

# ADT : A Generalized Algorithm and Program for Beyond Born-Oppenheimer Equations of ‘N’ Dimensional Sub-Hilbert Space

Koushik Naskar<sup>†,1</sup>, Soumya Mukherjee<sup>†,1</sup>, Bijit Mukherjee<sup>†</sup>, Satyam Ravi<sup>†</sup>, Saikat Mukherjee<sup>†</sup>,  
Subhankar Sardar<sup>†,‡</sup> and Satrajit Adhikari<sup>†,\*</sup>

<sup>†</sup>School of Chemical Sciences,

Indian Association for the Cultivation of Science, Jadavpur, Kolkata - 700032, India

<sup>‡</sup>Department of Chemistry, Bhatler College, Dantan, Paschim Medinipur - 721426, India

## USER MANUAL

---

\*Author for correspondence: e-mail: pcsa@iacs.res.in

<sup>1</sup>K. Naskar and S. Mukherjee contributed equally to this work.

# Contents

<b>1</b>	<b>Introduction</b>	<b>4</b>
<b>2</b>	<b>Files and folder structure</b>	<b>5</b>
<b>3</b>	<b>Prerequisites</b>	<b>7</b>
<b>4</b>	<b>Installation</b>	<b>7</b>
<b>5</b>	<b>Execution</b>	<b>8</b>
5.1	Analytic . . . . .	9
5.2	Numeric . . . . .	12
5.3	MOLPRO/ Gaussian/ Gamess: Optimization . . . . .	18
5.4	Molpro: Adiabatic PESs and NACTs . . . . .	20
<b>6</b>	<b>Instructions for Input Files of Numeric Section</b>	<b>23</b>
6.1	Input files for 2D calculation . . . . .	24
6.2	Input Files for 1D Calculation . . . . .	26
<b>7</b>	<b>Instructions for Input Files of optimization</b>	<b>27</b>
7.1	MOLPRO Configuration File . . . . .	27
7.2	Gaussian Configuration File . . . . .	29
7.3	Gamess Configuration File . . . . .	32
<b>8</b>	<b>Instructions for Input Files of Molpro Section for adiabatic PESs and NACTs</b>	<b>35</b>
8.1	MOLPRO Configuration File . . . . .	35
8.2	Atom Information File . . . . .	54
8.3	Equilibrium Geometry File . . . . .	54
8.4	Frequency File . . . . .	54
8.5	Wilson Matrix File . . . . .	55

## List of Tables

1	Possible values of <code>-anajob</code> and their significance . . . . .	10
2	Possible values of <code>-intpath</code> and their significance . . . . .	14
3	Sample input file for adiabatic PESs . . . . .	25
4	Sample input file for $\rho$ component of NACTs . . . . .	26
5	Sample input file for $\phi$ component of NACTs . . . . .	26
6	Description and possible inputs of MOLPRO configuration file ( <code>'mol.config'</code> ) . . . . .	28
7	Description of <code>'mol.config'</code> for NO <sub>2</sub> radical, NO <sub>3</sub> radical, C <sub>6</sub> H <sub>6</sub> and 1,3,5-C <sub>6</sub> H <sub>3</sub> F <sub>3</sub> for the test runs . . . . .	29
8	Description and possible inputs of Gaussian configuration file ( <code>'gauss.config'</code> ) . . . . .	30
9	Description of <code>'gauss.config'</code> for NO <sub>2</sub> radical, NO <sub>3</sub> radical, C <sub>6</sub> H <sub>6</sub> and 1,3,5-C <sub>6</sub> H <sub>3</sub> F <sub>3</sub> for the test runs . . . . .	32
10	Description and possible inputs of Gamess configuration file ( <code>'gms.config'</code> ) . . . . .	33
11	Description of <code>'gms.config'</code> for NO <sub>2</sub> radical, NO <sub>3</sub> radical, C <sub>6</sub> H <sub>6</sub> and 1,3,5-C <sub>6</sub> H <sub>3</sub> F <sub>3</sub> for the test runs . . . . .	35
12	Description and possible inputs of MOLPRO configuration file ( <code>'molpro.config'</code> ) for adiabatic PESs and NACTs . . . . .	51
13	Description of <code>'molpro.config'</code> for NO <sub>2</sub> , NO <sub>3</sub> , C <sub>6</sub> H <sub>6</sub> <sup>+</sup> and 1,3,5-C <sub>6</sub> H <sub>3</sub> F <sub>3</sub> <sup>+</sup> for the test runs . . . . .	52
14	Description of <code>'molpro.config'</code> for H <sub>3</sub> <sup>+</sup> and F+H <sub>2</sub> for the test runs . . . . .	53

## List of Figures

1	Files and folder structure of <code>'ADT-version0'</code> . . . . .	5
2	Files and folder structure of <code>'test_runs'</code> . . . . .	6
3	Working principles and command-line-arguments . . . . .	9
4	Jacobi coordinates of a triatomic system . . . . .	24

# 1 Introduction

Large and complicated functional form of adiabatic-to-diabatic transformation (ADT) equations is the major bottleneck of first principle based beyond Born-Oppenheimer (BBO) treatment, specially, when the number of coupled electronic states is high ( $N \geq 4$ ). Though explicit expressions for the ADT equations are already formulated for  $N = 3$  to 6 (See Ref. 24-28 of the main text), the algorithm ‘ADT’ generalizes this process of evaluating the expressions for any number of coupled electronic states ( $N$ ). This program package can be efficiently used to carry out the following operations:

- Formulation of analytic functional forms of ADT quantities, i.e., ADT matrix, ADT equations, coefficient matrix of gradient of ADT angles (see Eq. 14 of the main text), coefficient matrix of nonadiabatic coupling terms (NACTs) (see Eq. 14 of the main text) and diabatic potential energy matrix for any  $N$ .
- Numerical calculation of ADT quantities (ADT angles, residue of ADT angles, ADT matrices and diabatic potential energy matrices) for any number of coupled electronic states, if the input files [*ab initio* adiabatic Potential Energy Surfaces (PESs) and NACTs] are already available.
- Interfacing with standard *ab-initio* packages such as MOLPRO<sup>1</sup>, Gaussian<sup>2</sup> and Gamess<sup>3</sup> to obtain equilibrium geometry, frequency and wilson matrix for spectroscopic systems.
- Interfacing of MOLPRO quantum chemistry package<sup>1</sup> to compute *ab initio* data, namely, adiabatic PESs and NACTs, which will serve as the input for ADT calculation if the input files (adiabatic PESs and NACTs) are not available to the user.

Finally, the program package ‘ADT’ can be used for the construction of diabatic surfaces of spectroscopic as well as scattering processes. For spectroscopic systems, the nuclear configuration space (CS) is expressed in terms of normal mode coordinates. On the other hand, user can employ Jacobi or hyperspherical coordinates for scattering systems.

## 2 Files and folder structure

### Schematic Outline of Files present in 'ADT-version0' directory

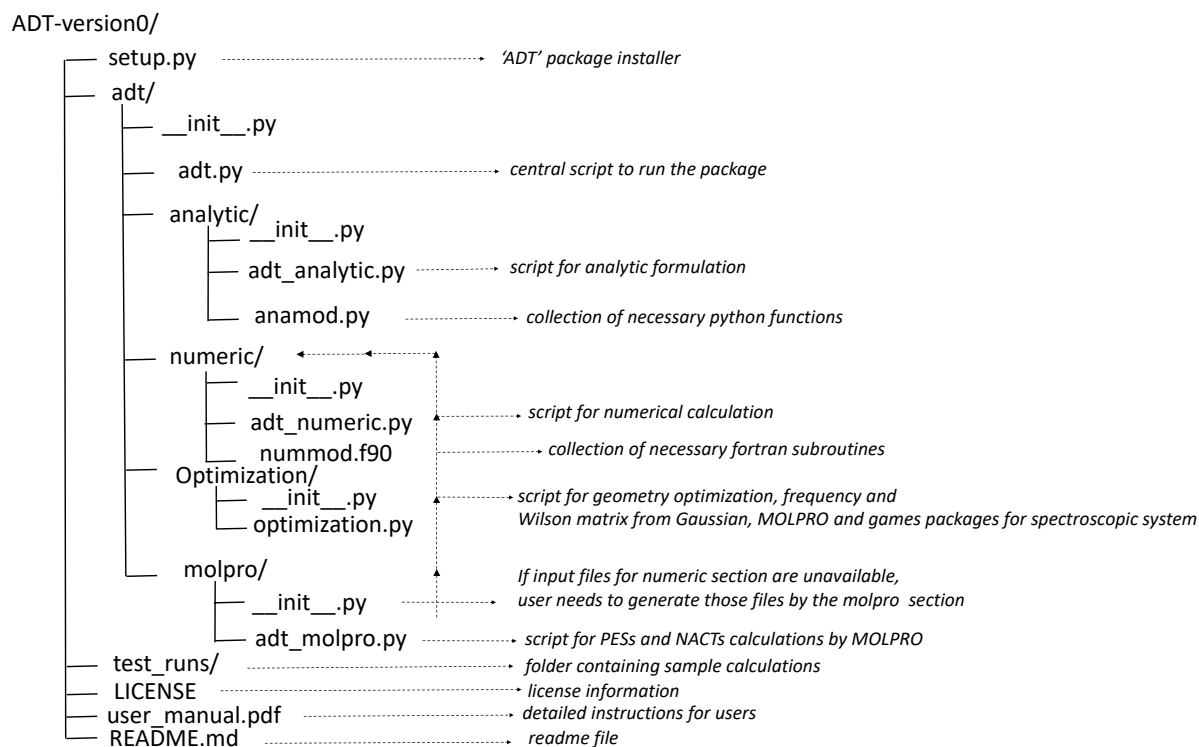


Figure 1: The above directory structure depicts the files and directories present in the parent folder, 'ADT-version0'.

