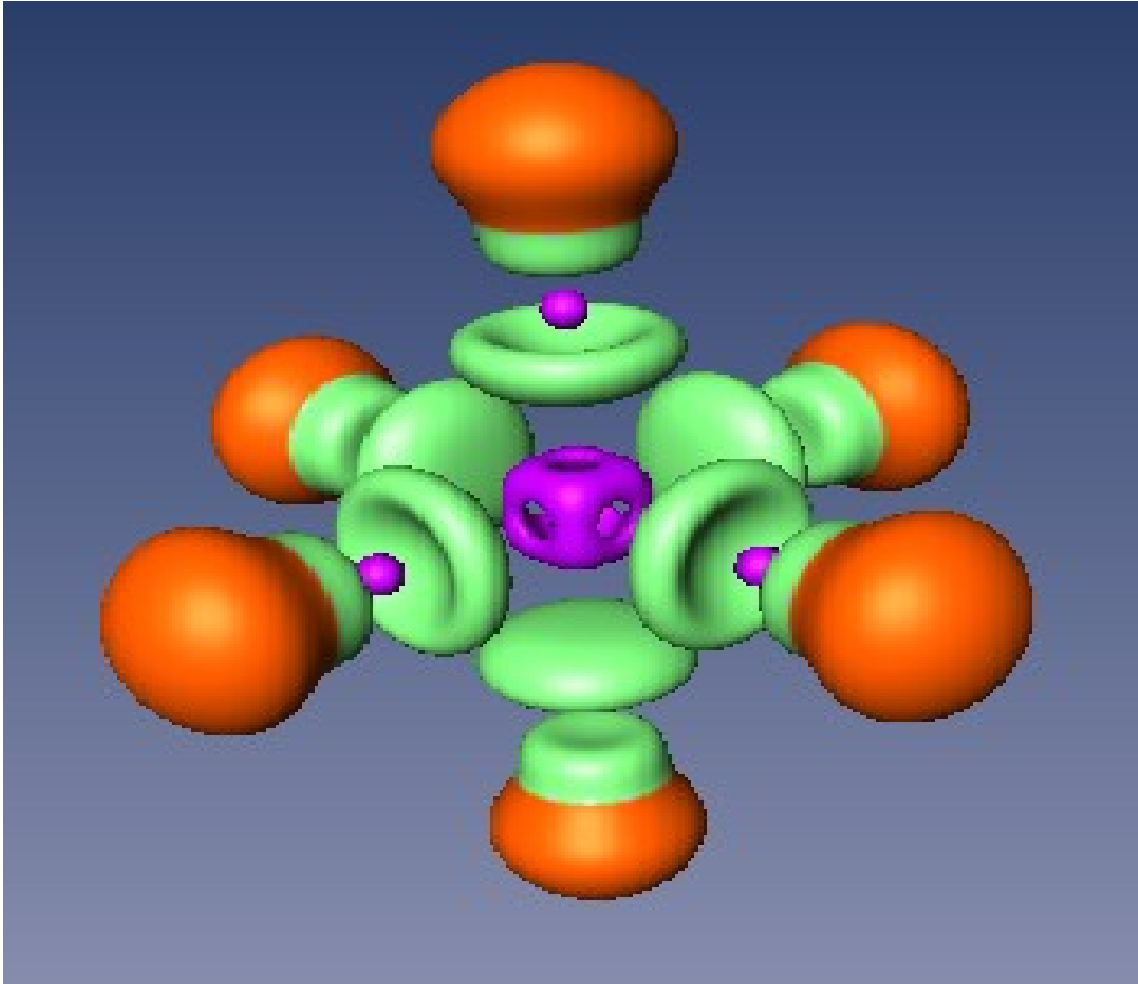


# The TopMoD

user's manual



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# 1 General Presentation

The ToPMoD package enables the calculation of the  $ELF$  function on a 3-dimensional grid, the assignment of the basins and the calculation of the basin populations and of their variance. It uses wave functions written in a `wfn` file which is available as output file generated by Gaussian92/94/98/03 and GAMESS *ab initio* softwares. The modules are written in FORTRAN 90 in order to enable dynamic memory allocation.

