

Stochastic modelling of licensed stem cell states

Rodrigo García-Tejera* and Luciana Luque

Here we explore the stochastic dynamics of the SL model previously discussed. The SL model captures the stochastic dynamics of a stem cell population composed of unlicensed (S species) and licensed (L species) states. The SL model is defined by the set of reactions



You can download the files from

<https://github.com/RodrigoGarciaTejera/EastBioClass/tree/main/Stochastic>. The file `run_SL_model.m` runs a stochastic simulation and shows the trajectories as well as the histograms for the statistics of the S and L species.

Play with the parameters and see what outcomes you get. Note that lines 27 and 28 set the homeostatic number of S and L cells, and then k_3 and k_4 are set such that the system yields that homeostatic state.

Questions

- change the value of the S -to- L switching rate k_2 , what happens when you increase it?
- try setting k_2 very slightly lower than k_1 , what do you observe? what could the reason be?
- what happens if we remove the reversible reaction? For example, you can add a “%” symbol at the beginning of line 32 to comment it, and define `k3=0` below.

* rodrigo.garcia@ed.ac.uk