```
1 : 1.7561 2.2540
2 : -1.7562 2.2541
3 : 0.0001 0.9800
```

The output from this calculation shows the expected results, with the local site spins being close to their ideal value 2.5 which would hold for a high-spin Fe(III) ion.

5.1.11 Population Analysis of UNOs

Population analyses for *UHF natural Orbitals* can be requested via the respective print options in the %output block:

```
%output
Print[ P_UNO_OccNum ] = 1;  # Occupation numbers
Print[ P_UNO_AtPopMO_M ] = 0;  # Mulliken atom pop. per UNO
Print[ P_UNO_OrbPopMO_M] = 0;  # Mulliken orbital pop. per UNO
Print[ P_UNO_ReducedOrbPopMO_M] = 0;  # Mulliken reduced orbital pop. per UNO
Print[ P_UNO_AtPopMO_L ] = 0;  # Loewdin atom pop. per UNO
Print[ P_UNO_OrbPopMO_L] = 0;  # Loewdin orbital pop. per UNO
Print[ P_UNO_ReducedOrbPopMO_L] = 0;  # Loewdin reduced orbital pop. per UNO
end
```

5.1.12 Keywords

A collection of useful keywords in the context of population analyses is given in Table 5.1, Table 5.2, Table 5.3, and Table 5.4.

Keyword	Description
AllPop	Activates all population analyses
NoPop	Deactivates all population analyses
ReducedPop	Prints reduced orbital populations per MO
NoReducedPop	Deactivates printing of reduced orbital populations
Mulliken	Activates Mulliken population analysis
NoMulliken	Deactivates Mulliken population analysis
Loewdin	Activates <i>Löwdin</i> population analysis
NoLoewdin	Deactivates Löwdin population analysis
Mayer	Activates Mayer population analysis
NoMayer	Deactivates Mayer population analysis
NPA	Activates NPA. NBO program required
NBO	Activates NPA and NBO analyses. NBO program required
NoNBO	Deactivates NBO
NoNPA	Deactivates NPA and NBO
MBIS	Activates MBIS population analysis
CHELPG	Requests calculation of CHELPG charges
CHELPG (LARGE)	Requests calculation of CHELPG charges with large grid

Table 5.1: Simple input keywords for population analyses.

Table 5.2: %output block input keywords and options for population analyses.

Keyword	Option	Description
Print[<option>]</option>	P_Mayer	Print Mayer population analysis (Default = on)
	P_NatPop	Print Natural population analysis (Default = off)

Table 5.2 - continued from previous page

Keyword	Option	Description
	P_NPA	Print Natural population analysis (Default = off)
	P_Hirshfeld	Print Hirshfeld population analysis (Default = off)
	P_MBIS	Print MBIS population analysis (Default = off)
	P_Mulliken	Print Mulliken population analysis (Default = on)
	P_AtCharges_M	Print Mulliken atomic charges
	P_OrbCharges_M	Print Mulliken orbital charges
	P_FragCharges_M	Print Mulliken fragment charges
	P_FragBondOrder_M	Print Mulliken fragment bond orders
	P_BondOrder_M	Print Mulliken bond orders
	P_ReducedOrbPop_M	Print Mulliken reduced orbital charges
	P_FragPopMO_M	Print Mulliken fragment population for each MO
	P_FragOvlMO_M	Print Mulliken overlap populations per fragment pair
	P_AtPopMO_M	Print Mulliken atomic charges in each MO
	P_OrbPopMO_M	Print Mulliken orbital population for each MO
	P_ReducedOrbPopMO_M	Print Mulliken reduced orbital population for each MO
	P_Loewdin	Print Loewdin population analysis (Default = on)
	P_AtCharges_L	Print Loewdin atomic charges
	P_OrbCharges_L	Print Loewdin orbital charges
	P_FragCharges_L	Print Loewdin fragment charges
	P_FragBondOrder_L	Print Loewdin fragment bond orders
	P_BondOrder_L	Print Loewdin bond orders
	P_ReducedOrbPop_L	Print Loewdin reduced orbital charges
	P_FragPopMO_L	Print Loewdin fragment population for each MO
	P_FragOvlMO_L	Print Loewdin overlap populations per fragment pair
	P_AtPopMO_L	Print Loewdin atomic charges in each MO
	P_OrbPopMO_L	Print Loewdin orbital population for each MO
	P_ReducedOrbPopMO_L	Print Loewdin reduced orbital population for each MO
	P_Fragments	Print fragment information
	P_GUESSPOP	Print initial guess populations
	P_UNO_FragPopMO_M	Print Mulliken fragment population per UNO
	P_UNO_OrbPopMO_M	Print Mulliken orbital pop. per UNO
	P_UNO_AtPopMO_M	Print Mulliken atomic charges per UNO
	P_UNO_ReducedOrbPopMO_M	Print Mulliken reduced orbital pop. per UNO
	P_UNO_FragPopMO_L	Print Loewdin fragment population per UNO
	P_UNO_OrbPopMO_L	Print Loewdin orbital pop. per UNO
	P_UNO_AtPopMO_L	Print Loewdin atomic charges per UNO
	P_UNO_ReducedOrbPopMO_L	Print Loewdin reduced orbital pop. per UNO
	P_UNO_OccNum	Print occupation numbers per UNO

Table 5.3: %method block input keywords and options for population analyses.

Keyword	Option	Description
LOEWDIN_BONDORDE	<real></real>	Sets printing threshold for Löwdin bond orders (default = 0.05)
MAYER_BONDORDERTI	<real></real>	Sets printing threshold for Mayer bond orders (default = 0.1)
MBIS_CHARGETHRESI	<real></real>	Sets convergence threshold for MBIS charges (default = 10^{-6})
MBIS_LARGEPRINT	true/false	Activate printing of <i>MBIS Quantities</i> (default = false)
MBIS_ORIGIN_MULT	CenterOfCoor	Sets origin for MBIS properties to center of coordinate system $(0,0,0)$
	CenterOfMass	Sets origin for MBIS properties to center of mass
	CenterOfNucCl	Sets origin for MBIS properties to center of nuclear charge
	CenterXYZ	Sets origin for MBIS properties to arbitrary position, set coordinates with MBIS_ORIMULT_XYZ
	CenterOfEachi	Sets origin for MBIS properties to center of each atom (default)
MBIS_ORIMULT_XYZ	<x,y,z></x,y,z>	Set the coordinates, otherwise 0,0,0 (unit: Angstrom)

Table 5.4: %chelpg block input keywords and options for CHELPG charges.

Keyword	Option	Description
GRID	<real></real>	Spacing of the regular grid in $Å$ (default = 0.3)
RMAX	<real></real>	Maximum distance of all atoms to any gridpoint in $Å$ (default = 2.8)
VDWRADII	COSMO	Activates usage of COSMO VdW radii of Atoms (default)
	BW	Activates usage of Breneman, Wiberg radii
DIPOLE	false/true	If true, the charges also reproduce the total dipole moment (default = false)

5.2 Natural Bond Orbital (NBO) Analysis

A popular and useful method for population analysis is the natural bond orbital analysis due to Weinhold and coworkers. It is implemented in the **NBO** program which is distributed in older versions via the CCL list and in newer versions via the University of Wisconsin/Madison. Information about the NBO program can be found at http://www.chem.wisc.edu/~nbo7. In order to use it together with ORCA you need a version of the stand-alone executable. Starting with version 3.1.x ORCA can only be used with NBO6 or NBO7. To specify the NBO executable the environment variable NBOEXE=/full/name/of/nbo7-executable has to be set. As the NBO part of the interface is not independent of the integer data-type width (i4 or i8), the NBO executable which will be used together with ORCA has to be compiled using i4!

