# Parallel Numerical Algorithms <br> Chapter 5 - Eigenvalue Problems <br> Section 5.2 - Eigenvalue Computation 

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## CS 554 / CSE 512

## Outline

(1) Basics
(2) Power Iteration
(3) QR Iteration
(4) Krylov Methods
(5) Other Methods

## Eigenvalues and Eigenvectors

- Given $n \times n$ matrix $\boldsymbol{A}$, find scalar $\lambda$ and nonzero vector $\boldsymbol{x}$ such that

$$
\boldsymbol{A x}=\lambda \boldsymbol{x}
$$

- $\lambda$ is eigenvalue and $\boldsymbol{x}$ is corresponding eigenvector
- $A$ always has $n$ eigenvalues, but they may be neither real nor distinct
- May need to compute only one or few eigenvalues, or all $n$ eigenvalues
- May or may not need corresponding eigenvectors


## Problem Transformations

- Shift: for scalar $\sigma$, eigenvalues of $\boldsymbol{A}-\sigma \boldsymbol{I}$ are eigenvalues of $\boldsymbol{A}$ shifted by $\sigma, \lambda_{i}-\sigma$
- Inversion: for nonsingular $\boldsymbol{A}$, eigenvalues of $\boldsymbol{A}^{-1}$ are reciprocals of eigenvalues of $\boldsymbol{A}, 1 / \lambda_{i}$
- Powers: for integer $k>0$, eigenvalues of $\boldsymbol{A}^{k}$ are $k$ th powers of eigenvalues of $\boldsymbol{A}, \lambda_{i}^{k}$
- Polynomial: for polynomial $p(t)$, eigenvalues of $p(\boldsymbol{A})$ are values of $p$ evaluated at eigenvalues of $\boldsymbol{A}, p\left(\lambda_{i}\right)$


## Similarity Transformations

- $\boldsymbol{B}$ is similar to $\boldsymbol{A}$ if there is nonsingular $\boldsymbol{T}$ such that

$$
\boldsymbol{B}=\boldsymbol{T}^{-1} \boldsymbol{A} \boldsymbol{T}
$$

- Then

$$
\boldsymbol{B} \boldsymbol{y}=\lambda \boldsymbol{y} \Rightarrow \boldsymbol{T}^{-1} \boldsymbol{A} \boldsymbol{T} \boldsymbol{y}=\lambda \boldsymbol{y} \Rightarrow \boldsymbol{A}(\boldsymbol{T} \boldsymbol{y})=\lambda(\boldsymbol{T} \boldsymbol{y})
$$

so $\boldsymbol{A}$ and $\boldsymbol{B}$ have same eigenvalues, and if $\boldsymbol{y}$ is eigenvector of $B$, then $\boldsymbol{x}=\boldsymbol{T} \boldsymbol{y}$ is eigenvector of $\boldsymbol{A}$

- Similarity transformations preserve eigenvalues, and eigenvectors are easily recovered


## Similarity Transformations

Forms attainable by similarity transformation

| $\boldsymbol{A}$ | $\boldsymbol{T}$ | $\boldsymbol{B}$ |
| :--- | :--- | :--- |
| distinct eigenvalues | nonsingular | diagonal |
| real symmetric | orthogonal | real diagonal |
| complex Hermitian | unitary | real diagonal |
| normal | unitary | diagonal |
| arbitrary real | orthogonal | real upper Hessenberg |
| arbitrary | unitary | upper triangular (Schur) |
| arbitrary | nonsingular | almost diagonal (Jordan) |

## Preliminary Reduction

- Eigenvalues easier to compute if matrix first reduced to simpler form by similarity transformation
- Diagonal or triangular most desirable, but cannot always be reached in finite number of steps
- Preliminary reduction usually to tridiagonal form (for symmetric matrix) or Hessenberg form (for nonsymmetric matrix)
- Preliminary reduction usually done by orthogonal transformations, using algorithms similar to QR factorization, but transformations must be applied from both sides to maintain similarity


## Parallel Algorithms for Eigenvalues

- Algorithms for computing eigenvalues and eigenvectors employ basic operations such as
- vector updates (saxpy)
- inner products
- matrix-vector and matrix-matrix multiplication
- solution of triangular systems
- orthogonal (QR) factorization
- In many cases, parallel implementations will be based on parallel algorithms we have already seen for these basic operations, although there will sometimes be new sources of parallelism


