

rhOver

User Manual v1.00

A FORTRAN Program to Determine Magnetic Anisotropy and Related Properties for Dysprosium(III) Single-Ion Magnets by Semi-Empirical Approaches utilizing Hartree–Fock Wave Functions

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- 1 Introduction to rhOver
- 1.1 What is rhOver?

1.2 License

rhover is released under the MIT license.

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1.3 Citation for rhOver

Please quote the usage of the rhover program and any scientific results in any form obtained with it by the following reference:

rhOver: Determination of Magnetic Anisotropy and Related Properties for Dysprosium(III)
 Single-Ion Magnets by Semi-Empirical Approaches utilizing Hartree–Fock Wave Functions. Michael Böhme, Winfried Plass, J. Comput. Chem. 2018, 39 (32), 2697–2712.

2 Installation Guide

rhover is distributed as FORTRAN source code and needs to be compiled in order to use the program.

Please note: Currently, rhover only supports MOLDEN files which have been generated by TURBOMOLE.¹ However, the support of other quantum computational software is planned for future releases of rhover. If you or your institution did not obtain a valid TURBOMOLE license, you can apply for a limited demo version of TURBOMOLE. Furthermore, it is recommended to install additional scalar-relativistic optimized basis sets (see section 2.4 Installation of Recommended Basis Sets for Turbomole).

2.1 Prerequisites

The following programs and software packages are required to compile rhover:

- a 64-bit FORTRAN compiler (with support for FORTRAN95 or later)
- GNU make
- BLAS development files and libraries
- LAPACK development files and libraries

rhover was successfully tested with the following FORTRAN compilers:

- GNU Fortran (gfortran)
- Intel ® Fortran Compiler (ifort)

Please use your package management software of your Linux distribution to install the necessary BLAS and LAPACK development files and libraries.

2.2 Building rhOver

```
tar xzf rhOver_v1.00.tar.gz
cd rhOver/
```

Table 1: Available targets for building rhover

TARGET	description
gfortran-omp	GNU gfortran with OpenMP parallelization
gfortran-serial	GNU gfortran
ifort-omp	Intel ® Fortran Compiler with OpenMP parallelization
ifort-serial	Intel ® Fortran Compiler

make -f Makefiles/TARGET

cp rhover \$HOME/bin/

2.3 Running rhOver

\$HOME/bin/rhover input.dy3 > output.log &

2.4 Installation of Recommended Basis Sets for Turbomole

For all calculations we strongly recommend the usage of the scalar-relativistic basis sets DZP-DKH/TZP-DKH² and SARC-DKH.³ These basis sets have been successfully tested in the original publication.⁴ Unfortunately, these basis sets are not redistributed in the default TURBOMOLE installation and thus, have to be installed manually. The usage of other basis sets than the tested ones especially smaller basis sets is not recommended.

Please note: For all calculations all-electron GTO basis sets must be used, since rhover does not support effective core potentials (ECPs).

The DZP-DKH/TZP-DKH² and SARC-DKH³ basis sets are provided within the distribution files of rhover and can be found in the DKH-BS subfolder. To install these basis sets for your TURBO-MOLE installation edit or create a file called ~/.definerc in your home folder. Add the following line stating the exact path to your rhover installation:

```
\verb|basis=/home/username/path_to_rhOver_installation/DKH-BS|
```

To check if the installation of the new basis set library was successfully, start TURBOMOLE's define, load a structure, and type 'lib' at the 'ATOMIC ATTRIBUTE DEFINITION MENU'. The new basis set library should be listened there as in this example:

```
AVAILABLE BASIS SET LIBRARIES:

#1 /share/apps/TURBOMOLE-7.2/basen/

#2 /share/apps/TURBOMOLE-7.2/basold/

#3 /home/michael/rhOver/DKH-BS/

INPUT NUMBER OF DESIRED LIBRARY (DEFAULT = 1)
```

Table 2: Recommended basis sets for rhover

Atom	Basis set	Ref.
Lu	SARC-DKH	3
Y	TZP-DKH	2
all donor atoms	TZP-DKH	2
non-donor atoms	DZP-DKH	2

- 3 User's Guide
- 3.1 Structure of a rhOver Input File
- 3.2 Basic Examples
- 3.3 Parallelization

rhover can take advantage of multi-processor CPUs and was programmed with OpenMP support. To use rhover with parallelization it is necessary to build it with OpenMP support (TARGET must have been set to gfortran-omp or ifort-omp; see section 2.2 Building rhover) By default the number of threads is set to the number of cores in your computer. The number of used threads can be adjusted by two different methods:

1. By setting/changing the following environment variable:

export OMP_NUM_THREADS=4

2. By adding the keyword PARA in the input file:

PARA 4

4 Advanced Examples

References

- [1] TURBOMOLE V7.2 2017, a development of University of Karlsruhe and Forschungszentrum Karlsruhe GmbH, 1989-2007, TURBOMOLE GmbH, since 2007; available from http://www.turbomole.com.
- [2] F. E. Jorge, A. C. Neto, G. G. Camiletti, S. F. Machado, J. Chem. Phys. 2009, 130, 064108.
- [3] D. A. Pantazis, F. Neese, J. Chem. Theory Comput. 2009, 5, 2229–2238.
- [4] M. Böhme, W. Plass, J. Comput. Chem. 2018, 39, 2697–2712.