A Fast and Adaptive SVD-free Algorithm for General Weighted Low-rank Recovery

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Abstract. This paper is devoted to proposing a general weighted low-rank recovery model and designs a fast SVD-free computational scheme to solve it. First, our generic weighted low-rank recovery model unifies several existing approaches in the literature. Moreover, our model readily extends to the non-convex setting. Algorithm-wise, most first-order proximal algorithms in the literature for low-rank recoveries require computing singular value decomposition (SVD). As SVD does not scale properly with the dimension of the matrices, these algorithms becomes slower when the problem size becomes larger. By incorporating the variational formulation of the nuclear norm into the sub-problem of proximal gradient descent, we avoid to compute SVD which results in significant speed-up. Moreover, our algorithm preserves the rank identification property of nuclear norm [33] which further allows us to design a rank continuation scheme that asymptotically achieves the minimal iteration complexity. Numerical experiments on both toy example and real-world problems including structure from motion (SfM) and photometric stereo, background estimation and matrix completion, demonstrate the superiority of our proposed algorithm.

Key words. Low-rank recovery, Weighted low-rank, Nuclear norm, Singular value decomposition, proximal gradient descent, alternating minimization, rank identification/continuation

AMS subject classifications. 49J52, 65K05, 65K10, 90C06, 90C30

1 Introduction

Low-rank matrix recovery is an important problem to study as it covers many interesting problems arising from diverse fields including machine learning, data science, signal/image processing, and computer vision, to name a few. The goal of low-rank recovery is to recover or approximate the targeted matrix $\hat{X} \in \mathbb{R}^{m \times n}$ whose rank is much smaller than its dimension. For example, matrix completion [54, 13], structure from motion [44], video segmentation [64, 28, 61], image processing and signal retrieval [26, 59] exploit the inherent low-rank structure of the data.

For many problems of interests, instead of accessing the data $\hat{X}$ directly, often we can only observe it through some agent (e.g. a linear operator) $\Psi$. A general observation model takes the following form

$$F = \Psi(\hat{X}) + \epsilon,$$

where $\Psi : \mathbb{R}^{m \times n} \to \mathbb{R}^{d \times \ell}$ is the (observation) operator which is assumed to be bounded linear. For example, in the compressed sensing scenario, $\Psi$ returns a linear measurement of $\hat{X}$ which is a $d$-dimensional vector [26]; for matrix completion $\Psi$ is a binary mask [13]. In the above model, variable $\epsilon \in \mathbb{R}^{d \times \ell}$ denotes additive noise (e.g. white Gaussian) and $F \in \mathbb{R}^{d \times \ell}$ is the obtained noise contaminated observation.

Over the years, numerous low-rank promoting recovery models are proposed in the literature, for example [39, 63, 59, 62, 61, 15], to mention a few. When the rank of $\hat{X}$ is available, one can consider the following rank
constrained weighted least square
\[
\min_{X \in \mathbb{R}^{m \times n}} \frac{1}{2} \left\| (\Psi(X) - F) \odot W \right\|^2 \quad \text{such that} \quad \text{rank}(X) \leq r,
\]
where \( r = \text{rank}(\hat{X}) \) is the rank of \( \hat{X}, W \in \mathbb{R}^{d \times l} \) is a non-negative weight matrix, and \( \odot \) is the Hadamard product. The motivation of considering a weight \( W \) is such that (2) can handle more general noise model \( \varepsilon \), rather than mere Gaussian noise [16, 22, 23]. A clear limitation of (2) is that, for many problems it is in general impossible to know \( \text{rank}(\hat{X}) \) \textit{a priori}. As a result, instead of using rank as constraint, one can penalize it to the objective which results in rank regularized recovery model
\[
\min_{X \in \mathbb{R}^{m \times n}} \frac{1}{2} \left\| (\Psi(X) - F) \odot W \right\|^2 + \tau \text{rank}(X),
\]
where \( \tau > 0 \) is the regularization parameter. Though avoids the estimation of \( r \), one needs to choose \( \tau \) properly. Moreover, due to \( \text{rank} \) function, (2) and (3) are non-convex, imposing challenges to both theoretical analysis and algorithmic design.

In literature, a popular approach to avoid non-convexity is to replace the \( \text{rank} \) function with its convex surrogate—the nuclear norm (a.k.a. trace norm) [35, 12]. Correspondingly, we obtain the following nuclear norm constrained form of (2)
\[
\min_{X \in \mathbb{R}^{m \times n}} \frac{1}{2} \left\| (\Psi(X) - F) \odot W \right\|^2 \quad \text{such that} \quad \|X\|_* \leq c,
\]
where \( c \) is a predefined constant, e.g. \( c = \|\hat{X}\|_* \) if possible. Consequently, for (3), we arrive at the following \textit{unconstrained} nuclear norm regularized recovery model
\[
\min_{X \in \mathbb{R}^{m \times n}} \frac{1}{2} \left\| (\Psi(X) - F) \odot W \right\|^2 + \tau \|X\|_*.
\]
Note that, besides the weighted \( \ell_2 \) loss, one can also consider the general loss function \( f(X, F, W) \) which gives the following recovery model (similar to [30, 51])
\[
\min_{X \in \mathbb{R}^{m \times n}} f(X, F, W) + \tau \|X\|_*,
\]
For the rest of the paper, we mainly focus on model (5) and only present a short discussion of (6) in Section 2.3.

1.1 Related work
Our recovery model (5) is connected with several established work in the literature, and moreover covers some as special cases. Therefore in what follows, we present a short overview of literature study.

\textbf{Trace LASSO} When the entries of \( W \) are all 1’s, problem (5) becomes the Trace LASSO considered in [25], \textit{i.e.} nuclear norm regularized least square. For this case, \( \varepsilon \) corresponds to additive white Gaussian noise. When \( \Psi \in \mathbb{R}^{d \times m}, F \in \mathbb{R}^{d \times n} \) and \( W = 1 \in \mathbb{R}^{d \times n} \), (5) becomes
\[
\min_{X \in \mathbb{R}^{m \times n}} \frac{1}{2} \|\Psi X - F\|^2 + \tau \|X\|_*,
\]
which is studied in [47]. Examples of (7) include multivariate linear regression, multi-class classification and multi-task learning [47]. As proposed in [29, 30], one can also consider a general loss function \( f(X, F) \) (as a special case of (6)) to solve
\[
\min_{X \in \mathbb{R}^{m \times n}} f(X, F) + \tau \|X\|_*,
\]
For different choice of \( f(X, F) \) in (8), one can recover affine-rank minimization [55, 39], regularized semi-definite linear least squares [51], etc.

\textbf{Weighted low-rank recovery} For the case \( W \) is a general non-negative weight, there is also a train of works in the literature. For most of them, \( \Psi \) is an identity operator. We start with rank constraint case
\[
\min_{X \in \mathbb{R}^{m \times n}} \frac{1}{2} \| (F - X) \odot W \|^2 \quad \text{such that} \quad \text{rank}(X) \leq r,
\]
which is well-studied in the literature under different settings [38, 52, 53, 42, 56]. In [41], instead of considering a generic weight $W$, the authors proposed a general matrix induced weighted norm

$$\min_{X \in \mathbb{R}^{mn \times n}} \|F - X\|_Q^2 \text{ such that } \text{rank}(X) \leq r,$$

where $Q \in \mathbb{R}^{mn \times mn}$ is symmetric positive definite and

$$\|F - X\|_Q^2 \overset{\text{def}}{=} \text{vec}(F - X)^\top Q \text{vec}(F - X)$$

with $\text{vec}(\cdot)$ being an operator which maps the entries of $\mathbb{R}^{m \times n}$ to vectors in $\mathbb{R}^{mn \times 1}$ by stacking the columns. We refer to [16, 41, 22, 23, 48, 20] and the references therein for more discussions. If we lift the constraint to the structure dependent analytically and empirically. We believe problem (5) with our dedicated analysis should be studied as a standalone problem to close the existing knowledge gap.

In this paper, we propose a general model (5) for low-rank recovery. Based on the variational formulation of nuclear norm, we propose an efficient algorithm which avoids computing SVD. More precisely, our contributions include the following aspects.

1.2 Contributions

In this paper, we propose a general model (5) for low-rank recovery. Based on the variational formulation of nuclear norm, we propose an efficient algorithm which avoids computing SVD. More precisely, our contributions include the following aspects.

