

A tutorial on the polaron transformed Redfield equation

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0.1 Correlation function in polaron frame

This is a tutorial for using the polaron transformed Redfield equation (PTRE) in HOQST. For more details on PTRE, [Xu and Cao](#) is a good reference.

In this example, we solve both the Redfield equation and PTRE for a single qubit model with system Hamiltonian

$$H_S = \epsilon\sigma_z + \Delta\sigma_x$$

coupling to an Ohmic bath via σ_z interaction

$$H = H_S + \sigma_z \otimes B + H_B .$$

Loosely speaking, the main difference between the Redfield equation and PTRE is that they have different bath correlation functions. For the Redfield equation, the bath correlation function is

$$C(t_1, t_2) = \langle B(t_1)B(t_2) \rangle .$$

In the polaron frame, however, the bath correlation function becomes

$$K(t_1, t_2) = \exp \left\{ -4 \int_0^t \int_{-\infty}^0 C(t_1, t_2) dt_1 dt_2 \right\} .$$

Again, interested readers can refer to [\[Amin and Averin\]](#) and [Leggett et al.](#) for more details.

0.1.1 Error bound on the second-order master equation

The simplest analysis is to compare the error bounds given in [Mozgunov and Lidar](#) between the Redfield equation and PTRE. We define the error scaling parameter as

$$error = \frac{\tau_B}{\tau_{SB}} .$$

Then we plot the error ratio between the Redfield equation and PTRE

$$R = \frac{error_{\text{Redfield}}}{error_{\text{PTRE}}} ,$$

vs. the system bath coupling strength ηg^2 while fixing other parameters in the Ohmic bath.

