

CASPT2 Method: Current Limitations and Benchmarks Valera Veryazov

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Multiconfigurational second-order perturbation method CASPT2 [1] is known as a reliable computational tool for the electronic structure calculations [2]. However, quite often this method is associated with calculations of very small molecules.

Recent development of CASSCF and CASPT2 codes in Molcas 7 package [3,4] allows to extend the limits for the calculations which can be produced with these methods. The purpose of this study is to demonstrate the possibility to use CASSCF/CASPT2 approach for relatively large molecules, and molecular systems.

Methods

The program system MOLCAS is a package for calculations of electronic and structural properties of molecular systems in gas, liquid, or solid phase. MOLCAS contains a number of modern quantum chemical methods for studies of the electronic structure in ground and exited electronic states, including Hartree-Fock, Density Functional Theory, Coupled-Cluster, multiconfigurational SCF (CASSCF) with dynamical electron correlation treated with multi-reference CI or second order perturbation theory (MS-CASPT2).Automatic geometry optimization using analytical gradient techniques are available for HF/DFT and CASSCF wave functions. CASPT2 structures may be obtained using a numerical gradient procedure. These procedure can be used to

Package

obtain equilibrium geometries, transition states, etc. both for ground and excited electronic states.

MOLCAS runs on almost all UNIX-like platforms and also on MS Windows and Mac OS X. It has a straightforward installation from a source code and a set of configuration files for different platforms and compilers. For Linux operating system MOLCAS could be compiled by g77, gfortran, Intel, PGI, NAG, PathScale compilers. The code can benefit from various BLAS/LAPACK

libraries - Atlas, MKL, ACML. MOLCAS 7.2 can use Cholesky decomposition for two-electron integrals [5] in HF/DFT, MP2, CASSCF and CASPT2 codes.

MOLCAS interfaced to various external programs: Tinker (QM/MM), EPCISO (spin-orbit CI), COLUMBUS (multi-reference CI), MOLSIM (MC/MD)

MOLCAS 7.2 contains Graphical User Interface to create input files (MING) and visually select active space for RASSCF/CASSCF calculations (GV).

On going applications within CASSCF/CASPT2 approach:

836

 D_{2h}

CASSCF: 12-on-13



Geometry and electronic structure of exciplex, formed by 1-cyanonaphthalene and pyridine C₁₆N₂H₁₂, Basis set ANO-S-VDZP

Scan of geometries from 1.4Å to 2.8Å between molecules CASSCF: 14-on-14, multistate calculation (4 roots)

Cr₂C₃₆N₄H₄₀, Basis set Cr:ANO-RCC-VTZP, ANO-RCC-VDZP

A study of the chemical bond in a compound with unusual short Cr-Cr distance



Applications





Interaction between two negatively charged C₆₀ molecules. Study of localization of atomic charges on fullerenes. C_{120}^{-n} , (n=6) Basis set ANO-S-VDZ CASSCF: 18-on-12 (for n=6)

Oxygen displacement in Cu-O plane in HTSC Cu₄O₂₀ (with a set of external charges), Basis Set ANO-RCC-VTZP CASSCF: 8-on-14

Calculations has been performed on two different machines Intel Core2 (2.4GHz, Cache 4Mb, 2Gb RAM), and Intel Xeon (2.83GHz, Cache 6Mb, 64Gb RAM). In test calculations the performance difference between these two machines is about 1.2. Xeon machine has been used for C120 calculations, which require 16Gb of memory. The Molcas code has been compiled by Intel compiler 10, with MKL libraries. The total time of a calculation is approximate, since it depends on an initial guess for CASSCF calculation.

Molecule	Symmery	Basis functions	Number of determinants	Active Orbitals	Frozen in CASPT2	CASPT2 wall time	Total time
C ₁₆ N ₂ H ₁₂	C_1	312	5,891,028	14/14	18	77h (20h for 1-root)	105h
Cr ₂ C ₃₆ N ₄ H ₄₀	C_{2v}	878	369,090	12/13	130	1h	6h
C ₁₂₀	C_i	1080	12,210	18/12	177	3.5h	11h

125,768

Benchmarks

Conclusions

References

There are two components which made possible to extend the limits of CASPT2 calculations in MOLCAS. One is the Cholesky decomposition of integrals, which reduces the amount of data transferred from the disk. Usage of MKL library allows to process BLAS computations (about 80% of the total time spent in CASSCF and CASPT2) on another core(s) of a multicore CPU.

The size of the active space is a hard limit for CASSCF/CASPT2 approach. However, it is now possible to make presize calculations of relatively large molecules even on a "normal" personal computer.

8/14

76

5h

8h

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Cu4O₂₀

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