## Power Iteration

- Simplest method for computing one eigenvalueeigenvector pair is power iteration, which in effect takes successively higher powers of matrix times initial starting vector
- $x_{0}=$ arbitrary nonzero vector for $k=1,2, \ldots$

$$
\begin{aligned}
& \boldsymbol{y}_{k}=\boldsymbol{A} \boldsymbol{x}_{k-1} \\
& \boldsymbol{x}_{k}=\boldsymbol{y}_{k} /\left\|\boldsymbol{y}_{k}\right\|_{\infty}
\end{aligned}
$$

end

- If $\boldsymbol{A}$ has unique eigenvalue $\lambda_{1}$ of maximum modulus, then power iteration converges to eigenvector corresponding to dominant eigenvalue


## Power Iteration

- Convergence rate of power iteration depends on ratio $\left|\lambda_{2} / \lambda_{1}\right|$, where $\lambda_{2}$ is eigenvalue having second largest modulus
- It may be possible to choose shift, $\boldsymbol{A}-\sigma \boldsymbol{I}$, so that ratio is more favorable and yields more rapid convergence
- Shift must then be added to result to obtain eigenvalue of original matrix


## Parallel Power Iteration

- Power iteration requires repeated matrix-vector products, which are easily implemented in parallel for dense or sparse matrix, as we have seen
- Additional communication may be required for normalization, shifts, convergence test, etc.
- Though easily parallelized, power iteration is often too slow to be useful in this form


## Inverse Iteration

- Inverse iteration is power iteration applied to $\boldsymbol{A}^{-1}$, which converges to eigenvector corresponding to dominant eigenvalue of $\boldsymbol{A}^{-1}$, which is reciprocal of smallest eigenvalue of $\boldsymbol{A}$
- Inverse of $\boldsymbol{A}$ is not computed explicitly, but only factorization of $\boldsymbol{A}$ (and only once) to solve system of linear equations at each iteration
- $x_{0}=$ arbitrary nonzero vector for $k=1,2, \ldots$

Solve $\boldsymbol{A} \boldsymbol{y}_{k}=\boldsymbol{x}_{k-1}$ for $\boldsymbol{y}_{k}$
$\boldsymbol{x}_{k}=\boldsymbol{y}_{k} /\left\|\boldsymbol{y}_{k}\right\|_{\infty}$
end

## Inverse Iteration

- Shifting strategy can greatly accelerate convergence
- Inverse iteration is especially useful for computing eigenvector corresponding to approximate eigenvalue, since it converges rapidly when approximate eigenvalue is used as shift
- Inverse iteration also useful for computing eigenvalue closest to any given value $\beta$, since if $\beta$ is used as shift, then desired eigenvalue corresponds to smallest eigenvalue of shifted matrix


## Parallel Inverse Iteration

- Inverse iteration requires initial factorization of matrix $\boldsymbol{A}$ and solution of triangular systems at each iteration, so it appears to be much less amenable to efficient parallel implementation than power iteration
- However, inverse iteration is often used to compute eigenvector in situations where
- approximate eigenvalue is already known, so using it as shift yields very rapid convergence
- matrix has previously been reduced to simpler form (e.g., tridiagonal) for which linear system is easy to solve


## Simultaneous Iteration

- To compute many eigenvalue-eigenvector pairs, could apply power iteration to several starting vectors simultaneously, giving simultaneous iteration

$$
\begin{aligned}
& \boldsymbol{X}_{0}=\text { arbitrary } n \times q \text { matrix of rank } q \\
& \text { for } k=1,2, \ldots \\
& \quad \boldsymbol{X}_{k}=\boldsymbol{A} \boldsymbol{X}_{k-1}
\end{aligned}
$$

end

- span $\left(\boldsymbol{X}_{k}\right)$ converges to invariant subspace determined by $q$ largest eigenvalues of $\boldsymbol{A}$, provided $\left|\lambda_{q}\right|>\left|\lambda_{q+1}\right|$
- Normalization is needed at each iteration, and columns of $\boldsymbol{X}_{k}$ become increasingly ill-conditioned basis for span $\left(\boldsymbol{X}_{k}\right)$


## Orthogonal Iteration

- Both issues addressed by computing QR factorization at each iteration, giving orthogonal iteration
$\boldsymbol{X}_{0}=$ arbitrary $n \times q$ matrix of rank $q$ for $k=1,2, \ldots$

