

# Quantum simulation with string-bond states: Joining PEPS and Monte Carlo

N. Schuch<sup>1</sup>, A. Sfondrini<sup>1,2</sup>, F. Mezzacapo<sup>1</sup>, J. Cerrillo<sup>1,3</sup>, M. Wolf<sup>1,4</sup>, F. Verstraete<sup>5</sup>, I. Cirac<sup>1</sup>

<sup>1</sup> Max-Planck-Institute for Quantum Optics, Garching, Germany
 <sup>2</sup> University of Padua, Italy
 <sup>3</sup> University of Barcelona, Spain
 <sup>4</sup> Niels-Bohr-Instititute, Copenhagen, Denmark
 <sup>5</sup> University of Vienna, Austria

#### The key to quantum simulations

• Aim: Simulation of quantum many-body systems



- Requirements on variational ansatz:
  - it should capture the states of interest
  - it should allow for efficient computation of energies etc.

#### **Matrix Product States**

- Matrix Product States (MPS):  $|\psi\rangle = \sum_{n_1,...,n_L} \operatorname{tr} \left[ A_{n_1}^1 A_{n_2}^2 \cdots A_{n_L}^L \right] |n_1,...,n_L \rangle$
- Alternatively:



MPS approximate well the ground states of gapped 1D Hamiltonians
 [Hastings, JSTAT '07]

#### **Computing expectation values on MPS**

- On MPS, expectation values can be computed efficiently
- Local observables *O*:



• Computation time  $\propto D^4$  ( $D^6$ ) for OBC (PBC) [can be improved to  $D^3$  ( $D^5$ )].

• Energies  $\langle \psi | H | \psi \rangle$ , corr. functions  $\langle \psi | O_i \otimes \tilde{O}_j | \psi \rangle$  can be computed efficiently.

#### Variational method with MPS: DMRG



•  $|\psi\rangle$  is linear in any  $A^{i} \equiv X : |\psi_{X}\rangle = \sum \operatorname{tr} \left[A_{n_{1}}^{1} \cdots X_{n_{k}} \cdots A_{n_{L}}^{L}\right] |n_{1}, \cdots, n_{L}\rangle$  $\Rightarrow$  The energy in state  $|\psi_{X}\rangle$  is  $E(\psi_{X}) = \frac{\langle \psi_{X} | H | \psi_{X} \rangle}{\langle \psi_{X} | \psi_{X} \rangle} = \frac{\vec{X}^{\dagger} M \vec{X}}{\vec{X}^{\dagger} N \vec{X}}$ 

 $\Rightarrow$  X which minimizes  $E(\psi_X)$  can be found efficiently.

 Density Matrix Renormalization Group (DMRG) method: [White, PRL '92] Sweep through the MPS and minimize tensors locally.



- DMRG algorithm performs extremely well! Accuracy is tuned by increasing D (computation time  $\propto D^3$ ).
- Idea extends to periodic boundary conditions (scales as  $D^5$ ).

#### **Extending DMRG to two dimensions**



• The MPS ansatz naturally generalizes to two dimensions.



 Tensor Product States (TPS), Projected Entangled Pair States (PEPS) [Nishino 90ies]

[Nishino 90ies] [Verstraete & Cirac, PRA '04]

- PEPS form a complete family.
- PEPS are well suited for ground and thermal states in higher dimensions.

[Hastings, PRB '06]

## Extending DMRG to two dimensions

- The MPS ansatz naturally generalizes to two dimensions.
  - $D^2$ 
    - **Problem:**

Computing expectation values on 2D PEPS seems to be hard: The size of the tensor grows exponentially in the perimeter!

[Schuch, Wolf, Verstraete & Cirac, PRL '07]

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## **PEPS algorithm with truncation**



Solution: proceed column-wise and truncate the bond dimension



- Corresponds to a DMRG-like optimization
- Works well in practice (due to symmetries of the system?)
- Known error

[Verstraete & Cirac, cond-mat/0407066] [Murg, Verstraete & Cirac, PRA '07] [Murg, Verstraete & Cirac '09]

- Works very well & outperforms other methods e.g. for frustrated systems
- Computation time scales as  $D^{12}(D^{18} \text{ for PBC})$  can be reduced to  $D^{10}(D^{16})$
- Up to D=5 (lattice up to  $20 \times 20$ ).
- Not practical for PBC or higher dim. lattices, or irregular geometries.

