

DiffuTrapKv2

Simulation of diffusion and trapping of a tetramer of size “molecule diameter” onto fixed interaction sites. In the tetramer there are “Cter” number of interacting subunits.

Reference: Paupiah et al (2025) *Epileptic encephalopathy-related Kv2.1 mutants impair channel clustering and membrane distribution but not neuronal excitability* bioRxiv <https://doi.org/10.1101/2025.10.13.681825>

Enter the parameters in the pop up window:

Number of Cter – number of molecules:

Enter the number of subunits bearing the entire Cter (with a PRC domain) and the number of molecules with these characteristics (two numbers, separated by a space):

Example: 2 100 -> means a simulation with 100 molecules (tetramers) with 2 PRC domains.

If you combine two groups, enter the values for one after the other:

Example: 2 100 4 100

Diffusion coefficient:

In $\mu\text{m}^2/\text{s}$ (mean value of a gaussian distribution of D).

Trajectory length:

Duration of the simulation, in “SPT” frames.

Circle diameter:

Circular space, in the middle of the simulation space, where interactions may occur.

Simulation space: Size of the bounding box where molecules diffuse.

Kon and koff: Effective constants for the scaffolding interaction (each PRC acting independently).

Time between images, Pixel size, Localization accuracy: The same than the SPT experiment that the simulation wants to reproduce.

Molecule diameter: Size of the diffusing molecule (circle).

Output :

- SPT-like trajectory files (.trc, a txt file), in \trc folder with:

1.000000e+00	3.000000e+00	5.969764e+00	6.562170e+00	1.000000e+00
1.000000e+00	4.000000e+00	5.335403e+00	6.836104e+00	1.000000e+00
1.000000e+00	5.000000e+00	4.689345e+00	7.000920e+00	1.000000e+00
molecule #	time (frame)	x position	y position	group # (# Cter)
1.000000e+00	8.000000e+00	4.202305e+00	6.835891e+00	1.000000e+00
1.000000e+00	9.000000e+00	4.378034e+00	5.516603e+00	1.000000e+00
1.000000e+00	1.000000e+01	4.876350e+00	4.450989e+00	1.000000e+00
1.000000e+00	1.100000e+01	5.716551e+00	3.920546e+00	1.000000e+00

Files **ctergroup#-simulation#.trc** : trajectories of each group separately

Files **sim-simulation#-all.trc** : all groups together

- A report file with the parameters (main folder)