z-Pares: A Complex Moment Based Parallel Sparse Eigenvalue Solver Package

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Motivation

For post-petascale supercomputers, we need to consider hierarchical structures:



New algorithm design: New concept of algorithm design to adapt to the change of architectures

Development of Parallel Eigensolver Software

 CREST Project: Development of an Eigen-Supercomputing Engine using a Post-Petascale Hierarchical Model (FY2011 – FY2015)

For both dense and sparse matrices

- In this talk, we focus on a sparse solver (z-Pares)
- Goal: Provide high performance eigensolver software for post-petascale architectures
 - Scalable and reliable parallel algorithms for eigenvalue computations
 - > High performance implementation for various applications
 - Provide practical and portable software

Implementation of z-Pares

A parallel eigensolver package for sparse matrices

- Implemented on Trilinos
- Uses Block Krylov methods and Shifted Kyrlov methods for spectral transformation



Our Approach for Parallel Scalability

Avoid recurrence calculations

Algorithm described by a recurrence relation (ex. Krylov methods):

$$a_{i+1} \rightarrow a_i \rightarrow a_{i+1} \rightarrow \cdots \rightarrow a_n$$

 $a_{i+1} = b_i a_i + c_i a_{i-1}$

> Algorithm described by a summation (ex. Numerical quadrature): $N = \frac{1}{N} = \sum_{w \in f} w \cdot f$

$$I_N = \sum_{j=1} w_j f_j$$

$$f_1 \qquad f_2 \qquad \dots \qquad f_{N-1} \qquad f_N$$

Basic Idea

Contour integral for a rational function





Spectral decomposition of the resolvent of A

$$(zI - A)^{-1} = \sum_{i=1}^{n} \frac{P_i}{z - \lambda_i}, \quad P_i = \boldsymbol{u}_i \boldsymbol{u}_i^{\mathrm{H}}$$

 λ_i : eigenvalue, u_i : eigenvector (for simplicity, we considered the case that A is real symmetric)

Contour integral for the resolvent of *A*

$$\frac{1}{2\pi \mathrm{i}} \oint_{\Gamma} (zI - A)^{-1} \mathrm{d}z = \sum_{\lambda_i \in G} P_i$$

Numerical Quadrature

Approximated by a numerical quadrature

$$\sum_{\lambda_i \in G} P_i = \frac{1}{2\pi i} \oint_{\Gamma} (zI - A)^{-1} dz \approx \sum_{j=1}^N w_j (z_j I - A)^{-1}$$

 z_j : quadrature point, w_j : quadrature wait

Apply for an arbitrary vector *v*

$$\sum_{\lambda_i \in G} P_i \boldsymbol{v} \approx \sum_{j=1}^N w_j (z_j I - A)^{-1} \boldsymbol{v}$$

Systems of linear equations at shift points z_j

$$(z_j I - A) \boldsymbol{x}_j = \boldsymbol{v}, \ j = 1, \dots, N$$

Eigensolver using Contour Integrals

SS method (Sakurai, et. al. (2003), Asakura, et al. (2009))
 Eigenvalue problem

$$T(\lambda)\boldsymbol{x} = \boldsymbol{0}$$

where $T(\lambda)$ is an analytic matrix function.

> Complex moments are calculated.

$$\boldsymbol{s}_{k} = \frac{1}{2\pi \mathrm{i}} \oint_{\Gamma} z^{k} T(z)^{-1} \mathrm{d} z \, \boldsymbol{v} \approx \sum_{j=1}^{N} w_{j} z_{j}^{k} T(z_{j})^{-1} \boldsymbol{v}$$

> When multiple input vectors $V = [v_1, v_2, ..., v_L]$ are used, systems with multiple right-hand sides are solved.

$$T(z_j)X_j = V, \ j = 1, \dots, N$$

Parallelization

The SS method requires no successive process in the subspace construction step



Hierarchical Parallel Structure

Hardware is grouped according to a hierarchical structure of the algorithm



Shifted Krylov Subspace Method for SEP

Shifted linear systems:

In case of standard eigenvalue problems, shifted linear systems are solved.

$$(z_j I - A)X_j = V, \ j = 1, \dots, N$$

- Reduction of number of Mat-Vec:
 - By using the shift invariance of the Krylov subspace, we can reduce the number of matrix-vector multiplications.



Shifted Krylov Subspace Method for SEP

In case of a lot of shift points:

If there are a lot of shifted systems, the computational cost for AXPY becomes dominant



Shifted Block Krylov method:

- AXPY for multiple RHSs can be implemented by GEMM
 - Improves performance

- Application for band calculation with real space density functional theory (RSDFT)
 - > An interior standard eigenvalue problem (SEP)
 - Eigenvalues around the band gap
 - Solved several matrices: $A(k) u = \lambda u, k = 0, \Delta k, 2\Delta k, ...$
- Material: Silicon nanowire
- Test Environment: T2K-Tsukuba System
 > 2,048 cores for 1,085 atoms
 > 8,192 cores for 1,654 atoms



Eigensolver:

- CG: Conjugate Gradient method
- SS: SS method with a shifted Block Krylov linear solver

| #atoms | Mat. dim. | #core | Time (sec) | | SS/CG |
|--------|-----------|-------|------------|-------|------------|
| | | | CG | SS | Time ratio |
| 1,085 | 576,000 | 2,048 | 2,373 | 1,534 | 0.65 |
| 1,654 | 1,453,248 | 8,192 | 1,074* | 563 | 0.52 |

*certain eigenvalues were not obtained in the interval to be computed

We have executed our code on the K computer with 6,144 cores for the case of silicon nanowire 9,924 atoms.



Application for molecular orbital (MO) computation of protein

- Generalized eigenvalue problem (GEP)
- Target protein: Epidamal Growth Factor Receptor 17,246 atoms
- Matrix dimension: 96,234, #nonzeros: 456,807,648
- Fest environment: T2k-Tsukuba System
- Linear solver: Block COCG method
- Preconditioner: MUMPS (sparse direct solver) for a dropped coefficient matrix



One contour path: 16 linear systems are solved with 128 cores



- Multiple contour paths
 - Eigenvalues from lowest orbital to highest occupied orbital are calculated for self consistent field (SCF) iteration
 - > 314 contour paths are set (128 cores for each contour path)
 - > 32,769 eigenvalues and corresponding eigenvectors are obtained



Conclusions

- Development of parallel eigensolver and its software for post-petascale computing environment
 - Contour integral based method
 - Scalable algorithm
 - Hierarchical parallel structure
 - Several number of linear systems with multiple RHSs
 - Block Krylov methods and Shifted Krylov methods
 - Time for solving linear systems is dominant
- Looking for
 - efficient linear solvers and preconditioners
 - > applications