SPARC

Simulation Package for Ab-initio Real-space Calculations

User guide

Material Physics & Mechanics Group, Georgia Tech
PI: Phanish Suryanarayana
Contributors
Citation
Acknowledgements

Introduction

SPARC is an open-source software package for the accurate, efficient, and scalable solution of the Kohn-Sham density functional theory (DFT) problem. The main features of SPARC currently include

- Applicable to isolated systems such as molecules as well as extended systems such as crystals, surfaces, and wires.
- Local, semilocal, and nonlocal (including hybrid) exchange-correlation functionals.
- Standard ONCV pseudopotentials, including nonlinear core corrections.
- Calculation of ground state energy, atomic forces, and stress tensor.
- Structural relaxation and ab initio molecular dynamics (NVE, NVT, and NPT).
- Spin polarized and unpolarized calculations.
- Spin-orbit coupling.
- Dispersion interactions through DFT-D3, vdW-DF1, and vdW-DF2.

Introduction

- Symmetry-adaption for cyclic and/or helical symmetries.
- Linear-scaling Spectral Quadrature (SQ) method.
- Soft and transferable table of SPMS pseudopotentials.
- On-the-fly machine-learned force field (MLFF) molecular dynamics (MD) simulations.

Contributors

- Phanish Suryanarayana (PI)
 - Qimen Xu: Code infrastructure, SCF, Energy, Force, LDA
 - Xin Jing: Code infrastructure, Hybrid functionals, SOC
 - **Abhiraj Sharma**: Code infrastructure, k-points, PBE, Stress, Non-orthogonal, Relaxation, NLCC
 - Boqin Zhang: vdW-DF, DFT-D3, meta-GGA (SCAN)
 - Shashikant Kumar: Testing framework, NLCC, MLFF
 - Mostafa Faghih Shojaei: SPMS table of pseudopotentials
 - Sayan Bhowmik¹: Atom code, DFT+U (Dudarev)
 - Swarnava Ghosh: Preliminary development
 - Deepa Phanish: Initial development
- John E. Pask (co-PI)
- Edmond Chow (co-PI)
 - Hua Huang: Subspace eigensolver, DP
 - Lucas Erlandson: Subspace eigensolver, DP
- Andrew J. Medford (co-PI)

Contributors

- Benjamin Comer: Code testing, Initial testing framework
- Sushree Jagriti Sahoo: Code testing
- Tian Tian: Socket communication layer, code testing, Python API
- Lucas R Timmerman: Socket communication layer, code testing, Python API
- Sayan Bhowmik¹: Atom code, DFT+U (Dudarev)

¹co-advised

Citation

If you publish work using/regarding SPARC, please cite some of the following articles, particularly those that are most relevant to your work:

- General: https://doi.org/10.1016/j.simpa.2024.100649 (v2), https://doi.org/10.1016/j.softx.2021.100709 (v1), https://doi.org/10.1016/j.cpc.2016.09.020 (initial developments, isolated systems), https://doi.org/10.1016/j.cpc.2017.02.019 (initial developments, extended systems)
- Non-orthogonal systems: https://doi.org/10.1016/j.cplett.2018.04.018
- Linear solvers: https://doi.org/10.1016/j.cpc.2018.07.007, https://doi.org/10.1016/j.jcp.2015.11.018
- Stress tensor/pressure: https://doi.org/10.1063/1.5057355

Citation

- Atomic forces: https://doi.org/10.1016/j.cpc.2016.09.020 (isolated systems), https://doi.org/10.1016/j.cpc.2017.02.019 (extended systems)
- Mixing: https://doi.org/10.1016/j.cplett.2016.01.033, https://doi.org/10.1016/j.cplett.2015.06.029 (restarting), https://doi.org/10.1016/j.cplett.2019.136983 (preconditioning)
- SPMS pseudopotentials: https://doi.org/10.1016/j.cpc.2022.108594
- Cyclic and/or helical symmetry:
 https://doi.org/10.1103/PhysRevB.103.035101,
 https://doi.org/10.1103/PhysRevB.100.125143 (initial developments, extended systems),
 https://doi.org/10.1016/j.jmps.2016.08.007 (initial developments, isolated systems)

Citation

 O(N) Spectral Quadrature method: https://doi.org/10.1016/j.cpc.2015.11.005 (initial implementation), https://doi.org/10.1016/j.cplett.2013.08.035 (formulation), https://doi.org/10.1007/978-3-031-22340-2_12 (detailed mathematical formulation)

```
On-the-fly MLFF: https://doi.org/10.1063/5.0180541,
https://doi.org/10.1063/5.0204229 (SQ),
https://doi.org/10.48550/arXiv.2408.07554 (Cyclix),
https://doi.org/10.48550/arXiv.2407.15290 (internal energy)
```

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Prerequisite: C compiler, MPI.

There are several options to compile SPARC, depending on the available external libraries.

- Option 1: Compile with BLAS and LAPACK.
 - Step 1: Install/Load OpenBLAS/BLAS and LAPACK.
 - Step 2: Go to src/ directory, there is an available makefile.
 - Step 3 (optional): Edit makefile. If the BLAS library path and LAPACK library path are not in the search path, edit the BLASROOT and LAPACKROOT variables, and add them to LDFLAGS. If you are using BLAS instead of OpenBLAS, replace all -lopenblas flags with -lblas.
 - Step 4 (optional): To turn on DEBUG mode, set DEBUG_MODE to 1 in the makefile.
 - Step 5: Within the src/ directory, compile the code by \$ make clean; make

- Option 2 (default): Compile with MKL.
 - Step 1: Install/Load MKL.
 - Step 2: Go to src/ directory, there is an available makefile.
 - Step 3: Edit makefile. Set USE_MKL to 1 to enable compilation with MKL. If the MKL library path is not in the search path, edit the MKLROOT variable to manually set the MKL path.
 - Step 4 (optional): For the projection/subspace rotation step, to use SPARC routines for matrix data distribution rather than ScaLAPACK (through MKL), set USE_DP_SUBEIG to 1. We found on some machines this option is faster.
 - Step 5 (optional): To turn on DEBUG mode, set DEBUG_MODE to 1 in the makefile.
 - Step 6: Within the src/ directory, compile the code by \$ make clean; make

- Option 3: Compile with BLAS, LAPACK, and ScaLAPACK.
 - Step 1: Install/Load OpenBLAS/BLAS, LAPACK, and ScaLAPACK.
 - Step 2: Go to src/ directory, there is an available makefile.
 - Step 3: Edit makefile. Set USE_SCALAPACK to 1 to enable compilation with ScaLAPACK. If the BLAS library path, LAPACK library path, and/or ScaLAPACK library path are not in the search path, edit the BLASROOT, LAPACKROOT, and/or SCALAPACKROOT variables accordingly, and add them to LDFLAGS. If you are using BLAS instead of OpenBLAS, replace all -lopenblas flags with -lblas.
 - Step 4 (optional): For the projection/subspace rotation step, to use SPARC routines for matrix data distribution rather than ScaLAPACK, set USE_DP_SUBEIG to 1. We found on some machines this option is faster.
 - Step 5 (optional): To turn on DEBUG mode, set DEBUG_MODE to 1 in the makefile.
 - Step 6: Within the src/ directory, compile the code by \$ make clean; make

Installation - lib

Once compilation is done, a binary named sparc will be created in the lib/ directory.

- Option 4: Install pre-compiled sparc binaries distributed by conda-forge
 - Pre-compiled sparc package can be installed on x86_64 or aarch64 Linux platforms with anaconda or miniconda installed. The binary is compiled with OpenBLAS and OpenMPI using flags USE_MKL=0 USE_SCALAPACK=1 USE_FFTW=1.
 - Step 1 (optional): create a conda environment (e.g. sparc-env)
 - \$ conda create -n sparc-env
 - \$ conda activate sparc-env
 - Step 2: install conda package sparc-x
 - \$ conda install -c conda-forge sparc-x
 - \$ echo sparc binary is located at: \$(which sparc)
 - \$ echo .psp files installed at: \$SPARC_PSP_PATH
 - \$ echo SPARC doc files installed at: \$SPARC_DOC_PATH

Input files

The required input files to run a simulation with SPARC are

- ".inpt" file User options and parameters.
- ".ion" file Atomic information.

It is required that the ".inpt" and ".ion" files are located in the same directory and share the same name. A detailed description of the input options is provided in this document. Examples of input files can be found in the directory SPARC/tests.

