# Manuscript Title\*

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In this paper, a numerical "decomposition method" is introduced to extract information in Green Function (or called Density Matrix) and construct a wave function in pseudo-BCS form. This method can be used in producing constraints for Quantum Monte Carlo (QMC) to restrict the random walk. This method is applied to Constrain Path Monte Carlo (CPMC) which need trial wave function constraints to control the sign problem in strongly correlated fermion systems. so that the new S.C.CPMC can get systematically improved constraints from the Green Function of last iteration. The behavior of this new S.C. CPMC is demonstrated in 2-D Hubbard Model. Some model become possible to be calculated by this new S.C. CPMC which are otherwise beyond reach in normal CPMC framework.

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#### I. INTRODUCTION

The hardest part to understand a quantum system, especially those strongly correlated systems, is how to get enough useful information from its exponential explosion Hilbert space. One of the key to solve this problem is Quanrum Monte Carlo (QMC)[?], which it's possible for classical computer to chase exponential scale and get the estimation of some physical quantities in principle.

However, phase problems in Fermion system prevent the convergence of QMC. The most effective approach to deal with phase problem is adding bias constraint which has been applied to many other fields. For example, Constrain Path Monte Carlo (CPMC)[?] can provide exact results without phase problem if its trial wave function "phiT" (constraint) is exact ground state.

But choosing a suitable constraint is hard and sometimes it leads to some fatal man-made bias. So an trial wave function source is needed. But in some situation, those "wave function results" methods (like Hartree-Fock (HF) or density-functional theory (DFT)) can't provide a good enough approximation of ground state and most importantly, they may lead CPMC run into an wrong way. And those "quantities results" methods (like AFQMC or DMRG) which can give a better approximation, are not able to provide a usable wave function without drastically changing its computational scaling or complexity.

In this paper, we are going to introduce an almost 'non-cost' algorithm which can be used to extract full information from output Green function (or called Density Matrix) to construct a wave function and input this wave function as trial wave function to CPMC calculation or in another view point, this new method now allow CPMC to use Green Function as a constrain rather than trial wave function. So that, CPMC, as an input-output algorithm, can be easily applied to self-consistent frame which can improve their results to local or even global optimized results with heat bath and luck by iteration.

For concreteness, we will use the Hubbard model to describe the self-consistent CPMC procedure:

$$H = \sum_{j,\delta,\sigma} -t_{j,j+\delta} c_{j,\sigma}^{\dagger} c_{j+\delta,\sigma} + U \sum_{j} n_{j,up} n_{j,dn}$$

where  $c_{j,\sigma}^{\dagger}$  ( $c_{j,\sigma}^{\dagger}$ ) creates (annihilates) an electron with spin  $\sigma$  ( $\sigma = up, dn$ ) at lattice site j and  $\delta$  connects two

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nearest neighbor sites or other sites which depends on detailed system,  $n_{j,\sigma} \equiv c_{j,\sigma}^{\dagger} c_{j,\sigma}$ .

### II. SELF CONSISTENT CPMC METHOD

To start the self-consistent procedure, we run some numerical calculation first (assume we do CPMC with free electrons trial wave function). Then with back propagation, we can get the Green Function G of Ground State.

Now, we want to construct a wave function which has the same Green Function with Ground State. If such a wave function can be found, this wave function must have almost the same behavior with this Ground State, so it can be a good approximation of Ground State and a good constraint in CPMC.

#### A. Theoretical derivation

Instead of using DET state (Slater Determinate) in normal CPMC, we choose pesudo-BCS state as trial wave function

BCS[?] state pairing wave function with  $N_{spin}$  paired particles can be written as

$$BCS(F) = \underbrace{\psi^{\dagger}...\psi^{\dagger}}_{N_{crip}} |0>$$

where

$$\psi^{\dagger} \equiv \sum_{i,j} f_{i,j} c_{i,up}^{\dagger} c_{j,dn}^{\dagger}$$

and  $N_{spin_{up}} = N_{spin_{dn}} \equiv N_{spin}$  is the number of pairs, |0> is empty state.

In principle,  $F = \{f_{i,j}\}$ , a Nsite by Nsite matrix (Nsite is the number of lattice sites), has some constraints in the definition of BCS state. For example  $f_{i,j} = f_{j,i}$ . Now, let's get rid of these constraints (F can be any matrix) and define it as pseudo-BCS state pBCS(F).