A generic low-rank recovery model We propose general low-rank recovery models which covers several existing works as special cases. We provide a detailed comparison of our problem with the existing ones, both analytically and empirically. We believe problem (5) with our dedicated structure dependent analysis should be studied as a standalone problem to close the existing knowledge gap.
An efficient SVD-free algorithm  In the literature, numerous numerical schemes can be applied to solve (5), since it is the sum of a smooth function and a non-smooth one. However, most of these algorithms require computing SVD, which does not scale properly with the dimension of the problem [17, 30, 47, 44]. To efficiently solve (5), we propose an SVD-free method (see Algorithm 1). By combining proximal gradient descent [36] and the variational characteristic of nuclear norm, we design a “proximal gradient & alternating minimization method” which we coin as ProGrAMMe. Our algorithm can also be applied to solve the general model (6) if the loss function \( f(X,F,W) \) is smoothly differentiable with gradient being Lipschitz continuous. Moreover, our algorithm can be easily extended to the non-convex loss function case.

A rank continuation strategy Based on the result of [33], we show that the sequence generated by Algorithm 1 can find the rank of the minimizer (to which the generated sequence converges) in finite number of iterations, which we call rank identification property. In turn, we design a rank continuation technique which leads to Algorithm 2. Compare to Algorithm 1, rank continuation is less sensitive to initial parameter, and asymptotically achieves the minimal per iteration complexity.

Numerical comparisons We evaluate our algorithms against 15 state-of-the-art weighted and unweighted low-rank approximation methods on various tasks, including structure from motion (SfM) and photometric stereo, background estimation from fully and partially observed data, and matrix completion. In these problems, different weights are used as deemed fit—from binary weights to random large weights. We observed in all the tasks our weighted low-rank algorithm performs either better or is as good as the other algorithms. This indicates that our algorithm is robust and scalable to both binary and general weights on a diverse set of tasks.

1.3 Notions and definitions
Throughout the paper, \( \mathbb{R}^m \) is a finite dimensional Euclidean space equipped with scalar product \( \langle \cdot, \cdot \rangle \) and induced norm \( \| \cdot \| \). We abuse the notation \( \| \cdot \| \) for the Frobenius norm when \( \cdot \) is a matrix. \( \text{Id}_m \) denotes the identity operator on \( \mathbb{R}^m \). Let \( S \subset \mathbb{R}^n \) be a non-empty close compact set, then \( \text{ri}(S) \) denotes its relative interior, and \( \text{par}(S) \) is the subspace which is parallel to \( \text{span}(S) \). The sub-differential of a proper closed convex function \( g : \mathbb{R}^m \to \mathbb{R} \cup \{ +\infty \} \) is a set-valued mapping defined by \( \partial g : \mathbb{R}^m \ni x \mapsto \{ v \in \mathbb{R}^m | g(x) \geq g(y) + \langle v, x - y \rangle, \forall y \in \mathbb{R}^m \} \).

**Definition 1.1.** The proximal mapping (or proximal operator) of a proper closed convex function \( g : \mathbb{R}^n \to \mathbb{R} \cup \{ +\infty \} \) is defined as: let \( \gamma > 0 \)

\[
\text{prox}_{\gamma g}(y) = \arg \min_x \{ \gamma g(x) + \frac{1}{2} \| x - y \|_2^2 \}.
\]

(15)

For nuclear norm, its proximal mapping is singular value thresholding (SVT) [12], which is the lifting of vector soft-shrinkage thresholding to matrix [7].

**Lemma 1.1 (Variational formulation of nuclear norm [49, 50]).** Let \( X \in \mathbb{R}^{m \times n} \) and \( U \in \mathbb{R}^{m \times r}, V \in \mathbb{R}^{r \times n} \) with \( r \geq \text{rank}(X) \). We can write

\[
\| X \|_* = \min_{X=UV, U \in \mathbb{R}^{m \times r}, V \in \mathbb{R}^{r \times n}} \| U \|_1 \| V \|_2 = \min_{X=UV, U \in \mathbb{R}^{m \times r}, V \in \mathbb{R}^{r \times n}} \frac{1}{2} (\| U \|^2 + \| V \|^2).
\]

Paper organization The rest of the paper is organized as following. In Section 2, we describe our proposed algorithm ProGrAMMe (Algorithm 1) and prove its global convergence property. In Section 3, we first prove the rank identification property of Algorithm 1 and then propose a rank continuation strategy in Algorithm 2. Extensive numerical experiments are provided in Section 4, followed by the conclusion of this paper.

2 An SVD-free algorithm
Problem (5) is the composition of a smooth term and a non-smooth term. In the literature, numerical methods for such a structured problem are well studied, such as proximal gradient descent [36] (a.k.a. Forward–Backward splitting) and its various variants including the celebrated FISTA [6, 14]. Indeed, our problem (5)
can be handled by proximal gradient descent. However, such a method requires repeated SVD computation, which can significantly slow down its performance in many practical scenarios where the data size is large. Therefore, in this section, by combining proximal gradient descent and the variational formulation of nuclear norm (c.f., Lemma 1.1), we propose a SVD-free method for solving (5).

2.1 Proposed algorithm

In this part, we provide a detailed derivation of our algorithm, which is a combination of proximal gradient descent, alternating minimization, and inertial acceleration. For convenience, denote \( f(X) = \frac{1}{2} \| (\Psi(X) - F) \circ W \|^2 \) and \( g(X) = \| X \|_* \).

**Step 1 - Inertial proximal gradient descent** The first step to derive our algorithm is applying an inertial proximal gradient descent [36, 33] to solve problem (5). Since \( \Psi \) is a bounded linear mapping, we have the following simple lemma.

**Lemma 2.1.** Let \( \tilde{W} = W \circ W \). The loss \( f(X) \) is smoothly differentiable with its gradient given by
\[
\nabla f(X) = \nabla \Psi(X) ( (\Psi(X) - F) \circ \tilde{W} ),
\]
which is \( L \)-Lipschitz continuous with \( L = \max_{i,j} \| \nabla \Psi(X) \|_i,j \).

Below we provide two examples of \( \nabla \Psi(X) \):
- In compressed sensing scenario, \( \Psi \in \mathbb{R}^{d \times mn} \) is a linear measurement matrix, \( \Psi(X) = \Psi \text{vec}(X) \) and \( \| \nabla \Psi(X) \| = \| \Psi \| \).
- For matrix completion problem, \( \Psi \in \mathbb{R}^{m \times n} \) is a binary mask, and \( \Psi(X) = \Psi \circ X \) and \( \| \nabla \Psi(X) \| = 1 \).

In the literature, a routine approach to solve (5) is inertial proximal gradient descent. Let \( X_0 \in \mathbb{R}^{m \times n} \) be an arbitrary starting point, we consider the following iteration
\[
Y_k = X_k + a_k (X_k - X_{k-1}),
\]
\[
Z_k = Y_k - \gamma \nabla \Psi(Y_k) ((\Psi(Y_k) - F) \circ \tilde{W}),
\]
\[
X_{k+1} = \arg \min_{X \in \mathbb{R}^{m \times n}} \tau \| X \|_* + \frac{1}{2\gamma} \| X - Z_k \|^2,
\]
where \( a_k \in [0, 1] \) is the inertial parameter, \( \gamma \in [0, 2/L] \) is the step-size and the last line of (16) is nothing but SVT of \( Z_k \). Iteration (16) is a special case of the general inertial scheme proposed in [33], and we refer to [33] and the references therein for more discussion on inertial schemes.

