



ELSI Interface Development Version

User's Guide

The ELSI Team

<http://elsi-interchange.org>

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1 Introduction

1.1 The Cubic Wall of Kohn-Sham Density-Functional Theory

In KS-DFT [1], the many-electron problem for the Born-Oppenheimer electronic ground state is reduced to a system of single particle equations known as the Kohn-Sham equations

$$\hat{h}^{\text{KS}}\psi_l = \epsilon_l\psi_l, \quad (1.1)$$

where ψ_l and ϵ_l are Kohn-Sham orbitals and their associated eigenenergies, and \hat{h}^{KS} denotes the Kohn-Sham Hamiltonian, which includes the kinetic energy, the average electrostatic potential of the electron density and of the nuclei (i.e. the Hartree potential), the exchange-correlation potential, and possible additional potential terms from external electromagnetic fields. These terms depend on the electron density \mathbf{n} , which is determined by the Kohn-Sham orbitals ψ_l . These terms also enter the Hamiltonian \hat{h}^{KS} , which determines the Kohn-Sham orbitals ψ_l .

Due to this circular dependency, the Kohn-Sham equations are in fact a non-linear optimization problem, and therefore must be solved iteratively. The most commonly used method is the self-consistent field (SCF) approach. It usually starts from an initial guess of the electron density, from which the kinetic energy, electrostatic potential, exchange-correlation potential, and external potential are computed, forming the Kohn-Sham Hamiltonian. Then, the Kohn-Sham orbitals (wavefunctions) are solved from the Hamiltonian, and new electron density is computed from the Kohn-Sham orbitals. To achieve self-consistency, the electron density is updated in every SCF iteration until converged to an acceptable level.

In almost all practical approaches, N_{basis} basis functions $\phi_i(\mathbf{r})$ are employed to approximately expand the Kohn-Sham orbitals:

$$\psi_l(\mathbf{r}) = \sum_{j=1}^{N_{\text{basis}}} c_{jl}\phi_j(\mathbf{r}). \quad (1.2)$$

The choice of basis set is one of the critical decisions in the design of an electronic structure code. Using non-orthogonal basis functions (e.g., Gaussian functions, Slater functions, numeric atom-centered orbitals) in 1.2 converts 1.1 to a generalized eigenvalue problem

$$\sum_j h_{ij}c_{jl} = \epsilon_l \sum_j s_{ij}c_{jl}, \quad (1.3)$$

where h_{ij} and s_{ij} are the elements of the Hamiltonian matrix \mathbf{H} and the overlap matrix \mathbf{S} , which can be computed through numerical integrations:

$$\begin{aligned} h_{ij} &= \int d^3r [\phi_i^*(\mathbf{r})\hat{h}^{\text{KS}}\phi_j(\mathbf{r})], \\ s_{ij} &= \int d^3r [\phi_i^*(\mathbf{r})\phi_j(\mathbf{r})]. \end{aligned} \quad (1.4)$$

1.3 can thus be expressed in the following matrix form:

$$\mathbf{H}\mathbf{c} = \epsilon\mathbf{S}\mathbf{c}. \quad (1.5)$$

Here, the matrix \mathbf{c} and diagonal matrix ϵ contain the eigenvectors and eigenvalues, respectively, of the eigensystem of the matrices \mathbf{H} and \mathbf{S} .

When using orthonormal basis sets (e.g. plane waves, multi-resolution wavelets), the eigenproblem described in 1.5 reduces to a standard form where $s_{ij} = \delta_{ij}$.

The explicit solution of 1.3 or 1.5 yields the Kohn-Sham orbitals ψ_i , from which the electron density $n(\mathbf{r})$ can be computed following an orbital-based method:

$$n(\mathbf{r}) = \sum_{l=1}^{N_{\text{basis}}} f_l \psi_l^*(\mathbf{r}) \psi_l(\mathbf{r}), \quad (1.6)$$

where f_l denotes the occupation number of each orbital. In an actual computation, it is sufficient to perform the summation only for the occupied ($f_l > 0$) orbitals. The ratio of occupied orbitals to the total number of basis functions can be below 1% for plane wave basis sets, whereas with some localized basis sets, fewer basis functions are required, leading to a larger fraction of occupied states typically between 10% and 50%.

An alternative method can be employed for localized basis functions:

$$n(\mathbf{r}) = \sum_{i,j}^{N_{\text{basis}}} \phi_i^*(\mathbf{r}) p_{ij} \phi_j(\mathbf{r}), \quad (1.7)$$

with p_{ij} being the elements of the density matrix \mathbf{P} that need to be computed before the density update:

$$p_{ij} = \sum_{l=1}^{N_{\text{basis}}} f_l c_{il} c_{jl}. \quad (1.8)$$

From a viewpoint of computational complexity, with localized basis functions, almost all standard pieces of solving the Kohn-Sham equations can be formulated in a linear scaling fashion with respect to the system size. The only remaining bottleneck for semilocal functionals is the eigenproblem described in Eqs. 1.3 and 1.5. The density matrix is directly accessible through methods other than diagonalization, therefore it is not always necessary to explicitly solve the eigenproblem. Which algorithm to use depends on many factors such as the choice of basis set, and the system and characters of the physical systems. In an SCF calculation, the eigenproblem needs to be tackled repeatedly. If this step is treated with the most efficient algorithm, the whole SCF calculation can be greatly accelerated.

1.2 ELSI, the ELectronic Structure Infrastructure

ELSI unifies the community effort in overcoming the cubic-wall problem of KS-DFT by bridging the divide between developers of electronic structure solvers and KS-DFT codes. Via a unified interface, ELSI gives KS-DFT developers easy access to multiple solvers that solve or circumvent the Kohn-Sham eigenproblem efficiently. Solvers are treated on equal footing within ELSI, giving solver developers a unified platform for implementation and benchmarking across codes and physical systems. Solvers may be switched dynamically in an SCF cycle, allowing the KS-DFT developer to mix-and-match strengths of different solvers. Solvers can work cooperatively with one another within ELSI, allowing for acceleration greater than either solver can achieve individually. Most importantly, ELSI exists as a community for KS-DFT and solver developers to interact and work together to improve performance of solvers, with monthly web meetings to discuss progress on code development, yearly on-site “connector meetings”, and planned webinars and workshops.

The current version of ELSI supports ELPA [2, 3], libOMM [4], PEXSI [5, 6], SLEPc-SIPs [7, 8], and NTPoly [9] solvers. Codes currently integrated with ELSI include DFTB+ [10], DGDFT [11], FHI-aims [12], and SIESTA [13].

Versatility: ELSI supports real-valued and complex-valued density matrix, eigenvalue, and eigenvector calculations. A unified software interface designed for rapid integration into a variety of electronic structure codes is provided. Fortran and C/C++ interfaces are provided.

Flexibility: ELSI supports both dense and sparse matrices as input/output. Supported matrix distribution layouts include 2D block-cyclic distribution, 1D block-cyclic distribution, and 1D block distribution. In situations where the input/output matrix format used by the electronic structure code and the format used internally by the requested solver are different, conversion and redistribution of matrices will be performed automatically.

Scalability: The solver libraries collected in ELSI are highly scalable. For instance, ELPA can scale to a hundred thousand CPU cores given a sufficiently large problem to solve, and PEXSI, with its efficient two-level parallelism, easily scales to tens of thousands of CPU cores.

Portability: ELSI and its redistributed library source packages have been confirmed to work on commonly-used HPC architectures (Cray, IBM, Intel, NVIDIA) using major compilers (Cray, GNU, IBM, Intel, PGI).

1.3 Kohn-Sham Solver Libraries Supported by ELSI

Solvers supported in the current version of ELSI are: ELPA [2, 3], libOMM [4], PEXSI [5, 6], SLEPc-SIPc [7, 8], and NTPoly [9]. The table below summarizes the supported data type, input/output matrix format, supported calculation type, and possible outputs of the solvers.

| Solver | Data type | Matrix format | Spin/ k -point | Output |
|----------------------------|--------------|---------------|------------------|---|
| ELPA | real/complex | dense/sparse | yes/yes | eigenvalues, eigenvectors, density matrix, energy-weighted density matrix, chemical potential, electronic entropy |
| libOMM | real/complex | dense/sparse | yes/yes | density matrix, energy-weighted density matrix |
| PEXSI | real/complex | dense/sparse | yes/yes | density matrix, energy-weighted density matrix, chemical potential |
| SLEPc-SIPs | real | dense/sparse | no/no | eigenvalues, eigenvectors, density matrix, energy-weighted density matrix, chemical potential, electronic entropy |
| NTPoly | real | dense/sparse | no/no | density matrix, energy-weighted density matrix, chemical potential |

What follows is a brief introduction of the solvers currently supported in ELSI. For detailed technical descriptions of the solvers, the reader is referred to the original publications of the solvers, e.g., those in the reference list of this document.

1.3.1 ELPA

The explicit solution of a generalized or standard eigenproblem is a well-studied task. The generalized eigenproblem in 1.5 is first transformed to the standard form, e.g., by Cholesky decomposition of the overlap matrix \mathbf{S} :

$$\mathbf{S} = \mathbf{L}\mathbf{L}^*, \quad (1.9)$$

where \mathbf{L} is a lower triangular matrix. Applying \mathbf{L} to \mathbf{H} and \mathbf{c} in the following way

$$\begin{aligned} \tilde{\mathbf{H}} &= \mathbf{L}^{-1}\mathbf{H}(\mathbf{L}^*)^{-1}, \\ \tilde{\mathbf{c}} &= \mathbf{L}^*\mathbf{c}, \end{aligned} \quad (1.10)$$

transforms 1.5 to a standard eigenproblem

$$\tilde{\mathbf{H}}\tilde{\mathbf{c}} = \epsilon\tilde{\mathbf{c}}. \quad (1.11)$$

This standard eigenproblem is solved by further transforming it to a tridiagonal form

$$\mathbf{T} = \mathbf{Q}\tilde{\mathbf{H}}\mathbf{Q}^*, \quad (1.12)$$

where \mathbf{Q} is a transformation matrix, and \mathbf{T} is a tridiagonal matrix whose eigenvalues and eigenvectors are computed by, e.g., the divide-and-conquer approach or the MRRR method. This procedure is called “diagonalization”, as the full matrix is reduced to a (tri)diagonal form.

The massively parallel direct eigensolver ELPA [2, 3] facilitates the direct solution of symmetric or Hermitian eigenproblems on high-performance computers by adopting a two-stage diagonalization algorithm, which first reduces the full

matrix to a banded intermediate form, then to the tridiagonal form:

$$\begin{aligned} \mathbf{B} &= \mathbf{Q}_1 \tilde{\mathbf{H}} \mathbf{Q}_1^*, \\ \mathbf{T} &= \mathbf{Q}_2 \mathbf{B} \mathbf{Q}_2^*. \end{aligned} \quad (1.13)$$

where \mathbf{Q}_1 and \mathbf{Q}_2 are transformation matrices used in the two-stage diagonalization; \mathbf{B} is a banded matrix; and \mathbf{T} is a tridiagonal matrix. Compared to the one-stage diagonalization (1.12), the two-stage approach introduces two additional steps. Still, the two-stage approach has been shown to enable faster computation and better parallel scalability on present-day computers. Specifically, the matrix-vector operations (BLAS level-2 routines) in 1.12 can be mostly replaced by more efficient matrix-matrix operations (BLAS level-3 routines) in 1.13. The computational workload associated with the back-transformation of the eigenvectors is greatly alleviated if only a small fraction of the eigenvectors representing the lowest eigenstates is required, and by architecture-specific linear-algebra “kernels” provided with the ELPa library.

1.3.2 libOMM

Instead of diagonalizing the $N_{\text{basis}} \times N_{\text{basis}}$ eigenproblem, the orbital minimization method (OMM) minimizes an unconstrained energy functional using a set of auxiliary Wannier functions. At the minimum of the OMM energy functional, the Wannier functions can be used to construct the density matrix. Specifically, N_{W} non-orthogonal Wannier functions χ_k are employed to represent the occupied subspace of a system with N_{electron} electrons:

$$\chi_k = \sum_{j=1}^{N_{\text{basis}}} W_{kj} \phi_j. \quad (1.14)$$

For non-spin-polarized systems, the index k runs from 1 to $N_{\text{W}} = N_{\text{electron}}/2$. Then the matrices \mathbf{H} and \mathbf{S} are transformed into the occupied subspace

$$\begin{aligned} \mathbf{H}_{\text{omm}} &= \mathbf{W}^* \mathbf{H} \mathbf{W}, \\ \mathbf{S}_{\text{omm}} &= \mathbf{W}^* \mathbf{S} \mathbf{W}, \end{aligned} \quad (1.15)$$

where \mathbf{W} is the coefficient matrix of the Wannier functions, whose dimension is $N_{\text{basis}} \times N_{\text{W}}$; \mathbf{H}_{omm} and \mathbf{S}_{omm} are $N_{\text{W}} \times N_{\text{W}}$ matrices. The OMM energy functional can then be evaluated from \mathbf{H}_{omm} and \mathbf{S}_{omm} :

$$E[\mathbf{W}] = 4\text{Tr}[\mathbf{H}_{\text{omm}}] - 2\text{Tr}[\mathbf{S}_{\text{omm}} \mathbf{H}_{\text{omm}}]. \quad (1.16)$$

This energy functional, when minimized with respect to the coefficients of Wannier functions \mathbf{W} , is equal to the “band structure” energy

$$E_{\text{BS}} = \sum_{l=1}^{N_{\text{basis}}} f_l \epsilon_l, \quad (1.17)$$

i.e. the sum of the energies of all eigenstates, weighted with their respective occupation numbers. Furthermore, the Wannier functions are driven towards perfect orthonormality at this minimum. The density matrix is then constructed from the Wannier functions that minimize $E[\mathbf{W}]$. Although this density matrix is sufficient for the electron density update following 1.7, without knowledge of individual eigenstates, the orbital minimization method cannot handle systems with fractional occupation numbers.