In addition, SPARC requires pseudopotential files of psp8 format which can be generated by D. R. Hamann's open-source pseudopotential code ONCVPSP. A large number of accurate and efficient pseudopotentials are already provided within the package. For access to more pseudopotentials, the user is referred to the SG15 ONCV potentials. Using the ONCVPSP input files included in the SG15 ONCV potentials, one can easily convert the SG15 ONCV potentials from upf format to psp8 format. Paths to the pseudopotential files are specified in the ".ion" file.

Execution

SPARC can be executed in parallel using the mpirun command. Sample PBS script files are available in "SPARC/tests" folder. It is required that the ".inpt" and ".ion" files are located in the same directory and share the same name. For example, to run a simulation with 8 processes with input files as "filename.inpt" and "filename.ion" in the root directory (SPARC/), use the following command:

\$ mpirun -np 8 ./lib/sparc -name filename

As an example, one can run one of the tests located in 'SPARC/tests/'. First go to 'SPARC/tests/Example_tests/' directory:

\$ cd tests/Example_tests/

There are a few input files available. Run a DC silicon system by

Execution

\$ mpirun -np 24 ../../lib/sparc -name Si8_kpt

The result is printed to output file "Si8_kpt.out", located in the same directory as the input files. If the file "Si8_kpt.out" is already present, the result will be printed to "Si8_kpt.out_1" instead. The max number of ".out" files allowed with the same name is 100. Once this number is reached, the result will instead overwrite the "Si8_kpt.out" file. One can compare the result with the reference out file named "Si8_kpt.refout".

In the tests/ directory, we also provide a suite of tests which are arranged in a hierarchy of folders. Each test system has its own directory. A python script is also provided which launches the suite of test systems. To run a set of four quick tests locally on the CPU, simply run:

\$ python test.py quick_run

Execution

The result is stored in the corresponding directory of the tests. A message is also printed in the terminal showing if the tests passed or failed. The tests can also be launched in parallel on a cluster by using the Python script. Detailed information on using the python script can be found in the 'ReadMe' file in the 'tests/' directory.

Output

Upon successful execution of the sparc code, depending on the calculations performed, some output files will be created in the same location as the input files.

Single point calculations

- ".out" file General information about the test, including input parameters, SCF convergence progress, ground state properties and timing information.
- ".static" file Atomic positions and atomic forces if the user chooses to print these information.

Structural relaxation calculations

• ".out" file - See above.

Output

- ".geopt" file Atomic positions and atomic forces for atomic relaxation, cell lengths and stress tensor for volume relaxation, and atomic positions, atomic forces, cell dimensions, and stress tensor for full relaxation.
- ".restart" file Information necessary to perform a restarted structural relaxation calculation. Only created if atomic relaxation is performed.

Quantum molecular dynamics (QMD) calculations

- ".out" file See above.
- ".aimd" file Atomic positions, atomic velocities, atomic forces, electronic temperature, ionic temperature and total energy for each QMD step.
- ".restart" file Information necessary to perform a restarted QMD calculation.

System

```
CELL | LATVEC_SCALE | LATVEC | FD_GRID | MESH_SPACING | ECUT | BC |
FD_ORDER | EXCHANGE_CORRELATION | SPIN_TYP | KPOINT_GRID |
KPOINT_SHIFT | ELEC_TEMP_TYPE | ELEC_TEMP | SMEARING | NSTATES |
D3_FLAG | D3_RTHR | D3_CN_THR | EXX_RANGE_FOCK | EXX_RANGE_PBE |
ATOM_TYPE | PSEUDO_POT | N_TYPE_ATOM | COORD | COORD_FRAC | RELAX |
SPIN | HUBBARD_FLAG | HUBBARD | U_ATOM_TYPE | U_VAL
```

SCF

```
CHEB_DEGREE | CHEFSI_BOUND_FLAG | RHO_TRIGGER | NUM_CHEFSI |
MAXIT_SCF | MINIT_SCF | TOL_SCF | SCF_FORCE_ACC | SCF_ENERGY_ACC
TOL_LANCZOS | MIXING_VARIABLE | MIXING_HISTORY |
MIXING_PARAMETER | MIXING_PARAMETER_SIMPLE
MIXING_PARAMETER_MAG | MIXING_PARAMETER_SIMPLE_MAG |
PULAY_FREQUENCY | PULAY_RESTART | MIXING_PRECOND |
MIXING_PRECOND_MAG | TOL_PRECOND | PRECOND_KERKER_KTF |
PRECOND_KERKER_THRESH | PRECOND_KERKER_KTF_MAG |
PRECOND_KERKER_THRESH_MAG | FIX_RAND | TOL_FOCK | MAXIT_FOCK |
MINIT_FOCK | TOL_SCF_INIT | EXX_ACC | EXX_METHOD | EXX_MEM |
EXX_FRAC | EXX_ACE_VALENCE_STATES | EXX_DOWNSAMPLING |
EXX_DIVERGENCE | READ_INIT_DENS
```

Electrostatics

```
TOL_POISSON | MAXIT_POISSON | TOL_PSEUDOCHARGE | REFERENCE_CUTOFF
```

Stress calculation

CALC_STRESS | CALC_PRES

QMD

```
MD_FLAG | MD_METHOD | MD_NSTEP | MD_TIMESTEP | ION_TEMP |
ION_TEMP_END | ION_VEL_DSTR | ION_VEL_DSTR_RAND | QMASS |
NPT_NH_QMASS | NPT_NH_BMASS | NPT_NP_QMASS | NPT_NP_BMASS |
NPT_SCALE_VECS | NPT_SCALE_CONSTRAINTS | TARGET_PRESSURE |
RESTART_FLAG | TWTIME
```

Structural relaxation

```
RELAX_FLAG | RELAX_METHOD | RELAX_NITER | TOL_RELAX |

TOL_RELAX_CELL | RELAX_MAXDILAT | NLCG_SIGMA | L_HISTORY |

L_FINIT_STP | L_MAXMOV | L_AUTOSCALE | L_LINEOPT | L_ICURV |

FIRE_DT | FIRE_MASS | FIRE_MAXMOV | RESTART_FLAG
```

Band structure

BAND_STRUCTURE | KPT_PATHS | KPT_PER_LINE | INPUT_DENS_FILE

Print options

```
PRINT_ATOMS | PRINT_FORCES | PRINT_MDOUT | PRINT_RELAXOUT |
PRINT_RESTART | PRINT_RESTART_FQ | PRINT_VELS | OUTPUT_FILE |
PRINT_EIGEN | PRINT_DENSITY | PRINT_ORBITAL |
PRINT_ENERGY_DENSITY
```

Parallelization options

```
NP_SPIN_PARAL | NP_KPOINT_PARAL | NP_BAND_PARAL |
NP_DOMAIN_PARAL | NP_DOMAIN_PHI_PARAL | EIG_SERIAL_MAXNS |
EIG_PARAL_BLKSZ | EIG_PARAL_ORFAC | EIG_PARAL_MAXNP
```

Orbital Free DFT

OFDFT_FLAG | TOL_OFDFT | OFDFT_LAMBDA

```
Socket communication
```

```
SOCKET_FLAG | SOCKET_HOST | SOCKET_PORT | SOCKET_INET | SOCKET_MAX_NITER |
```

System: .inpt file

CELL

Type

Double

Unit

Bohr

Default

None

Example

CELL: 10.20 11.21 7.58

Description

A set of three whitespace delimited values specifying the cell lengths in the lattice vector (LATVEC) directions, respectively.

Remark

Note that CELL ignores the lengths of the lattice vectors specified in the .inpt file and only treats them as unit vectors. LATVEC_SCALE and CELL cannot be specified simultaneously.

LATVEC_SCALE

Type
Double

Default
None

Unit
Bohr

Example
LATVEC_SCALE: 10.20 11.21 7.58

Description

A set of three whitespace delimited values specifying the scaling factors in the lattice vectors (LATVEC), respectively.

LATVEC_SCALE

Remark

The difference between LATVEC_SCALE and CELL is that CELL treats the lattice vectors as unit vectors, whereas LATVEC_SCALE scales the lattice vectors directly as specified by the user. LATVEC_SCALE and CELL cannot be specified simultaneously.

