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**Targeted excited state algorithms**

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To overcome the limitations of the traditional state-averaging approaches in excited state calculations, where one solves and represents all states between the ground state and excited state of interest, we have investigated a number of new excited state algorithms. Building on the work of van der Vorst and Sleijpen [SIAM J. Matrix Anal. Appl. 17, 401 (1996)], we have implemented harmonic Davidson and state-averaged harmonic Davidson algorithms within the context of the density matrix renormalization group (DMRG). We have assessed their accuracy and stability of convergence in complete-active-space DMRG calculations on the low-lying excited states in the acenes ranging from naphthalene to pentacene. We find that both algorithms offer increased accuracy over the traditional state-averaged Davidson approach, and, in particular, the state-averaged harmonic Davidson algorithm offers an optimal combination of accuracy and stability in convergence. © 2007 American Institute of Physics. [DOI: 10.1063/1.2768360]

**I. INTRODUCTION**

Many excited states possess complicated electronic structure which cannot be described by a single dominant electronic configuration. For such states, a reliable description requires a multireference quantum chemistry method.

Recently, the density matrix renormalization group (DMRG) has emerged as a new tool for multireference quantum chemistry problems. When applied to bond breaking, it achieves a balanced description across potential energy curves due to its reference-free nature. Reduced-scaling DMRG algorithms have also been developed and applied to large multireference problems in quasi-one-dimensional systems such as conjugated polyenes and acenes.

The DMRG ansatz can be written as a linear expansion in terms of many-body functions which are subsequently optimized with respect to internal nonlinear degrees of freedom \( \{ R \} \).

\[
| \Psi \rangle = \sum_{lr} \psi_{lr} | lr(\{ R \}) \rangle .
\]  

(1)

Note that if we choose the expansion functions \( | lr \rangle \) to be Slater determinants and the internal degrees of freedom \( \{ R \} \) to be their constituent orbitals, the above ansatz describes the complete-active-space self-consistent-field (CASSCF) wave function. In the DMRG, the expansion functions are instead complicated many-body basis states and the nonlinear degrees of freedom are renormalization matrices, which allows for a particularly compact and efficient expansion.

To obtain excited states in the DMRG we usually use the iterative Davidson algorithm to solve for eigenvectors \( | \Psi_i \rangle = \psi_{i} | lr \rangle \) ranging from the ground state to the excited state of interest. The nonlinear parameters \( \{ R \} \) for these states are subsequently optimized for a density matrix that is averaged over all the states \( | \Psi_i \rangle \). State averaging is necessary to improve the stability of the nonlinear optimization and to prevent root flipping, which occurs when the approximate wave function leaves the convergence basin of the target excited state and enters that of a different excited state.

The drawbacks of this conventional approach, which we shall refer to as the state-averaged Davidson (SA-D) algorithm, become clear if one is interested in higher regions of the spectrum because it becomes infeasible, both in terms of computational cost and accuracy, to solve for and adequately represent all the lower-lying eigenvectors in the state-averaged DMRG basis. Consequently, it is desirable to explore alternative algorithms that directly yield individual or a few excited state wave functions at a time. Any such algorithm should also retain the stability of the SA-D algorithm during nonlinear optimization, so as to be able to rapidly converge to the desired target excited state(s) without root flipping.

Iterative methods for linear algebra that work with shifted and inverted operators such as \((z^{-1} - H)^{-1}\) have long been used in numerical analysis to obtain the interior (i.e., excited state) eigenvalues of matrices. Sleijpen and van der Vorst proposed an efficient modification that used a shifted and inverted operator to directly calculate harmonic Ritz approximations to excited eigenvalues and eigenvectors. We shall refer to this variant as the harmonic Davidson (HD) algorithm to distinguish it from the original algorithm above. Aside from a demonstration for the one-electron Kohn-Sham equation in Ref. 24, we are not aware of the application of this technique elsewhere in quantum chemistry.

The purpose of this work is to investigate the harmonic Davidson algorithm as a mean to directly target individual excited states and regions of the spectrum within the DMRG. One area in which the current application to quantum chemistry differs from previous numerical applications is the presence of a subsequent nonlinear optimization step for the wave function. We investigate how combining the harmonic Davidson procedure with state averaging over nearby states in the spectrum [state-averaged harmonic Davidson...
(SA-HD)] can be used to confer stability in this nonlinear optimization. While we have focused on the DMRG method here, our findings are relevant to excited state algorithms for other quantum chemistry methods whose ansatz contains both linear and nonlinear parameters, such as in the CAS-SCT method.

The structure of this paper is as follows. In Sec. II, we briefly review the DMRG method and the Davidson and harmonic Davidson algorithms. In Sec. III, we present DMRG calculations on the excited states of acenes from naphthalene to pentacene using both direct targeting with the harmonic Davidson algorithm (in both state-averaged and non-state-averaged forms) as well as with the traditional (state-averaged) Davidson approach. We also compare our excited state spectrum with that obtained from the equation-of-motion coupled cluster theory. We summarize our findings in Sec. IV.

II. THEORY

A. DMRG

The quantum chemistry DMRG algorithm used in this work has been described fully elsewhere.11,25 As a detailed understanding is not necessary here, we shall restrict ourselves to only the essentials. As described above, the DMRG wave function may be written in the form of Eq. (1). The DMRG sweep algorithm then provides an iterative method through which the many-body basis functions |l⟩, |r⟩ may be optimized with respect to a set of internal nonlinear parameters R. For each orbital in the problem we can associate an R matrix, which describes a many-body renormalization transformation involving the orbital (i.e., not simply an orbital rotation). In a sweep to optimize the |l⟩ states (an analogous procedure holds for the |r⟩ states), R matrices are determined from the M eigenvectors of the many-particle reduced density matrix with the largest eigenvalues. In the ground-state case, the density matrix that determines the |l⟩ states is obtained by tracing out the |r⟩ states from the wave function, viz,

$$\Gamma_{lt} = \sum_r \phi_l \phi_t^*,$$

$$\Gamma_{lt} R_{tm} = \gamma R_{tm}, \quad m = 1, \ldots, M,$$  

where typically we choose equal weights for all the states of interest.

B. The Davidson algorithm

The Davidson algorithm provides an efficient iterative solver for the large number of linear coefficients in the expansion of the ground-state DMRG wave function [Eq. (1)].26,27 |Ψ⟩ is expressed in an auxiliary basis {η} (generated by the Davidson iterations),

$$|Ψ⟩ = \sum_i c_i |η⟩,$$

$$|η⟩ = η_i |r⟩.$$

The coefficients c_i are determined by left projection with ⟨η⟩,

$$\sum_i ⟨η | H − E | η⟩ c_i = 0,$$

where E is the approximate expectation value ⟨ψ|H|ψ⟩/⟨ψ|ψ⟩. Each iteration of the Davidson algorithm generates a new basis function |η⟩ from the current trial solution |ψ⟩ via

$$|η⟩ = (\text{diag}(H − E)^{-1}(H − E)|ψ⟩,$$

which is then orthogonalized against and added to the subspace {η}.

To obtain excited state eigenvectors, the simple generalization known as the block Davidson or Davidson-Liu algorithm26,28 is typically used. Here a residual vector is generated for each of the states from the ground state up to the target excited state. Solution of the subspace eigenvalue equation [Eq. (7)] then yields successive approximations to all eigenstates up to the excited state of interest. In the subsequent nonlinear optimization of the excited state in the DMRG algorithm, the eigenvectors obtained from the block Davidson algorithm (i.e., from the ground state to the target eigenvector of interest) are all averaged together in the density matrix [Eq. (4)]. We shall refer to this combined procedure as the SA-D algorithm.

From the above, we see that the primary drawbacks of the traditional SA-D approach are (i) computational cost—we must solve for all the states between the ground state and excited state of interest, and (ii) decreased accuracy—since a single set of nonlinear parameters must now represent multiple states rather than a single state.

C. The harmonic Davidson algorithm

To avoid the need to solve for the states below the excited state of interest as in the Davidson algorithm above, classic shift and invert methods map the target excited state of the Hamiltonian H onto the ground state of a shifted and inverted operator Ω,

$$Ω = H_w^{-1} = (ω − H)^{-1}.$$  

The harmonic Davidson algorithm introduced by Sleijpen and van der Vorst23 (see also Ref. 22 for a clear review) extends the Davidson algorithm to work with the operator Ω without the need to explicitly compute the operator inverse in Eq. (9). Each iteration generates a basis {η}, but now we expand the target excited state |Ψ⟩ in {H_w η}.
\[ |\Psi\rangle = \sum_i c_i |H_{\omega} \eta_i\rangle. \]  

(10)

Left projection with \( \langle \eta | H_{\omega} | \rangle \) yields a generalized eigenvalue problem,

\[ \langle \eta | H_{\omega} | (H_{\omega}^{-1} - E_{\omega}^{-1}) | H_{\omega} \eta \rangle c_i = 0 \]
\[ \Rightarrow \sum_i \left[ \langle \eta_i | H_{\omega} | \eta_i \rangle - E_{\omega}^{-1} \langle \eta | H_{\omega} | H_{\omega} \eta \rangle \right] c_i = 0, \]  

(11)

where \( E_{\omega}^{-1} \) is the current approximation to \((\omega - E)^{-1}\). \( E_{\omega} \) is known as a harmonic Ritz approximation to the corresponding eigenvalue of \( H_{\omega} \). From Eq. (11), we see that solving the eigenvalue equation for \( H_{\omega}^{-1} \) in the subspace \( \{ H_{\omega} \eta \} \) is equivalent to solving the eigenvalue equation for the noninverted operator \( H_{\omega} \), where the trial solution is expanded in the basis \( \{ |\eta\rangle \} \), and the coefficients are obtained by right projection using a different space \( \{ \eta | H_{\omega} \rangle \} \). This suggests that subspace \( \{ \eta \} \) for Eq. (11) can also be generated from the trial solution \( |\psi\rangle \) through a Davidson-type iteration,

\[ |\eta\rangle = (\text{diag}(H_{\omega}) - E_{\omega}'^{-1})(H_{\omega} - E_{\omega}'|\psi\rangle, \]

(12)

where here \( E_{\omega}' \) refers to the expectation value \( \langle \psi | H_{\omega} | \psi \rangle / \langle \psi | \psi \rangle \), which is distinct from \( E_{\omega} \) appearing in Eq. (11).

While we could obtain the excited state eigenvalues and eigenvectors directly from the generalized eigenvalue problem [Eq. (11)], in practice it is numerically more stable to consider a slightly different form. By Schmidt orthogonalization, we can construct an orthogonal decomposition \( \{ \tilde{\eta}_i \} \) of \( \{ H_{\omega} \eta \} \) such that \( \langle \tilde{\eta}_i | H_{\omega} | \tilde{\eta}_j \rangle = \delta_{ij} \). Reexpressing the eigenvalue problem in this basis gives

\[ \sum_i \left[ \langle \tilde{\eta}_i | H_{\omega} | \tilde{\eta}_i \rangle - E_{\omega}^{-1} \delta_{ij} \right] c_i = 0. \]  

(13)

From Eq. (13) we see that implementing the harmonic Davidson algorithm requires only minor alterations to the traditional Davidson routine relating to the change in the subspace from \( \{ \eta \} \) to \( \{ \tilde{\eta} \} \). In essence, there are only two additional steps: The subspace functions are first multiplied by \( H_{\omega} \) and second, they are Schmidt orthogonalized to yield \( \{ \tilde{\eta}_i \} \).

In our later DMRG calculations, we will refer to the use of the above iterative procedure to solve for the linear coefficients together with the nonlinear optimization of the many-body basis functions \( |\tilde{\eta}\rangle, |\tilde{\rho}\rangle \) without state averaging, collectively, as the HD algorithm.

While the operator \( H_{\omega} \) has the target excited state of interest as its ground-state eigenvector, stable convergence is not guaranteed in the nonlinear optimization. However, the formulation of the excited state problem as a ground-state minimization, albeit with a different operator \( \tilde{\Omega} \), illustrates that root flipping is really no different from the poor convergence that may be found in difficult ground-state DMRG calculations. Consequently, the same procedures may be used to eliminate the convergence difficulty: Either we can increase the size \( M \) of the DMRG basis or we can employ a state average over the competing states. While we do not know \textit{a priori} which states will cause convergence difficulties, it is reasonable to assume that they must lie energetically near our state of interest. We have thus implemented two types of state-averaged harmonic Davidson (SA-HD) algorithms. In the first (referred to as simply SA-HD) we average over the first \( n \) excited states of \( \Omega \). These correspond to the \( n \) excited states that lie immediately above our target excited state in the spectrum of \( \tilde{\Omega} \). In the second, we average over the \( n \) states which lie closest (on either side) to the target excited state in the \( \tilde{\Omega} \) spectrum. We refer to this variant algorithm as SA-HDa.

The second variant (SA-HDa) is particularly suited to an alternative way of using the shift \( \omega \). Rather than choosing a shift to target a specific excited state, we can instead choose to find the excited states around a given shift. If stable convergence is not achieved, then we simply increase the number of states used in the SA-HDa average until convergence is recovered. In this way, we can patch together the spectrum piece by piece by using successively higher shifts.

### III. APPLICATION TO ACENES

We have investigated the low-lying states of the acene series ranging from naphthalene (2-acene) to pentacene (5-acene). In the following subsections, we describe the details of the computations (Sec. III A), examine the excitation energies using the state-averaged, harmonic Davidson, and state-averaged harmonic Davidson DMRG algorithms (Sec. III B), and finally use the (near-exact) DMRG results to assess the accuracy of the excitation spectrum obtained from equation-of-motion coupled cluster (EOM-CC) theory (Sec. III C).

#### A. Computational Details

We used a model geometry for the acene with \( C_{2v} \) symmetry. The C–H bond lengths were 1.090 Å. Along the legs of the acene ladder, the alternate C–C bond lengths were 1.410 and 1.405 Å, respectively. Along the rungs of the acene ladder, the C–C bond length was 1.465 Å. An example geometry for naphthalene is shown in Fig. 1.

All calculations used the slater-type orbitals fitted to three Gaussians (STO-3G) minimal basis set, consisting of
**TABLE I.** RHF, CCSD, and DMRG(500) total energies of the acenes. All energies are in hartrees.

| Molecule | $E_{\text{RHF}}$ | CCSD | DMRG(500) |
|----------|----------------|------|-----------|
| C$_{2}$H$_{6}$ | $-378.66597$ | $-378.85130$ | $-378.85360$ |
| C$_{6}$H$_{10}$ | $-529.44420$ | $-529.70634$ | $-529.71032$ |
| C$_{8}$H$_{12}$ | $-680.21823$ | $-680.56059$ | $-680.56538$ |
| C$_{12}$H$_{14}$ | $-830.99045$ | $-831.41614$ | $-831.42016$ |

2$s$ $1p$ functions on C and 1$s$ functions on H. We obtained the atomic orbital integrals and restricted Hartree-Fock (RHF) orbitals from the PSI 3.2 package. The RHF energies are given in Table I. For the excited state calculations, we used a $\pi$ active space consisting of one $p_{z}$ orbital per carbon, i.e., $n$-acene would have a $(4n+2,4n+2)$ active space. In the DMRG calculations, we further symmetrically orthonormalized the $p_{z}$ orbitals with respect to the overlap S. This gave a local orthonormal basis which yields faster convergence in the DMRG calculations. The remaining nonactive orbitals from the RHF calculations were kept frozen in all calculations.

We calculated excitation energies with the SA-D, HD, and SA-HD algorithms described in Sec. II. Our calculations used the local quadractic-scaling DMRG algorithm described in Ref. 11. We employed a screening threshold of $10^{-8}$ hartree ($E_{0}$) with no spatial symmetry. The ordering of the orbitals for anthracene is shown in Fig. 2 and the other acenes were ordered similarly. In all of our sweeps, we added a small amount of random noise ($10^{-6}$–$10^{-7}$) to the density matrix so that we would not lose important quantum numbers. In the current algorithm it is difficult to converge DMRG energies beyond the intrinsic accuracy associated with the finite number $M$ of DMRG basis states. Thus DMRG energies were converged to within 1 millihartree ($mE_{0}$) ($M=50$), 0.5 $mE_{0}$ ($M=100$), 0.5 $mE_{0}$ ($M=250$), or 0.1 $mE_{0}$ ($M=500$), respectively. We note that our largest $M$ DMRG excitation energies are essentially exact (within the one-particle basis) to all reported digits. This is possible for the large active spaces used here because of the compact parametrization afforded by the DMRG wave function.

In the HD and SA-HD calculations, the shift $\omega$ for a specific root was obtained as follows. To begin, we guessed an initial shift (typically based on our previous SA calculations). In the case where the shift was too low or too high, the next guess for $\omega$ was obtained from the DMRG (block) iteration, where an undesired state first appeared as the ground state of the harmonic Davidson procedure. The shift $\omega$ was then taken to lie on the correct side of the desired state in this iteration. In this simple manner, we found that we could obtain a suitable shift for a given root with at most two to three guesses.

To determine the symmetries of the excitations in the DMRG calculations we used the following method. Firstly, spin symmetries were obtained from the expectation value of $S^{2}$. To obtain the spatial symmetries, we first assumed that the ground state $\Psi_{0}$ was of $A_{1}$ symmetry (as in experiment). For the excited states, we examined “dipole”-type matrix elements $\langle \Psi_{0}|n_{i}^{\alpha}+n_{i}^{\beta}-n_{i}^{\beta}-n_{i}^{\alpha}|\Psi_{i}\rangle$ (essentially a dipole transition element along the short axis of the acene; 0 and 1 refer to atom labels in Fig. 2.) For singlet excited states a nonvanishing dipole then implied $B_{1}$ symmetry, while a vanishing dipole implied $A_{1}$ symmetry. For the triplet excited states, all such matrix elements vanish. However, we could still determine the spatial symmetry through the expectation value $\langle \Psi_{0}|n_{i}^{\alpha}n_{i}^{\beta}|\Psi_{i}\rangle$ since $n_{i}^{\alpha}n_{i}^{\beta}$ does not preserve spin symmetry and creates a residual expectation value from which one can determine the spatial symmetry of the excited state.

To obtain the orbital character of the excitations, we calculated transition one-particle density matrices $\langle \Psi_{0}|a_{i}^{\dagger}a_{j}^{\dagger}|\Psi_{i}\rangle$, where $\Psi_{i}$ denotes the $i$th excited state and identified the largest matrix elements.

We further calculated the excitation spectrum (in the same $\pi$ active space as the DMRG calculations) with the equation-of-motion coupled cluster singles and doubles (EOM-CCSD) method using the DALTON package.

**B. Comparison of excited-state algorithms for DMRG by SA, HD, and SA-HD**

The ground-state DMRG energies for the acenes are given in Table I. Tables II–V contain the first seven $\pi-\pi^{*}$ excitation energies for each acene, while Fig. 3 displays them in graphical form. Under $C_{2v}$ symmetry, the only two possible representations of the $\pi-\pi^{*}$ excited states are 1,3$A_{1}$ and 1,1$B_{2}$. Experimentally, there are three well-documented singlet bands that appear in the visible spectrum. The a band and $\beta$ band correspond to a polarization along the long axis and the $p$ band corresponds to a transverse polarization. We observed the $\alpha$ transition as the lowest singlet excitation in each acene. Neither the $p$ band nor the $\beta$ band appeared within the first eight states of each acene. Instead, for the case of naphthalene, the $p$ band emerged at 8.42 eV (state 19). The $p$ band normally appears lower in the spectrum, but the absence of dynamic $\sigma-\pi$ correlations is responsible for its artificially high excitation energy here. This is consistent with previous studies of acenes using CASSCF and complete-active-space Moller-Plesset second order perturbation (CASMP2) theory. Triple excitations are somewhat harder to measure experimentally. We observe that the triplet excitation energies decrease in energy more rapidly with system size than the singlet excitations. Thus while in naphthalene and anthracene there is one triplet level between the first two singlet excitations, in naphthalene and pentacene there are two.

Comparing the accuracies of the SA-D, HD, and SA-HD calculations we observe that as expected (other than by the
TABLE II. DMRG excitation energies for naphthalene \((C_{10}H_8)\) obtained with the SA-D, HD, and SA-HD algorithms. All energies are in eV. State 0 refers to the ground state, and \(SA\{m\rightarrow n\}\) refers to a state average over all states from the \(m\)th to \(n\)th excited state. Numbers in parentheses give the number of DMRG states \(M\). The “Exact (HD(500))” numbers are the (near-exact) excitation energies, while other entries give the errors from this result. The “Excitation” row gives the character of the excitation, where 1 denotes HOMO, 2 denotes HOMO-1, 1’ denotes LUMO, 2’ denotes LUMO+1, and so on. The last column gives the mean improvement in the excitation energy over the \(SA\{0–7\}\) D result with the same \(M\). nc denotes no convergence.

| State | Mean improvement |
|-------|------------------|
| \(1\,^1A_1\) | \(1\,^1B_2\) | \(2\,^1A_1\) | \(2\,^1B_2\) | \(3\,^3B_1\) | \(2\,^3B_2\) | \(2\,^3A_1\) | \(3\,^3A_1\) |
| Excitation | 0.00 | 2.86 | 4.08 | 4.34 | 4.63 | 4.70 | 5.51 | 5.87 |
| \(SA\{0–7\}\) D (50) | 0.13 | 0.09 | 0.21 | 0.46 | 0.18 | 0.15 | 0.26 | 0.19 |
| \(SA\{3–3\}\) D (50) | 0.11 | 0.17 | 0.18 | 0.34 | 0.46 | 0.17 | 0.18 | 0.24 | 0.25 | 0.02 |
| \(SA\{3–7\}\) HD (50) | 0.04 | 0.05 | 0.08 | nc | nc | nc | nc | 0.08 | 0.09 |
| \(SD\{3–3\}\) HD (50) | 0.01 | 0.02 | 0.02 | 0.03 | 0.02 | 0.02 | 0.02 | 0.02 | 0.02 | 0.01 |
| \(SA\{0–7\}\) D (100) | 0.00 | 0.01 | 0.01 | 0.01 | 0.01 | 0.01 | 0.01 | 0.01 | 0.01 | 0.01 |
| \(SA\{0–3\}\) D (250) | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
| \(SA\{0–3\}\) (250) | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
| \(SA\{3–7\}\) HD (250) | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
| \(HD\) (250) | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
| \(SA\{3–7\}\) HD (500) | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |

TABLE III. DMRG excitation energies for anthracene \((C_{14}H_{9})\). Refer to Table II for details.

| State | Mean improvement |
|-------|------------------|
| \(1\,^1A_1\) | \(1\,^1B_2\) | \(2\,^1A_1\) | \(2\,^1B_2\) | \(3\,^3B_1\) | \(2\,^3B_2\) | \(2\,^3A_1\) | \(3\,^3A_1\) |
| Excitation | 0.00 | 2.08 | 3.57 | 3.71 | 3.85 | 4.46 | 4.73 | 4.80 |
| \(SA\{0–7\}\) D (50) | 0.40 | 0.45 | 0.75 | 0.65 | 1.28 | 0.77 | 0.82 | 0.91 |
| \(SA\{0–3\}\) D (50) | 0.29 | 0.24 | 0.46 | 0.41 | 0.73 | 0.58 | 0.60 | 0.47 | 0.69 | 0.27 |
| \(SA\{0–3\}\) HD (50) | 0.12 | 0.13 | 0.40 | nc | nc | nc | nc | nc | 0.32 |
| \(HD\) (50) | 0.01 | 0.05 | 0.05 | 0.04 | 0.04 | 0.73 | 0.58 | 0.60 | 0.47 | 0.69 | 0.27 |
| \(SA\{0–3\}\) HD (100) | 0.12 | 0.15 | 0.15 | 0.25 | 0.25 | 0.23 | 0.18 | 0.19 | 0.05 |
| \(HD\) (100) | 0.01 | 0.03 | 0.05 | 0.04 | 0.04 | 0.73 | 0.58 | 0.60 | 0.47 | 0.69 | 0.27 |
| \(SA\{0–3\}\) HD (500) | 0.12 | 0.15 | 0.15 | 0.25 | 0.25 | 0.23 | 0.18 | 0.19 | 0.05 |
| \(HD\) (500) | 0.01 | 0.03 | 0.05 | 0.04 | 0.04 | 0.73 | 0.58 | 0.60 | 0.47 | 0.69 | 0.27 |

size of the DMRG basis \(M\) the accuracy in the excitation energies is determined primarily by the number of eigenvectors in the state average. Consequently the traditional SA-D algorithm yielded the lowest accuracy (as it averages over all states between the ground state and excited state of interest), while the HD calculations were correspondingly the most accurate since they targeted a single state at a time. The accuracy of the SA-HD calculations lay somewhere in be-
In all cases, the differences between the various algorithms was most marked for the smaller sizes $M$ of the DMRG basis, as for larger $M$ all the wave functions become essentially exact. We would expect the differences to become more pronounced in larger systems, where we are unable to use a sufficiently large $M$ to reach exactness.

Regarding the stabilities of the various algorithms, we

### Table IV. DMRG excitation energies for naphthacene (C_{14}H_{12}). Refer to Table II for details.

| Method       | $1^1A_1$ | $1^3B_2$ | $2^1A_1$ | $2^3B_2$ | $3^1A_1$ | $3^3B_2$ | $2^3A_1$ | Mean improvement |
|--------------|----------|----------|----------|----------|----------|----------|----------|-----------------|
| **Excitation** |          |          |          |          |          |          |          |                 |
| Exact (HD(500)) | 0.00     | 0.09     |          |          |          |          |          |                 |
| SA [0–7] D (50) | 1.01     | 0.95     | 1.00     | 0.97     | 0.96     | 0.95     | 0.92     | 0.90            |
| SA [0–3] D (50) | 0.96     | 0.92     | 0.90     | 0.91     | 0.88     | 0.90     | 0.89     | 0.88            |
| SA [2–7] HD (50) |          |          |          |          |          |          |          |                 |
| SA [3–7] HD (50) |          |          |          |          |          |          |          |                 |
| **HD (50)** |          |          |          |          |          |          |          |                 |
| SA [1–2] HD (50) |          |          |          |          |          |          |          |                 |
| SA [2–3] HD (50) |          |          |          |          |          |          |          |                 |
| SA [0–7] D (100) |          |          |          |          |          |          |          |                 |
| SA [0–3] D (100) |          |          |          |          |          |          |          |                 |
| SA [2–7] HD (100) |          |          |          |          |          |          |          |                 |
| SA [3–7] HD (100) |          |          |          |          |          |          |          |                 |
| **HD (100)** |          |          |          |          |          |          |          |                 |
| SA [0–7] D (250) |          |          |          |          |          |          |          |                 |
| SA [0–3] D (250) |          |          |          |          |          |          |          |                 |
| SA [2–7] HD (250) |          |          |          |          |          |          |          |                 |
| SA [3–7] HD (250) |          |          |          |          |          |          |          |                 |
| **HD (250)** |          |          |          |          |          |          |          |                 |
| HD (50) |          |          |          |          |          |          |          |                 |
| HD (100) |          |          |          |          |          |          |          |                 |
| HD (250) |          |          |          |          |          |          |          |                 |
| HD (250) |          |          |          |          |          |          |          |                 |

### Table V. DMRG excitation energies for pentacene (C_{12}H_{14}). Refer to Table II for details.

| Method       | $1^1A_1$ | $1^3B_2$ | $2^1A_1$ | $2^3B_2$ | $3^1A_1$ | $3^3B_2$ | $2^3A_1$ | Mean improvement |
|--------------|----------|----------|----------|----------|----------|----------|----------|-----------------|
| **Excitation** |          |          |          |          |          |          |          |                 |
| Exact (HD(500)) | 0.00     | 0.09     |          |          |          |          |          |                 |
| SA [0–7] D (50) | 1.01     | 0.95     | 1.00     | 0.97     | 0.96     | 0.95     | 0.92     | 0.90            |
| SA [0–3] D (50) | 0.96     | 0.92     | 0.90     | 0.91     | 0.88     | 0.90     | 0.89     | 0.88            |
| SA [2–7] HD (50) |          |          |          |          |          |          |          |                 |
| SA [3–7] HD (50) |          |          |          |          |          |          |          |                 |
| **HD (50)** |          |          |          |          |          |          |          |                 |
| SA [1–2] HD (50) |          |          |          |          |          |          |          |                 |
| SA [2–3] HD (50) |          |          |          |          |          |          |          |                 |
| SA [0–7] D (100) |          |          |          |          |          |          |          |                 |
| SA [0–3] D (100) |          |          |          |          |          |          |          |                 |
| SA [2–7] HD (100) |          |          |          |          |          |          |          |                 |
| SA [3–7] HD (100) |          |          |          |          |          |          |          |                 |
| **HD (100)** |          |          |          |          |          |          |          |                 |
| HD (50) |          |          |          |          |          |          |          |                 |
| HD (100) |          |          |          |          |          |          |          |                 |
| HD (250) |          |          |          |          |          |          |          |                 |
| HD (250) |          |          |          |          |          |          |          |                 |

between depending on the number of states used in the average.
found that there were no difficulties in converging the DMRG sweeps to the correct states with the SA-D algorithm. The HD algorithm on the other hand exhibited the expected convergence difficulties characteristic of root flipping for certain higher excited states. As previously discussed, the stability of the HD algorithm would increase with the size of the DMRG many-body basis $M$. In naphthalene, we required $M=250$ to converge states 5–7 with the HD algorithm, while in pentacene, we required $M=500$ to converge states 4–7. While the HD algorithm exhibited root flipping, it was ameliorated with respect to simple eigenvector following because of the use of the shift $\omega$. For example, with $M=100$, the third excited state of naphthalene could not be converged with simple eigenvector following, but could be converged without difficulty using the HD algorithm. Including a sufficient number of states in the SA-HD algorithm restored the stability of the convergence. Certain “competing” states were particularly important for the state average, especially for smaller $M$. For all the acenes, the second and third excited states were examples of such states. Thus while the state averages SA$^{2}$–3 HD and SA$^{2}$–7 HD converged without difficulty, calculations using SA$^{3}$–7 HD did not, at least for smaller $M$.

As mentioned previously, rather than choosing a shift to target specific excited states, we could take the different approach of trying to find the excited states around the frequency of a given shift $\omega$. In this way, we could piece together a complete spectrum by performing, say, SA–HD or SA–HDa calculations with successively higher shifts. To demonstrate this, we computed the excitation energies for states 6–11 for naphthalene using the SA-HDa algorithm with a shift chosen slightly above the state 7 excitation energy as estimated from the previous SA-HD calculation. These are shown in Table VI.

### Table VI. Excitation Energies for Naphthalene, Anthracene, and Pentacene

| Acene        | State 4 | State 5 | State 6 | State 7 | State 8 | State 9 | State 10 | State 11 |
|--------------|---------|---------|---------|---------|---------|---------|----------|----------|
| Naphthalene  | (4.23)  | (5.00)  | (5.94)  | (6.85)  | (7.75)  | (8.66)  | (9.57)   | (10.48)   |
| Anthracene   | (4.20)  | (5.00)  | (5.94)  | (6.85)  | (7.75)  | (8.66)  | (9.57)   | (10.48)   |
| Pentacene    | (4.20)  | (5.00)  | (5.94)  | (6.85)  | (7.75)  | (8.66)  | (9.57)   | (10.48)   |

As mentioned previously, rather than choosing a shift to target specific excited states, we could take the different approach of trying to find the excited states around the frequency of a given shift $\omega$. In this way, we could piece together a complete spectrum by performing, say, SA–HD or SA–HDa calculations with successively higher shifts. To demonstrate this, we computed the excitation energies for states 6–11 for naphthalene using the SA-HDa algorithm with a shift chosen slightly above the state 7 excitation energy as estimated from the previous SA-HD calculation. These are shown in Table VI.

### C. Comparison of DMRG and EOM-CC excitation energies in the acenes

The ground-state EOM-CCSD energies for the acenes are summarized in Table I. We used our near-exact DMRG(500) excitation energies to examine the accuracy of the EOM-CC method in acenes. The EOM-CCSD and the DMRG symmetries and excitation energies are shown in Fig. 3. For the larger acenes, the EOM-CCSD excited states are in a qualitatively different order as compared to DMRG. Similarly, EOM-CCSD erroneously predicts a very small singlet-
triplet gap for the longer acenes. This points to the necessity of including relatively high order correlation effects to accurately describe excitations in the acenes.

IV. CONCLUSIONS

To overcome the computational and accuracy limitations of the traditional state-averaged Davidson algorithm, which requires both solving and representing all states between the ground state and excited state of interest, we have investigated a number of new excited state algorithms within the context of the density matrix renormalization group (DMRG). In the harmonic Davidson (HD) algorithm, using a shifted and inverted operator enabled us to directly solve for the excited state of interest. In the state-averaged harmonic Davidson (SA-HD) algorithm, we combined the HD method with an average over nearby excited states to confer greater stability and overcome problems of root flipping in the non-linear optimization of the wave function.

To assess the accuracy, stability, and computational cost of these new methods we calculated the low-lying excited states in the acenes ranging from naphthalene to pentacene. We found that as expected, in addition to the size of the states in the acenes ranging from naphthalene to pentacene.

TABLE VI. DMRG excitation energies for the higher excited states of naphthalene (C_{10}H_{8}). Refer to Table II for details.

| Method          | 2 1^A_1 | 3 1^A_1 | 4 1^B_2 | 3 1^A_1 | 1 1^B_2 | 4 1^A_1 | Mean improvement |
|-----------------|---------|---------|---------|---------|---------|---------|------------------|
| Excitation      | 4 → 1'  | 3 → 2'  | 4 → 2'  | 2 → 3'  | 1 → 3'  | 4 → 1'  |                 |
|                 | 1 → 4'  | 2 → 3'  | 2 → 4'  | 3 → 2'  | 3 → 1'  | 1 → 4'  |                 |
| Exact (HD(500)) | 5.51    | 5.87    | 6.28    | 6.48    | 6.84    | 6.84    |                 |
| SA [0–11] D (50)| 0.29    | 0.21    | 0.19    | 0.45    | 0.32    | 0.43    |                 |
| SA [6–11] HDa (50)| 0.29 | 0.20    | 0.17    | 0.46    | 0.35    | 0.43    | 0.00             |
| HD (50)         | nc      | 0.08    | nc      | nc      | nc      | nc      | 0.13             |
| SA DA [0–11] D (100) | 0.03 | 0.03    | 0.03    | 0.03    | 0.05    | 0.06    |                 |
| SA [6–11] HDa (100)| 0.03 | 0.02    | 0.03    | 0.03    | 0.05    | 0.06    | 0.00             |
| HD (100)        | 0.01    | 0.01    | 0.01    | 0.01    | 0.02    | nc      | 0.02             |
| SA [10–11] HDa (100)| 0.01 | 0.01    | 0.01    | 0.01    | 0.02    | nc      | 0.02             |
| SA [0–11] D (250)| 0.00    | 0.00    | 0.00    | 0.00    | 0.00    | 0.00    |                 |
| SA [6–11] HDa (250)| 0.00 | 0.00    | 0.00    | 0.00    | 0.00    | 0.00    | 0.00             |
| HD (250)        | 0.00    | 0.00    | 0.00    | 0.00    | 0.00    | nc      | 0.00             |
| SA DA [10–11] HDa (250)| 0.00 | 0.00    | 0.00    | 0.00    | 0.00    | 0.00    | 0.00             |
| SA [0–11] D (500)| 0.00    | 0.00    | 0.00    | 0.00    | 0.00    | 0.00    | 0.00             |
| SA [6–11] HDa (500)| 0.00 | 0.00    | 0.00    | 0.00    | 0.00    | 0.00    | 0.00             |

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