Different from the originally proposed linear scaling OMM method, the OMM implementation in the libOMM library [4] is a cubic scaling density matrix solver. Theoretically, this implementation has a smaller prefactor than the direct diagonalization method. In libOMM, the minimization of the OMM energy functional is carried out with the conjugate-gradient (CG) method, whose performance mainly depends on the convergence rate of the minimization.

1.3.3 PEXSI

The pole expansion and selected inversion (PEXSI) method [5, 6] expands the density matrix \mathbf{P} in 1.8 using a pole expansion:

$$\mathbf{P} = \sum \text{Im} \left(\frac{\omega_l}{\mathbf{H} - (z_l + \mu)\mathbf{S}} \right). \quad (1.18)$$

The shifts $\{z_l\}$ and weights $\{\omega_l\}$ of the poles are optimized to expand the Fermi operator. The number of terms needed by this expansion is proportional to $\log(\beta\Delta E)$, where $\beta = 1/(k_B T)$, k_B is the Boltzmann constant, T is the electronic temperature, and ΔE is the width of the eigenspectrum. This logarithmic scaling makes the pole expansion a highly efficient approach. In most cases, ~ 20 poles are already sufficient for the result obtained from PEXSI to be fully comparable to that obtained from diagonalization.

In PEXSI, only selected elements of the object $(\mathbf{H} - (z_l + \mu)\mathbf{S})^{-1}$ (and thus the density matrix), which correspond to the non-zero elements of \mathbf{H} and \mathbf{S} , are computed with the parallel selected inversion method. The computational cost scales at most as $O(N^2)$ for semilocal DFT. The actual complexity depends on the dimensionality of the system: $O(N)$, $O(N^{1.5})$, and $O(N^2)$ for 1D, 2D, and 3D systems, respectively. This favorable scaling hinges on the sparse character of the Hamiltonian and overlap matrices, but not on the existence of an energy gap. The PEXSI method is thus generally applicable to systems with and without a gap.

Designed in a multi-level parallelism structure, the PEXSI method is highly scalable, and can make efficient use of tens of thousands of processors on high performance computers.

1.3.4 SLEPc-SIPs

The shift-and-invert spectral transformation method, implemented in the SLEPc library [7], transforms the eigenproblem 1.5 by shifting the eigenspectrum:

$$(\mathbf{H} - \boldsymbol{\sigma}\mathbf{S}) = (\boldsymbol{\epsilon} - \boldsymbol{\sigma})\mathbf{S}\mathbf{c}, \quad (1.19)$$

where $\boldsymbol{\sigma}$ is a diagonal matrix with diagonal elements all equal to the shift σ . This shifted eigenproblem is converted to the standard form by inverting $(\mathbf{H} - \boldsymbol{\sigma}\mathbf{S})$ and $(\boldsymbol{\epsilon} - \boldsymbol{\sigma})$:

$$(\mathbf{H} - \boldsymbol{\sigma}\mathbf{S})^{-1}\mathbf{S}\mathbf{c} = (\boldsymbol{\epsilon} - \boldsymbol{\sigma})^{-1}\mathbf{c}, \quad (1.20)$$

which can be written in a form similar to 1.11:

$$\tilde{\mathbf{H}}\mathbf{c} = \tilde{\boldsymbol{\epsilon}}\mathbf{c}. \quad (1.21)$$

Here, the eigenvectors are not altered by the shift-and-invert transformation, and the eigenvalues of 1.21 relate to the original ones via

$$\tilde{\boldsymbol{\epsilon}} = (\boldsymbol{\epsilon} - \boldsymbol{\sigma})^{-1}. \quad (1.22)$$

If the shift can be chosen to be close to the target eigenvalue, 1.22 makes the magnitude of the transformed eigenvalues large, accelerating the convergence of the iterative Krylov-Schur eigensolver used in SLEPc.

On top of the basic shift-and-invert, the shift-and-invert parallel spectral transformation (SIPs) method [8] partitions the eigenspectrum of a given eigenproblem into N_{slice} slices. Accordingly, the processes involved in the calculation are split into N_{slice} groups, each of which solves one slice independently. Within the slices, carefully selected shifts are applied to the original problem. With this layer of parallelism across slices, the SLEPc-SIPs solver has the potential to exhibit enhanced scalability over direct diagonalization methods, especially when the load balance across slices can be guaranteed. Indeed, this has been reported to happen with very sparse Hamiltonian and overlap matrices out of density-functional tight-binding (DFTB) calculations [8].

1.3.5 NTPoly

Density matrix purification is an established way to achieve linear scaling KS-DFT. Assume orthogonal basis set, the density matrix \mathbf{P} should satisfy the following conditions:

$$\begin{aligned} \mathbf{P} &= \mathbf{P}^*, \\ \text{Tr}(\mathbf{P}) &= N_{\text{electron}}, \\ \mathbf{P} &= \mathbf{P}^2. \end{aligned} \quad (1.23)$$

An initial guess of such a density matrix can be obtained by scaling the Hamiltonian matrix to make sure its eigenvalues lie in between 0 and 1. Then, the “purification” method iteratively update the density matrix until it converges to a certain threshold. The converged density matrix satisfies the three conditions in 1.23.

The process of density matrix purification can be written in the general form

$$\mathbf{P}_{n+1} = f(\mathbf{P}_n), \quad (1.24)$$

where \mathbf{P}_n is the density matrix in the n^{th} purification iteration, \mathbf{P}_{n+1} is the density matrix in the $(n+1)^{\text{th}}$ iteration, and $f(\mathbf{P})$ is usually a matrix polynomial, which can be calculated by matrix-matrix multiplications.

Various algorithms have been developed to carry out the density matrix purification efficiently, such as the methods to purify the density matrix have been developed, such as the canonical purification [14], the trace resetting purification methods [15], and the generalized canonical purification [16]. These methods are implemented in the NTPoly library [9] using its sparse matrix-matrix multiplication kernel. Given sufficiently sparse matrices, the computational complexity of density matrix purification with NTPoly is $O(N)$ for insulating systems.

1.4 Citing ELSI

Key concepts of ELSI and the first version of its implementation are described in the following paper [17]:

V. W-z. Yu, F. Corsetti, A. García, W. P. Huhn, M. Jacquelin, W. Jia, B. Lange, L. Lin, J. Lu, W. Mi, A. Seifitokaldani, Á. Vázquez-Mayagoitia, C. Yang, H. Yang, and V. Blum, ELSI: A Unified Software Interface for Kohn-Sham Electronic Structure Solvers, *Computer Physics Communications*, 222, 267-285 (2018).

In addition, an incomplete list of publications describing the solvers supported in ELSI may be found in the bibliography of this document. Please consider citing these articles when publishing results obtained with ELSI.

1.5 Acknowledgments

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2 Installation of ELSI

2.1 Overview

The ELSI package contains the ELSI Interface software as well as redistributed source code for the solver libraries ELPA (version 2016.11.001), libOMM, PEXSI (version 1.0.3), and NTPoly (version 1.3). We highly encourage all users to request access to our GitLab server, where we regularly update ELSI between releases while preserving stability.

Starting from the May 2018 (version 2.0.0) release, the installation of ELSI makes use of the CMake software.

2.2 Prerequisites

To build ELSI, the minimum requirements are:

`CMake` [minimum version 3.0; newer version recommended]
`Fortran compiler` [with Fortran 2003]
`C compiler` [with C99]
`MPI`

Building the PEXSI solver (highly recommended) requires:

`C++ compiler` [with C++ 11]

Additionally, building the SLEPc-SIPs solver requires:

`SLEPc` [version 3.9.2 only]
`PETSc` [version 3.9.3 only, with SuperLU_DIST, MUMPS, ParMETIS, and PT-SCOTCH enabled]

Linear algebra libraries should be provided for ELSI to link against:

`BLAS`, `LAPACK`, `BLACS`, `ScaLAPACK`

By default, the redistributed ELPA, libOMM, and NTPoly solvers will be built. If PEXSI is enabled during configuration, the redistributed PEXSI library and its dependencies, namely the SuperLU_DIST and PT-SCOTCH libraries, will be built as well. Optionally, the redistributed ELPA, libOMM, SuperLU_DIST and PT-SCOTCH libraries may be substituted by user's optimized versions. Please note that in the current version of ELSI, an external version of PEXSI or NTPoly is not officially supported.

2.3 CMake Basics

This section covers some basics of using CMake. Users who are familiar with CMake may safely skip this section.

The typical workflow of using CMake to build ELSI looks like:

```
$ ls  
  
CMakeLists.txt  external/  src/  test/  ...
```

```

$ mkdir build
$ cd build
$ cmake [options] ..

...
...
-- Generating done
-- Build files have been written to: /current/dir

$ make [-j np]
$ make install

```

Whenever CMake is invoked, one of the command line arguments must point to the path where the top level CMakeLists.txt file exists, hence the “.” in the above example.

By default, CMake generates standard UNIX makefiles including specific rules to build the project with GNU make. Other build systems may be chosen with the “-G” (G for generator) option of CMake. We recommend Ninja in particular, which is a small build system with a focus on speed. A version of Ninja with Fortran support is freely available [here](#).

To build ELSI with Ninja:

```

$ ls

CMakeLists.txt  external/  src/  test/  ...

$ mkdir build
$ cd build
$ cmake -G Ninja [options] ..

...
...
-- Generating done
-- Build files have been written to: /current/dir

$ ninja
$ ninja install

```

Ninja also accepts the -j flag. Without this flag, Ninja runs on the number of available threads plus two by default (e.g., 10 on a machine with 8 threads). Thus, -j is typically not necessary.

An option may be defined by adding “-DKeyword=Value” to the command line when invoking CMake. If “Keyword” is of type boolean, its “Value” may be “ON” or “OFF”. If “Keyword” is a list of libraries or include directories, its items should be separated with “;” (semicolon) or “ ” (space).

For example,

```

-DCMAKE_INSTALL_PREFIX=/path/to/install/elsi
-DCMAKE_C_COMPILER=gcc
-DENABLE_TESTS=OFF
-DENABLE_PEXSI=ON
-DINC_PATHS="/path/to/include;/another/path/to/include"
-DLIBS="library1 library2 library3"

```

Available options for building ELSI with CMake are introduced in the next sections. Other options of CMake itself are available in its online documentation.

2.4 Configuration

2.4.1 Compilers

CMake automatically detects compilers. The choices made by CMake often work, but not necessarily lead to the optimal performance. In some cases, the compilers picked up by CMake may not be the ones desired by the user. To build ELSI, it is mandatory that the user explicitly sets the identification of the compilers:

```
-DCMAKE_Fortran_COMPILER=YOUR_MPI_FORTRAN_COMPILER
-DCMAKE_C_COMPILER=YOUR_MPI_C_COMPILER
-DCMAKE_CXX_COMPILER=YOUR_MPI_CXX_COMPILER
```

Please note that the C++ compiler is not needed when building ELSI without PEXSI.

In addition, it is highly recommended to specify the compiler flags, in particular the optimization flags:

```
-DCMAKE_Fortran_FLAGS=YOUR_FORTRAN_COMPILE_FLAGS
-DCMAKE_C_FLAGS=YOUR_MPI_C_COMPILE_FLAGS
-DCMAKE_CXX_FLAGS=YOUR_MPI_CXX_COMPILE_FLAGS
```

Note that with CMake versions older than 3.8.2, flags such as `-std=c99` and `-std=c++11` (or equivalents depending on the compilers) must be given in order to ensure compliance with the C99 and C++11 standards. Newer versions of CMake take care of this automatically.

2.4.2 Solvers

The ELPA, libOMM, PEXSI, and NTPoly solver libraries, as well as the SuperLU_DIST and PT-SCOTCH libraries (both required by PEXSI), are redistributed with the current ELSI package.

The redistributed version of ELPA comes with a few “kernels” specifically written to take advantage of processor architecture (e.g. vectorization instruction set extensions). A kernel may be chosen by the [ELPA2.KERNEL](#) keyword. Available options are:

```
-DELPA2_KERNEL=BGQ
-DELPA2_KERNEL=AVX
-DELPA2_KERNEL=AVX2
-DELPA2_KERNEL=AVX512
```

for the IBM Blue Gene Q, Intel AVX, Intel AVX2, and Intel AVX512 architectures, respectively. In ELPA, these kernels are employed to accelerate the calculation of eigenvectors, which is often a computational bottleneck when calculating a large percentage of eigenvectors. If this is the case in the user’s application, it is highly recommended that the user selects the kernel most suited to their system architecture.

