

Fast recovery from a union of subspaces

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Abstract

We address the problem of recovering a high-dimensional but structured vector from linear observations in a general setting where the vector can come from an arbitrary union of subspaces. This setup includes well-studied problems such as compressive sensing and low-rank matrix recovery. We show how to design more efficient algorithms for the union-of-subspace recovery problem by using *approximate* projections. Instantiating our general framework for the low-rank matrix recovery problem gives the fastest provable running time for an algorithm with optimal sample complexity. Moreover, we give fast approximate projections for 2D histograms, another well-studied low-dimensional model of data. We complement our theoretical results with experiments demonstrating that our framework also leads to improved time and sample complexity empirically.

1 Introduction

Over the past decade, exploiting low-dimensional structure in high-dimensional problems has become a highly active area of research in machine learning, signal processing, and statistics. In a nutshell, the general approach is to utilize a low-dimensional model of relevant data in order to achieve better prediction, compression, or estimation compared to a “black box” treatment of the ambient high-dimensional space. For instance, the seminal work on compressive sensing and sparse linear regression has shown how to estimate a sparse, high-dimensional vector from a small number of linear observations that essentially depends only on the small sparsity of the vector, as opposed to its large ambient dimension. Further examples of low-dimensional models are low-rank matrices, group-structured sparsity, and general union-of-subspaces models, all of which have found applications in problems such as matrix completion, principal component analysis, compression, and clustering.

These low-dimensional models have a common reason for their success: they capture important structure present in real world data with a formal concept that is suitable for a rigorous mathematical analysis. This combination has led to statistical performance improvements in several applications where the ambient high-dimensional space is too large for accurate estimation from a limited number of samples. However, exploiting the low-dimensional structure also comes at a cost: incorporating the structural constraints into the statistical estimation procedure often results in a more challenging algorithmic problems. Given the growing size of modern data sets, even problems that are solvable in polynomial time can quickly become infeasible. This leads to the following important question: Can we design efficient algorithms that combine (near)-optimal statistical efficiency with good computational complexity?

In this paper, we make progress on this question in the context of recovering a low-dimensional vector from noisy linear observations, which is the fundamental problem underlying both low-rank matrix recovery and compressive sensing / sparse linear regression. While there is a wide range of

algorithms for these problems, two approaches for incorporating structure tend to be most common: (i) convex relaxations of the low-dimensional constraint such as the ℓ_1 - or the nuclear norm [19], and (ii) iterative methods based on projected gradient descent, e.g., the IHT (Iterative Hard Thresholding) or SVP (Singular Value Projection) algorithms [5, 15]. Since the convex relaxations are often also solved with first order methods (e.g., FISTA or SVT [6]), the low-dimensional constraint enters both approaches through a structure-specific projection or proximal operator. However, this projection / proximal operator is often computationally expensive and dominates the overall time complexity (e.g., it requires a singular value decomposition for the low-rank matrix recovery problem).

In this work, we show how to reduce the computational bottleneck of the projection step by using *approximate* projections. Instead of solving the structure-specific projection exactly, our framework allows us to employ techniques from approximation algorithms without increasing the sample complexity of the recovery algorithm. While approximate projections have been used in prior work, our framework is the first to yield provable algorithms for general union-of-subspaces models (such as low-rank matrices) that combine *better running time* with *no loss in sample complexity* compared to their counterparts utilizing exact projections. Overall, we make three contributions:

1. We introduce an algorithmic framework for recovering vectors from linear observations given an arbitrary *union-of-subspaces* model. Our framework only requires approximate projections, which leads to recovery algorithms with significantly better time complexity.
2. We instantiate our framework for the well-studied *low-rank matrix recovery problem*, which yields a provable algorithm combining the optimal sample complexity with the best known time complexity for this problem.
3. We also instantiate our framework for the problem of recovering *2D-histograms* (i.e., piecewise constant matrices) from linear observations, which leads to a better empirical sample complexity than the standard approach based on Haar wavelets.

Our algorithmic framework generalizes recent results for structured sparse recovery [12, 13] and shows that approximate projections can be employed in a wider context. We believe that these notions of approximate projections are useful in further constrained estimation settings and have already obtained preliminary results for structured sparse PCA. For conciseness, we focus on the union-of-subspaces recovery problem in this paper.

Outline of the paper. In Section 2, we formally introduce the union-of-subspaces recovery problem and state our main results. Section 3 then explains our algorithmic framework in more detail and Section 4 instantiates the framework for low-rank matrix recovery. Section 5 concludes with experimental results. Due to space constraints, we address our results for 2D histograms mainly in Appendix C of the supplementary material.

2 Our contributions

We begin by defining our problem of interest. Our goal is to recover an unknown, structured vector $\theta^* \in \mathbb{R}^d$ from linear observations of the form

$$y = X\theta^* + e, \tag{1}$$

where the vector $y \in \mathbb{R}^n$ contains the linear observations / measurements, the matrix $X \in \mathbb{R}^{n \times d}$ is the design / measurement matrix, and the vector $e \in \mathbb{R}^n$ is an arbitrary noise vector. The formal goal is to find an estimate $\hat{\theta} \in \mathbb{R}^d$ such that $\|\hat{\theta} - \theta^*\|_2 \leq C \cdot \|e\|_2$, where C is a fixed, universal constant and $\|\cdot\|_2$ is the standard ℓ_2 -norm (for notational simplicity, we omit the subscript on the ℓ_2 -norm in the rest of the paper). The structure we assume is that the vector θ^* belongs to a *subspace model*:

Definition 1 (Subspace model). *A subspace model \mathbb{U} is a set of linear subspaces. The set of vectors associated with the subspace model \mathbb{U} is $\mathcal{M}(\mathbb{U}) = \{\theta \mid \theta \in U \text{ for some } U \in \mathbb{U}\}$.*

A subspace model is a natural framework generalizing many of the low-dimensional data models mentioned above. For example, the set of sparse vectors with s nonzeros can be represented with $\binom{d}{s}$ subspaces corresponding to the $\binom{d}{s}$ possible sparse support sets. The resulting problem of recovering θ^* from observations of the form (1) then is the standard compressive sensing / sparse linear regression problem. Structured sparsity is a direct extension of this formulation in which we only include a smaller set of allowed supports, e.g., supports corresponding to group structures.

Our framework also includes the case where the union of subspaces is taken over an infinite set: we can encode the low-rank matrix recovery problem by letting \mathbb{U} be the set of rank- r matrix subspaces, i.e., each subspace is given by a set of r orthogonal rank-one matrices. By considering the singular value decomposition, it is easy to see that every rank- r matrix can be written as the linear combination of r orthogonal rank-one matrices.

Next, we introduce related notation. For a linear subspace U of \mathbb{R}^d , let $P_U \in \mathbb{R}^{d \times d}$ be the orthogonal projection onto U . We denote the orthogonal complement of the subspace U with U^\perp so that $\theta = P_U\theta + P_{U^\perp}\theta$. We extend the notion of adding subspaces (i.e., $U + V = \{u + v \mid u \in U \text{ and } v \in V\}$) to subspace models: the sum of two subspace models \mathbb{U} and \mathbb{V} is $\mathbb{U} \oplus \mathbb{V} = \{U + V \mid U \in \mathbb{U} \text{ and } V \in \mathbb{V}\}$. We denote the k -wise sum of a subspace model with $\oplus^k \mathbb{U} = \mathbb{U} \oplus \mathbb{U} \oplus \dots \oplus \mathbb{U}$.

Finally, we introduce a variant of the well-known restricted isometry property (RIP) for subspace models. The RIP is a common regularity assumption for the design matrix X that is often used in compressive sensing and low-rank matrix recovery in order to decouple the analysis of algorithms from concrete sampling bounds.¹ Formally, we have:

Definition 2 (Subspace RIP). *Let $X \in \mathbb{R}^{n \times d}$, let \mathbb{U} be a subspace model, and let $\delta \geq 0$. Then X satisfies the (\mathbb{U}, δ) -subspace RIP if for all $\theta \in \mathcal{M}(\mathbb{U})$ we have $(1 - \delta)\|\theta\|^2 \leq \|X\theta\|^2 \leq (1 + \delta)\|\theta\|^2$.*

2.1 A framework for recovery algorithms with approximate projections

Considering the problem (1) and the goal of estimating under the ℓ_2 -norm, a natural algorithm is projected gradient descent with the constraint set $\mathcal{M}(\mathbb{U})$. This corresponds to iterations of the form

$$\hat{\theta}^{i+1} \leftarrow P_{\mathbb{U}}(\hat{\theta}^i - \eta \cdot X^T(X\hat{\theta}^i - y)) \quad (2)$$

where $\eta \in \mathbb{R}$ is the step size and we have extended our notation so that $P_{\mathbb{U}}$ denotes a projection onto the set $\mathcal{M}(\mathbb{U})$. Hence we require an oracle that projects an arbitrary vector $b \in \mathbb{R}^d$ into a subspace model \mathbb{U} , which corresponds to finding a subspace $U \in \mathbb{U}$ so that $\|b - P_U b\|$ is minimized. Recovery algorithms of the form (2) have been proposed for various instances of the union-of-subspaces recovery

¹Note that exact recovery from arbitrary linear observations is already an NP-hard problem in the noiseless case, and hence regularity conditions on the design matrix X are necessary for efficient algorithms. While there are more general regularity conditions such as the restricted eigenvalue property, we state our results here under the RIP assumption in order to simplify the presentation of our algorithmic framework.

problem and are known as Iterative Hard Thresholding (IHT) [5], model-IHT [1], and Singular Value Projection (SVP) [15]. Under regularity conditions on the design matrix X such as the RIP, these algorithms find accurate estimates $\hat{\theta}$ from an asymptotically optimal number of samples. However, for structures more complicated than plain sparsity (e.g., group sparsity or a low-rank constraint), the projection oracle is often the computational bottleneck.

To overcome this barrier, we propose two complementary notions of *approximate* subspace projections. Note that for an exact projection, we have that $\|b\|^2 = \|b - P_{\mathbb{U}}b\|^2 + \|P_{\mathbb{U}}b\|^2$. Hence minimizing the “tail” error $\|b - P_{\mathbb{U}}b\|$ is equivalent to maximizing the “head” quantity $\|P_{\mathbb{U}}b\|$. Instead of minimizing / maximizing these quantities exactly, the following definitions allow a *constant factor* approximation:

Definition 3 (Approximate tail projection). *Let \mathbb{U} and $\mathbb{U}_{\mathcal{T}}$ be subspace models and let $c_{\mathcal{T}} \geq 0$. Then $\mathcal{T} : \mathbb{R}^d \rightarrow \mathbb{U}_{\mathcal{T}}$ is a $(c_{\mathcal{T}}, \mathbb{U}, \mathbb{U}_{\mathcal{T}})$ -approximate tail projection if the following guarantee holds for all $b \in \mathbb{R}^d$: The returned subspace $U = \mathcal{T}(b)$ satisfies $\|b - P_U b\| \leq c_{\mathcal{T}} \|b - P_{\mathbb{U}}b\|$.*

Definition 4 (Approximate head projection). *Let \mathbb{U} and $\mathbb{U}_{\mathcal{H}}$ be subspace models and let $c_{\mathcal{H}} > 0$. Then $\mathcal{H} : \mathbb{R}^d \rightarrow \mathbb{U}_{\mathcal{H}}$ is a $(c_{\mathcal{H}}, \mathbb{U}, \mathbb{U}_{\mathcal{H}})$ -approximate head projection if the following guarantee holds for all $b \in \mathbb{R}^d$: The returned subspace $U = \mathcal{H}(b)$ satisfies $\|P_U b\| \geq c_{\mathcal{H}} \|P_{\mathbb{U}}b\|$.*

It is important to note that the two definitions are distinct in the sense that a constant-factor head approximation does not imply a constant-factor tail approximation, or vice versa (to see this, consider a vector with a very large or very small tail error, respectively). Another feature of these definitions is that the approximate projections are allowed to choose subspaces from a potentially larger subspace model, i.e., we can have $\mathbb{U} \subsetneq \mathbb{U}_{\mathcal{H}}$ (or $\mathbb{U}_{\mathcal{T}}$). This is a useful property when designing approximate head and tail projection algorithms as it allows for *bicriterion* approximation guarantees.

We now state the main result for our new recovery algorithm. In a nutshell, we show that using *both* notions of approximate projections achieves the same statistical efficiency as using exact projections. As we will see in later sections, the weaker approximate projection guarantees allow us to design algorithms with a significantly better time complexity than their exact counterparts. To simplify the following statement, we defer the precise trade-off between the approximation ratios to Section 3.

Theorem 5 (informal). *Let \mathcal{H} and \mathcal{T} be approximate head and tail projections with constant approximation ratios, and let the matrix X satisfy the $(\oplus^c \mathbb{U}, \delta)$ -subspace RIP for a sufficiently large constant c and a sufficiently small constant δ . Then there is an algorithm AS-IHT that returns an estimate $\hat{\theta}$ such that $\|\hat{\theta} - \theta^*\| \leq C\|e\|$. The algorithm requires $O(\log\|\theta\|/\|e\|)$ multiplications with X and X^T , and $O(\log\|\theta\|/\|e\|)$ invocations of \mathcal{H} and \mathcal{T} .*

Up to constant factors, the requirements on the RIP of X in Theorem 5 are the same as for exact projections. As a result, our sample complexity is only affected by a constant factor through the use of approximate projections, and our experiments in Section 5 show that the empirical loss in sample complexity is negligible. Similarly, the number of iterations $O(\log\|\theta\|/\|e\|)$ is also only affected by a constant factor compared to the use of exact projections [5, 15]. Finally, it is worth mentioning that using two notions of approximate projections is crucial: prior work in the special case of structured sparsity has already shown that only one type of approximate projection is not sufficient for strong recovery guarantees [13].

2.2 Low-rank matrix recovery

We now instantiate our new algorithmic framework for the low-rank matrix recovery problem. Variants of this problem are widely studied in machine learning, signal processing, and statistics, and are known under different names such as matrix completion, matrix sensing, and matrix regression. As mentioned above, we can incorporate the low-rank matrix structure into our general union-of-subspaces model by considering the union of all low-rank matrix subspaces. For simplicity, we state the following bounds for the case of square matrices, but all our results also apply to rectangular matrices. Formally, we assume that $\theta^* \in \mathbb{R}^d$ is the vectorized form of a rank- r matrix $\Theta^* \in \mathbb{R}^{d_1 \times d_1}$ where $d = d_1^2$ and typically $r \ll d_1$. Seminal results have shown that it is possible to achieve the subspace-RIP for low-rank matrices with only $n = O(r \cdot d_1)$ linear observations, which can be much smaller than the total dimensionality of the matrix d_1^2 . However, the bottleneck in recovery algorithms is often the singular value decomposition (SVD), which is necessary for both exact projections and soft thresholding operators and has a time complexity of $O(d_1^3)$.

