Analysis of stochastic Lanczos quadrature for spectrum approximation

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Abstract

The cumulative empirical spectral measure (CESM) $\Phi[\mathbf{A}]: \mathbb{R} \to [0,1]$ of a $n \times n$ symmetric matrix A is defined as the fraction of eigenvalues of A less than a given threshold, i.e., $\Phi[\mathbf{A}](x) := \sum_{i=1}^{n} \frac{1}{n} \mathbb{1}[\lambda_i[\mathbf{A}] \leq x].$ Spectral sums $tr(f[\mathbf{A}])$ can be computed as the Riemann– Stieltjes integral of f against $\Phi[\mathbf{A}]$, so the task of estimating CESM arises frequently in a number of applications, including machine learning. We present an error analysis for stochastic Lanczos quadrature (SLQ). We show that SLQ obtains an approximation to the CESM within a Wasserstein distance of $t |\lambda_{\max}[\mathbf{A}] - \lambda_{\min}[\mathbf{A}]|$ with probability at least $1 - \eta$, by applying the Lanczos algorithm for $\lceil 12t^{-1} + \frac{1}{2} \rceil$ iterations to $[4(n+2)^{-1}t^{-2}\ln(2n\eta^{-1})]$ vectors sampled independently and uniformly from the unit sphere. We additionally provide (matrix-dependent) a posteriori error bounds for the Wasserstein and Kolmogorov-Smirnov distances between the output of this algorithm and the true CESM. The quality of our bounds is demonstrated using numerical experiments.

1. Introduction

Given an $n \times n$ symmetric matrix \mathbf{A} , the cumulative empirical spectral measure (CESM) $\Phi[\mathbf{A}] : \mathbb{R} \to [0,1]$ gives the fraction of eigenvalues less than a given threshold. That is,

$$\Phi[\mathbf{A}](x) := \sum_{i=1}^{n} \frac{1}{n} \mathbb{I}[\lambda_i[\mathbf{A}] \le x],$$

where $\mathbb{1}[\cdot \leq x] : \mathbb{R} \to \{0,1\}$ is the indicator function defined by $\mathbb{1}[s \leq x] = 1$ if $s \leq x$ and $\mathbb{1}[s \leq x] = 0$ if s > x. The CESM contains all information about spectrum of **A** and can therefore be used to compute any quantity

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depending on just the spectrum. Conversely, computing the CESM requires exact knowledge of all the eigenvalues of **A**, which are expensive to compute.

For many applications, however, it suffices to provide a coarse estimate of the CESM. In machine learning, approximate CESMs have found use in facilitating backpropagation through implicit likelihoods (Ramesh & LeCun, 2018) as well as for for studying properties of Hessians during neural network training (Ghorbani et al., 2019; Papyan, 2019; Yao et al., 2020). This provides insight into differences between various training approaches and/or network architectures.

In data science more broadly, approximate CESMs have become a popular approach for exploring properties of graphs and networks as well as for approximating fundamental quantities such as matrix norms, log-determinants, number of eigenvalues in an interval, etc. (Avron, 2010; Napoli et al., 2016; Ubaru et al., 2017b; Han et al., 2017; Xi et al., 2018; Musco et al., 2019; Dong et al., 2019). Approximate CESMs have also long been used in computational physics and chemistry to study various observables (Ducastelle & Cyrot-Lackmann, 1970; Haydock et al., 1975; Wheeler & Blumstein, 1972) and remain widely used in these fields today (Weiße et al., 2006; Covaci et al., 2010; Sbierski et al., 2017; Schnack et al., 2020).

Finally, and as a result of their general usefulness, approximate CESMs have become the first stage of a range of algorithms for fundamental linear algebraic tasks including methods for computing matrix functions (Fan et al., 2019) and state of the art parallel eigensolvers (Polizzi, 2009; Li et al., 2019).

In this paper, we consider a well-known algorithm for computing an approximate CESM, which we refer to as stochastic Lanczos quadrature (SLQ). The algorithm described in this paper and closely related methods have been used for decades to estimate spectral densities (Haydock et al., 1972; 1975; Lambin & Gaspard, 1982; Benoit et al., 1992), and like the Lanczos algorithm (Lanczos, 1950) on which they are based, they remain highly relevant today (Lin et al., 2016; Ghorbani et al., 2019; Papyan, 2019).

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1.1. Stochastic Lanczos Quadrature

Using the standard definition of a matrix function, for a symmetric matrix \mathbf{A} , we denote by $\mathbb{1}[\mathbf{A} \leq x]$ the matrix with the same eigenvectors as \mathbf{A} , but whose eigenvalues are 0 or 1 depending on whether the corresponding eigenvalue of \mathbf{A} is below or above x; that is, $\mathbb{1}[\mathbf{A} \leq x]$ is the orthogonal projector onto the eigenspace associated to eigenvalues $\lambda_i[\mathbf{A}]$ such that $\lambda_i[\mathbf{A}] \leq x$. Thus, $\Phi[\mathbf{A}](x) = n^{-1} \operatorname{tr}(\mathbb{1}[\mathbf{A} \leq x])$.

With this definition in place, we define the weighted CESM,

$$\Psi[\mathbf{A}, \mathbf{v}](x) := \mathbf{v}^{\mathsf{T}} \mathbb{1}[\mathbf{A} \le x] \mathbf{v}.$$

If $\mathbf{v} \sim \mathcal{U}(\mathcal{S}^{n-1})$, where $\mathcal{U}(\mathcal{S}^{n-1})$ is the uniform distribution on the unit sphere, then the weighted CESM has the desirable properties (i) that it is an unbiased estimator for $\Phi[\mathbf{A}](x)$, and (ii) that it defines a cumulative probability distribution function; i.e. $\mathbb{E}[\Psi[\mathbf{A},\mathbf{v}](x)] = \Phi[\mathbf{A}](x)$ and $\Psi[\mathbf{A},\mathbf{v}]: \mathbb{R} \to [0,1]$ is weakly increasing, right continuous, and

$$\lim_{x\to -\infty} \Psi[\mathbf{A},\mathbf{v}](x) = 0, \qquad \lim_{x\to \infty} \Psi[\mathbf{A},\mathbf{v}](x) = 1.$$

Next, we consider the degree k Gaussian quadrature rule $[\Psi[\mathbf{A}, \mathbf{v}]]_k^{\mathrm{gq}}$ for $\Psi[\mathbf{A}, \mathbf{v}]$. In general, a Gaussian quadrature rule for a distribution function can be computed using the Stieltjes procedure, which for distributions of the form $\Psi[\mathbf{A}, \mathbf{v}]$, is equivalent to the Lanczos algorithm (Gautschi, 2004; Golub & Meurant, 2009). Specifically, if $[\mathbf{T}]_{:k,:k}$ is the symmetric tridiagonal matrix obtained by running Lanczos on \mathbf{A} and \mathbf{v} for k steps, then

$$[\Psi[\mathbf{A}, \mathbf{v}]]_k^{\mathrm{gq}} = \Psi([\mathbf{T}]_{:k.:k}, \hat{\mathbf{e}})$$

where $\hat{\mathbf{e}} = [1, 0, \dots, 0]^{\mathsf{T}}$.

By repeating this process over multiple samples and averaging, we arrive at SLQ, outlined in Algorithm 1.

Algorithm 1 Stochastic Lanczos Quadrature

$$\label{eq:continuous_problem} \begin{split} \overline{\textbf{input } \mathbf{A}, n_{\text{v}}, k} \\ \textbf{for } i &= 1, 2, \dots, n_{\text{v}} \ \textbf{do} \\ \text{Sample } \mathbf{v}_i &\sim \mathcal{U}(\mathcal{S}^{n-1}) \ (\text{and define } \Psi_i = \Psi(\mathbf{A}, \mathbf{v}_i)) \\ \text{Run Lanczos on } \mathbf{A}, \mathbf{v}_i \ \text{for } k \ \text{steps to compute } [\mathbf{T}_i]_{:k,:k} \\ \text{Define } [\Psi_i]_k^{\text{gq}} &= \Psi[[\mathbf{T}_i]_{:k,:k}, \hat{\mathbf{e}}] \\ \textbf{end for} \\ \textbf{return } \langle [\Psi_i]_k^{\text{gq}} \rangle := \frac{1}{n_{\text{v}}} \sum_{i=1}^{n_{\text{v}}} [\Psi_i]_k^{\text{gq}} \end{split}$$

SLQ is computationally efficient. In particular, all samples can be computed in parallel on separate machines or on a single machine using blocked linear algebra routines. Moreover, the algorithm is matrix free in that we only require a method to compute the map $\mathbf{v} \mapsto \mathbf{A}\mathbf{v}$, the cost of which we denote by T_{mv} . This is particularly important for large dense matrices, where the $O(n^2)$ storage required to keep every entry of \mathbf{A} may be intractable. In many cases, such as with Hessians from the training of neural networks, matrix-vector products can be computed implicitly using only O(n) storage (Pearlmutter, 1994). Similarly, if \mathbf{A} is sparse or structured, this map may be evaluated faster than the $O(n^2)$ computation cost for an arbitrary matrix vector product; e.g. $O(\mathrm{nnz}[\mathbf{A}])$ for a sparse matrix or $O(n\ln n)$ for a circulant matrix.

1.2. Discussion of results

Our first main result is a runtime guarantee for SLQ. In particular, we show that if $n_{\rm v} > 4(n+2)^{-1}t^{-2}\ln(2n\eta^{-1})$ and $k > 12t^{-1} + \frac{1}{2}$, then

$$\mathbb{P}\big[d_{\mathcal{W}}(\Phi[\mathbf{A}],\langle [\Psi_i]_k^{\mathrm{gq}}\rangle) > tI[\mathbf{A}]\big] < \eta,$$

where $I[\mathbf{A}] := |\lambda_{\max}[\mathbf{A}] - \lambda_{\min}[\mathbf{A}]|$ and $d_{\mathrm{W}}(\cdot, \cdot)$ denotes the Wasserstein distance between two distribution functions as in Definition 3 below. This implies that as $n \to \infty$, for $t \gg n^{-1/2}$, SLQ has a runtime of $O(T_{\mathrm{mv}}t^{-1}\log(t^{-2}\eta^{-1}))$. This bound is nearly tight in the sense that for any $t \in (0,1)$, there exists a matrix of size $\lceil (4t)^{-1} \rceil$ such that at least $(8t)^{-1}$ matrix vector products are required to obtain an output with Wasserstein distance less than $tI[\mathbf{A}]$.

The second main result is an a posteriori upper bound for Wasserstein and Kolmogorov–Smirnov (KS) distances, which take into account spectrum dependent features such as clustered or isolated eigenvalues.

Finally, in proving these results, we show that if $n_v > (n+2)^{-1}t^{-2}\ln(2\eta^{-1})$ then, for any $x \in \mathbb{R}$,

$$\mathbb{P}\left[\left|\Phi[\mathbf{A}](x) - \left(\frac{1}{n_{\mathbf{v}}} \sum_{i=1}^{n_{\mathbf{v}}} \mathbf{v}_{i}^{\mathsf{T}} \mathbb{1}[\mathbf{A} \leq x] \mathbf{v}_{i}\right)\right| > t\right] < \eta.$$
(1)

This is applicable to the analysis of a range of algorithms beyond SLQ.