In order to carry out a standard  $ELF$  analysis three programs have to be run in the following order:

- `grid90`: calculates  $ELF$  on a 3-D grid parallel to the standard axis defined in the MO calculation (this allows to exploit the factorization of the gaussian functions),
- `bas90`: assigns the grid points to basins,
- `pop90`: calculates the basin populations and variances.

The other modules do the following tasks:

- `top_syn`: exploits abelian symmetry operations,
- `search90`: localizes the critical points of the  $ELF$  gradient field,
- `mod_wfn`: enables to remove atoms in the case of large systems in order to focuss the calculation on the region of interest,
- `sym_wfn`: symmetrizes the `wfn` file for  $^2\Pi$  states of linear molecules
- `bas_to_syn`: assigns basin types from `bas.sbf` file,
- `sbf_to_cube`: converts `sbf` files to `cube` files for Molekel,
- `sbf_to_am`: converts `sbf` files to `iam` files for Amira,
- `wfn_to_line`: generates a wireframe, stick or ball and stick molecular skeletons for Amira.

## 1.1 Useful reading and references

### References

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## 2 Gaussian and GAMESS inputs

The wfn files are written by GAMESS and by Gaussian. It is not a standard option and therefore it must be indicated in the input.

### 2.1 GAMESS

Set AIMPA $\text{C} = .\text{TRUE}.$  in the \$ CONTRL line command.

#### Example

```
$ CONTRL SCFTYP=RHF RUNTYP=OPTIMIZE MPLEVL=0 ICHARG=0 MULT=1 COORD=ZMT
AIMPAC=.TRUE. $ END
$ SYSTEM TIMLIM=100 MEMORY=800000 $ END
$ BASIS GBASIS=n31 NGAUSS=6 NDFUNC=1 NPFUNC=1 $ END
$ GUESS GUESS=HUCKEL $ END
$ DATA
CH3OH
Cs

H
O 1 oh
C 2 co 1 coh
H 3 ch1 2 hco1 1 180.0
H 3 ch2 2 hco2 1 t
H 3 ch2 2 hco2 1 -t

oh=0.979
co=1.3
coh=110.0
ch1=1.0
hco1=109.0
ch2=1.0
hco2=109.0
t=60.0

$ END
```

## 2.2 Gaussian

Set `out=wfn` in the Route section and give the name of the `wfn` file at the end of the molecule specification. The title first word (contiguous non blank characters) are used to form the file names in, it might be dangerous to use non alpha numerical characters.

### 2.2.1 Hartree Fock or DFT calculations

#### Example: C<sub>4</sub>H<sub>4</sub> tetrahedrane

```
# P HF/6-311++G(2df,2p) opt out=wfn

tetrahedrane

0,1
C -0.524831 0.524831 0.524831
C 0.524831 -0.524831 0.524831
C -0.524831 -0.524831 -0.524831
C 0.524831 0.524831 -0.524831
H -1.134794 1.134794 1.134794
H 1.134794 -1.134794 1.134794
H -1.134794 -1.134794 -1.134794
H 1.134794 1.134794 -1.134794

tetrahedrane.wfn
```

To compute condensed Fukui functions over basins it is necessary to set `tt IOP(99/18=1)` such in the following example:

#### BeCl<sub>2</sub>: input for condensed Fukui function calculation

```
#P B3LYP/6-31G** popl iop(99/18=1) out=wfn

BeCl2

0, 1
Cl
Be 1 r
X 2 1.0 1 90.0
Cl 2 r 3 90.0 1 180.0

r=1.62

becl2.wfn
```



## 2.2.2 Post Hartree Fock calculations

The `NaturalOrbitals` or `NaturalSpinOrbitals` should be specified as options for the population keyword and `all` in the density one.

