

# CT-1/2-HYB-QMC: a new solver for time-dependent Anderson impurity

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## Motivation

Continuous-time quantum Monte Carlo methods (CT-QMC) are widely used to solve impurity models in equilibrium [1]. However, they suffer from an inherent dynamical sign problem while treating the nonequilibrium real-time dynamics [2, 3, 4]. Here we propose a method that alleviates the dynamical sign problem in case of the single-orbital Anderson impurity model (AIM).

## Model and method

The Hamiltonian of the time-dependent single-orbital AIM in a setup with two leads reads

$$H(t) = H_{\text{loc}}(t) + \sum_{\sigma} [H_{\text{bath},\sigma}(t) + H_{\text{hyb},\sigma}(t)]$$

$$H_{\text{loc}}(t) = \sum_{\sigma} \mathcal{E}_{d\sigma}(t) d_{\sigma}^{\dagger} d_{\sigma} + U(t) d_{\uparrow}^{\dagger} d_{\uparrow} d_{\downarrow}^{\dagger} d_{\downarrow}$$

$$H_{\text{bath},\sigma}(t) = \sum_{\alpha=-1,1} \sum_p^{N/2} \left( \varepsilon_{p\sigma}(t) + \alpha \frac{\mathcal{V}(t)}{2} \right) c_{\alpha p \sigma}^{\dagger} c_{\alpha p \sigma}$$

$$H_{\text{hyb},\sigma}(t) = \sum_{\alpha=-1,1} \sum_p (V_{p\sigma}(t) c_{\alpha p \sigma}^{\dagger} d_{\sigma} + \text{h.c.}).$$

The generating functional for all the time-dependent observables of the interest is the dynamical partition function  $\mathcal{Z}(\beta, t_{\text{max}})$

$$\mathcal{Z}(\beta, t_{\text{max}}) = \text{Tr} \left( e^{-\beta H(0)} \mathcal{T}_{-} e^{-i \int_0^{\beta} dt H(t)} \mathcal{T}_{+} e^{-i \int_0^{t_{\text{max}}} dt H(t)} \right)$$

$$\equiv \text{Tr} \left( \mathcal{T}_{\mathcal{C}} e^{-i \int_{\mathcal{C}} ds H(s)} \right) = \text{Tr} \left( \mathcal{T}_{\mathcal{C}} e^{-i \int_{\mathcal{C}} ds (H - H_{\text{hyb},\downarrow}(s))} e^{-i \int_{\mathcal{C}} ds H_{\text{hyb},\downarrow}(s)} \right).$$

In contrast to the standard CT-HYB-QMC algorithm we expand  $\mathcal{Z}$  perturbatively in the powers of  $H_{\text{hyb},\downarrow}(s)$  only

$$\mathcal{Z}(\beta, t_{\text{max}}) = \sum_{k=0}^{\infty} (-1)^k \int_{0+}^{-i\beta} ds_1 \cdots \int_{s_{k-1}}^{-i\beta} ds_k \int_{0+}^{-i\beta} ds'_1 \cdots \int_{s'_{k-1}}^{-i\beta} ds'_k$$

$$w_{\text{loc}+\text{bath}\uparrow}(\{s_m\}, \{s'_n\}) w_{\text{bath}\downarrow}(\{s_m\}, \{s'_n\}).$$

Terms of the expansion are then stochastically sampled within a Monte Carlo simulation. Each term's contribution may be split into the dressed local weight  $w_{\text{loc}+\text{bath}\uparrow}$  and the spin-down bath weight  $w_{\text{bath}\downarrow}$ , which is evaluated easily due to Wick's theorem

$$w_{\text{bath}\downarrow} = Z_{\text{bath}\downarrow} i^k \det \left[ \sum_{\alpha p} V_{p\downarrow}^*(s_m) g_{\alpha p \downarrow}(s_m, s'_n) V_{p\downarrow}(s'_n) \right]_{m,n=1,\dots,k},$$

where  $g_{\alpha p \downarrow}(s, s')$  is a non-interacting bath Green function.