**Step 2 - Alternating minimization** As computing the proximal mapping of nuclear norm requires SVD, the goal of second step is to avoid SVD in solving SVT of \( Z_k \) by incorporating the variational formulation of nuclear norm. To this end, the subproblem of (16) reads
\[
\min_{X \in \mathbb{R}^{m \times n}} \tau \| X \|_* + \frac{1}{2\gamma} \| X - Z_k \|^2.
\]
By plugging in the variational formulation of nuclear norm, we arrive at the following constrained minimization problem: let \( r > \text{SVT}(Z_k) \)
\[
\min_{X \in \mathbb{R}^{m \times n}, U \in \mathbb{R}^{m \times r}, V \in \mathbb{R}^{n \times r}} \frac{1}{2} \| X - Z_k \|^2 + \frac{\tau r}{2} (\| U \|^2 + \| V \|^2) \quad \text{such that} \quad X = UV.
\]
Instead of considering the augmented Lagrangian multiplier of the constraint [11], we directly substitute the constraint \( X = UV \) in the objective, which leads to
\[
\min_{U \in \mathbb{R}^{m \times r}, V \in \mathbb{R}^{n \times r}} \frac{1}{2} \| UV - Z_k \|^2 + \frac{\tau r}{2} (\| U \|^2 + \| V \|^2),
\]
which is a smooth, bi-convex optimization problem in each component \( U \) and \( V \).
Different from (17), problem (19) does not admits closed form solution. However, when either $U$ or $V$ is fixed, the problem becomes a simple least square. Hence we can solve (19) via a simple alternating minimization, namely a two block Gauss-Seidel iteration [3]: given $U_0 \in \mathbb{R}^{m \times r}, V_0 \in \mathbb{R}^{r \times n}$

\[
\begin{align*}
U_{i+1} &= Z_i V_i^\top (V_i V_i^\top + \tau \gamma \text{Id}_r)^{-1}, \\
V_{i+1} &= (U_i^\top U_{i+1} + \tau \gamma \text{Id}_r)^{-1} U_{i+1}^\top Z_i,
\end{align*}
\]

(20)

where $\text{Id}_r$ denotes the identity operator on $\mathbb{R}^r$. Substituting (20) into (16) as an inner loop, we obtain the following iterative scheme: let $I \in \mathbb{N}_+$.

\[
Z_k = Y_k - \gamma \nabla \Psi(Y_k)((\Psi(Y_k) - F) \odot \tilde{W}),
\]

Initialize $U_0, V_0$. For $i = 0, \ldots, I - 1$: \[\begin{align*}
U_{i+1} &= Z_i V_i^\top (V_i V_i^\top + \tau \gamma \text{Id}_r)^{-1}, \\
V_{i+1} &= (U_{i+1}^\top U_{i+1} + \tau \gamma \text{Id}_r)^{-1} U_{i+1}^\top Z_i,
\end{align*}\]

(21)

Assembling the above steps, we obtain our proposed algorithm, proximal gradient & alternating minimization method, which we call ProGrAMMe and is summarized below in Algorithm 1.

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**Algorithm 1: A Proximal Gradient & Alternating Minimization Method (ProGrAMMe)**

1: Compute $\tilde{W} = W \odot W, L$ and let $\gamma \in [0, 2/L]_0$; Choose $r > 0$ and $I \in \mathbb{N}_+$;
2: while not convergent do
3: $Y_k = X_k + a_k(X_k - X_{k-1}),$ \hspace{1cm} // inertial step //
4: $Z_k = Y_k - \gamma \nabla \Psi(Y_k)((\Psi(Y_k) - F) \odot \tilde{W}),$ \hspace{1cm} // gradient descent //
5: Initialize $U_0 \in \mathbb{R}^{m \times r}, V_0 \in \mathbb{R}^{r \times n}$,
6: for $i = 1, \ldots, I - 1$ do \hspace{1cm} // inner loop //
7: $U_{i+1} = Z_i V_i^\top (V_i V_i^\top + \tau \gamma \text{Id}_r)^{-1}$,
8: $V_{i+1} = (U_{i+1}^\top U_{i+1} + \tau \gamma \text{Id}_r)^{-1} U_{i+1}^\top Z_i$,
9: end for
10: $X_{k+1} = U_1 V_1$.
11: end while
12: return $X_{k+1}$

---

**Remark 2.2.**

- One highlight of our algorithm is that, via variational formulation of nuclear norm, we relaxed the convex subproblem (17) to a non-convex (19) problem.
- Every step of Algorithm 1 requires initializing $U_0, V_0$ for the inner step, and the simplest way is using the $U_i, V_i$ from the last step.
- For the controlling parameter $r$, theoretically it does not make any difference as long as it is larger than the rank of the solution of (5). However, practically it is crucial to the performance of Algorithm 1. Detailed discussion is provided in Section 3.

**Remark 2.3.** In the literature, several SVD-free approaches were proposed. For example, variational formulation was also considered in [11] and the resulted problem was solved by method of Lagrange multiplier while we directly plug the constraint into the objective. In [62], a dual characterization of the nuclear norm was used and a SVD-free gradient descent was designed. The benefits of our approach, as we shall see later, are simple convergence analysis (see Section 2.2) and extensions to more general settings (see Section 2.3).
Remark 2.4 (Per iteration complexity). Comparing Algorithm 1 to proximal gradient descent (16), the only difference is Line 5–9. For proximal gradient descent, since SVD is needed, the iteration complexity each step is \(O(mn\min\{m, n\})\). For Algorithm 1, suppose \(I = 1\), the complexity of Line 7–8 is \(O((m + n + r)r^2)\). It can be concluded that, the smaller the value of \(r\) (still larger than the rank of the solutions), the lower the per iteration complexity of Algorithm 1. As a result, the choice of \(r\) is crucial to the practical performance of Algorithm 1. Therefore, in Section 3, a detailed discussion is provided on how to choose \(r\).

Relation with existing work Our algorithm is closely related with proximal splitting method and its variants, as our first step to derive Algorithm 1 is the inertial proximal gradient descent (for example, proximal gradient descent [36] and its accelerated versions including FISTA [6, 14], as in [51, 55, 30] where FISTA was adopted to solving low-rank recovery problem).

Our model (5) and Algorithm 1 share similarities with those of [10, 15, 11], but there are some fundamental differences. First of all, all these works consider only the case \(\Psi = \text{Id}\), i.e. \(\Psi\) is an identity mapping

\[
\min_{X} \frac{1}{2} \|X - F \odot W\|^2 + \tau \|X\|_*. \tag{22}
\]

In [10, 11], the authors consider directly applying matrix factorization to (22) which results in: let \(r > 0\)

\[
\min_{U \in \mathbb{R}^{m \times r}, V \in \mathbb{R}^{n \times r}} \frac{1}{2} \|UV - F \odot W\|^2 + \frac{r}{2} \left(\|U\|^2 + \|V\|^2\right), \tag{23}
\]

which is a smooth and bi-convex optimization problem. Our approach, on the other hand, only consider applying matrix factorization for the subproblem of proximal gradient descent (16).

It is worth noting that (23) shares the same continuous property as (19), hence can by handled by alternating minimization algorithm. Both (23) and (19) are also special cases of the following non-convex problem

\[
\min_{U, V} f(U, V) + \tau_1 g_1(U) + \tau_2 g_2(V) \tag{24}
\]

where \(f(U, V)\) is differentiable with Lipschitz continuous gradient, \(\tau_1, \tau_2 > 0\) are regularization parameters and \(g_1(\cdot), g_2(\cdot)\) are (non-smooth) regularization terms for \(U, V\), respectively. Problem (24) was well studied in [2, 8]; for instance, the following algorithm was proposed in [2]:

\[
U_{k+1} = \arg \min_{U \in \mathbb{R}^{m \times r}} \left\{ f(U, V_k) + \tau_1 g_1(U) + \frac{1}{2\alpha} \|U - U_k\|^2 \right\},
\]

\[
V_{k+1} = \arg \min_{V \in \mathbb{R}^{n \times r}} \left\{ f(U_{k+1}, V) + \tau_2 g_2(V) + \frac{1}{2\gamma} \|V - V_k\|^2 \right\},
\]

where \(\alpha, \gamma > 0\) are parameters. Specializing to the case of (23), we get

\[
U_{k+1} = \arg \min \left\{ \frac{1}{2} \|UV_k - F \odot W\|^2 + \frac{r}{2} \|U\|^2 + \frac{1}{2\alpha} \|U - U_k\|^2 \right\},
\]

\[
V_{k+1} = \arg \min \left\{ \frac{1}{2} \|(U_{k+1}V - F) \odot W\|^2 + \frac{r}{2} \|V\|^2 + \frac{1}{2\gamma} \|V - V_k\|^2 \right\}.
\]

It can be observed that, though sharing similarities, our proposed algorithm is different from the above schemes. Same for the algorithm proposed in [8].

2.2 Global convergence of Algorithm 1

In this part, we provide global convergence analysis of Algorithm 1. The key of our proof is rewriting Algorithm 1 as an inexact version of inertial proximal gradient descent (16) whose convergence property is well established in the literature. Such an equivalence is obtained based on the result below from [11, Theorem 1].

Lemma 2.5 ([11, Theorem 1]). Let \(\hat{X}\) be the unique minimizer of (17) with rank \(\hat{r} = \text{rank}(\hat{X})\), and \((\hat{U}, \hat{V})\) a solution of (19) with \(r \geq \hat{r}\). There holds \(\hat{X} = \hat{U}\hat{V}\).