## Schematic Outline of Files and Folders in 'test\_runs'

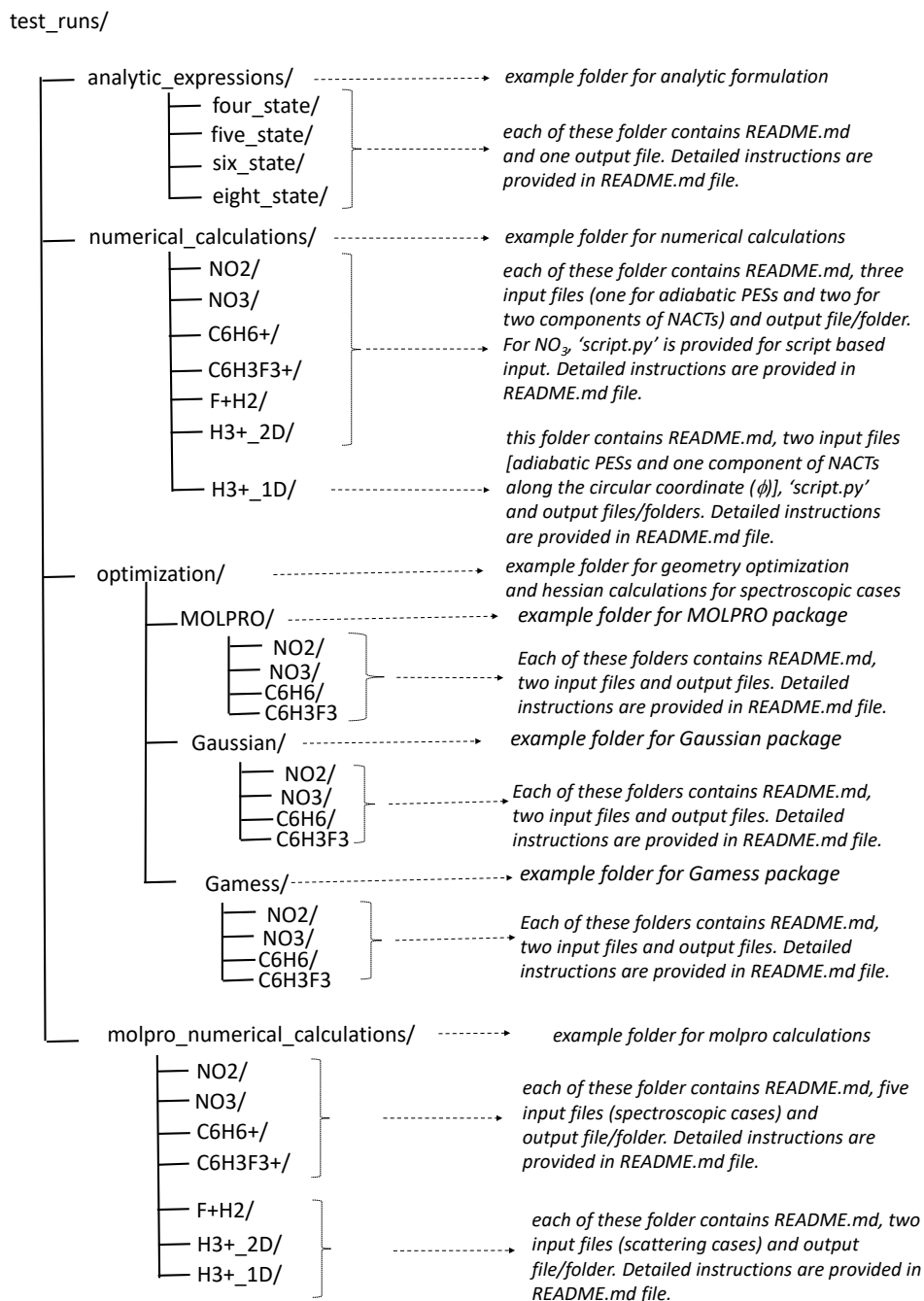


Figure 2: The above directory structure carries the information about the files and subdirectories inside the folder, 'test\_runs'.

### 3 Prerequisites

In order to install and execute the ‘ADT’ program package successfully, user needs a Fortran 90 compiler and Python (2.7 or 3.x), which are to be installed apriorly on the system. In addition, the following python dependencies are also required to use the package:

1. NumPy (v 1.13.0 or above)
2. setuptools
3. h5py

The first two modules are necessary for execution of this package, whereas the third one i.e. h5py is optional and is required only for HDF5 I/O. Alternatively, one can install these libraries through any python package manager (e.g., `pip`) or build them from source.

### 4 Installation

For installing the ‘ADT’ program package with default Fortran compiler for serial mode (no parallelization), user has to execute the following command in the terminal:

---

```
python setup.py install
```

---

After compilation of all the necessary files, the package is installed in the system and a command line executable `adt` is created. By default, the installation is performed in root of the system (usually in `/usr/local`) and therefore, root privilege may be required. If user is unable to access the root privilege or do not wish to install in the root of system, the ‘ADT’ package can be installed in a local package folder for python (usually in `~/local/`) by providing the `--user` flag during installation:

---

```
python setup.py install --user
```

---

Alternatively, if `pip` is available, user can install the package by executing the following command inside the source directory:

---

```
pip install .
```

---

On the contrary, for installing the package at any other location in the system, please visit the site [https://setuptools.readthedocs.io/en/latest/easy\\_install.html#custom-installation-locations](https://setuptools.readthedocs.io/en/latest/easy_install.html#custom-installation-locations).

In order to use the OpenMP parallelization scheme, user needs to link proper OpenMP libraries during the installation of the package by modifying the ‘fort\_args’ and ‘lib\_links’ variables in ‘setup.py’ script and execute it with corresponding Fortran compiler. As for example, uncomment the following lines in ‘setup.py’ to install ‘ADT’ using ‘gfortran’ compiler:

---

```
fort_args = ['-fopenmp']  
lib_links = ['-lgomp']
```

---

and execute the following command:

---

```
python setup.py config_fc --fcompiler=gnu95 install
```

---

whereas in order to install the package using ‘ifort’ compiler, uncomment the following lines in ‘setup.py’:

---

```
fort_args = ['-qopenmp']  
lib_links = ['-liomp5']
```

---

and execute the following command:

---

```
python setup.py config_fc --fcompiler=intelem install
```

---

During installation, the Fortran file is compiled to a python extension module by numpy distutils and to further configure the compilation process, user can visit the official NumPy distutils page (<https://docs.scipy.org/doc/numpy/reference/distutils.html#module-numpy.distutils>).

## 5 Execution

After installing the ‘ADT’ package, one has to execute the command line utility, **adt** to formulate analytic expressions or to perform numerical calculation or MOLPRO interfacing followed by numerical computation of ADT quantities employing the subcommands **ana**, **num** and **mol**, respectively. In order to perform geometry optimization and hessian calculations, user has to execute the subcommands **-opt** with any one of the packages’ names - **-mol** or **-gauss** or **-gamess**. The details



of the execution of these different sections are given in the following diagram:

## Working Principles and Command-line-arguments for Analytic, Numeric, geometry optimization and Molpro (adiabatic PESs and NACTs) Jobs

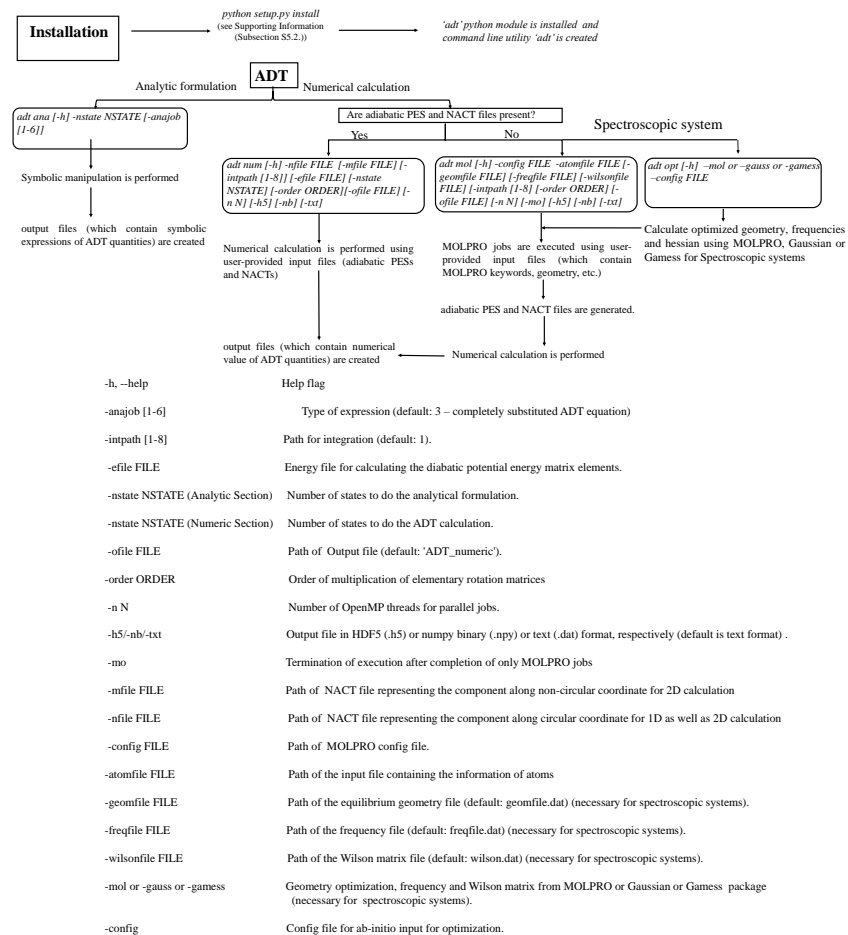


Figure 3: This diagram presents an overall schematic outline of the working principles for analytic expressions generation, numerical calculations for ADT, geometry optimization (only for spectroscopic cases) interfacing Molpro or Gaussian or Gamess and ab-initio calculations for adiabatic PESs and NACTs interfacing molpro. Detailed description of the arguments, `-anajob` and `-intpath` are presented in Tables 1 and 2.

### 5.1. Analytic

For analytic jobs, execute the following command line arguments,

---

```
adt ana [-h] -nstate NSTATE [-anajob [1-6]]
```

optional arguments:

-h, --help      show the help message  
 -anajob [1-6] Specify the type of expression (default: 3 -  
 completely substituted ADT equation)

Required arguments:

-nstate NSTATE Number of states

---

The possible values of **-anajob** and their significance are presented in the following table:

Table 1: This table depicts the correlation between the possible flags of the keyword argument, **-anajob** and the desired output

Keyword Argument (-anajob)	Analytical Quantities
1	elements of adiabatic to diabatic transformation (ADT) matrix
2	partially substituted forms of ADT equations [let say, for state=3, $\vec{\nabla}\theta_{12} = -\vec{\tau}_{12} - \sec\theta_{12}\tan\theta_{13}\vec{\tau}_{23}$ + $\tan\theta_{12}\tan\theta_{13}\vec{\nabla}\theta_{13}$ , ..]
3	complete forms of ADT equations (gradients in right hand side are replaced) [let say, for state=3, $\vec{\nabla}\theta_{12} = -\vec{\tau}_{12} - \sin\theta_{12}\tan\theta_{13}\vec{\tau}_{13}$ - $\cos\theta_{12}\tan\theta_{13}\vec{\tau}_{23}$ , ..]
4	elements of coefficient matrix of gradient of ADT angles (see Eq. 14 in the article for details)
5	elements of coefficient matrix of NACTs (see Eq. 14 in the article for details)
6	elements of diabatic potential energy matrix (detailed expressions)

## Examples:

1. Execute the following command to formulate completely substituted ADT equations for a five electronic state sub-Hilbert space:

---

```
adt ana -nstate 5 -anajob 3
```

---

The output file, ‘ADT\_EQUATIONS\_COMPLETE.DAT’ contains the analytic expressions of ADT equations.

2. Execute the following command to formulate diabatic potential energy matrix elements for a six electronic state sub-Hilbert space:

---

```
adt ana -nstate 6 -anajob 6
```

---

The output file is ‘W\_MATRIX.DAT’.

3. Execute the following command to formulate coefficient matrix of gradient of ADT angles (Eq. 14, described in the article) for a three electronic state sub-Hilbert space:

---

```
adt ana -nstate 3 -anajob 4
```

---

The output file is ‘GRADCOEFF.DAT’.

