Inchworm Monte Carlo method for open quantum systems

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Abstract

We investigate in this work a recently proposed diagrammatic quantum Monte Carlo method — the inchworm Monte Carlo method — for open quantum systems. We establish its validity rigorously based on resummation of Dyson series. Moreover, we introduce an integro-differential equation formulation for open quantum systems, which illuminates the mathematical structure of the inchworm algorithm. This new formulation leads to an improvement of the inchworm algorithm by introducing classical deterministic time-integration schemes. The numerical method is validated by applications to the spin-boson model.

Keywords: quantum Monte Carlo, open quantum system, diagrammatic methods, spin-boson model

1. Introduction

For realistic quantum systems, the system we are interested in is often coupled with an uninteresting environment to a non-negligible extent, which requires us to study the open quantum system, including the effects such as quantum decoherence and quantum dissipation. The application of open quantum system ranges in a wide variety of quantum fields, including quantum optical systems, nonlinear statistical mechanics, quantum computation, etc. Due to the existence of the quantum bath/environment, the evolution of the projected density matrix of the system is a non-Markovian process. One classical approach is to apply the Nakajima-Zwanzig projection operator technique to obtain an integro-differential master equation. By taking the weak-coupling limit, a Markovian approximation can be obtained, which is easier for numerical simulations. The Markovian approximation however breaks down for open systems with stronger coupling. Approaches that directly simulate the non-Markovian processes include the path-integral approaches such as the QuAPI (quasi-adiabatic propagator path integral) methods and the HEOM (hierarchical equations of motion) technique. These methods yield accurate numerical results, while the computational cost is huge, often unaffordable. To reduce the computational cost, one strategy is to replace the exact summation or numerical integration in these methods by the stochastic Monte Carlo methods. In this paper, we are going to study a specific type of path-integral methods called the diagrammatic quantum Monte Carlo method to solve the time-dependent open quantum systems. In particular, our study is largely motivated by the inchworm Monte Carlo method recently proposed in [4, 2] to reduce the variance in quantum Monte Carlo by diagrammatic resummation.

The basis of the diagrammatic quantum Monte Carlo method has been established as early as 1960s. However, as other quantum Monte Carlo methods, this type of methods also suffer from the notorious dynamical sign problem, meaning that the number of Monte Carlo samples is required to grow at least exponentially in time in order to keep the accuracy of the simulation. To relieve the dynamical sign problem, Stockburger and Grabert introduced stochastic unraveling of influence functionals in [31], and Makri [18, 21] proposed to assume a finite memory time of the bath-correlation function and apply an iterative procedure to efficiently implement the summation. The inchworm Monte Carlo method applies the idea of diagrammatic
resummation as in the bold diagrammatic Monte Carlo method [27] to the real-time evolution of the quantum systems. Similar to the bold-line diagrammatic Monte Carlo method proposed in [11, 12] (see also a more mathematical presentation [16]), the inchworm Monte Carlo method tackles the dynamical sign problem by lumping a large number of diagrams into a “bold line”, which effectively reduces the number of total diagrams to be summed to reach desired numerical accuracy. The inchworm method makes maximum use of the previous calculations, at the expense of higher memory cost for storing all Green’s functions. Despite its success in the application of spin-boson model [3] and the Anderson impurity model [7, 28], it requires better understanding to reveal the intrinsic mathematical structure of the inchworm method, and to further improve the method. In particular, it would be interesting to see how the bold lines are built on the basis of shorter bold lines, and how the bold lines propagate when the iterative procedure is precise. While the answers to these questions are not detailed in the original derivation of the inchworm method [4, 2], in this paper, we will show the validity of the inchworm method with mathematical rigor. The rigorous proof not only justifies the original algorithm, but also leads us to a new formulation of the open quantum system as an integro-differential equation, based on which more accurate and efficient numerical approaches can be developed.

The inchworm Monte Carlo method and the new integro-differential equation formulation will be proved to be applicable for the Ohmic spin-boson model, which is a simple open quantum system widely used as benchmark problems [34, 15, 8]. Based on the integro-differential equation, part of the Monte Carlo integration can be replaced by classical time-integration methods to achieve higher accuracy. The resulting new algorithm will be applied to the spin-boson model to show the numerical efficiency.

The rest of this paper is organized as follows. In Section 2, we introduce the basic formulation of the open quantum system and its Dyson series expansion. Section 3 gives a complete review of the inchworm Monte Carlo method and proves its validity. The integro-differential equation associated with the inchworm algorithm is derived in Section 4. As an application, we analyze the spin-boson model in Section 5. Our new numerical method is introduced in Section 6 and some numerical examples are given in Section 7. A simple summary is given in Section 8 as the end of the paper.

2. Dyson series expansion for open quantum systems

Before considering open quantum system, let us first recall the time-dependent perturbation theory and the associated Dyson series. Consider the von Neumann equation for quantum evolution (of a closed system)

\[ i \frac{d \rho}{dt} = [H, \rho], \]

where \( \rho(t) \) is the density matrix at time \( t \), and \( H \) is the Schrödinger picture Hamiltonian with the form

\[ H = H_0 + W. \]

Here \( H_0 \) is the unperturbed Hamiltonian and \( W \) is viewed as a perturbation. Following the convention, for any Hermitian operator \( A \), we define \( \langle A \rangle = \text{tr}(\rho(0)A) \). We are interested in the evolution of the expectation for a given observable \( O \), defined by

\[ \langle O(t) \rangle = \text{tr}(O\rho(t)) = \text{tr}(Oe^{-itH}\rho(0)e^{itH}) = \langle e^{itH}Oe^{-itH} \rangle. \]

Using standard time dependent perturbation theory, the unitary group \( e^{-itH} \) generated by \( H \) can be represented using a Dyson series expansion [9]

\[ e^{-itH} = \sum_{n=0}^{+\infty} \int_{t>n>\cdots>t_1>0} (-i)^n e^{-i(t-t_n)H_0} W e^{-i(t_{n-1}-t_n)H_0} W \cdots W e^{-i(t_2-t_1)H_0} W e^{-i(t_1)H_0} dt_1 \cdots dt_n, \]

where the integral should be interpreted as

\[ \int_{t>n>\cdots>t_1>0} dt_1 \cdots dt_n = \int_0^t \int_0^{t_1} \cdots \int_0^{t_2} dt_1 \cdots dt_{n-1} dt_n. \]
Inserting the Dyson series (3) into (2), one obtains

\[
\langle O(t) \rangle = \sum_{n=0}^{+\infty} \sum_{n'=0}^{+\infty} \int_{t>t_n>\cdots>t_1>0} \int_{t'>t_n'>\cdots>t'_1>0} (-i)^{n+n'} e^{-i(t-t_n)H_0} W e^{-i(t'-t_n')H_0} W \cdots W e^{-i(t-t_1)H_0} W e^{-i(t'-t'_1)H_0} O \times e^{-i(t-t_n)H_0} W e^{-i(t-n-1)H_0} W \cdots W e^{-i(t-t_1)H_0} W e^{-i(t'_1)H_0} dt'_1 \cdots dt'_n \, dt_1 \cdots dt_n.
\]

(5)

Since the unperturbed Hamiltonian \( H_0 \) is usually easier to solve, the above expansion provides the basis of a feasible approach to find \( \langle O(t) \rangle \) using Monte Carlo method.

For notational simplicity, the above integral is often denoted by the Keldysh contour plotted in Figure 1. The Keldysh contour should be read following the arrows in the diagram, and therefore has a forward (upper) branch and a backward (lower) branch. The symbols are interpreted as follows:

- Each line segment connecting two adjacent time points labeled by \( t_n \) and \( t_l \) means a propagator \( e^{-i(t-t_n)H_0} \). On the forward branch, \( t_f > t_n \), while on the backward branch, \( t_f < t_n \).
- Each black dot introduces a perturbation operator \( \pm iW \), where we take the minus sign on the forward branch, and the plus sign on the backward branch. At the same time, every black dot also represents an integral with respect to the label, whose range is from 0 to the adjacent label to its right.
- The cross sign at time \( t \) means the observable in the Schrödinger picture.

Note that according to the above interpretation, two Keldysh contours differ only when at least one of the values of \( n \), \( n' \) and \( t \) is different, while the positions of the labels on each branch do not matter. Thus, by taking the expectation \( \langle \cdot \rangle \) of this “contour”, we obtain the summand in (5). Therefore \( \langle O(t) \rangle \) can be understood as the sum of the expectations of all possible Keldysh contours.

\[
\text{Figure 1: Keldysh contour}
\]

Such an interpretation also shows that we do not need to distinguish the forward and backward branches when writing down the integrals. In fact, when the series (5) is absolutely convergent in the sense that

\[
\sum_{n=0}^{+\infty} \sum_{n'=0}^{+\infty} \int_{t>t_n>\cdots>t_1>0} \int_{t'>t_n'>\cdots>t'_1>0} \left| \langle e^{i(t-t_n)H_0} W e^{i(t'-t_n')H_0} W \cdots W e^{i(t-t_1)H_0} W e^{i(t'-t'_1)H_0} O \times e^{-i(t-t_n)H_0} W e^{-i(t-n-1)H_0} W \cdots W e^{-i(t-t_1)H_0} W e^{-i(t'_1)H_0} \rangle \right| dt'_1 \cdots dt'_n \, dt_1 \cdots dt_n < +\infty,
\]

we can reformulate (5) as

\[
\langle O(t) \rangle = \sum_{m=0}^{+\infty} \sum_{2t>s_m>\cdots>s_1>0} (-1)^{\# \{ s < t \}} l^m \times \left\langle G^{(0)}(2t, s_m) W G^{(0)}(s_m, s_{m-1}) W \cdots W G^{(0)}(s_2, s_1) W G^{(0)}(s_1, 0) \right\rangle \, ds_1 \cdots ds_m,
\]

(6)

where we use \( s \) as a short-hand for the decreasing sequence \( (s_m, \cdots, s_1) \) and \( \# \{ s < t \} \) is the number of elements in \( s \) which are less than \( t \), i.e., the number of \( s_k \) on the forward branch of the Keldysh contour.
For a given $t$, the propagator $G^{(0)}$ is defined as

$$
G^{(0)}(s_t, s_i) = \begin{cases} 
  e^{-i(s_t-s_i)H_0}, & \text{if } s_i \leq s_t < t, \\
  e^{i(s_t-s_i)H_0}, & \text{if } t \leq s_i \leq s_t, \\
  e^{i(s_t-s_i)H_0} O e^{-i(t-s_i)H_0}, & \text{if } s_i < t \leq s_t.
\end{cases}
$$

The integral (6) can also be understood graphically as the “unfolded Keldysh contour” plotted in Figure 2. In order to use only a single integral in (6), we set the range of the unfolded Keldysh contour to be $[0, 2t]$, and the mapping of time points from the unfolded Keldysh contour to the original Keldysh contour has been implied in the definition of $G^{(0)}(\cdot, \cdot)$. By comparing (6) with Figure 2, one can see that $G^{(0)}(\cdot, \cdot)$ can be considered as the unperturbed propagator on the unfolded Keldysh contour, with an action of observable $O$ at time $t$.