Experienced users are encouraged to link the ELSI interface against external, better optimized solver libraries. Relevant options for this purpose are:

```
-DUSE_EXTERNAL_ELPA=ON
-DUSE_EXTERNAL_OMM=ON
-DUSE_EXTERNAL_SUPERLU=ON
```

The external libraries and the include paths should be set via the following three keywords:

```
-DLIB_PATHS=DIRECTORIES_CONTAINING_YOUR_EXTERNAL_LIBRARIES
-DINC_PATHS=INCLUDE_DIRECTORIES_OF_YOUR_EXTERNAL_LIBRARIES
-DLIBS=NAMES_OF_YOUR_EXTERNAL_LIBRARIES
```

Each of the above keywords is a space-separated or semicolon-separated list. If an external library depends on additional libraries, [LIBS](#) should include all the relevant libraries. For instance, [LIBS](#) should include the ELPA library and CUDA libraries when using an external ELPA compiled with GPU (CUDA) support; [LIBS](#) should include the SuperLU_DIST library and the sparse matrix reordering library used to compile SuperLU_DIST when using an external SuperLU_DIST. Please note that in the current version of ELSI, an external version of PEXSI or NTPoly is not officially supported.

The PEXSI and SLEPc-SIPs solvers are not enabled by default. PEXSI may be activated by specifying:

`-DENABLE_PEXSI=ON`

if using redistributed SuperLU_DIST with PT-SCOTCH, or

```
-DENABLE_PEXSI=ON
-DUSE_EXTERNAL_SUPERLU=ON
-DINC_PATHS="/path/to/superlu_dist/include;/path/to/matrix/reordering/include"
-DLIB_PATHS="/path/to/superlu_dist/library;/path/to/matrix/reordering/include"
-DLIBS="superlu_dist;your_choice_of_matrix_reordering_library"
```

if using an externally compiled SuperLU_DIST. SuperLU_DIST 5.3.0 has been tested with this version of ELSI. Older/newer versions may or may not be compatible.

SLEPc-SIPs may be activated by specifying:

```
-DENABLE_SIPS=ON
-DUSE_EXTERNAL_SUPERLU=ON
-DINC_PATHS="/path/to/slepc/include;/path/to/slepc/${PETSC_ARCH}/include;
/path/to/petsc/include;/path/to/${PETSC_ARCH}/include"
-DLIB_PATHS="/path/to/slepc/${PETSC_ARCH}/library;/path/to/petsc/${PETSC_ARCH}/library"
-DLIBS="slepc;petsc;cmumps;dmumps;smumps;zmumps;mumps_common;pord;superlu_dist;parmetis;
metis;ptesmumps;ptscotchparmetis;ptscotch;ptscotcherr;esmumps;scotchmetis;scotch;scotcherr"
```

SLEPc 3.9.2 and PETSc 3.9.3 have been tested with this version of ELSI. Older/newer versions may or may not be compatible. The PETSc library must be compiled with MPI support, and (at least) with external packages SuperLU_DIST, MUMPS, ParMETIS, and PT-SCOTCH enabled. The SuperLU_DIST library redistributed through ELSI must be turned off by setting `USE_EXTERNAL_SUPERLU` to “ON”, as SuperLU_DIST is already present in the PETSc installation.

2.4.3 Build Targets

By default, a static library (libelsi.a) will be created as the target of the compilation. Building ELSI as a shared library may be enabled by:

`-DBUILD_SHARED_LIBS=ON`

Building ELSI test programs may be enabled by:

`-DENABLE_TESTS=ON`

In either case, linear algebra libraries, BLAS, LAPACK, BLACS, and ScaLAPACK, should be valid in the `LIB_PATHS` and `LIBS` keywords.

If test programs are turned on, the compilation of ELSI may be verified by

```
$ make test
```

or

```
$ ninja test
```

depending on the generator option “-G” used when invoking CMake. Alternatively, issue

```
$ ctest
```

to invoke the CTest program which performs all tests automatically.

Note that the tests may not run if launching MPI jobs is prohibited on the user’s working platform.

In order to install ELSI at the location specified by `CMAKE_INSTALL_PREFIX`, issue

```
$ make install
```

or

```
$ ninja install
```

depending on the CMake generator option “-G” used.

Among the files copied to the installation destinations is a CMake configuration file called `elsiConfig.cmake`. This file includes all the information about how the ELSI library and its dependencies should be included in an external CMake project. Please refer to 2.5 for information regarding linking a third-party package against ELSI.

2.4.4 List of All Configure Options

The options accepted by the ELSI CMake build system are listed here in alphabetical order. Some additional explanations are made below the table.

| Option | Type | Default | Explanation |
|--|---------|-------------|--|
| ADD_UNDERSCORE | boolean | ON | Suffix C functions with an underscore |
| BUILD_SHARED_LIBS | boolean | OFF | Build ELSI as a shared library |
| CMAKE_C_COMPILER | string | none | MPI C compiler |
| CMAKE_C_FLAGS | string | none | C flags |
| CMAKE_CXX_COMPILER | string | none | MPI C++ compiler |
| CMAKE_CXX_FLAGS | string | none | C++ flags |
| CMAKE_Fortran_COMPILER | string | none | MPI Fortran compiler |
| CMAKE_Fortran_FLAGS | string | none | Fortran flags |
| CMAKE_INSTALL_PREFIX | path | /usr/local | Path to install ELSI |
| ELPA2_KERNEL | string | none | ELPA2 kernel |
| ENABLE_C_TESTS | boolean | OFF | Build C test programs |
| ENABLE_PEXSI | boolean | OFF | Enable PEXSI support |
| ENABLE_SIPS | boolean | OFF | Enable SLEPc-SIPs support |
| ENABLE_TESTS | boolean | OFF | Build Fortran test programs |
| INC_PATHS | string | none | Include directories of external libraries |
| LIB_PATHS | string | none | Directories containing external libraries |
| LIBS | string | none | External libraries |
| MPIEXEC_NP | string | mpirun -n 4 | Command to run tests in parallel with MPI |
| MPIEXEC_1P | string | mpirun -n 1 | Command to run tests in serial with MPI |
| SCOTCH_LAST_RESORT | string | none | Command to invoke PT-SCOTCH header generator |
| USE_EXTERNAL_ELPA | boolean | OFF | Use external ELPA |
| USE_EXTERNAL_OMM | boolean | OFF | Use external libOMM and MatrixSwitch |
| USE_EXTERNAL_SUPERLU | boolean | OFF | Use external SuperLU_DIST |

Remarks

1) [ADD_UNDERSCORE](#): In the PEXSI and SuperLU_DIST code redistributed through ELSI, there are calls to functions of the linear algebra libraries, e.g. “dgemm”. If [ADD_UNDERSCORE](#) is “ON”, the code will call “dgemm_” instead of “dgemm”. Turn this keyword on if routines are suffixed with “_” in external linear algebra libraries. Turn it off if routines are not suffixed with “_”.

2) [ELPA2_KERNEL](#): There are a number of computational kernels available with the ELPA solver. Choose from “BGQ” (IBM Blue Gene Q), “AVX” (Intel AVX), “AVX2” (Intel AVX2), and “AVX512” (Intel AVX512). See 2.4.2 for more information.

3) [SCOTCH_LAST_RESORT](#): The compilation of the PT-SCOTCH library is a multi-step process. First, two auxiliary executables are created. Then, header files of the library are generated by running the two executables. Finally, the main source files of the library are compiled with the generated header files included. The header generation step may fail on platforms where directly running an executable is prohibited on a login/compile node. Often this can be circumvented by requesting an interactive session to a compute node and performing the compilation there, or by submitting the whole compilation as a job to the queuing system. However, this may still fail on platforms where an executable compiled with MPI must be launched by an MPI job launcher (aprun, mpirun, srun, etc). If the standard compilation of PT-SCOTCH fails due to this reason, the user may set [SCOTCH_LAST_RESORT](#) to the command that starts an MPI job with one

MPI task, e.g. “`mpirun -n 1`”. This command will be used to launch the auxiliary executables to generate necessary header files for PT-SCOTCH.

4) External libraries: ELSI redistributes source code of ELPA, libOMM, PEXSI, SuperLU_DIST, and PT-SCOTCH libraries, which by default will be built together with the ELSI interface. Experienced users are encouraged to link the ELSI interface against external, better optimized solver libraries. See 2.4.2 for more information.

2.4.5 “Toolchain” Files

It is sometimes convenient to edit the settings in a “toolchain” file that can be read by CMake:

```
-DCMAKE_TOOLCHAIN_FILE=YOUR_TOOLCHAIN_FILE
```

Example “toolchains” are provided in the “./toolchains” directory of the ELSI package, which the user may use as templates to create new ones.

2.5 Importing ELSI into Third-Party Code Projects

2.5.1 Linking against ELSI: CMake

A CMake configuration file called `elsiConfig.cmake` should be generated after ELSI is successfully installed (see 2.4.3). This file contains all the information about how the ELSI library and its dependencies should be included in an external project. For a project using CMake, only two lines are required to find and link to ELSI:

```
find_package(elsi REQUIRED)
target_link_libraries(my_project PRIVATE elsi::elsi)
```

If a minimum version of ELSI is required, this information may be passed to “`find_package`” by:

```
find_package(elsi 2.0 REQUIRED)
```

If the installed ELSI version is older than the requested minimum version, CMake stops with an appropriate error message. Other options of “`find_package`” are available in the documentation of CMake.

2.5.2 Linking against ELSI: Makefile

For a project using makefiles, an example set of compiler flags to link against ELSI would be:

```
ELSI_INCLUDE = -I/PATH/TO/BUILD/ELSI/include
ELSI_LIB      = -L/PATH/TO/BUILD/ELSI/lib -lelsi \
               -lfortjson -lomm -lMatrixSwitch -lelpa \
               -lNTPoly -lpexsi -lsuperlu_dist \
               -lptscotchparmetis -lptscotch -lptscotcherr \
               -lscotchmetis -lscotch -lscotcherr
```

Enabling/disabling PEXSI and SLEPc-SIPs or linking ELSI against preinstalled solver libraries will require the user modify these flags accordingly.

2.5.3 Using ELSI

ELSI may be used in an electronic structure code by importing the appropriate header file. For codes written in Fortran, this is done by using the ELSI module

```
USE ELSI
```

For codes written in C, the ELSI wrapper may be imported by including the header file

```
#include <elsi.h>
```

These import statements give the electronic structure code access to the ELSI interface. In the next chapter, we will describe the API for the ELSI interface.

3 The ELSI API

3.1 Overview of the ELSI API

In this chapter, we present the public-facing API for the ELSI Interface. We anticipate that fine details of this interface may change slightly in the future, but the fundamental structure of the interface layer is expected to remain consistent. While this chapter serves as a reference to the ELSI subroutines, the user is encouraged to explore the demonstration pseudo-codes of ELSI in 3.8.

To allow multiple instances of ELSI to co-exist within a single calling code, we define an `elsi_handle` data type to encapsulate the state of an ELSI instance, i.e., all runtime parameters associated with the ELSI instance. An `elsi_handle` instance is initialized with the `elsi_init` subroutine and is subsequently passed to all other ELSI subroutine calls.

ELSI provides a C interface in addition to the native Fortran interface. The vast majority of this chapter, while written from a Fortran-ic standpoint, applies equally to both interfaces. Information specifically about the C wrapper for ELSI may be found in 3.9.

3.2 Setting Up ELSI

3.2.1 Initializing ELSI

The ELSI interface must be initialized via the `elsi_init` subroutine before any other ELSI subroutine may be called.

`elsi_init`(handle, solver, parallel_mode, matrix_format, n_basis, n_electron, n_state)

| Argument | Data Type | in/out | Explanation |
|----------------------------|--------------------------------|--------|---|
| <code>handle</code> | <code>type(elsi_handle)</code> | out | Handle to ELSI. |
| <code>solver</code> | integer | in | Desired KS solver. Accepted values are: 1 (ELPA), 2 (LI-BOMM), 3 (PEXSI), 5 (SLEPc-SIPs), and 6 (NTPoly). |
| <code>parallel_mode</code> | integer | in | Parallelization mode. See remark 3. Accepted values are: 0 (SINGLE_PROC) and 1 (MULTIPROC). |
| <code>matrix_format</code> | integer | in | Matrix format. See remark 1. Accepted values are: 0 (BLACS_DENSE), 1 (PEXSI_CSC), and 2 (SIESTA_CSC). |
| <code>n_basis</code> | integer | in | Number of basis functions, i.e. global size of Hamiltonian. |
| <code>n_electron</code> | real double | in | Number of electrons. |
| <code>n_state</code> | integer | in | Number of states. See remark 2. |

Remarks

1) `matrix_format`: `BLACS_DENSE`(0) refers a dense matrix format in a 2-dimensional block-cyclic distribution, i.e. the BLACS standard. `PEXSI_CSC`(1) refers to a compressed sparse column (CSC) matrix format in a 1-dimensional block distribution. `SIESTA_CSC`(2) refers to a compressed sparse column (CSC) matrix format in a 1-dimensional block-cyclic distribution. As the Hamiltonian, overlap, and density matrices are symmetric (Hermitian), compressed sparse row (CSR) matrix format is effectively supported.

2) **n.state**: If ELPA or SLEPc-SIPs is the chosen solver, this parameter specifies the number of eigenstates to solve by the eigensolver. If libOMM is the chosen solver, **n.state** must be exactly the number of occupied states, as libOMM cannot handle fractional occupation numbers[4]. PEXSI and NTPoly do not make use of this parameter, thus a dummy value may be passed.

3) **parallel_mode**: The two allowed values of **parallel_mode**, 0 (**SINGLE_PROC**) and 1 (**MULTI_PROC**), allow for three parallelization strategies commonly employed by electronic structure codes. See below.

