

Ab Initio Geometry and Bright Excitation of Carotenoids: Quantum Monte Carlo and Many Body Green's Function Theory Calculations on Peridinin

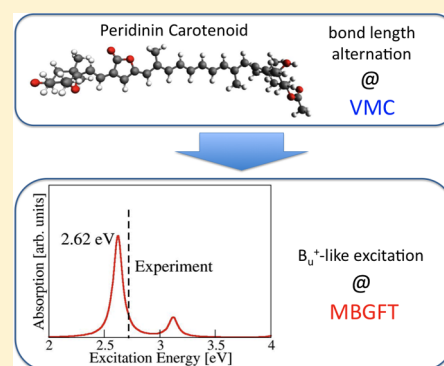
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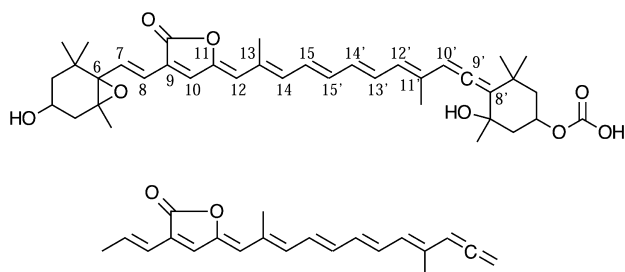
Supporting Information

ABSTRACT: In this letter, we report the singlet ground state structure of the full carotenoid peridinin by means of variational Monte Carlo (VMC) calculations. The VMC relaxed geometry has an average bond length alternation of 0.1165(10) Å, larger than the values obtained by DFT (PBE, B3LYP, and CAM-B3LYP) and shorter than that calculated at the Hartree–Fock (HF) level. TDDFT and EOM-CCSD calculations on a reduced peridinin model confirm the HOMO–LUMO major contribution of the B_u^+ -like (S_2) bright excited state. Many Body Green's Function Theory (MBGFT) calculations of the vertical excitation energy of the B_u^+ -like state for the VMC structure (VMC/MBGFT) provide an excitation energy of 2.62 eV, in agreement with experimental results in *n*-hexane (2.72 eV). The dependence of the excitation energy on the bond length alternation in the MBGFT and TDDFT calculations with different functionals is discussed.



Carotenoids are among the most abundant chemical species in biological systems. Similarly to chlorophylls, they play a fundamental role in light harvesting and energy transfer mechanisms in photosynthetic organisms.¹ In addition, carotenoids get the necessary photoprotective activity of quenching triplet chlorophylls and singlet oxygen molecules.² Although carotenoids are characterized by a large variety in terms of conjugation length of the polyenic chain and of substituents, they share some properties like the state ordering and the character of the singlet and triplet excitations. In the present letter, we consider the carotenoid peridinin (PID, Chart 1), present in the Peridinin-Chlorophyll-a Protein (PCP), a water-soluble complex deriving from marine dinoflagellate *Amphidinium carterae* containing the highest peridinin to chlorophyll-a ratio in nature, namely 4:1 for each domain.³ After a single-photon singlet excitation $S_0 \rightarrow S_2$ (B_u^+ -like, one-photon allowed), the chromophore couples with an

Chart 1. Sketch Representation of Peridinin (PID) and of the Reduced Peridinin Model (PID1)



adjacent chlorophyll-a according to a resonant energy transfer mechanism. Two energy transfer channels have been experimentally detected:^{2–7} the first, accounting for about 25% of the energy,⁶ is the direct transfer from S_2 to the Q_x state of the chlorophyll; the second very efficient route starts from a fast internal conversion from S_2 to S_1 (A_g^- -like, one-photon forbidden), eventually interacting with an intramolecular charge transfer (ICT) state and coupling with Q_y of the chlorophyll.^{8–13} Measurements in several solvents have been reported in the literature:¹⁴ values for S_1 and S_2 states in *n*-hexane (2.0–2.3 and 2.56 eV,^{1a,14} respectively) correspond to the spectral origin, the 0–0 transition, while the vertical Franck–Condon maximum for S_2 is given by the peak at 2.72 eV.^{1a,2,10}

