MSE-targeted Sampling of Bandlimited Graph Signals via Low-pass Graph Filtering Atoms

Fen Wang, Taihao Li, Minxiang Ye Institute of Artificial Intelligence, Zhejiang Lab Hangzhou, China fenwang@zhejianglab.com Gene Cheung Department of EECS, York University Toronto, Canada genec@yorku.ca

Abstract—Sampling is a fundamental problem in graph signal processing, which selects a node subset to collect samples, so that data in remaining nodes can be well recovered. In this paper, we propose a lightweight sampling algorithm to minimize MSE approximately for bandlimited graph signals via low-pass graph filtering dictionary atoms. Specifically, first we derive a greedy sampling objective without matrix inverse, based on a proxy of the MMSE criterion. We then define dictionary atoms of low-pass graph filters to accelerate computation in the reformulated problem. To further reduce complexity, we reuse partial results computed in the previous greedy step, so that candidate solutions can be evaluated via simple vector-vector multiplications. Extensive experiments show that our proposed method required the least sampling time compared to other MSEbased methods and achieved the best MSE performance overall.

Index Terms—Graph sampling, graph signal processing, greedy approach

I. INTRODUCTION

In the last decade, graph signal processing (GSP) [1]–[3] becomes a hotspot research field that that studies signals residing on graphs, such as sensed data in wireless communication networks [4] and movie ratings on social networks [5]. Signal samples on connected node pairs of a similarity graph tend to be alike; this translates to graph signals being (approximately) *bandlimited* in the graph Fourier domain. Specifically, the bandlimited (BL) graph signal has energies only in low graph frequencies, which are the first few eigenvectors of a variation operator like graph adjacency matrix, combinatorial Laplacian matrix, or their variants [2].

When the observation resources are limited, *sampling* of an assumed BL graph signal—select a node subset to collect samples such that the sought BL signal at unsampled nodes can be well estimated¹—is a fundamental problem in GSP [8]. In the noiseless case, BL graph signals can be recovered from partial samples without error by the *least square* (LS) method, if the corresponding partial eigenvector matrix is full column rank. Assuming that the signal is corrupted by additive independent and identically distributed (i.i.d.) noise, many graph sampling methods were proposed to *minimize the mean squared error* (MMSE) of the LS estimator.

Among MSE-based graph sampling schemes, [9] minimized the original MSE value directly, which is expensive computationally due to full eigen-decomposition (ED) and matrix inverse [10]. [11] leveraged graph spectral proxies (SP) to choose samples greedily using the first eigenvector² of a submatrix of a graph Laplacian raised to some power. [12] and [13] leveraged the Neumann series theorem to minimize a proxy to the MMSE criterion to mitigate matrix inverse, but still suffered from high complexity due to multiple matrixmatrix multiplications. None of the aforementioned graph sampling schemes are scalable to large graphs due to their heavy computation costs. Beyond MSE-based methods, there exist alternative low-complexity graph sampling approaches [14]–[16], but they tend to have sub-par MSE performance. Thus, a fast MSE-minimizing graph sampling scheme is still missing in the literature.

In this paper, we propose a MSE-targeted graph sampling method with low complexity leveraging *low-pass graph filter dictionary atoms* [17]—the ideal low-pass filtered outputs of impulse signals. Specifically, first, we define an equivalent sampling objective of a spectrum-shifted MMSE criterion for a sub-matrix of an ideal low-pass filter operator. Then, we define dictionary atoms of low-pass graph filters to accelerate computation of the reformulated problem. To reduce complexity, we reuse partial results computed in the previous greedy step to evaluate candidate solutions, resulting in simple vectorvector multiplications. Extensive experiments show that our proposed method had the least sampling time compared to existing MSE-based sampling schemes and achieved the best MSE performance.

II. PRELIMINARIES

A combinatorial graph can be expressed as $\mathcal{G} = (\mathcal{V}, \mathcal{E}, \mathbf{W})$ with N nodes in set $\mathcal{V} = \{1, \dots, N\}$. The connected edge set is denoted by \mathcal{E} , *i.e.*, $\exists (i, j) \in \mathcal{E}$ iff nodes i and j are connected. Entry $w_{ij} > 0$ in weight matrix \mathbf{W} is the edge weight of connected node pair (i, j) if $(i, j) \in \mathcal{E}$, and $w_{ij} = 0$ otherwise. Given weight matrix \mathbf{W} , we define a diagonal degree matrix \mathbf{D} , where $d_{ii} = \sum_{j} w_{ij}$. In this paper, we focus

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Corresponding Author: Taihao Li (lith@zhejianglab.com)

¹Sampling of graph signals can be divided into three categories: aggregation sampling [6], local measurement [7] and subset sampling [8]. In this paper, we focus on solving the subset sampling problem.

