# Hybridization expansion and non-crossing/one-crossing approximations

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# I. INTRODUCTION

This is an introduction to hybridization expansion and its approximation schemes, especially for non-crossing and one-crossing approximations. Detailed derivation are given in this note. In addition, several simple rules to determine tricky signs in any configurations and in any self-energy expressions are presented. Notations in this note follow from Ref. [1] and Ref. [2]. Details of hybridization expansion and approximation schemes are given in Sec. II and Sec. III, respectively. But before moving on, it would be beneficial to first get familiar with the notations and the general form of the Hamiltonian.

For a general quantum impurity model, the Hamiltonian  $H_{\text{QI}}$  consists of three terms:  $H_{\text{loc}}$ , which describes the "impurity",  $H_{\text{bath}}$ , which describes the "bath", and  $H_{\text{hyb}}$  which describes the coupling between the impurity and the bath. Thus,

$$H_{\rm QI} = H_{\rm loc} + H_{\rm bath} + H_{\rm hyb} . \tag{1}$$

Usually,  $H_{\rm loc}$  consists of two terms: hopping term  $H^0_{\rm loc}$  and interacting term  $H^1_{\rm loc}$ , both of which can be expressed in terms of fermionic operators  $d/d^{\dagger}$ . Explicit expressions for these terms are as follows:

$$H_{\rm loc} = H_{\rm loc}^0 + H_{\rm loc}^1 , \quad H_{\rm loc}^0 = \sum_{ab} E^{ab} d_a^{\dagger} d_b , \quad H_{\rm loc}^1 = \sum_{ijkl} I^{ijkl} d_i^{\dagger} d_j^{\dagger} d_k d_l + \cdots , \qquad (2)$$

where states are labeled by flavor indices a (including both orbital and spin degrees of freedom). As for  $H_{\text{bath}}$ , the general form is

$$H_{\text{bath}} = \sum_{p} \varepsilon_{p} c_{p}^{\dagger} c_{p} , \qquad (3)$$

in which flavor p is the combination of momentum, orbital and spin indices. Finally, the coupling between the impurity and the bath can be written as

$$H_{\rm hyb} = \sum_{pj} \left( V_p^j c_p^\dagger d_j + V_p^{j*} d_j^\dagger c_p \right) \,. \tag{4}$$

## **II. HYBRIDIZATION EXPANSION**

Hybridization expansion is a perturbative expansion regrading to  $H_{hyb}$ . Denote  $H_{loc} + H_{bath}$  and  $H_{hyb}$  by  $H_a$  and  $H_b$ , respectively. Then the partition function of the impurity model can be expressed as

$$Z = \operatorname{Tr}[e^{-\beta(H_a + H_b)}] = \operatorname{Tr}[e^{-\beta H_a}T_{\tau} \exp(-\int_0^\beta d\tau H_b(\tau))], \qquad (5)$$

where  $H_b(\tau) = e^{H_a \tau} H_b e^{-H_a \tau}$  and  $T_{\tau}$  is the time-ordering operator. After applying the Taylor expansion to the exponential, the partition function can be reexpressed as

$$Z = \sum_{k=0}^{+\infty} (-1)^k \frac{1}{k!} \int_0^\beta d\tau_1 \cdots \int_0^\beta d\tau_k \operatorname{Tr}[T_\tau e^{-\beta H_a} H_b(\tau_k) \cdots H_b(\tau_1)] .$$
(6)

