APPROXIMATING SPECTRAL DENSITIES OF LARGE MATRICES

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Abstract. In physics, it is sometimes desirable to compute the so-called Density Of States (DOS), also known as the spectral density, of a Hermitian (or symmetric) matrix $A$. The spectral density can be viewed as a probability density distribution that measures the likelihood of finding eigenvalues near some point on the real line. The most straightforward way to obtain this density is to compute all eigenvalues of $A$. But this approach is generally costly and wasteful, especially for matrices of large dimension. There exist alternative methods that allow us to estimate the spectral density function at much lower cost. The major computational cost of these methods is in multiplying $A$ with a number of vectors, which makes them appealing for large-scale problems where products of the matrix $A$ with arbitrary vectors are inexpensive. This paper defines the problem of estimating the spectral density carefully. It then surveys a few known methods for estimating the spectral density, and proposes some new variations of existing methods. All methods are discussed from a numerical linear algebra point of view.

Key words. spectral density, density of states, large scale sparse matrix, approximation of distribution, quantum mechanics

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1. Introduction. Given an $n \times n$ Hermitian (sparse real symmetric in what follows) matrix $A$, scientists in various disciplines often want to compute its Density Of States (DOS), or spectral density. Formally, the DOS is defined as

$$\phi(t) = \frac{1}{n} \sum_{j=1}^{n} \delta(t - \lambda_j),$$

(1.1)

where $\delta$ is the Dirac $\delta$-function or Dirac distribution, and the $\lambda_j$’s are the eigenvalues of $A$, assumed here to be labeled increasingly. We also denote by $u_j$ the eigenvector corresponding to $\lambda_j$, so $A$ has the eigen-decomposition

$$A = \sum_{j=1}^{n} \lambda_j u_j u_j^T.$$  (1.2)

The number of eigenvalues in an interval $[a,b]$ can be expressed as

$$\eta_{[a,b]} = \int_{a}^{b} \sum_{j} \delta(t - \lambda_j) \, dt = \int_{a}^{b} n\phi(t) \, dt.$$  (1.3)

Therefore, one can view $\phi(t)$ as a probability distribution function which gives the probability of finding eigenvalues of $A$ in a given infinitesimal interval near $t$. Ideally, if one has access to all the eigenvalues of $A$, computing the DOS would become a trivial task. However, in many scientific applications the matrix size is too large for eigenvalue solvers to be efficiently applied, especially when a large number of eigenvalue solvers to be efficiently applied, especially when a large number of

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eigenvalues are needed. Alternative algorithms that do not require all the eigenvalues of A are therefore of great interest in practice.

In solid state physics and electronic structure theory \[3, 40\], eigenvalues of a Hamiltonian matrix usually represent single-particle energy levels. The corresponding DOS gives the number of single-particle energy levels per unit energy. The DOS can also be used to characterize the so-called valence band (energy levels below the chemical potential) and the conduction band (energy levels beyond the chemical potential) along the energy spectrum. The gap between the valence band and the conduction band is called the band-gap, and the DOS inside the band-gap is zero.

In the study of excited state properties of electrons, the joint DOS associated with both the conduction and valence bands allows one to examine the interband transitions owing to the absorption and emission of light \[40, 35\].

At the molecular level, the DOS of the phonon spectrum or the normal modes of molecules, which are related to the eigenvalues of the Hessian of the potential function with respect to atomic coordinates, characterizes thermodynamic properties of solids or molecules. The DOS can also be used as a spectral measure for integrating certain quantities to yield the heat capacity and the entropy of a molecular system. For example, the heat capacity \(C_v\) of a molecular system has the expression \[58\]:

\[
C_v = k_B \int_0^\infty \frac{(\hbar \omega_c/k_B T)^2 e^{-\hbar \omega_c/k_B T} \phi(\omega)}{(1 - e^{-\hbar \omega_c/k_B T})^2} d\omega, \tag{1.4}
\]

where \(k_B\) is the Boltzmann constant, \(c\) is the speed of light, \(\hbar\) is Planck’s constant, \(T\) is the temperature and \(\omega = \sqrt{\lambda}\) is a vibration frequency.

Efficient algorithms for computing the DOS are also developed and employed for optical absorption spectra calculations \[55\], the transition state theory for chemical reactions \[28\], quantum transport simulations \[17\], dynamical correlation function calculations \[10\], and many other recent applications \[18, 27, 2, 1, 7, 47, 9, 4, 48\].

Recall that a Dirac \(\delta\)-function is not a proper function, it is a member of the adjoint space of a Hilbert space. It is formally defined through applications to a test function \(g\):

\[
\langle \delta(\cdot - \lambda), g \rangle = \int_{-\infty}^{\infty} \delta(t - \lambda) g(t) dt \equiv g(\lambda),
\]

where \(g\) is a rapidly decreasing \(C^\infty\) function with bounded support. More details on distributions can be found, e.g., in \[44\]. Because the spectral density is formally defined as the sum of Dirac distributions, it may be thought at first that it cannot be approximated by a smooth function. Indeed, when one tries to view this as a function, it appears to be highly discontinuous, and so it may seem unwise to try to approximate it by smooth functions, e.g., polynomials. However, a distribution can also be viewed as a discretized version of some smooth function after it is sampled at a finite number of points, and our goal is to approximate that original smooth function. Often in quantum mechanical applications, the spectrum of the Hamiltonian defined in the continuous space is continuous, and the function \((1.1)\) shown above can then be viewed as just a discrete version of a continuous function.

A rigorous approach to mitigate the issue, one that has both theoretical and practical importance, is to consider a continuous version of the distribution \(\phi\) obtained, in essence, by “blurring” it. For example, we can consider a function of the form:

\[
\phi_\sigma(t) = \frac{1}{n} \sum_{j=1}^n h_\sigma(t - \lambda_j), \tag{1.5}
\]
in which $h_{\sigma}(t)$ can be any infinitely differentiable function whose integral in $(-\infty, \infty)$ is one, and which is zero or very small outside of a narrow interval $[-C\sigma, C\sigma]$ where $C > 0$ is a small constant independent of $\sigma$. One example is the Gaussian function:

$$h_{\sigma}(t) = \frac{1}{(2\pi\sigma^2)^{1/2}} e^{-\frac{t^2}{2\sigma^2}}. \quad (1.6)$$

Such a setup may be appealing and can help establish some theory as well as develop algorithms as will be seen later.

A related question one might have is: How can we compare the original density $\phi$ with an approximate one $\tilde{\phi}$? The distribution $\phi$ is obtained as the limit of a sequence of functions $\phi_{\sigma} \in C^\infty$ with compact support, in the sense that

$$\lim_{\sigma \to 0} \langle \phi_{\sigma}, g \rangle = \langle \phi, g \rangle,$$

for every $C^\infty$ and compactly supported function $g$. Since

$$|\langle \phi_{\sigma} - \tilde{\phi}, g \rangle| \leq \|\phi_{\sigma} - \tilde{\phi}\|_1 \|g\|_\infty,$$
$$|\langle \phi_{\sigma} - \tilde{\phi}, g \rangle| \leq \|\phi_{\sigma} - \tilde{\phi}\|_2 \|g\|_2,$$
$$|\langle \phi_{\sigma} - \tilde{\phi}, g \rangle| \leq \|\phi_{\sigma} - \tilde{\phi}\|_\infty \|g\|_1,$$

where $\|\cdot\|_p$ stands for the $L^p$ norm, we will only need to first approximate the “true” DOS $\phi$ by a “surrogate” DOS $\phi_{\sigma}$, and then measure the distance between $\phi_{\sigma}$ and $\tilde{\phi}$ either in the $L^1, L^2$ or $L^\infty$ sense to quantify the error.

The goal then is to find a smooth approximation to the spectral density function $\phi$. Because calculating the spectral density is such an important problem in quantum mechanics, there is an abundant literature devoted to this problem and research in this area was extremely active in the 1970s and 1980s. Clever and powerful methods have been developed by physicists and chemists [15, 54, 13, 57] for this purpose and this paper will review a few of them from a linear algebra viewpoint. What is striking about research in this area is the wide variety of methods used as well as the mathematical depth of the various techniques.

The Kernel Polynomial Method [50, 56] expands the spectral density in a basis of orthogonal polynomials by resorting to the reproducing kernel property of orthogonal polynomials. The method, which is widely used in a variety of calculations that require the DOS [14], has continued to receive a tremendous amount of interest in the last few years [16, 5, 32, 39]. Haydock’s method [26] exploits the Lanczos algorithm [36] in an effort to evaluate functions of the form $f(t) = v^T(A - (t + i\eta)I)^{-1}v$ whose peaks reveal the presence of nearby poles, i.e. approximate eigenvalues of $A$. Its use for DOS calculation has recently been discussed in [5, 39, 6]. The Lanczos algorithm itself can be used to estimate the spectrum and the spectral density associated with each estimated eigenvalue. Lanczos also proposed a method [37], termed a “spectroscopic” approach, which turns out to be better suited for calculating spectral densities than actual spectra of Hermitian matrices. Lanczos’s technique also relies on detecting peaks of a certain function, that are indicative of the presence of eigenvalues.

In this paper we wish to take a careful look at the problem of computing the DOS and to describe a few common techniques used in the literature. One of our goals is to expose these techniques from a numerical linear algebra viewpoint. Not all methods described are new, but we will see a new way of developing the Kernel Polynomial Method, which may have a few advantages. In order to directly compute
the surrogate DOS $\phi_\sigma$, we will also describe a new algorithm based on orthogonal expansions of Gaussian functions, as well a method based on using the standard Lanczos algorithm. The next section will discuss spectral density functions in some detail.

2. Spectral densities as regular functions. One of the first questions one might ask when studying spectral densities is: how can we evaluate accuracy? This was briefly discussed in the introduction. A practical viewpoint is to evaluate $\|\phi_\sigma - \tilde{\phi}\|_p$ with the norm $p$ being 1, 2 or $\infty$ where $\phi_\sigma$ is a certain smoothed version of $\phi$. The function $\phi_\sigma$ can be viewed as a surrogate of $\phi$ which can be used for practical purposes. This viewpoint becomes useful for example for the purpose of plotting a spectral distribution for visually comparing it with some approximation.