 $^{^{2}}$ We call the eigenvector corresponding to the smallest eigenvalue by *first eigenvector*.

on connected, undirected positive graphs with no self-loops, and we adopt the symmetric *combinatorial graph Laplacian matrix* $\mathbf{L} = \mathbf{D} - \mathbf{W}$ as the variation operator³. Because \mathbf{L} is real, symmetric and *positive semi-definite* (PSD) for a positive graph [18], one can write the eigen-decomposition of \mathbf{L} as $\mathbf{L} = \mathbf{V} \mathbf{\Sigma} \mathbf{V}^{\mathsf{T}}$, where $\mathbf{V} = [\mathbf{v}_1, ..., \mathbf{v}_N]$ is an orthonormal eigenvector matrix, with corresponding non-decreasing eigenvalues $0 = \lambda_1 < \lambda_2 \leq \lambda_3 \leq ... \leq \lambda_N$ along the diagonal entries of diagonal matrix $\mathbf{\Sigma}$.

The graph Fourier transform (GFT) [1] of a graph signal $\mathbf{x} \in \mathbb{R}^N$ is defined as its projected vector onto the eigenvector space of \mathbf{L} , *i.e.*, $\tilde{\mathbf{x}} = \mathbf{V}^\top \mathbf{x}$, and the inverse GFT is $\mathbf{x} = \mathbf{V}\tilde{\mathbf{x}}$. A *K*-BL graph signal \mathbf{x} means that its GFT coefficients $\tilde{\mathbf{x}}$ are non-zero only for indices smaller than or equal to *K*, with bandwidth λ_K . A *K*-BL graph signal can be expressed as $\mathbf{x} = \mathbf{V}_K \tilde{\mathbf{x}}_K$, where \mathbf{V}_K is the first *K* columns of \mathbf{V} , and $\tilde{\mathbf{x}}_K$ is the first *K* elements of $\tilde{\mathbf{x}}$. To formulate the graph sampling problem, we first define a sampling operator as follows [11].

Definition 1: To sample M elements from \mathbf{x} , where $M \leq N$, and produce $\mathbf{x}_{S} = \mathbf{C}\mathbf{x} \in \mathbb{R}^{M}$ with a sample set $S \subseteq \mathcal{V}$ and |S| = M, we define a binary sampling matrix $\mathbf{C} \in \{0, 1\}^{M \times N}$ as

$$c_{ij} = \begin{cases} 1, & j = \mathcal{S}(i); \\ 0, & \text{otherwise.} \end{cases}$$
(1)

For noiseless BL graph signals, the sampled graph signal is $\mathbf{x}_{S} = \mathbf{C}\mathbf{V}_{K}\tilde{\mathbf{x}}_{K}$. Here, the LS reconstruction can interpolate original signal without error, assuming that matrix $\mathbf{C}\mathbf{V}_{K}$ is full column rank. However, if the sampled signal is corrupted by additive noise, then different sample sets of the same size would result in different reconstruction MSE. Thus, we can minimize the resulting MSE by carefully choosing sampling matrix **C**, as done in [12]:

$$\mathbf{C}^* = \operatorname*{arg\,min}_{\mathbf{C}\in\mathcal{F}} \operatorname{Tr}\left(\left[\left(\mathbf{C}\mathbf{V}_K \right)^\top \mathbf{C}\mathbf{V}_K \right]^{-1} \right), \qquad (2)$$

where \mathcal{F} is the set of sampling matrices defined in (1). The MMSE criterion (2) is also called the *A-optimality criterion* [19].

III. LIGHTWEIGHT MSE-BASED GRAPH SAMPLING

A. Shifted MMSE Criterion and Its Equivalence

Evaluating the objective in (2) for a candidate solution C requires matrix inverse. To reduce the evaluation complexity, in [13] the authors augmented the MMSE criterion using a constant spectrum shift:

$$\mathbf{C}^* = \operatorname*{arg\,min}_{\mathbf{C}\in\mathcal{F}} \operatorname{Tr}\left(\left[(\mathbf{C}\mathbf{V}_K)^\top \mathbf{C}\mathbf{V}_K + \mu \mathbf{I} \right]^{-1} \right), \quad (3)$$

where μ is a small weight (shift) parameter with $0 < \mu < 1$. The authors then proved the following theorem:

Proposition 1: The augmented objective (3) is equivalent to

$$\mathcal{S}^* = \operatorname*{arg\,min}_{\mathcal{S}:|\mathcal{S}|=M} \operatorname{Tr}\left[(\mathbf{V}_K \mathbf{V}_K^\top)_{\mathcal{S}} + \mu \mathbf{I} \right]^{-1}, \tag{4}$$

³Some existing methods [12], [14] are designed based on symmetric normalized graph Laplacian defined by $\mathbf{L}_n = \mathbf{I} - \mathbf{D}^{-\frac{1}{2}} \mathbf{W} \mathbf{D}^{-\frac{1}{2}}$. Our proposed methods are applicable for different graph variation operators.

if the selected matrix \mathbf{CV}_K is full column rank. See [13] for a proof.

Given that the sampling problem is combinatorial in nature, a greedy approach to optimally choose one sample at a time was adopted. Specifically, assuming that we have obtained set S_t after t iterations, to choose the (t+1)-th sample, we solve the following local optimization problem:

$$\min_{i \in \mathcal{S}_{t}^{c}} \quad \underbrace{\operatorname{Tr}\left[(\mathbf{V}_{K} \mathbf{V}_{K}^{\top} + \mu \mathbf{I})_{\mathcal{S}_{t} \cup \{i\}} \right]^{-1}}_{f(\mathcal{S}_{t} \cup \{i\})}. \tag{5}$$

Define $\mathbf{G} \triangleq \mathbf{V}_K \mathbf{V}_K^\top + \mu \mathbf{I}$ for notation simplicity. Equation (5) needs to compute inverse of $\mathbf{G}_{\mathcal{S}_t \cup \{i\}}$ for every possible candidate *i* in set \mathcal{S}_t^c . To reduce the computation cost of matrix inverse, the authors introduced the next *matrix inverse lemma* to compute value $\mathbf{G}_{\mathcal{S}_t \cup \{i\}}^{-1}$ for each *i* efficiently. *Lemma 1*: The inverse of matrix **M** can be computed using

Lemma 1: The inverse of matrix M can be computed using the inverse of sub-matrix A and the inverse of the Schur complement $\mathbf{H} \triangleq \mathbf{C} - \mathbf{V}\mathbf{A}^{-1}\mathbf{U}$ of sub-matrix A of matrix M, *i.e.*, M/A, as

$$\mathbf{M}^{-1} = \begin{bmatrix} \mathbf{A} & \mathbf{U} \\ \mathbf{V} & \mathbf{C} \end{bmatrix}^{-1}$$
(6)
$$= \begin{bmatrix} \mathbf{A}^{-1} + \mathbf{A}^{-1} \mathbf{U} \mathbf{H}^{-1} \mathbf{V} \mathbf{A}^{-1} & -\mathbf{A}^{-1} \mathbf{U} \mathbf{H}^{-1} \\ -\mathbf{H}^{-1} \mathbf{V} \mathbf{A}^{-1} & \mathbf{H}^{-1} \end{bmatrix}.$$

See [20] for a proof.

Note that matrix **G** is symmetric, and under some permutation, its sub-matrix $\mathbf{G}_{\mathcal{S}_t \cup \{i\}}$ can be expressed as

$$\mathbf{G}_{\mathcal{S}_t \cup \{i\}} = \begin{bmatrix} \mathbf{G}_{\mathcal{S}_t} & \mathbf{G}_{\mathcal{S}_t, \{i\}} \\ \mathbf{G}_{\{i\}, \mathcal{S}_t} & G_{ii} \end{bmatrix} = \begin{bmatrix} \mathbf{G}_{\mathcal{S}_t} & \mathbf{g}_{t,i} \\ \mathbf{g}_{t,i}^\top & G_{ii} \end{bmatrix}, \quad (7)$$

where $\mathbf{g}_{t,i}$ denotes the partial vector of *i*-th column of **G** indexed by S_t , and G_{ii} is the *i*-th diagonal entry in **G**.

