

# **Supplementary materials**

Stochastic simulation algorithm of beta-amyloid  
aggregation in Alzheimer Disease

Network Modeling and Simulation course report

**Carlo Aleotti, Anastasia Santo e Rayan Slatni**

Jan 2024

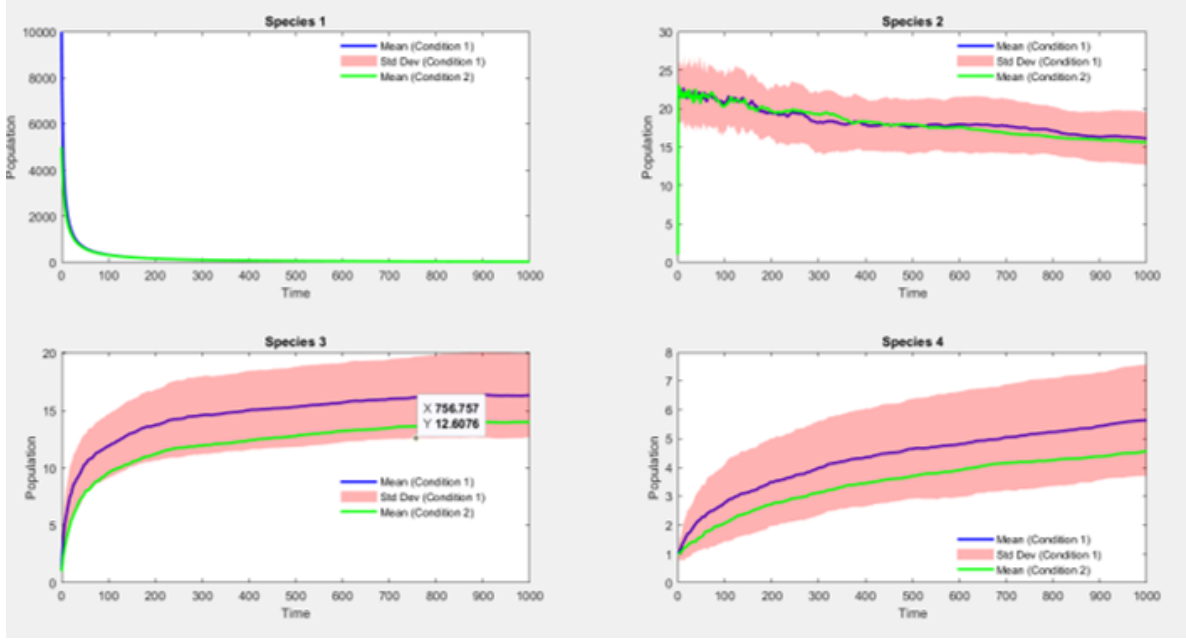


Figure 1: The evolution of amyloid beta aggregations for M1, M2, M3 and M4. Rate constants ( $k_0 \neq \dots \neq k_{53}$ ),  $k_0 = 0.00001$ ,  $k_2 = k_0/2, \dots$  Condition 1  $M1 = 10000$  and Condition 2  $M2 = 5000$ , and the other populations  $M2, \dots, M8 = 1$