ORCA features two methods to interface with the NBO program: ! NBO keyword and the %nbo-block. The following example illustrates the use for formaldehyde:

```
# Test the interface to the NBO program
! RHF SVP NBO
* xyz 0 1
 С
       0.000000
                 0.000000
                            0.000000
 0
       1.200000
                0.000000
                            0.000000
 Н
      -0.550000
                0.952628
                             0.000000
                  -0.952628
 Η
      -0.550000
                             -0.00000
```

This produces the following output:

```
Now starting NBO....
 NATURAL ATOMIC ORBITAL AND
                NATURAL BOND ORBITAL ANALYSIS
  **************** development version (D000000) *****************
   (c) Copyright 1996-2018 Board of Regents of the University of Wisconsin System
         on behalf of the Theoretical Chemistry Institute. All rights reserved.
                 Cite this program [NBO 7.0.0 (15-Nov-2018)] as:
                NBO 7.0. E. D. Glendening, J. K. Badenhoop, A. E. Reed,
                J. E. Carpenter, J. A. Bohmann, C. M. Morales, P. Karafiloglou,
                 C. R. Landis, and F. Weinhold, Theoretical Chemistry Institute,
                University of Wisconsin, Madison, WI (2018)
           /NPA / : Natural Population Analysis
                        / : Natural Bond Orbital Analysis
           /AONBO / : Checkpoint the AO to NBO transformation
           /ARCHIVE/ : Write the archive file to lfn47
 Job title: ORCA Job: NBO_1
 NATURAL POPULATIONS: Natural atomic orbital occupancies
  NAO Atom No lang Type (AO) Occupancy
                                                                               Energy

      1
      C
      1
      s
      Cor(1s)
      1.99997
      -11.34329

      2
      C
      1
      s
      Val(2s)
      1.01533
      -0.17540

      3
      C
      1
      s
      Ryd(3s)
      0.00701
      0.61376

      4
      C
      1
      px
      Val(2p)
      0.81697
      0.08822

      5
      C
      1
      px
      Ryd(3p)
      0.01268
      0.63900

      6
      C
      1
      py
      Val(2p)
      1.09795
      -0.01243

      7
      C
      1
      py
      Ryd(3p)
      0.00055
      0.80803

      8
      C
      1
      pz
      Val(2p)
      0.66003
      -0.03464

      9
      C
      1
      pz
      Ryd(3p)
      0.00283
      0.62824

      10
      C
      1
      dxy
      Ryd(3d)
      0.00576
      2.75039

      11
      C
      1
      dxz
      Ryd(3d)
      0.00375
      2.25746

      12
      C
      1
      dx2y
      Ryd(3d)
      0.00337
      2.74845

      14
      C
      1
      dz2
      Ryd(3d)
      0.00114
      2.406
  23 O 2 pz Ryd(3p)
                                                      0.00011
                                                                              1.30080
   24 O 2 dxy Ryd(3d)
                                                      0.00213
                                                                               3.26414
   25 O 2 dxz Ryd(3d)
                                                      0.00340
                                                                                3.20490
                                                     0.00000
0.00406
   26 O 2 dyz Ryd (3d)
                                                                               2.98918
                                                                             3.55008
   27 O 2 dx2y2 Ryd(3d)
   28 O 2 dz2 Ryd(3d)
                                                       0.00119
                                                                               3.17511

      29
      H
      3
      s
      Val(1s)
      0.88576
      0.07107

      30
      H
      3
      s
      Ryd(2s)
      0.00298
      0.41181

      31
      H
      3
      px
      Ryd(2p)
      0.00030
      2.18260

      32
      H
      3
      py
      Ryd(2p)
      0.00159
      2.49146

           H 3 py Ryd(2p)
```