From Lemma 1, for $t \ge 1$, the inverse of $\mathbf{G}_{\mathcal{S}_t \cup \{i\}}$ is

$$\mathbf{G}_{\mathcal{S}_{t}\cup\{i\}}^{-1} = \begin{bmatrix} \mathbf{G}_{\mathcal{S}_{t}}^{-1} + h_{i}^{-1}\mathbf{G}_{\mathcal{S}_{t}}^{-1}\mathbf{g}_{t,i}\mathbf{g}_{t,i}^{-1}\mathbf{G}_{\mathcal{S}_{t}}^{-1} & -h_{i}^{-1}\mathbf{G}_{\mathcal{S}_{t}}^{-1}\mathbf{g}_{t,i}\mathbf{g}_{t,i} \\ -h_{i}^{-1}\mathbf{g}_{t,i}^{-1}\mathbf{G}_{\mathcal{S}_{t}}^{-1} & h_{i}^{-1} \end{bmatrix}$$
(8)

where $h_i = G_{ii} - \mathbf{g}_{t,i}^\top \mathbf{G}_{\mathcal{S}_t}^{-1} \mathbf{g}_{t,i}$ is a scalar. Thus,

$$f(\mathcal{S}_t \cup \{i\}) = \operatorname{Tr}\left(\mathbf{G}_{\mathcal{S}_t \cup \{i\}}^{-1}\right)$$

= $\operatorname{Tr}\left(\mathbf{G}_{\mathcal{S}_t}^{-1}\right) + h_i^{-1} \operatorname{Tr}\left(\mathbf{G}_{\mathcal{S}_t}^{-1} \mathbf{g}_{t,i} \mathbf{g}_{t,i}^{\top} \mathbf{G}_{\mathcal{S}_t}^{-1}\right) + h_i^{-1} \qquad (9)$
= $f(\mathcal{S}_t) + h_i^{-1} \|\mathbf{G}_{\mathcal{S}_t}^{-1} \mathbf{g}_{t,i}\|_2^2 + h_i^{-1}.$

Because $f(S_t)$ is a constant not affected by the selection of candidate *i*, during the (t + 1)-th greedy step, given input S_t , the sampling problem (5) can be simplified to

$$\min_{i \in \mathcal{S}_{t}^{c}} g(i) = h_{i}^{-1} \| \mathbf{G}_{\mathcal{S}_{t}}^{-1} \mathbf{g}_{t,i} \|_{2}^{2} + h_{i}^{-1} \\
\text{s.t.} \quad h_{i} = G_{ii} - \mathbf{g}_{t,i}^{\top} \mathbf{G}_{\mathcal{S}_{t}}^{-1} \mathbf{g}_{t,i}; \ \mathbf{g}_{t,i} = \mathbf{G}_{\mathcal{S}_{t},\{i\}}.$$
(10)

Compared to problem (5), instead of computing matrix inverse $\mathbf{G}_{\mathcal{S}_t \cup \{i\}}^{-1}$ for each candidate $i \in \mathcal{S}^c$, (10) just requires computing matrix-vector product $\mathbf{G}_{\mathcal{S}_t}^{-1}\mathbf{g}_{t,i}$ using known $\mathbf{G}_{\mathcal{S}_t}^{-1}$ with complexity $\mathcal{O}(M^2)$ to evaluate one candidate. In this paper, we circumvent the explicit computation of the low-pass filter matrix **G** to obtain $\mathbf{G}_{S_t}^{-1}\mathbf{g}_{t,i}$, G_{ii} and $\mathbf{g}_{t,i}$ using low-pass graph filtering dictionary atoms.

B. Low-pass Graph Filtering Atoms (LPGFA) and Greedy Problem Reformulation

Denote by \mathbf{d}_i by

$$\mathbf{d}_i = \mathbf{V}_K \mathbf{V}_K^\top \boldsymbol{\delta}_i \tag{11}$$

where δ_i is the *i*-th column of identity matrix **I**, *i.e.*, the *impulse signal* with $\delta_i(i) = 1$ and $\delta_i(j) = 0, \forall j \neq i$. The atoms $\{\mathbf{d}_i\}_{i \in \mathcal{V}}$ are called LPGFA in this paper, which is a special case of atoms defined in [17]. The *local / squared graph coherence* at node *i* is defined by quantity $\|\mathbf{d}_i\|_2^2$ [15], [21]. Next, we propose a graph sampling algorithm to solve (10) using \mathbf{d}_i 's.