Let us consider this issue in some more detail. When drawing spectral density curves, we will be interested in some approximations to the density functions shown above. The first approximation that comes to mind is to exploit the “probabilistic” interpretation of $\phi$: we can just subdivide the interval $[a, b]$ into non-overlapping subintervals $I_j = (t_j - h/2, t_j + h/2)$ and define the value of $\phi$ in $I_j$ to be the constant $n_i / n$ where $n_i$ is the number of eigenvalues in $I_j$. This “histogram” approach will give a surrogate which is a piecewise constant function $\phi_{P,h}$ that depends on the interval width $h$. A slight variation of this scheme is to replace each of the Dirac functions $\delta(t - \lambda_i)$ in (1.3) by a piecewise constant function which has a value of $1/h$ in an interval of width $h$ centered at $\lambda_i$ and zero elsewhere. We would have:

$$\phi(t) \approx \phi_{P,h}(t) = \frac{1}{nh} \sum_{j=1}^{n} \chi\left(\frac{t - \lambda_i}{h/2}\right)$$

where $\chi(t)$ represents the indicator function for the interval $[-1, 1]$, which equals to one if $t$ is in the interval and zero otherwise. The integral of this function on the whole real line is one as required. We note that piecewise constant surrogates will tend to give us rough functions with big variations if the selected width $h$ is small.

A better alternative is to use Gaussians, replacing again each $\delta(t - \lambda_i)$ by the function $h_\sigma(t - \lambda_i)$ where $h_\sigma$ is defined by (1.6). The parameter $\sigma$ plays the role of $h$ in the above case of piecewise constant surrogates. Larger values of $\sigma$ will lead to smooth curves at the expense of accuracy. Too small values of $\sigma$ will again lead to rough curves that have peaks at the eigenvalues and zeros elsewhere. This is illustrated in Figure 2.1 where $\sigma$ takes 4 different values. We can see that as $\sigma$ increases, $\phi_\sigma$ becomes smoother. When $\sigma = 0.96$, which corresponds to a very smooth spectral density, we can still see the global profile of the eigenvalue distribution, although local variation of the spectral density is mostly averaged out.

One of the methods to be described later will consist of seeking an approximation of a smoothed version of $\phi$ obtained by discretizing the interval $[a, b]$ uniformly and placing Gaussians at each knot. In this way the function that is approximated is a regular function.

A heuristic criterion we adopt for choosing $\sigma$ in the case where uniform intervals are used is based on starting with some width $h$ that represents the discretization width of the interval. To obtain a compromise between smoothness and accuracy we select $\sigma$ so that the value of each Gaussian at its peak, is $\kappa$ times its value at the end point of each sub-interval, where $\kappa$ is a ratio larger than one. This means that

$$1 = \kappa e^{-(h/2)^2/(2\sigma^2)} \rightarrow \log(\kappa) = \frac{h^2}{8\sigma^2} \rightarrow \sigma = \frac{h}{2\sqrt{2\log(\kappa)}}$$

(2.1)
If $\kappa$ is close to one, then the overall curve, which is the sum of all the Gaussians will be very smooth. This is illustrated in Figure 2.2.

The criterion given by (2.1) is motivated by the case when the distribution is uniform, as Figure 2.2 illustrates. In this situation, we would like to have a function that has values not too far from one in the middle of the interval and at the interval intersections. One can decide on the level of smoothness desired in the final approx-
imation and select \( h \) and \( \kappa \) accordingly. One can choose \( h = \Delta E/N \), where \( \Delta E \) is the width of the spectrum, and \( N \) is the intended resolution of the DOS. Values of \( N \) between 30 and 100 and \( \kappa \) between 1.25 and 2.0 seem to serve as reasonable “default” selections.

When calculating these functions for the different \( \lambda_i \) we only need to evaluate the Gaussians at points nearby a given site \( \lambda_i \) because of the rapid decay of Gaussians. A calculation similar to the one given above will tell us which intervals are to be kept if we need the accuracy to be within a certain tolerance.

We remark that the optimal choice of \( \sigma \), and therefore the smoothness of the approximate DOS, is application dependent. Ideally, \( \sigma \) should be chosen to be as large as possible so that the surrogate DOS \( \phi_\sigma \) is easy to approximate numerically. However, increasing \( \sigma \) could cause an undesirable loss of detail and yield an erroneous result, e.g., when computing the heat capacity discussed in Section II. It is up to the user to select a value of \( \sigma \) that balances accuracy and efficiency.

3. The Kernel Polynomial Method. The Kernel Polynomial Method (KPM) was proposed by Silver and Röder [50] and Wang [56] in the mid-1990s to calculate the DOS. See also [51, 52, 14, 41] among others where similar approaches were also used.

The KPM method essentially determines the exact DOS of a matrix as a sum of Dirac \( \delta \)-functions by its expansion in Chebyshev polynomials. For simplicity we assume that the eigenvalues are in the interval \([-1, 1]\). As is the case for all methods which rely on Chebyshev expansions, a change of variables is first performed to map the interval \([\lambda_{\min}, \lambda_{\max}]\) into \([-1, 1]\). The goal is to estimate the spectral density function (1.1). For this, we will approximate \( \hat{\phi}(t) \) by a finite expansion in a basis of orthogonal polynomials, in this case, Chebyshev polynomials of the first kind. Following the Silver-Röder paper [50], we include, for convenience, the inverse of the weight function into the spectral density function, so we expand instead the distribution:

\[
\hat{\phi}(t) = \sqrt{1-t^2} \phi(t) = \sqrt{1-t^2} \times \sum_{j=1}^{n} \delta(t - \lambda_j).
\]

Then, we have the (full) expansion

\[
\hat{\phi}(t) = \sum_{k=0}^{\infty} \mu_k T_k(t).
\]

where the expansion coefficients \( \mu_k \) are formally defined by

\[
\mu_k = \frac{2 - \delta_{k0}}{\pi} \int_{-1}^{1} \frac{1}{\sqrt{1-t^2}} T_k(t) \hat{\phi}(t) dt
= \frac{2 - \delta_{k0}}{\pi} \int_{-1}^{1} \frac{1}{\sqrt{1-t^2}} T_k(t) \sqrt{1-t^2} \phi(t) dt
= \frac{2 - \delta_{k0}}{n\pi} \sum_{j=1}^{n} T_k(\lambda_j).
\]

Here \( \delta_{ij} \) is the Kronecker \( \delta \) symbol so that \( 2 - \delta_{k0} \) is equal to 1 when \( k = 0 \) and to 2 otherwise.

Thus, apart from the scaling factor \((2 - \delta_{k0})/(n\pi)\), \( \mu_k \) is the trace of \( T_k(A) \) and this can be estimated by various methods including, but not limited to, stochastic
approaches. There are variations on this idea starting from the use of different orthogonal polynomials, to alternative ways in which the traces can be estimated.

A common solution to the problem of estimating \( \text{Trace}(T_k(A)) \) is to use a stochastic argument as suggested in [50] and, e.g., also in [29, 53]. This entails generating a large number of random vectors \( v_0^{(1)}, v_0^{(2)}, \ldots, v_0^{(n_{\text{vec}})} \) with each component obtained from a normal distribution with zero mean and unit standard deviation, and each vector is normalized such that \( \|v_0^{(l)}\| = 1, l = 1, \ldots, n_{\text{vec}} \). The subscript 0 is added to indicate that the vector has not been multiplied by the matrix \( A \). Then we can estimate the trace of \( T_k(A) \) as follows:

\[
\text{Trace}(T_k(A)) \approx \frac{1}{n_{\text{vec}}} \sum_{l=1}^{n_{\text{vec}}} \left( v_0^{(l)} \right)^T T_k(A) v_0^{(l)}. \tag{3.4}
\]

Then this will lead to the desired estimate:

\[
\mu_k \approx 2 - \delta_k 0 = \frac{2}{n\pi n_{\text{vec}}} \sum_{l=1}^{n_{\text{vec}}} \left( v_0^{(l)} \right)^T T_k(A) v_0^{(l)}. \tag{3.5}
\]

Now we consider the computation of each term \( \left( v_0^{(l)} \right)^T T_k(A) v_0^{(l)} \). For simplicity we drop the superscript \( l \) and denote by \( v_0 \equiv v_0^{(l)} \). The 3-term recurrence of the Chebyshev polynomial is exploited to compute \( T_k(A)v_0 \):

\[
T_{k+1}(A)v_0 = 2AT_k(A)v_0 - T_{k-1}(A)v_0 \tag{3.6}
\]

so if we let \( v_k \equiv T_k(A)v_0 \), we have

\[
v_{k+1} = 2Av_k - v_{k-1}.
\]

Once the scalars \( \{\mu_k\} \) are determined, we would in theory get the expansion for \( \phi(t) = \frac{1}{\sqrt{1-t^2}} \phi(t) \). Practically however, the approximate density of states will be limited to Chebyshev polynomials of degree \( M \), so \( \hat{\phi} \) is approximated by:

\[
\hat{\phi}_M(t) = \frac{1}{\sqrt{1-t^2}} \sum_{k=0}^{M} \mu_k T_k(t). \tag{3.7}
\]

For a general matrix \( A \) whose eigenvalues are not necessarily in the interval \([-1,1]\], a linear transformation is first applied to \( A \) to bring its eigenvalues to the desired interval. Specifically, we will apply the method to the matrix

\[
B = \frac{A - cI}{d},
\]

where

\[
c = \frac{\lambda_{\min} + \lambda_{\max}}{2}, \quad d = \frac{\lambda_{\max} - \lambda_{\min}}{2}. \tag{3.8}
\]

It is important to ensure that the eigenvalues of \( B \) are within the interval \([-1,1]\). In an application requiring a similar approach [59], we obtain the upper and lower bounds of the spectrum from Ritz values provided by a standard Lanczos iteration.
We ran a few Lanczos steps but extended the interval $[\lambda_{\min}, \lambda_{\max}]$ by using the bounds obtained from the Lanczos algorithm.

Specifically, the upper bound for $\lambda_{\max}$, for example, is set to $[\tilde{\lambda}_n, \tilde{\lambda}_n + \eta]$ where $\eta = \| (A - \tilde{\lambda}_n I) \tilde{u}_n \|$ and $(\tilde{\lambda}_n, \tilde{u}_n)$ is the (algebraically) largest Ritz pair of $A$. This is plausible because we know that there is an eigenvalue in the interval $[\tilde{\lambda}_n, \tilde{\lambda}_n + \eta]$ (see e.g., [46].) Even though there is no theoretical guarantee that $[\tilde{\lambda}_n, \tilde{\lambda}_n + \eta]$ is indeed an upper bound for $\lambda_{\max}$, in practice it often is. Refined bounds which invoke the square of $\eta$ ([46]) can also be used. Note that $\eta$ can be computed without explicitly computing the eigenvector $\tilde{u}_n$. To summarize, we outline the major steps of the KPM for approximating the spectral density of a sparse matrix in Algorithm 1.