4. Execute the following command to formulate coefficient matrix of NACTs (Eq. 14, described in the article) for a eight electronic state sub-Hilbert space:

---

```
adt ana -nstate 8 -anajob 5
```

---

The output file is ‘TAUCOEFF.DAT’.

## 5.2. Numeric

User can carry out numerical calculations by using the subcommand `num`,

---

```
adt num [-h] -nfile FILE [-mfile FILE] [-intpath [1-8]]  
[-efile FILE] [-nstate NSTATE] [-order ORDER] [-ofile FILE]  
[-n N] [-h5] [-nb] [-txt]
```

optional arguments:

<code>-h, --help</code>	show the help message
<code>-mfile FILE</code>	Specify the path of the input NACT file representing the component of NACT for the non-circular coordinate (e.g. {0,pi/2} or {0,pi} or {0,infinity})
<code>-intpath [1-8]</code>	Specify the path for calculation (default: 1).
<code>-efile FILE</code>	Specify the path of the adiabatic PES file for calculating the diabatic potential energy matrix elements.
<code>-nstate NSTATE</code>	Specify the number of states to do the calculation. By default it includes all the data for calculation.
<code>-order ORDER</code>	Write the order of multiplication of the elementary rotation matrices.
<code>-ofile FILE</code>	Specify the path of the output file/folder (w/o extension) (default: ‘ADT_numeric’).

<code>-n N</code>	Specify the number of OpenMP threads for parallel jobs (default: 1).
<code>-h5</code>	Write results in a HDF5 file (.h5). Fast IO, smaller file size and hierarchical filesystem-like data format, preferable for saving and sharing large datasets in an organised way.
<code>-nb</code>	Write results in NumPy binary file (.npz). Preferable when working with NumPy for its much faster IO and easy portability.
<code>-txt</code>	Write results in a text file (default behaviour).

Required arguments:

<code>-nfile FILE</code>	Need to specify the path of the input NACT file, which is required both for 1D as well as 2D calculation. This file represents the component of NACT for the circular coordinate (e.g. {0,2pi}).
--------------------------	--

---

The probable values of `-intpath` and their significance are depicted in the following table:

Table 2: This table shows the relation between the possible flags of the keyword argument `-intpath` and the integration paths

Argument ( <code>-intpath</code> )	Path of Integration (R : first coordinate, P : second coordinate)
1	<ol style="list-style-type: none"> <li>1. At lowest value of R grid, the integration is performed for each positive step of P grid and the values of ADT angles are stored;</li> <li>2. With those values of the variables at each P grid, integration is carried out for each positive increment of R grid</li> </ol>
2	<ol style="list-style-type: none"> <li>1. At lowest value of R grid, the integration is performed for each negative step of P grid and the values of ADT angles are stored;</li> <li>2. With those values of the variables at each P grid, integration is carried out for each positive increment of R grid</li> </ol>
3	<ol style="list-style-type: none"> <li>1. At highest value of R grid, the integration is performed for each positive step of P grid and the values of ADT angles are stored;</li> <li>2. With those values of the variables at each P grid, integration is carried out for each negative increment of R grid</li> </ol>
4	<ol style="list-style-type: none"> <li>1. At highest value of R grid, the integration is performed for each negative step of P grid and the values of ADT angles are stored;</li> <li>2. With those values of the variables at each P grid, integration is carried out for each negative increment of R grid</li> </ol>
5	<ol style="list-style-type: none"> <li>1. At lowest value of P grid, the integration is performed for each positive step of R grid and the values of ADT angles are stored;</li> <li>2. With those values of the variables at each R grid, integration is carried out for each positive increment of P grid</li> </ol>
6	<ol style="list-style-type: none"> <li>1. At lowest value of P grid, the integration is performed for each negative step of R grid and the values of ADT angles are stored;</li> <li>2. With those values of the variables at each R grid, integration is carried out for each positive increment of P grid</li> </ol>
7	<ol style="list-style-type: none"> <li>1. At highest value of P grid, the integration is performed for each positive step of R grid and the values of ADT angles are stored;</li> <li>2. With those values of the variables at each R grid, integration is carried out for each negative increment of P grid</li> </ol>
8	<ol style="list-style-type: none"> <li>1. At highest value of P grid, the integration is performed for each negative step of R grid and the values of ADT angles are stored;</li> <li>2. With those values of the variables at each R grid, integration is carried out for each negative increment of P grid</li> </ol>

User can also employ the numerical segment of this package as a python module. While carrying out 1D and 2D calculation, one can use two functions, `adt1d` and `adt2d`, respectively, which take adiabatic PESs and NACTs as inputs, and return the ADT quantities (ADT angles, residue of ADT angles, ADT matrices and diabatic PESs).

## Examples

### 1. Using the command line executable `adt`

- (i) Execute the following command in terminal to carry out 2D ADT calculation, where ‘Tau\_Rho.dat’, ‘Tau\_Phi.dat’ and ‘Adiabatic\_PES.dat’ contain components of NACTs along non-circular coordinate, components of NACTs along circular coordinate and adiabatic PESs, respectively. The integration is done along path 6.

---

```
adt num -mfile Tau_Rho.dat -nfile Tau_Phi.dat
-efile Adiabatic_PES.dat -intpath 6
```

---

While running the command, ADT angles, residues of ADT angles, ADT matrices and diabatic potential energy matrices are computed and those are written in plain text files (default output format) inside a folder, ‘ADT\_numeric\_6’.

- (ii) Following command can be used to perform 2D ADT calculation, where ‘Tau\_Theta.npy’, ‘Tau\_Phi.npy’ and ‘Adiabatic\_PES.npy’ contain components of NACTs along non-circular coordinate, components of NACTs along circular coordinate and adiabatic PESs, respectively. The integration is done along path 4.

---

```
adt num -mfile Tau_Theta.npy -nfile Tau_Phi.npy
-efile Adiabatic_PES.npy -intpath 4 -nb
```

---

Output files will be saved in NumPy binary format inside a folder, ‘ADT\_numeric\_4’.

- (iii) Following command will carry out 2D ADT calculation, where ‘Tau\_Rho.h5’, ‘Tau\_Phi.h5’ and ‘Adiabatic\_PES.h5’ contain components of NACTs along non-circular coordinate, component of NACTs along circular coordinate and adiabatic PESs, respectively. The

integration is done along path 2.

---

```
adt num -mfile Tau_Rho.h5 -nfile Tau_Phi.h5  
-efile Adiabatic_PES.h5 -intpath 2 -h5
```

---

Output result will be saved as a HDF5 file, ‘ADT\_numeric.2.h5’ and the ADT quantities will be stored as different groups of dataset in that HDF5 file.

- (iv) If the ‘ADT’ package is installed with OpenMP enabled, user can employ `-n` flag to run the program in parallel threads. The following command executes the program package using 4 OpenMP threads.

---

```
adt num -mfile Tau_Rho.dat -nfile Tau_Phi.dat  
-efile Adiabatic_PES.dat -intpath 6 -n 4
```

---

The input and output file structures are similar as the serial ones.

- (v) If one wishes to perform 1D ADT, the following command can be executed with usual I/O files as the previous cases,

---

```
adt num -nfile Tau_Phi.dat -efile Adiabatic_PES.dat
```

---

The command will generate the ADT angle, residues of ADT angles, ADT matrices and diabatic potential energy matrices, and store the output in plain text files (default output format) inside a folder, ‘ADT\_numeric\_1D’.

- (vi) By default calculations are done with ADT matrix formed using the default order of multiplication of the elementary rotation matrices ( $A^{ij}(\Theta_{ij})$ ) where the matrices are multiplied in increasing order of the second index ( $j$ ) and then, for a fixed second index, in increasing order of the first index ( $i$ ). If user wants to use any other order of multiplication, the order can be provided using the `-order` keyword. For example, for a 3 state sub-Hilbert space to form the ADT matrix using the order ‘12,23,13’ i.e.

$$A(\Theta_{12}, \Theta_{23}, \Theta_{13}) = A^{12}(\Theta_{12}) \cdot A^{23}(\Theta_{23}) \cdot A^{13}(\Theta_{13})$$

execute the following command:



---

```
adt num -nfile Tau_Phi.dat -efile Adiabatic_PES.dat -order 12,23,13
```

---

The structure of input and output files are same as above.

## 2. Using python script

- (i) **1D:** The following sample python script shows the usage of the ‘ADT’ package as a python module to calculate 1D ADT. The files ‘Tau\_Phi.dat’ and ‘Adiabatic\_PES.dat’ are the NACT (along the circular coordinate) and adiabatic PES files. As for example, those two files can be taken from the ‘ADT\_version0/test\_runs /numerical\_calculation/H3+\_1D’.

---

```
import numpy as np
from adt.numeric import adt_numeric as an

nact = np.loadtxt('Tau_Phi.dat')
enr = np.loadtxt('Adiabatic_PES.dat')

angle, res, amat, db = an.adt1d(nact[:,0], nact[:,1:],
enr[:,1:])
```

---

- (ii) **2D:** This python script shows 2D calculation of ADT quantities along path 6, where the files ‘Tau\_Rho.npy’, ‘Tau\_Phi.npy’, ‘Adiabatic\_PES.npy’ contain components of NACTs along non-circular coordinate, components of NACTs along circular coordinate and adiabatic PESs, respectively. The input files contain *ab initio* data for five electronic manifold spanning  $50 \times 180$  grid points and can be taken from the ‘ADT\_version0/test\_runs /numerical\_calculation/NO3’.

---

```
import numpy as np
from adt.numeric import adt_numeric as an

nact1 = np.load('Tau_Rho.npy')
nact2 = np.load('Tau_Phi.npy')
enr = np.load('Adiabatic_PES.npy')
```

```

grid1 = np.unique(nact1[:,0])
grid2 = np.unique(nact1[:,1])

#If the number of grid points along the first and second
#coordinates are 50 and 180, respectively over 5
#electronic states with 10 NACT elements

enr = enr[:,2:].reshape(50, 180, 5)
nact1 = nact1[:,2:].reshape(50, 180, 10)
nact2 = nact2[:,2:].reshape(50, 180, 10)

#The integration is performed along path 6.

angle, res, amat,db = an.adt2d(grid1, grid2, nact1,
nact2, 6, enr)

```

---

### 5.3. MOLPRO/ Gaussian/ Gamess: Optimization

In spectroscopic cases, inputs files such as, equilibrium, frequency and wilson files can be directly obtained by optimization segment of ADT code, where geometry optimization and subsequent frequency calculations are performed by quantum chemistry packages, namely, MOLPRO, Gaussian and Gamess. The detailed discussion is given in Section 7 and then, the output files (`equigeom.dat`, `wilson.dat`, `frequency.dat`) can be further utilized for computing adiabatic PESs and NACTs in Molpro section of ADT package(see Figure 3).

---

```
adt opt [-h] -mol or -gauss or -gamess -config FILE
```

optional arguments:

```
-h, --help          show the help message
```

```
-mol                ab-initio calculations are performed by MOLPRO package
```

<code>-gauss</code>	ab-initio calculations are performed by Gaussian package
<code>-gamess</code>	ab-initio calculations are performed by Gamess package
<code>-config FILE</code>	the configuration files for respective quantum chemistry packages containing the necessary keywords of geometry optimization and hessian inputs.