#### The quest for other methods

MPQ

- DMRG works well in 1D.
- PEPS extend DMRG to 2D and give very good results.
- PEPS contain the relevant states (ground and thermal states)
- However: scaling of PEPS in *D* is unfavorable, in particular for PBC
  - truncation relies on lattice structure not applicable to irregular systems

Is it possible to find a smaller/different class of states

- for which expectation values can be computed efficiently
- which has a favorable scaling in the accuracy parameter
- which does not rely on the geometry of the system
- which well approximates the states of interest

#### **Variational Monte Carlo**

• Given a state  $|\psi
angle$  on *L* spins, how can we sample exp. values?

$$\langle \psi | H | \psi \rangle = \sum_{n} \langle \psi | n \rangle \langle n | H | \psi \rangle = \sum_{n} \underbrace{|\langle \psi | n \rangle|^{2}}_{\propto p(n)} \frac{\langle n | H | \psi \rangle}{\langle n | \psi \rangle}$$

 $n = (n_{1,}..., n_L)$  basis state in some local basis e.g.  $0 = \uparrow$ ,  $1 = \downarrow$ 

 $\Rightarrow \langle n | \psi \rangle$  and  $\langle n | H | \psi \rangle$  need to be efficiently computable!

• 
$$H = \sum h_i^{-1}$$
 local terms

$$\Rightarrow \langle \boldsymbol{n} | \boldsymbol{h}_{i} = \sum_{\text{few } \boldsymbol{n}'} \boldsymbol{\phi}(\boldsymbol{n}') \langle \boldsymbol{n}' | \qquad \Rightarrow \langle \boldsymbol{n} | \boldsymbol{H} | \boldsymbol{\psi} \rangle = \sum_{\text{few more } \boldsymbol{n}'} \boldsymbol{\phi}(\boldsymbol{n}') \langle \boldsymbol{n}' | \boldsymbol{\psi} \rangle$$

 $\langle n | \psi \rangle$  needs to be efficiently computable!

... the same works for products of Paulis, e.g.  $\sigma^+ \sigma^z \cdots \sigma^z \sigma^-$ 



#### **Classes for which this holds**

- $\langle \pmb{n} | \pmb{\psi} 
  angle$  needs to be efficiently computable!
- Examples of such states:
  - Matrix Product States:  $\langle \boldsymbol{n} | \psi \rangle = \operatorname{tr} \left[ A_{n_1}^1 \cdots A_{n_L}^L \right]$

(note: computation of expectation values scales as  $D^2$  [ $D^3$ ] instead of  $D^3$  [ $D^5$ ] – could be used to speed up DMRG or PEPS methods; see also Sandvik & Vidal, PRL '07)

- coherent version of classical thermal states:

$$|\psi
angle = \sum e^{-eta H(m{n})/2} |m{n}
angle$$
 for classical  $H$ 

$$\Rightarrow \langle \boldsymbol{n} | \boldsymbol{\psi} \rangle = e^{-\beta H(\boldsymbol{n})/2}$$

Can we find new classes of suitable states (by extending these classes)?





#### Generalization of "classical" states

What is the PEPS structure of "classical" states?

$$\underbrace{\stackrel{n_i}{\not}}_{\not} = - \underbrace{\sigma_{\not}}_{\not} \underbrace{\sigma_{-}}_{\not} \Rightarrow \langle n | \psi \rangle =$$

⇒  $\langle \boldsymbol{n} | \psi \rangle$  is product of **0-dimensional** objects! ⇒ It can be computed efficiently!

Generalize to 1D objects!



 $\Rightarrow \langle \mathbf{n} | \psi \rangle \text{ product of } \mathbf{1D} \text{ objects} - \\ \text{can still be efficiently computed!}$ 



- Generalization of "classical" states (or of Matrix Product States!)
- subclass of PEPS with eff. Monte Carlo sampling



 $\exp\left[-\beta h_{ii}(n_i, n_i)/2\right]$ 

#### String-bond states (SBS)



#### <u>"String-bond states" (SBS)</u>

 $\langle \boldsymbol{n} | \boldsymbol{\psi} \rangle = \prod f_i(\boldsymbol{n})$ 

• defined on string (subset) of spins  $(n_{i_1}, ..., n_{i_k})$ • efficiently computabe (e.g. matrix product trace)

# $\langle n | \psi \rangle$ is a product of efficiently computable functions defined on subsets of spins

- matrix product traces
- tree tensor networks
- on *small* subsets:
  - arbitrary state
  - mini-PEPS

- lines (horiz./vert./diagonal/...)
- 2x2 loops
- plaquettes (blocks) around each site
- ... and any combination thereof!