Given d_i defined in (11), we compute G_{ii} in (10) as

$$G_{ii} = \boldsymbol{\delta}_i^{\top} \mathbf{V}_K \mathbf{V}_K^{\top} \boldsymbol{\delta}_i + \mu = \mathbf{d}_i(i) + \mu = \|\mathbf{d}_i\|_2^2 + \mu, \quad (12)$$

where $\mathbf{d}_i(i)$ is the *i*-th element in vector \mathbf{d}_i . The last equation holds since $\|\mathbf{d}_i\|_2^2 = (\mathbf{V}_K \mathbf{V}_K^\top \boldsymbol{\delta}_i)^\top (\mathbf{V}_K \mathbf{V}_K^\top \boldsymbol{\delta}_i) = \boldsymbol{\delta}_i^\top \mathbf{V}_K \mathbf{V}_K^\top \boldsymbol{\delta}_i = \mathbf{d}_i(i).$

Similarly, $\mathbf{g}_{t,i}$ can be computed as

$$\mathbf{g}_{t,i} = \mathbf{G}_{\mathcal{S}_t,\{i\}} = \mathbf{C}_{\mathcal{S}_t} (\mathbf{V}_K \mathbf{V}_K^{\top} + \mu \mathbf{I}) \boldsymbol{\delta}_i = \mathbf{d}_i(\mathcal{S}_t) + \mu \mathbf{C}_{\mathcal{S}_t} \boldsymbol{\delta}_i$$
(13)

where C_{S_t} is the sampling matrix corresponding to set S_t , and $d_i(S_t)$ is a sub-vector of d_i specified by set S_t .

Given $i \in S_t^c$, we know that $\delta_i(S_t) = 0$. Hence,

$$\mu \mathbf{C}_{\mathcal{S}_t} \boldsymbol{\delta}_i = \mathbf{0}. \tag{14}$$

Thus, we can write

$$\mathbf{g}_{t,i} = \mathbf{d}_i(\mathcal{S}_t). \tag{15}$$

Using (12) to (15), we simplify the greedy optimization problem (10) to the following

$$\min_{i \in \mathcal{S}_{t}^{c}} \quad h_{i}^{-1} \|\mathbf{r}_{t,i}\|_{2}^{2} + h_{i}^{-1}$$
s.t.
$$h_{i} = \|\mathbf{d}_{i}\|_{2}^{2} + \mu - \mathbf{r}_{t,i}^{\top} \mathbf{d}_{i}(\mathcal{S}_{t})$$

$$\mathbf{r}_{t,i} = \mathbf{G}_{\mathcal{S}_{t}}^{-1} \mathbf{d}_{i}(\mathcal{S}_{t})$$

$$\mathbf{d}_{i} = \mathbf{V}_{K} \mathbf{V}_{K}^{\top} \boldsymbol{\delta}_{i}$$
(16)

To obtain $\mathbf{r}_{t,i}$, we need to compute **matrix-vector** multiplication. Next, we seek to obtain $\mathbf{r}_{t,i} = \mathbf{G}_{\mathcal{S}_t}^{-1} \mathbf{d}_i(\mathcal{S}_t)$ via simple **vector-vector** multiplications instead, reusing computed results in last greedy step.

C. Evaluation Complexity Reduction via Solution Reuse

Suppose that the optimal sample in step t is j_t^* . Then $S_t = S_{t-1} \cup \{j_t^*\}$. For candidate node in unsampled set $i \in S_t^c$, we can write

$$\mathbf{d}_{i}(\mathcal{S}_{t}) = \begin{bmatrix} \mathbf{d}_{i}^{\top}(\mathcal{S}_{t-1}) & \mathbf{d}_{i}(j_{t}^{*}) \end{bmatrix}^{\top} \in \mathbb{R}^{t}.$$
 (17)