1.3. Related work

A variety of algorithms for approximating the CESM have been developed; see (Fischer, 2011; Weiße et al., 2006; Lin et al., 2016; Adams et al., 2018; Cohen-Steiner et al., 2018) and the references therein for a more complete overview. By far, the most popular algorithms are the kernel polynomial method (KPM) (Weiße et al., 2006) and SLQ. The two algorithms differ primarily in how they approximate the weighted CESM $\Psi[\mathbf{A}, \mathbf{v}]$, and as a result, our analysis of

¹For a symmetric matrix \mathbf{A} with eigenvalue decomposition $\mathbf{A} = \mathbf{U} \boldsymbol{\Lambda} \mathbf{U}^\mathsf{T}$ and scalar function $f : \mathbb{R} \to \mathbb{R}$, the matrix function $f[\mathbf{A}]$ is defined by $f[\mathbf{A}] := \mathbf{U} f[\boldsymbol{\Lambda}] \mathbf{U}^\mathsf{T}$, where $f[\boldsymbol{\Lambda}]$ is the diagonal matrix with f applied entrywise to the diagonal entries of the matrix $\boldsymbol{\Lambda}$.

the estimator $\Psi[\mathbf{A}, \mathbf{v}]$ for $\Phi[\mathbf{A}]$ applies to the KPM as well. The main advantages of SLQ are that it is adaptive to the spectrum of \mathbf{A} (i.e. it automatically detects features such as gaps in the spectrum, isolated or clustered eigenvalues, etc.) and that it produces an output which is a probability distribution function. However, in doing so, SLQ requires the computation of inner products, which may be costly, especially in a distributed computing environment (Lin et al., 2016). Moreover, the behavior of the Lanczos method is not always straightforward in finite precision.

We remark that the KPM, both with exact and inexact matrix-vector products, has recently been analyzed (Braverman et al., 2021). The analysis for KPM with exact matrix-vector products yields similar rates to our analysis for SLQ, but the sample complexity is given in terms of unspecified universal constants and has a polynomially worse dependence on the accuracy parameter t. We believe this is an artifact of analysis not an inherent property of the KPM, but to be certain a full side-by-side comparison of the algorithms is needed.

Tail bounds similar to (1) have been derived in several contexts. First, while explicit constants are not specified, the same $n^{-1}t^{-2}\ln(\eta^{-1})$ dependence for $n_{\rm v}$ is implied by (Deift & Trogdon, 2020, Lemma 4.5). Second, if the ${\bf v}_i$ are replaced by *unnormalized* Gaussian vectors (with mean zero and variance n^{-1}), then well known bounds for trace estimation (Avron & Toledo, 2011; Roosta-Khorasani & Ascher, 2014) yield similar rates in terms of explicit constants. However, the weighed CESM corresponding to an unnormalized sample will not be a probability distribution function.

Finally, the algorithm studied in this paper is closely related to an algorithm, also commonly referred to as SLQ (but called gSLQ in this paper to avoid confusion), for approximating spectral sums $\operatorname{tr}(f[\mathbf{A}]) = \sum_{i=1}^n f(\lambda_i[\mathbf{A}])$ (Bai et al., 1996; Bai & Golub, 1996). Indeed, the SLQ studied here is a special case of gSLQ, with $f(s) = \mathbb{1}[s \leq x]$. However, SLQ is in fact equivalent to gSLQ the sense that the gSLQ approximation to $\operatorname{tr}(f[\mathbf{A}])$ can be obtained by computing a Riemann–Stieltjes integral of f against the output of Algorithm 1. More generally, matrix function trace estimation is closely related to CESM estimation due to the fact that

$$\operatorname{tr}(f[\mathbf{A}]) = n \int f(s) d\Phi(s).$$

In (Ubaru et al., 2017a), the convergence of gSLQ when f is smooth or analytic is studied. As a result, this analysis is not immediately applicable to CESM estimation itself, since $\mathbb{1}[\cdot \leq x]$ is discontinuous. One possibility is to solve a relaxed problem where $\mathbb{1}[\cdot \leq x]$ is replaced with a smoothed approximation such as a shifted hyperbolic tangent or the CDF of any continuous random variable with small enough

variance. This results in an approximation to the CESM equivalent to the convolution of the CESM with a smoothing kernel (Ubaru et al., 2017a; Han et al., 2017; Ghorbani et al., 2019).

If the CESM can be reasonably smoothed, then such an approach works well. However, it is often the case that the "variance" of the smoothing kernel, in order to preserve certain aspects of the CESM, such as large jumps due to clustered eigenvalues, has to be very small. In such cases, Gaussian quadrature bounds for smooth functions are often useless, necessitating bounds such as the ones presented in this paper; see Supplement C for a detailed discussion.

1.4. Preliminaries

Matrices are denoted by bold uppercase letters and vectors are denoted by bold lowercase letters. The first canonical unit vector $[1,0,\ldots,0]^\mathsf{T}$, of size determined by context, is denoted $\hat{\mathbf{e}}$. The set of all eigenvalues of a $d \times d$ symmetric matrix \mathbf{B} is denoted $\Lambda[\mathbf{B}]$, and the individual eigenvalues are $\lambda_{\min}[\mathbf{B}] = \lambda_d[\mathbf{B}] \leq \cdots \leq \lambda_1[\mathbf{B}] = \lambda_{\max}[\mathbf{B}]$. Unless otherwise stated, \mathbf{A} is $n \times n$ symmetric matrix.

We denote the *i*-th entry of a vector \mathbf{v} by $[\mathbf{v}]_i$ and the submatrix consisting of rows r to r' and columns c to c' by $[\mathbf{B}]_{r:r',c:c'}$. If any of these indices are equal to 1 or n, they may be omitted. If r=r' or c=c', then we will simply write this index once. Thus, $[\mathbf{B}]_{:,:2}$ denotes the first two columns of \mathbf{B} , and $[\mathbf{B}]_{3::}$ denotes the third row of \mathbf{B} .

For some positive integer $n_{\rm v}$ and a set of values $\{x_i\}_{i=1}^{n_{\rm v}}$, the sample average $\frac{1}{n_{\rm v}}\sum_{i=1}^{n_{\rm v}}x_i$ is denoted $\langle x_i\rangle$.

Definition 1. Let μ and ν be two probability distribution functions. We say the moments of μ and ν are equal up to degree k-1 if for all polynomials p of degree < k,

$$\int p(x)\mathrm{d}\mu(x) = \int p(x)\mathrm{d}\nu(x).$$

We also have the standard definition of Kolmogorov-Smirnov and Wasserstein distances.

Definition 2. Let μ and ν be two probability distribution functions. The Kolmogorov–Smirnov distance between μ and ν , denoted $d_{KS}(\mu, \nu)$, is defined by

$$d_{KS}(\mu, \nu) := \sup_{x} |\mu(x) - \nu(x)|.$$

Definition 3. Let μ and ν be two probability distribution functions. The Wasserstein (earth mover) distance between μ and ν , denoted $d_W(\mu, \nu)$, is defined by

$$d_{\mathbf{W}}(\mu,\nu) := \int |\mu(x) - \nu(x)| \mathrm{d}x.$$

2. The Lanczos algorithm

The primary computational cost of SLQ is due to the Lanczos algorithm (Lanczos, 1950). The Lanczos algorithm is typically viewed as a procedure for constructing an orthonormal basis $[\mathbf{Q}]_{:::k} := [\mathbf{q}_1, \dots, \mathbf{q}_k]$ for the Krylov subspace

$$\mathcal{K}_k(\mathbf{A}, \mathbf{v}) = \operatorname{span}(\mathbf{v}, \mathbf{A}\mathbf{v}, \dots, \mathbf{A}^{k-1}\mathbf{v}).$$

This can be done by a Gram–Schmidt-like process, where $\mathbf{A}\mathbf{q}_k$ is orthogonalized against previous basis vectors $\{\mathbf{q}_j\}_{j=1}^k$, which results in a factorization

$$\mathbf{A}[\mathbf{Q}]_{:,:k} = [\mathbf{Q}]_{:,:k}[\mathbf{T}]_{:k,:k} + \beta_k \mathbf{q}_{k+1} \mathbf{e}_k^\mathsf{T}$$

where $[\mathbf{T}]_{:k,:k}$ is upper-Hessenberg. However, since $[\mathbf{T}]_{:k,:k} = [\mathbf{Q}]_{:,:k}^\mathsf{T} \mathbf{A} [\mathbf{Q}]_{:,:k}$ is symmetric, then $[\mathbf{T}]_{:k,:k}$ is actually tridiagonal. Thus, $\mathbf{A}\mathbf{q}_k$ will be orthogonal to \mathbf{q}_j , j < k-1, so we only need to orthogonalize against \mathbf{q}_k and \mathbf{q}_{k-1} in each iteration. As a result, the runtime of Algorithm 2 is $O(k(T_{\mathrm{mv}} + n))$ and the required storage is O(n).

Algorithm 2 Lanczos

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\begin{split} & \overbrace{\mathbf{input} \ \mathbf{A}, \mathbf{v}, k} \\ & \mathbf{q}_0 = \mathbf{0}, \, \beta_{-1} = 0, \, \mathbf{q}_1 = \mathbf{v}/\|\mathbf{v}\| \\ & \mathbf{for} \ i = 0, 1, \dots, k-1 \ \mathbf{do} \\ & \quad \tilde{\mathbf{q}}_{i+1} = \mathbf{A}\mathbf{q}_i - \beta_{i-1}\mathbf{q}_{i-1} \\ & \quad \alpha_i = \langle \tilde{\mathbf{q}}_{i+1}, \mathbf{q}_i \rangle \\ & \quad \tilde{\mathbf{q}}_{i+1} = \tilde{\mathbf{q}}_{i+1} - \alpha_i \mathbf{q}_i \\ & \quad \mathbf{if} \ \text{`reorthogonlization'} \ \mathbf{then} \quad \text{orthogonalize} \ \tilde{\mathbf{q}}_{i+1} \\ & \quad \mathbf{against} \ \{\mathbf{q}_j\}_{j=1}^{i-1} \ \mathbf{end} \ \mathbf{if} \\ & \quad \beta_i = \|\mathbf{q}_{i+1}\| \\ & \quad \mathbf{q}_{i+1} = \tilde{\mathbf{q}}_{i+1}/\beta_i \\ & \quad \mathbf{end} \ \mathbf{for} \end{split}
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return $[\mathbf{T}]_{:k,:k}$ (diagonal $[\alpha_1, \ldots, \alpha_k]$ and the sub and super diagonal $[\beta_1, \ldots, \beta_{k-1}]$)

Remark 1. If Algorithm 2 is run for k=n iterations on any right hand side with nonzero projection onto each eigenvector, the tridiagonal matrix $[\mathbf{T}]_{:n,:n}$ produced will have the same eigenvalues as \mathbf{A} . Thus the CESM can be computed deterministically in time $O(nT_{\mathrm{mv}}+n^2)$.

Remark 2 (Gu & Eisenstat, 1995). The eigenvalues and first components of eigenvectors of a real symmetric tridiagonal matrix of size $k \times k$ can be computed in $O(k^2)$ operations.

Remark 3. Without reorthogonalization in Algorithm 2, the runtime of SLQ Algorithm 1 is $O(n_{\rm v}k(T_{\rm mv}+n))$ and the required storage is O(n). With reorthogonalization, the runtime is $O(n_{\rm v}k(T_{\rm mv}+nk))$ and the required storage is O(nk).