![Unfolded Keldysh contour](image)

Figure 2: Unfolded Keldysh contour

To proceed, we now assume that the von Neumann equation (1) describes an open quantum system coupled with a bath, which means that both $\rho$ and $H$ are Hermitian operators on the Hilbert space $\mathcal{H} = \mathcal{H}_s \otimes \mathcal{H}_b$, with $\mathcal{H}_s$ and $\mathcal{H}_b$ representing respectively the Hilbert spaces associated with the system and the bath. We consider the interaction picture and take $H_0$ to be the Hamiltonian without coupling:

$$
H_0 = H_s \otimes \text{Id}_b + \text{Id}_s \otimes H_b,
$$

where $H_s$ and $H_b$ are respectively the uncoupled Hamiltonians for the system and the bath, and $\text{Id}_s$ and $\text{Id}_b$ are respectively the identity operators for the system and the bath. Then the perturbation $W$ describes the coupling, and here we assume $W$ takes the form without loss of generality

$$
W = W_s \otimes W_b.
$$

Furthermore, we assume the initial density matrix has the separable form $\rho(0) = \rho_s \otimes \rho_b$, and we are concerned with observables acting only on the system $O = O_s \otimes \text{Id}_b$ (recall that physically the system is the interesting part). With these assumptions, (6) becomes

$$
\langle O(t) \rangle = \sum_{m=0}^{+\infty} \int_{2t>s_m, \ldots, s_1>0} (-1)^m(s<t) \operatorname{tr}_s(\rho U^{(0)}(2t, s, 0)) \mathcal{L}_b(s) \, ds_1 \cdots ds_m,
$$

where the integrand is separated into $U^{(0)}$ and $\mathcal{L}_b$ for the system and bath parts:

$$
U^{(0)}(s_t, s_i) = U^{(0)}(s_t, s_m, \ldots, s_1, s_i) = G_s^{(0)}(s_t, s_m) W_s G_b^{(0)}(s_m, s_{m-1}) W_s \cdots W_s G_s^{(0)}(s_2, s_1) W_s G_b^{(0)}(s_1, s_i),
$$

$$
\mathcal{L}_b(s) = \operatorname{tr}_b(\rho_b G_b^{(0)}(2t, s_m) W_b G_b^{(0)}(s_m, s_{m-1}) W_b \cdots W_b G_b^{(0)}(s_2, s_1) W_b G_b^{(0)}(s_1, 0)),
$$

where $\operatorname{tr}_s$ and $\operatorname{tr}_b$ take traces of the system and bath respectively. The propagators $G_s^{(0)}$ and $G_b^{(0)}$ are defined similarly to (7):

$$
G_s^{(0)}(s_t, s_i) = \begin{cases} 
  e^{-i(s_t-s_i)H_s}, & \text{if } s_i \leq s_t < t, \\
  e^{-i(s_t-s_i)H_s}, & \text{if } t \leq s_i \leq s_t, \\
  e^{-i(t-s_i)H_s} O_s e^{-i(t-s_i)H_s}, & \text{if } s_i < t \leq s_t.
\end{cases}
$$
There exists a function \( B \)

\[ G_b^{(0)}(s_t, s_i) = \begin{cases} 
  e^{-i(s_t-s_l)H_b}, & \text{if } s_i \leq s_t < t, \\
  e^{-i(s_t-s_l)H_b}, & \text{if } t \leq s_i \leq s_t, \\
  e^{-i(2s_t-s_l)H_b}, & \text{if } s_i < t \leq s_t.
\end{cases} \quad (12) \]

Note that the observable \( O \) is inserted into the propagator \( G_b^{(0)} \) to keep the expression in (9) concise.

### 3. Inchworm algorithm

In this section, we are going to study the inchworm algorithm introduced in [2], where the method was proposed for the spin-Boson model from a purely diagrammatic point of view. By matching the mathematical interpretation and the diagrammatic interpretation of the algorithm, we will establish rigorously the validity of the algorithm in a more general sense. The central idea of the algorithm is to consider the problem as an evolution problem and reuse as much previous information as possible, and we will start the introduction of the algorithm by introducing the “full propagators”, which are exactly the carriers of the information to be recycled.

#### 3.1. Full propagator and its Dyson series expansion

The inchworm algorithm proposed in [2] considers the following “full propagators”:

\[ G(s_t, s_i) = \begin{cases} 
  \text{tr}_b(\rho_b G_b^{(0)}(2t, s_t) e^{-i(s_t-s_l)H_b} G_b^{(0)}(s_l, 0)), & \text{if } s_i \leq s_l < t, \\
  \text{tr}_b(\rho_b G_b^{(0)}(2t, s_t) e^{-i(s_t-s_l)H_b} G_b^{(0)}(s_l, 0)), & \text{if } t \leq s_i \leq s_l, \\
  \text{tr}_b(\rho_b G_b^{(0)}(2t, s_t) e^{i(-s_t-H_b) G_b^{(0)}(s_l, 0)}), & \text{if } s_i < t \leq s_l.
\end{cases} \quad (13) \]

Here the trace is taken only on the space of the bath \( \mathcal{H}_b \), and hence \( G(s_t, s_i) \) is an operator on \( \mathcal{H}_b \). Following the same method from (3) to (7), we get the following Dyson series expansion for \( G(s_t, s_i) \):

\[ G(s_t, s_i) = \sum_{m=0}^{+\infty} \int_{s_i > s_m > \cdots > s_1 > s_t} (-1)^{\#\{s < t\} - m} \text{tr}_b \left( \rho_b G_b^{(0)}(2t, s_t) G_b^{(0)}(s_t, s_m) W G_b^{(0)}(s_m, s_{m-1}) W \cdots W G_b^{(0)}(s_2, s_1) \right) d s_1 \cdots d s_m. \quad (14) \]

To apply the inchworm algorithm, we need the following two hypotheses, which are abstracted from the spin-Boson model studied in [2]:

(H1) The initial density matrix for the bath \( \rho_b \) commutes with the Hamiltonian \( H_b \). Physically, this condition holds when the bath is initially at the thermal equilibrium associated with the Hamiltonian \( H_b \).

(H2) There exists a function \( B(\cdot, \cdot) \) such that the following Wick’s theorem holds:

\[ \mathcal{L}_b(s_m, \cdots, s_1) = \begin{cases} 
  0, & \text{if } m \text{ is odd}, \\
  \sum_{q \in \mathcal{Q}(s_m, \cdots, s_1)} \mathcal{L}(q), & \text{if } m \text{ is even},
\end{cases} \quad (15) \]

where the right hand side is given by all possible ordered pairings of the time points:

\[ \mathcal{L}(q) = \prod_{(\tau_1, \tau_2) \in q} B(\tau_1, \tau_2), \]

\[ \mathcal{Q}(s_m, \cdots, s_1) = \left\{ \left( s_{j_1, s_{k_1}}, \cdots, (s_{j_m/2, s_{k_m/2}}) \right) \mid \{ j_1, \cdots, j_{m/2}, k_1, \cdots, k_{m/2} \} = \{ 1, \cdots, m \}, \right. \]

\[ s_{j_l} \leq s_{k_l} \text{ for any } l = 1, \cdots, m/2 \} \right\}.

When \( m = 0 \), the value of \( \mathcal{L}(\emptyset) \) is defined as 1.
In hypothesis (H2), $Q(s_m, \ldots, s_1)$ is the set of all possible ordered pairings of $\{s_m, \ldots, s_1\}$. For example,

$$Q(s_2, s_1) = \{\{(s_1, s_2)\}\},$$

$$Q(s_4, s_3, s_2, s_1) = \{\{(s_1, s_2), (s_3, s_4)\}, \{(s_1, s_3), (s_2, s_4)\}, \{(s_1, s_4), (s_2, s_3)\}\}.$$

We can also represent these sets by diagrams:

$$Q(s_2, s_1) = \{\hspace{-1cm}\begin{array}{c}
\includegraphics[scale=0.5]{diag1.png} \\
\end{array}\hspace{-1cm}\},$$

$$Q(s_4, s_3, s_2, s_1) = \{\hspace{-1cm}\begin{array}{c}
\includegraphics[scale=0.5]{diag2.png} \\
\includegraphics[scale=0.5]{diag3.png} \\
\includegraphics[scale=0.5]{diag4.png} \\
\end{array}\hspace{-1cm}\}.$$

Manifestly, each arc stands for a pair formed by the labels on the two end points, and each diagram denotes a set of pairs.

As in the quantum field theory, Wick’s theorem (hypothesis (H2)) turns integrals into diagrams, which allows us to use diagrammatic quantum Monte Carlo methods in the simulation. The first hypothesis (H1) allows us to associate the full propagators with observables. Precisely, when $s_i < t < s_f$ and $s_i + s_f = 2t$,

$$\text{tr}_x(\rho_s G(s_f, s_i)) = \text{tr}\left(\rho_s e^{i(t-s_i)H} O e^{-i(t-s_f)H}\right) = \langle O(t-s_i) \rangle,$$

which shows that the evolution of the observable from time 0 to $t$ can be fully obtained once the propagator $G(s_f, s_i)$ is solved for every pair of $s_f$ and $s_i$. By splitting system and bath parts and applying Wick’s theorem (15), we get from (14) that

$$G(s_f, s_i) = \sum_{m=0}^{+\infty} \int_{s_f > s_m > \cdots > s_1 > s_i} \sum_{q \in Q(s)} (-1)^\#\{s<t\} i^m U(0)_{s_f, s_i} L(q) ds_1 \cdots ds_m. \tag{16}$$

Here the integral of $(-1)^\#\{s<t\} i^m U(0)_{s_f, s_i} L(q)$ can also be represented by a diagram like Figure 3, which is interpreted by

- Each line segment connecting two adjacent time points labeled by $t_s$ and $t_f$ means a propagator $G^{(0)}_{s_f}(t_f, t_s)$.
- Each black dot introduces a perturbation operator $\pm i W_s$, and we take the minus sign on the forward branch, and the plus sign on the backward branch. Here the label for time $t$, which separates the two branches of the Keldysh contour, is omitted. Additionally, each black dot also represents the integral with respect to the label over the interval from $s_i$ to its next label.
- The arc connecting two time points $t_s$ and $t_f$ stands for $B(t_s, t_f)$.

![Diagram](image.png)

**Figure 3:** Diagrammatic representation for the integral of $(-1)^\#\{s<t\} i^m U(0)_{s_f, s_i} L(q)$ when $m = 6$ and $q = \{(s_1, s_6), (s_2, s_4), (s_3, s_3)\}$.