33 H 3 pz Ryd(2p) 0.00002 1.85643 34 H 4 s Ryd(2p) 0.00298 0.41181 35 H 4 s Ryd(2p) 0.00298 0.41181 36 H 4 px Ryd(2p) 0.00159 2.49146 37 H 4 px Ryd(2p) 0.00159 2.49146 38 H 4 px Ryd(2p) 0.00159 2.49146 38 H 4 px Ryd(2p) 0.00002 1.85643 wmmary of Natural Population Analysis: Natural Population	34 H 4 s Val(1s) 0.88576 0.07107 35 H 4 s Ryd(2s) 0.00298 0.41181 36 H 4 px Ryd(2p) 0.00030 2.18260 37 H 4 py Ryd(2p) 0.00159 2.49146 38 H 4 pz Ryd(2p) 0.00159 2.49146 38 H 4 pz Ryd(2p) 0.00002 1.85643 ummary of Natural Population Analysis: Natural Population
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35 H 4 s Ryd(2s) 0.00288 0.41181 36 H 4 px Ryd(2p) 0.00030 2.18260 37 H 4 px Ryd(2p) 0.000159 2.49146 38 H 4 px Ryd(2p) 0.00002 1.85643 wnmary of Natural Population Analysis: Natural Population	135
Natural Population	37
Natural Population Natural Population Natural Population	Natural Population Natural
Natural Fopulation Natural	Natural Population Natural
Natural Population	Natural Population
Natural Population	Natural Population
Natural Charge	Atom No Charge Core Valence Rydberg Total C 1 0.37265 1.99997 3.59028 0.03709 5.62735 0 2 -0.59134 1.99998 6.57415 0.01720 8.59134 H 3 0.10934 0.00000 0.88576 0.00489 0.89066 H 4 0.10934 0.00000 0.88576 0.00489 0.89066 Total * -0.00000 3.99995 11.93596 0.06408 16.00000 Natural Population Core 3.99995 (99.9988% of 4) Valence 11.93596 (99.4664% of 12) Natural Minimal Basis 15.93592 (99.5995% of 16) Natural Rydberg Basis 0.06408 (0.4005% of 16) Atom No Natural Electron Configuration C 1 [core]2s(1.02)2p(2.57)3s(0.01)3p(0.02)3d(0.01) 0 2 [core]2s(1.71)2p(4.87)3d(0.01) H 3 1s(0.89) ATURAL BOND ORBITAL ANALYSIS: ATURAL BOND ORBITAL ANALYSIS: Document of the property of the
0 2 -0.59134 1.99998 6.57415 0.01720 8.59134 H 3 0.10934 0.00000 0.88576 0.00489 0.89066 H 4 0.10934 0.00000 0.88576 0.00489 0.89066 H 5 0.00498 0.00498 0.89066 H 5 0.00498 0.00498 0.00498 0.00498 0.89066 H 5 0.00498 0.004	0 2 -0.59134 1.99998 6.57415 0.01720 8.59134 H 3 0.10934 0.00000 0.88576 0.00489 0.89066 H 4 0.10934 0.00000 0.88576 0.00489 0.89066 H 5 0.00489 0.00489 0.89066 H 5 0.00489 0.00499 0.89066 H 5 0.00489 0.00489 0.89066 H 5 0.00489 0.00489 0.89066 H 5 0.00489 0.004
O 2 -0.59134 1.99998 6.57415 0.01720 8.59134 H 3 0.10934 0.00000 0.88576 0.00489 0.89066 H 4 0.10934 0.00000 0.88576 0.00489 0.89066 H 5 0.00498 0.00498 0.89066 H 5 0.00498 0	O 2 -0.59134
H 3 0.10934 0.00000 0.88576 0.00489 0.89066 H 4 0.10934 0.00000 0.88576 0.00489 0.89066 Total * -0.00000 3.99995 11.93596 0.06408 16.00000 Natural Population	H 3 0.10934 0.00000 0.88576 0.00489 0.89066 H 4 0.10934 0.00000 0.88576 0.00489 0.89066 H 4 0.10934 0.00000 0.88576 0.00489 0.89066
Natural Population	Natural Population
Natural Population	Natural Population Core 3.99995 (99.9988% of 4) Valence 11.93596 (99.4664% of 12) Natural Minimal Basis 15.93592 (99.5995% of 16) Natural Rydberg Basis 0.06408 (0.4005% of 16) Atom No Natural Electron Configuration C 1 [core]2s(1.02)2p(2.57)3s(0.01)3p(0.02)3d(0.01) 0 2 [core]2s(1.71)2p(4.87)3d(0.01) H 3 1s(0.89) H 4 1s(0.89) ATURAL BOND ORBITAL ANALYSIS: Occupancies Lewis Structure Low High Occupancy Lewis orbitals Core Occupancy Lewis orbitals Core 3.99995 (99.999% of 4) Valence Lewis 11.89676 (99.140% of 12) Total Lewis 15.89671 (99.354% of 16) Valence non-Lewis 0.07835 (0.490% of 16) Rydberg non-Lewis 0.02493 (0.156% of 16) Rydberg non-Lewis 0.02493 (0.156% of 16) Rydberg non-Lewis 0.02493 (0.156% of 16)
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Atom No Natural Electron Configuration C 1 [core]2s(1.02)2p(2.57)3s(0.01)3p(0.02)3d(0.01) 0 2 [core]2s(1.71)2p(4.87)3d(0.01) H 3 1s(0.89) H 4 1s(0.89) ATURAL BOND ORBITAL ANALYSIS: Occupancies Lewis Structure Low High occ occ occ occ occ occ occ occ occ oc	Atom No Natural Electron Configuration C 1 [core]2s(1.02)2p(2.57)3s(0.01)3p(0.02)3d(0.01) 0 2 [core]2s(1.71)2p(4.87)3d(0.01) H 3 1s(0.89) H 4 1s(0.89) ATURAL BOND ORBITAL ANALYSIS: Occupancies Lewis Structure Low High occ occ occ occ occ occ occ occ occ oc
Occupancies Lewis Structure Low High occ	Occupancies Lewis Structure Low High occ
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Core 3.99995 (99.999% of 4) Valence Lewis 11.89676 (99.140% of 12) Total Lewis 15.89671 (99.354% of 16) Valence non-Lewis 0.07835 (0.490% of 16) Rydberg non-Lewis 0.02493 (0.156% of 16)	Core 3.99995 (99.999% of 4) Valence Lewis 11.89676 (99.140% of 12) Total Lewis 15.89671 (99.354% of 16) Valence non-Lewis 0.07835 (0.490% of 16) Rydberg non-Lewis 0.02493 (0.156% of 16)
Core 3.99995 (99.999% of 4) Valence Lewis 11.89676 (99.140% of 12) Total Lewis 15.89671 (99.354% of 16) Valence non-Lewis 0.07835 (0.490% of 16) Rydberg non-Lewis 0.02493 (0.156% of 16)	Core 3.99995 (99.999% of 4) Valence Lewis 11.89676 (99.140% of 12) Total Lewis 15.89671 (99.354% of 16) Valence non-Lewis 0.07835 (0.490% of 16) Rydberg non-Lewis 0.02493 (0.156% of 16)
Valence non-Lewis 0.07835 (0.490% of 16) Rydberg non-Lewis 0.02493 (0.156% of 16)	Valence non-Lewis 0.07835 (0.490% of 16) Rydberg non-Lewis 0.02493 (0.156% of 16)
Rydberg non-Lewis 0.02493 (0.156% of 16)	Rydberg non-Lewis 0.02493 (0.156% of 16)

```
(Occupancy) Bond orbital / Coefficients / Hybrids
  ----- Lewis -----
1. (1.99997) CR (1) C 1
                                   s(100.00%)
                                     1.0000 0.0000 0.0000 0.0000 0.0000
                                     0.0000 0.0000 0.0000 0.0000 0.0000
0.0000 0.0000 0.0000
 2. (1.99998) CR (1) O 2
                                    s(100.00%)
                                     1.0000 0.0000 0.0000 0.0000 0.0000
0.0000 0.0000 0.0000 0.0000
0.0000 0.0000 0.0000 0.0000
3. (1.98853) LP (1) 0 2
                                   s(56.22%)p 0.78(43.73%)d 0.00(0.05%)
                                      0.0000 0.7496 -0.0170 0.6612 0.0069
                                     0.0000 0.0000 0.0000 0.0000 0.0000
                                     0.0000 0.0000 -0.0201 0.0100
4. (1.91757) LP (2) 0 2
                                    s( 0.00%)p 1.00( 99.89%)d 0.00( 0.11%)
                                     0.0000 0.0000 0.0000 -0.0000 -0.0000
                                     0.9994 -0.0098 0.0000 -0.0000 -0.0330
                                     -0.0000 0.0000 0.0000 -0.0000
 5. (1.99996) BD (1) C 1- O 2
             (33.33%) 0.5773* C 1 s( 0.00%)p 1.00( 99.44%)d 0.01( 0.56%)
                                     0.0000 -0.0000 -0.0000 0.0000 -0.0000
                                     -0.0000 0.0000 0.9951 -0.0652 -0.0000
                                      0.0750 0.0000 -0.0000 0.0000
             (66.67%) 0.8165* 0 2 s( 0.00%)p 1.00( 99.75%)d 0.00( 0.25%)
                                     0.0000 -0.0000 0.0000 0.0000 -0.0000
                                     -0.0000 0.0000 0.9987 -0.0090 0.0000
-0.0505 0.0000 0.0000 -0.0000
 6. (1.99975) BD (2) C 1- O 2
                        0.5709* C 1 s( 32.18%)p 2.09( 67.35%)d 0.01( 0.46%)
             ( 32.59%)
                                      0.0000 0.5628 0.0714 0.8149 0.0973
                                     -0.0000 -0.0000 -0.0000 -0.0000 -0.0000
-0.0000 0.0000 0.0618 -0.0286
             (67.41%) 0.8211* 0 2 s(43.84%)p 1.27(55.85%)d 0.01(0.31%)
                                      0.0000 0.6615 0.0284 -0.7470 -0.0215
                                     -0.0000 0.0000 -0.0000 -0.0000 0.0000
                                     0.0000 0.0000 0.0490 -0.0270
7. (1.99548) BD (1) C 1- H 3
             (56.63%)
                        0.7526* C 1 s( 33.98%)p 1.94( 65.86%)d 0.00( 0.16%)
                                      0.0000 0.5826 -0.0192 -0.3995 -0.0029
                                      0.7063 -0.0087 0.0000 -0.0000 -0.0318
                                     -0.0000 0.0000 -0.0180 -0.0153
             (43.37%) 0.6585* H 3 s(99.79%)p 0.00( 0.21%)
                                      0.9989 -0.0095 0.0184 -0.0416 0.0000
 8. (1.99548) BD (1) C 1- H 4
             (56.63%) 0.7526* C 1 s(33.98%)p 1.94(65.86%)d 0.00(0.16%)
                                     0.0000 0.5826 -0.0192 -0.3995 -0.0029
                                     -0.7063 0.0087 -0.0000 0.0000 0.0318
                                      0.0000 0.0000 -0.0180 -0.0153
             (43.37%) 0.6585* H 4 s(99.79%)p 0.00( 0.21%)
                                     0.9989 -0.0095 0.0184 0.0416 -0.0000
     ----- non-Lewis -----
 9. (0.00000) BD*(1) C 1-0 2
                        0.8165* C 1 s( 0.00%)p 1.00( 99.44%)d 0.01( 0.56%)
             (66.67%)
             ( 33.33%) -0.5773* O 2 s( 0.00%)p 1.00( 99.75%)d 0.00( 0.25%)
10. (0.00000) BD*(2) C 1-0 2
             (67.41%) 0.8211* C 1 s(32.18%)p 2.09(67.35%)d 0.01(0.46%)
             ( 32.59%) -0.5709* O 2 s( 43.84%)p 1.27( 55.85%)d 0.01( 0.31%)
11. (0.03918) BD*(1) C 1- H 3
             ( 43.37%) 0.6585* C 1 s( 33.98%)p 1.94( 65.86%)d 0.00( 0.16%)
                                     0.0000 -0.5826 0.0192 0.3995 0.0029
```

```
-0.7063 0.0087 -0.0000 0.0000 0.0318
                               0.0000 0.0000 0.0180 0.0153
           (56.63%) -0.7526* H 3 s(99.79%)p 0.00( 0.21%)
                              -0.9989 0.0095 -0.0184 0.0416 -0.0000
12. (0.03918) BD*(1) C 1- H 4
           ( 43.37%)
                  0.6585* C 1 s( 33.98%)p 1.94( 65.86%)d 0.00( 0.16%)
                               0.0000 -0.5826 0.0192 0.3995 0.0029
                              (56.63%) -0.7526* H 4 s(99.79%)p 0.00( 0.21%)
                              -0.9989 0.0095 -0.0184 -0.0416 0.0000
13. (0.00969) RY (1) C 1
                              s(29.83%)p 2.30(68.57%)d 0.05(1.60%)
                               0.0000 -0.0565 0.5432 -0.1169 0.8198
                               0.0000 -0.0000 0.0000 -0.0000 -0.0000
                              0.0000 0.0000 0.1087 -0.0648
                             s( 0.00%)p 1.00( 9.56%)d 9.46( 90.44%)
14. (0.00517) RY (2) C 1
                              0.0000 0.0000 -0.0000 0.0000 -0.0000
                              0.0465 0.3057 0.0000 -0.0000 0.9510
                              0.0000 0.0000 -0.0000 -0.0000
0.0000 -0.0000 -0.0000 0.0000 0.0000
                              0.0064 0.9948 -0.0000 0.0000 -0.1018
0.0000 0.0000 -0.0000 0.0000
23. (0.00014) RY (2) 0 2
                      s( 35.10%)p 1.44( 50.60%)d 0.41( 14.30%)
                              0.0000 -0.0178 0.5922 0.0556 -0.7091
                              0.0000 -0.0000 -0.0000 0.0000 -0.0000
                              0.0000 0.0000 0.3336 -0.1780
0.0096 0.9970 -0.0710 -0.0281 0.0000
0.0096 0.9970 -0.0710 0.0281 0.0000
38. (0.00001) RY (4) H 4
                              s( 0.57%)p99.99( 99.43%)
NHO DIRECTIONALITY AND BOND BENDING (deviation from line of nuclear centers at
                            the position of maximum hybrid amplitude)
      [Thresholds for printing: angular deviation > 1.0 degree]
                           p- or d-character > 25.0%
                           orbital occupancy > 0.10e
                 Line of Centers
                                  Hybrid 1
                                                   Hybrid 2
```