Concerning the nature of the low-lying singlet excitations, the S_1 state of PID is characterized by covalent configurations of double excitation character ($HOMO^2 \rightarrow LUMO^2$) which lower the excitation energy thanks to a favorable mixing with single-electron excitations of $HOMO-1 \rightarrow LUMO$ and $HOMO \rightarrow LUMO+1$ character^{11,15–17} and can be directly investigated by two-photon spectroscopic techniques.^{18,19} $HOMO \rightarrow LUMO$ transition instead dominates the S_2 state.^{11,17} The dark A_g^- state of polyenes with conjugation length $N > 4$ lies in energy below the bright B_u^+ state, thanks to a large singlet–triplet splitting:^{16a} for $N = 4$, the two states are near-degenerate at CASPT2 and CC3 levels,¹⁵ whereas TDDFT²⁰ calculations show an inverted ordering with the

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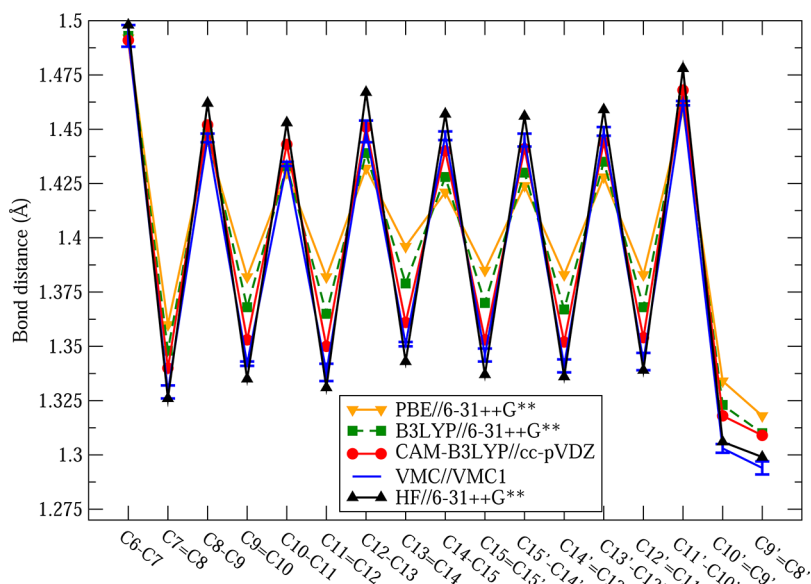


Figure 1. Comparison of the bond length alternation pattern of **PID** from ground state structures optimized at DFT/PBE, DFT/B3LYP, DFT/CAM-B3LYP, VMC, and Hartree–Fock levels. The VMC1 basis is defined in Table 1 of the SI and in refs 25c and 26.

dark A_g^- state higher in energy, independently of the chosen functional.¹⁵ The same occurs for the energies of B_u^+ -like and A_g^- -like states in carotenoids, **PID** included. The same problem, e.g., the difficulty of TDDFT in predicting the correct level ordering, is also found in the characterization of 1L_a and 1L_b excited states of nonlinear acenes.^{16b}

The accurate determination of the relaxed ground state geometry of carotenoids still represents a challenge for quantum chemistry calculations, due to the difficulties in the correct description of electronic correlation of the conjugated polyenic chain. Nevertheless, structural effects play an important role on the spectral tuning of carotenoids in the gas phase, in solution, or in their protein environment, and a reference high-level structure would be desirable. A key structural parameter in the spectral tuning of linear chromophores is the average bond length alternation (BLA), defined as the difference between the average of single bond and double bond distances, excluding (in the case of **PID**) the terminal double bond of the allene group. Differences of a few hundredths of an Ångström in average BLA may significantly alter vertical excitation energies of **PID**, since the molecular orbitals involved in the low-lying region of the absorption spectrum are delocalized along the polyenic chain. Geometries based on DFT calculations strongly depend on the choice of the functional: GGA and B3LYP functionals tend to overestimate the electronic delocalization on the carbon chain,^{21,22} whereas the CAM-B3LYP functional seems to give results in good agreement with experiments for linear all-*trans* polyenes.²² So far, accurate high-level quantum chemistry calculations (multireference or coupled cluster methods) are missing, due to their prohibitive computational cost on such a large system size. Thanks to its capability of exploiting the high performance computing facilities based on massively parallel petaflop machines, Monte Carlo methods²³ can be considered a valid and fully *ab initio* alternative for quantum chemical calculations. We use here the Variational Monte Carlo (VMC) approach^{23,24} to optimize the full **PID** ground state geometry by the Jastrow antisymmetrized geminal power²⁴ wave function