## Evaluation of the dressed local weight - discretized bath

The novel aspect is the evaluation of the dressed local weight, which incorporates both local and spin-up bath degrees of freedom. This auxiliary system is governed by the Hamiltonian  $H_0 = H_{\text{loc}} + H_{\text{bath}\uparrow} + H_{\text{hyb}\uparrow}$  and is effectively non-interacting since  $H_0$  conserves  $n_{\downarrow} = \langle d_{\downarrow}^{\dagger} d_{\downarrow} \rangle$ . Using the well-known result from determinant QMC [5] we get (for  $s_k > s'_k$ )

$$w_{\text{loc}+\text{bath}\uparrow} = \text{Tr}_{\text{loc},\text{bath}\uparrow} \left( e^{-\beta H_0(0)} d_{\downarrow}^{\dagger}(s_k) d_{\downarrow}(s'_k) \cdots d_{\downarrow}^{\dagger}(s_1) d_{\downarrow}(s'_1) \right)$$

$$= e^{-i \int_{s_k}^{-i\beta} ds \mathcal{E}_{d\downarrow}(s)} e^{-i \int_{s_{k-1}}^{s'_k} ds \mathcal{E}_{d\downarrow}(s)} \cdots e^{-i \int_{0+}^{s'_1} ds \mathcal{E}_{d\downarrow}(s)}$$

$$\det \left[ \mathbb{1} + u_{\downarrow}(-i\beta, s_k) u_0(s_k, s'_k) u_1(s'_k, s_{k-1}) \cdots u_0(s_1, s'_1) u_1(s'_1, 0+) \right],$$

where  $u_{n_{\downarrow}}$  is a single-particle time-evolution matrix

$$u_{n_{\downarrow}}(s, s') = \mathcal{T}_{\mathcal{C}} \exp \left( -i \int_{s'}^s d\bar{s} \begin{bmatrix} \mathcal{E}_{d\uparrow}(\bar{s}) + U(\bar{s}) n_{\downarrow} & V_{\uparrow}^*(\bar{s}) & \cdots & V_{\uparrow}^*(\bar{s}) \\ V_{\downarrow}(\bar{s}) & \varepsilon_{1\downarrow}(\bar{s}) & \cdots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ V_k(\bar{s}) & 0 & \cdots & \varepsilon_{k\downarrow}(\bar{s}) \end{bmatrix} \right).$$

The computational cost is  $\mathcal{O}(N^3)$ , where  $N$  is the size of the bath.

## Evaluation of the dressed local weight - discretized time

Formally one can integrate out the bath levels and work directly with the hybridization function

$$\Delta_{\uparrow}(s, s') = \sum_{\alpha p} V_{p\uparrow}^*(s) g_{\alpha p \uparrow}(s, s') V_{p\uparrow}(s')$$

The impurity spin-up Green function for a given MC configuration  $n_{\downarrow}(s)$  satisfies Dyson equation

$$G_{\uparrow}(s, s') = G_{0\uparrow}(s, s') + (G_{0\uparrow} \circ \Delta_{\uparrow} \circ G_{\uparrow})(s, s'),$$

where

$$G_{0\uparrow}^{-1}(s, s') = \left[ i \frac{\partial}{\partial s} - \mathcal{E}_{d\uparrow}(s) - U(s) n_{\downarrow}(s) \right] \delta(s, s').$$

One can discretize contour times  $s_i = \sum_{j=0}^{i-1} (\Delta s)_j$  and obtain the local weight as a determinant in the contour-time space in the limit  $\Delta s \rightarrow 0$

$$w_{\text{loc}+\text{bath}\uparrow} = \det [iG_{\uparrow}(s_i, s_j)]_{i,j} = \left( 1 + e^{-i \int_{0+}^{-i\beta} ds [\mathcal{E}_{d\uparrow}(s) + U(s) n_{\downarrow}(s)]} \right) \det [\delta_{ij} - (G_{0\uparrow} \circ \Delta_{\uparrow})(s_i, s_j)]_{i,j}.$$

The computational cost is  $\mathcal{O}(N_{\text{times}}^3)$  but does not depend on  $N$ . In principle, one can also introduce retarded interaction  $U_{\uparrow\downarrow}(s, s')$  at no additional cost.

