbf{B}_t\|_2 \gtrsim \delta^2/\epsilon$ as follows:

$$\frac{\|\mathbf{\Sigma}_t\|_2^2}{\|\mathbf{B}_t\|_2^2} \lesssim \frac{(\|\mathbf{B}_t\|_2 + 1)^2}{\|\mathbf{B}_t\|_2^2} \leq 2 + \frac{2}{\|\mathbf{B}_t\|_2^2} \lesssim 1 + \frac{1}{(\delta^2/\epsilon)^2} \;.$$

Regarding the samples required to achieve Equation (17), a sufficient number is

$$\begin{split} \tilde{n} &= C \log \left(\frac{2}{\tau}\right) \frac{1}{\xi^2} \\ &\lesssim \log \left(\frac{1}{\tau}\right) \max \left(1, \frac{1}{\epsilon'^2}\right) \\ &\lesssim \log \left(\frac{1}{\tau}\right) \max \left(1, \frac{1}{\gamma^2} \frac{\|\mathbf{\Sigma}_t\|_2^2}{\|\mathbf{B}_t\|_2^2}\right) \\ &\lesssim \log \left(\frac{1}{\tau}\right) \max \left(1, \frac{1}{\gamma^2} \max \left(1, \frac{\epsilon^2}{\delta^4}\right)\right) \\ &\lesssim \log \left(\frac{1}{\tau}\right) \max \left(1, p^2 \max \left(d, \frac{R^2 \epsilon^2}{\delta^2}\right) \max \left(1, \frac{\epsilon^2}{\delta^4}\right)\right) \\ &\lesssim \log \left(\frac{1}{\tau}\right) \max \left(p^2 d, \frac{p^2 R^2 \epsilon^2}{\delta^2}, \frac{p^2 d \epsilon^2}{\delta^4}, \frac{p^2 R^2 \epsilon^4}{\delta^6}\right) \;, \end{split}$$

which is smaller compared to the right-hand side of Equation (19).

D.3.3. ITEM 3

The following lemma establishes that the estimator of Item 3 is accurate when called once. By using a union bound on the maximum number of times that it can be called, we get the sample complexity requirement of $n = O\left((R^2/(\delta^2/\epsilon))\operatorname{polylog}\left(d,R,\frac{1}{\epsilon},\frac{1}{\tau}\right)\right)$.

Lemma D.16. Consider the context of Algorithm 5 and denote $T_t := c\widehat{\lambda}_t \|\mathbf{U}_t\|_F^2$. Given a weight function $w : \mathbb{R}^d \to [0,1]$, there exists an estimator f(w) on $n = O(\frac{R^2}{\delta^2/\epsilon} \log(1/\tau))$ samples such that, if $\mathbf{E}_{X \sim P}[w(X)\tilde{\tau}_t(X)] > T_t$, then with probability at least $1 - \tau$, $f > T_t/2$. Similarly, $\mathbf{E}_{X \sim P}[w(X)\tilde{\tau}_t(X)] < T_t$ implies $f < (3/2)T_t$. Moreover, the estimator uses $O(\log(1/\tau))$ memory and runs in O(nd) time.

Proof. We show the first direction; the other one has a symmetric proof. Suppose $\mathbf{E}_{X\sim P}[w(X)\tilde{\tau}_t(X)] > c\hat{\lambda}_t \|\mathbf{U}_t\|_F^2$. It suffices to show that with probability at least 0.9 we have that

$$\frac{1}{n} \sum_{i=1}^{N} w(X_i) \tilde{\tau}_t(X_i) > \frac{3}{4} \sum_{X \sim P} [w(X) \tilde{\tau}_t(X)] - \frac{1}{4} c \hat{\lambda}_t \| \mathbf{U}_t \|_F^2 , \qquad (20)$$

as we can repeat the procedure $O(\log(1/\tau))$ times and take the majority vote to boost the probability to $1-\tau$. By Chebyshev's inequality, we have that with probability 0.9 it holds that

$$\frac{1}{n} \sum_{i=1}^{n} w(X_i) \tilde{\tau}_t(X_i) > \underset{X \sim P}{\mathbf{E}}[w(X) \tilde{\tau}_t(X)] - \sqrt{\frac{10 \operatorname{Var}_{X \sim P}(w(X) \tilde{\tau}(X))}{n}}.$$

Therefore, it suffices to have $\sqrt{\frac{10\operatorname{Var}_{X\sim P}(w(X)\tilde{\tau}(X))}{n}} \leq \frac{1}{4}\operatorname{\mathbf{E}}_{X\sim P}[w(X)\tilde{\tau}_t(X)] + \frac{1}{4}c\widehat{\lambda}_t\|\mathbf{U}_t\|_F^2$, and thus we need n to be a sufficiently large multiple of $\operatorname{Var}_{X\sim P}(w(X)\tilde{\tau}_t(X))/(\mathbf{E}_{X\sim P}[w(X)\tilde{\tau}_t(X)] + c\widehat{\lambda}_t\|\mathbf{U}_t\|_F^2)^2$. For that, it suffices to choose

$$n = \Theta\left(\frac{\operatorname{Var}_{X \sim P}(w(X)\tilde{\tau}_t(X))}{\mathbf{E}_{X \sim P}[w(X)\tilde{\tau}_t(X)]c\hat{\lambda}_t \|\mathbf{U}_t\|_F^2}\right).$$

We now focus on bounding by above the right-hand side. Let $T'_t := C_3 \hat{\lambda}_t \|\mathbf{U}_t\|_F^2$ be the threshold used in the definition of $\tilde{\tau}_t(x) = \tilde{g}_t(x) \mathbb{1}\{\tilde{g}_t(x) \geq T'_t\}$. For the variance we have that

$$\operatorname{Var}(w(X)\tilde{\tau}_{t}(X)) \leq \underset{X \sim P}{\mathbf{E}}[((w(X)\tilde{\tau}_{t}(X))^{2}]$$

$$\leq \underset{X \sim P}{\mathbf{E}}[w(X)\tilde{\tau}_{t}^{2}(X)]$$

$$\lesssim R^2 \|\mathbf{U}_t\|_F^2 \sum_{X \sim P} [w(X)\tilde{\tau}_t(X)] , \qquad (21)$$

where the last inequality uses that $\mathbf{E}_{X \sim P_t}[\tilde{\tau}_t^2(X)] = \mathbf{E}_{X \sim P_t}[\tilde{g}_t^2(X)\mathbb{1}\{\tilde{g}_t(X) \geq T_t'\}]$ and bounds from above the one of the two factors of \tilde{g}_t as follows:

$$\tilde{g}_t(x) = \|\mathbf{U}_t(x - \mu_t)\|_2^2 \le \|\mathbf{U}_t\|_F^2 R^2 , \qquad (22)$$

where U_t is the matrix used in Line 26 of the algorithm. Using Equation (21), the number of samples that suffice can now be bounded as follows:

$$\frac{\operatorname{Var}_{X \sim P}(\tilde{\tau}_t(X))}{\widehat{\lambda}_t \operatorname{\mathbf{E}}_{X \sim P}[w(X)\tilde{\tau}_t(X)] \|\mathbf{U}_t\|_F^2} \lesssim \frac{R^2 \|\mathbf{U}_t\|_F^2}{\widehat{\lambda}_t \|\mathbf{U}_t\|_F^2} \lesssim \frac{R^2}{\delta^2/\epsilon} \;,$$

where we used that $\hat{\lambda}_t > C_2 \delta^2 / \epsilon$ from Line 20 of our algorithm.

E. Applications: Beyond Robust Mean Estimation

In this section, we develop robust streaming algorithms with near-optimal space complexity for more complex statistical tasks, specifically for robust covariance estimation and robust stochastic optimization. The main idea enabling these applications is that these tasks can be effectively reduced to robust mean estimation.

E.1. Robust Covariance Estimation

In this subsection, we study the problem of estimating the covariance matrix Σ of a distribution D, having access to ϵ -corrupted samples from D in the sense of Definition A.2. Let $X \sim D$ and the Kronecker product $Y = X \otimes X$. Note that $\mathbf{E}[Y] = \Sigma^{\flat}$, where \flat denotes the flattening operation. Then, using any robust mean estimation algorithm on this d^2 -dimensional distribution, one efficiently compute a vector close to Σ^{\flat} in ℓ_2 -norm, which translates to a Frobenius-norm guarantee for Σ . Of course, our mean estimator works as long as the distribution of Y is stable. If $\mathbf{Cov}[Y]$ is bounded from above by a multiple of the identity matrix, then Y is $(\epsilon, O(\sqrt{\epsilon}))$ -stable with respect to Σ^{\flat} , and thus we get the following as a corollary of Theorem D.2:

Theorem E.1 (Robust Covariance Estimation for Distributions with Bounded Moments). Let a distribution D with $\mathbf{Cov}_{X \sim D}[X \otimes X] \preceq \mathbf{I}_{d^2}$ and denote by Σ its covariance matrix. Let $d \in \mathbb{Z}_+$, $0 < \tau < 1$ and $0 < \epsilon < \epsilon_0$ for a sufficiently small constant ϵ_0 . There exists an algorithm that given ϵ, τ and a set of $n = (d^4/\epsilon) \operatorname{polylog}(d, 1/\epsilon, 1/\tau)$ samples in the single-pass streaming model of Definition A.1 from a distribution Q with $d_{\mathrm{TV}}(D, Q) \leq \epsilon$, runs in time $nd^2 \operatorname{polylog}(d, 1/\epsilon, 1/\tau)$, uses memory $d^2 \operatorname{polylog}(d, 1/\epsilon, 1/\tau)$, and outputs a matrix $\widehat{\Sigma}$ such that $\|\widehat{\Sigma} - \Sigma\|_F = O(\sqrt{\epsilon})$, with probability at least $1 - \tau$.

For the special case when D is Gaussian we have that the fourth moment tensor of D is bounded:

Fact E.2 (see, e.g., (Cheng et al., 2019b)). Let
$$X \sim \mathcal{N}(0, \Sigma)$$
 with $\Sigma \leq \mathbf{I}_d$ and $Y = X \otimes X$. Then, $\mathbf{Cov}[Y] \leq 2\mathbf{I}_{d^2}$.

Using the above fact, we have that the guarantees of Theorem E.1 hold in the Gaussian case, giving an algorithm for $O(\sqrt{\epsilon})$ -approximation in Frobenius norm. However, the information-theoretic lower bound for covariance estimation of the Gaussian distribution is of the order of ϵ . We can plug-in our streaming robust mean estimation algorithm to the covariance estimator given in (Cheng et al., 2019b), and achieve the nearly-optimal error of $O(\epsilon \log(1/\epsilon))$. This algorithm creates a series of estimates $\widehat{\Sigma}_i$. At the (i+1)-th step, all samples are multiplied by $\widehat{\Sigma}_i^{-1/2}$ thus, given that $\widehat{\Sigma}_i$ is a good approximation for Σ , this makes the distribution of the transformed samples closer to $\mathcal{N}(0, \mathbf{I}_d)$, which in turn allows us to produce a better approximation $\widehat{\Sigma}_{i+1}$ of Σ . The resulting guarantees are summarized in the following theorem.

Theorem E.3 (Robust Gaussian Covariance Estimation). Let Q be a distribution on \mathbb{R}^d with $d_{\mathrm{TV}}(Q, \mathcal{N}(0, \Sigma)) \leq \epsilon$ and assume that $\frac{1}{\kappa}\mathbf{I}_d \preceq \Sigma \preceq \mathbf{I}_d$, for some $\kappa > 0$. There is a single-pass streaming algorithm that uses $n = (d^4/\epsilon^2)\mathrm{polylog}(d, \kappa, 1/\epsilon, 1/\tau)$ samples from Q, runs in time $nd^2\mathrm{polylog}(d, \kappa, 1/\epsilon, 1/\tau)$, uses memory $d^2\mathrm{polylog}(d, \kappa, 1/\epsilon, 1/\tau)$, and outputs a matrix $\widehat{\Sigma}$ such that $\|\Sigma^{-1/2}\widehat{\Sigma}\Sigma^{-1/2} - \mathbf{I}_d\|_F = O(\epsilon \log(1/\epsilon))$, with probability at least $1 - \tau$.

The reader is referred to Appendix K for more details on using Algorithm 5 to obtain Theorem E.3.

Algorithm 7 Robust Gradient Descent

- 1: **Input:** $q(\cdot)$, τ
- 2: **for** t = 0 **to** T 1 **do**
- 3: $\theta^{t+1} = \arg\min_{\theta \in \Theta} \|\theta^t \eta g(\theta)\|_2^2$
- 4: end for

E.2. Stochastic Convex Optimization

Here we explore the implications of Algorithm 5 in outlier-robust stochastic convex optimization. This subsection crucially leverages the prior works (Prasad et al., 2020; Diakonikolas et al., 2019a), which apply robust mean estimation algorithms to perform robust stochastic optimization. In particular, we follow the framework of (Prasad et al., 2020).

Concretely, we study the following generic optimization problem: Let a parameter space Θ , sample space \mathcal{Z} , and a loss function $f(\theta;z):\Theta\times\mathcal{Z}\to\mathbb{R}^+$. For an unknown distribution D over \mathcal{Z} , the goal is to minimize the associated *risk* $\bar{f}(\theta)=\mathbf{E}_{z\sim D}[f(\theta;z)]$, given sample access to the distribution D. We will occasionally just write $f(\theta)$ instead of $f(\theta;z)$ when no confusion arises. This setup is central in machine learning, since it captures a plethora of learning tasks. For example, f can be a negative log-likelihood function for the learning problem of interest, e.g., square loss for linear regression and logistic loss for logistic regression. In the robust version of the problem, the algorithm has access only to an ϵ -corrupted version of D in the sense of Definition A.2.

We start by recalling a generic optimization algorithm that works whenever \bar{f} is τ_{ℓ} -strongly convex and τ_{u} -smooth, i.e., for all $\theta_{1}, \theta_{2} \in \Theta$, we have that

$$\frac{\tau_{\ell}}{2} \|\theta_1 - \theta_2\|_2^2 \le \bar{f}(\theta_1) - \bar{f}(\theta_2) - (\nabla \bar{f}(\theta_2))^T (\theta_1 - \theta_2) \le \frac{\tau_u}{2} \|\theta_1 - \theta_2\|_2^2.$$

We then give specific applications for robust linear regression and logistic regression.

The work of (Prasad et al., 2020) provides an analysis of projected gradient descent assuming oracle access to approximations of the gradient:

Definition E.4 $((\alpha, \beta)$ -gradient estimator). A function $g(\theta)$ is an (α, β) -gradient estimator for \bar{f} if $||g(\theta) - \nabla \bar{f}(\theta)||_2 \le \alpha ||\theta - \theta^*||_2 + \beta$, for every $\theta \in \Theta$.

Denoting by η the step size of gradient descent, define the following parameter:

$$\kappa := \sqrt{1 - \frac{2\eta \tau_{\ell} \tau_{u}}{\tau_{\ell} + \tau_{u}}} + \eta \alpha . \tag{23}$$

Theorem E.5 ((Prasad et al., 2020)). Let the domains $\Theta, \mathcal{Z} \subset \mathbb{R}^d$, a distribution D over \mathcal{Z} , and a loss function $f: \Theta \times \mathcal{Z} \to \mathbb{R}^+$ such that $\bar{f}(\theta) := \mathbf{E}_{z \sim D}[f(\theta; z)]$ is τ_ℓ -strongly convex and τ_u -smooth. Let g be an (α, β) -gradient estimator with $\alpha < \tau_\ell$. Let κ from Equation (23) and θ^* be the minimizer of \bar{f} . Then Algorithm 7, initialized at θ^0 with step size $\eta = 2/(\tau_\ell + \tau_u)$, after

$$T = \log_{\frac{1}{\kappa}} \left(\frac{(1-\kappa)\|\theta^0 - \theta^*\|_2}{\beta} \right) \tag{24}$$

iterations, returns a vector $\widehat{\theta}$ such that

$$\|\widehat{\theta} - \theta^*\|_2 \le \frac{2}{1 - \kappa} \beta. \tag{25}$$

If the distribution of the gradients has bounded covariance, then one can use the low-memory estimator of the previous sections in place of $g(\cdot)$. This bound on the covariance will not necessarily be known to the algorithm, thus we first need to strengthen the robust mean estimator so that it is adaptive to that unknown scale. This can be done using Lepski's method (Lepskii, 1991; Birgé, 2001) (the details are deferred to Appendix J). Having that version of the estimator at hand, we then obtain the following statement (see Appendix K for the proof):

Corollary E.6. In the setting of Theorem E.5, suppose that the distribution of gradients satisfies $\mathbf{Cov}[\nabla f(\theta)] \leq \sigma^2 \mathbf{I}_d$ with $\sigma^2 = \alpha^2 \|\theta - \theta^*\|_2^2 + \beta^2$ for all $\theta \in \Theta$, where $\alpha \sqrt{\epsilon} < \tau_\ell$. Assume that the radius of the domain Θ , $r := \max_{\theta \in \Theta} \|\theta\|_2$ is finite. There exists a single-pass streaming algorithm that given $O(T(d^2/\epsilon)\log(1 + \alpha r/\beta)\operatorname{polylog}(d, 1/\epsilon, T/\tau, 1 + \alpha r/\beta))$ samples, runs in time Tnd $\operatorname{polylog}(d, 1/\epsilon, T/\tau, 1 + \alpha r/\beta)$, uses memory d $\operatorname{polylog}(d, 1/\epsilon, T/\tau, 1 + \alpha r/\beta)$, and returns a vector $\widehat{\theta} \in \mathbb{R}^d$ such that $\|\widehat{\theta} - \theta^*\|_2 = O(\sqrt{\epsilon}\beta/(1 - \kappa))$ with probability at least $1 - \tau$.

We now proceed to more specific applications, where we work out the parameters α , β for some distributions of interest.