The above lemma implies that, although we relaxed the strongly convex problem (17) to a non-convex one (19), we can recover the unique minimizer of (17) via solving (19). In turn, we can cast Algorithm 1 back to the proximal gradient descent (16), possibly with approximation errors due to finite step inner loop, and then prove its convergence. To this end, we first propose the following inexact characterization of Algorithm 1.
Proposition 2.6 (Inexact inertial proximal gradient descent). For Algorithm 1, let $I \in \mathbb{N}_+$. Then Algorithm 1 is equivalent to the following inexact inertial proximal gradient descent
\begin{align*}
Y_k &= X_k + a_k (X_k - X_{k-1}), \\
Z_k &= Y_k - \gamma \nabla \psi \left( (\Psi(Y_k) - F) \circ \tilde{W} \right), \\
X_{k+1} &= \text{prox}_{\tau \gamma}(Z_k + e_k)
\end{align*}
where $e_k$ accounts for truncation error for finite-valued $I$, and has the form
\begin{equation}
e_k = P_I(S_I + s_k)Q_T^T - Z_k,
\end{equation}
with $P_I S_I Q_T^T$ being the SVD of $X_{k+1}$ and $s_k$ a diagonal matrix with all its diagonal elements $s_k(j,j), i = 1, \ldots, \min\{m,n\}$
\begin{equation}
s_k(j,j) = \begin{cases} 
\tau \gamma : S(j,j) \geq \tau \gamma, \\
0, \tau \gamma : S(j,j) < \tau \gamma.
\end{cases}
\end{equation}

Proof. Let the SVD $Z_k$ be $Z_k = PSQ_T$. Suppose the inner loop is ran for infinite steps of Algorithm 1, i.e. (19) is solved exactly, then we have
\begin{equation}
X_{k+1}' = U_{+\infty}V_{+\infty} = \text{prox}_{\tau \gamma}(Z_k) = P\mathcal{T}_{\tau \gamma}(S)Q_T^T,
\end{equation}
where $\mathcal{T}_{\tau \gamma}(S)$ is soft-thresholding operator [7]. Now we truncate the inner loop to $I$ finite steps, that is
\begin{equation}
X_{k+1} = U_I V_I = P_I S_I Q_T^T
\end{equation}
where $P_I S_I Q_T^T$ is the SVD of $X_{k+1}$ and $I$ is the number of iteration for inner loop. Apparently, we have
\begin{equation*}
I \to +\infty : P_I \to P, \quad S_I \to \mathcal{T}_{\tau \gamma}(S) \quad \text{and} \quad Q_I \to Q.
\end{equation*}
Let $e_k$ be such that
\begin{equation}
Z_k + e_k = P_I(S_I + s_k)Q_T^T - Z_k
\end{equation}
where $s_k$ is a diagonal matrix in (26). As a result, we get $e_k = P_I(S_I + s_k)Q_T^T - Z_k$ and conclude the proof. \qed

The inexact formulation of Algorithm 1 allows us to prove its convergence in a rather simple fashion, as (25) is nothing but a special case of the inertial proximal gradient descent discussed in [33]. As a consequence, we have the following result regarding the global convergence of Algorithm 1.

Proposition 2.7 (Convergence of Algorithm 1). For Algorithm 1, let the inertial sequences $\{a_k\}_{k \in \mathbb{N}}$ be such that
\begin{equation*}
\limsup_k a_k < 1 \quad \text{and} \quad \sum_{k \in \mathbb{N}} a_k \|X_k - X_{k-1}\|^2 < +\infty.
\end{equation*}
If, moreover, the error $e_k$ is such that
\begin{equation*}
\sum_{k \in \mathbb{N}} k \|e_k\| < +\infty.
\end{equation*}
Then, there exists $X^* \in \text{arg min}(f + \tau g)$ to which the sequence $\{X_k\}_{k \in \mathbb{N}}$ generated by Algorithm 1 converges.

The result is simply implied from [33, Theorem 3], hence we omit the proof here and refer to [33] for detailed discussions.

Remark 2.8. The condition on error $e_k$ implies that the inner problem needs to be solved with an increasing accuracy, i.e. an increasing number of inner iterations. However, in practice, fixed choice of $I$ works quite well; see our numerical examples later. The summability of $a_k \|X_k - X_{k-1}\|^2$ can be guaranteed by certain choices of $a_k$; See [33, Theorem 4]. One can also use an online approach to determine $a_k$ such that the summability condition holds. For instance, let $a \in [0,1]$ and $c > 0, \delta > 0$, then $a_k$ can be chosen as $a_k = \min\{a, \frac{c}{k^{1+a}\|X_k - X_{k-1}\|^2} \}$. 

8
Remark 2.9 (FISTA-like inertial parameter). If the error term $e_k$ can be carefully taken care of, such as gradually increase the value of $I$, then according to [4] FISTA rule for updating $a_k$ can be applied, e.g. $a_k = \frac{k-1}{k+d}$ for $d > 2$. One can then use the lazy-start strategy of [34] for further speed-up.

Remark 2.10. In [33], the authors also studied the local linear convergence of proximal gradient descent type methods, under the notion of “partial smoothness” (Definition 3.1). However, due to the existence of the error term $e_k$, the local linear convergence of (25) is more complicated than that of [33] where the analysis was obtained only for the exact case. Therefore, for this aspect, we forgo the theoretical analysis and only provide numerical discussions in Section 3.

### 2.3 Generalization of Algorithm 1

Throughout this paper, our main focus is (5) whose loss function is a simple weighted least square. In this scope, we discuss several generalization of Algorithm 1, including more general loss function (e.g. (6)), the non-convex setting and non-linear $\Psi$.

**General loss function** Algorithm 1 is loss function agnostic, as for proximal gradient descent type methods, the condition required for the smooth part is that the function should be smoothly differentiable with gradient being Lipschitz continuous. Therefore, we can apply Algorithm 1 to solve the more general model (6), and the only change we need to make to Algorithm 1 is Line 4 for which we now have

$$Z_k = Y_k - \gamma \nabla_1 f(Y_k, F, W),$$

where $\nabla_1$ denotes the gradient of $f(X, F, W)$ with respect to $X$. The global convergence result stays the same for the above update.

**Non-convex loss function** Algorithm 1 does not require convexity for the loss function. In the literature, the convergence properties of proximal gradient descent type methods for non-convex optimization are well studied, most of them are obtained under Kurdyka-Łojasiewicz inequality owing to the pioneered work [3]. As Algorithm 1 is a special case of inexact proximal gradient descent, it can also be applied to solve problems where the loss function $f(X, F, W)$ is non-convex. Parameter-wise, there are two main differences between the non-convex and convex cases:

- For step-size $\gamma$, different from the convex case whose upper bound is $2/L$, it reduces to $1/L$ for the non-convex case.
- The conditions on the error is different. For the non-convex case, the error should be such that a descent property of certain stability function (see e.g. [3]) should be maintained. As a result, line search might be needed for the number of inner loop iteration.

Remark 2.11. While keeping the loss function as least square, it still can be non-convex because of the operator $\Psi$, for which case $\Psi$ is a non-linear smooth mapping instead of being linear. For this case, as long as $\Psi$ is such that the gradient is Lipschitz continuous, global convergence of Algorithm 1 can be guaranteed.

### 3 Rank continuation

As we discussed above, the choice of parameter $r$ is crucial to the performance of Algorithm 1: let $r^*$ be the rank of a solution $X^*$ of the problem (5), then the closer the value of $r$ to $r^*$, the better practical performance of Algorithm 1, and the best performance can be obtained when $r = r^*$. However, in general it is impossible to know $r^*$ a priori, and usually an overestimation of $r^*$ is provided which damps the efficiency of the algorithm. In this section, we first discuss the rank identification property of Algorithm 1, and then discuss a rank continuation strategy which asymptotically achieves the optimal per iteration complexity.

#### 3.1 Rank identification

In [33], for proximal gradient descent type methods with non-smooth regularization, it was shown that the sequence generated by these methods has a so-called “finite activity identification property”. For nuclear norm,
this means that after a finite number of iterations, there holds \( \text{rank}(X_k) = \text{rank}(X^*) \), for all \( k \) large enough where \( X^* \) is the solution to which \( X_k \) converges. In the theorem below, we show that Algorithm 1 also has this finite rank identification property. We continue to use the notations \((f, g)\) used in subsection 2.1.