Since  $H_b$ , which can be separated as  $H_b = \sum_{pj} (V_p^j c_p^{\dagger} d_j + V_p^{j*} d_j^{\dagger} c_p) = \tilde{H}_{hyb} + \tilde{H}_{hyb}^{\dagger}$ , contains only one operator in each term which creates or annihilates an electron on the impurity, only even numbers of  $H_b$  with equal numbers of  $\tilde{H}_{hyb}$  and  $\tilde{H}_{hyb}^{\dagger}$  can make non-zero contribution. The partition function therefore becomes

$$Z = \sum_{k=0}^{+\infty} \frac{1}{(2k)!} \int_{0}^{\beta} d\tau_{1} \cdots \int_{0}^{\beta} d\tau_{2k} \operatorname{Tr}[T_{\tau}e^{-\beta H_{a}}H_{b}(\tau_{2k})\cdots H_{b}(\tau_{1})]$$

$$= \sum_{k=0}^{+\infty} \frac{1}{(2k)!} C_{2k}^{k}(k!)^{2} \iiint_{\tau_{1}^{1} < \cdots < \tau_{k}^{r}} d\tau_{1} \cdots d\tau_{k}^{r} \operatorname{Tr}[T_{\tau}e^{-\beta H_{a}}\tilde{H}_{hyb}(\tau_{k})\tilde{H}_{hyb}^{\dagger}(\tau_{k}^{r})\cdots \tilde{H}_{hyb}(\tau_{1})\tilde{H}_{hyb}^{\dagger}(\tau_{1}^{r})]$$

$$= \sum_{k=0}^{+\infty} \iiint_{\tau_{1}^{r} < \cdots < \tau_{k}^{r}} d\tau_{1} \cdots d\tau_{k}^{r} \operatorname{Tr}[T_{\tau}e^{-\beta H_{a}}\tilde{H}_{hyb}(\tau_{k})\tilde{H}_{hyb}^{\dagger}(\tau_{k}^{r})\cdots \tilde{H}_{hyb}(\tau_{1})\tilde{H}_{hyb}^{\dagger}(\tau_{1}^{r})], \qquad (7)$$

where  $C_{2k}^k$  is the number of choices to select  $k \ \tilde{H}_{hyb}$ 's from 2k candidates and  $(k!)^2$  comes from the time ordering of  $\tau$  and  $\tau'$  (i.e., let  $\tau_1 < \tau_2 < \cdots < \tau_k$  and  $\tau'_1 < \tau'_2 < \cdots < \tau'_k$ ).

Inserting the explicit expressions of  $\tilde{H}_{\rm hyb}$  and  $\tilde{H}_{\rm hyb}^{\dagger}$  yields

$$Z = \sum_{k=0}^{+\infty} \iiint_{\tau_{1}^{-} < \cdots < \tau_{k}^{+}} d\tau_{1} \cdots d\tau_{k}^{\prime} \sum_{\substack{j_{1}, \cdots, j_{k} \\ j_{1}^{\prime}, \cdots, j_{k}^{\prime}}} \sum_{p_{1}^{\prime}, \cdots, p_{k}^{\prime}} V_{p_{k}^{\prime}}^{j_{k}^{\prime}} V_{p_{k}}^{j_{k}} \cdots V_{p_{1}^{\prime}}^{j_{1}^{\prime}} V_{p_{1}}^{j_{1}} \\ \times \operatorname{Tr}[T_{\tau} e^{-\beta H_{a}} d_{j_{k}}(\tau_{k}) c_{p_{k}}^{\dagger}(\tau_{k}) c_{p_{k}^{\prime}}(\tau_{k}^{\prime}) d_{j_{k}^{\prime}}^{\dagger}(\tau_{k}^{\prime}) \cdots d_{j_{1}}(\tau_{1}) c_{p_{1}^{\prime}}(\tau_{1}) c_{p_{1}^{\prime}}(\tau_{1}^{\prime}) d_{j_{1}^{\prime}}^{\dagger}(\tau_{1}^{\prime})] \\ = \sum_{k=0}^{+\infty} \iiint_{\tau_{1}^{\prime} < \cdots < \tau_{k}^{\prime}} d\tau_{1} \cdots d\tau_{k}^{\prime} \sum_{\substack{j_{1}, \cdots, j_{k} \\ j_{1}^{\prime}, \cdots, j_{k}^{\prime}}} \sum_{p_{1}^{\prime}, \cdots, p_{k}^{\prime}} V_{p_{k}^{\prime}}^{j_{k}^{\prime}} V_{p_{k}^{\prime}}^{j_{k}^{\prime}} V_{p_{1}^{\prime}}^{j_{1}^{\prime}} W_{p_{1}^{\prime}}^{j_{1}^{\prime}} \\ \times \operatorname{Tr}_{d}[T_{\tau} e^{-\beta H_{\mathrm{loc}}} d_{j_{k}}(\tau_{k}) d_{j_{k}^{\prime}}^{\dagger}(\tau_{k}^{\prime}) \cdots d_{j_{1}}(\tau_{1}) d_{j_{1}^{\prime}}^{\dagger}(\tau_{1}^{\prime})] \times \operatorname{Tr}_{c}[T_{\tau} e^{-\beta H_{\mathrm{bath}}} c_{p_{k}^{\prime}}^{\dagger}(\tau_{k}) c_{p_{k}^{\prime}}(\tau_{k}^{\prime}) \cdots c_{p_{1}^{\prime}}^{\dagger}(\tau_{1}^{\prime})] , \qquad (8)$$