E.2.1. LINEAR REGRESSION

For linear regression, we assume the following generative model:

$$Y = X^T \theta^* + Z \,, \tag{26}$$

where $\theta^* \in \mathbb{R}^d$ belongs in the ball $\|\theta^*\|_2 \le r, X \sim D_x, Z \sim D_Z$ independently, and D_Z has zero mean. The loss function that we use in this case is $f(\theta) = \frac{1}{2}(Y - \theta^T X)^2$, and the risk function is

$$\bar{f}(\theta) = \underset{(X,Y)}{\mathbf{E}} [f(\theta)] = \frac{1}{2} (\theta - \theta^*)^T \underset{X \sim D_x}{\mathbf{E}} [XX^T] (\theta - \theta^*) + \frac{1}{2} \operatorname{Var}(Z) .$$

Letting $\lambda_{\max}(\mathbf{E}[XX^T])$ and $\lambda_{\min}(\mathbf{E}[XX^T])$ denote the largest and smallest eigenvalue of $\mathbf{E}[XX^T]$ respectively, it can be checked that for any $\tau_\ell \leq \lambda_{\min}(\mathbf{E}[XX^T])$ and $\tau_u \geq \lambda_{\max}(\mathbf{E}[XX^T])$, \bar{f} is τ_ℓ -strongly convex and τ_ℓ -smooth.

Since we want the distribution of gradients to be stable, we impose the following sufficient conditions on the distributions D_x and D_Z .

Assumption E.7. The random variables X, Z are independent and satisfy the following conditions:

- 1. $\mathbf{E}_{Z \sim D_Z}[Z] = 0$
- 2. $\operatorname{Var}_{Z \sim D_Z}[Z] \leq \xi^2$
- 3. $\gamma \mathbf{I}_d \leq \mathbf{E}_{X \sim D_x}[XX^T] \leq \sigma^2 \mathbf{I}_d$.
- 4. For some constant C > 0, for every $v \in \mathcal{S}^{d-1}$, $\mathbf{E}_{X \sim D_x}[(X^T v)^4] \leq C\sigma^4$.

As shown below, these assumptions imply that the resulting distribution of the gradients has bounded covariance (and thus is stable with respect to its mean).

Lemma E.8 (see, e.g., (Diakonikolas et al., 2019a)). For D_x , D_Z satisfying Assumption E.7, for every $\theta \in \Theta$, we have that $\mathbf{Cov}[\nabla f(\theta)] \leq (4\sigma^2 \xi^2 + 4C\sigma^4 \|\theta - \theta^*\|_2^2)\mathbf{I}_d$.

Having Lemma E.8 in hand, Corollary E.6 gives the following.

Theorem E.9 (Robust Linear Regression; full version of Theorem A.5). Consider the linear regression model of Equation (26) and suppose that Assumption E.7 holds. Let $0 < \epsilon < \epsilon_0$ for a sufficiently small constant ϵ_0 . Assume that $C\sigma^2\sqrt{\epsilon} < \gamma/2$. Let κ, T as in Equations (23) and (24) with $\tau_\ell = \gamma$, $\tau_u = \sigma^2$. There is an algorithm that uses $n = T \cdot (d^2/\epsilon) \log(1 + r\sigma/\xi)$ polylog $(d, 1/\epsilon, T/\tau, 1 + r\sigma/\xi)$ samples, runs in time Tnd polylog $(d, 1/\epsilon, T/\tau, 1 + r\sigma/\xi)$, uses memory d polylog $(d, 1/\epsilon, T/\tau, 1 + r\sigma/\xi)$, and returns a vector $\hat{\theta} \in \mathbb{R}^d$ such that $||\hat{\theta} - \theta^*||_2 = O(\sigma \xi \sqrt{\epsilon}/(1 - \kappa))$ with probability at least $1 - \tau$.

Proof. In our case, we have that $\tau_{\ell} = \gamma$ and $\tau_{u} = \sigma^{2}$. Given the bound of Lemma E.8, we use Corollary E.6 with $\alpha = 2C\sigma^{2}$ and $\beta = 2\sigma\xi$. The requirement from that corollary that $\alpha\sqrt{\epsilon} \leq \tau_{\ell}$ becomes $C\sigma^{2}\sqrt{\epsilon} < \gamma/2$. Moreover, $\alpha r/\beta = O(r\sigma/\xi)$.

E.2.2. LOGISTIC REGRESSION

We consider the joint distribution of $X \in \mathbb{R}^d$, $Y \in \{0,1\}$, where $X \sim D_x$ and Y given X is Bernoulli random variable:

$$Y|X \sim \text{Bernoulli}(p), \quad \text{with } p = \frac{1}{1 + e^{-x^T \theta^*}}.$$
 (27)

The loss function we are minimizing in this case is the negative log-likelihood, which eventually can be written as $f(\theta) = -(\theta^T x)y + \Phi(\theta^T x)$, where $\Phi(t) := \log(1 + e^t)$. Regarding the strong convexity parameters, the Hessian of \bar{f} can be shown to be

$$\nabla^2 \bar{f}(\theta) = \mathbf{E}_{X \sim D_x} \left[\frac{e^{\theta^T X}}{(1 + e^{\theta^T X})^2} X X^T \right] . \tag{28}$$

The parameter space Θ needs to be bounded in order for the eigenvalues of the Hessian to remain away from zero; we thus use $\Theta = \{\theta \in \mathbb{R}^d : \|\theta\|_2^2 \le r\}$ with r > 0 being a universal constant. We also impose the following assumptions on the covariates.

Assumption E.10. We assume the following for the distribution of X:

- 1. $\mathbf{E}[X] = 0$.
- 2. (concentration) For some constant C > 0, $\mathbf{E}[XX^T] \leq C^2\mathbf{I}_d$.
- 3. (anti-concentration) There exists constant $c_1 > 0$ and $c_2 \in (0,1/2)$ such that for every unit vector v, $\Pr_{X \sim D_x}[(v^T X)^2 > c_1 ||v||_2^2] \ge c_2$.

Under these assumptions, we have the following:

Lemma E.11 (Lemma 4 in (Prasad et al., 2020)). Supposing that Assumption E.10 holds, for every $\theta \in \Theta$, we have that $\mathbf{Cov}[\nabla f(\theta)] \leq O(1)\mathbf{I}_d$.

The above lemma shows that the distribution of $\nabla f(\theta)$ is $(\epsilon, O(\sqrt{\epsilon}))$ -stable, and thus using our robust mean estimation algorithm one can get an (α, β) -gradient estimator with $\alpha = 0$ and $\beta = O(\sqrt{\epsilon})$. This proves the following (see Appendix K for a detailed proof):

Theorem E.12 (Robust Logistic Regression; full version of Theorem A.6). Consider the logistic regression model of Equation (27) with the domain Θ of the unknown regressor being the ball of radius r, for some universal constant r > 0, and suppose that Assumption E.10 holds. Assume that $0 < \epsilon < \epsilon_0$ for a sufficiently small constant ϵ_0 . There is a single-pass streaming algorithm that uses $n = (d^2/\epsilon)$ polylog $(d, 1/\epsilon, 1/\tau)$ samples, runs in time nd polylog $(d, 1/\epsilon, 1/\tau)$, uses memory d polylog $(d, 1/\epsilon, 1/\tau)$, and returns a vector $\hat{\theta} \in \mathbb{R}^d$ such that $||\hat{\theta} - \theta^*||_2 = O(\sqrt{\epsilon})$ with probability at least $1 - \tau$.

E.3. Byzantine Adversary and Second-order Optimal Point

We now describe the application of our algorithm to the setting of robust distributed non-convex optimization. As before, for a parameter space $\Theta \subset \mathbb{R}^d$, a loss function $f:\Theta \times \mathcal{Z} \to \mathbb{R}^+$, and a distribution D over \mathcal{Z} , the goal is to approximately minimize $\bar{f}(\theta) = \mathbf{E}_{z \sim D}[f(\theta;z)]$. In this section, we consider the case when D is a uniform distribution over mn points $\{z_{i,j}: i \in [m], j \in [n]\}$ that are distributed over m machines (workers), with each machine having access to n samples. Furthermore, we do not impose convexity constraints on f, and thus would restrict ourselves to finding a second-order stationary point, i.e., a stationary point $\hat{\theta}$ such that the Hessian on $\hat{\theta}$ is not too negative in any direction.

We now explain the distributed setup in more detail. There are m workers who have their own private samples, and a single master machine which is responsible for collecting gradient estimates from the workers and updating the candidate vector iteratively. Concretely, the i-th worker has n samples $\{z_{ij}\}_{j=1}^n$. The master machine queries all workers with a parameter $\theta \in \Theta$, and each i-th worker responds with $g_i(\theta)$, where $g_i : \mathbb{R}^d \to \mathbb{R}^d$ is defined as follows: (i) if the i-th worker is honest, then $g_i(\theta)$ is the average of the gradients of f at θ of their samples, i.e., $g_i(\theta) := (1/n) \sum_{j=1}^n \nabla f(\theta; z_{ij})$, and (ii) if the i-th worker is dishonest, then $g_i(\cdot)$ is an arbitrary function. In our results, we require only that $(1 - \epsilon)$ -fraction of workers are honest. Recent work of (Yin et al., 2019) provided an algorithm that uses a robust mean estimation algorithm on the gradients as a black-box procedure. In particular, the algorithm of (Yin et al., 2019) requires only an access to the following oracle:

Definition E.13 (Δ -inexact gradient). We call the vector $v(\theta)$ a Δ -inexact gradient of \bar{f} at the point θ if $||v(\theta) - \nabla \bar{f}(\theta)||_2 \leq \Delta$.

We assume that each worker machine has access to its own samples throughout the optimization process, and our goal is to reduce the memory requirement of the master machine. Thus, we will use the algorithm from Corollary D.3 to calculate Δ -inexact gradient for the master machine, which requires only an oracle access to the gradient estimates $\{q_i(\theta) : i \in [m]\}$.

Assumption E.14. Let $\mathcal{I} \subseteq [m]$ be the set of honest workers with $|\mathcal{I}| \geq (1 - \epsilon)m$.

- 1. There exists δ with $0 \le \epsilon \le \delta \le \delta_0$, for some sufficiently small δ_0 , such that for every $\theta \in \Theta$, the set $\{g_i(\theta) \mid i \in \mathcal{I}\}$ is $(C\epsilon, \delta)$ -stable with respect to $\nabla \bar{f}(\theta)$ for a large enough constant C.
- 2. We assume that \bar{f} is L-smooth and ρ -Hessian Lipschitz on Θ , i.e., for every $\theta_1, \theta_2 \in \Theta$ we have that $\|\nabla \bar{f}(\theta_1) \nabla \bar{f}(\theta_2)\|_2 \le L\|\theta_1 \theta_2\|_2$ and $\|\nabla^2 \bar{f}(\theta_1) \nabla^2 \bar{f}(\theta_2)\|_2 \le \rho\|\theta_1 \theta_2\|_2$.

We note that if the samples of honest workers are sampled i.i.d. from a distribution P, then the set $\{g_i(\theta): i \in \mathcal{I}\}$ for a fixed $\theta \in \Theta$ will be stable with respect to $\nabla \bar{f}(\theta)$ with high probability, provided that the distribution of $\nabla f(\theta; Z)$ satisfies mild concentration under $Z \sim P$ and m is sufficiently large. Using a standard cover argument with the smoothness properties of f, this can be extended to all $\theta \in \Theta$. We thus obtain the following theorem, under Assumption E.14.

Theorem E.15. Suppose that Assumption E.14 holds. Let m denote the number of workers. Assume $0 < \tau < 1$, $\Delta := C'\delta < 1$, for C' a sufficiently large constant and define

$$Q := 2 \log \left(\frac{\rho(\bar{f}(\theta_0) - \inf_{\theta \in \mathbb{R}^d} \bar{f}(\theta))}{48L\tau(\Delta^{6/5}d^{3/5} + \Delta^{7/5}d^{7/10})} \right) \;, \quad T_{th} := \frac{L}{384(\rho^{1/2} + L(\Delta^{2/5}d^{1/5} + \Delta^{3/5}d^{3/10})} \;.$$

There is an algorithm where the master, if initialized at θ_0 , does $T = \frac{2(\bar{f}(\theta_0) - \inf_{\theta \in \mathbb{R}^d} \bar{f}(\theta))}{3\Delta^2} Q T_{th}$ iterations, each running in md polylog $(d, 1/\epsilon, T/\tau)$ time, uses d polylog $(d, 1/\epsilon, T/\tau)$ memory, and outputs a vector $\hat{\theta}$ such that, with probability $1 - \tau$, $\|\nabla \bar{f}(\hat{\theta})\|_2 \leq 4\Delta$ and $\lambda_{min}(\nabla^2 \bar{f}(\hat{\theta})) \geq -\Delta^{2/5} d^{1/5}$.

F. Discussion

In this work, we gave the first efficient streaming algorithm with near-optimal space complexity for outlier-robust high-dimensional mean estimation. As an application, we also obtained low-space streaming algorithms for a range of other robust estimation tasks. Our work is a first step towards understanding the space complexity of high-dimensional robust statistics in the streaming setting.

Our work suggests a number of open problems. First, the sample complexity of our mean estimation algorithm is $\tilde{O}(d^2/\epsilon^2)$, while the information-theoretic optimum (without space constraints!) is $\tilde{O}(d/\epsilon^2)$. What is the *optimal* sample-space tradeoff? A similar question can be asked for the broader tasks of covariance estimation and stochastic optimization. A more general goal is to characterize the tradeoff between space complexity, number of passes, and sample size/runtime for other robust high-dimensional statistics tasks, e.g., clustering and learning of mixture models.

Finally, another research direction concerns the considered contamination model. Throughout this paper, we focused on the TV-contamination model. One can consider an even stronger contamination model with an adaptive adversary, where the outliers can be completely arbitrary (i.e., not follow any distribution), and the adversary can additionally control the order in which the points are presented in the stream. Is it possible to obtain $\tilde{O}_{\epsilon}(d)$ -space single-pass streaming algorithms for robust mean estimation in the presence of such an adversary? While our algorithms can be shown to work in this model with a poly-logarithmic number of passes, it is not clear whether a single-pass algorithm with sub-quadratic space complexity exists in this setting.

G. Omitted Proofs from Appendix B: Technical Details Regarding Stability

Lemma B.11 (Certificate Lemma). Let G be an (ϵ, δ) -stable distribution with respect to $\mu \in \mathbb{R}^d$, for some $0 < \epsilon < 1/3$ and $\delta \geq \epsilon$. Let P be a distribution with $d_{TV}(P,G) \leq \epsilon$. Denoting by μ_P, Σ_P the mean and covariance of P, if $\lambda_{\max}(\Sigma_P) \leq 1 + \lambda$, for some $\lambda \geq 0$, then $\|\mu_P - \mu\|_2 = O(\delta + \sqrt{\epsilon \lambda})$.

Proof. Let $d_{\text{TV}}(P,G) = \alpha$. By Fact B.4 we can write $P = (1 - \alpha)G_0 + \alpha B$. We may assume without loss of generality $\alpha = \epsilon$, since we can always treat a part of the inliers as outliers. Denoting by μ_P, Σ_P the mean and covariance of P, and using $\mu_{G_0}, \mu_B, \Sigma_{G_0}, \Sigma_B$ for the corresponding quantities of the other two distributions, we have that

$$\Sigma_P = (1 - \epsilon) \Sigma_{G_0} + \epsilon \Sigma_B + \epsilon (1 - \epsilon) (\mu_{G_0} - \mu_B) (\mu_{G_0} - \mu_B)^T.$$

Letting v be the unit vector in the direction of $\mu_{G_0} - \mu_B$, we have that

$$1 + \lambda \ge v^T \mathbf{\Sigma}_P v = (1 - \epsilon) v^T \mathbf{\Sigma}_{G_0} v + \epsilon v^T \mathbf{\Sigma}_B v + \epsilon (1 - \epsilon) (v^T (\mu_{G_0} - \mu_B))^2.$$
(29)

The second term of the left-hand side is nonzero and the third one is just $\epsilon(1-\epsilon) \|\mu_{G_0} - \mu_B\|_2^2$. We now focus on the first term, which by adding and subtracting μ (the vector realizing the definition of stability for G) can be written as

$$(1 - \epsilon) \mathop{\mathbf{E}}_{X \sim G_0} [(v^T (X - \mu_{G_0}))^2] = (1 - \epsilon) \left(\mathop{\mathbf{E}}_{X \sim G_0} [(v^T (X - \mu))^2] - (v^T (\mu - \mu_{G_0}))^2 \right) . \tag{30}$$

We note that in the decomposition of Fact B.4, we can write $G_0(x) = w_0(x)G(x)$ with

$$w_0(x) = \frac{1}{1 - \epsilon} \begin{cases} P(x)/G(x) , & \text{if } G(x) > P(x) \\ 1 , & \text{otherwise }. \end{cases}$$

Letting $h(x) := (1 - \epsilon)w_0(x)$ we have that $h(x) \le 1$ for all x and $\mathbf{E}_{X \sim G}[h(X)] = 1 - \epsilon$, thus $G_0(x) = h(x)G(x)/(\int h(x)G(x)\mathrm{d}x) =: G_h(x)$. Returning to Equation (30), this means that

$$\mathbf{E}_{X \sim G_0}[(v^T (X - \mu))^2] = \mathbf{E}_{X \sim G_h}[(v^T (X - \mu))^2] = v^T \overline{\Sigma}_{h,G} v \ge 1 - \frac{\delta^2}{\epsilon} ,$$
 (31)

by applying stability. Similarly, the other term in Equation (30) is $(v^T(\mu - \mu_{G_0}))^2 \le \delta^2$. Putting everything together, Equation (29) becomes

$$1 + \lambda \ge (1 - \epsilon)(1 - \delta^2/\epsilon - \delta^2) + \epsilon(1 - \epsilon) \|\mu_{G_0} - \mu_B\|_2^2$$

$$\ge 1 - 3\delta^2/\epsilon + (\epsilon/2) \|\mu_{G_0} - \mu_B\|_2^2,$$

which yields $\|\mu_{G_0} - \mu_B\|_2 \lesssim \sqrt{\lambda/\epsilon} + \delta/\epsilon$. Then, writing $\mu_P = (1 - \epsilon)\mu_{G_0} + \epsilon\mu_B$ and using stability follows that $\|\mu_P - \mu\|_2 \lesssim \delta + \sqrt{\lambda\epsilon}$.