### Closed shell example: water molecule

```
#CISD/6-31G** popt pop=no density=all out=wfn

H2O

0 1
H
O 1 r
H 2 r 1 a

r=1.0
a=110.0

h2ono.wfn
```

### Open shell example: CH<sub>3</sub> radical

```
#P CISD/6-31G** popt pop=NaturalSpin density=all out=wfn

CH3no

0 2
H
C 1 r
H 2 r 1 120.0
H 2 r 1 120.0 3 180.0

r=1.07

ch3no.wfn
```

### 3 grid09 input

1. `filein(a40)` : wfn file name
2. `origin(3)` (free format) : coordinates of the origin of the box defining the grid
3. `edge(3)` (free format) : length of the edges of the box in Bohr
4. `intervalx, intervaly, intervalz` (free format) : number of intervals on each edge. Choose the number of intervals in order to have steps of about 0.1 a.u and to avoid grid points in a symmetry plane.

in output the program generates `title_elf.sbf`, `title_rho.sbf` and `itle_lap.sbf` files)in which title is the first non-blank characters of the title in gaussian input (here water)

#### 3.1 Input example: water molecule

The `h2o.wfn` file begins by the following lines:

```
H2O
GAUSSIAN 5 MOL ORBITALS 30 PRIMITIVES 3 NUCLEI
O 1 (CENTRE 1) 0.00000000 0.00000000 0.22212920 CHARGE = 8.0
H 2 (CENTRE 2) 0.00000000 1.43441917 -0.88851679 CHARGE = 1.0
H 3 (CENTRE 3) 0.00000000 -1.43441917 -0.88851679 CHARGE = 1.0
```

The  $x = 0$  and  $y = 0$  planes can be used as symmetry planes

##### 3.1.1 grid09 input without symmetry

```
h2o.wfn
-5.0 -6.0 -6.0
10.0 12.0 12.0
100 120 120
```

The program will write a 101 x 121 x 121 grid on the `H2O_elf.sbf` file to be directly used by `bas09` for the ELF analysis `H2O_rho.sbf` file to be directly used by `bas09` for the AIM analysis `H2O_lap.sbf` file to display the laplacian of the charge density.

### 3.1.2 grid09 input with 2 symmetry planes

```
h2o.wfn  
0.05 0.05 -6.0  
5.0 6.0 12.0  
50 60 120
```

The program will write a 51 x 61 x 121 grid on the H2O\_elf.sbf file to be updated by top\_sym.

## 4 bas09 input

1. `function (a3)` : type of function elf/rho
2. `filein (a40)` : input wfn file
3. `iacc` free format : accuracy level `iacc=0` no approximation in derivative evaluation, 1 and 2 use of a cutoff. 1 is recommended.
4. `external core shell option (a1)` : do not merge the external core shell basins (y/n)
5. for each atom bearing a pseudopotential
  - `ns1` (free format) : type of the pseudopotential 1 large core, 2 small core
6. `attractor search mode` (free format) :
  - `< 0` fast automatic from the current grid, finds only attractors within the box.
  - `0` full automatic, search in the whole molecular space (recommended if symmetry is used).
  - `> 0` number of expected non protonated valence basins given in input
  - skip if automatic search, for each attractor in input:
    - `xp, yp, zp` (free format) : cartesian coordinates of the attractor (in a. u.)
    - `type, ref_atom`  
  
    `type`: 2 point, 3 circle, 4 sphere.  
    `ref_atom` is the number of the atom used to define the  $C_\infty$  axis in case of circular attractor. For point and sphere attractor put any integer.
    - `s_order`: synaptic order of the basin.
    - `radius`: radius of the circle or of the sphere, for point attractors 0.5 can be used safely.
    - `basin_name (3a4)`: atomic symbols used to name the basin.
7. `assign grid points (a3)` : (y/n)

In output the program writes one file: `title_ebas.sbf` or `title_ras.sbf` according to the chosen function to be used by `pop09` and one input file for `bas09` named `temp.bas`. The `temp.bas` file corresponds to the input attractor mode, it can be generated with a different wave function (for example with a smaller basis set) which nevertheless yields the same overall topology and without grid point assignment. In this case it is not necessary to run `grid09` before. This procedure is recommended for large systems. Do not forget to update the wfn file name in `temp.bas` if another wfn file is used.

