minichem

Version 1.0

User Manual

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1 General considerations

minichem is a tiny quantum chemistry program written in educational purposes. It is based mainly on two textbooks:

- [1] A. Szabo, N. Ostlund, "Modern Quantim Chemistry";
- [2] T. Helgaker, P. Jorgensen, J. Olsen, "Molecular Electronic-Structure Theory".

The source code is written in the C99 programming language. Currently minichem is oriented on the Unix-like operation systems.

Features:

- single-point energy calculations;
- restricted (RHF) and unrestricted (UHF) Hartree-Fock theories;
- DIIS convergence technique;
- OpenMP parallelization;
- Available elements: all periodic table (but only non-relativistic treatment is available!);
- cartesian basis sets, arbitrary angular momentum of basis functions.

The source code is available on Github:

https://github.com/aoleynichenko/minichem

Authors is looking forward to any comments and questions! alexvoleynichenko@gmail.com.

2 Compilation and testing

Required libraries:

- libc the C standard library;
- MPI Message Passing Interface (higher-level routines for the distributed-memory communication environment);

- OpenMP Open Multi-Processing (API for shared memory multiprocessing programming);
- BLAS and LAPACK numerical linear algebra libraries.

Tools needed to compile the source code:

- CMake
- make
- C compiler (GNU or Intel are recommended)

How to compile:

```
$ mkdir build && cd build
$ cmake ..
$ make [-jN]
```

Test set can be found in the test folder. The testing system requires the Python2 programming language environment to be installed on your system. How to test:

```
$ cd test
$ python test.py
```

3 Getting started

How to run minichem:

```
$ minichem.x <input-file.inp>
```

Executable minichem.x sends the output to the stdout, hence it is more convenient to redirect it to file:

```
$ minichem.x <input-file.inp> | tee <output-file.out>
```

The input language design resembles the NWChem input files language [3]. As well as NWChem, minichem works as an interpreter: it executes input files written in a very simple scripting language. The minichem scripting language largely coincides with the NWChem language; wherever possible, the names of directives and sections have been chosen in that way that the input file can be interpreted by both quantum-chemical programs without some extra edits.

minichem input files consist of simple (top-level) directives and sections (compound directives). Directives are single-line commands; sections combine sets of directives. Usually sections specify control parameters for the separate modules (for example, convergence parameters of the SCF procedure). minichem reads the input file sequentially, line by line, and changes its internal variables according to the given values of the parameters specified in it. Once minichem reaches the task directive, the execution of the input file stops and calculation begins. Several task directives are allowed in one input file; between them the user can change any parameters, thus organizing a cascade of calculations.

The minichem language also supports single-line comments beginning with #. Language is case insensitive.

Let us consider the RHF/STO-3G calculation of a benzene molecule. Some explanations to all the directives encountered are given in the comments. Detailed descriptions of all possible directives and sections are given in Section 4. A few other examples can be found in Section 5.

```
# C6H6 single-point energy
# RHF molecular orbitals will be written to the molden-format
# file c6h6.mos
# name for the task
start C6H6
# allowed memory usage
memory 10 mb
# print input file before executing it
echo
# number of OpenMP threads
nproc 8
# geometry (cartesian)
# default: charge = 0
geometry
  C
       0.000
                 1.396
                           0.000
  С
       1.209
                 0.698
                           0.000
  C
       1.209
                -0.698
                           0.000
  С
       0.000
                -1.396
                           0.000
  C
                -0.698
      -1.209
                           0.000
  С
      -1.209
                 0.698
                           0.000
  Η
                 2.479
       0.000
                           0.000
                 1.240
  Η
       2.147
                           0.000
  Η
       2.147
                -1.240
                           0.000
                -2.479
                           0.000
  Η
       0.000
  Η
      -2.147
                -1.240
                           0.000
      -2.147
                 1.240
                           0.000
  Η
end
```

```
# basis set specification
# (keyword SPHERICAL -- only for compatibility with NWChem)
basis "ao basis" SPHERICAL
     3.42525091
                             0.15432897
                             0.53532814
      0.62391373
      0.16885540
                             0.44463454
C
     71.6168370
                           0.15432897
     13.0450960
                           0.53532814
                            0.44463454
     3.5305122
C
     2.9412494
                          -0.09996723
                            0.39951283
     0.6834831
     0.2222899
                             0.70011547
C
      2.9412494
                        0.15591627
      0.6834831
                           0.60768372
      0.2222899
                             0.39195739
end
# options for the SCF module
# by default: singlet
scf
 print "overlap"  # print AO overlap integrals
                 # enable DIIS, max subspace dim = 5
 diis 5
 maxiter 20 # max number of SCF iterations
end
# export calculated data (molecular orbitals)
 molden # to the MOLDEN .mos format
end
# do RHF calculation
task scf
```

4 Input files format

4.1 start

Syntax:

```
start <string name>
```

The start directive specifies a short task identifier (no whitespaces are allowed). This identifier is used in the names of some temporary and output files.

4.2 echo

Syntax:

echo

If the echo directive is specified in the input file, minichem redirects the contents of the input file to standard output. It is recommended always to use this directive.