Lemma B.12. For any $0 < \epsilon < 1/2$ and $\delta \ge \epsilon$, if a distribution G is $(2\epsilon, \delta)$ -stable with respect to $\mu \in \mathbb{R}^d$, and P is an ϵ -corrupted version of G in total variation distance, there exist distributions G_0 and B such that $P = (1 - \epsilon)G_0 + \epsilon B$ and G_0 is (ϵ, δ) -stable with respect to μ .

Proof. By Fact B.4, we have the decomposition $P = (1 - \epsilon)G_0 + \epsilon B$, where $G_0(x) = \min\{G(x), P(x)\}/(1 - \epsilon)$. We can write $G_0(x) = w_0(x)G(x)$, where

$$w_0(x) = \frac{1}{1-\epsilon} \begin{cases} P(x)/G(x) \;, & \text{if } G(x) > P(x) \\ 1 \;, & \text{otherwise} \;. \end{cases}$$

To see why the final claim is true, we consider a weight function $w : \mathbb{R}^d \to [0,1]$ such that $\mathbf{E}_{X \sim G_0}[w(X)] \ge 1 - \epsilon$ and examine the adjusted distribution G_{0w} . We have that

$$G_{0w}(x) = \frac{w(x)G_0(x)}{\int_{\mathbb{R}^d} w(x)G_0(x)dx} = \frac{(1-\epsilon)w(x)w_0(x)G(x)}{\int_{\mathbb{R}^d} (1-\epsilon)w(x)w_0(x)G(x)dx} = \frac{h(x)G(x)}{\mathbf{E}_{X\sim G}[h(X)]} = G_h(x) ,$$

where we let $h(x) := (1 - \epsilon)w(x)w_0(x)$. We have that $h(x) \le 1$ point-wise and $\int_{\mathbb{R}^d} h(x)G(x) dx = (1 - \epsilon) \mathbf{E}_{X \sim G_0}[w(X)] \ge (1 - \epsilon)^2 \ge 1 - 2\epsilon$. Recalling that G is $(2\epsilon, \delta)$ -stable, the conclusion follows.

Lemma B.13. Fix $0 < \epsilon < 1/2$ and $\delta \ge \epsilon$. Let $w : \mathbb{R}^d \to [0,1]$ such that $\mathbf{E}_{X \sim G}[w(X)] \ge 1 - \epsilon$ and let G be an (ϵ, δ) -stable distribution with respect to $\mu \in \mathbb{R}^d$. For any matrix $\mathbf{U} \in \mathbb{R}^{d \times d}$ and any vector $b \in \mathbb{R}^d$, we have that

$$\mathbf{E}_{X \sim G_{m}} \left[\|\mathbf{U}(X-b)\|_{2}^{2} \right] = \|\mathbf{U}\|_{F}^{2} (1 \pm \delta^{2}/\epsilon) + \|\mathbf{U}(\mu-b)\|_{2}^{2} \pm 2\delta \|\mathbf{U}\|_{F}^{2} \|\mu-b\|_{2}.$$

Proof. We can write

$$\mathbf{E}_{X \sim G_w} \left[\|\mathbf{U}(X - b)\|_2^2 \right] = \mathbf{E}_{X \sim G_w} \left[\|\mathbf{U}(X - \mu)\|_2^2 + \|\mathbf{U}(\mu - b)\|_2^2 + 2(X - \mu)^T \mathbf{U}^T \mathbf{U}(\mu - b) \right]
= \mathbf{E}_{X \sim G_w} \left[\|\mathbf{U}(X - \mu)\|_2^2 \right] + \|\mathbf{U}(\mu - b)\|_2^2 + 2(\mu_{w,G} - \mu)^T \mathbf{U}^T \mathbf{U}(\mu - b) .$$
(32)

We now focus on the first term. Let the spectral decomposition $\mathbf{U}^T\mathbf{U} = \operatorname{tr}(\mathbf{U}^T\mathbf{U}) \sum_{i=1}^d \alpha_i v_i v_i^T$, where $\sum_{i=1}^d \alpha_i = 1$ and $\alpha_i \geq 0$. We have that

$$\begin{split} \mathbf{E}_{X \sim G_w} \left[\| \mathbf{U}(X - \mu) \|_2^2 \right] &= \operatorname{tr} \left(\mathbf{U}^T \mathbf{U} \mathbf{E}_{X \sim G_w} [(X - \mu)(X - \mu)^T] \right) = \operatorname{tr}(\mathbf{U}^T \mathbf{U}) \sum_{i=1}^d \alpha_i \operatorname{tr}(v_i v_i^T \overline{\mathbf{\Sigma}}_{w,G}) \\ &= \operatorname{tr}(\mathbf{U}^T \mathbf{U}) \sum_{i=1}^d \alpha_i v_i^T \overline{\mathbf{\Sigma}}_{w,G} v_i = \operatorname{tr}(\mathbf{U}^T \mathbf{U}) (1 \pm \delta^2 / \epsilon) = \| \mathbf{U} \|_F^2 (1 \pm \delta^2 / \epsilon) , \end{split}$$

where the second from the end relation is due to stability. Regarding the last term of Equation (32), we have that

$$|(\mu_{w,G} - \mu)^T \mathbf{U}^T \mathbf{U}(\mu - b)| = |\operatorname{tr}(\mathbf{U}^T \mathbf{U}(\mu - b)(\mu_{w,G} - \mu)^T)| \le \operatorname{tr}(\mathbf{U}^T \mathbf{U}) ||(\mu - b)(\mu_{w,G} - \mu)^T||_2$$

= $||\mathbf{U}||_F^2 ||\mu - b||_2 ||\mu_{w,G} - \mu||_2 \le ||\mathbf{U}||_F^2 \delta ||\mu - b||_2$,

where the last inequality uses stability condition for the mean.

Corollary B.14. Fix $0 < \epsilon < 1/2$ and $\delta \ge \epsilon$. Let G be an (ϵ, δ) -stable distribution with respect to $\mu \in \mathbb{R}^d$. Let a matrix $\mathbf{U} \in \mathbb{R}^{d \times d}$ and a function $w : \mathbb{R}^d \to [0, 1]$ with $\mathbf{E}_{X \sim G}[w(X)] \ge 1 - \epsilon$. For the function $\tilde{g}(x) = \|\mathbf{U}(x - b)\|_2^2$, we have that

$$(1-\epsilon)\|\mathbf{U}\|_F^2(1-\delta^2/\epsilon-2\delta\|b-\mu\|_2) \leq \mathbf{E}_{X \sim G}[w(X)\tilde{g}(X)] \leq \|\mathbf{U}\|_F^2\left(1+\delta^2/\epsilon+\|b-\mu\|_2^2+2\delta\|b-\mu\|_2\right).$$

Proof. Beginning with the upper bound, we have the following inequalities:

$$\begin{split} \mathbf{E}_{X \sim G}[w(X)\tilde{g}(X)] &= \mathbf{E}_{X \sim G}[w(X)] \sum_{X \sim G_w} \mathbf{E}_{\tilde{g}}[\tilde{g}(X)] \\ &\leq \mathbf{E}_{X \sim G_w}[\tilde{g}(X)] \qquad \qquad (\text{Using } \tilde{g}(x) \geq 0 \text{ and } w(x) \leq 1) \\ &\leq \|\mathbf{U}\|_F^2 (1 + \delta^2/\epsilon) + \|\mathbf{U}(\mu - b)\|_2^2 + 2\delta \|\mathbf{U}\|_F^2 \|b - \mu\|_2 \\ &\leq \|\mathbf{U}\|_F^2 \left(1 + \delta^2/\epsilon + \|b - \mu\|_2^2 + 2\delta \|b - \mu\|_2\right) \;, \end{split}$$

where the second inequality from the end uses Lemma B.13. The lower bound is derived similarly:

$$\begin{split} \underset{X \sim G}{\mathbf{E}}[w(X)\tilde{g}(X)] &= \underset{X \sim G}{\mathbf{E}}[w(X)] \underset{X \sim G_w}{\mathbf{E}}[\tilde{g}(X)] \\ &\geq (1 - \epsilon) \underset{X \sim G_w}{\mathbf{E}}[\tilde{g}(X)] \\ &\geq (1 - \epsilon) \|\mathbf{U}\|_F^2 (1 - \delta^2/\epsilon - 2\delta \|b - \mu\|_2) \;, \end{split}$$

where we applied Lemma B.13 in the last step.

H. Omitted Proofs from Appendix C

H.1. Johnson-Lindenstrauss Sketch

Lemma C.5. Fix a set of n points $x_1, \ldots, x_n \in \mathbb{R}^d$. For $t \in [K]$, define $g_t(x) := \|\mathbf{M}_t(x - \mu_t)\|_2^2$ and let $\tilde{g}_t(x), v_{t,j}$ as in Algorithm 3. If C is a sufficiently large constant and $L = C \log((n+d)K/\tau)$, with probability at least $1 - \tau$, for every $t \in [K]$ we have the following:

1.
$$0.8g_t(x_i) \leq \tilde{g}_t(x_i) \leq 1.2g_t(x_i)$$
 for every $i \in [n]$,

2.
$$0.8 \|\mathbf{M}_t\|_F^2 \le \left(\frac{1}{L} \sum_{j=1}^L \|v_{t,j}\|_2^2\right) \le 1.2 \|\mathbf{M}_t\|_F^2$$
.

Proof. We show that the claim holds for a fixed iteration t with probability τ/K . Recall that $\tilde{g}_t(x)$ from Algorithm 3, can be written as

$$\tilde{g}_t(x) = \|\mathbf{U}_t(x - \mu_t)\|_2^2 = \frac{1}{L} \sum_{j \in [L]} (v_{t,j}^T(x - \mu_t))^2 = \frac{1}{L} \sum_{j \in [L]} (z_{t,j}^T \mathbf{M}_t(x_i - \mu_t))^2.$$

Applying Fact B.7 with $\gamma = \tau/K$ and $u_i = \mathbf{M}_t(x_i - \mu_t)$, gives that choosing $L = C \log(nK/\tau)$ suffices to guarantee that $\tilde{g}_t(x_i)/g_t(x_i) \in [0.8, 1.2]$ for every x_i with probability $1 - \tau$.

We now show the second claim. Again, fix a $t \in [K]$. Consider the orthonormal base $\{e_i\}_{i=1}^d$ of \mathbb{R}^d . We apply Fact B.7 with $\gamma = \tau/K$ and $u_i = \mathbf{M}_t e_i$, $i \in [d]$. This yields that choosing $L = C \log(dK/\tau)$, we get that for all $i \in [d]$:

$$\frac{1}{L} \sum_{j=1}^{L} \operatorname{tr}(z_{t,j} z_{t,j}^{T} \mathbf{M}_{t} e_{i} e_{i}^{T} \mathbf{M}_{t}^{T}) = \frac{1}{L} \sum_{j=1}^{L} (z_{t,j}^{T} u_{i})^{2} = [0.8, 1.2] \frac{1}{L} \sum_{j=1}^{L} ||u_{i}||^{2} = [0.8, 1.2] \operatorname{tr}(\mathbf{M}_{t}^{T} \mathbf{M}_{t} e_{i} e_{i}^{T}),$$

with probability $1 - \tau/K$. Summing all these inequalities for $i = 1, \ldots, d$ and noting that $\sum_{i=1}^{d} e_i e_i^T = \mathbf{I}_d$ gives that

$$\frac{1}{L} \sum_{i=1}^{L} \operatorname{tr}(z_{t,j} z_{t,j}^{T} \mathbf{M}_{t}^{T} \mathbf{M}_{t}) = [0.8, 1.2] \operatorname{tr}(\mathbf{M}_{t}^{T} \mathbf{M}_{t}) ,$$

which precisely means that $\frac{1}{L} \sum_{j=1}^{L} \|v_{t,j}\|^2 = [0.8, 1.2] \|\mathbf{M}_t\|_F^2$. To have both claims hold simultaneously, we can just apply Fact B.7 for all n+d points, giving the result.

H.2. Proof of Lemma C.6

It is more useful to think of the algorithm in the following equivalent form.

Algorithm 8 Downweighting Filter

- 1: **function** DownweightingFilter($P, w, \tilde{\tau}, R, T, \ell_{\text{max}}$)
- 2: $r \leftarrow CdR^{2+4\log d}$.
- 3: $w' \leftarrow w, \ell \leftarrow 1$.
- 4: while $\mathbf{E}_{X \sim P}\left[w'(X)\tilde{\tau}(X)\right] > 2T$ and $\ell \leq \ell_{\max}$ do
- 5: $\ell \leftarrow \ell + 1$
- 6: $w'(x) \leftarrow w(x)(1 \tilde{\tau}(x)/r)$
- 7: end while
- 8: **return** w'.
- 9: end function

Lemma C.6. Let $P = (1 - \epsilon)G + \epsilon B$ be the empirical distribution on n samples, as in Algorithm 3. If $(1 - \epsilon) \mathbf{E}_{X \sim G}[w(X)\tilde{\tau}(X)] \leq T$, $\|\tilde{\tau}\|_{\infty} \leq r$, and $\ell_{\max} > r/T$, then Algorithm 4 modifies the weight function w to w' such that

- 1. $(1 \epsilon) \mathbf{E}_{X \sim G}[w(X) w'(X)] < \epsilon \mathbf{E}_{X \sim B}[w(X) w'(X)],$
- 2. $\mathbf{E}_{X \sim P} \left[w'(X) \tilde{\tau}(X) \right] \leq 2T$,

and the algorithm terminates after $O(\log(\ell_{\max}))$ iterations, each of which takes O(n) time.

Proof. We show correctness of Algorithm 8. We denote by w_ℓ the weight function at the ℓ -th iteration of the filter, which is of the form $w_\ell(x) = w(x)(1 - \tilde{\tau}(x)/r)^\ell$ for every $x \in \mathbb{R}^d$. To show the first claim, we fix an iteration ℓ for which the algorithm has not stopped yet and examine the loss in weight between that iteration and the $(\ell+1)$ -th iteration. From the update rule $w_{\ell+1}(x) = w_\ell(x)(1 - \tilde{\tau}(x)/r)$ we get that $w_\ell(x) - w_{\ell+1}(x) = w_\ell(x)\tilde{\tau}(x)/r$. Thus, the weight removed in that iteration from the good distribution is

$$(1 - \epsilon) \underset{X \sim G}{\mathbf{E}} [w_{\ell}(X) - w_{\ell+1}(X)] = \frac{1 - \epsilon}{r} \underset{X \sim G}{\mathbf{E}} [w_{\ell}(X)\tilde{\tau}(X)] \le \frac{1}{r} T,$$

while the weight removed from the bad distribution is

$$\begin{split} \epsilon \mathop{\mathbf{E}}_{X \sim B}[w_{\ell}(X) - w_{\ell+1}(X)] &= \frac{1}{r} \epsilon \mathop{\mathbf{E}}_{X \sim B}[w_{\ell}(X) \tilde{\tau}(X)] \\ &= \frac{1}{r} \left(\mathop{\mathbf{E}}_{X \sim P}[w_{\ell}(X) \tilde{\tau}(X)] - (1 - \epsilon) \mathop{\mathbf{E}}_{X \sim G}[w_{\ell}(X) \tilde{\tau}(X)] \right) \\ &> \frac{1}{r} T \;, \end{split}$$

where the last inequality uses that $(1 - \epsilon) \mathbf{E}_{X \sim G}[w_{\ell}(X)\tilde{\tau}(X)] \leq (1 - \epsilon) \mathbf{E}_{X \sim G}[w(X)\tilde{\tau}(X)] \leq T$ and the fact that since the algorithm has not terminated in the ℓ -th iteration it must be true that $\mathbf{E}_{X \sim P}[w_{\ell}(X)\tilde{\tau}(X)] > 2T$. This completes the proof of the first claim.

Regarding runtime, it suffices to show that for any $\ell > \frac{r}{eT}$, $\mathbf{E}_{X \sim P} \left[w_{\ell}(X) \tilde{\tau}(X) \right] \leq 2T$. This follows from the inequalities

$$\begin{split} \underset{X \sim P}{\mathbf{E}} \left[w_{\ell}(X) \tilde{\tau}(X) \right] &= \underset{X \sim P}{\mathbf{E}} \left[w(X) (1 - \tilde{\tau}(X)/r)^{\ell} \tilde{\tau}(X) \right] \\ &\leq \underset{X \sim P}{\mathbf{E}} \left[w(X) \exp(-\ell \tilde{\tau}(X)/r) \tau(X) \right] \\ &\leq \frac{r}{e \cdot \ell} \underset{X \sim P}{\mathbf{E}} \left[w(X) \right] \leq \frac{r}{e \cdot \ell} \leq T \;, \end{split}$$

where we used the fact that $xe^{-\alpha x} \leq 1/(e \cdot \alpha)$ for all $x \in \mathbb{R}$. By noting that $w(x)(1-\tilde{\tau}(x)/r)^{\ell}$ is monotonically decreasing as ℓ grows, we can improve the running time by using a binary search implementation. This gives the logarithmic guarantee of our statement.

H.3. Proof of Lemma C.7

We state and prove a more general version of Lemma C.7 so that it can be also used in Appendix D. The difference is that we allow the scores to center points using a vector different from the true mean μ_t of P_t , so long as this vector is $O(\delta)$ -close to μ_t in Euclidean norm. Lemma C.7 is obtained by using Lemma H.1 below with $\hat{\mu}_t = \mu_t$.