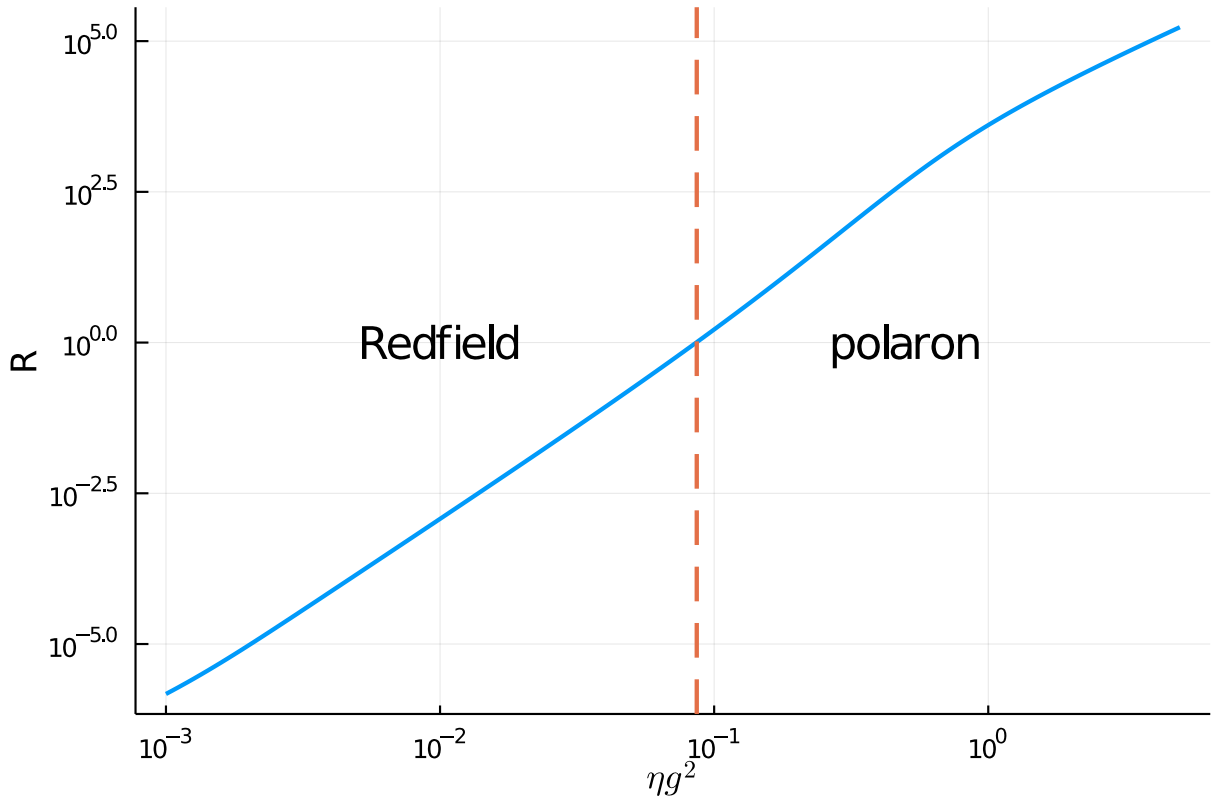
```

using OrdinaryDiffEq, OpenQuantumTools, Plots
using LaTeXStrings

function err_bound(tf, cfun)
    tsb, esb =  $\tau_{SB}$ (cfun)
    tb, eb =  $\tau_B$ (cfun, tf, tsb)
    tb / tsb
end

fc = 4; T = 12; tf = 1000;
 $\eta$ list = log_uniform(1e-3, 5, 1000)
err_ratio = []
err_clist = []
err_klist = []
for  $\eta$  in  $\eta$ list
    bath = Ohmic( $\eta$ , fc, T)
    cfun = (x)->correlation(x, bath)
    pfun = (x)->polaron_correlation(x, bath)
    err_c = err_bound(tf, cfun)
    err_k = err_bound(tf, pfun)
    push!(err_clist, err_c)
    push!(err_klist, err_k)
    push!(err_ratio, err_c/err_k)
end
idx = findfirst((x)->x>=1, err_ratio)
plot( $\eta$ list, err_ratio, xscale=:log10, yscale=:log10, label="", linewidth=2)
vline!([ $\eta$ list[idx]], label="", linestyle=:dash, linewidth=2)
annotate!([(0.5, 1.0, Plots.text("polaron")), (0.01, 1.0, Plots.text("Redfield"))])
xlabel!(L"\eta g^2")
ylabel!("R")

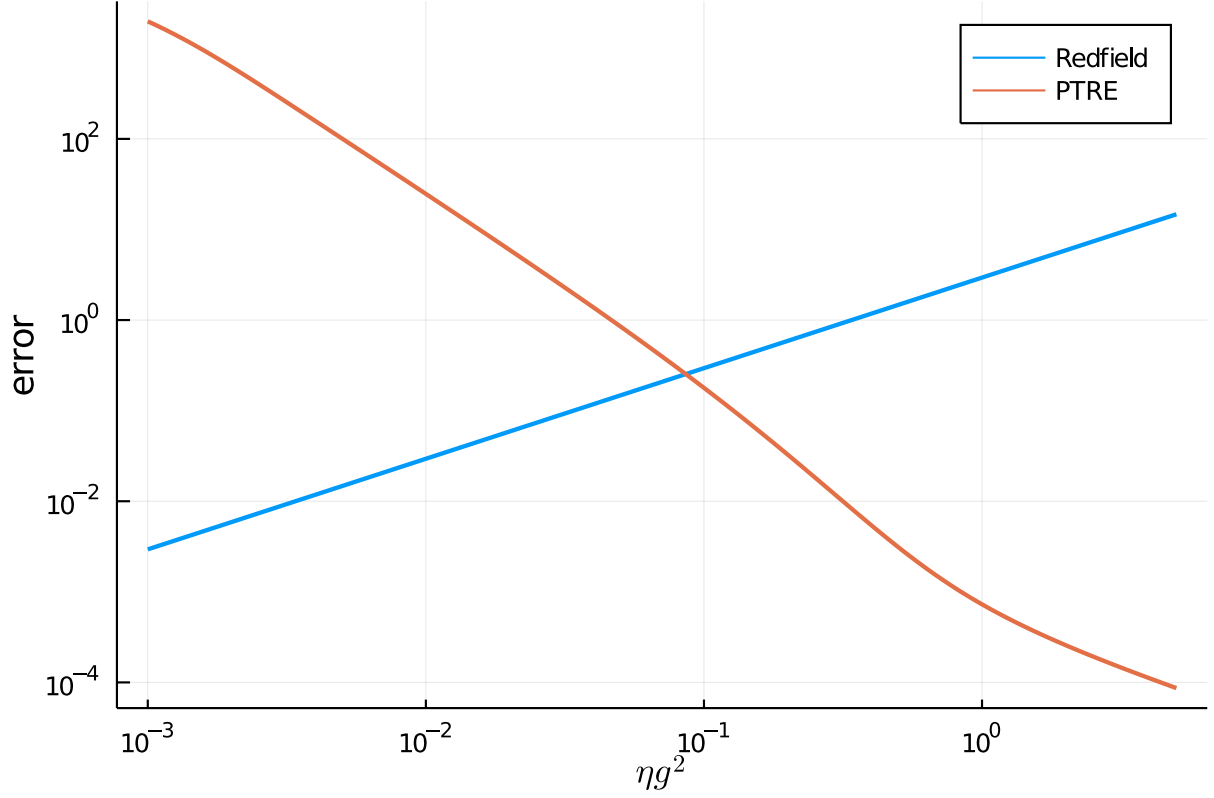
```



