Stochastic Iterative Hard Thresholding for Graph-structured Sparsity Optimization

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Abstract

Stochastic optimization algorithms update models with cheap per-iteration costs sequentially, which makes them amenable for large-scale data analysis. Such algorithms have been widely studied for structured sparse models where the sparsity information is very specific, e.g., convex sparsityinducing norms or ℓ^0 -norm. However, these norms cannot be directly applied to the problem of complex (non-convex) graph-structured sparsity models, which have important application in disease outbreak and social networks, etc. In this paper, we propose a stochastic gradient-based method for solving graph-structured sparsity constraint problems, not restricted to the least square loss. We prove that our algorithm enjoys a linear convergence up to a constant error, which is competitive with the counterparts in the batch learning setting. We conduct extensive experiments to show the efficiency and effectiveness of the proposed algorithms.

1. Introduction

Structured sparse learning models have received increasing attention. They can be formulated as follows

$$\min_{\boldsymbol{x} \in \mathcal{M}} F(\boldsymbol{x}), \quad F(\boldsymbol{x}) := \frac{1}{n} \sum_{i=1}^{n} f_i(\boldsymbol{x}). \tag{1}$$

Here, each $f_i(\boldsymbol{x})$ is convex and differentiable and the structured sparsity is reflected by the constraint set $\mathcal{M} \subseteq \mathbb{R}^p$ on \boldsymbol{x} . Typically, one can encode the sparsity by introducing sparsity-inducing penalties such as ℓ^1 -norm (Tibshirani, 1996; Chen et al., 2001), ℓ^1/ℓ^q mixed norms (Turlach et al.,

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2005; Yuan & Lin, 2006) and more structured norms built on either disjoint or overlapping groups of variables (Bach et al., 2012b; Jenatton et al., 2011; Obozinski & Bach, 2016; Morales et al., 2010). Such models of sparsity-inducing penalties are often convex and can be solved by convex optimization algorithms (Bach et al., 2012a).

There is a wide range of significant application, including the search of connected subgraphs in networks, where the constraint set \mathcal{M} cannot be encoded by sparsity-inducing norms. Notable application examples include disease outbreak as a connected cluster (Arias-Castro et al., 2011; Qian et al., 2014; Aksoylar et al., 2017), and social events as connected dense subgraphs (Rozenshtein et al., 2014). To capture these complex graph structures such as trees and connected graphs, recent works by Baraniuk et al. (2010) and Hegde et al. (2014; 2016; 2015b) have proposed to use structured sparsity model M to define allowed supports $\{S_1, S_2, \dots, S_k\}$ where \mathcal{M} can be represented by $\mathcal{M}(\mathbb{M}) = \{x : \operatorname{supp}(x) \subseteq S_i, \text{ for some } S_i \in \mathbb{M}\}.$ These complex models are non-convex and (batch) gradient descent algorithms involve a projection operator $P(\cdot, \mathcal{M}(\mathbb{M}))$ which is usually NP-hard. In a series of seminal work, Hegde et al. (2015b; 2016; 2015a) used two approximated projections (head and tail) without sacrificing too much precision. However, their work only focused on the least square loss, and computing full gradient per iteration resulted in expensive per-iteration cost $\mathcal{O}(mp)$ if m is the number of examples and p is the data dimension. This largely limits its practical application to big data setting where typically the data is contaminated by non-Gaussian noise and the data volume is huge (i.e., m or/and p is very large). Therefore, it is desirable to develop efficient optimization algorithms for graph-structured sparsity constrained problems scalable to large-scale data.

Stochastic optimization such as stochastic gradient descent (SGD) has become a standard tool for solving large-scale convex and non-convex learning problems (e.g., Bach & Moulines (2013); Bousquet & Bottou (2008); Jin et al. (2017); Kingma & Ba (2014); Li & Orabona (2018); Rakhlin et al. (2012); Shamir & Zhang (2013); Yang & Lin (2015); Ying & Zhou (2017; 2006)). SGD-type algorithms have a cheap per-iteration computation of the gradient over one or

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a few training examples, which makes them amenable for large-scale data analysis. Along this research line, SGD-type algorithms for sparse models have been proposed including stochastic proximal gradient methods (Rosasco et al., 2014; Duchi et al., 2011) and stochastic gradient hard thresholding (Nguyen et al., 2017; Zhou et al., 2018a; Murata & Suzuki, 2018; Shen & Li, 2017; Liu et al., 2017). However, these algorithms cannot address the important case of complex (non-convex) graph-structured sparsity constraint problems.

In this paper we leverage the recent success of SGD algorithms for non-convex problems and propose an efficient stochastic optimization algorithm for graph-structured sparsity constraint problems. In particular, we use the structure sparsity model $\mathcal{M}(\mathbb{M})$ to capture more complex graph-structured information of interest. Our main contributions can be summarized as follows:

- We propose a stochastic gradient-based algorithm for solving graph-structured sparsity constraint problems. To our best knowledge, our work is the first attempt to provide stochastic gradient descent-based algorithm for graph-structured sparsity constraint problems.
- The proposed algorithm enjoys linear convergence property under proper conditions.¹ It is proved applicable to a broad spectrum of loss functions. Specifically, for the least square loss, it enjoys a linear convergence rate which is competitive to the algorithm proposed in Hegde et al. (2016). For the logistic loss, we show that it also satisfies the linear convergence property with high probability. Our proofs could be easily applied to other types of projections.
- We conduct extensive experiments to validate the proposed stochastic algorithm. In particular, we focus on two applications: graph sparse linear regression and graph logistic regression on breast cancer data. Our experiment results show that the proposed stochastic algorithm consistently outperforms the deterministic ones.

Outline. We organize the rest of the paper as follows. In Section 2, we introduce the notations, definitions and key assumptions. Our proposed algorithm is presented in Section 3. Section 4 provides the theoretical analysis of the algorithm. Numerical experiments and discussion are respectively in Section 5 and 6. Due to the limited space, all of our formal proofs and further experiment results are provided in the Supplementary Material. The source code and datasets are accessible at: https://github.com/baojianzhou/graph-sto-iht.

2. Preliminary

Notation. We use boldface letters, e.g., X to denote matrices. The boldface lower-case letters, e.g., x, y are column vectors. The upper-case letters, e.g., H, T, S stand for subsets of $[p] := \{1, 2, \ldots, p\}$. $X_S \in \mathbb{R}^{|S| \times p}$ is the submatrix by keeping rows only in set S. Given the standard basis $\{e_i : 1 \le i \le p\}$ of \mathbb{R}^p , we also use H, T, S to represent subspaces. For example, the subspace induced by H is the subspace defined as $\text{span}\{e_i : i \in H\}$. We will clarify the difference only if confusion might occur. The restriction of x on S is the vector x_S such that i-th entry $(x_S)_i = x_i$ if $i \in S$; otherwise 0. We denote the support of x as $\text{supp}(x) := \{i : x_i \ne 0\}$. We use $\|x\|$ to denote ℓ^2 norm of x. The set complement is defined as $S^c = [p] \setminus S$.

Definition 1 (Subspace model (Hegde et al., 2016)). Given the space \mathbb{R}^p , a subspace model \mathbb{M} is defined as a family of linear subspaces of \mathbb{R}^p : $\mathbb{M} = \{S_1, S_2, \dots, S_k, \dots, \}$, where each S_k is a subspace of \mathbb{R}^p . The set of corresponding vectors in these subspaces is denoted as

$$\mathcal{M}(\mathbb{M}) = \{ x : x \in V, \text{ for some } V \in \mathbb{M} \}. \tag{2}$$

Equation (2) is general enough to include many sparse models. For example, the s-sparse set $\mathcal{M}(\mathbb{M}_s) = \{x : |\sup(x)| \leq s\}$ (Nguyen et al., 2017; Yuan et al., 2018; Jain et al., 2014; Bahmani et al., 2013), where \mathbb{M}_s contains subspaces spanned by s different standard basis vectors, i.e., $\mathbb{M}_s = \{\operatorname{span}\{e_{i_1}, e_{i_2}, \dots, e_{i_s}\} : \{i_1, i_2, \dots, i_s\} \subseteq [p]\}$. Although the proposed algorithm and theorems apply to any \mathbb{M} which admits efficient projections, in this paper, we mainly consider graph-structured sparsity constraint models such as the weighted graph model in Hegde et al. (2015b).

Definition 2 (Weighted Graph Model (Hegde et al., 2015b)). Given an underlying graph $\mathbb{G} = (\mathbb{V}, \mathbb{E})$ defined on the coefficients of the unknown vector \mathbf{x} , where $\mathbb{V} = [p]$ and $\mathbb{E} \subseteq \mathbb{V} \times \mathbb{V}$, then the weighted graph model (\mathbb{G}, s, g, C) -WGM can be defined as the following set of supports

$$\mathbb{M} = \{S : |S| \le s, \text{ there is an } F \subseteq \mathbb{V} \text{ with } \mathbb{V}_F = S, \gamma(F) = g, \text{ and } w(F) \le C\},$$

where C is the budget on weight of edges w, g is the number of connected components of F, and s is the sparsity.

 (\mathbb{G}, s, g, C) -WGM captures a broad range of graph structures such as groups, clusters, trees, and connected subgraphs. This flexibility requires an efficient projection operator onto $\mathcal{M}(\mathbb{M})$, i.e., $P(\cdot, \mathcal{M}(\mathbb{M})) : \mathbb{R}^p \to \mathbb{R}^p$ defined as

$$P(\boldsymbol{x}, \mathcal{M}(\mathbb{M})) = \underset{\boldsymbol{y} \in \mathcal{M}(\mathbb{M})}{\arg \min} \|\boldsymbol{x} - \boldsymbol{y}\|^{2}.$$
 (3)

The above operator is crucial for projected gradient descentbased methods, but to solve the operator is NP-hard in gen-

¹Linear convergence up to a tolerance error.

eral. To explore (3), one needs to solve the following equivalent minimization problem

$$\min_{\boldsymbol{y} \in \mathcal{M}(\mathbb{M})} \|\boldsymbol{x} - \boldsymbol{y}\|^2 \Leftrightarrow \min_{S \in \mathbb{M}} \|\boldsymbol{x} - P(\boldsymbol{x}, S)\|^2.$$

Since P(x, S) is an orthogonal projection operator that projects x onto subspace S, by projection theorem, it always has the following property

$$\|\boldsymbol{x}\|^2 - \|P(\boldsymbol{x}, S)\|^2 = \|\boldsymbol{x} - P(\boldsymbol{x}, S)\|^2.$$

By adding minimization to both sides with respect to subspace S, we obtain

$$\min_{S \in \mathbb{M}} \left\{ \|\boldsymbol{x}\|^2 - \|P(\boldsymbol{x}, S)\|^2 \right\} = \min_{S \in \mathbb{M}} \|\boldsymbol{x} - P(\boldsymbol{x}, S)\|^2
\|\boldsymbol{x}\|^2 + \max_{S \in \mathbb{M}} \|P(\boldsymbol{x}, S)\|^2 = \min_{S \in \mathbb{M}} \|\boldsymbol{x} - P(\boldsymbol{x}, S)\|^2.$$
(4)

The above observations lead to a key insight that the NPhard problem (3) can be solved either by maximizing $\|P(\boldsymbol{x},S)\|^2$ or by minimizing $\|\mathbf{x}-P(\boldsymbol{x},S)\|^2$ over S. Instead of minimizing or maximizing these two squared norms over S exactly, Hegde et al. (2015a;b) turned to approximated algorithms, and gave two important projectionshead projection and tail projection.

Assumption 1 (Head Projection (Hegde et al., 2016)). *Let* \mathbb{M} and $\mathbb{M}_{\mathcal{H}}$ be the predefined subspace models. Given any vector \mathbf{x} , there exists a $(c_{\mathcal{H}}, \mathbb{M}, \mathbb{M}_{\mathcal{H}})$ -Head-Projection which is to find a subspace $H \in \mathbb{M}_{\mathcal{H}}$ such that

$$\|\mathbf{P}(\boldsymbol{x}, H)\|^2 \ge c_{\mathcal{H}} \cdot \max_{S \in \mathbb{M}} \|\mathbf{P}(\boldsymbol{x}, S)\|^2, \tag{5}$$

where $0 < c_{\mathcal{H}} \leq 1$. We denote $P(\mathbf{x}, H)$ as $P(\mathbf{x}, M, M_{\mathcal{H}})$.

Assumption 2 (Tail Projection (Hegde et al., 2016)). Let \mathbb{M} and $\mathbb{M}_{\mathcal{T}}$ be the predefined subspace models. Given any vector x, there exists a $(c_T, \mathbb{M}, \mathbb{M}_T)$ -Tail-Projection which is to find a subspace $T \in \mathbb{M}_{\mathcal{T}}$ such that

$$\|P(\boldsymbol{x},T) - \boldsymbol{x}\|^2 \le c_{\mathcal{T}} \cdot \min_{S \in \mathbb{M}} \|\boldsymbol{x} - P(\boldsymbol{x},S)\|^2, \quad (6)$$

where $c_{\mathcal{T}} \geq 1$. We denote $P(\boldsymbol{x}, T)$ as $P(\boldsymbol{x}, \mathbb{M}, \mathbb{M}_{\mathcal{T}})$.

Intuitively, head projection keeps large magnitudes while tail projection discards small magnitudes. Take \mathbb{M}_s (a complete graph) as an example. Head projection is to find subspace $H = \text{span}\{e_{i_1}, e_{i_2}, \dots, e_{i_s}\}$ where $\{i_1, i_2, \ldots, i_s\} \subseteq [p]$ is the set of indices that keeps the largest s components of x, i.e, $|x_{i_1}| \ge |x_{i_2}| \ge \cdots \ge |x_{i_s}|$. As (3) could have multiple solutions, we could have $H \neq T$, but we must have $\|x - x_H\| = \|x - x_T\|$.

Definition 3 ($(\alpha, \beta, \mathcal{M}(\mathbb{M}))$ -RSC/RSS). We say a differentiable function $f(\cdot)$ satisfies the $(\alpha, \beta, \mathcal{M}(\mathbb{M}))$ -Restricted Strong Convexity (RSC)/Smoothness (RSS) property if there exist positive constants α and β such that

$$\frac{\alpha}{2} \|\boldsymbol{x} - \boldsymbol{y}\|^2 \le B_f(\boldsymbol{x}, \boldsymbol{y}) \le \frac{\beta}{2} \|\boldsymbol{x} - \boldsymbol{y}\|^2, \tag{7}$$

for all $x, y \in \mathcal{M}(\mathbb{M})$, where $B_f(x, y)$ is the Bregman divergence of f, i.e., $B_f(\mathbf{x}, \mathbf{y}) = f(\mathbf{x}) - f(\mathbf{y}) - \langle \nabla f(\mathbf{y}), \mathbf{x} - \mathbf{y} \rangle$ y). α and β are strong convexity parameter and strong smoothness parameter respectively.

RSC/RSS property was firstly introduced in Agarwal et al. (2012)². Since it captures sparsity of many functions, it has been widely used (e.g., Jain et al. (2014); Yuan et al. (2018); Elenberg et al. (2018); Johnson & Zhang (2013); Shen & Li (2017); Zhou et al. (2018a); Nguyen et al. (2017); Zhou et al. (2018b)). For example, if f is the least square loss, i.e., $f(x) = ||Ax - y||^2$, then RSC/RSS property can be reduced to the canonical subspace Restricted Isometry Property (RIP) (Candes & Tao, 2005). Next, we characterize the property of F and each f_i of (1) in Assumption 3.