---

For optimization, user has to supply two inputs files:

1. configuration file: contains `ab-initio` keywords for optimization (like memory, basis set, method of calculation, etc.), system information (initial geometry file name)
2. initial geometry file: contains initial geometry in XYZ format (in Å).

## Examples

1. By MOLPRO package, user can run following command in terminal

---

```
adt opt -mol -config mol.config
```

---

where, 'mol.config' is MOLPRO config file. This will execute optimization by Gaussian package to calculate the optimized geometry in 'XYZ' format, frequency and wilson file.

2. optimization by Gaussian package, one can run the following command in terminal,

---

```
adt opt -gauss -config gauss.config
```

---

where 'gauss.config' is the Gaussian configuration file.

3. Similarly, Gamess can be invoked for geometry optimization by executing following command

---

```
adt opt -gamess -config gms.config
```

---

where, gms.config is config file for `ab-initio` input of Gamess package.

## Limitations

- The ground state geometry optimization and hessian calculations are interfaced with only state-of-the-art single reference quantum chemistry methods such as HF, UHF, ROHF, MP2, UMP2, CCSD, CCSD(T), UCCSD, UCCSD(T) and DFT method.

#### 5.4. Molpro: Adiabatic PESs and NACTs

On the other hand, user can take the benefit of molpro section, where *ab initio* electronic structure calculations are performed according to the user provided input files (discussed in detail in Section 8) and then, the outputs (adiabatic PESs and NACTs) are used for subsequent ADT calculation (see Figure 3). User has to execute the following command,

---

```
adt mol [-h] -config FILE -atomfile FILE [-geomfile FILE]
[-freqfile FILE] [-wilsonfile FILE] [-intpath [1-8]] [-order ORDER]
[-ofile FILE] [-n N] [-mo] [-h5] [-nb] [-txt]
```

optional arguments:

```
-h, --help          show the help message

-geomfile FILE      Specify the geometry file containing the initial
                    grid point in 'xyz' format (in Angstrom)
                    (default: geomfile.dat).
                    (Ignore for scattering system).

-freqfile FILE      Specify the frequency information file, where
                    frequencies of normal modes are written in cm-1
                    (default: frequency.dat).
                    (Ignore for scattering system)

-wilsonfile FILE    Specify the filename containing the Wilson matrix
                    of a molecular species (default: wilson.dat).
                    (Ignore for scattering system).

-intpath [1-8]      Specify the path for calculation (default: 1).
```

-order ORDER      Write the order of multiplication of  
the elementary rotation matrices.

-ofile FILE        Specify the path of the output file/folder  
(w/o extension) (default: 'ADT\_numeric').

-n N                Specify the number of OpenMP threads for parallel  
jobs (default: 1).

-mo                Terminate the execution only after completion  
of MOLPRO jobs

-h5                Write results in a HDF5 file (.h5).  
Fast IO, smaller file size and  
hierarchical filesystem-like data format,  
preferable for saving and sharing large  
datasets in an organised way.

-nb                Write results in NumPy binary file (.npz).  
Preferable when working with NumPy for its  
much faster IO and easy portability.

-txt                Write results in a text file. (default  
behaviour)

#### Required arguments:

-config FILE        Specify the MOLPRO configuration file containing  
the necessary keywords of MOLPRO software.

-atomfile FILE      Specify the information file constituting atomic  
symbols and atomic masses.

---

For spectroscopic cases, user has to supply five input files:

1. MOLPRO configuration file: contains MOLPRO keywords (like memory, basis set, method of

calculation, etc.), system information (like symmetry, participating normal modes) and grid information

2. atom information file: contains atomic symbols and atomic masses
3. equilibrium geometry file: contains equilibrium geometry in xyz format (in Å)
4. frequency file: contains normal mode frequencies (in  $\text{cm}^{-1}$ )
5. wilson matrix file: contains mass-weighted second derivative matrix (Wilson matrix).

On the other hand, if user wishes to perform *ab initio* calculation of adiabatic PESs and NACTs for scattering systems, the first two files, i.e. MOLPRO configuration and atom information files are necessary and sufficient. The details of the above files are given in Section 8.

In order to store the geometry and MOLPRO output files for every successful and unsuccessful jobs, two folders, ‘CompleteJobs’ and ‘IncompleteJobs’ are generated, respectively during the execution of molpro section. One can use `-mo` argument to terminate the execution after completion of MOLPRO jobs to check the convergence and smoothness of adiabatic PESs and NACTs. Subsequently, user can remove/refine unphysical data points prior to ADT calculation (using the numeric section).

## Examples

1. For scattering case, one can run the following command in terminal,

---

```
adt mol -config molpro.config -atomfile atomfile.dat  
-intpath 6
```

---

where ‘molpro.config’ and ‘atomfile.dat’ are the MOLPRO configuration file and atom information file, respectively. This will execute MOLPRO to calculate the *ab initio* data (adiabatic PESs and NACTs) over the specified grid points defined in the configuration file and subsequently, carry out the ADT calculation along path 6.

2. In order to terminate the program after running the MOLPRO jobs (before calculating the ADT quantities), one can pass the `-mo` argument,

---

```
adt mol -config molpro.config -atomfile atomfile.dat -mo
```

---

3. For spectroscopic case, user has to run the following command in terminal,

---

```
adt mol -config molpro.config -atomfile atomfile.dat  
-geomfile geomfile.dat -freqfile frequency.dat  
-wilsonfile wilson.dat -intpath 4
```

---

where ‘geomfile.dat’, ‘frequency.dat’ and ‘wilson.dat’ are the equilibrium geometry file, frequency file and Wilson matrix file, respectively. Other two files, ‘molpro.config’ and ‘atomfile.dat’ are the same as discussed above.

## Limitations

- non-convergence of *ab initio* points originating from complete active space self-consistent field (CASSCF)/multi-reference configuration interaction (MRCI) methodology: The interpolation function in the python code cannot improve the output data if a large number of nuclear geometries are unrealistic.
- The swapping of *ab initio* computed NACTs due to the interchange of electronic states at CI point(s) are not considered in this software package due to molecule-specific characteristics.
- The *ab initio* adiabatic PESs can be computed only in `mrci` approach while calculating NACTs in finite difference method (`ddr`) (see Section 8.1).

## 6 Instructions for Input Files of Numeric Section

One can employ the numeric section of ‘ADT’ program package to solve the ADT equations along one dimensional as well as two dimensional contours of nuclear CS using adiabatic PES and NACT files. Those files can be provided either in HDF5 (.h5) or NumPy binary (.npz) or in text format. Detailed instructions related to the preparation of input files are given in the following paragraphs.

User can choose any one of the following coordinate systems to carry out ADT calculation:

- Spectroscopic systems:
  - (i) 2D ADT calculation: plane polar counterparts ( $\rho$  and  $\phi$ ) of a pair of normal modes ( $Q_i = \rho \cos \theta$  and  $Q_j = \rho \sin \theta$ )

(ii) 1D ADT calculation: circular coordinate,  $\phi$  for a fixed  $\rho$

- Scattering systems:

1. 2D ADT calculation:

- (a) hyperspherical coordinate ( $\rho$  and  $\phi$  for a fixed  $\theta$  or  $\theta$  and  $\phi$  for a fixed  $\rho$ )

- (b) Jacobi coordinate ( $q$  and  $\phi$ ) (see Figure 4)

2. 1D ADT calculation: circular coordinate,  $\phi$  for a fixed  $\rho$  and  $\theta$  (hyperspherical) or  $\phi$  for a fixed  $q$  (Jacobi).

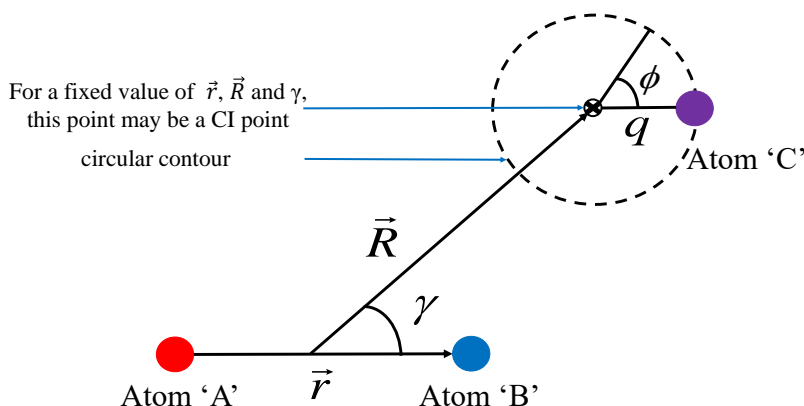


Figure 4: The above diagram briefly manifests the Jacobi coordinates of a triatomic system. The diatomic vector between Atom 'A' and Atom 'B' is represented as  $\vec{r}$  with distance  $r$ , the vector from diatomic center of mass to the third atom 'C' is symbolized as  $\vec{R}$  with distance  $R$ , and the angle between them is designated as  $\gamma$ . The circular contour is characterized by a radius ( $q$ ) and an angle ( $\phi$ ) along which *ab initio* calculation is performed. In this process, two Jacobi coordinates ( $R$  and  $\gamma$ ) are varying at a fixed value of  $r$  while performing such type of calculation.

### 6.1. Input files for 2D calculation

For the numeric section, three input files, namely, the adiabatic PES and components of NACTs are necessary.

- Input file for adiabatic PESs: The adiabatic PES input file must contain  $2 + N$  number of columns ( $N$  is number of coupled electronic states): first two columns represent grid points



of non-circular and circular coordinates, respectively and the rest of the columns contain the adiabatic PESs in increasing order of electronic states. User can supply scaled adiabatic energies for diabatization (scaled with respect to ground state minimum energy or asymptotic energy).

Example : The following table (Table 3) represents the arrangement of input file data points for adiabatic PESs of three electronic states ( $N = 3$ ).

Table 3: Sample input file for adiabatic PESs

First Coordinate	Second Coordinate	Adiabatic Energy (ground state)	Adiabatic Energy (first excited state)	Adiabatic Energy (second excited state)
0.01	0.000000	0.002	0.123	0.125
0.01	3.141592	0.003	0.125	0.126
0.01	6.283185	0.003	0.124	0.126
0.02	0.000000	-0.001	0.119	0.121
0.02	3.141592	-0.002	0.118	0.121
0.02	6.283185	-0.002	0.118	0.122

- Input files for NACTs: Two NACT files (**-mfile** and **-nfile**) are required containing the components of NACTs for the non-circular and circular coordinates, respectively. There should be  $2 + N * (N - 1)/2$  number of columns in each file, where the first two are non-circular and circular grids, respectively and the remaining ones signify one of the component of  $\tau$  values within  $N * (N - 1)/2$  pairs of electronic states. NACT components are represented as  $\tau_{\rho}^{ij}$  or  $\tau_{\theta}^{ij}$  or  $\tau_q^{ij}$  and  $\tau_{\phi}^{ij}$ . The columns of NACTs in the input files should be arranged in increasing order of first index ( $i$ ) for a fixed second index ( $j$ ) subjecting to the condition  $i < j$  and then, by increasing the second index.

