decoherence_times2rates(tau)

Decoherence rates from decoherence times. Zero decoherence rate for the exactly degenerate states and for the self-decoherence (diagonal elements)

$$r_{ij} = \begin{cases} \frac{1}{\tau_{ij}}, & if \ \tau_{ij} > 0\\ 0, otherwise \end{cases}$$

energy_gaps(Hvib)

Absolute values of energy gaps for a single trajectory at every timestep.

$$\Delta E_{ij}(t) = \left| Hvib_{ii}(t) - Hvib_{jj}(t) \right|$$

energy_gaps_ave(Hvib, itimes, nsteps)

Absolute values of energy gaps averaged over multiple trajectories at every timestep.

$$\Delta E_{ij}(t) = \frac{1}{N_{it}N_{tr}} \sum_{k=1}^{N_{it}} \sum_{n=1}^{N_{tr}} |Hvib_{ii}^{n}(t_k + t) - Hvib_{jj}^{n}(t_k + t)|$$

 N_{it} – the number of starting times (initial conditions)

 N_{tr} - the number of nuclear trajectories (datasets)

decoherence_times(Hvib, verbosity=0)

Decoherence times and rates for a single trajectory

$$\tau_{ij} = \begin{cases} \sqrt{\frac{12}{5}} \frac{1}{\delta E_{ij}} &, \delta E_{ij} > 0 \\ 10^{10}, otherwise \end{cases}, \delta E_{ij} > 0 \qquad r_{ij} = \begin{cases} \frac{1}{\tau_{ij}}, & if \ \tau_{ij} > 0 \\ 0, otherwise \end{cases}$$

Here,

$$\delta E_{ij} = \sqrt{\frac{1}{N_{steps}}} \sum_{p=1}^{N_{steps}} (\Delta E_{ij}(t_p) - \langle \Delta E_{ij} \rangle)^2,$$

$$\langle \Delta E_{ij} \rangle = \frac{1}{N_{steps}} \sum_{p=1}^{N_{steps}} \Delta E_{ij}(t_p),$$

$$\Delta E_{ij}(t_n) = |Hvib_{ij}(t_n) - Hvib_{ij}(t_n)|.$$

 N_{steps} – the number of MD steps in this single trajectory

 $t_p=p\Delta t$ – are the time-points along the MD trajectory, indexed by the integer subscript for clarity in practical implementation.

decoherence_times_ave_old(Hvib, itimes, nsteps, verbosity=0)

WARNING: this function should not be replaced by the next one!

Decoherence times and rates based on the averaged energy gap fluctuations (averaged by the magnitude)

$$\tau_{ij} = \begin{cases} \sqrt{\frac{12}{5}} \frac{1}{\delta E_{ij}} &, \delta E_{ij} > 0 \\ 10^{10}, otherwise \end{cases} , \delta E_{ij} > 0$$

$$r_{ij} = \begin{cases} \frac{1}{\tau_{ij}}, & if \ \tau_{ij} > 0 \\ 0, otherwise \end{cases}$$

Here,

$$\delta E_{ij} = \sqrt{\frac{1}{N_{steps}}} \sum_{p=1}^{N_{steps}} (\Delta E_{ij}(t_p) - \langle \Delta E_{ij} \rangle)^2,$$

$$\langle \Delta E_{ij} \rangle = \frac{1}{N_{steps}} \sum_{p=1}^{N_{steps}} \Delta E_{ij}(t_p),$$

$$\Delta E_{ij}(t_p) = \frac{1}{N_{ir}N_{tr}} \sum_{k=1}^{N_{it}} \sum_{n=1}^{N_{tr}} \left| Hvib_{ii}^n \left(t_k + t_p \right) - Hvib_{jj}^n \left(t_k + t_p \right) \right|.$$

 N_{it} – the number of starting times (initial conditions)

 N_{tr} - the number of nuclear trajectories (datasets)

 N_{steps} – the number of MD steps in this single trajectory

 $t_p=p\Delta t$ – are the time-points along the MD trajectory, indexed by the integer subscript for clarity in practical implementation.

 t_k – are the starting time-points of the MD trajectory.

decoherence_times_ave(Hvib, itimes, nsteps, verbosity=0)

Decoherence times and rates based on the averaged energy gap fluctuations (averaged by the magnitude).

$$\tau_{ij} = \begin{cases} \sqrt{\frac{12}{5}} \frac{1}{\delta E_{ij}} &, \delta E_{ij} > 0 \\ 10^{10} & otherwise \end{cases} \quad r_{ij} = \begin{cases} \frac{1}{\tau_{ij}}, & if \ \tau_{ij} > 0 \\ 0, otherwise \end{cases}$$

Here,

$$\delta E_{ij} = \sqrt{\frac{1}{N_{steps}N_{it}N_{tr}}} \sum_{p=1}^{N_{steps}} \sum_{k=1}^{N_{it}} \sum_{n=1}^{N_{tr}} \left(\Delta E_{ij}^{n} \left(t_k + t_p\right) - \langle \Delta E_{ij} \rangle \right)^2,$$

$$\langle \Delta E_{ij} \rangle = \frac{1}{N_{steps}N_{it}N_{tr}} \sum_{p=1}^{N_{steps}} \sum_{k=1}^{N_{it}} \sum_{n=1}^{N_{tr}} \Delta E_{ij}^n \big(t_k + t_p\big),$$

$$\Delta E_{ij}^n \big(t_k + t_p\big) = \big| Hvib_{ii}^n \big(t_k + t_p\big) - Hvib_{jj}^n \big(t_k + t_p\big) \big|.$$

 N_{it} – the number of starting times (initial conditions)

 N_{tr} - the number of nuclear trajectories (datasets)

 N_{steps} – the number of MD steps in this single trajectory

 $t_p=p\Delta t$ – are the time-points along the MD trajectory, indexed by the integer subscript for clarity in practical implementation.

 t_k – are the starting time-points of the MD trajectory.