Remark 4. In exact arithmetic, the reorthogonalization step of Algorithm 2 is unnecessary as $\tilde{\mathbf{q}}_{i+1}$ is already orthogonal to $\{\mathbf{q}_j\}_{j=1}^{i-1}$. However, in finite precision arithmetic this orthogonality may be lost. Our bounds, as well as the

bounds for gSLQ (Ubaru et al., 2017a), are derived based on exact arithmetic theory, so it must be assumed that $[T]_{:k,:k}$ is computed using some implementation which produces an output close to the exact arithmetic output. The easiest way to ensure this is with full reorthogonalization, although other more advanced schemes have been considered.

For the task of computing CESMs, some practitioners (Papyan, 2019) have noted the algorithm still works without reorthogonalization. In Supplement D, we provide an overview of existing analysis on the Lanczos algorithm in finite precision (Paige, 1976; 1980; Greenbaum, 1989) which provides a high level explanation as to why SLQ still works without reorthogonalization.

3. Results

We now state the main results.

Theorem 1. Given $0 < \eta < 1$ and t > 0, set $n_v > 4(n + 2)^{-1}t^{-2}\ln(2n\eta^{-1})$ and $k > 12t^{-1} + \frac{1}{2}$. Then, Algorithm 1 will output an estimate $\langle [\Psi_i]_k^{\rm gq} \rangle$ satisfying,

$$\mathbb{P}\big[d_{\mathcal{W}}(\Phi[\mathbf{A}], \langle [\Psi_i]_k^{\mathrm{gq}} \rangle) > tI[\mathbf{A}]\big] < \eta.$$

where
$$I[\mathbf{A}] := |\lambda_{\max}[\mathbf{A}] - \lambda_{\min}[\mathbf{A}]|$$
.

Theorem 1 is essentially a direct consequence of following theorem of the average of weighted CESMs and a straightforward bound on the Wasserstein distances of distribution functions with matching moments.

Theorem 2. Given a positive integer n_v , suppose $\{\mathbf{v}_i\}_{i=1}^{n_v} \overset{\text{iid}}{\sim} \mathcal{U}(\mathcal{S}^{n-1})$ and define $\Psi_i = \Psi(\mathbf{A}, \mathbf{v}_i)$. Then, for all $x \in \mathbb{R}$ and t > 0,

$$\mathbb{P}\left[\left|\Phi[\mathbf{A}](x) - \langle \Psi_i(x)\rangle\right| > t\right] \le 2\exp\left(-n_{\mathbf{v}}(n+2)t^2\right)$$
$$\mathbb{P}\left[d_{\mathrm{KS}}(\Phi[\mathbf{A}], \langle \Psi_i\rangle) > t\right] \le 2n\exp\left(-n_{\mathbf{v}}(n+2)t^2\right).$$

Proposition 1. Suppose μ and ν are two probability distribution functions constant on the complement of [a,b] whose moments are equal up to degree s. Then,

$$d_{\mathbf{W}}(\mu,\nu) \le 2(b-a)(1+\pi^2/2)s^{-1} < 12(b-a)s^{-1}.$$

We also provide a posteriori error guarantees which may be of practical use.

Theorem 3. Let $\{[d_i]_j\}_{j=1}^k$ and $\{[\theta_i]_j\}_{j=1}^k$ be the squares of the first component of eigenvectors and the eigenvalues respectively of $[\mathbf{T}_i]_{:k::k}$ from Algorithm 1. Then

$$\begin{aligned} d_{\mathrm{KS}}(\langle \Psi_i \rangle, \langle [\Psi_i]_k^{\mathrm{gq}} \rangle) &\leq \left\langle \max_{j=1,\dots,k} [d_i]_j \right\rangle, \\ d_{\mathrm{W}}(\langle \Psi_i \rangle, \langle [\Psi_i]_k^{\mathrm{gq}} \rangle) &\leq \left\langle \sum_{j=0}^n \max\{[d_i]_j, [d_i]_{j+1}\} ([\theta_i]_{j+1} - [\theta_i]_j) \right\rangle \end{aligned}$$

where, and for notational convenience, we have defined $[\theta_i]_0 = a$, $[\theta_i]_{n+1} = b$, and $[d_i]_0 = [d_i]_{n+1} = 0$ for some choice of a, b such that $a \le \lambda_{\min}[\mathbf{A}]$ and $b \ge \lambda_{\max}[\mathbf{A}]$.

Note that the Wasserstein distance bounds requires knowledge of points a,b such that $a \leq \lambda_{\min}[\mathbf{A}]$ and $b \geq \lambda_{\max}[\mathbf{A}]$. Such bounds can be computed rigorously, both a priori (Kuczyński & Woźniakowski, 1992) or a posteriori (Parlett et al., 1982). In practice, $\lambda_{\min}([\mathbf{T}_i]_{:k,:k}) \to \lambda_{\min}[\mathbf{A}]$ and $\lambda_{\max}([\mathbf{T}_i]_{:k,:k}) \to \lambda_{\max}[\mathbf{A}]$ rapidly, so the j=0 and j=k terms can be omitted with negligible effect.

As noted in Remark 1, the exact CESM can be computed with n matrix vector products. However, we also have the following lower bound for a specific class of matrices.

Theorem 4. For any $t \in (0,1)$, there exists a matrix **A** of size $\lceil (4t)^{-1} \rceil$ such that if Algorithm 1 uses fewer than $(8t)^{-1}$ matrix vector products, then Algorithm 1 will output an estimate $\langle |\Psi_i|_k^{gq} \rangle$ satisfying,

$$d_{\mathbf{W}}(\Phi[\mathbf{A}], \langle [\Psi_i]_k^{\mathrm{gq}} \rangle) > tI[\mathbf{A}].$$

While Theorems 3 and 4 involve random variables, the results hold surely and therefore with probability one.

4. Analysis and proofs

For notational convenience, we denote $\Phi[\mathbf{A}]$ by Φ in proofs.

4.1. Weighted CESM

We start with analyzing the weighted CESM. Note that this analysis is applicable to many algorithms for spectrum approximation, including the KPM.

Lemma 1. Suppose $\mathbf{v} \sim \mathcal{U}(\mathcal{S}^{n-1})$ and define $m = n\Phi[\mathbf{A}](x)$. Then,

$$\Psi[\mathbf{A}, \mathbf{v}](x) \sim \text{Beta}\left(\frac{m}{2}, \frac{n-m}{2}\right).$$

Proof. Let $\mathbf{U} = [\mathbf{u}_1, \dots, \mathbf{u}_n]$, where \mathbf{u}_i is the *i*-th normalized eigenvector of \mathbf{A} . Since \mathbf{U} is orthogonal, by the invariance of $\mathcal{U}(\mathcal{S}^{n-1})$ under orthogonal transforms, we have that $\mathbf{U}^\mathsf{T}\mathbf{v} \sim \mathcal{U}(\mathcal{S}^{n-1})$.

We may therefore assume $\mathbf{U}^\mathsf{T}\mathbf{v} \stackrel{\mathrm{d}}{=} \mathbf{x}/\|\mathbf{x}\|$, where $\mathbf{x} \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$. Recall that the *i*-th weight of $\Psi[\mathbf{A}, \mathbf{v}]$ is given by $w_i = (\mathbf{v}^\mathsf{T}\mathbf{u}_i)^2$. Thus, the w_i have joint distribution given by,

$$w_i \stackrel{\mathrm{d}}{=} \left(\frac{[\mathbf{x}]_i}{\|\mathbf{x}\|}\right)^2 = \frac{([\mathbf{x}]_i)^2}{([\mathbf{x}]_1)^2 + \dots + ([\mathbf{x}]_n)^2},$$

for $i = 1, \ldots, n$.

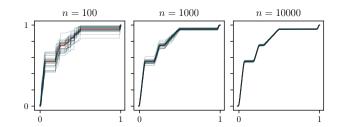


Figure 1. Concentration of 30 independent samples of the weighted CESM $\Psi[\mathbf{A}, \mathbf{v}]$ (—) about the CESM $\Phi[\mathbf{A}]$ (—) for matrices of different sizes constructed with qualitatively similar spectrums. *Remarks*: (i) the light lines are samples of a random variable with expectation given by the dark line, (ii) samples of this random variable define cumulative probability densities, and (iii) this random variable concentrates exponentially about the CESM as n increases.

Recall $m = n\Phi[\mathbf{A}](x)$. Then,

$$\Psi[\mathbf{A}, \mathbf{v}](x) = \sum_{j=1}^{m} w_j \stackrel{\mathrm{d}}{=} \frac{([\mathbf{x}]_1)^2 + \dots + ([\mathbf{x}]_m)^2}{([\mathbf{x}]_1)^2 + \dots + ([\mathbf{x}]_n)^2}.$$

It is well known that for independent chi-square random variables $Y \sim \chi_{\alpha}^2$ and $Z \sim \chi_{\beta}^2$ (see, for example, (Johnson et al., 1994, Section 25.2)),

$$\frac{Y}{Y+Z} \sim \operatorname{Beta}\left(\frac{\alpha}{2}, \frac{\beta}{2}\right).$$

Thus, since $([\mathbf{x}]_1)^2 + \cdots + ([\mathbf{x}]_m)^2$ and $([\mathbf{x}]_{m+1})^2 + \cdots + ([\mathbf{x}]_n)^2$ are independent chi-square random variables with m and n-m degrees of freedom respectively, $\Psi[\mathbf{A}, \mathbf{v}](x)$ is a beta random variable with parameters m/2 and (n-m)/2.

As seen in Figure 1, $\Psi[\mathbf{A}, \mathbf{v}](x)$ concentrates about its mean $\Phi[\mathbf{A}](x)$ as n increases. To understand this more precisely, we introduce the following definition and its consequences.

Definition 4. A random variable X is σ^2 -sub-Gaussian if

$$\mathbb{E}\left[\exp(\lambda(X - \mathbb{E}[X]))\right] \le \exp\left(\frac{\lambda^2 \sigma^2}{2}\right), \ \forall \lambda \in \mathbb{R}.$$

Lemma 2. Suppose X is σ^2 -sub-Gaussian. Let X_1, \ldots, X_{n_x} be iid samples of X. Then for all $t \geq 0$,

$$\mathbb{P}\big[|\langle X_i \rangle - \mathbb{E}[X]| > t\big] \le 2 \exp\left(-\frac{n_{\mathsf{v}}}{2\sigma^2}t^2\right).$$

Theorem 5 (Marchal & Arbel, 2017, Theorem 1). Suppose $X \sim \text{Beta}(\alpha, \beta)$. Then, $\mathbb{E}[X] = \alpha/(\alpha + \beta)$, and X is $(4(\alpha + \beta + 1))^{-1}$ -sub-Gaussian. If $\alpha = \beta$, then there is no smaller σ^2 such that X is σ^2 -sub-Gaussian.

With these results in place, the proof of Theorem 2 is straightforward.

Proof of Theorem 2. First note that the maximums exist because Φ and $\langle \Psi_i \rangle$ are right continuous and piecewise constant except at $\{\lambda_i[\mathbf{A}]\}_{i=1}^n$.