Example : For four coupled electronic states, elements of  $\tau_{\rho}$  can be ordered as  $\tau_{\rho}^{12}$ ,  $\tau_{\rho}^{13}$ ,  $\tau_{\rho}^{23}$ ,  $\tau_{\rho}^{14}$ ,  $\tau_{\rho}^{24}$  and  $\tau_{\rho}^{34}$  as one of the input files (Table 4), where  $\phi$  component of NACT will be given as  $\tau_{\phi}^{12}$ ,  $\tau_{\phi}^{13}$ ,  $\tau_{\phi}^{23}$ ,  $\tau_{\phi}^{14}$ ,  $\tau_{\phi}^{24}$  and  $\tau_{\phi}^{34}$  for the second input file of NACT (Table 5).

Table 4: Sample input file for  $\rho$  component of NACTs

$\rho$	$\phi$	$\tau_{\rho}^{12}$	$\tau_{\rho}^{13}$	$\tau_{\rho}^{23}$	$\tau_{\rho}^{14}$	$\tau_{\rho}^{24}$	$\tau_{\rho}^{34}$
0.01	0.000000	0.002	0.123	0.125	0.020	0.103	0.105
0.01	3.141592	0.003	0.125	0.126	0.021	0.105	0.101
0.01	6.283185	0.003	0.124	0.126	0.023	0.107	0.100
0.02	0.000000	-0.001	0.119	0.121	0.019	0.109	0.100
0.02	3.141592	-0.002	0.118	0.121	0.023	0.108	0.101
0.02	6.283185	-0.002	0.118	0.122	0.025	0.108	0.102

Table 5: Sample input file for  $\phi$  component of NACTs

$\rho$	$\phi$	$\tau_{\phi}^{12}$	$\tau_{\phi}^{13}$	$\tau_{\phi}^{23}$	$\tau_{\phi}^{14}$	$\tau_{\phi}^{24}$	$\tau_{\phi}^{34}$
0.01	0.000000	0.005	0.103	0.150	0.020	0.130	0.121
0.01	3.141592	0.007	0.105	0.156	0.030	0.133	0.128
0.01	6.283185	0.009	0.107	0.156	0.030	0.137	0.130
0.02	0.000000	-0.006	0.105	0.155	0.021	0.133	0.121
0.02	3.141592	-0.009	0.107	0.159	0.027	0.137	0.125
0.02	6.283185	-0.011	0.108	0.160	0.029	0.139	0.127

## 6.2. Input Files for 1D Calculation

For 1D ADT calculation, one should supply two input files, namely, the adiabatic PESs and angular component of NACT ( $\tau_{\phi}$ ) forming closed contour. There must be  $1 + N$  and  $1 + N * (N - 1)/2$  number of columns ( $N$  is number of coupled electronic states) in the adiabatic PES and NACT input files, respectively. The first column contains the grid points of the circular coordinate and the rest of the columns are exactly similar as the 2D input files for ADT calculation.

## 7 Instructions for Input Files of optimization

In ADT, geometry optimization and hessian calculations are interfaced with three quantum chemistry packages, MOLPRO, Gaussian and Gamess. User has to provide two input files with the flags `-opt` and `-mol` or `-gauss` or `-gamess` (quantum chemistry package's name). The details of config and geometry files are depicted below:

### 7.1. MOLPRO Configuration File

The MOLPRO configuration file, namely, 'mol.config' is also a standard INI configuration file, which contains two sections, namely, `optInfo`, `gInfo` to specify all necessary keywords to run MOLPRO.

- `optInfo` contains the following seven keywords:
  - `memory`: allotted memory for MOLPRO job
  - `processor` (optional): number of processors used for parallelization (default is 1)
  - `method`: specify electronic structure method, namely any single reference method such as MP2, CCSD, CCSD(t) or its unrestricted versions and DFT method.
  - `basis`: basis set keyword compatible with MOLPRO input.
  - `spin`: spin multiplicity ( $2s$ ) of a molecular species
  - `charge`: charge of a molecular species
  - `symmetry`(optional): any Abelian point group of the molecular species during *ab initio* calculation (default value is `nosym`).
- `gInfo` contains the following keyword:
  - `file`: name of initial geometry file. File must contain the first column as atom symbol and second column as the Cartesian coordinates of the atoms (in Å) in a  $n \times 3$  matrix form for a  $n$  atomic molecule.

The above description of 'mol.config' is presented in Table 6.

Table 6: Description and possible inputs of MOLPRO configuration file ('mol.config')

Sections	Keywords	Description
optInfo	memory	memory required for the MOLPRO job, e.g., 10,m
	processor	number of processors required for parallelization of the job, e.g., 1
	method	electronic structure method for geometry optimization and hessian calculation e.g. UMP2, B3LYP, etc.
	basis	basis for geometry optimization, namely: cc-pVDZ or 6-31G**
	spin	spin multiplicity (2s) of ground state of molecular system , namely: 1 for NO <sub>3</sub> radical
	charge	charge of molecular system, namely: 1 for C <sub>6</sub> H <sub>3</sub> F <sub>3</sub> <sup>+</sup>
	symmetry	point group of molecular system (default value: nosym)
gInfo	file	name of initial geometry file e.g., geom.dat

### Examples:

1. The 'mol.config' file shows important inputs for NO<sub>3</sub> radical, where geometry optimization and hessian calculation are performed at UCCSD with basis set cc-pVDZ (see Table 7):

---

```

#Configuration file for the optimization and frequency calculation by MOLPRO job of NO3
radical

#Info for running MOLPRO
[optInfo]
memory = 50,m

#for parallel MOLPRO job
processor = 1

#Info for running optimization and hessian

method = uccsd
basis = cc-pvdz
spin   = 2
charge = 0

```

```
symmetry = nosym
```

```
#Info of the molecular species
```

```
[gInfo]
```

```
file = geom.dat
```

---

2. The format of initial geometry file ('geom.dat') is as follows:

---

```
#Format of geom.dat file for NO3 radical job
```

```
N      0.00000000    0.00000000    0.00000000
O      1.23680091    0.00000000    0.00000000
O      -0.61840019  -1.07110018    0.00000000
O      -0.61840019    1.07110018    0.00000000
```

---

Table 7: Description of 'mol.config' for NO<sub>2</sub> radical, NO<sub>3</sub> radical, C<sub>6</sub>H<sub>6</sub> and 1,3,5-C<sub>6</sub>H<sub>3</sub>F<sub>3</sub> for the test runs

Sections	Keywords	NO <sub>2</sub> radical	NO <sub>3</sub> radical	C <sub>6</sub> H <sub>6</sub>	1,3,5-C <sub>6</sub> H <sub>3</sub> F <sub>3</sub>
optInfo	memory	50,m			
	processor	2			
	method	UCCSD	UCCSD	B3LYP	MP2
	basis	cc-pvdz			
	spin	2		0	
	charge	0			
gInfo	file	geom.dat			

## 7.2. Gaussian Configuration File

The Gaussian configuration file, namely, 'gauss.config' is a standard INI configuration file, which contains two sections, namely, **optInfo**, **gInfo** to specify all necessary keywords to run Gaussian.

- **optInfo** contains the following eight keywords:
  - **version** (optional): version of Gaussian package (default is g16)
  - **memory**: allotted memory for Gaussian job

- **processor** (optional): number of processors used for parallelization (default is 1)
- **method**: specify electronic structure method, namely any single reference method such as MP2, CCSD, CCSD(t) or its unrestricted versions and DFT method
- **basis**: basis set keyword compatible with Gaussian input.
- **spin**: spin multiplicity ( $2s + 1$ ) of a molecular species
- **charge**: charge of a molecular species
- **symmetry**(optional): any Abelian point group of the molecular species during *ab initio* calculation (default value is **nosym**).

- **gInfo** contains the following keyword:

- **file**: name of initial geometry file. File must contain the first column as atom symbols and second column as Cartesian coordinates of the atoms (in Å) in a  $n \times 3$  matrix form for a  $n$  atomic molecule.

The above description of ‘gauss.config’ is presented in Table 8.

Table 8: Description and possible inputs of Gaussian configuration file (‘gauss.config’)

Sections	Keywords	Description
optInfo	<b>version</b>	version of Gaussian package, namely: <b>g16</b> or <b>g09</b>
	<b>memory</b>	memory required for the Gaussian job, e.g., <b>1GB</b>
	<b>processor</b>	number of processors required for parallelization of the job, e.g., <b>1</b>
	<b>method</b>	electronic structure method for geometry optimization and hessian calculation e.g. UMP2, B3LYP, etc.
	<b>basis</b>	basis for geometry optimization, namely: <b>cc-pVDZ</b> or <b>6-31G**</b>
	<b>spin</b>	spin multiplicity ( $2s + 1$ ) of ground state of molecular system , namely: <b>2</b> for NO <sub>2</sub> radical
	<b>charge</b>	charge of molecular system, namely: <b>1</b> for C <sub>6</sub> H <sub>6</sub> <sup>+</sup>
	<b>symmetry</b>	point group of molecular system (default value: <b>nosymm</b> )
<b>gInfo</b>	<b>file</b>	name of initial geometry file e.g., <b>geom.dat</b>

## Examples:

1. The following sample ‘gauss.config’ file presents necessary inputs for NO<sub>2</sub> radical, where geometry optimization and hessian calculation are performed at UCCSD with basis set cc-pVDZ (see Table 9):

---

```
#Configuration file for the optimization and frequency calculation by Gaussian job of NO2

#Info for running Gaussian
[optInfo]
version = g16
memory = 1GB

#for parallel Gaussian job
processor = 2

#Info for running optimization and hessian

method = uccsd
basis = cc-pvdz
spin   = 2
charge = 0
symmetry = nosymm

#Info of the molecular species
[gInfo]
file = geom.dat
```

---

2. The format of initial geometry file (‘geom.dat’) is as follows:

---

```
#Format of geom.dat file for NO2 radical job
N  0.000000000  0.000000000 -0.324503807
O  0.000000000  1.104583330  0.142043685
O  0.000000000 -1.104583330  0.142043685
```

---

Table 9: Description of ‘gauss.config’ for NO<sub>2</sub> radical, NO<sub>3</sub> radical, C<sub>6</sub>H<sub>6</sub> and 1,3,5-C<sub>6</sub>H<sub>3</sub>F<sub>3</sub> for the test runs

Sections	Keywords	NO <sub>2</sub> radical	NO <sub>3</sub> radical	C <sub>6</sub> H <sub>6</sub>	1,3,5-C <sub>6</sub> H <sub>3</sub> F <sub>3</sub>
optInfo	version	g16			
	memory	1GB			
	processor	2			
	method	UCCSD	UCCSD	B3LYP	MP2
	basis	cc-pvdz			
	spin	2			
	charge	0			
	symmetry	nosymm			
gInfo	file	geom.dat			

### 7.3. Gamess Configuration File

The Gamess configuration file, namely, ‘gms.config’ is a standard INI configuration file, which contains two sections, namely, **optInfo**, **gInfo** to specify all necessary keywords to run Gamess.