**Algorithm 1: The Kernel Polynomial Method.**

**Input:** Real symmetric matrix $A$. A set of points $\{t_i\}$ at which DOS is to be evaluated, the degree $M$ of the expansion polynomial.

**Output:** Approximate DOS $\{\tilde{\phi}_M(t_i)\}$.

1: Set $\mu_k = 0$ for $k = 0, \cdots, M$;
2: for $l = 1 : n_{vec}$ do
3: Select a new random vector $v_0^{(l)}$;
4: for $k = 0 : M$ do
5: Compute $T_k(A)v_0^{(l)}$ using 3-term recurrence (3.6);
6: Update $\mu_k$ using (3.5);
7: end for
8: end for
9: Evaluate the average value of $\{\tilde{\phi}_M(t_i)\}$ at the given set of points $\{t_i\}$ using (3.7).

Because the KPM method amounts to approximating a discontinuous function, it may lead to some numerical difficulties. The Dirac $\delta$-function can be viewed as the derivative of the Heaviside step function $H(t)$ such that

$$H(t) = \begin{cases} 
1, & t \geq 0, \\
0, & t < 0.
\end{cases}$$

(3.9)

The discontinuity at 0 causes the polynomial approximation of $H(t)$ and $\delta(t)$ to exhibit rapid oscillations around $t = 0$. These so-called Gibbs oscillations can lead to loss of accuracy in the approximate DOS generated by KPM. A common approach used to damp these oscillations, is to use the Chebyshev-Jackson approximation [30, 45, 31], which modulates the coefficients $\mu_k$ with a damping factor $g_k^M$ defined by

$$g_k^M = \left(1 - \frac{k}{M+1}\right) \sin(\alpha_M) \cos(k\alpha_M) + \frac{1}{M+1} \cos(\alpha_M) \sin(\alpha_M) \sin(\alpha_M),$$

(3.10)

where $\alpha_M = \frac{\pi}{M+1}$. Consequently, the damped Chebyshev expansion has the form

$$\tilde{\phi}_M(t) = \sum_{k=0}^{M} \mu_k g_k^M T_k(t).$$

Approximations to the $\delta$-function $\delta(t)$ by Chebyshev polynomial expansions with and without the Jackson damping are compared in the left subfigure in Figure 3.1.
degrees of both polynomial approximations are set to 40. The right subfigure in Figure 3.1 shows similar approximations for the Heaviside function $H(t)$. Observe how the approximate function obtained with Jackson damping is smooth and free of the rapid oscillations seen in the Chebyshev expansion.

To illustrate how well this algorithm works, we now give an example on using KPM to approximate the spectral density of a relatively small matrix. The matrix we use for this example is a modified two-dimensional (2D) Laplacian operator with zero Dirichlet boundary condition discretized on a domain of size $[0,30] \times [0,30]$ using a five-point finite difference stencil. The modification involves adding a diagonal matrix as a discretized potential function. The diagonal matrix is generated by adding two Gaussians, one centered at the point $(4,5)$ of the domain and the other at the point $(25,15)$. In Figure 3.2 we plot both the approximate spectral density constructed by KPM with a 80-degree Chebyshev polynomial and 10 random vectors for estimating the trace in (3.4), and the smoothed “exact” spectral density constructed by Gaussian smoothing with the smoothing parameter $\sigma = 0.56$ (obtained by setting $\kappa = 1.25, h = 0.75$).

Fig. 3.2: The approximate spectral density constructed by KPM with a 80-degree Chebyshev polynomial for the modified 2D Laplacian example with Jackson damping (left) and without Jackson damping (right).
From a computational point of view, some savings in time can be achieved if we are willing to store more vectors. This is due to the formula:

\[ T_p(t)T_q(t) = \frac{1}{2} \left[ T_{p+q}(t) - T_{|p-q|}(t) \right], \]

from which we obtain

\[ T_{p+q}(t) = 2T_p(t)T_q(t) + T_{|p-q|}(t). \]

For a given \( k \) we can use the above formula with \( p = \lceil k/2 \rceil \) and \( q = k - p \). This requires that we compute and store \( v_r = T_r(A)v_0 \) for \( r \leq p \). Then the moments \( v_0^T T_r(A)v_0 \) for \( r \leq p \) can be computed in the usual way, and for \( r = p + q > p \) we can use the formula:

\[ v_0^T T_{p+q}(A)v_0 = 2v_p^T v_q + v_0^T v_{p-q}. \]

This saves 1/2 of the matrix-vector products at the expense of storing all the previous \( \{v_r\} \). It is not practical for high degree polynomials.

Chebyshev polynomials are not the only types of orthogonal polynomials that can be used in the expansion. We can use any other type of orthogonal polynomials. The only practical requirement is that we explicitly know the 3-term recurrence for the polynomials. For example, we can use the Legendre polynomials \( L_k(t) \) which obey the following 3-term recursion

\[ L_0(t) = 1, \quad L_1(t) = t, \quad (k + 1)L_{k+1}(t) = (2k + 1)tL_k(t) - kL_{k-1}(t). \]

See, for example [11], for three-term recurrences for a wide class of such polynomials, e.g., all those belonging to the Jacobi class, which include Legendre and Chebyshev polynomials as particular cases.

Figure 3.3 shows the quality of the approximation to the spectral density of the modified 2D Laplacian matrix obtained from the KPM with a Legendre polynomial expansion. The degree of the polynomial approximation is set to 80. We observe that the performance of the Legendre polynomial expansion is similar to that of a Chebyshev expansion without Jackson damping.

![Fig. 3.3](image-url)

Fig. 3.3: The approximate spectral density constructed by KPM with a 80-degree Legendre polynomial for the modified 2D Laplacian matrix.
4. Spectroscopic approaches. In his 1956 book titled “Applied Analysis” \[37\], Lanczos described a method for computing spectra of Hermitian matrices, which he termed “spectroscopic”. This approach, which also relies heavily on Chebyshev polynomials, is rather unusual in that it assimilates the spectrum of a matrix to a collection of frequencies and the goal is to detect these frequencies by Fourier analysis. Because it is not competitive with modern methods for computing eigenvalues, this technique has lost its appeal. However, it appears to be better suited for computing approximate spectral densities and so we consider here an adaptation of this approach for this purpose.

Assume that $B$ is the same matrix as the one in Section 3 which has eigenvalues in $[-1,1]$. Let $v_0$ be any initial vector and assume that it is expanded using the eigenvectors of $A$ as

$$v_0 = \sum_{j=1}^{n} \beta_j u_j,$$

where the coefficient $\beta_j = u_j^T v_0$. Consider the vector $v_k = T_k(A)v_0$, for $k = 0, \cdots, M$.

If we set $\lambda_j = \cos \theta_j$, then

$$v_0^T v_k = \sum_{j=1}^{n} \beta_j^2 T_k(\lambda_j) = \sum_{j=1}^{n} \beta_j^2 \cos(k\theta_j).$$

The number $v_0^T v_k$ can be viewed as a discretized version of the function

$$f(t) = \sum_{j=1}^{n} \beta_j^2 \cos(t\theta_j)$$

sampled at the integer values $t = 0, \cdots, M$. The problem is then to find all values of $\theta_j$. In order to find the $\theta_j$’s, Lanczos reasoned as follows. The function $f$ is a periodic function and we can compute its Fourier (in this case cosine) transform:

$$F(p) = \frac{1}{2} (f(0) + (-1)^p f(M)) + \sum_{k=1}^{M-1} f(k) \cos \frac{kp\pi}{M}, \quad p = 0, \cdots, M.$$  \hspace{1cm} (4.4)

Note that as is customary the end values are halved to account for the discontinuity of the data at the interval boundaries.

The idea then is that if $f$ has an eigenvalue $\lambda = \cos \theta$, there should be a component $\cos(p\theta)$, which should be revealed by a peak at the point

$$p = \frac{l\theta}{\pi}.$$  

Indeed, if we had exactly one term of the form $f(t) = \cos \left(\frac{lt\pi}{M}\right)$ where $0 \leq p \leq M$ is some integer, then only $F(M)$ would be nonzero. In the end, if we identify a peak value at $p_j$ then, the corresponding eigenvalue $\lambda_j$ corresponds to the angle $\theta_j = (p_j/M)\pi$, and so, $\lambda_j = \cos(\theta_j) = \cos(p_j\pi/M)$. In other words, after obtaining $F(p), p = 0, \cdots, M$, one can define an associated function

$$\hat{F}(\hat{p}) \equiv F \left( \frac{M}{\pi} \arccos \hat{p} \right), \quad \hat{p} = \cos(p\pi/M), p = 0, \cdots, M.$$  \hspace{1cm} (4.5)
Then $\hat{F}(\hat{p})$ can be interpolated on the entire interval of $[-1, 1]$, and finally the approximate DOS can be evaluated as

$$\tilde{\phi}_M(t) = C_M \hat{F} \left( \frac{t - c}{d} \right),$$

where $c, d$ are defined in Eq. (4.8), and $C_M$ is a normalization constant so that $\int \tilde{\phi}_M(t) dt = 1$.

Note that the quantity $\eta_k \equiv v_0^T v_k$ is based on a single vector $v_0$ and this will cause problems when one or more of the corresponding components $\beta_j$ in (4.1) are small. To avoid this we will proceed in the same way as with the KPM method by taking many vectors and averaging the above quantities.

The basic steps of the spectroscopic method are outlined in Algorithm 2. Fig. 4.1 illustrates the performance of the method for the modified 2D Laplacian matrix with the polynomial degrees set to 40 and 100, respectively. The performance of the spectroscopic method is comparable to that of the KPM method without Jackson damping. However, the usage of Jackson damping is more difficult to justify.

**Algorithm 2: Lanczos spectroscopic method**

**Input:** Real symmetric matrix $A$. A set of points $\{t_i\}$ at which the spectral density is to be evaluated and the degree of Chebyshev polynomial approximation $M$.