Lemma H.1. Consider the setting of Algorithm 3 and the deterministic Condition C.4. Moreover, let $\hat{\mu}_t$ be any vector in \mathbb{R}^d with $\|\hat{\mu}_t - \mu_t\| = O(\delta)$ and define the functions

$$f_{t}(x) := \|\mathbf{M}_{t}(x - \hat{\mu}_{t})\|_{2}^{2}, \qquad \qquad \tilde{f}_{t}(x) := \|\mathbf{U}_{t}(x - \hat{\mu}_{t})\|_{2}^{2}$$

$$h_{t}(x) := f_{t}(x)\mathbb{1}\{f_{t}(x) > C_{3}\|\mathbf{M}_{t}\|_{F}^{2}\lambda_{t}/\epsilon\}, \qquad \qquad \tilde{h}_{t}(x) := \tilde{f}_{t}(x)\mathbb{1}\{\tilde{f}_{t}(x) > C_{3}\|\mathbf{U}_{t}\|_{F}^{2}\hat{\lambda}_{t}/\epsilon\}.$$
(33)

We have that $\mathbf{E}_{X \sim G}[w_t(X)h_t(X)]$ and $\mathbf{E}_{X \sim G}[w_t(X)\tilde{h}_t(X)]$ are bounded from above by $c\lambda_t \|\mathbf{M}_t\|_F^2$ for some constant c of the form $c = C/C_2$, where C_2 is the constant used in Line 13 and C is some absolute constant.

We prove the result by using the facts from Appendix B.2. For brevity, we will prove the results for \tilde{h}_t by using the functions \tilde{f}_t and the matrix \mathbf{U}_t ; the results for h_t would follow by replacing \tilde{f}_t and \mathbf{U}_t with f_t and \mathbf{M}_t respectively and using that $\|\mathbf{U}_t\|_F$ is close to $\|\mathbf{M}_t\|_F$ (Item 2 of the deterministic condition). We begin with Lemma H.2, which is a generalization of the following implication of stability: The (ϵ, δ) -stability of a distribution G implies that $\mathbf{E}_{X \sim G}[(v^T(X - \mu))^2 \mathbb{1}\{X \in L\}] \leq 3\delta^2/\epsilon$ for any set L with mass $\Pr_{X \sim G}[X \in L] \leq \epsilon$ (see, for example, Proposition C.3 of (Pensia et al., 2020)). The following lemma generalizes this to having a matrix in place of v.

Lemma H.2. Under the setting of Algorithm 3, the deterministic Condition C.4, and using the notation of Equation (33), if $L_t \subseteq \mathbb{R}^d$ is a set with $\mathbf{E}_{X \sim G}[w_t(X)\mathbb{1}\{X \in L_t\}] \leq \epsilon$, then we have that

$$\underset{X \sim G}{\mathbf{E}}[w_t(X)\tilde{f}_t(X)\mathbb{1}\{X \in L\}] \le c\lambda_t \|\mathbf{U}_t\|_F^2 ,$$

for some constant c of the form $c = C'/C_2$, where C' is a sufficiently large constant.

Proof. Define the new weight function $w_t'(x) = w_t(x) \mathbb{1}\{x \notin L_t\}$. We have assumed that the distribution G is $(C'''\epsilon, \delta)$ -stable. Let $\epsilon' := C''\epsilon$ for brevity. We have the following inequalities, which we explain below.

$$\mathbf{E}_{X \in C}[w_t(X)\tilde{f}_t(X)\mathbb{1}\{X \in L_t\}] = \mathbf{E}_{X \in C}[w_t(X)\tilde{f}_t(X)] - \mathbf{E}_{X \in C}[w_t(X)\tilde{f}_t(X)\mathbb{1}\{X \notin L_t\}]$$

$$\begin{aligned}
&= \underset{X \sim G}{\mathbf{E}}[w_{t}(X)\tilde{f}_{t}(X)] - \underset{X \sim G}{\mathbf{E}}[w'_{t}(X)\tilde{f}_{t}(X)] \\
&\leq \|\mathbf{U}_{t}\|_{F}^{2} \left(1 + \frac{\delta^{2}}{\epsilon'} + \|\widehat{\mu}_{t} - \mu\|_{2}^{2} + 2\delta\|\widehat{\mu}_{t} - \mu\|_{2}\right) \\
&- (1 - 2\epsilon)\|\mathbf{U}_{t}\|_{F}^{2} \left(1 - \frac{\delta^{2}}{\epsilon'} - 2\delta\|\widehat{\mu}_{t} - \mu\|_{2}\right) \\
&\leq \|\mathbf{U}_{t}\|_{F}^{2} \left(3\frac{\delta^{2}}{\epsilon'} + 4\delta\|\widehat{\mu}_{t} - \mu\|_{2} + \|\widehat{\mu}_{t} - \mu\|_{2}^{2}\right) \\
&\leq \|\mathbf{U}_{t}\|_{F}^{2} \left(3\frac{\delta^{2}}{\epsilon'} + 4\delta(\|\widehat{\mu}_{t} - \mu_{t}\|_{2} + \|\mu_{t} - \mu\|_{2}) + 4\|\widehat{\mu}_{t} - \mu_{t}\|_{2}^{2} + 4\|\mu_{t} - \mu\|_{2}^{2}\right) \\
&\leq \|\mathbf{U}_{t}\|_{F}^{2} \left(3\frac{\delta^{2}}{\epsilon'} + \delta O(\delta + \sqrt{\epsilon'\lambda_{t}}) + O(\delta^{2} + \epsilon'\lambda_{t})\right) \\
&\leq c\|\mathbf{U}_{t}\|_{F}^{2} \lambda_{t}.
\end{aligned}$$

We note that the third line above follows by applying Corollary B.14 with $\mathbf{U} = \mathbf{U}_t$ and $b = \mu_t$ on both terms of the previous line. The fifth line uses the triangle inequality. The sixth line uses the assumption that $\|\widehat{\mu}_t - \mu_t\|_2 = O(\delta)$ as well as the certificate lemma (Lemma B.11). Note that the required assumption $d_{\text{TV}}(P_t, P) = O(\epsilon)$ from that lemma is satisfied because $\mathbf{E}_{X \sim G}[w_t'(X)] \ge \mathbf{E}_{X \sim G}[w_t(X)] - \epsilon \ge 1 - O(\epsilon)$ (see Claim C.13).

Regarding that last line, we recall Line 13 of Algorithm 3, which implies that $\lambda_t \gtrsim C_2 \delta^2 / \epsilon$. Thus the terms δ^2 / ϵ' can be bounded as $\delta^2 / \epsilon' \lesssim 1/C_2$. Using that, it can be seen that we can choose $c = C'/C_2$ for some constant C' > 0.

We are now ready to prove our result.

Proof of Lemma H.1. We first show the following.

Claim H.3. Consider the setting of Algorithm 3, the notation of Equation (33), and assume that the deterministic Condition C.4 holds. Let $L_t = \{x : \tilde{f}_t(x) > C_3 \|\mathbf{U}_t\|_F^2 \widehat{\lambda}_t/\epsilon \}$. We have that $\mathbf{E}_{X \sim G}[w_t(X)\mathbb{1}\{X \in L_t\}] \leq \epsilon$.

Proof. Let $u^* := \arg\max\{u : \mathbf{E}_{X \sim G}[w_t(X)\mathbb{1}\{\tilde{f}_t(X) > u\}] \ge \epsilon\}$ and the set $L_t^* = \{x : \tilde{f}_t(x) > u^*\}$. It suffices to show that $u^* \le C_3 \|\mathbf{U}_t\|_F^2 \hat{\lambda}_t / \epsilon$ (because this would mean that $L_t \subseteq L_t^*$).

By Lemma H.2, we have that $\mathbf{E}_{X \sim G}[w_t(X)\tilde{f}_t(X)\mathbb{1}\{X \in L_t^*\}] \leq (C'/C_2)\lambda_t \|\mathbf{U}_t\|_F^2$. If we define the new weights $w_t'(x) = w_t(x)\mathbb{1}\{x \in L_t^*\}$ and use them to normalize the distribution, we get that

$$\underset{X \sim G_{w'_t}}{\mathbf{E}} [\tilde{f}_t(X)] = \frac{1}{\mathbf{E}_{X \sim G}[w_t(X) \mathbb{1}\{X \in L_t^*\}]} \underset{X \sim G}{\mathbf{E}} [w_t(X) \tilde{f}_t(X) \mathbb{1}\{X \in L_t^*\}] \leq (C'/C_2) \|\mathbf{U}_t\|_F^2 \frac{\hat{\lambda}_t}{\epsilon} ,$$

where we used that the denominator is ϵ . The fact that $\mathbf{E}_{X \sim G_{w'_t}}[\tilde{f}_t(X)] \leq (C'/C_2) \|\mathbf{U}_t\|_F^2 \hat{\lambda}_t/\epsilon$ means that at least one point in L_t^* has $\tilde{f}_t(X) \leq (C'/C_2) \|\mathbf{U}_t\|_F^2 \hat{\lambda}_t/\epsilon$, which shows that $u^* \leq (C'/C_2) \|\mathbf{U}_t\|_F^2 \hat{\lambda}_t/\epsilon$. Since the algorithm uses the value $C_3 = (C'/C_2)$, the proof is completed.

Lemma H.1 now follows by combining Claim H.3 with Lemma H.2:

$$\mathbf{E}_{X \sim G}[w_t(X)\tilde{h}_t(X)] = \mathbf{E}_{X \sim G}\left[w_t(X)\tilde{f}_t(X)\mathbb{1}\left\{\tilde{f}_t(X) > C_3\|\mathbf{U}_t\|_F^2 \frac{\hat{\lambda}_t}{\epsilon}\right\}\right]$$

$$\leq (C'/C_2)\lambda_t\|\mathbf{U}_t\|_F^2 \leq 2(C'/C_2)\lambda_t\|\mathbf{M}_t\|_F^2,$$

where the last inequality is due to Item 2 of Condition C.4. Letting C=2C' we have that $\mathbf{E}_{X\sim G}[w_t(X)\tilde{h}_t(X)] \leq C/C_2$ as claimed. As mentioned earlier, the same techniques lead to a similar bound on $\mathbf{E}_{X\sim G}[w_t(X)h_t(X)]$.

H.4. Proof of Claim C.11

Claim C.11. Let the fraction of outliers be $\epsilon < 1/10$ and a parameter $0 < \tau < 1$. Let the distribution $P = (1 - \epsilon)G + \epsilon B$. Let R > 0, $\mu \in \mathbb{R}^d$ be such that $\Pr_{X \sim G}[\|X - \mu\|_2 > R] \le \epsilon$. There is an estimator $\widehat{\mu}$ on $k = O(\log(1/\tau))$ samples from P such that $\|\widehat{\mu} - \mu\|_2 \le 4R$ with probability at least $1 - \tau$. Furthermore, $\widehat{\mu}$ can be computed in time $O(k^2d)$ and memory O(kd).

We can use the following well-known fact (see, e.g., (Dong et al., 2019), for a proof):

Fact H.4. There is an algorithm NaivePrune with the following guarantees. Let $\epsilon' < 1/2$, and $\tau > 0$. Let $S \subset \mathbb{R}^d$ be a set of n points so that there exists a $\mu \in \mathbb{R}^d$ and a subset $S' \subseteq S$ so that $|S'| \ge (1 - \epsilon')n$, and $||x - \mu||_2 \le R$ for all $x \in S'$. Then NaivePrune (S, R, τ) runs in time $O(nd \log(1/\tau))$, uses memory O(nd), and with probability $1 - \tau$ outputs a set of points $T \subset S$ so that $S' \subseteq T$, and $||x - \mu||_2 \le 4R$ for all $x \in T$.

Proof of Claim C.11. The estimator draws a set $S = \{X_1, \dots, X_k\}$ of k samples from the distribution P. Letting $Y_i := \mathbb{1}\{\|X_i - \mu\|_2 > R\}$ we have that $\mathbf{E}[Y_i] \le 2\epsilon \le 1/5$. Thus, using Hoeffding bound,

$$\Pr\left[\frac{1}{k}\sum_{i=1}^{k}Y_{i} > 1/4\right] \le \Pr\left[\frac{1}{k}\sum_{i=1}^{k}Y_{i} < \mathbf{E}\left[\frac{1}{k}\sum_{i=1}^{k}Y_{i}\right] + 0.05\right] \le 2e^{-2k(0.05)^{2}}.$$

Choosing $k=200\log(2/\tau)$ makes this probability at most τ . Conditioning on that event, the fraction of points outside the ball is at most $\epsilon':=1/4$, thus running NaivePruning (S,R,τ) algorithm of Fact H.4 and outputting any point from the returned set yields the desired estimator.

H.5. Omitted Proofs from Appendix C.4

Claim C.13. Under Condition C.4, Algorithm 3 maintains the following invariant: $\mathbf{E}_{X \sim G}[w_t(X)] \ge 1 - 3\epsilon$. In particular, if $\epsilon \le 1/8$, then $d_{\mathrm{TV}}(P_t, P) \le 9\epsilon$.

Proof. For every iteration, we denote by $\Delta w_t = w_t - w_{t+1}$, that is for every point $x \in \mathbb{R}^d$, $\Delta w_t(x) = w_t(x) - w_{t+1}(x)$ is the difference between the weights for the two consecutive iterations t and t+1.

$$\begin{split} \mathbf{E}_{X \sim G}[w_t(X)] &= \mathbf{E}_{X \sim G}[w_1(X)] - \sum_{i=1}^{t-1} \mathbf{E}_{X \sim G}[\Delta w_i(X)] \\ &\geq 1 - \epsilon - \sum_{i=1}^{t-1} \mathbf{E}_{X \sim G}[\Delta w_i(X)] \\ &\geq 1 - \epsilon - \frac{\epsilon}{1 - \epsilon} \sum_{i=1}^{t-1} \mathbf{E}_{X \sim B}[\Delta w_i(X)] \\ &\geq 1 - \epsilon - \frac{\epsilon}{1 - \epsilon} \left(\mathbf{E}_{X \sim B}[w_1(X)] - \mathbf{E}_{X \sim B}[w_t(X)] \right) \\ &\geq 1 - \epsilon - \frac{\epsilon}{1 - \epsilon} \\ &\geq 1 - 3\epsilon \;, \end{split}$$
 (Claim C.12)

where the last line uses that $\epsilon \leq 1/2$. The proof of the second conclusion follows from Claim H.5 (stated below).

Claim H.5. Let
$$\epsilon \leq 1/8$$
. If $\mathbf{E}_{X \sim G}[w_t(X)] \geq 1 - 3\epsilon$, then $d_{\mathrm{TV}}(P_t, P) \leq 9\epsilon$.

Although we work with discrete distributions (the empirical distributions on the samples) in Appendix C, we prove the claim for continuous distributions because it will be useful in Appendix D.

Proof. By definition $P_t(x) = w_t(x)P(x)/\mathbf{E}_{X\sim P}[w_t(X)]$. Letting $L := \int_{\mathbb{R}^d} w_t(x)P(x)\mathrm{d}x = \mathbf{E}_{X\sim P}[w_t(x)]$, we have that

$$\int_{\mathbb{R}^d} |P_t(x) - P(x)| \mathrm{d}x = (1 - \epsilon) \int_{\mathbb{R}^d} G(x) \left| \frac{w_t(x) - L}{L} \right| \mathrm{d}x + \epsilon \int_{\mathbb{R}^d} B(x) \left| \frac{w_t(x) - L}{L} \right| \mathrm{d}x \ .$$

We note that $1 \ge L \ge (1 - \epsilon) \mathbf{E}_{X \sim G}[w_t(x)] \ge 1 - 4\epsilon$ using Claim C.13. The second term can be bounded as

$$\epsilon \int_{\mathbb{R}^d} B(x) \left| \frac{w_t(x) - L}{L} \right| dx \le \frac{\epsilon}{L} \int_{\mathbb{R}^d} B(x) (w_t(x) + L) \le \frac{2\epsilon}{1 - 4\epsilon} \le 4\epsilon.$$

For the first term, we have that

$$(1 - \epsilon) \int_{\mathbb{R}^d} G(x) \left| \frac{w_t(x) - L}{L} \right| dx \le \frac{1 - \epsilon}{L} \int_{\mathbb{R}^d} G(x) (|1 - w_t(x)| + |1 - L|) dx$$
$$\le \frac{1}{1 - 4\epsilon} \left(1 - \underset{X \sim G}{\mathbf{E}} [w_t(X)] + 4\epsilon \right) \le 14\epsilon .$$

Claim C.14. Under Condition C.4, if $C_1 \ge 22$, $\mathbf{B}_t \succeq (0.5C_1\delta^2/\epsilon)\mathbf{I}_d$ for every $t \in [K]$.