Then, for $i \in \mathcal{S}_t^c$,

$$\begin{aligned} \mathbf{r}_{t,i} &= \mathbf{G}_{\mathcal{S}_{t}}^{-1} \mathbf{d}_{i}(\mathcal{S}_{t}) = \mathbf{G}_{\mathcal{S}_{t-1} \cup \{j_{t}^{*}\}}^{-1} \mathbf{d}_{i}(\mathcal{S}_{t}) \\ &= \begin{bmatrix} \mathbf{G}_{\mathcal{S}_{t-1}}^{-1} + h_{j_{t}^{*}}^{-1} \mathbf{r}_{t-1,j_{t}^{*}} \mathbf{r}_{t-1,j_{t}^{*}}^{\top} & -h_{j_{t}^{*}}^{-1} \mathbf{r}_{t-1,j_{t}^{*}} \\ &-h_{j_{t}^{*}}^{-1} \mathbf{r}_{t-1,j_{t}^{*}}^{\top} & h_{j_{t}^{*}}^{-1} \end{bmatrix} \begin{bmatrix} \mathbf{d}_{i}(\mathcal{S}_{t-1}) \\ \mathbf{d}_{i}(j_{t}^{*}) \end{bmatrix} \\ &= \begin{bmatrix} \mathbf{r}_{t-1,i} + h_{j_{t}^{*}}^{-1} \mathbf{r}_{t-1,j_{t}^{*}} \mathbf{r}_{t-1,j_{t}^{*}}^{\top} \mathbf{d}_{i}(\mathcal{S}_{t-1}) - h_{j_{t}^{*}}^{-1} \mathbf{d}_{i}(j_{t}^{*}) \mathbf{r}_{t-1,j_{t}^{*}} \\ & -h_{j_{t}^{*}}^{-1} \mathbf{r}_{t-1,j_{t}^{*}}^{\top} \mathbf{d}_{i}(\mathcal{S}_{t-1}) + h_{j_{t}^{*}}^{-1} \mathbf{d}_{i}(j_{t}^{*}) \end{bmatrix} \\ &= \begin{bmatrix} \mathbf{r}_{t-1,i} + \alpha \mathbf{r}_{t-1,j_{t}^{*}} - \beta \mathbf{r}_{t-1,j_{t}^{*}} \\ & -\alpha + \beta \end{bmatrix} \end{aligned}$$

where

$$h_{j_t^*} = G_{j_t^*, j_t^*} - \mathbf{r}_{t-1, j_t^*}^\top \mathbf{d}_{j_t^*}(\mathcal{S}_{t-1})$$
(19)

$$\alpha = h_{j_t^*}^{-1} \mathbf{r}_{t-1,j_t^*}^\top \mathbf{d}_i(\mathcal{S}_{t-1})$$
(20)

$$\beta = h_{j_t^*}^{-1} \mathbf{d}_i(j_t^*). \tag{21}$$

Note that $\mathbf{r}_{t-1,i}$ was computed in the last greedy step for all $i \in \mathcal{S}_{t-1}^c$. Hence, we can reuse them to obtain value $\mathbf{r}_{t,i}$. Specifically, given computed $\mathbf{r}_{t-1,i}$ and \mathbf{r}_{t-1,j_t^*} , we obtain $\mathbf{r}_{t,i}$ by computing vector-vector products $\mathbf{r}_{t-1,j_t^*}^{\top} \mathbf{d}_{j_t^*}(\mathcal{S}_{t-1})$ and $\mathbf{r}_{t-1,j_t^*}^{\top} \mathbf{d}_i(\mathcal{S}_{t-1})$. The greedy evaluation complexity is thus reduced from $\mathcal{O}(M^2)$ to $\mathcal{O}(M)$ for each candidate *i*.

However, to evaluate all candidates in set S_t^c , we must compute \mathbf{d}_i 's for all $i \in \mathcal{V}$. Next, we leverage a strategy in [15] to avoid computing all \mathbf{d}_i 's.

It is known that

$$\mathbf{d}_{i}(j) = \boldsymbol{\delta}_{j}^{\top} (\mathbf{V}_{K} \mathbf{V}_{K}^{\top} \boldsymbol{\delta}_{i}) = \boldsymbol{\delta}_{i}^{\top} (\mathbf{V}_{K} \mathbf{V}_{K}^{\top} \boldsymbol{\delta}_{j}) = \mathbf{d}_{j}(i).$$
(22)

Thus, to obtain $d_i(S_t)$ in equation (16), we compute only

$$\mathbf{d}_{i}(\mathcal{S}_{t}) = \begin{bmatrix} \mathbf{d}_{i}(\mathcal{S}_{t}(1)) & \mathbf{d}_{i}(\mathcal{S}_{t}(2)) & \cdots & \mathbf{d}_{i}(\mathcal{S}_{t}(t)) \end{bmatrix}^{\top} \\ = \begin{bmatrix} \mathbf{d}_{\mathcal{S}_{t}(1)}(i) & \mathbf{d}_{\mathcal{S}_{t}(2)}(i) & \cdots & \mathbf{d}_{\mathcal{S}_{t}(t)}(i) \end{bmatrix}^{\top} \quad (23) \\ \triangleq \mathbf{d}_{\mathcal{S}_{t}}(i) \in \mathbb{R}^{t}$$

which consists of the *i*-th value of vectors $\mathbf{d}_{\mathcal{S}_t(k)}, k = 1, \dots, t. \mathbf{d}_{\mathcal{S}_t(k)}$'s are LPGFAs on nodes in sampled set \mathcal{S}_t . Thus, we compute at most M LPGFAs rather than N to obtain $\mathbf{d}_i(\mathcal{S}_t)$, where in practice sample size $M \ll N$. At step t+1, we only need to compute one new LPGFA \mathbf{d}_{j^*} . Combined results in (18) and (23) with problem (16), we obtain the optimal greedy solution via five vector-vector multiplications and at most M LPGFAs.