## 4.1 Example: H<sub>2</sub>O all electron calculation

Fast automatic search	from input
elf	elf
h2o.wfn	h2o.wfn
1	1
n	n
0	2
	1.0 0.0 0.7
	2 0
	1
	0.5
	O
	-1.0 0.0 0.0
	2 0
	1
	0.5
	O
y	y

## 4.2 Example: H<sub>2</sub>O pseudopotential (large core) calculation

Automatic search	from input
elf	elf
h2o.wfn	h2o.wfn
1	1
n	n
0	1
-1	1
	2
	1.0 0.0 0.7
	2 0
	1
	0.5
	O
	-1.0 0.0 0.0
	2 0
	1
	0.5
	O
y	y

## 5 top\_sym input

1. file\_in (a40) : sbf input file
2. nop (free format) : number of symmetry operations (  $\leq 3$  )
3. nop times
  - (a) name\_op (a3) : symmetry operation symbol (see table)
  - (b) xc, yc, zc (free format) : coordinates defining the operation (see table)

symbol	operation	xc, yc, zc
INV	inversion	center of inversion coordinates
PLX	$x = xc$ plane	xc, 0.0, 0.0
PLY	$y = yc$ plane	0.0, yc, 0.0
PLZ	$z = zc$ plane	0.0, 0.0, zc
PXY	$y = x + yc$ plane	0.0, yc, 0.0
PYX	$y = -x + yc$ plane	0.0, yc, 0.0
PXZ	$z = x + zc$ plane	0.0, 0.0, zc
PZX	$z = -x + zc$ plane	0.0, 0.0, zc
PYZ	$z = -y + zc$ plane	0.0, 0.0, zc
PZY	$z = y + zc$ plane	0.0, 0.0, zc
C2X	$C_2(x)$ axis	0.0, yc, zc intersection with $x = 0.0$ plane
C2Y	$C_2(y)$ axis	xc, 0.0, zc intersection with $y = 0.0$ plane
C2Z	$C_2(z)$ axis	xc, yc, 0.0 intersection with $z = 0.0$ plane
C4X	$C_4(x)$ axis	0.0, yc, zc intersection with $x = 0.0$ plane
C4Y	$C_4(y)$ axis	xc, 0.0, zc intersection with $y = 0.0$ plane
C4Z	$C_4(z)$ axis	xc, yc, 0.0 intersection with $z = 0.0$ plane
S4X	$S_4(x)$ axis	xc, yc, zc center of inversion coordinates
S4Y	$S_4(y)$ axis	xc, yc, zc center of inversion coordinates
S4Z	$S_4(z)$ axis	xc, yc, zc center of inversion coordinates
CTX	$C_4(x)$ screw axis	yc, zc intersection, xc translation with $x = xc$ plane
CTY	$C_4(y)$ screw axis	xc, zc intersection, yc translation with $y = yc$ plane
CTZ	$C_4(z)$ screw axis	xc, yc intersection, zc translation with $z = zc$ plane

Symmetry operations symbols and coordinates.

### 5.1 Example

```
H2O.bas.sbf
2
PLX
```

```
0.0 0.0 0.0
PLY
0.0 0.0 0.0
```

The program will write a 101 x 121 x 121 grid on the H2O\_bas.sbf which is now updated.

