

openLOWDIN

Release 0.1

openLOWDIN

CONTENTS:

1	About 1.1 Capabilities	3 3 3 3
2	Developers and contributors 2.1 Active developers (alphabetical order)	5 5 5
3	Installation	7
4	Getting started 4.1 Input file	9 9 10
5	Code	11
6	Input	13
7	CONTROL	15
8	Scratch	17
9	Outputs	19
10	Lib	21
11	Integrals	23
12	Hartree-Fock, HF	25
13	SCF	27
14	Properties	29
15	Density Functional Theory, DFT	31
16	Many-Body Perturbation Theory, MBPT	33
17	Propagathor Theory, PT	35
18	Configuration Interaction, CI	37

19	Non-Orthogonal Configuration Interaction, NOCI	39
20	Tutorials	41
21	Positronic systems	43
22	Positron covalent bond	45
23	Quantum Nuclei	51
24	Negative Muons	53
25	Quantum Drude Oscillators, QDOs	55
26	External Potentials	57
27	Indices and tables	59

openLOWDIN is a computational program that implements the Any Particle Molecular Orbital (APMO) method to study systems containing any type and number of quantum species, such as electrons, positrons, quantum nuclei, muons, or drude oscillators.

This manual is still in early construction stage! Thanks for your patience.

CONTENTS: 1

2 CONTENTS:

CHAPTER

ONE

ABOUT

openLOWDIN is a computational program that implements the Any Particle Molecular Orbital (APMO) method to study systems containing any type and number of quantum species, such as electrons, positrons, quantum nuclei, muons, or drude oscillators.

At present, openLOWDIN code is publicly available at https://github.com/efposadac/openLOWDIN

1.1 Capabilities

What can we do in Lowdin? The current version of the code encompasses the following quantum chemistry methods:

- HF
- DFT
- MP2
- CI (CIS, CISD, CIST, CISDTQ, FCI, CIPSI)
- PT (PT2, PT3, PP3, RENPP3, OVGF)
- NOCI

Check more details in the Code and Tutorials

1.2 Documentation

The online manual of openLOWDIN is available at https://github.com/openLOWDIN/openLOWDIN_manual A compiled pdf version of the manual can be found here openlowdin.pdf

1.3 How to cite:

Please cite the code as:

R.i Flores-Moreno, E. Posada, F. Moncada, J. Romero, J. Charry, M. Díaz-Tinoco, S.A. González, N.F. Aguirre, A. Reyes, LOWDIN: The any particle molecular orbital code. Int. J. Quantum Chem.. 114. (1), 50–56 (2014).

1.4 Acknowledgements:

This code results from an evolution of the APMO and LOWDIN software packages, both mainly developed at the Universidad Nacional de Colombia. Currently, the code is under an open-source initiative thanks to the support of some initial developers, who are presently spread around the world!

4 Chapter 1. About

DEVELOPERS AND CONTRIBUTORS

2.1 Active developers (alphabetical order)

- · Andrés Reyes
- Edwin Posada
- Jorge Charry
- · Félix Moncada

2.2 Contributors (alphabetic order)

- Jhonathan Romero (APMO/Propagators)
- Sergio Gonzalez (APMO: Older versions)
- Nestor Aguirre (APMO: Older versions)
- Danilo González Forero (APMO/COSMO)
- Jose Mauricio Rodas Rodriguez (APMO/(QM/MM))
- Carlos Ortiz-Mahecha (APMO/CC)
- Alejandro Peña Torres (APMO/CC)
- Laura Pedraza-González (APMO/core)
- Manuel Diaz (APMO/Propagators)
- Teresa Tamayo-Mendoza (Development of higher order propagator methods)
- Roberto Flores-Moreno (ADFT/ADPT/Propagators)

CHAPTER

THREE

INSTALLATION

Compile: (see below for a step-by-step example)

- run ./configure in LOWDIN root directory. Be sure that you have permissions to write in the installation directory and have properly exported the \$PATH environment.
- run make

Install:

• run make install

Uninstall

• run make uninstall

Documentation

• run make doc

The *make doc* command produces both latex and html documentation using doxygen program. Be sure you have installed doxygen, for instance in a debian-based distribution run:

apt-get install doxygen graphviz

To use latex documentation in doc/latex folder, run command:

pdflatex refman.tex

To visualize the html documentation use:

<web browser> doc/html/index.html

Clean the project

• run make clean and then make distclean

Step-by-step installation example: (replace apt-get with your preferred package manager)

sudo apt-get update sudo apt-get -y install wget git build-essential liblapack-dev libblas-dev libgsl0-dev autotools-dev automake libtool gfortran python3 gawk libeigen3-dev libgmp-dev libboost-all-dev # Define ENV Variables export WORKDIR=\$PWD/dependencies export PATH=\$PATH:\$WORKDIR/bin export C_INCLUDE_PATH=\$C_INCLUDE_PATH:\$WORKDIR/include:\$WORKDIR/include/libint2:/usr/include/eigen3 export CPLUS_INCLUDE_PATH=\$CPLUS_INCLUDE_PATH:\$WORKDIR/include:\$WORKDIR/include/libint2:/usr/include/export LIBRARY_PATH=\$LIBRARY_PATH:\$WORKDIR/lib export LD_LIBRARY_PATH=\$LD_LIBRARY_PATH:\$WORKDIR/lib # Create work directories mkdir \$WORKDIR/bin mkdir \$WORKDIR/bin mkdir \$WORKDIR/lib cd \$WORKDIR

Libint2 # If you have Ubuntu, you can get this precompiled Libint2 library wget https://www.dropbox.com/s/d3d44j238lkfwcr/libint-master-SEP052019.tgz tar xzvf libint-master-SEP052019.tgz

Otherwise, download and compile with minimal (default am), G12, fPIC options (libint2 commit 668b10c4bdca5876984058742d4212675eb93f3f) # git clone https://github.com/evaleev/libint.git # cd libint # git checkout 668b10c4bdca5876984058742d4212675eb93f3f # ./autogen.sh # mkdir ../build # cd ../build # ../libint/configure –prefix=\$WORKDIR –with-max-am=6 –enable-g12=4 –with-g12-max-am=4 –with-cxxgen-optflags # make -j 4 # make install # ../libint/configure –prefix=\$WORKDIR

cd -

Libint1 git clone https://github.com/evaleev/libint.git cd libint git checkout v1 aclocal -I lib/autoconf autoconf ./configure –prefix=\$WORKDIR make -j 4 make install make clean make distclean cd -

Libxc cd \$WORKDIR # If you have Ubuntu, you can get this precompiled Libxc library wget https://www.dropbox.com/s/6cja3zzhl1cq46i/libxc-master-MAY242023.tgz tar xzvf libxc-master-MAY242023.tgz # Otherwise, download and compile with default options (libxc commit 4bd0e1e36347c6d0a4e378a2c8d891ae43f8c951) # git clone https://gitlab.com/libxc/libxc.git # cd libxc # git checkout 4bd0e1e36347c6d0a4e378a2c8d891ae43f8c951 # autoreconf -i # ./configure –enable-shared –prefix=\$WORKDIR # make -j 4 # make install

cd.

Configure Lowdin ./configure -p \$WORKDIR/bin -s /tmp -1 "-lblas -llapack" # Build Lowdin make -j 4 # Install Lowdin make install # Run Tests make test

CHAPTER

FOUR

GETTING STARTED

Let's get ready to run openLOWDIN. Here you can find the basic information about the input and how to run the code. A more exhaustive description of all code keywords and files can be found in section

4.1 Input file

The code requires an text input file with extension .lowdin. Here is an example of a minimum input file

```
GEOMETRY
e-[0]
        6-31G
                                0.0000
                                                0.0000
                                                                 0.1173 multiplicity=2
→addParticles=-1
        6-31G
                                0.0000
                                                               -0.4692
e-[H]
                                                0.7572
e-[H]
        6 - 31G
                                0.0000
                                               -0.7572
                                                                -0.4692
                                  0.0000
          13S.ET.O.u.TF
                                                  0.0000
                                                                   0.1173
u-
         dirac
                                 0.0000
                                                 0.0000
                                                                  0.1173
                                                                 -0.4692 \text{ m} = 1836.15
H-a_1
        DZSPDN
                                 0.0000
                                                 0.7572
H-b 2
             DZSPDn
                                      0.0000
                                                      -0.7572
                                                                       0.4692 \text{ m} = 1836.15
END GEOMETRY
TASKS
        method = "UHF"
        propagatorTheoryCorrection = 2
END TASKS
CONTROL
        iterationScheme = 3
        convergenceMethod = 1
        readCoefficients = F
        hartreeProductGuess = T
END CONTROL
```

The minimum required blocks to run a calculation are GEOMETRY}, TASKS, and CONTROL.

The GEOMETRY block provides the information needed to build the molecular system. The first column declares the type of the quantum species. As shown in the above example, e-[H] and e-[0] define the electrons of a Hydrogen and a Oxygen atom respectively; U- defines a negative muon, 0_{16} , H_1 and H_2 define ^{16}O , 1H and 2H nuclei respectively.