that has been shown to provide accurate results for several molecular properties.^{25,26}

Figure 1 reports the bond lengths of the optimized geometry of the VMC structure (blue in the color version) as well as a comparison with structures obtained at DFT (PBE//6-31++G**, B3LYP//6-31++G**, and CAM-B3LYP//cc-pVDZ) and Hartree–Fock (HF//6-31++G**) levels. In terms of average BLA, the VMC value is 0.1165(10) Å, as expected smaller than the HF limit (0.135 Å) and higher than the PBE (0.066 Å), B3LYP (0.085 Å), and CAM-B3LYP (0.106 Å) values. Whereas the PBE functional produces the smallest difference between single and double bonds, as well-known from theoretical investigations on polyacetylenes,^{21b} the geometrical parameters of the CAM-B3LYP are quite similar to those obtained by VMC, as already observed in the gas phase structure of the retinal protonated Schiff base,²⁶ even though VMC double bonds are seen to be shorter.

The reduced model (**PID1** in Chart 1) is fully representative of the low-lying electronic excitations of **PID**, as already shown by Wagner et al.¹³ Despite standard TDDFT being known to fail in the prediction of the energy inversion between S_1 and S_2 states,¹⁵ it can provide a satisfactory description of the bright excitation. On the other hand, the well behaved performance of TDDFT at the LDA level using the Tamm–Dancoff approximation (TDA) for **PID**¹¹ has been demonstrated to derive from a fortuitous cancellation of errors.^{16a} TDDFT//6-31++G** calculations on **PID1** for the singlet excitations on the five structures give evidence (Tables 7 and 8 in Supporting Information (SI)) of a dominant HOMO → LUMO contribution in the B_u^+ -like state, lower in energy than the A_g^- -like state, independently of the chosen functional and geometry.

Semiempirical MNDO/PSDCI^{10,13,27,28} and EOM-CCSD²⁹ methods show a good agreement with experimental findings in terms of energy values and intensity of the transitions:^{10,13} using the D95 basis on a B3LYP geometry, the EOM-CCSD B_u^+ -like excitation energy has been found to be around 2.5 eV with the A_g^- -like energy close to 2.2 eV, if the MP2 ground state is used as a reference. However, Krylov asserts that doubly

excited dark states in polyenes (A_g^-) are poorly described by EOM-CCSD since the important configurations appear at different levels of excitation, providing an unbalanced description of the excitation.^{29d}

Basis sets and size of the active space can severely affect the convergence of EOM-CCSD excitation energies. Use of the PID1 model was needed in order to reduce the computational effort. An orbital window of 16 highest occupied orbitals and 16 lowest virtual orbitals has been selected, following the procedure of ref 13. A convergence study of the first two excited states with respect to the basis set is reported in Table 1; similar analysis for an 8 + 8 and a 32 + 32 orbital window can

Table 1. EOM-CCSD Excited States Energies (eV) of PID1, Calculated Using a 16 + 16 Orbital Window, As a Function of the Basis Sets^a

geometry	B3LYP	VMC
3-21G	3.84, <i>0.06</i>	4.42, <i>0.36</i>
	4.41, 2.99	4.80, 2.82
6-31G	3.82, <i>0.09</i>	4.38, <i>0.55</i>
	4.32, 3.04	4.72, 2.66
6-311G	3.81, 2.39	4.20, 3.26
	4.08, 1.68	4.69, 0.82
D95	3.84, <i>0.22</i>	4.37, 1.31
	4.20, 3.08	4.65, 2.06
cc-pVDZ	3.87, 2.53	4.26, 3.34
	4.11, 1.34	4.72, 0.54
cc-pVTZ	3.81, 3.51	4.19, 3.72
	4.18, <i>0.63</i>	4.83, 0.39

^aOscillator strengths (*f*) in italics.

be found in Tables 10 and 11 in the SI. At variance with the analysis in ref 13, the excitation energies are consistently referring to the EOM-CCSD ground state energy and not to the MP2 ground state energy.