in which the second line is based on the fact that impurity and bath in the trace are decoupled from each other  $([H_{\rm loc}, H_{\rm bath}] = 0, d/d^{\dagger}(\tau) = e^{H_{\rm loc}\tau} d/d^{\dagger} e^{-H_{\rm loc}\tau}$  and  $c/c^{\dagger}(\tau) = e^{H_{\rm bath}\tau} c/c^{\dagger} e^{-H_{\rm bath}\tau})$ .

Since  $H_{\text{bath}}$  is bilinear in creation and annihilation operators, Wick's theorem is valid and therefore can be utilized to simplify the trace over the bath. Thus, we have

$$\frac{1}{Z_{\text{bath}}} \sum_{\substack{p_1, \dots, p_k \\ p'_1, \dots, p'_k}} V_{p'_k}^{j'_k *} V_{p_k}^{j_k} \cdots V_{p'_1}^{j'_1 *} V_{p_1}^{j_1} \operatorname{Tr}_c[T_\tau e^{-\beta H_{\text{bath}}} c^{\dagger}_{p_k}(\tau_k) c_{p'_k}(\tau'_k) \cdots c^{\dagger}_{p_1}(\tau_1) c_{p'_1}(\tau'_1)] \\
= \sum_{\substack{p_1, \dots, p_k \\ p'_1, \dots, p'_k}} V_{p'_k}^{j'_k *} V_{p_k}^{j_k} \cdots V_{p'_1}^{j'_1 *} V_{p_1}^{j_1} \langle T_\tau c^{\dagger}_{p_k}(\tau_k) c_{p'_k}(\tau'_k) \cdots c^{\dagger}_{p_1}(\tau_1) c_{p'_1}(\tau'_1) \rangle \\
= \sum_{s} \sum_{\substack{p_1, \dots, p_k \\ p'_1, \dots, p'_k}} V_{p'_k}^{j'_k *} V_{p_k}^{j_k} \cdots V_{p'_1}^{j'_1 *} V_{p_1}^{j_1} \operatorname{sgn}(s) \langle T_\tau c^{\dagger}_{p_k}(\tau_k) c_{p'_{s(k)}}(\tau'_{s(k)}) \rangle \cdots \langle T_\tau c^{\dagger}_{p_1}(\tau_1) c_{p'_{s(1)}}(\tau'_{s(1)}) \rangle \\
= \sum_{s} \sum_{p_1, \dots, p_k} V_{p_k}^{j'_{s(k)} *} V_{p_k}^{j_k} \cdots V_{p_1}^{j'_{s(1)} *} V_{p_1}^{j_1} \operatorname{sgn}(s) \langle T_\tau c^{\dagger}_{p_k}(\tau_k) c_{p_k}(\tau'_{s(k)}) \rangle \cdots \langle T_\tau c^{\dagger}_{p_1}(\tau_1) c_{p_1}(\tau'_{s(1)}) \rangle , \tag{9}$$