LATVEC

Type

Double array

Default

1.0 0.0 0.0

0.0 1.0 0.0

0.0 0.0 1.0

Unit

No unit

Example

LATVEC:

0.5 0.5 0.0

0.0 0.5 0.5

0.5 0.0 0.5

Description

A set of three vectors in row major order specifying the lattice vectors of the simulation domain (CELL).

FD_GRID

Type

Integer

Unit

No unit

Default

None

Example

FD_GRID: 26 26 30

Description

A set of three whitespace delimited values specifying the number of finite-difference intervals in the lattice vector (LATVEC) directions, respectively.

FD_GRID

Remark

The convergence of results with respect to spatial discretization needs to be verified. ECUT, MESH_SPACING, FD_GRID cannot be specified simultaneously.

MESH_SPACING

Type

Double

Unit

Bohr

Default

None

Example

MESH_SPACING: 0.4

Description

Mesh spacing of the finite-difference grid.

Remark

The exact mesh-size will be determined by the size of the domain. MESH_SPACING, FD_GRID, ECUT cannot be specified simultaneously.

ECUT

Type Unit
Double Ha

Default Example
None ECUT: 30

Description

Equivalent plane-wave energy cutoff, based on which MESH_SPACING will be automatically calculated.

Remark

This is not exact, but rather an estimate. ECUT, MESH_SPACING, FD_GRID cannot be specified simultaneously.

Type
Character

Default
None

Example
BC: P D D

Description

A set of three whitespace delimited characters specifying the boundary conditions in the lattice vector directions, respectively. P represents periodic boundary conditions and D represents Dirichlet boundary conditions.

FD_ORDER

Type Intege

Integer

Default 12 Unit No unit

Example

FD_ORDER: 12

Description

Order of the finite-difference approximation.

Remark

Restricted to even integers since central finite-differences are employed. The default value of 12 has been found to be an efficient choice for most systems.

EXCHANGE_CORRELATION

Type
String

Unit
No unit

Default
No Default

Example
EXCHANGE_CORRELATION: LDA_PW

Description

Choice of exchange-correlation functional. Options are LDA_PW (Perdew-Wang LDA), LDA_PZ (Purdew-Zunger LDA), GGA_PBE (PBE GGA), GGA_RPBE (revised PBE GGA), and GGA_PBEsol (PBE GGA revised for solids), PBEO, HF (Hartree-Fock), HSE, vdWDF1 (van der Waals Density Functional developed by Dion et al.), vdWDF2 (vdW Density Functional modified by Lee et al), SCAN (SCAN metaGGA), RSCAN (rSCAN metaGGA), and R2SCAN (r2SCAN metaGGA).

EXCHANGE_CORRELATION

Remark

For spin-polarized calculation (SPIN_TYP = 1), LDA_PZ is not available. Currently SCAN, RSCAN and R2SCAN does not support nonlinear core correction pseudopotential.

SPIN_TYP

Type
Integer

Default
0

Unit
No unit

Example
SPIN_TYP: 1

Description

 ${\tt SPIN_TYP:\ 0\ performs\ spin\ unpolarized\ calculation.}$

SPIN_TYP: 1 performs unconstrained collinear spin-polarized calculation.

 ${\tt SPIN_TYP: 1\ performs\ unconstrained\ noncollinear\ spin-polarized\ calculation}.$

SPIN_TYP

Remark

SPIN_TYP can only take values 0, 1, 2. For collinear calculation, non-relativistic pseudopotential need to be used. For noncollinear calculation, fully relativistic pseudopotential need to be used.

KPOINT_GRID

Type

Integer array

Default

1 1 1

Unit

No unit

Example

KPOINT_GRID: 2 3 4

Description

Number of k-points in each direction of the Monkhorst-Pack grid for Brillouin zone integration.

Remark

Time-reversal symmetry is assumed to hold.

KPOINT_SHIFT

Type

Double array

Default

- 0.0 for odd k-point mesh
- 0.5 for even k-point mesh

Unit

No unit

Example

KPOINT_SHIFT: 0.5 0.5 0.5

Description

Shift of k-points with respect to k-point grid containing Γ -point.

Remark

The shift is in reduced coordinates.

ELEC_TEMP_TYPE

Type

String

Default

gaussian

Unit

No unit

Example

ELEC_TEMP_TYPE: fd

Description

Function used for the smearing (electronic temperature). Options are: fermi-dirac (or fd), gaussian.

Remark

Use ELEC_TEMP or SMEARING to set smearing value.

ELEC_TEMP

Type

Double

Default

2320.904 for gaussian 1160.452 for fermi-dirac Unit

Kelvin

Example

ELEC_TEMP: 315.775

Description

Electronic temperature.

Remark

This is equivalent to setting SMEARING (0.001 Ha = 315.775 Kelvin).

SMEARING

Type

Double

Unit

На

Default

0.007350 for gaussian 0.003675 for fermi-dirac Example

SMEARING: 0.001

Description

Value of smearing.

Remark

Equivalent to setting ELEC_TEMP (0.001 Ha = 315.775 Kelvin).

NSTATES

Type

Integer

Default

 $N_e/2 \times 1.2 + 5$

Unit

No unit

Example

NSTATES: 24

Description

The number of Kohn-Sham states/orbitals.

Remark

This number should not be smaller than half of the total number of valence electrons (N_e) in the system. Note that the number of additional states required increases with increasing values of ELEC_TEMP/SMEARING.

D3_FLAG

Type 0 or 1

Unit

No unit

Default

0

Example

D3_FLAG: 1

Description

Flag for adding Grimme's DFT-D3 correction on the result

Remark

Only active when using GGA-PBE, GGA-RPBE and GGA-PBEsol.

D3_RTHR

Type Double Unit

Bohr²

Default

1600

Example

D3_RTHR: 9000

Description

Square of cut-off radius for calculating DFT-D3 correction between two atoms

Remark

Only applicable when DFT-D3 correction D3_FLAG is used.

D3_RTHR should be larger or equal to cutoff radius of CN coefficient, D3_CN_THR.

D3_CN_THR

Type

Double

Unit

Bohr²

Default

625

Example

D3_CN_THR: 1600

Description

Square of cut-off radius for calculating CN value of every atom and DFT-D3 correction between three atoms

Remark

Only applicable when DFT-D3 correction D3_FLAG is used.

D3_CN_THR should be smaller or equal to cutoff radius of DFT-D3 correction cutoff radius, D3_RTHR.

EXX_RANGE_FOCK

Type

Double

Unit

1/Bohr

Default

0.1587

Example

EXX_RANGE_FOCK: 0.106

Description

Short range screen parameter of hartree-fock operator in HSE functional.

Remark

Default is using VASP's HSE03 value. Different code has different parameters. Be careful with the results.

EXX_RANGE_PBE

Type

Double

Unit

1/Bohr

Default

0.1587

Example

EXX_RANGE_PBE: 0.106

Description

Short range screen parameter of PBE in HSE functional.

Remark

Default is using VASP's HSE03 value. Different code has different parameters. Be careful with the results.

HUBBARD_FLAG

Type 0 or 1

Unit

No unit

Default

0

Example

HUBBARD_FLAG: 1

Description

Flag for adding hubbard correction on top of the EXCHANGE_CORRELATION.

Remark

Usually used along with LDA, GGA or SCAN type functionals.

System: .ion file

ATOM_TYPE

Type

String

Unit

No unit

Default

None

Example

ATOM_TYPE: Fe

Description

Atomic type symbol.

Remark

The atomic type symbol can be attached with a number, e.g., Fe1 and Fe2. This feature is useful if one needs to provide two different pseudopotential files (PSEUDO_POT) for the same element.

PSEUDO_POT

Type
String
Unit
No unit

Default
None
Example
PSEUDO_POT: ../psp/Fe.psp8

Description

Path to the pseudopotential file.

Remark

The default directory for the pseudopotential files is the same as the input files. For example, if a pseudopotential Fe.psp8 is put in the same directory as the input files, one can simply specify PSEUDO_POT: Fe.psp8.

N_TYPE_ATOM

Type
Integer

Default
None

Unit
No unit

Example
N_TYPE_ATOM: 2

Description

The number of atoms of a ATOM_TYPE specified immediately before this variable.

Remark

For a system with different types of atoms, one has to specify the number of atoms for every type.

COORD

Туре	Unit
Double	Bohr
Default	Example
None	COORD: 0.0 0.0 0.0 2.5 2.5 2.5

Description

The Cartesian coordinates of atoms of a ATOM_TYPE specified before this variable. If the coordinates are outside the fundamental domain (see CELL and LATVEC) in the periodic directions (see BC), it will be automatically mapped back to the domain.