3a) **SINGLE_PROC**: Solves the KS eigenproblem following a LAPACK-like fashion. This option may only be selected when ELPA is chosen as the solver. This allows the following parallelization strategy:

SINGLE_PROC Example:

Every MPI task independently handles a group of **k**-points uniquely assigned to it.

Example number of **k**-points: 16

Example number of MPI tasks: 4

MPI task 0 handles **k**-points 1, 2, 3, 4 sequentially;
 MPI task 1 handles **k**-points 5, 6, 7, 8 sequentially;
 MPI task 2 handles **k**-points 9, 10, 11, 12 sequentially;
 MPI task 3 handles **k**-points 13, 14, 15, 16 sequentially.

Pseudocode 1:

```
elsi_init(elsi_h, ..., parallel_mode=0, ...)
...
for i_kpt = 1, n_kpt_local, 1 do
    | elsi_ev_{real|complex}(elsi_h, ham_this_kpt, ovlp_this_kpt, eval_this_kpt, evvec_this_kpt)
end
```

3b) **MULTI_PROC**: Solves the KS eigenproblem following a ScaLAPACK-like fashion. This allows the usage of the following two parallelization strategies:

MULTI_PROC Example:

Groups of MPI tasks coordinate to handle the same **k**-point, uniquely assigned to that group.

Example number of **k**-points: 4

Example number of MPI tasks: 16

MPI tasks 0, 1, 2, 3 cooperatively handle **k**-point 1;
 MPI tasks 4, 5, 6, 7 cooperatively handle **k**-point 2;
 MPI tasks 8, 9, 10, 11 cooperatively handle **k**-point 3;
 MPI tasks 12, 13, 14, 15 cooperatively handle **k**-point 4.

Pseudocode 2:

```
elsi_init(elsi_h, ..., parallel_mode=1, ...)
elsi_set_mpi(elsi_h, my_mpi_comm)
...
elsi_ev_{real|complex}(elsi_h, my_ham, my_ovlp, my_eval, my_evec)
```

or

```
elsi_init(elsi_h, ..., parallel_mode=1, ...)
elsi_set_mpi(elsi_h, my_mpi_comm)
elsi_set_kpoint(elsi_h, n_kpt, my_kpt, my_weight)
elsi_set_mpi_global(elsi_h, mpi_comm_global)
...
elsi_dm_complex(elsi_h, my_ham, my_ovlp, my_dm, global_energy)
```

Please note that when there is more than one k -point, a global MPI communicator must be provided for inter- k -point communications. See 3.2.4 for [elsi_set_kpoint](#), [elsi_set_spin](#), and [elsi_set_mpi_global](#), which are used to set up a calculation with two spin channels and/or multiple k -points.

3.2.2 Setting Up MPI

The MPI communicator used by ELSI is passed into ELSI by the calling code via the [elsi_set_mpi](#) subroutine. When there is more than one k -point and/or spin channel, this communicator will be used only for solving one problem corresponding to one k -point and one spin channel. See 3.2.4 for details.

```
elsi_set_mpi(handle, mpi_comm)
```

| Argument | Data Type | in/out | Explanation |
|--------------------------|-------------------|--------|-------------------|
| handle | type(elsi_handle) | inout | Handle to ELSI. |
| mpi_comm | integer | in | MPI communicator. |

3.2.3 Setting Up Matrix Formats

When using the 2D block-cyclic distributed dense matrix format ([BLACS_DENSE](#)), BLACS parameters are passed into ELSI via the [elsi_set_blacs](#) subroutine. The matrix format used internally in the ELSI interface and the ELPA solver requires the block sizes of the 2-dimensional block-cyclic distribution are the same in the row and column directions. It is necessary to call this subroutine before calling any solver interface that makes use of the [BLACS_DENSE](#) matrix format.

```
elsi_set_blacs(handle, blacs_ctxt, block_size)
```

| Argument | Data Type | in/out | Explanation |
|----------------------------|-------------------|--------|--|
| handle | type(elsi_handle) | inout | Handle to ELSI. |
| blacs_ctxt | integer | in | BLACS context. |
| block_size | integer | in | Block size of the 2D block-cyclic distribution, specifying both row and column directions. |

When using the sparse matrix formats, namely 1D block distributed compressed sparse column ([PEXSI_CSC](#)) or 1D block-cyclic distributed compressed sparse column ([SIESTA_CSC](#)), the sparsity pattern should be passed into ELSI via the [elsi_set_csc](#) subroutine. It is necessary to call this subroutine before calling any solver interface that makes use of the sparse matrix formats.

`elsi_set_csc(handle, global_nnz, local_nnz, local_col, row_idx, col_ptr)`

| Argument | Data Type | in/out | Explanation |
|-------------------------|--------------------------------|--------|---|
| <code>handle</code> | <code>type(elsi_handle)</code> | inout | Handle to ELSI. |
| <code>global_nnz</code> | integer | in | Global number of non-zeros. |
| <code>local_nnz</code> | integer | in | Local number of non-zeros. |
| <code>local_col</code> | integer | in | Local number of matrix columns. |
| <code>row_idx</code> | integer, rank-1 array | in | Local row index array. Dimension: <code>local_nnz</code> . |
| <code>col_ptr</code> | integer, rank-1 array | in | Local column pointer array. Dimension: <code>local_col+1</code> . |

When using the 1D block distributed compressed sparse column (**PEXSI_CSC**) format, the block size of the distribution cannot be set by the user. This is because the PEXSI solver requires that the block size must be $\text{floor}(\text{N_basis}/\text{N_procs})$, where $\text{floor}(x)$ is the greatest integer less than or equal to x , `N_basis` and `N_procs` are the number of basis functions and the number of MPI tasks, respectively. When using the 1D block-cyclic distributed compressed sparse column (**SIESTA_CSC**) format, the block size of the 1D distribution must be explicitly set by calling `elsi_set_csc_blk`.

`elsi_set_csc_blk(handle, block_size)`

| Argument | Data Type | in/out | Explanation |
|-------------------------|--------------------------------|--------|---|
| <code>handle</code> | <code>type(elsi_handle)</code> | inout | Handle to ELSI. |
| <code>global_nnz</code> | integer | in | Block size of the 1D block-cyclic distribution. |

In most cases, input and output matrices should be distributed across all MPI tasks. The only exception happens when using the PEXSI solver, one of the sparse density matrix interfaces (`elsi_dm_real_sparse` or `elsi_dm_complex_sparse`), and the **PEXSI_CSC** matrix format. In this case, an additional parameter, `plexsi_np_per_pole`, must be set by the user. Input and output matrices should be 1D-block-distributed among the first `plexsi_np_per_pole` MPI tasks (not all the MPI tasks). Please also read the 2nd remark in 3.5.4 for more information.

3.2.4 Setting Up Multiple \mathbf{k} -points and/or Spin Channels

When there is more than one \mathbf{k} -point and/or spin channel in the simulating system, the ELSI interface can be set up to support parallel calculation of the \mathbf{k} -points and/or spin channels. The base case is an isolated system, e.g. atoms, molecules, clusters, without spin-polarization. In this case, in each SCF iteration, there is one KS eigenproblem (1.5) to solve. If the isolated system is spin-polarized, there are two eigenproblems:

$$\begin{aligned} \mathbf{H}_\alpha \mathbf{c}_\alpha &= \epsilon_\alpha \mathbf{S}_\alpha \mathbf{c}_\alpha, \\ \mathbf{H}_\beta \mathbf{c}_\beta &= \epsilon_\beta \mathbf{S}_\beta \mathbf{c}_\beta, \end{aligned} \quad (3.1)$$

where α and β denote the two spin channels. The overlap matrices are generally the same for the two spin channels, but the Hamiltonian matrices are not. These two eigenproblems can be solved one after another using all available processes, or can be solved concurrently using half of the processes for each spin channel.

If the system is periodically repeated in space, according to the Bloch theorem, the KS eigenproblem has an additional index \mathbf{k} :

$$\mathbf{H}_\mathbf{k} \mathbf{c}_\mathbf{k} = \epsilon_\mathbf{k} \mathbf{S}_\mathbf{k} \mathbf{c}_\mathbf{k}. \quad (3.2)$$

In practice, it is sufficient to study \mathbf{k} within a single primitive unit cell in the reciprocal space, usually the first Brillouin zone. The physical quantities, e.g. the electron density, are represented by Brillouin zone integrals:

$$n(\mathbf{r}) = \sum_{l=1}^{N_{\text{basis}}} \int_{BZ} f_{l\mathbf{k}} \psi_{l\mathbf{k}}^*(\mathbf{r}) \psi_{l\mathbf{k}}(\mathbf{r}) d^3k, \quad (3.3)$$

which is approximated by using a finite mesh of \mathbf{k} -points in the first Brillouin zone:

$$n(\mathbf{r}) \approx \sum_{n=1}^{N_{\text{kpt}}} w_n \sum_{l=1}^{N_{\text{basis}}} f_{l\mathbf{k}_n} \psi_{l\mathbf{k}_n}^*(\mathbf{r}) \psi_{l\mathbf{k}_n}(\mathbf{r}). \quad (3.4)$$

Here, w_n is the weight of the n^{th} \mathbf{k} -point; N_{kpt} is the number of \mathbf{k} -points. The weights of all \mathbf{k} -points add up to 1. Obviously, a denser grid of \mathbf{k} -points leads to a higher accuracy with higher computational cost. The Hamilton and overlap matrices for multiple \mathbf{k} -points are block-diagonal, such that each block on the diagonal corresponds to an eigenproblem of one \mathbf{k} -point. These eigenproblems can be solved separately.

The handling of spin-polarized case and periodic case in ELSI are more or less equivalent. The problems, either from two spin channels, or from multiple \mathbf{k} -points, are treated as equivalent “unit tasks”. If the chosen solver is an eigensolver (e.g. ELPA), all the unit tasks are solved independently, returning separate eigensolutions to the electronic structure code. The electronic structure code can then assemble the pieces of the solutions and construct the electron density. When computing density matrices, the unit tasks are coupled together by the normalization condition of the number of electrons:

$$N_{\text{electron}} = \sum_{n=1}^{N_{\text{kpt}}} \sum_{m=1}^{N_{\text{spin}}} \sum_{l=1}^{N_{\text{basis}}} w_n f_{lmn}, \quad (3.5)$$

where N_{kpt} , N_{spin} , and N_{basis} are the number of \mathbf{k} -points, the number of spin channels, and the number of basis functions, respectively. w_n is again the weight of the n^{th} \mathbf{k} -point. f_{lmn} is the occupation number of the l^{th} state in the m^{th} spin channel and the n^{th} \mathbf{k} -point. To determine the occupation numbers, the eigenvalues at each unit task need to be collected across all the tasks. With the correct occupation numbers, density matrices can be computed by 1.8.

If the PEXSI solver is chosen, the pole expansion in 1.18 is performed for all the unit tasks in parallel, with the same trial chemical potential μ . The resulting number of electrons needs to be determined in order to refine the chemical potential. The chemical potential yielding the correct number of electrons is used to construct the density matrices on the unit tasks. If the OMM solver is chosen, the orbital minimization in 1.16 is performed for all the unit tasks to obtain density matrices. Again, OMM cannot handle systems with fractional occupation numbers.

At present, the SLEPc-SIPs and NTPoly solvers are not compatible with spin-polarized and/or periodic calculations.

To set up the ELSI interface for a calculation with more than one \mathbf{k} -point and/or more than one spin channel, the `elsi_set_kpoint` and/or `elsi_set_spin` subroutines are called to pass the required information into ELSI. The MPI communicator for each unit task is passed into ELSI by calling `elsi_set_mpi`. In addition, a global MPI communicator for all tasks is passed into ELSI by calling `elsi_set_mpi_global`. Note that the current ELSI interface only supports the case where the eigenproblems for all the \mathbf{k} -points and spin channels are fully parallelized, i.e., there is no MPI task handling more than one \mathbf{k} -point and/or more than one spin channel. Another limitation is that the two spin channels are always coupled by the normalization condition 3.5, with a uniform chemical potential for the two channels. The distribution of electrons among the two channels, and thus the net spin moment of the system, is solely determined by 3.5. Future work will enable calculations with a fixed, user-specified spin moment.

`elsi_set_kpoint(handle, n_kpt, i_kpt, weight)`

| Argument | Data Type | in/out | Explanation |
|---------------------|--------------------------------|--------|---|
| <code>handle</code> | <code>type(elsi_handle)</code> | inout | Handle to ELSI. |
| <code>n_kpt</code> | integer | in | Total number of \mathbf{k} -points. |
| <code>i_kpt</code> | integer | in | Index of the \mathbf{k} -point handled by this MPI task. |
| <code>weight</code> | integer | in | Weight of the \mathbf{k} -point handled by this MPI task. |

`elsi_set_spin(handle, n_spin, i_spin)`

| Argument | Data Type | in/out | Explanation |
|---------------------|--------------------------------|--------|---|
| <code>handle</code> | <code>type(elsi_handle)</code> | inout | Handle to ELSI. |
| <code>n_spin</code> | integer | in | Total number of spin channels. |
| <code>i_spin</code> | integer | in | Index of the spin channel handled by this MPI task. |

`elsi_set_mpi_global(handle, mpi_comm_global)`

| Argument | Data Type | in/out | Explanation |
|------------------------------|--------------------------------|--------|---|
| <code>handle</code> | <code>type(elsi_handle)</code> | inout | Handle to ELSI. |
| <code>mpi_comm_global</code> | integer | in | Global MPI communicator used for communications among all \mathbf{k} -points and spin channels. |

3.2.5 Finalizing ELSI

When an ELSI instance is no longer needed, its associated handle should be cleaned up by calling [elsi_finalize](#).