where s is a permutation of  $(1, 2, \dots, k)$ ,  $Z_{\text{bath}} = \text{Tr}[e^{-\beta H_{\text{bath}}}] = \prod_p \prod_{\sigma} (1 + e^{-\beta \varepsilon_p})$  and relation  $\langle T_{\tau} c_{p_i}^{\dagger}(\tau) c_{p'_j}(\tau') \rangle = \delta_{p_i = p'_j} \langle T_{\tau} c_{p_i}^{\dagger}(\tau) c_{p_i}(\tau') \rangle$  has been used. Define the hybridization function as

$$\Delta_{lm}(\tau_l' - \tau_m) = -\sum_p V_p^{j_l'*} V_p^{j_m} \langle T_\tau c_p(\tau_l') c_p^{\dagger}(\tau_m) \rangle = \sum_p \frac{V_p^{j_l^**} V_p^{j_m}}{e^{\varepsilon_p \beta} + 1} \begin{cases} -e^{-\varepsilon_p(\tau - \beta)}, \ 0 < \tau < \beta, \\ e^{-\varepsilon_p \tau}, \ -\beta < \tau < 0, \end{cases}$$
(10)

where  $\tau = \tau'_l - \tau_m$  is the time difference. Then equ (9) can be simplified as  $\sum_s \operatorname{sgn}(s) \Delta_{s(1)1} \Delta_{s(2)2} \cdots \Delta_{s(k)k} = \det \Delta$ , where  $\operatorname{sgn}(s) = (-1)^n$  and *n* is the number of exchanges to obtain  $(1, \cdots, k)$  from  $(s(1), \cdots, s(k))$ .

Unfortunately, since the interaction term makes  $H_{loc}$  not bilinear, the trace over impurity cannot be simplified in this way. As a result, the final form of the partition function is

$$Z = Z_{\text{bath}} \sum_{k=0}^{+\infty} \iiint_{\substack{\tau_1 < \dots < \tau_k \\ \tau_1' < \dots < \tau_k'}} d\tau_1 \cdots d\tau_k' \sum_{\substack{j_1, \dots, j_k \\ j_1', \dots, j_k'}} \operatorname{Tr}_d[T_\tau e^{-\beta H_{\text{loc}}} d_{j_k}(\tau_k) d_{j_k'}^{\dagger}(\tau_k') \cdots d_{j_1}(\tau_1) d_{j_1'}^{\dagger}(\tau_1')] \det \Delta .$$
(11)

In some specific cases, this expression can be simplified further, e.g., some models can be expressed in the so-called segment representation. But this is beyond the scope of this note.

One way to do simulations is to sample configurations stochastically by using continuous-time Monte Carlo method, called CT-HYB in this representation. According to equ (11), the configuration space is described by  $\boldsymbol{x} = (k, \tau, \boldsymbol{j})$ , where  $\boldsymbol{\tau} = (\tau_1, \cdots, \tau_k, \tau'_1, \cdots, \tau'_k)$  and  $\boldsymbol{j} = (j_1, \cdots, j_k, j'_1, \cdots, j'_k)$ . Denote the trace in equ (11) by  $w_{\text{loc}}$ . CT-HYB samples each configuration  $\boldsymbol{x}$  according to its weight  $w_{\text{loc}}(\boldsymbol{x}) \det \Delta(\boldsymbol{x})$ . Note that the expression of hybridization expansion is exact as long as  $H_{\text{b}}(\tau)$  is not infinite large (this will never happen in real simulations), so CT-HYB is always numerically accurate. In general, however,  $w_{\text{loc}}(\boldsymbol{x}) \det \Delta(\boldsymbol{x})$  is not always non-negative. So this method may suffer from severe sign problem.

