# A Parallel Solver for Extreme Eigenpairs<sup>1</sup>

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Abstract. In this paper a parallel algorithm for finding a group of extreme eigenvalues is presented. The algorithm is based on the well known Davidson method for finding one eigenvalue of a matrix. Here we incorporate knowledge about the structure of the subspace through the use of an arrowhead solver which allows more parallelization in both the original Davidson and our new version. In our numerical results various preconditioners (diagonal, multigrid and ADI) are compared. The performance results presented are for the Paragon but our implementation is portable to machines which provide MPI and BLAS.

# 1 Introduction

A large number of scientific applications rely on the computation of a few eigenvalues for a given matrix A. Typically they require the lowest or highest eigenvalues. Our algorithm (DSE) is based on the Davidson algorithm, but calculates various eigenvalues through implicit shifting. DSE was first presented in [10] under the name RDME to express its ability to identify eigenvalues with multiplicity bigger than one. The choice of preconditioner is an important issue in eliminating convergence to the wrong eigenvalue [14] In the next section, we describe the Davidson algorithm and our version for computing several eigenvalues. In [9] Oliveira presented convergence rates for Davidson type algorithm dependent on the type of preconditioner. These results are summarized here in Section 3. Section 4 addresses parallelization strategies discussing the data distribution in a MIMD architecture, and a fast solver for the projected subspace eigenproblem. In Section 5 we present numerical and performance results for the parallel implementation on the Paragon. Further results about the parallel algorithm and other numerical results are presented in [2].

# 2 The Davidson Algorithm

Two of the most popular iterative methods for large symmetric eigenvalue problems are Lanczos and Davidson algorithms. Both methods solve the eigenvalue

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problem  $Au = \lambda u$  by constructing an orthonormal basis  $V_k = [v_1, \ldots, v_k]$ , at each  $k^{th}$  iteration step, and then finding an approximation for the eigenvector u of A by using a vector  $u_k$  from the subspace spanned by  $V_k$ . Specifically, the original problem is projected onto the subspace which reduces the problem to a smaller eigenproblem  $S_k y = \tilde{\lambda} y_k$ , where  $S_k = V_k^T A V_k$ . Then the eigenpair  $(\tilde{\lambda}_k, y_k)$  can be obtained by applying a efficient procedure for small matrices. To complete the iteration, the eigenvector  $y_k$  is mapped back as  $u_k = V_k y_k$ , which is an approximation to the eigenvector u of the original problem. The difference between the two algorithms consists on the way that basis  $V_k$  is built. The attractiveness of the Lanczos algorithm results from the fact that each projected matrix  $S_k$  is tridiagonal. Unfortunately, sometimes this method may require a large number of iterations. The Davidson algorithm defines a dense matrix  $S_k$  on the subspace, but since we can incorporate a preconditioner in this algorithm the number of iterations can be much lower than for Lanczos. In Davidson type algorithms, a preconditioner  $M_{\lambda_k}$  is applied to the current residual,  $r_k = Au_k - \lambda_k u_k$ , and the preconditioned residual  $t_k = M_{\lambda_k} r_k$  is orthonormalized against the previous

columns of  $V_k = [v_1, v_2, \ldots, v_k]$ . Although in the original formulation  $M_{\lambda}$  is the diagonal matrix  $(diag(A) - \lambda I)^{-1}$  [6], the Generalized Davidson (GD) algorithm allows the incorporation of different operators for  $M_{\lambda}$ . The DSE algorithm can be summarized as follows.

Algorithm 1 – Restarted Davidson for Several Eigenvalues Given a matrix A, a normalized vector  $v_1$ , number of eigenpairs p, restart index q, and the minimal dimension m for the projected matrix S (m > p), compute approximations  $\lambda$  and u for the p smallest eigenpairs of A.

