Interior state computation of nano structures

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Abstract. We are concerned with the computation of electronic and optical properties of quantum dots. Using the Energy SCAN (ESCAN) method with empirical pseudopotentials, we compute interior eigenstates around the band gap which determine their properties. Numerically, this interior Hermitian eigenvalue problem poses several challenges, both with respect to accuracy and efficiency. Using these criteria, we evaluate several state-of-the art preconditioned iterative eigensolvers on a range of CdSe quantum dots of various sizes. All the iterative eigensolvers are seeking for the minimal eigenvalues of the folded operator with reference shift in the band-gap. The tested methods include standard Conjugate-Gradient (CG)-based Rayleigh-Quotient minimization, Locally Optimal Block-Preconditioned CG (LOBPCG) and two variants of the Jacobi Davidson method: JDQMR and GD+1. Our experimental results conclude that the Jacobi Davidson method is often the fastest.

Keywords: comput. nanotechnology, parallel eigensolvers, quantum dots

1 Introduction

The computation of electronic properties of large nano structures such as quantum dots is an important field of current research. In this paper, we are interested in electronic structure calculations based on the Kohn-Sham approximations and leading to the solution of the effective single-particle Schrödinger equations

$$H\Psi_i \equiv \left[-\frac{1}{2}\nabla^2 + V \right] \Psi_i = E_i \Psi_i. \tag{1}$$

Here H denotes the approximate Hamiltonian. The atomic system is described by the potential V which we assume as externally given empirical pseudopotential [1]. The set $\{\Psi_i\}$ denotes the orthogonal wave-functions (eigenstates) and $\{E_i\}$ their corresponding energies. If the system has an electron with energy E_i , then $\Psi_i(r)$ describes the spatial probability distribution for the electron.

In the context of the Self-Consistent Field iteration [8], a large number of eigenstates of (1) need to be computed [11]. On the other hand, only a *small* number of eigenstates of (1) are often relevant for determining certain optical

and electronic properties. Our task is to compute these states at the top of the valence and at the bottom of the conduction band to find the band gap. These states typically lie in the middle of the spectrum with too large a distance from the lowest (ground) state to efficiently compute all of them from the lowest up.

In a discrete finite plane-wave basis, Equation (1) directly translates into a Hermitian eigenvalue problem and we use the same notation H for the matrix in question. The discrete Laplacian is commonly computed in the plane wave basis because it is diagonal. On the other hand, the discretized potential is commonly available only in real space. Thus the resulting H becomes only implicitly available through matrix-vector products and via the Fast Fourier Transform (FFT). This together with the system size requires the use of efficient iterative eigenvalue methods.

Our parallel ESCAN method [1] uses a folded spectrum approach [17] to find the interior eigenstates that we are looking for. Based on physical knowledge about the system, a reference energy E_{ref} is chosen and then the smallest eigenvalues of the system $(H-E_{ref})^2$ currently are computed via PCG Rayleigh-Quotient minimization, see [17], the references in [5] and also [7, 3]. In our study, we compare this method with two other state-of-the art iterative eigensolvers, the LOBPCG [4] and some variants of the Jacobi-Davidson method [12].

The rest of this paper is organized as follows. Section 2 gives a brief overview of the mathematical and computational properties of the iterative eigensolvers being compared. Section 3 describes the setting for our experiments and the results of our evaluation. Our conclusions are given in Section 4.

2 Overview of eigenvalue methods

The collection of iterative methods studied in this paper is subdivided in two parts, the CG-based methods and the Jacobi-Davidson based methods.

The CG-based techniques successively minimize the Rayleigh quotient function $f(x_i) = (x_i^* A x_i)/(x_i^* x_i)$, where the gradient is given by $\nabla f(x_i) = A x_i - x_i (x_i^* A x_i)/(x_i^* x_i)$. Note that the gradient is the residual of the approximate eigenvectors x_i ($r_i = \nabla f(x_i)$). Together with a suitable preconditioner, this approach (PCG) has been used to solve a number of problems of practical interest. In a blocked form the minimization is simultaneously applied to a set of orthogonal vectors X_i . The LOBPCG method [16, 4] extends PCG by applying Rayleigh Ritz on $span(W_i, X_i, X_{i-1})$, where $W_i = P\nabla f(X_i) = Pr_i$ for an appropriate preconditioner P.