# **III. APPROXIMATION SCHEMES**

### A. Expanding the configuration space

Another way to do simulations is to use self-consistent resummations of diagrams, which are popular approximations of the hybridization expansion. Generally speaking, we construct the Dyson equation of the dressed propagator, which will be defined later, and find the relation between this propagator and the self-energy based on some self-similar structures of the configurations. Then we just iterate these self-consistent equations to obtain convergent results. To achieve this goal, it is beneficial to expand the configuration space from  $\boldsymbol{x}$  to  $\boldsymbol{x}' = (\boldsymbol{x}, s)$ . In other words, in each fixed  $\boldsymbol{x}$ , there are k! distinct configurations resulting from different pairings between  $(\tau_1, \dots, \tau_k)$  and  $(\tau'_1, \dots, \tau'_k)$ , which can be represented by some curves, called hybridization lines, connecting the corresponding pairs. Then it is easy to figure out the corresponding weight is  $\operatorname{sgn}(s)w_{\operatorname{loc}}(\boldsymbol{x})\Delta_{s(1)1}\Delta_{s(2)2}\cdots\Delta_{s(k)k}$ . Define the bare propagator  $R_0(\tau) = e^{-H_{\operatorname{loc}}\tau}$  and the dress propagator

$$R(\tau) = \sum_{\mathbf{x}'} \operatorname{sgn}(s) \Delta_{s(1)1} \Delta_{s(2)2} \cdots \Delta_{s(k)k} T_{\tau} e^{-\beta H_{\operatorname{loc}}} d_{j_k}(\tau_k) d_{j'_k}^{\dagger}(\tau'_k) \cdots d_{j_1}(\tau_1) d_{j'_1}^{\dagger}(\tau'_1) .$$
(12)

Then it is easy to verify that

$$Z/Z_{\text{bath}} = \text{Tr}[R(\beta)] . \tag{13}$$

And the expectation value of an arbitrary local operator O can be computed easily:

$$\langle O \rangle = \operatorname{Tr}[R(\beta)O]/\operatorname{Tr}[R(\beta)] . \tag{14}$$

After some exploration, it is found that any configuration of  $R(\tau)$  can be constructed by the following steps:

- 1. Choose an order  $k \in \mathbb{N}$ ;
- 2. Specify values of  $\boldsymbol{\tau}$  fulfilling the restrictions  $0 < \tau_1 < \cdots < \tau_k < \tau$  and  $0 < \tau'_1 < \cdots < \tau'_k < \tau$ ;
- 3. Pair  $(\tau_1, \dots, \tau_k)$  and  $(\tau'_1, \dots, \tau'_k)$ , represented by hybridization lines connecting the corresponding pairs.

A typical configuration is shown in Fig. 1. The contribution of each configuration can be written down by the overall sign "sgn" times the contribution calculated from the rules shown in Fig. 2. The overall sign comes from two places: (i) the time-ordering of  $d/d^{\dagger}$  operators and (ii) the pairing between  $(\tau_1, \dots, \tau_k)$  and  $(\tau'_1, \dots, \tau'_k)$ . For fig. 1, the former is +1 and the later is -1, so that sgn = -1 and the contribution is

$$-R_0(\tau-\tau_3')d_{j_3'}^{\dagger}R_0(\tau_3'-\tau_3)d_{j_3}R_0(\tau_3-\tau_2')d_{j_2'}^{\dagger}R_0(\tau_2'-\tau_1')d_{j_1'}^{\dagger}R_0(\tau_1'-\tau_2)d_{j_2}R_0(\tau_2-\tau_1)d_{j_1}R_0(\tau_1).$$

(Note: in diagrams, right direction corresponds to time increasing while in the expression of  $R(\tau)$ , right direction corresponds to time decreasing because of the convention of the time-ordering operator.)

#### **B.** Dyson equation

By analyzing the structure of all diagrams, it is easy to formulate the Dyson equation:

$$R(\tau) = R_0(\tau) + \int_0^\tau d\tau_2 \int_0^{\tau_2} d\tau_1 R(\tau - \tau_2) S(\tau_2 - \tau_1) R_0(\tau_1) , \qquad (15)$$



FIG. 1: A typical configuration of  $R(\tau)$ .