Our new algorithmic framework for approximate projections allows us to leverage recent results on *approximate* SVDs. We show that it is possible to compute both head and tail projections for low-rank matrices in $\tilde{O}(r \cdot d_1^2)$ time, which is significantly faster than the $O(d_1^3)$ time for an exact SVD in the relevant regime where $r \ll d_1$. Overall, we get the following result.

Theorem 6. *Let $X \in \mathbb{R}^{n \times d}$ be a matrix with subspace-RIP for low-rank matrices, and let T_X denote the time to multiply a d -dimensional vector with X or X^T . Then there is an algorithm that recovers an estimate $\hat{\theta}$ such that $\|\hat{\theta} - \theta^*\| \leq C\|e\|$. Moreover, the algorithm runs in time $\tilde{O}(T_X + r \cdot d_1^2)$.*

In the regime where multiplication with the matrix X is fast, the time complexity of the projection dominates the time complexity of the recovery algorithms. For instance, structured observations such as a subsampled Fourier matrix achieve $T_X = \tilde{O}(d_1^2)$; see Appendix D for details. Here, our algorithm runs in time $\tilde{O}(r \cdot d_1^2)$, which is the first provable running time faster than the $O(d_1^3)$ bottleneck given by a single exact SVD. While prior work has suggested the use of approximate SVDs in low-rank matrix recovery [9], our results are the first that give a provably better time complexity for this combination of projected gradient descent and approximate SVDs. Hence Theorem 6 can be seen as a theoretical justification for the heuristic use of approximate SVDs.

Finally, we remark that Theorem 6 does not directly cover the low-rank matrix completion case because the subsampling operator does not satisfy the low-rank RIP [9]. To clarify our use of approximate SVDs, we focus on the RIP setting in our proofs, similar to recent work on low-rank matrix recovery [7, 22]. We believe that similar results as for SVP [15] also hold for our algorithm, and our experiments in Section 5 show that our algorithm works well for low-rank matrix completion.

2.3 2D-histogram recovery

Next, we instantiate our new framework for 2D-histograms, another natural low-dimensional model. As before, we think of the vector $\theta^* \in \mathbb{R}^d$ as a matrix $\Theta \in \mathbb{R}^{d_1 \times d_1}$ and assume the square case for simplicity (again, our results also apply to rectangular matrices). We say that Θ is a k -histogram if the coefficients of Θ can be described as k axis-aligned rectangles on which Θ is constant. This definition is a generalization of 1D-histograms to the two-dimensional setting and has found applications in several areas such as databases and density estimation. Moreover, the theoretical computer science community has studied sketching and streaming algorithms for histograms, which is essentially the problem of recovering a histogram from linear observations. While the wavelet tree model with Haar

wavelets give the correct sample complexity of $n = O(k \log d)$ for 1D-histograms, the wavelet tree approach incurs a *suboptimal* sample complexity of $O(k \log^2 d)$ for 2D-histograms. It is possible to achieve the optimal sample complexity $O(k \log d)$ also for 2D-histograms, but the corresponding exact projection requires a complicated dynamic program (DP) with time complexity $O(d_1^5 k^2)$, which is impractical for all but very small problem dimensions [18].

We design significantly faster *approximate* projection algorithms for 2D histograms. Our approach is based on an approximate DP [18] that we combine with a Lagrangian relaxation of the k -rectangle constraint. Both algorithms have parameters for controlling the trade-off between the size of the output histogram, the approximation ratio, and the running time. As mentioned above, the bicriterion nature of our approximate head and tail guarantees becomes useful here. In the following two theorems, we let \mathbb{U}_k be the subspace model of 2D histograms consisting of k -rectangles.

Theorem 7. *Let $\zeta > 0$ and $\varepsilon > 0$ be arbitrary. Then there is an $(1 + \varepsilon, \mathbb{U}_k, \mathbb{U}_{c,k})$ -approximate tail projection for 2D histograms where $c = O(1/\zeta^2 \varepsilon)$. Moreover, the algorithm runs in time $\tilde{O}(d^{1+\zeta})$.*

Theorem 8. *Let $\zeta > 0$ and $\varepsilon > 0$ be arbitrary. Then there is an $(1 - \varepsilon, \mathbb{U}_k, \mathbb{U}_{c,k})$ -approximate head projection for 2D histograms where $c = O(1/\zeta^2 \varepsilon)$. Moreover, the algorithm runs in time $\tilde{O}(d^{1+\zeta})$.*

Note that both algorithms offer a running time that is *almost linear*, and the small polynomial gap to a linear running time can be controlled as a trade-off between computational and statistical efficiency (a larger output histogram requires more samples to recover). While we provide rigorous proofs for the approximation algorithms as stated above, we remark that we do not establish an overall recovery result similar to Theorem 6. The reason is that the approximate head projection is competitive with respect to k -histograms, but not with the space $\mathbb{U}_k \oplus \mathbb{U}_k$, i.e., the sum of two k -histogram subspaces. The details are somewhat technical and we give a more detailed discussion in Appendix C.3. However, under a natural structural conjecture about sums of k -histogram subspaces, we obtain a similar result as Theorem 6. Moreover, we experimentally demonstrate that the sample complexity of our algorithms already improves over wavelets for k -histograms of size 32×32 .

Finally, we note that our DP approach also generalizes to γ -dimensional histograms for any constant $\gamma \geq 2$. As the dimension of the histogram structure increases, the gap in sample complexity between our algorithm and the prior wavelet-based approach becomes increasingly wide and scales as $O(k \log d)$ vs $O(k \log^\gamma d)$. For simplicity, we limit our attention to the 2D case described above.

2.4 Related work

Recently, there have been several results on approximate projections in the context of recovering low-dimensional structured vectors. (see [12, 13] for an overview). While these approaches also work with approximate projections, they only apply to less general models such as dictionary sparsity [12] or structured sparsity [13] and do not extend to the low-rank matrix recovery problem we address. Among recovery frameworks for general union-of-subspaces models, the work closest to ours is [4], which also gives a generalization of the IHT algorithm. It is important to note that [4] addresses approximate projections, but requires *additive error* approximation guarantees instead of the weaker *relative error* approximation guarantees required by our framework. Similar to the structured sparsity case in [13], we are not aware of any algorithms for low-rank or histogram projections that offer additive error guarantees faster than an exact projection. Overall, our recovery framework can be seen as a generalization of the approaches in [13] and [4].

Low-rank recovery has received a tremendous amount of attention over the past few years, so we refer the reader to the recent survey [9] for an overview. When referring to prior work on low-rank recovery, it is important to note that the fastest known running time for an exact low-rank SVD (even for rank 1) of a $d_1 \times d_2$ matrix is $O(d_1 d_2 \min(d_1, d_2))$. Several papers provide rigorous proofs for low-rank recovery using exact SVDs and then refer to Lanczos methods such as PROPACK [16] while accounting a time complexity of $O(d_1 d_2 r)$ for a rank- r SVD. While Lanczos methods can be faster than exact SVDs in the presence of singular value gaps, it is important to note that all rigorous results for Lanczos SVDs either have a polynomial dependence on the approximation ratio or singular value gaps [20, 17]. No prior work on low-rank recovery establishes such singular value gaps for the inputs to the SVD subroutines (and such gaps would be necessary for *all* iterates in the recovery algorithm). In contrast, we utilize recent work on gap-independent approximate SVDs [17], which enables us to give rigorous guarantees for the entire recovery algorithm. Our results can be seen as justification for the heuristic use of Lanczos methods in prior work.

The paper [2] contains an analysis of an approximate SVD in combination with an iterative recovery algorithm. However, [2] only uses an approximate tail projection, and as a result the approximation ratio $c_{\mathcal{T}}$ must be very close to 1 in order to achieve a good sample complexity. Overall, this leads to a time complexity that does not provide an asymptotic improvement over using exact SVDs.

Recently, several papers have analyzed a non-convex approach to low-rank matrix recovery via factorized gradient descent [3, 7, 23, 24, 22]. While these algorithms avoid SVDs in the iterations of the gradient method, the overall recovery proofs still require an exact SVD in the initialization step. In order to match the sample complexity of our algorithm or SVP, the factorized gradient methods require multiple SVDs for this initialization [7, 22]. As a result, our algorithm offers a better provable time complexity. We remark that [7, 22] use SVP for their initialization, so combining our faster version of SVP with factorized gradient descent might give the best overall performance.

As mentioned earlier, 1D and 2D histograms have been studied extensively in several areas such as databases [14, 8] and density estimation. They are typically used to summarize “count vectors”, with each coordinate of the vector θ corresponding the number of items with a given value in some data set. Computing linear sketches of such vectors, as well as efficient methods for recovering histogram approximations from those sketches, became key tools for designing space efficient dynamic streaming algorithms [11, 10, 21]. For 1D histograms it is known how to achieve the optimal sketch length bound of $n = O(k \log d)$: it can be obtained by representing k -histograms using a tree of $O(k \log d)$ wavelet coefficients as in [10] and then using the structured sparse recovery algorithm of [1]. However, applying this approach to 2D histograms leads to a sub-optimal bound of $O(k \log^2 d)$.

3 An algorithm for recovery with approximate projections

We now introduce our algorithm for recovery from general subspace models using only approximate projections. The pseudo code is formally stated in Algorithm 1 and can be seen as a generalization of IHT [5]. Similar to IHT, we give a version without step size parameter here in order to simplify the presentation (it is easy to introduce a step size parameter in order to fine-tune constant factors). To clarify the connection with projected gradient descent as stated in Equation (2), we use $\mathcal{H}(b)$ (or $\mathcal{T}(b)$) as a function from \mathbb{R}^d to \mathbb{R}^d here. This function is then understood to be $b \mapsto P_{\mathcal{H}(b)} b$, i.e., the orthogonal projection of b onto the subspace identified by $\mathcal{H}(b)$.

The main difference to “standard” projected gradient descent is that we apply a projection to

Algorithm 1 Approximate Subspace-IHT

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1: function AS-IHT( $y, X, t$ )
2:    $\hat{\theta}^0 \leftarrow 0$ 
3:   for  $i \leftarrow 0, \dots, t$  do
4:      $b^i \leftarrow X^T(y - X\hat{\theta}^i)$ 
5:      $\hat{\theta}^{i+1} \leftarrow \mathcal{T}(\hat{\theta}^i + \mathcal{H}(b^i))$ 
6:   return  $\hat{\theta} \leftarrow \hat{\theta}^{t+1}$ 

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both the gradient step and the new iterate. Intuitively, the head projection ensures two points: (i) The result of the head projection on b^i still contains a constant fraction of the residual $\theta^* - \hat{\theta}^i$ (see Lemma 13 in Appendix A). (ii) The input to the tail approximation is close enough to the constraint set \mathbb{U} so that the tail approximation does not prevent the overall convergence. In a nutshell, the head projection “denoises” the gradient so that we can then safely apply an approximate tail projection (as pointed out in [13], only applying an approximate tail projection fails precisely because of “noisy” updates). Formally, we obtain the following theorem for each iteration of AS-IHT (see Appendix A.1 for the corresponding proof):

Theorem 9. *Let $\hat{\theta}^i$ be the estimate computed by AS-IHT in iteration i and let $r^{i+1} = \theta^* - \hat{\theta}^{i+1}$ be the corresponding residual. Moreover, let \mathbb{U} be an arbitrary subspace model. We also assume:*

- $y = X\theta^* + e$ as in Equation (1) with $\theta^* \in \mathcal{M}(\mathbb{U})$.
- \mathcal{T} is a $(c_{\mathcal{T}}, \mathbb{U}, \mathbb{U}_{\mathcal{T}})$ -approximate tail projection.
- \mathcal{H} is a $(c_{\mathcal{H}}, \mathbb{U} \oplus \mathbb{U}_{\mathcal{T}}, \mathbb{U}_{\mathcal{H}})$ -approximate head projection.
- The matrix X satisfies the $(\mathbb{U} \oplus \mathbb{U}_{\mathcal{T}} \oplus \mathbb{U}_{\mathcal{H}}, \delta)$ -subspace RIP.

Then the residual error of the next iterate, i.e., $r^{i+1} = \theta^* - \hat{\theta}^{i+1}$ satisfies

$$\|r^{i+1}\| \leq \eta \|r^i\| + \rho \|e\| ,$$

$$\begin{aligned} \text{where} \quad \eta &= (1 + c_{\mathcal{T}}) \left(\delta + \sqrt{1 - \eta_0^2} \right) , & \rho &= (1 + c_{\mathcal{T}}) \left(\frac{\eta_0 \rho_0}{\sqrt{1 - \eta_0^2}} + \sqrt{1 + \delta} \right) , \\ \eta_0 &= c_{\mathcal{H}}(1 - \delta) - \delta , & \text{and} \quad \rho_0 &= (1 + c_{\mathcal{H}}) \sqrt{1 + \delta} . \end{aligned}$$

The important conclusion of Theorem 9 is that AS-IHT still achieves linear convergence when the approximation ratios $c_{\mathcal{T}}, c_{\mathcal{H}}$ are sufficiently close to 1 and the RIP-constant δ is sufficiently small. For instance, our approximation algorithms for both low-rank matrices offer such approximation guarantees. We can also achieve a sufficiently small value of δ by using a larger number of linear observations in order to strengthen the RIP guarantee (see Appendix D). Hence the use of approximate projections only affects the theoretical sample complexity bounds by constant factors. Moreover, our experiments show that approximate projections achieve essentially the same empirical sample complexity as exact projections (see Section 5).

Given sufficiently small / large constants $c_{\mathcal{T}}, c_{\mathcal{H}}$, and δ , it is easy to see that the linear convergence implied by Theorem 9 directly gives the recovery guarantee and bound on the number of iterations stated in Theorem 5 (see Appendix A.1). However, in some cases it might not be possible to design approximation algorithms with constants $c_{\mathcal{T}}$ and $c_{\mathcal{H}}$ sufficiently close to 1 (in contrast, increasing the sample complexity by a constant factor in order to improve δ is usually a direct consequence

of the RIP guarantee or similar statistical regularity assumptions). In order to address this issue, we show how to “boost” an approximate head projection so that the new approximation ratio is arbitrarily close to 1. While this also increases the size of the resulting subspace model, this increase usually affects the sample complexity only by constant factors as before. Note that for any fixed $c_{\mathcal{T}}$, setting $c_{\mathcal{H}}$ sufficiently close to 1 and δ sufficiently small leads to a convergence rate $\eta < 1$ (c.f. Theorem 9). Hence head boosting enables a linear convergence result for *any* initial combinations of $c_{\mathcal{T}}$ and $c_{\mathcal{H}}$ while only increasing the sample complexity by a constant factor (see Appendix A.3). Formally, we have the following theorem for head boosting, the proof of which we defer to Appendix A.2.