More recently, variants of Davidson's method [2] proved to be very effective for solving a number of applications. Davidson's original algorithm targeted mainly the smallest eigenvalues of (strongly diagonal dominant) matrices arising from applications in Chemistry. It relies on the fact that in the case of strongly diagonal dominant matrices the inverse of the diagonal entries of this matrix lead to a good approximation for the inverse of the matrix, i.e. $diag(A)^{-1} \approx A^{-1}$. In this case, given an approximate eigenpair $(\hat{x}, \hat{\theta})$ and the residual $r = (\hat{\theta}I - A)\hat{x}$, the vector obtained from $t = (\hat{\theta}I - diag(A))^{-1}r$ can be used to generate a new

vector to be added to the search subspace S. However, depending on A and $\hat{\theta}$, the problem $t = (\hat{\theta} - diag(A))^{-1}r$ may be ill-conditioned (in particular note that this problem is singular if $\hat{\theta}$ is an eigenvalue of A. To remedy this situation, an orthogonal component correction has been proposed in conjunction with Davidson's algorithm [12]. This correction replaces $t = (\hat{\theta} - diag(A))^{-1}r$ with an approximate solution of the correction equation

$$(I - \hat{x}\hat{x}^*)(A - \hat{\theta}I)(I - \hat{x}\hat{x}^*)t = -r, \ x \perp t.$$
 (2)

This equation can be solved by using the QMR algorithm, for example. This allows for the generation of more general search subspaces than the original algorithm. In addition, whereas Davidson's algorithm requires preconditioners that accurately approximate the inverse of the given operator and that may in fact lead to ill-conditioned problems, the solver for the correction equation (2) can take great advantage of such preconditioners. We derive our Jacobi-Davidson based method for ESCAN from the JaDa software (§2.5).

2.1 Folded spectrum method

Currently, all our methods are only able to find eigenvalues at the extreme of the spectrum. Since we are interested only in the interior eigenvalues, we need to apply a spectral transformation such that the interior eigenvalues become extreme eigenvalues of another operator, then find these extreme eigenvalues and transform them back to the original problem.

This approach is classically performed using a polynomial transformation, for a detailed discussion see for example [10,16]. In our case we simply choose the polynomial of second order $(x - E_{ref})^2$. The eigenvectors corresponding to the minimum eigenvalues of $(H - E_{ref}I)^2$ are the eigenvectors of H corresponding to the eigenvalues that are closest to E_{ref} . This approach is called folded spectrum.

Other polynomials can be used for example, the shift-and-invert technique [9] considers the polynomial $(x-E_{ref})^{-1}$, where the inverse operation is done by an inner iterative linear solver at each step. In this case, the Lanczos and Arnoldi algorithms [6] can be applied to the shifted and inverted operator $(H-E_{ref}I)$, for a fixed E_{ref} of interest. Although finding the extreme eigenvalues of $(H-E_{ref}I)^{-1}$ requires very few iterations, applying the operator $(H-E_{ref}I)^{-1}$ at each step is extremely expensive and we have not found this method effective in practice.

Even though the polynomial $(x - E_{ref})^2$ transforms interior eigenvalues to minimum eigenvalues, it also clusters quadratically those eigenvalues around E_{ref} , unfortunately. Typically the convergence with the operator $(H - E_{ref}I)^2$ is much slower than the one with H.

Finally we note that in theory, the Jacobi-Davidson method is able to aim effectively and directly at interior eigenvalues using an inner iteration scheme (e.g. QMR). We have not this feature available in the current version of the Jacobi-Davidson solver. This is work for the near future (see also [15]).

2.2 Preconditioner

All iterative methods studied in this paper allow for the use of a preconditioner. The goal of the preconditioner in the case of the folded spectrum method is to approximate the inverse of the matrix $(H - E_{ref}I)^2$ where H is given in Eq. 1. The preconditioner we use is diagonal and is applied in the Fourier space as

$$P = (I + (-\frac{1}{2}\nabla^2 + V_{avg} - E_{ref})/E_k)^2$$

where $-\frac{1}{2}\nabla^2$ is the Laplacian (diagonal in the Fourier space), E_{ref} is the shift used in the folded spectrum, V_{avg} is the average potential and E_k is the average kinetic energy of a given initial approximation of a wave function ψ_{init} [18].