3. End For.

The core ideas of DSE (Algorithm 1) are based on the projection of A into the subspace spanned by the columns of  $V_k$ . The interation number k is not necessarily equal to  $\dim S_k$ , since we have incorporated implicit restarts. The matrix  $S_k$  is obtained by adding one more column and row  $V_k^T A v_k$  to matrix  $S_{k-1}$  (step 2.a). Other important aspects of the DSE algorithm are:

(1) the eigenvalue solver for the subspace matrix  $S_k$  (step 2.c); (2) the use of an auxiliary matrix  $W_k = [w_1, \ldots, w_k]$  to provide a residual calculation  $r_k = Au_k - \lambda_k u_k = w_k y_k - \lambda_k u_k$  with less computational work (step 2.e); the choice of a preconditioner M (step 2.f); and the use of modified Gram-Schmidt orthonormalization (step 2.g) which preserves numerical stability when updating the orthonormal basis  $V_{k+1}$ . At each iteration, the algorithm expands the matrix S either until all the first p eigenvalues have been converged, or S reaches a maximum dimension m + q; In the latter case, restarting is applied by using the orthonormal decomposition  $S_k = Y_k^T A_k Y_k$  of S. It corresponds to step 2.b in the algorithm. Because of our choice for m, note that in step 2.c dim S will be always bigger or equal to j.

### 3 Convergence Rate

A proof of convergence (but without a rate estimate) for the Davidson algorithm is given in Crouzeix, Philippe and Sadkane [5]. A bound on the convergence rate was first presented in [10]. The complete proof is shown in Oliveira [9]. Let A be the given matrix whose eigenvalues and eigenvectors are wanted. The preconditioner M is given for one step, and Davidson's algorithm is used with  $u_k$  being the current computed approximate eigenvector. The current eigenvalue estimate is the Rayleigh quotient  $\hat{\lambda}_k = \rho_A(u_k) = (u_k^T A u_k)/(u_k^T u_k)$ . Let the exact eigenvector with the smallest eigenvalue of A be u, and

$$Au = \lambda u$$

(If  $\lambda$  is a repeated eigenvalue of A, then we can let u be the normalized projection of  $u_k$  onto this eigenspace.)

**Theorem 1.** Let P be the orthogonal projection onto  $ker(A - \lambda I)^{\perp}$ . Suppose that A and M are symmetric positive definite. If

$$\|P - PMP(A - \lambda I)\|_2 \le \sigma < 1,$$

then for almost any starting value  $x_1$ , the convergence of the eigenvalue estimates  $\hat{\lambda}_k$  converge to  $\lambda$  ultimately geometrically with convergence factor bounded by  $\sigma^2$ , and the angle between the computed eigenvector and the exact eigenspace goes to zero ultimately geometrically with convergence factor bounded by  $\sigma$ .

A geometric convergence rate can be found for DSE (which obtains eigenvalues beyond the smallest (or largest) eigenvalue) by modifying Theorem 1. In the following theorem assume that

$$\sigma' = ||P' - P'MP'P'(A - \lambda_p I)P'||_2$$

where P' is the orthogonal projection onto the orthogonal complement of the span of the first p-1 eigenvectors. Then we can shown, in a similar way to Theorem 1 that the convergence factor for the new algorithm is bounded by

 $(\sigma')^2$  To prove Theorem 2 we use the fact that  $P's_k = s_k$ , as  $s_k$  is orthogonal to the bottom p eigenvectors. and that although  $(A - \lambda_p I)$  is no longer positive semi-definite,  $P'(A - \lambda_p I)P'$  is.

**Theorem 2.** Suppose that A and M are symmetric positive definite and that the first p-1 eigenvectors have been found exactly. Let P' be the orthogonal projection onto the orthogonal complement of the span of the first p-1 eigenvectors of A. If

$$||P' - P'MP'(A - \lambda_p I)P'||_2 \le \sigma' < 1,$$

then for almost any starting value  $x_1$ , the eigenvalue estimates  $\widehat{\lambda}_k$  obtained by our modified Davidson algorithm for several eigenvalues converges to  $\lambda_p$  ultimately geometrically with convergence factor is bounded by  $(\sigma')^2$ , and the angle between the exact and computed eigenvector goes to zero ultimately geometrically with convergence factor bounded by  $\sigma'$ .