Theorem 10. *Let \mathcal{H} be a $(c_{\mathcal{H}}, \mathbb{U}, \mathbb{U}_{\mathcal{H}})$ -approximate head projection running in time $O(T)$, and let $\varepsilon > 0$. Then there is a constant $c = c_{\varepsilon, c_{\mathcal{H}}}$ that depends only on ε and $c_{\mathcal{H}}$ such that we can construct a $(1 - \varepsilon, \mathbb{U}, \oplus^c \mathbb{U}_{\mathcal{H}})$ -approximate head projection running in time $O(c(T + T'_1 + T'_2))$ where T'_1 is the time needed to apply a projection onto a subspace in $\oplus^c \mathbb{U}_{\mathcal{H}}$, and T'_2 is the time needed to find an orthogonal projector for the sum of two subspaces in $\oplus^c \mathbb{U}_{\mathcal{H}}$.*

We note that the idea of head boosting has already appeared in the context of structured sparse recovery [13]. However, the proof of Theorem 10 is more involved because the subspace in a general subspace model can have arbitrary angles (for structured sparsity, the subspaces are either parallel or orthogonal in each coordinate).

4 Low-rank matrix recovery

We now instantiate our framework for recovery from a subspace model to the low-rank matrix recovery problem. Since we already have proposed the top-level recovery algorithm in the previous section, we only have to provide the problem-specific head and tail approximation algorithms here. We use the following result from prior work on approximate SVDs.

Fact 11 ([17]). *There is an algorithm APPROXSVD with the following guarantee. Let $A \in \mathbb{R}^{d_1 \times d_2}$ be an arbitrary matrix, let $r \in \mathbb{N}$ be the target rank, and let $\varepsilon > 0$ be the desired accuracy. Then with probability $1 - \psi$, APPROXSVD(A, r, ε) returns an orthonormal set of vectors $z_1, \dots, z_r \in \mathbb{R}^{d_1}$ such that for all $i \in [r]$, we have*

$$|z_i^T A A^T z_i - \sigma_i^2| \leq \varepsilon \sigma_{r+1}^2, \quad (3)$$

where σ_i is the i -th largest singular value of A . Furthermore, let $Z \in \mathbb{R}^{d_1 \times r}$ be the matrix with columns z_i . Then we also have

$$\|A - Z Z^T A\|_F \leq (1 + \varepsilon) \|A - A_r\|_F, \quad (4)$$

where A_r is the best rank- r Frobenius-norm approximation of A . Finally, the algorithm runs in time $O\left(\frac{d_1 d_2 r \log(d_2/\psi)}{\sqrt{\varepsilon}} + \frac{d_1 r^2 \log^2(d_2/\psi)}{\varepsilon} + \frac{r^3 \log^3(d_2/\psi)}{\varepsilon^{3/2}}\right)$.

It is important to note that the above results hold for *any* input matrix and do not require singular value gaps. The guarantee (4) directly gives a tail approximation guarantee for the subspace corresponding to the matrix $Z Z^T A$. Moreover, we can convert the guarantee (3) to a head approximation guarantee (see Theorem 18 in Appendix B for details). Since the approximation ε only enters the running time in the approximate SVD, we can directly combine these approximate

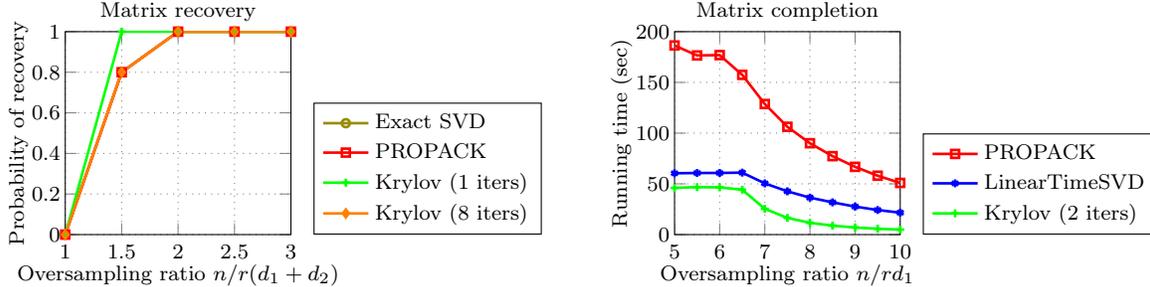


Figure 1: Left: Results for a low-rank matrix recovery experiment using subsampled Fourier measurements. SVP / IHT with one iteration of a block Krylov SVD achieves the same phase transition as SVP with an exact SVD. Right: Results for a low-rank matrix completion problem. SVP / IHT with a block Krylov SVD achieves the best running time and is about 4 – 8 times faster than PROPACK.

projections with Theorem 9, which then yields Theorem 6 (see Appendix B.1 for details).² Empirically, we show in the next section that a very small number of iterations in APPROXSVD already suffices for accurate recovery.

5 Experiments

We now investigate the empirical performance of our proposed algorithms. We refer the reader to Appendix E for more details about the experiments and results for 2D histograms.

Considering our theoretical results on approximate projections for low-rank recovery, one important empirical question is how the use of approximate SVDs such as [17] affects the sample complexity of low-rank matrix recovery. For this, we perform a standard experiment and use several algorithms to recover an image of the MIT logo from subsampled Fourier measurements (c.f. Appendix D). The MIT logo has also been used in prior work [19, 15]; we use an image with dimensions 200×133 and rank 6 (see Appendix E). We limit our attention here to variants of SVP because the algorithm has good empirical performance and has been used as baseline in other works on low-rank recovery. Figure 1 shows that SVP / IHT combined with a single iteration of a block Krylov SVD [17] achieves the same phase transition as SVP with exact SVDs. This indicates that the use of approximate projections for low-rank recovery is not only theoretically sound but can also lead to practical algorithms. In Appendix E we also show corresponding running time results demonstrating that the block Krylov SVD also leads to the fastest recovery algorithm.

We also study the performance of approximate SVDs for the matrix completion problem. We generate a symmetric matrix of size 2048×2048 with rank $r = 50$ and observe a varying number of entries of the matrix. The approximation errors of the various algorithms are again comparable and reported in Appendix E. Figure 1 shows the resulting running times for several sampling ratios. Again, SVP combined with a block Krylov SVD [17] achieves the best running time. Depending on

²We remark that our definitions require head and tail projections to be *deterministic*, while the approximate SVD is *randomized*. However, the running time of APPROXSVD depends only logarithmically on the failure probability, and it is straightforward to apply a union bound over all iterations of AS-IHT. Hence we ignore these details here to simplify the presentation.

the oversampling ratio, the block Krylov approach (now with two iterations) is 4 to 8 times faster than SVP with PROPACK.

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A Proofs for our recovery framework using approximate projections

In this appendix, we provide the convergence proof of our recovery algorithm AS-IHT and related results. Before we begin with the analysis of AS-IHT, we first establish useful consequences of the subspace RIP. For convenience, we recall the definition of the subspace RIP:

Definition 2 (Subspace RIP). *Let $X \in \mathbb{R}^{n \times d}$, let \mathbb{U} be a subspace model, and let $\delta \geq 0$. Then X satisfies the (\mathbb{U}, δ) -subspace RIP if for all $\theta \in \mathcal{M}(\mathbb{U})$ we have $(1 - \delta)\|\theta\|^2 \leq \|X\theta\|^2 \leq (1 + \delta)\|\theta\|^2$.*

Note that the above definition implies that $\|XP_U\| \leq \sqrt{1 + \delta}$ for all $U \in \mathbb{U}$, where $\|XP_U\|$ denotes the spectral norm of XP_U . The following lemma summarizes further consequences of the subspace RIP.

Lemma 12 (Needell, Tropp 2008). *Let $X \in \mathbb{R}^{n \times d}$ be a matrix satisfying the (\mathbb{U}, δ) -subspace RIP. Moreover, let $U \in \mathbb{U}$ be a subspace in the model. Then the following properties hold for all $\theta \in \mathbb{R}^d$ and $y \in \mathbb{R}^n$:*

$$\|P_U X^T y\| \leq \sqrt{1 + \delta} \|y\|, \quad (5)$$

$$\|P_U X^T X P_U \theta\| \leq (1 + \delta) \|\theta\|, \quad (6)$$

$$\|(I - P_U X^T X P_U) \theta\| \leq \delta \|\theta\|. \quad (7)$$

Proof. Equations (5) and (6) follow directly from the bound on the spectral norm of $X P_U$ (which has the same spectral norm as $P_U X^T$).

For Equation (7), consider the eigendecomposition of the symmetric matrix $P_U X^T X P_U$. All eigenvalues are in the interval $[1 - \delta, 1 + \delta]$. Hence forming $I - P_U X^T X P_U$ shifts all eigenvalues into the interval $[-\delta, \delta]$, which implies the spectral norm bound in Equation (7). \square

A.1 Convergence of AS-IHT

We first prove an important lemma for the convergence proof of our algorithm AS-IHT. In a nutshell, the lemma shows that the approximate head projection captures a significant fraction of the residual vector.

Lemma 13. *Let \mathbb{U} , $\mathbb{U}_{\mathcal{T}}$, and $\mathbb{U}_{\mathcal{H}}$ be subspace models, and let $\theta^* \in \mathbb{U}$ and $\hat{\theta}^i \in \mathbb{U}_{\mathcal{T}}$ be vectors such that $y = X\theta^* + e$ as in Equation (1) and $\hat{\theta}^i$ is arbitrary. We also assume that the matrix X satisfies the $(\mathbb{U} \oplus \mathbb{U}_{\mathcal{T}} \oplus \mathbb{U}_{\mathcal{H}}, \delta)$ -subspace RIP. Furthermore, let \mathcal{H} be a $(c_{\mathcal{H}}, \mathbb{U} \oplus \mathbb{U}_{\mathcal{T}}, \mathbb{U}_{\mathcal{H}})$ -approximate head projection. Finally, we define the residual $r^i = \theta^* - \hat{\theta}^i$, the update vector $b^i = X^T(y - X\hat{\theta}^i) = X^T X r^i + X^T e$, and the subspace $U = \mathcal{H}(b^i)$. Then we have*

$$\|P_{U^\perp} r^i\| \leq \sqrt{1 - \eta_0^2} \|r^i\| + \frac{\eta_0 \rho_0}{\sqrt{1 - \eta_0^2}} \|e\|, \quad (8)$$

where

$$\eta_0 = c_{\mathcal{H}}(1 - \delta) - \delta \quad \text{and} \quad \rho_0 = (1 + c_{\mathcal{H}}) \sqrt{1 + \delta}.$$

We assume that $c_{\mathcal{H}}$ and δ are such that $\eta_0 < 1$.

Proof. We first give a lower bound on the part of the residual $\|P_U r\|$ that is “captured” by the approximate head projection. We establish this lower bound via the norm of the update vector $\|P_U b^i\|$. Let $V \in \mathbb{U} \oplus \mathbb{U}_{\mathcal{T}}$ be a subspace such that $r^i \in V$ (note that this is always possible because $\theta \in \mathcal{M}(\mathbb{U})$ and $\hat{\theta}^i \in \mathcal{M}(\mathbb{U}_{\mathcal{T}})$). Using the approximate head projection property of \mathcal{H} , we get

$$\begin{aligned} \|P_U b^i\| &\geq c_{\mathcal{H}} \|P_V b^i\| \\ &= c_{\mathcal{H}} \|P_V X^T X r^i + P_V X^T e\| \\ &\geq c_{\mathcal{H}} \|P_V X^T X P_V r^i\| - c_{\mathcal{H}} \|P_V X^T e\| \end{aligned} \quad (9)$$

$$\geq c_{\mathcal{H}}(1 - \delta) \|r^i\| - c_{\mathcal{H}} \sqrt{1 + \delta} \|e\|. \quad (10)$$

Equation (9) follows from the triangle inequality and the definition of V , which implies $P_V r^i = r^i$. Equation (10) uses Lemma 12 twice.

We now establish an upper bound on $\|P_U b^i\|$:

$$\begin{aligned} \|P_U b^i\| &= \|P_U X^T X r^i + P_U X^T e\| \\ &= \|P_U X^T X r^i - P_U r^i + P_U r^i + P_U X^T e\| \\ &\leq \|P_U(X^T X r^i - r^i)\| + \|P_U r^i\| + \|P_U X^T e\| \\ &\leq \|P_{U+V}(X^T X r^i - r^i)\| + \|P_U r^i\| + \sqrt{1+\delta}\|e\| \end{aligned} \quad (11)$$

$$= \|P_{U+V} X^T X P_{U+V} r^i - r^i\| + \|P_U r^i\| + \sqrt{1+\delta}\|e\| \quad (12)$$

$$\leq \delta\|r^i\| + \|P_U r^i\| + \sqrt{1+\delta}\|e\|. \quad (13)$$

Equation (11) uses Lemma 12 and $U \subseteq U + V$, which implies that projecting onto the subspace $U + V$ instead of U cannot decrease the norm. Equation (12) follows from the definition of V , which implies $r^i \in V$ and hence $P_{U+V} r^i = r^i$. Equation (13) uses Lemma 12 again.

Combining Equations (10) and (13) gives

$$\|P_U r^i\| \geq \eta_0\|r^i\| - \rho_0\|e\|,$$

where η_0 and ρ_0 are as defined in the statement of the lemma. Since we also have $\|P_{U^\perp} r^i\|^2 = \|r^i\|^2 - \|P_U r^i\|^2$, we can now establish the desired upper bound. To simplify notation, we complete our proof with the following claim.

Claim 14. *Let $w, x, y, z \in \mathbb{R}$ be such that $x \geq \eta_0 z - w$ and $y^2 = z^2 - x^2$. Then*

$$y \leq \sqrt{1 - \eta_0^2} z + \frac{\eta_0 w}{\sqrt{1 - \eta_0^2}}.$$

Instantiating Lemma 14 with $w = \rho_0\|e\|$, $x = \|P_U r^i\|$, $y = \|r^i\|$, and $z = \|P_{U^\perp} r^i\|$ then directly implies Equation (8). So it only remains to prove Claim 14, which we accomplish by completing the square:

$$\begin{aligned} y^2 &= z^2 - x^2 \\ &\leq z^2 - (\eta_0 z - w)^2 \\ &= (1 - \eta_0^2)z^2 + 2\eta_0 z w - w^2 \\ &= (1 - \eta_0^2)z^2 + 2\eta_0 z w + \frac{\eta_0^2 w^2}{1 - \eta_0^2} - \frac{\eta_0^2 w^2}{1 - \eta_0^2} - w^2 \\ &= \left(\sqrt{1 - \eta_0^2} z + \frac{\eta_0 w}{\sqrt{1 - \eta_0^2}} \right)^2 - \frac{\eta_0^2 w^2}{1 - \eta_0^2} - w^2. \end{aligned}$$

Since $\eta_0 < 1$, this proves Claim 14. □

Next, we prove that the iterates $\hat{\theta}^i$ of AS-IHT converge linearly.