Compute reduced QR factorization

$$
\hat{\boldsymbol{Q}}_{k} \boldsymbol{R}_{k}=\boldsymbol{X}_{k-1}
$$

$$
\boldsymbol{X}_{k}=\boldsymbol{A} \hat{\boldsymbol{Q}}_{k}
$$

end

- Converges to block triangular form, and blocks are triangular where moduli of consecutive eigenvalues are distinct


## Parallel Orthogonal Iteration

- Orthogonal iteration requires matrix-matrix multiplication and QR factorization at each iteration, both of which we know how to implement in parallel with reasonable efficiency


## QR Iteration

- If we take $\boldsymbol{X}_{0}=\boldsymbol{I}$, then orthogonal iteration should produce all eigenvalues and eigenvectors of $\boldsymbol{A}$
- Orthogonal iteration can be reorganized to avoid explicit formation and factorization of matrices $\boldsymbol{X}_{k}$
- Instead, sequence of unitarily similar matrices is generated by computing QR factorization at each iteration and then forming reverse product, giving $Q R$ iteration
$\boldsymbol{A}_{0}=\boldsymbol{A}$
for $k=1,2, \ldots$
Compute QR factorization
$\boldsymbol{Q}_{k} \boldsymbol{R}_{k}=\boldsymbol{A}_{k-1}$
$\boldsymbol{A}_{k}=\boldsymbol{R}_{k} \boldsymbol{Q}_{k}$
end


## QR Iteration

- In simple form just given, each iteration of QR method requires $\Theta\left(n^{3}\right)$ work
- Work per iteration is reduced to $\Theta\left(n^{2}\right)$ if matrix is in Hessenberg form, or $\Theta(n)$ if symmetric matrix is in tridiagonal form
- Preliminary reduction is usually done by Householder or Givens transformations
- In addition, number of iterations required is reduced by preliminary reduction of matrix
- Convergence rate also enhanced by judicious choice of shifts


## Parallel QR Iteration

- Preliminary reduction can be implemented efficiently in parallel, using algorithms analogous to parallel QR factorization for dense matrix
- But subsequent QR iteration for reduced matrix is inherently serial, and permits little parallel speedup for this portion of algorithm
- This may not be of great concern if iterative phase is relatively small portion of total time, but it does limit efficiency and scalability


## Reduction to Hessenberg/Tridiagonal

- To reduce to Hessenberg can execute QR on each column
- If $\boldsymbol{H}_{i}$ is the Householder transformation used to annihilate $i$ th subcolumn, perform similarity transformation

$$
\boldsymbol{A}^{(i+1)}=\boldsymbol{H}_{i} \boldsymbol{A}^{(i)} \boldsymbol{H}_{i}^{T}
$$

- More generally, run QR on $b$ subcolumns of $\boldsymbol{A}$ to reduce to band-width $b$

$$
\boldsymbol{B}^{(i+1)}=\boldsymbol{Q}_{i} \boldsymbol{B}^{(i)} \boldsymbol{Q}_{i}^{T}
$$

- To avoid fill $\boldsymbol{Q}_{i}^{T}$ must not operate on the $b$ columns which $Q_{i}$ reduces
- Once reduction completed to a band-width subsequent eliminations introduce fill (bulges) but can be done


## Reduction from Banded to Tridiagonal



- Two-sided successive orthogonal reductions generate bulges of fill when applied to a banded matrix
- Bulges can be chased with pipelined parallelism


## Krylov Subspace Methods

- Krylov subspace methods reduce matrix to Hessenberg (or tridiagonal) form using only matrix-vector multiplication
- For arbitrary starting vector $x_{0}$, if

$$
\boldsymbol{K}_{k}=\left[\begin{array}{llll}
\boldsymbol{x}_{0} & \boldsymbol{A} \boldsymbol{x}_{0} & \cdots & \boldsymbol{A}^{k-1} \boldsymbol{x}_{0}
\end{array}\right]
$$

then

$$
\boldsymbol{K}_{n}^{-1} \boldsymbol{A} \boldsymbol{K}_{n}=\boldsymbol{C}_{n}
$$

where $C_{n}$ is upper Hessenberg (in fact, companion matrix)