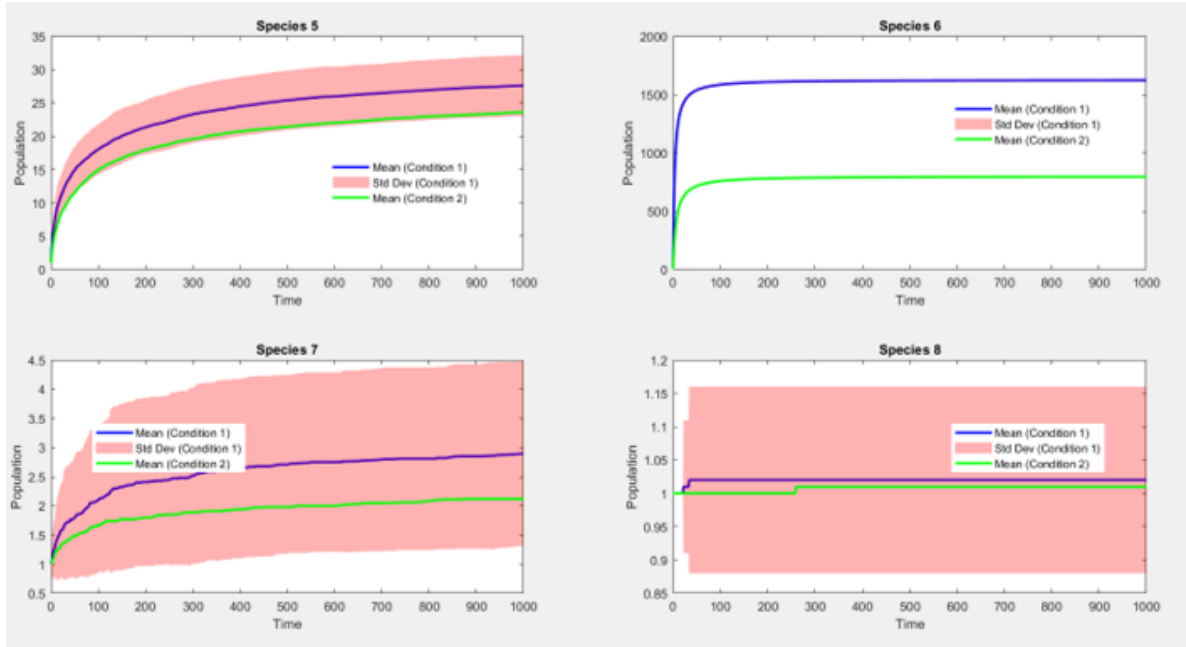


Figure 2: The evolution of amyloid beta aggregations for M5, M6, M7 and M8. Rate constants ( $k_0 \neq \dots \neq k_{53}$ ),  $k_0 = 0.00001$ ,  $k_2 = k_0/2, \dots$  Condition 1  $M1 = 10000$  and Condition 2  $M2 = 5000$ , and the other populations  $M2, \dots, M8 = 1$

Function Name	Calls	Total Time (s) ↓	Self Time* (s)	Total Time Plot (dark band = self time)
<a href="#">modelwithoutfragmentation</a>	1	46.602	0.637	
<a href="#">MonteCarloSimulation</a>	2	44.515	0.071	
<a href="#">GillespieAlgorithm</a>	200	44.329	25.212	
<a href="#">randsample</a>	754845	16.146	4.692	
<a href="#">RandStream.RandStream&gt;RandStream.getGlobalStream</a>	754845	5.958	0.829	
<a href="#">RandStream.RandStream&gt;localGetSetGlobalStream</a>	754845	5.128	5.128	
<a href="#">histcounts</a>	754845	3.172	2.195	
<a href="#">exprnd</a>	754845	2.970	2.970	
<a href="#">RandStream.rand</a>	754845	2.325	2.325	

Figure 3: Profiling of the Monte Carlo-Gillespie algorithm using aggregation.

Function Name	Calls	Total Time (s) ↓	Self Time* (s)	Total Time Plot (dark band = self time)
<a href="#">modelwithoutfragmentation</a>	1	46.948	0.709	
<a href="#">MonteCarloSimulation</a>	2	44.814	0.070	
<a href="#">NextReactionMethod</a>	200	44.629	25.033	
<a href="#">randsample</a>	754906	16.584	4.894	
<a href="#">RandStream.RandStream&gt;RandStream.getGlobalStream</a>	754906	6.060	0.839	
<a href="#">RandStream.RandStream&gt;localGetSetGlobalStream</a>	754906	5.221	5.221	
<a href="#">histcounts</a>	754906	3.292	2.273	
<a href="#">exprnd</a>	754906	3.012	3.012	
<a href="#">RandStream.rand</a>	754906	2.337	2.337	

Figure 4: Profiling of the FRM algorithm using aggregation.