From above figure we can see that, as the system-bath coupling strength is bigger than 10^{-1} ,

PTRE should have better error scaling than the usual form of Redfield equation. We also plot the corresponding error values for both Redfield and PTRE:

```
plot(ηlist, err_clist, xscale=:log10, yscale=:log10, label="Redfield", linewidth=2)
plot!(ηlist, err_klist, xscale=:log10, yscale=:log10, label="PTRE", linewidth=2)
xlabel!(L"\eta g^2")
ylabel!("error")
```



The above figure confirms that the range of applicability of the Redfield equation and PTRE are weak and strong coupling regimes, respectively.

0.1.2 Solving PTRE

Since PTRE and the Redfield equation have identical forms, `solve_redfield` can also be used for PTRE. To see this, let's first write down the PTRE for our example.

$$\dot{\rho}_S = \epsilon \sigma_z + [\sigma_i, \Lambda_i(t) \rho_S(t)] + h.c.$$

where $i, j \in [+, -]$, $i \neq j$ and

$$\Lambda_i(t) = \Delta^2 \int_0^t K(t - \tau) U(t, \tau) \sigma_j U^\dagger(t, \tau) d\tau .$$

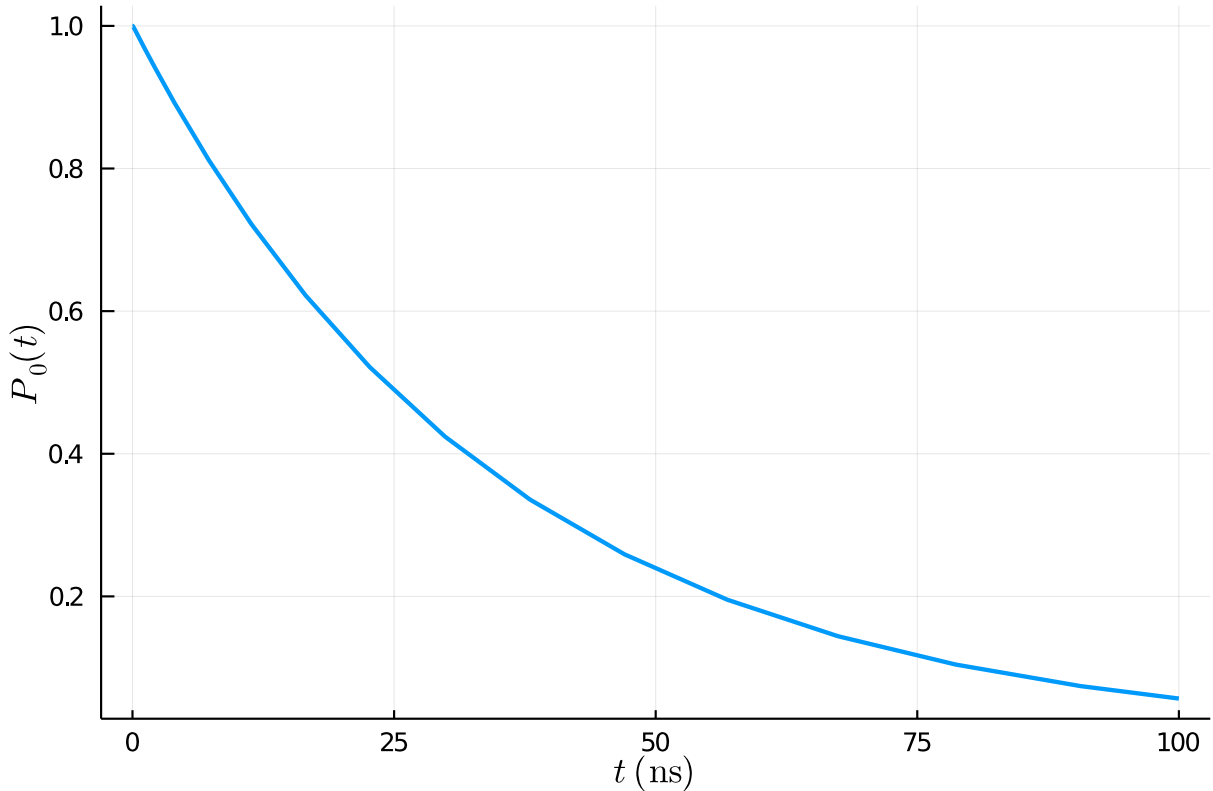
From the above equations, it is clear that the following steps are needed to define an annealing process in the polaron frame:

1. define a new Hamiltonian $H = \epsilon \sigma_z$;
2. define new coupling operators σ_- and σ_+ ;

- define new correlated bath with two-point correlation $K_{i,j}(t_1, t_2)$;

The following code block illustrates how these can be done in HOQST:

```
# assume  $\epsilon = 1$ 
const  $\Delta = 0.1$ 
# define the Ohmic bath in polaron transformed frame
 $\eta = 0.5$ ; bath = Ohmic( $\eta$ , fc, T)
K(t1, t2) =  $\Delta^2$  * polaron_correlation(t1-t2, bath)
cfun = [nothing K; K nothing]
pbath = CorrelatedBath(((1,2),(2,1)), correlation=cfun)
# define coupling as  $\sigma+$  and  $\sigma-$  operators
 $\sigma_p = \begin{bmatrix} 0 & 1 \\ 0 & 0.0im \end{bmatrix}$ ;  $\sigma_m = \begin{bmatrix} 0 & 0 \\ 1 & 0.0im \end{bmatrix}$ 
coupling = ConstantCouplings([ $\sigma_p$ ,  $\sigma_m$ ])
# manually define the unitary operator
U(t) = exp(-2.0im *  $\pi$  *  $\sigma_z$  * t)
H = DenseHamiltonian([(s)->1.0], [ $\sigma_z$ ])
u0 = PauliVec[3][1]
annealing = Annealing(H, u0, coupling = coupling, bath = pbath)
tf = 100
sol_ptre = solve_redfield(annealing, tf, U, alg=Tsit5(), Ta=2, reltol=1e-5)
pop_e = [real(s[1,1]) for s in sol_ptre.u]
plot(sol_ptre.t, pop_e, xlabel=L" $t \ (\mathrm{ns})$ ", ylabel=L" $P_0(t)$ ", label="",
linewidth = 2)
```