## Krylov Subspace Methods

- To obtain better conditioned basis for span $\left(\boldsymbol{K}_{n}\right)$, compute QR factorization

$$
\boldsymbol{Q}_{n} \boldsymbol{R}_{n}=\boldsymbol{K}_{n}
$$

so that

$$
\boldsymbol{Q}_{n}^{H} \boldsymbol{A} \boldsymbol{Q}_{n}=\boldsymbol{R}_{n} \boldsymbol{C}_{n} \boldsymbol{R}_{n}^{-1} \equiv \boldsymbol{H}
$$

with $\boldsymbol{H}$ upper Hessenberg

## Krylov Subspace Methods

- Equating $k$ th columns on each side of equation $\boldsymbol{A} \boldsymbol{Q}_{n}=\boldsymbol{Q}_{n} \boldsymbol{H}$ yields recurrence

$$
\boldsymbol{A} \boldsymbol{q}_{k}=h_{1 k} \boldsymbol{q}_{1}+\cdots+h_{k k} \boldsymbol{q}_{k}+h_{k+1, k} \boldsymbol{q}_{k+1}
$$

relating $\boldsymbol{q}_{k+1}$ to preceding vectors $\boldsymbol{q}_{1}, \ldots, \boldsymbol{q}_{k}$

- Premultiplying by $\boldsymbol{q}_{j}^{H}$ and using orthonormality,

$$
h_{j k}=\boldsymbol{q}_{j}^{H} \boldsymbol{A} \boldsymbol{q}_{k}, \quad j=1, \ldots, k
$$

- These relationships yield Arnoldi iteration, which produces upper Hessenberg matrix column by column using only matrix-vector multiplication by $\boldsymbol{A}$ and inner products of vectors


## Arnoldi Iteration

```
\mp@subsup{x}{0}{}}=\mathrm{ arbitrary nonzero starting vector
\mp@subsup{q}{1}{}}=\mp@subsup{\boldsymbol{x}}{0}{}/|\mp@subsup{\boldsymbol{x}}{0}{}\mp@subsup{|}{2}{
for }k=1,2,\ldots
    \mp@subsup{u}{k}{}}=\boldsymbol{A}\mp@subsup{\boldsymbol{q}}{k}{
    for j=1 to k
        hjk}=\mp@subsup{\boldsymbol{q}}{j}{H}\mp@subsup{\boldsymbol{u}}{k}{
        u
    end
    hk+1,k}=|\mp@subsup{\boldsymbol{u}}{k}{}\mp@subsup{|}{2}{
    if }\mp@subsup{h}{k+1,k}{}=0\mathrm{ then stop
    \mp@subsup{\boldsymbol{q}}{k+1}{}=\mp@subsup{\boldsymbol{u}}{k}{}/\mp@subsup{h}{k+1,k}{}
end
```


## Arnoldi Iteration

- If

$$
\boldsymbol{Q}_{k}=\left[\begin{array}{lll}
\boldsymbol{q}_{1} & \cdots & \boldsymbol{q}_{k}
\end{array}\right],
$$

then

$$
\boldsymbol{H}_{k}=\boldsymbol{Q}_{k}^{H} \boldsymbol{A} \boldsymbol{Q}_{k}
$$

is upper Hessenberg matrix

- Eigenvalues of $\boldsymbol{H}_{k}$, called Ritz values, are approximate eigenvalues of $\boldsymbol{A}$, and Ritz vectors given by $\boldsymbol{Q}_{k} \boldsymbol{y}$, where $\boldsymbol{y}$ is eigenvector of $\boldsymbol{H}_{k}$, are corresponding approximate eigenvectors of $\boldsymbol{A}$
- Eigenvalues of $\boldsymbol{H}_{k}$ must be computed by another method, such as QR iteration, but this is easier problem if $k \ll n$


## Arnoldi Iteration

- Arnoldi iteration expensive in work and storage because each new vector $\boldsymbol{q}_{k}$ must be orthogonalized against all previous columns of $Q_{k}$, which must be stored
- So Arnoldi process usually restarted periodically with carefully chosen starting vector
- Ritz values and vectors produced are often good approximations to eigenvalues and eigenvectors of $\boldsymbol{A}$ after relatively few iterations
- Work and storage costs drop dramatically if matrix symmetric or Hermitian, since recurrence then has only three terms and $\boldsymbol{H}_{k}$ is tridiagonal (so usually denoted $\boldsymbol{T}_{k}$ ), yielding Lanczos iteration