Such a diagrammatic representation allows us to rewrite (16) as

$$\begin{array}{c}
\includegraphics[scale=0.5]{diag5.png} \\
= \hspace{-1cm}\begin{array}{c}
\includegraphics[scale=0.5]{diag6.png} \\
\includegraphics[scale=0.5]{diag7.png} \\
\end{array}\hspace{-1cm}\end{array} + \hspace{-1cm}\begin{array}{c}
\includegraphics[scale=0.5]{diag8.png} \\
\includegraphics[scale=0.5]{diag9.png} \\
\end{array} + \cdots \tag{17}
\end{array}$$
where the bold line on the left side represents the full propagator $G(s_f, s_i)$. The Monte Carlo method based on (17) is referred to as “bare diagrammatic quantum Monte Carlo” method in [2], which is essentially identical to the Monte Carlo method based on the Dyson series expansion (3), except that the bath function $L_0(s)$ is also evaluated by the Monte Carlo method.

3.2. Description of the inchworm algorithm

The inchworm algorithm uses another series expansion of $G(s_f, s_i)$ that leads to efficient use of results of previous time steps for future computations on the contour. Before introducing the inchworm series, we need the following definitions:

**Definition 1** (Linked pairs and linked set of pairs). Two pairs of real numbers $(s_1, s_2)$ and $(\tau_1, \tau_2)$ satisfying $s_1 \leq s_2$ and $\tau_1 \leq \tau_2$ are linked if either of the following two statements holds:

1. $s_1 \leq \tau_1 \leq s_2$ and $\tau_1 \leq s_2 \leq \tau_2$.
2. $\tau_1 \leq s_1 \leq \tau_2$ and $s_1 \leq \tau_2 \leq s_2$.

For two sets of pairs $q_1$ and $q_2$, we say the two sets $q_1$ and $q_2$ are linked if there exists $(s_1, s_2) \in q_1$ and $(\tau_1, \tau_2) \in q_2$ such that $(s_1, s_2)$ and $(\tau_1, \tau_2)$ are linked.

Given a set of pairs $q$, we say $q$ is a linked set of pairs if it cannot be decomposed into the union of two sets of pairs that are not linked. We define

$$Q_\downarrow(s_m, \ldots, s_1) = \{q \in Q(s_m, \ldots, s_1) \mid q \text{ is linked}\}.$$  

For example, according to the above definition, when $s_1 < s_2 < s_3 < s_4$, the pairs $(s_1, s_3)$ and $(s_2, s_4)$ are linked, while $(s_1, s_4)$ and $(s_2, s_3)$ are not, which can be also clearly seen from the diagrams:

It is also clear that the set of pairs in the left diagram is a linked set, while the other one is not.

**Definition 2** (Linked component decomposition). For any $q \in Q(s_m, \ldots, s_1)$, there exists a collection of its disjoint subsets $q_1, q_2, \ldots, q_n$ such that

1. $q_1, q_2, \ldots, q_n$ are all linked sets of pairs;
2. $q = q_1 \cup q_2 \cup \cdots \cup q_n$;
3. $q_{n_1}$ and $q_{n_2}$ are not linked if $n_1 \neq n_2$.

We call $q = q_1 \cup q_2 \cup \cdots \cup q_n$ the linked component decomposition of $q$.

An example of the linked component decomposition is given below by digrammatic representations:

**Definition 3** (Inchworm properness). Given a decreasing sequence of real numbers $s_m, \ldots, s_1$ and a real number $s_\uparrow$, a set of integer pairs $q \in Q(s_m, \ldots, s_1)$ is called inchworm proper, if in its linked component decomposition $q = q_1 \cup q_2 \cup \cdots \cup q_n$, each $q_k$ contains at least one point greater than or equal to $s_\uparrow$. Below we use $Q_{s_\uparrow}(s_m, \ldots, s_1)$ to denote the collection of all inchworm proper pair sets.

The structure of the set $Q_{s_\uparrow}(s_m, \ldots, s_1)$ is determined by the relative position of $s_\uparrow$ in the decreasing sequence $s_m, \ldots, s_1$. For example, if $m = 4$ and $s_\uparrow \in (s_2, s_3)$, then

$$Q_{s_\uparrow}(s_4, s_3, s_2, s_1) = \{(s_1, s_3), (s_2, s_4)\}, \{(s_1, s_4), (s_2, s_3)\} = \begin{array}{c}
\quad \\
\quad \\
\quad \\
\quad \\
\end{array}.$$  

if $s_\uparrow \in (s_3, s_4)$, then $Q_{s_\uparrow}(s_4, s_3, s_2, s_1)$ contains only one element:

$$Q_{s_\uparrow}(s_4, s_3, s_2, s_1) = \{(s_1, s_3), (s_2, s_4)\} = \begin{array}{c}
\quad \\
\quad \\
\quad \\
\end{array}.$$
In fact, if \( s_\uparrow \in (s_{m-1}, s_m] \), the number of sets of pairs in the linked component decomposition of an inchworm proper set \( q \) must be 1, which means \( q \) is linked. Therefore we have

\[
\mathcal{Q}_{s_\uparrow}(s_m, \cdots, s_1) = \mathcal{Q}_q(s_m, \cdots, s_1), \quad \text{if } s_{m-1} < s_\uparrow \leq s_m. \tag{18}
\]

Based on the definition of inchworm proper set of pairs, we have the following theorem:

**Theorem 1.** Suppose the Dyson series (16) is absolutely convergent in the sense that

\[
\sum_{m=0}^{+\infty} \int_{s_f > s_m > \cdots > s_1 > s_t} \mathcal{U}(s_f, s, s_t) \left\| L(q) \right\| ds \cdots ds_m < +\infty, \quad \forall s_f, s_t \in [0, 2t], \tag{19}
\]

where \( \left\| \cdot \right\| \) is the operator norm in the Hilbert space \( \mathcal{H}_s \). For any \( s_\uparrow \in (s_f, s_t) \), we have

\[
G(s_f, s_i) = G_{s_\uparrow}(s_f, s_i) + \sum_{m=2}^{+\infty} \int_{s_f > s_m > \cdots > s_1 > s_t} (-1)^\# \{s < t\} \left( \sum_{q \in \mathcal{Q}_q(s)} L(q) \right) \times \]

\[
G_{s_\uparrow}(s_f, s_m) W_s G_{s_\uparrow}(s_m, s_{m-1}) W_s \cdots W_s G_{s_\uparrow}(s_2, s_1) W_s G_{s_\uparrow}(s_1, s_i) ds_1 \cdots ds_m,
\]

where

\[
G_{s_\uparrow}(s_f, s_i) = \begin{cases} 
G(s_f, s_i), & \text{if } s_i \leq s_f \leq s_\uparrow, \\
G^{(0)}(s_f, s_i), & \text{if } s_\uparrow < s_i \leq s_f, \\
G^{(0)}(s_f, s_\uparrow) G(s_\uparrow, s_i), & \text{if } s_i \leq s_\uparrow < s_f.
\end{cases}
\]

Note that the right-hand side of (20) has a very similar structure to (16), except that the bare Green’s function \( G^{(0)} \) is replaced by \( G_{s_\uparrow} \), and the sum consists only of “inchworm proper diagrams”, which will be further discussed below. This theorem shows that \( G(s_f, s_i) \) can be evaluated by the Monte Carlo simulation of the right-hand side of (20) based on the knowledge of \( G(\tau_2, \tau_1) \) for all \( s_i < \tau_1 < \tau_2 < s_\uparrow \). Therefore the algorithm can be designed as follows:

Choose a time step \( \Delta t = t/N \)
for \( s_f \) from 0 to 2\( t \) with step \( \Delta t \)
for \( s_i \) from \( s_f \) to 0 with step \( -\Delta t \)
end for
end for

This algorithm computes all \( G(s_f, s_i) \) when \( s_i \) and \( s_f \) are multiples of \( \Delta t \). When evaluating the right-hand side of (20), interpolation might be needed to get \( G(\tau_2, \tau_1) \) when \( \tau_1 \) or \( \tau_2 \) is not an integer multiple of \( \Delta t \).

Details about the interpolation will be given in Section 6.

A rigorous proof of Theorem 1 will be given in the next section. Here we would like to provide the diagrammatic understanding of this algorithm, following [2, 3]. As already mentioned, on the right-hand side of (20), only “inchworm proper diagrams” appear, for example,

\[
\text{(21)}
\]

These diagrams are interpreted as follows:
3.3. Proof of Theorem 1

This section is devoted to the proof of (20), which is to equate the right-hand side of (20) and the series (16). To see this, we will first introduce $s_t$ to the expansion of $G(s_t, s_t)$ by the following lemma:

**Lemma 1.** When the Dyson series (16) is absolutely convergent in the sense of (19), for any $s_t \in [s, s_i)$, it holds that

$$G(s_t, s_i) = G^{(0)}(s_t, s_t)G(s_t, s_i) + \sum_{m=2}^{+\infty} \sum_{p=0}^{m-1} \int_{s_i>s_1>s_2>\cdots>s_{p+1}>s_t} \int_{s_1>s_2>\cdots>s_{p+1}>s_t} \int_{s_p} \int_{s_3} \cdots \int_{s_{p+1}} \sum_{q \in Q(s)} (-1)^{\#(s_t, s_i)} \mathcal{U}^{(0)}(s_t, s_i, s_q) L(q) \, ds_1 \cdots ds_p \, ds_{p+1} \cdots ds_m. \tag{23}$$
Proof. For any positive integer \( m \), it holds that

\[
\int_{s_t > s_{m+1} > \cdots > s_1 > s_t} \varphi(s) \, ds_1 \cdots ds_m = \sum_{p=0}^{m} \int_{s_t > s_{m+1} > \cdots > s_p+1 > s_t} \int_{s_t > s_{p+1} > s_t} \varphi(s) \, ds_1 \cdots ds_p \, ds_{p+1} \cdots ds_m
\]

for any function \( \varphi \). This can be proven by mathematical induction since when the above equation holds for some \( m \), we have

\[
\int_{s_t > s_{m+1} > \cdots > s_1 > s_t} \varphi(s) \, ds_1 \cdots ds_{m+1} = \int_{s_t} \left( \int_{s_{m+1} > s_m > \cdots > s_1 > s_t} \varphi(s) \, ds_1 \cdots ds_m \right) \, ds_{m+1}
\]

\[
= \int_{s_t} \left( \int_{s_{m+1} > s_m > \cdots > s_1 > s_t} \varphi(s) \, ds_1 \cdots ds_m \right) \, ds_{m+1} + \int_{s_t} \left( \int_{s_{m+1} > s_m > \cdots > s_1 > s_t} \varphi(s) \, ds_1 \cdots ds_m \right) \, ds_{m+1}
\]

\[
= \int_{s_t > s_{m+1} > \cdots > s_1 > s_t} \varphi(s) \, ds_1 \cdots ds_{m+1}
\]

\[
+ \sum_{p=0}^{m} \int_{s_t > s_{m+1} > \cdots > s_p+1 > s_t} \int_{s_t > s_{p+1} > s_t} \varphi(s) \, ds_1 \cdots ds_p \, ds_{p+1} \cdots ds_{m+1}
\]

\[
= \sum_{p=0}^{m} \int_{s_t > s_{m+1} > \cdots > s_p+1 > s_t} \int_{s_t > s_{m+1} > \cdots > s_p+1 > s_t} \varphi(s) \, ds_1 \cdots ds_p \, ds_{p+1} \cdots ds_{m+1},
\]