**Output:** Approximate DOS $\{\tilde{\phi}_M(t_i)\}$.

1: Set $\eta_k = 0$ for $k = 0, \ldots, M$;
2: for $l = 1 : n_{\text{vec}}$ do
3: Select a new random vector $v_0^{(l)}$;
4: for $k = 0 : M$ do
5: Compute $v_{k+1}^{(l)}$ via the three-term recurrence $v_{k+1}^{(l)} = 2Av_k^{(l)} - v_{k-1}^{(l)}$ (for $k = 0$, $v_1^{(l)} = Av_0^{(l)}$);
6: Compute $\eta_k \leftarrow \eta_k + v_0^{(l)T} v_k^{(l)}$;
7: end for
8: end for
9: Set $\eta_k = \eta_k/n_{\text{vec}}$ for all $k = 0, 1, \ldots, M$;
10: Take the discrete cosine transform of $\{\eta_k\}$ according to Eq. (4.4) to obtain $F(p)$ for $p = 0, 1, \ldots, M$;
11: Evaluate $\tilde{\phi}_M(t_i)$ according to Eq. (4.6) for each $t_i$.

5. The Delta-Chebyshev expansion algorithm. In this section, we will use the insight provided by the spectroscopic method seen in the previous section to derive a new algorithm which has some of the features of the spectroscopic method and which will turn out to be mathematically equivalent to KPM. The main idea relies on an expansion of the $\delta$-function at a given set of points $\{t_i\}$.

A $\delta$-function defined at $t_i$ can be viewed as a spectral probe at $t_i$. In theory, the presence of an eigenvalue at $t_i$ can be detected by integrating the $\delta$-function over the entire spectrum of $A$ with respect to a spectral point measure defined at eigenvalues only, i.e., $\int \delta(t - t_i) \, dt \equiv \sum_{i=1}^{\infty} \delta(\lambda_i - t_i)/n$. The integral returns $+\infty$ if $t_i$ is an eigenvalue of $A$ and 0 otherwise. However, in practice, this integration cannot be performed without knowing the eigenvalues of $A$ in advance.
A practical probing scheme can be devised by replacing the $\delta$-function with a polynomial approximation, and approximating the spectral measure using the same stochastic approach we introduced earlier for the KPM.

For example, if the problem has been scaled and shifted so that the spectrum of the transformed problem has been mapped to the interval $[-1, 1]$, we may approximate the $\delta$-function centered at $t_i$ by using an $M_i$-degree Chebyshev polynomial expansion of the form

$$\delta(t - t_i) \approx \rho_{M_i}(t) \equiv \sum_{k=0}^{M_i} \mu_{k_i}(t_i) T_{k_i}(t),$$

(5.1)

where the expansion coefficient $\mu_{k_i}(t_i)$ can be determined from

$$\mu_{k_i}(t_i) = 2 - \delta_{k0} \frac{1}{\pi} \int_{-1}^{1} \frac{1}{\sqrt{1 - t^2}} T_{k_i}(t) \delta(t - t_i) dt = 2 - \delta_{k0} \frac{T_{k_i}(t_i)}{\sqrt{1 - t_i^2}}$$

(5.2)

If $v_0$ is some normalized random vector, whose expansion in the eigenbasis is given by (4.1), then $v_{M_i} \equiv \rho_{M_i}(A)v_0$ will have the expansion

$$v_{M_i} = \sum_{j=1}^{n} \beta_j \rho_{M_i}(\lambda_j) u_j.$$  

(5.3)

Taking the Euclidean inner product between $v_0$ and $v_{M_i}$ yields

$$\langle v_{M_i}, v_0 \rangle = \sum_{j=1}^{n} \beta_j^2 \rho_{M_i}(\lambda_j).$$

(5.4)

Since $\sum_{j=1}^{n} \beta_j^2 = 1$, (5.4) can be viewed as an integral of $\rho_{M_i}(t)$ associated with a point measure $\{\beta_j^2\}$ defined at eigenvalues of the transformed matrix $A$. If $\beta_j^2 = 1/n$ for all $j$, we can then simply rewrite the integral as $\text{Trace}[\rho_{M_i}(A)]/n$. As we have already shown in previous sections, such a trace can be approximated by choosing multiple random vectors $v_0$ and averaging $\langle v_0, \rho_{M_i}(A)v_0 \rangle$ for all these vectors.
Formally, the ideas described above can be organized as in Algorithm 3 which we will refer to as the Delta-Chebyshev expansion algorithm. First, we select a number of nodes $t_i$ at which the spectral density of $A$ is to be evaluated. At each of these points, we evaluate the expansion coefficients $\mu_k$ in (5.1). We then loop through a large number of random vectors and compute the averages of the estimates (5.4) for each of the sample points $t_i$. These steps are outlined more precisely in Algorithm 3.

Algorithm 3: Multi-point Delta-Chebyshev expansion

\begin{algorithm}
\begin{algorithmic}
\State \textbf{Input:} Real symmetric matrix $A$. A set of points $\{t_i\}$, and associated Chebyshev polynomial degree $\{M_i\}$, and $M_{\text{max}}$ is the maximum degree employed for all the points.
\State \textbf{Output:} Approximate DOS $\{\tilde{\phi}_M(t_i)\}$.
\For {each $t_i$}
\State Compute and store the expansion coefficients $\mu_k$, for $k = 0, \cdots, M_i$;
\EndFor
\For {$l = 1 : n_{\text{voc}}$}
\State Select a new random vector $v_0^{(l)}$;
\State For $k = 0 : M_{\text{max}}$
\State \hspace{1em} Compute $v_{k+1}^{(l)}$ via the three-term recurrence $v_{k+1}^{(l)} = 2 Av_k^{(l)} - v_k^{(l)}$ (for $k = 0$, $v_1^{(l)} = Av_0^{(l)}$);
\State \hspace{1em} Compute $\eta_k \leftarrow \eta_k + (v_0^{(l)})^T v_k^{(l)}$;
\EndFor
\State Set $\eta_k = \eta_k/n_{\text{voc}}$ for all $k = 0, 1, \cdots, M_{\text{max}}$;
\State Evaluate $\tilde{\phi}_M(t_i)$ using $\eta_k$ and the stored $\mu_k$'s;
\end{algorithmic}
\end{algorithm}

Note that this algorithm would have been an unacceptably expensive procedure if it were not for the fact that the same Chebyshev sequence $v_k = T_k(A)v_0$ can be used for all points $t_i$ at the same time. Since the average value $\langle v_0, p_M(A)v_0 \rangle$ can be viewed as an approximation to the trace of $p_M(A)$, we can rewrite this quantity, which serves as an approximation to the spectral density at $t_i$, as

$$\phi(t_i) = \frac{1}{n} \sum_{k_i = 0}^{M_i} \mu_{k_i}(t_i) \sum_{j=1}^{n} T_{k_i}(\lambda_j) = \frac{1}{n} \sum_{k_i = 0}^{M_i} \frac{2 - \delta_{k_i,0}}{\pi} T_{k_i}(t_i) \sqrt{1 - t_i^2} \text{Trace}(T_{k_i}(A)) = \sum_{k_i = 0}^{M_i} \left[ \frac{2 - \delta_{k_i,0}}{n\pi} \text{Trace}(T_{k_i}(A)) \right] T_{k_i}(t_i) \sqrt{1 - t_i^2}. \quad (5.5)$$

Note that the coefficients within the square bracket in (5.5) are exactly the same coefficients that appear in the expansion (3.2) of the function $\phi(t) = \sqrt{1 - t^2}\phi(t)$ used in the KPM. Therefore, when $M_i = M$ for all $i$, the Delta-Chebyshev expansion method is identical to the KPM. Hence, the cost of this approach is the same as that of KPM if polynomials of the same degree and the same number of sampling vectors are used at each $t_i$.

However, when $M_i$ is allowed to vary with respect to $i$, there is a slight advantage of using the Delta-Chebyshev method in terms of flexibility. We can use polynomials
of different degrees in different parts of the spectrum to obtain a more accurate approximation. Note that in this situation, if $M_{\text{max}}$ is the maximum degree employed for all the points, the number of matrix-vector products employed remains the same and equal to $M_{\text{max}}$, since we will need to compute for each random vector $v_0$, the vectors $T_k(A)v_0$ for $k = 0, \cdots, M_{\text{max}}$ as these are needed by the points requiring the highest degree. However, some of the other calculations (inner products) required to obtain the spectral density can be avoided, though in most cases applying $T_k(A)$ to $v_0$ dominate the computational cost in the DOS calculation. The Delta-Gauss-Legendre expansion to be developed in the next section shares similar property in this aspect.

We should point out that it is also possible to apply the Delta-Chebyshev method for approximating the accumulated spectral density, which is simply the integral of the spectral density:

$$\psi(t) = \int_{-\infty}^{t} \phi(s)ds$$  \hspace{1cm} (5.6)

Consider the step function shown on the right side of Figure 5.1. Let $H(t)$ be the Heaviside function defined in Eq. (3.9). Then the ideal step function $H(t - s)$ which changes value at certain $t$ and we have:

$$\psi(t) = \int_{-\infty}^{t} \phi(s)ds = \int_{-\infty}^{\infty} H(t - s)\phi(s)ds$$

So, $\psi(t_0)$ can also be estimated by an expression like (5.4) where now the polynomial is an approximation of the step function $H(t)$ instead of the $\delta$-function.

![Fig. 5.1: The approximate spectral densities constructed by the Delta-Chebyshev method with Jackson damping (left) and without Jackson damping (right) for the modified 2D Laplacian matrix. The polynomial degree is 80.](image)

6. The Delta-Gauss-Legendre expansion approach. In Section 2 we invoked the idea of comparing the approximation $\tilde{\phi}$ of the spectral density not with the spectral density $\phi$ directly but with a smoothed version of $\phi$ which we denoted by $\phi_{\sigma}$, where $\sigma$ is a smoothing parameter. We argued that this approach is sensible because it may be easier in practice to compare smooth functions than to compare a distribution with a function. In this section, we will take the idea a little further. Instead of approximating $\phi$ directly we will first select a representative $\phi_{\sigma}$ of $\phi$ for a given $\sigma$ and then will approximate $\phi_{\sigma}$ by expanding it in orthogonal polynomials. In this way,
comparisons will be relative to the same objects and the function to approximate can be as smooth as desired.