Then we are going to prove we can find a pseudo-BCS state which has any Green Function we want.

Theory: For any Green Function

$$G_{i,j,\delta} = \frac{<\psi|C_{i,\delta}^{\dagger}C_{j,\delta}|\psi>}{<\psi|\psi>}$$

with  $N_{spin_{up}} = N_{spin_{dn}}$  and 'part of spin symmetry', there exist a pseudo-BCS state pBCS(F) which has Green function G' such that G' can approach or equal to G

Proof: For any Green Function G, by eigenvalue decomposition, we can get

$$G_{i,j,\delta} = \frac{\langle \psi | C_{i,\delta}^{\dagger} C_{j,\delta} | \psi \rangle}{\langle \psi | \psi \rangle} = \{ T_{\delta} \lambda_{\delta} T_{\delta}^{\dagger} \}_{i,j}$$

where  $\lambda_{up} = \lambda_{dn} \equiv \lambda$  (This additional requirement called "part of spin symmetry") is diagonalized eigenvalue matrix and T is eigenvectors.

Then we claim: for any diagonalized Green function, there is a diagonslized matrix F, such that the Green Function G' of this pseudo-BCS state pBCS(F) is G or at least the error can be minimized.

For diagonalized matrix F,

$$pBCS(F) = \prod (\sum_{i,j} f_{i,j} C_{i,up}^{\dagger} C_{j,dn}^{\dagger}) =$$

$$\sum_{\{k\}}(\lambda_{k_1}...\lambda_{k_{Nspin}}C^{\dagger}_{k_1,up}C^{\dagger}_{k_1,dn}...C^{\dagger}_{k_{Nspin},up}C^{\dagger}_{k_{Nspin},dn})|0>$$

where  $\{k\}$  is a set of non-repetitive sample of 1, 2, ... N site and  $\lambda_{k_*} \equiv f_{k_*, k_*}$ .

it's easy to see that  $C_{k_1,up}^{\dagger}C_{k_1,dn}^{\dagger}...C_{k_{Nspin},up}^{\dagger}C_{k_{Nspin},dn}^{\dagger}$  is a DET state with DET matrix  $D_{i,j}=1$  for every  $i=k_j$  and  $D_{i,j}=0$  for others.

Now, the Green Function G' of pBCS(F) is:

$$G'_{ij,\delta} \equiv \frac{< pBCS | C^{\dagger}_{i,\delta}C_{j,\delta}| pBCS >}{< pBCS | pBCS >}$$

If  $i \neq j$ , then  $< pBCS|C_{i,\delta}^{\dagger}C_{j,\delta}|pBCS> = 0$ , since this new paired state

$$C_{i,\delta}^{\dagger}C_{j,\delta}(C_{k_1,up}^{\dagger}C_{k_1,dn}^{\dagger}...C_{k_{Nspin},up}^{\dagger}C_{k_{Nspin},dn}^{\dagger})|0>$$

can't be found in < pBCS|. If i = j, then

$$<\!\!pBCS|C_{i,\delta}^{\dagger}C_{i,\delta}(C_{k_1,up}^{\dagger}C_{k_1,dn}^{\dagger}...C_{k_{Nspin},up}^{\dagger}C_{k_{Nspin},dn}^{\dagger})|0>\neq 0$$

if and only if there exist some  $k_j = i$ .

$$G_{ii,\delta} = \frac{N_{spin} \sum_{\{k\}, \exists k_j = i} (\lambda_{k_1} ... \lambda_{k_{Nspin}})^2}{\sum_{\{k\}} (\lambda_{k_1} ... \lambda_{k_{Nspin}})^2}$$

For diagonal matrix G and G',  $G_{i,i} = G'_{i,i}$  gives us a "Nsite variable equation set". This equation set may have analytic solution with low computation complexity, but is hard to find out. Two 'almost non-cost' approximate solutions are given which let G' approach to G good enough at most of the cases .

By applying linear transformation T to basis  $C^{\dagger}$  and C, this results can be generalized exactly to any Green Function G with  $Nspin_{up} = Nspin_{dn}$  and 'part of spin symmetry'.

# B. Concrete Steps

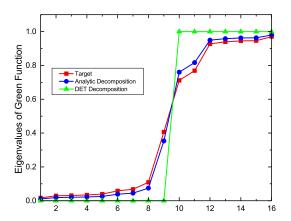
For any Green Function G we can follow steps below to construct a pseudo-BCS state which has nearly the same Green Function G' with G.