and it is obvious that (24) holds for \( m = 1 \). By (24), we can rewrite (16) as

\[
G(s_{t}, s_{t}) = \sum_{m=0}^{\infty} \sum_{\text{m is even}} \int_{s_t > s_{m+1} > \cdots > s_1 > s_t} \sum_{q \in Q(s)} (-1)^{\#(s < t) \cdot m} \mathcal{U}^{(0)}(s_t, s, s_t) \mathcal{L}(q) \, ds_1 \cdots ds_p \, ds_{p+1} \cdots ds_m
\]

\[
= \sum_{m=0}^{\infty} \int_{s_t > s_{m+1} > \cdots > s_1 > s_t} \sum_{q \in Q(s)} (-1)^{\#(s < t) \cdot m} \mathcal{U}^{(0)}(s_t, s, s_t) \mathcal{L}(q) \, ds_1 \cdots ds_m \tag{25}
\]

\[
+ \sum_{m=2}^{\infty} \sum_{m=0}^{m-1} \int_{s_t > s_{m+1} > \cdots > s_1 > s_t} \int_{s_t > s_{m+1} > \cdots > s_1 > s_t} \sum_{q \in Q(s)} (-1)^{\#(s < t) \cdot m} \mathcal{U}^{(0)}(s_t, s, s_t) \mathcal{L}(q) \, ds_1 \cdots ds_m
\]

where we have used the absolute convergence (19) to ensure the validity of the second equality. When \( s_m < s_t \), we have \( \mathcal{U}^{(0)}(s_t, s, s_t) = G^{(0)}_{s_t}(s_t, s_t) \mathcal{U}^{(0)}(s_t, s_t, s_t) \). Hence

\[
\sum_{m=0}^{\infty} \int_{s_t > s_{m+1} > \cdots > s_1 > s_t} \sum_{q \in Q(s)} (-1)^{\#(s < t) \cdot m} \mathcal{U}^{(0)}(s_t, s, s_t) \mathcal{L}(q) \, ds_1 \cdots ds_m
\]

\[
= G^{(0)}_{s_t}(s_t, s_t) \sum_{m=0}^{\infty} \int_{s_t > s_{m+1} > \cdots > s_1 > s_t} \sum_{q \in Q(s)} (-1)^{\#(s < t) \cdot m} \mathcal{U}^{(0)}(s_t, s, s_t) \mathcal{L}(q) \, ds_1 \cdots ds_m = G^{(0)}_{s_t}(s_t, s_t) \mathcal{U}^{(0)}(s_t, s_t, s_t).
\]
Inserting this equation into (25) yields our conclusion (23).

By the above lemma, we have extracted the first diagram in (22) from the definition of the bold line (17). Our next step is to introduce the inchworm proper pairings to the series expansion. The basic idea is to decompose every set of pairs into the union of one inchworm proper subset and several other unlinked subsets. The result reads

Lemma 2. When the Dyson series (16) is absolutely convergent in the sense of (19), for any \( s^+ \in (s_1, s_2) \), it holds that

\[
G(s_1, s_i) = G^{(0)}(s_1, s_i)G(s^+, s_1) + \sum_{\tilde{m}=2}^{+\infty} \sum_{\tilde{n}=2}^{\tilde{m}-1} \int_{s_1}^{s_2} \cdots \int_{s_{n+1}}^{s_1} \frac{\prod_{p=0}^{\tilde{p}} \prod_{n_p=0}^{\tilde{p}-p-n_{p-1}} \prod_{n_{p-1}=0}^{\tilde{p}-p-n_{p-2}} \cdots \prod_{n_1=0}^{\tilde{p}-p-n_1} q \in Q_{s^+}(s) \prod_{q \in Q(s^{(p)})} \prod_{q_0 \in Q(s^{(0)})}}{(-1)^{\#(\tilde{s}<\tilde{t})} \prod_{\tilde{m}} U^{(0)}(s_1, \tilde{s}, s_1) \mathcal{L}(\tilde{q})} d\tilde{s}_1 \cdots d\tilde{s}_{\tilde{m}-1} d\tilde{s}_{\tilde{m}},
\]

(26)

where the notations are

- \( \tilde{s} = (\tilde{s}_{\tilde{m}}, \ldots, \tilde{s}_1) \);
- \( \tilde{q} = q \cup q_p \cup \cdots \cup q_0 \);
- \( s, s^{(p)} , \ldots, s^{(0)} \) are all subsequences of \( \tilde{s} \), defined by

\[
\tilde{s} = (s_m, s_{p+1}, s^{(p)}, s_p, s^{(p-1)}, s_{p-1}, \ldots, s^{(1)}, s_1, s^{(0)}),
\]

where \( m = \tilde{m} + p - \tilde{p} \) and \( s = (s_m, \ldots, s_1) \).

Proof. By comparing (26) with (23), we construct the following map for given \( \tilde{m}, \tilde{p} \) and a decreasing sequence \( \tilde{s} \):

\[
\mathcal{P} \quad \rightarrow \quad Q(\tilde{s})
\]

\[
(p, n_p, \ldots, n_1, q, q_p, \ldots, q_0) \quad \mapsto \quad q \cup q_p \cup \cdots \cup q_0
\]

(27)

where

\[
\mathcal{P} = \{(p, n_p, \ldots, n_1, q, q_p, \ldots, q_0) \mid p \in \{\tilde{p}, \tilde{p} - 2, \ldots, \tilde{p} - 2[\tilde{p}/2]\}; n_1 + \cdots + n_p \leq \tilde{p} - p;
\]

\[
n_p, \ldots, n_1 \text{ are even}; q \in Q_{s^+}(s); q_p \in Q(s^{(p)}); \cdots; q_0 \in Q(s^{(0)})\}
\]

The subsequences \( s, s^{(p)} , \ldots, s^{(0)} \) are defined as in the lemma. If (27) is a bijection, then (26) is a rearrangement of the series (23). The equality (26) then follows by the absolute convergence of the Dyson series.

Now it remains only to show that (27) is one-to-one. The definition of \( \mathcal{P} \) shows that in a map \( (p, n_p, \ldots, n_1, q, q_p, \ldots, q_0) \mapsto \tilde{q} \), if \( q \) and \( \tilde{q} \) are given, other parameters are automatically determined by

\[
\begin{align*}
p & = 2|q| - (\tilde{m} - \tilde{p}), \quad q_0 = \{(\tau_1, \tau_2) \in \tilde{q} \mid \tau_1 < \tau_2 < s_1\}, \\
q_k & = \{(\tau_1, \tau_2) \in \tilde{q} \mid s_k < \tau_1 < \tau_2 < s_{k+1}\}, \quad k = 1, \ldots, p, \\
n_k & = 2|q_k|, \quad k = 1, \ldots, p.
\end{align*}
\]

(28)

Here \( 2|q| \) (\( 2|q_k| \)) is actually the number of time points in \( q \) (\( q_k \)). Meanwhile, the set of pairs \( q \) satisfies

1. \( q \) is not linked to any other pair in \( \tilde{q} \);
2. \( q \in Q_{s^+}(s) \) and therefore contains \( \tilde{s}_{\tilde{p}+1}, \ldots, \tilde{s}_{\tilde{m}} \).
Now we consider an arbitrary set of pairs \( \tilde{q} \in \mathcal{Q}(\tilde{s}) \) with its linked component decomposition being \( \tilde{q} = \tilde{q}_1 \cup \cdots \cup \tilde{q}_n \). The only subset of \( \tilde{q} \) satisfying the above two conditions is
\[
\tilde{q} = \bigcup \{ \tilde{q}_k \mid \tilde{q}_k \text{ contains a pair } (\tau_1, \tau_2) \text{ with } \tau_2 > \tilde{s}_\tilde{p} \}.
\]
Taking such a \( \tilde{q} \) and applying (28) to find other parameters, we obtain an inverse image of \( \tilde{q} \). This inverse image is unique due to the uniqueness of \( q \), which yields that (27) is a bijection.

**Lemma 3.** When the Dyson series (16) is absolutely convergent in the sense of (19), for any \( s_\tau \in (s, s_\tau) \), it holds that
\[
G(s_\tau, s_i) = G^{(p)}(s_\tau, s_\tau)G(s_\tau, s_i) + \sum_{m=2}^{+\infty} \sum_{p=0}^{m-1} \sum_{n_p=0}^{+\infty} \int_{s_\tau > \tilde{s}_m > \cdots > \tilde{s}_{p+1} > s_\tau} \int_{s_\tau > \tilde{s}_p > \cdots > \tilde{s}_1 > s_i} (-1)^{\#(s < t)} \tilde{u}^{(0)}(s_\tau, \tilde{s}, s_1) \mathcal{L}(\tilde{q}) d\tilde{s}_1 \cdots d\tilde{s}_{p+1} \cdots d\tilde{s}_m,
\]
where \( \tilde{m} = m + n_0 + \cdots + n_p \), and other parameters are defined in the same way as in Lemma 2.

**Proof.** It is not difficult to see that the map
\[
\mathcal{P} \quad \mapsto \quad \tilde{\mathcal{P}}
\]
with \( \tilde{m} = m + n_0 + \cdots + n_0 \) and \( \tilde{p} = p + n_0 + \cdots + n_0 \) is a bijection. Here
\[
\mathcal{P} = \left\{ (m, p, n_p, \cdots, n_0) \mid m, n_p, \cdots, n_0 \text{ are positive and even; } p \in \{0, \cdots, m-1\} \right\},
\]
\[
\tilde{\mathcal{P}} = \left\{ (\tilde{m}, \tilde{p}, n_p, \cdots, n_1) \mid m, n_p, \cdots, n_1 \text{ are positive and even; } \tilde{p} \in \{0, \cdots, \tilde{m}-1\}; p \in \{0, 2, \cdots, 2[\tilde{p}/2]\}; p + n_0 + \cdots + n_1 \leq \tilde{p} \right\}.
\]
Hence (29) is a rearrangement of the series (26), and therefore (29) follows by the absolute convergence of the Dyson series.