2.3 Banded PCG Method

The Banded-PCG method [17] is the original method used in ESCAN (see also [7, 3] and Algorithm 1). This algorithm is well-suited for this kind of problem and has proven to be extremely efficient. The parameters of the algorithm are niter and nline. numEvals is the number of eigenvalues we are looking for and A is the folded matrix $(H - E_{ref}I)^2$.

The loop in j from line 5 to line 16 is simply nline iterations of the nonlinear Conjugate Gradient method applied to the Rayleigh Ritz minimization problem on the operator $(I-X(m)X(m)^*)A$ where X represents the already converged eigenvalues. The process is repeated for all the eigenvalues sought (loop in m from line 2 to 17) and a Rayleigh Ritz step and restart is performed as long as the convergence is not obtained (loop in i from line 1 to 19). niter represents the maximum number of outer iterations the algorithm is allowed to perform.

2.4 LOBPCG Method

LOBPCG stands for Locally Optimal Block-Preconditioned CG [4]. This methods is a block extension of the PCG method described in §2.3 with Rayleigh Ritz minimization (instead of conjugate gradient minimization). Briefly, the LOBPCG method can be described with the pseudo-code in Algorithm 2.

Note that the m and j loops from Algorithm 1 are replaced with just the blocked computation of the preconditioned residual, and the Rayleigh-Ritz on $\operatorname{span}\{X_i\}$ with Rayleigh-Ritz on $\operatorname{span}\{X_{i-1},X_i,R\}$. The direct implementation of this algorithm becomes unstable as X_{i-1} and X_i become closer and closer, and therefore special care and modifications have to be taken (see [4]). In the experiments presented in Section 3, we are using our own version of LOBPCG with a special treatment to work on the folded spectrum operator $(H - E_{ref}I)^2$ but to control the accuracy using the Hamiltonian matrix H.

Algorithm 1 PCG Eigensolver

```
1: for i = 1, niter do
 2:
        for m = 1, numEvals do
            orthonormalize X(m) to X(1:m-1)
 3:
 4:
            ax = A X(m)
 5:
            for j = 1, nline do
               \lambda(m) = X(m) \cdot ax
 6:
               if (||ax - \lambda(m) X(m)||_2 < tol .or. j == nline) exit
 7:
               r_{j+1} = (I - X(m) \ X(m)^*) \ ax
 8:
               \beta = \frac{r_{j+1} \cdot Pr_{j+1}}{r_j \cdot Pr_j}
d_{j+1} = -P \ r_{j+1} + \beta \ d_j
 9:
10:
                d_{j+1} = (I - X(m)X(m)^*)d_{j+1}
11:
               \gamma = ||d_{j+1}||_2^{-1}
12:
               \gamma = ||a_{j+1}||_2
\theta = 0.5 \left| atan \frac{2 \gamma d_{j+1} \cdot ax}{\lambda(m) - \gamma^2 d_{j+1} \cdot A d_{j+1}} \right|
13:
                X(m) = cos(\theta) X(m) + sin(\theta) \gamma d_{i+1}
14:
15:
               ax = cos(\theta) \ ax + sin(\theta) \ \gamma \ A \ d_{i+1}
16:
            end for
         end for
17:
18:
         [X, \lambda] = \text{Rayleigh-Ritz on span}\{X\}
19: end for
```

Algorithm 2 LOBPCG Eigensolver

```
1: for i = 1, niter do

2: R = P (A X_i - \lambda X_i)

3: check convergence criteria

4: [X_i, \lambda] = \text{Rayleigh-Ritz on span}\{X_i, X_{i-1}, R\}

5: end for
```

2.5 Jacobi Davidson Method

The Jacobi-Davidson method [12] used in our experiments is based on the JaDa software. The JDQMR code is a JDQR version using QMR based on the classical Jacobi-Davidson [12] while GD+k is based on [14]. The code is in C with additional functionality of the early block Fortran code described in [13].