Theorem 9. *Let $\hat{\theta}^i$ be the estimate computed by AS-IHT in iteration i and let $r^{i+1} = \theta^* - \hat{\theta}^{i+1}$ be the corresponding residual. Moreover, let \mathbb{U} be an arbitrary subspace model. We also assume:*

- $y = X\theta^* + e$ as in Equation (1) with $\theta^* \in \mathcal{M}(\mathbb{U})$.
- \mathcal{T} is a $(c_{\mathcal{T}}, \mathbb{U}, \mathbb{U}_{\mathcal{T}})$ -approximate tail projection.

- \mathcal{H} is a $(c_{\mathcal{H}}, \mathbb{U} \oplus \mathbb{U}_{\mathcal{T}}, \mathbb{U}_{\mathcal{H}})$ -approximate head projection.
- The matrix X satisfies the $(\mathbb{U} \oplus \mathbb{U}_{\mathcal{T}} \oplus \mathbb{U}_{\mathcal{H}}, \delta)$ -subspace RIP.

Then the residual error of the next iterate, i.e., $r^{i+1} = \theta^* - \hat{\theta}^{i+1}$ satisfies

$$\|r^{i+1}\| \leq \eta \|r^i\| + \rho \|e\| ,$$

$$\begin{aligned} \text{where} \quad \eta &= (1 + c_{\mathcal{T}}) \left(\delta + \sqrt{1 - \eta_0^2} \right) , & \rho &= (1 + c_{\mathcal{T}}) \left(\frac{\eta_0 \rho_0}{\sqrt{1 - \eta_0^2}} + \sqrt{1 + \delta} \right) , \\ \eta_0 &= c_{\mathcal{H}}(1 - \delta) - \delta , & \text{and} \quad \rho_0 &= (1 + c_{\mathcal{H}}) \sqrt{1 + \delta} . \end{aligned}$$

Proof. We start by applying the tail projection property of \mathcal{T} on the input vector $a = \hat{\theta}^i + \mathcal{H}(b^i)$:

$$\begin{aligned} \|r^{i+1}\| &= \|\theta^* - \hat{\theta}^{i+1}\| = \|\theta^* - T(a)\| \\ &\leq \|\theta^* - a\| + \|a - T(a)\| \\ &\leq (1 + c_{\mathcal{T}}) \|\theta^* - a\| \\ &= (1 + c_{\mathcal{T}}) \|r^i - \mathcal{H}(X^T X r^i + X^T e)\| . \end{aligned} \tag{14}$$

Intuitively, the quantity on the right hand side of (14) is small for two reasons: first, the matrix $X^T X$ behaves close to an isometry on the vector r^i because r^i is in the subspace model $\mathbb{U} \oplus \mathbb{U}_{\mathcal{T}}$. Second, as we have shown in Lemma 13, the subspace identified by the approximate head projection \mathcal{H} captures a good fraction of the residual r^i , and hence $r^i - \mathcal{H}(b^i)$ is small.

More formally, let the subspaces U and V be defined as before in Lemma 13, i.e., $U = \mathcal{H}(b^i)$ and $V \in \mathbb{U} \oplus \mathbb{U}_{\mathcal{T}}$ is such that $r^i \in V$. Then we get

$$\begin{aligned} \|r^i - \mathcal{H}(X^T X r^i + X^T e)\| &= \|P_U r^i + P_{U^\perp} r^i - P_U X^T X r^i - P_U X^T e\| \\ &\leq \|P_U (X^T X r^i - r^i)\| + \|P_{U^\perp} r^i\| + \|P_U X^T e\| \\ &\leq \|P_{U+V} (X^T X r^i - r^i)\| + \|P_{U^\perp} r^i\| + \|P_U X^T e\| \\ &= \|P_{U+V} X^T X P_{U+V} r^i - r^i\| + \|P_{U^\perp} r^i\| + \|P_U X^T e\| \end{aligned} \tag{15}$$

$$\leq \delta \|r^i\| + \|P_{U^\perp} r^i\| + \sqrt{1 + \delta} \|e\| . \tag{16}$$

Equation (15) uses that $r^i \in V$, and in Equation (16) we invoke consequences of the RIP (see Lemma 12). Combining Equations (14), (16), and Lemma 13 then gives

$$\|r^{i+1}\| \leq (1 + c_{\mathcal{T}}) \left(\delta \|r^i\| + \sqrt{1 - \eta_0^2} \|r^i\| + \frac{\eta_0 \rho_0}{\sqrt{1 - \eta_0^2}} \|e\| + \sqrt{1 + \delta} \|e\| \right) .$$

Rearranging this inequality yields the statement of the theorem. \square

While Theorem 9 only gives a guarantee from one iteration of AS-IHT to the next, it is straightforward to extend this to a guarantee for the entire algorithm.

Corollary 15. *We adopt the setting of Theorem 9, i.e.,*

- $y = X\theta^* + e$ as in Equation (1) with $\theta^* \in \mathcal{M}(\mathbb{U})$.
- \mathcal{T} is a $(c_{\mathcal{T}}, \mathbb{U}, \mathbb{U}_{\mathcal{T}})$ -approximate tail projection.

- \mathcal{H} is a $(c_{\mathcal{H}}, \mathbb{U} \oplus \mathbb{U}_{\mathcal{T}}, \mathbb{U}_{\mathcal{H}})$ -approximate head projection.
- The matrix X satisfies the $(\mathbb{U} \oplus \mathbb{U}_{\mathcal{T}} \oplus \mathbb{U}_{\mathcal{H}}, \delta)$ -subspace RIP.

Furthermore, assume that $c_{\mathcal{T}}$, $c_{\mathcal{H}}$, and δ are such that $\eta < 1$. Set the number of iterations to

$$t = \left\lceil \frac{\log \frac{\|\theta^*\|}{\|e\|}}{\log \frac{1}{\eta}} \right\rceil.$$

Then AS-IHT(y, X, t) returns an estimate $\hat{\theta}$ such that

$$\|\theta^* - \hat{\theta}\| \leq \left(1 + \frac{\rho}{1 - \eta}\right) \|e\|. \quad (17)$$

Proof. Note that $\|r^0\| = \|\theta^*\|$ due to our initialization $\hat{\theta}^0 = 0$. Invoking Theorem 9 and a straightforward induction then yields

$$\|\theta^* - \hat{\theta}\| = \|\theta^* - \hat{\theta}^{t+1}\| = \|r^{t+1}\| = \eta^t \|\theta^*\| + \rho \|e\| \sum_{i=0}^t \eta^i.$$

We can bound the first term on the RHS because we have $\eta^t \|\theta^*\| \leq \|e\|$ for t as defined above. In the second term on the RHS, we bound the geometric series by $\frac{1}{1-\eta}$. Combining these bounds yields Equation (17). \square

Note that Corollary 15 is essentially the formal version of Theorem 5 stated in Section 2. For completeness, we first repeat Theorem 5:

Theorem 5 (informal). *Let \mathcal{H} and \mathcal{T} be approximate head and tail projections with constant approximation ratios, and let the matrix X satisfy the $(\oplus^c \mathbb{U}, \delta)$ -subspace RIP for a sufficiently large constant c and a sufficiently small constant δ . Then there is an algorithm AS-IHT that returns an estimate $\hat{\theta}$ such that $\|\hat{\theta} - \theta^*\| \leq C\|e\|$. The algorithm requires $O(\log\|\theta\|/\|e\|)$ multiplications with X and X^T , and $O(\log\|\theta\|/\|e\|)$ invocations of \mathcal{H} and \mathcal{T} .*

Let c_1 and c_2 be fixed constants. When \mathcal{T} is a $(c_{\mathcal{T}}, \mathbb{U}, \oplus^{c_1} \mathbb{U})$ -approximate tail projection and \mathcal{H} is a $(c_{\mathcal{H}}, \oplus^{c_1} \mathbb{U} \oplus \mathbb{U}, \oplus^{c_2} \mathbb{U})$ -approximate head projection, Theorem 5 is the special case where

- $\mathbb{U}_{\mathcal{T}} = \oplus^{c_1} \mathbb{U}$
- $\mathbb{U}_{\mathcal{H}} = \oplus^{c_2} \mathbb{U}$
- $c = 1 + c_1 + c_2$.

The iteration bound from Corollary 15 implies the bound on the number of multiplications with X and X^T , and the bound on the number of invocations of \mathcal{H} and \mathcal{T} .

A.2 Boosting approximate projections

In some cases, it is hard to design efficient approximate projection algorithms that satisfy the stringent conditions on $c_{\mathcal{T}}$ and $c_{\mathcal{H}}$ in Theorem 9. To overcome this difficulty, we now show how to “boost” the approximation ratio of an approximate head projection to be arbitrarily close to 1.

First, we start with a single iteration of boosting.

Theorem 16. Let $\mathcal{H} : \mathbb{R}^d \rightarrow \mathbb{U}_{\mathcal{H}}$ be a $(c_{\mathcal{H}}, \mathbb{U}, \mathbb{U}_{\mathcal{H}})$ -approximate head projection running in time $O(T)$. Then we can construct a $(2c_{\mathcal{H}} - 2c_{\mathcal{H}}^{3/2} + c_{\mathcal{H}}^2, \mathbb{U}, \mathbb{U}_{\mathcal{H}} \oplus \mathbb{U}_{\mathcal{H}})$ -approximate head projection running in time $O(T + T'_1 + T'_2)$, where T'_1 is the time needed to apply a projection onto a subspace in $\mathbb{U}_{\mathcal{H}}$, and T'_2 is the time needed to find an orthogonal projector for the sum of two subspaces in $\mathbb{U}_{\mathcal{H}}$.

Proof. Consider Algorithm 2. The running time bound follows directly from the definition of BOOSTEDHEAD1. It is also easy to see that the returned subspace is in $\mathbb{U}_{\mathcal{H}} \oplus \mathbb{U}_{\mathcal{H}}$. Hence it remains to show that BOOSTEDHEAD1 satisfies the desired approximation ratio.

In the following, let $OPT = \max_{U' \in \mathbb{U}} \|P_{U'} b\|^2$ be the best possible head approximation, and let $U^* \in \mathbb{U}$ be a subspace achieving OPT . Moreover, let $\tilde{c}_{\mathcal{H}}$ be the head-approximation ratio achieved by the subspace U , i.e.,

$$\|P_U b\|^2 = \tilde{c}_{\mathcal{H}} OPT .$$

Let W be the subspace returned by the algorithm. Then we have

$$\|P_W b\|^2 = \|P_U P_W b\|^2 + \|P_{U^\perp} P_W b\|^2 . \quad (18)$$

We can write $P_U = B^T B$ for an orthogonal basis B of the subspace U , and $P_W = [B^T | D^T] \begin{bmatrix} B \\ D \end{bmatrix}$, where D is an orthonormal basis of the orthogonal complement of U in W (it is easy to see that such a pair of bases always exists, e.g., by following the Gram-Schmidt procedure). Basic linear algebra then shows that $P_U P_W = P_U = P_W P_U$. We can use this fact to bound the first term above:

$$\|P_U P_W b\|^2 = \|P_U b\|^2 = \tilde{c}_{\mathcal{H}} OPT . \quad (19)$$

Next, we consider the second term in Equation (18). We have

$$P_{U^\perp} P_W = (I - P_U) P_W = P_W - P_U P_W = P_W - P_W P_U = P_W (I - P_U) .$$

Since $(I - P_U)b = r$, this gives

$$\|P_{U^\perp} P_W b\|^2 = \|P_W (I - P_U) b\|^2 = \|P_W r\|^2 \geq \|P_V r\|^2 , \quad (20)$$

where the last equality follows from the fact that the subspace W contains the subspace V .

From the head-approximation guarantee of the oracle \mathcal{H} , we know that

$$\|P_V r\|^2 \geq c_{\mathcal{H}} \max_{U' \in \mathbb{U}} \|P_{U'} r\|^2 \geq c_{\mathcal{H}} \|P_{U^*} r\|^2 . \quad (21)$$

Next, we bound $\|P_{U^*} r\|$ (note that we omitted the square).

$$\begin{aligned} \|P_{U^*} r\| &= \|P_{U^*} (I - P_U) b\| = \|P_{U^*} b - P_{U^*} P_U b\| \\ &\geq \|P_{U^*} b\| - \|P_{U^*} P_U b\| \\ &\geq \|P_{U^*} b\| - \|P_U b\| \\ &= \sqrt{OPT} - \sqrt{\tilde{c}_{\mathcal{H}}} \sqrt{OPT} \\ &= \left(1 - \sqrt{\tilde{c}_{\mathcal{H}}}\right) \sqrt{OPT} . \end{aligned}$$

The second line uses the triangle inequality, the third line uses the fact that P_U is an orthogonal projection, and the fourth line uses the optimality of the subspace U^* and the approximation guarantee of the subspace U , respectively. Squaring both sides then yields

$$\|P_{U^*} r\|^2 \geq \left(1 - \sqrt{\tilde{c}_{\mathcal{H}}}\right)^2 OPT . \quad (22)$$

We can now combine Equations (18) to (22) and get

$$\begin{aligned}\|P_W b\|^2 &\geq \tilde{c}_{\mathcal{H}} OPT + c_{\mathcal{H}} \left(1 - \sqrt{\tilde{c}_{\mathcal{H}}}\right)^2 OPT \\ &= (\tilde{c}_{\mathcal{H}} + c_{\mathcal{H}} - 2c_{\mathcal{H}}\sqrt{\tilde{c}_{\mathcal{H}}} + c_{\mathcal{H}}\tilde{c}_{\mathcal{H}}) OPT.\end{aligned}\tag{23}$$

We know that $c_{\mathcal{H}} \leq \tilde{c}_{\mathcal{H}} \leq 1$. In order to get a uniform bound, we analyze the factor in front of OPT . Let $x = \sqrt{\tilde{c}_{\mathcal{H}}}$, then we can write the approximation ratio as

$$f(x) = (1 + c_{\mathcal{H}})x^2 - 2c_{\mathcal{H}}x + c_{\mathcal{H}}.$$

Computing the derivative and setting it to zero yields

$$\begin{aligned}f'(x) &= 2(1 + c_{\mathcal{H}})x - 2c_{\mathcal{H}} \\ x &= \frac{c_{\mathcal{H}}}{1 + c_{\mathcal{H}}}.\end{aligned}$$

So the unconstrained minimum is achieved for a value of $x \leq c_{\mathcal{H}}$, i.e., $\tilde{c}_{\mathcal{H}} \leq c_{\mathcal{H}}^2 \leq c_{\mathcal{H}}$ because $0 \leq c_{\mathcal{H}} \leq 1$. Since the quadratic function f is increasing for $x \geq c_{\mathcal{H}}$, the constrained minimum is achieved for $\tilde{c}_{\mathcal{H}} = c_{\mathcal{H}}$, which gives

$$\|P_W b\|^2 \geq 2c_{\mathcal{H}} - 2c_{\mathcal{H}}^{3/2} + c_{\mathcal{H}}^2. \quad \square$$

Algorithm 2 Boosted head projection

```

1: function BOOSTEDHEAD1( $\mathcal{H}, b$ )
2:    $U \leftarrow \mathcal{H}(b)$ 
3:    $r \leftarrow b - P_U b$ 
4:    $V \leftarrow \mathcal{H}(r)$ 
5:   return an orthogonal projection onto the subspace  $U + V$ 

6: function BOOSTEDHEAD( $\mathcal{H}, b, t$ )
7:   if  $t = 1$  then
8:     return  $\mathcal{H}(b)$ 
9:   else
10:    return BOOSTEDHEAD1(BOOSTEDHEAD( $\mathcal{H}, \cdot, t - 1$ ),  $b$ )

```

Next, we extend one iteration of boosting to several rounds. In our final applications of head approximation boosting, we are mainly interested in boosting a constant head approximation ratio $c_{\mathcal{H}}$ to an improved head approximation ratio $c'_{\mathcal{H}}$ that is close to one but still a constant. Hence it suffices to state a boosting result without explicit dependence between $c_{\mathcal{H}}$ and $c'_{\mathcal{H}}$, which simplifies the argument in the following theorem.