Assumption 3. Given the objective function F(x) in (1), we assume that F(x) satisfies α -RSC in subspace model $\mathcal{M}(\mathbb{M} \oplus \mathbb{M}_{\mathcal{H}} \oplus \mathbb{M}_{\mathcal{T}})$. Each function $f_i(\boldsymbol{x})$ satisfies β -RSS in $\mathcal{M}(\mathbb{M} \oplus \mathbb{M}_{\mathcal{H}} \oplus \mathbb{M}_{\mathcal{T}})$, where \oplus of two models \mathbb{M}_1 and \mathbb{M}_2 is defined as $M_1 \oplus M_2 := \{S_1 \cup S_2 : S_1 \in M_1, S_2 \in M_2\}.$

3. Algorithm: GRAPHSTOIHT

Algorithm 1 GRAPHSTOIHT

- 1: **Input**: $\eta_t, F(\cdot), \mathbb{M}, \mathbb{M}_{\mathcal{H}}, \mathbb{M}_{\mathcal{T}}$
- 2: **Initialize**: x^0 such that supp $(x) \in \mathbb{M}$ and t = 0
- 3: **for** $t = 0, 1, 2, \dots$ **do**
- Choose ξ_t from [n] with probability $Pr(\xi_t)$
- $\mathbf{b}^{t} = P(\nabla f_{\xi_{t}}(\mathbf{x}^{t}), \mathbb{M} \oplus \mathbb{M}_{\mathcal{T}}, \mathbb{M}_{\mathcal{H}})$ $\mathbf{x}^{t+1} = P(\mathbf{x}^{t} \eta_{t}\mathbf{b}^{t}, \mathbb{M}, \mathbb{M}_{\mathcal{T}})$
- 7: end for
- 8: Return \boldsymbol{x}^{t+1}

The proposed algorithm is named GRAPHSTOIHT presented in Algorithm 1, a stochastic-based method for graphstructured sparsity problems. Initially, $x^0 = 0$ and t = 0. At each iteration, it works as the following three steps:

- Step 1: in Line 4, it randomly selects ξ_t from [n] with probability mass function $Pr(\xi_t)$, that is, the current loss $f_{\xi_t}(\cdot)$ has been chosen with probability $Pr(\xi_t)$;
- Step 2: in Line 5, the head projection inputs the gradient of $f_{\mathcal{E}_t}(\boldsymbol{x})$ and outputs the projected vector \boldsymbol{b}^t so that $\operatorname{supp}(\boldsymbol{b}^t) \in \mathcal{M}(\mathbb{M}_{\mathcal{H}});$
- Step 3: in Line 6, the next estimated x^{t+1} is then updated by using the tail projection.

The algorithm repeats the above three steps until stop condition is satisfied. The difference between GRAPHSTOIHT and STOIHT (Nguyen et al., 2017), the latter of which is

²This property is related with the theory of paraconvexity (Van Ngai & Penot, 2008) as pointed out by Agarwal et al. (2012).

essentially a stochastic projected gradient descent, is twofold:

- 1) instead of directly using the gradient $\nabla f_{\xi_t}(\cdot)$ at each iteration as STOIHT did, GRAPHSTOIHT uses a thresholded gradient. Thresholding $\nabla f_{\xi_t}(\cdot)$ at the first stage could be helpful under sparse setting, because most nonzero entries of $\nabla f_{\xi_t}(\cdot)$ are irrelevant.
- 2) instead of projecting on $\mathcal{M}(\mathbb{M}_s)$ as STOIHT did, the tail projection projects each estimation onto graph-structured subspaces $\mathcal{M}(\mathbb{M}_T)$.

We call a batch version (n=1) of this algorithm as GRAPHIHT, where we simply apply the head and tail projection to AS-IHT in Hegde et al. (2016) by considering a general loss function.

Time complexity Analysis. Due to the NP-hardness of the projection, there is a trade-off between the time cost and the quality of projection. The time complexity of two projections depends on the graph size p and the number of edges $|\mathbb{E}|$. As proved in Hegde et al. (2015b), the running time of both head projection and tail projection is bounded by $\mathcal{O}(|\mathbb{E}|\log^3(p))$. If the graph is sparse as common in real-world applications, i.e., $|\mathbb{E}| = \mathcal{O}(p)$, two projections have nearly-linear time complexity: $\mathcal{O}(p\log^3(p))$ with respect to the feature dimension p. Hence, per-iteration cost $\mathcal{O}(p\log^3(p))$, is still cheaper than that of the deterministic GD algorithms GRAPHIHT in which the computation of the full gradient will be of cost $\mathcal{O}(np)$.

4. Convergence Analysis of GRAPHSTOIHT

In this section, firstly we give the convergence analysis of GRAPHSTOIHT by characterizing the *estimation error* between x^{t+1} and x^* , i.e, $||x^{t+1} - x^*||$, where x^* is an optimal solution of (1). Then we analyze two commonly used objective functions and prove our algorithm achieves a linear convergence up to a constant error. Our analysis is applicable not only to graph-structured sparsity model but also to any other head and tail projection.

We denote $\xi_{[t]}=(\xi_0,\xi_1,\ldots,\xi_t)$ as the history of stochastic process ξ_0,ξ_1,\ldots up to time t, and all random variables ξ_t are independent of each other. Define the *inverse condition number* $\mu:=\alpha/\beta$. To simplify the analysis, we define the probability mass function $Pr(\xi_t=i)=1/n, 1\leq i\leq n$. Before presenting our main Theorem 1, let's take a look at the following key lemma.

Lemma 1. If each $f_{\xi_t}(\cdot)$ and $F(\mathbf{x})$ satisfy Assumption 3 and, given head projection model $(c_{\mathcal{H}}, \mathbb{M} \oplus \mathbb{M}_{\mathcal{T}}, \mathbb{M}_{\mathcal{H}})$ and tail projection model $(c_{\mathcal{T}}, \mathbb{M}, \mathbb{M}_{\mathcal{T}})$, then we have the following inequality

$$\mathbb{E}_{\xi_t} \| (\boldsymbol{x}^t - \boldsymbol{x}^*)_{H^c} \| \le \sqrt{1 - \alpha_0^2} \mathbb{E}_{\xi_t} \| \boldsymbol{x}^t - \boldsymbol{x}^* \| + \sigma_1, (8)$$

where

$$\sigma_{1} = \left(\frac{\beta_{0}}{\alpha_{0}} + \frac{\alpha_{0}\beta_{0}}{\sqrt{1 - \alpha_{0}^{2}}}\right) \mathbb{E}_{\xi_{t}} \|\nabla_{I} f_{\xi_{t}}(\boldsymbol{x}^{*})\|,$$

$$H = supp(P(\nabla f_{\xi_{t}}(\boldsymbol{x}^{t}), \mathbb{M} \oplus \mathbb{M}_{\mathcal{T}}, \mathbb{M}_{\mathcal{H}})),$$

$$\alpha_{0} = c_{\mathcal{H}}\alpha\tau - \sqrt{\alpha\beta\tau^{2} - 2\alpha\tau + 1}, \quad \beta_{0} = (1 + c_{\mathcal{H}})\tau,$$

$$I = \underset{S \in \mathbb{M} \oplus \mathbb{M}_{\mathcal{T}} \oplus \mathbb{M}_{\mathcal{H}}}{\arg \max} \mathbb{E}_{\xi_{t}} \|\nabla_{S} f_{\xi_{t}}(\boldsymbol{x}^{*})\|, \text{ and } \tau \in (0, 2/\beta).$$

Inequality (8) means that the head projection always excludes a small fraction of the residual vector $oldsymbol{x}^t - oldsymbol{x}^*$ at each iteration. That is to say, most of the large magnitudes in $x^t - x^*$ are captured by the head projection. More specifically, the head projection thresholds small magnitude entries of $\nabla f_{\mathcal{E}_t}(\boldsymbol{x}^t)$ to zeros. These small magnitude entries could lead the algorithm to a wrong direction. When F(x) is the least square loss, Lemma 1 here is similar to Lemma 13 in Hegde et al. (2016). However, there are two important differences: 1) our Lemma 1 can be applied to any functions that satisfy Assumption 3 above, while the RIP condition used in Hegde et al. (2016) can be only applied to the least square loss; 2) since each $f_{\xi_t}(x)$ is not strongly convex, the proof in Hegde et al. (2016) cannot be directly used. Instead, we use co-coercivity in Nguyen et al. (2017) and then obtain the main theorem below.

Theorem 1. Let x^0 be the start point of Algorithm 1. If we choose a constant learning rate with $\eta_t = \eta$ and use Lemma 1, then the solution x^{t+1} of Algorithm 1 satisfies

$$\mathbb{E}_{\xi_{[t]}} \| \boldsymbol{x}^{t+1} - \boldsymbol{x}^* \| \le \kappa^{t+1} \| \boldsymbol{x}^0 - \boldsymbol{x}^* \| + \frac{\sigma_2}{1 - \kappa}, \quad (9)$$

where

where
$$\kappa = (1 + c_{\mathcal{T}}) \Big(\sqrt{\alpha \beta \eta^2 - 2\alpha \eta + 1} + \sqrt{1 - \alpha_0^2} \Big),$$

$$\sigma_2 = \sigma_1 + \eta \mathbb{E}_{\mathcal{E}_{\epsilon}} \|\nabla_I f_{\mathcal{E}_{\epsilon}}(\boldsymbol{x}^*)\|, \text{ and } \eta, \tau \in (0, 2/\beta).$$

Theorem 1 shows that Algorithm 1 still possibly enjoys linear convergence even if the full gradient is not available. In order to make more detailed analysis of Theorem 1, we call κ in (9) the *contraction factor*. A natural question to ask is: under what condition can we get an effective contraction factor, κ , i.e., $\kappa<1$. When $\kappa<1$, the estimation error is dominated by the term $\frac{\sigma_2}{1-\kappa}$. Suppose the random distribution of ξ_t is uniform and the approximation factor of the head projection $c_{\mathcal{H}}$ can be arbitrarily boosted close to 1. Taking $\eta=1/\beta$, in order to get an effective contraction factor, $\kappa<1$, the inverse of the condition number μ and tail projection factor $c_{\mathcal{T}}$ need to satisfy

$$\kappa = (1 + c_{\mathcal{T}})(1 + 2\sqrt{\mu})\sqrt{1 - \mu} < 1.$$
(10)

To be more specific, if κ is a function of η , then it takes minimum at $\eta = 1/\beta$. By letting $\eta = 1/\beta$, then κ is simplified as $(1 + c_T)(\sqrt{1 - \mu} + \sqrt{1 - \alpha_0^2})$. Furthermore,

 α_0 is a concave function with respect to τ (recall that τ is a free parameter in $(0,2/\beta)$), and then we have its maximum at $\tau=\frac{1}{\beta}(1+c_{\mathcal{H}}\sqrt{\frac{\beta-\alpha}{\beta-c_{\mathcal{H}}^2\alpha}})$ after calculation. By taking $c_{\mathcal{H}}\to 1^-$, we present κ as $(1+c_{\mathcal{T}})(1+2\sqrt{\mu})\sqrt{1-\mu}$

One of the assumptions of (10) is that $c_{\mathcal{H}}$ can be arbitrarily close to 1. One can boost the head projection by using the method proposed in Hegde et al. (2016). In our experiment, we find that it is not necessary to boost the head projection because executing the head projection once is sufficient enough to obtain good performance. In the remainder of the section, we consider two popular loss functions to discuss the conditions of getting effective contraction factors.

Graph sparse linear regression. Given a design matrix $A \in \mathbb{R}^{m \times p}$ and corresponding observed noisy vector $y \in \mathbb{R}^m$ that are linked via the linear relationship

$$y = Ax^* + \epsilon, \tag{11}$$

where $\epsilon \sim \mathcal{N}(\mathbf{0}, \sigma^2 \mathbf{I})$. The graph sparse linear regression is to estimate the underlying sparse vector \mathbf{x}^* . The underlying graph $\mathbb{G}(\mathbb{V}, \mathbb{E})$ is defined on \mathbf{x}^* . For example, $\operatorname{supp}(\mathbf{x}^*)$ induces a connected subgraph with s nodes, showcased in Figure 1 in the experiment section. To estimate \mathbf{x}^* , naturally we consider the least square loss and formulate it as

$$\underset{\operatorname{supp}(\boldsymbol{x}) \in \mathcal{M}(\mathbb{M})}{\arg \min} F(\boldsymbol{x}) := \frac{1}{n} \sum_{i=1}^{n} \frac{n}{2m} \|\boldsymbol{A}_{B_i} \boldsymbol{x} - \boldsymbol{y}_{B_i}\|^2, \quad (12)$$

where m observations have been partitioned into n blocks, B_1, B_2, \ldots, B_n . Each block B_i is indexed by i with block size b = m/n. In Corollary 1, we show that it is possible to get a linear convergence rate under RSC/RSS assumption.

Corollary 1. If F(x) in (12) satisfies the α -RSC property and each $f_i(x) = \frac{n}{2m} \|\mathbf{A}_{B_i} x - y_{B_i}\|^2$ satisfies the β -RSS property, then we have the following condition

$$\frac{\alpha}{2} \|x\|^2 \le \frac{1}{2m} \|Ax\|^2 \le \frac{\beta}{2} \|x\|^2.$$

Let the strong convexity parameter $\alpha=1-\delta$ and strong smoothness parameter $\beta=1+\delta$, where $0<\delta<1$. The condition of effective contraction factor is as the following

$$(1+c_{\mathcal{T}})\left(\sqrt{\frac{2}{1+\delta}} + \frac{2\sqrt{2(1-\delta)}}{1+\delta}\right)\sqrt{\delta} < 1.$$

Table 1 gives a contrast of contraction factor κ in our method and in Hegde et al. (2016). Both conditions are obtained by letting $c_{\mathcal{H}} \to 1^-$. Table 1 shows that κ of the two algorithms could be effective if $\delta \to 0$. By using a sufficiently large number of observations m, we always have $\delta \to 0$. According to the preceding analysis, GRAPHIHT uses a constant learning rate $\eta = 1$ while GRAPHSTOIHT

Table 1. κ of two algorithms

Algorithm	κ
GRAPHIHT	$(1+c_{\mathcal{T}})\Big(\sqrt{\delta}+2\sqrt{1-\delta}\Big)\sqrt{\delta}$
GRAPHSTOIHT	$(1+c_{\mathcal{T}})\left(\sqrt{\frac{2}{1+\delta}}+\frac{2\sqrt{2(1-\delta)}}{1+\delta}\right)\sqrt{\delta}$

uses the learning rate $\eta=1/(1+\delta)$. However, the difference between the two learning rates disappears when $\delta\to 0$. To be more specific, κ of Graphith is controlled by $\mathcal{O}(\sqrt{\delta}\cdot 2(1+c_{\mathcal{T}}))$ while for Graphstoiht, κ is controlled by $\mathcal{O}(\sqrt{\delta}\cdot 3\sqrt{2}(1+c_{\mathcal{T}}))$. Consequently, the contraction factor κ of Graphstoiht is competitive with that of Graphith while Graphstoiht has the advantage of cheaper per-iteration cost $\mathcal{O}(mp/n)$, if the size of blocks n is large, compared to $\mathcal{O}(mp)$ of Graphith. To obtain $\kappa<1$, $\delta\leq0.0527$ for Graphith while $\delta\leq0.0142$ for Graphstoiht. The gap between the two κ is mainly due to the randomness introduced in our algorithm.

The error term σ_2 in Equation (9) mainly depends on b and σ . The gradient of f_{ξ_t} at \boldsymbol{x}^* is $\boldsymbol{A}_{B_{\xi_t}}^{\top} \boldsymbol{\epsilon}_{B_{\xi_t}}/b$. As shown in Nguyen et al. (2017), it can be bounded as $C\sqrt{\sigma^2|I|\log p/b}$, where C is a constant independent of b and σ .

Though GRAPHSTOIHT needs more observations than GRAPHIHT theoretically, our experiment results demonstrate that GRAPHSTOIHT uses less observations than GRAPHIHT to estimate \boldsymbol{x}^* accurately under the same condition. Such kind of interesting experiment phenomenon is reported in Nguyen et al. (2017) but for sparsity constraint.

Graph logistic regression. Given a dataset $\{a_i, y_i\}_{i=1}^m$ ($\{a_i\}_{i=1}^m$ is a set of i.i.d random vectors), the graph logistic regression is formulated as the following problem

$$\underset{\text{supp}(\boldsymbol{x}^*) \in \mathcal{M}(\mathbb{M})}{\arg \min} F(\boldsymbol{x}) := \frac{1}{n} \sum_{i=1}^{n} f_i(\boldsymbol{x}), \tag{13}$$

where each $f_i(x)$ is defined as

$$f_i(\boldsymbol{x}) = \frac{n}{m} \sum_{i=1}^{m/n} \left[\log(1 + \exp(-y_{ij} \boldsymbol{a}_{ij}^{\mathsf{T}} \boldsymbol{x})) \right] + \frac{\lambda}{2} \|\boldsymbol{x}\|^2.$$
 (14)

Here λ is the regularization parameter. Again we divide [m] into n blocks, i.e., B_1, B_2, \ldots, B_n and assume each block size b is the same, i.e., b = m/n. Problem (13) has many important applications. For example, in its application to breast cancer classification based on gene mirco-array dataset and gene-gene network, each a_i is the i-th sample representing the gene expression records of patient i, and y_i is associated label with value -1 or 1. $y_i = 1$ means metastatic growth on patient i while $y_i = -1$ non-metastatic growth on patient i. Assume that we have more prior information such as

gene-gene network, the goal here is to minimize the objective function F(x) meanwhile to find a connected subgraph induced by $\operatorname{supp}(x)$ which is highly relevant with breast cancer-related genes. This kind of application is theoretically based on the following corollary.