**Theorem 3.1 (Rank identification).** For Algorithm 1, suppose the conditions of Proposition 2.7 hold, then \( X_k \) converges to \( X^* \in \text{Arg min}(f + \tau g) \). If, moreover, the following non-degeneracy condition holds

\[
-\nabla \Psi(X^*)(\Psi(X^*) - F) \cap W) \in \tau \text{ri}(\partial \|X^*\|_1),
\]

then there exists a \( K > 0 \) such that for all \( k \geq K \) there holds \( \text{rank}(X_k) = \text{rank}(X^*) \).

To prove the result, we need the help of 
partly smoothness, which was first introduced in [32]. Let \( \mathcal{M} \) be a \( C^2 \)-smooth embedded submanifold of \( \mathbb{R}^n \) around a point \( x \). To lighten notation, henceforth, we use \( C^2 \)-manifold instead of \( C^2 \)-smooth embedded submanifold of \( \mathbb{R}^n \). The natural embedding of a submanifold \( \mathcal{M} \) into \( \mathbb{R}^n \) permits to define a Riemannian structure on \( \mathcal{M} \), and we simply say \( \mathcal{M} \) is a Riemannian manifold. \( \mathcal{T}_\mathcal{M}(x) \) denotes the tangent space to \( \mathcal{M} \) at any point near \( x \) in \( \mathcal{M} \).

**Definition 3.1 (Partial smoothness).** Let \( g \) be proper closed and convex, \( g \) is said to be \textit{partly smooth at \( x \) relative to a set} \( \mathcal{M} \) containing \( x \) if \( \partial g(x) \neq \emptyset \). We also define the smoothness, sharpness, and continuity of \( g \) at \( x \) relative to a set \( \mathcal{M} \) as follows:

- **Smoothness:** \( \mathcal{M} \) is a \( C^2 \)-manifold around \( x \), \( g \) restricted to \( \mathcal{M} \) is \( C^2 \) around \( x \);
- **Sharpness:** The tangent space \( \mathcal{T}_\mathcal{M}(x) \) coincides with \( T_x = \text{par}(\partial g(x))^{-1} \);
- **Continuity:** The set-valued mapping \( \partial g \) is continuous at \( x \) relative to \( \mathcal{M} \).

For nuclear norm, it is partly smooth along the set of fixed-rank matrices [32]. Other examples of partly smooth functions including \( \ell_1 \)-norm for sparsity, \( \ell_{1,2} \)-norm for group sparsity, etc; We refer to [33] and the references therein for more examples of partly smooth functions.

**Proof of Theorem 3.1.** Since \( f \) locally is \( C^2 \)-smooth around \( X^* \), the smooth perturbation rule of partly smooth functions [32, Corollary 4.7], ensures that \( f + \tau g \) is partial smoothness at \( X^* \) relative to \( \mathcal{M}_{X^*} \) def \( \{X \in \mathbb{R}^{m \times n} : \text{rank}(X) = \text{rank}(X^*) \} \).

By assumption, the sequence \( X_k \) created by Algorithm 1 converges to \( X^* \in \text{arg min}(f + \tau g) \). The non-degeneracy condition (27) is equivalent to \( 0 \in \text{ri}(\partial (f + \tau g)(X^*)) \). Now (25) is equivalent to

\[
Y_k - \gamma \nabla f(Y_k) - X_{k+1} + e_k \in \gamma \tau \partial g(X_{k+1})
\]

\[
\Leftrightarrow (Y_k - \gamma \nabla f(Y_k)) - (X_{k+1} - \gamma \nabla f(X_{k+1})) + e_k \in \gamma \partial (f + \tau g)(X_{k+1}).
\]  

By Baillon-Haddad theorem [5], \( \text{Id} - \gamma \nabla f \) is non-expansive, whence we get

\[
\text{dist}(0, \partial (f + \tau g)(X_{k+1})) \leq \|(\text{Id} - \gamma \nabla f)(Y_k) - (\text{Id} - \gamma \nabla f)(X_{k+1})\| + \|e_k\|
\]

\[
\leq \|Y_k - X_{k+1}\| + \|e_k\|
\]

\[
\leq \|X_k - X_{k+1}\| + a_k \|X_k - X_{k+1}\| + \|e_k\|.
\]

Since \( X_k \) is convergent and \( \|e_k\| \to 0 \), we have that \( \text{dist}(0, \partial (f + \tau g)(X_k)) \to 0 \). Owing to our assumptions, \( f + \tau g \) is sub-differentially continuous at every point in its domain, and in particular at \( X^* \) for 0, which in turn entails \( (f + \tau g)(X_k) \to (f + \tau g)(X^*) \). Altogether, this shows that the conditions of [27, Theorem 5.3] are fulfilled, and the rank identification result follows.

To demonstrate the rank identification property of Algorithm 1, the following low-rank recovery problem is considered as an illustration:

\[
\min_{X \in \mathbb{R}^{100 \times 100}} \frac{1}{2} \|(\Psi(X) - F) \cap W\|^2 + \tau \|X\|_1,
\]

where we have \( \Psi \in \mathbb{R}^{2352 \times 10000} \) and \( F = \Psi \text{vec}(\hat{X}) + \epsilon \).
with \( \text{rank} (\hat{X}) = 4 \) and \( \varepsilon \) being random Gaussian noise. Moreover, we choose \( \tau = 2\|\varepsilon\| \).

The problem is solved with ProGrAMMe with \( a_k \equiv 0 \) and standard proximal gradient descent (PGD) \([36]\), and the observation is shown below in Figure 1. We observed that, both schemes have rank identification property, as the rank of \( X_k \) for both schemes eventually becomes constant.

![Figure 1: Rank identification of Algorithm 1.](image-url)

### 3.2 Rank continuation

As we remarked above that the choice of \( r \) in Algorithm 1 is crucial to the practical performance of the method. To overcome the difficulty of a tight estimation of the rank of minimizers. In this section, we introduce a rank continuation strategy, see Algorithm 2, which adaptively adjusts the rank of the output and asymptotically attains the optimal per iteration complexity.

**Algorithm 2:** ProGrAMMe with Rank Continuation

1. Compute \( \tilde{W} = W \odot W \), \( L \) and \( \gamma \in ]0, 2/L[ \); Choose \( r > 0 \) and \( I \in \mathbb{N}_+ \);
2. while not convergent do
3. \( Y_k = X_k + a_k (X_k - X_{k-1}) \), \hspace{1cm} //inertial step//
4. \( Z_k = Y_k - \gamma \nabla \Psi (Y_k) \left( (\Psi (Y_k) - F) \odot \tilde{W} \right) \), \hspace{1cm} //gradient descent//
5. Initialize \( U_0 \in \mathbb{R}^{m \times r}, V_0 \in \mathbb{R}^{r \times n} \),
6. for \( i = 1, \ldots, I - 1 \) do \hspace{1cm} //inner loop//
7. \( U_{i+1} = Z_k V_i^\top (V_i V_i^\top + \tau \gamma I_r)^{-1} \)
8. \( V_{i+1} = (U_{i+1}^\top U_{i+1} + \tau \gamma I_r)^{-1} U_{i+1}^\top Z_k \)
9. end for
10. \( X_{k+1} = U_I V_I \),
11. \( r = \text{rank}(U_I) \), \hspace{1cm} //rank continuation//
12. end while
13. return \( X_{k+1} \)

**Remark 3.2.** In Line 11, instead of using \( \text{rank}(X_{k+1}) \) to update \( r \), we choose to use \( \text{rank}(U_I) \), since \( \text{rank}(X_{k+1}) \leq \min \{ \text{rank}(U_I), \text{rank}(V_I) \} \) and it is less computational demanding to evaluate the rank of \( U_I \) than that of \( X_{k+1} \).

**Remark 3.3.** Though the initial value of \( r \) is no longer as important as that of Algorithm 1 whose \( r \) is fixed, it is still beneficial to have a relatively good estimate of \( \text{rank}(X^*) \) as it can further reduce the computational cost.
of the algorithm. Also, it is not desirable to compute \( \text{rank}(U_I) \) every iteration and a practical approach is to do it in a regular interval.