We will proceed in the same way as with the Delta-Chebyshev approach of the previous section, but instead of expanding the Dirac function at the point \( t \) we will now expand its smoothed version or surrogate defined by

\[
h_\sigma(\lambda - t) = \frac{1}{(2\pi \sigma^2)^{1/2}} \exp \left[-\frac{(\lambda - t)^2}{2\sigma^2}\right].
\]  
(6.1)

This is a Gaussian centered at \( t \), and we wish to expand it in Legendre polynomials. We will drop the normalization coefficient for now and seek the expansion parameters

\[
\gamma_k(t) = \int_{-1}^{1} L_k(s)e^{-\frac{1}{2}(s-t)^2}/\sigma^2} ds,
\]  
(6.2)

where \( L_k(\lambda) \) is the Legendre polynomial of degree \( k \). In the following we consider one fixed value \( t \) and so we will drop the variable \( t \) and use \( \gamma_k \) to denote \( \gamma_k(t) \). We point out that Legendre polynomials do not have unit norms. Therefore, they should be properly scaled in the expansion. Specifically, since the norm of \( L_k \) is \( \sqrt{2/(2k+1)} \), the expansion is actually:

\[
h_\sigma(\lambda - t) = \frac{1}{(2\pi \sigma^2)^{1/2}} \exp \left[-\frac{(\lambda - t)^2}{2\sigma^2}\right] = \frac{1}{(2\pi \sigma^2)^{1/2}} \sum_{k=0}^{\infty} \left(\frac{k+1}{2}\right) \gamma_k L_k(\lambda).
\]  
(6.3)

We now calculate the \( \gamma_k \)'s starting with \( \gamma_0 \). Since \( L_0(\lambda) = 1 \), a change of variable \( t = (s-t)/\sqrt{2\sigma^2} \) yields

\[
\gamma_0 = \sigma \sqrt{\frac{\pi}{2}} \left[ \text{erf} \left(\frac{1-t}{\sqrt{2\sigma}}\right) - \text{erf} \left(-\frac{1-t}{\sqrt{2\sigma}}\right) \right] = \sigma \sqrt{\frac{\pi}{2}} \left[ \text{erf} \left(\frac{1-t}{\sqrt{2\sigma}}\right) + \text{erf} \left(\frac{1+t}{\sqrt{2\sigma}}\right) \right],
\]  
(6.4)

where we have used the standard error function:

\[
\text{erf}(x) = \frac{2}{\sqrt{\pi}} \int_{0}^{x} e^{-t^2} dt.
\]

Now consider a general coefficient \( \gamma_{k+1} \) with \( k \geq 0 \). There does not seem to exist a closed form formula for \( \gamma_k \) for a general \( k \). However, these coefficients can be obtained by a recurrence relation. To this end we will need to determine concurrently the sequence:

\[
\psi_k = \int_{-1}^{1} L_k(s)e^{-\frac{1}{2}(s-t)^2}/\sigma^2} ds.
\]  
(6.5)

From the 3-term recurrence of the Legendre polynomials:

\[
(k + 1)L_{k+1}(\lambda) = (2k + 1)\lambda L_k(\lambda) - kL_{k-1}(\lambda)
\]  
(6.6)

we get by integration:

\[
(k + 1)\gamma_{k+1} = (2k + 1) \int_{-1}^{1} sL_k(s)e^{-\frac{1}{2}(s-t)^2}/\sigma^2} ds - k\gamma_{k-1}.
\]  
(6.7)
A useful observation is that the above formula is valid for \( k = 0 \) by setting \( \gamma_{-1} \equiv 0 \). This comes from (6.6), which is valid for \( k = 0 \) by setting \( L_{-1}(\lambda) \equiv 0 \). Next we expand the integral term in the above equality:

\[
\int_{-1}^{1} s e^{-\frac{i}{2}(s-t)/\sigma^2} L_k(s) ds = \sigma^2 \int_{-1}^{1} \frac{s - t}{\sigma^2} e^{-\frac{i}{2}(s-t)/\sigma^2} L_k(s) ds + t\gamma_k \quad (6.8)
\]

\[
= \sigma^2 \int_{-1}^{1} \frac{d}{ds}[-e^{-\frac{i}{2}(s-t)/\sigma^2}] L_k(s) ds + t\gamma_k. \quad (6.9)
\]

The next step is to proceed with integration by parts for the integral in the above expression:

\[
\int_{-1}^{1} \frac{d}{ds}[-e^{-\frac{i}{2}(s-t)/\sigma^2}] L_k(s) ds = -L_k(s)e^{-\frac{i}{2}(s-t)/\sigma^2} \bigg|_{-1}^{1} + \int_{-1}^{1} e^{-\frac{i}{2}(s-t)/\sigma^2} L'_k(s) ds. \quad (6.10)
\]

Noting that \( L_k(1) = 1 \) and \( L_k(-1) = (-1)^k \) for all \( k \), we get

\[
\int_{-1}^{1} \frac{d}{ds}[-e^{-\frac{i}{2}(s-t)/\sigma^2}] L_k(s) ds = -e^{-\frac{i}{2}(1-t)/\sigma^2} + (-1)^k e^{-\frac{i}{2}(1+t)/\sigma^2} + \psi_k \quad (6.11)
\]

\[
= -e^{-\frac{i}{2}(1+t^2)/\sigma^2} \left[ e^{t/\sigma^2} - (-1)^k e^{-t/\sigma^2} \right] + \psi_k \quad (6.12)
\]

\[
\equiv \psi_k - \zeta_k, \quad (6.13)
\]

where we have defined

\[
\zeta_k = e^{-\frac{i}{2}(1-t)/\sigma^2} - (-1)^k e^{-\frac{i}{2}(1+t)/\sigma^2} \quad (6.14)
\]

We note in passing that according to (6.12), \( \zeta_k \) can be written as

\[
\zeta_k = \begin{cases} 
2e^{-\frac{i}{2}(1+t^2)/\sigma^2} \sin(t/\sigma^2) & \text{for } k \text{ even} \\
2e^{-\frac{i}{2}(1+t^2)/\sigma^2} \sinh(t/\sigma^2) & \text{for } k \text{ odd} 
\end{cases}
\]

Substituting (6.11) into (6.9) and the result into (6.7) yields

\[
(k + 1)\gamma_{k+1} = (2k + 1) \left[ \sigma^2(\psi_k - \zeta_k) + t\gamma_k \right] - k\gamma_{k-1} \quad (6.15)
\]

The only thing that is left to do is to find a recurrence for the \( \psi_k \)'s. Here we use the elegant formula which can be found in, e.g., [38] p. 47]

\[
L'_{k+1}(\lambda) = (2k + 1) L_k(\lambda) + L'_{k-1}(\lambda). \quad (6.16)
\]

Integrating in \([-1,1]\) yields the relation:

\[
\psi_{k+1} = (2k + 1) \gamma_k + \psi_{k-1} \quad (6.17)
\]

Note that initial values of \( \psi_k \) are \( \psi_0 = 0, \psi_1 = \gamma_0 \). In the end, we obtain the following recurrence relations:

\[
\begin{align*}
\gamma_{k+1} &= \frac{2k+1}{k+1} \left[ \sigma^2(\psi_k - \zeta_k) + t\gamma_k \right] - \frac{k}{k+1} \gamma_{k-1} \\
\psi_{k+1} &= (2k + 1) \gamma_k + \psi_{k-1}. \quad (6.18)
\end{align*}
\]
It can be noted that the above formulas work for $k = 0$ by setting $\gamma - 1 = \psi - 1 = 0$.

The recurrence starts with $k = 0$, using the initial values $\gamma_0$ given by (6.4), $\psi_1 = \gamma_0$, and $\psi_0 = 0$. The result is now summarized in the following proposition.

**Proposition 6.1.** The Gaussian function (6.1) admits the following expansion in Legendre polynomials:

$$h_\sigma(\lambda - t) = \frac{1}{(2\pi \sigma^2)^{1/2}} \sum_{k=0}^{\infty} \left( k + \frac{1}{2} \right) \gamma_k L_k(\lambda),$$  \hspace{1cm} (6.19)

where $\gamma_0$ is given by (6.4), and the coefficients $\gamma_k$ for $k \geq 1$ are defined by the recurrence (6.18), starting with $k = 0$ and using the initial values $\gamma_{-1} = 0$, $\psi_0 = \psi_{-1} = 0$.

An important remark here is that one has to be careful about the application of the recurrence (6.18). The perceptive reader may notice that such a recurrence runs the risk of being unstable. In fact we observe the following behavior. For large values of $\sigma$ the function (6.1) can be very smooth and as a result a very small degree of polynomials may be needed, i.e., the value of $\gamma_k$ drop to small values quite rapidly as $k$ increases. If we ask for a high degree polynomial and continue the recurrence (6.18) beyond the point where the expansion has converged (indicated by small value of $\gamma_k$) we will essentially iterate with noise. As it turns out, this noise is amplified by the recurrence. This is because the coefficient $\psi_k - \zeta_k$ becomes just noise and this causes the recurrence to diverge. An easy remedy is to just stop iterating (6.18) as soon as two consecutive $\gamma_k$’s are small. This takes care of two issues at the same time. First, it determines a sort of optimal degree to be used. Second, it avoids the unstable behavior observed by continuing the recurrence. Specifically, a test such as the following is performed:

$$|\gamma_{k-1}| + |\gamma_k| \leq k \cdot \text{tol},$$  \hspace{1cm} (6.20)

where tol is a small tolerance, which can be set to $10^{-6}$ for example. The factor $k$ reflects the scaling $k + 1/2$ in (6.19).

With this we can now easily develop an adaptation of Algorithm 3 which we will call the Delta-Gauss-Legendre (DGL) expansion algorithm. In the DGL algorithm, we will refer to formula (5.4). But now $p_M$ is the $M$-degree polynomial

$$p_M(\lambda) = \frac{1}{(2\pi \sigma^2)^{1/2}} \sum_{k=0}^{M} \left( k + \frac{1}{2} \right) \gamma_k L_k(\lambda),$$  \hspace{1cm} (6.21)

obtained by truncating the sum (6.19) to $M + 1$ terms.

In Figure 6.1 we show the approximate spectral density for the modified 2D Laplacian matrix constructed by the DGL method. The expansion coefficient $\gamma_k(t)$ are obtained by setting $\sigma = 0.56$ (obtained by setting $\kappa = 1.25, h = 0.75$).

**7. Use of the Lanczos Algorithm.** Because finding a highly accurate DOS essentially amounts to computing all eigenvalues $A$, any method that can provide approximations to the spectrum of $A$ can be used to construct an approximate DOS also. Since the Lanczos algorithm yields good approximations to extreme eigenvalues, it is a good candidate for computing localized spectral densities at both ends of the spectrum. In this section, we show that it is also possible to combine the Lanczos algorithm with multiple randomly generated starting vectors to construct a good approximation to the complete DOS.

Algorithm 4: Multi-point Delta-Gauss-Legendre expansion.

Input: Real symmetric matrix $A$. A set of points $\{t_i\}$ at which the DOS is to be evaluated, and $M_{\text{max}}$ is the maximum degree employed for all the points.

Output: Approximate DOS $\{\tilde{\phi}_M(t_i)\}$.