COORD

Remark

For a system with different types of atoms, one has to specify the coordinates for every ATOM_TYPE. One can also specify the coordinates of the atoms using COORD_FRAC.

COORD_FRAC

Туре	Unit
Double	None
Default	Example
None	COORD_FRAC:
	0.5 0.5 0.0
	0.0 0.5 0.5

Description

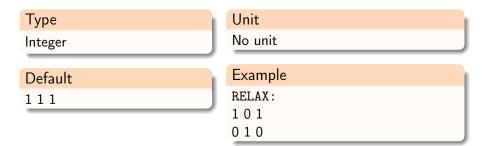
The fractional coordinates of atoms of a ATOM_TYPE specified before this variable. COORD_FRAC $(i,j) \times \text{CELL}(j)$, (j=1,2,3) gives the coordinate of the i^{th} atom along the j^{th} LATVEC direction. If the coordinates are outside the fundamental domain (see CELL and LATVEC) in the periodic directions (see BC), it will be automatically mapped back to the domain.

COORD_FRAC

Remark

For a system with different types of atoms, one has to specify the coordinates for every ATOM_TYPE. One can also specify the coordinates of the atoms using COORD.

RELAX



Description

Atomic coordinate with the corresponding RELAX value 0 is held fixed during relaxation/QMD.

SPIN

Туре	Unit	
Double	No unit	
Default	Example	
0.0	SPIN:	
	0 0 1.0	
	0 0 -1.0	

Description

Specifies the net initial spin on each atom for a spin-polarized calculation. If collinear spin used, user could use either 1 column of data for z-direction of each atom, or 3 columns of data with 0 on the first 2 columns (x,y-directions). For noncollinear spin, use need to use 3 columns of data for all directions.

HUBBARD

Type
None
Unit
No unit

Default
None
Example
HUBBARD:

Description

Start of the DFT+U parameter block. Must be followed by specifying the atoms on which a U correction is desired along with value of U as per Dudarev's scheme.

Remark

HUBBARD_FLAG must be specified to 1 in .inpt file.

U_ATOM_TYPE

Type
String
Unit
No unit

Default
None

Example
U_ATOM_TYPE: Ni

Description

Atomic type symbol for which a U correction is desired. Must be specified within the HUBBARD block. The atomic type should already have been specified above in the .ion file.

U_VAL

Type
Double

Default
None

Unit
Hartree

Example
U_VAL: 0 0 0.1 0

Description

Value of effective U for every azimuthal quantum number (s, p, d, and f orbitals) of the atom on which the correction is applied per Dudarev's scheme. The corrections will only be applied to the outermost states available in the corresponding PSEUDO_POT. Must be specified after specifying the U_ATOM_TYPE.

SCF

CHEB_DEGREE

Type

Integer

Default

Automatically set.

Unit

No unit

Example

CHEB_DEGEE: 25

Description

Degree of polynomial used for Chebyshev filtering.

Remark

For larger mesh-sizes, smaller values of CHEB_DEGREE are generally more efficient, and vice-versa.

CHEFSI_BOUND_FLAG

Type
Integer

Default
0

Unit
No unit

Example
CHEFSI_BOUND_FLAG: 1

Description

Flag to recalculate the bounds for Chebyshev filtering. If set to 0, then only for the very first SCF will the upper bound be evaluated based on the maximum eigenvalue using Lanczos algorithm, and the upper bound will be the same for the rest steps. If set to 1, the upper bound will be reevaluated for every SCF.

RHO_TRIGGER

Type

Integer

Default

4, 6, or 10

Unit

No unit

Example

RHO_TRIGGER: 5

Description

The number of times Chebyshev filtering is repeated before updating the electron density in the very first SCF iteration.

RHO_TRIGGER

Remark

Values smaller than the default value of 4 can result in a significant increase in the number of SCF iterations. Larger values can sometimes reduce the number of SCF iterations. For non-collinear spin calculation, default is 6 otherwise 4. For band structure calculations, default is 10.

NUM_CHEFSI

Type Integer

nteger

Default

Ŧ

Unit

No unit

Example

NUM_CHEFSI: 2

Description

The number of times ChefSI algorithm is repeated in SCF iteration except the first one, which is controlled by RHO_TRIGGER.

Remark

For non-collinear spin calculation, it might helped SCF convergence in some cases.

MAXIT_SCF

Type

Integer

Unit

No unit

Default

100

Example

MAXIT_SCF: 50

Description

Maximum number of SCF iterations.

Remark

Larger values than the default of 100 may be required for highly inhomogeneous systems, particularly when small values of SMEARING/ELEC_TEMP are chosen.

MINIT_SCF

Type
Integer

Default

MAXIT_SCF: 5

Description

Minimum number of SCF iterations.

TOL_SCF

Type
Double

Default
see description

Unit
No unit

Example
TOL_SCF: 1e-5

Description

In case of single point calculation, TOL_SCF is set for 10^{-5} Ha/atom energy accuracy. In case of MD, TOL_SCF is set for 10^{-3} Ha/Bohr force accuracy. In case of relaxation, TOL_SCF is set for TOL_RELAX/5 Ha/Bohr force accuracy.

The tolerance on the normalized residual of the effective potential or the electron density for convergence of the SCF iteration.

TOL_SCF

Remark

Only one of TOL_SCF, SCF_ENERGY_ACC, or SCF_FORCE_ACC can be specified.

SCF_FORCE_ACC

Type

Double

Unit

Ha/Bohr

Default

None

Example

SCF_FORCE_ACC: 1e-4

Description

The tolerance on the atomic forces for convergence of the SCF iteration.

Remark

Only one of SCF_FORCE_ACC, TOL_SCF or SCF_ENERGY_ACC can be specified.

SCF_ENERGY_ACC

Type

Double

Unit

Ha/atom

Default

None

Example

SCF_ENERGY_ACC: 1e-5

Description

The tolerance on the free energy for the convergence of the SCF iteration.

Remark

Only one of SCF_ENERGY_ACC, SCF_FORCE_ACC, or TOL_SCF can be specified.

TOL_LANCZOS

Type

Double

Unit

No unit

Default

1e-2

Example

TOL_LANCZOS: 1e-3

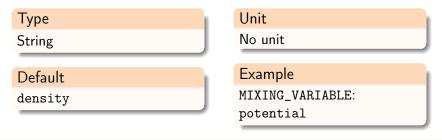
Description

The tolerance within the Lanczos algorithm for calculating the extremal eigenvalues of the Hamiltonian, required as part of the CheFSI method.

Remark

Typically, the Lanczos tolerance does not need to be very strict.

MIXING_VARIABLE



Description

This specifies whether potential or density mixing is performed in the SCF iteration. Available options are: potential and density.

MIXING_HISTORY

Type Integer

nteger

Default

1

Unit

No unit

Example

MIXING_HISTORY: 40

Description

The mixing history used in Pulay mixing.

Remark

Too small values of MIXING_HISTORY can result in poor SCF convergence.

MIXING_PARAMETER

Type

Double

Unit

No unit

Default

0.3

Example

MIXING_PARAMETER: 0.1

Description

The value of the relaxation parameter used in Pulay/simple mixing.

Remark

Values larger than the default value of 0.3 can be used for insulating systems, whereas smaller values are generally required for metallic systems, particularly at small values of SMEARING or ELEC_TEMP.

MIXING_PARAMETER_SIMPLE

Type

Double

Default

Automatically set to the same as MIXING_PARAMETER

Unit

No unit

Example

MIXING_PARAMETER_SIMPLE: 0.1

Description

The value of the relaxation parameter used in the simple mixing step in the periodic Pulay scheme.

MIXING_PARAMETER_MAG

Type

Double

Default

Automatically set to the same as MIXING_PARAMETER.

Unit

No unit

Example

MIXING_PARAMETER_MAG: 4.0

Description

The mixing parameter for the magnetization density in Pulay mixing for spin-polarized calculations.

MIXING_PARAMETER_MAG

Remark

For spin-polarized calculations, when SCF has difficulty to converge, increasing the mixing parameter to magnetization density might help. For example, setting it to 4.0, while turning off the preconditioner applied to the magnetization density (by setting MIXING_PRECOND_MAG to 'none') is a good choice.