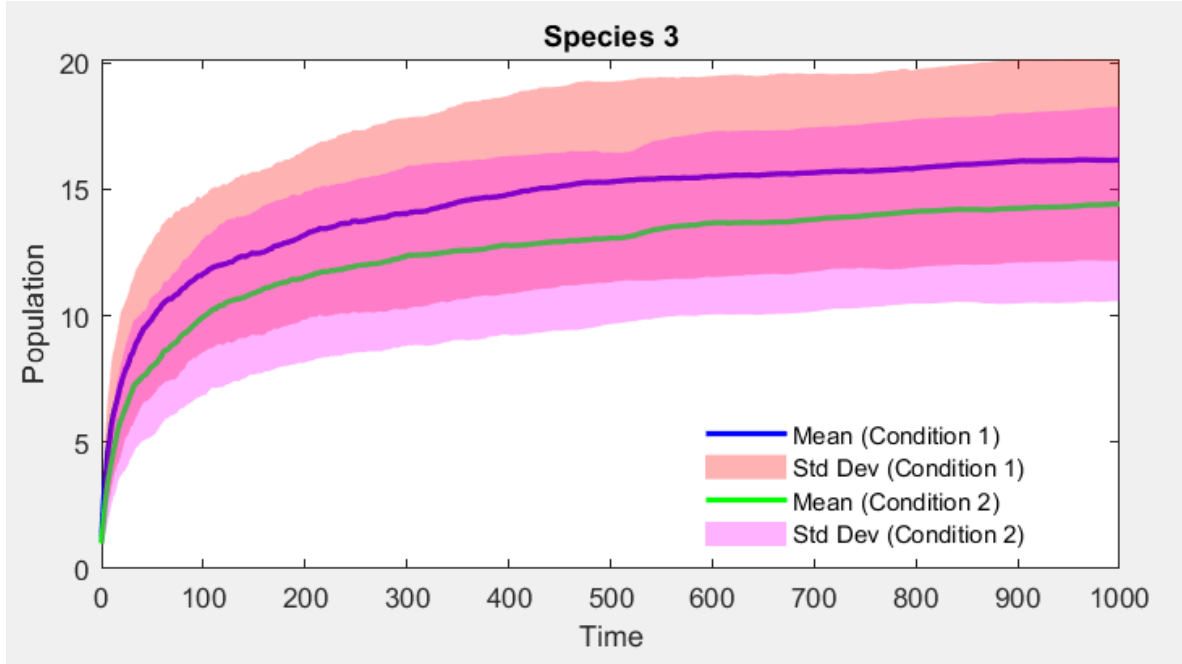


Figure 5: The evolution of amyloid beta aggregations and fragmentation for M3. Rate constants ( $k_0 \neq \dots \neq k_{53}$ ),  $k_0 = 0.00001$ ,  $k_2 = k_0/2, \dots$  Condition 1  $M1 = 10000$  and Condition 2  $M2 = 5000$ , and the other populations  $M2, \dots, M8 = 1$