**Case of  $T_d$  group** In the case of the  $T_d$  group, the best way to exploit symmetry is to put the vertices of the tetrahedron at the vertices of a cube like in the following example:

```
C 1 (CENTRE 1) 0.98523954 0.98523954 0.98523954 CHARGE = 6.0
C 2 (CENTRE 2) -0.98523954 -0.98523954 0.98523954 CHARGE = 6.0
C 3 (CENTRE 3) -0.98523954 0.98523954 -0.98523954 CHARGE = 6.0
C 4 (CENTRE 4) 0.98523954 -0.98523954 -0.98523954 CHARGE = 6.0
```

The input for grid09 corresponds to a quarter of a cube, i.e.:

```
0.05 0.05 -7.25
7.20 7.20 14.5
```

Then run top\_sym with two  $C_2$  rotation, one around the  $x$  axis, the other around  $y$ :

```
tetrahedrane_ebas.sbf 2
C2X
0.0 0.0 0.0
C2Y
0.0 0.0 0.0
```

## 6 pop09 and fast\_pop input

1. `filein(a40)`: input wfn file
2. `tol` (free format): integration threshold (recommended value 7)
3. `n_elf, n_aim` (free format): numbers of ELF and AIM basins considered (if 0 all the basins are taken into account in the calculation)
4. skip if `n_elf=0`: (`elf_ind(i), i=1, n_elf`): label of the considered ELF basins in the order of the bas09 output.
5. skip if `n_aim=0`: (`aim_ind(i), i=1, n_aim`): label of the considered AIM basins in the order of the bas09 output.

### 6.1 Example: H<sub>2</sub>O all electron calculation

Full calculation	only V(H <sub>2</sub> O) and Atom(O)	
<code>h2o.wfn</code>	<code>h2o.wfn</code>	
7	7	
0 0	1 1	The variance calculation is not valid
	3	
	2	

for MP2 wave functions use `fast_pop`. The variance calculation is not valid on a part of the molecule use `fast_pop` (asymmetric unit).

### 6.2 pop09 output: definition of calculated properties.

The following symbols have been used to define the calculated properties:

1.  $\Omega_i$ : basin labelled by  $i$
2.  $\rho(\mathbf{r})$ : spinless electron density.
3.  $\rho^\alpha(\mathbf{r})$ :  $\alpha$ -spin electron density.
4.  $\rho^\beta(\mathbf{r})$ :  $\beta$ -spin electron density.
5.  $\pi(\mathbf{r}, \mathbf{r}')$ : spinless electron pair density.
6.  $\pi^{\alpha\beta}(\mathbf{r}, \mathbf{r}')$ :  $\alpha\beta$  contribution to electron pair density.
7.  $\pi^{\alpha\alpha}(\mathbf{r}, \mathbf{r}')$ :  $\alpha\alpha$  contribution to electron pair density.
8.  $\pi^{\beta\beta}(\mathbf{r}, \mathbf{r}')$ :  $\beta\beta$  contribution to electron pair density.



### 6.2.1 Basin properties.

1. basin volume  $V(\Omega_i) = \int_{\Omega_i} d\mathbf{r}$ , for  $ELF \geq 0.02$
2. basin population  $\bar{N}(\Omega_i) = \int_{\Omega_i} \rho(\mathbf{r}) d\mathbf{r}$ .
3.  $\alpha\beta$  pair population  $\bar{P}^{\alpha\beta}(\Omega_i, \Omega_i) = \int_{\Omega_i} \int_{\Omega_i} \pi^{\alpha\beta}(\mathbf{r}, \mathbf{r}') d\mathbf{r} d\mathbf{r}'$ .
4.  $\alpha\alpha$  pair population  $\bar{P}^{\alpha\alpha}(\Omega_i, \Omega_i) = \int_{\Omega_i} \int_{\Omega_i} \pi^{\alpha\alpha}(\mathbf{r}, \mathbf{r}') d\mathbf{r} d\mathbf{r}'$ .
5.  $\beta\beta$  pair population  $\bar{P}^{\beta\beta}(\Omega_i, \Omega_i) = \int_{\Omega_i} \int_{\Omega_i} \pi^{\beta\beta}(\mathbf{r}, \mathbf{r}') d\mathbf{r} d\mathbf{r}'$ .
6. variance  $\sigma^2(\Omega_i) = \int_{\Omega_i} \int_{\Omega_i} \pi(\mathbf{r}, \mathbf{r}') d\mathbf{r} d\mathbf{r}' + \bar{N}\Omega_i - \bar{N}^2(\Omega_i)$ .
7. relative fluctuation  $\lambda(\Omega_i) = \sigma^2(\Omega_i) / \bar{N}\Omega_i$
8. integrated spin density  $S_z(\Omega_i) = \frac{1}{2} \int_{\Omega_i} (\rho^\alpha(\mathbf{r}) - \rho^\beta(\mathbf{r})) d\mathbf{r}$ .
9. condensed Fukui functions  $f^+, f^0, f^-$