FIG. 2: Contribution for each segment of the diagram. Right direction corresponds to the increasing of the imaginary time and  $\tau = \tau_b - \tau_a$  is the corresponding time increment.

which can be expressed graphically as

 $S(\tau)$  is the set of all one-particle irreducible diagrams, corresponding to the "self-energy" of  $R(\tau)$ . However, because the structure of  $S(\tau)$  is still complicated, we need do some approximations to get a relatively simple expression.

# C. Non-crossing approximation

One popular approximation is the so-called non-crossing approximation (NCA), which neglects any configurations in which any two hybridization lines have a crossover point (the hybridization line is required to be "regular" so that it cannot have more than one crossover point with another line). In this situation, the self-similar structure of  $S(\tau) = S_0(\tau)$  gives another relation with R:

$$S_{0}(\tau) = \sum_{ab} \left[ \operatorname{sgn}_{1} d_{b}^{\dagger} R(\tau) d_{a} \Delta_{ba}(\tau) + \operatorname{sgn}_{2} d_{b} R(\tau) d_{a}^{\dagger} \Delta_{ab}(-\tau) \right] .$$

$$(17)$$

 $sgn_1$  and  $sgn_2$  are signs of corresponding diagrams, which will be analyzed later.

## D. One-crossing approximation

One improvement to NCA is the so-called one-crossing approximation (OCA), which neglects any configurations in which any one hybridization line has more than one crossover points in total with all other lines. OCA can be considered as a more accurate approximation compared to NCA, because it includes more diagrams. For example, Fig. 1 is included in OCA but not in NCA. By similar analysis, a self-similar structure can be found:  $S(\tau) = S_0(\tau) + S_1(\tau)$  and

$$S_{1}(\tau) = \int_{0}^{\tau_{1}} \frac{\tau_{2}}{\tau_{2}} \tau + \int_{0}^{\tau_{1}} \frac{\tau_{2}}{\tau_{1}} \tau_{2} \tau + \int_{0}^{\tau_{1}} \frac{\tau_{2}}{\tau_{1}} \tau_{2} \tau + \int_{0}^{\tau_{1}} \frac{\tau_{2}}{\tau_{1}} \tau_{2} \tau + \int_{0}^{\tau_{1}} \frac{\tau_{2}}{\tau_{1}} \frac{\tau_{2}}{\tau_{2}} \tau + \int_{0}^{\tau_{1}} \frac{\tau_{2}}{\tau_{2}} \frac{\tau_{2}}{\tau_{2}} \tau + \int_{0}^{\tau_{1}} \frac{\tau_{2}}{\tau_{2}}$$

## E. Determining the overall sign

In this subsection, two useful theorems are presented to determine the overall signs in any configurations and in any self-energy expressions. As a result, one can determine signs by just observing the diagrams.

**Theorem 1** Moving a hybridization line across an arbitrary complete diagram D (any hybridization lines starting within D will also end within D) will not change its sign. It has the following graphical expressions:

$$\operatorname{sgn}( \underbrace{\begin{array}{c} & & \\ & & \\ & & \\ \end{array} }) = \operatorname{sgn}( \underbrace{\begin{array}{c} & & \\ & & \\ \end{array} })$$
 $\operatorname{sgn}( \underbrace{\begin{array}{c} & & \\ & & \\ \end{array} }) = \operatorname{sgn}( \underbrace{\begin{array}{c} & & \\ & & \\ \end{array} })$ 

As an exercise,  $sgn_1$  in NCA can be obtained conveniently:

$$\operatorname{sgn}(\overbrace{D}) = \operatorname{sgn}(\overbrace{D}) = \operatorname{sgn}(\overbrace{D}) \times \operatorname{sgn}(\overbrace{D}) \times \operatorname{sgn}(\overbrace{D}) .$$
(19)

