

***N*-electron valence state perturbation theory based on a density matrix renormalization group reference function, with applications to the chromium dimer and poly-p-phenylene vinylene**

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(Dated: 18 December 2015)

The strongly-contracted variant of second order *N*-electron valence state perturbation theory (NEVPT2) is an efficient perturbative method to treat dynamic correlation without the problems of intruder states or level shifts, while the density matrix renormalization group (DMRG) provides the capability to tackle static correlation in large active spaces. We present a combination of the DMRG and strongly-contracted NEVPT2 (DMRG-SC-NEVPT2) that uses an efficient algorithm to compute high order reduced density matrices from DMRG wave functions. The capabilities of DMRG-SC-NEVPT2 are demonstrated on calculations of the chromium dimer potential energy curve at the basis set limit, and the excitation energies of poly-p-phenylene vinylene (PPV).

I. INTRODUCTION

In many quantum chemistry applications, such as to excited states, non-equilibrium geometries, and transition-metal chemistry, the mean-field description of electron interactions is insufficient. The deviation from mean-field behaviour of the electrons, i.e. electron correlation, is usually classified into two types: static or non-dynamic correlation, and dynamic correlation. Conventionally, static correlation is the correlation between near-degenerate valence orbitals, and can be treated by considering the set of possible configurations of electrons in the set of active valence orbitals. If all such configurations are included and the orbitals are simultaneously optimized, one arrives at the complete active space self-consistent-field (CASSCF) model¹⁻³, the most widely used method to describe static correlation. Dynamic correlation is the correlation between valence orbitals and the remaining empty (virtual) or doubly occupied (core) states. This effect is thought of as captured by low particle-rank excitations, such as used in perturbation theories (PT)⁴, configuration interaction singles and doubles (CISD)^{5,6}, or coupled cluster theory (CC)^{4,7-11}. The combined treatment of static and dynamic correlation is the domain of multireference correlation theories, such as multi-reference perturbation theories (MRPT)¹², multi-reference configuration interaction (MRCI)¹³⁻¹⁵ or multi-reference coupled cluster theories (MRCC)¹⁶⁻²².

The density matrix renormalization group (DMRG)²³⁻⁴³ has made it possible to employ large active spaces to describe static correlation^{27,29,30,44-57} and it has become straightforward to use the DMRG as a robust numerical solver in a variety of frontier applications, including benchmark solutions of small molecules^{29,30,50,58}, multi-center transition metal clusters^{49,52,56,59}, as well as to molecular crystals⁶⁰. There have been several efforts to further include dynamic correlation on top of the DMRG reference. For example, DMRG-CT^{61,62}, DMRG-CASPT2⁶³,

DMRG-cu(4)-MRCI⁶⁴ (in an internally contracted (IC) formulation) and MPS-PT⁶⁵ and MPS-LCC⁶⁶ (in an uncontracted formulation) are all attempts in this direction. There have also been efforts to combine DMRG with density functional theory (DFT)⁶⁷. In the internally contracted multireference methods, high order reduced density matrices (RDM) of the DMRG active space wavefunction are required. Such density matrices, although simple in a theoretical sense, are computationally non-trivial to obtain.

In this work, we have developed a general algorithm to efficiently compute high order reduced density matrices from DMRG wave functions. Using these RDM calculations, we may then implement a variety of methods to treat electron correlation external to the DMRG active space. As an example of this, we have implemented the strongly contracted variant of second order *N*-electron valence state perturbation theory (NEVPT2)⁶⁸⁻⁷⁰, an intruder-state-free multireference perturbation theory, on top of the DMRG reference wavefunction, to yield a method we call DMRG-SC-NEVPT2. To demonstrate the potential of this approach, we apply the method to study the potential energy curve of the chromium dimer, and to compute the excitation energies of the quasi-one dimensional conjugated poly(p-phenylene vinylene) (PPV). The chromium dimer is a particularly demanding small molecule that has been widely studied with many techniques^{63,66,71-76}, and thus forms a good test bed to demonstrate the performance of DMRG-SC-NEVPT2. Using an active space with 22 orbitals and basis sets extrapolated to the complete basis set (CBS) limit, we show that we can compute both the spectroscopic constants and the full curve, with an accuracy that compares quite favourably to earlier calculations. PPV on the other hand, is of interest as a prototypical light-emitting polymer⁷⁷⁻⁷⁹. Light emission relies on the correct ordering of the first optically bright 1^1B_u state and dark 2^1A_g state, and describing this ordering requires an appropriate treatment of electron correlation⁸⁰⁻⁸⁵. With

DMRG-SC-NEVPT2, we show that we can compute the excitation energies and energy ordering of the different states accurately, starting from the full-valence active π space of the molecule.

II. THEORIES

A. DMRG wavefunction and optimization algorithm

As with other wavefunction methods in quantum chemistry, DMRG is based on a wavefunction ansatz, namely the matrix product state (MPS). An MPS is a non-linear wavefunction, built from a contraction of tensors for each orbital in the basis. If an MPS is composed of tensors with a limited dimension (called the bond dimension, M) it explores only a physically motivated, but restricted, subset of the Hilbert space. By increasing M , the MPS ansatz will then converge to the full configuration interaction (FCI) result. The DMRG algorithm provides a combination of renormalization and truncation steps that efficiently find a variational and optimal MPS. In practice, two slightly different types of MPS are used in practical DMRG calculations: the one-site and two-site MPS.