- 1. Get the target Green Function G and its eigenvectors  $T_{up}$ ,  $T_{dn}$ , eigenvalue matrix  $\lambda$  (in the system with "part of spin symmetry",  $\lambda_{up} = \lambda_{dn} \equiv \lambda$ ).
- 2. The new pseudo-BCS trial wave function F is  $F = T_{up}^{\dagger} A T_{dn}$  where  $A \equiv f(\lambda)$  is diagonalized matrix.
  - 3. There are two kind of f:

1st, DET decomposition:  $A_{ii} = 1$  if  $\lambda_{ii}$  is larger or equal to the Nspin-th largest value in  $\lambda$  or  $A_{ii} = 0$  (when this decomposition is applied, the pseudo-BCS state reduced to DET state and this is the S.C. CPMC mentioned by Mingpu)

2nd, Analytic decomposition:  $A_{ii} = \sqrt{\lambda_{ii}/(1 - \lambda_{ii})}$ . This is the exact decomposition for pseudo-HFB state (with the same definition as pseudo-BCS state), which means in some grand canonical ensemble system, this decomposition is the exact solution for G = G'.

FIG. 1. The comparison of two different decomposition methods in trying to fit a Green Function. (From the steps above, they all have the same eigenvectors, so we show their difference in eigenvalues.) This is a real situation for 4 by 4 nearest-hopping Hubbard system. The eigenvalues of green function vs. the order sorted by corresponding eigenvalues.



The decomposition method is the key of this selfconsistent CPMC procedure, if we choose "DET decomposition", the pesudo-BCS reduce to DET state and it is easy to see why DET state is not good to be used as trial wave function. Since there is no analytic methods or efficient numerical methods to get an exact decomposition, "Analytic Decomposition" (though is not analytic) is the best decomposition we have now. But it definitely can be improved when this method applied to some detailed system. In FIG.1, we give a briefly comparison for these two decomposition method to give a brief idea about what those decomposition methods are going to do and how it works to improve the approximation. And In TABLE.III, we shows this improved approximation indeed improve the results of CPMC with Exact Green Function input.

Another thing we want to mention is that due to the

proof above, this new psudo-BCS S.C. CPMC only work in  $Nspin_{up} = Nspin_{dn}$  and 'part of spin symmetry' system.

#### C. Technical detail

If pesudo-BCS state is applied as trial wave function, there are some changes from "DET trial wave function CPMC" to "pseudo-BCS trial wave function CPMC":

$$\langle pBCS | \phi \rangle = det(A) \equiv det[\phi_{up}^T \cdot F \cdot \phi_{dn}^T]$$

$$\frac{< pBCS|c_{(i,up)}c_{(j,up)}^{\dagger}|\phi>}{< pBCS|\phi>} = \delta_{i,j} - [F \cdot \phi_{dn} \cdot A^{-1} \cdot \phi_{up}^T]_{j,i}$$

and Back propagation,  $\langle pBCS|n_in_j|\phi \rangle$  is mentioned in Ettore's recent work[?]. ( $\phi$  is the DET state, F is the matrix in pBCS(F))

#### III. RESULTS

There are some results to show the behavior of pesudo-BCS S.C. CPMC with Analytic Decomposition.

First we use the one-band two-dimensional Hubbard model at density n=0.875 as a general test case. The hopping matrix element  $\delta$  is t for nearest neighbors and 0 otherwise. These parameters regime suffer a severe fermion sign problems and its ground state in the thermodynamic limit still remains unknown.

TABLE.I, we shows our results in different system size, U, boundary condition and initial trial wave function where Exact Diagonalization (ED) for 4\*4 system and DMRG for 4\*8, 4\*12 system can be done to produce a effective comparison. The 4\*4 systems are in Twist B.C. and 4\*8, 4\*12 are in the B.C. called "closeopen" which is the same B.C. used in Mingpu recent work. Closeopen B.C. is a cylindrical geometry, i.e., with Periodic B.C. in the x-direction and Open B.C. in the y-direction, where DMRG can give very accurate benchmark results. Pinning fields with fields strengths t/4 are also applied at left and right side (the open edges) in this Closeopen B.C.. For most of the situation, especially in small U system, these iterations give a significant improvement with no need to worry about trial wave function and some high accuracy has never been achieved before.

In FIG.2, we zoom in a typical iteration procession which shows the behaviors of this two decomposition methods. In the case, Analytic Decomposition converge faster than DET Decomposition.