The second column declares the basis sets. When the dirac basis is chosen, the particle is treated as a classical point charge. The third, fourth and fifth columns declare the \$x,y,z\$ coordinates of the particle basis set center.

The sixth column provides additional information via keywords addParticles and multiplicity. These keywords are used to change the default values. addParticles is used to modify the number of particles of a quantum species. As shown in the provided example, one electron is removed from the system. multiplicity defines the multiplicity for open shell calculations. In the example, an electronic multiplicity of 2 was chosen.

4.2 How to run

To run LOWDIN

lowdin2 -i inputname.lowdin

This will generate a plain text output file called inputname.out

CHAPTER

FIVE

CODE

This sections summarizes the multicomponent version of the common quantum chemistry methods and capabilities implemented in openLOWDIN, as well as of all available input code keywords.

- Input
- CONTROL
- Scratch
- Outputs
- Lib
- Integrals
- Hartree-Fock, HF
- SCF
- Properties
- Density Functional Theory, DFT
- Many-Body Perturbation Theory, MBPT
- Propagathor Theory, PT
- Configuration Interaction, CI
- $\bullet \ \textit{Non-Orthogonal Configuration Interaction, NOCI}$

12 Chapter 5. Code

СНАРТЕ	R
SIX	K

INPUT

14 Chapter 6. Input

CHAPTER SEVEN

CONTROL

CHAPTER **EIGHT**

SCRATCH

18 Chapter 8. Scratch

CHAPTER	
NINE	
MINE	

OUTPUTS

20 Chapter 9. Outputs

CHAPTE	R
TEI	1

LIB

22 Chapter 10. Lib

CHAPTER **ELEVEN**

INTEGRALS

CHAPTER TWELVE

HARTREE-FOCK, HF

CHAPTER THIRTEEN

SCF

28 Chapter 13. SCF

CHAPTER FOURTEEN

PROPERTIES

СНАРТЕЯ	
FIFTEEN	

DENSITY FUNCTIONAL THEORY, DFT

CHAPTER	
SIXTEEN	

MANY-BODY PERTURBATION THEORY, MBPT

CHAPTER SEVENTEEN		
	QUARTER	
SEVENTEEN	CHAPTER	
	SEVENTEEN	

PROPAGATHOR THEORY, PT

CHAPTER

EIGHTEEN

CONFIGURATION INTERACTION, CI

The APMO/CI wave function is written as a linear combination of CI configurations between all quantum species

$$|\Phi_{0}\rangle = c_{0}|\Psi_{0}\rangle + \sum_{\alpha} \sum_{ia \in \alpha} c_{i}^{a}|\Psi_{i}^{a}\rangle + \sum_{\alpha,\beta} \sum_{\substack{ia \in \alpha \\ jb \in \beta}} c_{ij}^{ab}|\Psi_{ij}^{ab}\rangle + \sum_{\alpha,\beta} \sum_{\substack{ia \in \alpha \\ jb \in \alpha \\ kc \in \beta}} c_{ijk}^{abc}|\Psi_{ijk}^{abc}\rangle + \cdots$$

$$(18.1)$$

CHAPTER	
NINETEEN	

NON-ORTHOGONAL CONFIGURATION INTERACTION, NOCI

CHAPTER TWENTY

TUTORIALS

In this section you can find multiples examples for openLOWDIN

POSITRONIC SYSTEMS

This is an example on how to run a Configuration Interaction, CI calculations for PsH

```
GEOMETRY
        e-(H)
                SHARON-E-6S2P
                                      0.00
                                                   0.00
                                                                0.00 addParticles=1
       Н
                 dirac
                                                           0.00
                                                                        0.00
                                               0.00
                  SHARON-E+6S2P
        e+
                                       0.00
                                                    0.00
                                                                0.00
END GEOMETRY
TASKS
        method = "UHF"
        configurationInteractionLevel ="FCI"
        !configurationInteractionLevel ="CISD"
END TASKS
CONTROL
readCoefficients=F
numberOfCIstates=3
CINaturalOrbitals=T
        CIStatesToPrint = 1
        !CIdiagonalizationMethod = "DSYEVX"
        CIdiagonalizationMethod = "JADAMILU"
        !CIPrintEigenVectorsFormat = "NONE"
        CIPrintEigenVectorsFormat = "OCCUPIED"
        !CIPrintEigenVectorsFormat = "ORBITALS"
        CIPrintThreshold = 5e-2
       buildTwoParticlesMatrixForOneParticle=T
END CONTROL
INPUT_CI
        species="E-ALPHA" core=0 active=0
        species="E-BETA" core=0 active=0
        species="E+" core=0 active=0
END INPUT CI
```