- **optInfo** contains the following nine keywords:
  - **path** (optional): Path of Gamess script (default is **rungms**)
  - **memddi**: allotted memory for **distributed data interface** (DDI) in Gamess run
  - **memory**: allotted memory for Gamess job
  - **processor** (optional): number of processors used for parallelization (default is 1)
  - **method**: specify electronic structure method, namely any single reference method such as MP2, CCSD, CCSD(t) or its unrestricted versions and DFT method
  - **basis**: basis set keyword compatible with Gamess input.
  - **spin**: spin multiplicity ( $2s + 1$ ) of a molecular species
  - **charge**: charge of a molecular species
  - **symmetry**(optional): any Abelian point group of the molecular species consistent with Gamess input (default value is **C<sub>1</sub>**).



- **gInfo** contains the following keyword:
  - **file**: name of initial geometry file. File must contain the first column as atom symbol and second column as the atomic number of atoms and third column as Cartesian coordinates of the atoms (in Å) in a  $n \times 3$  matrix form for a  $n$  atomic molecule.

The above description of ‘gms.config’ is presented in Table 10.

Table 10: Description and possible inputs of Gamess configuration file (‘gms.config’)

Sections	Keywords	Description
optInfo	<b>path</b>	path of Gamess script, e.g., /home/bijit/SATYAM/gamess/rungms
	<b>memddi</b>	memory required for DDI storage, e.g., 1
	<b>memory</b>	memory required for the Gamess job, e.g., 10
	<b>processor</b>	number of processors required for parallelization of the job, e.g., 1
	<b>method</b>	electronic structure method for geometry optimization and hessian calculation e.g. UMP2, B3LYP, etc.
	<b>basis</b>	basis for geometry optimization compatible with Gamess input, e.g., ccd or accd
	<b>spin</b>	spin multiplicity ( $2s + 1$ ) of ground state of molecular system , namely: 2 for NO <sub>2</sub>
	<b>charge</b>	charge of molecular system, namely: 1 for C <sub>6</sub> H <sub>3</sub> F <sub>3</sub> <sup>+</sup>
	<b>symmetry</b>	point group of molecular system in Gamess’s input, e.g., C1 or Cnv 2
gInfo	<b>file</b>	name of initial geometry file e.g., geom.dat

## Examples:

1. The ‘gms.config’ file shows important inputs for C<sub>6</sub>H<sub>6</sub> molecule, where geometry optimization and hessian calculation are performed at DFT/B3LYP method with basis set cc-pVDZ (see Table 11):

---

#Configuration file for the optimization and frequency calculation by Gamess job of C6H6

[optInfo]

# path of Gamess script

path = /home/bijit/SATYAM/gamess/rungms

```

# memory required to Gamess job
memddi = 30
memory = 50

#for parallel Gamess job
processor = 1

#Info for running optimization and hessian
method = b3lyp
basis = ccd
spin   = 1
charge = 0

[gInfo]
#Info for geometry file name
file = geom.dat

```

---

2. The format of initial geometry file ('geom.dat') is as follows:

---

```

#Format of geom.dat file for C6H6 job
C   6   1.3820899072   0.0000000000   0.0
C   6   0.6910449531  -1.1969249688   0.0
C   6  -0.6910449531  -1.1969249688   0.0
C   6  -1.3820899072   0.0000000000   0.0
C   6  -0.6910449531   1.1969249688   0.0
C   6   0.6910449531   1.1969249688   0.0
H   1   2.4568717910   0.0000000000   0.0
H   1   1.2284358943  -2.1277133837   0.0
H   1  -1.2284358943  -2.1277133837   0.0
H   1  -2.4568717910   0.0000000000   0.0
H   1  -1.2284358943   2.1277133837   0.0
H   1   1.2284358943   2.1277133837   0.0

```

---

Table 11: Description of ‘gms.config’ for NO<sub>2</sub> radical, NO<sub>3</sub> radical, C<sub>6</sub>H<sub>6</sub> and 1,3,5-C<sub>6</sub>H<sub>3</sub>F<sub>3</sub> for the test runs

Sections	Keywords	NO <sub>2</sub> radical	NO <sub>3</sub> radical	C <sub>6</sub> H <sub>6</sub>	1,3,5-C <sub>6</sub> H <sub>3</sub> F <sub>3</sub>
	path	/home/bijit/SATYAM/gamess/rungms			
optInfo	memddi	30	30	30	30
	memory	50	100	50	50
	processor	1			
	method	UCCSD	UCCSD	B3LYP	MP2
	basis	ccd			
	spin	2		1	
	charge	0			
gInfo	file	geom.dat			

## 8 Instructions for Input Files of Molpro Section for adiabatic PESs and NACTs

In order to interface MOLPRO by the software package, ‘ADT’ for calculation of energy and NACTs calculations, user has to provide five input files for spectroscopic problems and two input files for scattering cases. The MOLPRO configuration and the atom information files are common for both cases. The details of such files are depicted below:

### 8.1. MOLPRO Configuration File

The MOLPRO configuration file, namely, ‘molpro.config’ is a standard INI configuration file, which contains six sections, namely, `molInfo`, `sysInfo`, `eInfo`, `nInfo`, `mInfo` and `gInfo` to specify all necessary keywords to run MOLPRO.

- `molInfo` contains the following three keywords:
  - `screddir` (optional): path of scratch directory required during MOLPRO execution and storage of wavefunction (`.wfu`) files.

- **memory** (optional): allotted memory for MOLPRO job
- **processor** (optional): number of processors used for parallelization (default is 1)
- **sysInfo** contains the following keyword:
  - **type**: type of the system for electronic structure calculation (**spec** for spectroscopic systems in normal mode coordinates, **scat\_hyper** for scattering systems in hyperspherical coordinates and **scat\_jacobi** for scattering cases in Jacobi coordinates)
  - **symmetry** (optional): any Abelian point group of the molecular species during *ab initio* calculation (default value is **nosym**).
- **eInfo** contains the following keywords for adiabatic PESs calculation:
  - **method**: method of adiabatic PESs calculation, namely, **multi** or **mrci**
  - **basis**: basis set
  - **cas**: active space
  - **electron**: number of electrons
  - **spin**: spin multiplicity ( $2s$ ) of a molecular species
  - **charge**: charge of a molecular species
  - **state**: number of electronic states
  - **scale**(optional): nuclear geometry at which the ground adiabatic PES is set to zero.
    - \* spectroscopic cases:  $\rho, \phi$
    - \* scattering hyperspherical cases:  $\rho, \theta, \phi$
    - \* scattering Jacobi cases :  $r, R, \gamma$
  - **restricted** (optional): type of the initial hartree-fock calculation. **true** for **rhf** and **false** for **uhf**, default is **false**
  - **uhf\_extra** (optional): any extra keyword for initial Hartree-Fock calculation
  - **multi\_extra** (optional): extra keywords for **multi** calculation
  - **mrci\_extra** (optional): extra keywords for **mrci** calculation
- **nInfo** contains necessary keywords to calculate NACTs:

- **method**: method for NACT calculation, namely, **cpmcscf** or **ddr**
  - **basis**: basis set
  - **nact\_extra** (optional): extra keywords regarding the NACT calculation.
- **mInfo** (only for spectroscopic cases) contains the following keywords:
    - **varying**: indexes of the variable normal modes. The indexes are assigned to the normal modes in the ascending order of the corresponding frequencies provided in the frequency file. [not similar as the standard nomenclature, e.g., indexes of  $Q_{4x}$  and  $Q_{4y}$  normal modes (degenerate components of asymmetric bending mode) for  $\text{NO}_3$  radical will be 1,2 as the corresponding frequencies acquire the lowest values.]
  - **gInfo** contains the following keywords:
    - For normal mode coordinates:
      - **rho**:  $\rho_{min}, \rho_{max}, d\rho$ . The grid is defined from  $\rho_{min}$  to  $\rho_{max}$  in stepsize  $d\rho$ , where  $0 \leq \rho_{min} < \rho_{max}$  and  $d\rho > 0$ .  $\rho$  is expressed in mass-weighted dimensionless coordinates.
      - **phi**:  $\phi_{min}, \phi_{max}, d\phi$ . The grid is defined from  $\phi_{min}$  to  $\phi_{max}$  in stepsize  $d\phi$ , where  $\phi_{min} = 0, \phi_{max} = 2\pi$  and  $d\phi > 0$ .  $\phi$  is in degrees.
      - **drho (dphi)**: geometry increment in  $\rho$  ( $\phi$ ) coordinate used for finite difference DDR method for NACT calculation. It is not required for NACT calculation using coupled-perturbed multi-configuration self-consistent field (CP-MCSCF) method.
    - For hyperspherical coordinate :
      - **rho**:  $\rho_{min}, \rho_{max}, d\rho$ . The grid is defined from  $\rho_{min}$  to  $\rho_{max}$  in stepsize  $d\rho$ , where  $0 < \rho_{min} < \rho_{max}$  and  $d\rho > 0$ .  $\rho$  is expressed in atomic units.
      - **theta**:  $\theta_{min}, \theta_{max}, d\theta$ . The grid is defined from  $\theta_{min}$  to  $\theta_{max}$  in stepsize  $d\theta$ , where  $0 < \theta_{min} < \theta_{max}$  and  $d\theta > 0$ .  $\theta$  is in degrees.
      - **phi**:  $\phi_{min}, \phi_{max}, d\phi$ . The grid is defined from  $\phi_{min}$  to  $\phi_{max}$  in stepsize  $d\phi$ , where  $\phi_{min} = 0, \phi_{max} = 2\pi$  and  $d\phi > 0$ .  $\phi$  is in degrees.
      - **drho (dtheta/dphi)**: geometry increment in  $\rho$  ( $\theta/\phi$ ) coordinate used for finite difference DDR method for NACT calculation. It is not required for NACT calculation using CP-MCSCF method.

– For Jacobi coordinate:

- ▶ **small\_r**:  $r$  (diatom bond length) in atomic units.
- ▶ **capital\_r**:  $R$  (distance of the third atom from the center of the diatom) in atomic units.
- ▶ **gamma**:  $\gamma$  (angle between  $\vec{r}$  and  $\vec{R}$  in degrees).
- ▶ **q**:  $q_{min}, q_{max}, dq$ . The grid is defined from  $q_{min}$  to  $q_{max}$  in stepsize  $dq$ , where  $0 < q_{min} < q_{max}$  and  $dq > 0$ .  $q$  is expressed in atomic units. One can provide a single value of  $q$  to calculate 1D PECs and NACTs along the  $\phi$  grid.
- ▶ **phi**:  $\phi_{min}, \phi_{max}, d\phi$ . The grid is defined from  $\phi_{min}$  to  $\phi_{max}$  in stepsize  $d\phi$ , where  $\phi_{min} = 0, \phi_{max} = 2\pi$  and  $d\phi > 0$ .  $\phi$  is in degrees.
- ▶ **dq (dphi)**: geometry increment in  $q$  ( $\phi$ ) coordinate used for finite difference DDR method for NACT calculation. It is not required for NACT calculation using CP-MCSCF method.

The above illustration of ‘molpro.config’ is compactly presented in Table 12.