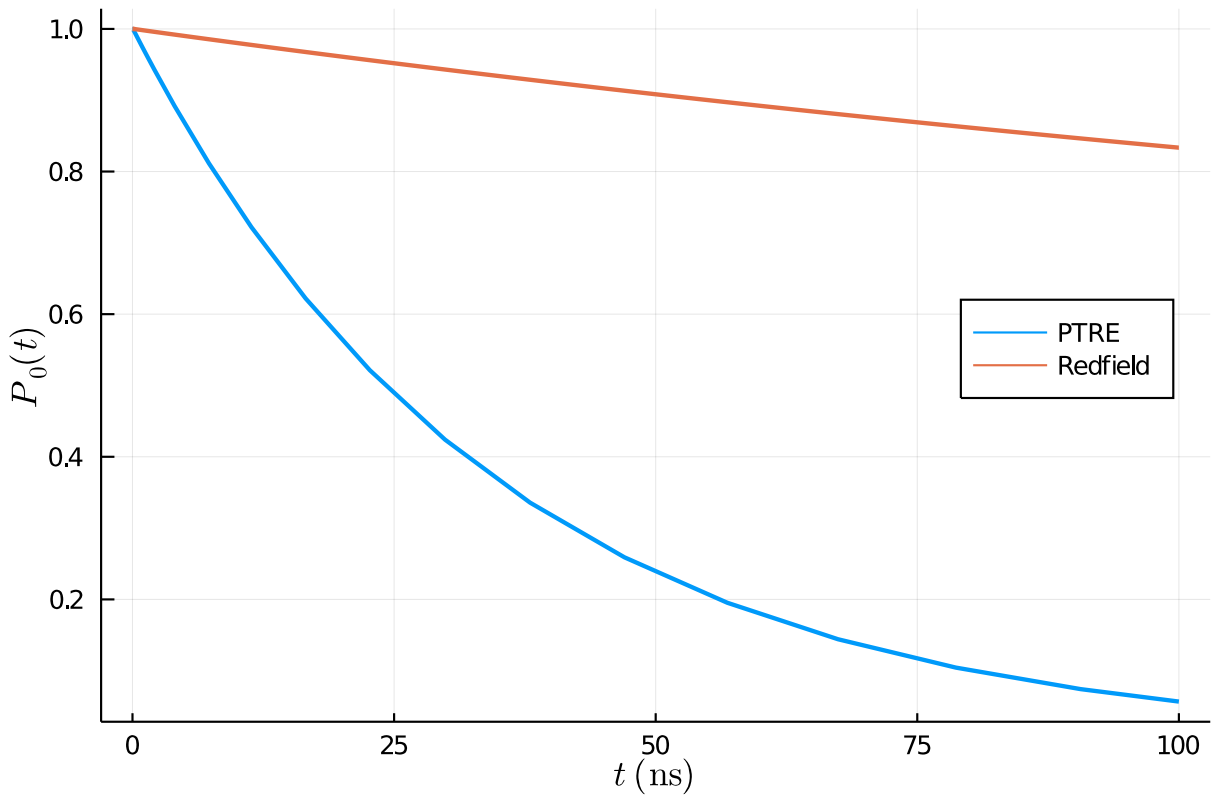


For historical reasons, this is known as an example of "incoherent tunneling". The off-diagonal elements of the density matrix in computational bases(Z-bases) during the entire evolution are 0(shown in the next section).

0.1.3 Redfield equation

What would happen to the Redfield equation in this regime? We can also try:

```
H = DenseHamiltonian([(s)->1.0], [σz+0.1*σx])
coupling = ConstantCouplings(["Z"])
annealing = Annealing(H, u0, coupling = coupling, bath = bath)
tf = 100
sol_redfield = solve_redfield(annealing, tf, U, alg=Tsit5(), Ta=40, reltol=1e-5,
callback=PositivityCheckCallback())
pop_e_redfield = [real(s[1,1]) for s in sol_redfield.u]
plot(sol_ptre.t, pop_e, xlabel=L"t\ (\mathrm{ns})", ylabel=L"P_0(t)", label="PTRE",
linewidth = 2, legend = :right)
plot!(sol_redfield.t, pop_e_redfield, xlabel=L"t\ (\mathrm{ns})", ylabel=L"P_0(t)",
label="Redfield", linewidth = 2)
```



PTRE gives a much stronger decay than the Redfield equation for the parameters chosen in this example. One can also verify the amplitude of the off-diagonal elements during the evolution. Unlike PTRE, the solution of the Redfield equation has non-vanishing off-diagonal elements of the density matrix.

```
t_axis = range(0, 5, length=100)
off_diag_ptre = [abs(sol_ptre(t)[1,2]) for t in t_axis]
off_diag_redfield = [abs(sol_redfield(t)[1,2]) for t in t_axis]
plot(t_axis, off_diag_ptre, xlabel=L"t\ (\mathrm{ns})", ylabel=L"\lvert \rho_{12} \rvert(t)", label="PTRE", linewidth = 2, legend=:right)
plot!(t_axis, off_diag_redfield, xlabel=L"t\ (\mathrm{ns})", ylabel=L"\lvert \rho_{12} \rvert(t)", label="Redfield", linewidth = 2)
```