D. Algorithm and Complexity Analysis

The procedure for problem (10) is illustrated in Algorithm 1, called LPGFA-based graph sampling. The subscript t is abbreviated for clarity. When $S = \emptyset$, to find the first node, we need to solve the next problem based on (5):

$$\min_{i \in \mathcal{V}} \operatorname{Tr} \left[(\mathbf{V}_K \mathbf{V}_K^\top + \mu \mathbf{I})_{ii} \right]^{-1} = \frac{1}{G_{ii}}$$
(24)

which has the same optimal solution as

$$\max_{i \in \mathcal{V}} \quad G_{ii} = \|\mathbf{d}_i\|_2^2 + \mu \tag{25}$$

based on equation (12).

Algorithm 1 Low-pass graph filtering atoms (LPGFA)-based sampling of bandlimited graph signals

| Inp | ut: L, $\mathcal{S} = \emptyset$ and μ | |
|-----|---|---------------------------------|
| 1: | Compute squared graph coherence $\ \mathbf{d}_i\ _2^2$, $\forall i$ | $\in \mathcal{V}$ |
| 2: | Select the first node by $i^* = \operatorname{argmax}_{i \in \mathcal{V}} \ \mathbf{d}_i\ $ | 2 2 |
| 3: | Update $\mathcal{S} \leftarrow \mathcal{S} \cup \{i^*\}$ | |
| 4: | While $ \mathcal{S} < M$ | $\triangleright \mathcal{O}(M)$ |
| 5: | Compute one LPGFA d_{i^*} | |
| 6: | $\forall i \in S^c$, compute | $\triangleright \mathcal{O}(N)$ |
| | If $ \mathcal{S} = 1$ | |
| | $\mathbf{r}_i = rac{1}{\ \mathbf{d}_{i^*}\ _2^2 + \mu} \mathbf{d}_{i^*}(i)$ | |
| | else | |
| | $\alpha = h_{i^*}^{-1} \mathbf{r}_{i^*}^\top \mathbf{d}_{\mathcal{S} \setminus \{i^*\}}(i)$ | $\triangleright \mathcal{O}(M)$ |
| | $\beta = h_{i^*}^{-1} \mathbf{d}_{i^*}(i)$ | |
| | $\mathbf{r}_{i} = \left[\begin{array}{c} \mathbf{r}_{i} + \alpha \mathbf{r}_{i^{*}} - \beta \mathbf{r}_{i^{*}} \\ -\alpha + \beta \end{array} \right]$ | |
| | end If | |
| | Compute $h_i = \ \mathbf{d}_i\ _2^2 + \mu - \mathbf{r}_i^\top \mathbf{d}_{\mathcal{S}}(i)$ | |
| 7: | Select $i^* = \underset{i \in S^c}{\arg \min} h_i^{-1} \mathbf{r}_i _2^2 + h_i^{-1}$ | |
| | Update $\mathcal{S} \leftarrow \mathcal{S} \cup \{i^*\}$ | |
| 8: | end While | |

9: Return S

TABLE I Impact of μ on Reconstruction MSE of The Proposed Method

| μ | 0.02 | 0.06 | 0.10 | 0.14 | 0.18 | 0.22 | 0.26 |
|-----------------|------|------|------|------|------|------|-------|
| \mathcal{G}_1 | 19.1 | 19.4 | 19.9 | 21.4 | 21.1 | 22.6 | 129 |
| \mathcal{G}_2 | 20.3 | 20.7 | 21.3 | 21.9 | 23.0 | 51.2 | 1.1e3 |
| \mathcal{G}_3 | 22.0 | 22.1 | 22.7 | 23.2 | 24.1 | 24.9 | 57.6 |

After one node is sampled, *i.e.*, $S = \{i^*\}$, there is no memorized \mathbf{r}_i for greedy update using (18). Based on equation (16), for $i \in S^c$,

$$\mathbf{r}_{i} = G_{i^{*}i^{*}}^{-1} \mathbf{d}_{i}(i^{*}) = \frac{\mathbf{d}_{i^{*}}(i)}{\|\mathbf{d}_{i^{*}}\|_{2}^{2} + \mu}$$
(26)

which is easy to compute. When the sample size $|S| \ge 2$, \mathbf{r}_i can be greedily updated using equation (18).