Calculations on the B3LYP and VMC structures confirm within the EOM-CCSD framework that the B_u^+ -like bright excited state has a dominant HOMO–LUMO contribution, whereas the A_g^- -like state, together with HOMO–1 → LUMO and HOMO → LUMO+1 transitions, has a significant HOMO² → LUMO² double-excitation character. All the excitation energies are, anyway, systematically overestimated, and the use of larger basis sets, like cc-pVDZ and cc-pVTZ, results in an incorrect level ordering, with B_u^+ -like state lower in energy than the A_g^- -like one. When the excitation operators are applied to a larger (32 + 32) orbital window (Table 11 in the SI), EOM-CCSD calculations find the right level ordering for all the considered basis sets for the B3LYP geometry, even if the energies remain too high. The same effect is not observed for the VMC structure, evidence that the geometrical parameters play a fundamental role in the theoretical characterization of low-lying singlet states of peridinin and carotenoids. The failure in finding a robust and clear convergence as a function of the basis set complexity may be ascribed to the small orbital space, to the unbalanced description of configurations with the same excitations within the EOM-CCSD scheme, and to the use of a mean-field (HF) reference. In light of our results, the agreement between the calculations and the experimental excitation reported in ref 13 might be due to a combination of several aspects: the size of the orbital window, the choice of the basis set, the BLA pattern of the ground state geometry, and the fact that the reported

excitations are calculated as energy differences between the EOM-CCSD excited state and MP2 ground state energies.

Moving to the excited states of the full chromophore, Table 2 summarizes the energy and oscillator strengths of the first two

Table 2. TDDFT Excited States Energies (eV) of PID, Calculated Using SVWN, B3LYP, and CAM-B3LYP Functionals^a

geometry	PBE	B3LYP	CAM-B3LYP	VMC	HF
TD-SVWN	2.02, 2.95	2.08, 2.61	2.13, 2.20	2.18, 2.08	2.23, 1.85
	2.22, <i>0.39</i>	2.34, <i>0.61</i>	2.47, <i>0.81</i>	2.55, <i>0.78</i>	2.65, <i>0.91</i>
TD-B3LYP	2.20, 3.45	2.31, 3.29	2.44, 3.03	2.52, 2.95	2.63, 2.79
	2.76, <i>0.20</i>	2.91, <i>0.31</i>	3.09, <i>0.48</i>	3.19, <i>0.52</i>	3.32, <i>0.62</i>
TD-CAM-B3LYP	2.42, 3.66	2.61, 3.62	2.82, 3.55	2.93, 3.53	3.10, 3.51
	3.65, <i>0.09</i>	3.86, <i>0.09</i>	4.07, <i>0.02</i>	4.18, <i>0.00</i>	4.30, <i>0.03</i>

^aThe 6-31++G** basis set has been used for all the calculations. Oscillator strengths (*f*) in italics.

electronic excitations investigated at the TDDFT//6-31++G** level, using the Gaussian 09 package³⁰ employing three different functionals (SVWN, B3LYP, and CAM-B3LYP) on five different geometries (PBE, B3LYP, CAM-B3LYP, VMC, and HF). TDDFT calculations allowed us to investigate the performance of several functionals (LDA, GGA, hybrid, and long-range corrected hybrids) on both geometry and excited states of PID.