#### An example: Lines





#### **String-Bond States – Examples**

MPQ

 String-bond states form a complete family as they encompass MPS



- Any (generalized, weighted) graph state is a SBS:
  - the cluster state
  - any stabilizer state



 The toric code state and the quantum double models are SBS:
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for even parity for odd parity

#### **String-bond states: Properties**

- SBS form a hierarchy of states which can be enlarged by
  - increasing *D* for the matrix products
  - adding new strings
- SBS include various common classes of states
- Computational resources scale favorably:  $D^2$  (OBC) or  $D^3$  (PBC) [cf. PEPS:  $D^{12}$  (2D OBC),  $D^{18}$  (2D PBC)]
- SBS can thus be used for PBC or 3D systems, and much higher *D*'s can be used
- no underlying geometry/locality necessary: also suitable for simulation of molecules, systems with non-local interactions, etc.
- each SBS is a PEPS, but PEPS-D exponential in #strings/site



#### Variational algorithm



• Idea: Determine M and N by MC sampling:

$$E(\psi_{X}) = \frac{\sum \langle \psi_{X} | n \rangle \langle n | H | \psi_{X} \rangle}{\sum \langle \psi_{X} | n' \rangle \langle n' | \psi_{X} \rangle} = \frac{\sum p_{0}(n) \vec{X}^{\dagger} \cdot \vec{b}_{n} \vec{a}_{n}^{\dagger} \cdot \vec{X}}{\sum p_{0}(n') \vec{X}^{\dagger} \cdot \vec{b}_{n'} \cdot \vec{b}_{n'}^{\dagger} \cdot \vec{X}} \qquad \vec{b}_{n}^{\dagger} \cdot \vec{X} := \frac{\langle n | \psi_{X} \rangle}{\langle n | \psi_{X_{0}} \rangle}$$
$$\vec{a}_{n}^{\dagger} \cdot \vec{X} := \frac{\langle n | H | \psi_{X} \rangle}{\langle n | \psi_{X_{0}} \rangle}$$
$$\vec{a}_{n}^{\dagger} \cdot \vec{X} := \frac{\langle n | H | \psi_{X} \rangle}{\langle n | \psi_{X_{0}} \rangle}$$
$$\vec{a}_{n}^{\dagger} \cdot \vec{X} := \frac{\langle n | H | \psi_{X} \rangle}{\langle n | \psi_{X_{0}} \rangle}$$
$$p_{0}(n) := |\langle n | \psi_{X_{0}} \rangle|^{2}$$

 $\rightarrow$  Reweighting: one distribution samples all

#### However ...



$$M = \sum p_0(\boldsymbol{n}) \vec{b}_n \vec{a}_n^{\dagger}$$

$$N = \sum p_0(\boldsymbol{n}) \vec{b}_n \vec{b}_n^{\dagger}$$
find  $\vec{X}$  which minimizes  $E(\psi_X) = \frac{\vec{X}^{\dagger} M \vec{X}}{\vec{X}^{\dagger} N \vec{X}}$ 

However, there are some **problems** :

- *M* and *N* are **very large**:  $(dD^2)^2$  sampling rel. slow
- *M* and *N* less accurate far from  $\vec{X}_0$  due to reweighting
- sampling error  $\propto \sqrt{MC}$  sample length

 $\rightarrow$  small errors in kernel of N can have **big effect** 

 $\Rightarrow$  method **unstable!** 

#### **Gradient flow**

• Solution: Move along gradient (in small steps), from initial value  $X = X_0$ :

$$\nabla_{X} E(\psi_{X}) \Big|_{X=X_{0}} = 2 [M - E(\psi_{X_{0}})N] \vec{X}_{0}$$

- $\rightarrow$  well-behaved in N and M
- Gradient can be **sampled directly**:

$$\nabla_{X} E(\psi_{X}) \Big|_{X=X_{0}} = 2 \sum p_{0}(\mathbf{n}) \vec{b}_{\mathbf{n}} [E_{n} - E(\psi_{X_{0}})]$$

• Gradients indep. to first order: All tensors can be updated simultaneously (also allows for more eff. sampling)

$$E(\psi_{X}) = \frac{\vec{X}^{\dagger} M \vec{X}}{\vec{X}^{\dagger} N \vec{X}}$$
$$\vec{X}_{0}^{\dagger} N \vec{X}_{0} = 1$$

$$E_{n} = \frac{\left\langle n \left| H \right| \psi_{X_{0}} \right\rangle}{\left\langle n \right| \psi_{X_{0}} \right\rangle}$$
$$\vec{b}_{n}^{\dagger} \cdot \vec{X} := \frac{\left\langle n \left| \psi_{X_{0}} \right\rangle}{\left\langle n \right| \psi_{X_{0}} \right\rangle}$$

$$E_{n} = \frac{\langle n | H | \psi_{X_{0}} \rangle}{\langle n | \psi_{X_{0}} \rangle}$$
$$\vec{b}^{\dagger} \cdot \vec{X} \cdot = \frac{\langle n | \psi_{X} \rangle}{\langle n | \psi_{X} \rangle}$$

#### The full algorithm

- Fix a string pattern & bond dimensions *D*.
- Choose initial configuration for the strings (random, guess, ...)
- Repeat: 1) Monte Carlo sample gradients (& Energy)
   2) Update tensors according to gradient and stepwidth *ε* (to this end: normalize gradients)

3) Stop if energy has converged, otherwise go to 1)

- Increase precision: either
  - decrease stepwidth  $\epsilon$
  - increase D
  - add new strings

and restart algorithm from obtained optimum, until energy does not improve any more



#### **Testing the algorithm**

• How can we benchmark SBS?

