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Zhuoran He and Andrew J. Millis Phys. Rev. B **96**, 085107 — Published 7 August 2017 DOI: 10.1103/PhysRevB.96.085107

Entanglement entropy and computational complexity of the Anderson impurity model out of equilibrium I: quench dynamics

Zhuoran He¹ and Andrew J. Millis¹

¹Department of Physics, Columbia University, New York, New York 10027, USA

We study the growth of entanglement entropy in density matrix renormalization group calculations of the real-time quench dynamics of the Anderson impurity model. We find that with appropriate choice of basis, the entropy growth is logarithmic in both the interacting and noninteracting singleimpurity models. The logarithmic entropy growth is understood from a noninteracting chain model as a critical behavior separating regimes of linear growth and saturation of entropy, which correspond respectively to an overlapping and gapped energy spectra of the set of bath states. We find that a logarithmic entropy growth is the generic behavior of quenched impurity models when the bath orbitals in the matrix product state are ordered in energy. A noninteracting calculation of the doubleimpurity Anderson model supports the conclusion in the multi-impurity case. The logarithmic growth of entanglement entropy enables studies of quench dynamics to very long times.

PACS numbers: 71.10-w, 71.15-m

I. INTRODUCTION

It has been a long-standing challenge to develop efficient real-time impurity solvers to study the dynamics of a system out of equilibrium. The Anderson impurity model [1] (AIM), a single spin-degenerate orbital with an intra-orbital Hubbard interaction U coupled to a bath of noninteracting orbitals, is of fundamental importance in its own right as a nontrivial but solvable [2, 3] interacting electron model and as an auxiliary problem for dynamical mean-field theory [4, 5]. The nonequilibrium properties of this model [6, 7] provide an important laboratory for the development of real-time methods [8–11].

The density matrix renormalization group (DMRG) [12] is a powerful numerical technique for solving lowdimensional electron problems including the Anderson impurity model. In DMRG, the wave function of the system is represented by a matrix product state (MPS). Every matrix in the MPS corresponds to a local degree of freedom. If these degrees of freedom are labeled by an index n, then the logarithm of the bond dimension D_n between the *n*-th and (n + 1)-th matrices (which equals the column dimension of the *n*-th and the row dimension of the (n + 1)-th matrix) of the MPS is greater than or of the order of the entanglement entropy S_n between orbitals $1, 2 \dots n$ and $n+1 \dots N$. The rate of growth with time of the entanglement entropy across the maximum entropy cut controls the maximum time achievable in a real-time DMRG simulation.

In this paper we study the growth of entanglement entropy in DMRG calculations, focusing primarily on the single-impurity Anderson model. Our work is motivated by the recent results of Wolf *et al.* [13], who found that a particular "star geometry" arrangement of bath orbitals provided a favorable growth of entanglement entropy. We consider a variety of bath Hamiltonians and find that the growth of entanglement entropy S_n across the cut at n is controlled by whether the spectra of MPS orbitals $1, 2 \dots n$ overlaps that of the $n+1 \dots N$ orbitals. The star geometry with energy-ordered bath orbitals in the MPS ensures no energy overlap at every bond, which leads to a logarithmic growth of maximum entanglement entropy with time. This method of controlling entropy growth using energy separation is hypothesized to work for multiimpurity models as well; the hypothesis is supported by a calculation of a non-interacting double-impurity Anderson model.

The rest of the paper is organized as follows. Section II describes the single-impurity Anderson model (SIAM) and Sec. III describes our implementation of DMRG, which differs slightly from the standard implementation. In Secs. IV–VI, we show results obtained using our method for the SIAM, discuss the logarithmic growth of entanglement entropy and also present a few results for the quenched double-impurity Anderson model. Section VII is a conclusion and summary.

II. THEORY

We focus on the single-impurity Anderson model [1] (SIAM) defined by the Hamiltonian

$$H = H_d + H_{\text{bath}} + H_{\text{mix}},\tag{1}$$

$$H_d = \sum_{\sigma} \epsilon_d \, d^{\dagger}_{\sigma} d_{\sigma} + U \, d^{\dagger}_{\uparrow} d_{\uparrow} d^{\dagger}_{\downarrow} d_{\downarrow}, \qquad (2)$$

$$H_{\text{bath}} = \sum_{k\sigma} \epsilon_k \, c_{k\sigma}^{\dagger} c_{k\sigma}, \qquad (3)$$

$$H_{\rm mix} = \sum_{k\sigma} V_k \, d^{\dagger}_{\sigma} c_{k\sigma} + \text{h.c.}. \tag{4}$$

The model has one d orbital with a Hubbard U interaction in its Hamiltonian H_d and $\mathcal{N} \to \infty$ noninteracting bath orbitals with Hamiltonian H_{bath} . The two systems hybridize via the one-body hopping terms in H_{mix} . The bath orbitals are labeled by k and the two spins \uparrow and \downarrow of electrons are labeled by σ . For simplicity, we take

the impurity-bath coupling amplitudes $V_k = V/\sqrt{N}$ to be k-independent. We define the bath density of states as $\text{DOS}(\epsilon) = \frac{1}{N} \sum_k \delta(\epsilon - \epsilon_k)$ and consider a semicircle DOS with a half band width E. The initial state that we consider is a product state

$$|\Psi_{t=0}\rangle = |\Psi_0\rangle_d \otimes |\text{FS}\rangle_{\text{bath}},\tag{5}$$

where the Fermi-sea state $|\text{FS}\rangle_{\text{bath}}$ of the bath is initially half-filled. In the studies we present here, the *d*-orbital energies ϵ_d and $\epsilon_d + U$ are chosen to be symmetric about the Fermi level at 0. The situation is depicted in Fig. 1. The formalism generalizes to a mixed initial state with a product density matrix $\rho_{t=0} = (\rho_0)_d \otimes (\rho_0)_{\text{bath}}$, where $(\rho_0)_{\text{bath}}$ satisfies the Wick's theorem, but here we will focus on a pure initial state.

The numerical methods we use require a truncation of the bath to a finite number N of bath orbitals. To choose the best truncation, we first calculate the hybridization function $\Delta_{\sigma}(t,t') = -i \sum_{k} |V_k|^2 \langle \mathcal{T}_{\mathcal{C}} c_{k\sigma}(t) c_{k\sigma}^{\dagger}(t') \rangle_{\text{bath}}$ on the Keldysh-contour [6]. The contour \mathcal{C} goes from t = 0to a sufficiently late time $t_{\text{max}} > \max(t,t')$ and then back to t = 0. $\mathcal{T}_{\mathcal{C}}$ is the contour-ordering symbol. The operators $c_{k\sigma}(t)$ and $c_{k\sigma}^{\dagger}(t')$ are in the Heisenberg picture of H_{bath} and the mean value $\langle \ldots \rangle_{\text{bath}}$ is taken with respect to $|\text{FS}\rangle_{\text{bath}}$. With our choice of the semicircular DOS and constant V_k , the hybridization function can be analytically found for $\mathcal{N} \to \infty$ to be

$$\Delta_{\sigma}(t,t') = \begin{cases} -\frac{V^2}{E\tau} [H_1(E\tau) + iJ_1(E\tau)], \ t \succ_{\mathcal{C}} t', \\ -\frac{V^2}{E\tau} [H_1(E\tau) - iJ_1(E\tau)], \ t \prec_{\mathcal{C}} t', \end{cases}$$
(6)

where $\tau = t - t'$, H_1 is the 1st-order Struve function and J_1 is the 1st-order Bessel function. The symbols $\succ_{\mathcal{C}}$ and $\prec_{\mathcal{C}}$ refer to Keldysh-contour ordering. Then we fit the hybridization function to that of a finite bath with only N orbitals, i.e.,

$$\Delta_{\sigma}(t,t') \approx -i \sum_{j=1}^{N} V_j^2 \langle \mathcal{T}_{\mathcal{C}} c_{j\sigma}(t) c_{j\sigma}^{\dagger}(t') \rangle_{\text{bath}}$$
$$= -i \sum_{j=1}^{N} V_j^2 \left[\Theta_{\mathcal{C}}(t,t') - n_{j\sigma}^0 \right] e^{-i\epsilon_j(t-t')}, \quad (7)$$

where $\Theta_{\mathcal{C}}(t,t') = 1$ if $t \succ_{\mathcal{C}} t'$ and 0 if $t \prec_{\mathcal{C}} t'$. In fitting Eq. (6) with Eq. (7), all 2N real parameters ϵ_j and V_j are varied to minimize the least-square error up to a maximum time. The occupancies $n_{j\sigma}^0$ are chosen to be either 0 or 1 to fit the $t \succ_{\mathcal{C}} t'$ and $t \prec_{\mathcal{C}} t'$ parts independently and to make the initial state of the finite bath a Slater determinant. This is possible even if the original bath was at a nonzero temperature. Since our bath is particle-hole symmetric, we choose N to be even to preserve this symmetry.

The number N of bath orbitals controls the maximum time $t_N \leq 2\pi N/|E_{\text{max}} - E_{\text{min}}| = \pi N/E$ up to which



FIG. 1. The density of states of the bath orbitals. We consider a semicircle DOS with a half band width E. The bath is initially half-filled, and the *d*-orbital energy ϵ_d and $\epsilon_d + U$ are symmetric about the Fermi level at 0.

the exact hybridization function is reproduced with good accuracy. For example, N = 40 bath orbitals are enough to reach $Et \lesssim 100$ and N = 170 orbitals can reach $Et \lesssim 500$. Adding more orbitals increases the maximal time that can be reached, but does not significantly improve the accuracy of the fit at shorter times.

III. METHOD

We use DMRG/MPS methods to carry out the time evolution. We represent the SIAM wave function $|\Psi(t)\rangle$ as an entangled state between the impurity d orbital and the bath, i.e.,

$$|\Psi(t)\rangle = \sum_{i} c_{i}(t)|i\rangle_{d} \otimes |\Psi_{i}(t)\rangle_{\text{bath}}, \qquad (8)$$

where *i* sums over the 4 impurity states $|0\rangle$, $|\uparrow\rangle$, $|\downarrow\rangle$ and $|\uparrow\downarrow\rangle$. Every bath state $|\Psi_i(t)\rangle_{\text{bath}}$ is a normalized matrix product state (MPS). The coefficients $c_i(t)$ are real and nonnegative. Eq. (8) is a Schmidt decomposition of $|\Psi(t)\rangle$ between the *d* orbital and the bath if $|\Psi(t)\rangle$ is a simultaneous eigenstate of N_{\uparrow} and N_{\downarrow} , the total numbers of spin-up and spin-down electrons. This representation differs from the conventional DMRG in that it removes the *d* orbital from the MPS, enabling analysis of the entanglement among the bath orbitals.

We evolve the wave function $|\Psi(t)\rangle$ in the interaction picture of $H_0 = H_d + H_{\text{bath}}$ according to

$$|\Psi(t)\rangle = \mathcal{T}e^{-i\int_0^t dt' \hat{H}_{\min}(t')} |\Psi_{t=0}\rangle, \tag{9}$$

where \mathcal{T} is the time-ordering symbol and

$$\hat{H}_{\min}(t) = e^{iH_0 t} H_{\min} e^{-iH_0 t}$$
$$= \sum_{j\sigma} V_j e^{i(Un_{d\sigma} + \epsilon_d - \epsilon_j)t} d^{\dagger}_{\sigma} c_{j\sigma} + \text{h.c.}, \qquad (10)$$

where $\bar{\sigma}$ is the opposite spin of σ . The main advantage of the interaction picture is that $\hat{H}_{\text{mix}}(t)$ typically has a

narrower spectral radius than H_0 (bath bandwidth ~ E large compared with impurity level width ~ V^2/E).

We evaluate Eq. (9) by discretizing the time evolution into time steps Δt . The Hamiltonian used during the time step centered at t is

$$\tilde{H}_{\min}(t) = \frac{1}{\Delta t} \int_{t-\Delta t/2}^{t+\Delta t/2} \hat{H}_{\min}(t') dt'$$
$$= \sum_{j\sigma} \tilde{V}_{j\sigma}(t) d_{\sigma}^{\dagger} c_{j\sigma} + \text{h.c.}, \qquad (11)$$

with the coupling amplitudes

$$\tilde{V}_{j\sigma}(t) = V_j \, e^{i(Un_{d\bar{\sigma}} + \epsilon_d - \epsilon_j)t} \operatorname{sinc}(\frac{Un_{d\bar{\sigma}} + \epsilon_d - \epsilon_j}{2} \Delta t).$$
(12)

The errors of both the mid-point Hamiltonian $\hat{H}_{mix}(t)$ and the time-averaged Hamiltonian $\tilde{H}_{mix}(t)$ are $\mathcal{O}(\Delta t^2)$. The latter choice is preferred if the bath bandwidth is large compared with the *d*-level width, because the very high and very low-energy bath orbitals are suppressed by the sinc function.

To apply the Hamiltonian $\tilde{H}_{\text{mix}}(t)$ to the wave function $|\Psi(t)\rangle$ in Eq. (8), we work in the Jordan-Wigner transformed representation with the *d* orbital being the first orbital (*d* and d^{\dagger} having no Jordan-Wigner signs). The Hamiltonian in Eq. (11) is rewritten as

$$\tilde{H}_{\rm mix}(t) = \sum_{\sigma} (-1)^{n_{d\bar{\sigma}}} d^{\dagger}_{\sigma} \tilde{c}_{\sigma}(t) + \text{h.c.}, \qquad (13)$$

$$\tilde{c}_{\sigma}(t) = \sum_{j} \tilde{V}_{j\sigma}(t) \, (-1)^{n_1 + \dots + n_{j-1}} \tilde{c}_{j\sigma}, \qquad (14)$$

where the $\tilde{c}_{j\sigma}$ is the Jordan-Wigner transformed form of $c_{j\sigma}$. The two operators are related by

$$c_{j\sigma} = (-1)^{n_d + n_1 + \dots + n_{j-1}} \tilde{c}_{j\sigma}, \qquad (15)$$

so that the operators $\tilde{c}_{j\sigma}$ and $\tilde{c}_{j'\sigma'}$ for $j \neq j'$ commute. We do the same Jordan-Wigner transform for the two spins of the same orbital, so that $\tilde{c}_{j\uparrow}$ and $\tilde{c}_{j\downarrow}$ still anticommute. But this is easy to handle with a local 4×4 matrix. The bath operator $\tilde{c}_{\sigma}(t)$ in Eq. (14) is then represented by a matrix-product operator (MPO)

$$\tilde{c}_{\sigma}(t) = [0,1] \prod_{j=1}^{N} \begin{bmatrix} I & 0\\ \tilde{V}_{j\sigma}(t)\tilde{c}_{j\sigma} & (-1)^{n_j} \end{bmatrix} \begin{bmatrix} 1\\ 0 \end{bmatrix}, \qquad (16)$$

where the j = 1 matrix is left-multiplied by [0, 1] to pick the second row, and the j = N matrix is right-multiplied by $[1, 0]^T$ to pick the first column. The MPO has a bond dimension of 2. We can similarly express $\tilde{c}^{\dagger}_{\sigma}(t)$ in terms of $\tilde{c}^{\dagger}_{j\sigma}$. The Hamiltonian $\tilde{H}_{\text{mix}}(t)$ can then act on $|\Psi(t)\rangle$ following DMRG routines [12].

The final evolution scheme is given by

$$|\Psi(t + \Delta t)\rangle \approx e^{-i\tilde{H}_{\min}\left(t + \frac{\Delta t}{2}\right)\Delta t} |\Psi(t)\rangle$$
(17)

with the exponential factor Taylor expanded into a 4thorder polynomial of $\tilde{H}_{\rm mix}(t+\Delta t/2)$. The narrow spectral radius of $\tilde{H}_{\rm mix}\Delta t$ ensures good unitarity of the 4th-order truncation. Since the bath operators \tilde{c}_{σ} and $\tilde{c}_{\sigma}^{\dagger}$ are longrange, we cannot locally exponentiate the Hamiltonian as in the time-evolving block decimation (TEBD) [14] method and have to Taylor expand the exponential.

We adjust the truncation error tolerance of the singular value decomposition (SVD) in DMRG according to the MPS norm so that the higher-order terms of $e^{-i\tilde{H}_{\rm mix}\Delta t}$ do not take much time to calculate. The error tolerance in our code for a norm-1 MPS is set to $10^{-6}/N$ (in terms of probability loss) per SVD truncation. This number is multiplied by (norm)⁻² for MPSs with smaller norms (coefficients). If the adjusted error tolerance becomes greater than 1 (which happens if the MPS norm is very small), the MPS is truncated to a product state. The coefficients $c_i(t)$ in $|\Psi(t)\rangle$ in Eq. (8) are normalized at the end of every time step. The *d*-occupancy produced for U = 0 is found to agree in $3 \sim 4$ decimal places with a Slater-determinant based noninteracting code.

We parallelize the calculations of the 4 MPSs in Eq. (8) on 4 processors and also use the total numbers N_{\uparrow} and N_{\downarrow} of spin-up and spin-down electrons as symmetries to further speed up the calculation.

IV. RESULTS

In this section we show some results obtained for the interacting SIAM with U/E = 1 using the method and other model parameters described in Secs. II and III. The impurity-bath coupling $V/E = 0.1 \sim 0.5$. This is the parameter range of interest. The impurity level width $\sim V^2/E$ remains smaller than the band width $\sim E$ while the Kondo temperature $T_K \approx 0.4V e^{-\pi E^2/16V^2}$ (see [15]) can change by orders of magnitudes. The Kondo temperature T_K is a Hubbard U induced energy scale that measures the spin relaxation rate in the near-equilibrium regime of an interacting SIAM in the small V/E (or Kondo) limit.

Fig. 2a shows the charge relaxation dynamics, obtained by starting from an initially empty d orbital $|\Psi_0\rangle_d = |0\rangle_d$ and a half-filled Fermi-sea state $|\text{FS}\rangle_{\text{bath}}$ for the bath. Our choice of particle-hole symmetric parameters ensures that $n_d = \langle n_{d\uparrow} \rangle + \langle n_{d\downarrow} \rangle$ always equilibrates to 1 so long as the impurity-bath coupling V is not big enough to form a bound state on the impurity. We see in agreement with previous work [13, 16] that the charge equilibration proceeds relatively rapidly. The reciprocal of the time $t_{0.5}$ it takes to reach $n_d = 0.5$ is plotted in Fig. 3a. At small $V/E \leq 0.1$, $t_{0.5} \sim V^{-2}$ is inversely proportional to the d-level width $\sim V^2/E$. For $V/E \gtrsim 0.15$, the rate $1/t_{0.5}$ crosses over to approximately linear in V and the equilibration process in Fig. 2a becomes more oscillatory as we are approaching the formation of a bound state on the impurity. The variation of charge equilibration rates with V can be seen in calculations performed for a



FIG. 2. (Color online) The charge and spin dynamics of the SIAM. (a) The occupancy n_d v.s. t starting from $|\Psi_0\rangle_d = |0\rangle_d$ with impurity-bath coupling V/E = 0.2, 0.25, 0.3, 0.35, 0.4 from bottom to top; (b) The magnetic moment m_d v.s. t starting from $|\Psi_0\rangle_d = |\uparrow\rangle_d$ with the same values of V/E from top to bottom. Dashed lines show the linear fits used to obtain the long-time relaxation rates in Fig. 3b. Hubbard U/E = 1. The number of bath orbitals we used was N = 20 in (a) and N = 130 in (b).

noninteracting SIAM (U = 0), suggesting that the charge relaxation physics is essentially due to hybridization. The Hubbard U does not change the behavior of the model qualitatively.

Fig. 2b shows the spin relaxation dynamics obtained by starting from $|\uparrow\rangle_d \otimes |\text{FS}\rangle_{\text{bath}}$, a fully spin-polarized dorbital and the same half-filled Fermi-sea state $|\text{FS}\rangle_{\text{bath}}$ of the bath. The magnetization $m_d = \langle n_{d\uparrow} \rangle - \langle n_{d\downarrow} \rangle$ relaxes much more slowly than the charge, again in agreement with previous results [17, 18]. The asymptotic behavior of m_d v.s. t shows approximately an exponential tail, with the relaxation rate $\Gamma_{\infty} \equiv d \ln m_d/dt|_{t\to\infty}$ plotted in Fig. 3b. Γ_{∞} is estimated by fitting $\ln m_d(t)$ v.s. tto a straight line for $t_{\max}/2 < t < t_{\max}$, where t_{\max} is the maximum time reached in the simulation for the slope. The solid red line is a trend line. We also show as the dashed green line the analytical result — the Kondo temperature T_K calculated using the formula in [15] and interpreted as a relaxation rate.



FIG. 3. (Color online) The charge equilibration rate $1/t_{0.5}$ in (a) and the spin relaxation rate $\Gamma_{\infty} \equiv d \ln m_d/dt|_{t\to\infty}$ in (b) obtained from $n_d(t)$ and $m_d(t)$ (partly shown in Figs. 2a and 2b). Γ_{∞} is estimated using $m_d(t)$ up to $Et \leq 600$. Hubbard interaction U/E = 1.

The Kondo result has a similar magnitude and Vdependence to the calculated results. The numerical differences at large V arise from relaxation processes associated with valence fluctuations not included in the Kondo limit, while the more pronounced differences at small V are an intermediate asymptotics effect. For small V, even at the very long times ($Et \leq 600$) accessible to our method, the magnetization m_d is still substantial, so the Kondo-limit expression, which gives the linear response relaxation for small magnetization $(m_d \rightarrow 0)$, is not applicable. Evidently, the nonlinear response (relaxation of a finite m_d) is stronger than the linear response. Developing a theory of the relaxation in the small V and intermediate m_d regime is an interesting open question. For intermediate $V/E \simeq 0.25$, the theoretical result is within a factor of 2 of the numerical one with the differences likely arising from the convention used for the Kondo temperature T_K .

V. LOGARITHMIC GROWTH OF ENTROPY

A remarkable feature of the simulations reported here is the long time scales that can be reached; these time scales are necessary to reveal, for example, the magnetization decay. As we show in this section, this is possible because the maximum entanglement entropy of the 4 bath MPSs in Eq. (8) grows only logarithmically during the simulation, which means the long times are not exponentially hard to reach, but are of only polynomialtime complexity.

A. Entanglement entropy growth in SIAM

In this section, we compare the maximum entanglement entropy of the interacting SIAM (U/E = 1) with a noninteracting SIAM (U = 0) with $\epsilon_d = 0$ at the Fermi level. Both models start from the same initial condition $|0\rangle_d \otimes |FS\rangle_{bath}$ with an empty d orbital and a half-filled bath in Fig. 1. Results of the entanglement entropy are shown in Fig. 4. The entropy growth starting from a spinpolarized impurity $|\uparrow\rangle_d \otimes |FS\rangle_{bath}$ is also logarithmic but takes smaller values.

The curves in Figs. 4a and 4b are obtained in slightly



FIG. 4. (Color online) The logarithmic growth of entanglement entropy. (a) The entanglement entropy of the initially occupied part of the bath with the rest of the system at U = 0 and $V/E = 0.1, 0.2, \ldots, 0.6$ from bottom to top. (b) The maximum entanglement entropy encountered in the interacting SIAM simulation v.s. time t at U/E = 1 and V/E = 0.15, 0.2, 0.25, 0.3, 0.35 from bottom to top.

different ways. Fig. 4a shows the results obtained in a noninteracting simulation (using a Slater-determinant based code) of N = 1000 bath orbitals all coupled to one empty d orbital at the Fermi level. We then plot the entanglement entropy between the 500 bath orbitals below the Fermi level with the other 501 orbitals up to Et = 1000. The Fermi level is found to be close to the maximum entropy cut of the system. The data shows a logarithmic growth of entanglement entropy at all values of the impurity-bath coupling V. The slopes of the curves at long times are the same; only the offset and the transient entropy growth depend on V.

Fig. 4b shows the results of the interacting SIAM $(U/E = 1, \epsilon_d = -U/2)$ simulated using the method of Sec. III. Every point on a curve is obtained in a different simulation with a different bath size N. In a simulation up to time t, the hybridization function is first fitted up to t with the minimal number of bath orbitals N(typically between $10 \sim 200$) needed to keep the rootmean-square error (RMSE) of the fit under 3×10^{-4} . Then the maximum entanglement entropy S_{max} seen on all bonds of the 4 bath MPSs encountered during the simulation from 0 to t is plotted v.s. t. Notice that S_{max} may be encountered before t due to the finite bath effect. So Fig. 4b takes into account the possibility of using the finite bath effect to limit entropy growth. But still the logarithmic growth of entropy and the independence of the steady-state slope of S v.s. log t on the impurity-bath coupling V are the same as in the noninteracting SIAM in Fig. 4a. These two properties mean $S \leq c \ln t$, and therefore the bond dimension $D \sim e^S < t^c$. Hence, the interacting SIAM can be simulated in polynomial time $\mathcal{O}(D^3) = \mathcal{O}(t^{3c})$ of t.

B. Analysis of entropy growth

To understand the logarithmic growth of entropy, we consider a noninteracting chain model, as is shown in Fig. 5. In this model, the impurity is coupled to two semiinfinite chains. We choose a constant hopping amplitude between the bath sites in each chain. By adjusting the on-site energy difference of the two chains, we can vary the densities of states as shown in the lower panel of Fig. 5, obtaining either overlapping, gapped or just touching spectra. Our computation of the entanglement entropy S_{occ} across the impurity site shows that we have linear growth, logarithmic growth, and saturation, respectively. In the numerical test we did, chain awas initially empty and chain b was initially full. But the conclusion is found to hold for randomized initial occupancies, too. A similar noninteracting model with two chains directly connected via a modified hopping amplitude was studied in [19, 20] in the formalism of conformal field theory.

The logarithmic growth of S_{occ} is seen at a critical $E = V_a + V_b$, at which the density of states (DOS) of the two semi-infinite chains touch at only one energy



FIG. 5. (Color online) The two semi-infinite chain model (lower panel) and its critical behavior at $E = V_a + V_b$ (upper panel) with $\epsilon_d = 0$, $V_a = V_b \equiv V$, $V_c = 0.5V$ and E/V = 1, 1.5, 2, 2.5, 3 from top to bottom. S_{occ} is the entanglement entropy between chain *b* (initially occupied) with the rest of the system (initially empty impurity and chain *a*).

point. When $E > V_a + V_b$, the system is gapped and entropy growth saturates. This can be explained by the lack of energy eigenstates that are extended in both regions a and b, which then means that particles (or holes) that are originally in a cannot go into b and vice versa beyond a penetration depth determined by the gap, which then puts an upper bound on the entanglement entropy between a and b. This energy barrier works for a general initial occupancy. Starting from any product state, so long as the semi-chains a and b are gapped, the entropy must saturate.

When $E < V_a + V_b$, there is a finite overlap of the DOS of the two semi-infinite chains and we see a linear growth of entanglement entropy in Fig. 5. In rare cases this does not happen. For example, for a uniform chain $V_a = V_b = V_c$ and $E = \epsilon_d = 0$, the entropy growth is logarithmic rather than linear. But this behavior depends on the initial occupancy. If the occupied sites are randomized, or if the model parameters are slightly modified to deviate from a uniform chain, the expected behavior of a linear growth of the entanglement entropy is seen between a and b. The energy criterion guarantees that particles do not enter the forbidden regions of a noninteracting bath. But once the energy barrier is not at work, it is difficult in general, though not impossible, to

organize the migrated particles into a low entanglement entropy state to make the MPS matrices small.

The logarithmic growth of entropy in Fig. 4b can be understood as the result of arranging the bath orbitals in the MPS in energy order, so that at any bond of the MPS, the left and right parts of the bath degrees of freedom always have touching energy spectra. This argument applies to an interacting model, too, because the bath is still noninteracting, and the Hubbard U only reduces the chance for the impurity – the only bridge via which the bath orbitals can indirectly hop to one another – to be doubly occupied, thus reducing its bridging efficiency. The bath entanglement entropy of an interacting SIAM is therefore upper bounded by that of a noninteracting SIAM from this picture.

C. Bath in chain geometry

So far we have been working in the star geometry of the bath. Bath orbitals do not hop to each other directly. They only do so via the impurity. The diagonalization of bath orbitals in energy space leads to a logarithmic growth of entanglement entropy, according to the energy criterion in the previous section. In this section, we would like to emphasize again that the energy criterion is a sufficient but not necessary condition for the entropy to grow slowly. The example to give here is the evolution of the quenched SIAM in the chain geometry of the bath. The impurity is the head of the chain, which is directly connected to only one bath orbital, which in turn is connected to another bath orbital, and so on so forth. One can go from the star geometry to the chain geometry via Lanczos tridiagonalization starting from the impurity orbital, and from the chain back to



FIG. 6. (Color online) The entropy profiles at different times $Et = 0, 20, 40, \ldots 300$ in the chain geometry starting from $|0\rangle_d \otimes |\text{FS}\rangle_{\text{bath}}$ with $|\text{FS}\rangle_{\text{bath}}$ given in Fig. 1. Hubbard U = 0 and impurity-bath coupling V/E = 0.25. The number of bath orbitals N = 2000. S_n is the entanglement entropy of the bath orbitals $1, 2 \ldots n$ with the rest of the system.



FIG. 7. (Color online) Entanglement entropy of the noninteracting SIAM in the chain and star geometries. The impurity-bath coupling V/E = 0.25. At t = 0, the impurity is empty and the bath is half-filled. The initial occupancy is ordered if the occupied bath orbitals are n = 1, 2, ..., N/2, and random if the N/2 occupied bath orbitals are randomly shuffled. The bath orbital energies in the star geometry are ordered if they are in ascending order of n and random if they are randomly shuffled. $S_{N/2}$ is the entanglement entropy between the $n \leq N/2$ bath orbitals and the rest of the system. The number of bath orbitals is N = 2000. The random results are averaged over 10 simulations.

the star by diagonalizing the bath. More details of the two geometries can be found in [13].

Starting from the initial state $|0\rangle_d \otimes |FS\rangle_{bath}$ with $|FS\rangle_{bath}$ being the same filled Fermi-sea state as in Fig. 1 transformed to the chain geometry, the maximum entropy on the chain (the entanglement entropy between the left and right parts of the chain at the maximum entropy cut) is still found to grow logarithmically. Fig. 6 shows the result of a noninteracting calculation. The initial occupancies on the chain are spatially uniform. Every site has an occupancy of 0.5 per spin except the empty impurity. The entanglement entropy S_n between sites $1, 2, \ldots n$ and $n + 1, \ldots N$ on the chain are then plotted in Fig. 6 as a function of n at equal intervals of time. On top of the logarithmic background of S_n of the equilibrium state $|FS\rangle_{bath}$, an entropy peak propagates like a soliton from the impurity down the chain at a speed $\propto E$. The maximum entanglement entropy (height of the peak) therefore increases with time logarithmically, even though there is no separation of energy spectrum on the chain, i.e., partition of the bath into different regions with different energies like in the star geometry.

Starting from an inverted half-filled Fermi-sea state with the highest energies initially occupied, the same logarithmic growth of entropy in the chain geometry is seen due to particle-hole symmetry. But starting from a product state with random 0-1 initial occupancies of the bath orbitals in the star geometry, the entanglement profile transformed to the chain geometry becomes very high $(\max(S_n) \propto N)$ even at t = 0. Also, a linear growth of entropy is seen starting from a product state in the chain geometry with randomized 0-1 initial bath occupancies (see Fig. 7, blue line), while in the energyordered star geometry, the entropy growth (green line) is still logarithmic under the same condition. These results demonstrate that the logarithmic entropy growth in Fig. 6 is not guaranteed by the MPS basis of the Hamiltonian, but is due to the initial filled Fermi-sea state. For such a special initial state, the star geometry does not have a big advantage over the chain geometry, as they both give a logarithmic growth of maximum entanglement entropy. The benefit of the star geometry is its good behavior for more general initial states.

It is important to point out, as is shown in Fig. 7, that the star geometry alone does not guarantee a logarithmic entropy growth. The order of the bath orbitals in the MPS matters. The initial occupancies affect the transient growth of entropy, while the asymptotic entropy growth is determined by the ordering of the bath orbital energies. The steady-state growth of $S_{N/2}$ is logarithmic if the bath orbitals in the MPS are energy-ordered and linear if the bath orbital energies are randomly shuffled.

VI. DOUBLE-IMPURITY MODEL

In this section, we show that the logarithmic growth of entanglement entropy is not limited to the singleimpurity Anderson model by doing a noninteracting simulation of a double-impurity Anderson model. The most general noninteracting double-impurity Anderson model can be pictorially represented in Fig. 8. Fig. 8a is in the basis in which the 2 impurity orbitals and all bath orbitals are diagonal, which is the double-impurity version of the star geometry. Fig. 8b shows the doubleimpurity version of the chain geometry by Lanczos tridiagonalizing the star geometry in Fig. 8a starting



FIG. 8. (Color online) The general noninteracting doubleimpurity Anderson model in (a) the star geometry and (b) the chain geometry. Every orbital energy and every hopping line is an independent parameter. Panel (c) shows the doubleimpurity generalization of the model in Fig. 5.



FIG. 9. (Color online) The logarithmic growth of entanglement entropy in a noninteracting double-impurity Anderson model. The bath DOS and filling are the same as Fig. 1. The two *d* orbital energies $\epsilon_{d1,2}/E = \pm 0.2$ and $d_1 - d_2$ hopping $V_{d1,d2}/E = 0.15$. Both d_1 and d_2 are uniformly coupled to all bath orbitals with coupling amplitude V/\sqrt{N} each. The coupling $V/E = 0.1, 0.15, \ldots, 0.4$ from bottom to top. Number of bath orbitals N = 1000.

from the two impurities. One can also tridiagonalize the bath orbitals above and below the Fermi level separately (Fig. 8c) to obtain the double-impurity generalization of the two semi-infinite chain model in Fig. 5. Since the left and right semi-chains have touching energy spectra, a logarithmic growth of entropy is expected as a critical behavior between linear growth and saturation of entropy, as is discussed in Sec. V B.

Figure 9 shows a sample result. We chose a half-filled bath with a semicircle DOS the same as Fig. 1, and put two d orbitals at $\pm 0.2E$ (E is the half band width) with d - d hopping 0.15E to mimic typical crystal field splitting. The two d orbitals are equally coupled to all bath orbitals. In the basis in which the two d orbitals are diagonalized, their orbital energies are $\pm 0.25E$ and the original d-d hopping makes the two d orbitals now couple to the bath differently, which is more realistic. Then we plot the entanglement entropy $S_{\rm occ}$ between the initially occupied bath orbitals and the rest of the system.

The double-impurity model has a richer dynamics than SIAM. Since both impurities are initially empty, the one below the Fermi level leaks a hole into the bath, leading to a short-term entropy peak. The steady-state growth of $S_{\rm occ}$ is still logarithmic, but the slope of $S_{\rm occ}$ v.s. log t is not constant. This is because the two d orbitals are not at the Fermi level (one is above and one is below). Their distances in energy to the Fermi level $|\epsilon_{d1,2}|$ relative to the impurity-bath coupling V determine the slope, which approaches a maximum for the case of a d-orbital at the Fermi level ($\epsilon_d = 0$) as V gets large.

The logarithmic growth of entropy again shows that

the quenched multi-impurity model is not exponentially hard in DMRG simulations, but is of only polynomialtime complexity. Whether the conclusion still holds for interacting models needs further investigation in DMRG, especially for those multi-impurity models with nondensity-density (spin flipping and pair-hopping) terms, whose entanglement entropies need not be bounded by the corresponding noninteracting models.

VII. CONCLUSION

We have studied the growth of entanglement entropy in quenched Anderson impurity models. It is found that the growth of entropy is determined by the representation of the bath orbitals in the matrix product state (MPS). The Hubbard U on the impurity orbital does not change the qualitative behavior of the steady-state growth of entanglement entropy of the bath MPS. The crucial feature controlling the growth of entanglement entropy is the overlap in energy of the density of states of the two parts of the maximum entanglement partition. In the star geometry of energy-ordered bath orbitals, the touching-spectra condition is satisfied at every bond, so the maximum bond dimension is power law in t. The power is upper bounded by the case of a half-filled dorbital at the Fermi level and does not grow with the impurity-bath coupling, which allows a simulation of the long-time dynamics of the quenched impurity models in polynomial time. The conclusion is likely to generalize to multi-impurity models.

The growth of entanglement entropy of an interacting quantum system and the associated computational cost has been studied previously [21, 22] in terms of the integrability of the quantum model. Our study looks at the problem from a different perspective. We focus on a special class of quantum models — the impurity models — and think of the growth of entanglement entropy among the bath orbitals. Because of the sparsity of interactions in the model, the entropy growth in the noninteracting bath is controlled by the energy partitioning of the bath and the localization of bath electrons to the energies they belong to. Since the new criterion of energy-partitioning the bath is not related in obvious ways to the integrability of the whole model (bath + impurity), hopefully this new view of entropy growth of complexity can help us find new polynomial-time solvable models, parameter ranges, and/or special initial conditions that are not covered by the previously established integrability criterion.

Acknowledgments: We thank Dr. Dante Kennes for helpful discussions. This research is supported by the Department of Energy under grant DE-SC0012375.

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