In FIG.3, we give a comparison to Mingpu's work[?] in the 4 by 16 system with Closeopen B.C. [?] where DMRG are supposed to give an exact results about this system's magnetic order. DET decomposition and Mingpu's methods converge to each other but still can't

L	$(N_{up},N_{dn})$	U/t	B.C.	initial input	CPMC	S.C. CPMC	STDEV	ED/DMRG	Error %
4*4	(7,7)	4	(0.01, 0.02)	FS	-15.8491	-15.7344	0.0093	-15.7660	-0.2002
4*4	(7,7)	4	(0,0)	FS	-15.5905	-15.8022	0.0057	-15.744	0.3702
4*4	(7,7)	8	(0.01, 0.02)	FS	-12.2185	-12.0588	0.0390	-11.8752	1.5456
4*4	(7,7)	12	(0.01, 0.02)	FS	-10.5287	-9.9309	0.0365	-10.0543	-1.2272
4*8	(14,14)	4	closeopen	FS	-31.7378	-31.8412	0.0081	-31.8609	-0.0619
4*8	(14,14)	6	closeopen	FS	-26.8088	-26.8501	0.0196	-26.8243	0.0963
4*8	(14,14)	8	closeopen	FS	-23.7089	-23.4801	0.0485	-23.5659	-0.3640
4*12	(21,21)	4	closeopen	FS	-48.3804	-48.4456	0.0115	-48.4397	0.0123
4*12	(21,21)	6	closeopen	FS	-40.877	-40.9293	0.0278	-40.779	0.3686
4*12	(21,21)	8	closeopen	FS	-36.119	-35.7647	0.0574	-35.8133	-0.1356

TABLE I. The comparison of ground state energy between "Exact" (For 4\*4 system, results from ED. For 4\*8, 4\*12 system, results from DMRG), normal CPMC (first step results of S.C. CPMC which is the normal CPMC start with initial input trial wave function), pesudo-BCS S.C. CPMC with Analytic Decomposition in different system size, U, twist Boundary Condition 2D nearest hopping one band Hubbard model.

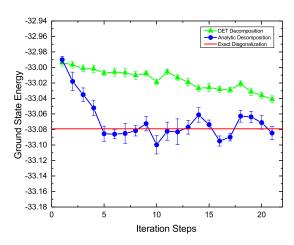


FIG. 2. 4\*8, U=4,  $Nspin_{up}=Nspin_{dn}=14$ , P.B.C., nearest-hopping Hubbard Model. The comparison between two detail iteration curves from pesudo-BCS S.C. CPMC with two different decomposition methods. Ground state energy vs. the number of iteration steps.

gives a good approximation for the saddle part of charge density. Analytic Decomposition improve the results and match this saddle point well.

In FIG.4 we apply our methods to  $t^\prime$  Hubbard model[?] (t for nearset-hopping and  $t^\prime$  for second-nearest hopping) which can't be touched by CPMC before (can't even give a description of what state it is). The consensus about its phase transition behavior in the thermodynamic limit still going on. DMRG is supposed to give a good approximation in small one-Dimension-like system, so results of DMRG is used as Exact results, and we compare it with Analytic Decomposition, DET decomposition to shows it is now possible for us to use S.C. CPMC in dealing with this problem. The following formal research about this  $t^\prime$  problem with S.C. CPMC will come out later. The ground state energy comparison is showed in TABLE.II.

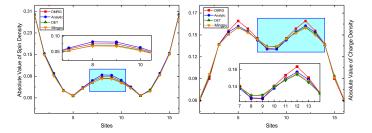


FIG. 3. 4\*16, U=4,  $Nspin_{up}=Nspin_{dn}=28$ , Pinning Field B.C., nearest-hopping Hubbard Model. The comparison between pesudo-BCS S.C. CPMC and Mingpu's results. Absolute value of spin density  $(< n_{i,up} - n_{i,dn} > /2)$  along the y-direction vs. site label. The right panel plots the corresponding charge density  $(1-< n_{i,up} + n_{i,dn} >)$ .