							(cc	ontinued fro	m previous
NBO	Theta	 Phi 	Thet		 ni 	-	Theta		Dev
3. LP (1) O 2 4. LP (2) O 2			90.	О С	0.0				
5. BD (1) C 1- O 2	90.0	0.0	3.	О С	0.0	87.0	178.7	180.0	88.7
COND ORDER PERTURBATI	ON THEORY	Y ANALY:	SIS OF	FOCK	MAT	RIX IN 1	NBO BA	SIS	
Threshold for print	ing: 0	.50 kca	l/mol			E / 2	V E (NIT) _E (I)	F(L,NL)
Donor (L) NBO		Accept	or (NL) NBO					
thin unit 1									
3. LP (1) 0 2	13	3. RY (1) C	1		8.5	58	1.40	0.098
4. LP (2) 0 2	1.3	l. BD*(1) C	1- H	3	26.3	11 :	1.16	0.156
4. LP (2) 0 2	12	2. BD*(1) C	1- H	4	26.3	11 :	1.16	0.156
4. LP (2) 0 2	14	1. RY (2) C	1		5.	72	3.06	0.118
4. LP (2) 0 2		6. RY (0.		3.75	
7. BD (1) C 1- H		2. BD*(4	0.	74	1.42	
7. BD (1) C 1- H		2. RY (2.2		2.12	
8. BD (1) C 1- H		l. BD*(3				0.029
8. BD (1) C 1- H	4 22	2. RY (1) 0	2		2.2	25 :	2.12	0.062
NBO lecular unit 1 (CH2	====== O)		=====			geminal, ======			
1. CR (1) C 1		 L . 99997							
2. CR (1) 0 2	, -	1.99998	-20	.56485	5				
3. LP (1) 0 2	-	1.98853	-0	.81352	2 1	3(v)			
4. LP (2) 0 2	-	1.91757				1(v),12	(v),14	(v),26((g)
5. BD (1) C 1- O		1.99996		.53505					
6. BD (2) C 1- 0		1.99975					, ,		
7. BD (1) C 1- H 8. BD (1) C 1- H		1.99548				2 (v),12			
o. BD (1) C 1- H	4 . 	L.99548 		. /2/03	o Z.	2(v),11	(9)		
9. BD*(1) C 1- O	2 (0.00000	0	.20704	1				
10. BD*(2) C 1- O	2 (0.00000	0	.95146	5				
11. BD*(1) C 1- H	3 (0.03918	0	.69317	7				
12. BD*(1) C 1- H	4 (0.03918	0	.69317	7				
13. RY (1) C 1	(0.00969	0	.58169	9				
14. RY (2) C 1		0.00517		.58837					
15. RY (3) C 1		0.00001		.75652					
16. RY (4) C 1		0.00000		.96046					
17. RY (5) C 1		0.00000		.64510					
18. RY (6) C 1		0.00000		.49223 .24615					
19 RV (7) C 1		0.00000		.08566					
19. RY (7) C 1				.49414					
20. RY (8) C 1				· 1711.					
20. RY (8) C 1 21. RY (9) C 1		0.00000	1	.39676	5				
20. RY (8) C 1 21. RY (9) C 1 22. RY (1) O 2	(0.00368		.39676 .56107					
20. RY (8) C 1 21. RY (9) C 1 22. RY (1) O 2 23. RY (2) O 2	(1	.56107	7				
20. RY (8) C 1 21. RY (9) C 1 22. RY (1) O 2	((0.00368	1 2		7				
20. RY (8) C 1 21. RY (9) C 1 22. RY (1) O 2 23. RY (2) O 2 24. RY (3) O 2	(((0.00368 0.00014 0.00000	1 2 1	.56107 .18160	7) 2				

```
27. RY ( 6) O 2 0.00000 3.20505
28. RY ( 7) O 2 0.00000 2.98918
  29. RY (8) 0 2
                             0.00000
                                        2.69359
                                        3.12898
  30. RY (9) 0 2
                             0.00000
                                        0.41874
  31. RY (1) H 3
                             0.00308
  32. RY (2) H 3
                             0.00002
                                        2.57996
  33. RY (3) H 3
                             0.00002
                                         1.85643
  34. RY (4) H 3
                             0.00001
                                         2.06898
  35. RY (1) H
                4
                             0.00308
                                         0.41874
  36. RY (2) H
                4
                             0.00002
                                         2.57996
  37. RY (3) H
                4
                             0.00002
                                         1.85643
  38. RY (4) H 4
                             0.00001
                                         2.06898
               Total Lewis 15.89671 (99.3545%)
          Valence non-Lewis 0.07835 ( 0.4897%)
          Rydberg non-Lewis 0.02493 ( 0.1558%)
             Total unit 1 16.00000 (100.0000%)
             Charge unit 1 0.00000
$CHOOSE
  LONE 2 2 END
  BOND D 1 2 S 1 3 S 1 4 END
SEND
NBO analysis completed in 0.05 CPU seconds (0 wall seconds)
Maximum scratch memory used by NBO was 297106 words (2.27 MB)
Stopping NBO...Storing NBOs: NBO_1.nbo
                              *** returned from NBO program ***
```