[elsi_finalize](#)(handle)

| Argument | Data Type | in/out | Explanation |
|------------------------|-------------------|--------|-----------------|
| handle | type(elsi_handle) | inout | Handle to ELSI. |

3.3 Solving Eigenvalues and Eigenvectors

The following subroutines return all the eigenvalues and a subset of eigenvectors of the provided generalized eigenproblem defined by H and S matrices. For standard eigenproblems, please see [elsi_set_unit_ovlp](#) in 3.5.1. Only ELPA and SLEPc-SIPs may be selected as the desired solver when using these subroutines.

[elsi_ev_real](#)(handle, ham, ovlp, eval, evec)

| Argument | Data Type | in/out | Explanation |
|------------------------|---------------------------|--------|---|
| handle | type(elsi_handle) | inout | Handle to ELSI. |
| ham | real double, rank-2 array | inout | Real Hamiltonian matrix in 2D block-cyclic dense format. See remark 1. |
| ovlp | real double, rank-2 array | inout | Real overlap matrix (or its Cholesky factor) in 2D block-cyclic dense format. See remark 1. |
| eval | real double, rank-1 array | inout | Eigenvalues. See remark 2. |
| evec | real double, rank-2 array | out | Real eigenvectors in 2D block-cyclic dense format. See remark 3. |

[elsi_ev_complex](#)(handle, ham, ovlp, eval, evec)

| Argument | Data Type | in/out | Explanation |
|------------------------|------------------------------|--------|--|
| handle | type(elsi_handle) | inout | Handle to ELSI. |
| ham | complex double, rank-2 array | inout | Complex Hamiltonian matrix in 2D block-cyclic dense format. See remark 1. |
| ovlp | complex double, rank-2 array | inout | Complex overlap matrix (or its Cholesky factor) in 2D block-cyclic dense format. See remark 1. |
| eval | real double, rank-1 array | inout | Eigenvalues. See remark 2. |
| evec | complex double, rank-2 array | out | Complex eigenvectors in 2D block-cyclic dense format. See remark 3. |

[elsi_ev_real_sparse](#)(handle, ham, ovlp, eval, evec)

| Argument | Data Type | in/out | Explanation |
|------------------------|---------------------------|--------|--|
| handle | type(elsi_handle) | inout | Handle to ELSI. |
| ham | real double, rank-1 array | inout | Real Hamiltonian matrix in 1D block or block-cyclic CSC sparse format. |
| ovlp | real double, rank-1 array | inout | Real overlap matrix in 1D block or block-cyclic CSC sparse format. |
| eval | real double, rank-1 array | inout | Eigenvalues. See remark 2. |
| evec | real double, rank-2 array | out | Real eigenvectors in 2D block-cyclic dense format. See remark 3. |

`elsi_ev_complex_sparse(handle, ham, ovlp, eval, evec)`

| Argument | Data Type | in/out | Explanation |
|---------------------|--------------------------------|--------|---|
| <code>handle</code> | <code>type(elsi_handle)</code> | inout | Handle to ELSI. |
| <code>ham</code> | complex double, rank-1 array | inout | Complex Hamiltonian matrix in 1D block or block-cyclic CSC sparse format. |
| <code>ovlp</code> | complex double, rank-1 array | inout | Complex overlap matrix in 1D block or block-cyclic CSC sparse format. |
| <code>eval</code> | real double, rank-1 array | inout | Eigenvalues. See remark 2. |
| <code>evec</code> | complex double, rank-2 array | out | Complex eigenvectors in 2D block-cyclic dense format. See remark 3. |

Remarks

- 1) The Hamiltonian matrix will be destroyed by ELPA during computation. ELPA will overwrite the overlap matrix in its initial execution with the Cholesky factor, which will be reused by subsequent subroutine calls to `elsi_ev_real` or `elsi_ev_complex`. When using `elsi_ev_real_sparse`, the Cholesky factor (which is not sparse) is stored internally in the BLACS_DENSE format. Starting from the second call to `elsi_ev_real_sparse`, the input sparse overlap matrix will not be referenced.
- 2) When using the ELPA solver, `elsi_ev_real`, `elsi_ev_complex`, `elsi_ev_real_sparse`, and `elsi_ev_complex_sparse` always compute all the eigenvalues, regardless of the choice of `n_state` specified in `elsi_init`. The dimension of `eval` thus should always be `n_basis`.
- 3) When using the ELPA solver, `elsi_ev_real`, `elsi_ev_complex`, `elsi_ev_real_sparse`, and `elsi_ev_complex_sparse` compute a subset of all eigenvectors. The number of eigenvectors to compute is specified by the keyword `n_state` in `elsi_init`. However, the local `eigenvectors` array should always be initialized to correspond to a global array of size `n_basis` \times `n_basis`, whose extra part is used as working space in ELPA. Note that when using `elsi_ev_real_sparse` and `elsi_ev_complex_sparse`, the eigenvectors are returned in a dense format (BLACS_DENSE), as they are in general not sparse.

3.4 Computing Density Matrices

The following subroutines return the density matrix computed from the provided H and S matrices, as well as the energy corresponding to the occupied eigenstates. When the selected solver is ELPA, ELSI will internally construct the density matrix using the eigenvalues and eigenvectors returned by ELPA.

`elsi_dm_real(handle, ham, ovlp, dm, bs_energy)`

| Argument | Data Type | in/out | Explanation |
|---------------------|--------------------------------|--------|---|
| <code>handle</code> | <code>type(elsi_handle)</code> | inout | Handle to ELSI. |
| <code>ham</code> | real double, rank-2 array | inout | Real Hamiltonian matrix in 2D block-cyclic dense format. |
| <code>ovlp</code> | real double, rank-2 array | inout | Real overlap matrix (or Cholesky factor) in 2D block-cyclic dense format. See remark 1. |
| <code>dm</code> | real double, rank-2 array | out | Real density matrix in 2D block-cyclic dense format. |
| <code>energy</code> | real double | out | Energy corresponding to the occupied eigenstates ("band structure energy"). |

[elsi_dm_complex](#)(handle, ham, ovlp, dm, energy)

| Argument | Data Type | in/out | Explanation |
|------------------------|------------------------------|--------|--|
| handle | type(elsi_handle) | inout | Handle to ELSI. |
| ham | complex double, rank-2 array | inout | Complex Hamiltonian matrix in 2D block-cyclic dense format. |
| ovlp | complex double, rank-2 array | inout | Complex overlap matrix (or its Cholesky factor) in 2D block-cyclic dense format. See remark 1. |
| dm | complex double, rank-2 array | out | Complex density matrix in 2D block-cyclic dense format. |
| energy | real double | out | Energy corresponding to the occupied eigenstates (“band structure energy”). |

[elsi_dm_real_sparse](#)(handle, ham, ovlp, dm, energy)

| Argument | Data Type | in/out | Explanation |
|------------------------|---------------------------|--------|--|
| handle | type(elsi_handle) | inout | Handle to ELSI. |
| ham | real double, rank-1 array | inout | Non-zero values of the real Hamiltonian matrix in 1D block or block-cyclic CSC format. |
| ovlp | real double, rank-1 array | inout | Non-zero values of the real overlap matrix in 1D block or block-cyclic CSC format. |
| dm | real double, rank-1 array | out | Non-zero values of the real density matrix in 1D block or block-cyclic CSC format. |
| energy | real double | out | Energy corresponding to the occupied eigenstates (“band structure energy”). |

[elsi_dm_complex_sparse](#)(handle, ham, ovlp, dm, energy)

| Argument | Data Type | in/out | Explanation |
|------------------------|------------------------------|--------|---|
| handle | type(elsi_handle) | inout | Handle to ELSI. |
| ham | complex double, rank-1 array | inout | Non-zero values of the complex Hamiltonian matrix in 1D block or block-cyclic CSC format. |
| ovlp | complex double, rank-1 array | inout | Non-zero values of the complex overlap matrix in 1D block or block-cyclic CSC format. |
| dm | complex double, rank-1 array | out | Non-zero values of the complex density matrix in 1D block or block-cyclic CSC format. |
| energy | real double | out | Energy corresponding to the occupied eigenstates (“band structure energy”). |

Remarks

1) If the chosen solver is ELPA or libOMM, the Hamiltonian matrix will be destroyed during the computation. ELPA will overwrite the overlap matrix in its initial execution with the Cholesky factor, which will be reused by subsequent calls to [elsi_dm_real](#).

3.5 Customizing ELSI

In ELSI, reasonable default values have been provided for a number of parameters used in the ELSI interface the the supported solvers. However, no set of default parameters can adequately cover all use cases. Parameters that can be overridden are described in the following subsections.

3.5.1 Customizing the ELSI Interface

In all the subroutines listed below, the first argument (input and output) is an `elsi_handle`. The second argument (input) of each subroutine is the name of parameter to set.

Note that logical variables are not used in all ELSI API. Integers are used to represent logical, with 0 being false and any positive integer being true.

`elsi_set_output(handle, out_level)`
`elsi_set_output_log(handle, out_log)`
`elsi_set_write_unit(handle, write_unit)`
`elsi_set_unit_ovlp(handle, unit_ovlp)`
`elsi_set_zero_def(handle, zero_def)`
`elsi_set_sing_check(handle, sing_check)`
`elsi_set_sing_tol(handle, sing_tol)`
`elsi_set_sing_stop(handle, sing_stop)`
`elsi_set_mu_broaden_scheme(handle, mu_broaden_scheme)`
`elsi_set_mu_mp_order(handle, mu_mp_order)`
`elsi_set_mu_broaden_width(handle, mu_broaden_width)`
`elsi_set_mu_tol(handle, mu_tol)`

| Argument | Data Type | Default | Explanation |
|--------------------------------|-------------|------------|---|
| <code>out_level</code> | integer | 0 | Output level of the ELSI interface. 0: no output. 1: standard ELSI output. 2: 1 + info from the solvers. 3: 2 + additional debug info. |
| <code>out_log</code> | integer | 0 | If not 0, a separate log file in JSON format will be written out. |
| <code>write_unit</code> | integer | 6 | The unit used in ELSI to write out information. |
| <code>unit_ovlp</code> | integer | 0 | If not 0, the overlap matrix will be treated as an identity (unit) matrix in ELSI and the solvers. See remark 1. |
| <code>zero_def</code> | real double | 10^{-15} | When converting a matrix from dense to sparse format, values below this threshold will be discarded. |
| <code>sing_check</code> | integer | 0 | If not 0, the singularity check of the overlap matrix will be performed. See remark 2. |
| <code>sing_tol</code> | real double | 10^{-5} | Eigenfunctions of the overlap matrix with eigenvalues smaller than this threshold will be removed to avoid ill-conditioning. See remark 1. |
| <code>sing_stop</code> | integer | 0 | If not 0, the code always stops if the overlap matrix is detected to be singular. See remark 1. |
| <code>mu_broaden_scheme</code> | integer | 0 | The broadening scheme employed to compute the occupation numbers and the Fermi level. 0: Gaussian. 1: Fermi-Dirac. 2: Methfessel-Paxton. 4: Marzari-Vanderbilt. |
| <code>mu_mp_order</code> | integer | 0 | The order of the Methfessel-Paxton broadening scheme. No effect if Methfessel-Paxton is not the chosen broadening scheme. |
| <code>mu_broaden_width</code> | real double | 0.01 | The broadening width employed to compute the occupation numbers and the Fermi level. See remark 3. |
| <code>mu_tol</code> | real double | 10^{-13} | The convergence tolerance (in terms of the absolute error in electron count) of the bisection algorithm employed to compute the occupation numbers and the Fermi level. |

Remarks

- 1) If the overlap matrix is set to be an identity matrix, all settings related to the singularity (ill-conditioning) check take no effect. The `ovlp` argument passed into `elsi_ev_real`, `elsi_ev_complex`, `elsi_ev_real_sparse`, `elsi_ev_complex_sparse`, `elsi_dm_real`, `elsi_dm_complex`, `elsi_dm_real_sparse`, and `elsi_dm_complex_sparse` will not be referenced.
- 2) If the singularity check is not disabled, in the first iteration of each SCF cycle, possible singularity of the overlap matrix is checked by computing all its eigenvalues. If there is any eigenvalue smaller than `sing_tol`, the matrix is considered to be singular.
- 3) In all supported broadening schemes, there is a term $(\epsilon - E_F)/W$ in the distribution function, where ϵ is the energy of an eigenstate, and E_F is the Fermi level. The `broadening_width` parameter should be set to W , in the unit of ϵ and E_F .

