

Fast evaluation of imaginary-time Feynman diagrams and robust bath fitting

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Two software tools: **ppsc-soe** and **adapol** .

1. **ppsc-soe**

- ▶ **P**seudo-**p**article self-consistent calculations via **s**um-**o**f-**e**xponentials.
- ▶ **F**ast, *deterministic*, and **p**arallel impurity solver.
- ▶ Written in C++, with a Python interface, compatible with TRIQS.

2. **adapol**

- ▶ **A**daptive **p**ole fitting for Matsubara functions.
- ▶ Applications: bath fitting in DMFT, analytic continuation.
- ▶ Written in Python, with a TRIQS interface.

Acknowledgement

Impurity solver (ppsc-soe)



Jason Kaye
CCQ & CCM



Hugo Strand
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Denis Golež
Jozef Stefan

Pole fitting (Adapol)



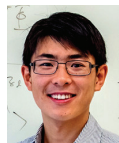
Chia-nan Yeh
CCQ



Nils Wentzell
CCQ



Jason Kaye
CCQ & CCM



Lin Lin
UC Berkeley



Emanuel Gull
U Michigan

How?

A shared ingredient of both softwares are *sum-of-exponentials approximation* of the Matsubara functions, also known as the *pole fitting* problem.

Pole fitting of Matsubara functions

We'd like to find an approximation to $G(i\nu_k)$ in the following form:

$$G(i\nu_k) \approx \sum_{j=1}^p \frac{W_j}{i\nu_k - \lambda_j}, \quad \forall k \in \mathbb{Z}.$$

Here λ_j are the poles, and W_j are the weight matrices ($n \times n$). This is equivalent to the following approximation to $G(\tau)$:

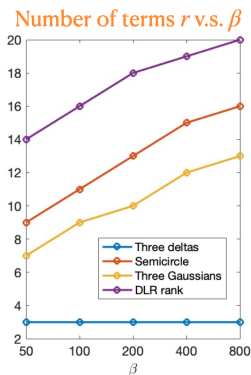
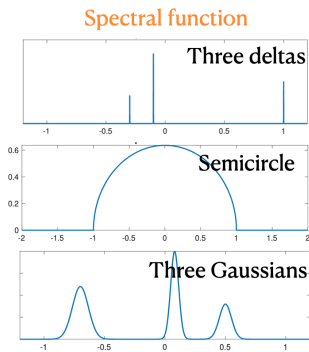
$$G(\tau) \approx - \sum_{j=1}^p \widetilde{W}_j e^{-\lambda_j \tau}, \quad \forall \tau \in [0, \beta].$$

Here $\widetilde{W}_j = \frac{W_j}{1 \mp e^{-\beta E}}$.

We hope that the number of terms p is *as small as possible*.

Fewer poles than DLR?

- ▶ Let spectral width $\leq \Lambda$. Given $\beta\Lambda$ and accuracy ε , DLR provides a **basis** of exponentials for **all** Green's functions.
- ▶ Why do we want another pole fitting method?
- ▶ If we only need poles for a specific $G(\tau)$, could we find a more compact basis (poles)?
- ▶ The answer is YES!



AAA algorithm¹ for rational approximation

Barycentric formula for rational interpolants ($z \in \mathbb{C}$):

$$f(z) = p(z)/q(z) = \left(\sum_{l=1}^m \frac{w_l f_l}{z - z_l} \right) / \left(\sum_{l=1}^m \frac{w_l}{z - z_l} \right).$$

This satisfies $f(z_j) = f_j$ automatically at support points z_j .

AAA algorithm: an **iterative** procedure that selects the next support point in a **greedy** fashion. At the j -th iteration:

- ▶ Select the next sample point z_j , at which the error of the current barycentric approximant is maximal.
- ▶ Choose w_l 's by minimizing $\sum_{z \neq z_j} |f(z)q(z) - p(z)|^2$. This is a linear problem and boils down to compute an SVD.
- ▶ Continue until desired accuracy.

The poles of $f(z)$ are the zeros of $q(z)$. This reduces to a polynomial root finding problem.

Pole fitting procedure

Given $i\nu_j$ and $G(i\nu_j)$,

Step 1: Use AAA algorithm to obtain poles $\{\lambda_k\}_{k=1}^p$.

Step 2: Obtain weights W_j using linear least square fitting.

$$\min_{W_j} \sum_k \left\| G(i\nu_k) - \sum_{j=1}^p \frac{W_j}{i\nu_k - \lambda_j} \right\|^2.$$

Note that this is a linear least squares problem with respect to W_j . (But not linear for the poles λ_j).