1: for each $t_i$ do
2: Compute and store the expansion coefficients $\gamma_{k,i}$ for $k = 0, \ldots, M$;
3: end for
4: for $l = 1 : n_{\text{vec}}$ do
5: Select a new random vector $v_0^{(l)}$;
6: for $k = 0 : M_{\text{max}}$ do
7: Compute $v_{k+1}^{(l)}$ via the three-term recurrence $v_{k+1}^{(l)} = 2Av_k^{(l)} - v_{k-1}^{(l)}$ (for $k = 0$,
$v_1^{(l)} = Av_0^{(l)}$);
8: Compute $\eta_k \leftarrow \eta_k + \left(v_0^{(l)}\right)^T v_k^{(l)}$;
9: end for
10: end for
11: Set $\eta_k = \eta_k/n_{\text{vec}}$ for all $k = 0, 1, ..., M_{\text{max}}$;
12: Evaluate $\tilde{\phi}_M(t_i)$ using $\eta_k$ and the stored $\gamma_{k,i}$'s;

Fig. 6.1: The approximate spectral density constructed by the Delta-Gauss-Legendre method for the modified 2D Laplacian matrix example. The same value $\sigma = 0.56$ as that used in the Gaussian smearing of the exact density is used in DGL. The polynomial degree is 80.

7.1. Lanczos approximation to the spectral density. For a given starting vector $v_0$, an $M$-step Lanczos procedure for a real symmetric matrix $A$ can be succinctly described by the following equations

$$AV_M = V_MT_M + f e_{M+1}^T, \quad V_M^TV_M = I_M, \quad V_M^Tf = 0, \quad V_Me_1 = v_0,$$

here $T_M$ is an $(M + 1) \times (M + 1)$ tridiagonal matrix, $V_M$ is an $n \times (M + 1)$ orthogonal matrix, and $I_M$ is an $(M + 1) \times (M + 1)$ identity matrix. It is well known that the $k$-th column of $V_M$ can be expressed as

$$V_Me_k = p_{k-1}(A)v_0, \quad k = 1, \ldots, M + 1.$$
where \( \{ p_k(t) \} \), \( k = 0, 1, 2, \ldots, M \) is a set of orthogonal polynomials with respect to the weighted spectral distribution \( \phi_{v_0}(t) \) taking the form

\[
\phi_{v_0}(t) = \sum_{j=1}^{n} \beta_j^2 \delta(t - \lambda_j). \quad (7.2)
\]

Here \( v_0 \) is assumed to be expanded in the basis of the eigenvectors of \( A \) as in Eq. (4.1).

It is also well known that these orthogonal polynomials can be generated by a three-term recurrence whose coefficients are defined by the matrix elements of \( T_M \) [22].

If \( (\theta_k, y_k) \), \( k = 0, 1, 2, \ldots, M \) are eigenpairs of the tridiagonal matrix \( T_M \), and \( \tau_k \) is the first entry of \( y_k \), then the distribution function defined by

\[
\sum_{k=0}^{M} \tau_k^2 \delta(t - \theta_k), \quad (7.3)
\]

serves as an approximation to the weighted spectral density function \( \phi_{v_0}(t) \), in the sense that

\[
\sum_{j=1}^{n} \beta_j^2 p_q(\lambda_j) = \sum_{k=0}^{M} \tau_k^2 p_q(\theta_k), \quad (7.4)
\]

for all polynomials of degree \( 0 \leq q \leq 2M + 1 \). The moment matching property described by (7.4) is well known [24]. It is the basis of Gaussian quadrature rules [23, 25].

Since in most cases, we are interested in the standard spectral density defined by (1.1), we would like to choose a starting vector \( v_0 \) such that \( \beta_j^2 = 1/n \). However, this is generally not possible without knowing the eigenvectors \( \{ u_j \} \) of \( A \) in advance. To address this issue, we resort to the same stochastic approach we used in previous sections.

We repeat the Lanczos process with multiple randomly generated starting vectors \( v_0^{(l)}, l = 1, 2, \ldots, n_{\text{vec}} \). We assume the average

\[
\frac{1}{n_{\text{vec}}} \sum_{l=1}^{n_{\text{vec}}} \left( v_0^{(l)} \right)^T \delta(tI - A) v_0^{(l)} = \frac{1}{n_{\text{vec}}} \sum_{l=1}^{n_{\text{vec}}} \sum_{j=1}^{n} \left( v_0^{(l)} \right)^T u_j \delta(t - \lambda_j) u_j^T v_0^{(l)}
\]

\[
= \frac{1}{n} \sum_{j=1}^{n} \left( \frac{1}{n_{\text{vec}}} \sum_{l=1}^{n_{\text{vec}}} \left( \beta_j^{(l)} \right)^2 \delta(t - \lambda_j) \right)
\]

\[
= \frac{1}{n} \sum_{j=1}^{n} \left( \beta_j \right)^2 \approx 1/n. \quad (7.5)
\]

is indeed a good approximation to the standard spectral density \( \phi(t) \) in Eq. (1.1), i.e.

\[
\frac{1}{n_{\text{vec}}} \sum_{l=1}^{n_{\text{vec}}} \left( \beta_j^{(l)} \right)^2 \approx 1/n.
\]

Since each distribution (7.3) generated by the Lanczos procedure is a good approximation to (7.3), then taking the average of (7.3) over \( l \), i.e.

\[
\tilde{\phi}(t) = \frac{1}{n_{\text{vec}}} \sum_{l=1}^{n_{\text{vec}}} \left( \frac{1}{n} \sum_{k=0}^{M} \tau_k^{(l)} \delta(t - \theta_k^{(l)}) \right)
\]

\[
= \frac{1}{n_{\text{vec}}} \sum_{l=1}^{n_{\text{vec}}} \left( \frac{1}{n} \sum_{k=0}^{M} \tau_k^{(l)} \delta(t - \theta_k^{(l)}) \right) \quad (7.6)
\]
should yield a good approximation to the standard spectral density (1.1).

Clearly, the resolution of the approximation to (7.3) given by (7.6) becomes better as $M$ increases. We may also replace $\delta$-functions in (7.6) with its surrogate version using Gaussian functions, i.e.

$$
\tilde{\phi}_\sigma(t) = \frac{1}{n_{\text{vec}}} \sum_{l=1}^{n_{\text{vec}}} \left( \frac{1}{n} \sum_{k=0}^{M} \left( \tau_k^{(l)} \right)^2 \phi_\sigma(t - \theta_k^{(l)}) \right).
$$

This will also allow us to evaluate the spectral density at some point that is different from a Ritz value $\theta_k$. The method using Eq. (7.7) will be referred to as the Lanczos method for computing DOS in the following discussion. In Figure 7.1, we show the approximate density of states of the modified 2D Laplacian obtained from Eq. (7.7) running $M = 80$ Lanczos steps.

Fig. 7.1: The approximate spectral density constructed by the Lanczos method for the modified 2D Laplacian matrix. The same $\sigma = 0.56$ is used in the Gaussian smearing of both the exact DOS, and the approximation in (7.7).

An alternative way to refine the Lanczos based DOS approximation from a $M$-step Lanczos run is to first construct an approximate cumulative spectral density or cumulative density of states (CDOS) defined in (5.6). The approximate CDOS can be written as

$$
\tilde{\psi}(t) = \int_{-\infty}^{t} H(t - s) \tilde{\phi}(s) ds = \sum_{k=0}^{M} \eta_k^2 \delta(t - \theta_k),
$$

where $\eta_k^2 = \sum_{i=1}^{k} \tau_i^2$, and $\theta_k$ and $\tau_k$ are eigenvalues and the first components of the eigenvectors of the tridiagonal matrix $T_M$ defined in (7.4). This approximation is plotted as a staircase function in Figure 7.2 for the modified 2D Laplacian. Note that both $\psi(t)$ and $\tilde{\psi}(t)$ are monotonically non-decreasing functions. Furthermore, it can been shown [33] that $\psi(t) - \tilde{\psi}(t)$ has precisely $2M - 1$ sign changes within the spectrum of $A$. A sign change occurs when $\psi(t)$ crosses either a vertical or horizontal step of $\tilde{\psi}(t)$. These properties allow us to construct an “interpolated” CDOS that matches $\psi(t)$ and $\tilde{\psi}(t)$ at the points where $\psi(t)$ crosses $\tilde{\psi}(t)$.

Since we do not know $\psi(t)$, we do not know exactly where it crosses $\tilde{\psi}(t)$ in advance. However, it can be seen from the left subfigure in Figure 7.2 that these crossing points are often near the midpoint of a vertical or horizontal step. Therefore, we may
use these midpoints to construct an interpolating function that is monotonically non-decreasing. For many practical problems, $\psi(t)$ approaches a smooth function as the dimension of the problem increases. Therefore, we can impose some constraints on the smoothness of the interpolating function in addition to monotonicity. One way to do this is to use the monotone splines introduced in [20, 21], and other choices of interpolating functions are also possible. The interpolation procedure allows us to evaluate the CDOS at any point within $[\lambda_1, \lambda_n]$. A recipe for constructing such an approximation is also given in [19]. The right subfigure in Figure 7.2 shows an interpolated approximate CDOS evaluated at 20 points evenly spaced points in $[\lambda_1, \lambda_n]$ for the modified Laplacian matrix. We observe that the interpolated CDOS is very close to the true CDOS at this resolution.

From the smoothly interpolated CDOS approximation, we can obtain an approximate DOS at any point in $[\lambda_1, \lambda_n]$ by taking the derivative of the interpolated $\tilde{\psi}(t)$ with respect to $t$. We can use either the analytic derivative of the spline or simply perform a finite difference calculation. Figure 7.3 shows the approximate DOS obtained in this fashion. We observe that the approximate DOS captures the basic profile of the surrogate DOS smeared by Gaussian functions.

The quality of the approximation also improves systematically as the number of starting vectors $n_{\text{vec}}$ increases. Figure 7.3 shows the averaged DOS approximations produced from 100 Lanczos runs looks much better than the approximation obtained from a single Lanczos run.

Furthermore, by carefully choosing $v_0$, we can also use this technique to obtain a localized DOS approximation. For example, if we are interested in the DOS at the low end of the spectrum, we can apply a low pass polynomial filter in $A$ first to a randomly chosen $v_l$. Figure 7.4 shows that this technique indeed provide a good approximation to the local DOS.