3.5.2 Customizing the ELPA Solver

`elsi_set_elpa_solver(handle, elpa_solver)`

`elsi_set_elpa_n_single(handle, elpa_n_single)`

`elsi_set_elpa_gpu(handle, elpa_gpu)`

`elsi_set_elpa_gpu_kernels(handle, elpa_gpu_kernels)`

`elsi_set_elpa_autotune(handle, elpa_autotune)`

| Argument | Data Type | Default | Explanation |
|-------------------------------|-----------|---------|---|
| <code>elpa_solver</code> | integer | 2 | 1: ELPA 1-stage solver. 2: ELPA 2-stage solver. The latter is usually faster and more scalable. |
| <code>elpa_n_single</code> | integer | 0 | Number of SCF steps using single precision ELPA to solve standard eigenproblems. See remark 1. |
| <code>elpa_gpu</code> | integer | 0 | If not 0, try to enable GPU-acceleration in ELPA. See remark 2. |
| <code>elpa_gpu_kernels</code> | integer | 0 | If not 0, try to enable GPU-acceleration and GPU kernels in ELPA. See remark 2. |
| <code>elpa_autotune</code> | integer | 1 | If not 0, try to enable auto-tuning of runtime parameters in ELPA. See remark 3. |

Remarks

- 1) `elpa_n_single`: If single precision arithmetic is available in an externally compiled ELPA library, it may be enabled by setting `elpa_n_single` to a positive integer, then the standard eigenproblems in the first `elpa_n_single` SCF steps will be solved with single precision. The transformations between generalized eigenproblem and the standard form are always performed with double precision. Although this keyword accelerates the solution of standard eigenproblems, the overall SCF convergence may be slower, depending on the physical system and the SCF settings used in the electronic structure code. This keyword is ignored if single precision calculations are not available, which is the case if the internal version of ELPA is used, or if an external ELPA has not been compiled with single precision support.
- 2) `elpa_gpu` and `elpa_gpu_kernels`: If GPU-acceleration is available in an externally compiled ELPA library, it may be enabled by setting `elpa_gpu` to a non-zero integer. Note that by setting `elpa_gpu`, the GPU kernels for eigenvector back-transformation will not be used. To enable the GPU kernels, `elpa_gpu_kernels` should be set to a non-zero value. These two keywords are ignored if GPU-acceleration is not available, which is the case if the internal version of ELPA is used, or if an external ELPA has not been compiled with GPU support.
- 3) `elpa_autotune`: If auto-tuning of runtime parameters is available in an externally compiled ELPA library, it may be enabled by setting `elpa_autotune` to a nonzero integer. This keyword is ignored if auto-tuning is not available, which is the case if the internal version of ELPA is used.

3.5.3 Customizing the libOMM Solver

`elsi_set_omm_flavor(handle, omm_flavor)`

`elsi_set_omm_n_elpa(handle, omm_n_elpa)`

`elsi_set_omm_tol(handle, omm_tol)`

| Argument | Data Type | Default | Explanation |
|-------------------------|-------------|------------|--|
| <code>omm_flavor</code> | integer | 0 | Method to perform OMM minimization. See remark 1. |
| <code>omm_n_elpa</code> | integer | 6 | Number of SCF steps using ELPA. See remark 2. |
| <code>omm_tol</code> | real double | 10^{-12} | Convergence tolerance of orbital minimization. See remark 3. |

Remarks

1) `omm_flavor`: Allowed choices are 0 for a basic minimization of a generalized eigenproblem and 2 for a Cholesky factorization of the overlap matrix transforming the generalized eigenproblem to the standard form. Usually 2 (Cholesky) leads to a faster convergence of the OMM energy functional minimization, at the price of transforming the eigenproblem. When using sufficiently many steps of ELPA to stabilize the SCF cycle, 0 (basic) is probably a better choice to finish the remaining SCF cycle. See also remark 2 below.

2) `omm_n_elpa`: It has been demonstrated that OMM is optimal at later stages of an SCF cycle where the electronic structure is closer to its expected local minimum, requiring only one CG iteration to converge the minimization of the OMM energy functional. Accordingly, it is recommended to use ELPA initially, then switching to libOMM after `omm_n_elpa` SCF steps.

3) `omm_tol`: A large minimization tolerance of course leads to a faster convergence, however unavoidably with a lower accuracy. `omm_tol` should be tested and chosen to balance the desired accuracy and computation time of the calling code.

3.5.4 Customizing the PEXSI Solver

`elsi_set_pexsi_n_pole(handle, pexsi_n_pole)`

`elsi_set_pexsi_n_mu(handle, pexsi_n_mu)`

`elsi_set_pexsi_np_per_pole(handle, pexsi_np_per_pole)`

`elsi_set_pexsi_np_symbo(handle, pexsi_np_symbo)`

`elsi_set_pexsi_temp(handle, pexsi_temp)`

`elsi_set_pexsi_gap(handle, pexsi_gap)`

`elsi_set_pexsi_delta_e(handle, pexsi_delta_e)`

`elsi_set_pexsi_mu_min(handle, pexsi_mu_min)`

`elsi_set_pexsi_mu_max(handle, pexsi_mu_max)`

`elsi_set_pexsi_inertia_tol(handle, pexsi_inertia_tol)`

| Argument | Data Type | Default | Explanation |
|--------------------------------|-------------|---------|---|
| <code>pexsi_n_pole</code> | integer | 20 | Number of poles used by PEXSI. See remark 1. |
| <code>pexsi_n_mu</code> | integer | 2 | Number of mu points used by PEXSI. See remark 1. |
| <code>pexsi_np_per_pole</code> | integer | - | Number of MPI tasks assigned to each mu point. See remark 2. |
| <code>pexsi_np_symbo</code> | integer | 1 | Number of MPI tasks for symbolic factorization. See remark 3. |
| <code>pexsi_temp</code> | real double | 0.002 | Temperature. See remark 4. |
| <code>pexsi_gap</code> | real double | 0.0 | Spectral gap. See remark 5. |
| <code>pexsi_delta_e</code> | real double | 10.0 | Spectral radius. See remark 6. |
| <code>pexsi_mu_min</code> | real double | -10.0 | Minimum value of mu. See remark 7. |
| <code>pexsi_mu_max</code> | real double | 10.0 | Maximum value of mu. See remark 7. |
| <code>pexsi_inertia_tol</code> | real double | 0.05 | Stopping criterion of inertia counting. See remark 7. |

Remarks

1) In PEXSI, 20 poles are usually sufficient to get an accuracy that is comparable with the result obtained from diagonalization. The chemical potential is determined by performing Fermi operator expansion at several chemical potential values (referred to as “points” by PEXSI developers) in an SCF step, then interpolating the results at all points to the final answer. The `pexsi.n.mu` parameter controls the number of chemical potential “points” to be evaluated. 2 points followed by a simple linear interpolation often yield reasonable results.

In short, we recommend `pexsi.n.pole` = 20 and `pexsi.n.mu` = 2.

2) `pexsi.np.per.pole`: PEXSI has, by construction, a 3-level parallelism: the 1st level independently handles all the poles in parallel; within each pole, the 2nd level evaluates the Fermi operator at all the chemical potential points in parallel; finally, within each point, parallel selected inversion is performed as the 3rd level. The value of `pexsi.np.per.pole` is the number of MPI tasks assigned to a single chemical potential point, for the parallel selected inversion at that point. Ideally, the total number of MPI tasks should be `pexsi.np.per.pole` \times `pexsi.n.mu` \times `pexsi.n.pole`, i.e., all the three levels of parallelism are fully exploited. In case that this is not feasible, PEXSI can also process the poles in serial, whereas all the chemical potential points must be evaluated simultaneously. The user should make sure that the total number of MPI tasks is divisible by the product of the number of MPI tasks per pole and the number of points. The code will stop if this requirement is not fulfilled.

When using the `BLACS_DENSE` or `SIESTA_CSC` matrix formats, `pexsi.np.per.pole` is automatically determined to balance the three levels of parallelism in PEXSI. Input and output matrices should be distributed across all MPI tasks in either a 2D block-cyclic distribution (`BLACS_DENSE`) or a 1D block-cyclic distribution (`SIESTA_CSC`).

Note that when using the `PEXSI_CSC` matrix format together with the PEXSI solver, input and output matrices should be distributed among the first `pexsi.np.per.pole` MPI tasks (not all the MPI tasks) in a 1D block distribution. The block size of the distribution must be $\text{floor}(\text{N.basis}/\text{N.procs})$, where $\text{floor}(x)$ is the greatest integer less than or equal to x , `N.basis` and `N.procs` are the number of basis functions and the number of MPI tasks, respectively.

when using the `PEXSI_CSC` matrix format with the ELPA, libOMM, or SLEPc-SIPs solver, input and output matrices should be distributed across all the MPI tasks in a 1D block distribution. Again, the block size of the distribution must be $\text{floor}(\text{N.basis}/\text{N.procs})$.

3) `pexsi.np.symbo`: Unless there is a memory bottleneck, using 1 MPI task for matrix reordering and symbolic factorization is favorable. When running in serial, the matrix reordering in PT-SCOTCH or ParMETIS introduces a minimal number of “fill-ins” to the factorized matrices. Using more MPI tasks introduces more fill-ins. As the matrix reordering and symbolic factorization are performed only once per SCF cycle (with a fixed overlap matrix), using 1 MPI task should not affect the overall timing too much. On the other hand, more fill-ins lead to slower numerical factorization in every SCF step. In addition, the number of MPI tasks used for matrix reordering and symbolic factorization cannot be too large. Otherwise, the symbolic factorization may fail. Therefore, the default number of MPI tasks for symbolic factorization is 1. It is worth testing and increasing this number for large-scale calculations.

4) `pexsi.temp`: This value corresponds to the $1/k_B T$ term (not T) in the Fermi-Dirac distribution function.

5) `pexsi.gap`: The PEXSI method does not require a gap. If an estimate of the gap is unavailable, the default value usually works.

6) `pexsi.delta.e`: This is the spectral width of the eigensystem, i.e., the difference between the largest and smallest eigenvalues. Use the default value if no access to a better estimate.

7) The chemical potential determination in PEXSI relies on inertia counting to narrow down the chemical potential searching interval in the first few SCF steps. The `pexsi.inertia.tol` parameter controls the stopping criterion of the inertia counting procedure. With a small interval obtained from the inertia counting step, PEXSI then selects a number of points in this interval to perform Fermi operator calculations, based on which a final chemical potential will be determined. The trick of this algorithm is that the chemical potential interval of the current SCF step can be used as a descent guess in the next SCF step. Therefore, the mechanism to choose input values for `pexsi.mu.min` and `pexsi.mu.max` is two-fold. For the first SCF iteration, they should be set to safe values that guarantee the true chemical potential lies

in this interval. Then, for the n^{th} SCF step, `pexsi_mu_min` should be set to $(\mu_{\text{min}}^{n-1} + \Delta V_{\text{min}})$, `pexsi_mu_max` should be set to $(\mu_{\text{max}}^{n-1} + \Delta V_{\text{max}})$. Here, μ_{min}^{n-1} and μ_{max}^{n-1} are the lower bound and the upper bound of the chemical potential that are determined by PEXSI in the $(n-1)^{\text{th}}$ SCF step. They can be retrieved by calling `elsi_get_pexsi_mu_min` and `elsi_get_pexsi_mu_max`, respectively (see 3.6.2. Suppose the effective potential (Hartree potential, exchange-correlation potential, and external potential) is stored in an array V , whose dimension is the number of grid points. From one SCF iteration to the next, ΔV denotes the potential change, and ΔV_{min} and ΔV_{max} are the minimum and maximum values in the array ΔV , respectively. The whole process is summarized in the following pseudo-code.

```

mu_min = -10.0
mu_max = 10.0
ΔV_min = 0.0
ΔV_max = 0.0

while SCF not converged do
    Update Hamiltonian

    elsi_set_pexsi_mu_min(elsi_h, mu_min + ΔV_min)
    elsi_set_pexsi_mu_max(elsi_h, mu_max + ΔV_max)

    elsi_dm_{real|complex}(elsi_h, ham, ovlp, dm, bs_energy)

    elsi_get_pexsi_mu_min(elsi_h, mu_min)
    elsi_get_pexsi_mu_max(elsi_h, mu_max)

    Update electron density
    Update potential

    ΔV_min = minval(V_new - V_old)
    ΔV_max = maxval(V_new - V_old)

    Check SCF convergence
end

```

3.5.5 Customizing the SLEPc-SIPs Solver

```

elsi_set_sips_interval(handle, sips_lower, sips_upper)

elsi_set_sips_n_elpa(handle, sips_n_elpa)

elsi_set_sips_n_slice(handle, sips_n_slice)

```

| Argument | Data Type | Default | Explanation |
|---------------------------|-------------|---------|---|
| <code>sips_lower</code> | real double | -2.0 | Lower bound of eigenspectrum. See remark 1. |
| <code>sips_upper</code> | real double | 2.0 | Upper bound of eigenspectrum. See remark 1. |
| <code>sips_n_elpa</code> | integer | 0 | Number of SCF steps using ELPA. See remark 2. |
| <code>sips_n_slice</code> | integer | 1 | Number of slices. See remark 3. |

Remarks

1) `sips_lower` and `sips_upper`: SLEPc-SIPs relies on some inertia counting steps to estimate the lower and upper bounds of the spectrum. Only eigenvalues within this interval, and their associated eigenvectors, will be solved. The inertia-counting-based eigenvalue searching starts from the interval determined by `sips_lower` and `sips_upper`. Depending on the results of inertia counting, this interval may expand or shrink to make sure that the 1st to the `n.state`th eigenvalues are all within this interval. If a good estimate of the lower and upper bounds of the eigenspectrum is available, it should be set by `elsi_set_sips_interval`.

2) `sips_n_elpa`: The performance of SLEPc-SIPs mainly depends on the load balance across slices. Optimal performance is expected if the desired eigenvalues are evenly distributed across slices. In an SCF calculation, eigenvalues obtained in the current SCF step can be used as an approximated distribution of eigenvalues in the next SCF step. This approximation should become better as the SCF cycle approaches its convergence. On the other hand, at the beginning of an SCF cycle, the load balance is only coarsely checked by inertia calculations. Using the direct eigensolver ELPA in the first `sips_n_elpa` SCF steps can circumvent the load imbalance of spectrum slicing in the initial SCF steps.

