Stochastic Model of Prion's Disease: Nucleation-Polymerisation Model Part II

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Modelling each reaction

Precise reactions step and parameters Deterministic rate equations Stochastic algorithm

Further...

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- n: is the size of the nucleus.

Nucleation

$$M \stackrel{1}{\underset{c_0}{\rightleftharpoons}} f_1$$
 (spontaneous conformation) (1)

$$f_1 + f_1 \stackrel{c_f}{\underset{\frac{c_f}{d_1}}{\longleftarrow}} f_2 \text{ (dimerization)}$$
 (2)

$$f_{k-1} + f_1 \stackrel{c_f}{\rightleftharpoons} f_k$$
 ((k)-mer formation) (3)

$$f_{n-1} + f_1 \xrightarrow{c_f} u_n$$
 (nucleus formation) (4)

Polymerization

$$u_{i} + f_{1} \xrightarrow{k_{pf}} u_{i+1}, (i \geq n), \text{ (elongation)}$$

$$u_{i} \xrightarrow{\frac{k_{pf}}{d_{2}}} u_{i-1} + f_{1}, (i \geq n), \text{ (shortening)}$$

$$u_{i} \xrightarrow{k_{b}} u_{i-j} + u_{j}, (i \geq n), \text{ (polymer break)}$$

$$\text{and } u_{k} \xrightarrow{\infty} k * f_{1}, \text{ if } k < n-1, \text{ (oligomer instability)}$$

$$(5)$$

$$M=c_0*f_1$$

▶ c_0 : equilibrium constant between inactive monomers M and active monomers f_1 . We suppose (fast variable)

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- ightharpoonup M(0): initial quantity of monomers.
- n: nucleus size.



General Behavior: sigmoid

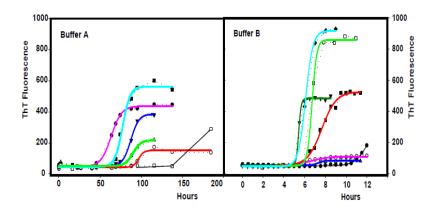
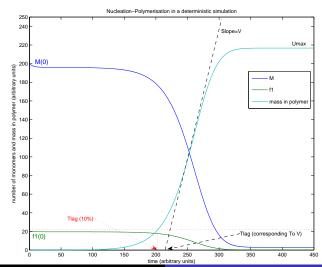


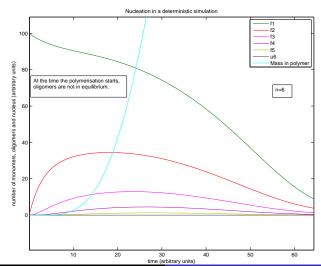
Figure: Several polymerisation experiments



General Behavior: key definitions



General Behavior: nucleation dynamics

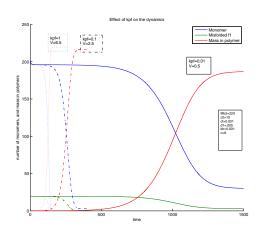


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- c_f: Only changes the Lag time.
- ▶ d_1 : Influences the nucleation time. Can also limit the polymerisation if the oligomers don't disappears quickly enough (ie if d1 high, which is unlikely to be).

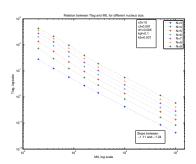
▶ k_{pf} and k_b : Influence both polymerization and nucleation in the same way. Change also the final equilibrium.



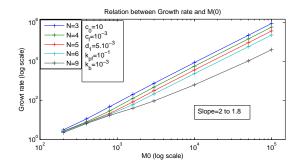


▶ M(0) and n: M(0) reduces the lag time, while n increases it. Up to a constant, Tlag follows the same relation with M(0), whatever the nucleus size is. In agreement with experimental date. We have

$$log(Tlag) \simeq -1.07 * log(M(0))$$
 (8)



▶ M(0) and n: M(0) speeds up the polymerization, while n slightly decreases it. This contradicts experiments.

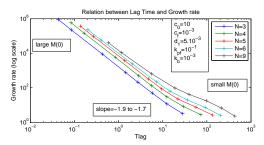


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- ► $M_{tot} * k_{pf} \gg k_b$ guarantee an initially growing polymer.
- ▶ Tlag versus V: For large M(0), they are clearly inversely correlated. This contradicts experiments.