Theorem 10. *Let \mathcal{H} be a $(c_{\mathcal{H}}, \mathbb{U}, \mathbb{U}_{\mathcal{H}})$ -approximate head projection running in time $O(T)$, and let $\varepsilon > 0$. Then there is a constant $c = c_{\varepsilon, c_{\mathcal{H}}}$ that depends only on ε and $c_{\mathcal{H}}$ such that we can construct a $(1 - \varepsilon, \mathbb{U}, \oplus^c \mathbb{U}_{\mathcal{H}})$ -approximate head projection running in time $O(c(T + T'_1 + T'_2))$ where T'_1 is the time needed to apply a projection onto a subspace in $\oplus^c \mathbb{U}_{\mathcal{H}}$, and T'_2 is the time needed to find an orthogonal projector for the sum of two subspaces in $\oplus^c \mathbb{U}_{\mathcal{H}}$.*

Proof. Consider the algorithm BOOSTEDHEAD. If BOOSTEDHEAD(\mathcal{H}, b, t) achieves head approximation ratio $c_{\mathcal{H},t}$, then BOOSTEDHEAD($\mathcal{H}, b, t + 1$) achieves head approximation ratio $c_{\mathcal{H},t+1} = 2c_{\mathcal{H},t} - 2c_{\mathcal{H},t}^{3/2} + c_{\mathcal{H},t}^2$ (see Theorem 16). Hence it suffices to show that the sequence $c_{\mathcal{H},t}$ converges to 1 for any starting value $c_{\mathcal{H},0} = c_{\mathcal{H}}$.

Consider the function $f(x) = 2x - 2x^{3/2} + x^2$ and note that $c_{\mathcal{H},t+1} = f(c_{\mathcal{H},t})$. An elementary calculation shows that $f(x) > x$ for $0 < x < 1$. Hence the sequence $x_{i+1} = f(x_i)$ converges to 1 for any $0 < x_0 < 1$. For a proof by contradiction, let $x' < 1$ be the smallest value such that $x_i \leq x'$ for all i . Let $\delta = f(x') - x' > 0$. Since f is continuous, we can find a point x_{i^*} close to x' such that $f(x_{i^*}) > f(x') - \delta$ and hence $f(x_{i^*}) > x'$, a contradiction. So for any $\varepsilon > 0$, there is a $c = c_{\varepsilon, c_{\mathcal{H}}}$ such that $c_{\mathcal{H},c} \geq 1 - \varepsilon$. \square

A.3 A boosted recovery algorithm

We now combine our convergence result for AS-IHT with the boosting technique to prove a general result that holds for *any* constant head and tail approximation ratios.

Theorem 17. *We make the following assumptions:*

- $y = X\theta^* + e$ as in Equation (1) with $\theta^* \in \mathcal{M}(\mathbb{U})$.
- \mathcal{T} is a $(c_{\mathcal{T}}, \mathbb{U}, \mathbb{U}_{\mathcal{T}})$ -approximate tail projection.
- \mathcal{H} is a $(c_{\mathcal{H}}, \mathbb{U} \oplus \mathbb{U}_{\mathcal{T}}, \mathbb{U}_{\mathcal{H}})$ -approximate head projection.
- The matrix X satisfies the $(\oplus^c(\mathbb{U} \oplus \mathbb{U}_{\mathcal{T}} \oplus \mathbb{U}_{\mathcal{H}}), \delta)$ -subspace RIP for c sufficiently large and δ sufficiently small, where c and δ depend only on $c_{\mathcal{T}}$ and $c_{\mathcal{H}}$.

Then there is an algorithm BOOSTED-AS-IHT that returns an estimate $\hat{\theta}$ such that

$$\|\theta^* - \hat{\theta}\| \leq C\|e\|,$$

where C depends only on δ , $c_{\mathcal{T}}$, and $c_{\mathcal{H}}$. Moreover, the algorithm requires $O(\log\|\theta\|/\|e\|)$ iterations.

Proof. The algorithm BOOSTED-AS-IHT is AS-IHT (Algorithm 1) with BOOSTEDHEAD (Algorithm 2) in place of the approximate head projection.

In order to invoke Corollary 15, we need to show that

$$\eta = (1 + c_{\mathcal{T}}) \left(\delta + \sqrt{1 - \eta_0^2} \right)$$

is less than 1, where η_0 is given by (c.f. Theorem 9)

$$\eta_0 = c_{\mathcal{H}}(1 - \delta) - \delta.$$

Note that by making δ sufficiently small and $c_{\mathcal{H}}$ sufficiently close to 1, we can achieve η_0 arbitrarily close to 1 and hence η arbitrarily small for any *fixed* tail approximation ratio $c_{\mathcal{T}}$.

While the assumption in the current theorem allows for small δ as long as δ only depends on $c_{\mathcal{H}}$ and $c_{\mathcal{T}}$, we need to handle arbitrarily small, fixed $c_{\mathcal{H}}$. In order to do so, we invoke Theorem 10, which allows us to get a boosted head approximation with approximation ratio $c'_{\mathcal{H}}$ arbitrarily close to 1. The invocation of BOOSTEDHEAD leads to a larger output set $\oplus^c \mathbb{U}_{\mathcal{H}}$. As a result, we require the subspace-RIP over the set $\mathbb{U} \oplus \mathbb{U}_{\mathcal{T}} \oplus \oplus^c \mathbb{U}_{\mathcal{H}}$. The current theorem provides this subspace-RIP by assumption. \square

B Proofs for low-rank matrix recovery

We now formally show how to convert the approximate SVD guarantees of [17] to approximate head and tail projections for low-rank matrices. For convenience, we first repeat the main result of [17].

Fact 11 ([17]). *There is an algorithm APPROXSVD with the following guarantee. Let $A \in \mathbb{R}^{d_1 \times d_2}$ be an arbitrary matrix, let $r \in \mathbb{N}$ be the target rank, and let $\varepsilon > 0$ be the desired accuracy. Then with probability $1 - \psi$, $\text{APPROXSVD}(A, r, \varepsilon)$ returns an orthonormal set of vectors $z_1, \dots, z_r \in \mathbb{R}^{d_1}$ such that for all $i \in [r]$, we have*

$$|z_i^T A A^T z_i - \sigma_i^2| \leq \varepsilon \sigma_{r+1}^2, \quad (3)$$

where σ_i is the i -th largest singular value of A . Furthermore, let $Z \in \mathbb{R}^{d_1 \times r}$ be the matrix with columns z_i . Then we also have

$$\|A - Z Z^T A\|_F \leq (1 + \varepsilon) \|A - A_r\|_F, \quad (4)$$

where A_r is the best rank- r Frobenius-norm approximation of A . Finally, the algorithm runs in time $O\left(\frac{d_1 d_2 r \log(d_2/\psi)}{\sqrt{\varepsilon}} + \frac{d_1 r^2 \log^2(d_2/\psi)}{\varepsilon} + \frac{r^3 \log^3(d_2/\psi)}{\varepsilon^{3/2}}\right)$.

As mentioned before, Equation (4) directly gives a tail approximation. We now show how to convert Equation (3) to a head approximation guarantee. In the following, we let \mathbb{U}_r be the subspace model of rank- r matrices.

Theorem 18. *There is an algorithm APPROXLOWRANK with the following property. For an arbitrary input matrix $A \in \mathbb{R}^{d_1 \times d_2}$ and a target rank r , APPROXLOWRANK produces a subspace of rank- r matrices U and a matrix $Y = P_U A$, the projection of A onto U . With probability $99/100$, the output satisfies both an $(1 - \varepsilon, \mathbb{U}_r, \mathbb{U}_r)$ -approximate head projection guarantee and an $(1 + \varepsilon, \mathbb{U}_r, \mathbb{U}_r)$ -approximate tail projection guarantee. Moreover, APPROXLOWRANK runs in time*

$$O\left(\frac{d_1 d_2 r \log d_2}{\sqrt{\varepsilon}} + \frac{d_1 r^2 \log^2 d_2}{\varepsilon} + \frac{r^3 \log^3 d_2}{\varepsilon^{3/2}}\right).$$

Proof. Let z_1, \dots, z_r be the vectors returned by $\text{APPROXLOWRANK}(A, r, \varepsilon)$. Then APPROXLOWRANK returns the matrix $Y = Z Z^T A$ and the subspace U spanned by the vectors z_i and $z_i^T A$ (it is easy to see that Y is indeed the projection of A onto U). Both operations can be performed in time $O(d_1 d_2 r)$. Hence the overall running time is dominated by the invocation of APPROXSVD, which leads to the running time stated in the theorem.

It remains to prove the desired head and tail approximation ratios. The tail approximation guarantee follows directly from Equation (4). For the head approximation, first note that Equation (3) implies

$$z_i^T A A^T z_i \geq (1 - \varepsilon) \sigma_i^2.$$

We now apply this inequality by rewriting the head quantity $\|ZZ^T A\|_F^2$ as follows:

$$\begin{aligned}
\|ZZ^T A\|_F^2 &= \text{tr}(A^T ZZ^T ZZ^T A) \\
&= \text{tr}(A^T ZZ^T A) \\
&= \text{tr}\left(A^T \left(\sum_{i=1}^r z_i z_i^T\right) A\right) \\
&= \text{tr}\left(\sum_{i=1}^r A^T z_i z_i^T A\right) \\
&= \sum_{i=1}^r \text{tr}(A^T z_i z_i^T A) \\
&= \sum_{i=1}^r \text{tr}(z_i^T A A^T z_i) \\
&= \sum_{i=1}^r z_i^T A A^T z_i \\
&\geq (1 - \varepsilon) \sum_{i=1}^r \sigma_i^2 \\
&= (1 - \varepsilon) \|A_r\|_F^2
\end{aligned}$$

where the matrix A_r is the best rank- r approximation of the matrix A . This proves the desired head approximation guarantee. \square

B.1 The final recovery algorithm

We now prove our overall result for low-rank matrix recovery.

Theorem 6. *Let $X \in \mathbb{R}^{n \times d}$ be a matrix with subspace-RIP for low-rank matrices, and let T_X denote the time to multiply a d -dimensional vector with X or X^T . Then there is an algorithm that recovers an estimate $\hat{\theta}$ such that $\|\hat{\theta} - \theta^*\| \leq C\|e\|$. Moreover, the algorithm runs in time $\tilde{O}(T_X + r \cdot d_1^2)$.*

Proof. We assume that X satisfies the low-rank RIP for matrices of rank $4r$ and RIP constant $\delta \leq 0.1$. We remark that it is possible to fine-tune these constants, but our focus here is on the scaling with the problem dimensions.

Instantiating Theorem 18 gives us approximate head and tail projections with the following guarantees:

- \mathcal{T} is a $(1.1, \mathbb{U}_r, \mathbb{U}_r)$ -approximate tail projection.
- \mathcal{H} is a $(0.9, \mathbb{U}_{2r}, \mathbb{U}_{2r})$ -approximate tail projection.

Note that $\mathbb{U}_r \oplus \mathbb{U}_r \subseteq \mathbb{U}_{2r}$, so \mathcal{T} and \mathcal{H} satisfy the conditions of Theorem 9. Moreover, $\mathbb{U} \oplus \mathbb{U}_{\mathcal{T}} \oplus \mathbb{U}_{\mathcal{H}} \subseteq \mathbb{U}_{4r}$, and therefore the matrix X also satisfies the RIP condition of Theorem 9. Substituting $c_{\mathcal{T}} = 1.1$, $c_{\mathcal{H}} = 0.9$, and $\delta = 0.1$ into Theorem 9 then yields $\eta < 0.9$, so we can invoke Corollary 15.

Corollary 15 direct implies the desired recovery guarantee $\|\theta^* - \hat{\theta}\| \leq C\|e\|$. Moreover, the corresponding bound on the number of iterations is $O(\log\|\theta^*\|/\|e\|)$. This has two consequences: (i) The total number of multiplications with X or X^T is $\tilde{O}(1)$. (ii) The total number of invocations

of the approximate head and tail projections is $\tilde{O}(1)$. Recall that each matrix multiplication with X takes T_X time, and that the time complexity of the approximate projections is $\tilde{O}(r \cdot d_1^2)$, where we again assume the square case for simplicity. Combining these results gives the stated time complexity. \square

We remark that for fast design matrices (e.g., structured observations such as a subsampled Fourier matrix), we have $T_X = \tilde{O}(d_1^2)$ and the total running time becomes $\tilde{O}(r \cdot d_1^2)$. See Appendix D for such a construction.

C Approximation algorithms for 2D histograms

We now describe our approximate head and tail projections for histograms. One key ingredient in our algorithms are *hierarchical* histograms. Overall, our goal is to approximate arbitrary 2D histograms, i.e., arbitrary partitions of a $\sqrt{d} \times \sqrt{d}$ matrix with k non-overlapping rectangles (for simplicity, we limit our attention to the case of square matrices). Such histograms are also known as *tiling* histograms. However, tiling histograms are hard to work with algorithmically because they do not allow a clean decomposition for a dynamic program. Instead, work in histogram approximation has utilized hierarchical histograms, which are also partitions of a matrix into k non-overlapping rectangles. The additional restriction is that the partition can be represented as a tree in which each rectangle arises through a vertical or horizontal split of the parent rectangle. We refer the reader to [18] for a more detailed description of different histogram types.

An important result is that every tiling histogram consisting of k rectangles can be simulated with a hierarchical histogram consisting of at most $4k$ rectangles (d’Amore and Franciosa, 1992). Since Theorems 7 and 8 provide bicriterion guarantees for the output space, i.e., projections into a space of histograms consisting of $O(k)$ rectangles, we focus our attention on approximation algorithms for hierarchical histograms in the following. These results can then easily be converted into statements for tiling histograms by increasing the number of histogram tiles by 4.

Next, we introduce some histogram-specific notation. For a histogram subspace U , we denote the number of histogram pieces in U with $\gamma(U)$. We denote the set of hierarchical histogram subspaces with \mathcal{H}_h . When we have an upper bound on the number of histogram pieces, we write $\mathcal{H}_{h,k}$ for the set of hierarchical histogram subspaces U with $\gamma(U) \leq k$.