To illustrate the performance of rank continuation, we consider a low-rank recovery problem with randomly missing entries:

\[
\min_{X \in \mathbb{R}^{2000 \times 2000}} \frac{1}{2} \| (\Psi \odot X - F) \odot W \|^2 + \tau \| X \|_*,
\]

where \( \Psi \in \mathbb{R}^{2000 \times 2000} \) is a random binary mask with 50% entries equal to 0 and

\[ F = \Psi \odot \hat{X} + \epsilon \]

with \( \text{rank}(\hat{X}) = 10 \) and \( \epsilon \) being random Gaussian noise. For this case we set \( \tau = \| \epsilon \| \).

**Comparison to SVD based PGD/FISTA** We first compare the performances of PGD/FISTA, Algorithm 1 (ProGrAMMe) and Algorithm 2 (ProGrAMMe-RC), with the following settings:

- The step-size \( \gamma \) for all these schemes are chosen as \( \gamma = 1.25/L \). Note that this choice of \( \gamma \) exceeds the upper bound of step-size of FISTA, however for this example FISTA converges.
- For FISTA, we adopt the “lazy start” strategy, e.g. \( a_k = k - 1 \), proposed in [34] which is faster than the standard FISTA scheme.
- For both Algorithm 1 (ProGrAMMe) and Algorithm 2 (ProGrAMMe-RC), we choose \( I = 1 \) and \( a_k \equiv 0 \). We set \( r = 500 \) for ProGrAMMe which is also the initial value of \( r \) for Algorithm 2.

All schemes are stopped when the relative error \( \| X_k - X_{k-1} \| \) reaches \( 10^{-10} \), and we note the the average wall clock time of 10 runs for these schemes as:

| Schemes            | PGD   | FISTA | ProGrAMMe | ProGrAMMe-RC |
|--------------------|-------|-------|-----------|--------------|
| Time (in seconds)  | 127.77| 104.70| 14.68     | 8.39         |

As illustrated in Figure 2 (a), we observe:

- PGD/FISTA is much slower than Algorithm 1 and Algorithm 2. In particular, the rank continuation scheme is about an order faster than PGD/FISTA.
- ProGrAMMe-RC is about 40% faster than ProGrAMMe which indicates the advantage of rank continuation under the considered setting.

For the magenta line in Figure 2 (a) for ProGrAMMe-RC, it has several jumps which is due to the update of \( r \).

**Remark 3.4.** For FISTA scheme, one can also consider the adaptive restarted scheme of [46], which can further improve the performance of FISTA. However, as long as SVD is involved, it cannot compete with ProGrAMMe for the considered relative large-scale problem.

**Effect of different starting rank** To further understand the advantage of rank continuation over the static one, we conduct a comparison of Algorithm 2 under different initial values for \( r \). Precisely, we considered:

- four different values of \( r = 500, 300, 100, 50 \).
- \( r \) is updated every 10 steps.

We use “nRC” to denote ProGrAMMe without rank continuation and “RC” with rank continuation, and the result is shown in Figure 2 (b). We observed:

- Without rank continuation, the smaller the value of \( r \), the better the performance of ProGrAMMe.
- For \( r = 500, 300 \), the red and black lines in the figure, rank continuation (dashed lines) shows clear advantage over the standard scheme (solid lines).
- While for \( r = 100, 50 \), rank continuation actually becomes slower than the static scheme, and the extra time is mainly the overhead of computing \( \text{rank}(U_I) \).

From the above observations, we conclude:

- For problems where a tight estimation of the rank of the solution can be obtained, one can simply consider Algorithm 1;
• When the rank of solutions is difficult to estimate, then rank continuation can be applied to achieve acceleration.

We leave the comparison of ProGrAMMe with inertial to the next section.

4 Numerical experiments

To understand the effects of inertial acceleration, validate the strengths and flexibility of our recovery model and algorithm, in this section we perform numerical experiments on several low-rank recovery problems. Throughout this section, we typically use two different versions of ProGrAMMe—(i) ProGrAMMe-1 which terminates the inner loop in each iteration and (ii) ProGrAMMe-ε which terminates the inner loop when the relative error of the inner iterates reach an ε precision or maximum inner iteration is achieved, whichever occurs first. Throughout the section, for ProGrAMMe-ε, we use $\varepsilon = 10^{-4}$ and maximum number of inner iteration is set to 20, unless otherwise specified.

4.1 Effects of inertial acceleration

We continue the matrix completion problem (30) to study the effect of inertial acceleration. Both Algorithm 1 and 2 are tested, the setting of the tests are

• For both algorithms, we initialize $r$ with value of 500; In terms of step-size, we keep the previous choice which is $\gamma = 1.25/L$.
• In total, 5 different choices of inertial parameter $a_k$ are considered

$$a_k \equiv 0, \ a_k \equiv \frac{1}{4}, \ a_k \equiv \frac{1}{2}, \ a_k \equiv \frac{3}{4} \quad \text{and} \quad a_k = \frac{k-1}{k+20}.$$ 

The results are shown in Figure 3, whose left figure is the comparison of Algorithm 1 without rank continuation

• In general, inertial schemes demonstrate faster performance than the standard scheme. For example, $a_k \equiv \frac{3}{4}$ is about 30% faster than $a_k \equiv 0$.
• For $a_k \equiv \frac{1}{1}, \frac{1}{2}, \frac{3}{4}$, the larger the value of $a_k$, the faster the performance of inertial scheme.
• Lazy-start FISTA performs no better than $a_k \equiv \frac{3}{4}$.

The comparison for rank continuation scheme is provided in Figure 3 (b), where similar observation as above can be obtained.
4.2 Low-rank recovery experiments on synthetic data

For these experiments, we generate the low-rank matrix, $L$, as a product of two independent full-rank matrices of size $m \times r$ with $r < m$ such that elements are independent and identically distributed (i.i.d.) and sampled from a normal distribution — $\mathcal{N}(0, 1)$. We used two different types of sparse noise—Gaussian noise and arbitrary large noise. For each case, we generate the sparse matrix, $S$, such that for a sparsity level $\alpha \in (0, 1)$, the sparse support is created randomly.

- For random Gaussian noise, we construct the sparse matrix $S_{\text{random}}$ whose elements are i.i.d. $\mathcal{N}(0, 1)$ random variables and form $F$ as: $F = L + \eta S_{\text{random}}$, where $\eta$ controls the noise level and we set $\eta = 0.2 \max_{i,j}(L)_{ij}$.
- For large noise, we generate the sparse matrix, $S_{\text{sparse}}$, such that its elements are chosen from the interval $[-50, 50]$ and construct $F$ as $F = L + S_{\text{sparse}}$.

We fix $m = 100$, define $\rho_r = \text{rank}(L)/m$, where $\text{rank}(L)$ varies and set the sparsity level $\alpha \in (0, 1)$. For each $(\rho_r, \alpha)$ pair, we apply RPCA GD [63], NCF [19], and ProGrAMMe to recover a low-rank matrix $X$. We consider RMSE, $\|F - X\|/\sqrt{mn}$, as performance measure.

For each class of noise, we run the experiments for 10 times and plot the average RMSEs for each $(\rho_r, \alpha)$. Note that, RPCA GD and NCF use an operator $\mathcal{T}_{\eta}[S]$ that does not perform an explicit Euclidean projection onto the sparse support of $S$, as the exact projection on $S$ is expensive [19, 63, 18]. Inspired by this, for sparse noise, we design our weight matrix such that it has large weights for the sparse support of $S$ and 1 otherwise. However, for random noise, we simply use a random weight matrix. From Figure 4 we see that for both random and sparse noise, ProGrAMMe has the least average RMSEs. Moreover, while two PCP algorithms show significant differences in their RMSE diagrams, our ProGrAMMe produce almost similar RMSE and obtain lower values compare to PCP algorithms in both types of noises.

Effect of the condition number of $W$ on the convergence of ProGrAMMe

The problem (22) is tricky, as the condition number, $\kappa_W$ of the weight matrix, $W$ plays an important role in convergence [47, 17]. We perform a detailed empirical convergence analysis of ProGrAMMe-\(\epsilon\) and ProGrAMMe-1 on synthetic data in Appendix A.2 by varying $\kappa_W$ and compared with proximal algorithms. We observe that proximal algorithms are sensitive to $\kappa_W$ (a higher $\kappa_W$, translates to slower convergence), but both ProGrAMMe-\(\epsilon\) and ProGrAMMe-1 are less sensitive $\kappa_W$, maintains a stable convergence profile for different $\kappa_W$, and converge faster than proximal algorithms in almost all cases.
Figure 4: RMSE: $\|F - X\|/\sqrt{mn}$ for different methods. Top row is for random noise, bottom row is for sparse large noise. For both cases, $\min_{i,j} W_{ij} = 5, \max_{i,j} W_{ij} = 10$.