Step 3: (optional) postprocessing: further refinement of the poles using bilevel optimizations.

Application: bath fitting

- ▶ In (equilibrium) DMFT with Hamiltonian-based impurity solvers, bath fitting is a crucial subroutine for the self-consistent iterations.
- ▶ Given $\Delta(iw)$, we want to obtain a fitting in the following form:

$$\Delta(iw) \approx \sum_{k=1}^{N_b} \frac{V_k V_k^\dagger}{iw - \omega_k}.$$

- ▶ This could be achieved via **pole fitting** + *semidefinite programming (SDP)* + *bilevel optimization*.
- ▶ Also applicable to the problem of analytic continuation ², especially for noisy data.

Software for hybridization fitting: Adapol

<https://github.com/flatironinstitute/adapol>

Documentation: <https://flatironinstitute.github.io/adapol/>

Adapol

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Python reference manual

Documentation

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Tutorials

- Matsubara fitting for data generated by discrete poles
- Matsubara fitting for data with continuous spectrum (semicircular density)

Python reference manual

- Hybridization fitting
 - `hyffit()`
- Analytic continuation
 - `anacont()`
- TRIQS interface
 - `hyffit_triqs()`
 - `anacont_triqs()`

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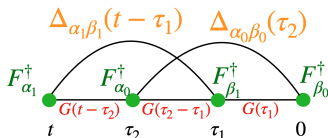
Built with Sphinx using a theme provided by Read the Docs.

Application: fast diagram evaluation

- ▶ Using the same setting as in H. U. R. Strand's previous talk: strong coupling expansion and pseudo-particle representation as an example.
- ▶ The most time-consuming part is to calculate the self-energy Feynman diagrams and single-particle Green's functions Feynman diagrams.

Pseudo-particle self energy Feynman diagrams

OCA (2nd order) diagrams:



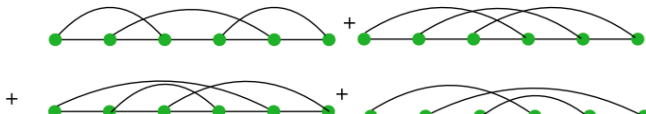
$$= \sum_{\alpha_1, \beta_1=1}^n \sum_{\alpha_0, \beta_0=1}^n \int_0^t d\tau_2 \int_0^{\tau_2} d\tau_1 \Delta_{\alpha_1\beta_1}(t-\tau_1) \Delta_{\alpha_0\beta_0}(\tau_2) F_{\alpha_1}^\dagger G(t-\tau_2) F_{\alpha_0}^\dagger G(\tau_2-\tau_1) F_{\beta_1} G(\tau_1) F_{\beta_0}$$

n : impurity size, Δ : $n \times n$ p.p. hybridization function,

G : $N \times N$ p.p. Green's function,

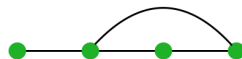
$F_\alpha, F_\beta^\dagger$: $N \times N$: ED of impurity operators $\hat{c}_\alpha, \hat{c}_\beta^\dagger$. ($\alpha, \beta = 1, \dots, n$.)

TCA (third order) diagrams:



Why is the calculation of these diagrams difficult: *crossings of hybridization lines*

- ▶ An m -th order diagram is a $(2m - 2)$ -dimensional integral.
- ▶ Direct calculation cost: $O(n^{2m} \beta^{2m-2})$ for an n -th order diagram. (n is the impurity size.)
- ▶ If there are no crossing of hybridization lines:



$$= \int_0^t dt_2 \int_0^{t_2} dt_1 \Delta(t_2) F^\dagger G(t - t_2) F^\dagger G(t_2 - t_1) F G(t_1) F$$

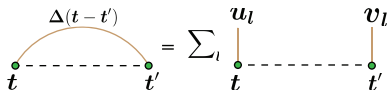
$$= \int_0^t dt_2 F^\dagger G(t - t_2) F^\dagger \Delta(t_2) F \int_0^{t_2} dt_1 G(t_2 - t_1) F G(t_1) F.$$

- ▶ The 2D integral becomes two 1D integrals.

Cutting the hybridization

Decomposition of the hybridization function:

$$\Delta(t - t') = \sum_{l=1}^r c_l e^{-\lambda_l t} e^{\lambda_l t_1} =: \sum_{l=1}^r u_l(t) v_l(t_1).$$



Take OCA diagrams as an example:

Consider $\int_0^t dt_2 \int_0^{t_2} dt_1 \Delta(t - t_1) \Delta(t_2) G(t - t_2) G(t_2 - t_1) G(t_1)$.