An important subroutine in our approximate projections is the following notion of a hierarchical histogram oracle.

Definition 19. *An (α, ζ) -hierarchical histogram oracle is an algorithm with the following guarantee: given any $b \in \mathbb{R}^{\sqrt{d} \times \sqrt{d}}$ and $\lambda \in \mathbb{R}$ as input, the algorithm returns a hierarchical histogram subspace U such that*

$$\|P_U b\|^2 - \frac{\lambda}{\alpha} \gamma(U) \geq \max_{U' \in \mathcal{H}_h} \|P_{U'} b\|^2 - \lambda \gamma(U'). \quad (24)$$

Moreover, the algorithm runs in time $O(d^{1+\zeta})$.

An algorithm with the following guarantee directly follows from the hierarchical dynamic programming techniques introduced in [18]. In particular, Theorem 3 of [18] implies a dependence of $\alpha = O(1/\zeta^2)$.

Equation (24) has the flavor of a head approximation (a max-quantified guarantee). As a direct consequence of Equation (24), we also get the following “tail approximation” variant.

Algorithm 3 Tail projection for hierarchical histograms

```

1: function HISTOGRAMTAIL( $b, k, \nu, \xi$ )
2:    $\Delta \leftarrow \min \{ |b_{i,j} - b_{i',j'}| \mid b_{i,j} - b_{i',j'} \neq 0 \}$ 
3:    $\varepsilon_{\min} \leftarrow \frac{\Delta^2}{d^2}$ 
4:    $\lambda_0 \leftarrow \frac{\varepsilon_{\min}}{2k}$ 
5:    $U_0 \leftarrow \text{HISTOGRAMORACLE}(b, \lambda_0)$ 
6:   if  $\|b - P_{U_0}\| = 0$  and  $\gamma(U_0) \leq \alpha k$  then
7:     return  $U_0$ 
8:    $\lambda_l \leftarrow 0$ 
9:    $\lambda_r \leftarrow 2\alpha \|b\|$ 
10:   $\varepsilon \leftarrow \frac{\varepsilon_{\min} \xi}{k}$ 
11:  while  $\lambda_r - \lambda_l \geq \varepsilon$  do
12:     $\lambda_m \leftarrow \frac{\lambda_l + \lambda_r}{2}$ 
13:     $U_m \leftarrow \text{HISTOGRAMORACLE}(b, \lambda_m)$ 
14:    if  $\gamma(U_m) \geq \alpha k$  and  $\gamma(U_m) \leq \nu \alpha k$  then
15:      return  $U_m$ 
16:    if  $\gamma(U_m) \geq \nu \alpha k$  then
17:       $\lambda_l \leftarrow \lambda_m$ 
18:    else
19:       $\lambda_r \leftarrow \lambda_m$ 
20:  return  $\text{HISTOGRAMORACLE}(b, \lambda_r)$ 

```

Lemma 20. *The solution U returned by an (α, ζ) -hierarchical histogram oracle also satisfies*

$$\|b - P_U b\|^2 + \frac{\lambda}{\alpha} \gamma(U) \leq \min_{U' \in \mathcal{H}_h} \|b - P_{U'} b\|^2 + \lambda \gamma(U'). \quad (25)$$

Proof. Multiplying both sides of Equation (24) with -1 and pulling the negative sign into the max gives

$$-\|P_U b\|^2 + \frac{\lambda}{\alpha} \gamma(H) \leq \min_{U' \in \mathcal{H}_h} -\|P_{U'} b\|^2 + \lambda \gamma(U').$$

Adding $\|b\|^2$ to both sides and using that P_U and $P_{U'}$ are orthogonal projections then gives Equation (25) via the Pythagorean Theorem. \square

However, note that neither Equation (24) nor (25) give direct control over the number of histogram pieces k . In the following, we give algorithms that convert these guarantees into approximate projections. In a nutshell, we show that carefully choosing the trade-off parameter λ , combined with a postprocessing step of the corresponding solution, yields head and tail approximations.

C.1 Approximate tail projection

We now show how to construct an approximate tail projection from a hierarchical histogram oracle. In the following, we assume that $\text{HISTOGRAMORACLE}(b, \lambda)$ is an (α, ζ) -hierarchical histogram oracle.

First, we establish a lower bound on the approximation error $\|b - P_U\|^2$ if b is not in the histogram subspace U .

Lemma 21. Let $b \in \mathbb{R}^d$ and U be a histogram subspace. If $b \notin U$, then we have $\|b - P_U\|^2 \geq \varepsilon_{\min}$ where ε_{\min} is as defined in Algorithm 3.

Proof. If $b \notin U$, there is a histogram piece in U on which b is not constant. Let R be the set of indices in this piece. We now give a lower bound on the projection error based on the histogram piece R (recall that the projection of b onto U averages b in each histogram piece):

$$\|b - P_U\|^2 \geq \sum_{(i,j) \in R} (b_{i,j} - \bar{b}_R)^2 \quad \text{where} \quad \bar{b}_R = \frac{1}{|R|} \sum_{(i,j) \in R} b_{i,j}.$$

Let (i^*, j^*) be the index of the largest coefficient in the histogram piece R (ties broken arbitrarily). Then we bound the sum on the right hand side above with the term corresponding to (i^*, j^*) :

$$\|b - P_U\|^2 \geq \left(b_{i^*, j^*} - \frac{1}{|R|} \sum_{(i,j) \in R} b_{i,j} \right)^2.$$

Let Δ_R be the smallest non-zero difference between coefficients in R . Note that $\Delta_R > 0$ because b is not constant on R . Moreover, we have $\Delta_R \leq \max_{(i,j) \in R} b_{i^*, j^*} - b_{i,j}$. Hence we get

$$\left(b_{i^*, j^*} - \frac{1}{|R|} \sum_{(i,j) \in R} b_{i,j} \right)^2 \geq \left(b_{i^*, j^*} - \frac{|R| - 1}{|R|} b_{i^*, j^*} - \frac{1}{|R|} (b_{i^*, j^*} - \Delta_R) \right)^2$$

because b_{i^*, j^*} is one of the largest coefficients in R and at least one coefficient is smaller than b_{i^*, j^*} by at least Δ_R . Combining the inequalities above and simplifying then yields

$$\|b - P_U\|^2 \geq \frac{\Delta_R^2}{|R|^2} \geq \frac{\Delta^2}{d^2} = \varepsilon_{\min}. \quad \square$$

Next, we prove that the histogram oracle returns roughly a k -histogram if the input is a k -histogram and we set the parameter λ correctly.

Lemma 22. Let ε_{\min} and λ_0 be defined as in Algorithm 3. If b is a hierarchical k -histogram, then $\text{HISTOGRAMORACLE}(b, \lambda_0)$ returns a hierarchical histogram subspace U_0 such that $b \in U_0$ and $\gamma(U_0) \leq \alpha k$.

Proof. First, we show that $b \in U_0$, i.e., that $\|b - P_{U_0}\| = 0$. Since $b \in \mathcal{H}_{h,k}$, we know that there is a hierarchical histogram subspace U' such that $\|b - P_{U'}\| = 0$ and $\gamma(U') \leq k$. Substituting this histogram subspace U' and λ_0 into Equation (25) gives

$$\|b - P_{U_0}\|^2 \leq \lambda_0 \gamma(U') \leq \frac{\varepsilon_{\min}^2}{2}$$

where we also used that $\frac{\lambda_0}{\alpha} \gamma(U_0) \geq 0$. Since $\varepsilon_{\min} > 0$, the contrapositive of Lemma 21 shows that $b \in U_0$.

Next, we prove that $\gamma(U_0) \leq \alpha k$. Substituting into Equation (25) again and using $\|b - P_{U_0}\| = 0$ now gives the desired bound on the number of histogram pieces:

$$\frac{\lambda_0}{\alpha} \gamma(U_0) \leq \lambda_0 k. \quad \square$$

With these preliminaries in place, we now show the main result for our tail approximation algorithm.

Theorem 23. *Let $b \in \mathbb{R}^d$, $k \in \mathbb{N}$, $\nu > 1$, and $\xi > 0$. Then $\text{HISTOGRAMTAIL}(b, k, \nu, \xi)$ returns a histogram subspace U such that $\gamma(U) \leq \nu\alpha k$ and*

$$\|b - P_U b\|^2 \leq \left(1 + \frac{1}{\nu - 1} + \xi\right) \min_{U' \in \mathcal{H}_{h,k}} \|b - P_{U'} b\|^2.$$

Moreover, the algorithm runs in time

$$O\left(n^{1+\zeta} \log\left(\frac{\alpha d \|b\|}{\xi \Delta}\right)\right)$$

where Δ is as defined in Algorithm 3.

Proof. We analyze the three cases in which HISTOGRAMTAIL returns separately. First, consider Line 7. In this case, U_0 clearly satisfies the conditions of the theorem. So in the following, we condition on the algorithm not returning in Line 7. By the contrapositive of Lemma 22, this implies that $b \notin \mathcal{M}(\mathcal{H}_{h,k})$.

Next, consider the case that HISTOGRAMTAIL returns in Line 15. This directly implies that $\gamma(U_m) \leq \nu\alpha k$. Moreover, substituting into Equation 25 and restricting the right hand side to histogram subspaces with at most k pieces gives

$$\begin{aligned} \|b - P_{U_m}\|^2 + \frac{\lambda_m}{\alpha} \gamma(U_m) &\leq \min_{U' \in \mathcal{H}_{h,k}} \|b - P_{U'} b\|^2 + \lambda_m \gamma(U') \\ \|b - P_{U_m}\|^2 &\leq \min_{U' \in \mathcal{H}_{h,k}} \|b - P_{U'} b\|^2 + \lambda_m \gamma(U') - \lambda_m k \\ \|b - P_{U_m}\|^2 &\leq \min_{U' \in \mathcal{H}_{h,k}} \|b - P_{U'} b\|^2 \end{aligned}$$

where we used that $\gamma(U_m) \geq \alpha k$ and $\gamma(U') \leq k$.

For the remaining case (Line 20), we use the following shorthands in order to simplify notation: Let U_l and U_r be the histogram subspaces returned by HISTOGRAMORACLE with parameters λ_l and λ_r , respectively. We denote the corresponding tail errors with $t_l = \|b - P_{U_l}\|^2$ and $t_r = \|b - P_{U_r}\|^2$. Moreover, we denote the optimal tail error with $t^* = \min_{U' \in \mathcal{H}_{h,k}} \|b - P_{U'} b\|^2$. Finally, let $\gamma_l = \gamma(U_l)$ and $\gamma_r = \gamma(U_r)$ be the number of histogram pieces in the respective histogram subspaces. Rewriting Equation 25 in terms of the new notation gives

$$t_l + \frac{\lambda_l}{\alpha} \gamma_l \leq t^* + \lambda_l k \tag{26}$$

$$t_r + \frac{\lambda_r}{\alpha} \gamma_r \leq t^* + \lambda_r k \tag{27}$$

We will use Equation 26 in order to bound our tail projection error t_l . For this, we establish an upper bound on λ_r . Note that $\lambda_r \leq \lambda_l + \varepsilon$ when the algorithm reaches Line 20. Moreover, the binary search over λ is initialized so that we always have $\gamma_l > \nu\alpha k$ and $\gamma_r < \alpha k$. Combining these facts

with Equation (27) leads to an upper bound on λ_l :

$$\begin{aligned} t_l + \frac{\lambda_l}{\alpha} \gamma_l &\leq t^* + \lambda_l k \\ \frac{\lambda_l}{\alpha} \nu \alpha k &\leq t^* + \lambda_l k \\ \lambda_l &\leq \frac{t^*}{(\nu - 1)k}. \end{aligned}$$

We use these facts in order to establish an upper bound on t_r . Substituting into Equation (27) gives

$$\begin{aligned} t_r &\leq t^* + (\lambda_l + \varepsilon)k \\ t_r &\leq t^* + \frac{t^*}{\nu - 1} + \frac{\varepsilon_{\min} \xi}{k} k \\ t_r &\leq \left(1 + \frac{1}{\nu - 1} + \xi\right) t^* \end{aligned}$$

where we used that $t^* \geq \varepsilon_{\min}$ because b is not a hierarchical k -histogram if the algorithm reaches Line 20 (see Lemma 21). Combined with the fact that $\gamma_r \leq \alpha k$, this proves the statement of the theorem.

Finally, we consider the running time bound. It is straightforward to see that the overall running time is dominated by the invocations of HISTOGRAMORACLE, each of which takes $O(d^\zeta)$ time. The number of iterations of the binary search is bounded by the initial gap between λ_l and λ_r and the final gap ε , which gives an iteration bound of

$$\left\lceil \log \frac{\lambda_r^{(0)} - \lambda_l^{(0)}}{\varepsilon} \right\rceil = O\left(\log\left(\frac{\alpha d^2 k \|b\|}{\xi \Delta^2}\right)\right).$$

Simplifying and multiplying this iteration bound with the running time of HISTOGRAMORACLE leads to the running time bound stated in the theorem. \square

Theorem 7 now follows directly from Theorem 23. We first restate Theorem 7:

Theorem 7. *Let $\zeta > 0$ and $\varepsilon > 0$ be arbitrary. Then there is an $(1 + \varepsilon, \mathbb{U}_k, \mathbb{U}_{c,k})$ -approximate tail projection for 2D histograms where $c = O(1/\zeta^2 \varepsilon)$. Moreover, the algorithm runs in time $\tilde{O}(d^{1+\zeta})$.*

Setting $\xi = O(\varepsilon)$ and $\nu = O(1/\varepsilon)$ gives the $1 + \varepsilon$ guarantee in Theorem 7. Moreover, we use the $\alpha = O(1/\zeta^2)$ dependence from Theorem 3 of [18].

C.2 Approximate head projection

Next, we show how to construct an approximate head projection from a hierarchical histogram oracle. Similar to the approximate tail projection above, we perform a binary search over the parameter λ in order to achieve a good trade-off between sparsity and approximation. In contrast to the tail case, we now need an additional subroutine for extracting a ‘‘high-density’’ sub-histogram of a given hierarchical histogram. We reduce this task of extracting a sub-histogram to a problem on trees. Formally, we build on the following lemma about the subroutine FINDSUBTREE.