For any x, let $m = m(x) = n\Phi(x)$. Using Lemmas 1 and 2 and Theorem 5 we have that for any x,

$$\begin{split} \mathbb{P}\big[\, |\Phi(x) - \langle \Psi_i(x) \rangle| &> t \, \big] \\ &\leq 2 \exp\left(-\frac{n_{\text{\tiny V}}}{2(4(\frac{m}{2} + \frac{n-m}{2} + 1))^{-1}} t^2 \right). \end{split}$$

We also have

$$\sup_{x \in \mathbb{R}} |\Phi(x) - \langle \Psi_i(x) \rangle|
= \max_{i=1,\dots,n-1} |\Phi(\lambda_i[\mathbf{A}]) - \langle \Psi_i(\lambda_i[\mathbf{A}]) \rangle|.$$

The second result follows by applying a union bound to the events that the maximum is attained at $\lambda_i[\mathbf{A}]$ for each $i=1,\ldots,n$.

4.2. Gaussian quadrature

We now shift our attention to the approximation of the weighted CESM by a Gaussian quadrature rule.

Definition 5. Let μ be a distribution function with finite moments up to degree 2k-1. The k-point Gaussian quadrature rule for μ , is the distribution

$$\nu(x) = \sum_{j=1}^{k} d_j \mathbb{1}[\theta_j \le x]$$

corresponding to nodes $\{\theta_j\}_{j=1}^k$ and weights $\{d_j\}_{j=1}^k$ such that the moments of μ and ν are equal up to degree 2k-1. We denote such a distribution function by $[\mu]_{\nu}^{\text{gq}}$.

This definition implies the total mass of a Gaussian quadrature rule must agree with the original distribution, which in the context of computing approximations to the weighted CESM, means that the SLQ approximation remains a probability distribution function. This property is not retained by other approaches to approximating the weighted CESM such as the KPM. More generally, Proposition 1 asserts that the Wasserstein distance decays inversely with the number of matching moments.

Since, Proposition 1 holds uniformly for all probability distribution functions constant on the complement of [a,b], we also recall an a posteriori characterization of the closeness of distribution functions with matching moments due to (Karlin & Shapley, 1972) but known implicitly far earlier (Stieltjes, 1918). Before stating this theorem, we introduce a definition and a resulting lemma.

Definition 6. A function γ has a sign change at x if there exists x' < x such that $\gamma(x') \neq 0$ and $x = \inf\{t > x' : \gamma(t)\gamma(x') < 0\}$.

Lemma 3. Suppose γ is weakly increasing on an interval (a,b). Then γ and has a sign change at x if and only if there exists x' < x such that $\gamma(x') < 0$, $\gamma(y) \le 0$ for all $y \in (a,x)$ and $\gamma(y) > 0$ for all $y \in (x,b)$.

Theorem 6 (Karlin & Shapley, 1972, Theorem 22.1). Suppose μ and ν are two probability distribution functions constant on the complement of [a,b] whose moments are equal up to degree s. Define $\gamma:[a,b] \to [0,1]$ by $\gamma(x) = \mu(x) - \nu(x)$. Then γ is identically zero or changes sign at least s times.

Note that for a probability distribution function, $[\mu]_k^{\text{gq}}$ is piecewise constant with k points of discontinuity. Using the fact that $[\mu]_k^{\text{gq}}$ and μ share moments up to degree 2k-1 along with Theorem 6, we immediatley obtain the following bounds on $[\mu]_k^{\text{gq}}$ (proved in Supplement A for completeness).

Corollary 1. Suppose μ is a probability distribution function constant on the complement of [a,b]. Let $\{\theta_j\}_{j=1}^k$ and $\{d_j\}_{j=1}^k$ respectively be the nodes and weights of the Guassian quadrature rule $[\mu]_k^{gq}$. Define $[\mu]_k^{\downarrow}$ and $[\mu]_k^{\uparrow}$ by

$$[\mu]_k^{\downarrow}(x) := \sum_{j=1}^{k-1} d_j \mathbb{1} [\theta_{j+1} \le x],$$

$$[\mu]_k^{\uparrow}(x) := d_1 + \sum_{j=2}^k d_j \mathbb{1} [\theta_{j-1} \le x].$$

Then, for all $x \in [a, b]$,

$$[\mu]_k^{\downarrow}(x) \leq \mu(x) \leq [\mu]_k^{\uparrow}(x)$$

In turn, Corollary 1 implies bounds on the Wasserstein and Kolmogorov–Smirnov distances between μ and $[\mu]_k^{gq}$.

Corollary 2. Suppose μ is a probability distribution function constant on the complement of [a,b]. Let $\{\theta_j\}_{j=1}^k$ and $\{d_j\}_{j=1}^k$ respectively be the nodes and weights of the Guassian quadrature rule $[\mu]_{\mathfrak{g}}^{\mathfrak{g}\mathfrak{q}}$. Then

$$\begin{split} d_{\text{KS}}(\mu, [\mu]_k^{\text{gq}}) &\leq \max_{j=1, \dots, k} d_j \\ d_{\text{W}}(\mu, [\mu]_k^{\text{gq}}) &\leq \sum_{j=0}^k \max\{d_j, d_{j+1}\} (\theta_{j+1} - \theta_j) \end{split}$$

where we define $\theta_0 = a$, $\theta_{k+1} = b$, and $d_0 = d_{k+1} = 0$.

Finally, we note the classical result that Lanczos algorithm computes a Gaussian quadrature rule for $\Psi[\mathbf{A}, \mathbf{v}]$ (Gautschi, 2004; Golub & Meurant, 2009).

Proposition 2. Let $[\mathbf{T}]_{:k,:k}$ be the output of Algorithm 2 run on \mathbf{A} , \mathbf{v} for k steps. Then the eigenvalues of $[\mathbf{T}]_{:k,:k}$ and the square of the first components of the eigenvectors of $[\mathbf{T}]_{:k,:k}$ form a degree k Gaussian quadrature rule for μ . That is, $[\Psi[\mathbf{A}, \mathbf{v}]]_k^{gq} = \Psi[[\mathbf{T}]_{:k,:k}, \hat{\mathbf{e}}]$.

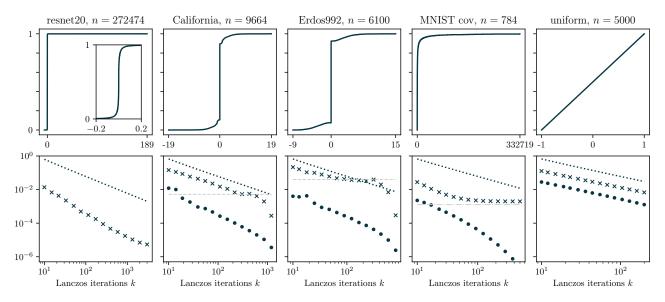


Figure 2. Top: distribution functions Bottom: Wasserstein error and error bounds. All problems are scaled so that $I[\mathbf{A}] = 1$ for easier comparison. From left to right, $n_v = 2, 6, 9, 68, 11$ chosen so that n_v is roughly of size $O(n^{-1})$. Legend: $d_W(\Phi[\mathbf{A}], \langle [\Psi_i]_k^{gq} \rangle)$ (•), bound $\sum_{j=0}^n \max\{[d_i]_j, [d_i]_{j+1}\}([\theta_i]_{j+1} - \theta_i]_j)$ (×), bound $12I[\mathbf{A}](2k-1)^{-1}$ (·····), $(\Phi(d^-) - \Phi(c)) | d-c |$ described in (2) (---), $I[\mathbf{A}]n^{-1}$ (····).

Since k is typically much smaller than n, and since $[\mathbf{T}]_{:k,:k}$ is tridiagonal, then the exact weighted CESM $\Psi[[\mathbf{T}]_{:k,:k}, \hat{\mathbf{e}}]$ can be computed directly. This allows for the efficient computation of Gaussian quadrature rules for $\Psi[\mathbf{A}, \mathbf{v}]$.

4.3. Remaining proofs

Proof of Theorem 1. Note that for any probability distribution functions μ and ν constant on the complement of [a, b],

$$d_{\mathbf{W}}(\mu, \nu) \leq (b - a) d_{\mathbf{KS}}(\mu, \nu).$$

For $i=1,\ldots,n_{\rm v}$, define Ψ_i as in Algorithm 1. Then, using Theorem 2,

$$\mathbb{P}[d_{\mathrm{KS}}(\Phi, \langle \Psi_i \rangle) > t/2] \le 2n \exp(-(n+2)n_{\mathrm{v}}t^2/4),$$

so since $\langle \Psi_i \rangle$ and Φ are constant on the complement of $[\lambda_{\min}[\mathbf{A}], \lambda_{\max}[\mathbf{A}]]$,

$$\mathbb{P}[d_{\mathbf{W}}(\Phi, \langle \Psi_i \rangle) > tI[\mathbf{A}]/2] \le 2n \exp(-n_{\mathbf{v}}(n+2)t^2/4)$$
.

By Proposition 1 and the definition of Gaussian quadrature rule we have, with probability one,

$$d_{\mathbf{W}}([\Psi_i]_k^{\mathrm{gq}}, \Psi_i) < 12I[\mathbf{A}](2k-1)^{-1}$$

for $i = 1, ..., n_v$. Thus, by the triangle inequality, again with probability one,

$$d_{\mathbf{W}}(\langle [\Psi_i]_k^{\mathrm{gq}} \rangle, \langle \Psi_i \rangle) < 12I[\mathbf{A}](2k-1)^{-1}.$$

Finally, we apply the triangle inequality to obtain,

$$d_{\mathbf{W}}(\Phi, \langle [\Psi_i]_k^{\mathrm{gq}} \rangle \le d_{\mathbf{W}}(\Phi, \langle \Psi_i \rangle) + d_{\mathbf{W}}(\langle \Psi_i \rangle, \langle [\Psi_i]_k^{\mathrm{gq}} \rangle)$$

Setting $n_{\rm v} > 4(n+2)^{-1}t^{-2}\log(2n\eta^{-1})$ and $k > 12t^{-1} + \frac{1}{2}$ ensures the sum of the two terms is at most $I[\mathbf{A}]t$ with probability at least $1-\eta$.

Proof of Theorem 3. This is a direct consequence of Corollary 2 and the triangle inequality. \Box

Proof of Theorem 4. Let $\Upsilon(x) = x$ on [0,1] be the probability distribution function for a uniform density on [0,1].