Some key factor: general scaling laws

 $\kappa = \sqrt{2*M_{tot}*k_{pf}*k_b}$: In Knowles et al. (2009), using the nucleated equilibrium hypothesis, and a analytical approximation of the solution, the authors show a relation between the normalized growth (V/M(0)) and κ , and between T_{lag} and κ

$$\frac{V}{M_{tot}} = C_{M(0),n}^{te} * \kappa \tag{9}$$

$$T_{lag} = \frac{log(C_{M(0),n}^{te})}{\kappa}$$
 (10)



experimental variability

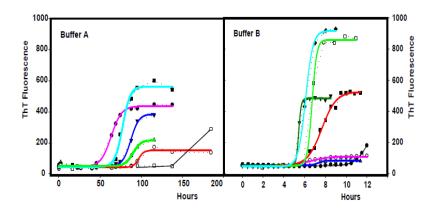


Figure: Several polymerisation experiments



Variability in Lag Time and Growth rate (Controversial)

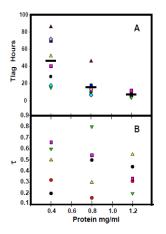


Figure: Relation between M(0) and Tlag and V



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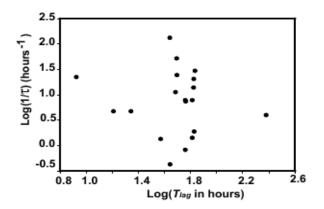
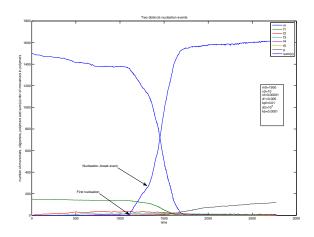


Figure: Relation between Tlag and V

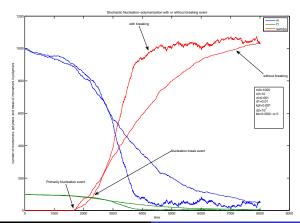


Stochastic algorithm - global dynamic



Break events or Secondary nucleation pathway

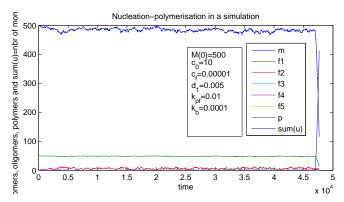
 One single nucleation event then elongation and break events can explain the sigmoid form





Stochasticity in the oligomer

 One can have many (small) fluctuations in the oligomer species, before reaching nucleation

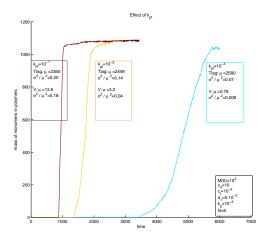


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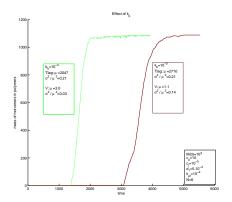
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- $ightharpoonup d_1$: Has no evident effect on variability.

▶ k_{pf} : Large k_{pf} will increase the variability of the growth rate.

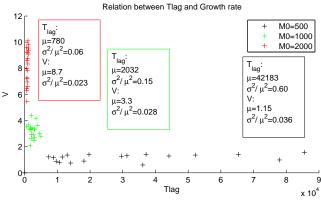




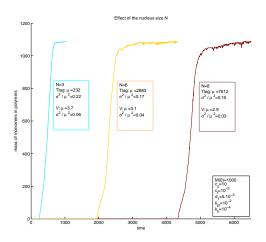
k_b: Influences on the respective numbers of nucleation events/ break events. Small k_b appears to give higher variability on the growth rate.



▶ M(0): Small M(0) will increase variability in both Lag time and polymerization speed.

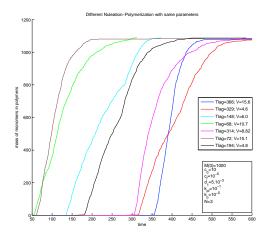


n: Small nucleus increases the variability.





algorithm results



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- Introduction of structure variability.