Proof. Using the simple fact that for random variables X, Y it holds $Var(Y) \ge \mathbf{E}_X[Var(Y|X)]$, we get that

$$\Sigma_{t} \succeq (1 - \epsilon) \underset{X \sim G_{w_{t}}}{\mathbf{E}} \left[(X - \mu_{G_{w_{t}}})(X - \mu_{G_{w_{t}}})^{T} \right] \\
= (1 - \epsilon) \left(\underset{X \sim G_{w_{t}}}{\mathbf{E}} \left[(X - \mu)(X - \mu)^{T} \right] - (\mu_{G_{w_{t}}} - \mu)(\mu_{G_{w_{t}}} - \mu)^{T} \right) \\
\succeq (1 - \epsilon) \left((1 - \delta^{2}/\epsilon) \mathbf{I}_{d} - \delta^{2} \mathbf{I}_{d} \right) \qquad \text{(by stability and Claim C.13)} \\
\succeq (1 - \epsilon)(1 - 2\delta^{2}/\epsilon) \mathbf{I}_{d} \\
\succeq (1 - 3\delta^{2}/\epsilon) \mathbf{I}_{d} ,$$

where we used that $\epsilon \leq \delta$. We recall the definition $\mathbf{B}_t = (\mathbf{E}_{X \sim P_t}[w_t(X)])^2 \mathbf{\Sigma}_t - (1 - C_1 \delta^2 / \epsilon) \mathbf{I}_d$ and bound the first term as follows

$$\left(\sum_{X \sim P_t} [w_t(X)]\right)^2 \mathbf{\Sigma}_t \succeq (1 - 3\epsilon)^2 (1 - 3\delta^2/\epsilon) \mathbf{I}_d \qquad (Claim C.13)$$

$$\succeq (1 - 4\delta^2/\epsilon - 6\epsilon - 27\delta^2\epsilon) \mathbf{I}_d$$

$$\succeq (1 - 11\delta^2/\epsilon) \mathbf{I}_d,$$

where the last line uses $\epsilon < 1/6$ and $\epsilon \le \delta$. Therefore, if we choose $C_1 \ge 22$, we get that $\mathbf{B}_t \succeq (0.5C_1\delta^2/\epsilon)\mathbf{I}_d$.

Claim C.10. In the setting of Algorithm 3 and under the Condition C.4, if $x \in S$, we have that $\tau_t(x) \leq 1.25\tilde{\tau}_t(x) + 3C_3(\lambda_t/\epsilon)\mathrm{tr}(\mathbf{M}_t^2)$, where C_3 is the constant used in Algorithm 3.

Proof. By Condition C.4 we have that for all the n samples, $\tilde{g}_t(x) \geq 0.8g_t(x)$. Recall the definitions $\tilde{\tau}_t(x) = \tilde{g}(x)\mathbb{1}\{\tilde{g}(x) > C_3\|\mathbf{U}_t\|_F^2\hat{\lambda}_t/\epsilon\}$ and $\tau_t(x) = g(x)\mathbb{1}\{g(x) > C_3\|\mathbf{M}_t\|_F^2\lambda_t/\epsilon\}$. We split into cases based on whether each of g_t , \tilde{g}_t has been zeroed by its thresholding operation:

- If $\tau_t(x)$ has been zeroed, (i.e., $g_t(x) < C_3 \|\mathbf{M}_t\|_F^2 \lambda_t/\epsilon$), the claim trivially holds since the left-hand side is zero.
- If none of $\tilde{\tau}_t(x)$, $\tau_t(x)$ has been zeroed, then $\tilde{\tau}_t(x) = \tilde{g}_t(x)$ and $\tau_t(x) = g_t(x)$, thus the claim holds by the aforementioned fact that $\tilde{g}_t(x) \geq 0.8g_t(x)$.
- If $\tilde{\tau}_t(x)$ has been zeroed but $\tau_t(x)$ has not, then the worst case is $g_t(x) = (1/0.8)\tilde{g}_t(x)$. This means that in this case:

$$\tau_t(x) \le \frac{1}{0.8} C_3 \frac{\hat{\lambda}_t}{\epsilon} |\mathbf{U}_t||_F^2 < 3C_3 \frac{\lambda_t}{\epsilon} |\mathbf{M}_t||_F^2 ,$$

where we used that $\hat{\lambda} \leq 1.2\lambda_t$ and $\|\mathbf{U}_t\|_F^2 \leq 1.2\|\mathbf{M}_t\|_F^2$, due to Condition C.4.

I. Omitted Proofs from Appendix D

I.1. Omitted Proofs from Appendix D.2

Lemma D.6. In the context of Algorithm 5, if $(1 - \epsilon) \mathbf{E}_{X \sim G}[w(X)\tilde{\tau}(X)] \leq T$, $\|\tilde{\tau}\|_{\infty} \leq r$, and $\ell_{\max} > r/T$, then Algorithm 6 modifies the weight function w to w' such that $(i) (1 - \epsilon) \mathbf{E}_{X \sim G}[w(X) - w'(X)] < \epsilon \mathbf{E}_{X \sim B}[w(X) - w'(X)]$, and (ii) upon termination we have $\mathbf{E}_{X \sim P}[w'(X)\tilde{\tau}(X)] \leq 54T$. Furthermore, if the estimator of Line 3 is set to be that of Lemma D.16, the algorithm terminates after $O(\log(\ell_{\max}))$ iterations, each of which uses $O((R^2\epsilon/\delta^2)\log(1/\tau))$ samples, takes O(nd) time and memory $O(\log(1/\tau))$.

Proof. Let the set $L^* = \{\ell \in [\ell_{\max}] : 6T \leq \mathbf{E}_{X \sim P}[w(X)(1-\tilde{\tau}(X)/r)^{\ell}\tilde{\tau}(X)] \leq 18T\}$. The invariant is that throughout Algorithm 6, the set L maintained has non-zero intersection with L^* . This can be seen by examining cases about ℓ in Line 6. If $\ell \in L^*$, then ℓ is kept in L. If $\ell \notin L^*$, then all elements discarded are not members of L^* (for example if $\mathbf{E}_{X \sim P}[w(X)(1-\tilde{\tau}(X)/r)^{\ell}\tilde{\tau}(X)] > 18T$, then by Lemma D.16 $f(\ell) > 9T$ and we discard the lower half of L). Thus, at the end, L has at most two elements with at least one of them belonging in L^* . This element would satisfy $3T \leq f(\ell) \leq 27T$. Thus the algorithm will definitely return some element. On the other hand, any element returned will satisfy $2T < \mathbf{E}_{X \sim P}[w(X)(1-\tilde{\tau}(X)/r)^{\ell}\tilde{\tau}(X)] < 54T$. This has already shown part (ii) of the lemma. For part (i), it is more convenient to imagine let ℓ increased by one at each step until it reaches the value finally returned by the algorithm and consider the loss in weight between that and the next iteration, exactly as in the proof of Lemma C.6. That proof was using only the facts that for all $\ell' \leq \ell$, $2T < \mathbf{E}_{X \sim P}[w(X)(1-\tilde{\tau}(X)/r)^{\ell'}\tilde{\tau}(X)]$ (which we just showed above) and $\mathbf{E}_{X \sim G}[w(X)(1-\tilde{\tau}(X)/r)^{\ell'}\tilde{\tau}(X)] < T$ (which is true by assumption). The reason why $\ell_{\max} = r/T$ suffices is also shown identically to the proof of Lemma C.6.

I.2. Omitted Proofs from Appendix D.2.1

We now focus on showing Lemma D.9. In order to avoid confusion with the fraction of outliers ϵ , we use ϵ' for our accuracy parameter. We will use a uniform convergence result from (Anthony & Bartlett, 1999) combined with a powerful VC-dimension bound from (Goldberg & Jerrum, 1995) for the class of functions that are computable by a small number of arithmetic operations. (Goldberg & Jerrum, 1995) considers the class of concepts parameterized by k real numbers, $\mathcal{F} = \{h_a\}_{a \in \mathbb{R}^k}$, for which there exists an algorithm \mathcal{A} for calculating $h_a(x)$ that takes as input x, a and each line of \mathcal{A} is one of the following:

- an arithmetic operation $+, -, \times$, and / on two inputs or previously computed values,
- a jump to a different line of the algorithm conditioned on whether an input or a previously calculated value is greater than or equal to zero,
- · output zero or one.

The parameters a and the inputs x consist of real numbers, and the model of computation assumed allows for arithmetic operations and the comparisons between reals to be done in constant time. We refer the reader to (?)Section 2]gold-berg1995bounding for more details and relation with algebraic decision trees with bounded depths. The result from (Goldberg & Jerrum, 1995) is that $VCdim(\mathcal{F}) = O(mk)$ where k is the size of the parameterization and m is the runtime of the algorithm \mathcal{A} . Using the bound on the VC dimension, we have the following result for the uniform convergence:

Proposition I.1 ((Goldberg & Jerrum, 1995; Anthony & Bartlett, 1999)). Let \mathcal{F} be a class of functions of the form $\mathcal{F} = \{h_a : \mathbb{R}^d \to [0,1] \mid a \in \mathbb{R}^k\}$, where for any $(a,x) \in \mathbb{R}^k \times \mathbb{R}^d$, $h_a(x)$ can be computed by an algorithm \mathcal{A} with runtime m that takes as input a,x and is allowed to perform conditional jumps (conditioned on equality and inequality of real values) and execute the standard arithmetic operations on real numbers $(+,-,\times,/)$ in constant time. For any distribution D on \mathbb{R}^d and any $\epsilon' \in (0,1)$, there exist $N = O\left(\frac{1}{(\epsilon')^2}(\log(km) + km\log(1/\epsilon'))\right)$ points x_1,\ldots,x_N in \mathbb{R}^d such that

$$\sup_{h \in \mathcal{F}} \left| \underset{X \sim D}{\mathbf{E}} [h(X)] - \frac{1}{N} \sum_{i=1}^{N} h(x_i) \right| \le \epsilon'.$$

For completeness, we show at the end of this section how this is derived from the statements of (Anthony & Bartlett, 1999) and (Goldberg & Jerrum, 1995). We now apply this result to our case. We need to specify a family \mathcal{F} of functions broad

enough to capture every $w_{t+1}\tilde{\tau}_t$ and $w_{t+1}\tau_t$ that could be encountered during the execution of Algorithm 5. The factor r used in the statement below is a normalization factor to make sure that the functions are in [0, 1].

Lemma I.2. Consider the setting of Algorithm 5. Let $r' := (CdR^2 + 1 + C_1\delta^2/\epsilon)^{C\log d}$. There exists a family \mathcal{F} of functions from \mathbb{R}^d to [0,1] such that:

- 1. For every iteration t of Algorithm 5, we have that $\frac{1}{r'}w_{t+1}\tau_t \in \mathcal{F}$ and $\frac{1}{r'}w_{t+1}\tilde{\tau}_t \in \mathcal{F}$.
- 2. Functions of \mathcal{F} are parameterized by at most $k = O(dK \max(L, d))$ real numbers, that is, \mathcal{F} has the form $\mathcal{F} = \{h_a : \mathbb{R}^d \to [0, 1] \mid a \in \mathbb{R}^k\}$.
- 3. For every $h_a \in \mathcal{F}$ and $x \in \mathbb{R}^d$, $h_a(x)$ can be in $m = dK \max(L, d) (dR\epsilon/\delta^2)^{O(\log d)}$ steps in the model that allows conditional jumps and standard arithmetic operations on real numbers.

Proof. Let $L_2' := \max(L,d)$. Every function in our family will be parameterized by 2K+1 scalars, $\{u_t \in \mathbb{R} : t \in [K+1]\} \cup \{\ell_t \in \mathbb{R} : t \in [K]\}$, and $(K+1)(L_2'+1)$ vectors in \mathbb{R}^d , $\{a_t : t \in [K+1]\} \cup \{v_{t,j} : t \in [K+1], j \in [L_2']\}$. For brevity, we denote by \mathbf{V} the tensor in $\mathbb{R}^{(K+1) \times L_2' \times d}$ having all the vectors $\mathbf{V}_{t,j} = v_{t,j}$ and by \mathbf{A} the tensor in $\mathbb{R}^{(K+1) \times d}$ with $\mathbf{A}_t = a_t, t \in [K+1]$. Similarly, denote by u the vector (u_1, \dots, u_{K+1}) and let $\ell = (\ell_1, \dots \ell_K)$. We define our class to be

$$\mathcal{F} = \left\{ h_{\ell,u,\mathbf{V},\mathbf{A}} : \mathbb{R}^d \to [0,1] : u \in \mathbb{R}^{K+1}, \ell \in \mathbb{R}^K, \mathbf{V} \in \mathbb{R}^{(K+1) \times L_2' \times d}, \mathbf{A} \in \mathbb{R}^{(K+1) \times d} \right\} ,$$

which includes all functions of the form $h_{\ell,u,\mathbf{V},\mathbf{A}}(x) = \tilde{h}_{\ell,u,\mathbf{V},\mathbf{A}} \mathbb{1}\{\tilde{h}_{\ell,u,\mathbf{V},\mathbf{A}}(x) \in (0,1)\}$, where

$$\tilde{h}_{\ell,u,\mathbf{V},\mathbf{A}}(x) = \mathbb{1} \left\{ \|x - \hat{\mu}\|_{2} \le 5R \right\} \frac{1}{r} \cdot \frac{\sum_{j=1}^{L'_{2}} (v_{K+1,j}^{T}(x - a_{K+1}))^{2}}{L'_{2}} \mathbb{1} \left\{ \frac{\sum_{j=1}^{L'_{2}} (v_{K+1,j}^{T}(x - a_{K+1}))^{2}}{L'_{2}} > u_{K+1} \right\}
\cdot \prod_{t=1}^{K} \left(1 - \frac{1}{r} \frac{\sum_{j=1}^{L'_{2}} (v_{t,j}^{T}(x - a_{t}))^{2}}{L'_{2}} \mathbb{1} \left\{ \frac{\sum_{j=1}^{L'_{2}} (v_{t,j}^{T}(x - a_{t})))^{2}}{L'_{2}} > u_{t} \right\} \right)^{\ell_{t}} .$$
(34)

We note that the radius $r' := \left(CdR^2 + 1 + C_1\delta^2/\epsilon\right)^{C\log d}$ is an upper bound on the value that the functions $w_{t+1}\tau_t$ and $w_{t+1}\tilde{\tau}_t$ in Algorithm 5 can take: For $\tau_t(x)$ we have

$$\tau_t(x) \le \|\mathbf{M}_t(x - \mu_t)\|_2^2 \le \|\mathbf{M}_t\|_2^2 \|x - \mu_t\|_2^2 \le \|\mathbf{\Sigma}_t\|_2^{2\log d} R^2 = O(dR^{2+4\log d}),$$

while for $\tilde{\tau}(x)$ we have the bounds

$$\tilde{\tau}_{t}(x) \leq \tilde{g}_{t}(x) \leq \|\mathbf{U}_{t}(x - \mu_{t})\|_{2}^{2} \lesssim R^{2} \|\mathbf{U}_{t}\|_{2}^{2} \lesssim R^{2} \|\mathbf{U}_{t}\|_{F}^{2}
\leq R^{2} \frac{1}{L} \sum_{j=1}^{L} \|\mathbf{M}_{t} z_{t,j}\|_{2}^{2} \leq dR^{2} \|\mathbf{M}_{t}\|_{2}^{2} \leq dR^{2} \|\mathbf{B}_{t}\|_{2}^{2 \log d}
\leq dR^{2} (\|\mathbf{\Sigma}_{t}\|_{2} + 1 + C_{1} \delta^{2} / \epsilon)^{2 \log d} \leq dR^{2} (CR^{2} + 1 + C_{1} \delta^{2} / \epsilon)^{2 \log d}
\leq (CdR^{2} + 1 + C_{1} \delta^{2} / \epsilon)^{O(\log d)}.$$

We check that \mathcal{F} can indeed implement the functions $w_{t+1}\tilde{\tau}_t$ used in Algorithm 5 for any $t\in[K]$: Note that the scores \tilde{g}_t used in the algorithm are means of the form $\frac{1}{L}\sum_{j=1}^L(v_{t,j}^T(x-a_t))^2$. Thus, the first line of Equation (34) implements $\mathbb{1}\{\|x-\hat{\mu}\|_2\leq 5R\}\frac{1}{r}\tilde{\tau}_t$. The purpose of the second line in Equation (34) is to match the operation of the Downweighting filter, which, in the t-th round multiplies w_t with $(1-\tilde{\tau}_t(x)/r)^{\ell_t}$ for some power ℓ_t . Finally, we note that $w_{t+1}\tau_t$ are implemented in \mathcal{F} by taking $v_{t,j}$ to be the rows of the matrix \mathbf{M}_t (this is why we need the sums to be on $L'=\max(L,d)$ terms in Equation (34)).

We need to specify the arithmetic complexity m and the dimension of the parameterization k of our family \mathcal{F} . For the first, we have that for any $h \in \mathcal{F}$ and $x \in \mathbb{R}^d$, the value h(x) can be computed using $O(KdL'\ell_{\max})$ standard arithmetic

operations and jumps, where ℓ_{\max} is the maximum exponent that ℓ_t can have and is set to be $\ell_{\max} := \left(\frac{dR}{\delta^2/\epsilon}\right)^{C\log d}$ in Line 19 of Algorithm 5. The dL' comes from the computation of the means $(1/L')\sum_{j=1}^{L'}(v_{t,j}^T(x-a_t))^2$ and the K comes from the fact that we have K factors in the expression of h.

Regarding the other parameter k, we have that every $h \in \mathcal{F}$ is parameterized by O(K) scalars and O(KL') d-dimensional vectors. Thus, k = O(KL'd).

We are now ready to prove Lemma D.9.

Lemma D.9. Consider the setting of Algorithm 5, where B is the distribution of outliers supported in a ball of radius R around μ . Let $r' := \left(CdR^2 + 1 + C_1\delta^2/\epsilon\right)^{C\log d}$ for sufficiently large constant C. Denote by ϵ the contamination rate and let an arbitrary $\epsilon' \in (0,1)$. There exists a set S_{cover} of $N = \frac{1}{\epsilon'^3}d^4K^2L^2(dR\epsilon/\delta^2)^{O(\log d)}$ points x_1,\ldots,x_N lying in the ball of radius R around μ , such that for all $t \in [K]$, for all choices of the vectors $z_{t,j}$ of Line 22 of Algorithm 5 it holds

$$\left| \sum_{X \sim B} \left[\frac{1}{r'} w_{t+1}(X) \tilde{\tau}_t(X) \right] - \frac{1}{N} \sum_{i=1}^N \frac{1}{r'} w_{t+1}(x_i) \tilde{\tau}_t(x_i) \right| \le \epsilon'$$
and
$$\left| \sum_{X \sim B} \left[\frac{1}{r'} w_{t+1}(X) \tau_t(X) \right] - \frac{1}{N} \sum_{i=1}^N \frac{1}{r'} w_{t+1}(x_i) \tau_t(x_i) \right| \le \epsilon'.$$

Proof of Lemma D.9. We use Proposition I.1 for the family \mathcal{F} of Lemma I.2 and plug the upper bounds for the arithmetic complexity m and the dimension of the parameters k. Proposition I.1 states that N can be chosen to be a multiple of

$$\frac{1}{\epsilon'^2}(\log(km) + km\log(1/\epsilon')) .$$

Taking the much looser bound $N = \Theta(\frac{km}{\epsilon'^3})$ suffices for our purposes. Plugging in $k = O(dK \max(L,d))$, $m = dK \max(L,d)(dR\epsilon/\delta^2)^{O(\log d)}$ from Lemma I.2, we get $km = d^2K^2 \max(d^2,L^2)(dR\epsilon/\delta^2)^{O(\log d)} \lesssim d^4K^2L^2(dR\epsilon/\delta^2)^{O(\log d)}$.