POSITRON COVALENT BOND

This is an example on how to compute the binding energy of a dihydride positron-bound system, using *Configuration Interaction*, CI calculations, as was done in https://doi.org/10.1002/anie.201800914

$$H^-e^+H^- - - > PsH + H^-$$

This input computes the energy of the dihydride system

```
!The goal of this calculation is to compute the binding energy of a positron bound.
!Reported:
!E(e+H2^2-): -1.279680 a.u.
SYSTEM_DESCRIPTION='e+H2^2- from Charry 2018 (10.1002/anie.201800914)'
!add two electrons (one for each hydrogen anion)
!remove one positron
GEOMETRY
                AUG-CC-PVDZ
                                    0.00
                                                0.00
                                                             -1.6 addParticles=1
        e-(H)
                AUG-CC-PVDZ
                                    0.00
                                                0.00
                                                             1.6 addParticles=1
        e-(H)
                  E+-H-AUG-CC-PVDZ
                                                                  -1.6
                                          0.00
                                                      0.00
        6+
                                                                  1.6 addParticles=-1
                  E+-H-AUG-CC-PVDZ
                                          0.00
                                                      0.00
       Н
                 dirac
                                      0.00
                                                  0.00
                                                               -1.6
       Η
                 dirac
                                      0.00
                                                  0.00
                                                               1.6
END GEOMETRY
!method to solve the SCF - CI only works for unrestricted reference
!CI level strings chooses the desired excitations to be included. FCI is all possible...
→excitations
TASKS
        method = "UHF"
        configurationInteractionLevel ="FCI"
        !configurationInteractionLevel ="CIS", "CISD", "CISDT", "CISDTQ"
END TASKS
!Compute only the "numberOfCIstates" states. Here we only need the ground state
!Compute the density matrix for "CIstatesToPrint" states, for density outputs
!Generate the natural orbitals, for visualization in molden files
!The Davidson diagonalization implemented in JADAMILU is the recomended method.
!For small systems, full matrix diagonalization with DSYEVX is possible
!CI EigenVectors with coefficient higher than "CIPrintThreshold" are printed
```

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```
!Printing format "OCCUPIED" shows the coefficients, "ORBITALS" shows the strings, "NONE" ...
→skips printing
!Strict SCF convergence improves the quality of the CI results (not required for the FCI)
CONTROL
        numberOfCIstates=1
        CIStatesToPrint=1
        CINaturalOrbitals=T
        CIdiagonalizationMethod = "JADAMILU"
        !CIdiagonalizationMethod = "DSYEVX"
        CIPrintEigenVectorsFormat = "OCCUPIED"
        !CIPrintEigenVectorsFormat = "NONE", "ORBITALS"
        CIPrintThreshold = 5e-2
        !totalEnergyTolerance=1E-12
END CONTROL
!INPUT_CI block help us define the frozen core and active virtuals orbitals. Here we are.
→not restricting the excitation space
INPUT_CI
        species="E-ALPHA" core=0 active=0
        species="E-BETA" core=0 active=0
        species="POSITRON" core=0 active=0
END INPUT_CI
!With CI, moldenFiles, 1D and 2D density slices and density cubes are good ways to...
⇒visualize the density results
OUTPUTS
        moldenFile state=1
        densityPlot dimensions=2 point1=0.0 0.0 -6.0 point2=0.0 0.0 6.0 state=1
        densityPlot dimensions=3 point1=0.0 -6.0 -6.0 point2=0.0 -6.0 6.0 point3=0.0 6.0
\rightarrow-6.0 state=1
END OUTPUTS
```

Then, we have to subtract the energy obtained from calculations of the dissoc