Thus, in this example the NBO analysis of formaldehyde shows that a single Lewis structure is dominant with single bonds between C and H, a double bond between C and O and two lone pairs at the oxygen – just as ordinary chemical arguments would imply. In addition, the program produces the four corresponding valence antibonds. The remaining components of the basis set span the "Rydberg" space and lead to semilocalized, orthogonal orbitals that are assigned to single atoms (Note the nomenclature: BD = bond, BD* = antibond, LP = lone pair, CR = core orbital, RY= Rydberg orbital). The NPA analysis shows a patially negative oxygen and partially positive carbon and hydrogen atoms.

Additionally, the NBO orbitals are stored in the ORCA .gbw file format as jobname.nbo. This file can be used for further analysis and usage with ORCA e.g. for plotting orbitals via orca_plot.

The NBO program has many additional features and analysis tools. The features that are implemented in ORCA can be controlled via the %nbo-block

```
%nbo
NBOKEYLIST = "$NBO
                      ... $END"
DELKEYLIST = "$DEL
                      ... $END"
COREKEYLIST = "$CORE
                       ... $END"
NRTSTRKEYLIST = "$NRTSTR ... $END"
NPEPAKEYLIST = "$NPEPA
                       ... $END"
end
```

The syntax of the respective keylists is given by the NBO6.x/NBO7.x manual.

Specifying the single! NBO keyword corresponds to the %nbo-block

```
%nbo
NBOKEYLIST
              = "$NBO NBO NPA AONBO=C ARCHIVE $END"
end
```

The full set of features beyond those which can be give via the %nbo block can be accessed using the file FILE. 47,

which is generated by the NBO program. This is an ascii file that can be edited with a text editor. Add or remove keywords in the corresponding blocks as needed and call the gennbo program like

gennbo < FILE.47 >jobname.nboout

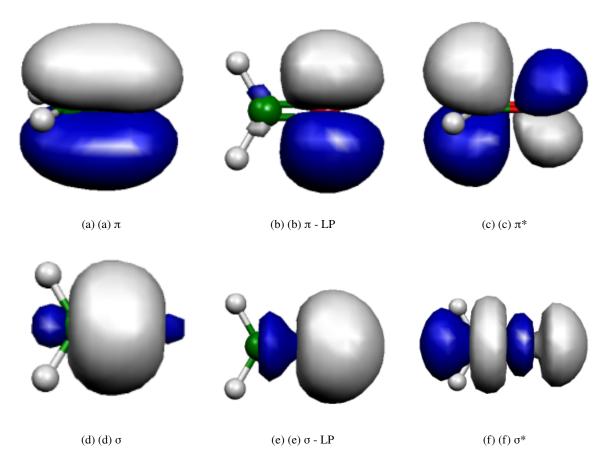


Fig. 5.1: Six NBOs of the H_2CO molecule. Shown are the occupied bonding π and σ orbitals (left) for C and O, the two oxygen lone pairs (middle) and the two π^* and σ^* antibonding orbitals (right).

The FILE.47 file looks like:

	NATOMS= \$END	=4 NBAS=38	UPPER BODM	FORMAT	\$END
\$COORD					
ORCA Jo	ob: check	ζ			
6	6	0.000000	0.00000	0	0.000000
8	8	2.267671	0.00000	0	0.000000
1	1	-1.039349	1.80020	6	0.000000
1	1	-1.039349	-1.80020	6	0.000000
\$END					
\$BASIS					
SBASIS					

If you have no need for this (rather large) file, then you have to delete it manually!

5.2.1 NBO Deletions

An advanced feature, which has been implemented via the ORCA-NBO interface, is the possibility of using deletions.

```
! RHF 3-21G BOHRS TightSCF
%nbo
nbokeylist="$nbo nbo npa aonbo=c archive $end"
delkeylist="$del lewis delete 1 element 3 11 $end"
*xyz 0 1
C
              1.4089705283
                                 0.0210567401
                                                    0.0000000000
N
             -1.3645072652
                               -0.1355759321
                                                   0.000000000
             1.9849776453
                                 1.9986808971
                                                   0.0000000000
                                                   1.6818209547
Н
             2.1492280974
                               -0.9096841007
             2.1492280974
-2.0504340036
-2.0504340036
                               -0.9096841007
                                                   -1.6818209547
                                0.7268536543
                                                  -1.5583845544
Н
                                                    1.5583845544
```