## Lanczos Iteration

$$
\begin{aligned}
& \boldsymbol{q}_{0}=\mathbf{0} \\
& \beta_{0}=0 \\
& \boldsymbol{x}_{0}=\text { arbitrary nonzero starting vector } \\
& \boldsymbol{q}_{1}=\boldsymbol{x}_{0} /\left\|\boldsymbol{x}_{0}\right\|_{2} \\
& \text { for } k=1,2, \ldots \\
& \quad \boldsymbol{u}_{k}=\boldsymbol{A} \boldsymbol{q}_{k} \\
& \quad \alpha_{k}=\boldsymbol{q}_{k}^{H} \boldsymbol{u}_{k} \\
& \quad \boldsymbol{u}_{k}=\boldsymbol{u}_{k}-\beta_{k-1} \boldsymbol{q}_{k-1}-\alpha_{k} \boldsymbol{q}_{k} \\
& \beta_{k}=\left\|\boldsymbol{u}_{k}\right\|_{2} \\
& \text { if } \beta_{k}=0 \text { then stop } \\
& \boldsymbol{q}_{k+1}=\boldsymbol{u}_{k} / \beta_{k} \\
& \text { end }
\end{aligned}
$$

## Lanczos Iteration

- $\alpha_{k}$ and $\beta_{k}$ are diagonal and subdiagonal entries of symmetric tridiagonal matrix $\boldsymbol{T}_{k}$
- As with Arnoldi, Lanczos iteration does not produce eigenvalues and eigenvectors directly, but only tridiagonal matrix $\boldsymbol{T}_{k}$, whose eigenvalues and eigenvectors must be computed by another method to obtain Ritz values and vectors
- If $\beta_{k}=0$, then algorithm appears to break down, but in that case invariant subspace has already been identified (i.e., Ritz values and vectors are already exact at that point)


## Lanczos Iteration

- In principle, if Lanczos algorithm is run until $k=n$, resulting tridiagonal matrix is orthogonally similar to $A$
- In practice, rounding error causes loss of orthogonality, invalidating this expectation
- Problem can be overcome by reorthogonalizing vectors as needed, but expense can be substantial
- Alternatively, can ignore problem, in which case algorithm still produces good eigenvalue approximations, but multiple copies of some eigenvalues may be generated


## Krylov Subspace Methods

- Virtue of Arnoldi and Lanczos iterations is ability to produce good approximations to extreme eigenvalues for $k \ll n$
- Moreover, they require only one matrix-vector multiplication by $\boldsymbol{A}$ per step and little auxiliary storage, so are ideally suited to large sparse matrices
- If eigenvalues are needed in middle of spectrum, say near $\sigma$, then algorithm can be applied to matrix $(\boldsymbol{A}-\sigma \boldsymbol{I})^{-1}$, assuming it is practical to solve systems of form $(\boldsymbol{A}-\sigma \boldsymbol{I}) \boldsymbol{x}=\boldsymbol{y}$


## Parallel Krylov Subspace Methods

- Krylov subspace methods composed of
- vector updates (saxpy)
- inner products
- matrix-vector multiplication
- computing eigenvalues/eigenvectors of tridiagonal matrices
- Parallel implementation requires implementing each of these in parallel as before
- For early iterations, Hessenberg or tridiagonal matrices generated are too small to benefit from parallel implementation, but Ritz values and vectors need not be computed until later


## Jacobi Method

- Jacobi method for symmetrix matrix starts with $\boldsymbol{A}_{0}=\boldsymbol{A}$ and computes sequence

$$
\boldsymbol{A}_{k+1}=\boldsymbol{J}_{k}^{T} \boldsymbol{A}_{k} \boldsymbol{J}_{k}
$$

where $\boldsymbol{J}_{k}$ is plane rotation that annihilates symmetric pair of off-diagonal entries in $\boldsymbol{A}_{k}$

- Plane rotations are applied repeatedly from both sides in systematic sweeps through matrix until magnitudes of all off-diagonal entries are reduced below tolerance
- Resulting diagonal matrix is orthogonally similar to original matrix, so diagonal entries are eigenvalues, and eigenvectors given by product of plane rotations