MIXING_PARAMETER_SIMPLE_MAG

Type

Double

Unit

No unit

Default

Automatically set to the same as MIXING_PARAMETER_MAG

Example

MIXING_PARAMETER_SIMPLE_MAG: 4.0

Description

The value of the relaxation parameter for the magnetization density used in the simple mixing step in the periodic Pulay scheme for spin-polarized calculations.

PULAY_FREQUENCY

Type
Integer

Default
1

Unit
No unit

Example
PULAY_FREQUENCY: 4

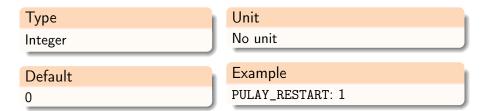
Description

The frequency of Pulay mixing in Periodic Pulay.

Remark

The default value of 1 corresponds to Pulay mixing.

PULAY_RESTART



Description

The flag for restarting the 'Periodic Pulay' mixing. If set to 0, the restarted Pulay method is turned off.

MIXING_PRECOND

Type

String

Default

kerker

Unit

No unit

Example

MIXING_PRECOND: none

Description

This specifies the preconditioner used in the SCF iteration. Available options are: none, kerker.

MIXING_PRECOND_MAG

Type
String

Unit
No unit

Default
none

Example
MIXING_PRECOND_MAG: kerker

Description

This specifies the preconditioner used for the magnetization density in the SCF iteration for spin-polarized calculations. Available options are: none, kerker.

TOL_PRECOND

Type

Double

Unit

No unit

Default

 $h^2 \times 0.001$

Example

TOL_PRECOND: 1e-4

Description

The tolerance on the relative residual for the linear systems arising during the real-space preconditioning of the SCF.

Remark

The linear systems do not need to be solved very accurately. h is the mesh spacing.

PRECOND_KERKER_KTF

Type
Double

Default

1.0

Unit
Bohr⁻¹

Example
PRECOND_KERKER_KTF: 0.8

Description

The Thomas-Fermi screening length appearing in the kerker preconditioner (MIXING_PRECOND).

PRECOND_KERKER_THRESH

Type
Double

Default
0.1

Unit
No unit

Example
PRECOND_KERKER_THRESH: 0.0

Description

The threshold for the kerker preconditioner (MIXING_PRECOND).

Remark

This threshold will be scaled by the MIXING_PARAMETER. If the threshold is set to 0, the original kerker preconditioner is recovered.

PRECOND_KERKER_KTF_MAG

Type
Double

Default
1.0

Unit
Bohr⁻¹

Example
PRECOND_KERKER_KTF_MAG: 0.8

Description

The Thomas-Fermi screening length appearing in the kerker preconditioner for the magnetization density (MIXING_PRECOND_MAG).

PRECOND_KERKER_THRESH_MAG

Type
Double

Default
0.1

Unit
No unit

Example
PRECOND_KERKER_THRESH_MAG: 0.0

Description

The threshold for the kerker preconditioner the magnetization density (MIXING_PRECOND_MAG).

Remark

This threshold will be scaled by the MIXING_PARAMETER_MAG. If the threshold is set to 0, the original kerker preconditioner is recovered.

FIX_RAND

Туре	Unit
Integer	No unit
Default	Example
0	FIX_RAND: 1

Description

Flag to fix the random seeds for setting initial guesses. Once set to 1, the random seeds will be fixed for different runs and for different numbers of processors. This option will make sure the answers will be exactly the same (up to machine precision) when SPARC is executed with different numbers of processors.

TOL_FOCK

Type

Double

Default

Automatically set for 10^{-3} Ha/Bohr accuracy in force

Unit

No unit

Example

TOL_FOCK: 1e-6

Description

The tolerance on the Hartree-Fock outer loop, measured by the difference between the band energy in 2 consecutive outer loops.

Remark

Only active when using hybrid functionals, like PBE0 and HSE.

MAXIT_FOCK

Type Intege

Integer

Default 20 Unit

No unit

Example

MAXIT_FOCK: 50

Description

The maximum number of iterations for Hartree-Fock outer loop.

Remark

Only active when using hybrid functionals, like PBE0 and HSE.

MINIT_FOCK

Type
Integer

Default

MINIT_FOCK: 3

Description

The minimum number of iterations for Hartree-Fock outer loop.

Remark

Only active when using hybrid functionals, like PBE0 and HSE.

TOL_SCF_INIT

Type

Double

Unit

No unit

Default

 $max(TOL_FOCK \times 10, 0.001)$

Example

TOL_SCF_INIT: 1e-6

Description

The initial SCF tolerance for PBE iteration when using hybrid functionals.

Remark

Only active when using hybrid functionals, like PBE0 and HSE. Change the TOL_SCF_INIT to change the initial guess for Hartree Fock outer loop.

EXX_ACC

Type Integer

Default

1

Unit

No unit

Example

EXX_ACC: 0

Description

Use accelerated method for exact exchange. For low-T, use ACE operator to accelarte the hybrid calculation, while in high-T, use the local projection method.

Remark

Without acceleration, the hybrid functional will be extremely slow.

EXX_METHOD

Type	Unit
String	No unit
	For mark
Default	Example
KRON	EXX_METHOD: FFT

Description

Method to solve the Poisson's equations in exact exchange. Options include KRON (Laplacian using kronecker product) and FFFT (FFT to solve in Fourier space). Now, KRON is only available for orthogonal periodic cell on gamma point and dirichlet molecule using PBE0. Generally, KRON way is faster than FFT, while FFT is faster on certain type of grid with even number of grid points each direction etc.

EXX_MEM

Туре	Unit
Integer	No unit
Default	Example
20	EXX_MEM: 0

Description

Number of Poisson's equations to be solved in each process at a time when creating exact exchange operator or ACE operator. Typically, when EXX_MEM is larger than 20, the speed of code is barely affected. When it is 0, all Poisson's equations are solved together and it hits the fastest speed but largest memory requirement. Only effective on low-T calculation.

EXX_FRAC

Type

Double

Unit

No unit

Default

0.25 for PBE0 and HSE

Example

EXX_FRAC: 0.3

Description

Fraction of exact exchange in hybrid functional, e.g. PBE0 and HSE, while the fraction of PBE is $1\text{-}EXX_FRAC$

EXX_ACE_VALENCE_STATES

Type

Integer

Default

3

Unit

No unit

Example

EXX_ACE_VALENCE_STATES: 1

Description

Control of number of unoccupied states used to construct ACE operator. Only effective on low-T calculation.

Remark

Only active when using hybrid functionals with ACE operator.

EXX_DOWNSAMPLING

Type Integer

Unit No unit

Default

Example

EXX_DOWNSAMPLING: 1 2 3

Description

Down-sampling of k-points grids. There should be 3 nonnegative integers. 0 means using 0 k-point in that direction, requiring 0 is one of the k-point after time-reversal symmetry in that direction. Positive value should be a factor of the number of grid points in that direction. Only effective on low-T calculation.

EXX_DOWNSAMPLING

Remark

One could only do downsampling on directions of periodic boundary condition.

EXX_DIVERGENCE

Type

String

Unit

No unit

Default

SPHERICAL

Example

EXX_DIVERGENCE: AUXILIARY

Description

Treatment of divergence in exact exchange. Options are SPHERICAL (spherical truncation), AUXILIARY (auxiliary function method) and ERFC (erfc screening). Only effective on low-T calculation.

EXX_DIVERGENCE

Remark

For systems with cube-like geometry, both methods converge fast. For slab and wire, auxiliary function method is a better option. ERFC screening is the default option for HSE in bulk and molecule simulation.

READ_INIT_DENS

Type
Integer

Default
0

Unit
No unit

Example
READ_INIT_DENS: 1

Description

Flag for reading initial density guess for SCF calculation.

Remark

Please use it together with INPUT_DENS_FILE

Electrostatics

TOL_POISSON

Type

Integer

Unit

No unit

Default

 $TOL_SCF \times 0.01$

Example

TOL_POISSON: 1e-6

Description

The tolerance on the norm of the relative residual for the Poisson equation.

Remark

The tolerance for poisson solver should not be worse than TOL_SCF, otherwise it might seriously affect the convergence of the SCF iteration.

MAXIT_POISSON

Type Integer

Default 3000 Unit

No unit

Example

MAXIT_POISSON: 1000

Description

The maximum number of iterations for solving the Poisson equation using an iterative linear solver.