Algorithm 1 has three parts of complexities: (1) squared graph coherence $\|\mathbf{d}_i\|_2^2$ for all nodes $i \in \mathcal{V}$; (2) LPGFA \mathbf{d}_i on sampled nodes $i \in \mathcal{S}$, and (3) sample selection. For sample selection, from Algorithm 1, we evaluate all candidates in set \mathcal{S}^c , whose size is at most N. For each evaluation, we need to compute vector-vector multiplication $\mathbf{r}_i^{\top} \mathbf{d}_{\mathcal{S}}(i)$ with complexity $\mathcal{O}(M)$. Combined with sample budget M, the sample selection complexity is $\mathcal{O}(NM^2)$. This is lower than the method proposed in [13], which is $\mathcal{O}(NM^3)$ during sampling.

IV. EXPERIMENTAL RESULTS

We evaluate the efficacy of our proposed LPGFA sampling scheme for BL graph signals via extensive simulations. All experiments were performed in MATLAB R2019a, running on a desktop with Intel Core i7-9700K CPU with 3.6 GHz CPU and 64 GB RAM. First, we generated three graphs using code in [22], with parameters listed as follows:

 \mathcal{G}_1 : community graph with N = 1000 nodes and 10 communities. Edge weights are computed based on generated coordinates \mathbf{c}_i with function $w_{ij} = \exp\left(-(\mathbf{c}_i - \mathbf{c}_j)^2\right)$; \mathcal{G}_2 : random sensor graph with 1000 nodes;

 \mathcal{G}_3 : hyper-cube graph with 1002 nodes in three dimensions.

The graph signals were assumed to be BL with bandwidth K = 50 on the eigen-space of those graphs. Their first K GFT coefficients were generated randomly using distribution $\mathcal{N}(1, 0.5^2)$, and the remaining GFT coefficients were set to be zeros. Graph signals were then computed via inverse GFT of the generated spectral coefficients. Gaussian noise was added into graph signals in the vertex domain based on different signal to noise ratios (SNRs).

First, we compared our proposed methods with five other deterministic MSE-targeted graph sampling algorithms on \mathcal{G}_1 , \mathcal{G}_2 and \mathcal{G}_3 : A-optimal [9], E-optimal [23], spectral proxies [11], GFS [13] and MIA [12]. We did not simulate other low-complexity methods since their goal is not targeting on MSE performance. We used combinatorial graph Laplacian matrix ($\mathbf{L} = \mathbf{D} - \mathbf{W}$) as the variation operator for all simulated methods. The parameters of those approaches were the same as that used in [13]. The shift parameter μ was set to 0.01 for the proposed LPGFA sampling. In this paper, we choose explicit ED to obtain the exact \mathbf{d}_i 's.

Average reconstruction MSE and execution time as functions of sample size are shown in Fig. 1, where SNR is 0dB. In this figure, the proposed LPGFA had almost the same performance as the A-optimal sampling, which optimized the original A-optimality problem using greedy approach directly, considered as the performance benchmark. Further, compared with all simulated MSE-targeted methods, our method LPGFA had the least sampling time. Based on the above analysis, we conclude that among MSE-based methods, the speed superiority of our proposed method was significant, and it also achieves the best performance. We conducted experiments on $\mathcal{G}_1, \mathcal{G}_2$ and \mathcal{G}_3 with different μ to investigate its impact on the MSE value of LPGFA method, where the sample size is 100 and SNR is 0dB. The simulation results are shown in Table. I, which indicates that the proposed LPGFA has the best performance when μ is small.

V. CONCLUSION

In this paper, we proposed a low-pass graph filtering atoms (LPGFA)-based sampling method for bandlimited graph signals targeting the MMSE criterion. We first introduced a spectrum shifted MMSE criterion and its equivalent version, which is a function of a submatrix of an ideal low-pass filter. Then, we avoided inverse computation of a greedily enlarged submatrix by using a matrix inverse lemma. We defined graph filtering atoms—the low-pass filtered impulse signal—to reformulate the problem for acceleration. For fast sampling, we reused results from the previous greedy step to speed up computation of candidates' scores. Experiments had



Fig. 1. Experimental results of different MMSE-based graph sampling algorithms in terms of sample size. From left to right: \mathcal{G}_1 , \mathcal{G}_2 and \mathcal{G}_3 .

validated the efficiency of our proposed method in both speed and MSE performance.

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