$$\begin{aligned} \langle f^- \rangle &= \int_{\Omega_i} [\rho_N(\mathbf{r}) - \rho_{N-1}(\mathbf{r})] d\mathbf{r} \\ \langle f^+ \rangle &= \int_{\Omega_i} [\rho_{N+1}(\mathbf{r}) - \rho_N(\mathbf{r})] d\mathbf{r} \\ \langle f^0 \rangle &= \frac{1}{2} (\langle f^- \rangle + \langle f^+ \rangle) \end{aligned}$$

To get condensed Fukui set IOP (99/18=1) in Gaussian03 input file

### 6.2.2 variance related properties.

1.  $\alpha$ -spin entropy:  $\sum_{i,j} p^{\alpha\alpha}(\Omega_i, \Omega_j) \log \frac{p^{\alpha\alpha}(\Omega_i, \Omega_j)}{n^\alpha(\Omega_i)n^\alpha(\Omega_j)}$ .
2.  $\beta$ -spin entropy:  $\sum_{i,j} p^{\beta\beta}(\Omega_i, \Omega_j) \log \frac{p^{\beta\beta}(\Omega_i, \Omega_j)}{n^\beta(\Omega_i)n^\beta(\Omega_j)}$ .
3. total entropy:  $\sum_{i,j} p(\Omega_i, \Omega_j) \log \frac{p(\Omega_i, \Omega_j)}{n(\Omega_i)n(\Omega_j)}$ .
4. covariance  $B_{i,j} = \bar{N}(\Omega_i)\bar{N}(\Omega_j) - P(\Omega_i, \Omega_j)$ .
5. relative covariance  $B_{i,j} / \bar{N}(\Omega_j)$ .

6. fluctuation contribution:  $B_{i,j} / \sum_{j \neq i} B_{i,j}$

where  $p(\Omega_i, \Omega_j) = P(\Omega_i, \Omega_j) / N(N - 1)$ ,  $n(\Omega_i) = \bar{N}(\Omega_j) / N$ .

## 7 search09 input

1. `filein(a40)`: input wfn file
2. type of critical point (free format): 0 any, 1 attractor, -1 end
3. `x`, `y`, `z` (free format): guessed location

### 7.1 Example

```
h2o.wfn
0
1.0 -1.  0.5
1
0.3 0.2 0.1
-1
```

## 8 wfn file manipulations

For some specific applications the `wfn` file must be modified because GAUSSIAN contains a bug in the case of ROHF calculations (all the orbitals are doubly occupied!). There are two `wfn` manipulations:

1. ROHF: change occupation number of singly occupied orbitals

```
MO 5 MO 0.0 OCC NO = 2.0000000 ORB. ENERGY = 0.0
MO 5 MO 0.0 OCC NO = 1.0000000 ORB. ENERGY = 0.0
```

2. ROHF and CASSCF: specify orbital contribution to  $M_S$  (in fact twice). The 0.0 after MO should be changed to -1 ( $\beta$ ), 0 (mixing true singlet), 1 ( $\alpha$ ), example

```
MO 5 MO 1.0 OCC NO = 1.0000000 ORB. ENERGY = 0.0
```