TOL_PSEUDOCHARGE

Type

Double

Unit

No unit

Default

 $TOL_SCF \times 0.001$

Example

TOL_PSEUDOCHARGE: 1e-6

Description

The error in the net enclosed charge for the pseudocharge density of each atom.

REFERENCE_CUTOFF

Type

Double

Unit

Bohr

Default

0.5

Example

REFERENCE_CUTOFF: 1.0

Description

The cutoff radius of the reference potential used for evaluating the electrostatic correction arising from overlapping pseudocharge densities.

Remark

This number should be smaller than half the smallest interatomic distance.

Stress calculation

CALC_STRESS

Type Integer

Unit

No unit

Example

CALC_STRESS: 1

Default

Description

Flag for calculation of the Hellmann-Feynman stress tensor (in cartesian coordinates).

CALC_PRES

Type

Integer

Unit

No unit

Default

U

Example

CALC_PRES: 1

Description

Flag for calculation of the pressure.

Remark

Pressure is directly calculated, without calculation of the stress tensor.

QMD

MD_FLAG

Type Integer

Unit

No unit

Default

0

Example

MD_FLAG: 1

Description

QMD simulations are performed if the flag is set to 1.

Remark

MD_FLAG and RELAX_FLAG both cannot be set to a value greater than 0.

MD_METHOD

Type

String

Default

 NVT_NH

Unit

No unit

Example

MD_METHOD: NVE

Description

Type of QMD to be performed. Currently, NVE (microcanonical ensemble), NVT_NH (canonical ensemble with Nose-Hoover thermostat), NVK_G (isokinetic ensemble with Gaussian thermostat), NPT_NH (isothermal-isobaric ensemble with Nose-Hoover thermostat) and NPT_NP (isothermal-isobaric ensemble with Nose-Poincare thermostat) are supported.

MD_NSTEP

Type

Integer

Unit

No unit

Default

1e7

Example

MD_NSTEP: 100

Description

Specifies the number of QMD steps.

Remark

If MD_NSTEP = N, the QMD runs from 0 to $(N-1) \times MD_TIMESTEP$ fs.

MD_TIMESTEP

Type Double

Default

1

Unit

Femtosecond

Example

MD_TIMESTEP: 0.1

Description

QMD time step.

Remark

Total QMD time is given by: $MD_TIMESTEP \times (MD_NSTEP - 1)$.

ION_TEMP

Type

Double

Unit

Kelvin

Default

No Default

Example

ION_TEMP: 315

Description

Starting ionic temperature in QMD, used to generate initial velocity distribution.

Remark

Must be specified if MD_FLAG is set to 1. It is also the target temperature in MD_METHOD NPT_NH and NPT_NP.

ION_TEMP_END

Type

Double

Unit

Kelvin

Default

ION_TEMP

Example

ION_TEMP_END: 100

Description

Specifies the final temperature of the thermostat. Thermostat temperature is varied linearly from ION_TEMP to ION_TEMP_END with respect to time.

Remark

Available for NVT_NH quantum molecular dynamics only. Not supported in NPT_NH and NPT_NP.

ION_VEL_DSTR

Type

Integer

Unit

No unit

Default

2

Example

ION_VEL_DSTR: 1

Description

Specifies the type of distribution for the initial velocity of atoms based on their initial temperature. Choose 1 for uniform velocity distribution and 2 for Maxwell-Boltzmann distribution.

Remark

Currently, the code supports only two options for the variable.

ION_VEL_DSTR_RAND

Type
Integer

Default
0

Unit
No unit

Example
ION_VEL_DSTR_RAND: 1

Description

Flag to reseed the initial velocities of atoms in a QMD simulation. Set this flag to 1 to shuffle (change the random seed for) the initial velocities for different runs. Set this flag to 0 to maintain the same initial velocities.

Remark

This option is convenient for parallel statistics calculations.

QMASS

Type

Double

Unit

atomic unit

Default

1653.654933459720

Example

QMASS: 100000

Description

Gives the inertia factor for Nose Hoover thermostat.

Remark

Applicable to NVT_NH MD_METHOD only.

NPT_NH_QMASS

Type

1st number int; others double

Default

No default value

Unit

atomic unit

Example

NPT_NH_QMASS: 2 700.0 700.0

Description

Gives the amount (first number) and inertia masses (others) of thermostats in NPT_NH. The maximum amount of thermostat variables of the Nose-Hoover chain is 60

Remark

Applicable to NPT_NH $\underline{\text{MD_METHOD}}$ only. Program will exit if NPT_NH is selected but NPT_NH_QMASS is not input

NPT_NH_BMASS

Type

Double

Unit

atomic unit

Default

No default value

Example

NPT_NH_BMASS: 5000

Description

Gives the inertia mass for the barostat variable in NPT_NH.

Remark

Applicable to NPT_NH ${\tt MD_METHOD}$ only. Program will exit if NPT_NH is selected but NPT_NH_BMASS is not input

NPT_NP_QMASS

Type

Double

Unit

atomic unit

Default

No default value

Example

NPT_NP_QMASS: 100

Description

Gives the inertia mass for the thermostat variable in NPT_NP.

Remark

Applicable to NPT_NP $\underline{\text{MD_METHOD}}$ only. Program will exit if NPT_NP is selected but NPT_NP_BMASS is not input

NPT_NP_BMASS

Type

Double

Unit

atomic unit

Default

No default value

Example

NPT_NP_BMASS: 20

Description

Gives the inertia mass for the barostat variable in NPT_NP.

Remark

Applicable to NPT_NP ${\tt MD_METHOD}$ only. Program will exit if NPT_NP is selected but NPT_NP_BMASS is not input

NPT_SCALE_VECS

Type Unit No unit Int Example **Default**

NPT_SCALE_VECS: 2

Description

123

Specify which lattice vectors can be rescaled in NPT NH and NPT NP. The cell will only expand or shrink in the specified directions. Rescaled vectors can be specified for orthogonal systems if NPT NP thermostat is used.

NPT_SCALE_VECS

Remark

Only three numbers 1, 2 and 3 can be accepted. For example, if "2 3" is the input, the cell will only expand or shrink in the directions of lattice vector 2 and lattice vector 3.

If it is set in NPT_NH, the expansion or shrinkage on designated lattice vector will try to keep the total pressure to oscillate near the target pressure.

If it is set in NPT_NP, the expansion or shrinkage on designated lattice vector will only try to keep the normal stress at their direction to oscillate near the target pressure.

NPT_SCALE_CONSTRAINTS

Type
Double

Default
none

Unit
No unit

Example
NPT_SCALE_CONSTRAINTS: 12

Description

Set the scale constraint for lattice vectors in NPT_NP. The length ratio between the designated lattice vector keeps constant in NPT_NP thermostat. For example, if "12" is set, then the length ratio between 1st and 2nd lattice vectors will keep constant.

NPT_SCALE_CONSTRAINTS

Remark

Applicable to orthogonal system using NPT_NP MD_METHOD only. There are 4 types of available constraints. "12" or "21"; "13" or "31"; "23" or "32"; "123" or "132" or "213" or "231" or "312" or "321".

TARGET_PRESSURE

Type

Double

Unit

GPa

Default

0.0

Example

TARGET_PRESSURE: 40.9611

Description

Gives the outer pressure in NPT_NH and NPT_NP.

Remark

Applicable to NPT_NH and NPT_NP MD_METHOD only.

RESTART_FLAG

Type
Integer

Default
0

Unit
No unit

Example
RESTART_FLAG: 0

Description

Flag for restarting quantum molecular dynamics and structural relaxation. Stores last three histories for quantum molecular dynamics simulations in .restart, .restart-0 and .restart-1 files, respectively.

RESTART_FLAG

Remark

Restarts from the previous configuration which is stored in a .restart file. Currently, code provides restart feature for atomic relaxation and QMD only.

TWTIME

Type
Double

Default
1e9

Unit
min

Example
TWTIME: 1000

Description

Gives the upper bound on the wall time for quantum molecular dynamics.

Structural relaxation

RELAX_FLAG

Type
Integer

Default
0

Unit
No unit

Example
RELAX_FLAG: 1

Description

Flag for performing structural relaxation. 0 means no structural relaxation. 1 represents relaxation of atom positions. 2 represents optimization of volume with the fractional coordinates of the atoms fixed. 3 represents full optimization of the cell i.e., both atoms and cell volume are relaxed

Remark

This flag should not be specified if MD_FLAG is set to 1.