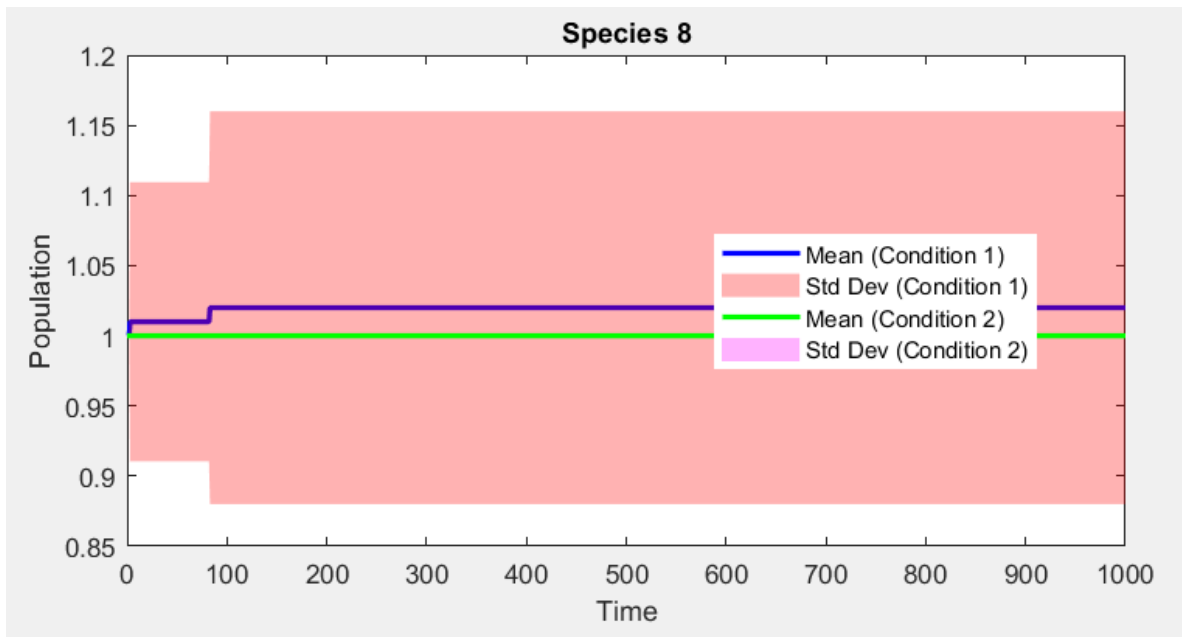


Figure 6: The evolution of amyloid beta aggregations and fragmentation for M3. Rate constants ( $k_0 \neq \dots \neq k_{53}$ ),  $k_0 = 0.00001$ ,  $k_2 = k_0/2, \dots$ . Condition 1  $M1 = 10000$  and Condition 2  $M2 = 5000$ , and the other populations  $M2, \dots, M8 = 1$

$$S_a = \begin{bmatrix} -2 & 1 & 0 & 0 & 0 & 0 & 0 & 0 \\ -1 & -1 & 1 & 0 & 0 & 0 & 0 & 0 \\ -1 & 0 & -1 & 1 & 0 & 0 & 0 & 0 \\ 0 & -2 & 0 & 1 & 0 & 0 & 0 & 0 \\ -1 & 0 & 0 & -1 & 1 & 0 & 0 & 0 \\ -1 & -2 & 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & -1 & -1 & 0 & 1 & 0 & 0 & 0 \\ -1 & 0 & 0 & 0 & -1 & 1 & 0 & 0 \\ 0 & -1 & 0 & -1 & 0 & 1 & 0 & 0 \\ 0 & 0 & -2 & 0 & 0 & 1 & 0 & 0 \\ -2 & -2 & 0 & 0 & 0 & 1 & 0 & 0 \\ -1 & 0 & 0 & 0 & 0 & -1 & 1 & 0 \\ 0 & -1 & 0 & 0 & -1 & 0 & 1 & 0 \\ -1 & 0 & -2 & 0 & 0 & 0 & 1 & 0 \\ 0 & 0 & -1 & -1 & 0 & 0 & 1 & 0 \\ -1 & -3 & 0 & 0 & 0 & 0 & 1 & 0 \\ -1 & 0 & 0 & 0 & 0 & 0 & -1 & 1 \\ 0 & -1 & 0 & 0 & 0 & -1 & 0 & 1 \\ 0 & 0 & -1 & 0 & -1 & 0 & 0 & 1 \\ 0 & 0 & 0 & -2 & 0 & 0 & 0 & 1 \\ -1 & -1 & 0 & 0 & -1 & 0 & 0 & 1 \\ -1 & 0 & -1 & -1 & 0 & 0 & 0 & 1 \\ 0 & -2 & 0 & -1 & 0 & 0 & 0 & 1 \\ 0 & -1 & -2 & 0 & 0 & 0 & 0 & 1 \\ -2 & 0 & -2 & 0 & 0 & 0 & 0 & 1 \\ 0 & -4 & 0 & 0 & 0 & 0 & 0 & 1 \\ -2 & -1 & 0 & -1 & 0 & 0 & 0 & 1 \end{bmatrix}$$

Figure 7: Stechiomatrix without fragmentation