Algorithm 4 Head projection for hierarchical histograms

```

1: function HISTOGRAMHEAD( $b, k, \tau$ )
2:    $b_{\max} \leftarrow \max_{b_{i,j}} |b_{i,j}^2|$ 
3:    $\lambda_l \leftarrow \frac{b_{\max}\tau}{k}$ 
4:    $U_l \leftarrow \text{HISTOGRAMORACLE}(b, \lambda_l)$ 
5:   if  $\gamma(U_l) \leq \frac{2\alpha}{\tau}k$  then
6:     return  $U_l$ 
7:    $\lambda_r \leftarrow 2\alpha\|b\|^2$ 
8:    $\varepsilon \leftarrow \frac{b_{\max}\tau}{2k}$ 
9:   while  $\lambda_r - \lambda_l > \varepsilon$  do
10:     $\lambda_m \leftarrow \frac{\lambda_l + \lambda_r}{2}$ 
11:     $U_m \leftarrow \text{HISTOGRAMORACLE}(b, \lambda_m)$ 
12:    if  $\gamma(U_m) > \frac{2\alpha}{\tau}k$  then
13:       $\lambda_l \leftarrow \lambda_m$ 
14:    else
15:       $\lambda_r \leftarrow \lambda_m$ 
16:     $U_l \leftarrow \text{HISTOGRAMORACLE}(b, \lambda_l)$ 
17:     $U_r \leftarrow \text{HISTOGRAMORACLE}(b, \lambda_r)$ 
18:     $U'_l \leftarrow \text{FINDSUBHISTOGRAM}(b, U_l, \frac{2\alpha}{\tau}k)$ 
19:    if  $\|P_{U'_l}b\|^2 \geq \|P_{U_r}b\|^2$  then
20:      return  $U'_l$ 
21:    else
22:      return  $U_r$ 

```

Lemma 24. *Let $T = (V, E)$ be a tree with node weights $w : V \rightarrow \mathbb{R}$. Moreover, let $s \leq |V|$ be the target subtree size. Then $\text{FINDSUBTREE}(T, w, s)$ returns a node subset $V' \subseteq V$ such that V' forms a subtree in T , its size is at most $2s$, and it contains a proportional fraction of the node weights, i.e., $\sum_{i \in V'} w(i) \geq \frac{s}{|V|} \sum_{i \in V} w(i)$.*

Proof. Let w' and i be defined as in FINDSUBTREE . An averaging argument shows that there must be a contiguous subsequence S as defined in FINDSUBTREE with

$$\sum_{j=i}^{i+2s-1} w'(j) \geq \frac{2s}{2|V|-1} \sum_{j=1}^{2|V|-1} w'(j) \geq \frac{s}{|V|} \sum_{j \in V} w(j)$$

where the first inequality holds because S contains $2s$ nodes, and the second inequality holds by the construction of the tour W .

Let V' be the nodes in S . Note that we have defined w' such that every node weight is used only once, and hence we get

$$\sum_{j \in V'} w(j) \geq \sum_{j=i}^{i+2s-1} w'(j) \geq \frac{s}{|V|} \sum_{j \in V} w(j)$$

as desired. Finally, since S is contiguous in the tour W , the nodes V' form a subtree in T of size at most $2s$. \square

Algorithm 5 Subroutines for the head projection

```

1: function FINDSUBHISTOGRAM( $b, U, s$ )
2:   Let  $T_U = (V_U, E_U)$  be a tree corresponding to the histogram subspace  $U$ .
3:   Let  $w : V_U \rightarrow \mathbb{R}$  be the node weight function corresponding to  $U$  and  $b$ .
4:   Let  $T_U^*$  be the tree  $T_U$  with an additional root node  $r$ .
5:   Let  $w^*$  be defined as  $w$  with the root node weight  $w^*(r) = \|P_{R_0} b\|^2$ .
6:    $V' \leftarrow$  FINDSUBTREE( $T_U^*, w^*, s$ )
7:   if  $r \in V'$  then
8:     return the sub-histogram defined by the splits in  $V'$ 
9:   else
10:    Let  $r'$  be the root node in the subtree defined by  $V'$ .
11:    Let  $U''$  be a 4-piece hierarchical histogram such that one of the leaf rectangles is  $R_{r'}$ .
12:    return the composition of  $U''$  and the sub-histogram defined by  $V'$ 

13: function FINDSUBTREE( $T, w, s$ )
14:   Let  $W = (v_1, \dots, v_{2|V|-1})$  be a tour through the nodes of  $T$ .  $\triangleright T = (V, E)$ 
15:   Let  $w'(j) = \begin{cases} w(v_j) & \text{if position } j \text{ is the first appearance of } v_j \text{ in } W \\ 0 & \text{otherwise} \end{cases}$ 
16:   Let  $S = (v_i, \dots, v_{i+2s-1})$  be a contiguous subsequence of  $W$  with  $\sum_{j=i}^{i+2s-1} w'(j) \geq \frac{s}{|V|} \sum_{j=1}^{2|V|-1} w(j)$ 
17:   return the set of nodes in  $S$ .

```

Utilizing Lemma 24, we now show how to extract a “good” sub-histogram from a given hierarchical histogram. More precisely, our goal is to find a sub-histogram U' with a bounded number of histogram pieces that still achieves a comparable “density” $\frac{\|P_{U'} b\|^2}{\gamma(U')} \approx \frac{\|P_U b\|^2}{\gamma(U)}$. In order to precisely state our algorithm and proof, we now formalize the connection between hierarchical histograms and tree graphs.

For a given histogram subspace U , let $T_U = (V_U, E_U)$ be the tree defined as follows: First, every split in the hierarchical histogram corresponds to a node in V_U . For each split, we then add an edge from the split to the split directly above it in the histogram hierarchy. For a histogram subspace with $\gamma(U)$ pieces, this leads to a tree with $\gamma(U) - 1$ nodes. We also associate each node v in the tree with three rectangles. Specifically, let $R(v)$ be the rectangle split at v , and let $R_l(v)$ and $R_r(v)$ be the left and right child rectangles resulting from the split, respectively.

Next, we define the node weight function $w : V_U \rightarrow \mathbb{R}$. The idea is that the weight of a node corresponds to the “projection refinement”, i.e., the gain in preserved energy when projected onto the finer histogram. More formally, for a rectangle R , let $P_R b$ the projection of b onto the rectangle R , i.e.,

$$(P_R b)_{i,j} = \begin{cases} 0 & \text{if } (i, j) \notin R \\ \frac{1}{|R|} \sum_{(u,v) \in R} b_{u,v} & \text{otherwise} \end{cases}.$$

Then we define the weight of a node v as

$$w(v) = \|P_{R_l(v)} b\|^2 + \|P_{R_r(v)} b\|^2 - \|P_{R(v)} b\|^2.$$

Let $R_1, \dots, R_{\gamma(U)}$ be the rectangles in the hierarchical histogram U , and let R_0 be the $\sqrt{d} \times \sqrt{d}$ “root” rectangle. Since the rectangles are non-overlapping, we have

$$\sum_{i=1}^{\gamma(U)} P_{R_i} b = P_U b.$$

Note that the rectangles $R_1, \dots, R_{\gamma(U)}$ are exactly the child rectangles of the leaves in the tree T_U . Moreover, by the construction of the weight function w , we have

$$\|P_{R_0} b\|^2 + \sum_{v \in V_U} w(v) = \|P_U b\|^2$$

because the contributions from intermediate nodes in the tree T_U cancel out.

Lemma 25. *Let $b \in \mathbb{R}^{\sqrt{d} \times \sqrt{d}}$, let U be a hierarchical histogram subspace, and let $s \leq \gamma(U)$ be the target number of histogram pieces. Then $\text{FINDSUBHISTOGRAM}(b, U, s)$ returns a hierarchical histogram subspace U' such that $\gamma(U') \leq 2s + 4$ and $\|P_{U'} b\|^2 \geq \frac{s}{\gamma(U)} \|P_U b\|^2$. Moreover, the algorithm runs in time $O(d)$.*

Proof. Note that by construction, the tree T_U^* defined in FINDSUBHISTOGRAM has k nodes and the node weights w^* satisfy

$$\sum_{v \in V_{T_U^*}} w(v) = \|P_U b\|^2.$$

Lemma 24 then shows that the subtree defined by the set of nodes V' satisfies $|V'| \leq 2s$ and

$$\sum_{v \in V'} w(v) \geq \frac{s}{\gamma(U)} \sum_{v \in V_{T_U^*}} w(v) \geq \frac{s}{\gamma(U)} \|P_U b\|^2.$$

Let $R'_1, \dots, R'_{|V'|}$ be the leaf rectangles of the subtree V' . The above lower bound on the sum of the node weights implies that

$$\sum_{i=1}^{|V'|} \|P_{R'_i} b\|^2 \geq \frac{s}{\gamma(U)} \|P_U b\|^2.$$

because the rectangles R'_i are non-overlapping and the weights of the inner tree nodes in V' cancel as before. Hence any hierarchical histogram containing the rectangles $R'_1, \dots, R'_{|V'|}$ satisfies the desired head projection bound. It remains to show that we can convert the subtree defined by V' into a hierarchical histogram.

If the set V' contains the root node of T_U^* , the subtree V' directly gives a valid sub-histogram of U . On the other hand, if the root node of T_U^* is not in V' , we can construct a simple 4-piece hierarchical histogram U'' that contains the root rectangle $R_{r'}$ of V' as one of its leaf nodes. The histogram subspace U'' is given by four splits corresponding to the boundaries of the root rectangle $R_{r'}$. We can then combine the hierarchical histogram U'' with the subtree V' by adding the splits in V' to the hierarchical histogram in U'' (by construction, all these splits are valid). The resulting hierarchical histogram then has at most $4 + |V'| \leq 4 + 2s$ pieces.

The running time bound is straightforward: all pre-processing can be accomplished in linear time by computing partial sums for the vector b (projections onto a rectangle can then be computed in constant time). The subroutine FINDSUBTREE also runs in linear time because it requires only a single pass over the tree of size $O(\gamma(U))$. \square

We can now state our approximate head projection algorithm.

Theorem 26. *Let $b \in \mathbb{R}^d$, $k \in \mathbb{N}$, and $0 < \tau < 1$. Then $\text{HISTOGRAMHEAD}(b, k, \tau)$ returns a histogram subspace U such that $\gamma_U \leq \frac{4\alpha}{\tau}k + 4$ and*

$$\|P_U b\|^2 \geq (1 - \tau) \max_{U' \in \mathcal{H}_{h,k}} \|P_{U'} b\|^2.$$

Moreover, the algorithm runs in time $O(d^{1+\zeta} \log \frac{\alpha d}{\tau})$.

Proof. First, we introduce a few shortands to simplify notation. Let the histogram subspace U_l be the solution returned by $\text{HISTOGRAMORACLE}(b, \lambda_l)$. We then write $h_l = \|P_{U_l} b\|^2$ for the head approximation of U_l and $\gamma_l = \gamma(U_l)$ for the number of histogram pieces in the histogram subspace U_l . We adopt a similar convention for h_r and γ_r (corresponding to the solution for parameter λ_r). Finally, let h^* be the optimal head approximation achievable with a k -histogram, i.e., $h^* = \max_{U' \in \mathcal{H}_{h,k}} \|P_{U'} b\|^2$.

Rearranging Equation (24), using the new notation, and substituting the optimal k -histogram solution for the max-quantifier gives

$$h_l \geq h^* - \lambda_l \left(k - \frac{\gamma_l}{\alpha} \right). \quad (28)$$

We now consider the case that the algorithm returns in Line 6. We clearly have $\gamma(U_l) \leq \frac{4\alpha}{\tau}k + 4$ when reaching Line 6. Moreover, substituting for λ_l in Equation 28 gives

$$\begin{aligned} h_l &\geq h^* - \frac{b_{\max} \tau}{k} \left(k - \frac{\gamma_l}{\alpha} \right) \\ &\geq (1 - \tau) h^* \end{aligned}$$

where the second line follows from $h^* \geq b_{\max}$. This inequality holds because any histogram with at least 4 pieces can always create a rectangle that isolates the largest element in b (for simplicity, we assume that $k \geq 4$ and $b \neq 0$). Hence U_l satisfies the conditions of the theorem.

Next, we consider the case that the algorithm reaches the binary search. Note that the binary search is initialized and performed such that we have $\lambda_l \leq \lambda_r \leq \lambda_l + \varepsilon$ when it terminates. Moreover, we have $\gamma_r \leq \frac{2\alpha}{\tau}k$ and $\gamma_l > \frac{2\alpha}{\tau}k$. We now distinguish two sub-cases based on the “density” $\frac{h_l}{\gamma_l}$ of the solution U_l corresponding to λ_l . Let $\phi = \frac{\tau(1-\tau/2)}{2\alpha}$ be the density threshold compared to the optimal solution density $\frac{h^*}{k}$.

Sub-case 1: $\frac{h_l}{\gamma_l} \leq \phi \frac{h^*}{k}$. This inequality allows us to establish an upper bound on λ_l . Rearranging Equation (28) gives (note that $k - \frac{\lambda_l}{\alpha}$ is negative):

$$\begin{aligned} \lambda_l &\leq \frac{h_l - h^*}{\gamma_l/\alpha - k} \\ &\leq \frac{\alpha h_l}{\gamma_l - \alpha k}. \end{aligned}$$

We now use $\gamma_l \geq \frac{2\alpha}{\tau}k$:

$$\begin{aligned}
\lambda_l &\leq \frac{\alpha h_l}{\gamma_l - \tau\gamma_l/2} \\
&\leq \frac{h_l}{\gamma_l} \cdot \frac{\alpha}{1 - \tau/2} \\
&\leq \phi \frac{h^*}{k} \frac{\alpha}{1 - \tau/2} \\
&\leq \frac{\tau}{2} \cdot \frac{h^*}{k}.
\end{aligned}$$

where we used the density upper bound for U_l valid in this subcase and the definition of ϕ . Next, we derive a lower bound on h_r . Instantiating Equation (28) with U_r instead of U_l gives

$$\begin{aligned}
h_r &\geq h^* - \lambda_r \left(k - \frac{\gamma_r}{\alpha} \right) \\
&\geq h^* - \lambda_r k \\
&\geq h^* - (\lambda_l + \varepsilon)k \\
&= h^* - \lambda_l k - \varepsilon k \\
&\geq h^* - \frac{\tau}{2} h^* - \frac{\tau}{2} b_{\max} \\
&\geq (1 - \tau)h^*
\end{aligned}$$

where we again used $b_{\max} \leq h^*$. So in this sub-case, U_r satisfies the conditions of the theorem.

Sub-case 2: $\frac{h_l}{\gamma_l} \geq \phi \frac{h^*}{k}$. In this subcase, the solution U_l has a good density, so FINDSUBHISTOGRAM can extract a good solution with a bounded number of histogram pieces. More formally, since $\gamma_l \geq \frac{2\alpha}{\tau}k$, we can invoke Lemma 25 and get

$$\begin{aligned}
\|P_{U'_l} b\|^2 &\geq \frac{\frac{2\alpha}{\tau}k}{\gamma_l} h_l \\
&\geq \frac{2\alpha k}{\tau} \phi \frac{h^*}{k} \\
&\geq \left(1 - \frac{\tau}{2}\right) h^*.
\end{aligned}$$

Moreover, the output of FINDSUBHISTOGRAM satisfies $\gamma(U'_l) \leq \frac{4\alpha}{\tau}k + 4$, and hence U'_l satisfies the conditions of the theorem.