Now we are ready to carry out the proof of the inchworm series (20):

**Proof of Theorem 1.** Following the notations in Lemma 2 and 3, we have the following identities:
\[
\tilde{u}^{(0)}(s_\tau) = u^{(0)}(s_\tau, s_m, \cdots, s_{p+1}, s_1)u^{(0)}(s_\tau, s^{(p)}(s_p, s_p)w_{s_\tau}u^{(0)}(s_p, s^{(p-1)}_{s_\tau-1}, s_{p-1})w_{s_p} \cdots w_{s_1}u^{(0)}(s_1, s^{(0)}_1, s_1),
\]
\[
\#(s < t) = \#(s < t) + \#(s^{(p)}(s_p, s_p)w_{s_\tau}u^{(0)}(s_1, s^{(0)}_1, s_1), L(\tilde{q}) = L(q) L(q_p) \cdots L(q_0).
\]
Substituting these equalities to (29) yields
\[
G(s_\tau, s_i) = G^{(p)}(s_\tau, s_\tau)G(s_\tau, s_i) + \sum_{m=2}^{+\infty} \sum_{p=0}^{m-1} \sum_{n_p=0}^{+\infty} \sum_{n_0=0}^{+\infty} \int_{s_\tau}^{s_m} \int_{s_\tau}^{s_{p+1}} \int_{s_\tau}^{s_p} \int_{s_\tau}^{s_1} \int_{s_\tau}^{s} \int_{s_\tau}^{s(s_p, s_p)} \int_{s_\tau}^{s(s_{p-1}, s_{p-1})} \cdots \int_{s_\tau}^{s(s^{(1)}_1, s^{(1)}_1)} \int_{s_\tau}^{s(s^{(0)}_1, s^{(0)}_1)} (-1)^{\#(s < t) + \#(s^{(p)}(s_p, s_p)w_{s_\tau}u^{(0)}(s_1, s^{(0)}_1, s_1), y)(s_p, s_p)w_{s_p} \cdots w_{s_1}u^{(0)}(s_1, s^{(0)}_1, s_1), L(\tilde{q}) = L(q) L(q_p) \cdots L(q_0).
\]

Substituting these equalities to (29) yields
\[
G(s_\tau, s_i) = G^{(p)}(s_\tau, s_\tau)G(s_\tau, s_i) + \sum_{m=2}^{+\infty} \sum_{p=0}^{m-1} \sum_{n_p=0}^{+\infty} \sum_{n_0=0}^{+\infty} \int_{s_\tau}^{s_m} \int_{s_\tau}^{s_{p+1}} \int_{s_\tau}^{s_p} \int_{s_\tau}^{s_1} \int_{s_\tau}^{s} \int_{s_\tau}^{s(s_p, s_p)} \int_{s_\tau}^{s(s_{p-1}, s_{p-1})} \cdots \int_{s_\tau}^{s(s^{(1)}_1, s^{(1)}_1)} \int_{s_\tau}^{s(s^{(0)}_1, s^{(0)}_1)} (-1)^{\#(s < t) + \#(s^{(p)}(s_p, s_p)w_{s_\tau}u^{(0)}(s_1, s^{(0)}_1, s_1), y)(s_p, s_p)w_{s_p} \cdots w_{s_1}u^{(0)}(s_1, s^{(0)}_1, s_1), L(\tilde{q}) = L(q) L(q_p) \cdots L(q_0).
\]
where the integrals with respect to $s^{(k)}$ are interpreted by

$$
\int_{s^{(k)} \subset (a,b)} ds^{(k)} = \int_{b>s^{(k)}_a \ldots > s^{(k)}_1>a} ds^{(k)}_1 \ldots ds^{(k)}_n.
$$  \hfill (32)

Hence the integrals in the second line of (31) is actually the same as the integrals in (29). Due to the absolute convergence, we are allowed to safely interchange sums and integrals, which can give us the following factors:

\begin{align}
\sum_{n_p=0}^{+\infty} \int_{s^{(p)} \subset (s_p,s')} \sum_{q_p \in \mathcal{Q}(s^{(p)})} (-1)^{\#(s^{(p)}<t)} \{ n_s \mathcal{U}^{(0)}(s^p,s,p) \mathcal{L}(q_p) \} ds^{(p)}, \\
\sum_{n_k=0}^{+\infty} \int_{s^{(k)} \subset (s_k,s_{k+1})} \sum_{q_k \in \mathcal{Q}(s^{(k)})} (-1)^{\#(s^{(k)}<t)} \{ n_s \mathcal{U}^{(0)}(s_{k+1},s^{(k)},s_k) \mathcal{L}(q_k) \} ds^{(k)}, \quad k = 0, \ldots, p - 1.
\end{align}

This quantity can be replaced, respectively, by $G(s_p,s^\tau)$ and $G(s_{k+1},s_k)$, $k = 0, \ldots, p - 1$ according to (16). Such replacement turns (31) into

$$
G(s^\tau,s_i) = G^{(0)}(s^\tau,s^\tau)G(s^\tau,s_i) + \sum_{m=2}^{+\infty} \sum_{m \text{ is even}}^{m-1} \int_{s^\tau>s_m>\ldots>s_{p+1}>s^\tau} \int_{s^\tau>s_p>\ldots>s_{1}>s^\tau} \sum_{q \in \mathcal{Q}_{s^\tau}(s)} (-1)^{\#(s^\tau<q)} \times

\mathcal{U}^{(0)}(s^\tau,s_m,\ldots,s_{p+1},s^\tau)G(s^\tau,s_p)W_sG(s_p,s_{p-1})W_s \ldots W_sG(s_2,s_1)W_sG(s_1,s_i) \mathcal{L}(q) \ ds_1 \ldots ds_p \ ds_{p+1} \ldots ds_m.
$$  \hfill (34)

Noting that

$$
\mathcal{U}^{(0)}(s^\tau,s_m,\ldots,s_{p+1},s^\tau)G(s^\tau,s_p)W_sG(s_p,s_{p-1})W_s \ldots W_sG(s_2,s_1)W_sG(s_1,s_i) = G_{s^\tau}(s_1,s_m)W_sG_{s^\tau}(s_m,s_{m-1})W_s \ldots W_sG_{s^\tau}(s_2,s_1)W_sG_{s^\tau}(s_1,s_i),
$$

we see that the integrand in (34) is actually the same as the integrand in (20). Furthermore, the sum over $p$ in (34) can be replaced by $\sum_{p=0}^{m}$, because when $p = m$, the set $\mathcal{Q}_{s^\tau}(s)$ is empty. By doing this, we can apply the identity (24) and obtain the equality (20), which completes the proof of Theorem 1.

4. Integro-differential equations

To better understand the algorithm, we are going to derive the limiting equation of the inchworm algorithm by considering the cases in which $s_1 - s^\tau$ is infinitesimal. Suppose $\Delta t = s_1 - s^\tau$ and $s^\tau$ and $s_1$ are both on the forward branch of the Keldysh contour, i.e. $s^\tau < s_1 < t$. By (7) and (9), we can find that

$$
G^{(0)}_{s^\tau}(s^\tau,s^\tau) = e^{-i\Delta t H_s} = I - i\Delta t H_s + O(\Delta t^2),
$$

$$
\mathcal{U}^{(0)}(s^\tau,s^\tau,s^\tau) = G^{(0)}_{s^\tau}(s^\tau,s^\tau) W_s G^{(0)}_{s^\tau}(s^\tau,s^\tau) = W_s + O(\Delta t).
$$  \hfill (35)

In the inchworm method (34), the domain of the integral

$$
\int_{s^\tau>s_m>\ldots>s_{p+1}>s^\tau} \text{(integrand)} \ ds_{p+1} \ldots ds_{m-1} \ ds_m
$$
has volume $\frac{1}{(m-p)!}\Delta^{m-p}$. Therefore all the terms with $p < m - 1$ are higher order in $\Delta t$, and (34) can be rewritten as

$$G(s_t, s_i) = G^{(0)}(s_t, s_t)G(s_t, s_i) + \sum_{m=2}^{+\infty} \int_{s_t}^{+\infty} \int_{s_t > s_m > \cdots > s_1 > s_i} \sum_{q \in \mathcal{Q}(s)} (-1)^{\#\{s < t\}} \mathcal{L}(q) \times U(s_t, s_m \uparrow) \mathcal{U}(s_t, s_m, s_{m-1}, \cdots, s_1, s_i) \text{d}s_1 \cdots \text{d}s_m - \text{d}s_{m-1} + O(\Delta t^2)$$

where we have used the short hand

$$U(s_t, s_m, s_{m-1}, \cdots, s_1, s_i) = G(s_t, s_m)W_sG(s_m, s_{m-2})W_s \cdots G(s_2, s_1)W_sG(s_1, s_i).$$

Our assumption that $s_1 < s_t < t$ shows that $\#\{s < t\} = m$. Also, by (10) and (18), one sees that

$$\sum_{q \in \mathcal{Q}(s)} \mathcal{L}(q) = \sum_{q \in \mathcal{Q}(s, s_{m-1}, \cdots, s_1)} \mathcal{L}(q) + O(\Delta t) = \sum_{q \in \mathcal{Q}(s, s_{m-1}, \cdots, s_1)} \mathcal{L}(q) + O(\Delta t).$$

Thus on the right-hand side of (36), the first order term of the integrand is actually independent of $s_m$, which can then be integrated out. We write the result by moving $G(s_t, s_i)$ to the left-hand side and divide both sides by $\Delta t$:

$$\frac{G(s_t, s_i) - G(s_t, s_i)}{\Delta t} = -iH_sG(s_t, s_i) - \sum_{m=2}^{+\infty} \int_{s_t > s_m > \cdots > s_1 > s_i} \sum_{q \in \mathcal{Q}(s)} (-1)^{m-1} \mathcal{L}(q)W_sU(s_t, s_m, s_{m-1}, \cdots, s_1, s_i) \text{d}s_1 \cdots \text{d}s_{m-1} + O(\Delta t).$$

Taking the limit as $\Delta t \to 0$ and renaming $m$ to $m + 1$, we obtain the integro-differential equations for the propagator $G(s_t, s_i)$:

$$\frac{\partial G(s_t, s_i)}{\partial s_t} = -iH_sG(s_t, s_i)$$

$$- \sum_{m=1}^{+\infty} \int_{s_t > s_m > \cdots > s_1 > s_i} \sum_{q \in \mathcal{Q}(s)} (-1)^{\#\{s < t\}} \mathcal{L}(q)W_sU(s_t, s_m, s_{m-1}, \cdots, s_1, s_i) \text{d}s_1 \cdots \text{d}s_{m-1}. $$

Here we have again used the fact that all the components of $s$ are less than $t$. The equation (38) gives an integro-differential equation for the full propagator $G(s_t, s_i)$ when $s_t < t$. If $s_t > t$, we can use the same method to derive a similar integro-differential equation:

$$\frac{\partial G(s_t, s_i)}{\partial s_t} = iH_sG(s_t, s_i)$$

$$+ \sum_{m=1}^{+\infty} \int_{s_t > s_m > \cdots > s_1 > s_i} \sum_{q \in \mathcal{Q}(s)} (-1)^{\#\{s < t\}} \mathcal{L}(q)W_sU(s_t, s_m, s_{m-1}, \cdots, s_1, s_i) \text{d}s_1 \cdots \text{d}s_{m-1}. $$

When $s_t = t$, one can see from (13) that $G(\cdot, s_i)$ is discontinuous, and it satisfies

$$\lim_{s_t \to t^-} G(s_t, s_i) = O_s \lim_{s_t \to t^-} G(s_t, s_i).$$

Combing (38) and (39), we get the following theorem:
Theorem 2. When the Dyson series (16) is absolutely convergent in the sense of (19), the full propagator $G(\cdot, \cdot)$ satisfies the integro-differential equation

$$\sgn(s_t - t) \frac{\partial G(s_t, s_i)}{\partial s_t} = iH_s G(s_t, s_i) + \sum_{m=0}^{\infty} i^{m+1} \int_{s_t > s_m > \cdots > s_1 > s_i} \sum_{q \in Q_{(s_m, s_i)}} (-1)^{\#(s_t < q)} W_t \mathcal{U}(s_t, q, s_i) \, ds_1 \cdots ds_m, \quad \forall s_i \in [0, 2t] \setminus \{t\}, \quad s_t \in [s_i, 2t] \setminus \{t\},$$

with the jump condition (40) and the “initial condition” $G(s_t, s_t) = 1$.