## 4 Parallel Implementation

Previous implementations for the Davidson algorithm solve the eigenvalue problem in subspace S by using algorithms for dense matrices: early works [3, 4, 17] adopt EISPACK [12] routines, and later implementations [13, 15] use LAPACK [1] or reductions to tridiagonal form. Partial parallelization is obtained through the matrix-vector operations and sparse format storage for matrix A [13, 15]. Here we explore the relationship between two successive matrices  $S_k$  which allows us to represent  $S_k$  through an arrowhead matrix. The arrowhead structure is extremely sparse and the associated eigenvalue problem can be solved by a highly parallelizable method.

#### 4.1 Data Distribution

Data partitioning significantly affects the performance of a parallel system by determining the actual degree of concurrency of the processors. Matrices are partitioned along distinct processors so that the program exploits all the best possible data parallelism: The final distribution is well balanced, and most of the computational work can be performed without communication. These two conditions make the parallel program very suited for distributed memory architectures. Both computational workload and storage requirements are the same for all processors. Communication overhead is kept as low as possible. Matrix A is split into row blocks  $A^i$ ,  $i = 1, \ldots, N$ , each one containing  $\leq \lceil n/N \rceil$  rows of A. Thus processor i,  $i = 1, \ldots, N$  stores  $A^i$ , the  $i^{th}$  row block of A. Matrices  $V_k$  and  $W_k$  are stored in the same fashion. This data distribution allow us to perform many of the matrix-vector computations in place.

The orthonormalization strategy is also an important aspect in parallel environments. Recall that the modified Gram Schmidt (MGS) algorithm will be applied to the extended matrix  $[V_k, t_k]$  where the current basis  $V_k$  has been previously orthonormalized. This observation reduces the computational work by eliminating the outer loop from the two nested loops in the full MGS algorithm.

#### 4.2 The Arrowhead Relationship Between Matrices $S_k$

As pointed in [2, 10], the relationship between  $S_k$  and  $S_{k-1}$  can be used to show that  $S_k$  is explicitly similar to an arrowhead matrix  $\tilde{S}_k$  of the form

$$\tilde{S}_k = \begin{bmatrix} \Lambda_{k-1} & \tilde{s}_k \\ \tilde{s}_k^T & s_{kk} \end{bmatrix} \quad , \tag{1}$$

where  $\tilde{s}_k = Y_{k-1}^T V_{k-1}^T w_k$ ,  $s_{kk} = v_k^T w_k$ , and the diagonal matrix  $\Lambda_{k-1}$  corresponds to the orthonormal decomposition  $S_{k-1} = Y_{k-1}\Lambda_{k-1}Y_{k-1}^T$ . In practice, the matrix  $S_k$  does not need to be stored: only a vector for  $\Lambda_k$  and a matrix for  $Y_k$  are required from one iteration to the next. Thus, given the eigenvalues  $\Lambda_{k-1}$  and eigenvectors  $Y_{k-1}$  of  $S_{k-1}$ , matrix  $\tilde{S}_k$  can be used to find the eigenvalues  $\Lambda_k$  of  $S_k$ . Arrowhead eigensolvers [8,11] are highly parallelizable and typically perform  $\mathcal{O}(k^2)$  operations, instead of the usual  $\mathcal{O}(k^3)$  effort of algorithms for dense matrices S.

# 5 Numerical Results

In our numerical results we employ three kind of preconditiners: diagonal preconditioner (as in the original Davidson), multigrid and ADI. A preconditioner can be expressed as the matrix which solves Ax = b by applying an iterative method to MAx = Mb instead. In the case of a Diagonal preconditioner this would correspond to scaling the system and then solving. Multigrid and ADI preconditioners are more complex and for that we refer the reader to [16, 18, 19]. In our implementation level 1, 2 and 3 BLAS and the Message Passing Interface (MPI) library were used for easy portability.