JaDa is a multi-method code with a lot of parameters that enable us to simulate a number of methods. Among the useful parameters are $\mathtt{maxBasisSize}$ which represents the maximum size of the search space, $\mathtt{minRestartSize}$ which represents the number of vectors among the search space that are kept at each restart, $\mathtt{maxBlockSize}$ which represents the block size used in the method, $\mathtt{maxPrevRetain}$ which represents the number of vectors that are kept from the previous restart (useful to simulate $\mathtt{GD+k}$), $\mathtt{maxInnerIterations}$ and $\mathtt{relTolBase}$ which give stopping criteria for the inner iterations with QMR. The inner iterations in the QMR solve are stopped when the number of iterations is greater than $\mathtt{maxInnerIterations}$ or when the residual is less than $\mathtt{relTolBase}^{\#}$ iterations.

Typically in our experiments, for the Jacobi-Davidson method, we will take

The block size will always be set to one, and we always will keep one vector from the previous restart (GD+1), i.e. $\mathtt{maxBlockSize} = 1$, $\mathtt{maxPrevRetain} = 1$.

Case maxInnerIterations = 0 gives rise to GD+1 (see [14]). When maxInnerIterations = 30, relTolBase = 2.0 the method is called JDQMR (see [12]). It is interesting to note that by setting the parameters of JaDa to

```
maxBasisSize = 3 * maxBlockSize
minRestartSize = maxBlockSize
maxBlockSize = numEvals
maxPrevRetain = maxBlockSize
```

(and having maxInnerIterations set to 0) then the obtained method is very close to LOBPCG. We call this variant JD-LOBPCG and our numerical experiments confirm that JD-LOBPCG behaves like a CG-method. If we want to use LOBPCG but not on the whole block of eigenvectors (when a large number of eigenvalues is requested for example), one can set maxBlockSize to a smaller value than numEvals, and then (hard-)locking will be used to to find all the eigenvalues. We call this methods JD-LOBPCG+L.

3 Test results

We perform experiments on a set of four quantum dots of Cadmium-Selenium of increasing size (see Figure 1 for a description of the quantum dots).

test case	# atoms	n	Time matvec (s)	
Cd20.Se19	39	11,331	0.005	
Cd83.Se81	164	34,143	0.014	
Cd232.Se235	467	75,645	0.043	
Cd534.Se527	1061	141,625	0.105	

Fig. 1. Time for a matrix-vector product as the number of atoms grows, see Fig. 2 for a graphic visualization. n is the size of the matrix in the Fourier space.

The goal of the experiments is to compare the different iterative methods in terms of time to solution and robustness.

The different methods compared are a set of four Conjugate-Gradient like methods: Banded-PCG, LOBPCG, JD-LOBPCG, JD-LOBPCG+L, and two Jacobi-Davidson like methods: GD+1 and JDQMR. See Section 2 for a description of the methods.

The default parameters used for JaDa are:

```
restarting.scheme='thick restart'
restart.target='smallest'
robustShifts='on'
```

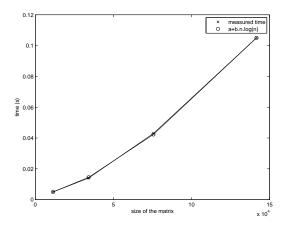


Fig. 2. Time for a matrix-vector product for some quantum dots of Cadmium Selenium of different size. The time for a matrix-vector product is growing as $n \log(n)$ where n is the size of the linear system as expected.

In terms of robustness, it is important to note that all of these methods are extremely robust. They all converge to the correct solutions. For our applications this also means that they find degenerate states without any particular problem.

All methods are looking for the numEvals = 10 eigenstates of highest energy in the valence band maximum (VBM). The shift for the folded spectrum is in -4.8eV and we use the diagonal preconditioner described in Section 2.2.

All the runs are performed on one node of IBM SP3 (seaborg.nersc.gov) which represents 16 processors in shared memory.

We use two stopping criteria. For the banded PCG and LOBPCG, the stopping criterion is based on H. This means that iterations are stopped when for all $i=1,\ldots,mx$ $\|H\psi_i-\psi_iE_i\|\leq tol$. For JaDa (JD-LOBPCG, JD-LOBPCG+L, JD-QMR, GD+1), the stopping criterion used is based on the folded matrix $(H-E_{ref}I)^2$. This means that iterations are stopped when for all $i=1,\ldots,mx$ $\|(H-E_{ref})^2\psi_i-\psi_i\lambda_i\|\leq tol$. These two stopping criteria differ slightly in practice and our experiments show that stopping the iteration with $(H-E_{ref}I)^2$ implied that the error with H is always smaller than $5 \cdot tol$.