First, for any K, non-negative weights $\{d_i\}_{i=1}^K$ summing to one and ordered points $\{\theta_i\}_{i=1}^K$ in [0,1], define $\{D_i\}_{i=0}^K$ by $D_i = \sum_{j=1}^i d_i$ (where $D_0 = 0$) and consider the functions

$$\varphi(x) = \sum_{i=1}^{K} d_i \mathbb{1}[\theta_i \le x],$$

$$\tilde{\varphi}(y) = \theta_1 + \sum_{i=1}^{K-1} (\theta_{i+1} - \theta_i) \mathbb{1}[D_i \le y].$$

Note that,

$$d_{\mathbf{W}}(\Upsilon,\varphi) = \int_{0}^{1} |\varphi(x) - x| \mathrm{d}x = \int_{0}^{1} |\tilde{\varphi}(y) - y| \mathrm{d}y.$$

Next, define

$$C = \{ \psi : \psi(0) = 0, \psi(1) = 1, \psi'(y) = 1 \,\forall y \in (D_i, D_{i+1}), i = 0, \dots, K \}$$

and observe that, because the contribution on each subinterval is independent of other subintervals,

$$d_{\mathbf{W}}(\Upsilon, \varphi) \ge \min_{\psi \in \mathcal{C}} \int_{0}^{1} |\tilde{\varphi}(y) - \psi(y)| dy$$

$$= \min_{\psi \in \mathcal{C}} \sum_{i=0}^{K-1} \int_{D_{i}}^{D_{i+1}} |\tilde{\varphi}(y) - \psi(y)| dy$$

$$= \sum_{i=0}^{K-1} \min_{\psi \in \mathcal{C}} \int_{D_{i}}^{D_{i+1}} |\tilde{\varphi}(y) - \psi(y)| dy$$

$$= \sum_{i=0}^{K-1} \left(\frac{D_{i+1} - D_{i}}{2}\right)^{2}.$$

Thus, by the Cauchy-Schwarz inequality,

$$\frac{1}{4} = \left(\sum_{i=0}^{K-1} \frac{d_i}{2}\right)^2 \le \left(\sum_{i=0}^{K-1} 1^2\right) \left(\sum_{i=0}^{K-1} \left(\frac{D_{i+1} - D_i}{2}\right)^2\right)$$
$$= K \sum_{i=0}^{K-1} \left(\frac{D_{i+1} - D_i}{2}\right)^2$$

so, since $y \mapsto y \in \mathcal{C}$, we have that

$$\begin{split} d_{\mathbf{W}}(\Upsilon,\varphi) &= \int_{0}^{1} |\tilde{\varphi}(y) - y| \mathrm{d}y \\ &\geq \min_{\psi \in \mathcal{C}} \int_{0}^{1} |\tilde{\varphi}(y) - \psi(y)| \mathrm{d}y \geq \frac{1}{4K}. \end{split}$$

We now construct a matrix whose CESM has small Wasserstein distance to Υ . Let $n = \lceil (4t)^{-1} \rceil$ and define a matrix **A** with eigenvalues $\{(2n)^{-1} + kn^{-1} : k = 0, 1, \dots n - 1\}$. By the above argument, noting the Cauchy–Schwarz inequality is an equality of all terms in the sum are equal, it is clear that,

$$d_{\mathbf{W}}(\Upsilon, \Phi) = \frac{1}{4n} < t.$$

Now, note that $\langle [\Psi_i]_k^{\text{gq}} \rangle$ is of the form of φ with $K = n_{\text{v}}k$. Thus, for any n_{v} , k such that $n_{\text{v}}k < (8t)^{-1}$, with probability one,

$$d_{\mathbf{W}}(\Upsilon, \langle [\Psi_i]_k^{\mathrm{gq}} \rangle) \ge \frac{1}{4n..k} > 2t.$$

Then, using the triangle inequality, again with probability one.

$$d_{\mathcal{W}}(\Phi, \langle [\Psi_i]_k^{gq} \rangle) \ge d_{\mathcal{W}}(\Upsilon, \langle [\Psi_i]_k^{gq} \rangle) - d_{\mathcal{W}}(\Upsilon, \Phi) > t.$$

Since $n_v k$ is the number of matrix vector products required

by Algorithm 1, and $I(\mathbf{A}) < 1$, the result holds.

Note that this proof constructs two distribution functions with matching moments up to degree k whose Wasserstein distance is $\Omega(k^{-1})$. This immediately implies that if the output of an algorithm used to approximate distribution functions depends only on the first k moments, there exist inputs on which the output error has Wasserstein distance $\Omega(k^{-1})$.

5. Numerical verification and discussion

We demonstrate the effectiveness of our bounds on several test problems. The convergence of $\langle \Psi_i \rangle$ to $\Phi[\mathbf{A}]$ is straightforward, so we focus on the convergence of the Gaussian quadrature rules $[\Psi_i]_k^{\mathrm{gq}}$ of Ψ_i .

Here, "resnet20" is a Hessian for the ResNet20 network (He et al., 2016) trained on the Cifar-10 dataset. To apply the Lanczos algorithm to this example, we use a slightly modified version of PyHessian (Yao et al., 2020). The "California" and "Erdos992" examples are graph adjacency matrices from the sparse matrix suite (Davis & Hu, 2011), the "MNIST cov" example is the covariance matrix of the MNIST training data, and "uniform" is a synthetic problem with 5000 eigenvalues evenly spaced between -1 and 1.

Our first example studies the global convergence of $\langle [\Psi_i]_k^{\mathrm{gq}} \rangle$ to $\langle \Psi_i \rangle$ as the number of Lanczos iterations k increases. Specifically, we consider the upper bounds $d_{\mathrm{W}}(\langle \Psi_i \rangle, \langle [\Psi_i]_k^{\mathrm{gq}} \rangle) \leq 12I[\mathbf{A}](2k-1)^{-1}$ from the proof of Theorem 1 and the bound $d_{\mathrm{W}}(\langle \Psi_i \rangle, \langle [\Psi_i]_k^{\mathrm{gq}} \rangle) \leq \langle \sum_{j=0}^k \max\{[d_i]_j, [d_i]_{j+1}\}([\theta_i]_{j+1} - [\theta_i]_j) \rangle$ from Theorem 3. These bounds, with the true extreme eigenvalues of \mathbf{A} used for $[\theta_i]_0$ and $[\theta_i]_{k+1}$ (except for on the "resnet20" example where these terms are omitted), are shown illustrated in Figure 2 for several test problems. Qualitatively, we observe several types of behavior in both the true Wasserstein distance and the bounds.

First, when k is small relative to n, the convergence rate is similar to $O(k^{-1})$. This behavior is especially visible on the "uniform" example, where the true CESM is relatively "smooth" and aligns with the intuition behind our lower bound Theorem 4. On the other hand, as observed on the "MNIST cov" example, when k becomes sufficiently large, the convergence accelerates past $O(k^{-1})$. This is also unsurprising since when k=n, $[\Psi_i]_{i=1}^{\mathbb{R}^d}=\Psi_i$.

Second, we observe the Wasserstein distance bound from Theorem 3 sometimes stagnates. There are two causes for this. The first cause of stagnation, observed on the "MNIST cov" example, is due to the fact that the bound from Theorem 3 will never be smaller than $I[\mathbf{A}]n^{-1}$. The second source of stagnation, due to many tightly clustered eigenvalues, is observed in the "California" and "Erdos992"

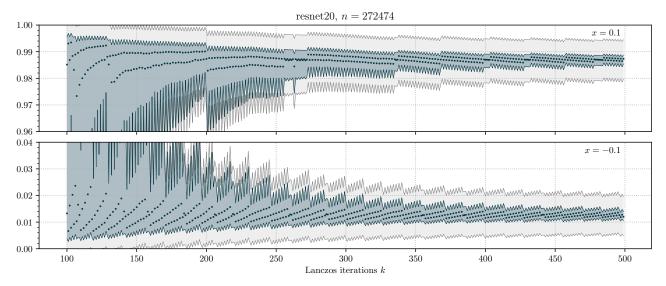


Figure 3. Bounds for $\Phi[\mathbf{A}](x)$ with $\eta=0.01$. The average weighted CESM $\langle \Psi_i(x)\rangle = \langle \Psi(\mathbf{A},\mathbf{v}_i)(x)\rangle$ is bounded between the averaged lower and upper bounds $\langle [\Psi_i]_k^{\downarrow}\rangle$ and $\langle [\Psi_i]_k^{\uparrow}\rangle$ (considered CESM $\langle \Psi_i(x)\rangle = \langle \Psi(\mathbf{A},\mathbf{v}_i)(x)\rangle$ is bounded between the averaged lower and upper bounds $\langle [\Psi_i]_k^{\downarrow}\rangle$ and $\langle [\Psi_i]_k^{\uparrow}\rangle$ (considered CESM with probability at least $1-\eta$, and therefore lies between $\langle [\Psi_i]_k^{\downarrow}(x)\rangle - t$ and $\langle [\Psi_i]_k^{\uparrow}(x)\rangle + t$ (considered CESM) with this same probability. The output of Algorithm 1 $\langle [\Psi_i]_k^{gq}\rangle$ (•) is shown for reference.

examples. To understand this stagnation, suppose there are many eigenvalues clustered near x; i.e. in the interval $(x - \epsilon, x + \epsilon)$ for some small ϵ . Let,

$$c := \max_{\lambda \in \Lambda[\mathbf{A}]} \{\lambda < x - \epsilon\}, \quad d := \min_{\lambda \in \Lambda[\mathbf{A}]} \{\lambda > x + \epsilon\}.$$

so that there are no eigenvalues in the $(c,d) \setminus (x-\epsilon,x+\epsilon)$. Then, if $|d-c| \gg \epsilon$ and k is not too large, the Gaussian quadrature rule will have only one node in (c,d) located very near to x. As a result, the bound will stagnate near

$$(\Phi(d^{-}) - \Phi(c)) |d - c|.$$
 (2)

If the cluster of eigenvalues are all exactly equal so ϵ can be taken to be zero, this stagnation will persist for all k. If they are not, then eventually the Gaussian quadrature rule will place more nodes in this interval and the bound will recover, as observed in both examples. In Supplement B.1, we discuss a heuristic approach to address this stagnation.

Our second example studies bounds for $\Phi[\mathbf{A}](x)$ for a fixed value of x. These can be used to obtain upper and lower bounds for the number of eigenvalues in an interval. Specifically, we consider the lower and upper bounds $\langle [\Psi_i]_k^{\uparrow}(x) \rangle - t$ and $\langle [\Psi_i]_k^{\uparrow}(x) \rangle + t$, which provide probabilistic upper and lower bounds according to Theorem 2 and Corollary 2. In Figure 3, for fixed values of x, we plot the bounds as a function of the number of Lanczos iterations k for the "resnet20" example and as a function of x. Together, these plots imply that with probability 99/100, roughly 94-99% of the eigenvalues are in the interval [-0.1, 0.1].

6. Outlook

The analysis in this paper gives rigorous bounds on the accuracy of SLQ for spectrum approximation. These bounds are suited to the parameter ranges encountered in practice and demonstrate that SLQ is a viable method for spectrum approximation in many applications. As a result, we hope our analysis will allow practitioners to obtain more precise and theoretically justifiable insights about their applications without the need for heuristics.

More broadly, SLQ and KPM fall into a larger class of spectrum approximation algorithms which approximate iid samples of the weighted CESM using information from Krylov subspaces. However, the exact relationship between these algorithms has not been fully described, making the tradeoffs between the algorithms murky at best. In order to shed light on the tradeoffs between these algorithms, a comprehensive treatment providing a *unified* perspective is needed.

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A. Omitted proofs

In this section, we provide several of the proofs omitted in the main text.

A.1. Proof of results

We provide proofs of Corollaries 1 and 2. These both follow from Theorem 6.