## References

- [1] E. Gull, A. J. Millis, A. I. Lichtenstein, A. N. Rubtsov, M. Troyer, P. Werner, Continuous-time Monte Carlo methods for quantum impurity models, *Rev. Mod. Phys.* **83**, 349 (2011)
- [2] L. Mühlbacher, E. Rabani, Real-Time Path Integral Approach to Nonequilibrium Many-Body Quantum Systems, *Phys. Rev. Lett.* **100**, 176403 (2008)
- [3] P. Werner, T. Oka, A. J. Millis, Diagrammatic Monte Carlo simulation of nonequilibrium systems, *Phys. Rev. B* **79**, 035320 (2009)
- [4] M. Schiró, Real-time dynamics in quantum impurity models with diagrammatic Monte Carlo, *Phys. Rev. B* **81**, 085126 (2010)
- [5] R. Blankenbecler, D. J. Scalapino, R. L. Sugar, Monte Carlo calculations of coupled boson-fermion systems. I, *Phys. Rev. D* **24**, 2278 (1981)

## Measurement of current - discretized bath

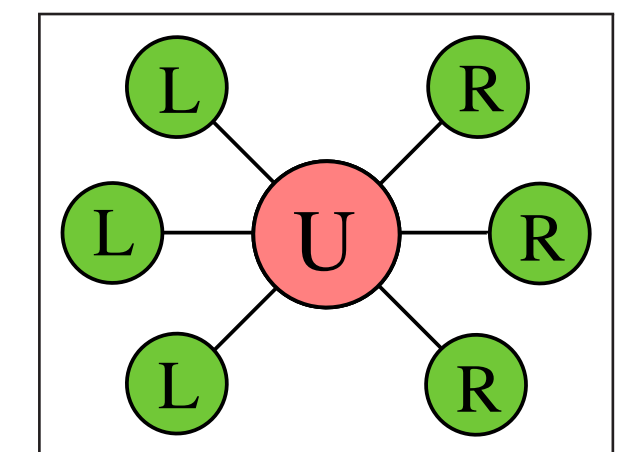
The current to the left ( $\alpha = -1 \equiv L$ ) lead is given by

$$I_{\sigma}(t) = -2 \text{Im} \sum_p V_{p\sigma}(t) \langle c_{Lp\sigma}^{\dagger} d_{\sigma} \rangle(t).$$