The DELKEYLIST provides NBO with the task to perform certain deletions of orbitals/interactions. Per deletion ORCA calculates a new Fock matrix on basis of an NBO density corresponding to the deletions:

```
Stopping NBO...Starting NBO again for $del instructions...
 LEWIS: Delete all non-Lewis NBOs
 Deletion of the following orbitals from the NBO Fock matrix:
   10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28
 Orbital occupancies:
           Orbital
                                                                       No deletions This deletion Change

      1.99978
      2.00000
      0.00022

      1.99983
      2.00000
      0.00017

      1.97796
      2.00000
      0.02204

      1.99846
      2.00000
      0.00154

      1.99858
      2.00000
      0.00594

      1.99406
      2.00000
      0.00594

      1.99440
      2.00000
      0.00560

      1.99440
      2.00000
      0.00560

      0.0009
      0.00000
      -0.00560

      0.00763
      0.00000
      -0.00763

      0.00763
      0.00000
      -0.00763

      0.00424
      0.00000
      -0.00424

      0.00424
      0.00000
      -0.00424

      0.00034
      0.00000
      -0.00034

      0.00020
      0.00000
      -0.00020

      0.00014
      0.00000
      -0.00014

      0.00034
      0.00000
      -0.00014

      0.00034
      0.00000
      -0.0001

      0.00163
      0.00000
      -0.00016

      0.00163
      0.00000
      -0.00079

                                                                            1.999782.000000.000221.999832.000000.00017
       1. CR (1) C 1
       2. CR (1) N 2
       3. LP (1) N 2
       4. BD (1) C 1- N 2
       5. BD (1) C 1- H 3
       6. BD (1) C 1- H 4
       7. BD (1) C 1- H 5
       8. BD (1) N 2- H 6
      9. BD (1) N 2- H 7
     10. BD*(1) C 1- N 2
     11. BD*(1) C 1- H 3
     12. BD*(1) C 1- H 4
     13. BD*(1) C 1- H 5
     14. BD*(1) N 2- H 6
     15. BD*(1) N 2- H 7
     16. RY (1) C 1
     17. RY (2) C 1
     18. RY (3) C
                                1
     19. RY (4) C
                                1
     20. RY (1) N
                                 2
     21. RY (2) N
                                 2
     22. RY (3) N
     23. RY (4) N
                                 2
     24. RY (1) H
                                                                                                       0.00000
                                 3
                                                                             0.00163
                                                                                                                                 -0.00163
     25. RY (1) H
                                                                                                       0.00000
                                                                             0.00079
                                                                                                                                 -0.00079
                                                                             0.00079
                                                                                                      0.00000
     26. RY (1) H 5
                                                                                                                                 -0.00079
     27. RY (1) H 6
                                                                                             0.00000
                                                                                                                                  -0.00117
                                                                             0.00117
```

```
0.00117 0.00000
  28. RY (1) H 7
                                                                  -0.00117
NEXT STEP: Perform one SCF cycle to evaluate the energy of the new density
           matrix constructed from the deleted NBO Fock matrix.
Copying NBO density...
Calculating new Fock-Matrix...
Calculating Fock-Matrix...done!
New NBO energy via Fock-Matrix: -94.629937
Starting NBO again for $del/return energy instructions...
  Energy of deletion :
Total SCF energy :
                            -94.629936711
                            -94.679444929
                         0.049508 a.u.,
                                                     31.067 kcal/mol
      Energy change :
```

Multiple deletions can also be specified, as can be seen for this example. The output then also contains the additional energy values:

```
Starting NBO again for $del instructions...
Deletion of the following NBO Fock matrix elements:
    3, 11;
Orbital occupancies:
     Orbital
                                    No deletions This deletion Change
                                      ______
   1. CR (1) C 1
   2. CR (1) N
                2
   3. LP (1) N
                2
   4. BD (1) C 1- N 2
   5. BD (1) C
                 1- H
                      3
   6. BD (1) C 1- H
                       4
   7. BD (1) C 1- H 5
   8. BD (1) N 2- H 6
   9. BD (1) N 2- H 7
  10. BD*(1) C 1- N 2
  11. BD*(1) C 1- H 3
  12. BD*(1) C 1- H 4
  13. BD*(1) C 1- H 5
  14. BD*(1) N 2- H 6
  15. BD*(1) N 2- H 7
  16. RY (1) C 1
  17. RY (2) C 1
  18. RY (3) C 1
  19. RY (4) C 1
  20. RY (1) N 2
  21. RY (2) N 2
  22. RY (3) N
                2
  23. RY (4) N
                 2
  24. RY (1) H
                 3

      0.00163
      0.00022

      0.00079
      0.00083

      0.00079
      0.00083

      0.00117
      0.00118

  25. RY (1) H
                                                                   0.00005
                 4
  26. RY (1) H
                                                                   0.00005
                5
  27. RY (1) H 6
                                                                   0.00000
```

```
0.00117 0.00118
  28. RY (1) H 7
                                                                   0.00000
NEXT STEP: Perform one SCF cycle to evaluate the energy of the new density
            matrix constructed from the deleted NBO Fock matrix.
Copying NBO density...
Calculating new Fock-Matrix...
Calculating Fock-Matrix...done!
New NBO energy via Fock-Matrix: -94.668383
Starting NBO again for $del/return energy instructions...
                            -94.668383268
  Energy of deletion :
    Total SCF energy:
                            -94.679444929
                                                      6.941 kcal/mol
      Energy change :
                              0.011062 a.u.,
```



Note that deletions are only implemented for SCF methods!

5.2.2 NBO for Post-HF Densities

NBO analysis can be performed on all methods producing a density. In some methods the density generation has to be specified explicitly, e. g. for MP2 calculations this would be:

```
! MP2 3-21G TightSCF BOHRS NBO
%MP2
density relaxed
end
*xyz 0 1

      1.4089705283
      0.0210567401
      0.0000000000

      -1.3645072652
      -0.1355759321
      0.0000000000

      1.9849776453
      1.9986808971
      0.0000000000

      2.1492280974
      -0.9096841007
      1.6818209547

      2.1492280974
      -0.9096841007
      -1.6818209547

  С
  Ν
  Н

      2.1492280974
      -0.9096841007

      2.1492280974
      -0.9096841007

      -2.0504340036
      0.7268536543

      -2.0504340036
      0.7268536543

  Н
  Н
   Н
                                                                                                                                                                                 -1.5583845544
                                                                                                                                                                                  1.5583845544
   Η
```

The output will contain both the NBO analysis of the SCF density as well as of the MP2 relaxed density. An NBO analysis of a density generated by the MDCI module can be specified as follows:

```
! CISD 3-21G TightSCF BOHRS NBO
%mdci
    density linearized
end
*xyz 0 1
C    1.4089705283    0.0210567401    0.000000000
```

N	-1.3645072652	-0.1355759321	0.000000000
Н	1.9849776453	1.9986808971	0.000000000
Н	2.1492280974	-0.9096841007	1.6818209547
Н	2.1492280974	-0.9096841007	-1.6818209547
Н	-2.0504340036	0.7268536543	-1.5583845544
Н	-2.0504340036	0.7268536543	1.5583845544
*			

Again, the output will contain both the NBO analysis of the SCF density as well as of the CISD linearized density.

5.2.3 Natural Chemical Shielding Analysis (NCS)

For closed-shell calculations of NMR chemical shielding at the SCF level (see sections *NMR Chemical Shifts* and *EPRNMR - keywords for magnetic properties*), the NCS analysis can be requested by adding NCS to the NBOKEYLIST. The NCS keyword accepts the arguments U, I, CSA, XYZ, and MO to analyze the "unperturbed", "induced", anisotropic, Cartesian, and canonical MO contributions to the shielding tensors, respectively, as well as a decimal number for the printing threshold (in ppm). For more information, consult the NBO manual and the original publication [587].

```
! PBE def2-TZVP NMR
%nbo
 NBOKeyList = "$NBO NCS=0.01, I, U, XYZ $END"
end
* xyz 0 1
 Н
        0.00
                  0.00
                           0.00
  С
        1.06
                  0.00
                           0.00
 Ν
        2.23
                  0.00
                            0.00
```

```
Summary of isotropic NMR chemical shielding
Total Lewis (L) and non-Lewis (NL) contributions: (ppm)
                  H 1 C 2 N 3
  1. C 2(cr) L
                  -0.18 200.26 0.18
              NL
                 -0.02 0.02
                               0.00
              L -0.03 -0.12 235.18
  2. N 3(cr)
                 0.00 0.01 0.02
              NL
                  1.02 -33.00 -151.92
  3. N 3(lp)
              T.
                               12.18
              NL
                   -1.04 1.81
                  25.75
     H 1- C 2
              L
                        -49.28
                               -20.30
                   -1.24
              NL
                         6.10
                                2.26
                         15.40
     C 2- N 3
              L
                   2.29
                                13.66
                   0.02
                          0.00
              NL
                                -0.00
                         15.40
     C 2- N 3
              L
                   2.29
                                13.66
              NL
                   0.02
                         0.00
                                -0.00
                        -77.94 -151.00
  7. C 2- N 3
              L
                   0.46
              NT.
                   0.05
                         -4.41
                                0.95
                  31.59
                         70.70 -60.53
        Lewis
                  -2.21 3.53 15.42
     non-Lewis
        Total 29.38 74.23 -45.11
```