### 8.1 sym\_wfn

Molecular *ab initio* programs use basis real functions, in the case of linear molecules these computed MOs are not eigenfunctions of the  $L_z$  operator and therefore the cylindrical symmetry can be lost. This is particularly the case for open shell systems in a  $\Pi$  state. `sym_wfn` has been written to correct this drawback for  $^2\Pi$  states calculated with ROHF. In order to get symmetrized figures run `sym_wfn` before performing the `grid09` calculation. The program prompts the user to get the input `wfn` file name and the name of the symmetrized `sym_wfn` file.

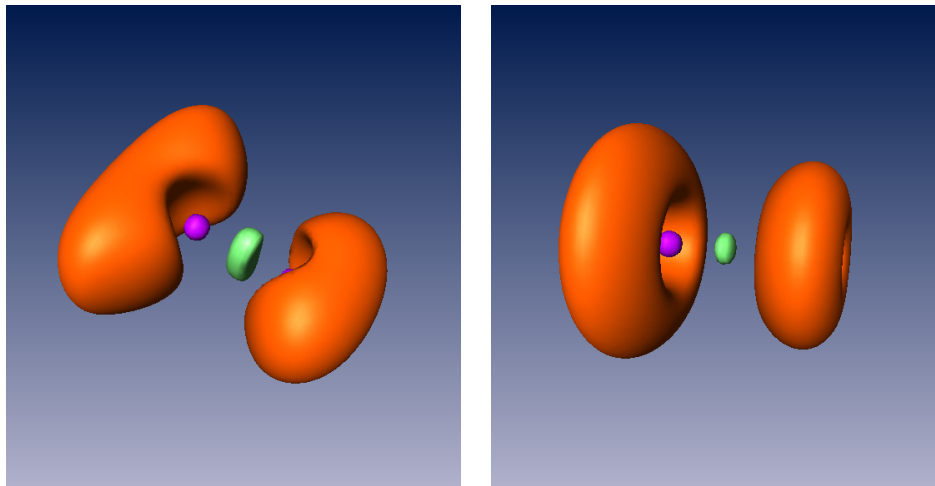


Figure 1: NO ( ${}^2\Pi$ ): left unsymmetrized, right symmetrized

## 8.2 mod\_wfn

For large molecular systems `mod_wfn` enables to restrict the analysis to a subpart of the system. The atoms of the subpart of interest and their first neighbours are entered as data of `mod_wfn`

### 8.2.1 mod\_wfn input

1. `filein (a40)` : wfn file name
2. `natoms2 (free format)` : number of selected atoms
3. `iatom(i), i=1, natoms2` : list of the selected atoms
4. `fileout (a40)` : modified wfn file name

**Example: CH2ClCH2COOH** The selected subpart is the functional group. The atoms in the wfn appear in the following order:

C	1	(CENTRE	1)	1.52388347	0.82729056	0.60836370	CHARGE =	6.0
C	2	(CENTRE	2)	-0.36562060	-1.02000099	-0.54662559	CHARGE =	6.0
Cl	3	(CENTRE	3)	4.74167762	-0.16320636	-0.05397264	CHARGE =	17.0
C	4	(CENTRE	4)	-3.10102364	-0.30181750	-0.07030471	CHARGE =	6.0
O	5	(CENTRE	5)	-4.78491297	-1.77008583	0.40373635	CHARGE =	8.0
O	6	(CENTRE	6)	-3.48476780	2.22384618	-0.29263556	CHARGE =	8.0
H	7	(CENTRE	7)	1.35335617	0.91475856	2.66069875	CHARGE =	1.0
H	8	(CENTRE	8)	1.30958103	2.72008831	-0.16895560	CHARGE =	1.0