RELAX_METHOD

Type

String

Default

LBFGS

Unit

No unit

Example

RELAX_METHOD: NLCG

Description

Specifies the algorithm for structural relaxation. The choices are 'LBFGS' (limited-memory BFGS), 'NLCG' (Non-linear conjugate gradient), and 'FIRE' (Fast inertial relaxation engine).

Remark

LBFGS is typically the best choice.

RELAX_NITER

Type

Integer

Unit

No unit

Default

300

Example

RELAX_NITER: 25

Description

Specifies the maximum number of iterations for the structural relaxation (RELAX_FLAG).

Remark

If RESTART_FLAG is set to 1, then relaxation will restart from the last atomic configuration and run for maximum of RELAX_NITER iterations.

TOL_RELAX

Type
Double
Unit
Ha/Bohr

Default
5e-4

Unit
Type
ToL_RELAX: 1e-3

Description

Specifies the tolerance for termination of the structural relaxation. The tolerance is defined on the maximum force component (in absolute sense) over all atoms.

TOL_RELAX_CELL

Туре

Double

Default

1e-2

Unit

GPa

Example

TOL_RELAX: 1e-3

Description

Specifies the tolerance for termination of the cell relaxation. The tolerance is defined on the maximum principle stress component.

RELAX_MAXDILAT

Type
Double

Default
1.06

Unit
No unit

Example
RELAX_MAXDILAT: 1.4

Description

The maximum scaling of the volume allowed with respect to the initial volume defined by CELL and LATVEC. This will determine the upper-bound and lower-bound in the bisection method (Brent's method) for the volume optimization.

NLCG_SIGMA

Type

Double

Unit

No unit

Default

0.5

Example

NLCG_SIGMA: 1

Description

Parameter in the secant method used to control the step length in NLCG (RELAX_METHOD).

Remark

L_HISTORY

Type Integer

Unit

No unit

Example

L_HISTORY: 15

Description

20

Size of history in LBFGS (RELAX_METHOD).

Remark

L_FINIT_STP

Type

Double

Unit

Bohr

Default

5e-3

Example

L_FINIT_STP: 0.01

Description

Step length for line optimizer in LBFGS (RELAX_METHOD).

Remark

L_MAXMOV

Type

Double

Default

0.2

Unit

Bohr

Example

L_MAXMOV: 1.0

Description

The maximum allowed step size in LBFGS (RELAX_METHOD).

Remark

L_AUTOSCALE

Type Intege

Integer

Default

1

Unit

No unit

Example

L_AUTOSCALE: 0

Description

Flag for automatically determining the inverse curvature that is used to determine the direction for next iteration in LBFGS (RELAX_METHOD).

Remark

L_LINEOPT

Type
Integer

Default

L_LINEOPT: 0

Description

Flag for atomic force based line minimization in LBFGS (RELAX_METHOD).

Remark

Required only if L_AUTOSCALE is 0.

L_ICURV

Type

Double

Unit

No unit

Default

1.0

Example

L_ICURV: 0.1

Description

Initial inverse curvature, used to construct the inverse Hessian matrix in LBFGS (RELAX_METHOD).

Remark

Needed only if L_AUTOSCALE is 0. Default value works well in most cases.

FIRE_DT

Type
Double

Default

Femto second

Example

FIRE DT: 0.1

Description

Time step used in FIRE (RELAX_METHOD).

Remark

FIRE_MASS

Type

Double

Unit

Atomic mass unit

Default

1.0

Example

FIRE_MASS: 2.5

Description

Pseudomass used in FIRE (RELAX_METHOD).

Remark

FIRE_MAXMOV

Type Double

Unit

Bohr

Default

0.2

Example

FIRE_MAXMOV: 1.0

Description

Maximum movement for any atom in FIRE (RELAX_METHOD).

Remark

Band structure

BAND_STRUCTURE

Type
Integer

Default
0

Unit
No unit

Example
BAND_STRUCTURE: 1

Description

Flag for calculation of the band structure.

KPT_PATHS

Туре	Unit
Double	No unit
Default	Example
None	KPT_PATHS: 2 0.0 0.0 0.0 # Gamma 0.5 0.5 0.5 # A
	0.5 0.5 0.5 # A 0.5 0.5 0.0 # M

Description

The k-point paths (high-symmetry lines) for band structure calculations. It starts with the total number of high-symmetry lines, followed by fractional coordinates indicating the start and end points of each line.

KPT_PER_LINE

Type

Integer

Unit

No unit

Default

10

Example

KPT_PER_LINE: 15

Description

The number of k-points on each high-symmetry line.

Remark

This value accounts for both endpoints of the high-symmetry line.

INPUT_DENS_FILE

Type

String

Unit

No unit

Default

None

Example

INPUT_DENS_FILE: Si.dens Si.densUp Si.densDwn

Description

The name of the input density file(s) in cube format.

Remark

For spin-unpolarized systems, only the total density file is required. For spin-polarized systems, three density files are required, which are the total electron density, spin-up density, and spin-down density, respectively.

Print options

PRINT_ATOMS

Type
0 or 1

Default
0
PRINT_ATOMS: 1

Description

Flag for writing the atomic positions. For ground-state calculations, atom positions are printed to a '.static' output file. For structural relaxation calculations, atom positions are printed to a '.geopt' file. For QMD calculations, atom positions are printed to a '.aimd' file.

PRINT_FORCES

Type
0 or 1

Default

Default

PRINT_FORCES: 1

Description

Flag for writing the atomic forces. For ground-state calculations, forces are printed to a '.static' output file. For structural relaxation calculations, forces are printed to a '.geopt' file. For QMD calculations, forces are printed to a '.aimd' file.

PRINT_MDOUT

Type
0 or 1

Default
1

Default
1

Unit
No unit

Example
PRINT_MDOUT: 0

Description

Flag for printing the the QMD output into the .aimd file.

PRINT_RELAXOUT

Type 0 or 1 Unit

No unit

Default

Example

PRINT_RELAXOUT: 0

Description

Flag for printing the structural relaxation data in a .geopt file.

Remark

Required only if RELAX_FLAG is greater than 0.

PRINT_RESTART

Type

0 or 1

Unit

No unit

Default

1

Example

PRINT_RESTART: 0

Description

Flag for writing the .restart file, used to restart QMD and structural relaxation simulations.

Remark

Relevant only if either MD_FLAG is 1 or RELAX_FLAG is 1.

PRINT_RESTART_FQ

Type

Integer

Unit

No unit

Default

1

Example

PRINT_RESTART_FQ: 10

Description

Frequency at which .restart file is written in QMD and structural optimization simulations.

Remark

Relevant only if either MD_FLAG is 1 or RELAX_FLAG is 1.

PRINT_VELS

Type
0 or 1

Default
1

Unit
No unit

Example
PRINT_VELS: 0

Description

Flag for printing the ion velocities in an QMD simulation into the .aimd file.

Remark

Relevant only if MD_FLAG is set to 1.

OUTPUT_FILE

Type

String

Default

Same as the input file name

Unit

No unit

Example

OUTPUT_FILE: myfname

Description

The name of the output files. The output files are attached with a suffix ('.out', '.static', '.geopt' and '.aimd').

OUTPUT_FILE

Remark

If an output file with the same name already exist, the results will be written to a file with a number attached, e.g., 'myfname.out $_1$ '. The maximum number of output files with the same name allowed is 100. After that the output files will be overwritten in succession.

PRINT_EIGEN

Type
int

Default

Default

PRINT_EIGEN: 1

Description

Flag for writing eigenvalues and occupations into .eigen file.

PRINT_DENSITY

Туре	Unit
int	No unit
Default	Example
Delault	•
0	PRINT_DENSITY: 1

Description

Flag for writing electron density into cube format. For spin-unpolarized calculation, electron density is printed into .dens file. For collinear spin calculation, total, spin-up and spin-down electron density are printed into .dens, .densUp and .densDwn file, respectively.

Type
int

Default

Default

PRINT_ORBITAL: 1

Description

Flag for writing Kohn-Sham orbitals into a binary file.