L	t'	CPMC	S.C. CPMC	Exact
4*8	0.3	-27.360(9)	-27.67(1)	-27.6924
4*8	0.35	-27.839(9)	-27.95(1)	-27.9755
4*8	1.15	-41.586(9)	-41.68(1)	-41.709
4*8	1.2	-43.130(8)	-43.16(1)	-43.156
4*8	1.5	-53.029(7)	-53.02(1)	-53.065
4*16	0.3	-56.04(3)	-56.25(4)	-56.236
4*16	0.4	-57.78(2)	-57.98(2)	-57.930
4*16	1.1	-83.66(2)	-83.67(1)	-83.8
4*16	1.2	-89.77(2)	-89.83(3)	-89.9
4*16	1.5	-109.60(1)	-109.87(1)	-109.98

TABLE II. 4\*8 and 4\*16, U=4,  $Nspin_{up}=Nspin_{dn}$  half-filling, Pinning Field B.C., t' Hubbard Model. The comparison of ground state energy between "Exact" (For 4\*8, 4\*12 system, results from DMRG), normal CPMC (first step results of S.C. CPMC) and pesudo-BCS S.C. CPMC with Analytic Decomposition.

# IV. DICUSSION

When this method be applied into large U system, the most important reason why it won't work well like small u system is because CPMC can't give an exact enough estimation of Green Function in large U system which in

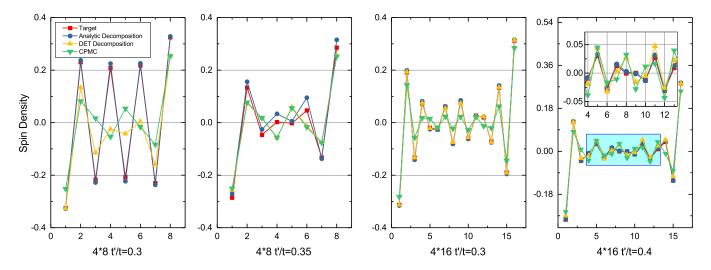


FIG. 4. 4\*8 and 4\*16, U = 4,  $Nspin_{up} = Nspin_{dn}$  half-filling, Pinning Field B.C., t' Hubbard Model. Spin density treated by pesudo-BCS S.C. CPMC with different decomposition method and DMRG. Spin density along the y-direction vs. site label.

most of this situation, some of the eigenvalues may be larger than 1. In some of this large U system, DET decomposition work better than Analytical decomposition

And since it is not a exact decomposition, in TA-BLE.II, an exact input Green Function won't give an exact result. So error always exist for this iteration.

Fortunately, there are many ways to improve this S.C. CPMC if it is applied in some detailed system, for example: releasing the constrain for a while, modifying Green Function due to certain symmetry or other information and finding a better decomposition method.

# V. SUMMERY

In summary, we have developed a new "Green Function to wave function" method which allowed many "wave

function input-output algorithm" to be applied to a self-consistent frame. By doing this to CPMC, many results has been improved and this new S.C. CPMC allowed us to touch farther field which we can't do before.

The decomposition method is not optimal, which means many other optimal method like Machine Learning may improve it and lead to better results. So this paper also gives a optimal problem, solving this problem will give a meaningful improvement to many numerical algorithm.

# A. Citations and References

A citation in text uses the command \cite{#1} or \onlinecite{#1} and refers to an entry in the bibliography. An entry in the bibliography is a reference to another document.

L	$(N_{up},N_{dn})$	U/t	Twist B.C.	Decomposition Method	First Step Energy	S.C. CPMC	Exact Energy
4*4	(7,7)	4	(0.01, 0.02)	Analytic	-15.79(1)	-15.706(9)	-15.766
4*4	(7,7)	4	(0.01, 0.02)	DET	-15.68(1)	-15.679(5)	-15.766
4*4	(7,7)	4	(0,0)	Analytic	-15.76(2)	-15.806(5)	-15.744
4*4	(7,7)	4	(0,0)	DET	-15.65(1)	-15.648(4)	-15.744
4*4	(7,7)	8	(0.01, 0.02)	Analytic	-11.87(7)	-12.05(4)	-11.875
4*4	(7,7)	8	(0.01, 0.02)	DET	-11.75(3)	-11.97(1)	-11.875
4*4	(7,7)	12	(0.01, 0.02)	Analytic	-10.0(1)	-9.85(3)	-10.054

TABLE III. "First Step Energy" are the results from the first step CPMC calculation in S.C. CPMC with pBCS state as the trial wave function which decomposed from Exact Ground State Green Function by different "Decomposition Method" and "S.C. CPMC" is the converged results in this S.C. calculation. A small modification at Exact Green Function is needed to break the degeneracy.