The computational results in this section were obtained with a finite difference approximation for

$$-\Delta u + gu = f \tag{2}$$

on a unit square domain. Here g is null inside a  $0.2 \times 0.2$  square on the center of the rectangle and g = 100 for the remaining domain.

To compare the performance delivered by distinct preconditioners we observe the total timing and number of iterations required for the sequential DSE for finding the ten smallest eigenpairs (p = 10) assuming convergence for residual norms less or equal to  $10^{-7}$ . The restart indexes were q = 10 and m = 15. This corresponds to apply restarting every time that the projected matrix  $S_k$ achieves order 25, reducing its order to 15. Table 1 presents the actual running tIMINGS In a single processor of the Intel Paragon, running three grid sizes:  $31 \times 31$ ,  $63 \times 63$ , and  $127 \times 127$  (matrices of orders 961, 3969 and 16129, respectively.). It reflects the tradeoff between preconditioning strategies: although the diagonal preconditioner (DIAG) is the easiest and fastest to compute, it requires an increasing number of iterations for larger matrices. Multigrid preconditioners (MG) are more expensive than DIAG, but they turn to be more effective for larger matrices. Finally, the ADI method aggregate the advantages of the previous preconditioners in the sense that it is more effective and less expensive than MG. More details about the preconditioners used here can be found in [2] and its references.

	matrix order 961		matrix order 3969		matrix order 16129	
	iterations	time (sec)	iterations	time $(sec)$	iterations	time (sec)
ADI	29	4.7	26	16.2	27	80.7
MG	34	8.8	40	43.0	40	254.1
DIAG	174	8.6	319	43.2	700	386.0

Table 1. Sequential times and number of iterations for three preconditioners.

The overall behavior of the DSE algorithm (with a multigrid preconditioner) is shown in Figure 1 for matrices sizes 3969 and 16129, as a function of the number of processors. Note that the estimated optimal number of processors is not far from the actual optimal. The model for our estimates is presented in [2].

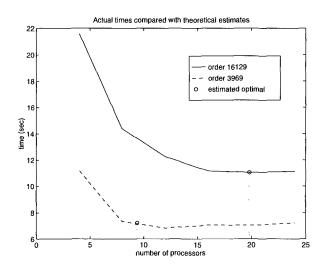


Fig. 1. Actual and estimated times for equation (2). Results for two different matrix sizes performance on the Paragon are shown.

To conclude, we compare the performance of the parallel DSE with PAR-PACK [7], a parallel implementation of ARPACK <sup>1</sup>. Figure 2 presents the total running times for both algorithms for the problem described above. For these runs, DSE used our parallel implementation of the ADI as its preconditioner.

<sup>&</sup>lt;sup>1</sup> ARPACK implements an Implicitly Restarted Arnoldi Method (IRAM) which in the symmetric case corresponds to the Implicitly Restarted Lanczos algorithm. We used the regular mode when running PARPACK.

The problem was solved by using 4, 8, and 16 processors to obtain relative residuals  $||Au - \lambda u||/||u||$  of order less than  $10^{-5}$ . We show our theoretical analysis for the parallel algorithm in [2]. Other numerical results for the sequential DSE algorithm, including examples showing the behavior of the algorithm for eigenvalues with multiplicity greater than one, were presented in [10].

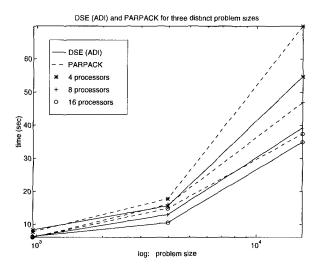


Fig.2. Running times for DSE and PARPACK using 4, 8, and 16 processors on the Paragon.

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