For completeness, we provide the proof of Proposition I.1.

Proof of Proposition I.1. We derive the result from the statements of (Anthony & Bartlett, 1999) without explaining all of the definitions of the notions involved. Please see (Anthony & Bartlett, 1999) for more details. Applying Theorem 17.7 (Anthony & Bartlett, 1999) with the loss function $\ell_h(x,y) = h(x)$ we obtain

$$\Pr\left[\sup_{h\in\mathcal{F}}\left|\frac{\mathbf{E}}{X\sim D}[h(X)] - \frac{1}{N}\sum_{i=1}^{N}h(X_i)\right| > \epsilon'\right] \le 4\mathcal{N}_1(\epsilon'/8, \mathcal{F}, 2N)\exp(-\epsilon'^2N/32),$$
(35)

where the probability is taken over a set of N i.i.d. points X_1, \ldots, X_N drawn from D.

To bound from above the covering number $\mathcal{N}_1(\epsilon'/8, \mathcal{F}, 2N)$, we use Theorem 18.4 from (Anthony & Bartlett, 1999) which gives that $\mathcal{N}_1(\epsilon'/8, \mathcal{F}, 2N) \leq e(d'+1)(16e/\epsilon')^{d'}$ where $d' = \operatorname{Pdim}(\mathcal{F})$ is the pseudo-dimension of \mathcal{F} . From that, we conclude that choosing any

$$N > \frac{32}{\epsilon'^2} \log(4e(d'+1)) + \frac{32d'}{\epsilon'^2} \log(16e/\epsilon')$$

makes the probability in Equation (35) less than 1.

It remains to bound d' from above, which can be done as follows. Define the *subgraph class* associated to the family \mathcal{F}

$$\mathcal{B}_{\mathcal{F}} := \{ B_h \mid h \in \mathcal{F} \} .$$

where for any $h \in \mathcal{F}$, $B_h : \mathbb{R}^{d+1} \to \{0,1\}$ is defined as $B_h(x,y) = \mathbb{1}\{h(x) \ge y\}$. The pseudo-dimension is defined to be $\operatorname{Pdim}(\mathcal{F}) = \operatorname{VCdim}(\mathcal{B}_{\mathcal{F}})$ (see Section 11.2 in (Anthony & Bartlett, 1999)). By Theorem 2.3 in (Goldberg & Jerrum,

1995), we have that $\operatorname{VCdim}(\mathcal{B}_{\mathcal{F}}) = O(km)$ since it $\mathcal{B}_{\mathcal{F}}$ functions that are parametrized by vectors of \mathbb{R}^k (same as for family \mathcal{F}) and the functions of $B_h(x,y)$ can be computed using at most m+2 operations (m to compute h and two to do the comparison with y and threshold). Putting everything together, it suffices to choose

$$N = C \frac{1}{\epsilon'^2} (\log(km) + km \log(1/\epsilon'))$$

in order to make the probability in Equation (35) less than 1. In that case, by probabilistic argument, there exists at least one set of N points satisfying the desired event.

Claim D.10. Let S be the cover of Lemma D.9 with r' and ϵ' as defined above. Suppose that the deterministic condition Condition D.5 holds. If $x \in S_{cover}$, then $\tau_t(x) \leq 5\tilde{\tau}_t(x) + (18C_3 + 12/C_2)(\lambda_t/\epsilon) \|\mathbf{M}_t\|_F^2$, where C_3 and C_2 are the constants used in Algorithm 5.

Proof. By Condition D.5 we have that for all the N samples of the cover, $\tilde{g}_t(x) \geq 0.2g_t(x) - 0.8(\delta^2/\epsilon^2) \|\mathbf{M}_t\|_F^2$. Recall the definitions $\tilde{\tau}_t(x) = \tilde{g}(x) \mathbb{1}\{\tilde{g}(x) > C_3 \|\mathbf{U}_t\|_F^2 \hat{\lambda}_t/\epsilon\}$, $\tau_t(x) = g(x) \mathbb{1}\{g(x) > C_3 \|\mathbf{M}_t\|_F^2 \lambda_t/\epsilon\}$. We split into cases based on whether each of g_t , \tilde{g}_t has been zeroed by their thresholding operation:

- If $\tau_t(x)$ has been zeroed, (i.e., $g_t(x) < C_3 \|\mathbf{U}_t\|_F^2 \lambda_t / \epsilon$), the claim trivially holds since the left-hand side is zero.
- If none of $\tilde{\tau}_t(x)$, $\tau_t(x)$ has been zeroed, then $\tilde{\tau}_t(x) = \tilde{g}_t(x)$ and $\tau_t(x) = g_t(x)$, thus the claim holds by the aforementioned fact that $\tilde{g}_t(x) \geq 0.2g_t(x) 0.8(\delta^2/\epsilon^2) \|\mathbf{M}_t\|_F^2$.
- If $\tilde{\tau}_t(x)$ has been zeroed but $\tau_t(x)$ has not, then the worst case is $\tilde{g}_t(x) = 0.2g_t(x) 0.8(\delta^2/\epsilon^2) \|\mathbf{M}_t\|_F^2$. This means that in this case:

$$\tau_{t}(x) \leq \frac{1}{0.2} C_{3} \frac{\hat{\lambda}_{t}}{\epsilon} \|\mathbf{U}_{t}\|_{F}^{2} + 4 \frac{\delta^{2}}{\epsilon^{2}} \|\mathbf{M}_{t}\|_{F}^{2}$$

$$< 18 C_{3} \frac{\lambda_{t}}{\epsilon} \|\mathbf{M}_{t}\|_{F}^{2} + 4 \frac{\delta^{2}}{\epsilon^{2}} \|\mathbf{M}_{t}\|_{F}^{2}$$

$$\leq 18 C_{3} \frac{\lambda_{t}}{\epsilon} \|\mathbf{M}_{t}\|_{F}^{2} + \frac{12}{C_{2}} \frac{\lambda_{t}}{\epsilon} \|\mathbf{M}_{t}\|_{F}^{2},$$

where in the second inequality we used that $\hat{\lambda}_t \leq 3\lambda_t$ and $\|\mathbf{U}_t\|_F^2 \leq 1.2\|\mathbf{M}_t\|_F^2$ due to Condition D.5, and in the last inequality we used that $\delta^2/\epsilon < \hat{\lambda}_t/C_2$ and $\hat{\lambda}_t \leq 3\lambda_t$ (Condition D.5 again).

Remark I.3 (On the choice of K and L). We comment on how the values for the number of iterations K and L that are used in Algorithm 5 are derived. First, the derivation of $K = C \log d \log(dR/(\delta^2/\epsilon))$ for large enough constant C is identical to that of Appendix C.4 (see Equation (4)). We will thus focus on L. We note that in the proof of Lemma D.7 we use Lemma D.9 with $\epsilon' \gtrsim \frac{(\delta^2/\epsilon)^{2\log d}}{\epsilon (CdR^2+1+C_1\delta^2/\epsilon)^{C\log d}}$. This means that the cover S_{cover} of that lemma has size bounded by

$$|S_{cover}| \leq \frac{1}{\epsilon'^3} d^4 K^2 L^2 \left(\frac{dR}{\delta^2/\epsilon}\right)^{O(\log d)} \lesssim \frac{(C dR^2 + 1 + C_1 \delta^2/\epsilon)^{O(\log d)}}{(\delta^2/\epsilon)^{O(\log d)}} L^2 \ .$$

The analog of Lemma C.5 thus requires that L is multiple of $\log\left(\frac{|S_{cover}|+d}{\tau}\right)$, where τ is the desired probability of failure. Note that we have the following (rough) bounds

$$\begin{split} \log \left(\frac{|S_{cover}| + d}{\tau} \right) &\lesssim \log \left(\frac{(L(CdR^2 + 1 + C_1 \delta^2 / \epsilon)^{O(\log d)}}{\tau (\delta^2 / \epsilon)^{O(\log d)}} \right) \\ &\lesssim \log^2(d) \log(CdR^2 + 1 + C_1 \delta^2 / \epsilon) \log \left(\frac{1}{\tau \epsilon} \right) \log(L) \;. \end{split}$$

Thus, we want to choose L such that it holds $L \ge C \log^2(d) \log(CdR^2 + 1 + C_1\delta^2/\epsilon) \log\left(\frac{1}{\tau\epsilon}\right) \log L$. Using the basic fact that for any a > 0, $x \ge 2a \log a \Rightarrow x \ge a \log x$ with $a = C \log^2(d) \log(CdR^2 + 1 + C_1\delta^2/\epsilon) \log\left(\frac{1}{\tau\epsilon}\right)$, it suffices to choose any L satisfying the following

$$L \ge C \log^2(d) \log \left(C dR^2 + 1 + C_1 \frac{\delta^2}{\epsilon} \right) \log \left(\frac{1}{\tau \epsilon} \right) \log \left(\log^2(d) \log \left(C dR^2 + 1 + C_1 \frac{\delta^2}{\epsilon} \right) \log \left(\frac{1}{\tau \epsilon} \right) \right) .$$

We see that the choice in Algorithm 5 satisfies this condition.

I.3. Omitted Proofs from Appendix D.3

Lemma D.14. Let $\mathbf{A}, \mathbf{B}, \mathbf{B}_1, \dots, \mathbf{B}_p$ be symmetric $d \times d$ matrices and define $\mathbf{M} = \mathbf{B}^p, \mathbf{M}_S = \prod_{i=1}^p \mathbf{B}_i$. If $\|\mathbf{B}_i - \mathbf{B}\|_2 \le \delta \|\mathbf{B}\|_2$, then $\|\mathbf{M}_S - \mathbf{B}^p\|_2 \le p\delta(1+\delta)^p \|\mathbf{B}\|_2^p$.

Proof. We have the following:

$$\mathbf{B}^p - \prod_{i=1}^p \mathbf{B}_i = \sum_{i=0}^{p-1} \left(\left(\prod_{j=1}^i \mathbf{B}_j \right) \mathbf{B}^{p-i} - \left(\prod_{j=1}^{i+1} \mathbf{B}_j \right) \mathbf{B}^{p-i-1} \right) = \sum_{i=0}^{p-1} \left(\left(\prod_{j=1}^i \mathbf{B}_j \right) (\mathbf{B} - \mathbf{B}_{i+1}) \mathbf{B}^{p-i-1} \right) .$$

Using that $\|\mathbf{B}_i\|_2 \le (1+\delta)\|\mathbf{B}\|_2$, we obtain the following bound:

$$\begin{aligned} \left\| \mathbf{B}^{p} - \prod_{i=1}^{p} \mathbf{B}_{i} \right\|_{2} &\leq \sum_{i=0}^{p-1} \left\| \left(\prod_{j=1}^{i} \mathbf{B}_{j} \right) (\mathbf{B} - \mathbf{B}_{i+1}) \mathbf{B}^{p-i-1} \right\|_{2} \\ &\leq \sum_{i=0}^{p-1} \left(\left(\prod_{j=1}^{i} \| \mathbf{B}_{j} \| \right) \| \mathbf{B} - \mathbf{B}_{i+1} \|_{2} \| \mathbf{B} \|_{2}^{p-i-1} \right) \\ &\leq \sum_{i=0}^{p-1} \| \mathbf{B} \|_{2}^{p} (1+\delta)^{i} \delta \leq p \delta (1+\delta)^{p} \| \mathbf{B} \|_{2}^{p}. \end{aligned}$$

I.4. Omitted Proofs from Appendix D.3.1

Lemma I.4. For any $\delta, \tau \in (0,1)$ and any distribution D on \mathbb{R}^d with mean μ and covariance matrix Σ , there exists an estimator $\widehat{\mu}$ on $n = O\left((\operatorname{tr}(\Sigma)/\delta^2)\log(1/\tau)\right)$ i.i.d. samples from D, such that $\|\widehat{\mu} - \mu\|_2 = O(\delta)$. Moreover, this $\widehat{\mu}$ can be computed in time $O(nd\log(1/\tau))$ and using memory $O(d\log(1/\tau))$.

Proof. Let X_1, \ldots, X_m be independent samples from D. We first show that the empirical mean $Y := (1/m) \sum_{i=1}^m X_i$, is δ -accurate with constant probability.

$$\mathbf{E}[\|Y - \mu\|_2^2] = \sum_{j=1}^d \mathbf{E}[(Y_j - \mu_j)^2] = \frac{1}{m} \sum_{j=1}^d \mathbf{\Sigma}_{jj} = \frac{\operatorname{tr}(\mathbf{\Sigma})}{m} .$$

By Markov's inequality, we get that

$$\Pr[\|Y - \mu\|_2^2 > \delta^2] \le \frac{\operatorname{tr}(\Sigma)}{m\delta^2} \le \frac{1}{20}$$
, (36)

where the last inequality is true if we choose $m=20\mathrm{tr}(\Sigma)/\delta^2$. Having Equation (36) at hand, the probability of success of the above estimator can be boosted to $1-\tau$ by using Claim C.11. We use that claim with G being the distribution of Y, B=G, $\epsilon=1/20$ and $R=\delta$. This completes the proof.

As a corollary of the above, we obtain the estimators $\hat{\mu}_t$ of Algorithm 5.

Lemma D.11. In the setting of Algorithm 5, there exist estimators $\widehat{\mu}_t$ such that, with probability at least $1 - \tau$, for all $t \in [K]$ we have that $\|\widehat{\mu}_t - \mu_t\|_2 \le \delta/100$. Furthermore, each $\widehat{\mu}_t$ can be computed on a stream of $n = O\left(\frac{R^2}{\delta^2/\epsilon}\log(K/\tau) + \frac{d(1+\delta^2/\epsilon)}{\delta^2}\log(K/\tau)\right)$ independent samples from P_t , in time $O(nd\log(K/\tau))$ and using memory $O(d\log(K/\tau))$.

Proof. We use the estimator of Lemma I.4 with τ/K in place of τ . It remains to bound $\operatorname{tr}(\Sigma_t)$. We have that $d_{\mathrm{TV}}(P_t,G)=1-O(\epsilon)$, thus, by Fact B.4 we can write $P_t=(1-\alpha)G_0+\alpha B$, with $\alpha=O(\epsilon)$ and $G_0(x)=h(x)G(x)/(\int h(x)G(x)\mathrm{d}x)$ some weighted version of the inlier's distribution with $\mathbf{E}_{X\sim G}[h(X)]=1-\alpha$ (same argument that we have used before in the proof of Lemma B.11). We have that

$$\mathbf{\Sigma}_t = (1 - \alpha)\mathbf{\Sigma}_{G_0} + \alpha\mathbf{\Sigma}_B + \alpha(1 - \alpha)(\mu_{G_0} - \mu_B)(\mu_{G_0} - \mu_B)^T.$$

Due to stability, the first term has $\Sigma_{G_0} \leq (1 + \delta^2/\epsilon) \mathbf{I}_d$. For the second term we use that

$$\operatorname{tr}(\mathbf{\Sigma}_B) = \underset{X \sim B}{\mathbf{E}} [\operatorname{tr}((X - \mu_B)(X - \mu_B)^T)] = \underset{X \sim B}{\mathbf{E}} [\|X - \mu_B\|_2^2] \le O(R^2) .$$

We also bound the trace of the last term by $O(\epsilon R^2)$. Therefore, we obtain that $\operatorname{tr}(\Sigma_t) \lesssim d(1+\delta^2/\epsilon) + \epsilon R^2$.

J. Adaptive Choice of Upper Bound on Covariance

In this section, we show that a simple procedure can be used to make the algorithm adaptive to the scale of covariance (such a procedure is useful for some of our applications in Appendix E).

As noted earlier, the definition of stability that we have used so far (Definitions B.8 and B.9) was designed for distributions with covariance matrix comparable to \mathbf{I}_d . In particular, if inliers satisfy $\mathbf{Cov}[X] \leq \mathbf{I}_d$, then our algorithms result in error $O(\sqrt{\epsilon})$. In many practical cases, some of which are encountered in Appendix E, the inliers are much better concentrated, satisfying $\mathbf{Cov}[X] \leq \sigma \mathbf{I}_d$, with σ much smaller than 1. In that case, the optimal asymptotic error is $\Theta(\sigma\sqrt{\epsilon})$. If σ is known beforehand, then a simple preprocessing step allows our algorithms to obtain the error $O(\sigma\sqrt{\epsilon})$. We now describe a procedure using Lepski's method (Lepskii, 1991; Birgé, 2001) that can adapt to the setting when σ is unknown. Concretely, we consider the task of robustly estimating the mean μ of a distribution where inliers have bounded covariance, $\mathbf{Cov}[X] \leq \sigma^2 \mathbf{I}_d$, but σ is unknown to the algorithm.

Let RobustMean $(\tilde{\sigma}, \gamma)$ be any black-box robust mean estimation algorithm, where $\tilde{\sigma}$ is a guess for an upper bound on the covariance of inliers (ideally, we would like to use $\tilde{\sigma} = \sigma$) and γ is the probability of failure. The procedure below tries different values for $\tilde{\sigma}$ in order to find a vector that is as good as the output of RobustMean when run with the best choice of $\tilde{\sigma} = \sigma$. The assumption made here is that σ belongs in some known interval [A, B].