H	9	(CENTRE 9)	-0.08789824	-1.09332176	-2.59892348	CHARGE =	1.0
H	10	(CENTRE 10)	-0.09155624	-2.92840929	0.18473731	CHARGE =	1.0
H	11	(CENTRE 11)	-5.27799149	2.49847708	0.00257117	CHARGE =	1.0

The selected atoms are C(2), C(4), O(5), O(6) and H(11). The input is :

```
ch2clch2cooh.wfn
5
2 4 5 6 11
ghost.wfn
```

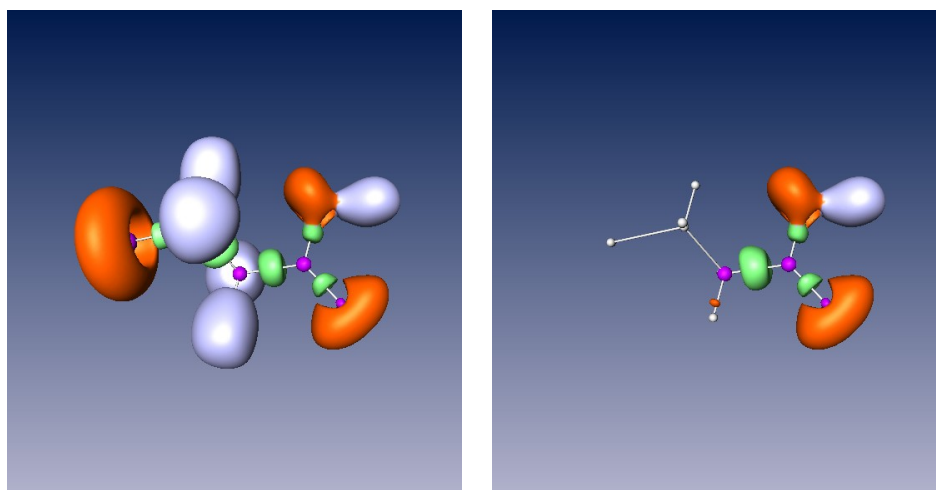


Figure 2: Display of  $\text{CH}_2\text{ClCH}_2\text{COOH}$  localization domains. Left: total, right: selection.

## 9 Visualization

In order to visualize, first create a color code file with the `bas_to_syn` utility program. According to your data analyzer, we recommend Amira and Molekel (version 4), generate either `title_elf.cube` and `title_esynf.cube` files with the `sbf_to_cube` utility for Molekel. The `sbf_to_am` utility converts the `sbf` files to the `am` format readable by Amira.

Molekel is an Advanced Interactive 3D-graphics package for visualizing molecular and electronic structure data from output of various Quantum Chemistry softwares such as GAUSSIAN94/98, GAMESS-US, ADF, HONDO. It works under UNIX, LINUX and also WINDOWS. It uses the Open/GL, Mesa, GLUT and GLUI libraries. MOLEKEL was designed and developed at University of Geneva and CSCS / ETHZ by Peter F. Flükiger in the early 90ties and later by Stefan Portmann.

Molekel is available at the following URL:

<http://www.cscs.ch/molekel>

A ppt file explains how to use Molekel. The `syn.rgbfile` in the SYN directory provides the texture and color codes.

Amira was originally distributed by TGS. The following URL provides useful piece of information and enable to download a documentation brochure and a free trial

<http://www.amiravis.com/>

The `syn.am` file in the SYN directory provides the texture and color codes.

## **9.1 sbf\_to\_cube and sbf\_to\_am inpiuts**

### **9.1.1 sbf\_to\_cube**

1. `filein(a40)`: wfn file name
2. `filesbf(a40)`: sbf file to be converted
3. `ix, iy, iz`: step size in each direction (standard 1 1 1)

Run `sbf_to_cube` for the elf and syn sbf files.

### **9.1.2 sbf\_to\_am**

1. `filesbf(a40)`: sbf file to be converted

Run `sbf_to_am` for the elf and syn sbf files.