The one-site MPS is defined as

$$|\Psi\rangle = \sum_{n_1, n_2, \dots, n_p, \dots, n_k} \mathbf{A}^{n_1} \mathbf{A}^{n_2} \dots \mathbf{A}^{n_p} \dots \mathbf{A}^{n_k} \quad (1)$$

Here, n_i is the occupancy of orbital i , one of $\{|\rangle, |\uparrow\rangle, |\downarrow\rangle, |\uparrow\downarrow\rangle\}$, and k is the number of orbitals. For a given n_i , \mathbf{A}^{n_i} is an $M \times M$ matrix, except for the first and last ones; the dimensions of \mathbf{A}^{n_1} and \mathbf{A}^{n_k} are $1 \times M$ and $M \times 1$ respectively. This ensures that for a given occupancy string $n_1 n_2 \dots n_k$ the $\mathbf{A}^{n_1} \mathbf{A}^{n_2} \dots \mathbf{A}^{n_k}$ product yields a scalar, namely, the coefficient of the determinant $|\mathbf{n}_1 \mathbf{n}_2 \dots \mathbf{n}_k\rangle$.

Similarly, the two-site MPS is defined as

$$|\Psi\rangle = \sum_{n_1, n_2, \dots, n_p, \dots, n_k} \mathbf{A}^{n_1} \mathbf{A}^{n_2} \dots \mathbf{A}^{n_p n_{p+1}} \dots \mathbf{A}^{n_k} \quad (2)$$

The only difference with the one-site MPS, is that one of the tensors ($\mathbf{A}^{n_p n_{p+1}}$) is associated with two sites at a time. The two-site MPS thus has slightly more variational freedom than the one-site MPS, and further exists in different non-equivalent forms, depending on which sites are chosen for the special two-site tensor.

In the DMRG sweep algorithm, $\langle \Psi | H | \Psi \rangle / \langle \Psi | \Psi \rangle$ is variationally optimized, and a single tensor (\mathbf{A}^{n_i} in a one-site MPS or $\mathbf{A}^{n_i, n_{i+1}}$ in a two-site MPS) is updated at site i during the sweep. This update leads to solving an eigenvalue problem of an effective “superblock” Hamiltonian, which is a $4M^2 \times 4M^2$ matrix. The multiplication between this effective Hamiltonian and a MPS is usually computed using $O(k^2)$ “normal” and “complementary” operators built from different sets (blocks) of the active orbitals. The cost of this eigenvalue problem

is $O(k^2 M^3)$ for a single tensor update, and $O(k^3 M^3)$ for a whole sweep optimization.

As discussed, a two-site MPS has more variational freedom than a one-site MPS at the site where the special tensor is placed. This allows the DMRG optimization to change quantum numbers, i.e. conserved local symmetries, during the sweep, and thus helps avoid local minima. Further, the discarded weight associated with the renormalization step in a two-site MPS sweep optimization can be used to extrapolate away the residual error due to finite bond dimension M . However, as the two-site MPS wavefunction at each step of the sweep is slightly different (because the special tensor is moved along the sweep), the two-site formalism does not provide a unique definition of the DMRG expectation values. Thus, to evaluate the reduced density matrix, we complete the DMRG sweep optimization in the one-site representation, to obtain a single DMRG wavefunction. This is referred to as the “two-site to one-site” algorithm (a two-site MPS optimization followed by a one-site one)⁵⁸. From here on when we refer to an MPS, we will mean the one-site MPS.

B. Reduced density matrix (RDM) evaluation algorithm

High order RDM’s are common in post-active space methods that are used to capture the “dynamic” correlation outside of the active space, such as multireference configuration interaction¹⁶ and multireference perturbation theory^{12,70}. The expression of quantities in terms of RDM’s is a feature of the internally contracted formulations, which avoid the expansion of the external correction to the active space wavefunction explicitly in terms of many-particle basis states (such as Slater determinants or configuration state functions). Internally contracted CASPT2¹² and NEVPT2⁷⁰ (both strongly and partially contracted variants) require the four-particle RDM, while higher levels of correlation can require even higher order RDM’s.

In principle, the evaluation of RDM’s in DMRG is straightforward; the evaluation of a matrix element such as $\langle \Psi | a_i^\dagger a_j | \Psi \rangle$ where Ψ is an MPS, is an operation of $O(kM^3)$ cost and can be carried out within any standard DMRG implementation. However, a naive algorithm which evaluates each RDM element separately would lead to far too high a computational cost. Instead, the efficient evaluation of the RDM’s of an MPS requires forming suitable DMRG renormalized operators which can be reused in *all* RDM elements $\langle a_i^\dagger a_j \rangle$. Such efficient algorithms to build the 2-RDM were proposed some years ago by several groups^{86,87}, and require only $O(k^3 M^3)$ cost, as opposed to the $O(k^5 M^3)$ cost arising in a naive element by element evaluation. The efficient algorithm to build the 3-RDM was introduced by Kurashige and Yanai.⁶³ Here we introduce a more general algorithm that automatically organises the necessary DMRG renormalized operators for any order of RDM. The RDM’s we will generally be interested in will be the spin-summed

quantities, e.g. for the 2-RDM, this is

$$\gamma_{ijkl} = \sum_{\sigma, \tau} \langle \Psi | a_{i, \sigma}^\dagger a_{j, \tau}^\dagger a_{k, \tau} a_{l, \sigma} | \Psi \rangle \quad (3)$$

1. General loop over orbital type patterns and number patterns

To organize the efficient and non-redundant evaluation of the RDM elements, we label each element in terms of two classifiers: its “type pattern” (which describes the pattern of creation and annihilation operators, ordered by the sites on the lattice), and its “number pattern” (which describes how to partition the creation and annihilation operators). The complete set of RDM elements can be then be obtained efficiently with proper reuse of intermediates in a DMRG sweep, by looping over all type patterns, and all indices corresponding to given number patterns, at each step of the sweep (with a few restrictions associated with edge cases near the ends of the sweep, described in the appendix), building the DMRG renormalized operators corresponding to the given type and number patterns, and forming the appropriate contractions with the DMRG renormalized wavefunction.

The type pattern is precisely defined as follows. Consider an RDM element $\langle a_i^\dagger a_j^\dagger \dots a_m a_n \dots \rangle$. We reorder the indices such that the indices of the creation and annihilation operators are arranged successively in the order as they appear in the DMRG site ordering: $o_{i'} o_{j'} \dots o_{m'} o_{n'}$, with $i' \leq j' \leq \dots \leq m' \leq n'$ and where o is either a or a^\dagger . The corresponding binary pattern of $2N$ (where N is the rank of the RDM) creation and annihilation operators appearing in the above operator string is the type pattern of the RDM matrix element.

The number pattern encodes how to partition the labels of the expectation value in $\langle a_i^\dagger a_j^\dagger \dots a_m a_n \dots \rangle$ onto different blocks in the DMRG sweep, so as to maximize the reuse of DMRG renormalized operators. Recall that a DMRG sweep consists of a sequence of operations over the orbitals; at each step in the DMRG sweep, there is the set of orbitals preceding the current orbital (the system block (S)), the current orbital (the dot block (D)), and the set of orbitals whose associated site tensors have yet to be optimized (the environment block (E)). The intermediates in the construction of an RDM matrix element are DMRG renormalized operators built and stored on the three blocks. To determine the renormalized operators to build, we partition the $2N$ orbital labels of the operator O among the 3 blocks. The most efficient partitioning (when generating the full set of RDM elements) is to compute a given element from renormalized operators with the maximum number of orbital labels appearing on the dot block, and an equal number of orbital labels on the system and environment blocks. For typical RDM elements (i.e. not for elements such as γ_{0000}), at least one orbital label can be placed on D and at most $N - 1$ orbital labels on E by choosing an appropriate position

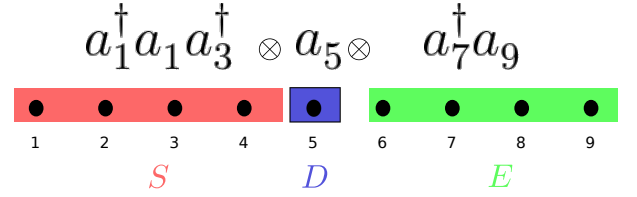


FIG. 1. Evaluation of a three RDM element $\gamma_{137;159}$. $a_1^\dagger a_1 a_3^\dagger$ is on S , a_5 is on D , and $a_7^\dagger a_9$ is on E . This elements belongs to the (3,1,2) “number pattern” and the $(a^\dagger a a^\dagger a a^\dagger a)$ “type pattern”.

of the dot site (i.e. the step in the DMRG sweep). The numbers $\{n_S, n_D, n_E\}$ then denote the numbers of orbital labels distributed on the different blocks, and this is the number pattern. One can generate all valid number patterns by considering all cases where the number of orbital labels on S is no more than N , that on D is no more than 4 and that on E is no more than $N - 1$, and the total is $2N$ except for some edge cases for the first step and last step in the sweep, as discussed in the appendix. The number pattern and type pattern for a representative RDM element is illustrated in Fig. IIB 1.

With the above process, the renormalized operators that need to be constructed on each block S , D , and E in the DMRG sweep, are enumerated by the corresponding set of type patterns and number patterns of the RDM that is being built. In total this leads to $O(k^N)$ renormalized operators that need to be built, most of which are constructed on the system block S at different steps in the sweep. The cost to build these operators is $O(k^N M^3)$ at each step of the sweep and $O(k^{N+1} M^3)$ in total.

2. Computing the expectation values

After building the renormalized operators needed on the three blocks, $[O]^S$, $[O]^D$, $[O]^E$ at site i of the sweep, they are traced with the renormalized DMRG wavefunction, i.e. the site tensor \mathbf{A}^{n_i} . This corresponds to the computation

$$\gamma = \sum_{s n e s' n' e'} A_{se}^{n_i p} [O]_{ss'}^S [O]_{nn'}^D [O]_{ee'}^E A_{s'e'}^{n'_i} \quad (4)$$

where the above tensor contraction is performed in stages (e.g. contracting s , n , e indices separately) and intermediates such as $\sum_{s n e s' n' e'} A_{se}^{n_i} [O]_{ss'}^S A_{s'e'}^{n'_i}$ are reused to minimize the computational cost, and p is a parity operation that inserts appropriate minus signs for fermions²⁹. With appropriate intermediates, the total cost of this step for an N -RDM computation is $O(k^{N+1} M^3 + k^{2N} M^2)$.

3. Spin recoupling

Through the above steps, an expectation value such as $\langle a_0^\dagger a_1^\dagger a_2 a_3 \rangle$ is obtained. For a non-spin-adapted DMRG algorithm, where a^\dagger, a are simple creation and annihilation operators, we can then obtain γ_{0132} by permuting a_2 and a_3 with appropriate signs. For spin-adapted DMRG implementations, the a^\dagger, a are spin tensor creation and annihilation operators, and $\langle a_0^\dagger a_1^\dagger a_2 a_3 \rangle$ is instead an element such as $\langle \{[(a_0^\dagger a_1^\dagger)_{S_1}^{S_1} (a_2)_{D}]^{S_2} (a_3)_E\}^{S_3} \rangle$ with different spins S_1, S_2, S_3 and spin couplings. The spin tensor creation and annihilation operators can be expanded as a linear combination of spin orbital creation and annihilation operators, leading to a set of linear equations to obtain the spin orbital RDM elements from a single spin reduced matrix element. Using singlet embedding^{53,88}, expectation values for the spin tensor combinations with non-zero total spin are zero, significantly decreasing the number of expectation values to compute. The coefficients appearing in the linear equations are the same for all operators with the same pattern and can thus be generated automatically and reused. The cost of this process is independent of the DMRG bond dimension M , and is thus negligible compared to the other parts of the RDM calculation.

C. N -electron valence state perturbation theory

We now briefly review the strongly contracted NEVPT2^{69,70}, a second-order multireference perturbation theory. The target zeroth-order wave function is defined as

$$P_{CAS} H P_{CAS} |\Psi^{(0)}\rangle = E^{(0)} |\Psi^{(0)}\rangle \quad (5)$$

where P_{CAS} is the projector onto the CAS space. The zeroth order Hamiltonian is chosen in the form given by Dyall⁸⁹:

$$H^D = H_i + H_v + C \quad (6)$$

where H_i is a one-electron (diagonal) operator in the non-active (core and external) subspace:

$$H_i = \sum_{i,\sigma}^{core} \epsilon_i a_{i,\sigma}^\dagger a_{i,\sigma} + \sum_{r,\sigma}^{virt} \epsilon_r a_{r,\sigma}^\dagger a_{r,\sigma} \quad (7)$$

and ϵ_i and ϵ_r are orbital energies; H_v is a two-electron operator confined to the active space:

$$H_v = \sum_{ab,\sigma}^{act} h_{ab}^{eff} a_{a,\sigma}^\dagger a_{b,\sigma} + \sum_{abcd,\sigma,\eta}^{act} \langle ab|cd \rangle a_{a,\sigma}^\dagger a_{b,\eta}^\dagger a_{d,\eta} a_{c,\sigma} \quad (8)$$

where $h_{ab}^{eff} = h_{ab} + \sum_i^{core} (2 \langle ai|bi \rangle - \langle ai|ib \rangle)$; and C is a constant to ensure that H^D is equivalent to the full Hamiltonian within the CAS space.

The zeroth order wave functions external to the CAS space are referred to as the “perturber functions”. A perturber function is written as $|\Psi_{l,\mu}^{(k)}\rangle$. It belongs to a CAS space is denoted $P_{S_l^{(k)}}$, where k is the number of electrons promoted to ($k > 0$) or removed from ($k < 0$) the active space, l denotes the occupancy of nonactive orbitals and μ enumerates the various perturber functions in $S_l^{(k)}$. In strongly contracted NEVPT2, only one perturber function for each $S_l^{(k)}$ subspace is used. The perturber function is defined as $|\Psi_l^{(k)}\rangle = S_l^{(k)} H |\Psi^{(0)}\rangle$, where $P_{S_l^{(k)}}$ is the projector onto the space. The energy of the perturber function is

$$E_l^{(k)} = \frac{\langle \Psi_l^{(k)} | H^D | \Psi_l^{(k)} \rangle}{\langle \Psi_l^{(k)} | \Psi_l^{(k)} \rangle} \quad (9)$$

The zeroth order Hamiltonian then becomes

$$H_0 = \sum_{kl} |\Psi_l^{(k)'}\rangle E_l^{(k)} \langle \Psi_l^{(k)'}| + |\Psi^{(0)}\rangle E^{(0)} \langle \Psi^{(0)}| \quad (10)$$

where $|\Psi_l^{(k)'}\rangle$ corresponds to the normalized $|\Psi_l^{(k)}\rangle$.

The bottleneck in SC-NEVPT2 is the evaluation of the energies of the perturber functions, where up to the 4RDM appears. This 4RDM is subsequently contracted with two electron integrals in the active space to form auxiliary matrices.⁷⁰ Computing the 4RDM requires $O(k^8 M^2 + k^5 M^3)$ cost, much higher than that of a standard DMRG energy optimization. To minimize the cost, it is generally necessary to evaluate the 4RDM using a lower bond dimension than is used in the DMRG energy optimization of $|\Psi^{(0)}\rangle$. To obtain an MPS approximation of lower bond dimension for the reference, we “compress” the converged reference MPS of a larger bond dimension. This is done in practice by carrying out a “reverse schedule” set of sweeps in a DMRG code, where M is decreased⁵⁸. All of our DMRG-SC-NEVPT2 calculations use reference MPS that have been compressed in this way to compute the RDM’s. The compressed bond dimension is denoted M' .

Further, to avoid storage problems in our implementation of DMRG-SC-NEVPT2, the 4-RDM is computed on the fly. In principle, it is possible to completely avoid generating the 4-RDM by directly computing the auxiliary matrices (i.e. with the integrals already contracted). This is the approach taken, for example, in Kurashige and Yanai’s original implementation of DMRG-CASPT2⁶³, where the auxiliary matrices can be computed with a cost scaling similar to that of the 3-RDM, using special kinds of DMRG renormalized operators as intermediates. However, the Dyall Hamiltonian in NEVPT2 involves all the active-space two-electron integrals, and this is more complicated to handle than the diagonal Fock energies that appear in the CASPT2 zeroth Hamiltonian (in its canonical representation). Thus, although the scaling of such an algorithm may be lower, it is complicated to implement and the computational prefactor is large due to

the many different intermediates that need to be considered. We have implemented such an algorithm only in our reference NEVPT2 code that uses CASCI type reference wavefunctions, and not in our DMRG-SC-NEVPT2 code.

III. APPLICATIONS

A. Chromium dimer

The chromium dimer has been a challenging molecule for quantum chemistry for many years. This is because a relatively large active (12e, 12o) space is needed to describe the minimal spin-recoupling of the electrons involved in the nominal sextuple bond, while, at the same time, a large amount of dynamic correlation is required even to yield a bound potential energy curve.^{63,66,71,73,74,90–92}

The minimal (12e,12o) CAS, derived from the 3d and 4s atomic orbitals, has been widely employed in earlier chromium dimer calculations.^{66,73,74,90–92} Although these earlier calculations could reproduce the general shape of the potential energy curve, they also have a number of shortcomings. For example, both CASPT2^{90–92} and NEVPT2⁶⁹ based on the CASSCF(12e,12o) reference function significantly overestimate the dissociation energy of the molecule, especially when large basis sets (e.g. those including g-, h-, or i- type functions) are used.^{72,73} Further, with NEVPT3,⁷³ starting from the CAS(12e,12o) wavefunction results in a large correction to the NEVPT2 result and an unphysical potential energy curve. These various results suggest that (at least when used with low orders of perturbation theory and internally contracted formulations which have a limited ability to relax the reference) the minimal (12e,12o) CAS is too small for a quantitative description.

Further, when using CASPT2, the results are also sensitive to the choice of the zeroth-order Hamiltonian and the level shift employed.^{72,75} This is especially true for the (12e,12o) active space, but sensitivity can still be seen even in DMRG-CASPT2 calculations with a large (12e,28o) CAS derived from the 3d, 4s, 4p, 4d atomic orbitals.⁶³ While the large active space DMRG-CASPT2 curves are generally close to the experimentally derived curve, the different choices of zeroth-order level shifts still yielded a variation in D_e of about 0.2 eV.⁶³ Since NEVPT2 does not require level shifts to avoid intruders, the above provides further incentive to recompute the chromium dimer potential energy curve using DMRG-SC-NEVPT2, and with a large active space.

For our DMRG-SC-NEVPT2 calculations, we used a suite of cc-pwCVXZ (X = T, Q, 5) basis sets. No basis set superposition error (BSSE) corrections were applied due to the large basis sets used (which included h- and i-type functions). The X2C Hamiltonian was used to describe scalar relativistic effects. The DMRG reference and corresponding RDM and DMRG-SC-NEVPT2 calculations

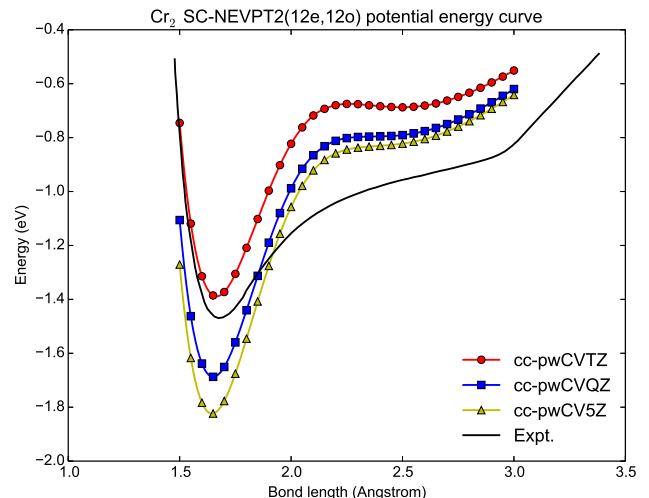


FIG. 2. SC-NEVPT2(12e,12o) potential energy curve with a suite of cc-pwCVXZ (X=T, Q, 5) basis sets.

were performed using the BLOCK program⁵³. All other calculations were performed in PYSCF⁹³. In the figures, the energy of the isolated atoms is set to zero.

We first show the SC-NEVPT2(12e,12o) potential energy curve of Cr₂ in Figure 2 to demonstrate the basis set convergence with the cc-pwCVXZ bases. The (12e,12o) active space was obtained from a CASSCF calculation. We find that using larger basis sets together with this small active space indeed yields far too deep a curve, in agreement with earlier studies using atomic natural orbital (ANO) basis sets.⁷³

We then extended the (12e,12o) active space by adding another set of σ , π , π' , δ , δ' orbitals and their corresponding anti-bonding orbitals to obtain a (12e,22o) active space. We selected the orbitals from the converged (12e,12o) CASSCF calculation based on symmetry and their orbital energies, and the orbitals were then further optimized using DMRG-CASSCF ($M=1000$, no frozen core). Although the additional active orbitals contained both 4p and 4d character at first, the fully optimized additional active orbitals were mainly of 4d character. This is not surprising, as a 4d double shell is well known to greatly improve transition metal CAS results⁹⁴. From the DMRG-CASSCF optimized orbitals, we then carried out a final larger DMRG calculation with $M = 4000$ to obtain an accurate reference wavefunction. The resulting DMRG reference energy in the active space is accurate to better than 0.1 mE_h. To compute the RDM's for the DMRG-SC-NEVPT2 calculation, the large bond dimension reference MPS was compressed down to an MPS with $M' = 800$. Finally, the (TZ/QZ/5Z) energies were extrapolated to the CBS limit using an exponential formula for the DMRG-CASSCF energy, and an l^{-3} formula for the DMRG-SC-NEVPT2 correction, where l is the maximum angular momentum of the basis.

Figure 3 shows the potential energy curves obtained

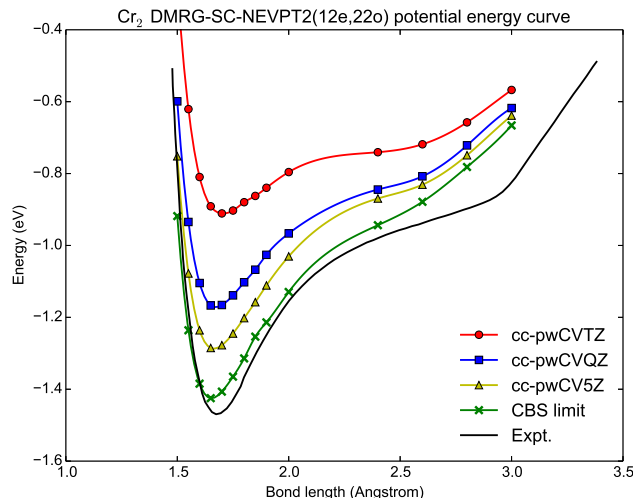


FIG. 3. Extended active space DMRG-SC-NEVPT2(12e,22o)($M'=800$) potential energy curves with a suite of cc-pwCVXZ ($X=T, Q, 5$) basis sets, and the extrapolated CBS limit. The extended active space includes a double d shell. The experimental curve is taken from Ref. 95.

with the cc-pwCVXZ ($X=T, Q, 5$) basis sets and the X2C Hamiltonian, as well as the extrapolated CBS curve. A comparison of the CBS curve to curves obtained with other correlation methods is shown in Fig. 4. We see that larger basis sets using this active space yield systematically improved curves, and that the CBS potential energy curve now agrees very well with the experimental curve, over a wide range of distances. The only significant deviation occurs near the bend at about 2.8 Å, where the reliability of the RKR potential has previously been questioned.⁷¹ The spectroscopic constants are shown in Table I. Our obtained CBS D_e of 1.43 eV is very close to the inferred experimental binding energy of 1.47 eV from the negative ion photoelectron spectroscopy measurements of Casey and Leopold⁹⁵, while the bond-length and vibrational frequency also agree well. It is remarkable to see that even with the largest cc-pwCV5Z basis, the binding energy is still 0.14 eV away from the CBS limit, demonstrating the very large basis set effects in this system.

Compared to earlier “accurate” calculations, the computed DMRG-SC-NEVPT2 curve here compares quite favourably. For example, near the equilibrium distance, the DMRG-SC-NEVPT2(12e,22o) curve is almost identical to the DMRG-CASPT2(12e,28o) curve⁶³. However, in the stretched 4s bonding (“shoulder”) region, the DMRG-SC-NEVPT2(12e,22o) curve is much closer to the experimental one. This seems consistent with the increased correlation strength in the 4s bonding region, which could be harder to describe with the simpler zeroth order Hamiltonian employed in CASPT2. Across the full range of distances, the DMRG-SC-NEVPT2 curve

TABLE I. Spectroscopic constants for the ground state of Cr₂ using different methods.

	$D_e(\text{eV})$	$R_0(\text{\AA})$	$\omega_e(\text{cm}^{-1})$
DMRG-SC-NEVPT2(12e,22o) ^a			
M=800	1.425	1.654	482
M=1200	1.429	1.656	471
DMRG-CASPT2(12e,28o) ^b	1.610	1.681	480
MR-AQCC(12e,12o) ^c	1.355	1.685	459
FP-AFQMC ^d	1.63(5)	1.65(2)	552(93)
experiment	1.47(5) ^e	1.679 ^f	480.6(5) ^e
	1.47 ± 0.1 ^g		

^a From fitting the energies at five points, $R=1.60, 1.65, 1.70, 1.75$, and 1.80 Å

^b DMRG-CASPT2(12e,28o)/ $g_1(M=\infty)$ with the cc-pwCV5Z basis set in Ref. 63

^c TZP/QZP CBS limit in Ref. 74

^d Ref. 96

^e Ref. 95

^f Ref. 97

^g Ref. 98

also appears considerably more accurate than the MR-AQCC(12e,12o) curve, likely due to the larger active space employed.

The (12e,28o) CAS used in the earlier DMRG-CASPT2 calculations contained 4p orbitals. However, the accuracy of the DMRG-SC-NEVPT2(12e,22o) curve as compared to the DMRG-CASPT2(12e,28o), together with the fact that only 4d orbitals remain in the active space after orbital optimization, suggests that the 4d orbitals are more important than the 4p orbitals in the correlations of this system. It has also been suggested in some studies that semi-core correlations (involving the 3s and 3p orbitals) should also have a large effect. In Fig. 5 we show the corresponding curves computed using a (28e,20o) active space containing the semi-core 3s, 3p orbitals and 3d, 4s valence shell (but no 4d shell). We see that the curves are in fact much worse than the curves obtained with the (12e,22o) active space above. This indicates double shell correlations are more important than the semi-core correlations in this system, and that the latter are well accounted for at the second-order perturbation theory level.

B. Poly(p-phenylene vinylene)

Poly(p-phenylene vinylene) is a well-known light emitting conjugated polymer. It has several low-lying states that participate in its photophysics and has been extensively investigated with many theoretical methods, for example various semi-empirical methods⁸⁰, DMRG based on the Pariser-Parr-Pople (PPP) model Hamiltonian^{81,82,85}, time-dependent density functional theory⁸³ and symmetry-adapted cluster configuration interaction (SAC-CI)⁸⁴. Here we examine the excited states and their energies using DMRG-SC-NEVPT2. We limit our calculations to PPV3 (three phenyl rings) to demonstrate

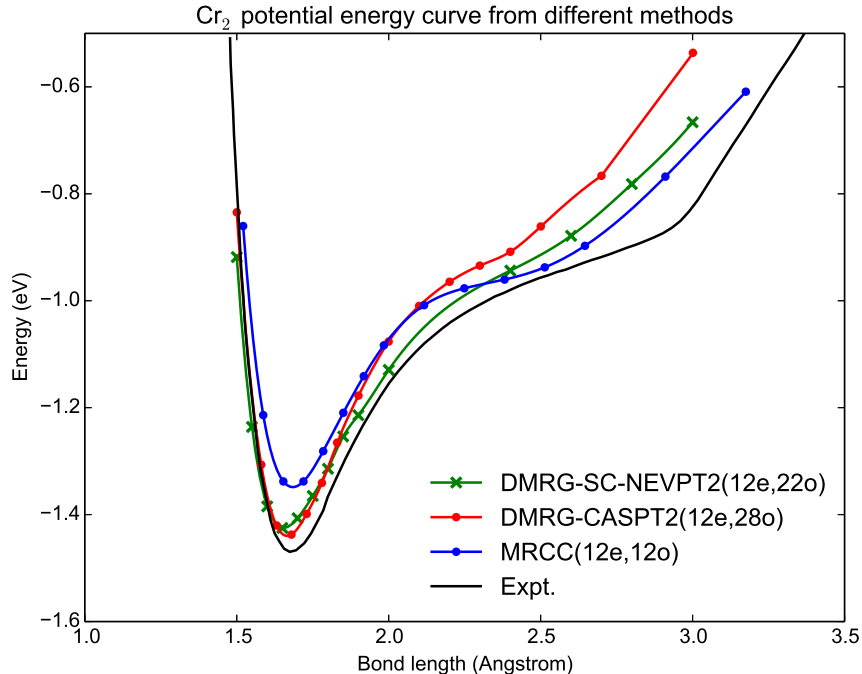


FIG. 4. Cr_2 potential energy curves from different methods. The DMRG-CASPT2(12e,28o) curve is from Ref. 63 and the MR-AQCC(12e,12o) curve is from Ref. 74.

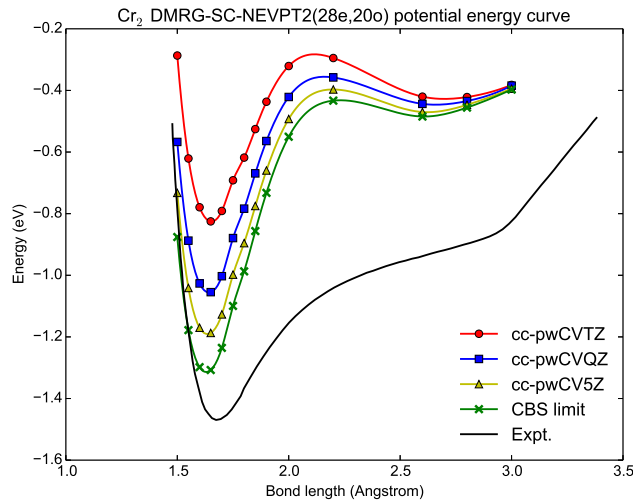


FIG. 5. Extended semi-core DMRG-SC-NEVPT2(28e,20o) ($M'=800$) potential energy curve with a suite of cc-pwCVXZ ($X=T, Q, 5$) basis sets, and the extrapolated CBS limit. The semi-core space includes the 3s and 3p shells. The experimental curve is taken from Ref. 95.

the capabilities of DMRG-SC-NEVPT2 for low-lying excited states in conjugated molecules.

The cc-pVDZ basis set⁹⁹ and a (22e, 22o) active space containing the 22 conjugated π, π^* orbitals were used.

The ground state equilibrium geometry of PPV3 was obtained via DMRG-CI geometry optimization¹⁰⁰, which is implemented within an interface between BLOCK and ORCA¹⁰¹, starting from the DFT/B3LYP optimized geometry. The DMRG-CI geometry optimization was first run at $M = 1000$, and near convergence with $M = 2000$. Canonical Hartree-Fock orbitals were used in the DMRG calculations in the geometry optimization.

Compared to canonical orbitals, localized orbitals usually give better convergence in DMRG calculations.⁵⁸ Therefore, further state-averaged DMRG-CI calculations were carried out for the 6 lowest singlet states, using split-localized orbitals (where the occupied and unoccupied π, π^* orbitals are separately localized with the Pipek-Mezey method¹⁰²). The energies were converged to $0.2mEh$ with a bond dimension of $M = 3200$. The excited states were identified according to their excitation signature from the ground state through the transition 1-RDM and 2-RDM in the canonical Hartree-Fock basis. The symmetry of the states was also determined by the excitation signature: as the ground-state is an A_g state, excitations between two orbitals of different symmetries lead to a B_u excited state, and excitations between two orbitals of the same symmetry lead to an A_g state.

Due to the lack of $\pi - \sigma$ interaction and dynamical correlation, the energy ordering of excitations in DMRG-CI is qualitatively different from experiment, and the state with the HOMO \rightarrow LUMO excitation signature was not the lowest one in energy. To obtain the correct state

TABLE II. Vertical excitation energies of PPV3 in vacuum from different methods.

State	DMRG-SC-NEVPT2	DMRG/PPP	SAC-CI	expt
1^1B_u	3.86	3.52 ^a , 3.91 ^b , 3.751 ^c	3.57 ^d	3.69 ^e
2^1A_g	4.66	4.06 ^a , 4.618 ^c		

^a Ref. 81

^b Ref. 82

^c Ref. 85

^d Ref. 84

^e Ref. 103

ordering, dynamic correlation was further included using DMRG-SC-NEVPT2. The DMRG-CI wavefunctions were compressed to a smaller bond dimension of $M'=500$ and 750. The difference in excitation energy from these two different M' bond dimensions in the DMRG-SC-NEVPT2 calculation was less than $0.6meV$ (Table III), and thus the error from incomplete M' can effectively be ignored. The excitation energy of the first bright state as computed by DMRG-SC-NEVPT2 ($3.86eV$) now agrees well with the experimental number from fluorescence in vacuo by extrapolation the linear fits to $n_{solv} = 1$ ($3.69eV$)¹⁰³ (Table II) and the first bright (B_u state) is found to lie below the $2A_g$ state, as required for efficient light emission. We also find a second B_u state slightly below the $2A_g$ state in energy. The electronic signatures of the states are shown in Table III.

IV. CONCLUSION

In this work we described an efficient and general algorithm to compute high order RDM's from a DMRG wave function. These RDM's may be used as the starting point for many kinds of internally contracted dynamic correlation methods. Using up to the 4th order RDM, we combined DMRG and strongly contracted NEVPT2 to obtain an intruder state free second order multireference perturbation theory that can be used with large active spaces.

To demonstrate the capability of the DMRG-SC-NEVPT2 method, we calculated the potential energy curve of the chromium dimer and the excited states of PPV3. For the chromium dimer, the best extended active space included the so called “double d” shell. Our obtained curve compares quite favourably with earlier calculations and tracks the experimental curve very closely, except at very long bond-lengths where the experimental inversion may be suspect. We find that the semi-core orbitals are much less important for an accurate description of this curve. For PPV3, using an active space containing the full set of π, π^* orbitals, we found that the dynamic correlation included through NEVPT2 allowed us to obtain a correct ordering of the low-lying excited states, and the lowest excitation energy is in good agreement with the experimental data. In summary, these examples paint an optimistic picture for the potential of ap-

plying the DMRG-SC-NEVPT2 method developed here to a wide range of challenging problems.

ACKNOWLEDGEMENTS

APPENDIX A: EDGE CASES FOR NUMBER PATTERNS IN RDM

All number patterns in the N -RDM, $\{n_S, n_D, n_E\}$, need to be computed if

$$0 \leq n_S \leq N \quad (11)$$

$$0 \leq n_E \leq N - 1 \quad (12)$$

$$1 \leq n_D \leq 4 \quad (13)$$

$$n_S + n_D + n_E = 2N \quad (14)$$

However, there are some RDM elements, whose indices cannot be arranged as described above, because too many of the indices lie on the first or last site. We need additional number patterns to compute these elements.

For the first step of the DMRG sweep, additional patterns are added. These patterns satisfy

$$N + 1 \leq n_S \leq \min(4, 2N) \quad (15)$$

$$0 \leq n_D \leq 4 \quad (16)$$

$$n_S + n_D + n_E = 2N \quad (17)$$

For the last step of the DMRG sweep, additional patterns are also added. These patterns satisfy

$$N \leq n_E \leq \min(4, 2N) \quad (18)$$

$$0 \leq n_D \leq 4 \quad (19)$$

$$n_S + n_D + n_E = 2N \quad (20)$$

Because S (or E) is composed of a single site in the first or last step of the sweep, the expectation value of an operator with more than two a 's or two a^\dagger 's is zero. Finally, the number of indices on single sites should not exceed 4.

APPENDIX B: RESTRICTION ON TYPE PATTERNS

It is a waste of resources to compute RDM elements which are zero or redundant. Further RDM elements can be computed through several choices of combinations of operators. We implemented restrictions to skip the calculations of zero and redundant RDM elements.

The operators a_i and a_j^\dagger are first permuted, so that the orbital indices do not decrease. Only permutations between two operators with different orbital indices are needed. If a_i precedes a_j^\dagger and $i = j$, the corresponding compound operators are not required in the RDM calculation. For a single site block, the compound operator with a ahead of a^\dagger is omitted. For other blocks, the loop over $o_i o_j o_k \dots$ should satisfy $i \leq j \leq k \dots$ ('=' is only

TABLE III. Excitation energy (eV) calculated from DMRG-SC-NEVPT2 ($M'=500, 750$) with localized orbitals. Excitations energies of DMRG-CI ($M=3200$) also shown for comparison. The transition signatures are calculated from the DMRG-CI wave function with $M = 3200$. The excitation coefficient of the transition $i \rightarrow j$ ($i, j \rightarrow k, l$) is given by $\frac{1}{\sqrt{2}} \langle \Psi | a_j^\dagger a_i | 1^1 A_g \rangle$ ($\frac{1}{2} \langle \Psi | a_i^\dagger a_k^\dagger a_j a_l | 1^1 A_g \rangle$). The transition labels $n \rightarrow m'$ are as follows: 1, 2, 3 ... denote HOMO, HOMO-1, HOMO-2 ... canonical orbitals, while 1', 2', 3' ... denote LUMO, LUMO+1, LUMO+2 canonical orbitals.

symmetry	excitation signature	$M'=500$	$M'=750$	DMRG-CI($M=3200$)
$1^1 B_u$	0.83 (1 \rightarrow 1')	3.86	3.86	5.13
$2^1 B_u$	0.51 (5 \rightarrow 1') + 0.47 (1 \rightarrow 3')	4.50	4.50	4.50
$2^1 A_g$	0.35 (2 \rightarrow 1') + 0.30 (1 \rightarrow 2') + 0.47 (1, 1 \rightarrow 1', 1')	4.66	4.66	4.85
$3^1 B_u$	many singly excited terms	4.83	4.83	4.75
$4^1 B_u$	many singly excited terms	4.84	4.84	4.76

valid when the two adjacent operators are in normal order).

Further, the RDM (but not the transition RDM) is a Hermitian matrix. Using the 3-RDM as an example

$$\langle a_i^\dagger a_j^\dagger a_k^\dagger a_l a_m a_n \rangle = \langle (a_i^\dagger a_j^\dagger a_k^\dagger a_l a_m a_n)^\dagger \rangle = \langle a_n^\dagger a_m^\dagger a_l^\dagger a_k a_j a_i \rangle \quad (21)$$

We can always use transposes and permutations to ensure that the operator string begins with a^\dagger rather than a . Therefore, all type patterns beginning with a are omitted.

Similarly, if an operator begins with $a_i^\dagger a_i$, $a_i^\dagger a_i a_j^\dagger a_j$ or $a_i^\dagger a_i a_j^\dagger a_j a_k^\dagger a_k$, the remaining set of operators also has transpose and permutation symmetry. e.g.:

$$\begin{aligned} & \langle a_i^\dagger a_i (o_j o_k \dots) \rangle \\ &= \langle (a_i^\dagger a_i (o_j o_k \dots))^\dagger \rangle \\ &= \langle (o_j o_k \dots)^\dagger (a_i^\dagger a_i)^\dagger \rangle \\ &= \langle a_i^\dagger a_i (o_j o_k \dots)^\dagger \rangle \end{aligned} \quad (22)$$

where $i < j \leq k \dots$. Thus, there is no need to compute type patterns where a follows $a_i^\dagger a_i$.

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