5.3 Fractional Occupation Number Weighted Density (FOD)

Many approximate QC methods do not yield reliable results for systems with significant static electron correlation (SEC) and moreover it is often difficult to predict if the system in question suffers from SEC or not. Existing scalar SEC diagnostics (e.g., the T_1 diagnostic) do not provide any information where the SEC is located in the molecule. Furthermore, often quite expensive calculations have to be performed first (e.g., CCSD) in order to judge the reliability of the results based on a single value. Molecular systems with strong SEC (e.g. covalent bond-breaking, biradicals, open-shell transition metal complexes) are usually characterized by small energy gaps between frontier orbitals, and hence, the existence of many equally important determinants in their electronic wavefunction. This finding is used in the fractional occupation number weighted density (FOD) analysis proposed by Hansen and Grimme.[588] The FOD analysis is based on *finite temperature KS-DFT* where the *fractional occupation numbers* are determined from the Fermi distribution ("Fermi smearing").

$$f_i = \frac{1}{e^{(\varepsilon_i - E_F)/kT_{el}} + 1} \tag{5.25}$$

The central quantity of the FOD analysis is the fractional occupation number weighted electron density (ρ^{FOD}), a real-space function of the position vector r:

$$\rho^{FOD}(r) = \sum_{i}^{N} (\delta_1 - \delta_2 f_i) |\varphi_i(r)|^2$$
(5.26)

 $(\delta_1 \text{ and } \delta_2 \text{ are unity if the level is lower than } E_F \text{ while they are } 0 \text{ and } -1, \text{ respectively, for levels higher than } E_F).$ The f_i represent the fractional occupation numbers $(0 \le f_i \le 1; \text{ sum over all electronic single-particle levels obtained by solving self-consistently the KS-SCF equations minimizing the <math>free$ -electronic energy).

 $\rho^{FOD}(r)$ can be plotted using a pre-defined contour surface value (see Section *FOD Plots*). The integration of ρ^{FOD} over all space yields as additional information a single size-extensive number termed N_{FOD} which correlates well with other scalar SEC diagnostics and can be used to globally quantify SEC effects in the molecule. Accordingly, FOD analysis and plots represent a cost-effective and robust way to identify the 'hot' (strongly correlated) electrons in a molecule and to choose appropriate approximate quantum chemical methods for a subsequent computational study of the systems in question. Based on our experience, the following rules of thumb can be derived:

- no significant ρ^{FOD} : use (double)-hybrid functionals or (DLPNO-)CCSD(T) (single-reference electronic structure)
- significant but rather localized ρ^{FOD} : use semi-local GGA functionals (or hybrid functional with low Fock-exchange, avoid HF or MP2; slight multi-reference character)
- significant and delocalized ρ^{FOD} : use multi-reference methods (or finite temperature DFT; strong multi-reference character)

A Warning

Even though the FOD is intuitive and easy-to-use, it is not a 100% safe diagnostic for MR character and should only be used as a first indication. We recommend to always combine the FOD analysis with more sophisticated multi-reference diagnostics.

🗘 Tip

Mulliken reduced orbital charges based on $\rho^{FOD}(r)$ (see *Mulliken Population Analysis*) offer a fast alternative to get the information of the FOD plot (cf. *Example: p-Benzyne*).

5.3.1 Basic Usage

The FOD analysis is a very efficient and practicable tool to get information about the amount and localization of SEC in the system of question and can be invoked via the !FOD simple input keyword.

```
! FOD
```

This will perform the analysis at the default TPSS/def2-TZVP (TightSCF) level that was chosen since it is fast and robust. Besides the standard analysis output (cf. Section 5.3.3), ρ^{FOD} is stored in the file basename.scfp_fod file which is included in the general basename.densities container (for plotting the FOD, see Section 5.3.2).

The FOD analysis can be customized via simple input and the %scf block.

```
! B3LYP def2-TZVP TightSCF

%scf
SmearTemp 9000
end
```

Important

• ρ^{FOD} (and N_{FOD}) strongly depend on the orbital energy gap which itself depends almost linearly on the amount of the non-local Fock exchange admixture a_x (cf. hybrid DFT). The following (empirical) function of the optimal electronic temperature T_{el} on a_x

$$T_{el} = 20000 \,\mathrm{K} \times a_x + 5000 \,\mathrm{K} \tag{5.27}$$

is used to ensure that similar results of the FOD analysis are obtained with various functionals. For example, the SmearTemp has to be 5000 K for TPSS (a_x = 0), 9000 K for B3LYP (a_x = 20%), 10000 K for PBE0 (a_x = 25%), and 15800 K for M06-2x (a_x = 54%).

1 Some Notes on FOD

- The FOD analysis is not strongly dependent on the employed basis set (see supplementary information of the original publication[588]).
- The FOD analysis will be always printed (including Mulliken reduced orbital charges based on ρ^{FOD}) if SmearTemp > 0 K.
- Since the \hat{S}^2 expectation value is not defined for fractional occupation numbers, its printout is omitted.
- The FOD analysis may also be useful for finding a suitable active space for e.g. in CASSCF calculations.

5.3.2 FOD Plots

The fractional occupation number weighted electron density (ρ^{FOD}), see *Fractional Occupation Number Weighted Density (FOD)*) can be plotted in 3D for a pre-defined contour surface value which, after extensive testing, was set to the default value of $\sigma=0.005$ e/Bohr³. In order to allow comparison of various systems this value should be kept fix (in critical cases, one may also check the FOD plot with a a smaller value of $\sigma=0.002$ e/Bohr³ for comparison). The FOD is strictly positive in all space and resembles orbital densities (e.g., π -shape in large polyenes) or the total charge density for an ideal 'metal' with complete orbital degeneracy in simple cases.

Basically, ρ^{FOD} can be plotted analogously to an electron density using the <code>orca_plot</code> utility program and the generated <code>basename.scfp_fod</code> density file that is stored in the <code>basename.scfp_fod</code> is stored in the <code>basename.densities</code> container.