### 4.3 Real-world applications

To validate the strengths and flexibility of our proposed algorithms, we use three real world problems—structure from motion (SfM), matrix completion with noise, and background estimation from fully and partially observed datasets. We compared our algorithms against 15 state-of-the-art weighted and unweighted low-rank approximation algorithms (see Table 3 in Appendix).

**Structure from motion and photometric stereo** SfM uses local image features without a prior knowledge of locations or pose and infers a three dimensional structure or motion. For these experiments, we used three popular datasets\(^1\): non-rigid occluded motion of a giraffe in the background (for nonrigid SfM), a toy dinosaur (for affine SfM), and the light directions and surface normal of a static face with a moving light source (for photometric stereo) (See more details in Figure 5). The datasets have 69.35%, 23.08%, and 58.28% observable entries, respectively. Therefore, we use a binary mask as weight $W$ such that $W_{ij} = 1$ if the data has an entry at the $(i,j)$th position, otherwise, $W_{ij} = 0$. With this setup, our formulation works as a matrix completion.

\(^1\)http://www.robots.ox.ac.uk/abm/
Figure 5: Sample frame from the static Face, Giraffe and toy dinosaur sequences. The data matrices, $F$ are of size $2944 \times 20$, $240 \times 167$, and $319 \times 72$, respectively and the prior rank of the sequences are 4, 6, and 4, respectively.

problem. We compared ProGrAMMe-1 and ProGrAMMe-$\epsilon$ with respect to the damped Newton algorithm in [10]. Admittedly, [10] obtains the best factorization pair, $X = UV$ such that it gives minimum loss (within the observable entry), $\frac{\| (F - XV)W \|}{\| W \|}$ for all cases. Additionally, we also calculated the loss outside the observable entries, that is, $\frac{\| (F - XV)W \|}{\| (I - W) \|}$ where the performance of ProGrAMMe is better in all cases. See results provided in Table 2 below.

| Dataset     | $\frac{\| (F - XV)W \|}{\| W \|}$ | $\frac{\| (F - XV)W \|}{\| (I - W) \|}$ |
|-------------|-----------------------------------|---------------------------------|
| Giraffe     | 0.5085 (ProGrAMMe-1)              | 284.1509 (ProGrAMMe-1)          |
|             | 0.5399 (ProGrAMMe-$\epsilon$)    | **271.4828** (ProGrAMMe-$\epsilon$) |
|             | **0.3228** [10]                   | 364.2476 [10]                   |
| Toy Dino    | 4.4081 (ProGrAMMe-1)             | 244.5437 (ProGrAMMe-1)          |
|             | 5.2453 (ProGrAMMe-$\epsilon$)    | **239.6335** (ProGrAMMe-$\epsilon$) |
|             | **1.0847** [10]                   | 318.666 [10]                    |
| Face        | 0.023 (ProGrAMMe-1)              | **0.7143** (ProGrAMMe-1)        |
|             | 0.0232 (ProGrAMMe-$\epsilon$)    | 0.7338 (ProGrAMMe-$\epsilon$)   |
|             | **0.0223** [10]                   | 0.98 [10]                       |

Table 2: Comparisons between ProGrAMMe-$\epsilon$, ProGrAMMe-1, and damped Newton method on structure from motion and photometric stereo datasets.

Matrix completion with noise on power-grid data Matrix completion is one of the important special cases of weighted low-rank estimation problems as for this problem, the weights are reduced to $\{0,1\}$. This set of experiments are inspired by [15]. The dataset and the codes for BIRSVD are collected from author’s website\(^2\). In this experiment, the test dataset contains 48 hours of temperature data sampled every 30 minutes over 20 European cities. The $(i,j)^{th}$ entry of the matrix represents the $i^{th}$ temperature measurement for the $j^{th}$ city. As a prior information, we use the fact that temperature varies smoothly over time and the data matrix is low-rank.

For this data set, we use a rank 2 approximation. We run each algorithm with random initialization for 10 times and plot the average results in Figure 6. Each algorithm is run for 50 global iterations or tolerance set to machine precision, whichever attained first. For BIRSVD, ProGrAMMe-1, and BLF (with $\ell_2$ loss) we use $\tau = 0.006$. Additionally, for BIRSVD another regularizer is set to 0 (as in [15]). FPCA [39] computes a low rank approximation by using a prior information that the nuclear norm of the matrix is bounded. As it turns out, the EM algorithm [53] has a slower convergence and the performance of ProGrAMMe is albeit better. For

\(^2\)https://homepage.univie.ac.at/saptarshi.das/index.html
Background estimation

Background estimation and moving object detection from a video sequence is a classic problem in computer vision and it plays an important role in human activity recognition, tracking, and video analysis from surveillance cameras. In the conventional matrix decomposition framework used for background estimation, the video frames are concatenated in a data matrix $F$, and the background matrix, $X$, is of low-rank [45], as the background frames are often static or close to static. However, the foreground is usually sparse. The desired target rank of the background is hard to determine due to some inherent challenges, such as, changing illumination, occlusion, dynamic foreground, reflection, and other noisy artifacts. Therefore, robust PCA algorithms such as, iEALM [35], APG [61], ReProCS [26] overcome the rank challenge robustly. However, in some cases, the target rank and the sparsity level can be part of the user-defined hyperparameters. Therefore, instead one might use a different approach as in [64, 19, 63].

For these set of experiments, we use ProGrAMMe-$\epsilon$ and compared with two different types robust PCP formulations\(^3\). Note that, the PCP formulations have a way to detect the sparse outliers but our formulation does not. We overcome this challenge by using large weights. Similar to the heuristic used in the experiments for synthetic data, we choose a random subset of entries from the set $[m] \times [n]$ and use a large range of weight at those elements. This is similar to the idea used in [19, 63, 18], as for specific sparsity percentage $\alpha$, the operator $T_\alpha[S]$ performs an approximate projection onto the sparse support of $S$. We argue that randomly selecting $\alpha\%$ of elements from $[m] \times [n]$ and hitting them by large weights, we obtain the same artifact. Indeed our empirical evidence in Figure 7 justifies that.

In our experiments, we use eight different video sequences: (i) the Basic sequence from Stuttgart synthetic dataset [9], (ii) four video sequences from CDNet2014 datasets [57], and (iii) three video sequences from SBI dataset [40, 1]. We extensively use the Stuttgart video sequence as it is equipped with foreground ground truth for each frame. For iEALM and APG, we set $\lambda = 1/\sqrt{m}$ and use $\mu = 1.25/\|F\|_2$ and $\rho = 1.5$, where $\|F\|_2$ is the spectral norm (maximum singular value) of $F$. For Best pair RPCA, RPCA GD, NCF, GoDec, and our ProGrAMMe-$\epsilon$, we set $r = 2$, target sparsity 10% and additionally, for GoDec, we set $q = 2$. For GRATSA, we set the parameters the same as those mentioned in the authors’ website\(^4\). The qualitative analysis on the background and foreground recovered suggest that our method recovers a visually similar or better quality background and foreground compare to the other methods. Note that, RPCA GD and ReProCS recover a

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\(^3\)See [59] for an overview.

\(^4\)https://sites.google.com/site/hejunzz/grasta
fragmentary foreground with more false positives; moreover, GRASTA, iEALM, and APG cannot remove the static foreground object. See Section A.4 in Appendix for more qualitative results (Figure 13) and detailed quantitative results (Figure 14) of MSSIM and PSNR.

**Background estimation from partially observed/missing data** We randomly select the set of observable entries in the data matrix $F$ and perform our experiments on Stuttgart Basic video. For these set of experiments, we use ProGrAMMe-1. As this is a missing data case, for ProGrAMMe-1, we use a binary mask as the weight. Figure 8 shows the qualitative results on different subsampled video. We provide a detailed quantitative evaluation of our ProGrAMMe with respect to the $\varepsilon$-proximity metric $d_\varepsilon(X, Y)$ as in [19] in recovering the foreground objects and show the execution time for different missing data cases in Figure 15 in Section A.5 of Appendix.

Figure 8: Sample video frames from the Stuttgart Basic video sequence for the missing data case. From top to bottom we use $\Omega = 0.8, 0.7, \text{and } 0.6$, respectively.