Corollary 2. Suppose we have the logistic loss, F(x) in (13) and each sample a_i is normalized, i.e., $||a_i|| = 1$. Then F(x) satisfies λ -RSC and each $f_i(x)$ in (14) satisfies $(\alpha + (1 + \nu)\theta_{max})$ -RSS. The condition of getting effective contraction factor of GRAPHSTOIHT is as the following

$$\frac{\lambda}{\lambda + n(1+\nu)\theta_{max}/4m} \ge \frac{243}{250},\tag{15}$$

with probability $1 - p \exp(-\theta_{max}\nu/4)$, where $\theta_{max} = \lambda_{max}(\sum_{j=1}^{m/n} \mathbb{E}[\boldsymbol{a}_{i_j}\boldsymbol{a}_{i_j}^{\mathsf{T}}])$ and $\nu \geq 1$. $\mathbb{E}[\boldsymbol{a}_{i_j}\boldsymbol{a}_{i_j}^{\mathsf{T}}]$ is the expectation with respect to the random variable \boldsymbol{a}_{i_j} .

Given a sufficiently large number of observations m and a suitable n, it is still possible to find a bound such that the inequality (15) is valid. In Bahmani et al. (2013), they show when $\mu > 1/(1+\frac{(1+\nu)\theta_{\rm max}}{4\lambda})$, it is sufficient for GRASP (Bahmani et al., 2013) to get a linear convergence rate. Our corollary shares the similar spirit.

5. Experiments

We conduct experiments on both synthetic and real datasets to evaluate the performance of GRAPHSTOIHT³. We consider two applications, graph sparse linear regression and breast cancer metastatic classification. All codes are written in Python and C language. All experiments are tested on 56 CPUs of Intel Xeon(R) E5-2680 with 251GB of RAM.

5.1. Graph sparse linear regression

Experimental setup. The graph sparse linear regression is to recover a graph-structured vector $x^* \in \mathbb{R}^p$ by using a Gaussian matrix $A \in \mathbb{R}^{m \times p}$, and the observation vector $oldsymbol{y}$ where $oldsymbol{y}$ is measured by $oldsymbol{y} = oldsymbol{A} oldsymbol{x}^* + oldsymbol{\epsilon}$ as defined in (11). The entries of the design matrix A are sampled from $\mathcal{N}(0,1/\sqrt{m})$ independently and nonzero entries of x^* from $\mathcal{N}(0,1)$ independently for the simulation study. ϵ is potentially a Gaussian noise vector and $\epsilon = 0$ the noiseless case. We mainly follow the experimental settings in Nguyen et al. (2017). All results are averaged on 50 trials of trimmed results by excluding the best 5% and the worst 5%. All methods terminate when $\|Ax^{t+1} - y\| \le 10^{-7}$ (corresponding to convergence) or t/n > 500 (corresponding to the maximum number of epochs allowed). For GRAPH-STOIHT, one epoch contains n iterations. We recall that A has been partitioned into n blocks with block size b. We say x^* is successfully recovered if the *estimation error* $||x^{t+1} - x^*|| < 10^{-6}$.

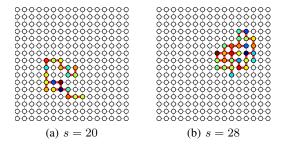


Figure 1. Two simulated graphs. Subgraphs (colored nodes and edges) are generated by $random\ walk$. Each node v_i is associated with x_i . Colored-face nodes have values sampled from $\mathcal{N}(0,1)$ while white-face nodes have values 0.0.

To simulate a graph structure on $\boldsymbol{x}^* \in \mathbb{R}^p$ that mimics realistic subgraph patterns (e.g., malicious activities) in a network (Yu et al., 2016), we fix p=256 and generate its s nonzero entries by using $random\ walk$ (Lovász et al., 1993) on a 16×16 grid graph so that $\|\boldsymbol{x}^*\|_0$ forms a connected subgraph. Each edge has unit weight 1.0. Specifically, the procedure of $random\ walk$ follows three main steps: 1) select an initial node (we choose the center of the grid); 2) move to its neighbor with probability $1/d(v_t)$, where $d(v_t)$ is the degree of node v_t ; 3) repeat 2) until s different nodes have been visited. Figure 1 presents two $random\ walks$ of s=20 (on the left) and s=28 (on the right).

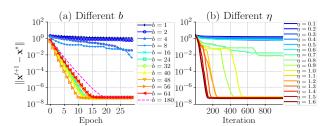


Figure 2. Choice of b and η . The left part illustrates the estimation error as a function of epochs for different choices of b. When b=180, it degenerates to GRAPHIHT (the dashed line). The right part shows the estimation error as a function of iterations for different choices of η .

Choice of b and η . We first consider how block size b and learning rate η affect the performance of GraphStoIHT. We fix the sparsity s=8 and n=180 and try different b from set $\{1,2,4,8,16,24,32,40,48,56,64,180\}$. When b=1, only 1 observation has been used at each iteration; when b=180, all measurements used, corresponding to GraphIHT. Results are presented on the left of Figure 2. In order to successfully recover x^* within 30 epochs, the block size should be at least s. A potential explanation is that, since b < s, the Hessian matrix of f_{ξ_t} , i.e., $A_{B_{\xi_t}}^{\top} A_{B_{\xi_t}}$ is not positive definite. Thus it is hard to converge to x^* in a short time. Another interesting finding is that GraphStoIHT converges faster than GraphIHT when block size

³Implementation details of the head and tail projection are provided in Supplementary material.

is suitable, say between 32 and 64. As shown in Appendix D, the run time is dominated by the time of calculating the gradient when p and m are both increasing.

To further explore learning rate η , we use the similar setting (s=8) above except b=8, m=80 (when m=80, Graphihit successfully recovers x^* with high probability as shown in Figure 4 (a). We consider 16 different η from set $\{0.1, 0.2, \ldots, 1.5, 1.6\}$. The right part of Figure 2 shows that Graphstoiht converges even when the learning rate is relatively large say $\eta \geq 1.5$. It means we can choose relatively larger learning rate so that the algorithm can achieve optimal solution faster. By Theorem 1, the optimal learning rate η is chosen by $\eta = 1/\beta$, so a potential explanation is that the strong smoothness parameter β is relatively small due to the head projection inequality $\|\nabla_H f_{\mathcal{E}_t}(x^{t+1}) - \nabla_H f_{\mathcal{E}_t}(x^*)\| \leq \beta \|x^{t+1} - x^*\|$.

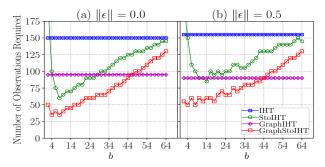


Figure 3. Robustness to noise ϵ . The number of observations required is a function of different block sizes.

Robustness to noise ϵ . To explore the performance under noise setting, we consider two noise conditions: $\|\epsilon\| = 0.0$ (without noise) and $\|\epsilon\| = 0.5$, where $\epsilon \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$. When $\|\epsilon\| = 0.5$, the sparse vector is successfully recovered if $\|x^{t+1} - x^*\| \le 0.5$. We try the block sizes b from the set $\{2, 4, 6, 8, 10, \dots, 62, 64\}$, and then measure the number of observations m required by the algorithms (so that x^* can be successfully recovered with probability 1.0). The minimum number of observations is required such that the error $\|\boldsymbol{x}^{t+1} - \boldsymbol{x}^*\| \leq 10^{-6}$ for $\|\boldsymbol{\epsilon}\| = 0$ and $\|\boldsymbol{x}^{t+1} - \boldsymbol{x}^*\| \leq$ 0.5 for $\|\boldsymbol{\epsilon}\| = 0.5$ in all trials. The results are reported in Figure 3. We can see both GRAPHSTOIHT and STOIHT are robust to noise. In particular, when block size is between 4 and 8, the number of observations required is the least for our method. Compared with the noiseless (the left), the performance of noise case (the right) degrades gracefully.

Results from synthetic dataset. We explore the performance of GraphstoIHT on the *probability of recovery*, which is defined as the total number of successful trials divided by the total number of trimmed trials. Recall that \boldsymbol{x}^* is successfully recovered if the *estimation error* $\|\boldsymbol{x}^{t+1} - \boldsymbol{x}^*\| \leq 10^{-6}$. We compare GraphstoIHT with three baseline methods, i.e., Iterative Hard Thresholding (IHT) (Blumensath & Davies, 2009), Stochastic Iterative

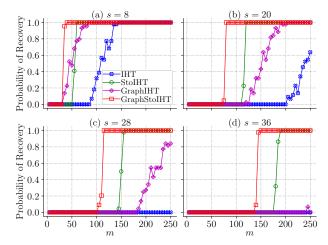


Figure 4. Probability of recovery on synthetic dataset. The probability of recovery is a function of number of observations m.

Hard Thresholding (STOIHT) (Nguyen et al., 2017), and GRAPHIHT (Hegde et al., 2016). We consider four different sparsity levels, i.e., $s \in \{8, 20, 28, 36\}$. For each trial, x^* is generated by random walk. To be consistent with the setting in Nguyen et al. (2017), the block size is set by $b = \min(s, m)$ with the number of observations m chosen from $\{5, 10, \dots, 245, 250\}$. All of the four methods, including ours, use a constant learning rate $\eta = 1$. Our method uses less observations to successfully recover x^* due to the randomness (compared with IHT and GRAPHIHT) and the graph-structured projection (compared with IHT and STOIHT) as shown in Figure 4. It indicates that GRAPH-STOIHT outperforms the other three baselines in terms of probability of recovery. It also shows that SGD-based methods are more stable than batch methods with respect to small perturbation of data as recently shown by the works of Hardt et al. (2016) and Charles & Papailiopoulos (2018).

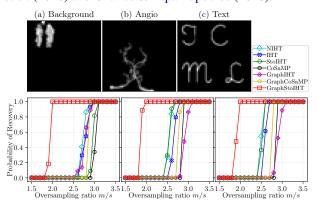


Figure 5. Probability of recovery on three 50×50 resized real images: (a) Background, (b) Angio, and (c) Text (Hegde et al., 2015b). The probability of recovery is a function of the oversampling ratio m/s.

Results from three real-world images. In order to further demonstrate the merit of GRAPHSTOIHT, we compare it with another three popular methods: NIHT (Blumen-

Table '	2 ΔΙ	IC	score -	+ standard	Ldeviation	on the	breast	cancer dataset.

Folding ID	ℓ^1 -Pathway	ℓ^1/ℓ^2 -Pathway	ℓ^1 -Edge	ℓ^1/ℓ^2 -EDGE	IHT	STOIHT	GRAPHIHT	GRAPHSTOIHT
Folding 00	0.705 ± 0.09	0.715 ± 0.07	0.726 ± 0.07	0.724 ± 0.07	0.726 ± 0.07	0.731 ±0.06	0.716 ± 0.02	0.718 ± 0.03
Folding 01	0.665 ± 0.11	0.718 ± 0.03	0.688 ± 0.04	0.703 ± 0.07	0.680 ± 0.07	0.683 ± 0.07	0.710 ± 0.06	0.704 ± 0.05
Folding 02	0.640 ± 0.07	0.729 ± 0.06	0.714 ± 0.04	0.717 ± 0.04	0.716 ± 0.06	0.723 ± 0.06	0.724 ± 0.07	0.720 ± 0.08
Folding 03	0.691 ± 0.05	0.705 ± 0.05	0.720 ± 0.05	0.722 ± 0.04	0.699 ± 0.05	0.703 ± 0.05	0.687 ± 0.04	0.687 ± 0.04
Folding 04	0.680 ± 0.05	0.690 ± 0.05	0.705 ± 0.06	0.721 ± 0.05	0.694 ± 0.05	0.695 ± 0.06	0.687 ± 0.05	0.688 ± 0.03
Folding 05	0.648 ± 0.07	0.694 ± 0.04	0.658 ± 0.06	0.683 ± 0.07	0.671 ± 0.07	0.671 ± 0.07	0.675 ± 0.07	0.712 ± 0.05
Folding 06	0.682 ± 0.04	0.733 ± 0.04	0.701 ± 0.05	0.701 ± 0.06	0.712 ± 0.07	0.733 ± 0.06	0.742 ± 0.06	0.741 ± 0.06
Folding 07	0.674 ± 0.04	0.682 ± 0.07	0.695 ± 0.04	0.704 ± 0.03	0.711 ± 0.08	0.704 ± 0.07	0.725 ± 0.07	0.714 ± 0.08
Folding 08	0.686 ± 0.06	0.705 ± 0.06	0.696 ± 0.07	0.691 ± 0.07	0.724 ± 0.07	0.720 ± 0.07	0.703 ± 0.03	0.729 ± 0.03
Folding 09	0.671 ± 0.07	0.690 ± 0.07	0.660 ± 0.05	0.687 ± 0.03	0.712 ± 0.06	0.712 ± 0.06	0.703 ± 0.06	0.704 ± 0.06
Folding 10	0.693 ± 0.09	0.735 ± 0.09	0.706 ± 0.10	0.718 ± 0.06	0.701 ± 0.08	0.717 ± 0.09	0.710 ± 0.08	0.707 ± 0.07
Folding 11	0.669 ± 0.04	0.697 ± 0.07	0.711 ± 0.05	0.704 ± 0.05	0.707 ± 0.05	0.706 ± 0.04	0.733 ± 0.06	0.733 ± 0.06
Folding 12	0.670 ± 0.05	0.716 ± 0.06	0.703 ± 0.05	0.701 ± 0.03	0.714 ± 0.06	0.715 ± 0.07	0.711 ± 0.07	0.711 ± 0.08
Folding 13	0.678 ± 0.07	0.688 ± 0.04	0.703 ± 0.04	0.697 ± 0.04	0.699 ± 0.06	0.701 ± 0.06	0.703 ± 0.04	0.715 ± 0.05
Folding 14	0.653 ± 0.02	0.700 ± 0.02	0.703 ± 0.03	0.692 ± 0.03	0.695 ± 0.04	0.699 ± 0.04	0.721 ± 0.06	0.721 ± 0.06
Folding 15	0.663 ± 0.07	0.682 ± 0.07	0.697 ± 0.08	0.687 ± 0.07	0.724 ± 0.07	0.712 ± 0.06	0.725 ± 0.07	0.716 ± 0.07
Folding 16	0.687 ± 0.07	0.720 ± 0.04	0.697 ± 0.06	0.729 ± 0.06	0.721 ± 0.05	0.723 ± 0.05	0.719 ± 0.04	0.715 ± 0.04
Folding 17	0.684 ± 0.05	0.703 ± 0.05	0.677 ± 0.07	0.716 ± 0.05	0.712 ± 0.04	0.712 ± 0.04	0.730 ± 0.03	0.720 ± 0.04
Folding 18	0.660 ± 0.06	0.714 ± 0.06	0.675 ± 0.08	0.685 ± 0.08	0.713 ± 0.06	0.706 ± 0.06	0.735 ± 0.05	0.735 ± 0.05
Folding 19	0.694 ± 0.02	0.692 ± 0.04	0.727 ± 0.05	0.713 ± 0.04	0.719 ± 0.02	0.703 ± 0.03	0.725 ± 0.05	0.715 ± 0.05
Averaged	0.675 ± 0.06	0.705 ± 0.06	0.698 ± 0.06	0.705 ± 0.06	0.707 ± 0.06	0.708 ± 0.06	0.714 ± 0.06	0.715 ±0.06

sath & Davies, 2010), CoSAMP (Needell & Tropp, 2009), GRAPHCOSAMP (Hegde et al., 2015b). We test all of them on three 50×50 resized real images: Background, Angio, and Text provided in Hegde et al. (2015b), where the first two images have one connected component while Text has four. NIHT and CoSAMP have sparsity s as input parameter. GRAPHSTOIHT shares the same head and tail projection as GRAPHCOSAMP. The learning rates η of IHT, STOIHT, GRAPHIHT and GRAPHSTOIHT are tuned from the set $\{0.2, 0.4, 0.6, 0.8\}$, and the block sizes b of STOIHT and GRAPHSTOIHT are tuned from the set $\{m/5, m/10\}$. We tune b and η on an additional validation dataset with 100 observations. To clarify, the design matrix A used here is Gaussian matrix, different from the Fourier Matrix used in Hegde et al. (2015b). Our method outperforms the others consistently.