7.2. Rational transformation. In addition to approximating the spectrum of $A$ directly, the Lanczos algorithm can also be used to provide a rational approximation to the DOS, through the use of a well known technique for approximating a $\delta$-function by a rational function. It follows from the Sokhotski-Plemelj formula [33]

$$\lim_{t \to 0} \frac{1}{t + iy} = P(1/t) - i\pi\delta(t)$$
Fig. 7.3: The approximate DOS of the modified 2D Laplacian matrix obtained from the derivative of the CDOS approximation produced by a 20-step Lanczos procedure. The left subfigure is obtained from a single Lanczos run. The right subfigure is obtained from the average of 100 Lanczos runs with different random starting vectors.

Fig. 7.4: The approximation to the local DOS at the low end of the spectrum of the modified 2D Laplacian using a filtered initial starting vector.

that a δ-function can be expressed as the imaginary part of the rational function $1/(t + i\eta)$ in the limit that $\eta \to 0$, and $\mathcal{P}$ represents the Cauchy principal value. As a result, we may express the DOS of $A$ by

$$
\phi(t) = \lim_{\eta \to 0^+} -\frac{1}{n\pi} \text{Im} \sum_{j=1}^{n} \frac{1}{t + i\eta - \lambda_j}
$$

$$
= \lim_{\eta \to 0^+} -\frac{1}{n\pi} \text{Im} \text{Tr} [(tI - A + i\eta)^{-1}].
$$

It follows from the above discussion that estimating the spectral density of $A$ is equivalent to computing the trace of $(tI - A + i\eta I)^{-1}$. We can use the same type of stochastic approach used previously in the KPM to compute the approximate trace by

$$
\frac{1}{n_{\text{vec}}} \sum_{l=1}^{n_{\text{vec}}} (v_0^{(l)})^T (tI - A + i\eta I)^{-1} v_0^{(l)},
$$

for several randomly sampled vectors $v_0$. 

A direct calculation of (7.10) requires solving linear systems $[A - (t_i + i\eta)I]z = v_0$ repeatedly for each point $t_i$ at which the spectral density is to be evaluated. This approach can be prohibitively expensive. We now describe an alternative approach that allows us to approximate $v_0^T(t_i I - A + i\eta I)^{-1}v_0$ for multiple $t_i$’s at the cost of performing a single Lanczos factorization and some additional calculations that are much lower in complexity. If $v_0$ is used as the starting vector of the Lanczos procedure, then it follows from the shift-invariant property of the Lanczos algorithm that

$$
[A - (t_i + i\eta)I]v_M = V_M [T_M - (t_i + i\eta)I] + f c_{M+1}^T,
$$

(7.11)

where $V_M$ and $T_M$ are the same orthonormal and tridiagonal matrices respectively that appear in (7.11). After multiplying (7.11) from the left by $[A - (t_i + i\eta)I]^{-1}$, from the right by $[T_M - (t_i + i\eta)I]^{-1}$ and rearranging terms, we obtain

$$
[A - (t_i + i\eta)I]^{-1}V_M = V_M [T_M - (t_i + i\eta)I]^{-1} - [A - (t_i + i\eta)I]^{-1}f c_{M+1}^T [T_M - (t_i + i\eta)I]^{-1}.
$$

It follows that

$$
v_0^T [A - (t_i + i\eta)I]^{-1}v_0 = e_1^T V_M^T [A - (t_i + i\eta)I]^{-1} V_M e_1
\quad = e_1^T [T_M - (t_i + i\eta)I]^{-1} e_1 + \xi,
$$

where $\xi = - (v_0^T [A - (t_i + i\eta)I]^{-1} f) (c_{M+1}^T [T_M - (t_i + i\eta)I]^{-1} e_1)$. If $\xi$ is sufficiently small, computing $v_0^T(t_i I - A + i\eta I)^{-1}v_0$ reduces to computing the $(1,1)$-th entry of the inverse of $T_M - (t_i + i\eta)I$. Because $T_M$ is tridiagonal with $\alpha_1, \alpha_2, ..., \alpha_{M+1}$ on the diagonal and $\beta_2, \beta_3, ..., \beta_{M+1}$ on the sub-diagonals and super-diagonals, $e_1^T(zI - T_M)^{-1}e_1$ can be computed in a recursive fashion using a continued fraction formula

$$
e_1^T(zI - T_M)^{-1}e_1 = \frac{1}{z - \alpha_1 + \frac{\beta_2}{z - \alpha_2 + \cdots}}.
$$

(7.12)

This formula can be verified from the identity

$$
(T_M)^{-1}_{1,1} \equiv \frac{\det(zI - T_M)}{\det(z - T_M)}
$$

where $T_M$ is the trailing submatrix starting from the $(2, 2)$ entry of $T_M$, (7.12) and tridiagonal structure of both $T_M$ and $T_M$ matrices.

This approach is often referred to as Haydock’s method, and was first suggested by Haydock, Heine, and Kelly [26]. It is also related to the generation of Sturm sequences which is used in bisection methods for computing eigenvalues of tridiagonal matrices [12]. The computational cost of computing $e_1^T(T_M - z_iI)^{-1}e_1$ is $O(M + 1)$ for each shift $z_i = t_i + i\eta$ regardless whether this is done by applying a sparse direct method to compute $(T_M - z_iI)^{-1}e_1$ or through the use of a continued fraction. For most problems, this cost is small compared to that required to perform the Lanczos procedure to obtain $T_M$. Even in exact arithmetic, Haydock’s method produces an exact $v_0^T(z_i I - A)^{-1}v_0$ only when the last term on the right-hand side of (7.11) is zero. This term vanishes when $f = 0$. In this case, columns of $V_M$ form an invariant subspace of $A$. However, for most problems, $f$ rarely becomes zero even for large $M$. But when $\eta$ is relatively large, the off-diagonal elements of $(z_i I - T_M)^{-1}$ decreases sufficiently fast, away from the diagonal. Therefore, the $v_0^T(z_i I - A)^{-1}v_0$ can be evaluated accurately in this case.
The performance of Haydock's approach for computing the spectral density for the modified 2D Laplacian example is shown in Fig. 7.5. The parameter \( \eta \) is chosen to be comparable to the resolution of the spectral density we display. Specifically, we choose

\[
\eta = \frac{\varepsilon_{\text{max}} - \varepsilon_{\text{min}}}{2n_x},
\]

where \( \varepsilon_{\text{max}} \) and \( \varepsilon_{\text{min}} \) are the largest and smallest eigenvalues of \( A \) respectively, and \( n_x \) is the number of uniformly distributed grid points at which the spectral density is evaluated. In Fig. 7.5 we choose \( n_x = 55 \), which corresponds to \( \varepsilon = 0.187 \). This is roughly one third of the value of \( \sigma = 0.56 \). We observe that running 40 Lanczos steps produces a qualitatively different spectral density approximation. The quality of the approximation becomes much better when 100 Lanczos steps are taken in Haydock's approach.

8. Numerical Results. In this section, we compare all methods discussed above for approximating the spectral density of \( A \) through numerical examples. To compare these methods quantitatively, we compute the \( L^1 \), \( L^2 \) and \( L^\infty \) distances between the approximate spectral density \( \phi(t) \) and the surrogate spectral density \( \phi_\sigma(t) \), and the latter is the true spectral density blurred by a Gaussian of the form (1.6) with a fixed \( \sigma \) value. Both \( \phi_\sigma(t) \) and \( \hat{\phi}(t) \) are normalized to have \( \|\phi_\sigma(t)\|_1 = \|\hat{\phi}(t)\|_1 = 1 \). It is well known that both the \( L^2 \) and \( L^\infty \) norms are sensitive to outliers in the spectral density approximation whereas the \( L^1 \) norm is relatively less sensitive to outliers.

8.1. Modified 2D Laplacian. In Table 8.1 we list the \( L^1 \), \( L^2 \) and \( L^\infty \) errors associated with different types of approximations to the spectral density of the modified Laplacian matrix we used in the previous sections to illustrate how these methods behave. The dimension of the matrix is 750, which is relatively small. We set \( \sigma \) to 0.56. This corresponds to setting \( \kappa \) to 1.25 and \( h = 0.75 \) in (2.1). The degree of Chebyshev or Legendre polynomials used in the KPM and DGL methods is set to 80. Similarly, the number of iterations in the Lanczos method is also set to 80. We also present the results for the spectroscopic method and Haydock’s method with the number of iterations being 40 and 100, respectively. We observe that the DGL, Lanczos and KPM with Jackson damping yield the most accurate approximation among
all methods in $L^1$, $L^2$ and $L^\infty$ norms, followed by the KPM method without Jackson damping. The spectroscopic and Haydock’s methods do not yield sufficiently accurate approximations if only 40 steps of Lanczos iterations are taken. In order to reach the same level of accuracy achieved by DGL and Lanczos in $L^1$ norm, at least 100 Lanczos iterations are required in these methods to further reduce the approximation error. In all test cases, we use 10 random vectors.

| Method                        | $L^1$ error | $L^2$ error | $L^\infty$ error |
|-------------------------------|-------------|-------------|------------------|
| KPM w/ Jackson, deg=80        | 2.994e-02   | 9.2190e-03  | 6.036e-03        |
| KPM w/o Jackson, deg=80       | 1.057e-01   | 3.3897e-02  | 2.409e-02        |
| KPM Legendre, deg=80          | 1.123e-01   | 3.4598e-02  | 2.628e-02        |
| Spectroscopic, deg=40         | 8.724e-02   | 2.6650e-02  | 1.566e-02        |
| Spectroscopic, deg=100        | 6.818e-02   | 2.1735e-02  | 1.395e-02        |
| DGL, deg=80                   | 3.031e-02   | 9.8346e-03  | 5.180e-03        |
| Lanczos, deg=80               | 2.047e-02   | 5.7810e-03  | 3.344e-03        |
| Haydock, deg=40               | 1.583e-01   | 5.9017e-02  | 4.631e-02        |
| Haydock, deg=100              | 8.074e-02   | 2.3214e-02  | 1.367e-02        |

Table 8.1: $L^1$, $L^2$, and $L^\infty$ error compared with the normalized “surrogate” DOS for modified 2D Laplacian matrix, using the same set of parameters as described in this section.