#### **Quantum Monte Carlo**

- + PBC, 3D possible
- no frustrated systems

#### PEPS

- + can do frustr. systems
- only 2D OBC

#### • Note: Aim – simulating systems neither QMC nor PEPS can do!



#### **Implemented patterns**



#### **Comparison to QMC: transverse Ising model**





QMC vs. lines and lines+loops

#### **Comparison to PEPS: frustrated systems**

- first comparisons: frustrated XX model SBS can outperform PEPS (D = 12 vs. D = 4).
- simulation of 2D frustrated PBC XX model w. transv. field:



• extensive comparison (and extension to PBC):  $J_1 - J_2$  model





#### **Projected String Bond States**

- Can we incorporate symmetries of *H* in the ansatz?
- SU(2) invariant Hamiltonian:

There exists ground state  $|\psi_0
angle$  with  $\hat{M}|\psi_0
angle$ =0 (where  $\hat{M}$ = $\sum_i \sigma_i^z$ )

- For any candidate  $|\Psi_0\rangle$ ,  $|\bar{\Psi}_0\rangle = \Pi_0 |\Psi_0\rangle$  is at least as good!
- Implementation:

(this is a SBS –  $\langle \pmb{n} | \pmb{\psi} \rangle$  can be computed efficiently)

 $\rightarrow$  enforce by sampling only from the M = 0 subspace.

• In practice: start from configuration with M=0and swap a random pair of spins



#### **Molecules**



• Simulation of molecules:



• Problem: no natural ordering (geometry) of modes

## Molecules

• Ansatz: put strings between all pairs of modes

$$|\psi\rangle = \prod_{\alpha>\beta} f_{\alpha,\beta}(n_{\alpha}, n_{\beta}) \underbrace{(c_{L}^{\dagger})^{n_{L}} \cdots (c_{1}^{\dagger})^{n_{1}} |\operatorname{vac}\rangle}_{\equiv |n_{1}, \dots, n_{L}\rangle}$$



- And: sample only in subspace with  $N_e$  fermions
- Advantages:
  - all modes treated on equal footing
  - ansatz invariant under relabeling of modes
- gradient flow optimization of the  $f_{\alpha\beta}$
- possible to include canonical transformations  $\tilde{c}_i = \sum O_{ij} c_j$ in the gradient optimization
- extension: use  $f_{\alpha\beta\gamma\delta}(n_{\alpha}, n_{\beta}, n_{\gamma}, n_{\delta})$  on quadruples of modes
- results soon!



# Speeding up DMRG/PEPS with Monte Carlo

- The computational cost is  $D^2 / D^3$  times the number of strings (each string can be evaluated by sequence of matrix multiplications)
- This is better than the scaling of DMRG (=a special SBS)  $D^3$  /  $D^5$  .
- <u>Reason</u>: We don't have to contract the "sandwich"  $\langle \psi | O | \psi \rangle$ but only  $\langle n | \psi \rangle$  and  $\langle n | O | \psi \rangle$ .



• Monte Carlo sampling can thus be used to speed up DMRG and, more importantly, the PEPS algorithm ( $D^6$  instead of  $D^{12}$ ).

[see also: Sandvik and Vidal, PRL '07]

# Summary

- Monte Carlo sampling can help in tensor contraction
- String-bond states:  $\langle \pmb{n} | \pmb{\psi} \rangle$  prod. of eff. computable "local" objects
- String pattern can reflect geometry/ent. structure of system
- Enlarge family by larger D or more strings
- Cost of contraction:  $D^3 \times \#$ strings
- SBS allow for efficient simulation of frustrated systems also with 2D PBC and in 3D
- Monte Carlo can speed up any tensor network based method
- <u>Current & future direction:</u>
  - better understand & characterize class of SBS
  - simulate molecules
  - string patterns using plaquettes
  - (imag.) time evolution
  - smarter sampling

[N.Schuch, M. Wolf, F. Verstraete, I. Cirac, PRL '08] [A. Sfondrini *et al.*, in preparation]