5.2. Graph sparse logistic regression

To further test our method on real-world dataset, we apply it to the breast cancer dataset in Van De Vijver et al. (2002), which contains 295 training samples including 78 positives (metastatic) and 217 negatives (non-metastatic). Each a_i in (13) is the training sample of patient i with dimension p = 8,141 (genes). Label $y_i = 1$ if patient i has metastatic growth; otherwise -1. We use the Protein-Protein Interaction (PPI) network in Jacob et al. (2009)⁴. There are 637 pathways in this PPI network. We restrict our analysis on 3,243 genes (nodes) which form a connected graph with 19,938 edges. Due to lack of edge weights, all weights have been set to 1.0. We fold the dataset uniformly into 5 subfolds as done by Jenatton et al. (2011) to make comparison later. All related parameters are tuned by 5-fold-cross-validation on each training dataset. More experiment details are available in the Supplementary Material. We repeat the folding

strategy 20 times, and Table 2 reports AUC scores with standard deviation.

Metastasis classification. We compare our algorithm with the three aforementioned non-convex based methods and four ℓ^1/ℓ^2 mixed norm-based algorithms, ℓ^1 -PATHWAY, ℓ^1/ℓ^2 -PATHWAY, ℓ^1 -EDGE, and ℓ^1/ℓ^2 -EDGE⁵. ℓ^1 -PATHWAY and ℓ^1/ℓ^2 -PATHWAY use pathways as groups while ℓ^1 -EDGE and ℓ^1/ℓ^2 -EDGE use edges as groups. On average, GRAPHSTOIHT achieves 0.715 AUC score, the highest among the eight methods.

Gene identification. We also investigate different numbers of breast cancer-related genes identified by each method. Out of 25 breast cancer-related genes, GRAPH-STOIHT finds 24% of them, more than GRAPHIHT (20%), STOIHT (16%) and IHT (8%). The mixed norm-based methods ℓ^1 -PATHWAY, ℓ^1/ℓ^2 -PATHWAY, ℓ^1 -EDGE, and ℓ^1/ℓ^2 -EDGE find 16%, 8%, 12%, and 12% respectively. Our method outperforms other methods in finding cancer-related genes.

6. Conclusion and Future Work

In this paper, we have proposed GRAPHSTOIHT, a stochastic gradient-based method for solving graph-structured sparsity constraint problems. We proved that it enjoys a linear convergence property. Experimental evaluation shows our method consistently outperforms other algorithms for both graph sparse linear regression and graph logistic regression on real-world datasets. In future, it would be interesting to see if one can apply the variance reduction techniques such as SAGA (Defazio et al., 2014) and SVRG (Johnson & Zhang, 2013) to GRAPHSTOIHT.

⁴This network was originally proposed by Chuang et al. (2007).

⁵The code of these four methods is sourced from http://cbio.ensmp.fr/~ljacob/documents/overlasso-package.tgz.

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A. Proofs

Before proving Lemma 1, we first prove Lemma 2, 3, 4, 5, 6. We prove the main Theorem 1 and three Corollaries after proving Lemma 1. In the following proofs, given any \mathbb{M} , $\overline{\mathbb{M}}$ is the closure of \mathbb{M} under taking subsets, which is defined as $\overline{\mathbb{M}} = \{\Omega : \Omega \subseteq S, \text{ for some } S \in \mathbb{M}\}.$

Lemma 2 (co-coercivity). *If* f *is convex differentiable and satisfies the* $(\alpha, \beta, \mathcal{M}(\mathbb{M}))$ -RSS/RSC property, then we have

$$\|\nabla_{\Omega} f(\boldsymbol{x}) - \nabla_{\Omega} f(\boldsymbol{y})\|^{2} \le \beta \langle \boldsymbol{x} - \boldsymbol{y}, \nabla f(\boldsymbol{x}) - \nabla f(\boldsymbol{y}) \rangle, \tag{16}$$

where $\|\boldsymbol{x}\|_0 \cup \|\boldsymbol{y}\|_0 \subseteq \Omega$ and $\Omega \in \overline{\mathbb{M}}$.

Proof. By the definition of $(\alpha, \beta, \mathcal{M}(\mathbb{M}))$ -RSS/RSC property, we have

$$B_f(\boldsymbol{x}, \boldsymbol{y}) := f(\boldsymbol{x}) - f(\boldsymbol{y}) - \langle \nabla f(\boldsymbol{y}), \boldsymbol{x} - \boldsymbol{y} \rangle \le \frac{\beta}{2} \|\boldsymbol{x} - \boldsymbol{y}\|^2.$$

Let $x_0 \in \mathcal{M}(\mathbb{M})$ be any fixed vector. We define a surrogate function $\varphi(y) := f(y) - \langle \nabla f(x_0), y \rangle$. φ is also β -smoothness in $\mathcal{M}(\mathbb{M})$. To see this, notice that

$$B_{\varphi}(\boldsymbol{x}, \boldsymbol{y}) = \varphi(\boldsymbol{x}) - \varphi(\boldsymbol{y}) - \langle \nabla \varphi(\boldsymbol{y}), \boldsymbol{x} - \boldsymbol{y} \rangle$$

$$= f(\boldsymbol{x}) - f(\boldsymbol{y}) - \langle \nabla f(\boldsymbol{x}_0), \boldsymbol{x} - \boldsymbol{y} \rangle - \langle \nabla f(\boldsymbol{y}) - \nabla f(\boldsymbol{x}_0), \boldsymbol{x} - \boldsymbol{y} \rangle$$

$$= f(\boldsymbol{x}) - f(\boldsymbol{y}) - \langle \nabla f(\boldsymbol{y}), \boldsymbol{x} - \boldsymbol{y} \rangle$$

$$= B_f(\boldsymbol{x}, \boldsymbol{y}) \leq \frac{\beta}{2} \|\boldsymbol{x} - \boldsymbol{y}\|^2.$$

Since $\nabla \varphi(x_0) = \mathbf{0}$, φ gets its minimum at x_0 . By using the above inequality on φ with x replaced by $y - 1/\beta \nabla_{\Omega} \varphi(y)$ (noticing that $\|y - 1/\beta \nabla_{\Omega} \varphi(y)\|_0 \subseteq \Omega$), we get

$$\varphi(\boldsymbol{x}) - \varphi(\boldsymbol{y}) - \langle \nabla \varphi(\boldsymbol{y}), \boldsymbol{x} - \boldsymbol{y} \rangle \leq \frac{\beta}{2} \|\boldsymbol{x} - \boldsymbol{y}\|^{2}$$

$$\Leftrightarrow \varphi(\boldsymbol{y} - \frac{1}{\beta} \nabla_{\Omega} \varphi(\boldsymbol{y})) - \varphi(\boldsymbol{y}) - \langle \nabla \varphi(\boldsymbol{y}), -\frac{1}{\beta} \nabla_{\Omega} \varphi(\boldsymbol{y}) \rangle \leq \frac{\beta}{2} \| -\frac{1}{\beta} \nabla_{\Omega} \varphi(\boldsymbol{y}) \|^{2}$$

$$\Leftrightarrow \varphi(\boldsymbol{x}_{0}) \leq \varphi(\boldsymbol{y} - \frac{1}{\beta} \nabla_{\Omega} \varphi(\boldsymbol{y})) \leq \varphi(\boldsymbol{y}) - \frac{1}{2\beta} \|\nabla_{\Omega} \varphi(\boldsymbol{y})\|^{2}.$$

By using the definition of φ on the above inequality, we finally have

$$f(\boldsymbol{x}_{0}) - \langle \nabla f(\boldsymbol{x}_{0}), \boldsymbol{x}_{0} \rangle \leq f(\boldsymbol{y}) - \langle \nabla f(\boldsymbol{x}_{0}), \boldsymbol{y} \rangle - \frac{1}{2\beta} \|\nabla_{\Omega} f(\boldsymbol{y}) - \nabla_{\Omega} f(\boldsymbol{x}_{0})\|^{2}$$

$$\Leftrightarrow f(\boldsymbol{x}_{0}) + \langle \nabla f(\boldsymbol{x}_{0}), \boldsymbol{y} - \boldsymbol{x}_{0} \rangle + \frac{1}{2\beta} \|\nabla_{\Omega} f(\boldsymbol{y}) - \nabla_{\Omega} f(\boldsymbol{x}_{0})\|^{2} \leq f(\boldsymbol{y})$$

$$\Leftrightarrow \frac{1}{2\beta} \|\nabla_{\Omega} f(\boldsymbol{y}) - \nabla_{\Omega} f(\boldsymbol{x}_{0})\|^{2} \leq f(\boldsymbol{y}) - f(\boldsymbol{x}_{0}) + \langle \nabla f(\boldsymbol{x}_{0}), \boldsymbol{x}_{0} - \boldsymbol{y} \rangle$$
(*)

Let $x_0 = x$. By adding two copies of (*) with x and y interchanged, we prove the lemma.

Remark 1. The proof of Lemma 2 mainly follows Nesterov (2013) where the function considered is convex differentiable and has a Lipschitz continuous gradient in \mathbb{R}^p . Instead of proving the property in \mathbb{R}^p , we prove it in the subspace model $\mathcal{M}(\mathbb{M})$. Similar uses of this property can be found in Nguyen et al. (2017); Shen & Li (2017). One should notice that this co-coercivity property does not depend on the strong-convexity parameter α , i.e., $\alpha \geq 0$.

Lemma 3. For any convex differentiable function f satisfies $(\alpha, \beta, \mathcal{M}(\mathbb{M}))$ -RSS/RSC property with strongly convex parameter α and strongly smoothness parameter β , we have the following inequalities

$$\alpha \|\boldsymbol{x} - \boldsymbol{y}\| \le \|\nabla_{\Omega} f(\boldsymbol{x}) - \nabla_{\Omega} f(\boldsymbol{y})\| \le \beta \|\boldsymbol{x} - \boldsymbol{y}\|$$
(17)

where $\|\boldsymbol{x}\|_0 \cup \|\boldsymbol{y}\|_0 \subseteq \Omega$ and $\Omega \in \overline{\mathbb{M}}$.

Proof. Since f(x) satisfies $(\alpha, \beta, \mathcal{M}(\mathbb{M}))$ -RSS/RSC property, by using left inequality of (7) and exchanging x, y and summing them together, we can get

$$\alpha \| \boldsymbol{x} - \boldsymbol{y} \|^{2} \leq \langle \nabla f(\boldsymbol{x}) - \nabla f(\boldsymbol{y}), \boldsymbol{x} - \boldsymbol{y} \rangle$$

$$= \langle \nabla_{\Omega} f(\boldsymbol{x}) - \nabla_{\Omega} f(\boldsymbol{y}), \boldsymbol{x} - \boldsymbol{y} \rangle$$

$$\leq \| \nabla_{\Omega} f(\boldsymbol{x}) - \nabla_{\Omega} f(\boldsymbol{y}) \| \cdot \| \boldsymbol{x} - \boldsymbol{y} \|,$$
(*)

where the first equality follows by $\|x\|_0 \cup \|y\|_0 \subseteq \Omega$ and the second inequality follows by the Cauchy-Schwarz inequality. Therefore, the LHS of inequality (16) is obtained by eliminating $\|x - y\|$ on both sides(Notice that when x = y, two inequalities stated in this lemma are trivially true). Furthermore, $\|\nabla_{\Omega} f(x) - \nabla_{\Omega} f(y)\|^2$ can be upper bounded as

$$\begin{split} \|\nabla_{\Omega} f(\boldsymbol{x}) - \nabla_{\Omega} f(\boldsymbol{y})\|^2 &= \|\nabla_{\Omega} f(\boldsymbol{x}) - \nabla_{\Omega} f(\boldsymbol{y})\|^2 \\ &\leq \beta \langle \nabla f(\boldsymbol{x}) - \nabla f(\boldsymbol{y}), \boldsymbol{x} - \boldsymbol{y} \rangle \\ &= \beta \langle \nabla_{\Omega} f(\boldsymbol{x}) - \nabla_{\Omega} f(\boldsymbol{y}), \boldsymbol{x} - \boldsymbol{y} \rangle \\ &\leq \beta \|\nabla_{\Omega} f(\boldsymbol{x}) - \nabla_{\Omega} f(\boldsymbol{y})\| \cdot \|\boldsymbol{x} - \boldsymbol{y}\|, \end{split}$$

where the first inequality follows by Lemma 2. Hence, the RHS of (16) is obtained by eliminating $\|\nabla_{\Omega} f(x) - \nabla_{\Omega} f(y)\|$ on both sides. We prove the lemma.

Lemma 4 (Nguyen et al. (2017)). Let ξ_t be a discrete random variable defined on [n] and its probability mass function is defined as $Pr(\xi_t = i) := 1/n$, which means the probability of ξ_t selects ith block at time t. For any fixed sparse vectors \mathbf{x} , \mathbf{y} and $0 < \tau < \frac{2}{\beta}$, we have

$$\mathbb{E}_{\xi_t} \left\| \boldsymbol{x} - \boldsymbol{y} - \tau \left(\nabla_{\Omega} f_{\xi_t}(\boldsymbol{x}) - \nabla_{\Omega} f_{\xi_t}(\boldsymbol{y}) \right) \right\| \le \sqrt{\alpha \beta \tau^2 - 2\alpha \tau + 1} \|\boldsymbol{x} - \boldsymbol{y}\|, \tag{18}$$

where Ω be such set that $\|\boldsymbol{x}\|_0 \cup \|\boldsymbol{y}\|_0 \subseteq \Omega$ and $\Omega \in \overline{\mathbb{M}}$.

Proof. We first try to obtain an upper bound $\mathbb{E}_{\xi_t} \| \boldsymbol{x} - \boldsymbol{y} - \tau (\nabla_{\Omega} f_{\xi_t}(\boldsymbol{x}) - \nabla_{\Omega} f_{\xi_t}(\boldsymbol{y})) \|^2$ as the following

$$\mathbb{E}_{\xi_{t}} \left\| \boldsymbol{x} - \boldsymbol{y} - \tau \left(\nabla_{\Omega} f_{\xi_{t}}(\boldsymbol{x}) - \nabla_{\Omega} f_{\xi_{t}}(\boldsymbol{y}) \right) \right\|^{2} \\
= \|\boldsymbol{x} - \boldsymbol{y}\|^{2} - 2\tau \mathbb{E}_{\xi_{t}} \left\langle \boldsymbol{x} - \boldsymbol{y}, \nabla_{\Omega} f_{\xi_{t}}(\boldsymbol{x}) - \nabla_{\Omega} f_{\xi_{t}}(\boldsymbol{y}) \right\rangle + \tau^{2} \mathbb{E}_{\xi_{t}} \left\| \nabla_{\Omega} f_{\xi_{t}}(\boldsymbol{x}) - \nabla_{\Omega} f_{\xi_{t}}(\boldsymbol{y}) \right\|^{2} \\
= \|\boldsymbol{x} - \boldsymbol{y}\|^{2} - 2\tau \left\langle \boldsymbol{x} - \boldsymbol{y}, \mathbb{E}_{\xi_{t}} \left[\nabla f_{\xi_{t}}(\boldsymbol{x}) - \nabla f_{\xi_{t}}(\boldsymbol{y}) \right] \right\rangle + \tau^{2} \mathbb{E}_{\xi_{t}} \|\nabla_{\Omega} f_{\xi_{t}}(\boldsymbol{x}) - \nabla_{\Omega} f_{\xi_{t}}(\boldsymbol{y}) \|^{2} \\
\leq \|\boldsymbol{x} - \boldsymbol{y}\|^{2} - 2\tau \left\langle \boldsymbol{x} - \boldsymbol{y}, \mathbb{E}_{\xi_{t}} \left[\nabla f_{\xi_{t}}(\boldsymbol{x}) - \nabla f_{\xi_{t}}(\boldsymbol{y}) \right] \right\rangle + \tau^{2} \beta \left\langle \boldsymbol{x} - \boldsymbol{y}, \mathbb{E}_{\xi_{t}} \left[\nabla f_{\xi_{t}}(\boldsymbol{x}) - \nabla f_{\xi_{t}}(\boldsymbol{y}) \right] \right\rangle \\
= \|\boldsymbol{x} - \boldsymbol{y}\|^{2} + (\tau^{2}\beta - 2\tau) \left\langle \boldsymbol{x} - \boldsymbol{y}, F(\boldsymbol{x}) - F(\boldsymbol{y}) \right\rangle \\
\leq (\alpha\beta\tau^{2} - 2\alpha\tau + 1) \|\boldsymbol{x} - \boldsymbol{y}\|^{2},$$

where the second equality uses the fact that $\|x\|_0 \cup \|y\|_0 \subseteq \Omega$, the first inequality follows from Lemma 2, the third equality is obtained by using the fact that $\mathbb{E}_{\xi_t}[\nabla f_{\xi_t}(x) - \nabla f_{\xi_t}(y)] = \nabla F(x) - \nabla F(y)$ and the last inequality is due to the restricted strong convexity of F(x) on $\mathcal{M}(\mathbb{M})$. We complete the proof by taking the square root of both sides and using the fact: for any random variable X, we have $(\mathbb{E}[X])^2 \leq \mathbb{E}X^2$.