As a small note, a more explicit notation would be $\operatorname{RobustMean}(S, \tilde{\sigma}, \gamma)$, where S is the dataset used, but we omit S because this depends on the data-access model: If a streaming model is assumed, then S necessarily has to be different in each call of the algorithm, otherwise there is no need for using different datasets.

Algorithm 9 Adaptive search for σ

```
1: input: A, B, \gamma, r(\cdot)

2: Denote \tilde{\sigma}_j := B/2^j for j = 0, 1, \ldots, \log(B/A) and set \gamma' := \gamma/\log(B/A).

3: J \leftarrow 0

4: \hat{\mu}^{(0)} \leftarrow \text{RobustMean}(\tilde{\sigma}_0, \gamma')

5: while \tilde{\sigma}_j \geq A and \|\hat{\mu}^{(J)} - \hat{\mu}^{(j)}\|_2 \leq r(\tilde{\sigma}_J) + r(\tilde{\sigma}_j) for all j = 0, 1, \ldots, J-1 do

6: J \leftarrow J+1.

7: \hat{\mu}^{(J)} \leftarrow \text{RobustMean}(\tilde{\sigma}_J, \gamma')

8: end while

9: \hat{J} \leftarrow J-1

return \hat{\mu}^{(\hat{J})}
```

Theorem J.1. Let $\mu \in \mathbb{R}^d$ A, B > 0, $\sigma \in [A, B]$, and a non-decreasing function $r : \mathbb{R}^+ \to \mathbb{R}^+$. Suppose that RobustMean $(\tilde{\sigma}, \gamma)$ is a black-box algorithm which is guaranteed to return a vector $\hat{\mu}$ such that $\|\hat{\mu} - \mu\|_2 \leq r(\tilde{\sigma})$ with probability $1 - \gamma$, whenever $\tilde{\sigma} \geq \sigma$. Then, Algorithm 9 returns $\hat{\mu}^{(\hat{J})}$ such that, with probability at least $1 - \gamma$, we have that $\|\hat{\mu}^{(\hat{J})} - \mu\|_2 \leq 3r(2\sigma)$. Moreover, Algorithm 9 calls RobustMean $O(\log(B/A))$ times with desired failure probability set to $\gamma/\log(B/A)$ and using at most $O(d\log(B/A))$ additional memory.

Proof. For $j=0,1,\ldots,\log(B/A)$, denote by \mathcal{E}_j the event that $\|\widehat{\mu}^{(j)}-\mu\|_2 \leq r(\tilde{\sigma}_j)$. Let J be the index corresponding to the value of the unknown parameter σ , i.e., $\tilde{\sigma}_{J+1} \leq \sigma \leq \tilde{\sigma}_J$. Conditioned on the event $\cap_{j=0}^J \mathcal{E}_j$, we have that $\|\widehat{\mu}^{(j)}-\mu\|_2 \leq r(\tilde{\sigma}_j)$ for all $j=0,1,\ldots,J$. Using the triangle inequality, this gives that $\|\widehat{\mu}^{(J)}-\widehat{\mu}^{(j)}\|_2 \leq r(\tilde{\sigma}_J)+r(\tilde{\sigma}_j)$. This means that the stopping condition of Line 5 is satisfied on round J and thus, if $\widehat{\mu}^{(\widehat{J})}$ denotes the vector returned by the algorithm, we have that

$$\|\widehat{\mu}^{(\widehat{J})} - \widehat{\mu}^{(J)}\|_2 \le r(\widetilde{\sigma}_{\widehat{J}}) + r(\widetilde{\sigma}_J) \le 2r(\widetilde{\sigma}_J) \le 2r(2\sigma)$$
,

where the first inequality uses that the condition of Line 5, the second uses that r is non-decreasing and $\tilde{\sigma}_{\widehat{J}} \leq \tilde{\sigma}_J$, and the last one uses that J was defined to be such that $\tilde{\sigma}_{J+1} \leq \sigma \leq \tilde{\sigma}_J$ so multiplying σ by 2 makes it greater than $\tilde{\sigma}_J$. Using the triangle inequality once more, we get $\|\hat{\mu}^{(\widehat{J})} - \mu\|_2 \leq 3r(2\sigma)$. Finally, by union bound on the events \mathcal{E}_j , the probability of error is upper bounded by $\sum_{j=0}^J \gamma' \leq \gamma$. The additional memory requirement of this algorithm is to store $\{\hat{\mu}_j : j \in \{0, \dots, \log(B/A)\}\}$.

We now state the implications that Theorem J.1 has for Algorithms 3 and 5, given in Appendices C and D:

Corollary J.2. Let A, B > 0. In the setting of Corollary D.3, let $\sigma > 0$ be such that the scaled version $S' = \{x/\sigma : x \in S\}$ of the dataset S is $(C\epsilon, \delta)$ -stable with respect to μ/σ . Assuming that $\sigma \in [A, B]$, there exists an algorithm that given $S, \epsilon, \delta, \tau, A, B$ (but not σ), accesses each point of S at most polylog $(d, 1/\epsilon, 1/\tau, B/A)$ times, runs in time nd polylog $(d, 1/\epsilon, 1/\tau, B/A)$, uses additional memory d polylog $(d, 1/\epsilon, 1/\tau, B/A)$, and outputs a vector $\widehat{\mu}$ such that, with probability at least $1 - \tau$, it holds $\|\mu - \widehat{\mu}\|_2 = O(\sigma\delta)$.

Proof. In order to use the search method of Algorithm 9, we define the procedure RobustMean $(\tilde{\sigma}, \gamma)$ to be the following:

- Let $\tilde{S} = \{x/\tilde{\sigma} : x \in S\}.$
- Let $\tilde{\mu}$ be the vector found by the estimator of Corollary D.3 on \tilde{S} using γ for the desired probability of failure.
- Return $\tilde{\sigma}\tilde{\mu}$.

Theorem J.1 with $r(\tilde{\sigma}) = C'\sigma\delta$, for a sufficiently large C' > 0, implies the correctness. In terms of resources used, Algorithm 9 calls the robust mean estimation algorithm at most $\log(B/A)$ times, and thus the running time gets multiplied by $\log(B/A)$. We also need to store one vector for each call, thus $d\log(B/A)$ additional memory suffices.

Corollary J.3. Let A, B > 0. In the setting of Theorem D.2, let $\sigma > 0$ be such that the distribution D' of the points X/σ , $X \sim D$ is $(C\epsilon, \delta)$ -stable with respect to μ . Assuming that $\sigma \in [A, B]$, there exists an algorithm that given

$$n = O\left(R^2 \max\left(d, \frac{\epsilon}{\delta^2}, \frac{(1 + \delta^2/\epsilon)d}{\delta^2 R^2}, \frac{\epsilon^2 d}{\delta^4}, \frac{R^2 \epsilon^2}{\delta^2}, \frac{R^2 \epsilon^4}{\delta^6}\right) \text{polylog}\left(d, \frac{1}{\epsilon}, \frac{1}{\tau}, R, \frac{B}{A}\right)\right)$$
(37)

samples in a stream according to the model of Definition A.1, and given the parameters $\epsilon, \delta, \tau, A, B$ (but not σ), runs in time $nd \operatorname{polylog}(d, 1/\epsilon, 1/\tau, R, B/A)$, uses additional memory $d \operatorname{polylog}(d, 1/\epsilon, 1/\tau, R, B/A)$, and returns a vector $\widehat{\mu}$ such that, with probability at least $1 - \tau$, it holds $\|\mu - \widehat{\mu}\|_2 = O(\sigma \delta)$.

Finally, we note that a similar search procedure can be used for designing algorithms that are adaptive to the parameter δ when σ is known. However, we will not need this generalization for our applications.

K. Omitted Details from Appendix E

K.1. Proof Sketch of Theorem E.3

We describe how Algorithm 5 can be plugged in the algorithm of (Cheng et al., 2019b). We outline the analysis and describe in more detail only the parts from (Cheng et al., 2019b) that need to be changed. The algorithm is Algorithm 1 from (Cheng et al., 2019b), which remains unchanged. This uses Algorithm 2 as a subroutine, which we replace by our estimator of Algorithm 5.

Regarding the analysis, the proof in (Cheng et al., 2019b) uses two claims that state correctness of the black-box robust mean estimator: Lemma 3.4 and Lemma 3.5. For our case, Lemma 3.4 is replaced by our Theorem D.2 specialized to bounded covariance distributions (also see part 2 of Theorem A.3 which says that the sample complexity of Algorithm 5 for that case is $\tilde{O}(d^2/\epsilon)$).

Lemma 3.5 in (Cheng et al., 2019b) also holds when using our estimator. We restate this as a claim below and provide a proof:

Claim K.1. Let D be a distribution supported on \mathbb{R}^d with unknown mean μ^* and covariance Σ . Let $0 < \gamma < 1$, $0 < \epsilon < \epsilon_0$ for some universal constant ϵ_0 and $\delta = O(\sqrt{\tau \epsilon} + \epsilon \log(1/\epsilon))$ for some $\tau = O(\sqrt{\epsilon})$. Suppose that D has exponentially decaying tails and Σ is close to the identity matrix $\|\Sigma - \mathbf{I}_d\|_2 \le \tau$. Denote $R := \sqrt{(d/\epsilon)(1 + \delta^2/\epsilon)}$. Algorithm 5 uses

$$n = \tilde{O}\left(R^2 \max\left(d, \frac{\epsilon}{\delta^2}, \frac{(1 + \delta^2/\epsilon)d}{\delta^2 R^2}, \frac{\epsilon^2 d}{\delta^4}, \frac{R^2 \epsilon^2}{\delta^2}, \frac{R^2 \epsilon^4}{\delta^6}\right)\right)$$
(38)

samples drawn from D and outputs a hypothesis vector $\hat{\mu}$ such that $\|\hat{\mu} - \mu^*\|_2 = O(\delta)$, with probability $1 - \gamma$. Moreover, this is done in $nd \operatorname{polylog}(d, 1/\epsilon, 1/\gamma)$ time and $d \operatorname{polylog}(d, 1/\epsilon, 1/\gamma)$ space.

Proof. Since D has exponentially decaying tails, we know that D is stable with respect to its mean μ^* and covariance $\Sigma \leq O(1)\mathbf{I}_d$ with parameter $\delta = O(\epsilon \log(1/\epsilon))$ (this follows from the tails of the distribution and Definition B.8). That is, for any weight function $w : \mathbb{R}^d \to [0,1]$ with $\mathbf{E}_{X \sim D}[w(X)] \geq 1 - \epsilon$ we have that

$$\left\|\mu_{w,D} - \mu \right\|_2 \leq \delta \quad \text{and} \quad \left\|\overline{\mathbf{\Sigma}}_{w,D} - \mathbf{\Sigma} \right\|_2 \leq \frac{\delta^2}{\epsilon} \; .$$

We claim that D is $(\epsilon, O(\sqrt{\tau\epsilon} + \epsilon \log(1/\epsilon)))$ -stable in the sense of Definition B.8 (the difference from what written above is that Definition B.8 uses identity matrix in place of Σ). This can be seen by using triangle inequality:

$$\left\|\overline{\Sigma}_{w,D} - \mathbf{I}_{d}\right\|_{2} \leq \left\|\overline{\Sigma}_{w,D} - \Sigma\right\|_{2} + \left\|\Sigma - \mathbf{I}_{d}\right\|_{2} \leq \frac{1}{\epsilon} \left(\delta + \sqrt{\epsilon\tau}\right)^{2}.$$
 (39)

The proof is concluded by recalling the guarantee of Algorithm 5 for $(\epsilon, O(\sqrt{\tau\epsilon} + \epsilon \log(1/\epsilon)))$ -stable distributions and using Claim C.12 for the value of R.

We also note that (Cheng et al., 2019b) uses a fast matrix inversion and multiplication procedure for calculating the rotated versions $Y = \hat{\Sigma}_i^{-1/2} X$ of the samples X. In our case, the run-time of our robust mean-estimation procedure exceeds that of these methods, thus we do not need to use them. We can instead use any numerically stable method that has running time up to $\tilde{O}(d^6)$ and approximates the result within error $\operatorname{poly}(\epsilon \kappa/d)$ (see, e.g., (Banks et al., 2020)). Finally, since Claim K.1 is essentially used for the d^2 -dimensional distributions of the points $Y \otimes Y$, we get the d^4 factor in the final sample complexity, as well as the d^2 factors in the time and space complexity.

K.2. Omitted Proofs from Appendix E.2

Corollary E.6. In the setting of Theorem E.5, suppose that the distribution of gradients satisfies $\mathbf{Cov}[\nabla f(\theta)] \leq \sigma^2 \mathbf{I}_d$ with $\sigma^2 = \alpha^2 \|\theta - \theta^*\|_2^2 + \beta^2$ for all $\theta \in \Theta$, where $\alpha \sqrt{\epsilon} < \tau_\ell$. Assume that the radius of the domain Θ , $r := \max_{\theta \in \Theta} \|\theta\|_2$ is finite. There exists a single-pass streaming algorithm that given $O(T(d^2/\epsilon)\log(1 + \alpha r/\beta)\operatorname{polylog}(d, 1/\epsilon, T/\tau, 1 + \alpha r/\beta))$ samples, runs in time Tnd $\operatorname{polylog}(d, 1/\epsilon, T/\tau, 1 + \alpha r/\beta)$, uses memory d $\operatorname{polylog}(d, 1/\epsilon, T/\tau, 1 + \alpha r/\beta)$, and returns a vector $\widehat{\theta} \in \mathbb{R}^d$ such that $\|\widehat{\theta} - \theta^*\|_2 = O(\sqrt{\epsilon}\beta/(1 - \kappa))$ with probability at least $1 - \tau$.

Proof. This follows by using the estimator of Corollary J.3 in place of $g(\cdot)$ in Algorithm 7. The known bounds for σ , $A \le \sigma \le B$ are $A = \beta$ and $B = 2\alpha r + \beta$, thus $B/A \le 1 + 2\alpha r/\beta$. The distribution of the scaled gradients $\frac{1}{\sigma}\nabla f(\theta)$ is $(C\epsilon, O(\sqrt{\epsilon}))$ -stable. For these parameters, n from Equation (37) gives $n = (d^2/\epsilon)\operatorname{polylog}(d, 1/\epsilon, \tau', 1 + \alpha r/\beta)$, where τ' is the desired probability of failure for each call of the estimator. Setting $\tau' = \tau/T$ ensures that the estimates of all rounds are successful with probability $1 - \tau$. Successful estimates of the gradients are within $O(\sigma\delta) = O((\alpha \|\theta - \theta^*\|_2 + \beta)\sqrt{\epsilon})$ from the true one in Euclidean norm, thus in every round we have an $(\sqrt{\epsilon}\alpha, \sqrt{\epsilon}\beta)$ -gradient estimation (in the sense of Definition E.4). Finally, Theorem E.5 requires the condition $\alpha\sqrt{\epsilon} < \tau_\ell$. Assuming that this is true, that theorem concludes the proof.

Theorem E.12 (Robust Logistic Regression; full version of Theorem A.6). Consider the logistic regression model of Equation (27) with the domain Θ of the unknown regressor being the ball of radius r, for some universal constant r > 0, and suppose that Assumption E.10 holds. Assume that $0 < \epsilon < \epsilon_0$ for a sufficiently small constant ϵ_0 . There is a single-pass streaming algorithm that uses $n = (d^2/\epsilon)$ polylog $(d, 1/\epsilon, 1/\tau)$ samples, runs in time nd polylog $(d, 1/\epsilon, 1/\tau)$, uses memory d polylog $(d, 1/\epsilon, 1/\tau)$, and returns a vector $\widehat{\theta} \in \mathbb{R}^d$ such that $\|\widehat{\theta} - \theta^*\|_2 = O(\sqrt{\epsilon})$ with probability at least $1 - \tau$.

Proof. The algorithm is that of Theorem E.5 using the estimator of Algorithm 5 in place of $g(\cdot)$ in Algorithm 7. The distribution of the gradients is $(C\epsilon, O(\sqrt{\epsilon}))$ -stable because of Lemma E.11. For these stability parameters, a sufficient number of samples is (d^2/ϵ) polylog $(d, 1/\epsilon, T/\tau)$ (see Equation (9) with $\delta = O(\sqrt{\epsilon})$ and $R = O(\sqrt{d})$), where T is the number of iterations over which take a union bound. It thus remains to specify the parameters τ_ℓ, τ_u, k, T .

Using Assumption E.10, we can calculate bounds on the parameters τ_{ℓ}, τ_{u} . For τ_{ℓ} , let v be a unit vector from \mathbb{R}^{d} . Let the event $\mathcal{E}_{v,\theta} := \{(v^{T}X)^{2} \geq c_{1} \text{ and } |\theta^{T}X| \leq 2rC^{2}/c_{2}\}$, where c_{1}, c_{2}, C are the constants from Assumption E.10. The probability of the complement of this event is

$$\Pr[\mathcal{E}_{v,\theta}^c] \le \Pr[(v^T X)^2 < c_1] + \Pr[|\theta^T X| > 2rC^2/c_2] \le 1 - c_2 + c_2/2 \le 1 - c_2/2$$
,

where the first term is bounded using the anti-concentration property and the second is bounded using the concentration property along with Markov's inequality. Thus, using the formula of Equation (28) for the Hessian, we have that

$$v^T \nabla^2 \bar{f}(\theta) v \ge \Pr[\mathcal{E}_{v,\theta}] \underset{X \sim D_x}{\mathbf{E}} \left[\frac{e^{\theta^T X}}{(1 + e^{\theta^T X})^2} (v^T X)^2 \middle| \mathcal{E}_{v,\theta} \right] \ge 0.5 c_2 \frac{e^{2rC^2/c_2}}{(1 + e^{2rC^2/c_2})^2} c_1 .$$

Regarding the upper bound τ_u , using the bounded covariance property we get that $v^T \nabla^2 \bar{f}(\theta) v \leq C^2 \sup_{a \in \mathbb{R}} e^a/(1+e^a)^2 = C^2/4$. Therefore, we can choose the values

$$\tau_{\ell} = 0.5c_2 \frac{e^{2rC^2/c_2}}{(1 + e^{2rC^2/c_2})^2} c_1$$
 and $\tau_u = C^2/4$,

for Algorithm 7. The guarantees of our mean estimator imply that we have an $(0,O(\sqrt{\epsilon}))$ -gradient estimator (in the sense of Definition E.4). Regarding the value of κ , we use Equation (23) with a=0. Since that τ_ℓ,τ_u are positive constants, this means that κ is bounded away from 1. Therefore, we have that the factor $1/(1-\kappa)$ appearing in the final error (Equation (25)) is O(1) and the number of iterations from Equation (24) are upper bounded by $T\lesssim \log_2(\|\theta_0-\theta^*\|_2/\sqrt{\epsilon})\lesssim \log(1/\epsilon)$, where we used that the radius of the domain Θ is r=O(1).