Although the integro-differential equation (41) has been derived from the inchworm method, the infinitesimal terms $O(\Delta t)$ or $O(\Delta t^2)$ have not been rigorously verified. Below we are going to provide a rigorous proof of Theorem 2 starting from the definition of $G(\cdot, \cdot)$.

Proof of Theorem 2. We will start the proof by deriving the dynamics of the propagator with a more straightforward method, and then show that the result is equivalent to (41). Again we consider the case $s_t < t$, and take derivative of the definition of $G(13)$ to get

$$\frac{\partial G(s_t, s_i)}{\partial s_t} = -iH_s G(s_t, s_i) - \sum_{m=0}^{\infty} i^{m+1} \int_{s_t > s_m > \cdots > s_1 > s_i} \sum_{q \in Q_{(s_m, s_i)}} (-1)^{\#(s_t < q)} W_t \mathcal{U}(s_t, q, s_i) \, ds_1 \cdots ds_m. \quad (42)$$

The propagator $e^{-i(s_t - s_i)H}$ can be expanded into Dyson series as (3), which turns (42) into

$$\frac{\partial G(s_t, s_i)}{\partial s_t} = -iH_s G(s_t, s_i) - \sum_{m=0}^{\infty} i^{m+1} \int_{s_t > s_m > \cdots > s_1 > s_i} \sum_{q \in Q_{(s_m, s_i)}} (-1)^{\#(s_t < q)} W_t \mathcal{U}(s_t, q, s_i) \, ds_1 \cdots ds_m.$$

Since $G_b(0) G_b(0, s_i) = 0$, we can use the definitions (10) and (9) to simplify the above equation:

$$\frac{\partial G(s_t, s_i)}{\partial s_t} = -iH_s G(s_t, s_i) - \sum_{m=0}^{\infty} i^{m+1} \int_{s_t > s_m > \cdots > s_1 > s_i} \sum_{q \in Q_{(s_m, s_i)}} (-1)^{\#(s_t < q)} W_t \mathcal{U}(s_t, q, s_i) \, ds_1 \cdots ds_m. \quad (43)$$

To see that the above equation is identical to the “inchworm equation” (38), we can mimic (31) and replace the propagators $G(\cdot, \cdot)$ in the integral of (38) by its Dyson series expansion. Following the same way as in the previous section, we obtain a result similar to (26):

$$\frac{\partial G(s_t, s_i)}{\partial s_t} = -iH_s G(s_t, s_i) - \sum_{m=0}^{\infty} \sum_{n_{m}=0}^{\infty} \sum_{n_{m+1}=0}^{\infty} i^{m+1} \int_{s_t > \tilde{s}_m > \cdots > \tilde{s}_1 > s_i} \sum_{q \in Q_{(s_m, s_i)}} (-1)^{\#(s_t < q)} W_t \mathcal{U}(s_t, q, s_i) \, d\tilde{s}_1 \cdots d\tilde{s}_m, \quad (44)$$

where $\tilde{s}_m$ is the $m$th largest variable in $\{s_1, s_2, \ldots, s_t\}$.
where
\[ \tilde{q} = q \cup q_m \cup \cdots \cup q_0, \quad \tilde{m} = m + n_m + \cdots + n_0, \quad \tilde{s} = (s_{\tilde{m}}, \cdots, \tilde{s}_1), \]
and \( s, s^{(m)}, \cdots, s^{(0)} \) can be determined by
\[ \tilde{s} = (s^{(m)}, s_m, s^{(m-1)}, s_{m-1}, \cdots, s_1, s^{(0)}), \quad s = (s_m, \cdots, s_1). \]
Now we need to use the following equivalence of sums:
\[
\sum_{m=1}^{+\infty} \sum_{n_m=0}^{+\infty} \cdots \sum_{n_0=0}^{+\infty} = \sum_{m=0}^{+\infty} \sum_{n_m=0}^{+\infty} \cdots \sum_{n_0=0}^{+\infty}.
\]

Due to the absolute convergence of the Dyson series, the equation (45) becomes
\[
\frac{\partial G(s_1, s_i)}{\partial s_{\uparrow}} = -iH_\uparrow G(s_1, s_i) - \sum_{\tilde{m} \equiv \text{odd}}^{+\infty} \int_{s_{\uparrow} > \tilde{s}_{\uparrow} > \cdots > \tilde{s}_1 > s_1} \sum_{\tilde{m} \equiv \text{even}}^{+\infty} \sum_{n_m=0}^{+\infty} \cdots \sum_{n_0=0}^{+\infty} \sum_{q \in Q(s_1, s)} \sum_{q_m \in Q(s^{(m)})} \cdots \sum_{q_0 \in Q(s^{(0)})} (-1)^#(s < \tilde{s}) W(s, \tilde{s}, q, s) \mathcal{L}(\tilde{q}) \, d\tilde{s}_{\uparrow} \cdots d\tilde{s}_{\uparrow}.
\]

The last step is to verify that the right-hand side of the above equation equals the right-hand side of (44). By comparison, we just need to show that for any given positive odd integer \( \tilde{m} \) and a sequence \( \tilde{s} = (s_{\tilde{m}}, \cdots, s_1) \) satisfying \( s_1 > s_{\tilde{m}} > \cdots > s_1 > s_i \), the following map is a bijection:
\[
\tilde{P} \quad \text{to} \quad Q_{s_1}(\tilde{s})
\]
where
\[
\tilde{P} = \{(m, n_m, \cdots, n_0, q, q_p, \cdots, q_0) \mid p \in \{\tilde{m}, \tilde{m} - 2, \cdots, 1\}; \ n_0 + \cdots + n_m = \tilde{m} - m; \ n_m, \cdots, n_0 \text{ are even} \text{; } q \in Q_c(s_{\uparrow}, \tilde{s}); \ q_m \in Q(s^{(p)}); \ \cdots; \ q_0 \in Q(s^{(0)})\}.
\]

This map can actually be considered as a special case of (27) when \( \tilde{p} = \tilde{m} - 1 \), and hence is one-to-one.

Till now, the inchworm algorithm (41) is confirmed to be identical to (42) when \( s_{\uparrow} < t \), which describes the correct dynamics of the full propagator. The case \( s_{\uparrow} > t \) can be dealt with following the same procedure, which is omitted for conciseness.

Since the equation in the above theorem is firstly derived by setting \( s_{\uparrow} = s_\uparrow \) to be infinitesimal in the inchworm algorithm, the algorithm can actually be considered as an iterative scheme for the equation. From this point of view, we can improve the numerical method by solving (41) with a higher-order scheme. Before introducing the details of the method, we will first present the spin-boson model, and show that it satisfies all the conditions needed for the inchworm method.

5. Spin-boson model

To demonstrate the algorithm in a specific model, we consider the spin-boson model in which the system is a single spin and the bath is given by a large number of harmonic oscillators. In detail, we have
\[
\mathcal{H}_s = \text{span}\{|1\rangle, |2\rangle\}, \quad \mathcal{H}_b = \bigotimes_{l=1}^{L} (L^2(\mathbb{R}^3)),
\]
where $L$ is the number of harmonic oscillators. The corresponding Hamiltonians are

$$H_s = \epsilon \hat{\sigma}_z + \Delta \hat{\sigma}_x, \quad H_b = \sum_{l=1}^{L} \frac{1}{2} (\hat{p}_l^2 + \omega_l^2 \hat{q}_l^2).$$

The notations are described as follows:

- $\epsilon$: energy difference between two spin states.
- $\Delta$: frequency of the spin flipping.
- $\hat{\sigma}_x, \hat{\sigma}_z$: Pauli matrices satisfying $\hat{\sigma}_x |1\rangle = |2\rangle, \hat{\sigma}_x |2\rangle = |1\rangle, \hat{\sigma}_z |1\rangle = |1\rangle, \hat{\sigma}_z |2\rangle = -|2\rangle$.
- $\omega_l$: frequency of the $l$th harmonic oscillator.
- $\hat{q}_l$: position operator for the $l$th harmonic oscillator defined by $\psi(q_1, \cdots, q_L) \mapsto q_l \psi(q_1, \cdots, q_L)$.
- $\hat{p}_l$: momentum operator for the $l$th harmonic oscillator defined by $\psi(q_1, \cdots, q_L) \mapsto -i \nabla_{q_l} \psi(q_1, \cdots, q_L)$.

The coupling between system and bath is assumed to be linear:

$$W = W_s \otimes W_b, \quad W_s = \hat{\sigma}_z, \quad W_b = \sum_{l=1}^{L} c_l \hat{q}_l,$$

where $c_l$ is the coupling intensity between the $l$th harmonic oscillator and the spin. Suppose the initial state of the bath is in the thermal equilibrium with inverse temperature $\beta$, i.e. $\rho_b = Z^{-1} \exp(-\beta H_b)$, and $Z$ is chosen such that $\text{tr}(\rho_b) = 1$. Thus the hypothesis (H1) is fulfilled, and Wick’s theorem (15) holds for

$$B(\tau_1, \tau_2) = \sum_{l=1}^{L} \frac{c_l^2}{2\omega_l} \left[ \coth \left( \frac{\beta \omega_l}{2} \right) \cos \omega_l (\tau_2 - \tau_1) - i \sin \omega_l (\tau_2 - \tau_1) \right]. \quad (48)$$

In order to apply the inchworm algorithm to the spin-boson model, we need to show the absolute convergence (19). By the definition of $U$ (9), we immediately have

$$\|U(s_1, s, s_1)\|_s \leq \|W_s\|_s^m \max\{\|O_s\|_s, 1\} = \|\hat{\sigma}_z\|_s^m \max\{\|O_s\|_s, 1\} = \max\{\|O_s\|_s, 1\}.$$  

And from (48), we see that

$$|B(\tau_1, \tau_2)| \leq \sum_{l=1}^{L} \frac{c_l^2}{2\omega_l} \sqrt{\coth^2 \left( \frac{\beta \omega_l}{2} \right) \cos^2 \omega_l (\tau_2 - \tau_1) + \sin^2 \omega_l (\tau_2 - \tau_1)} \leq \sum_{l=1}^{L} \frac{c_l^2}{2\omega_l} \coth \left( \frac{\beta \omega_l}{2} \right). \quad (49)$$

Let $C_b$ be the right-hand side of the above inequality. Then when $m$ is even,

$$|\mathcal{L}(q)| \leq C_b^{m/2}, \quad \forall q \in \mathcal{Q}(s_m, \cdots, s_1).$$

Since the number of pair sets in $\mathcal{Q}(s_m, \cdots, s_1)$ is $(m - 1)!!$, we have

$$\sum_{m=0}^{\infty} \int_{m \text{ is even}}^{\infty} \|U(0)(s_l, s, s_1)\|_s \sum_{q \in \mathcal{Q}(s)} \mathcal{L}(q) \, ds_l \cdots ds_1 \cdots ds_m \leq \sum_{m=0}^{\infty} \int_{m \text{ is even}}^{\infty} \max\{\|O_s\|_s, 1\} \cdot (m - 1)!! C_b^{m/2} ds_l \cdots ds_m \quad (50)$$

$$= \max\{\|O_s\|_s, 1\} \sum_{m=0}^{\infty} \frac{(s_l - s_1)^m}{m!!} C_b^{m/2} = \max\{\|O_s\|_s, 1\} \exp \left( \frac{C_b (s_l - s_1)^2}{2} \right).$$
Since $\mathcal{H}$ is a finite-dimensional space, the observable $O_s$ is always a bounded operator. Therefore the right-hand side of the above equation is finite, which shows the absolute convergence. In the spin-boson model, people are usually interested in the population of the spin on each of the two spin states, meaning that we can take $O_s = \hat{\sigma}_z$.