```
!The goal of this calculation is to compute the binding energy of a positron bound.
!Reported:
!E(PsH): -0.734559
SYSTEM_DESCRIPTION='PsH from Charry 2018 (10.1002/anie.201800914)'
GEOMETRY
               AUG-CC-PVDZ
                                   0.00
                                               0.00
                                                             0.00 addParticles=1
        e-(H)
                 E+-H-AUG-CC-PVDZ
                                         0.00
                                                     0.00
                                                                 0.00
        e+
                                                 0.00
                                                               0.00
       Η
                 dirac
                                     0.00
END GEOMETRY
```

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```
!method to solve the SCF - CI only works for unrestricted reference
!CI level strings chooses the desired excitations to be included. FCI is all possible...
→excitations
TASKS
        method = "UHF"
        configurationInteractionLevel ="FCI"
        !configurationInteractionLevel ="CIS", "CISD", "CISDT", "CISDTQ"
END TASKS
!Compute only the "numberOfCIstates" states. Here we only need the ground state
!Compute the density matrix for "CIstatesToPrint" states, for density outputs
!Generate the natural orbitals, for visualization in molden files
!The Davidson diagonalization implemented in JADAMILU is the recomended method.
!For small systems, full matrix diagonalization with DSYEVX is possible
!CI EigenVectors with coefficient higher than "CIPrintThreshold" are printed
!Printing format "OCCUPIED" shows the coefficients, "ORBITALS" shows the strings, "NONE".
→skips printing
!Strict SCF convergence improves the quality of the CI results (not required for the FCI)
CONTROL
        numberOfCIstates=1
        CIStatesToPrint=1
        CINaturalOrbitals=T
        CIdiagonalizationMethod = "JADAMILU"
        !CIdiagonalizationMethod = "DSYEVX"
        !CIdiagonalizationMethod = "ARPACK"
        CIPrintEigenVectorsFormat = "OCCUPIED" !"NONE","ORBITALS"
        CIPrintThreshold = 5e-2
        !totalEnergyTolerance=1E-12
END CONTROL
!INPUT_CI block help us define the frozen core and active virtuals orbitals. Here we are.
→not restricting the excitation space
INPUT_CI
        species="E-ALPHA" core=0 active=0
        species="E-BETA" core=0 active=0
        species="POSITRON" core=0 active=0
END INPUT CI
!With CI, moldenFiles, 1D and 2D density slices and density cubes are good ways to...
→visualize the density results
OUTPUTS
        moldenFile state=1
        densityPlot dimensions=2 point1=0.0 0.0 -6.0 point2=0.0 0.0 6.0 state=1
END OUTPUTS
```

```
!The goal of this calculation is to compute the binding energy of a positron bound.
!Reported:
!E(H-): -0.524029
SYSTEM_DESCRIPTION='H- from Charry 2018 (10.1002/anie.201800914)'
GEOMETRY
        e-(H)
                AUG-CC-PVDZ
                                    0.00
                                                0.00
                                                              0.00 addParticles=1
        Η
                 dirac
                                      0.00
                                                   0.00
                                                                0.00
END GEOMETRY
!method to solve the SCF - CI only works for unrestricted reference
!CI level strings chooses the desired excitations to be included. FCI is all possible...
→excitations
TASKS
        method = "UHF"
        configurationInteractionLevel ="FCI"
        !configurationInteractionLevel ="CIS", "CISD", "CISDT", "CISDTQ"
END TASKS
!Compute only the "numberOfCIstates" states. Here we only need the ground state
!Compute the density matrix for "CIstatesToPrint" states, for density outputs
!Generate the natural orbitals, for visualization in molden files
!The Davidson diagonalization implemented in JADAMILU is the recomended method.
!For small systems, full matrix diagonalization with DSYEVX is possible
!CI EigenVectors with coefficient higher than "CIPrintThreshold" are printed
!Printing format "OCCUPIED" shows the coefficients, "ORBITALS" shows the strings, "NONE"
→skips printing
!Strict SCF convergence improves the quality of the CI results (not required for the FCI)
CONTROL
       numberOfCIstates=1
        CIStatesToPrint=1
        CINaturalOrbitals=T
        CIdiagonalizationMethod = "JADAMILU"
        !CIdiagonalizationMethod = "DSYEVX"
        !CIdiagonalizationMethod = "ARPACK"
        CIPrintEigenVectorsFormat = "OCCUPIED" !"NONE","ORBITALS"
        CIPrintThreshold = 5e-2
        !totalEnergyTolerance=1E-12
END CONTROL
!INPUT_CI block help us define the frozen core and active virtuals orbitals. Here we are.
→not restricting the excitation space
INPUT CI
        species="E-ALPHA" core=0 active=0
        species="E-BETA" core=0 active=0
END INPUT_CI
!With CI, moldenFiles, 1D and 2D density slices and density cubes are good ways to...
\rightarrowvisualize the density results
                                                                            (continues on next page)
```

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OUTPUTS

moldenFile state=1

densityPlot dimensions=2 point1=0.0 0.0 -6.0 point2=0.0 0.0 6.0 state=1 END OUTPUTS

CHAPTER TWENTYTHREE

QUANTUM NUCLEI

CHAPTER	
TWENTYFOUR	

NEGATIVE MUONS

CHAPTER
TWENTYFIVE

QUANTUM DRUDE OSCILLATORS, QDOS

СНАРТЕЯ	ER
TWENTYSIX	IX

EXTERNAL POTENTIALS

CHAPTER

TWENTYSEVEN

INDICES AND TABLES

- genindex
- modindex
- search