Elevia Manual

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1 Input File Format

Elevia assumes a new input format:

```
[method] [basis] [the number of atoms] [charge] [spin multiplicity]

[atom symbol] [Rx] [Ry] [Rz]

[atom symbol] [Rx] [Ry] [Rz]

[atom symbol] [Rx] [Ry] [Rz]

... ...
```

For instance, an input file for test is

Listing 1: test.vie

```
RHF STO-3G 3 0 1
1
2
3
  0
        0.01984457
                     -0.19931037
                                     0.0000000
4
  Η
        0.97984454
                     -0.19906927
                                     0.0000000
5
       -0.30038273
                      0.70570591
                                     0.00000000
```

2 Method

Only support RHF.

3 Basis Set

Currently supported basis sets:

- STO-3G
- 3-21G
- 6-31G
- 6-31G(d)
- 6-31G(d,p)
- 6-311G(d,p)

Basis must strictly be typed in these displayed forms, only for that could Elevia recognize. The source code about reading basis data (dat/basis/*.g94) into the program is included in "intro_basis.cpp".

4 SCF Procedure

The SCF procedure is carried out as follows:

- 1. Specify a system (a set of nuclear coordinates $\{\mathbf{R}_a\}$, atomic numbers $\{Z_a\}$, and the total number of electrons N) and a basis set $\{\chi_s\}$.
- 2. Calculate all required molecular integrals S_{rs} , h_{rs} , and (rp|sq), (rp|qs).
- 3. Diagonalize the overlap matrix **S** and obtain a transformation matrix $\mathbf{X} = \mathbf{U}\mathbf{s}^{-1/2}$.
- 4. Assume an initial bond-order matrix P (often in the first iteration, we put P = 0, as if there were no electron repulsion).
- 5. Find the \mathcal{F} matrix using matrix **P**.
- 6. Calculate the transformed Fock matrix $\mathcal{F}' = \mathbf{X}^{\dagger} \mathcal{F} \mathbf{X}$.
- 7. Diagonalize \mathcal{F}' to obtain \mathbf{C}' and $\boldsymbol{\epsilon}$.
- 8. Calculate C = XC'.
- 9. Form a new **P** from **C**.
- 10. Calculate the total energy E.
- 11. Determine whether the procedure has converged, i.e., determine whether the difference between two successive total energy is less than a threshold (10⁻⁸). If the procedure has not converged, return to step (5) with the new bond-order matrix. If the procedure has converged, then calculate and output the quantities of interest.

References