## Examples:

1. The following sample 'molpro.config' file presents necessary inputs for NO<sub>2</sub> radical, where adiabatic PESs are evaluated in CASSCF approach and analytic NACTs are calculated in CP-MCSCF method (see Table 13):

---

```
#Configuration file for the MOLPRO job of NO2

#Info for running molpro
[molInfo]
screedir = /tmp/adtpprogram
memory = 10,m

#for parallel molpro job
processor = 2

#Info of the molecular species
[sysInfo]
type = spec
#C_s point group (yz is plane of symmetry)
symmetry = x

#Info for energy calculation
[eInfo]
method = multi
basis = {spdf,n,cc-pvtz;spdf,o,cc-pvtz}
cas    = occ,11,3;closed,3,0
electron = 23
spin   = 1
charge = 0
state  = 2,0

#defining extra keyword required to run molpro for this system
#format should be same as in MOLPRO
multi_extra = maxiter,40
```

```

#defining scaling geometry for the adiabatic PESs
#ground state minima of NO2 w.r.t normal modes
scale = 0.0,0.0

#Info for NACT calculation
[nInfo]
method = cpmcscf
basis = {spdf,n,cc-pvtz;spdf,o,cc-pvtz}

#Info for varying modes
#put the indexes of the normal modes that will be varying
#The following two numbers represent Q_1 and Q_3 normal
#modes (bending and asymmetric stretching modes)
[mInfo]
varying = 1,3

#Put grid in format of <start>, <end>, <step>
#for spectroscopic put the 2D grid as rho and phi
[gInfo]
rho = 0.2, 12.0, 0.2
phi = 0, 360, 1
#Qx = rho*cos(phi) and Qy = rho*sin(phi)

```

---

2. The following sample ‘molpro.config’ file represents necessary inputs for  $\text{H}_3^+$  (1D PECs and NACTs calculation in Jacobi coordinate using  $C_s$  symmetry), where adiabatic energies are calculated in MRCI level of theory and analytic NACTs are evaluated in CP-MCSCF method (see Table 14):

---

```

#Configuration file for H3+ in Jacobi coordinates

#Info for running molpro
[molInfo]
screedir = /tmp/adtprogram
memory = 10,m

```



```

#for parallel molpro job
processor = 2

#Info of the molecular species
[sysInfo]
type = scat_jacobi
#C_s point group (yz is plane of symmetry)
symmetry = x

#Info for energy calculation
[eInfo]
method = mrci
basis = cc-pv5z
cas    = occ,10,1;closed,0,0;
electron = 2
spin = 0
charge = 1
state = 3,0

#Here scaling of PECs is not necessary as Jacobi coordinate
#is used only for locating the conical intersections (CIs)
#small_r = 1.4, capital_r = 10.0, gamma = 90.0
#minima:H2 at the asymptote of H2+H+
#scale = 1.4,10.0,90.0

#defining extra keyword required to run molpro for this system
#format should be same as in MOLPRO
restricted = true
uhf_extra = wf,2,1,0,1;
multi_extra = maxiter,40

#Info for NACT calculation
#method should be either ddr or cpmcscf
[nInfo]
method = cpmcscf

```

```

basis = 6-311++G**

#put a fixed value of small_r, capital_r, gamma and q
#and provide a grid of phi
[gInfo]
small_r = 2.0
capital_r = 1.732050808
gamma = 90
#q must be smaller than capital_r
q = 0.5
#Put grid in format of <start>, <end>, <step>
phi = 0, 360, 3

```

---

3. The following sample ‘molpro.config’ file depicts necessary inputs for  $\text{H}_3^+$  (2D ADT calculation in hyperspherical coordinate using  $C_s$  symmetry), where adiabatic PESs are evaluated in MRCI level of theory and NACTs are computed in DDR method (see Table 14):

---

```

#Configuration file for H3+ in hyperspherical coordinates

#Info for running molpro
[molInfo]
smdir = /tmp/adtprogram
memory = 10,m

#for parallel molpro job
processor = 2

#Info of the molecular species
[sysInfo]
type = scat_hyper
#C_s point group (yz is plane of symmetry)
symmetry = x

#Info for energy calculation
[eInfo]

```

```

method = mrci
basis = cc-pv5z
cas    = occ,10,1;closed,0,0
electron = 2
spin = 0
charge = 1
state = 3,0

#defining extra keyword required to run molpro for this system
#format should be same as in MOLPRO
restricted = true
uhf_extra = wf,2,1,0,1;
multi_extra = maxiter,40;ACCURACY,GRADIENT=1.d-4,
ENERGY=1.d-6,STEP=1.d-2

#defining scaling geometry for the adiabatic PESs
#ground state minima of H3+
#rho = 1.65, theta = 0.0, phi = 0.0 (D3h point)
scale = 1.65,0.0,0.0

#Info for NACT calculation
#method should be either ddr or cpmcscf
[nInfo]
method = ddr
basis = cc-pv5z

#Put grid in format of <start>, <end>, <step>
#for scattering put theta, phi as the 2d grid, with a given
#fixed value for rho
[gInfo]
rho  = 4.5
theta = 3,90,3
phi = 0, 360, 3

#following options are only required for ddr nact calculation
dtheta = 0.03

```

dphi = 0.03

---

4. The following sample ‘molpro.config’ file contains necessary MOLPRO keywords and other inputs for F+H<sub>2</sub> (2D ADT calculation in hyperspherical coordinate), where adiabatic PESs are generated in MRCI level of theory and analytic NACTs are calculated using CP-MCSCF method (see Table 14):
- 

```
#Configuration file for F+H2 in hyperspherical coordinates

#Info for running molpro
[molInfo]
screedir = /tmp/adtprogram
memory = 10,m

#for parallel molpro job
processor = 2

#Info of the molecular species
[sysInfo]
type = scat_hyper

#Info for energy calculation
[eInfo]
method = mrci
basis = aug-cc-pvtz
cas = occ,10;closed,2;core,2
electron = 11
spin = 1
charge = 0
state = 3

#defining extra keyword required to run molpro for this system
#format should be same as in MOLPRO
multi_extra = maxiter,40
mrci_extra = maxiter,200,200
```

```

#defining scaling geometry for the adiabatic PESs
#minima of H2 at assymptotic F+H2 entrance channel
#rho = 25.0, theta = 90.0, phi = 175.4

scale = 25.0,90.0,175.4

#Info for NACT calculation
#method should be either ddr or cpmcscf
[nInfo]
method = cpmcscf
basis = 6-311++G**

#Put grid in format of <start>, <end>, <step>
#for scattering put theta, phi as the 2d grid, with a given
#fixed value for rho
[gInfo]
rho = 5.0
theta = 0,90,5
phi = 0,360,4

```

---

5. The following sample ‘molpro.config’ file depicts necessary *ab initio* information for NO<sub>3</sub> radical, where adiabatic PESs are evaluated in CASSCF level of theory and analytic NACTs are calculated in CP-MCSCF method (see Table 13):
- 

```

#Configuration file for the MOLPRO job of NO3

#Info for running MOLPRO
[molInfo]
screedir = /tmp/adtprogram
memory = 10,m

#for parallel MOLPRO job
processor = 2

```

```

#Info of the molecular species
[sysInfo]
type = spec

#If symmetry is not specified, by default it will take 'nosym'
#C_s point group (yz is plane of symmetry)
#symmetry = x

#Info for energy calculation
[eInfo]
method = multi
basis = 6-31G**
cas = occ,19;closed,11
electron = 31
spin = 1
charge = 0
state = 5

#defining extra keyword required to run molpro for this system
#format should be same as in MOLPRO
uhf_extra = accu,5
multi_extra = maxiter,40

#defining scaling geometry for the adiabatic PESs
#rho = 0.0, phi = 0.0
#ground state minima of NO3 PES w.r.t normal modes
scale = 0.0,0.0

#Info for NACT calculation
[nInfo]
method = cpmcscf
basis = 6-31G**
nact_extra = accu=1.d-10

#Info for varying modes
#put the indexes of the normal modes that will be varying

```

```

#The following two numbers represent the lowest frequency
#modes, but actually they are Q_4x and Q_4y normal modes
#(degenerate asymmetric bending modes)
[mInfo]
varying = 1,2

#Put grid in format of <start>, <end>, <step>
#for spectroscopic put the 2D grid as rho and phi
[gInfo]
rho = 0.1, 0.3, 0.1
phi = 0, 360, 3
#Qx = rho*cos(phi) and Qy = rho*sin(phi)

```

---

6. The following sample ‘molpro.config’ file represents necessary inputs for  $\text{C}_6\text{H}_6^+$  radical cation, where adiabatic PESs are computed in CASSCF level of theory and analytic NACTs are evaluated in CP-MCSCF method (see Table 13):

---

```

#Configuration file for the MOLPRO job of C6H6+

#Info for running molpro
[molInfo]
smdir = /tmp/adtpprogram
memory = 100,m

#for parallel MOLPRO job
processor = 1

#Info of the molecular species
[sysInfo]
type = spec

#Info for energy calculation
[eInfo]
method = multi

```

```

basis = 6-31G**
cas    = occ,21;closed,6
electron = 41
spin = 1
charge = 1
state = 5

#defining extra keyword required to run molpro for this system
#format should be same as in MOLPRO
uhf_extra = wf,41,1,1,1
multi_extra = maxiter,40;

#defining scaling geometry for the adiabatic PESs
#ground state minima of C6H6+ w.r.t normal modes
scale = 0.0,0.0

#Info for NACT calculation
[nInfo]
method = cpmcscf
basis = 6-31G**
nact_extra = accu=1.d-10

#Info for varying modes
#put the indexes of the normal modes that will be varying
#The following two numbers represent Q_16x and Q_16y normal
#modes (degenerate asymmetric stretching modes)
[mInfo]
varying = 23,24

#Put grid in format of <start>, <end>, <step>
#for spectroscopic put the 2D grid as rho and phi
[gInfo]
rho = 0.1, 0.3, .1
phi = 0, 360, 3
#Qx = rho*cos(phi) and Qy = rho*sin(phi)

```

---



7. The following sample ‘molpro.config’ file depicts relevant inputs for 1,3,5-C<sub>6</sub>H<sub>3</sub>F<sub>3</sub><sup>+</sup> radical cation, where adiabatic PESs are computed in CASSCF level of theory and analytic NACTs are evaluated in CP-MCSCF method (see Table 13):

---

```
#Configuration file for the MOLPRO job of 1,3,5-C6H3F3+

#Info for running molpro
[molInfo]
screedir = /tmp/adtpprogram
memory = 100,m

#for parallel molpro job
processor = 1

#Info of the molecular species
[sysInfo]
type = spec

#Info for energy calculation
[eInfo]
method = multi
basis = cc-pvdz
cas    = occ,36;closed,27
electron = 65
spin = 1
charge = 1
state = 6

#defining extra keyword required to run molpro for this system
#format should be same as in MOLPRO
uhf_extra = wf,65,1,1,1;accu,5
multi_extra = maxiter,40

#defining scaling geometry for the adiabatic PESs
#ground state minima of 1,3,5-C6H3F3+ w.r.t normal modes
scale = 0.0,0.0
```

```

#Info for NACT calculation
[nInfo]
method = cpmcscf
basis = 6-31G**

#Info for varying modes
#put the indexes of the normal modes that will be varying
#The following two numbers represent Q_12x and Q_12y normal
#modes (degenerate C-C-C scissoring modes)
[mInfo]
varying = 16,17

#Put grid in format of <start>, <end>, <step>
#for spectroscopic put the 2D grid as rho and phi
[gInfo]
rho = 0.02, 0.06, 0.02
phi = 0, 360, 3
#Qx = rho*cos(phi) and Qy = rho*sin(phi)