3) `sips_n_slice`: SLEPc-SIPs partitions the eigenspectrum into slices and solves the slices in parallel. The `sips_n_slice` parameter controls the number of slices to use in SLEPc-SIPs. The default value, 1, should always work, but by no means leads to the optimal performance of the solver. There are some general rules to set this parameter. Firstly, as a requirement of the SLEPc library, the total number of MPI tasks must be divisible by `sips_n_slice`. Secondly, setting `sips_n_slice` to be equal to the number of computing nodes (not MPI tasks) usually yields better performance, as the communication between nodes is minimized in this case. The optimal value of `sips_n_slice` depends on the actual problem as well as the computing hardware.

3.5.6 Customizing the NTPoly Solver

`elsi_set_ntpoly_method`(handle, ntpoly_method)

`elsi_set_ntpoly_filter`(handle, ntpoly_filter)

`elsi_set_ntpoly_tol`(handle, ntpoly_tol)

| Argument | Data Type | Default | Explanation |
|----------------------------|-------------|------------|--|
| <code>ntpoly_method</code> | integer | 0 | Method to perform density matrix purification. See remark 1. |
| <code>ntpoly_filter</code> | real double | 10^{-15} | When performing sparse matrix multiplications, values below this filter will be discarded. See remark 2. |
| <code>ntpoly_tol</code> | real double | 10^{-8} | Convergence tolerance of purification. See remark 2. |

Remarks

1) `ntpoly_method`: Allowed choices are 0 for the canonical purification, 1 for the trace correcting purification, 2 for the 4th order trace resetting purification, and 3 for the generalized hole-particle canonical purification.

2) `ntpoly_filter` and `ntpoly_tol` control the accuracy and computational cost of the density matrix purification methods. Tight choices of `ntpoly_filter` and `ntpoly_tol`, e.g. the default values here, lead to highly accurate results that are comparable to the results obtained from diagonalization. However, linear scaling can only be achieved with a relatively large `ntpoly_filter` such as 10^{-6} . Correspondingly, `ntpoly_tol` may be set to 10^{-3} . Note that the purification may not converge if `ntpoly_filter` is too large relative to `ntpoly_tol`. Setting `ntpoly_filter` to be $\leq 10^{-3} \times \text{ntpoly_tol}$ is safe in most cases.

3.6 Getting Additional Results from ELSI

In 3.3 and 3.4, the interfaces to compute and return the eigensolutions and the density matrices have been introduced. Internally, ELSI and the solvers perform additional calculations whose results may only be useful at a certain stage of an SCF calculation. One example is the energy-weighted density matrix that is employed to evaluate the Pulay forces during a geometry optimization calculation. The subroutines introduced in the following subsections are used to retrieve such additional results from ELSI.

3.6.1 Getting Results from the ELSI Interface

In all the subroutines listed below, the first argument (input and output) is an `elsi_handle`. The second argument (output) of each subroutine is the name of parameter to get.

`elsi_get_initialized`(handle, handle_init)

[elsi_get_version](#)(handle, major, minor, patch)
[elsi_get_datestamp](#)(handle, date_stamp)
[elsi_get_n_sing](#)(handle, n_sing)
[elsi_get_mu](#)(handle, mu)
[elsi_get_entropy](#)(handle, ts)
[elsi_get_edm_real](#)(handle, edm_real)
[elsi_get_edm_complex](#)(handle, edm_complex)
[elsi_get_edm_real_sparse](#)(handle, edm_real_sparse)
[elsi_get_edm_complex_sparse](#)(handle, edm_complex_sparse)

| Argument | Data Type | Explanation |
|------------------------------------|------------------------------|---|
| handle_init | integer | 0 if the ELSI handle has not been initialized; 1 if initialized. |
| major | integer | Major version number. |
| minor | integer | Minor version number. |
| patch | integer | Patch level. |
| date_stamp | integer | Date stamp of ELSI (yyyymmdd). |
| n_sing | integer | Number of eigenvalues of the overlap matrix that are smaller than the singularity tolerance. See 3.5.1. |
| mu | real double | Chemical potential. See remark 1. |
| ts | real double | Entropy. See remark 1. |
| edm_real | real double, rank-2 array | Real energy-weighted density matrix in 2D block-cyclic dense format. See remark 2. |
| edm_complex | complex double, rank-2 array | Complex energy-weighted density matrix in 2D block-cyclic dense format. See remark 2. |
| edm_real_sparse | real double, rank-1 array | Non-zero values of the real density matrix in 1D block CSC format. See remark 2. |
| edm_complex_sparse | complex double, rank-1 array | Non-zero values of the complex density matrix in 1D block CSC format. See remark 2. |

Remarks

1) In ELSI, the chemical potential will only be available if one of the density matrix solver interfaces has been called, with ELPA, PEXSI, or NTPoly being the chosen solver. The chemical potential can be retrieved by calling [elsi_get_mu](#). The entropy will only be available if one of the density matrix solver interfaces has been called with ELPA being the chosen solver. The user should avoid calling the subroutine when the chemical potential or the entropy is not ready.

2) In general, the energy-weighted density matrix is only needed in a late stage of an SCF cycle to evaluate forces. It is, therefore, not calculated when any of the density matrix solver interface is called. When the energy-weighted density matrix is actually needed, it can be requested by calling the [elsi_get_edm](#) subroutines. Note that these subroutines all have the requirement that the corresponding [elsi_dm](#) subroutine must have been invoked. For instance, [elsi_get_edm_real_sparse](#) only makes sense if [elsi_dm_real_sparse](#) has been successfully executed.

3.6.2 Getting Results from the PEXSI Solver

[elsi_get_pexsi_mu_min](#)(handle, pexsi_mu_min)
[elsi_get_pexsi_mu_max](#)(handle, pexsi_mu_max)

| Argument | Data Type | Explanation |
|------------------------------|-------------|------------------------------------|
| pexsi_mu_min | real double | Minimum value of mu. See remark 1. |
| pexsi_mu_max | real double | Maximum value of mu. See remark 1. |

Remarks

- 1) Please refer to the 7th remark in 3.5.4 for the chemical potential determination algorithm in PEXSI and ELSI.

3.7 Parallel Matrix I/O

To test the solvers in ELSI, it is convenient to use matrices generated from actual electronic structure calculations. There exist a number of libraries invented for high-performance parallel I/O that are particularly capable of reading and writing a large amount of data with hierarchical structures and complex metadata. However, the I/O task in ELSI is very simple in terms of the complexity of the data to manipulate. The data structure is simply arrays that represent matrices, with a few integers to define the dimension of the matrices. In order to circumvent the unavoidable development and performance overhead associated with a high level I/O library, the parallel I/O functionality defined in the MPI standard is directly used to read and write matrices in ELSI.

When ELSI runs in parallel with multiple MPI tasks, the matrices are distributed across tasks. The choice of writing the distributed matrices into N_{procs} separate files, where N_{procs} is the number of MPI tasks, is not promising due to the difficulty of managing and post-processing a large number of files, especially with a different number of MPI tasks. The implementation of matrix I/O in ELSI adopts collective MPI I/O routines to write data to (read data from) a single binary file, as if the data was gathered onto a single MPI task then written to one file (read from one file by one MPI task then scattered to all tasks). The optimal I/O performance, both with MPI I/O and in general, is often obtained by making large and contiguous requests to access the file system, rather than small, non-contiguous, or random requests. Therefore, before being written to file, matrices are always redistributed to a 1D block distribution. This guarantees that each MPI task writes a contiguous trunk of data to a contiguous piece of file. Similarly, matrices read from file are in a 1D block distribution, and can be redistributed automatically if needed.

A matrix is always stored in the CSC format in an ELSI matrix file. A dense matrix is automatically converted to the CSC format before writing to file, and can be converted back after reading from file.

Next, we present the API for parallel matrix I/O.

3.7.1 Setting Up Matrix I/O

An `elsi_rw_handle` must be initialized via the `elsi_init_rw` subroutine before any other matrix I/O subroutine may be called. This `elsi_rw_handle` is subsequently passed to all other matrix I/O subroutine calls.

```
elsi_init_rw(handle, task, parallel_mode, n_basis, n_electron)
```

| Argument | Data Type | in/out | Explanation |
|----------------------------|-----------------------------------|--------|---|
| <code>handle</code> | <code>type(elsi_rw_handle)</code> | out | Handle to matrix I/O instance. |
| <code>task</code> | integer | in | Matrix I/O task to perform. Accepted values are: 0 (READ_MATRIX) and 1(WRITE_MATRIX). |
| <code>parallel_mode</code> | integer | in | Parallelization mode. The only accepted value is 1 (MULTIPROC) for now. |
| <code>n_electron</code> | real double | in | Number of electrons. See remark 1. |
| <code>n_basis</code> | integer | in | Number of basis functions, i.e. global size of matrix. |

Remarks

- 1) `n_electron`: Many matrices written out with ELSI matrix I/O are from real electronic structure calculations. Having the information of the number of electrons available makes the matrix file useful for testing density matrix solvers such as PEXSI. Therefore, it is recommended to set the correct number of electrons when initializing an matrix I/O handle, although setting it to an arbitrary number will not affect the matrix I/O operation.

2) `n.basis`: This can be set to an arbitrary value if `task` is 0 (`READ_MATRIX`). Its value will be read from file when calling `elsi_read_mat_dim` or `elsi_read_mat_dim_sparse`.

The MPI communicator which encloses the MPI tasks to perform the matrix I/O operation needs to be passed into ELSI via the `elsi_set_rw_mpi` subroutine.

`elsi_set_rw_mpi`(handle, mpi_comm)

| Argument | Data Type | in/out | Explanation |
|-----------------------|-----------------------------------|--------|--------------------------------|
| <code>handle</code> | <code>type(elsi_rw_handle)</code> | inout | Handle to matrix I/O instance. |
| <code>mpi_comm</code> | integer | in | MPI communicator. |

When reading or writing a dense matrix, BLACS parameters are passed into ELSI via the `elsi_set_rw_blacs` subroutine.

`elsi_set_rw_blacs`(handle, blacs_ctxt, block_size)

| Argument | Data Type | in/out | Explanation |
|-------------------------|-----------------------------------|--------|--|
| <code>handle</code> | <code>type(elsi_rw_handle)</code> | inout | Handle to matrix I/O instance. |
| <code>blacs_ctxt</code> | integer | in | BLACS context. |
| <code>block_size</code> | integer | in | Block size of the 2D block-cyclic distribution, specifying both row and column directions. |

When writing a sparse matrix, its dimensions are passed into ELSI via the `elsi_set_rw_csc` subroutine. The only sparse matrix format currently supported by ELSI matrix I/O is the `PEXSL_CSC` format. When reading a sparse matrix, there is no need to call this subroutine. The relevant parameters will be read from file when calling `elsi_read_mat_dim` or `elsi_read_mat_dim_sparse`.

`elsi_set_rw_csc`(handle, global_nnz, local_nnz, local_col)

| Argument | Data Type | in/out | Explanation |
|-------------------------|-----------------------------------|--------|---------------------------------|
| <code>handle</code> | <code>type(elsi_rw_handle)</code> | inout | Handle to matrix I/O instance. |
| <code>global_nnz</code> | integer | in | Global number of non-zeros. |
| <code>local_nnz</code> | integer | in | Local number of non-zeros. |
| <code>local_col</code> | integer | in | Local number of matrix columns. |

When a matrix I/O instance is no longer needed, its associated handle should be cleaned up by calling `elsi_finalize_rw`.

`elsi_finalize_rw`(handle)

| Argument | Data Type | in/out | Explanation |
|---------------------|-----------------------------------|--------|--------------------------------|
| <code>handle</code> | <code>type(elsi_rw_handle)</code> | inout | Handle to matrix I/O instance. |

3.7.2 Writing Matrices

The following two subroutines write a dense matrix to file. Before writing a dense matrix, MPI and BLACS should be set up properly using `elsi_set_rw_mpi` and `elsi_set_rw_blacs`.

`elsi_write_mat_real`(handle, filename, mat)

| Argument | Data Type | in/out | Explanation |
|-----------------------|-----------------------------------|--------|---|
| <code>handle</code> | <code>type(elsi_rw_handle)</code> | in | Handle to matrix I/O instance. |
| <code>filename</code> | string | in | Name of file to write. |
| <code>mat</code> | real double, rank-2 array | in | Local matrix in 2D block-cyclic dense format. |

[elsi_write_mat_complex](#)(handle, filename, mat)

| Argument | Data Type | in/out | Explanation |
|--------------------------|------------------------------|--------|---|
| handle | type(elsi_rw_handle) | in | Handle to matrix I/O instance. |
| filename | string | in | Name of file to write. |
| mat | complex double, rank-2 array | in | Local matrix in 2D block-cyclic dense format. |

The following two subroutines write a sparse matrix to file. Before writing a sparse matrix, MPI and CSC matrix format should be set up properly using [elsi_set_rw_mpi](#) and [elsi_set_rw_csc](#).

