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spontaneousORIGINAL (Calls: 1, Time: 142.298 s) Generated 15-Jul-2024 15:16:58 using performance time. script in file D:\Aalto\2324\BScThesis\OLD\spontaneousORIGINAL.m Copy to new window for comparing multiple runs

```
Function listing
 time
         Calls
                  line
                  28 close all
                  29 profile on
                  30
< 0.001
                   31 N = 1500;
< 0.001
              1
                   32 Nr = 25;
< 0.001
                   33 w = 1; %The qubit frequency (taken to be normalized to 1)
< 0.001
                   34 include mutual = 1:
              1
< 0.001
              1
                   35 gamma = w/(5*sqrt(2));
< 0.001
                   37 te_results = zeros(1, N); %The array for collecting the results of long time evolution
                   38 gge_results = zeros(1, N+1); %The array for collecting the results of the GGE prediction
< 0.001
                  39 %%
                   40 dia1 = sort(2*w*rand(N,1));
< 0.001
                   41 for idx = 1:Nr
              1
< 0.001
                  42
                  43 %{
                  44 %In this section we first generate the bath Hamiltonian H1
                  45 according to the rule Hij = [-gamma/sqrt(N),gamma/sqrt(N)] and Hii =
                  46 [0,2*Omega]. Then we expand it to include the initially exited site to form
                  47 the total Hamiltonian H.
                  48 %}
                  49
                   50 a = -(gamma/sqrt(N)) + 2*(gamma/sqrt(N))*rand(N);
  1.172
             25
  0.381
                   51 a = include_mutual*triu(a);
                   52 H1 = a+a';
  0.437
             25
                   53 H1 = H1 - diag(diag(H1)) + diag(dia1);
  0.595
             25
                  54
                  55 %Diagonalize the bath Hamiltonian
 25,479
             25
                   56 [vek1, ek1] = eig(H1);
                  57
                  58 %vek1 = \lceil vek1; zeros(N+3,N) \rceil;
                  59 %vek2 = [zeros(N+3,N); vek2];
                  60 \text{ %vek} = [\text{vek1 vek2}]
                  61
                  62 %Generate the couplings from the baths to the resonators
  0.004
             25
                  63 lambda1 = -(gamma/sqrt(N)) + 2*(gamma/sqrt(N))*rand(N,1);
                  64
                  65 %Build the total Hamiltonian
  0.309
             25
                   66 H = blkdiag(H1, w);
  0.001
             25
                   67 \text{ H}(1:N,N+1) = \text{lambda1};
  0.003
                   68 H(N+1,1:N) = lambda1';
                  69 %%
                  70 %{
                  71 In this section, we diagonalize the total Hamiltonian H and set the intial state.
                  72 %}
 25.155
                   74 [vel, el] = eig(H); %Diagonalize the total Hamiltonian to eigenvectors vel and eigevalues el
  0.575
             25
                   75 plot(diag(el), 'o')
                   76 xi02 = zeros(N+1);
  0.002
< 0.001
             25
                   77 xi02(N+1, N+1) = 1;
                   78 xi0 = xi02; %The initial state of the system
  0.057
                  79
                  80 %%
                  81 %{
                  82 In this section, we time-evolve the intial density matrix and calculate the resulting populations.
                  83 %}
< 0.001
                   84 tmax = 8000000000; "The final time at which the populations are calculated
                   85 t = linspace(0,tmax,2); %Vector for times for which to calculate the time evolution
  0.005
             25
                  86 %E1 = zeros(1, N);
< 0.001
             25
                   88 for i = 1:length(t)
                          Ul = expm(1i*t(i)*el); %Time-evolution operator exp(iHt) in the eigenbasis of H (easy to calculate)
  1.578
             50
                   89
 36.118
             50
                   90
                          Uj = vel*Ul*(vel'); %Convert it to the basis of the sites
 46,059
             50
                   91
                          xi = (Uj')*xi0*Uj; %\rho(t) = exp(-iHt)\rho(t=0)exp(iHt)
                   92 end
  0.002
                  93
                  94 %{
                  95 for j = 1:N
                  96
                           Opi = zeros(N+1):
                           Opj(j,j) = 1; %Defining the number operator Opj of the j:th site
                  97
                           E1(j) = trace(Opj*xi); %Calculating the expectation value of it
                  99 end
```

```
100 %}
  0.005
                  101 e1 = diag(xi);
                  102 E1 = e1(1:N);
  0.001
             25
                  103
  0.106
                104 el = diag(el);
                 105 %%
                 106 %{
                  107 In this section, we calculate the numerical GGE prediction for the populations, which is to compared
                  108 the long-time evolution. Basically, we just implement a convolution
                 109 formula.
                 110 %}
  0.002
             25
                 111 nau = zeros(1, N+1);
  0.003
             25
                 112 ujt = abs(vel(N+1,:)).^2;
                 113
< 0.001
             25
                 114 for k = 1:(N+1)
  1.171
          37525
                  115
                          uki = abs(vel(k,:)).^2;
  0.052
          37525
                  116
                          nau(k) = dot(ujt, uki);
          37525
                  117 end
  0.008
                  118
  0.053
             25 119 te_results = te_results+E1;
             25 120 gge_results = gge_results+nau;
< 0.001
                 121
                 122
                 123 end
  0.001
                 124 %%
                 125 %{
                  126 Finally, we average over the set number of iterations.
                 127 %}
  0.006
              1 128 te result = te results/Nr;
< 0.001
                  129 gge result = gge results/Nr;
< 0.001
              1 130 epsilon = sort(dia1);
                 131 %%
                 132 %{
                 133 In this section, we plot the results of both the numerical long-time
                 134 evolution and the GGE prediction and compare an analytical formula.
                 135 %}
                 136
  0.005
                  137 omega = linspace(0,2*w,1000000);
                  138 gavg = (gamma^2)/(3*N);
< 0.001
< 0.001
              1
                  139 \text{ Omega} = w;
< 0.001
                  140 nu0 = N/(2*Omega);
< 0.001
                 141 g2 = (gamma^2)/(3*N);
                  142 rate = pi*nu0*gavg;
< 0.001
              1
                 143 nl = 2*gavg./((1-omega).^2+(2*rate)^2); %The analytical prediction for the populations
  0.002
              1
                  145 %-The final plotting of the results:
                  146 % (i) Numerical long-time evolution (ii) Numerical GGE (iii) Analytical
                 147 % -%
  1.643
                  148 a1 = semilogy(dia1, te_result, 'o', "Color", 'b');
              1 149 hold on
  0.008
                 150
                 151 a2 = plot(dia1, gge_result(1:N), 'x', "LineWidth", 1.1, "Color", "g");
  0.004
  0.004
              1 152 a3 = plot(omega, nl, "LineWidth", 1.2, "Color", "r");
                 153
                 154 xlabel("$\omega\\Omega$", 'Interpreter',"latex", 'FontSize',18)
155 ylabel("$n$", 'Interpreter',"latex", 'FontSize',18)
  0.020
  0.007
                  156 legend([a1(1), a2(1), a3(1)], 'Long-time evolution', 'Numerical GGE', 'Analytical GGE', 'location', "northwest")
  1.160
                  157 ylim([0.5*10^(-5),10^(-1)])
  9.997
  0.003
              1 158 hold off
                 159 %%
                 160 %{
                 161 Uncomment the line below and change the path to your desired location to save the resulting data.
                  162 %}
                 163
                  164 %save("mutual_off", "te_result", "gge_result", "dia1")
                  165
  0.090
              1 166 profile viewer
```

Other subfunctions in this file are not included in this listing.