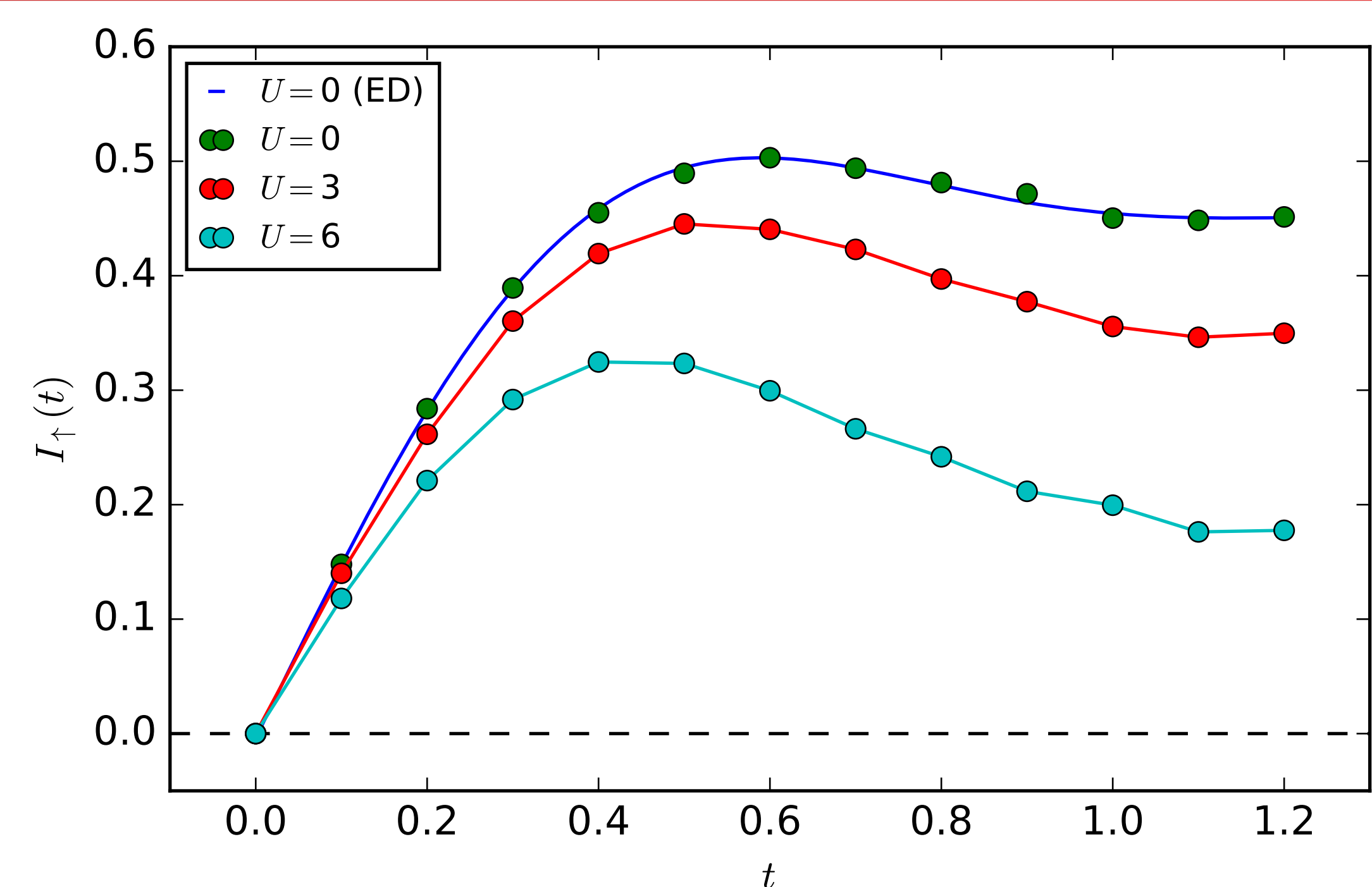
Since the spin-up subsystem is solved explicitly for each MC configuration, one has a direct access to the relevant expectation values

$$\langle c_{\alpha p \uparrow}^{\dagger} d_{\uparrow} \rangle(t) = \langle - [\mathbb{1} + u_{0/1}(t, \tilde{s}_i) \cdots u_1(\tilde{s}_1, 0+) u_1(-i\beta, \tilde{s}_{2k}) \cdots u_{0/1}(\tilde{s}_{i+1}, t)]_{\alpha p, d}^{-1} \rangle_{\text{QMC}}.$$