When the Dyson series expansion (16) is directly used in the Monte Carlo simulation, the fast growth of the variance as $s_t - s_i$ increases causes great numerical difficulties. Similar to (50), the expectation of $\|U(0)(s_t, s_s, s_i)L(q)\|^2$ can be estimated by

$$
\sum_{m=0}^{\infty} \int_{s_t > s_m > \cdots > s_1 > s_i} \sum_{q \in \mathcal{Q}(s)} \|U(0)(s_t, s_s, s_i)\|^2 \|L(q)\|^2 \, ds_1 \cdots ds_m
$$

$$
\leq \sum_{m=0}^{\infty} \int_{s_t > s_m > \cdots > s_1 > s_i} (m-1)!(\max\{\|O_s\|, 1\})^2 C_m^2 \| s\| \, ds_1 \cdots ds_m
$$

$$
= (\max\{\|O_s\|, 1\})^2 \exp(C^2(s_t - s_i)^2).
$$

Since $\|G(s_t, s_i)\| \leq \max\{\|O_s\|, 1\}$, the growth of the variance is characterized by

$$
\exp(C^2(s_t - s_i)^2) - 1.
$$

This is the well-known “dynamical sign problem” in the quantum Monte Carlo simulations. The inchworm method relieves the dynamical sign problem by lumping a number of samples based on the simulation results of shorter bold lines, which pushes the simulation time significantly longer.

Different from the inchworm methods presented in [2], which directly applies the Monte Carlo method to (20), we will design our numerical method based on the integro-differential equation (41), and apply the idea of classical Runge-Kutta methods for temporal discretization to solve the full propagators. As will be presented, this can both enhance the numerical efficiency and simplify the implementation. The algorithm will be detailed in the following section.

6. Numerical method

In order to find the full propagator $G(s_t, s_i)$ numerically for all $s_i \in [0, 2t]$ and $s_t \in [s_i, 2t]$. Below we are going to develop a second-order method in analogous to Heun’s method for general ordinary differential equations. For the general initial value problem

$$
u'(t) = f(t, u(t)), \quad t > 0$$

with initial condition $u(0) = u_0$, Heun’s method reads

$$
U_k^* = U_{k-1} + \Delta t f(t_{k-1}, U_{k-1}),
$$

$$
U_k = \frac{1}{2}(U_{k-1} + U_k^*) + \frac{1}{2} \Delta t f(t_k, U_k^*),
$$

where $\Delta t$ is the time step, $t_k = t_{k-1} + \Delta t$, and $U_k$ is the numerical approximation of $u(t_k)$. The method is second-order if the solution is third-order continuously differentiable. In our case, the full propagator $G(\cdot, \cdot)$ is known to be discontinuous on the line segments $[0, t] \times \{t\}$ and $\{t\} \times [0, t]$ due to the presence of the observable $O_s$. Therefore in order to keep the second-order convergence rate, special care needs to be taken for these discontinuities.

In our implementation, we take a uniform time step $\Delta t = t/N$, and compute the numerical solutions $G(s_t, s_i)$ only when $s_t$ and $s_i$ are multiples of $\Delta t$. This corresponds to a two-dimensional triangular mesh.
shown in Figure 4 (for $N = 5$). Let $t_k = k \Delta t$ and $G_{jk}^{\Delta t}$ be the numerical approximation of $G(t_j, t_k)$. Due to the discontinuity, when $j = N$ (green line in Figure 4) or $k = N$ (blue line in Figure 4), $G_{jk}^{\Delta t}$ is considered to be multiple-valued, and we will use the notation $j, k = N^+, N^-$ to define the left and right limits. Precisely,

- For $k < N$, $G_{N-k}^{\Delta t}$ and $G_{N+k}^{\Delta t}$ are respectively the approximation of $\lim_{s \to t^-} G(s, k \Delta t)$ and $\lim_{s \to t^+} G(s, k \Delta t)$.
- For $j > N$, $G_{N-j}^{\Delta t}$ and $G_{N+j}^{\Delta t}$ are respectively the approximation of $\lim_{s \to t^-} G(j \Delta t, s)$ and $\lim_{s \to t^+} G(j \Delta t, s)$.
- $G_{N-N}^{\Delta t} = G_{N+N}^{\Delta t} = \text{Id}$ and $G_{N+N}^{\Delta t} = O_s$.

Now we are ready to sketch our numerical algorithm. In general, the values of $G_{jk}^{\Delta t}$ are obtained in the following order:

$$
G_0^{\Delta t},
G_{11}^{\Delta t}, G_{10}^{\Delta t},
\ldots
\ldots
G_{N-1,N-1}^{\Delta t}, \ldots, G_{N-1,0}^{\Delta t},
G_{N-N-1}^{\Delta t}, \ldots, G_{N-0}^{\Delta t},
G_{N,N+1}^{\Delta t}, G_{N,N-1}^{\Delta t}, G_{N-1,N}^{\Delta t}, \ldots, G_{N-0}^{\Delta t},
G_{N+1,N+1}^{\Delta t}, G_{N+1,N}^{\Delta t}, G_{N+1,N-1}^{\Delta t}, \ldots, G_{N+1,0}^{\Delta t},
\ldots
\ldots
\ldots
\ldots
G_{2N,2N}^{\Delta t}, \ldots, G_{2N,N+1}^{\Delta t}, G_{2N,N}^{\Delta t}, G_{2N,N-1}^{\Delta t}, \ldots, G_{2N,0}^{\Delta t}.
$$

This corresponds to computing the values of $G(\cdot, \cdot)$ column by column in Figure 4 with the green line and the blue line split to two lines due to the discontinuity. We first list out the three special cases:

- If $j = k$ ($N^+$ is considered to be not equal to $N^-$), we set $G_{jk}^{\Delta t}$ to be $\text{Id}$. This corresponds to the nodes on the red line in Figure 4.
- If $j = N^+$ and $k \neq j$, we set $G_{jk}^{\Delta t}$ to be $O_s G_{N-k}^{\Delta t}$. This is applied to the nodes on the green line in Figure 4.
- If $k = N^-$ and $j \neq k$, we set $G_{jk}^{\Delta t}$ to be $G_{jN}^{\Delta t} O_s$. This is applied to the nodes on the blue line in Figure 4.
For all other cases, we follow Heun’s method and find the values of \( G_{jk}^{\Delta t} \) as follows:

1. Let

\[
G_{jk}^* = G_{jk-1,k}^{\Delta t} + \text{sgn}(t_j - t)\Delta t \left[ iH_s G_{jk-1,k}^{\Delta t} + \sum_{m=1}^{\infty} \int_{t_{j-1} > s_m > \cdots > s_1 > t_k} \sum_{q \in \mathbb{Q}(t_j, s)} (-1)^{\# \{ s < t \}} \right.
\]

\[
\times \mathcal{L}(q) W_s G_I(t_{j-1}, s_m) W_s G_I(s_m, s_{m-1}) W_s \cdots W_s G_I(s_2, s_1) W_s G_I(s_1, t_k) \, ds_1 \cdots ds_m \right] ,
\]

(53)

where \( G_I(\cdot, \cdot) \) is the interpolated function satisfying

\[
G_I(t_j, t_k) = G_{jk}^{\Delta t} , \quad \text{for all integers } j', k' \text{ satisfying } k \leq k' \leq j - 1 .
\]

(54)

In our implementation, piecewise linear interpolation is adopted, and the function \( G_I(\cdot, \cdot) \) is linear on each triangle in Figure 4.

2. Set \( G_{jk}^{\Delta t} \) to be

\[
\frac{1}{2} G_{jk-1,k}^{\Delta t} + \frac{1}{2} G_{jk}^* + \frac{1}{2} \text{sgn}(t_j - t)\Delta t \left[ iH_s G_{jk}^* + \sum_{m=1}^{\infty} \int_{t_{j-1} > s_m > \cdots > s_1 > t_k} \sum_{q \in \mathbb{Q}(t_j, s)} (-1)^{\# \{ s < t \}} \right.
\]

\[
\times \mathcal{L}(q) W_s G_I^*(t_j, s_m) W_s G_I^*(s_m, s_{m-1}) W_s \cdots W_s G_I^*(s_2, s_1) W_s G_I^*(s_1, t_k) \, ds_1 \cdots ds_m \right] ,
\]

(55)

where \( G_I^*(\cdot, \cdot) \) is the interpolated function satisfying \( G_I^*(t_j, t_k) = G_{jk}^* \) and

\[
G_I^*(t_{j'}, t_{k'}) = G_{jk'}^{\Delta t} , \quad \text{for all integers } j', k' \text{ satisfying } k \leq k' \leq j \text{ and } (j', k') \neq (j, k) .
\]

(56)

Again, the same piecewise linear interpolation is adopted.

When applying (53) and (55), we follow the rules as below:

- When \( j = N^- \), \( j - 1 \) is interpreted as \( N - 1 \); when \( j = N + 1 \), \( j - 1 \) is interpreted as \( N^+ \).
- \( \text{sgn}(t_{N^-} - t) = -1 \).
- The interpolation of \( G_I \) and \( G_I^* \) should respect such discontinuities. Therefore when \( k = N \) or \( l = N \), the equation (54) should be interpreted by

\[
\lim_{s \to t^L} \lim_{s' \to t^L} G_I(t_k, s) = G_{N^+}^{\Delta t}, \quad \lim_{s \to t^L} \lim_{s' \to t^L} G_I(s, t_l) = G_{N^+}^{\Delta t},
\]

\[
\lim_{s \to t^L} \lim_{s' \to t^-} G_I(s, \tilde{s}) = \lim_{s \to t^-} \lim_{s' \to t^-} G_I(s, \tilde{s}) = \text{Id}, \quad \lim_{s \to t^L} \lim_{s' \to t^-} G_I(s, \tilde{s}) = O_s.
\]

The equation (56) should be similarly interpreted. Especially, when \( j = N^+ \), the term \( G_I(t_{N^+}, s_m) \) in the integral should be interpreted as \( O_s G_I(t_{N^-}, s_m) \).