L. Bit Complexity of Algorithm 5

Until this point, we have assumed that our algorithms could save real numbers exactly in a single memory cell and perform calculations involving reals in O(1) time. Thus, by saying that Algorithm 5 uses extra memory at most $d \operatorname{polylog}(d,R,1/\epsilon,1/\tau)$, we meant that it needs to store only that many real numbers. We now describe how the algorithm would work in the most realistic word RAM model, where finite precision numbers can be stored in registers of predetermined word size and operations like addition, subtraction and multiplication are performed in O(1) time. We now show that the previous bound of $d \operatorname{polylog}(d,R,1/\epsilon,1/\tau)$, worsened only by another poly-logarithmic factor, holds for the total number of bits that need to be stored. We begin by clarifying how the input is given to the algorithm.

Definition L.1 (Single-Pass Streaming Model with Oracle Access for Real Inputs). Let S be a fixed set of points in \mathbb{R}^d . The elements of S are revealed one at a time to the algorithm as follows: For each point of S that is about to be revealed,

the algorithm is allowed to query as many bits as it wants from that point with whatever order it wants. The process then continues with the next point in the stream. Each point of S is presented only once to the algorithm in the aforementioned way.

In the reminder of this section, we use the same notation as in Theorem D.2. We assume $R \leq M$ and that $\|\mu\|_2 \leq M$, for some $M = (d/\epsilon)^{\operatorname{polylog}(d/\epsilon)}$ (otherwise, the estimation of the mean with extra memory of the order $d\operatorname{polylog}(d/\epsilon)$ becomes impossible). The modified algorithm for this model is the following: Every input point X is ignored if found to have norm greater than 2M. Otherwise, it is deterministically rounded to an X' so that their difference $X - X' := \eta(X)$ has norm at most η , for some $\eta \leq O(\min\{\delta, \frac{\delta^2}{\epsilon R}, R\})$ (see below for more on this choice of η). The exact same algorithm as Algorithm 5 is run on these rounded points.

Correctness First, we note the rejection step removes less than an ϵ -fraction of the input, thus the resulting distribution has not changed more than ϵ in total variation distance from the original one. Moreover, the distribution of the rounded points has essentially the same stability property required by our theorem. Concretely, if we choose the rounding error η to be $\eta = O(\min\{\delta, \frac{\delta^2}{\epsilon M}, M\})$, then it can be seen (Lemma L.2 below) that the distribution of the rounded points is $(\epsilon, O(\delta))$ -stable and $\Pr_{X'}[\|X' - \mu\|_2 = O(R)] \ge 1 - \epsilon$, which are the only assumptions needed for Algorithm 5 to provide an accurate estimate up to $O(\delta)$ error.

Lemma L.2. Fix $0 < \epsilon < 1/2$ and $\delta \ge \epsilon$. Let G be an (ϵ, δ) -stable distribution with respect to some vector $\mu \in \mathbb{R}^d$ and assume G is a distribution such that $\|X - \mu\|_2 \le M$ almost surely for some M > 0. For any deterministic function $\eta : \mathbb{R}^d \to \mathbb{R}^d$ with $\|\eta(x)\|_2 \le \eta$ for all x in the support of G, if G' denotes the distribution of the points $X' = X + \eta(X)$, where $X \sim G$, then G' is $(\epsilon, O(\delta + \eta + \sqrt{\epsilon \eta M}))$ -stable with respect to μ .

Proof. We check the two conditions for stability. Let a weight function $w : \mathbb{R}^d \to [0,1]$ with $\mathbf{E}_{X \sim G}[w(X)] \ge 1 - \epsilon$ and let $\delta' = O(\delta + \eta + \sqrt{\epsilon \eta M})$. We have that

$$\|\mu_{w,G'} - \mu\|_{2} \leq \|\mu_{w,G'} - \mu_{w,G}\|_{2} + \|\mu_{w,G} - \mu\|_{2}$$

$$\leq \left\| \int_{\mathbb{R}^{d}} (x + \eta(x)) \frac{w(x)G(x)}{\mathbf{E}_{X \sim G}[w(X)]} dx - \mu_{w,G} \right\|_{2} + \delta$$

$$\leq \left\| \int_{\mathbb{R}^{d}} \eta(x) \frac{w(x)G(x)}{\mathbf{E}_{X \sim G}[w(X)]} dx \right\|_{2} + \delta$$

$$\leq \eta + \delta \leq \delta',$$

where first inequality uses the triangle inequality and the second one uses the stability of G. Regarding the second stability condition, we have the following:

$$\begin{split} \left\| \overline{\mathbf{\Sigma}}_{w,G'} - \mathbf{I}_{d} \right\|_{2} &\leq \left\| \overline{\mathbf{\Sigma}}_{w,G'} - \overline{\mathbf{\Sigma}}_{w,G} \right\|_{2} + \left\| \overline{\mathbf{\Sigma}}_{w,G} - \mathbf{I}_{d} \right\|_{2} \\ &\leq \left\| \int_{\mathbb{R}^{d}} (x - \mu + \eta(x))(x - \mu + \eta(x))^{T} \frac{w(x)G(x)}{\mathbf{E}_{X \sim G}[w(X)]} \mathrm{d}x \right\|_{2} + \frac{\delta^{2}}{\epsilon} \\ &\leq \left\| \int_{\mathbb{R}^{d}} (x - \mu)\eta(x)^{T} \frac{w(x)G(x)}{\mathbf{E}_{X \sim G}[w(X)]} \mathrm{d}x \right\|_{2} + \left\| \int_{\mathbb{R}^{d}} (x - \mu)^{T} \eta(x) \frac{w(x)G(x)}{\mathbf{E}_{X \sim G}[w(X)]} \mathrm{d}x \right\|_{2} \\ &+ \left\| \int_{\mathbb{R}^{d}} \eta(x)\eta(x)^{T} \frac{w(x)G(x)}{\mathbf{E}_{X \sim G}[w(X)]} \mathrm{d}x \right\|_{2} + \frac{\delta^{2}}{\epsilon} \\ &\leq 2M\eta + \eta^{2} + \frac{\delta^{2}}{\epsilon} \leq \frac{\delta'^{2}}{\epsilon} , \end{split}$$

where we used stability of G, triangle inequality and the bounds $||x - \mu||_2 \le M$, $||\eta(x)||_2 \le \eta$.

Total Bits of Memory Used In order for the differences $X-X':=\eta(X)$ to have $\|\eta(X)\|_2 \leq \eta$ for all X, it is sufficient to round every coordinate to absolute error $O(\eta/\sqrt{d})$. Recall that by our assumption on the a priori bound on the norm of the true mean and the way that we reject input points of large norm, we know that all points surviving will have norm at most 2M. Thus, each coordinate of such points can be stored in a word of $O(\log(Md/\eta))$ bits after being rounded to accuracy η . Therefore, each d-dimensional point that the algorithm will need to manipulate can be stored using d registers

of size $O(\log(Md/\eta))$. However, we need to show that the results of all intermediate calculations can be calculated in low memory. We show the following result to this end:

Claim L.3. In the context of Theorem D.2, given a stream of d-dimensional points, where each coordinate has bit complexity B, Algorithm 5 can be implemented in a word RAM machine using $d \operatorname{polylog}(d, 1/\epsilon, 1/\tau, R)$ many of registers of size $B \operatorname{polylog}(d, 1/\epsilon, 1/\tau, R)$.

Proof Sketch. Multiplying two numbers of bit complexity B_1 and B_2 may result in bit complexity $B_1 + B_2$. Adding k numbers of bit complexity B, may make the resulting bit complexity $B + \log(k)$. We need to check that every step of the algorithm performs calculations that cannot cause the bit complexity to grow by more than poly-log factors.

Line 7 performs only comparisons and counting. Regarding Line 23: As pointed out at the beginning of Appendix D, the vector $v_{t,j} \leftarrow \widehat{\mathbf{M}}_t z_{t,j}$ is calculated by multiplying $z_{t,j}$ by $\widehat{\mathbf{B}}_{t,k}$ for $k=1,\ldots,\log d$ iteratively. Consider a single iteration, say the first one. Performing $\widehat{\mathbf{B}}_{t,k} z_{t,j}$ involves calculating $\frac{1}{n} \left(\sum_x x x^T \right) z_{t,j}$ (see Appendix D.3.2), which can be done as $\frac{1}{n} \sum_x x (x^T z_{t,j})$, i.e., calculating the inner products $x^T z_{t,j}$ first). A single inner product of that form is just a sum of d numbers of bit complexity B with appropriate signs, thus the bit complexity increases only by $O(\log d)$. Finally, multiplying by x and taking the mean over for all of the x's can add only another $O(B + \log(n))$. Since the number of iterations of such calculations is $\log d$, the final result has the claimed bit complexity.

Regarding the Downweighting filter (Algorithm 6), there are a couple of places where the weights w_t are involved in calculations. We note that Algorithm 4 stores only the counts ℓ_t , which fit in registers of size $\log(\ell_{\max}) = \operatorname{polylog}(d, 1/\epsilon, R)$. These counts are used to calculate $w_t(x)$ as $w_t(x) = \prod_{t' \leq t} (1 - \tilde{\tau}_{t'}(x)/r)^{\ell_{t'}}$ whenever there is such a need. An exact calculation would require operations of the form x^y , for some $x \in [0,1]$ and $y \in [\ell_{\max}]$, i.e., exponentiation of a real number. In fact, as we will show later on, instead of calculating $w_t(x)$ with perfect accuracy, it suffices to use an approximate value of $w_t(x) \pm \eta$ for some error $|\eta| < \operatorname{poly}(1/d, 1/R, \epsilon, \tau)^{\log d}$. This will allow us to calculate a good enough approximation in $\operatorname{polylog}(d, R, 1/\epsilon, 1/\tau)$ bits as follows: we can use exponentiation by squaring algorithm for calculating $w_t(x)$'s and round the result in each step to make it fit into our registers. We first explain this in more detail below.

Claim L.4. Let $x \in [0,1]$, $y \in \mathbb{Z}_+$, and assume that both x,y have bit complexity B. The power x^y can be calculated up to a rounding error of $2^{-\Omega(B)}$ in the word RAM model that uses registers of size 2B. Furthermore, this can be done in O(B) standard arithmetic operations.

Proof. We can use exponentiation by squaring: This consists of writing x in binary as $b_k \cdots b_0$ for $k = \log y$ and calculating the sequence r_{k+1}, \ldots, r_0 as $r_{k+1} = 1$, $r_i = r_{k+1}^2 x^{b_i}$ for $i = k, \ldots, 0$. We assume every r_i gets rounded to 2B bits. Because of the rounding, we incur error 2^{-2B} in each round. However, the error of the previous rounds gets amplified, since the result of that round (true value plus error) gets squared. We consider one such iteration to see how that sequence of errors grows: In the t-th iteration, let res_{t-1} denote the true result (before rounding) from the previous round and η_t the rounding error of that round (i.e., $r_t = \operatorname{res}_t + \eta_t$). Then, we have that

$$\operatorname{res}_t + \eta_t := (\operatorname{res}_{t-1} + \eta_{t-1})^2 + 2^{-2B} \le \operatorname{res}_{t-1}^2 + \eta_{t-1}^2 + 2\eta_{t-1} + 2^{-2B}$$

where w.l.o.g. we assume that $\operatorname{res}_t \leq 1$ always. Thus, the rounding error grows as $\eta_t \leq \eta_{t-1}^2 + 2\eta_{t-1} + 2^{-2B} \leq 3\eta_{t-1} + 2^{-2B}$. In the first round, we start with rounding error of 2^{-2B} . Thus, after $k = \log(y) = B$ rounds, the final error is $\eta_k \leq 2^{-\Omega(B)}$.

We continue with examining how fine approximations for w_t are needed. First, in Appendix D.3.2, we use the estimator $\widehat{W}_t = \mathbf{E}_{X \sim \mathcal{U}(S_0)}[w_t(X)]$, which we require to be η -close to $\mathbf{E}_{X \sim P}[w_t(X)]$ (Equation (17)) for some $\eta > \text{poly}(1/d, 1/R, \epsilon, \tau)$. Therefore, when calculating w_t , it suffices to round the intermediate results to error η . This would mean using Claim L.4 with $B = O(\log((1/d, 1/R, \epsilon, \tau))$.

Second, the weights w_t are also used in evaluating the stopping condition of the Downweighting filter. Line 3 of that filter is implemented using the estimator of Lemma D.16. As it can be seen in Equation (20), it suffices to use rounded versions of w_t in $(1/n)\sum_{i=1}^N w_t(X_i)\tilde{\tau}_t(X_i)$, as long as it does not change the result by an additive factor of $c\hat{\lambda}_t \|\mathbf{U}_t\|_F^2$, for a small constant c. Since $\tilde{\tau}_t(X_i) = O(dR^{2+4\log d})$ (Equation (2)) and $\hat{\lambda}_t \|\mathbf{U}_t\|_F^2 > (\delta^2/\epsilon)^{\Theta(\log d)}$ (otherwise, the algorithm has terminated), we can again round w_t up to error $\operatorname{poly}(1/d, 1/R, \epsilon)^{\log(d)}$. This means that the results of these calculations fit into $\operatorname{polylog}(d, R, 1/\epsilon, R)$ bits.

Finally, there are two places in Algorithm 5 where we need to simulate samples from the weighted distribution P_{w_t} : (i) Line 17, whose implementation is outlined in Appendix D.3 and (ii) Line 31. We focus on the first one since the argument for the other case is identical. To simulate P_{w_t} , we use rejection sampling, as described at the beginning of Appendix D.3, with the only difference that we use the rounded versions of the weights w_t . We are thus essentially simulating samples from a slightly different distribution $P_{\widehat{w}_t}$. However, this is close to P_{w_t} in total variation distance, as shown below.

Claim L.5. Let P be a distribution on \mathbb{R}^d and let P_w denote the weighted version of P according to the function $w: \mathbb{R}^d \to [0,1]$, i.e., $P_w(x) = w(x)P(x)/\int_{\mathbb{R}^d} w(x)P(x)\mathrm{d}x$. For any $w, \widehat{w}: \mathbb{R}^d \to [0,1]$ such that $\int_{\mathbb{R}^d} w(x)P(x)\mathrm{d}x \geq 1/2$ and $\sup_{x \in \mathbb{R}^d} |\widehat{w}(x) - w(x)| \leq \xi$ with $\xi \leq 1/8$, it holds that $d_{\mathrm{TV}}(P_{\widehat{w}}, P_w) \leq 8\xi$.

Proof. First, letting the normalization factors $\widehat{C} := \int_{\mathbb{R}^d} \widehat{w}(x) P(x) dx$ and $C := \int_{\mathbb{R}^d} w(x) P(x) dx$, we note that $|C - \widehat{C}| \le \xi$. Letting $\Delta w(x) := \widehat{w}(x) - w(x)$ and $\Delta C := \widehat{C} - C$, we have that

$$d_{\text{TV}}(P_{\widehat{w}}, P_w) = \frac{1}{2} \int_{\mathbb{R}^d} |P_{\widehat{w}}(x) - P_w(x)| \, \mathrm{d}x = \frac{1}{2} \int_{\mathbb{R}^d} \left| \frac{w(x) + \Delta w(x)}{C + \Delta C} - \frac{w(x)}{C} \right| P(x) \, \mathrm{d}x$$

$$\leq \frac{1}{2} \int_{\mathbb{R}^d} \left| \frac{Cw(x) + C\Delta w(x) - Cw(x) - \Delta Cw(x)}{C^2 + C\Delta C} \right| P(x) \, \mathrm{d}x$$

$$\leq 4 \int_{\mathbb{R}^d} |C\Delta w(x) - \Delta Cw(x)| P(x) \, \mathrm{d}x \leq 8\xi ,$$

where in the last line, we first use that $C^2 + C\Delta C \ge 1/4 - \xi \ge 1/8$ (since $1/2 \le C \le 1$ and $0 \le \xi \le 1/8$) and then we use that $|C\Delta w(x) - \Delta Cw(x)| \le |\Delta w(x)| + |\Delta C| \le 2\xi$.

As N (more than one) samples are drawn from P_t in the t-th iteration (see Appendix D.3), we require the joint distribution of these N samples from P_{w_t} and $P_{\hat{w}_t}$ to be within total variation τ (the probability under which the conclusion of Lemma D.13 holds true). This bound on the total variation distance implies that Lemma D.13 continues to hold for $P_{\hat{w}_t}$ with an additional failure probability of τ . To do this, we use Claim L.5 with $\xi = \Theta(\tau/N)$, which means that these rounded weights have bit complexity $\Theta(\log \xi) = \operatorname{polylog}(d, R, 1/\epsilon, 1/\tau)$.