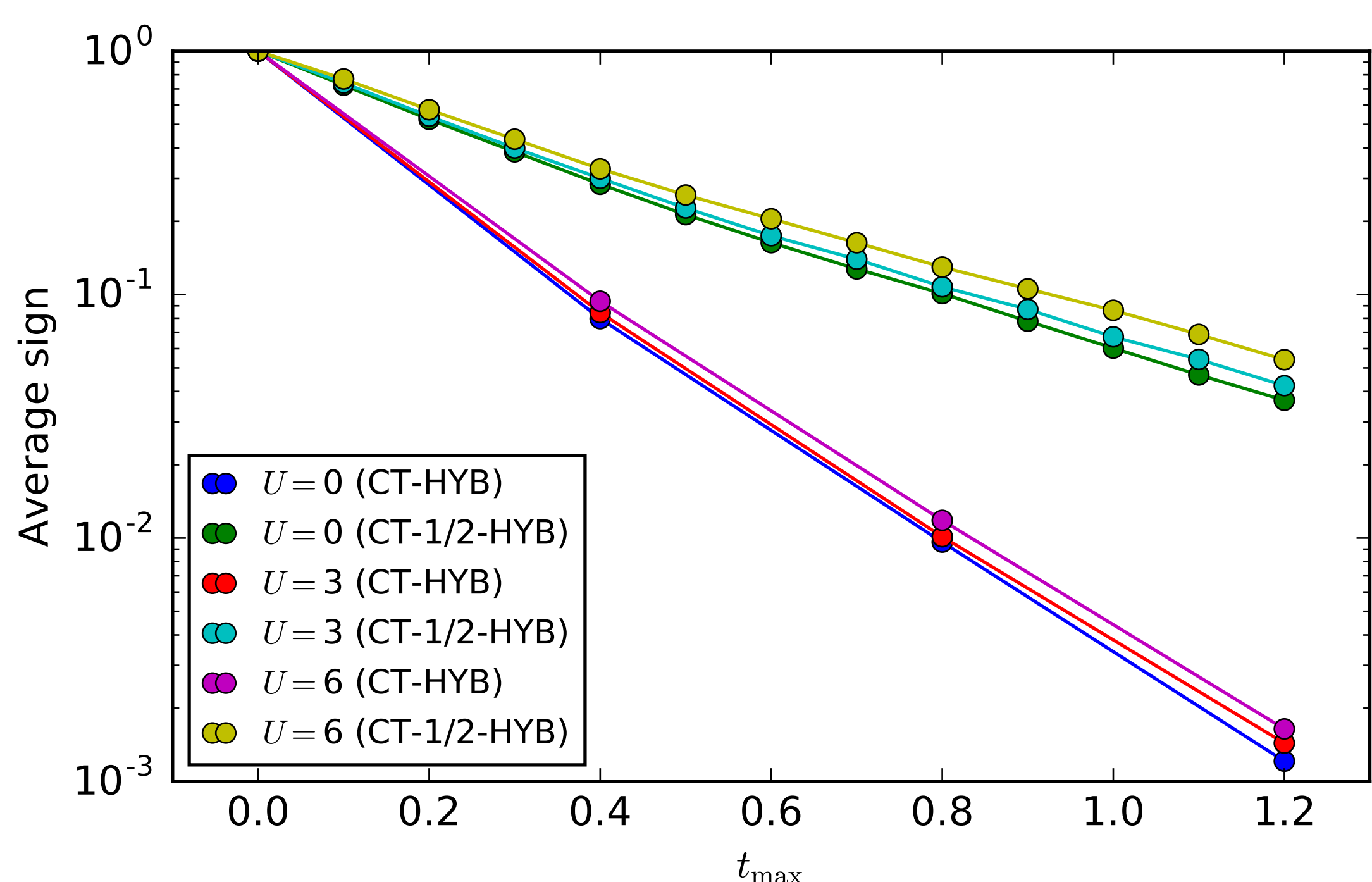
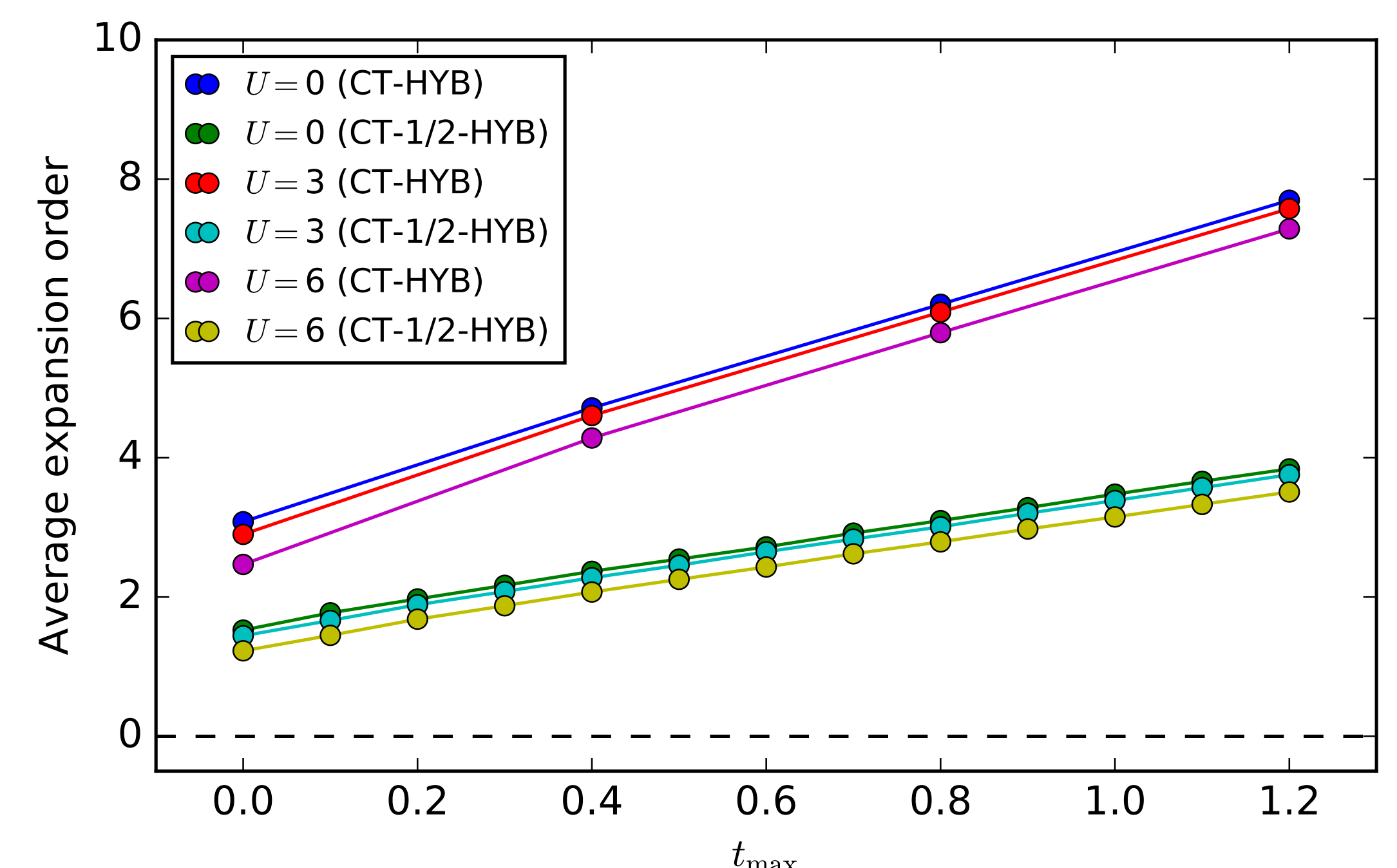
Here we consider the voltage quench from  $\mathcal{V}(0) = 0$  to  $\mathcal{V}(t > 0) = 4$  for  $U(t) = 3$  and 6. Each lead is composed of  $N/2 = 15$  bath states  $\varepsilon_{\alpha p \sigma}$  equidistantly spaced around 0 in the interval of half-width  $D = 3$ . We use the integrated impurity-bath coupling  $\Gamma = \frac{\pi N |V_p|^2}{4D}$  as energy unit. The inverse temperature  $\beta = 2$ .



## Current from CT-1/2-HYB



## Average expansion order



## Conclusions

- CT-1/2-HYB-QMC allows one to solve the finite-size single-orbital Anderson impurity model exactly on twice as long time scales as within the standard CT-HYB-QMC.
- The average expansion order is reduced by a factor of 2, which decreases the computational burden of the bath weight evaluations with respect to CT-HYB by 4.
- The computational complexity of the *discretized bath* algorithm is  $\mathcal{O}(N^3)$  as opposed to  $\mathcal{O}(e^N)$  for exact-diagonalization methods.
- The computational complexity of the *discretized time* algorithm is  $\mathcal{O}(N_{\text{times}}^3)$  and the extrapolation  $\Delta s \rightarrow 0$  shall be performed.
- CT-1/2-HYB can be extended to multiorbital problems with density-density interactions. However, the increase in the time span that can be simulated is only  $\frac{n}{n-1}$ , where  $n$  is the number of electronic flavors ( $n = 2$  for the single-orbital AIM).

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