Lemma 5 (Matrix Chernoff (Tropp (2012))). Consider a finite sequence M_1, M_2, \dots, M_r of $d \times d$, independent, random, self-adjoint matrices that satisfy $M_i \succeq 0$ and $\lambda_{max}(M_i) \leq R$ almost surely. Let $\theta_{max} = \lambda_{max}(\sum_{i=1}^r \mathbb{E}[M_i])$. Then for $\nu \geq 0$,

$$Pr\left\{\lambda_{max}\left(\sum_{i=1}^{r} \mathbf{M}_{i}\right) \ge (1+\nu)\theta_{max}\right\} \le de^{\frac{\theta_{max}}{R}h(\nu)},\tag{19}$$

where $h(\nu) = \nu - (1 + \nu) \log(1 + \nu)$.

Lemma 6. Given the matrix $\mathbf{A}_{B_i}^{\mathsf{T}} \mathbf{A}_{B_i} = \sum_{j=1}^{m/n} \mathbf{a}_{i_j} \mathbf{a}_{i_j}^{\mathsf{T}}$ and $\|\mathbf{a}_{i_j}\| = 1$. $\mathbf{a}_{i_1}, \mathbf{a}_{i_2}, \ldots$ are independent and random. The matrix norm of $\mathbf{A}_{B_i}^{\mathsf{T}} \mathbf{A}_{B_i}$ can be bounded as $\|\mathbf{A}_{B_i}^{\mathsf{T}} \mathbf{A}_{B_i}\|_2 < (1+\nu)\theta_{max}$ with probability $1-p\exp\left(-\theta_{max}\nu/4\right)$, where $\theta_{max} = \lambda_{max}(\sum_{j=1}^{n/m} \mathbb{E}[\mathbf{a}_{i_j} \mathbf{a}_{i_j}^{\mathsf{T}}])$, and $\nu \geq 1$.

Proof. Let $M_j = a_{i_j} a_{i_j}^T$. Then the finite sequence $M_1, M_2, \dots, M_{m/n}$ is independent, random, and self-adjoint matrices. We write the matrix as a summation as the following

$$\mathbf{A}_{B_i}^\mathsf{T} \mathbf{A}_{B_i} = \sum_{j=1}^{m/n} oldsymbol{a}_{i_j} oldsymbol{a}_{i_j}^\mathsf{T} = \sum_{j=1}^{m/n} \mathbf{M}_j.$$

Given $\nu \geq 1$, we know the fact that $\nu - (1+\nu)\log(1+\nu) \leq -\frac{\nu\log(1+\nu)}{2}$. Furthermore, $-\log(1+\nu) \leq -\frac{\nu}{1+\nu}$, which means $\nu - (1+\nu)\log(1+\nu) \leq \frac{-\nu^2}{2(1+\nu)}$. As $\frac{\nu}{1+\nu} \geq \frac{1}{2}$, $\nu - (1+\nu)\log(1+\nu) \leq -\frac{\nu}{4}$. Therefore, we use inequality (19) of Lemma 5 by replacing r with m/n and R with 1 (by noticing that $\boldsymbol{a}_{i_j}\boldsymbol{a}_{i_j}^\mathsf{T}$ is normalized).

$$Pr\left\{\lambda_{max}\left(\sum_{j=1}^{m/n}\mathbf{M}_{j}\right) \geq (1+\nu)\theta_{max}\right\}$$

$$\leq p\exp\left(\frac{\theta_{max}}{R}\left(\nu-(1+\nu)\log(1+\nu)\right)\right)$$

$$\leq p\exp\left(\frac{\theta_{max}}{R}\frac{-\nu}{4}\right) = p\exp\left(-\frac{\theta_{max}\nu}{4}\right).$$

In other words, the probability of $Pr\left\{\lambda_{max}\left(\sum_{j=1}^{m/n}\mathbf{M}_{j}\right)<(1+\nu)\theta_{max}\right\}$ is greater than $1-p\exp\left(-\frac{\theta_{max}\nu}{4}\right)$. We prove the lemma.

Remark 2. Lemma 6 shows that the maximum eigenvalue of matrix $A_{B_i}^{\top} A_{B_i}$ can be upper bounded properly. Our Corollary 2 depends on Lemma 6.

Lemma 1. Denote $\xi_{[t]} = (\xi_0, \xi_1, \dots, \xi_t)$ as the history of the stochastic process ξ_0, ξ_1, \dots , up to time t and all random variables ξ_t are independent of each other. Define the probability mass function $Pr(\xi_t = i) = 1/n, 1 \le i \le n$. Given an optimal solution $\mathbf{x}^* \in \mathcal{M}(\mathbb{M})$, head projection model $(c_{\mathcal{H}}, \mathbb{M} \oplus \mathbb{M}_{\mathcal{T}}, \mathbb{M}_{\mathcal{H}})$, tail projection model $(c_{\mathcal{T}}, \mathbb{M}, \mathbb{M}_{\mathcal{T}})$ and $f_{\xi_t}(\cdot)$ and $F(\mathbf{x})$ satisfies Assumption 3, then we have

$$\mathbb{E}_{\xi_t} \| (\boldsymbol{x}^t - \boldsymbol{x}^*)_{H^c} \| \le \sqrt{1 - \alpha_0^2} \mathbb{E}_{\xi_t} \| \boldsymbol{x}^t - \boldsymbol{x}^* \| + \left(\frac{\beta_0}{\alpha_0} + \frac{\alpha_0 \beta_0}{\sqrt{1 - \alpha_0^2}} \right) \mathbb{E}_{\xi_t} \| \nabla_I f_{\xi_t}(\boldsymbol{x}^*) \|, \tag{20}$$

where

$$\begin{split} H := \| \mathrm{P}(\nabla f_{\xi_t}(\boldsymbol{x}^t), \mathbb{M} \oplus \mathbb{M}_{\mathcal{T}}, \mathbb{M}_{\mathcal{H}}) \|_0, \quad \alpha_0 := c_{\mathcal{H}} \alpha \tau - \sqrt{\alpha \beta \tau^2 - 2\alpha \tau + 1}, \quad \beta_0 := (1 + c_{\mathcal{H}})\tau, \\ I := \underset{S \in \mathbb{M} \oplus \mathbb{M}_{\mathcal{T}} \oplus \mathbb{M}_{\mathcal{H}}}{\max} \mathbb{E}_{\xi_t} \| \nabla_S f_{\xi_t}(\boldsymbol{x}^*) \|, \text{ and } 0 < \tau < 2/\beta. \end{split}$$

Proof. Let $r^t = x^t - x^*$ and $\Omega := ||r^t||_0 \in \overline{\mathbb{M} \oplus \mathbb{M}_T}$. The component $\mathbb{E}_{\xi_t} ||\nabla_H f_{\xi_t}(x^t)||$ can be lower bounded as

$$\mathbb{E}_{\xi_{t}} \| \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{t}) \| \geq c_{\mathcal{H}} \mathbb{E}_{\xi_{t}} \max_{S \in \mathbb{M} \oplus \mathbb{M}_{\mathcal{T}}} \| \nabla_{S} f_{\xi_{t}}(\boldsymbol{x}^{t}) \|$$

$$= c_{\mathcal{H}} \mathbb{E}_{\xi_{t}} \max_{S' \in \mathbb{M} \oplus \mathbb{M}_{\mathcal{T}}} \| \nabla_{S'} f_{\xi_{t}}(\boldsymbol{x}^{t}) \|$$

$$\geq c_{\mathcal{H}} \mathbb{E}_{\xi_{t}} \| \nabla_{\Omega} f_{\xi_{t}}(\boldsymbol{x}^{t}) \|$$

$$\geq c_{\mathcal{H}} \mathbb{E}_{\xi_{t}} (\| \nabla_{\Omega} f_{\xi_{t}}(\boldsymbol{x}^{t}) - \nabla_{\Omega} f_{\xi_{t}}(\boldsymbol{x}^{*}) \| - \| \nabla_{\Omega} f_{\xi_{t}}(\boldsymbol{x}^{*}) \|_{2})$$

$$\geq c_{\mathcal{H}} (\| \nabla_{\Omega} F(\boldsymbol{x}^{t}) - \nabla_{\Omega} F(\boldsymbol{x}^{*}) \| - \mathbb{E}_{\xi_{t} | \xi_{[t-1]}} \| \nabla_{\Omega} f_{\xi_{t}}(\boldsymbol{x}^{*}) \|_{2})$$

$$\geq c_{\mathcal{H}} \alpha \| \boldsymbol{r}^{t} \|_{2} - c_{\mathcal{H}} \mathbb{E}_{\xi_{t}} \| \nabla_{I} f_{\xi_{t}}(\boldsymbol{x}^{*}) \|,$$

where the first inequality follows by the definition of the head projection and the last inequality follows by (16) using

 $(\alpha, \beta, \mathbb{M} \oplus \mathbb{M}_T \oplus \mathbb{M}_H)$ -RSC/RSS property. The component $\|\nabla_H f_{\xi_t}(x^t)\|$ can also be upper bounded as

$$\mathbb{E}_{\xi_{t}} \| \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{t}) \| \leq \frac{1}{\tau} \mathbb{E}_{\xi_{t}} \| \tau \left(\nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{t}) - \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{*}) \right) \| + \mathbb{E}_{\xi_{t}} \| \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{*}) \|$$

$$\leq \frac{1}{\tau} \mathbb{E}_{\xi_{t}} \| \tau \left(\nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{t}) - \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{*}) \right) - \boldsymbol{r}_{H}^{t} \|_{2} + \frac{1}{\tau} \mathbb{E}_{\xi_{t}} \| \boldsymbol{r}_{H}^{t} \| + \mathbb{E}_{\xi_{t}} \| \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{*}) \|$$

$$\leq \frac{1}{\tau} \mathbb{E}_{\xi_{t}} \| \tau \left(\nabla_{H \cup \Omega} f_{\xi_{t}}(\boldsymbol{x}^{t}) - \nabla_{H \cup \Omega} f_{\xi_{t}}(\boldsymbol{x}^{*}) \right) - \boldsymbol{r}_{H \cup \Omega}^{t} \|_{2} + \frac{1}{\tau} \mathbb{E}_{\xi_{t}} \| \boldsymbol{r}_{H}^{t} \| + \mathbb{E}_{\xi_{t}} \| \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{*}) \|$$

$$= \frac{1}{\tau} \mathbb{E}_{\xi_{t}} \| \boldsymbol{r}^{t} - \tau \left(\nabla_{H \cup \Omega} f_{\xi_{t}}(\boldsymbol{x}^{t}) - \nabla_{H \cup \Omega} f_{\xi_{t}}(\boldsymbol{x}^{*}) \right) \| + \frac{1}{\tau} \mathbb{E}_{\xi_{t}} \| \boldsymbol{r}_{H}^{t} \| + \mathbb{E}_{\xi_{t}} \| \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{*}) \|$$

$$\leq \frac{\sqrt{\alpha \beta \tau^{2} - 2\alpha \tau + 1}}{\tau} \| \boldsymbol{r}^{t} \| + \frac{1}{\tau} \| \boldsymbol{r}_{H}^{t} \| + \mathbb{E}_{\xi_{t}} \| \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{*}) \|,$$

where the last inequality follows from (18) of Lemma 4 by using the fact that $H \cup \Omega$ is in $\mathbb{M} \oplus \mathbb{M}_{\mathcal{T}} \oplus \mathbb{M}_{\mathcal{H}}$. Combining the two bounds, we obtain the inequality:

$$\|\boldsymbol{r}_H^t\| \ge \alpha_0 \|\boldsymbol{r}^t\| - \beta_0 \mathbb{E}_{\xi_t} \|\nabla_I f_{\xi_t}(\boldsymbol{x}^*)\|,$$

where $\alpha_0 = c_{\mathcal{H}}\alpha\tau - \sqrt{\alpha\beta\tau^2 - 2\alpha\tau + 1}$ and $\beta_0 = (1 + c_{\mathcal{H}})\tau$. In order to obtain an upper bound of $\|\boldsymbol{r}_{H^c}^t\|$, we assume $\boldsymbol{r}^t \neq \boldsymbol{0}$. Otherwise, our statement is trivially true. Two possible cases exist. The first case is that if $\alpha_0 \|\boldsymbol{r}^t\| - \beta_0 \mathbb{E}_{\xi_t} \|\nabla_I f_{\xi_t}(\boldsymbol{x}^*)\| \leq 0$, i.e. $\alpha_0 \|\boldsymbol{r}^t\| \leq \beta_0 \mathbb{E}_{\xi_t} \|\nabla_I f_{\xi_t}(\boldsymbol{x}^*)\|$, then we have $\|\boldsymbol{r}_{H^c}^t\| \leq \|\boldsymbol{r}^t\| \leq \frac{\beta_0}{\alpha_0} \mathbb{E}_{\xi_t} \|\nabla_I f_{\xi_t}(\boldsymbol{x}^*)\|$. The second case is that if $\alpha_0 \|\boldsymbol{r}^t\| - \beta_0 \mathbb{E}_{\xi_t} \|\nabla_I f_{\xi_t}(\boldsymbol{x}^*)\| > 0$, i.e. $\alpha_0 \|\boldsymbol{r}^t\| > \beta_0 \mathbb{E}_{\xi_t} \|\nabla_I f_{\xi_t}(\boldsymbol{x}^*)\|$, then we have

$$\|\boldsymbol{r}_{H}^{t}\| \geq \left(\alpha_{0} - \frac{\beta_{0}\mathbb{E}_{\xi_{t}}\|\nabla_{I}f_{\xi_{t}}(\boldsymbol{x}^{*})\|}{\|\boldsymbol{r}^{t}\|}\right)\|\boldsymbol{r}^{t}\|.$$

Moreover, notice that $\|\boldsymbol{r}_H^t\|^2 = \|\boldsymbol{r}^t\|^2 - \|\boldsymbol{r}_{H^c}^t\|_2^2$. We obtain

$$\|\boldsymbol{r}_{H^c}^t\| \leq \|\boldsymbol{r}^t\| \sqrt{1 - \left(\alpha_0 - \frac{\beta_0 \mathbb{E}_{\xi_t} \|\nabla_I f_{\xi_t}(\boldsymbol{x}^*)\|}{\|\boldsymbol{r}^t\|}\right)^2}.$$

Denote $x_0 := \alpha_0 - \beta_0 \mathbb{E}_{\xi_t} \|\nabla_I f_{\xi_t}(\boldsymbol{x}^*)\| / \|\boldsymbol{r}^t\|$. Notice that $x_0 \in [0,1)$. Define function $\varphi(x) = 1/\sqrt{1-x^2} - xx_0/\sqrt{1-x^2}$ on (0,1). The first derivative of φ is $\varphi'(x) = (x-x_0)/(1-x^2)\sqrt{1-x^2}$ and φ gets its minimum at x_0 , i.e., $\varphi(x_0) \leq \varphi(x)$. Therefore, substituting into the bound for $\|\boldsymbol{r}_{H^c}^t\|$, we get

$$\begin{aligned} \| \boldsymbol{r}_{H^{c}}^{t} \| &\leq \| \boldsymbol{r}^{t} \| \varphi(x_{0}) \leq \| \boldsymbol{r}^{t} \| \varphi(x) \\ &= \| \boldsymbol{r}^{t} \| \left(\frac{1}{\sqrt{1 - x^{2}}} - \frac{x}{\sqrt{1 - x^{2}}} \left(\alpha_{0} - \frac{\beta_{0} \mathbb{E}_{\xi_{t}} \| \nabla_{I} f_{\xi_{t}}(\boldsymbol{x}^{*}) \|}{\| \boldsymbol{r}^{t} \|} \right) \right) \\ &= \frac{1 - x \alpha_{0}}{\sqrt{1 - x^{2}}} \| \boldsymbol{r}^{t} \| + \frac{x \beta_{0}}{\sqrt{1 - x^{2}}} \mathbb{E}_{\xi_{t}} \| \nabla_{I} f_{\xi_{t}}(\boldsymbol{x}^{*}) \|. \end{aligned}$$

In order to make the overall convergence rate as small as possible, we should make $(1-x\alpha_0)/\sqrt{1-x^2}$ as small as possible. Define the φ function again as $\varphi(x)=(1-x\alpha_0)/\sqrt{1-x^2}$, we obtain the minimum at $x=\alpha_0$. Therefore, by combining the two cases, we obtain

$$\|\boldsymbol{r}_{H^c}^t\| \leq \sqrt{1-\alpha_0^2} \|\boldsymbol{r}^t\| + \left[\frac{\beta_0}{\alpha_0} + \frac{\alpha_0 \beta_0}{\sqrt{1-\alpha_0^2}}\right] \mathbb{E}_{\xi_t} \|\nabla_I f_{\xi_t}(\boldsymbol{x}^*)\|,$$

which proves the lemma.