Corollary 1. Suppose μ is a probability distribution function constant on the complement of [a,b]. Let $\{\theta_j\}_{j=1}^k$ and $\{d_j\}_{j=1}^k$ respectively be the nodes and weights of the Guassian quadrature rule $[\mu]_k^{gq}$. Define $[\mu]_k^{\downarrow}$ and $[\mu]_k^{\uparrow}$ by

$$[\mu]_k^{\downarrow}(x) := \sum_{j=1}^{k-1} d_j \mathbb{1} [\theta_{j+1} \le x],$$

$$[\mu]_k^{\uparrow}(x) := d_1 + \sum_{j=2}^k d_j \mathbb{1} [\theta_{j-1} \le x].$$

Then, for all $x \in [a, b]$,

$$[\mu]_k^{\downarrow}(x) \le \mu(x) \le [\mu]_k^{\uparrow}(x).$$

Proof. Suppose $[\mu]_k^{\text{gq}} \neq \mu$ and define $\gamma(x) = \mu(x) - [\mu]_k^{\text{gq}}(x)$. Observe that for any $j = 1, \ldots, k-1$, $[\mu]_k^{\text{gq}}$ is constant on (θ_j, θ_{j+1}) , so γ is weakly increasing on this interval. Lemma 3 states that if γ change signs at some point $y_j \in (\theta_j, \theta_{j+1})$ then $\gamma(x) > 0$ for all $x \in (y_j, \theta_{j+1})$, $j = 1, \ldots, k-1$ and $\gamma(x) \leq 0$ for all $x \in (\theta_j, y_j)$, so γ cannot change signs at any other point in (θ_j, θ_{j+1}) .

Observe further that on (a, θ_1) , $[\mu]_k^{\mathrm{gq}}(x) = 0 \le \mu(x)$ so $\gamma(x) \ge 0$, and on (θ_n, b) , $[\mu]_k^{\mathrm{gq}}(x) = 1 \ge \mu(x)$ so $\gamma(x) \le 0$. Thus, by Lemma 3, no sign changes can occur on these intervals.

As a result, the only possible locations for sign changes of γ on (a,b) are $\{\theta_j\}_{j=1}^k$ and $\{y_j\}_{j=1}^{k-1}$. This is exactly 2k-1 possible sign changes, so by Theorem 6, a sign change must occur at each of these points. In particular, since γ has a sign change at θ_j ,

$$[\mu]_k^{\mathrm{gq}}(\theta_j^-) \le \mu(\theta_j) \le [\theta_j]_k^{\mathrm{gq}}.$$

Therefore, for $x \in (\theta_i, \theta_{i+1})$,

$$[\mu]_{k}^{\downarrow}(x) = [\mu]_{k}^{gq}(\theta_{i}^{-}) \leq \mu(\theta_{i}) \leq \mu(x) \leq \mu(\theta_{i+1}) \leq [\mu]_{k}^{gq}(\theta_{i+1}) = [\mu]_{k}^{\uparrow}(x).$$

Corollary 2. Suppose μ is a probability distribution function constant on the complement of [a,b]. Let $\{\theta_j\}_{j=1}^k$ and $\{d_j\}_{j=1}^k$ respectively be the nodes and weights of the Guassian quadrature rule $[\mu]_k^{gq}$. Then

$$d_{KS}(\mu, [\mu]_k^{gq}) \le \max_{j=1,...,k} d_j$$

$$d_{W}(\mu, [\mu]_k^{gq}) \le \sum_{j=0}^k \max\{d_j, d_{j+1}\} (\theta_{j+1} - \theta_j)$$

where we define $\theta_0 = a$, $\theta_{k+1} = b$, and $d_0 = d_{k+1} = 0$.

Proof. Note that

$$|\mu(x) - [x]_k^{gq}| \le \max\{|[\mu]_k^{gq}(x) - [\mu]_k^{\uparrow}(x)|, |[\mu]_k^{gq}(x) - [\mu]_k^{\downarrow}(x)|\}$$

= \text{max}\{d_j, d_{j+1}\}\mathbf{1}[x \in [\theta_j, \theta_{j+1})].

Thus,

$$d_{KS}(\mu, [\mu]_k^{gq}) = \sup_x |\mu(x) - [x]_k^{gq}| \le \max_{j=1,\dots,k} d_j.$$

and

$$d_{\mathbf{W}}(\mu, [\mu]_{k}^{\mathrm{gq}}) = \int_{a}^{b} |\mu(s) - [\mu]_{k}^{\mathrm{gq}}(s)| \mathrm{d}s \le \sum_{j=0}^{k} \max\{d_{j}, d_{j+1}\}(\theta_{j+1} - \theta_{j}).$$

A.2. Definitions of random variables

Definition 7. The beta distribution with parameters α, β , denoted Beta (α, β) , is defined by density function on [0, 1],

$$x \mapsto \frac{\Gamma(\alpha+\beta)}{\Gamma(\alpha)\Gamma(\beta)} x^{\alpha-1} (1-x)^{\beta-1}.$$

Definition 8. The chi-square distribution with positive integer parameter k, denoted χ_k^2 , is defined by density function on $[0,\infty)$,

$$x \mapsto \frac{1}{2^{k/2}\Gamma(k/2)} x^{k/2-1} \exp(-x/2).$$

A.3. Other proofs

For reader convenience, we also provide proofs for several of the technical results cited throughout the main portion of the text

To begin, we provide a standard moment generating function argument to bound the tails of random variables.

Lemma 2. Suppose X is σ^2 -sub-Gaussian. Let X_1, \ldots, X_{n_v} be iid samples of X. Then for all $t \geq 0$,

$$\mathbb{P}\big[|\langle X_i \rangle - \mathbb{E}[X]| > t\big] \le 2 \exp\left(-\frac{n_{\text{v}}}{2\sigma^2}t^2\right).$$

Proof. WLOG assume $\mathbb{E}[X] = 0$.

$$\begin{split} \mathbb{P}[n_{\mathbf{v}}\langle X_{i}\rangle \geq n_{\mathbf{v}}t] &= \mathbb{P}[\exp(\lambda n_{\mathbf{v}}\langle X_{i}\rangle) \geq \exp(\lambda n_{\mathbf{v}}t)] \\ &\leq \exp(-n_{\mathbf{v}}\lambda t)\mathbb{E}[\exp(\lambda n_{\mathbf{v}}\langle X_{i}\rangle)] \\ &= \exp(-n_{\mathbf{v}}\lambda t)\mathbb{E}[\exp(\lambda X)]^{n_{\mathbf{v}}} \\ &\leq \exp(-n_{\mathbf{v}}\lambda t)\exp(n_{\mathbf{v}}\lambda^{2}\sigma^{2}/2) \\ &= \exp(-n_{\mathbf{v}}\lambda t + n_{\mathbf{v}}\lambda^{2}\sigma^{2}/2). \end{split} \tag{Sub-Gaussian}$$

This expression is minimized when $\lambda = t/\sigma^2$ from which we obtain,

$$\mathbb{P}[\langle X_i \rangle \ge t] \le \exp\left(-\frac{n_{\text{v}}}{2\sigma^2}t^2\right).$$

Proposition 1. Suppose μ and ν are two probability distribution functions constant on the complement of [a,b] whose moments are equal up to degree s. Then,

$$d_{\mathbf{W}}(\mu,\nu) \le 2(b-a)(1+\pi^2/2)s^{-1} < 12(b-a)s^{-1}.$$

Proof. Define $f:[a,b]\to\mathbb{R}$ by

$$f(x) = -\int_{a}^{x} \operatorname{sign}(\mu(s) - \nu(s)) ds.$$

Then, f is continuous and piecewise linear with slope ± 1 , and as a result, f is of bounded variation and $|f(y)-f(z)| \le |y-z|$ for all $y, z \in [a, b]$. Moreover, since μ and ν are weakly increasing bounded functions, they are each of bounded variation, as is the difference $\mu - \nu$. Therefore, we can therefore integrate by parts over the closed interval [a, b], see for instance

(Folland, 1999, Theorem 3.36), to obtain

$$\begin{split} d_{\mathbf{W}}(\mu,\nu) &= \int_{a}^{b} |\mu(s) - \nu(s)| \mathrm{d}s = \int_{a}^{b} \mathrm{sign}(\mu(s) - \nu(s))(\mu(s) - \nu(s)) \mathrm{d}s \\ &= \left[-f(s)(\mu(s) - \nu(s)) \right]_{a^{-}}^{b} + \int_{a}^{b} f(s) \mathrm{d}(\mu(s) - \nu(s)) \\ &= \int_{a}^{b} f(s) \mathrm{d}(\mu(s) - \nu(s)). \end{split}$$

Next, note that for any polynomial p of degree < k,

$$\int_{a}^{b} f(s) d(\mu(s) - \nu(s)) = \int_{a}^{b} (f(s) - p(s)) d(\mu(s) - \nu(s))$$

$$\leq \int_{a}^{b} \|f - p\|_{\infty} |d\mu(s) - \nu(s)|$$

$$\leq 2\|f - p\|_{\infty}.$$

Finally, Jackson's theorem, see for example (Han & Atkinson, 2009, Theorem 3.7.2), asserts that if f satisfies $|f(y)-f(z)| \le |y-z|$ for all $y,z \in [a,b]$, then over [a,b],

$$\min_{\deg(p) \le s} \|f - p\|_{\infty} \le (b - a)(1 + \pi^2/2)s^{-1}.$$

Combining these statements and minimizing over polynomials p of degree $\leq s$ yields the result.

Lemma 3. Suppose γ is weakly increasing on an interval (a,b). Then γ and has a sign change at x if and only if there exists x' < x such that $\gamma(x') < 0$, $\gamma(y) \le 0$ for all $y \in (a,x)$ and $\gamma(y) > 0$ for all $y \in (x,b)$.

Proof. Suppose that γ has a sign change at x and let x' be as in the definition. Then, for any t>x', since γ is weakly increasing, $\gamma(x') \leq \gamma(t)$. This means that $\gamma(t)\gamma(x') < 0$ implies $\gamma(x') < 0$ and $\gamma(t) > 0$. Since $x = \inf\{t > x' : \gamma(t)\gamma(x') < 0\}$, we have $\gamma(y) > 0$ for all $y \in (x,b)$ and $\gamma(y) \leq 0$ for all $y \in (x',x)$ and $\gamma(x') < 0$

The reverse direction follows directly from the definition of sign change.

Finally, we recall the proof of Theorem 6.

Theorem 6 (Karlin & Shapley, 1972, Theorem 22.1). Suppose μ and ν are two probability distribution functions constant on the complement of [a,b] whose moments are equal up to degree s. Define $\gamma:[a,b]\to [0,1]$ by $\gamma(x)=\mu(x)-\nu(x)$. Then γ is identically zero or changes sign at least s times.

Proof. Suppose $\gamma \neq 0$. For the sake of contradiction, suppose also that γ has fewer than s sign changes. Then, there exists a degree at most s-1 polynomial r such that for all $x \in [a,b]$, $r(x)\gamma(x) \geq 0$; i.e. pick r to have a sign change at every sign change of γ .

Thus, since $\gamma \neq 0$ is right continuous and r is continuous,

$$\int_{a}^{b} r(x)\gamma(x)\mathrm{d}x > 0.$$

Let R be an antiderivative of r. Then, by integrating by parts over the closed interval [a, b],

$$\int_{a}^{b} r(x)\gamma(x)dx = \left[R(x)\gamma(x)\right]_{a^{-}}^{b} - \int_{a}^{b} R(x)d\gamma(x).$$

Since μ and ν are equal on the compliment of [a, b],

$$[R(x)\gamma(x)]_a^b = R(b)(\mu(b) - \nu(b)) - R(a)(\mu(a^-) - \nu(a^-)) = 0$$

and, since μ and ν share moments up to degree s,

$$\int_a^b R(x)d\gamma(x) = \int_a^b R(x)(d\mu(x) - d\nu(x)) = 0.$$

This contradicts the earlier assertion that this integral is non-zero.