Remark

It consists of headers with system information and the orbitals. First define a few variables and their types.

name	type, length	description
Nx Ny Nz	int, 1	Number of FD nodes in x,y,z directions
Nd	int, 1	Total number of FD nodes
dx, dy,dz	double, 1	mesh size in x,y,z directions
dV	double, 1	unit Volume
Nspinor	int, 2	Number of spinor in orbitals
isReal	int, 1	Flag for orbitals being real or complex
nspin	int, 1	Number of spin channel printed
nkpt	int, 1	Number of k-point printed

Remark - cont.

nband	int, 1	Number of bands printed
name	type, length	description
spin_index	int, 1	spin index of specific orbital
kpt_index	int, 1	k-point index of specific orbital
kpt_vec	double, 3	k-point in reduced coordinates
band_index	int, 1	band index of specific orbital
psi_real	double, Nd	real Kohn-Sham orbitals
psi_complex	k double complex, Nd	complex Kohn-Sham orbitals

The header is organized as: Nx, Ny, Nz, Nd, dx, dy, dz, dV, Nspinor, isReal, nspin, nkpt, nband, followed by the data for Kohn-sham orbital. Below is the pseudo-code to read orbitals after reading variables in headers.

Remark - cont.

```
for ispin = 1:nspin do
   for ikpt = 1:nkpt do
      for iband = 1:nband do
          spin index, kpt_index, kpt_vec, band_index
          if isReal == 1 then
             psi real
          else
             psi complex
          end if
      end for
   end for
end for
```

PRINT_ENERGY_DENSITY

Type Unit No unit

Default Example

PRINT_ENERGY_DENSITY: 1

Description

Flag for writing a few energy densities into cube format. Currently, only kinetic energy density, exchange correlation energy density (without exact exchange contribution) and exact exchange energy density (if any) are implemented.

PRINT_ENERGY_DENSITY

Remark

For spin-unpolarized calculation, kinetic energy density is written into .kedens, exchange correlation energy density is written into .xcedens, and exact exchange energy density is written into .exxedens. For collinear spin calculation, total, spin-up, spin-down kinetic energy density are written into .kedens, kedensUp, kedensDwn files, total, spin-up, spin-down exact exchange energy density are writted into .exxedens, .exxedensUp, .exxedensDwn files, respectively.

Parallelization options

NP_SPIN_PARAL

Type

Integer

Default

Automatically optimized

Unit

No unit

Example

NP_SPIN_PARAL: 2

Description

Number of spin groups.

Remark

NP_KPOINT_PARAL

Type

Integer

Default

Automatically optimized

Unit

No unit

Example

NP_KPOINT_PARAL: 5

Description

Number of k-point groups.

Remark

NP_BAND_PARAL

Type

Integer

Default

Automatically optimized

Unit

No unit

Example

NP_BAND_PARAL: 5

Description

Number of band groups.

Remark

NP_DOMAIN_PARAL

Type

Integer

Default

Automatically optimized

Unit

No unit

Example

NP_DOMAIN_PARAL: 3 3 2

Description

Dimensions of the 3D Cartesian topology embedded in each band group.

Remark

NP_DOMAIN_PHI_PARAL

Type

Integer

Default

Automatically optimized

Unit

No unit

Example

NP_DOMAIN_PHI_PARAL: 1 1 2

Description

Dimensions of the 3D Cartesian topology embedded in the global communicator.

Remark

EIG_SERIAL_MAXNS

Type

Integer

Unit

No unit

Default

2000

Example

EIG_SERIAL_MAXNS: 1000

Description

Maximum NSTATES value up to which a serial algorithm will be used to solve the subspace eigenproblem.

Remark

If one wants to use a parallel algorithm to solve the subspace eigenproblem for all cases, simply set EIG_SERIAL_MAXNS to 0. Alternatively, set EIG_SERIAL_MAXNS to a very large value to always use serial algorithm.

EIG_PARAL_BLKSZ

Type Integer

Default

128

Unit

No unit

Example

EIG_PARAL_BLKSZ: 64

Description

Block size for the distribution of matrix in block-cyclic format in a parallel algorithm for solving the subspace eigenproblem.

EIG_PARAL_ORFAC

Type Double

Unit No unit

Default

Example

0.0

EIG_PARAL_ORFAC: 0.001

Description

Specifies which eigenvectors should be reorthogonalized when using the parallel eigensolver p?syevx or p?sygvx for solving the subspace eigenproblem. The parallel eigensolvers can be turned on using the EIG_SERIAL_MAXNS flag. No reorthogonalization will be done if EIG_PARAL_ORFAC equals zero. A default value of 0.001 is used if EIG_PARAL_ORFAC is negative. Note that reorthogonalization of eigenvectors is extremely time-consuming.

EIG_PARAL_MAXNP

Type

Integer

Default

Generated by a linear model.

Unit

No unit

Example

EIG_PARAL_MAXNP: 36

Description

Maximum number of processors used in parallel eigensolver. The number is machine dependent. Users could provide their own value for best performance.

Orbital Free DFT

Comments

The code will fail with the following options if OFDFT is activated and the related input options are listed.

- Polarized calculation: SPIN_TYP.
- K-point calculation: KPOINT_GRID, KPOINT_SHIFT.
- Dirichlet boundary condition in any direction: BC
- Define number of states/orbitals: NSTATES
- Define SCF related variables: TOL_SCF, CHEB_DEGREE
- CS and SQ3 method: CS_FLAG, SQ3_FLAG
- Hybrid functionals: EXCHANGE_CORRELATION
- Stress and pressure calculation: CALC_STRESS, CALC_PRES
- Print eigenvalues into file: PRINT_EIGEN

OFDFT_FLAG

Type
Integer

Default
0

Unit
No unit

Example
0

OFDFT_FLAG: 1

Description

Flag for Orbital-free DFT calculation.

TOL_OFDFT

Type
Double

Default
10⁻³

Unit
No unit

Example
TOL_OFDFT: 1E-6

Description

The tolerance on the normalized residual (l2-norm of residual/sqrt(N), N is number of nodes) for convergence of NLCG.

OFDFT_LAMBDA

Type
Double

Default
0.2

Unit
No unit

Example
0FDFT_LAMBDA: 0.3

Description

The coefficient in front of Laplacian operator is -lambda/2.

Socket communication in SPARC

Brief introduction

The socket communication layer allows SPARC to be controlled by a socket server that is compatible with i-PI protocol. The SPARC source code should be compiled with the USE_SOCKET=1 option.

To start a SPARC program with socket interface, use either option:

- Option 1: Specify socket address in command line
 - To start an INET socket, use
 - \$ mpirun -n 8 ./lib/sparc -socket localhost:12345 -name
 filename
 - To start a UNIX socket, use
 \$ mpirun -n 8 ./lib/sparc -socket /tmp/sparc.socket:unix -name filename
- Option 2: Provide .inpt parameters
 Visit the following pages for parameter specifications. Any parameter
 SPARC_* in the .inpt file will overwrite the settings from command line.

Output files

The following files will be generated in the socket mode and may slightly differ from standard SPARC outputs:

- .out: All the SCF steps will be written to .out file, similar to a relaxation / MD calculation.
- .static: It is concatenated from all single point steps. In addition to normal .static output, the lattice information are also recorded.

SOCKET_FLAG

Type
Integer

Default
0

Unit
No unit

Example
SOCKET_FLAG: 1

Description

Flag for starting the socket communication layer. It is equivalent to the -socket switch of command line.

Setting SOCKET_FLAG: 1 will disable MD_FLAG and RELAX_FLAG.

SOCKET_HOST

Type

String

Default

localhost

Unit

No unit

Example

SOCKET_HOST: 127.0.0.1

Description

Host name of the socket address that SPARC listens to. If it's an INET socket, it is the address of the interface. For a UNIX socket, it is the filename of the socket file (e.g. /tmp/sparc.socket).

SOCKET_PORT

Type

Integer

Default

None

Unit

No unit

Example

SOCKET_PORT: 12345

Description

When SPARC connects to an INET socket server, it is the port number. The SOCKET_PORT has no effect for a UNIX socket.

SOCKET_INET

Type
Integer

Default
None

Unit
No unit

Example
SOCKET_INET: 0

Description

1 for INET socket, 0 for UNIX socket. If no set in .inpt file, its value is determined by the -socket address:port command line switch.

SOCKET_MAX_NITER

Type

Integer

Unit

No unit

Default

10000

Example

SOCKET_MAX_NITER: 10000

Description

Maximum number of ionic SCF steps in the socket mode. As a socket client, SPARC will terminate after SOCKET_MAX_NITER steps are called.