The order of computation (52) ensures that all information needed in the two stages has been obtained beforehand.

Now we consider the numerical computation of the infinite sums in (53) and (55). The numerical results in [3] show that in the original inchworm method (20), we can truncate the series at \( m = M \) for some
positive even integer $M$, and obtain results with sufficient quality. In our method, the integer $M$ needs to be odd and we perform the similar truncation by replacing the infinite sums in (53) and (55) with

$$
\sum_{m=1}^{M} \int_{t_{m+1} > s_{m} > \cdots > s_{1} > t_{k}} \sum_{q \in Q_{c}(t_{1}, s)} S(q, s) \, ds_{1} \cdots ds_{m},
$$

(57)

where $S(q, s)$ is the summand in (53) or (55), and $t_{k}$ is the corresponding $t_{j}$ or $t_{j+1}$. For every $m$, the high-dimensional integral over $s$ and the sum over $q$ are evaluated using the Monte Carlo method. The sampling of $s$ can be done by sampling a uniform distribution in $[t_{k}, t_{j}]^m$, and then sort the time points. As for $q$, we first list out all the elements in $Q_{c}(t_{1}, s)$, and then pick a random one for each sample. Obviously the number of pair sets in $Q_{c}(t_{1}, s)$ depends only on the value of $m$, and it has been given in [29] that this number can be evaluated by

$$
N_{1} = 1, \quad N_{m} = \frac{m - 1}{2} \sum_{j=1}^{m-2} N_{j} N_{m-1-j}.
$$

In [30], it is proven that $N_{m}$ grows asymptotically as $m!!$. In our numerical experiments, we are only concerned about a small $m$, and therefore such a strategy is feasible.

As the end of this section, we will discuss briefly the difference between this numerical method and the original inchworm method based on the Monte Carlo simulation of (20). Both numerical methods are time-stepping methods which can be regarded as a numerical approximation of

$$
G(s_{t_{1}}, s_{1}) = G(s_{t_{1}}, s_{1}) + \int_{s_{t_{1}}}^{s} \text{RHS}(s) \, ds,
$$

(58)

where $\text{RHS}(s)$ is the right-hand side of (38) or (39), with $s_{t_{1}}$ replaced by $s$. From the diagrammatic equation (22), one can see that the original inchworm algorithm in principle allows to put an arbitrary number of points between $s_{t}$ and $s_{t_{1}}$ to evaluate the integral in (58), and all these points are stochastic, which implies that in the time-stepping process, the integral between two time steps is approximated using a Monte Carlo simulation; while in our method, this is replaced by a numerical integration of second-order convergence, which can be expected to be more efficient. One possible benefit of the Monte Carlo integration from $s_{t}$ to $s_{t_{1}}$ is that when a sufficient number of samples are used, this integral can be evaluated up to arbitrary precision; while in the Runge-Kutta integration, an error of $O((s_{t} - s_{t_{1}})^{m})$, where $\alpha$ depends on the order of the method, is always there. However, this does not mean that the numerical error will vanish as the number of samples increases in the original inchworm algorithm, since another part of numerical error comes from the approximation of $\text{RHS}(s)$, where the interpolation on the lattice shown in Figure 4 is inevitable in both methods. Such error will only vanish as the grid size tends to zero, but will not vanish as the number of samples tends to infinity. Usually, it is sufficient to match the order of accuracy for evaluating the integral in (58) and the interpolation. This can be achieved easier and cheaper by using the Runge-Kutta integration strategy. Another benefit of this new method is that we can draw the samples for $q$ more easily. Consider a diagram with four points. The set $Q_{c}(s_{1}, s_{2}, s_{3}, s_{4})$ contains only one set of pairs $\{(s_{1}, s_{3}), (s_{2}, s_{4})\}$, while the inchworm proper set of pairs has 9 possibilities:

- $Q_{s_{1}}(s_{1}, s_{2}, s_{3}, s_{4}) = \{(s_{1}, s_{3}), (s_{2}, s_{4})\}$ if $s_{3} < s_{t} \leq s_{4}$;
- $Q_{s_{2}}(s_{1}, s_{2}, s_{3}, s_{4}) = \{(s_{1}, s_{3}), (s_{2}, s_{4})\}, \{(s_{1}, s_{4}), (s_{2}, s_{3})\}$ if $s_{2} < s_{t} \leq s_{3}$;
- $Q_{s_{3}}(s_{1}, s_{2}, s_{3}, s_{4}) = \{(s_{1}, s_{3}), (s_{2}, s_{4})\}, \{(s_{1}, s_{4}), (s_{2}, s_{3})\}, \{(s_{1}, s_{2}), (s_{3}, s_{4})\}$ if $s_{1} < s_{t} \leq s_{2}$;
- $Q_{s_{4}}(s_{1}, s_{2}, s_{3}, s_{4}) = \{(s_{1}, s_{3}), (s_{2}, s_{4})\}, \{(s_{1}, s_{4}), (s_{2}, s_{3})\}, \{(s_{1}, s_{2}), (s_{3}, s_{4})\}$ if $s_{t} \leq s_{1}$.

For the more time points, finding out all the inchworm proper pair sets is even harder than just finding out $Q_{c}(s)$. This indicates that the implementation of the new method is easier.
7. Numerical experiments

In our numerical experiments, we consider the spin-Boson model with a bath with Ohmic spectral density, for which the frequency $\omega_l$ are distributed in $[0, \omega_{\text{max}}]$ as introduced in [20]:

$$\omega_l = \omega_c \ln \left( 1 - \frac{l}{L} [1 - \exp(\omega_{\text{max}}/\omega_c)] \right), \quad l = 1, \ldots, L,$$

where $\omega_c$ is the primary frequency to be specified later. The coupling intensity $c_l$ is

$$c_l = \omega_l \sqrt{\frac{\xi \omega_c}{L} [1 - \exp(\omega_{\text{max}}/\omega_c)]}, \quad l = 1, \ldots, L,$$

with $\xi$ being the Kondo parameter. To compare our results with reference solutions, we adopt the parameters provided in [14] where $L = 200$ and $\beta = 5\Delta^{-1}$. Different settings of bias, coupling intensity and nonadiabaticity will be considered in our experiments. In all the numerical tests, the maximum frequency $\omega_{\text{max}}$ is set to be $4\omega_c$, and the time step is chosen to be $\Delta t = 0.1$ if not otherwise specified. Numerical results obtained by the QuAPI method [22, 23] will be used as reference solutions.

7.1. Experiments with changing bias

We first choose $\omega_c = 2.5\Delta$ and $\xi = 0.2$, for which the amplitude of the bath correlation function $B(\cdot, \cdot)$ is plotted in Figure 5. In our implementation, we precompute the values of $B(\cdot, \cdot)$ up to a very high precision on a very fine grid, and in the inchworm algorithm, we retrieve its value by linear interpolation when necessary. The numerical results for $\epsilon = 0$, $\Delta$ and $2\Delta$ are given in Figure 6. It turns out that $M = 3$ in (57) can already provide satisfying numerical results up to time $t = 5\Delta^{-1}$, and the general behavior of the observable has been well captured by the results of $M = 1$.

![Figure 5: The amplitude of the bath correlation function $B(\cdot, \cdot)$](image)

The order of convergence of our numerical method is also verified using the test case with $\epsilon = 0$. In general, the stochastic error and the “deterministic error” caused by Runge-Kutta and interpolation cannot be separated. In order to cast off the stochastic error, we only consider the truncation $M = 1$, for which the sum (57) contains only a one-dimensional integral, and thus can be evaluated by the composite mid-point rule. As a result, the whole scheme is deterministic and the order of convergence is still expected to be $O(\Delta t^2)$. The reference solution is obtained by choosing $\Delta t = 1/320\Delta^{-1}$, and the numerical error of $\langle \hat{\sigma}_z(t) \rangle$ is tabulated in Table 1, which clearly shows the numerical error is second order of $\Delta t$.

7.2. Experiments with changing coupling intensity

Now we fix the values of $\omega_c$ and $\epsilon$ to be $2.5\Delta$ and $\Delta$ respectively, and consider the coupling intensity $\xi = 0.1$ and $\xi = 0.2$. It can be expected that the convergence of the inchworm series gets slower when $\xi$ increases, as is confirmed in our numerical tests shown in Figure 7. In spite of this, very good matching with the QuAPI results can still be obtained using $M = 1$ for both parameters, and further improvement is indeed achieved by using $M = 3$.
Figure 6: Evolution of $\langle \sigma_z(t) \rangle$ under different settings of the electronic bias (from left to right: $\epsilon = 0$, $\epsilon = \Delta$ and $\epsilon = 2\Delta$), with other parameters $\omega_c = 2.5\Delta$ and $\xi = 0.2$. QuAPI results are plotted as a reference.

<table>
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<th>$h$</th>
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Table 1: Numerical error of $\langle \sigma_z(t) \rangle$ and the order of accuracy

As a reference, the numerical results for a direct summation of the truncated Dyson series

$$G(s_r, s_i) \approx \sum_{m=0}^{M} \int s_{r_1} > s_{m_1} > \cdots > s_{r_1} > s_i \sum_{q \in \mathcal{Q}(s)} (-1)^{\#(s < t)} i^{m |q|} L(q) d s_1 \cdots d s_m,$$

are also provided in Figure 7 with label “bare dQMC”. The results show that the convergence of the Dyson series (16) is much slower than the inchworm series, and therefore require a much larger $M$ to get the same quality of the solutions for large $t$.

Figure 7: Evolution of $\langle \sigma_z(t) \rangle$ under different settings of the coupling intensity (left: $\xi = 0.1$, right: $\xi = 0.2$), with other parameters $\omega_c = 2.5\Delta$ and $\xi = \Delta$. QuAPI results are plotted as a reference.

### 7.3. Experiments with changing nonadiabaticity

To show the role of the parameter $\omega_c$, we fix $\xi$ to be 0.4 and $\epsilon$ to be $\Delta$, and consider the cases $\omega_c = 0.25\Delta$ and $\omega_c = 2.5\Delta$. Since the upper bound of $|B(\tau_1, \tau_2)|$ in (49) gets larger when $\omega_c$ increases, we can expect slower convergence in terms of $M$ for larger $\omega_c$. The evolution of the observable is plotted in Figure 8. In the right figure, due to the large value of both $\xi$ and $\omega_c$, even when $M = 3$ is used, some discrepancy between our results and QuAPI can still be observed.
8. Summary

We have studied the inchworm Monte Carlo method introduced in [4, 2], and proven rigorously that the method converges to the solution of the open quantum system whose bath configuration satisfies Wick’s theorem. By assuming the iterative step length to be infinitesimal, we have derived a new continuous model for the open quantum system, and have proposed an improvement of the inchworm method to achieve better numerical efficiency and simpler implementation. In this new method, the stochastic error and the deterministic error are coupled together, and one needs to apply numerical analysis to find optimal combinations of parameters. This will be left for future works.

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