B. Additional numerical experiments

In Figure 4 we illustrate the bounds on $\Phi[\mathbf{A}](x)$ for varying x and fixed k. These are the same quantities as shown in Figure 3, which showed them for fixed x and varying k.

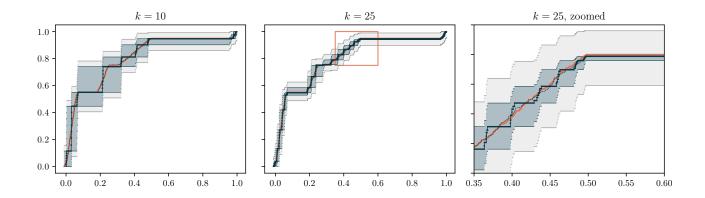


Figure 4. Bounds for $\Phi[\mathbf{A}](x)$ () with $n_v = 7$. The average weighted CESM $\langle \Psi_i(x) \rangle = \langle \Psi(\mathbf{A}, \mathbf{v}_i)(x) \rangle$ is bounded between the averaged lower and upper bounds $\langle [\Psi_i]_k^{\downarrow} \rangle$ and $\langle [\Psi_i]_k^{\uparrow} \rangle$ () from Corollary 1. By Theorem 2, the CESM $\Phi[\mathbf{A}](x)$ is within t of the average weighted CESM with probability at least $1 - \eta$, and therefore lies betwen $\langle [\Psi_i]_k^{\downarrow}(x) \rangle - t$ and $\langle [\Psi_i]_k^{\uparrow}(x) \rangle + t$ () with this same probability. The output of Algorithm 1 $\langle [\Psi_i]_k^{gq} \rangle$ () is shown for reference.

We make several remarks. First, observe that both the average Gaussian quadrature rule $\langle [\Psi_i]_k^{\mathfrak{gq}} \rangle$ and the upper and lower bounds $\langle [\Psi_i]_k^{\uparrow} \rangle$ and $\langle [\Psi_i]_k^{\downarrow} \rangle$ are piecewise constant with kn_v points of discontinuity. Importantly, however, the points of discontinuity corresponding to different samples concentrate when k is relatively small compared to n. This is most visible in the k=10 plot, which at first glance, appears to only have 10 points of discontinuity. This concentration is essentially due to the fact that the weighted CESMs Ψ_i concentrate about $\Phi[\mathbf{A}]$, so moments and therefore the Gaussian quadrature nodes and weights also concentrate. For further discussions on this behavior we refer readers to (Kuijlaars, 2000; Garza-Vargas & Kulkarni, 2020). Second, observe that the CESM $\Phi[\mathbf{A}](x)$ is not necessarily contained between $\langle [\Phi_i]_k^{\downarrow}(x) \rangle$ and $\langle [\Phi_i]_k^{\uparrow}(x) \rangle$. Rather, the average weighted CESM $\langle \Phi_i \rangle$ is containe between these values, and $\Phi[\mathbf{A}](x)$ is within t of $\langle \Phi_i \rangle$ with some probability η .

In Figure 5 we illustrate the same bounds as Figure 3 along with a plot of the eigenvalues of $[\mathbf{T}_i]_{:k,:k}$. We make several remarks. First, concentration of the Gaussian quadrature nodes is clearly visible. Second, the position of these nodes as the iterations change provide insight into the sawtooth behavior of the bounds on $\langle [\Psi_i]_k^{\mathrm{gq}} \rangle$. Moreover, we observe that the value of $\langle [\Psi_i]_k^{\mathrm{gq}}(x) \rangle$ is very near to the limiting value of $\langle [\Psi_i]_k^{\mathrm{gq}}(x) \rangle$ (which we estimate by computing $\langle [\Psi_i]_{3000}^{\mathrm{gq}}(x) \rangle$) whenever there are Guassian quadrature nodes near to x. Further study of this phenomena may yield useful heuristic CESM estimation.

B.1. Added points

In Section 5, we remarked that on spectrums with many clustered eigenvalues, our bounds sometimes encounter plateaus. In this section, we discuss a heuristic approach to address this issue by introducing eigenvalues with known weights to \mathbf{A} .

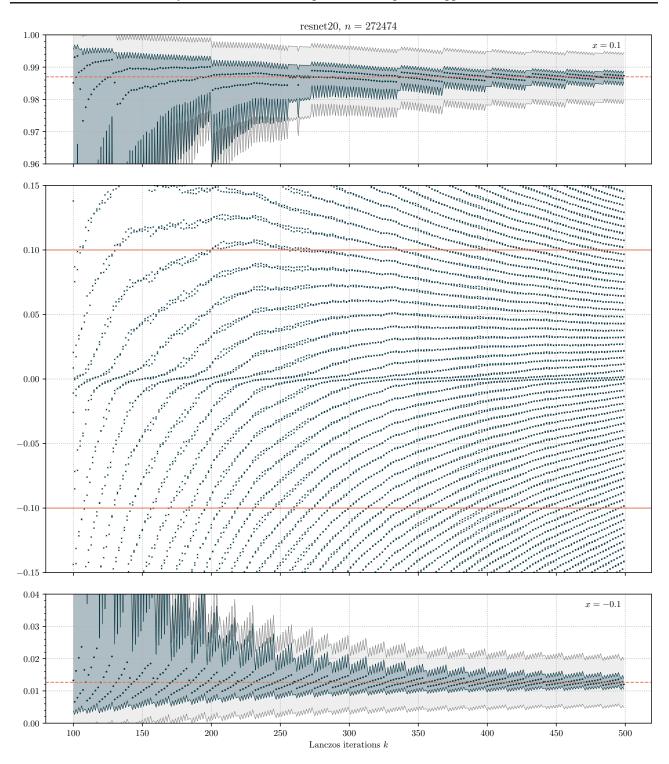
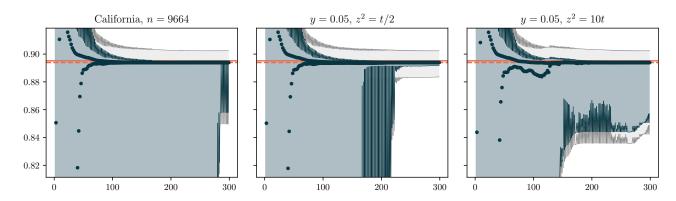


Figure 5. Top/Bottom: Bounds for $\Phi[\mathbf{A}](x)$ when $x=\pm-.01$ and $n_v=2$. The average weighted CESM $\langle \Psi_i(x) \rangle = \langle \Psi(\mathbf{A},\mathbf{v}_i)(x) \rangle$ is bounded between the averaged lower and upper bounds $\langle [\Psi_i]_k^{\downarrow} \rangle$ and $\langle [\Psi_i]_k^{\uparrow} \rangle$ ((\blacksquare) from Corollary 1. By Theorem 2, the CESM $\Phi[\mathbf{A}](x)$ is within t of the average weighted CESM with probability at least $1-\eta$, and therefore lies betwen $\langle [\Psi_i]_k^{\downarrow}(x) \rangle - t$ and $\langle [\Psi_i]_k^{\uparrow}(x) \rangle + t$ ((\blacksquare)) with this same probability. The output of Algorithm 1 $\langle [\Psi_i]_k^{gq} \rangle$ (\bullet) is shown for reference. Middle: Eigenvalues of $[\mathbf{T}_i]_{:k,:k}$ for i=1,2. Remarks: In the middle plot, the eigenvalues of $[\mathbf{T}_i]_{:k,:k}$ (\times) are very close for both samples. This aligns with the expected concentration. We also observe that when there is an eigenvalues of $[\mathbf{T}_i]_{:k,:k}$ near the value of x of interest (\blacksquare), $\langle [\Psi_i(x)]_k^{gq} \rangle \rangle$ is near its limiting value $\langle \Psi_i(x) \rangle$ (estimated by $\langle [\Psi_i]_{3000}^{gq} \rangle$ (\bullet).



Towards this end, define,

$$ar{\mathbf{A}} = egin{bmatrix} \mathbf{A} & & & \\ & y \end{bmatrix}, & ar{\mathbf{v}}_i = egin{bmatrix} \mathbf{v}_i & & & \\ z & & & \end{bmatrix}$$

for scalars y and z. By increasing the value of z, we can introduce an Gaussian quadrature node near y. Thus, we may introduce nodes near to locations suspected of having large jumps in the spectrum. Often, the origin is such a point, since many matrices may be low rank or close to low rank.

As before, we have that,

$$\langle \Psi[\bar{\mathbf{A}}, \bar{\mathbf{v}}_i](x) \rangle \ge \langle [\Psi(\bar{\mathbf{A}}, \bar{\mathbf{v}}_i)]_k^{\downarrow}(x) \rangle$$

$$\langle \Psi[\bar{\mathbf{A}}, \bar{\mathbf{v}}_i](x) \rangle \le \langle [\Psi(\bar{\mathbf{A}}, \bar{\mathbf{v}}_i)]_k^{\uparrow}(x) \rangle$$
(3)

Similarly, $\Psi(\bar{\mathbf{A}}, \bar{\mathbf{v}}_i)$ will concentrate strongly around its mean. Note, however, that the mean is not $\Phi(\bar{\mathbf{A}})$, since the weight on y is z^2 rather than 1/(n+1). However, we do have that with probability at least $1-2\exp(-n_v(n+2)t^{-2})$,

$$\mathbb{E}[\Psi[\bar{\mathbf{A}}, \bar{\mathbf{v}}_i](x)] \ge \langle [\Psi[\bar{\mathbf{A}}, \bar{\mathbf{v}}_i]]_k^{\downarrow}(x) \rangle - t$$

$$\mathbb{E}[\Psi[\bar{\mathbf{A}}, \bar{\mathbf{v}}_i](x)] \le \langle [\Psi[\bar{\mathbf{A}}, \bar{\mathbf{v}}_i]]_h^{\downarrow}(x) \rangle + t$$

Note that

$$\Phi[\mathbf{A}](x) = \mathbb{E}[\Psi[\bar{\mathbf{A}}, \bar{\mathbf{v}}_i](x)] - z^2 \mathbb{1}[y \le x].$$

so

$$\Phi[\mathbf{A}](x) \ge \langle [\Psi[\bar{\mathbf{A}}, \bar{\mathbf{v}}_i]]_k^{\uparrow}(x) \rangle - z^2 \mathbb{1}[y \le x] - t$$

$$\Phi(\mathbf{A})(x) \le \langle [\Psi[\bar{\mathbf{A}}, \bar{\mathbf{v}}_i]]_k^{\uparrow}(x) \rangle - z^2 \mathbb{1}[y \le x] + t.$$
(4)

In Figure 6, we illustrate the effect of introducing a node at 0.05, between the large cluster of eigenvalues at the origin, and our point of evaluation, x = 0.1. The larger the value of z, the earlier in the iteration the Gaussian quadrature node near y appears. However, the larger value of z also weakens the quality of the lower bounds.

