

TP 4 : Kinetic Monte-Carlo (*KMC*)

Cours de modélisation numérique

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Introduction

In this TP, we will implement the chemical reaction seen as an example during the lessons, through the *Kinetic Monte-Carlo KMC* method.

Model

Consider two interacting chemical species A and B of initial quantities A_0 and B_0 , and let the reaction rates be $k_1 \geq 0$ and $k_2 \geq 0$ such that :



It can be shown that the evolution of these quantities is given by the following equations :

$$A(t) = \frac{k_2(A_0 + B_0)}{k_1 + k_2} + \frac{A_0k_1 - B_0k_2}{k_1 + k_2}e^{-(k_1+k_2)t} \qquad (2)$$

$$B(t) = \frac{k_1(A_0 + B_0)}{k_1 + k_2} - \frac{A_0k_1 - B_0k_2}{k_1 + k_2}e^{-(k_1+k_2)t} \qquad (3)$$

We propose to simulate the evolution of the system using a Monte Carlo method. For this purpose, A and B will be discretised for example in a number of molecules. The algorithm will have to, at each time step Δt , iterate over the number $N = A + B$ of molecules present in the system and :

- Draw with probability $p(A) = \frac{A}{N}$ a molecule A , or $p(B) = \frac{B}{N}$ a molecule B .
- If species A is drawn, one carries out reaction $A \rightarrow B$ with probability $k_1\Delta t$.
- If species B is drawn, one carries out reaction $B \rightarrow A$ with probability $k_2\Delta t$.

Note that the choice of parameters must be such that $k_1\Delta t \leq 1$ and $k_2\Delta t \leq 1$ are satisfied.

Write a `Python` code implementing this algorithm and test it for different values of $A_0, B_0, k_1, k_2, \Delta t$. You can use the `random.uniform()` function, which returns a random number between 0 and 1.

Compare your results with the expected value for the analytical solution when $t \rightarrow \infty$.