A systematic blue shift in the excitation energies is observed when increasing the average BLA, i.e., moving from PBE to HF geometry (Figure 2 and Table 2), as also described by a very

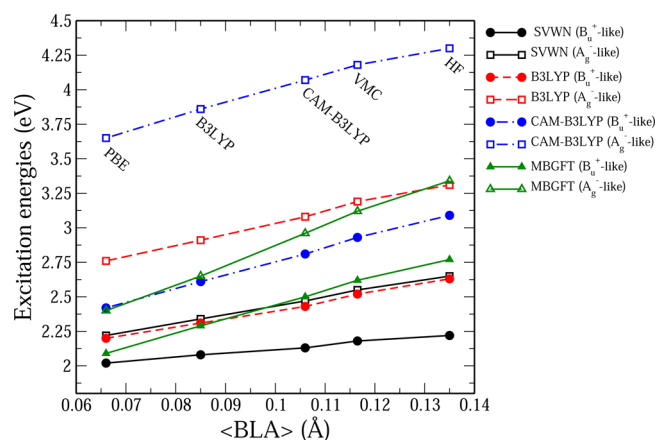


Figure 2. B_u^+ -like and A_g^- -like excited state energies (eV) of PID calculated by TDDFT//6-31++G** with three different functionals (SVWN, B3LYP, and CAM-B3LYP) as a function of the average BLA (Å). MBGFT results are also shown.

recent DFT/MRCI investigation on PID.³¹ In all cases, the order of states appears wrong, the A_g^- -like being, independently of the specific combination of functional and geometry, too high in energy, as already found for the reduced model. Excitations, as expected, are also strongly dependent on the chosen functional. Looking at the B_u^+ -like state (the lowest in energy in our calculations) in Table 2, the SVWN functional tends to underestimate the vertical energy by ~0.5–0.7 eV. TDDFT/B3LYP and TDDFT/CAM-B3LYP on the VMC structure (2.52 and 2.93 eV) differ from the experimental 2.72 eV by ~0.2 eV, respectively underestimating and over-

estimating the excitation energy. None of the TDDFT vertical energies does lead to a satisfactory agreement when the VMC structure is used. Since the choice of the functional influences both the BLA and the excited state energy, different combinations of functionals for the ground state geometry optimization and for the TDDFT calculations can match the experimental value, although the agreement has to be considered fortuitous, as in the case of the CAM-B3LYP excitation on the B3LYP geometry (2.61 eV) and the B3LYP excitation on HF geometry (2.63 eV). On the other hand, the bright excitation energy obtained by using the CAM-B3LYP functional for both geometry optimization and excited states calculation (2.82 eV) is comparable to the experimental absorption, with a discrepancy of 0.1 eV. In spite of the good performance of CAM-B3LYP for the excited state calculations of PID, the same functional seems to overestimate the S_1 state energy of retinal and other positively charged retinal and GFP-like models, as reported in refs 26 and 32.

It is interesting to note that the oscillator strength of B_u^+ -like (A_g^- -like) slightly decreases (increases) for the VMC and HF structures with respect to the DFT geometries; no switch between the two states is however observed. Table 9 in the Supporting Information (SI) reports the main components of the excitations in terms of molecular orbitals: (i) the B_u^+ -like state is dominated by a HOMO \rightarrow LUMO transition; (ii) HOMO-1 \rightarrow LUMO and HOMO \rightarrow LUMO+1 components characterize the A_g^- -like state; (iii) structural changes do affect the transition intensity but not the character of the first two states.

TD-B3LYP estimation of the adiabatic transition to the S_2 state performed on the B3LYP structures corresponds to 2.21 eV, 0.10 eV smaller than the vertical one (2.31 eV, Table 2), fully consistent with the shift of 0.16 eV experimentally observed.²