8.2. The Kohn-Sham Hamiltonian associated with the Benzene molecule.

To demonstrate that the methods presented in this paper are applicable to matrices other than the modified Laplacian, we now show the $L^1$, $L^2$ and $L^\infty$ error associated with different approximations to the spectral density of a matrix, which represents a finite difference approximation of the Kohn-Sham Hamiltonian associated with a benzene molecule. This matrix is generated from the PARSEC software [34]. The dimension of this matrix is 8,219. The matrix can be obtained from the Florida Sparse Matrix repository [12]. For this problem, we also set the $\sigma$ parameter for the smoothed exact spectral density to 0.56. We use 10 random Gaussian test vectors in all methods. Table 8.2 shows that the accuracy of different methods in $L^1$, $L^2$ and $L^\infty$ norms exhibits a similar pattern to that observed for the modified Laplacian matrix. The DGL, Lanczos and the KPM with Jackson damping still yield the most accurate results. Haydock’s method is less accurate compared to other methods for this example. The quality of the approximation provided by different methods can also be seen from Figures 8.1 to 8.5.

8.3. Convergence with respect to the number of random vectors. An interesting observation we made from the experimental comparison of the DGL method and the KPM is that DGL can produce reasonable results even for a very small number of random vectors. In contrast, the accuracy of KPM can vary significantly when only a few vectors are sampled. This is illustrated in Figure 8.6 which repeats the previous experiment using only one random vector. The curve on the left corresponds to KPM. It is one of many possible ones that vary quite a bit in shape. The one on the right corresponds to DGL. It remains more stable than KPM for multiple vectors used and captures the overall true spectral density quite well.

While using one random vector in KPM, DGL or Lanczos to obtain an approximate spectral density is an extreme case, we should expect the accuracy of the approx-
Table 8.2: $L^1$, $L^2$, and $L^\infty$ error compared with the normalized “surrogate” DOS for benzene matrix, using the same set of parameters as used in the pictures in this section.

| Item                        | $L^1$ error | $L^2$ error | $L^\infty$ error |
|-----------------------------|-------------|-------------|-------------------|
| KPM w/ Jackson, deg=80     | 2.592e-02   | 5.0329e-03  | 2.785e-03         |
| KPM w/o Jackson, deg=80    | 2.634e-02   | 4.4542e-03  | 2.002e-03         |
| KPM Legendre, deg=80       | 2.504e-02   | 3.7881e-03  | 1.174e-03         |
| Spectroscopic, deg=40      | 5.589e-02   | 8.6527e-03  | 2.871e-03         |
| Spectroscopic, deg=100     | 4.624e-02   | 7.5829e-03  | 2.447e-03         |
| DGL, deg=80                | 1.998e-02   | 3.3793e-03  | 1.149e-03         |
| Lanczos, deg=80            | 2.755e-02   | 4.1785e-03  | 1.599e-03         |
| Haydock, deg=40            | 6.951e-01   | 1.3023e-01  | 6.176e-02         |
| Haydock, deg=100           | 2.581e-01   | 4.6530e-02  | 1.420e-02         |

Fig. 8.1: The approximate spectral density constructed by KPM for the benzene Hamiltonian. Chebyshev polynomials of degree 80 with Jackson damping (left) and without Jackson damping (right) are used in KPM.

Fig. 8.2: The approximate spectral density for the benzene Hamiltonian obtained from KPM. A 80-degree Legendre polynomial is used in KPM.
Fig. 8.3: The approximate spectral densities obtained from the Lanczos spectroscopic method in which 40 (left) and 100 (right) steps of the Lanczos algorithm are taken respectively.

Fig. 8.4: The approximate spectral densities for the benzene Hamiltonian obtained from the Delta-Gauss-Legendre method and the Lanczos-Delta-Gauss fit. The $\sigma$ value used in the Gaussian approximation of the $\delta$-function in both methods is chosen to be the same as that used in the Gaussian smearing of the exact density.

Fig. 8.5: The approximate spectral densities for the benzene Hamiltonian obtained from Haydock’s method. The number of Lanczos step taken are 40 (left) and 100 (right). The smoothing parameter $\eta$ is set to $\eta = 0.178$. 
Fig. 8.6: KPM with degree 80 (left) and, Delta-Gauss-Legendre expansion with degree 80 ($\sigma = 0.56$) (right) for perturbed 2D Laplacian operator. In contrast with Figure 6.1 here only one random vector is used.

imation to improve systematically as we average approximations obtained from $n_{\text{vec}}$ random vectors. This should be the case at least for the DGL and Lanczos methods, since these two methods are indeed based on the expansion of the surrogate spectral density. If the variance of the DGL or Lanczos approximation produced by sampling one random vector is denoted by $\text{var}$, then averaging $n_{\text{vec}}$ approximations produced by $n_{\text{vec}}$ independent random vectors can reduce the variance of the approximation to $\text{var}/n_{\text{vec}}$, and the standard deviation to $\sqrt{\text{var}/n_{\text{vec}}}$. In other words, the $L^1$, $L^2$ and $L^\infty$ error should be proportional to the standard deviation and decreases as $n_{\text{vec}}^{-\frac{1}{2}}$. Fig. 8.4 shows that, for the modified 2D Laplacian matrix with $\sigma = 0.56$, the $L^1$, $L^2$ and $L^\infty$ errors associated with DGL, Lanczos and KPM (Chebyshev expansion with Jackson damping) all decrease as the number of random vectors used ($n_{\text{vec}}$) increases. We observe that the convergence with respect to $n_{\text{vec}}$ is roughly on the order of $O(n_{\text{vec}}^{-\frac{1}{2}})$. For methods (such as the KPM) that do not directly expand the surrogate DOS $\phi_{\sigma}$, there is some positive distance between the result obtained from KPM and that obtained from the surrogate DOS even when $n_{\text{vec}}$ becomes large, although the absolute value of the error can be small as $10^{-3} \sim 10^{-2}$.

8.4. DGL and Lanczos for other matrices. We now show that in addition to the modified Laplacian and the Kohn-Sham Hamiltonian for the benzene molecule, the DGL and Lanczos methods work equally well for a variety of problems listed in Table 8.3. The $\text{Ga}_{10}\text{As}_{10}\text{H}_{30}$ matrix arises from a Kohn-Sham density functional theory calculation of a Ga and As cluster. It has a relatively large dimension. The PE3K matrix arises from the molecular vibration analysis of a polyethylene molecule with 3000 atoms [58]. The CFD1 matrix arises from a computational fluid dynamics calculation. The SHWATER matrix arises from another computational fluid dynamics application involving weather shallow water equations. In order to evaluate the accuracy of the DGL and Lanczos methods, we compute all eigenvalues of these matrices in advance using the ScaLAPACK software. The algebraically smallest and largest eigenvalues are also listed in Table 8.3. Table 8.4 compares the $L^1$, $L^2$ and $L^\infty$ errors for the DGL and Lanczos methods for these problems.

As can be seen from the table, both the DGL and the Lanczos methods yield accurate approximations to the spectral densities of these matrices. We overlay the
Fig. 8.7: The $L^1$, $L^2$ and $L^\infty$ errors for the DGL, Lanczos methods and the KPM (Chebyshev expansion with Jackson damping) with respect to the number of random vectors used ($n_{vec}$). The matrix chosen is the modified 2D Laplacian matrix with $\sigma = 0.56$.

Table 8.3: Description of the size and the spectrum range of the test matrices.

| Matrix          | $n$  | $\lambda_1$ | $\lambda_n$ |
|-----------------|------|--------------|--------------|
| Ga$_{10}$As$_{10}$H$_{30}$ | 113,081 | $-1.2$       | $1.3 \times 10^3$ |
| PE3K            | 9,000 | $8.1 \times 10^{-6}$ | $1.3 \times 10^2$ |
| CFD1            | 70,656 | $2.0 \times 10^{-5}$ | 6.8          |
| SHWATER         | 81,920 | 5.8          | $2.0 \times 10^1$ |

computed spectral density (by DGL) on top of the “exact” spectral density for both the CFD1 and SHWATER matrices in Figure 8.8. As can be seen, the computed spectral densities are nearly indistinguishable from the exact ones. The same observation is made for Ga$_{10}$As$_{10}$H$_{30}$, which shows a similar spectral density profile as that associated with the benzene molecule, and the PE3K molecule, which we will show in the next section.

Fig. 8.8: The approximate spectral densities of the CFD1 and SHWATER matrices obtained from the DGL method match well with the corresponding smoothed “exact” spectral density.
8.5. Heat capacity calculation. As we mentioned in section 1, one of the utilities of the DOS is that it provides an appropriate measure for integrating certain physical quantities of interest. One good example is the heat capacity defined in (1.4). Figure 8.9 shows that, the approximate spectral density allows us to captures the variation of the heat capacity with respect to temperature very well.

Fig. 8.9: The spectral density of the PE3K matrix and the variation of the computed heat capacity with respect to temperature change. The computed heat capacities are obtained from using the exact (red) and the DGL approximate (blue) spectral densities respectively.

9. Conclusion. We presented a number of algorithms for estimating the spectral density of a Hermitian matrix $A$ from a numerical linear algebra perspective. Some of these algorithms (KPM, DC and DGL) are based on constructing polynomial approximations to $\delta$-functions. In these approaches, estimating the spectral density essentially amounts to approximating the trace of $A$ or $p(A)$ where $p(t)$ is a polynomial. An important technique for approximating the trace of a Hermitian matrix is stochastic sampling and averaging of the Rayleigh quotient $v_0^T p(A)v_0$, with $\|v_0\|_1 = 1$.

A few other methods make use of the Lanczos procedure to either directly estimate the spectrum of $A$ or to construct a rational approximation of the $\delta$-function. Stochastic sampling of the starting vector of the Lanczos process often improve the
fidelity of the approximation. However, if the number of Lanczos steps used in these methods is sufficiently large, even one starting vector may be sufficient to produce a reliable estimation of the spectral density.

We suggest that it is more appropriate to approximate a smoothed version of the spectral density, which we sometimes call a surrogate spectral density, instead of the exact spectral density given by \( \delta \) directly. We can view this approach as a regularization technique to attenuate the numerical difficulties associated with approximating \( \delta \)-functions directly. The smoothness of the surrogate spectral density can be controlled by one parameter \( \sigma \). However, there is a trade-off between the amount of smoothness of the surrogate (hence the cost of the approximation) and the accuracy of the approximation. When \( \sigma \) is chosen to yield a very smooth surrogate spectral density, accurate approximations to this surrogate can often be obtained easily with a relatively low degree polynomial or a small number of Lanczos iterations. However, the surrogate itself may miss some important features of the exact spectral density required in some applications. On the other hand, if \( \sigma \) is chosen to minimize the difference between the surrogate and \( \delta \), then more work is required obtain a good approximation. In the extreme case when the exact spectral density given by \( \delta \) is to be obtained, then all eigenvalues of the matrix \( A \) must be computed.

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