4.3 memory

Syntax:

```
memory <integer> <string units>
# <units>: one of b (bytes), kb, mb, mw (megawords), gb
```

The maximum amount of RAM that can be allocated and used by the program.

4.4 nproc

Syntax:

```
nproc <integer>
```

Number of OpenMP threads (for parallel execution).

4.5 task

Syntax:

```
task <string theory>
```

The directive specifies which quantum chemical method is to be used to solve an electronic problem.

Since currently only the HF method is implemented, the task directive can be called only with the scf argument:

task scf

4.6 geometry

Syntax:

```
geometry [units <string units default angstroms>]
  [charge <integer default 0>]
  [mult <integer default 1>]
  <string elem> <real x y z>
    . . .
end
```

The geometry compound directive specifies the cartesian coordinates of the atoms, distance units (boron or angstrom), total system charge spin multiplicity. Given the charge and multiplicity values, the program can automatically choose which version of the Hartree-Fock method should be used.

4.7 charge

Syntax:

```
charge <integer>
```

The directive sets the total charge of the system (a.u.). Added for compatibility with NWChem.

4.8 basis

Syntax:

Gaussian basis set definition. Currently only cartesian basis sets are implemented (the cartesian keyword), so the spherical keyword is useful only for compatibility with NWChem.

The coefficients of the contracted basis functions are arranged in columns; the first column contains the exponents of primitive gaussian functions.

The angular momentum (<string shell_L>) of the block of basic functions is denoted, as usual, by the letters S, P, D ... The SP notation (one compressed function of type s, the second is of type p) is also allowed. Basis functions with arbitrary L are supported.

4.9 scf

Syntax:

```
[rhf | uhf]
[singlet | doublet | triplet | quartet | quintet]
[guess (core | eht)]
[direct | nodirect]
[maxiter <integer max_no_of_iterations default 50>]
[diis [<integer subspace_dim default 5>] | nodiis]
[print <string what>]
[noprint <string what>]
end
```

The scf compound directive is used to configure the SCF procedure. It may contain one or more directives from the following list:

guess MO initial guess. Possible values:

- core bare nuclei guess: all two-electron (electron-repulsion) integrals are neglected. Works well only for small systems with few electrons;
- eht (default) extended Hückel theory [4] (uses the Wolfsberg-Helmholtz formula [5]).

rhf/uhf Hartee-Fock theory type (rhf - restricted, uhf - unrestricted).

singlet/doublet/triplet/quartet/quintet spin multiplicity.

maxiter max number of SCF iterations. Default value: 50.

enable DIIS convergence technique [6, 7]. The optional argument of the directive is the maximum dimension of the iterative subspace that is used for extrapolation (default value: 5). DIIS can be recommended in almost all cases and is therefore included by default.

nodiis disable DIIS.

enable direct SCF method (two-electron AO integrals are recomputed on each iteration without storage on disk). Direct SCF can work an order of magnitude slower than the conventional one. May be useful in case of lack of disk space.

nodirect conventinal SCF – read precomputed two-electron AO integrals from disk (default).

print

what additional information should be printed (but not printed by default). The type of the argument is a string in double quotes. Possible arguments of the print directive are listed below:

"final vectors analysis"	MO expansion coefficients
"overlap"	AO overlap integrals
"kinetic"	AO kinetic energy integrals
"potential"	AO nuclear-attraction integrals
"eri"	AO two-electron (repulsion) integrals

noprint

disables printing of additional information. Arguments are the same as for print (see above).

4.10 out

Syntax:

```
out
  [molden]
end
```

Export data for further processing by other programs. May contain one or more directives from the following list:

molden

export information about the basis set and molecular orbitals to the MOLDEN format (.mos) [8, 9, 10]. At the moment, is implemented only for the RHF method.

5 Typical calculations

Li atom, UHF/STO-3G:

```
start Li
echo
geometry units atomic
 Li 0 0 0
basis "ao basis" SPHERICAL
Li
     16.1195750
                                0.15432897
                                0.53532814
      2.9362007
      0.7946505
                                0.44463454
Li
      SP
                               -0.09996723
                                                         0.15591627
      0.6362897
```

```
0.1478601 0.39951283 0.60768372 0.0480887 0.70011547 0.39195739 end scf uhf diis doublet end task scf
```

CH₃ radical, UHF/STO-3G

```
start CH3
echo
geometry
      0.08745162 -0.08744725 -0.08742186
H
      0.52503428
                    0.78909888
                                   -0.52508604
Η
     -0.78884450
                    -0.52530342
                                   -0.52536318
      0.52530257
Н
                    -0.52529218
                                   0.78892712
end
# STO-3G
basis "ao basis" SPHERICAL
      3.42525091
                            0.15432897
                             0.53532814
      0.62391373
     0.16885540
                            0.44463454
С
     71.6168370
                             0.15432897
     13.0450960
                             0.53532814
                             0.44463454
      3.5305122
С
      2.9412494
                           -0.09996723
      0.6834831
                            0.39951283
      0.2222899
                             0.70011547
С
      2.9412494
                             0.15591627
      0.6834831
                             0.60768372
      0.2222899
                             0.39195739
end
scf
  uhf
  doublet
end
```

task scf

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