If $\Delta(t - t_1) = \sum_{l=1}^r u_l(t) v_l(t_1)$, then the original integral becomes

$$\sum_{l=1}^r \int_0^t dt_2 \int_0^{t_2} dt_1 u_l(t) v_l(t_1) \Delta(t_2) G(t - t_2) G(t_2 - t_1) G(t_1) = \sum_l$$

$$= \sum_{l=1}^r u_l(t) \left(\int_0^t dt_2 \Delta(t_2) G(t - t_2) \left(\int_0^{t_2} dt_1 v_l(t_1) G(t_2 - t_1) G(t_1) \right) \right) = \sum_l$$

Computational complexity

A pole fitting with r poles enables a hybridization of decomposition with nr terms:

$$\Delta(t - t_1) = \sum_{l=1}^{nr} u_l(t)v_l(t_1).$$

Computational cost for m -th order diagram:

$$O((nr)^{m-1}mr_0^2), \quad r_0 \text{ is the DLR rank.}$$

Comparison:

$$n^{2m}\beta^{2m-2} \rightarrow (nr)^{m-1}mr_0^2.$$

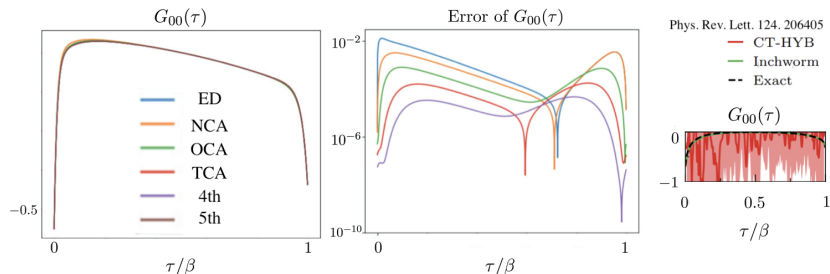
$$r_0 = \log(\beta), \quad r < r_0,$$

Examples

Spinless dimer model: strongly correlated, exact solvable.

$$H = U c_0^\dagger c_0 c_1^\dagger c_1 - v (c_0^\dagger c_1 + c_1^\dagger c_0) - t \sum_{k=0}^1 \sum_{i=0}^1 (c_i^\dagger b_{ik} + \text{h.c.}) - t' \sum_{k=0}^1 (b_{0k}^\dagger b_{1k} + \text{h.c.}),$$

Plot of $G_{00}(\tau)^3$. (For $\beta = 16$.)



Spinless dimer ($n = 2$):

ppsc-soe (TCA)	ppsc-soe (4th order)	ppsc-soe (5th order)	Inchworm	CT-hyb
7.64 seconds	1 hour	90 hours	500 hours	12000 hours

Anderson two-band model ($n = 4$):

ppsc-soe (TCA)	ppsc-soe (4th order)	Inchworm	CT-hyb
0.2 hours	22 hours	1000 hours	3000 hours


DMFT calculation for toy model of Ca_2RuO_4 ($n = 6$):

TCA: 59.7 core hours (for $\beta = 10$).

It takes 6 iterations to reach accuracy $\varepsilon = 10^{-6}$.

There are $4 \times 2^3 \times 6^6 = 1492992$ 5-dim integrals.

⁴Z. Huang, J. Kaye, H. Strand, D. Golez, in preparation.

⁵Multiorbital Quantum Impurity Solver for General Interactions and Hybridizations, E. Eidelstein, E. Gull and G. Cohen, Phys. Rev. Lett., 124, 206405. 

Implementation

- ▶ Code is written in C++, based on cppdlr.
- ▶ Parallelizable: the evaluation of $(np)^{m-1}$ terms could be done simultaneously.
- ▶ Easy to use: python interface.
- ▶ TRIQS compatible.

Conclusion and outlook

Fast impurity solver:

- ▶ Robust, high-order accurate diagram evaluation possible beyond typical lowest-order approximations. Free of sign problem.
- ▶ Access to low temperature.
- ▶ Code ready to be tested on applications.

Bath fitting scheme:

- ▶ Python package (Adapol) available.
- ▶ TRIQS interface.

Ongoing work:

- ▶ Real-time bath fitting, real-time Feynman diagrams, real-time impurity solvers.
- ▶ Analytic continuation of continuous spectrum.
- ▶ Connections to Tensor Cross Interpolation (TCI).
- ▶ More software development ...

Thank you for your attention!

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