[elsi_write_mat_real_sparse](#)(handle, filename, row_idx, col_ptr, mat)

| Argument | Data Type | in/out | Explanation |
|--------------------------|---------------------------|--------|---|
| handle | type(elsi_rw_handle) | in | Handle to matrix I/O instance. |
| filename | string | in | Name of file to write. |
| row_idx | integer, rank-1 array | in | Local row index array. |
| col_ptr | integer, rank-1 array | in | Local column pointer array. |
| mat | real double, rank-1 array | in | Local non-zero values in 1D block CSC format. |

[elsi_write_mat_complex_sparse](#)(handle, filename, row_idx, col_ptr, mat)

| Argument | Data Type | in/out | Explanation |
|--------------------------|------------------------------|--------|---|
| handle | type(elsi_rw_handle) | in | Handle to matrix I/O instance. |
| filename | string | in | Name of file to write. |
| row_idx | integer, rank-1 array | in | Local row index array. |
| col_ptr | integer, rank-1 array | in | Local column pointer array. |
| mat | complex double, rank-1 array | in | Local non-zero values in 1D block CSC format. |

When writing a dense matrix to file, values smaller than a predefined threshold will be discarded. The default value of this threshold is 10^{-15} . It can be overridden via [elsi_set_rw_zero_def](#).

[elsi_set_rw_zero_def](#)(handle, zero_def)

| Argument | Data Type | in/out | Explanation |
|--------------------------|----------------------|--------|---|
| handle | type(elsi_rw_handle) | inout | Handle to matrix I/O instance. |
| zero_def | real double | in | When writing a dense matrix to file, values below this threshold will be discarded. |

An array of eight user-defined integers can be optionally set up via [elsi_set_rw_header](#). This array will be attached to the matrix file written out by the above subroutines. When reading a matrix file, this array may be retrieved via [elsi_get_rw_header](#).

[elsi_set_rw_header](#)(handle, header)

| Argument | Data Type | in/out | Explanation |
|------------------------|-----------------------|--------|--------------------------------|
| handle | type(elsi_rw_handle) | inout | Handle to matrix I/O instance. |
| header | integer, rank-1 array | in | An array of eight integers. |

3.7.3 Reading Matrices

The following subroutines read a dense or sparse matrix from file. While writing a matrix to file can be done in one step, it is easier to read a matrix from file in two steps, i.e., first read the dimension of the matrix and allocate memory accordingly, then read the actual data of the matrix.

The following three subroutines read a dense matrix from file. Before reading a dense matrix, MPI and BLACS should be set up properly using [elsi_set_rw_mpi](#) and [elsi_set_rw_blacs](#). [elsi_read_mat_dim](#) is used to read the dimension of a matrix, including the number of electrons in the physical system (for testing purpose), the global size of the matrix, and the local size of the matrix. Memory needs to be allocated according to the return values of [local_row](#) and [local_col](#). Then [elsi_read_mat_real](#) or [elsi_read_mat_complex](#) may be called to read a real or complex matrix, respectively.

[elsi_read_mat_dim](#)(handle, filename, n_electron, n_basis, local_row, local_col)

| Argument | Data Type | in/out | Explanation |
|----------------------------|----------------------|--------|--|
| handle | type(elsi_rw_handle) | inout | Handle to matrix I/O instance. |
| filename | string | in | Name of file to read. |
| n_electron | real double | out | Number of electrons. |
| n_basis | integer | out | Number of basis functions, i.e. global size of matrix. |
| local_row | integer | out | Local number of matrix rows. |
| local_col | integer | out | Local number of matrix columns. |

[elsi_read_mat_real](#)(handle, filename, mat)

| Argument | Data Type | in/out | Explanation |
|--------------------------|---------------------------|--------|---|
| handle | type(elsi_rw_handle) | inout | Handle to matrix I/O instance. |
| filename | string | in | Name of file to read. |
| mat | real double, rank-2 array | out | Local matrix in 2D block-cyclic distribution. |

[elsi_read_mat_complex](#)(handle, filename, mat)

| Argument | Data Type | in/out | Explanation |
|--------------------------|------------------------------|--------|---|
| handle | type(elsi_rw_handle) | inout | Handle to matrix I/O instance. |
| filename | string | in | Name of file to read. |
| mat | complex double, rank-2 array | out | Local matrix in 2D block-cyclic distribution. |

The following three subroutines read a sparse matrix from file. Before reading a sparse matrix, MPI should be set up properly using [elsi_set_rw_mpi](#). [elsi_read_mat_dim_sparse](#) is used to read the dimension of a matrix, including the number of electrons in the physical system (for testing purpose), the global size of the matrix, and the local size of the matrix. Memory needs to be allocated according to the return values of [local_nnz](#) and [local_col](#). Then [elsi_read_mat_real_sparse](#) or [elsi_read_mat_complex_sparse](#) may be called to read a real or complex matrix, respectively.

[elsi_read_mat_dim_sparse](#)(handle, filename, n_electron, n_basis, global_nnz, local_nnz, local_col)

| Argument | Data Type | in/out | Explanation |
|----------------------------|----------------------|--------|--|
| handle | type(elsi_rw_handle) | inout | Handle to matrix I/O instance. |
| filename | string | in | Name of file to read. |
| n_electron | real double | out | Number of electrons. |
| n_basis | integer | out | Number of basis functions, i.e. global size of matrix. |
| global_nnz | integer | out | Global number of non-zeros. |
| local_nnz | integer | out | Local number of non-zeros. |
| local_col | integer | out | Local number of matrix columns. |

[elsi_read_mat_real_sparse](#)(handle, filename, row_idx, col_ptr, mat)

| Argument | Data Type | in/out | Explanation |
|--------------------------|---------------------------|--------|---|
| handle | type(elsi_rw_handle) | inout | Handle to matrix I/O instance. |
| filename | string | in | Name of file to read. |
| row_idx | integer, rank-1 array | out | Local row index array. |
| col_ptr | integer, rank-1 array | out | Local column pointer array. |
| mat | real double, rank-1 array | out | Local non-zero values in 1D block CSC format. |

`elsi_read_mat_complex_sparse(handle, filename, row_idx, col_ptr, mat)`

| Argument | Data Type | in/out | Explanation |
|-----------------------|-----------------------------------|--------|---|
| <code>handle</code> | <code>type(elsi_rw_handle)</code> | inout | Handle to matrix I/O instance. |
| <code>filename</code> | string | in | Name of file to read. |
| <code>row_idx</code> | integer, rank-1 array | out | Local row index array. |
| <code>col_ptr</code> | integer, rank-1 array | out | Local column pointer array. |
| <code>mat</code> | complex double, rank-1 array | out | Local non-zero values in 1D block CSC format. |

An array of eight user-defined integers can be optionally set up via `elsi_set_rw_header`. This array will be attached to the matrix file written out by the above subroutines. When reading a matrix file, this array may be retrieved via `elsi_get_rw_header`.

`elsi_get_rw_header(handle, header)`

| Argument | Data Type | in/out | Explanation |
|---------------------|-----------------------------------|--------|--------------------------------|
| <code>handle</code> | <code>type(elsi_rw_handle)</code> | inout | Handle to matrix I/O instance. |
| <code>header</code> | integer, rank-1 array | out | An array of eight integers. |

3.8 Demonstration Pseudo-Code

The typical workflow of ELSI within an electronic structure code is demonstrated by the following pseudo-code.

3.8.1 2D Block-Cyclic Distributed Dense Matrix + ELSI Eigensolver Interface

SCF initialize

```
elsi_init(elsi_h, ELPA, MULTIPROC, BLACS_DENSE, n_basis, n_electron, n_state)
elsi_set_mpi(elsi_h, mpi_comm)
elsi_set_blacs(elsi_h, blacs_ctxt, block_size)
```

while *SCF not converged* **do**

Update Hamiltonian

```
    elsi_ev_{real|complex}(elsi_h, ham, ovlp, eval, evec)
```

Update electron density

Check SCF convergence

end

```
elsi_finalize(elsi_h)
```

3.8.2 1D Block Distributed CSC Sparse Matrix + ELSI Eigensolver Interface

SCF initialization

```
elsi_init(elsi_h, ELPA, MULTIPROC, PEXSI_CSC, n_basis, n_electron, n_state)
elsi_set_mpi(elsi_h, mpi_comm)
elsi_set_blacs(elsi_h, blacs_ctxt, block_size)
elsi_set_csc(elsi_h, global_nnz, local_nnz, local_col, row_idx, col_ptr)

while SCF not converged do
    Update Hamiltonian

    elsi_ev_{real|complex}_sparse(elsi_h, ham, ovlp, eval, evec)

    Update electron density
    Check SCF convergence
end

elsi_finalize(elsi_h)
```

Remarks

- 1) The calculated eigenvectors are returned in the BLACS_DENSE format, which is required to be properly set up.

3.8.3 1D Block-Cyclic Distributed CSC Sparse Matrix + ELSI Eigensolver Interface

SCF initialization

```
elsi_init(elsi_h, ELPA, MULTIPROC, SIESTA_CSC, n_basis, n_electron, n_state)
elsi_set_mpi(elsi_h, mpi_comm)
elsi_set_blacs(elsi_h, blacs_ctxt, block_size)
elsi_set_csc(elsi_h, global_nnz, local_nnz, local_col, row_idx, col_ptr)
elsi_set_csc_blk(elsi_h, block_size)

while SCF not converged do
    Update Hamiltonian

    elsi_ev_{real|complex}_sparse(elsi_h, ham, ovlp, eval, evec)

    Update electron density
    Check SCF convergence
end

elsi_finalize(elsi_h)
```

Remarks

- 1) The calculated eigenvectors are returned in the BLACS_DENSE format, which is required to be properly set up.

3.8.4 2D Block-Cyclic Distributed Dense Matrix + ELSI Density Matrix Interface

SCF initialization

```
elsi_init(elsi_h, LIBOMM, MULTIPROC, BLACS_DENSE, n.basis, n.electron, n.state)
elsi_set_mpi(elsi_h, mpi_comm)
elsi_set_blacs(elsi_h, blacs_ctxt, block_size)

while SCF not converged do
    Update Hamiltonian

    elsi_dm_{real|complex}(elsi_h, ham, ovlp, dm, bs.energy)
    elsi_get_edm_{real|complex}(elsi_h, edm)

    Update electron density
    Check SCF convergence
end

elsi_finalize(elsi_h)
```

3.8.5 1D Block Distributed CSC Sparse Matrix + ELSI Density Matrix Interface

SCF initialization

```
elsi_init(elsi_h, PEXSI, parallel_mode, PEXSI_CSC, n.basis, n.electron, n.state)
elsi_set_mpi(elsi_h, mpi_comm)
elsi_set_csc(elsi_h, global_nnz, local_nnz, local_col, row_idx, col_ptr)

while SCF not converged do
    Update Hamiltonian

    elsi_dm_{real|complex}_sparse(elsi_h, ham, ovlp, dm, bs.energy)
    elsi_get_edm_{real|complex}_sparse(elsi_h, edm)

    Update electron density
    Check SCF convergence
end

elsi_finalize(elsi_h)
```

Remarks

- 1) Refer to the 7th remark in 3.5.4 for the chemical potential determination algorithm in PEXSI.

3.8.6 1D Block-Cyclic Distributed CSC Sparse Matrix + ELSI Density Matrix Interface

SCF initialization

```
elsi_init(elsi_h, PEXSI, parallel_mode, SIESTA_CSC, n_basis, n_electron, n_state)
elsi_set_mpi(elsi_h, mpi_comm)
elsi_set_csc(elsi_h, global_nnz, local_nnz, local_col, row_idx, col_ptr)
elsi_set_csc_blk(elsi_h, block_size)

while SCF not converged do
    Update Hamiltonian

    elsi_dm_{real|complex}_sparse(elsi_h, ham, ovlp, dm, bs_energy)
    elsi_get_edm_{real|complex}_sparse(elsi_h, edm)

    Update electron density
    Check SCF convergence
end

elsi_finalize(elsi_h)
```

Remarks

- 1) Refer to the 7th remark in 3.5.4 for the chemical potential determination algorithm in PEXSI.

3.8.7 Multiple k -points Calculations

SCF initialization

```
elsi_init(elsi_h, ELPA, parallel_mode, BLACS_DENSE, n_basis, n_electron, n_state)
elsi_set_mpi(elsi_h, mpi_comm)
elsi_set_blacs(elsi_h, blacs_ctxt, block_size)

elsi_set_kpoint(elsi_h, n_kpt, i_kpt, weight)
elsi_set_mpi_global(elsi_h, mpi_comm_global)

while SCF not converged do
    Update Hamiltonian

    elsi_dm_{real|complex}(elsi_h, ham, ovlp, dm, bs_energy)
    elsi_get_edm_{real|complex}(elsi_h, edm)

    Update electron density
    Check SCF convergence
end

elsi_finalize(elsi_h)
```

Remarks

- 1) When there are multiple k -points, other than setting up the k -points and a global MPI communicator, there is no change in the way ELSI solver interfaces are called.

- 2) The electronic structure code needs to assemble the real-space density from the density matrices returned for the \mathbf{k} -points. The returned band structure energy, however, is already summed over all \mathbf{k} -points with respect to the weight of each \mathbf{k} -point. Refer to 3.2.4 for more information.
- 3) Calculations with two spin channels can be set up similarly.

3.9 C/C++ Interface

ELSI is written in Fortran. A C interface around the core Fortran code is provided, which can be called from a C or C++ program. Each C wrapper function corresponds to a Fortran subroutine, where we have prefixed the original Fortran subroutine name with `c_` for clarity and consistency. Argument lists are identical to the associated native Fortran subroutine. For the complete definition of the C interface, the user is encouraged to look at the `elsi.h` header file directly.

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