Remark 3. We should point out that we mainly follow the proof of Lemma 10 in Hegde et al. (2015a) where they consider the least square as the objective function with the model-RIP property. However, there are two main differences: 1) we assume a more general case, that is, our objective F(x) is strongly convex and each $f_{\xi_t}(x)$ is strong smoothness. The above lemma can be applied to any function that satisfies this property; 2) The inequality (8) satisfies in stochastic setting.

Theorem 1. Denote $\xi_{[t]} = (\xi_0, \xi_1, \dots, \xi_t)$ as the history of the stochastic process ξ_0, ξ_1, \dots up to time t, and all random variables ξ_t are independent of each other. Define the probability mass function $Pr(\xi_t = i) = 1/n$. Let \mathbf{x}^* be an optimal solution of (1) and \mathbf{x}^0 be the start point of Algorithm 1. If we choose a constant learning rate with $\eta_t = \eta$, then the solution \mathbf{x}^{t+1} generated by Algorithm 1 satisfies

$$\mathbb{E}_{\xi_{[t]}} \| \boldsymbol{x}^{t+1} - \boldsymbol{x}^* \| \le \kappa^{t+1} \| \boldsymbol{x}^0 - \boldsymbol{x}^* \| + \frac{1}{1 - \kappa} \left(\frac{\beta_0}{\alpha_0} + \frac{\alpha_0 \beta_0}{\sqrt{1 - \alpha_0^2}} + \eta \right) \mathbb{E}_{\xi_t} \| \nabla_I f_{\xi_t}(\boldsymbol{x}^*) \|,$$

where

$$\kappa = (1 + c_{\mathcal{T}}) \Big(\sqrt{\alpha \beta \eta^2 - 2\alpha \eta + 1} + \sqrt{1 - \alpha_0^2} \Big), \qquad \alpha_0 = c_{\mathcal{H}} \alpha \tau - \sqrt{\alpha \beta \tau^2 - 2\alpha \tau + 1},$$

$$\beta_0 = (1 + c_{\mathcal{H}}) \tau, I = \underset{S \in \mathbb{M} \oplus \mathbb{M}_{\mathcal{T}} \oplus \mathbb{M}_{\mathcal{H}}}{\arg \max} \mathbb{E}_{\xi_t} \|\nabla_S f_{\xi_t}(\boldsymbol{x}^*)\|, \text{ and } \tau, \eta \in (0, 2/\beta).$$

Proof. Since \boldsymbol{x}^{t+1} is completely determined by the realizations of the independent random variables $(\xi_0, \xi_1, \dots, \xi_t)$, the total expectation of the approximation error $\|\boldsymbol{x}^{t+1} - \boldsymbol{x}\|$ can be taken as $\mathbb{E}\|\boldsymbol{x}^{t+1} - \boldsymbol{x}^*\| := \mathbb{E}_{\xi_{[t]}}\|\boldsymbol{x}^{t+1} - \boldsymbol{x}^*\| = \mathbb{E}_{\xi_{[t]}}\|\boldsymbol{x}^{t+1} - \boldsymbol{x}\|$. Without loss of generality, $\mathbb{E}_{\xi_{[-1]}}\|\boldsymbol{x}^0 - \boldsymbol{x}^*\|^2 = \|\boldsymbol{x}^0 - \boldsymbol{x}^*\|^2$. Define $\boldsymbol{r}^{t+1} := \boldsymbol{x}^{t+1} - \boldsymbol{x}^*$, $\Omega := \sup(\boldsymbol{r}^{t+1})$, $H := \|P(\nabla f_{\xi_t}(\boldsymbol{x}^t), \mathbb{M} \oplus \mathbb{M}_{\mathcal{T}}, \mathbb{M}_{\mathcal{H}})\|_0$, $T := \|P(\boldsymbol{x}^t - \eta \boldsymbol{b}^t, \mathbb{M}, \mathbb{M}_{\mathcal{T}})\|_0$ as the support of head projection and tail projection respectively. Firstly, we try to get an upper bound of $\|\boldsymbol{r}^{t+1}\|$ as the following

$$\begin{split} \mathbb{E}_{\xi_{[t]}} \| \boldsymbol{x}^{t+1} - \boldsymbol{x}^* \| &= \mathbb{E}_{\xi_t | \xi_{[t-1]}} \| \boldsymbol{x}^{t+1} - \boldsymbol{x}^* \| \\ &= \mathbb{E}_{\xi_t | \xi_{[t-1]}} \| P(\boldsymbol{x}^t - \eta \boldsymbol{b}^t, \mathbb{M}, \mathbb{M}_{\mathcal{T}}) - \boldsymbol{x}^* \| \\ &\leq \mathbb{E}_{\xi_t | \xi_{[t-1]}} \| P(\boldsymbol{x}^t - \eta \boldsymbol{b}^t, \mathbb{M}, \mathbb{M}_{\mathcal{T}}) - (\boldsymbol{x}^t - \eta \boldsymbol{b}^t) \| + \mathbb{E}_{\xi_t | \xi_{[t-1]}} \| (\boldsymbol{x}^t - \eta \boldsymbol{b}^t) - \boldsymbol{x}^* \| \\ &\leq (1 + c_{\mathcal{T}}) \mathbb{E}_{\xi_t | \xi_{[t-1]}} \| \boldsymbol{x}^t - \eta \boldsymbol{b}^t - \boldsymbol{x}^* \| \\ &= (1 + c_{\mathcal{T}}) \mathbb{E}_{\xi_t | \xi_{[t-1]}} \| \boldsymbol{x}^t - \eta \nabla_H f_{\xi_t}(\boldsymbol{x}^t) - \boldsymbol{x}^* \| \\ &= (1 + c_{\mathcal{T}}) \mathbb{E}_{\xi_t | \xi_{[t-1]}} \| \boldsymbol{r}^t - \eta \nabla_H f_{\xi_t}(\boldsymbol{x}^t) \|, \end{split}$$

where the first inequality follows by the triangle inequality and the second inequality follows by the definition of the tail projection. To be more specific, denote $\Gamma := \|\boldsymbol{x}^*\|_0 \in \mathbb{M}$ and given any vector \boldsymbol{a} , we always have $\|\boldsymbol{a} - \boldsymbol{a}_{\Gamma}\| \leq \|\boldsymbol{a} - \boldsymbol{x}_{\Gamma}^*\|$. By the definition of the tail projection, one can get $\|\boldsymbol{a} - \boldsymbol{a}_T\| \leq c_T \min_{S \in \mathbb{M}} \|\boldsymbol{a} - \boldsymbol{a}_S\| \leq c_T \|\boldsymbol{a}^t - \boldsymbol{a}_{\Gamma}^t\|$. We get the second inequality by replacing \boldsymbol{a} with $\boldsymbol{x}^t - \eta \boldsymbol{b}^t$. The third equality follows by Line 5 of Algorithm 1. In the rest of the proof, we just need to bound the term, $\|\boldsymbol{r}^t - \eta \nabla_H f_{\xi_t}(\boldsymbol{x}^t)\|$. Indeed, we can further bound it as

$$\begin{split} & \mathbb{E}_{\xi_{t}|\xi_{[t-1]}} \| \boldsymbol{r}^{t} - \eta \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{t}) \| = \mathbb{E}_{\xi_{t}|\xi_{[t-1]}} \| \boldsymbol{r}^{t}_{H^{c}} + \boldsymbol{r}^{t}_{H} - \eta \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{t}) \| \\ & \leq \mathbb{E}_{\xi_{t}|\xi_{[t-1]}} \left[\left\| \boldsymbol{r}^{t}_{H} - \eta \left(\nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{t}) - \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{*}) \right) \right\| + \eta \| \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{*}) \| + \| \boldsymbol{r}^{t}_{H^{c}} \| \right] \\ & \leq \mathbb{E}_{\xi_{t}|\xi_{[t-1]}} \left[\left\| \boldsymbol{r}^{t}_{H \cup \Omega} - \eta \left(\nabla_{H \cup \Omega} f_{\xi_{t}}(\boldsymbol{x}^{t}) - \nabla_{H \cup \Omega} f_{\xi_{t}}(\boldsymbol{x}^{*}) \right) \right\| + \eta \| \nabla_{H} f_{\xi_{t}}(\boldsymbol{x}^{*}) \| + \| \boldsymbol{r}^{t}_{H^{c}} \| \right] \\ & \leq \sqrt{\alpha \beta \eta^{2} - 2\alpha \eta + 1} \mathbb{E}_{\xi_{[t-1]}} \| \boldsymbol{r}^{t} \| + \eta \mathbb{E}_{\xi_{t}} \| \nabla_{I} f_{\xi_{t}}(\boldsymbol{x}^{*}) \| + \mathbb{E}_{\xi_{t}|\xi_{[t-1]}} \| \boldsymbol{r}^{t}_{H^{c}} \| \\ & \leq \left(\sqrt{\alpha \beta \eta^{2} - 2\alpha \eta + 1} + \sqrt{1 - \alpha_{0}^{2}} \right) \mathbb{E}_{\xi_{[t-1]}} \| \boldsymbol{r}^{t} \| + \left(\frac{\beta_{0}}{\alpha_{0}} + \frac{\alpha_{0}\beta_{0}}{\sqrt{1 - \alpha_{0}^{2}}} + \eta \right) \mathbb{E}_{\xi_{t}} \| \nabla_{I} f_{\xi_{t}}(\boldsymbol{x}^{*}) \|, \end{split}$$

where the first three inequalities follow by the triangle inequality and the fourth inequality uses Lemma (3). Combine these two bounds together, we obtain

$$\begin{split} \mathbb{E}_{\xi_{[t]}} \| \boldsymbol{x}^{t+1} - \boldsymbol{x}^* \| &\leq (1 + c_{\mathcal{T}}) \mathbb{E}_{\xi_t | \xi_{[t-1]}} \| \boldsymbol{r}^t - \eta \nabla_H f_{\xi_t}(\boldsymbol{x}^t) \| \\ &\leq \kappa \mathbb{E}_{\xi_{[t-1]}} \| \boldsymbol{x}^t - \boldsymbol{x}^* \| + \Big(1 + c_{\mathcal{T}} \Big) \Bigg(\frac{\beta_0}{\alpha_0} + \frac{\alpha_0 \beta_0}{\sqrt{1 - \alpha_0^2}} + \eta \Bigg) \mathbb{E}_{\xi_{[t]}} \| \nabla_I f_{\xi_t}(\boldsymbol{x}^*) \|, \end{split}$$

where $\kappa = (1 + c_T) \left(\sqrt{\alpha \beta \eta^2 - 2\alpha \eta + 1} + \sqrt{1 - (\alpha_0)^2} \right)$. We finish the proof by applying the above inequality recursively over t iterations:

$$\mathbb{E}_{\xi_{[t]}} \| \boldsymbol{x}^{t+1} - \boldsymbol{x}^* \|_2 \leq \kappa^{t+1} \| \boldsymbol{x}^0 - \boldsymbol{x}^* \|_2 + \frac{1}{1 - \kappa} \Big(1 + c_{\mathcal{T}} \Big) \left(\frac{\beta_0}{\alpha_0} + \frac{\alpha_0 \beta_0}{\sqrt{1 - \alpha_0^2}} + \eta \right) \mathbb{E}_{\xi_t} \| \nabla_I f_{\xi_t}(\boldsymbol{x}^*) \|.$$

Corollary 0. Suppose the random distribution is uniform, i.e., $nPr(\xi_t) = 1$ and the approximation factor of the head projection can be arbitrary boosted close to 1. Taking $\eta = 1/\beta$, in order to get an effective linear convergence rate, i.e., $\kappa < 1$, the inverse of the condition number $\mu = \alpha/\beta$ and tail projection factor c_T need to satisfy

$$(1+c_{\mathcal{T}})(1+2\sqrt{\mu})\sqrt{1-\mu} < 1.$$

Proof. To analyze κ , we write κ as a function of η , i.e., $\kappa(\eta) = (1 + c_T)(\sqrt{\alpha\beta\eta^2 - 2\alpha\eta + 1} + \sqrt{1 - \alpha_0^2})$. We claim that $\kappa(\eta)$ takes its minimum at $\eta = 1/\beta$. To see this, we have

$$\kappa(\eta) = (1 + c_{\mathcal{T}}) \left(\sqrt{\alpha \beta (\eta - 1/\beta)^2 + 1 - \mu} + \sqrt{1 - \alpha_0^2} \right)$$
$$\geq (1 + c_{\mathcal{T}}) \left(\sqrt{1 - \mu} + \sqrt{1 - \alpha_0^2} \right).$$

Therefore, when $\eta = 1/\beta$, $\kappa(\eta)$ takes its minimum. Recall $\alpha_0 = c_H \alpha \tau - \sqrt{\alpha \beta \tau^2 - 2\alpha \tau + 1}$, $\tau \in (0, 2/\beta)$. Again, we redefine α_0 as a function of τ

$$\alpha_0(\tau) = c_{\mathcal{H}}\alpha\tau - \sqrt{\alpha\beta\tau^2 - 2\alpha\tau + 1}, \tau \in (0, 2/\beta).$$

 $\alpha_0(\tau)$ is a concave function with respect to τ . To see this, the first and second derivative of $\alpha_0(\tau)$ are

$$\alpha_0'(\tau) = c_{\mathcal{H}}\alpha - \frac{\alpha\beta\tau - \alpha}{\sqrt{\alpha\beta\tau^2 - 2\alpha\tau + 1}}, \qquad \alpha_0''(\tau) = \left(\alpha\beta\tau^2 - 2\alpha\tau + 1\right)^{-3/2} \left(\alpha^2 - \alpha\beta\right).$$

Hence, $\alpha_0^{-\prime\prime}(\tau) < 0$ for $\tau \in (0, 2/\beta)$. To get the maximum of $\alpha_0(\tau)$, let $\alpha_0^{\prime\prime}(\tau) = 0$, and solve the following equation

$$c_{\mathcal{H}}\alpha = \frac{\alpha\beta\tau - \alpha}{\sqrt{\alpha\beta\tau^2 - 2\alpha\tau + 1}}.$$

We get two solutions

$$\tau_1 = \frac{1}{\beta} \left(1 + c_{\mathcal{H}} \sqrt{\frac{\beta - \alpha}{\beta - c_{\mathcal{H}}^2 \alpha}} \right), \qquad \tau_2 = \frac{1}{\beta} \left(1 - c_{\mathcal{H}} \sqrt{\frac{\beta - \alpha}{\beta - c_{\mathcal{H}}^2 \alpha}} \right).$$

Since we assume that the approximation factor c_H of head-oracle can be boosted to any arbitrary constant value close to 1, i.e., $c_H \to 1^-$. Therefore,

$$\lim_{c_{\mathcal{H}} \to 1^{-}} \alpha_0(\tau_1) = 2\mu - 1, \lim_{c_{\mathcal{H}} \to 1^{-}} \alpha_0(\tau_2) = -1.$$

As we require $\alpha_0 > 0$, we only take $\alpha_0(\tau_1)$. Finally, the upper bound of κ can be simplified as

$$\lim_{c_{\mathcal{H}} \to 1^{-}} \kappa = (1 + c_{\mathcal{T}}) \left(\sqrt{1 - \mu} + \sqrt{1 - \lim_{c_{\mathcal{H}} \to 1^{-}} \alpha_{0}(\tau_{1})^{2}} \right)$$

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

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

$$= (1 + c_{\mathcal{T}}) (1 + 2\sqrt{\mu}) \sqrt{1 - \mu} < 1.$$

Hence we prove the corollary.

Corollary 1. If F(x) in (11) satisfies the α -RSC property and each $f_i(x) = \frac{n}{2m} \|\mathbf{A}_{B_i} x - \mathbf{y}_{B_i}\|^2$ satisfies the β -RSS property, then we have the following condition

$$\frac{\alpha}{2} \|\boldsymbol{x}\|^2 \le \frac{1}{2m} \|\boldsymbol{A}\boldsymbol{x}\|^2 \le \frac{\beta}{2} \|\boldsymbol{x}\|^2.$$

Let the strong convexity parameter $\alpha = 1 - \delta$ and strong smoothness parameter $\beta = 1 + \delta$, where $0 < \delta < 1$. The condition of effective contraction factor is as the following

$$(1+c_{\mathcal{T}})\left(\sqrt{\frac{2}{1+\delta}}+\frac{2\sqrt{2(1-\delta)}}{1+\delta}\right)\sqrt{\delta}<1.$$

To obtain the condition of getting effective contraction factor, we need to have $\delta = \frac{1-\mu}{1+\mu} \le 7/493 \approx 0.014199$.