## Parallel Jacobi Method

- Jacobi method, though slower than QR iteration serially, parallelizes better
- Parallel implementation of Jacobi method performs many annihilations simultaneously, at locations chosen so that rotations do not interfere with each other (analogous to parallel Givens QR factorization)


## Divide-and-Conquer Method

- One method for computing eigenvalues and eigenvectors of real symmetric tridiagonal matrix is based on divide-and-conquer
- Express symmetric tridiagonal matrix $T$ as

$$
\boldsymbol{T}=\left[\begin{array}{ll}
\boldsymbol{T}_{1} & \boldsymbol{O} \\
\boldsymbol{O} & \boldsymbol{T}_{2}
\end{array}\right]+\beta \boldsymbol{u} \boldsymbol{u}^{T}
$$

- Can now compute eigenvalues and eigenvectors of smaller matrices $T_{1}$ and $T_{2}$
- To relate these back to eigenvalues and eigenvectors of original matrix requires solution of secular equation, which can be done reliably and efficiently


## Divide-and-Conquer Method

- Applying this approach recursively yields divide-and-conquer algorithm that is naturally parallel
- Parallelism in solving secular equations grows as parallelism in processing independent tridiagonal matrices shrinks, and vice versa
- Algorithm is complicated to implement and difficult questions of numerical stability, eigenvector orthogonality, and load balancing must be addressed


## Polar Decomposition

- The polar decomposition uses spectrum slicing to obtain the symmetric eigenvalue decomposition via $O(\log n)$ QR factorizations and matrix multiplications and $O\left(n^{3}\right)$ work
- The polar decomposition of a matrix $\boldsymbol{A}$ is composed of an orthogonal matrix $\boldsymbol{U}$ and symmetric positive semi-definite matrix $\boldsymbol{H}, \boldsymbol{A}=\boldsymbol{U} \boldsymbol{H}$
- It can be computed via the QR-based Dynamically Weighted Halley (QDWH) algorithm, an iterative scheme that starts with $\boldsymbol{X}_{0}=\boldsymbol{A} / \alpha$ and computes

$$
\left[\begin{array}{c}
\sqrt{c_{k}} \boldsymbol{X}_{k} \\
\boldsymbol{I}
\end{array}\right]=\left[\begin{array}{c}
\boldsymbol{Q}_{1} \\
\boldsymbol{Q}_{2}
\end{array}\right] \boldsymbol{R}, \quad \boldsymbol{X}_{k+1}=\frac{b_{k}}{c_{k}} \boldsymbol{X}_{k}+\frac{1}{\sqrt{c_{k}}}\left(a_{k}-b_{k} / c_{k}\right) \boldsymbol{Q}_{1} \boldsymbol{Q}_{2}^{T}
$$

- With appropriate choices of $\alpha, a_{k}, b_{k}, c_{k}$, QDWH converges within 6 iterations for double precision


## Eigenvalues via Polar Decomposition

- When $\boldsymbol{A}$ is symmetric, the orthogonal matrix $\boldsymbol{U}$ in the polar decomposition is given by the sign function,

$$
\boldsymbol{U}=\operatorname{sign}(\boldsymbol{A})=\boldsymbol{A}\left(\boldsymbol{A}^{2}\right)^{-1 / 2}
$$

- If the eigenvalue decomposition of $\boldsymbol{A}$ is $\boldsymbol{A}=\boldsymbol{V} \boldsymbol{D} \boldsymbol{V}^{T}$, then

$$
\boldsymbol{U}=\boldsymbol{V} \operatorname{sign}(\boldsymbol{D}) \boldsymbol{V}^{T}=\left[\begin{array}{ll}
\boldsymbol{V}_{1} & \boldsymbol{V}_{2}
\end{array}\right]\left[\begin{array}{ll}
\boldsymbol{I}_{k} & \\
& -\boldsymbol{I}_{n-k}
\end{array}\right]\left[\begin{array}{ll}
\boldsymbol{V}_{1} & \boldsymbol{V}_{2}
\end{array}\right]^{T}
$$

- Consequently, given an estimate $\sigma$ of a median eigenvalue of $\boldsymbol{A}$, the polar decomposition of $\boldsymbol{A}-\sigma \boldsymbol{I}$ allows one to partition the spectrum by finding $V_{1}$ from subspace iteration on $\boldsymbol{U}+\boldsymbol{I}=2 \boldsymbol{V}_{1} \boldsymbol{V}_{1}^{T}$


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