Because  $sgn(-\bigcirc -)$  has been incorporated into R, we have  $sgn_1 = sgn(\frown -) = -1$ . Using similar procedure, we get  $sgn_2 = +1$ . So the final from of NCA self-energy is

$$S_0(\tau) = -\sum_{ab} \left[ d_b^{\dagger} R(\tau) d_a \Delta_{ba}(\tau) + d_b R(\tau) d_a^{\dagger} \Delta_{ab}(\beta - \tau) \right] \,. \tag{20}$$

Because in the configurations of  $R_1(\tau)$ , hybridization lines may have a crossover point. To deal with this situation, it is beneficial to introduce another theorem:

**Theorem 2** Switching two adjacent crossing lines will contribute a minus sign. It has the following graphical expressions:



As another exercise, we calculate  $sgn_1$  in OCA.

$$\operatorname{sgn}(\overbrace{D_1}^{\bullet}, \overbrace{D_2}^{\bullet}, \overbrace{D_3}^{\bullet}) = \operatorname{sgn}(\overbrace{)}^{\bullet} \times \operatorname{sgn}(\overbrace{D_1}^{\bullet}, \overbrace{D_2}^{\bullet}, \overbrace{D_3}^{\bullet})$$
$$= -\operatorname{sgn}(\overbrace{)}^{\bullet} \times \operatorname{sgn}(\overbrace{D_1}^{\bullet}, \overbrace{D_2}^{\bullet}, \overbrace{D_3}^{\bullet}). \quad (21)$$

So  $sgn_1 = -sgn(2)$  = -1. Similarly, we get  $sgn_2 = sgn_3 = -sgn_4 = +1$ . So the final from of  $S_1(\tau)$  in OCA is

$$S_{1}(\tau) = -\sum_{abcd} \int_{0}^{\tau} d\tau_{2} \int_{0}^{\tau_{2}} d\tau_{1} \left[ d_{d}^{\dagger} R(\tau - \tau_{2}) d_{c}^{\dagger} R(\tau_{2} - \tau_{1}) d_{b} R(\tau_{1}) d_{a} \Delta_{db}(\tau - \tau_{1}) \Delta_{ca}(\tau_{2}) \right. \\ \left. + d_{d} R(\tau - \tau_{2}) d_{c}^{\dagger} R(\tau_{2} - \tau_{1}) d_{b}^{\dagger} R(\tau_{1}) d_{a} \Delta_{bd}(\beta - \tau + \tau_{1}) \Delta_{ca}(\tau_{2}) \right. \\ \left. + d_{d}^{\dagger} R(\tau - \tau_{2}) d_{c} R(\tau_{2} - \tau_{1}) d_{b} R(\tau_{1}) d_{a}^{\dagger} \Delta_{db}(\tau - \tau_{1}) \Delta_{ac}(\beta - \tau_{2}) \right. \\ \left. + d_{d} R(\tau - \tau_{2}) d_{c} R(\tau_{2} - \tau_{1}) d_{b}^{\dagger} R(\tau_{1}) d_{a}^{\dagger} \Delta_{bd}(\beta - \tau + \tau_{1}) \Delta_{ac}(\beta - \tau_{2}) \right] \,.$$

Using theorem 1 and theorem 2, it is now straightforward to obtain two corollaries:

**Corollary 1**  $S(\tau)$  can be obtained by: i) replace R by  $R_0$  in the self-energy expression; ii) write down its contribution, including sgn; iii) replace  $R_0$  by R.

**Corollary 2** For any configuration, its sign equals  $(-1)^n$ , where n equals the total number of hybridization lines flowing rightward plus the total number of crossover points. In other words,

$$n = \# \text{ of } (\frown) + \# \text{ of } (\bigstar).$$

It is straightforward to verify these corollaries are consistent with the results we obtained before. Using these corollaries, we can easily write a computer program to calculate the analytic expressions for higher-order approximations, which might be beneficial in the future.

[2] A. Rüegg, E. Gull, G. A. Fiete, and A. J. Millis, Phys. Rev. B 87, 075124 (2013).

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