Proof. The derivative of $f_i(\boldsymbol{x})$ is $\nabla f_i(\boldsymbol{x}) = \frac{n}{m} \boldsymbol{A}_{B_i}^\mathsf{T} (\boldsymbol{A}_{B_i} \boldsymbol{x} - \boldsymbol{y}_{B_i})$. $\|\nabla f_i(\boldsymbol{x}) - \nabla f_i(\boldsymbol{y})\|$ can be upper bounded as

$$\|\nabla f_i(\boldsymbol{x}) - \nabla f_i(\boldsymbol{y})\| = \frac{n}{m} \|\boldsymbol{A}_{B_i}^\mathsf{T} \boldsymbol{A}_{B_i}(\boldsymbol{x} - \boldsymbol{y})\| \le \beta \|\boldsymbol{x} - \boldsymbol{y}\|,$$

where the inequality follows by β -RSS of $f_i(\boldsymbol{x})$, i.e. $\|\nabla f_i(\boldsymbol{x}) - \nabla f_i(\boldsymbol{y})\| \le \beta \|\boldsymbol{x} - \boldsymbol{y}\|$. Sum above n inequalities together and divided by 2/n, then it is equivalent to say that $F(\boldsymbol{x})$ satisfies $\frac{1}{2m} \|\boldsymbol{A}\boldsymbol{x}\|^2 \le \frac{\beta}{2} \|\boldsymbol{x}\|^2$, where $\boldsymbol{x} \in \mathcal{M}(\mathbb{M})$. In order to satisfy α -RSC, the Hessian matrix $\frac{1}{m}\boldsymbol{A}^{\mathsf{T}}\boldsymbol{A}$ also needs to satisfy

$$\frac{\alpha}{2} \|\boldsymbol{x}\|^2 \leq \frac{1}{2} \boldsymbol{x}^\mathsf{T} \frac{\boldsymbol{A}^\mathsf{T} \boldsymbol{A}}{m} \boldsymbol{x}.$$

Combine above inequalities, we prove the condition. Therefore, under the RIP requirement, F(x) is $(1 - \delta)$ -RSC and $(1 + \delta)$ -RSS. In order to satisfy Theorem 1, we require that

$$\kappa = (1 + c_T)(1 + 2\sqrt{\mu})\sqrt{1 - \mu} < 1.$$

By replacing $\mu=\frac{1-\delta}{1+\delta}$, we prove the condition of the effective contraction factor. If $\mu\geq\frac{243}{250}$ and $c_{\mathcal{T}}$ is enough close to 1, then $\kappa<1$. Finally, $\delta=\frac{1-\mu}{1+\mu}\leq 7/493\approx 0.014199$.

Corollary 2. Suppose we have the logistic loss, F(x) in (13) and each sample a_i is normalized, i.e., $||a_i|| = 1$. Then F(x) satisfies λ -RSC and each $f_i(x)$ in (14) satisfies ($\alpha + (1 + \nu)\theta_{max}$)-RSS. The condition of getting effective contraction factor of GRAPHSTOIHT is as the following

$$\frac{\lambda}{\lambda + n(1+\nu)\theta_{max}/4m} \ge \frac{243}{250},$$

with probability $1 - p \exp\left(-\theta_{max}\nu/4\right)$, where $\theta_{max} = \lambda_{max}(\sum_{j=1}^{m/n} \mathbb{E}[\boldsymbol{a}_{i_j}\boldsymbol{a}_{i_j}^\mathsf{T}])$ and $\nu \geq 1$.

Proof. The derivative and the Hessian of $f_i(x)$ are

$$\nabla f_i(\boldsymbol{x}) = \frac{n}{m} \sum_{i=1}^{m/n} \left(-\frac{y_{i_j} \boldsymbol{a}_{i_j}}{1 + \exp(y_{i_j} \boldsymbol{a}_{i_j}^\mathsf{T} \boldsymbol{x})} + \lambda \boldsymbol{x} \right), \qquad \boldsymbol{H}(\boldsymbol{x}) = \frac{n}{4m} \boldsymbol{A}_{B_i}^\mathsf{T} \boldsymbol{Q} \boldsymbol{A}_{B_i} + \lambda \boldsymbol{I}, \tag{21}$$

where Q is an $b \times b$ diagonal matrix whose diagonal entries are $Q_{i_j i_j} = sech^2(\frac{1}{2}y_{i_j} \boldsymbol{a}_{i_j}^\mathsf{T} \boldsymbol{x})$ and sech is defined as

 $sech(x) = \frac{2}{\exp{(x)} + \exp{(-x)}}$. To see this, the Hessian of $f_i(x)$ can be calculated as

$$\begin{split} \boldsymbol{H}(\boldsymbol{x}) &= \frac{n}{m} \sum_{j=1}^{m/n} \left(\frac{y_{i_j}^2 \exp\left(y_{i_j} \boldsymbol{a}_{i_j}^\mathsf{T} \boldsymbol{x}\right) \boldsymbol{a}_{i_j} \boldsymbol{a}_{i_j}^\mathsf{T}}{(1 + \exp\left(y_{i_j} \boldsymbol{a}_{i_j}^\mathsf{T} \boldsymbol{x}\right))^2} + \lambda \boldsymbol{I} \right) \\ &= \frac{n}{m} \sum_{j=1}^{m/n} \left(\frac{\boldsymbol{a}_{i_j} \boldsymbol{a}_{i_j}^\mathsf{T}}{(1 + \exp\left(-y_{i_j} \boldsymbol{a}_{i_j}^\mathsf{T} \boldsymbol{x}\right)\right) (1 + \exp\left(y_{i_j} \boldsymbol{a}_{i_j}^\mathsf{T} \boldsymbol{x}\right))} + \lambda \boldsymbol{I} \right) \\ &= \frac{n}{m} \sum_{j=1}^{m/n} \left(\frac{\boldsymbol{a}_{i_j} \boldsymbol{a}_{i_j}^\mathsf{T}}{(\exp\left(\frac{1}{2} y_{i_j} \boldsymbol{a}_{i_j}^\mathsf{T} \boldsymbol{x}\right) + \exp\left(-\frac{1}{2} y_{i_j} \boldsymbol{a}_{i_j}^\mathsf{T} \boldsymbol{x}\right))^2} + \lambda \boldsymbol{I} \right) \\ &= \frac{n}{4m} \sum_{j=1}^{m/n} \left(\boldsymbol{a}_{i_j} \left(\frac{2}{\exp\left(\frac{1}{2} y_{i_j} \boldsymbol{a}_{i_j}^\mathsf{T} \boldsymbol{x}\right) + \exp\left(-\frac{1}{2} y_{i_j} \boldsymbol{a}_{i_j}^\mathsf{T} \boldsymbol{x}\right)} \right)^2 \boldsymbol{a}_{i_j}^\mathsf{T} + 4\lambda \boldsymbol{I} \right) \\ &= \frac{n}{4m} \boldsymbol{A}_{B_i}^\mathsf{T} \boldsymbol{Q} \boldsymbol{A}_{B_i} + \lambda \boldsymbol{I}. \end{split}$$

Since $Q_{i_i i_j} \leq 1$ and $A_{B_i}^{\mathsf{T}} Q A_{B_i}$ is symmetric matrix, we have

$$\lambda \mathbf{I} \leq \mathbf{H}(\mathbf{x}) \leq \lambda \mathbf{I} + \frac{n}{4m} \mathbf{A}_{B_i}^{\mathsf{T}} \mathbf{A}_{B_i},$$
 (22)

where $0 \leq M$ means M is a positive semi-definite matrix. Hence, F(x) is λ -RSC. Given the derivative in (21), we aim at getting the upper bound of $\|\nabla f_i(x) - \nabla f_i(y)\|$ as follows

$$\begin{split} \|\nabla f_i(\boldsymbol{x}) - \nabla f_i(\boldsymbol{y})\| &= \left\| \lambda(\boldsymbol{x} - \boldsymbol{y}) + \frac{n}{m} \sum_{j=1}^{m/n} y_{i_j} \boldsymbol{a}_{i_j} (\frac{1}{1 + \exp\left(y_{i_j} \boldsymbol{a}_{i_j}^{\mathsf{T}} \boldsymbol{y}\right)} - \frac{1}{1 + \exp\left(y_{i_j} \boldsymbol{a}_{i_j}^{\mathsf{T}} \boldsymbol{x}\right)}) \right\| \\ &= \left\| \lambda(\boldsymbol{x} - \boldsymbol{y}) - \frac{n}{m} \sum_{j=1}^{m/n} \frac{y_{i_j}^2 \exp\left(y_{i_j} \boldsymbol{a}_{i_j}^{\mathsf{T}} \boldsymbol{\theta}\right)}{(1 + \exp\left(y_{i_j} \boldsymbol{a}_{i_j}^{\mathsf{T}} \boldsymbol{\theta}\right))^2} \boldsymbol{a}_{i_j} \boldsymbol{a}_{i_j}^{\mathsf{T}} (\boldsymbol{x} - \boldsymbol{y}) \right\| \\ &= \left\| \left(\lambda \boldsymbol{I} - \frac{n}{m} \sum_{j=1}^{m/n} \frac{\exp\left(y_{i_j} \boldsymbol{a}_{i_j}^{\mathsf{T}} \boldsymbol{\theta}\right)}{(1 + \exp\left(y_{i_j} \boldsymbol{a}_{i_j}^{\mathsf{T}} \boldsymbol{\theta}\right))^2} \boldsymbol{a}_{i_j} \boldsymbol{a}_{i_j}^{\mathsf{T}} \right) (\boldsymbol{x} - \boldsymbol{y}) \right\| \\ &= \left\| \left(\lambda \boldsymbol{I} - \frac{n}{4m} \boldsymbol{A}_{B_i}^{\mathsf{T}} \boldsymbol{Q} \boldsymbol{A}_{B_i} \right) (\boldsymbol{x} - \boldsymbol{y}) \right\|, \end{split}$$

where the first equality is from the derivative of $f_i(\boldsymbol{x})$ and $f_i(\boldsymbol{y})$, the second equality follows by using the mean-value theorem for $\phi(\boldsymbol{x}) = 1/(1 + \exp{(y_i \boldsymbol{a}_i^\mathsf{T} \boldsymbol{x})})$, i.e. $\phi(\boldsymbol{y}) - \phi(\boldsymbol{x}) = \nabla \phi(\boldsymbol{\theta})(\boldsymbol{y} - \boldsymbol{x})$, $\boldsymbol{\theta} \in \{(1 - t)\boldsymbol{x} + t\boldsymbol{y} : 0 \le t \le 1\}$, and the third equality follows by $y_i^2 = 1$.

We define $\|M\|_2$ as a matrix norm, i.e., $\|M\|_2 = \sup\{\|Mx\|/\|x\| : x \in \mathbb{R}^p \text{ with } x \neq 0\}$, then we always have $\|Mx\| \leq \|M\|_2 \cdot \|x\|$. Hence, $\|\nabla f_i(x) - \nabla f_i(y)\|$ can be upper bounded as

$$\|\nabla f_i(\boldsymbol{x}) - \nabla f_i(\boldsymbol{y})\| \le \|\lambda \boldsymbol{I} - \frac{n}{4m} \boldsymbol{A}_{B_i}^\mathsf{T} \boldsymbol{Q} \boldsymbol{A}_{B_i}\|_2 \|\boldsymbol{x} - \boldsymbol{y}\|$$

$$\le \left(\lambda + \frac{n}{4m} \|\boldsymbol{A}_{B_i}^\mathsf{T} \boldsymbol{Q} \boldsymbol{A}_{B_i}\|_2\right) \|\boldsymbol{x} - \boldsymbol{y}\|$$

$$\le \left(\lambda + \frac{n}{4m} \|\boldsymbol{A}_{B_i}^\mathsf{T} \boldsymbol{A}_{B_i}\|_2\right) \|\boldsymbol{x} - \boldsymbol{y}\|,$$

where the second inequality follows by the triangle inequality of matrix norm, i.e., $\|\boldsymbol{A} + \boldsymbol{B}\|_2 \leq \|\boldsymbol{A}\|_2 + \|\boldsymbol{B}\|_2$. Therefore, $f_i(\boldsymbol{x})$ satisfies $(\lambda, \lambda + \frac{n}{4m} \|\boldsymbol{A}_{B_i}^\mathsf{T} \boldsymbol{A}_{B_i}\|_2, \mathcal{M}(\mathbb{M}))$ -RSS/RSC property. Furthermore, suppose the data is normalized, i.e., $\|\boldsymbol{a}_i \boldsymbol{a}_i^\mathsf{T}\|_2 = \|\boldsymbol{a}_i\|^2 = 1$, by using Lemma 6, we can bound $\|\boldsymbol{A}_{B_i}^\mathsf{T} \boldsymbol{A}_{B_i}\|_2$ as $\|\boldsymbol{A}_{B_i}^\mathsf{T} \boldsymbol{A}_{B_i}\|_2 \leq (1+\nu)\theta_{max}$, where $\nu \geq 1$ and θ_{max} is defined above. In order to satisfy Theorem 1, we require

$$\kappa = (1 + c_{\mathcal{T}})(1 + 2\sqrt{\mu})(\sqrt{1 - \mu}) < 1,$$

where the inverse condition number of logistic regression is $\mu = \frac{\lambda}{\lambda + n(1+\nu)\theta_{max}/4m}$. After calculation, $\mu \geq \frac{243}{250}$. We prove the corollary.

B. Results from benchmark datasets

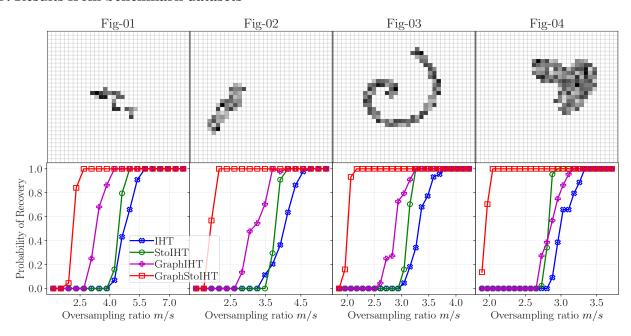


Figure 6. The probability of recovery as a function of oversampling ratio. The oversampling ratio is defined as the number of observations m divided by sparsity s, i.e., m/s. These four public benchmark graphs (a), (b), (c), and (d) in the upper row are from Arias-Castro et al. (2011).

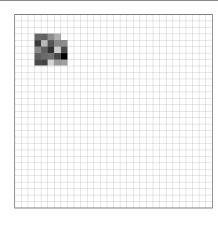
Implementation details of GRAPHSTOIHT. The head and tail projection of GRAPHSTOIHT are implemented in C language. We exactly follow the original implementation⁶. The two projections are essentially two binary search algorithms. Each iteration of the binary search executes the Prize Collecting Steiner Tree (PCST) algorithm (Johnson et al., 2000) on the target graph. Both projections have two main parameters: a lower bound sparsity s_l and an upper bound sparsity s_u . In all of the experiments, two sparsity parameters have been set to $s_l = p/2$ and $s_u = s_l * (1 + \omega)$ for the head projection, where the ω is the tolerance parameter set to 0.1. For the tail projection, we set $s_l = s$ and $s_u = s_l * (1 + \omega)$. The binary search algorithm terminates when it reaches 50 maximum iterations.

In order to demonstrate GraphStoIHT can handle different shapes of graph structure, we consider four public benchmark graphs in Arias-Castro et al. (2011). In the upper row of Figure 6, all of the subgraphs are embedded in 33×33 grid graph. The sparsity of these four graphs is 26, 46, 92, and 132 respectively. All settings keep consistent with Section 5.1. The learning rate η and block size are tuned from $\{0.2, 0.4, 0.6, 0.8\}$ and $\{m/5, m/10\}$ on additional 100 observations. The results are reported in the bottom row of Figure 6. It shows that when the sparsity s increases, the number of observations required is also increasing. This is consistent with the lower bound of the number of observations $\mathcal{O}(s\log(p/s))$, which increases as s increases. Our method is consistent with the bound $\mathcal{O}(s\log(d(v_i)) + \log p)$ shown in Equation (5) of Hegde et al. (2015b) where the weight-degree $d(v_i)$ is 4 and the budget is s-1. To further compare our method with GraphCosamp (Hegde et al., 2015b), we repeat the experiment conducted in Appendix A of Hegde et al. (2015b) and report the results in Figure 7. Thanks to the introduced randomness, GraphStoIHT outperforms GraphCosamp.