C. Gaussian quadrature convergence for smooth and analytic functions

Most existing analysis for SLQ for spectrum approximation has been in terms of smoothed approximations to the empertical spectral measure (ESM) $d\Phi[\mathbf{A}](x)/dx$ or the CESM $\Phi[\mathbf{A}](x)$ of \mathbf{A} . Such smoothed approximations can be obtained by the

integral of certain functions against $d\Phi[\mathbf{A}](x)$ (Lin et al., 2016; Han et al., 2017; Ghorbani et al., 2019). Specifically, if v approximates a delta function and Υ approximates a step function, then

$$\frac{\mathrm{d}\Phi[\mathbf{A}](x)}{\mathrm{d}x} \approx \int \upsilon(x) \mathrm{d}\Phi[\mathbf{A}](x), \qquad \qquad \Phi[\mathbf{A}](x) \approx \int \Upsilon(x) \mathrm{d}\Phi[\mathbf{A}](x).$$

Defining $\Psi_i = \Psi(\mathbf{A}, \mathbf{v}_i)$ for $\{\mathbf{v}_i\} \stackrel{\text{iid}}{\sim} \mathcal{U}(\mathcal{S}^{n-1})$, the smoothed ESM/CESM can therefore respectively be approximated by

$$\left\langle \int \upsilon(x) \mathrm{d}[\Psi_i]_k^{\mathrm{gq}}(x) \right\rangle, \qquad \left\langle \int \upsilon(x) \mathrm{d}[\Psi_i]_k^{\mathrm{gq}}(x) \right\rangle.$$

If these v and Υ are smooth or analytic, the following well known bounds apply.

Theorem 7 (Golub & Meurant, 2009, Section 6.2). Let μ be a probability distribution function constant on the compliment of [a,b] and let $p_k(s)$ be the k-th monic orthogonal polynomial of μ . Then, if f is 2k times differentiable, for some $\xi \in [a,b]$,

$$\int f(s) d\mu(s) - \int f(s) d[\mu]_k^{gq}(s) = \frac{f^{(2k)}(\xi)}{(2k)!} \int \prod_{i=1}^k p_k(s)^2 d\mu(s).$$

Theorem 8 (Ubaru et al., 2017a, Theorem 4.2). Let a function f be analytic in [-1,1] and analtically continuable in the open Bernstein ellipse E_{ρ} with foci ± 1 and sum of major and minor axis equal to $\rho > 1$, where it satisfies $|g(z)| \leq M_{\rho}$. Then, the k-step Lanczos quadrature satisfies,

$$\left| \int f(s) d\Psi(s) - \int f(s) d[\Psi]_k^{gq}(s) \right| \le \frac{4M_\rho}{(\rho^2 - 1)\rho^{2(k-1)}}$$

where $\Psi = \Psi[\mathbf{A}, \mathbf{v}]$ for some \mathbf{A} with eigenvalues in [-1, 1] and a possibly deterministic right hand size \mathbf{v} .

At first glance, Theorems 7 and 8 suggest much faster convergence than Theorem 1. However, in practice, constants hidden by big-O notation are important. Typical choices for v, such as Gaussian densities, grow rapidly away from the real axis, meaning M_{ρ} grows as ρ becomes larger. Similarly, the width of the interval over which Υ goes from being near 0 to being near 1 is closely related to ρ and M_{ρ} . Indeed, in order to closely approximate a discontinuous function, the analytic approximations are necessarily poorly behaved in the complex plane. This typically manifests as singularities which approach the real axis as the width over which the jump occurs decreases, which in turn, forces $\rho \to 1$ and $M_{\rho}/(\rho^2-1) \to \infty$.

As a result, even though SLQ approximations to the smoothed EMS/CESM have exponential convergence, the number of iterations predicted by the bounds is typically not useful for the smoothing parameters used in practice. As an explicit example, suppose $v(s) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp(-(s-x)^2/2\sigma^2)$ is a Gaussian density function, and that we wish to evaluate the smoothed ESM at x=0. Then, using that $i(\rho+\rho^{-1})/2$ is the uppermost point of E_ρ ,

$$M_{\rho} \ge \left| \upsilon \left(i \frac{\rho + \rho^{-1}}{2} \right) \right| \ge \left| \upsilon \left(i \frac{\rho}{2} \right) \right| = \frac{1}{\sqrt{2\pi\sigma^2}} \exp \left(\frac{\rho^2}{8\sigma^2} \right).$$

Thus, to reduce the bound from Theorem 8 to a fraction c of the maximum possible value of the smoothed density, v(0), we require

$$\frac{4M_{\rho}}{(\rho^2 - 1)\rho^{2(k-1)}} < c\upsilon(0).$$

Solving for k we have,

$$k > 1 + \frac{\ln\left(\frac{4M_{\rho}}{cv(0)(\rho^2 - 1)}\right)}{2\ln(\rho/2)} = 1 + \frac{\rho^2/8\sigma^2 + \ln(4) - \ln(c(\rho^2 - 1))}{2\ln(\rho)}$$

If $\sigma^2 = 10^{-5}$ as suggested in (Ghorbani et al., 2019), then k must be exceedingly large for the bound to provide any meaningful convergence gurantee; even for an extremely weak bound of c = 1/2, we require k > 33900 iterations. On the other hand, our bounds, while only inversely dependent on k, are applicable to the parameter ranges found in practice; i.e. for a small number of Lanczos iterations.

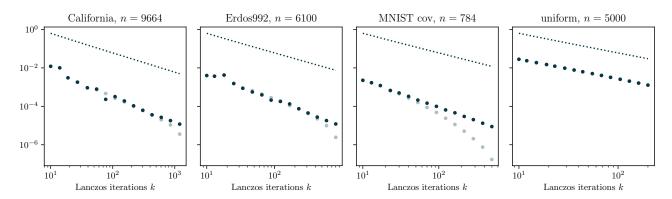


Figure 7. Wasserstein error $d_W(\Phi[\mathbf{A}], \langle [\Psi_i]_k^{gq} \rangle$ (in finite preicsion without reorthogonalization) (•). All problems are scaled so that $I(\mathbf{A}) = 1$ for easier comparison. For comparison, the errors with reorthogonalization (•) from Figure 2 and the bound $12I[\mathbf{A}](2k-1)^{-1}(\cdots)$, are also included.

D. The Lanczos algorithm in finite precision

In finite precision, Lanczos may behave very differently than in exact arithmetic (Greenbaum, 1997; Meurant & Strakoš, 2006). The most visible effects are a loss of orthogonality in the basis vectors $[\mathbf{Q}]_{:,:k}$ and the appearence of multiple tightly clustered eigenvalues in $[\mathbf{T}]_{:k,:k}$ all approximating isolated eigenvalues of \mathbf{A} (in exact arithmetic, each eigenvalue of \mathbf{A} would be approximated by at most one eigenvalue of $[\mathbf{T}]_{:k,:k}$). This is due primarily to a loss of orthogonality in the basis vectors $\{\mathbf{q}_i\}_{i=1}^k$ caused by orthogonalizing $\tilde{\mathbf{q}}_{i+1}$ against only \mathbf{q}_i and \mathbf{q}_{i-1} which occurs precisely when an eigenvalue of $[\mathbf{T}]_{:k,:k}$ has converged to an eigenvalue of \mathbf{A} (Paige, 1976; 1980).

The most straightforward solution is to reorthogonalize against all previous basis vectors at each step. This maintains orthogonality to near machine precision, albeit at the cost of increasing the runtime to $O(k(T_{\rm mv}+n+k))$ and the required storage to O(nk). Other more computationally efficient schemes, which orthogonalize selevatively against the basis vectors corresponding to converged eigenvalues of $[T]_{:k,:k}$ have also been studied (Parlet & Scott, 1979).

In the context of iterative algorithms, the repeated eigenvalues found by Lanczos often manifests as a delay of convergence; i.e. and increase in the number of iterations required to reach a given accuracy. This behavior is observed in SLQ and illustrated in Figure 7. Note that the delay is worse on problems where there are isolated eigenvalues, especically in the upper spectrum. This is because, in exact arithmetic, Lanczos will quickly find these eigenvalues, orthogonality will be lost, and multiple iterations will be wasted finding repeated copies of the eigenvalue. On problems such as "uniform", where there are no isolated eigenvalues and $k \ll n$, Lanczos behaves nearly identically in finite precision and exact arithmetic.

As noted by (Papyan, 2019), even in finite precision SLQ still seems to converge. We now give a high level explanation for this phenomenon.

Under certain conditions, (Greenbaum, 1989) shows that the matrix $[T]_{:k,:k}$ obtained by an implementation of the Lanczos algorithm run in finite precision can be viewed as the output the Lanczos algorithm run in exact arithmetic on a certain "nearby" problem. Loosely speaking, the necessary conditions for the analysis of (Greenbaum, 1989) to apply are that for all $i \le k$,

- (i) the finite precision Lanczos vectors are nearly unit length: $\|\mathbf{q}_i\| \approx 1$
- (ii) that they are close locally orthogonal: $\beta_i \mathbf{q}_i^\mathsf{T} \mathbf{q}_{i-1} \approx 1$
- (iii) that they approximately satisfy the three term recurrence: $\mathbf{A}\mathbf{q}_i \approx \beta_i \mathbf{q}_{i+1} + \alpha_i \mathbf{q}_i + \beta_{i-1} \mathbf{q}_{i-1}$.

Such conditions are satisfied by Algorithm 2 (without reorthogonalization) in finite precision (Paige, 1976; 1980).

If these conditions are met, (Greenbaum, 1989) shows that there exists a $N \times N$ matrix $\bar{\mathbf{A}}$ and vector $\bar{\mathbf{v}}$ such that Lanczos run on $\bar{\mathbf{A}}$, $\bar{\mathbf{v}}$ in exact arithmetic for k steps produces $[\mathbf{T}]_{:k,:k}$ and, that

(i) Eigenvalues of $\bar{\mathbf{A}}$ clustered near to those of \mathbf{A} : for any $j \in \{1, \dots, N\}$, there exists $i \in \{1, \dots, n\}$ such that

$$\lambda_i(\bar{\mathbf{A}}) \approx \lambda_i(\mathbf{A}).$$

(ii) The sum of square of first components of eigenvectors of clusters of $\bar{\bf A}$ are near to the square of the projections of ${\bf v}$ onto the eigenvectors of ${\bf A}$: for an eigenvalue $\lambda_i({\bf A})$

$$w_i \approx \sum_{j \in S} \bar{w}_j$$

where S_i is the set of indices such that $\lambda_j(\bar{\mathbf{A}}) \approx \lambda_i(\mathbf{A})$ for all $j \in S$.

Together, these conditions imply that

$$\Psi[\mathbf{A}, \mathbf{v}] \approx \Psi(\bar{\mathbf{A}}, \bar{\mathbf{v}}). \tag{5}$$

Since the $[\mathbf{T}]_{:k,:k}$ produced by the finite precision computation corresponds to an exact Gaussian quadrature rule for $\Psi(\bar{\mathbf{A}}, \bar{\mathbf{v}})$, the theory developed in this paper applies to the exact computation with $\bar{\mathbf{A}}$ and $\bar{\mathbf{v}}$. Of course, what the " \approx " in (5) means depend on the details of both the analysis in (Greenbaum, 1989) and implementation of the Lanczos algorithm used. If, for instance, the Wasserstein distance between $\Psi[\mathbf{A}, \mathbf{v}]$ and $\Psi(\bar{\mathbf{A}}, \bar{\mathbf{v}})$ is not reasonably small for the precision used, then knowing that the $[\mathbf{T}]_{:k::k}$ corresponds to an exact computation is with $\bar{\mathbf{A}}, \bar{\mathbf{v}}$ is not useful in terms of our bounds.