We can now conclude the proof of the theorem: always, one of sub-case 1 and sub-case 2 holds. Since HISTOGRAMHEAD always returns the best of the two choices U_r and U'_l , the overall result has the desired head approximation guarantee.

The overall running time is dominated by the invocations of HISTOGRAMORACLE in the binary search. Each invocation takes $O(n^{1+\zeta})$ time and the number of invocations is the number of iterations of the binary search, i.e., bounded by

$$\left\lceil \log \frac{\lambda_r^{(0)} - \lambda_l^{(0)}}{\varepsilon} \right\rceil \leq \left\lceil \frac{\lambda_r^{(0)}}{\varepsilon} \right\rceil \leq \left\lceil \log \frac{4\alpha k \|b\|^2}{b_{\max} \tau} \right\rceil.$$

Since $k \leq d$ and $\frac{\|b\|^2}{b_{\max}} \leq d$, the running time bound in the theorem follows. \square

As before, Theorem 8 follows as a direct consequence of Theorem 26. For completeness, we repeat the statement of Theorem 8:

Theorem 8. *Let $\zeta > 0$ and $\varepsilon > 0$ be arbitrary. Then there is an $(1 - \varepsilon, \mathbb{U}_k, \mathbb{U}_{c \cdot k})$ -approximate head projection for 2D histograms where $c = O(1/\zeta^2\varepsilon)$. Moreover, the algorithm runs in time $\tilde{O}(d^{1+\zeta})$.*

Setting $\tau = O(\varepsilon)$ gives the $1 - \varepsilon$ guarantee in Theorem 8. Moreover, we use the $\alpha = O(1/\zeta^2)$ dependence from Theorem 3 of [18].

C.3 Recovery of 2D histograms

While we have approximate projections for 2D histograms, they do not suffice to state an overall recovery guarantee in the current form. The issue is that Theorem 9 requires an approximate head projection that is competitive with respect to the sum of subspaces $\mathbb{U} \oplus \mathbb{U}_{\mathcal{T}}$. While this is easy to satisfy for low-rank matrices (the sum of two rank- r subspace models is contained in the rank- $2r$ subspace model), adding histogram subspace models is more subtle. For instance, consider two k -histogram subspaces corresponding to k rows and columns of a $k \times k$ matrix, respectively. The sum of the two subspaces then contains k^2 individual rectangles (a chessboard pattern). While these k^2 rectangles are not independent (the dimension of the space is only $2k$), the chessboard pattern is not directly contained in the set of $2k$ -histogram subspaces. As a result, a head approximation that is competitive with respect to $2k$ -histograms is not immediately competitive with respect to the sum of two k -histograms.

While head boosting is not directly helpful to overcome this issue, we believe that 2D histograms are “well-behaved” in the sense that boosting is still helpful. In particular, we believe that the sum of two k -histograms still allows a constant-factor head approximation with a single $O(k)$ -histogram subspace. More formally, we state the following conjecture.

Conjecture 1. *Let $c > 0$ be fixed. Then there are universal constants $c_1 > 0$ and $c_2 > 0$ depending on c such that the following holds. For any $b \in \mathbb{R}^d$, there is a $c_1 k$ -histogram subspace U such that we have*

$$\|P_U b\| \geq c_2 \|P_{\oplus^c \mathbb{U}_k} b\|.$$

If the above conjecture is true, Theorem 26 yields an approximate head projection that is competitive to $\oplus^c \mathbb{U}_k$. Combining this with the boosted version of our recovery framework (see Appendix A.3) then yields an overall recovery algorithm.

D Sample complexity of subspace recovery

Here, we establish bounds on the sample complexity of subspace recovery for some particular instances. In particular, our focus is on *fast* sampling operators, i.e., operators that support matrix-vector multiplications with a running time that is *nearly-linear* in the size of the vector. Our results follow from a standard concatenation of previously existing results.

D.1 Low-rank matrices

Consider the case where the subspace model \mathbb{U} corresponds to the set of rank- r matrices of size $d_1 \times d_1$. Then, the subspace RIP corresponds to the (somewhat) more well-known *rank- r restricted isometry property*, first introduced in Recht, Fazel, and Parillo. We obtain the following result:

Theorem 27. *Let $d = d_1^2$. Then, there exists a randomized construction of a matrix $X \in \mathbb{R}^{n \times d}$, with parameters $n = O(\text{rd polylog } d)$, such that X satisfies the rank- r RIP with high probability. Moreover, X supports matrix-vector multiplications with complexity $O(d \log d)$.*

Proof. We begin by considering matrices that satisfy the *standard* RIP for s -sparse vectors, as well as support fast matrix-vector multiplication. To the best of our knowledge, the sharpest such bounds have been recently obtained by Haviv and Regev (SODA 2016). They show that with high probability, a matrix formed by randomly subsampling $n = O(\delta^{-2} s \log^2(s/\delta) d)$ rows of the discrete Fourier Transform (DFT) matrix satisfies the standard RIP (with isometry constant δ) over the set of s -sparse vectors.

Next, we invoke a well-known result by Ward and Krahmer ("New and Improved Johnson-Lindenstrauss Embeddings via the RIP"). Consider a diagonal matrix D_ξ , where the diagonal ξ is a Rademacher sequence uniformly distributed over $\{-1, 1\}^d$. Also consider any fixed set of vectors B with $|B| = m$ where $s > O(\log \frac{m}{\eta})$. If X' is any $n \times d$ matrix that satisfies the standard RIP over the set of s -sparse vectors with constant $\delta < \varepsilon/4$, then high probability the matrix $X = X'D_\xi$ is a *Johnson-Lindenstrauss* embedding for E . Formally, the following is true with probability exceeding $1 - \eta$:

$$(1 - \varepsilon)\|\beta\|_2^2 \leq \|X\beta\|_2^2 \leq (1 + \varepsilon)\|\beta\|_2^2.$$

uniformly for all $\beta \in B$.

Next, we invoke Lemma 3.1 of Candes and Plan ("Tight Oracle Bounds for Matrix Recovery"), who show that the set of vectors corresponding to rank- k matrices, S_k , exhibits an ε -net \bar{S}_k (with respect to the Euclidean norm) such that

$$|\bar{S}_k| \leq (9/\varepsilon)^{(d_1+d_2+1)k}.$$

Also from Candes and Plan, we have that if X is a Johnson-Lindenstrauss embedding with isometry constant ε for an \bar{S}_k , then X satisfies the rank- k RIP with constant $\delta = O(\varepsilon)$. Plugging in $s = O(k(d_1 + d_2))$ and $m = O(s \text{ polylog } d)$ and adjusting constants, we get the stated result. \square

D.2 Histograms

Now, consider the case where the subspace model \mathbb{U} corresponds to the set of (hierarchical or tiling) histograms. Since either type of histogram can be modeled as superpositions of sub-rectangles of the domain $\sqrt{d} \times \sqrt{d}$, we can simply model the histogram subspace model \mathbb{U} as a subset of *dictionary-sparse* vectors $\{x | x = D\alpha, \|\alpha\|_0 \leq k\}$. Here, D is a dictionary of size $d \times \binom{d^2}{2}$ where each column of D corresponds to a single tile (normalized to unit ℓ_2 -norm).

Therefore, any matrix that satisfies the RIP with respect to the dictionary D (abbreviated sometimes as the D -RIP) also suffices for reliable histogram subspace recovery. The following result is folklore, and a formal proof can be found in the appendix of Hegde, Indyk, and Schmidt ("Nearly Linear-Time Model-Based Compressive Sensing").

Theorem 28. *There exists a randomized construction of a matrix $X \in \mathbb{R}^{n \times d}$, with parameters $n = O(k \log d/k)$, such that with high probability, X satisfies the subspace RIP for the histogram subspace model. Moreover, X supports matrix-vector multiplications with complexity $O(d \log d + k^2 \text{ polylog } d)$.*

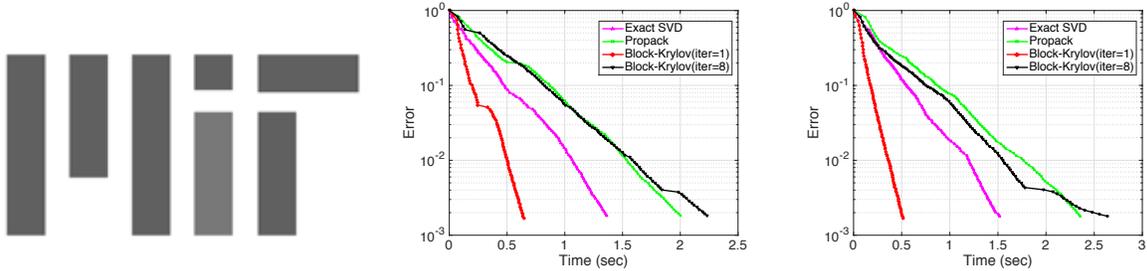


Figure 2: (left) Example low-rank matrix of size $d = 133 \times 200, r = 6$. (right) Recovery error of various algorithms as a function of time (2 independent trials).

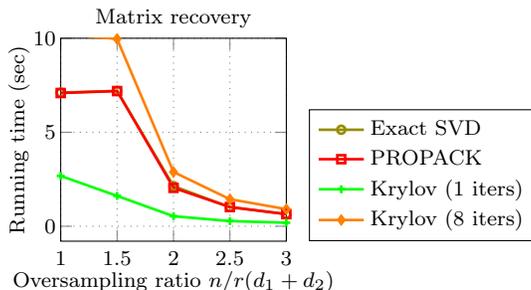


Figure 3: Running times corresponding to the low-rank matrix recovery experiment in Figure 1. The block Krylov variant of IHT with one iteration has the best running time.

E Supplemental experiments

We begin with a description of the experimental setup. All experiments were conducted on an iMac desktop computer with an Intel Core i5 CPU (3.2 GHz) and 16 GB RAM. With the exception of the dynamic program (DP) for 2D histograms, all code was written in Matlab. We chose C++ for the 2D histogram DP because it heavily relies on for-loops, which tend to be slow in Matlab. Since the Krylov SVD of [17] is only available as a Matlab routine, we also chose the Matlab version of PROPACK [16] so that the implementations are comparable. Unless reported otherwise, all reported data points were averaged over at least 10 trials.

E.1 Low-rank matrix recovery experiments

Figure E shows an image of the MIT logo used in the low-rank matrix recovery experiments [19, 15]. For our first experiment, we record $n = 3.5(d_1 + d_2)r = 6994$ linear measurements of the image. The measurement operator is constructed by subsampling m rows of a Fourier matrix and multiplying its columns by a randomly chosen Bernoulli vector, similar to the RIP matrix given in Appendix D. The goal is to recover the image from these observations.

We adapt the Singular Value Projection (SVP) algorithm of [15] by replacing the exact SVD step with approximate SVDs (some of which are very coarse), and demonstrate that we can still achieve efficient matrix recovery from few observations. As alternatives to Matlab's in-built `svd` function, we include the PROPACK [16] numerical linear algebra package, which implements a Lanczos-type method. We also include an implementation of the recent Block-Krylov SVD algorithm of [17], which

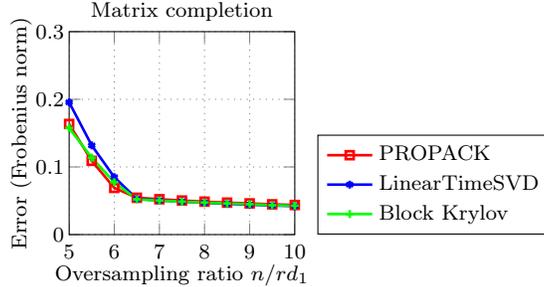


Figure 4: Average approximation errors for the low-rank matrix completion experiment in Figure 1. As for low-rank matrix recovery, the different SVDs achieve essentially the same error.

offers a nice tradeoff between approximation ratio and running time. We test this method with 1 and 8 Krylov subspace iterations (8 is the default provided in the code of [17]).

Figure 3 shows the running times corresponding to the phase transition plot in Figure 1. The only stopping criteria we used were based on a small residual and a maximum number of iterations, so the running times of the algorithms are slowest in the regime where they do not recover the signal.

The subspace IHT algorithm is iterative, i.e., it produces a sequence of matrix estimates $\{\hat{\theta}^0, \hat{\theta}^1, \dots, \hat{\theta}^t\}$. Figure E displays the estimation error, $\frac{\|\theta^* - \hat{\theta}^t\|}{\|\theta^*\|}$, as a function of wall-clock time, on two different trial runs. We observe from the plots that PROPACK and the Block Krylov method (with 8 iterations) perform similar to the exact SVD due to the small problem size. Interestingly, a *very coarse* approximate SVD (a single Block Krylov subspace iteration) provides the fastest convergence. Overall, using approximate SVDs within SVP / IHT does not only yield computational speed-ups, but also offers competitive statistical performance.

We also report results of using the SVP / IHT algorithm with approximate projections on a larger matrix completion problem. We generate a matrix of size $d = 2048 \times 2048$ with rank $r = 50$. We only sample n randomly chosen entries of this matrix and attempt to reconstruct the matrix from these entries using SVP with approximate low-rank projections. We vary n and obtain error curves as well as running times. Figure 4 shows the approximation errors for the matrix completion experiment in Figure 1. As for the matrix recovery experiments, all SVDs achieve essentially the same error. We note that the error floor of about 0.05 is a result of our stopping criterion.

E.2 2D histogram recovery

Finally, we show our results for recovering a 2D histogram from linear observations. As before, we use subsampled Fourier measurements. Our test vector is a 32×32 hierarchical histogram consisting of 4 rectangles. Hierarchical histograms are essentially 2D piecewise constant functions over a 2D domain where the constant pieces (or tiles) are generated by starting with the entire domain as a single tile and recursively partitioning tiles by making horizontal or vertical splits. We compare three approaches: (i) “Standard” sparsity in the Haar wavelet domain. (ii) Tree sparsity in the Haar wavelet domain [10, 1]. (iii) Our approximate projection algorithm. The focus in our experiments is on sample complexity, so we have implemented only one “level” of the DP in [18]. Figure 5 shows the corresponding phase transitions. The 2D histogram DP does indeed offer the best empirical sample complexity.

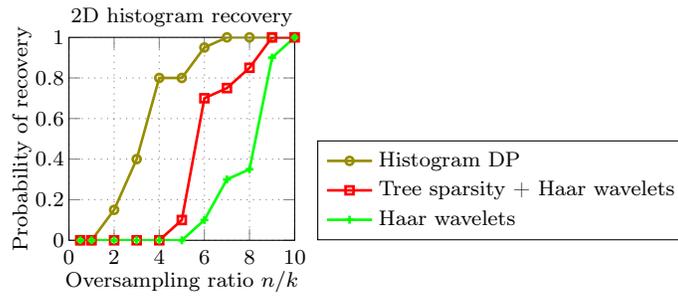


Figure 5: Results for recovering a hierarchical histogram from subsampled Fourier measurements. As predicted by our theoretical argument, the 2D histogram DP has the best sample complexity.