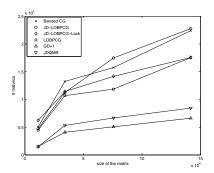
Various sets of parameters have been tested for GD+1 and JDQMR in all the cases, but we can see that there are very few influences on the final time.

Figure 3 gives a summary of our results. As the size of the quantum dots grows we see that the problems become harder and need more matrix-vector products. This result was expected. The Figure also clearly shows two trends represented correspondingly by the class of CG-based methods and the class of Jacobi-based methods. The Jacobi-Davidson based methods clearly outperform the CG-based methods either in terms of number of matrix-vector products or in time to solution.

The slope in terms of matrix-vector products (see Figure 3, Left) for the Jacobi-Davidson is almost flat, which shows that the methods scale numerically very well as the size of the quantum dots increases. A general remark is that if one is concerned only with the number of matrix-vector products until convergence then the best methods in our study is always GD+1.

The slope for the Jacobi-Davidson methods in terms of time is not flat, but one has to remember that those experiments are performed on a fixed size machine. Since the time for a matrix-vector product increases linearly with the size of the problem (see Figure 2), we recover this linear increase in the slope of our methods.

It is noteworthy that the matrix-vector product operation is performed without blocking, the reason being that the blocked version we have is not more efficient on these size problems than the nonblocked version. Such behavior is a major drawback for methods like LOBPCG that take full advantage of the block matrix-vector product.



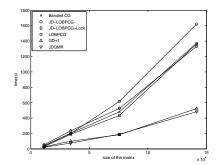


Fig. 3. Number of matrix-vector products (Left) and time in seconds (Right) for finding the 10 highest energies (and states) in the VBM for some quantum dots of Cadmium Selenium of various sizes. The folded spectrum method is used for all the methods with shift on $E_{ref} = -4.8 \text{eV}$. Figure 1 has a description of the quantum dots.

An interesting quantity to look at is the ratio of time that these methods are spending in the matrix-vector products. The remaining part of the time is mostly due to orthogonalization procedure. Therefore we can have a rough estimate on the ratio orthogonalization/matrix-vector product those methods are using. These ratios differ a lot from one method to the others (see Figure 4 for the results on the Cd83Se81 quantum dots). In the case of Figure 4, although the method with the lesser matrix-vector products is GD+1, it is not the fastest one. For this type of problem, matrix-vector products are relatively fast compared to orthogonalization schemes and thus a method with more matrix-vector products but far less orthogonalization (JDQMR) is faster than the optimal method in terms of matrix-vector products (GD+1).

Method	Time(s)	# matvecs	Time in	% in
			matvec(s)	matvec
Banded CG	236.13	15096	201.46	85.3%
LOBPCG	190.37	10688	146.11	76.8%
JDQMR	75.65	5314	73.20	96.7%
GD+1	100.87	4084	57.17	56.6%

Fig. 4. Time spent in the matrix-vector multiply operations for Cd83Se81.

4 Conclusions and possible extensions

This paper presents a comparison of eigensolvers in the context of the computation of electronic and optical properties of quantum dots using the ESCAN method with empirical pseudopotentials.

An evaluation of several state-of-the-art preconditioned iterative eigensolvers on a range of CdSe quantum dots of various sizes leads us to the conclusion that the Jacobi-Davidson type solvers were superior in terms of time and in terms of matrix-vector products for this range of problems. GD+1 is always the method that leads to the less matrix-vector products when the basis size is fixed. The new solvers are almost three times faster than the original method.

The current Jacobi Davidson method can be improved in several ways. In particular, we would like

- 1. to avoid the need to fold the spectrum when seeking for the interior eigenvalues,
- 2. to have a better control of the eigenvalues we get from the methods (the one on the left or the one on the right of E_{ref}),
- 3. to have a better stopping criterion in the inner QMR loop,
- 4. to be able to pass the shift to the diagonal preconditioner for a better preconditioning in the inner iterations,
- 5. and finally to stop the iterations of the Jacobi Davidson method with the original matrix instead with the folded one.

All these new features should lead us to a better Jacobi Davidson solver.

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