To do so call orca_plot in interactive mode via

```
orca_plot basename.gbw -i
```

The interface will now show various plot options.

```
... MO-PLOT
PlotType
MO/Operator ... 0 0
Output file ... (null)
Format ... Grid3D/B3
Resolution ... 40 40 40
Boundaries ... -22.359
               ... Grid3D/Binary
                     -22.359392 16.311484 (x direction)
                . . .
                      -12.628859 13.574504 (y direction)
                       -7.001812 7.002006 (z direction)
       1 - Enter type of plot
       2 - Enter no of orbital to plot
       3 - Enter operator of orbital (0=alpha,1=beta)
       4 - Enter number of grid intervals
       5 - Select output file format
       6 - Plot CIS/TD-DFT difference densities
       7 - Plot CIS/TD-DFT transition densities
       8 - Set AO(=1) vs MO(=0) to plot
       9 - List all available densities
      10 - Perform Density Algebraic Operations
      11 - Generate the plot
      12 - exit this program
Enter a number:
```

To check all available densities in the basename.densities container one can use the 9 - List all available densities option.

```
List of density names

Index:

Name of Density

O:

basename.scfp_fod <--- required for FOD plot

1:

basename.scfp

2:

basename.scfr
```

The most general way to create the FOD plot is now to create a . cube file that can be visualized with many external programs like ChimeraX or Chemcraft (cf. *Graphical User Interfaces*).

To do so we use the following subsequent user inputs to orca_plot:

```
1 (type of plot)
2 (electron density)
n (default name: no)
basename.scfp_fod (name of the FOD file)
4 (number of grid intervals)
120 (NGrid)
5 (output file format)
7 (cube)
10 (generate plot)
11 (exit)
```

Warning

Note that producing .cube files can become very large and their generation with orca_plot may take a considerable amount of time for larger molecules. This is particularly the case if high quality plots using tight

grid resolution settings (i.e., 120x120x120 resolution) are used for publication purposes.

It is also possible to generate *.cube files from ρ^{FOD} (analogously to electron density plots) with other programs that can read ORCA's baseame.gbw and electron density files by simply using the basename.scfp_fod file instead of the basename.scfp file.

An example FOD plot for Ni(bis-dithiolene) is shown below (Fig. 5.2). It was generated with ChimeraX and a . cube file generated by $orca_plot$ and clearly shows the strong FOD on the metal center and the adjacent ligands. This observation is in line with various studies on the multi-reference character of this complex.

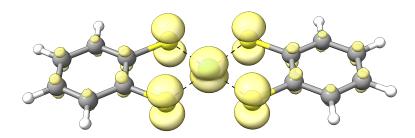


Fig. 5.2: FOD plot at $\sigma = 0.005$ e/Bohr³ (TPSS/def2-TZVP (T = 5000 K) level) for the NiBDT complex (FOD depicted in yellow).

More examples of FOD plots generated with the same setup can be found in the original publication and corresponding supplementary information. [588]

5.3.3 Example: p-Benzyne

In this example, the FOD analysis for the ground state of p-Benzyne is performed.

```
! FOD

* xyz 0 1
C 0.0000000 1.2077612 0.7161013
C 0.0000000 0.0000000 1.3596219
C 0.0000000 -1.2077612 0.7161013
C 0.0000000 -1.2077612 -0.7161013
C 0.0000000 0.0000000 -1.3596219
C 0.0000000 1.2077612 -0.7161013
H 0.0000000 2.1606260 1.2276695
H 0.0000000 -2.1606260 -1.2276695
H 0.0000000 2.1606260 -1.2276695
H 0.0000000 2.1606260 -1.2276695
**
```

The respective output reads:

```
ORCA LEAN-SCF
memory conserving SCF solver

(continues on next page)
```

```
-----D-T-T-S-----
          Energy (Eh)
                              Delta-E RMSDP
                                                  MaxDP DIISErr Damp _
Iteration
→Time(sec)
             *** Starting incremental Fock matrix formation ***
       -230.8982516003082139 0.00e+00 5.01e-03 1.01e-01 1.12e-01 0.700 _
→1.7
Warning: op=0 Small HOMO/LUMO gap ( -0.021) - skipping pre-diagonalization
        Will do a full diagonalization
        -230.9463607195993120 -4.81e-02 1.15e-03 2.59e-02 4.06e-02 0.700 L
→1.6
                            ***Turning on AO-DIIS***
 ... etc.
                               -5.02e-09 3.33e-07 7.37e-06 7.95e-06 0.000 _
  12
      -231.0033984839932089
\hookrightarrow 1.1
               **** Energy Check signals convergence ****
FOD:
Fermi smearing: E(HOMO(Eh)) = -0.201252 \text{ MUE} = -0.179318 \text{ gap} = 1.119 \text{ eV}
N_FOD = 0.920364
```

The high N_{FOD} already indicates strong SEC and checking the Mulliken reduced orbital charges based on $\rho^{FOD}(r)$ (see *Mulliken Population Analysis*) gives a first impression about the localization of hot electrons in the molecule. The printout for the first carbon atom is given below:

```
FOD BASED MULLIKEN REDUCED ORBITAL CHARGES
        s : 0.006371 s: 0.006371 pz : 0.016375 p: 0.030785 px : 0.004516 dz2 : 0.004248 d: 0.010308 dxz : 0.004254 dyz : 0.004855 dx2y2 : 0.004855 dx2y2 : 0.000860 dxy : 0.000091 f0 : 0.000006 f: 0.000378
   0 C s
                            0.000014
         f+1
                      :
         f-1
                             0.000002
                     :
         f+2
                            0.000006
         f-2
         f+3
                             0.000010
                     :
         f-3 : 0.000032
```

If other population analysis printouts are wanted the user is referred to the Löwdin analysis (*Löwdin Population Analysis*) which is turned on by default using the total SCF density of the calculation, also in the case of finite electronic temperature.

The plot the ρ^{FOD} for p-Benzyne clearly shows the significant and rather delocalized FOD (^1A_g) , thus indicating that multi-reference methods would be needed for reliable computational study of this molecule.

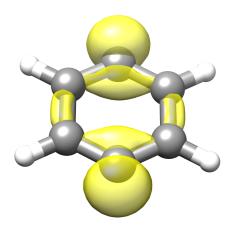


Fig. 5.3: FOD plot at $\sigma = 0.005$ e/Bohr³ (TPSS/def2-TZVP (T = 5000 K) level) for the 1A_g ground state of p-benzyne (FOD depicted in yellow).

5.4 Excited States Calculations

A plethora of methods to compute excited states exists in ORCA. Here we give a brief overview of the main methods, with focus on the basic usage and current capabilities of each. Detailed documentation for each method is indicated in each subsection.

Multi-reference methods, such as NEVPT2 or MRCI, are described elsewhere (*N-Electron Valence State Perturbation Theory (NEVPT2)*, *Multireference Configuration Interaction and Pertubation Theory (uncontracted)*) in the manual.

5.4.1 Excited States with RPA, CIS, TD-DFT and SF-TDA

ORCA features a module to perform TD-DFT, single-excitation CI (CIS) and RPA. The module works with either closed-shell (RHF or RKS) or unrestricted (UHF or UKS) reference wavefunctions. For DFT models the module automatically chooses TD-DFT and for HF wavefunctions the CIS model. If the RI approximation is used in the SCF part it will also be used in the excited states calculation. A detailed documentation is provided in section *Excited States via RPA*, CIS, TD-DFT and SF-TDA

General Use

In its simplest form it is only necessary to provide the number of roots sought:

```
! BP86 def2-SVP TightSCF
%tddft
 Nroots
         1.0
 triplets true
end
*xyz 0 1
 С
       0.016227
                   -0.000000
                               0.000000
                               -0.00000
 0
       1.236847
                   0.000000
 Н
       -0.576537
                   0.951580
                               -0.000000
 Η
       -0.576537
                   -0.951580
                               -0.00000
```

Which gives the following output: