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The fine structure of the entanglement entropy in the O(2) model

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We compare two calculations of the particle density in the superfluid phase of the O(2) model with a chemical potential μ in 1+1 dimensions. The first relies on exact blocking formulas from the Tensor Renormalization Group (TRG) formulation of the transfer matrix. The second is a worm algorithm. We show that the particle number distributions obtained with the two methods agree well. We use the TRG method to calculate the thermal entropy and the entanglement entropy. We describe the particle density, the two entropies and the topology of the world lines as we increase μ to go across the superfluid phase between the first two Mott insulating phases. For a sufficiently large temporal size, this process reveals an interesting fine structure: the average particle number and the winding number of most of the world lines in the Euclidean time direction increase by one unit at a time. At each step, the thermal entropy develops a peak and the entanglement entropy increases until we reach half-filling and then decreases in a way that approximately mirrors the ascent. This suggests an approximate fermionic picture.

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I. INTRODUCTION

The O(2) model, also called the classical XY model, in one space and one Euclidean time (1+1) dimensions plays an important role in the field theoretical approach of condensed matter phenomena and is prominently featured in standard textbooks[1–4]. This model provides the simplest example of a Berezinski-Kosterlitz-Thouless transition [5, 6] and it may be used as an effective theory for the Bose-Hubbard model [7]. More generally, its associated quantum Hamiltonian of Abelian rotors appears in many different contexts such as the formulation of Abelian lattice gauge theories [8], Josephson junctions arrays [9] and cold atom simulators [10, 11].

When a chemical potential μ is introduced, the model displays a rich phase diagram depicted in Ref. 10. For sufficiently small β , the inverse temperature of the classical model, if we increase μ , we go from a Mott insulating (MI) phase where the average particle number ρ remains zero until it reaches a superfluid (SF) phase where ρ starts increasing with μ . This proceeds until ρ reaches one per site and we enter in a new MI phase where it stabilizes at this value. For β small enough, this alternation of MI and SF phases repeats several times. In suitable coordinates [10], the phase diagram is similar to what is found for the one-dimensional quantum Bose-Hubbard model [12, 13].

A simple way to locate approximately the SF phase consists in calculating the thermal entropy, defined precisely in Eq. (16), on a lattice with a large enough temporal size. An example is shown in Fig. 1. The central line at β =0.1, where we do calculations hereafter, covers the first SF phase and the MI phases with ρ = 0 and 1. As we increase μ the thermal entropy goes through a sequence of peaks culminating around ln 2, signaling level crossings that will be explained below. The number of peaks equals the number of sites in the spatial direction. This is clearly a finite size feature, while the notion of phase used above should be understood in the limit of an infinite number of sites.



FIG. 1. (Color online). Intensity plot for the thermal entropy of the O(2) model on a 4×128 lattice in the β - μ plane. The dark (blue) regions are close to zero and the light (yellow ochre) regions peak near ln 2. The MI phase with ρ =0 is below the lowest light band, the MI phase with ρ =1 is above the highest light band and there is a single SF phase in between the two MI phases.

In this article, we describe microscopically the rich sequence of changes which occurs as we move across the

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SF phase along a line of constant β as described above and illustrated in Fig. 1. The figure can be embedded in the phase diagrams of Ref. 10: the lower and upper light lines ending at the tips of the the two MI phases. The fine structure that we report here is first studied for a small spatial size of four sites and then for larger sizes. At infinite spatial size, there are three phases in Fig. 1, the SF phase approximately covering the four light bands with larger values of the thermal entropy and the three darker regions in between. We also report about the numerical methods that we developed in this process.

The specific O(2) model used in this article is a planar version of the Ising model with a O(2) symmetry, on a $L_x \times L_t$ lattice. The notations used later to characterize the model and the basic numerical methods are provided in Sec. II. Most of the calculations done in this article will rely on the tensor renormalization group (TRG) method [14] which can be used to write exact blocking formulas for the model considered [15–17]. In Sec. II B, we remind how the method can be applied to the calculation of the transfer matrix [10].

It should be noticed that the presence of μ causes the action to be complex. This sign problem prevents the use of Monte Carlo simulations when μ is too large to rely on reweighing methods. However, using a Fourier expansion of the Boltzmann weights [18], it is possible to find a formulation of the partition function in terms of world lines with a positive weight as long as μ is real. This positivity allows statistical sampling [19] using a classical version of the worm algorithm [20]. The computational methods used for the worm algorithm are briefly reviewed in Sec. II C.

The TRG method can be used for arbitrary complex values [21] of β and μ . The only sources of error are the truncations of the infinite sum to finite ones as required for the numerical treatment. These truncations take place in the original formulation of the partition function and also at each step of the coarse-graining process. A first check of the agreement between the TRG and the worm methods is provided in Sec. III where we compare particle number density histograms and show that they agree well.

In the SF phase, the model is gapless in the the limit $L_x \to \infty$. At finite L_x and small β , the gap is expected to scale like $1/L_x^2$. This follows from a non-relativistic dispersion relation and can be justified using degenerate perturbation theory. An invaluable method to study the long range correlations in near gapless situations is to calculate the entanglement entropy. Interestingly, it seems possible to measure the entanglement entropy in many-body systems implemented in cold atoms [22–28]. This includes studies of rather small L_x systems. The entanglement entropy needs to be distinguished from the thermal entropy [25]. For this reason, in Sec. IV we discuss these two entropies using the transfer matrix formalism.

We use a lattice version of the setup of Calabrese and Cardy [29, 30] for 1+1 dimensional nonlinear sigma models. We consider the case of finite, but often large, L_t . A finite L_t introduces a temperature proportional to $1/L_t$, distinct from $1/\beta$ used in the classical formulation, and therefore we have a thermal density matrix. In this context, the relation between the two entropies is a rather open topic of investigation[31]. Numerical calculations of the related Renyi entropy of the classical XY model, without chemical potential, were presented in Ref. 32.

In the theoretical framework developed in Refs. 29 and 30, it is common to move back and forth between the classical and quantum formulation using the transfer matrix. An interesting example is given in Ref. 33 for the O(N) model. In our article, the discussion is focused on the classical formulation. A quantum Hamiltonian corresponding to the model considered here is provided in Eq. (30) of Ref. 11. Reasonable approximations are obtained for a spin-1 truncation (3 states per sites). In general, the spin-1 quantum XY model with an extra $(\mathbf{S}^{z})^{2}$ term in the Hamiltonian is closely related to the model studied here. In addition, the relationship between these quantum models and a family of Bose-Hubbard models with large on-site repulsion, possibly implementable on optical lattices, is discussed in Refs. 10 and 11. The approximate description in terms of hardcore bosons suggests an underlying fermionic description of our findings. Furthermore, for μ corresponding to the superfluid region, the state $|\mathbf{S}^z = -1\rangle$ plays a negligible role and an approximate connection with the spin-1/2quantum XY model with a magnetic field emerges. This can be explained by the fact that \mathbf{S}^{z} and $(\mathbf{S}^{z})^{2}$ have the same effect on the remaining subspace of with eigenvalues 0 and 1.

With the TRG method, we approximate the reduced density matrix by a finite dimensional matrix which can be diagonalized numerically and we do not need to use the replica trick as in Refs. 29, 30, and 32. We then show how to use the TRG method to express the entanglement entropy for a bipartition of the system. We show that for small L_t , the thermal entropy is larger than the entanglement entropy but the entanglement entropy becomes larger as L_t is increased.

In Sec. V, we use degenerate perturbation theory to get an approximate idea of the large structure (location of SF phases) and fine structure (changes of ρ and entanglement entropy across one SF phase) of the phase diagram. We relate in some approximate way, the eigenvectors of the transfer matrix with particle number n to world-line configurations with a winding number which is also n. We then calculate numerically the average particle number density, the thermal entropy and the entanglement entropy for values of μ spanning the first SF phase as explained above.

For sufficiently large L_t , the results show an interesting fine structure: the particle number and the winding number of (most of) the world lines increase by one unit at a time as we keep increasing μ . At each step, the thermal entropy develops a peak and the entanglement entropy increases with μ until ρ reaches half-filling. As we keep increasing μ beyond this value, the entanglement entropy decreases in a way that approximately mirrors its ascent. This approximate symmetry can be justified by noticing that if we interchange the occupied links with the unoccupied links in the world lines, we transform a configuration with particle number n to one with a particle number $L_x - n$. This approximate particlehole transformation can be reformulated in the context of degenerate perturbation theory and is similar to an exact symmetry of the spin-1/2 quantum XY model in an external magnetic field briefly explained in the conclusions.

Our results are summarized in Sec. VI where we also briefly discuss work in progress. We suggest ways to reduce the small truncation errors reported in Sec. III and to interpret the approximate particle-hole symmetry found in Sec. V. We also briefly discuss the relationship of our work with Polyakov's loop studies [34, 35] and with recently proposed cold atom experiments [36].

II. NOTATIONS AND NUMERICAL METHODS

In this section, we introduce the O(2) model with a chemical potential. We describe the two numerical methods used in this article. The first is the TRG which relies on coarse-graining, the second is a worm algorithm which relies on sampling. This section contains important concepts used in the next sections, such as the representation of the partition function in terms of integers labeling Fourier modes attached to the links, also called bonds, of the lattice, their use in the transfer matrix formulation and their graphical illustration as world lines. The section also contains technical details about the numerical implementations that are given for completeness.

A. The model

We consider the O(2) model, sometimes called the classical XY model, with one space and one Euclidean time direction, and a chemical potential μ . One can interpret μ as the imaginary part of a constant gauge field in the temporal direction. The sites of the rectangular lattice are labelled (x, t) and the unit vectors denoted \hat{x} and \hat{t} . The line segments joining two nearest neighbor sites are called links. The total number of sites is $L_x \times L_t$. In the following, we are typically interested in the case $L_t >> L_x$. We assume periodic boundary conditions in space and time. The partition function reads

$$Z = \int \prod_{(x,t)} \frac{d\theta_{(x,t)}}{2\pi} e^{-S}$$
(1)

with

$$S = -\beta_{\hat{t}} \sum_{(x,t)} \cos(\theta_{(x,t+1)} - \theta_{(x,t)} - i\mu)$$

$$-\beta_{\hat{x}}\sum_{(x,t)}\cos(\theta_{(x+1,t)}-\theta_{(x,t)}).$$
(2)

In all the numerical calculations done in this article, we consider space-time isotropic couplings $\beta_{\hat{x}} = \beta_{\hat{t}} = \beta$. However, if we set $\beta_{\hat{x}}$ to zero, the model becomes a collection of decoupled solvable models. For analytical purposes, when β is small, it is sometimes convenient to first consider the solvable case $\beta_{\hat{x}}=0$ and then restore the isotropic situation $\beta_{\hat{x}} = \beta_{\hat{t}} = \beta$ perturbatively (see Sec. V).

As explained in Refs. 10, 18, and 19, one can use the Fourier expansion of $e^{\beta \cos \theta}$ in terms of modified Bessel function of the first kind and then integrate out the $\theta_{(x,t)}$ variables. The Fourier indices associated with the links coming out of the site (x,t) in the space and time directions are denoted $n_{(x,t),\hat{x}}$ and $n_{(x,t),\hat{t}}$ respectively. They can be graphically interpreted as currents passing through the links. The partition function can then be expressed as a sum of product of Bessel functions:

$$Z = \sum_{\{n\}} \prod_{(x,t)} I_{n_{(x,t),\hat{x}}}(\beta_{\hat{x}}) I_{n_{(x,t),\hat{t}}}(\beta_{\hat{t}}) e^{\mu n_{(x,t),\hat{t}}} \times \delta_{n_{(x-1,t),\hat{x}} + n_{(x,t-1),\hat{t}}, n_{(x,t),\hat{x}} + n_{(x,t),\hat{t}}} .$$
(3)

The Kronecker delta function in Eq. (3) can be interpreted as a current conservation and the terms in the partition function as current loops (also called world lines) which can then be statistically sampled [19].

The action given in Eq. (2) has a manifest global O(2) symmetry which corresponds to shifting all the angles $\theta_{(x,t)}$ by the same arbitrary amount. This symmetry is the same as the U(1) symmetry of a relativistic complex scalar field, where the invariance under multiplication by a global phase implies the particle number conservation. The sum of the $n_{(x,t),\hat{t}}$ on the time links between two time slices is conserved and is the particle number. In Refs. 10 and 11, we discuss Bose-Hubbard models related to the O(2) model discussed here where the particle number is actually related to numbers of bosonic atoms.

For a system with periodic boundary condition in both space and time directions, as considered in this paper, the world line can wind around in both directions. The winding numbers are important to understand the superfluid properties of the system [37].

An typical allowed configuration for $\beta=0.1$ and $\mu=3$ is shown in Fig. 2 for $L_x=4$ and $L_t=32$. Sites at the boundary should be identified as explained in the figure caption. The uncovered links on the grid have n=0, the more pronounced dark lines have |n| = 1 and the wider lines have n=2. A discussion regarding the sign convention and the spatial winding number of Fig. 2 can be found in Appendix A.

This configuration can be used to visualize a transfer matrix that connects consecutive time slices. For instance, in Fig. 2, the time slice 5 represents a transition between $|1100\rangle$ and $|0200\rangle$ and its relative statistical weight can be obtained from a transfer matrix that will be discussed in Sec. II B. The configuration was



FIG. 2. (Color online). Graphical representation of an allowed configuration of $\{n\}$ for a 4 by 32 lattice, $\beta = 0.1$ and $\mu = 3.0$. The uncovered links on the grid have n=0, the more pronounced dark lines have |n|=1 and the wider lines have n=2. The signs of the links with |n|=1 lines are discussed in the Appendix A. The large dots on the vertical boundaries (red) need to be identified in pairs with the same t coordinates. Similarly, the slightly smaller dots (blue) on the horizontal boundaries have to be identified in pairs with the same x coordinate.

generated using a sampling method designed in Ref. 19 and briefly discussed in Sec. II C.

B. TRG approach of the transfer matrix

As explained in Ref. 10, the partition function can be expressed in terms of a transfer matrix:

$$Z = \operatorname{Tr} \mathbb{T}^{L_t} . \tag{4}$$

The matrix elements of \mathbb{T} can be expressed as a product of tensors associated with the sites of a time slice (fixed t) and traced over the space indices. To make the equations easier to read, we use the notations n_x for the time indices in the past $(n_{(x,t-1),\hat{t}})$, the primed symbols n'_x for the time indices in the future $(n_{(x,t),\hat{t}})$ and \tilde{n}_x for the space indices $(n_{(x,t),\hat{x}})$. The matrix elements of \mathbb{T} have the explicit form

$$\mathbb{I}_{(n_{1},n_{2},\dots,n_{L_{x}})(n'_{1},n'_{2},\dots,n'_{L_{x}})} = \sum_{\tilde{n}_{1}\tilde{n}_{2}\dots\tilde{n}_{L_{x}}} T^{(1,t)}_{\tilde{n}_{L_{x}}\tilde{n}_{1}n_{1}n'_{1}} T^{(2,t)}_{\tilde{n}_{1}\tilde{n}_{2}n_{2}n'_{2}\dots} \\ \dots T^{(L_{x},t)}_{\tilde{n}_{L_{x}-1}\tilde{n}_{L_{x}}n_{L_{x}}n'_{L_{x}}},$$
(5)

with

$$T_{\tilde{n}_{x-1}\tilde{n}_{x}n_{x}n'_{x}}^{(x,t)} = \sqrt{I_{n_{x}}(\beta_{\hat{t}})I_{n'_{x}}(\beta_{\hat{t}})\exp(\mu(n_{x}+n'_{x}))} \sqrt{I_{\tilde{n}_{x-1}}(\beta_{\hat{x}})I_{\tilde{n}_{x}}(\beta_{\hat{x}})} \delta_{\tilde{n}_{x-1}+n_{x},\tilde{n}_{x}+n'_{x}} .(6)$$

This construction can be represented graphically in a way that helps understanding the allowed configurations such as the one depicted in Fig. 2. The building blocks are the microscopic tensors of Eq. (6) which are represented as crosses with spacial half links in the horizontal direction representing the \tilde{n} indices and vertical half links representing the n (bottom) and n'(top) indices. The traces over the spacial indices \tilde{n} in Eq. (5) are represented in Fig. 3 as joined half links forming a complete horizontal link. The product of transfer matrices are obtained by tracing the time indices and can be represented by piling up copies of Fig. 3 in the vertical direction. The trace in Eq. (4) can be visualized as a cylinder, or a torus if periodic boundary conditions in space are imposed.



FIG. 3. Graphical representation of the transfer matrix $\mathbb{T}_{(n_1,n_2,\ldots,n_{L_x})(n'_1,n'_2,\ldots,n'_{L_x})}$ as defined in Eq. (5). The horizontal indices are summed over and the vertical indices are left open as explained in the text.

The Kronecker delta function in Eq. (6) is the same as in Eq. (3) and reflects the existence of a conserved current as discussed above, that we will call "particle number" in the following. For periodic (spatial trace) or open (zero spatial indices at both ends) boundary condition, the local conservation law implies that the transfer matrix elements are zero unless the sum of the two sets of indices (respectively denoted n and n' in Eq. (5)) are equal. In other words, the transfer matrix is block diagonal in each particle number sector. The particle number that reflects the above conservation law is defined as

$$n = \sum_{x=1}^{L_x} n_x = \sum_{x=1}^{L_x} n'_x . \tag{7}$$

When the chemical potential μ is zero, there is a charge conjugation symmetry which allows us to change the sign of all the *n*'s in all the temporal sums without affecting the final results. Given the rapid decay of the modified Bessel function when the index *n* increases, good approximations can be obtained by replacing the infinite sums by sums restricted from $-n_{max}$ to n_{max} . When the chemical potential is nonzero, it is more efficient to shift the range in the same direction as the sign of the chemical potential. In general, we call D_s the number of states kept after truncation. In the following, we will use a coarse-graining procedure for the transfer matrix where we repeatedly apply a truncation at the same value D_s .

For numerical purposes, we reduce the size of the transfer matrix by using a blocking procedure [14, 16]. It consists in iteratively replacing blocks of size two in the transfer matrix by a single site using a higher order singular value decomposition. During the truncation, for a given pair of sites, we replace the direct product $D_s^2 \times D_s^2$ matrix by a $D_s \times D_s$ matrix. The process is illustrated in Fig. 4. The Y-shaped parts in the bottom part of the figure and their mirror versions in the top part, as the truncating matrix $D_s^2 \times D_s$ [14] in the tensor product, schematically represent this coarse-graining process. The full technical details can be found in Refs. 14 and 16.



FIG. 4. Graphical representation of the coarse graining truncation of the transfer matrix described in Fig. 3. The blocking procedure of replacing blocks of two sites in the transfer matrix into a single site represented by the Y-shaped structure.

C. The worm algorithm

Because of the conservation law in Eq. (6), the terms of the partition function can be interpreted as current loops which can be statistically sampled [19]. The sampling procedure for one complete worm algorithm update goes as follows. We pick randomly a site on the lattice, then pick randomly a neighboring direction in the (positive or negative) spatial or time direction, change the current n to $n \pm 1$ with a given probability(the details of the accept-reject procedure are given in the Appendix A of Ref. 19). We repeat the above procedure until we come back to the original random site. Because of the conservation law, a particle number can be attributed to a configuration. It can be calculated by summing the temporal n's between any two time slices. The particle number distribution can be calculated by generating a large number of configurations according to the above procedure.

An easily controllable source of error is the limited statistics. There is also an unavoidable truncation error. Following Ref. 19, |n| is constrained to be not larger than 20 in Eq. (3). However, given that $I_{20}(0.1) \exp(3 \times 20)/I_0(0.1) \simeq 4 \times 10^{-19}$, this is negligible for our calculations. By construction, there is a single worm and the algorithm cannot generate disconnected loops. This is not believed to be a significant source of error. Comparisons at small volumes where the truncation errors are controllable [38] indicate that the worm algorithm is statistically exact.

III. PARTICLE DENSITY CALCULATIONS

In this section, we show how to calculate the average particle density using the TRG formulation. We then apply the method numerically and compare the results with the ones obtained with the worm algorithm. The particle number conservation can also be exploited in the TRG approach. For the initial one-site tensor $T^{(x,t)}$ in Eq. (6), we have

$$T'^{(x,t)} \equiv \partial T^{(x,t)} / \partial \mu = \frac{1}{2} (n_x + n'_x) T^{(x,t)} .$$
 (8)

Here, n_x and n'_x are the particle number associated with the time indices of the original tensor $T^{(x,t)}$ in Eq. (6) with the tensor indices omitted. Consequently, we can associate a particle number n(i) with each eigenvalue λ_i of the transfer matrix \mathbb{T} . The average particle number density is an extensive quantity defined as

$$\rho = \frac{1}{L_x L_t} \frac{\partial \ln Z}{\partial \mu}.$$
(9)

From the expression of Z in terms of the transfer matrix and the cyclicity of the trace, we have

$$\partial Z/\partial \mu = L_t \operatorname{Tr}(\mathbb{T}'\mathbb{T}^{L_t - 1}), \qquad (10)$$

where $\mathbb{T}' = \partial \mathbb{T}/\partial \mu$ can be calculated by applying the chain rule to Eq. (5). This can be achieved iteratively by defining an "impurity" tensor initialized with the derivative of the initial tensor and then blocked and symmetrized with the original "pure" tensor. This

guarantees the recursive replacement of $T^{(x,t)}$ by $T'^{(x,t)}$ in the transfer matrix as prescribed by the chain rule. This procedure can be simplified if we know the particle number n(i) associated with each eigenvalue λ_i as discussed above. We can then write

$$\frac{1}{L_t} \frac{\partial \ln Z}{\partial \mu} = \frac{\sum_i \lambda_i^{L_t} n(i)}{\sum_i \lambda_i^{L_t}}.$$
(11)

In practice, finding the particle number associated with the eigenvalues is not completely straightforward. It requires to keep track of the particle number in the projected basis or to write the blocking algorithm sector by sector. The enforcement of the conservation law in the coarse-graining process makes the numerical calculation more stable. The tensor elements that violating the conservation law are exactly zero. If we only handle nonzero elements, we can reach relatively larger values of the dimension D_s in the truncation procedure and get more accurate results. We enforce the conservation law in TRG calculations done below.

Knowing the n(i) associated with λ_i also allows us to define a probability P(n) for the particle number n:

$$P(n) = \sum_{i:n(i)=n} \lambda_i^{L_t} / \sum_i \lambda_i^{L_t}.$$
 (12)

These probabilities can also be calculated directly from histograms obtained with the worm algorithm. Both methods can be used to calculate and compare the average particle number density using

$$\rho = \frac{1}{L_x} \sum_n nP(n) . \tag{13}$$

With the TRG method, we studied the distribution of P(n) for various values of μ spanning a range covering the boundary of the MI phase and the SF phase. The other parameters are kept fixed at $D_s = 201, \beta = 0.1,$ $L_x = 32$, and $L_t = 128$. When $\mu = 2.8$ and 2.85, the distribution bears one-bin structure with n = 0, corresponding to the MI phase. When $\mu = 2.9, 2.95$ and 3, the distribution carries more bins, corresponding to the SF phase, which shows that the phase transition occurs in the range $\mu = (2.85, 2.9)$ as illustrated in Fig. 5. We then compared the distributions obtained with the TRG and the worm algorithm for $\mu=3$, which is near the middle to the SF phase. The results are shown in Fig. 6. To the best of our knowledge, the errors associated with the worm calculations are purely statistical. The errors made with the TRG are due to the repeated truncation from D_s^2 to D_s indices in the coarse-graining process. We have used $D_s=101$, 201 and 301 and the variations are comparable to the worm errors. Overall, the results from $D_s = 301$ are very close to the worm calculation. The particle number distribution calculated from the worm histograms are from averaging over configurations generated with twelve different initial random seeds and one million configurations per seed. When the truncation



FIG. 5. (Color online). The particle number distribution P(n) with β =0.1 when μ varies between the MI and SF phases. For μ = 2.80 and 2.85, we only have n = 0. For μ = 2.90, 2.95 and 3.0, there are three visible groups of bins. With μ increasing, the distribution shifts to the right with larger most probable particle number.



FIG. 6. (Color online). Comparison of the particle number distribution P(n) between the worm algorithm and TRG with different D_s .

dimension D_s is fixed, the accuracy becomes worse with increasing L_x . When the time dimension L_t becomes large enough, the particle density distribution is almost centralized in one bin and therefore varies less with increasing D_s .

IV. CALCULATION OF THE THERMAL AND ENTANGLEMENT ENTROPY

In this section, we explain how to compute the thermal entropy and the entanglement entropy in a consistent way. We construct the reduced density matrix following Refs. 29 and 30. More specifically, we consider the path integral representation of the thermal density matrix (Eq. (6) in Ref. 29) for the O(2) model. However, we use the $(n_1, n_2, \dots, n_{L_x})$ representation and the corresponding transfer matrix introduced in Sec. II B rather than the original spin variables. In the following, the eigenstates of the transfer matrix will be treated as quantum states.

We consider a system, denoted AB, and subdivide it into two parts denoted A and B. We first define the thermal density matrix $\hat{\rho}_{AB}$ for the whole system

$$\hat{\rho}_{AB} \equiv \mathbb{T}^{L_t} / Z \ . \tag{14}$$

We have the usual normalization Tr $\hat{\rho}_{AB} = 1$. If the largest eigenvalue of the transfer matrix is non degenerate with an eigenstate denoted $|\Omega\rangle$, we have the pure state limit

$$\lim_{L_t \to \infty} \hat{\rho}_{AB} = |\Omega\rangle \langle \Omega| \,. \tag{15}$$

In the following, we will work at finite L_t and will deal with the entanglement of thermal states [39]. In general, the eigenvalue spectrum $\{\rho_{AB_i}\}$ of $\hat{\rho}_{AB}$ can then be used to define the thermal entropy

$$S_T = -\sum_i \rho_{AB_i} \ln(\rho_{AB_i}). \tag{16}$$

The subdivision of AB into A and B refers to a subdivision of the spatial indices. We define the reduced density matrix $\hat{\rho}_A$ as

$$\hat{\rho}_A \equiv \mathrm{Tr}_B \hat{\rho}_{AB}.\tag{17}$$

We define the entanglement entropy of A with respect to B as the von Neumann entropy of this reduced density matrix $\hat{\rho}_A$. The eigenvalue spectrum $\{\rho_{A_i}\}$ of the reduced density matrix can then be used to calculate the entanglement entropy

$$S_E = -\sum_i \rho_{A_i} \ln(\rho_{A_i}). \tag{18}$$

The computation of the thermal entropy can be performed using the eigenvalues of the transfer matrix λ_i discussed in Sec. III together with the normalization $\rho_i = \lambda_i^{L_t} / \sum_j \lambda_j^{L_t}$. Note that if we introduce a temperature T and energy levels by identifying $\lambda_i^{L_t}$ with $\exp(-E_i/T)$, we recover the standard relation $S_T = \langle E \rangle / T + \ln Z$. This identification makes clear that $T \propto 1/L_t$. Later, in Eq.(19), we will use units where $T = 1/L_t$.

For the computation of the entanglement entropy , we assume that A has 2^{ℓ_A} sites and B has 2^{ℓ_B} sites and then perform the ℓ_A and ℓ_B blockings for the subsystems. The coarse-graining along the spatial direction ends with two sites, one for A and the other for B. We can then contract the indices from the two sites in the time direction without further truncation. Tracing over the



FIG. 7. (Color online). Illustration of the entanglement entropy calculation. The horizontal lines represent the traces over the space indices. There are L_t of them, the missing ones being represented by dots. The two vertical lines represent the traces over the blocked time indices in A and B.

space indices linking A and B, we obtain the transfer matrix $T_{(\{n_A\},\{n'_A\},\{n_B\},\{n'_B\})}$ for the whole AB system. Taking the L_t power, tracing over n_B , and normalizing, we obtain the reduced density matrix $\hat{\rho}_A(\{n_A\},\{n'_A\})$. This is illustrated in Fig. 7.

In the following, we consider the case of $\ell_A = \ell_B$. There is no numerical difficulty in considering the system with unequal ℓ_A and ℓ_B since the space indices are not renormalized in the transfer matrix approach.

As a first study, we have considered the cases $\beta = 0.1$, $L_x = 4$ and $L_t = 16$, 32, 64 and 128 with μ between 2.8 and 3.2. Both the thermal entropy and entanglement entropy develop a peak over the SF phase. We see that for small L_t , the thermal entropy is larger than the entanglement entropy, but as we increase L_t , the entanglement entropy becomes larger than the thermal entropy. The results are shown in Fig. 8. For both the thermal entropy and the entanglement entropy, a fine structure appears for large L_t . This is discussed in the next section.

V. THE FINE STRUCTURE OF THE SF PHASE

In this section, we first give an idea of the large structure of the phase diagram (where the different SF and MI phases are located) and then explain in more details the fine structure of a single SF phase, more specifically along a line of constant β and where μ is varied in order to interpolate between the the two MI phases with successive values of ρ .

At small β , we have mostly MI phases with small SF phases in between them. A good qualitative picture can be obtained by temporarily considering the anisotropic case $\beta_{\hat{x}} = 0$ where the problem is exactly solvable and then restoring $\beta_{\hat{x}} = \beta_{\hat{t}} = \beta$ perturbatively. When $\beta_{\hat{x}} = 0$, we have only onsite interactions and in the large L_t limit, the problem reduces to finding the value n^* of n which



FIG. 8. (Color online). Entanglement entropy (EE, dash line) and thermal entropy (TE, solid line) for $\beta = 0.1$, $L_x = 4$ and $L_t = 16, 32, 64$ and 128.

maximizes $I_n(\beta_{\hat{t}})e^{\mu n}$ for given μ and $\beta_{\hat{t}}$. The largest eigenvalue of \mathbb{T} is then $(I_{n^*}(\beta_{\hat{t}})e^{\mu n^*})^{L_x}$. The quantum picture is that for large L_t , the relevant state has all L_x sites in the n^* state and we are in the MI phase with $\rho = n^*$. In summary, the approximate large structure at small β is obtained by increasing μ from zero and going through the MI phases with $n^* = 0, 1, 2, \ldots$.

The fine structure of the SF phase between the MI phases can be approached by restoring $\beta_{\hat{x}} = \beta_{\hat{t}}$ perturbatively. The SF phases are approximately located near values of μ where n^* changes. To be specific, we will consider the example of $\beta = 0.1$ and $L_x = 4$, where the transition occurs near $\mu_c = 2.997\cdots$ when $\beta_{\hat{x}} = 0$. In this limit, we have 16 degenerate states $|0000\rangle$, $|1000\rangle$, ..., $|1111\rangle$ which can be organized in "bands" with n = 0 (1 state), n = 1 (4 states), etc. Below, we call the approximation where the indices inside the kets are only 0 or 1 the "two-state approximation".

The effect of $\beta_{\hat{x}}$ is to give these bands a width and lift the degeneracy. The energy levels are defined in terms of the eigenvalues of the transfer matrix as

$$E_i = -\ln(\lambda_i). \tag{19}$$

If we plot the energy levels versus μ , we see that we have successive crossings corresponding to states of increasing n. This is illustrated with the two lowest energy levels in Fig. 9. Notice the piecewise linear behavior with slopes corresponding to the particle number 0, -1, -2, -3 and -4. As the levels cross the thermal entropy rises to ln 2.

We observe that near $\mu = 2.90$, the lowest energy state changes from $|0000\rangle$ to a state with n = 1

$$|\Omega, n = 1\rangle = \frac{1}{2}(|1000\rangle + |0100\rangle + |0010\rangle + |0001\rangle)$$
. (20)

It is easy to calculate the reduced density matrix for A defined as first two sites and B as the last two sites in



FIG. 9. (Color online). The two lowest energy levels (EL, solid lines) as a function of μ for $L_x = 4$, $L_t = 256$, $\beta = 0.1$ and $D_s = 101$. As μ increases, lines of successive slopes 0, -1, -2, -3, and -4 are at the lowest level. At each crossing, the thermal entropy(dash dot line) jumps. The values of thermal entropy are shifted vertically by -0.5 to make the figure readable.

the limit where L_t becomes infinite and for values of μ where $|\Omega, n = 1\rangle$ is the unique ground state

$$\hat{\rho}_A = \operatorname{Tr}_B |\Omega, n = 1\rangle \langle \Omega, n = 1|$$

$$= \frac{1}{4} (|10\rangle + |01\rangle) (\langle 10| + \langle 01|) + \frac{1}{2} |00\rangle \langle 00| .$$
(21)

The eigenvalues of $\hat{\rho}_A$ are 1/2, 1/2, 0 and 0 and the entanglement entropy of this reduced density matrix is ln 2.

A n = 2 state becomes the ground state near $\mu = 2.95$. It is in good approximation a linear superposition of the 6 states with two 0's and two 1's. The two states $|1010\rangle$ and $|0101\rangle$ have a slightly larger coefficient suggesting weak repulsive interactions. In the numerical expression of this ground state, we also have contributions from states such as $|2000\rangle$ but with a small coefficient. In general, for small β , the two state approximation is good (the corrections are small).

A n = 3 state becomes the ground state near $\mu = 3.03$. The ground state can in good approximation be described as

$$|\Omega, n = 3\rangle = \frac{1}{2}(|0111\rangle + |1011\rangle + |1101\rangle + |1110\rangle) , (22)$$

which is $|\Omega, n = 1\rangle$ with 0's and 1's interchanged and one can interpret the 0 as "holes". Finally, near $\mu=3.10$, $|1111\rangle$ becomes the ground states (with again many small corrections). In general, there is approximate mirror symmetry about the "half-filling" situation.

A similar approximate two state description is valid in the next SF phase located between the MI phases with ρ 1 and 2. One just needs to replace 0 by 1 and 1 by 2. We now follow the same path in the phase diagram but from the point of view of the world lines generated with the worm algorithm with $L_t=256$. Typical world lines are displayed in Fig. 2. Given that $L_t=256$ is relatively large, by taking μ approximately in the middle of the density plateaus (discussed below, see Fig. 11), we obtain that most configurations have *n* corresponding to the average value. Figure 10 shows typical results for $\mu = 2.93$ (n=1), 3.00 (n=2), 3.07 (n=3) and 3.14 (n=4) using a graphical representation similar to Fig. 2. Their spatial winding number are discussed in Appendix B. The most



FIG. 10. Worm configurations for $\mu = 2.93$, n=1 (a); $\mu = 3.00$, n=2 (b); $\mu = 3.07$, n=3 (c) and 3.14, n=4 (d). In all cases, $\beta = 0.1$, $L_x = 4$ and $L_t = 256$.

important feature of these configurations is that almost all the |n|'s are 0 or 1. For a given particle number n, between most time slices we have n time links carrying a current 1 and $L_x - n$ time links carrying no current. In rare occasions, we observe that the lines merge or cross. Overall, we can think of these configurations as a set of weakly interacting loops carrying a current 1. This is in line with the dominance of the states like $|1010\rangle$ over states like $|2000\rangle$ discussed above.



FIG. 11. (Color online). Fine structure of the entanglement entropy (EE, blue circle), thermal entropy (TE, green square) and particle density ρ (red triangle) spanning the MI($\rho = 0$), SF and MI($\rho = 1$) phases successively with μ increasing at fixed $\beta = 0.1$. There are three different system sizes: $L_x =$ 4 with $L_t = 256$, $L_x = 8$ with $L_t = 512$, and $L_x = 16$ with $L_t = 1024$, respectively. The thermal entropy has L_x peaks culminating near $\ln 2 \simeq 0.69$; ρ goes from 0 to 1 in L_x steps and the entanglement entropy has an approximate mirror symmetry near half filling where it peaks.

We now proceed to calculate the thermal entropy and the entanglement entropy for $L_x = 4$ with $L_t = 256$ (as discussed above), and also for larger lattices $L_x = 8$ with $L_t = 512$, and $L_x = 16$ with $L_t = 1024$. Figure 11 shows the fine structure in the SF phase for increasing sizes. In each case, we go through the MI($\rho = 0$), SF, and MI($\rho = 1$) phases successively as we keep increasing μ while keeping the other parameters fixed, as already illustrated in Fig. 1. We kept the same truncation dimension $D_s = 101$ for the three cases.

From Fig. 11, we observe an oscillating structure in the entanglement entropy and the thermal entropy. The number of sites in the spatial direction L_x dictates the fine structure. There are L_x transition points in the entanglement entropy and L_x peaks in the thermal entropy. With L_x increasing, the transition points close to the two boundaries become difficult to resolve as shown in the case of $L_x = 16$, $L_t = 1024$ in Fig. 11(c). Higher μ resolution and larger D_s are needed to obtain a clearer picture of the oscillations. The approximate mirror symmetry of the entanglement entropy with respect to the half-filling point persists for larger lattices. As far as the thermal entropy is concerned, the height of the peaks are all close to $\ln 2$, which corresponds to the two-fold degeneracy in the ground state. The peaks are located at the place where the energy levels from the ground state and the first excited state cross as shown in Fig. 9.

As L_x increases, the fine structure near the two boundaries connected to the MI phases is not so prominent as the middle regime and a better resolution is needed to discern the fine structure. A similar discussion applies to the situation when the system experience the $MI(\rho = i)$, SF, $MI(\rho = i + 1)$ phases successively and we have checked that the same type of fine structure appears. Note also that for small L_x and large enough L_t , ρ has significant plateaus that could be qualified as incompressible regions. However, the width of these regions shrinks like $1/L_x$ as L_x increases.

VI. CONCLUSIONS

In summary, we have used the TRG method with the conservation law implemented explicitly to calculate the particle number density, the thermal entropy and the entanglement entropy for the O(2) model with a chemical potential. The particle number distributions agree well but not perfectly with the results obtained with the worm algorithm. The discrepancies seem to increase with L_x . This can probably be explained by the fact that at small β , the spectrum has bands with L_x , $\frac{L_x(L_x-1)}{2}$ etc ... states with increasing n. Some of these bands merge when μ is increased enough to get in the SF phase. As L_x increases, the bands become denser and we will typically keep the lowest energy states irrespectively of their particle number. Taking into account the particle number in the truncation process may help getting more

accurate distributions. The energy bands are illustrated in Fig. 12. More generally, the decomposition of states in charge sectors could help understanding truncation errors [40] more systematically. This question is under active investigation.



FIG. 12. (Color online). Energy levels for $\beta = 0.1, L_x = 4$.

Besides the large structure associated with the alternation between SF and MI phases with integer particle number density, we found that in the SF regime, there is an interesting fine structure controlled by the spatial dimension L_x . The entanglement entropy, the thermal entropy and the particle number density vary in a way that is consistent with each other. The thermal entropy shows L_x peaks located at where the energy levels from the ground state and first excited state cross. Degenerate perturbation theory explains why the energy levels cross while varying the chemical potential. As a result, the step-wise structure occurs in the particle number density in the SF regime, as already observed in Ref. 19. The particle number and the winding number of the world lines increase by one unit at a time. The entanglement entropy shows L_x steps and an approximate mirror symmetry with respect to the halffilling point. The details of the fine structure depend on the ratio L_x/L_t and the infinite volume limit of the twodimensional classical model needs to be defined carefully.

It is clear that increasing L_t generally emphasizes the sharp features. The dependence on the aspect ratio L_x/L_t in the infinite volume limit has been studied for a hard-core boson Hamiltonian at half filling, which is equivalent to the quantum S = 1/2 XY model [41] and for two solvable models at the conformal critical point [42]. We are planning to carry on a similar study for the model considered here.

In the approximation where the eigenstates of the transfer matrix are made of states with only 0 and 1 for all sites, or correspondingly if the world lines have mostly links carrying currents with |n| equal to 0 or 1, this symmetry corresponds to interchanging 0 and 1

and the particle number n with $L_x - n$ which explains the approximate mirror symmetry of the entanglement entropy.

From Refs. 10 and 11, the restriction to states with only 0 and 1 magnetic quantum number simplifies the Hamiltonian (\mathbf{S}^z and (\mathbf{S}^z)² have the same effect on the restricted subspace) and suggests the correspondence with the spin-1/2 quantum XY model on a line with L_x sites and a magnetic field h (in suitable units). One can use the exact equivalence to a quadratic hamiltonian for a spinless fermion and show that the fermion number is $n = L_x(1 - \arccos(h)/\pi)$. By changing $h \to -h$, one changes $n \to L_x - n$. Here h corresponds to $\mu - \mu_{h.f.}$ in the O(2) problem, with $\mu_{h.f.}$ the chemical potential for $n = L_x/2$.

As a general summary, in the SF phase, the chemical potential determines the relevant particle number sector. In each sector, the sum of the indices of the transfer matrix adds up to a given number n, however there are n + 1 ways to distribute the total particle number over the two subsystems A and B. The entanglement entropy defined by Eq. (18) characterizes this uncertainty. We are in the process of developing new numerical methods for a better general understanding of this question.

We believe that the fine structure of the entanglement entropy described in our article can be observed in quantum models such as the Bose-Hubbard, as studied in Refs. 12 and 13. We also would like to mention some analogies. As the chemical potential can be interpreted as an imaginary gauge field in the time direction, it is not surprising that studies of Polyakov's loop [34, 35] (Wilson loops closing in the periodic time direction) show similar fine structure. Note also that the crossing pattern of the energy levels as a function of μ found here resembles the energy crossing found in a study of the spectrum of rotating tubes as a function of the rotation rate for a proposed cold atom experiments [36].

Appendix A: Remarks about Fig. 2

In this appendix, we discuss the sign convention and the spatial winding number of Fig. 2. We also explain why this configuration is typical.

The sign of the n's associated with links with |n|=1 can be figured out from the following information. All the (vertical) temporal link indices are positive. The sign of the (horizontal) spatial links can be obtained using the conservation law. On time slices 5, 7 and 17, the current moves to the right and the sign is (by convention) positive. On time slices 21 and 22, the current moves to the left and the sign is negative. In this configuration, the winding number in the temporal direction is 2 and the winding number in the spatial direction is 0 (there are as many positive as negative spatial n's).

The fact that the configuration of Fig. 2 is typical for $\beta=0.1$ and $\mu=3$ can be understood from the numerical values of the weights. Because of the large value for μ , the

weight for the temporal links with n=0 ($I_0(0.1) \simeq 1.0025$) and n=1 ($I_1(0.1) \exp(3) \simeq 1.00553$) are almost the same, while the weight for n=2 ($I_2(0.1) \exp(6) \simeq 0.5047$) is smaller. The weight for n=-1 ($I_1(0.1) \exp(-3) \simeq$ 0.002492) is very small and there are no temporal links with negative values of n in the configuration. For the spatial links, the relative cost of a lateral move is $I_1(0.1)/I_0(0.1) \simeq 0.05$ and there are only 6 lateral moves. The fact that there are only 2 temporal links with n=2can be understood from the fact that the merging of two n=1 lines into one n=2 line requires one lateral move in addition of a weight about twice smaller.

Appendix B: Spatial winding numbers in Fig. 10

In this appendix, we discuss the spatial winding numbers of the configurations of Fig. 10. For $\mu=2.93$, all the time (vertical) links have n=1. For the spatial links there are 20 right movers and 16 left movers (so the spatial winding number is 1). For $\mu = 3.00$ (n=2), between most time slices, there are two vertical links carrying a n=1 current, and the two lines only merge 4 times into a single n=2 line for a small number of time steps (the total is 5 vertical links with n=2). There are 26 right movers and 34 left movers (so the spatial winding number is -2). For $\mu=3.07$ (n=3), between most time slices, there are three vertical links carrying a n=1 current, There are 5 occurrences where two n=1 merge into a single n=2 line for a small number of time steps (the total is 10 vertical links with n=2). There are also 3 crossing (points where the four lines attached all have |n|=1). There are 19 right movers and 15 left movers (so the spatial winding number is +1). For $\mu=3.14$ (n=4), $\beta = 0.1$, $L_x = 4$ and $L_t = 256$, there are essentially four parallel vertical lines each carrying a n=1 current except for 4 occurrences where they briefly merge $(11 \ n=2 \text{ vertical lines}, 4 \text{ right})$ movers, 4 left movers, no spatial winding number).

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