```

---

Table 12: Description and possible inputs of MOLPRO configuration file ('molpro.config') for adiabatic PESs and NACTs

Sections	Keywords	Description
molInfo	screadir	path of scratch directory for wavefunction file, namely: /tmp/adtprogram
	memory	memory required for the job, e.g., 10,m
	processor	number of processors required for parallelization of the job, e.g., 2
sysInfo	type	spec or scat.hyper or scat.jacobi
	symmetry	symmetry of the molecule (default value: nosym) optional, e.g., x for H <sub>3</sub> <sup>+</sup> (C <sub>s</sub> point group: yz is the plane of symmetry)
eInfo	method	potential energy surface: multi or mrci
	basis	basis for energy calculation, namely: 6-31G** for NO <sub>3</sub>
	cas	complete active space, e.g., occ,19;closed,11 for NO <sub>3</sub>
	electron	number of electrons, namely: 31 for NO <sub>3</sub>
	spin	spin multiplicity (2s), namely: 1 for NO <sub>3</sub>
	charge	charge of a molecular species, namely: 1 for H <sub>3</sub> <sup>+</sup>
	state	number of electronic states, e.g., 5 for NO <sub>3</sub> , 3,0 for H <sub>3</sub> <sup>+</sup> (two values due to C <sub>s</sub> symmetry)
	scale	spectroscopic: Q <sub>i</sub> and Q <sub>j</sub> ⇒ ρ and ϕ, namely, 0.0,0.0 for NO <sub>3</sub> scattering (hyperspherical): ρ, θ and ϕ, namely, 25.0,90.0,175.4 for F+H <sub>2</sub> scattering (Jacobi): r, R and γ, namely, 1.0,24.0,0.0 for F+H <sub>2</sub>
	restricted	true for rhf and false for uhf (default is false)
	uhf_extra	extra specification for uhf, namely: wf,2,1,0,1; for H <sub>3</sub> <sup>+</sup>
	multi_extra	extra specification for multi jobs, namely: maxiter,40 for NO <sub>3</sub>
	mrci_extra	extra specification for mrci jobs, namely: maxiter,200,200 for F+H <sub>2</sub>
nInfo	method	NACT calculation: cpmcscf for analytic gradient and ddr for numerical finite difference method.
	basis	basis set for NACT, e.g., 6-31G** for NO <sub>3</sub>
	nact_extra	extra specification for NACT jobs, namely: accu=1.d-10 for NO <sub>3</sub>
mInfo	varying	specification for pairwise normal modes (Q <sub>i</sub> , Q <sub>j</sub> ), e.g., 1,2 (only required for spectroscopic calculation)
gInfo	rho	ρ grid for spectroscopic or scattering calculation in {start, end, step} format, namely: 0.1, 0.3, 0.1 or 1.0, 20.0, 0.25, respectively
	theta	θ grid (hyperspherical) for scattering calculation, e.g., 3, 90, 3
	phi	ϕ grid for spectroscopic/scattering calculation, e.g., 0, 360, 3
	small_r	diatomic bond length, e.g, 2.0
	capital_r	distance of third atom from diatomic center of mass, e.g, 1.73205
	gamma	angle between small_r and capital_r, e.g., 90.0
	q	radius of the circular path (see Figure 4) e.g., 0.5
	drho*	ρ spacing in ddr method, namely: 0.01
	dtheta*	θ spacing in ddr method, namely: 0.03
	dphi*	ϕ spacing in ddr method, namely: 0.03
	dq*	q spacing in ddr method, namely: 0.02

\*The values of drho, dtheta, dphi and dq require adjustment based on the convergence of the calculated result.

Table 13: Description of ‘molpro.config’ for NO<sub>2</sub>, NO<sub>3</sub>, C<sub>6</sub>H<sub>6</sub><sup>+</sup> and 1,3,5-C<sub>6</sub>H<sub>3</sub>F<sub>3</sub><sup>+</sup> for the test runs

Sections	Keywords	NO <sub>2</sub>	NO <sub>3</sub>	C <sub>6</sub> H <sub>6</sub> <sup>+</sup>	1,3,5-C <sub>6</sub> H <sub>3</sub> F <sub>3</sub> <sup>+</sup>
molInfo	screedir	/tmp/adtpprogram			
	memory	10,m		100,m	
	processor	2		1	1
sysInfo	type	spec			
	symmetry	x	—	—	—
eInfo	method	multi	multi	multi	multi
	basis	spdf,n,cc-pvtz; spdf,o,cc-pvtz	6-31G**	6-31G**	cc-pvdz
	cas	occ,11,3; closed,3,0;	occ,19; closed,11	occ,21; closed,6	occ,36; closed,27
	electron	23	31	41	65
	spin	1			
	charge	0	0	1	1
	state	2,0	5	5	6
	scale	0.0,0.0			
	uhf_extra	—	accu,5	wf,41,1,1,1	wf,65,1,1,1;accu,5
	multi_extra	maxiter,40			
nInfo	method	cpmcscf			
	basis	spdf,n,cc-pvtz; spdf,o,cc-pvtz	6-31G**	6-31G**	6-31G**
	nact_extra	—	accu=1.d-10	accu=1.d-10	—
mInfo	varying	1,3	1,2	23,24	16,17
gInfo	rho	0.2,12.0,0.2	0.1,0.3,0.1	0.1,0.3,0.1	0.02,0.06,0.02
	phi	0, 360, 1	0, 360, 3		

Table 14: Description of ‘molpro.config’ for  $\text{H}_3^+$  and  $\text{F}+\text{H}_2$  for the test runs

Sections	Keywords	$\text{H}_3^+$ (1D calculation)	$\text{H}_3^+$ (2D calculation)	$\text{F}+\text{H}_2$
molInfo	scrdir	/tmp/adtprogram		
	memory	10,m		
	processor	2	2	2
sysInfo	type	scat_jacobi	scat_hyper	
	symmetry	x		—
eInfo	method	mrci		
	basis	cc-pv5z		aug-cc-pvtz
	cas	occ,10,1;closed,0,0		occ,10;closed,2; core,2
	electron	2		11
	spin	0		1
	charge	1		0
	state	3,0		3
	scale	—	1.65,0.0,0.0	25.0,90.0,175.4
	restricted	true	true	—
	uhf_extra	wf,2,1,0,1;		—
	multi_extra	maxiter,40	maxiter,40;ACCURACY, GRADIENT=1.d-4, ENERGY=1.d-6, STEP=1.d-2	maxiter,40
	mrci_extra	—	—	maxiter,200,200
nInfo	method	cpmcsf	ddr	cpmcsf
	basis	6-311++G**	cc-pv5z	6-311++G**
gInfo	rho	—	4.5	5.0
	theta	—	3, 90, 3	0, 90, 5
	phi	0, 360, 3		0, 360, 4
	small_r	2.0	—	—
	capital_r	1.73205	—	—
	gamma	90.0	—	—
	q	0.5	—	—
	dtheta	—	0.03	—
	dphi	—	0.03	—

## 8.2. Atom Information File

- necessary file both for spectroscopic and scattering calculations
- First column: atomic symbols
- Second column: atomic masses in amu

**Example:** Sample file for NO<sub>3</sub> radical, ‘atomfile.dat’:

---

N	14.006700
O	15.999400
O	15.999400
O	15.999400

---

## 8.3. Equilibrium Geometry File

- necessary file for spectroscopic calculations
- contains the equilibrium geometry of the system
- Cartesian coordinates of the constituent atoms (in Angstroms) must be in a  $n \times 3$  matrix form for a  $n$  atomic molecule
- order of atoms must be same as in atom information file

**Example:** Sample file for NO<sub>3</sub> radical, ‘geomfile.dat’:

---

0.00000000	0.00000000	0.00000000
1.23680091	0.00000000	0.00000000
-0.61840019	-1.07110018	0.00000000
-0.61840019	1.07110018	0.00000000

---

## 8.4. Frequency File

- necessary file for spectroscopic calculations

- contains frequencies of normal modes (in  $\text{cm}^{-1}$ ) of a species arranged in ascending order (in a single row)

**Example:** Sample file for  $\text{NO}_3$  radical, ‘frequency.dat’:

---

476.88	476.91	745.13	1047.52	1185.60	1185.85
--------	--------	--------	---------	---------	---------

---

## 8.5. Wilson Matrix File

- necessary file for spectroscopic systems
- provides the Wilson matrix of a molecular species
- the dimension of Wilson matrix will be  $3n \times 3n - 5$  for linear molecules (vibrational DOFs:  $3n - 5$ ) and  $3n \times 3n - 6$  for non-linear ones (vibrational DOFs:  $3n - 6$ )
- columns are written as the increasing order of normal mode frequencies
- rows are sorted as per the order of atoms in atom information file

**Example:** Sample file for  $\text{NO}_3$  radical, ‘wilson.dat’:

---

0.4681243	0.0000000	0.0000000	0.0000102	0.0000000	0.7449584
0.0000000	-0.4682375	0.0000000	0.0000000	0.7448872	0.0000000
0.0000000	0.0000000	0.8798314	0.0000000	0.0000000	0.0000000
0.3427462	0.0000000	0.0000000	0.5774593	0.0000000	-0.5394732
0.0000000	0.6348331	0.0000000	0.0000000	0.0749380	0.0000000
0.0000000	0.0000000	-0.2744083	0.0000000	0.0000000	0.0000000
-0.3903747	-0.4233146	0.0000000	-0.2887344	-0.2660973	-0.0787757
0.4234383	-0.0983620	0.0000000	-0.4999028	-0.3859480	-0.2660775
0.0000000	0.0000000	-0.2744055	0.0000000	0.0000000	0.0000000
-0.3903747	0.4233146	0.0000000	-0.2887344	0.2660973	-0.0787757
-0.4234383	-0.0983620	0.0000000	0.4999028	-0.3859480	0.2660775
0.0000000	0.0000000	-0.2744055	0.0000000	0.0000000	0.0000000

---

The first three (3) rows signify 'x, y, z' coordinates of one (1) nitrogen atom and the following nine (9) rows represent Cartesian coordinates of three (3) oxygen atoms.



## References

- [1] Werner, H.-J.; et. al., MOLPRO, version 2010.1, a package of ab initio programs. 2010; <http://www.molpro.net>.
- [2] Frisch, M. J. et al. Gaussian~16 Revision C.01. 2016; Gaussian Inc. Wallingford CT.
- [3] Gordon, M. S.; Schmidt, M. W. *Theory and applications of computational chemistry*; Elsevier, 2005; pp 1167–1189.