## 5 Conclusions

In this paper, we proposed a generic weighted low-rank recovery model and designed an SVD-free fast algorithm for solving the model. Our model covers several existing low-rank approaches in the literature as special cases and can easily be extended to the non-convex setting. Our proposed algorithm combines proximal gradient descent method and the variational formulation of nuclear norm, which does not require to compute the SVD in each step. This makes the algorithm highly scalable to larger data and enjoys a lower per iteration complexity than those who require SVD. Moreover, based on a rank identification property, we designed a rank continuation scheme which asymptotically achieves the minimal per iteration complexity. Numerical experiments on various problems and settings were performed, from which we observe superior performance of our proposed algorithm compared to a vast class of weighted and unweighted low-rank algorithms.
In this section, we added some extra numerical experiments that complement our experiments and other claims in the main paper.

### A.1 Table of baseline methods
In Table 3 we summarize all the methods compared in this paper.

### A.2 Convergence behavior
In this section, we demonstrate the convergence of our algorithm(s). For this purpose, we generate the low-rank matrix, $L$, as a product of two independent full-rank matrices of size $m \times r$ with $r < m$ such that elements are independent and identically distributed (i.i.d.) and sampled from a normal distribution—$\mathcal{N}(0, 1)$. We add Gaussian noise and generate $E$ as a noise matrix whose elements are i.i.d. $\mathcal{N}(0, 1)$ random variables as well. We constructed $F$ as: $A = L + E$. We fixed $\min_{i,j} W_{ij} = 1$ but choose $\max_{i,j} W_{ij}$ from a set $\Lambda = [5, 10, 50, 100, 500, 1000, 5000, 10^4, 5 \times 10^4, 10^5, 5 \times 10^6, 10^7]$. At each instance, $i$, we create an $m \times m$ weight matrix $W$ by fusing $\min_{i,j} W_{ij}$ with $\Lambda_i$ by using MATLAB function `randperm`.

We compare our ProGrAMMe-ε (Algorithm 1), its inexact counterpart ProGrAMMe-1, proximal gradient (PG) algorithm and its accelerated version—accelerated proximal gradient (APG) for these experiments. For different condition number ($\kappa_W$) of the weight matrix $W$, we plotted the functional value $\Phi(X_k)$ versus iterations in Figure 9 and difference between consecutive iterates, $\|X_{k+1} - X_k\|$, versus iterations in Figure 10. Note that by construction, $\kappa_W$ ranges between 1238.021 to 21043.1574. The convergence plots justify our claims, that, although problem (5) belongs to the class of problems (8), the general algorithms used in [30, 51, 55, 37] fail to provide good convergence results when $\kappa_W$ is large; that, our approaches are faster compare to those general approaches; and, that the performance of both exact and inexact ProGrAMMe are the same.

### A.3 Matrix completion with missing data
In this part, we conduct extensive tests of the power-grid missing data problem. For first set of experiments, we ran the methods until the relative error, $\|X_{k+1} - X_k\|/\|X_k\|$ or consecutive iteration error, $\|X_{k+1} - X_k\|$ is less than the machine precision or a maximum number of iterations (500) is reached, whichever happens
Figure 9: Convergence in terms of functional value of proximal gradient applied to original problem (5) (Direct PG), accelerated proximal gradient-II (APG-II), ProGrAMMe-1, and ProGrAMMe-ε. For all problems $\tau = 10^{-2}/\lambda_{\max}^2$.

first. Eventually, Figure 11 shows that the performance of ProGrAMMe-1 improves as it runs for more global iterations although it has a fractions of the execution time compare to the other methods. Next, in Figure 12 we ran more samples (50) with the same stopping criteria as before but with an increased number of iterations (1000). From Figure 12 we find that for fidelity inside the sample, ProGrAMMe-1 performs better that the two previously best performing methods—EM and BIRSVD. However, for fidelity outside the sample BIRSVD is still the best, but the behavior of ProGrAMMe improves compared to the previous cases.

A.4 Background estimation

We show the qualitative results on 7 video sequences from the CDNet 2014 and SBI datasets in Figure 13. In almost all sequences, our ProGrAMMe-ε performs consistently well compare to the other state-of-the-art methods. We do not include iEALM or APG due to their higher execution time.

In Figure 14, we use two robust quantitative measures for the background estimation experiments on Stuttgart Basic video: peak signal to noise ratio (PSNR) and mean structural similarities index measure (SSIM) [58]. PSNR is defined as $10\log_{10}$ of the ratio of the peak signal energy to the mean square error (MSE) between the processed video signal and the ground truth. Let $F(;i) - \hat{X}(;i)$ be the reconstructed vectorized foreground frame and $G(;i)$ be the corresponding ground truth frame, then PSNR is defined as $10\log_{10} \frac{M^2}{\text{MSE}}$, where $\text{MSE} = \frac{1}{mn} \|F(;i) - \hat{X}(;i) - G(;i)\|^2$ and $M_I = 255$ is the maximum possible pixel value of the image, as the pixels are represented using 8 bits per sample. For a reconstructed image with 8 bits bit depth, the PSNR are between 30 and 50 dB, where the higher is the better as we minimize the MSE between images with respect the maximum signal value of the image.

For both measures, we perceive the information how the high-intensity regions of the image are coming through the noise, and we pay much less attention to the low-intensity regions. We remove the noisy components from the recovered foreground, $F(;i) - \hat{X}(;i)$, by using the threshold $\epsilon'$, such that we set the components
Figure 10: Convergence in terms of difference between consecutive iterates ($\|X_{k+1} - X_k\|$) of proximal gradient applied to original problem (5) (Direct PG), accelerated proximal gradient-II (APG-II), ProGrAMMe-$\varepsilon$ (Algorithm 1), and ProGrAMMe-I. For all problems $\tau = 10^{-2}/\lambda_{\text{max}}$.

Figure 11: Fidelity within and out of the data, ran for more number of iterations (500). Here $\Omega$ denotes the percentage of missing data. The last bar diagram shows the execution time of different algorithms for different subsample $\Omega$. Although BLF with $\ell_2$ loss has the least execution time, its performance is not so good for this set of experiments.

below $\varepsilon'$ in $E$ to 0. In our experiments, we set $\varepsilon' = 10^{-4}$. In order to calculate the SSIM of each recovered foreground video frame, we consider an $11 \times 11$ Gaussian window with standard deviation ($\sigma$) 1.5 and consider the corresponding ground truth as the reference image. Among the methods tested, ProGrAMMe-$\varepsilon$ has the highest average SSIM (or MSSIM). To compare PSNR of recovered foreground frames, we use ProGrAMMe-$\varepsilon$, GRASTA [28], recursive projected compressive sensing (ReProCS)[26], inexact ALM (iEALM) [35], and RPCA GD [63]. iEALM has the highest average PSNR, 30.05 dB among all the methods, whereas our ProGrAMMe-$\varepsilon$ has average PSNR 29.45 dB. However, ProGrAMMe-$\varepsilon$ needs an average 9.48 seconds to produce the results, compare to the average execution time of iEALM is 183 seconds.
Figure 12: Fidelity within and out of the sample, ran for more samples (50) over a higher number of iterations (1000). Here $\Omega$ denotes the percentage of missing data.

Figure 13: Sample video frames from the CDNet 2014 and SBI datasets. ProGrAMMe provides a visually high quality background for almost all sequences. The red bounding boxes in recovered BG denote shadows, ghosting effects of FG objects, static FG object etc.

A.5 Background estimation from partially observed/missing data

For background estimation on partially observed data we used the $\varepsilon$-proximity metric—$d_\varepsilon(X,Y)$ proposed in [19] on Stuttgart Basic video. The performance of RPCA feasibility [19] with respect to $d_\varepsilon(X,Y)$ stays stable for all subsample $\Omega$. The performance of the best pair RPCA is also stable except for $\Omega = 0.3$. The performance of RPCA GD [63] keeps downgrading as we decrease $\Omega$. Surprisingly, the performance of ProGrAMMe-1 gets
better for this typical experiment as we decrease $\Omega$. Furthermore, the average execution time for ProGrAMMe-1 is stable for different $\Omega$, and is around 8 seconds. While the next best average execution time 19.67 seconds is recorded for RPCA GD. The average execution time of NCF and best pair are 44 and 43 seconds, respectively.

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Figure 15: Quantitative comparison between different algorithms on Stuttgart Basic sequence for different levels of partially observed/missing data case with respect to the $d_\epsilon(X,Y)$ metric. Figure (h) the bar diagram shows the execution time of different algorithms for different subsample $\Omega$. ProGrAMMe-1 has the least average execution time in all scenarios.

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