Since the application of several combinations of functionals (for TDDFT) and ground state structures gives rise to a wide range of values for the bright excitation energy, a step further in the characterization of the B_u^+ -like state is possible by means of the use of the Many Body Green's Function Theory (MBGFT) methods for excited states.³³ Such techniques have been successfully used to describe with high accuracy quasiparticle energies and optical excitations in several materials,³³ including polyenic chains where TDDFT with local and semilocal approximation dramatically fails.³⁴ Due to the high computational cost, very few calculations are present in the literature on biomolecules,³⁵ showing a remarkable agreement with experiments. The MBGFT excitation energies have been calculated using the plane waves Yambo code.³⁶ The absorption spectra have been calculated considering a Kohn–Sham system within the Local Density Approximation (LDA) as a zero order noninteracting Hamiltonian using the Quantum Espresso package.³⁷ The quasiparticle corrections have been evaluated within the G_0W_0 approximation for the self-energy operator, and the electron–hole screened interaction is included by solving the Bethe–Salpeter (BS) equation, where the static screening is calculated within the random-phase approximation. As observed in ref 35 for other biomolecules, we have also found that TDA is not suitable for the PID system. Details on the implementation of the GW/BS calculations can be found in ref 36, and the computational details are reported in the SI. In Table 3, electronic gaps at the Kohn–Sham level and G_0W_0 corrected are reported for all the studied structures, together with the excitation energies calculated at the BS level.

Table 3. Kohn–Sham Gap (LDA), G_0W_0 Gap, and Excited States Energies (for the First Two States) by Solving BS Equation for PID^a

geometry	PBE	B3LYP	CAM-B3LYP	VMC	HF
Kohn–Sham gap	1.20	1.34	1.49	1.56	1.69
G_0W_0 gap	4.16	4.41	4.71	4.89	5.11
excited states (BS)	2.09	2.29	2.50	2.62	2.77
	2.40	2.65	2.96	3.12	3.34

^aAll the energies are in eV.

Quasiparticle energies of the frontier orbitals are shown in Figure 1 of the SI. We observe that the electronic gap, already at the DFT level, increases with the increasing of the BLA, and the effect is enhanced when quasiparticle corrections are included. The blue shift in the excitation energies is also observed at the BS level going through increasing BLA values. Noticeably, the dependence of the B_u^+ -like energy on the average BLA calculated by MBGFT methods is enhanced with respect to TDDFT calculations indicating an even larger role of structural effects in the tuning of absorption properties of PID.

The BS absorption spectra are collected in Figure 2 of the SI. The bright excitation of 2.62 eV on the VMC structure is mainly due to the HOMO \rightarrow LUMO transition (88%), showing good agreement with the B_u^+ -like vertical experimental energy (2.72 eV) measured in *n*-hexane. The latter value can be taken as a reference for gas phase calculations of the vertical excitation to the bright state, assuming small electrostatic coupling between *n*-hexane and PID and zero point energy contributions. Nevertheless, it is worth it to mention that vibronic terms could have an appreciable weight in the applied theoretical model,³⁸ and a deeper and more quantitative insight on these effects requires further investigation. In summary, the use of high-level geometry optimization such as VMC in combination with a converged treatment by first principles of electronic excitations by MBGFT is able to provide an accurate estimation of the bright electronic excitation in PID. In order to correctly describe the double excitation character of the A_g^- -like state in the MBGFT framework, the use of a dynamic kernel in the BS equation is required.³⁹

In this Letter, we have reported the ground state optimized geometry of a carotenoid at the Quantum Monte Carlo level of theory, which is becoming an affordable many-body technique for the study of large molecular systems, where electron correlation plays an important role. The combined use of the VMC structure and MBGFT methods (VMC/MBGFT) represents a fully *ab initio* approach to obtain excitation energies in meaningful agreement with the experimental findings, overcoming the difficulties in the choice of the proper functional in the TDDFT framework and avoiding inclusion of parameters, as in the case of the hybrid CAM-B3LYP. This work can be considered as a needed step for the quantitative investigation of the effects of the protein environment of peridinin molecules and other carotenoids in photosynthetic complexes.

■ ASSOCIATED CONTENT

📄 Supporting Information

VMC wave function; details on MBGFT calculations; bond distances for DFT, VMC, and HF structures; transition components for TDDFT calculations on PID and PID1; MBGFT quasiparticle energies and absorption spectra; convergence tables as a function of basis set and active space

for EOM-CCSD calculations on PID1. This material is available free of charge via the Internet at <http://pubs.acs.org>

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Author Contributions

The manuscript was written through contributions of all authors. All authors have given approval to the final version of the manuscript.

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Notes

The authors declare no competing financial interest.

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