C. Results from breast cancer dataset

Parameter tuning. All parameters are selected by using 5-fold-cross-validation. The sparsity s is tuned from $\{10, 15, 20, 25, \ldots, 90, 95, 100\}$. The regularization parameter λ is tuned from $\{10^{-3}, 10^{-4}\}$. The block size b is tuned from $\{m, m/2\}$. Since we need to find a connected subgraph, the number of connected component is g=1. The 4 non-convex methods all use backtracking line search (ArmijoGoldstein condition) to obtain the learning rate at each iteration and stop at the maximum iteration of 40. The dataset has been normalized by using z-score, i.e., each sample a_i is subtracted by mean and divided by the standard deviation of $\{a_i\}_{i=1}^{295}$.

⁶The original implementation is available at: https://github.com/ludwigschmidt/cluster_approx.



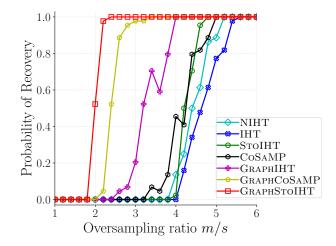


Figure 7. The probability of recovery as a function of oversampling ratio. This grid graph is from Hegde et al. (2015b).

We first measure the performance by using the balanced classification error in Jacob et al. (2009). The results are reported in Table 4. Grapht produces the lowest balanced classification error while the results of GraphstoIHT are still competitive. We argue that this may be caused by insufficient samples.

Gene identification. We pooled 25 genes from Phaedra et al. (2009); Couch et al. (2017); Rheinbay et al. (2017); Györffy et al. (2010). These genes are highly related with breast cancer, namely, MKI67, CDKN1A, ATM, TFF3, FBP1, XBP1, DSC2, CDH3, CHEK2, TOP2A, BRCA1, BRCA2, BARD1, NAT1, CA12, AR, TK2, RAD51D, GATA3, TOP2B, CCND3, CCND2, AGR2, FOXA1, and FOXC1. GRAPHSTOIHT identifies more breast cancer-related genes but use less gene features (see Table 5) than ℓ^1/ℓ^2 -PATHWAY and GRAPHIHT.

Table 3. Breast cancer-related genes identified by different algorithms.

		, ,		
Algorithm Number of genes		Cancer related genes		
GRAPHSTOIHT	6	BRCA2, CCND2, CDKN1A, ATM, AR, TOP2A		
GRAPHIHT	5	ATM, CDKN1A, BRCA2, AR, TOP2A		
ℓ^1 -Pathway	4	BRCA1, CDKN1A, ATM, DSC2		
STOIHT	4	MKI67, NAT1, AR, TOP2A		
STOIHT	4	MKI67, NAT1, AR, TOP2A		
ℓ^1/ℓ^2 -Edge	3	CCND3, ATM, CDH3		
ℓ^1 -EDGE	3	CCND3, AR, CDH3		
ℓ^1/ℓ^2 -Pathway	2	BRCA1, CDKN1A		
IHT	2	NAT1, TOP2A		

<i>Table 4.</i> Balanced Classification Error on the breast cancer dataset.									
Folding ID	ℓ^1 -Pathway	ℓ^1/ℓ^2 -Pathway	ℓ^1 -Edge	ℓ^1/ℓ^2 -EDGE	IHT	STOIHT	GRAPHIHT	GRAPHSTOIHT	
Folding 00	0.349 ± 0.06	0.349 ± 0.07	0.339 ± 0.04	0.358 ± 0.06	0.352 ± 0.08	0.350 ± 0.08	0.322 ± 0.04	0.380 ± 0.05	
Folding 01	0.419 ± 0.09	0.321 ± 0.04	0.341 ± 0.05	0.364 ± 0.06	0.369 ± 0.08	0.359 ± 0.08	0.358 ± 0.08	0.383 ± 0.07	
Folding 02	0.398 ± 0.07	0.314 ± 0.05	0.340 ± 0.05	0.315 ± 0.05	0.353 ± 0.05	0.348 ± 0.07	0.311 ± 0.06	0.322 ± 0.06	
Folding 03	0.396 ± 0.03	0.342 ± 0.04	0.336 ± 0.07	0.352 ± 0.07	0.358 ± 0.06	0.342 ± 0.06	0.365 ± 0.03	0.365 ± 0.03	
Folding 04	0.380 ± 0.06	0.366 ± 0.06	0.356 ± 0.08	0.358 ± 0.05	0.349 ± 0.08	0.368 ± 0.08	0.357 ± 0.06	0.350 ± 0.05	
Folding 05	0.418 ± 0.04	0.330 ± 0.07	0.386 ± 0.11	0.368 ± 0.08	0.353 ± 0.04	0.347 ± 0.05	0.354 ± 0.09	0.310 ± 0.06	
Folding 06	0.376 ± 0.03	0.307 ± 0.04	0.384 ± 0.07	0.353 ± 0.04	0.343 ± 0.05	0.367 ± 0.08	0.321 ± 0.03	0.321 ± 0.03	
Folding 07	0.394 ± 0.04	0.363 ± 0.08	0.360 ± 0.06	0.357 ± 0.05	0.352 ± 0.06	0.334 ± 0.05	0.327 ± 0.04	0.363 ± 0.09	
Folding 08	0.408 ± 0.05	0.336 ± 0.05	0.313 ± 0.04	0.366 ± 0.07	0.340 ± 0.07	0.340 ± 0.07	0.355 ± 0.05	0.357 ± 0.05	
Folding 09	0.366 ± 0.05	0.374 ± 0.05	0.394 ± 0.03	0.359 ± 0.03	0.343 ± 0.07	0.343 ± 0.07	0.335 ± 0.06	0.335 ± 0.05	
Folding 10	0.353 ± 0.07	0.313 ± 0.08	0.362 ± 0.08	0.350 ± 0.05	0.358 ± 0.07	0.334 ± 0.07	0.356 ± 0.05	0.340 ± 0.07	
Folding 11	0.410 ± 0.06	0.376 ± 0.05	0.370 ± 0.08	0.342 ± 0.06	0.369 ± 0.05	0.384 ± 0.07	0.329 ± 0.08	0.329 ± 0.08	
Folding 12	0.371 ± 0.04	0.355 ± 0.07	0.371 ± 0.07	0.377 ± 0.04	0.327 ± 0.05	0.331 ± 0.07	0.310 ± 0.06	0.357 ± 0.08	
Folding 13	0.405 ± 0.06	0.385 ± 0.04	0.383 ± 0.04	0.384 ± 0.05	0.362 ± 0.06	0.386 ± 0.08	0.355 ± 0.04	0.325 ± 0.05	
Folding 14	0.417 ± 0.04	0.358 ± 0.05	0.374 ± 0.03	0.354 ± 0.06	0.360 ± 0.05	0.346 ± 0.06	0.297 ± 0.06	0.297 ± 0.06	
Folding 15	0.416 ± 0.08	0.381 ± 0.04	0.402 ± 0.07	0.398 ± 0.07	0.332 ± 0.09	0.349 ± 0.08	0.329 ± 0.07	0.331 ± 0.06	
Folding 16	0.382 ± 0.07	0.357 ± 0.07	0.363 ± 0.05	0.345 ± 0.07	0.341 ± 0.06	0.334 ± 0.07	0.322 ± 0.06	0.331 ± 0.06	
Folding 17	0.385 ± 0.08	0.344 ± 0.09	0.347 ± 0.08	0.355 ± 0.04	0.331 ± 0.10	0.331 ± 0.10	0.290 ±0.07	0.327 ± 0.09	
Folding 18	0.401 ± 0.04	0.309 ± 0.06	0.382 ± 0.04	0.372 ± 0.02	0.341 ± 0.05	0.394 ± 0.08	0.302 ± 0.03	0.318 ± 0.03	
Folding 19	0.391 ± 0.03	0.341 ± 0.07	0.377 ± 0.03	0.367 ± 0.06	0.352 ± 0.04	0.360 ± 0.03	0.355 ± 0.06	0.343 ± 0.04	
Averaged	0.392 ± 0.06	0.346 ± 0.06	0.364 ± 0.07	0.360 ± 0.06	0.349 ± 0.07	0.352 ± 0.07	0.332 ±0.06	0.339 ± 0.07	

<i>Table 5.</i> Number of nonzeros on the breast cancer dataset.								
Folding ID	ℓ^1 -Pathway	ℓ^1/ℓ^2 -Pathway	ℓ^1 -Edge	ℓ^1/ℓ^2 -Edge	IHT	STOIHT	GRAPHIHT	GRAPHSTOIHT
Folding 00	075.4 ± 12.92	208.6 ± 71.37	053.2 ± 13.76	081.6 ± 28.88	082.0 ± 36.00	088.0 ± 24.00	028.0 ± 04.00	037.6±10.97
Folding 01	081.6 ± 04.13	113.0 ± 27.62	003.8 ± 02.14	059.8 ± 39.75	051.0 ± 22.00	057.0 ± 22.27	095.4 ± 00.80	066.2 ± 36.55
Folding 02	072.2 ± 10.57	108.8 ± 45.66	044.6 ± 22.54	054.2 ± 19.77	075.0 ± 00.00	040.0 ± 18.44	040.6 ± 00.49	024.6 ± 08.69
Folding 03	068.0 ± 21.38	117.8 ± 54.22	041.2 ± 12.58	063.6 ± 21.84	083.0 ± 06.00	040.0 ± 08.94	051.6 ± 08.33	051.6 ± 08.33
Folding 04	044.4 ± 17.97	135.8 ± 68.41	046.6 ± 14.68	049.2 ± 17.35	035.0 ± 00.00	043.0 ± 17.20	028.8 ± 17.60	065.0 ± 11.33
Folding 05	075.6 ± 20.37	121.0 ± 50.32	051.6 ± 12.97	069.0 ± 15.95	085.0 ± 00.00	071.0 ± 28.00	049.4 ± 12.21	032.2 ± 12.06
Folding 06	058.2 ± 10.15	147.6 ± 47.86	042.2 ± 13.14	016.2 ± 22.45	010.0 ± 00.00	044.0 ± 16.85	079.6 ± 09.20	079.8 ± 09.60
Folding 07	040.2 ± 14.55	115.8 ± 70.68	022.8 ± 09.83	050.6 ± 11.99	040.0 ± 00.00	044.0 ± 03.74	098.4 ± 04.27	080.4 ± 35.45
Folding 08	049.0 ± 23.73	137.4 ± 62.11	009.2 ± 02.48	020.4 ± 07.26	095.0 ± 00.00	095.0 ± 00.00	032.2 ± 32.90	040.2 ± 33.17
Folding 09	052.0 ± 29.54	135.8 ± 34.07	024.8 ± 15.84	057.2 ± 26.11	049.0 ± 02.00	049.0 ± 02.00	052.0 ± 14.00	047.2 ± 04.40
Folding 10	052.8 ± 25.07	137.0 ± 67.58	038.6 ± 17.35	027.2 ± 13.53	095.0 ± 00.00	072.0 ± 30.27	083.0 ± 05.06	074.8 ± 18.09
Folding 11	063.8 ± 14.69	202.2 ± 40.56	055.4 ± 13.88	043.8 ± 32.08	044.0 ± 02.00	037.0 ± 09.27	098.4 ± 09.60	098.4 ± 09.60
Folding 12	057.2 ± 23.96	217.4 ± 28.99	040.8 ± 15.59	059.2 ± 06.08	046.0 ± 12.00	036.0 ± 17.72	068.2 ± 24.10	041.2 ± 16.67
Folding 13	068.0 ± 18.44	177.4 ± 91.56	056.6 ± 34.55	073.8 ± 22.13	077.0 ± 26.00	061.0 ± 12.00	088.6 ± 23.80	031.0 ± 03.74
Folding 14	066.6 ± 19.89	148.4 ± 27.38	038.6 ± 12.56	040.8 ± 15.52	070.0 ± 10.00	066.0 ± 14.28	045.0 ± 00.00	045.0 ± 00.00
Folding 15	062.8 ± 09.99	215.2 ± 51.84	037.0 ± 08.41	063.0 ± 14.21	056.0 ± 18.00	056.0 ± 22.00	029.4 ± 02.33	063.0 ± 28.57
Folding 16	051.2 ± 23.59	075.2 ± 39.45	045.8 ± 28.60	060.8 ± 16.09	088.0 ± 06.00	077.0 ± 16.91	029.6 ± 01.85	029.2 ± 02.14
Folding 17	050.6 ± 15.53	147.8 ± 18.14	043.8 ± 32.50	039.4 ± 23.10	094.0 ± 12.00	094.0 ± 12.00	034.2 ± 02.14	029.2 ± 09.83
Folding 18	069.6 ± 11.45	116.8 ± 53.54	037.8 ± 22.43	044.2 ± 21.33	088.0 ± 06.00	041.0 ± 13.19	038.6 ± 03.83	050.6 ± 13.65
Folding 19	065.0 ± 30.55	172.4 ± 20.71	064.4 ± 04.45	053.0 ± 29.57	095.0 ± 00.00	082.0 ± 11.66	032.4 ± 23.80	046.0 ± 10.20
Averaged	061.2 ± 22.12	147.6±64.84	039.9 ± 23.09	051.4 ± 27.34	067.9 ± 27.28	059.6 ± 25.70	055.2 ± 28.88	051.7±26.65

D. Run time analysis

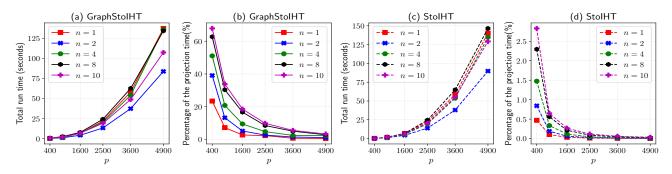


Figure 8. The run time of GRAPHSTOIHT and STOIHT as a function of p. (a) shows the total run time of GRAPHSTOIHT for recovering \boldsymbol{x}^* (Recall that \boldsymbol{x}^* has been recovered if $\|\boldsymbol{A}\boldsymbol{x}^{t+1}-\boldsymbol{y}^*\| \leq 10^{-7}$). (b) illustrates the percentage of the projection time of GRAPHSTOIHT with respect to p. (c) and (d) show the total run time and percentage of the projection time for STOIHT respectively. For GRAPHSTOIHT, the percentage of the projection time is the run time of head and tail projection divided by the total run time. For STOIHT, the percentage of the projection time is the run time of hard thresholding divided by the total run time. All results are averaged on 50 trials.

In this section, we explore the run time of GraphStoIHT on the task of graph sparse linear regression. The experimental setup follows in Section 5.1. In order to consider high dimensional and large number of measurement setting, the dimension p is chosen from $\{400, 900, 1600, 2500, 3600, 4900\}$ and the number of measurements m is set to p, i.e., m=p so that all methods have sufficient number of measurements to recover x^* . The sparsity s is set to s0 and the number of blocks s1 is from s1, s3. The learning rate is s3 shown in Figure 8 (a) and (b), when the number of measurements s4 and dimension s5 are increasing, the run time of the algorithm is dominated by calculating the gradient for both GraphStoIHT and StoIHT. For comparison, we also show the run time and the percentage of projection time in Figure 8 (c) and (d).

Figure 9 illustrates the run time of head and tail projection. It has been shown in Section 3 that the time complexity of head/tail projection is $\mathcal{O}(p\log^3(p))$ when the graph is sparse. The results of Figure 9 are consistent with the time complexity where the run time of these two projections is nearly-linear increasing with respect to the dimension p.

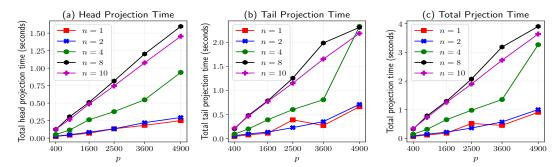


Figure 9. The run time of projections as a function of dimension p. (a) shows the total run time of the head projection for recovering x^* . (b) illustrates the total run time of tail projection of GRAPHSTOIHT with respect to p. The total projection time has been shown in (c). All results are averaged on 50 trials.

⁷To clarify, in order to be consistent with Nguyen et al. (2017), when n = 1, the learning rate of IHT algorithm proposed in Blumensath & Davies (2009) should be $\eta = 0.5$ in our implementation. It is exactly corresponding to Equation (10) in Blumensath & Davies (2009).