

On The Asymptotic Distribution of Nucleation Times of Polymerization Processes

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Joint work with Philippe Robert



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Overview

Nucleation phenomenon

Observations from biological experiments

Mathematical literature

Model

A Markovian model for nucleation

Basic Assumptions

The math problem

Results

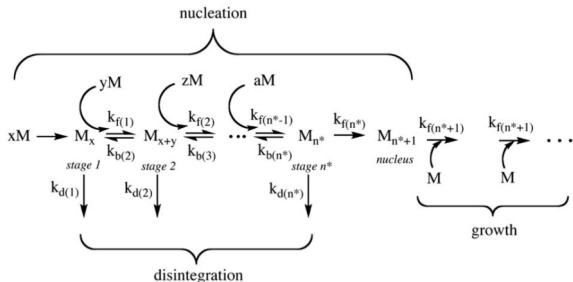
Main Results

Sketch of proofs

Future work and references

Polymerization & Nucleation

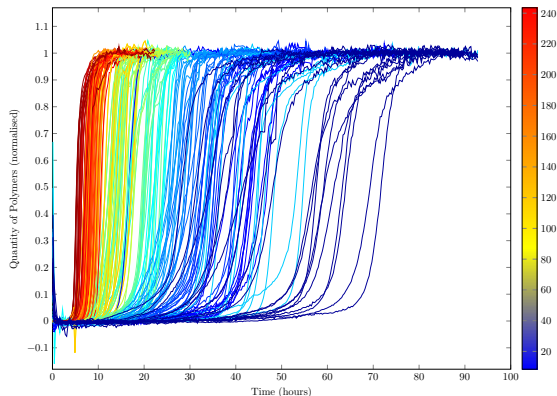
small particles $\xrightleftharpoons[\text{decomposition}]{\text{synthesis}}$ Big (stable) clusters.



- ▶ Physics:
Aerosols...
- ▶ Chemistry:
Polymers/monomers
- ▶ Biology:
Protein/Peptide
v.s. amino acid
monomers

Figure: Flyvbjerg, Jobs, and Leibler's model (96' PNAS)
for the self-assembly of microtubules, retrieved from
Morris et al. (09' Biochimica et Biophysica Acta)

Experiments: large variability in nucleation



Observations:

- ▶ sharp curve;
- ▶ huge variance in time.

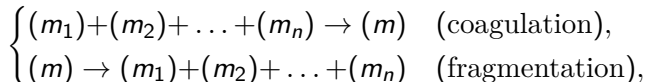
Figure: Experiments for the evolution of polymerized mass.
From data published in Xue et al.(08' PNAS).

Goals of our study

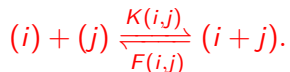
- ▶ Explain **sharp phase transition** in nucleation;
- ▶ Explain **high variance** of the transition moment.

Literature: coagulation and fragmentation models

- ▶ Particles are identified by their sizes.
- ▶ Reactions: for $m = \sum_{i=1}^n m_i$,



- ▶ Binary reaction: **Smoluchowski Model**



where $(F(i,j))$, $(K(i,j))$ are reaction rates.

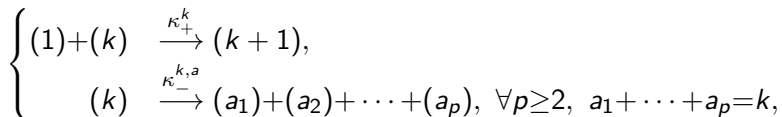
Literature: coagulation and fragmentation models

- ▶ Deterministic studies:
Oosawa et al. (75), Ball et al. (86'), Penrose (89',08'),
Jabin et al. (03'), Niethammer (04') ...
- ▶ Stochastic studies:
Jeon (98'), Durrett et al. (99'), Norris (99'), Ranjbar et al. (10'),
Bertoin (06',17'), Calvez et al. (12'), Sun (18') ...
- ▶ Survey:
Aldous (99'), Hingant & Yvinec (16')

Can not explain the high variance observed in the experiments! (CLT is not enough!)

Model with the nucleus

- ▶ Reaction:



- ▶ Critical Nucleus size: n_c
- ▶ Polymers larger than the nucleus are more stable than the smaller polymers: $\forall s < n_c < l,$

$$\frac{\kappa_-^s}{\kappa_+^s} \gg \frac{\kappa_-^l}{\kappa_+^l}.$$

where $\kappa_-^k = \sum_a \kappa_-^{k,a}$.

Assumptions & Markovian description

- ▶ Only monomers at $t = 0$ with total mass N ;
- ▶ **Scaling assumption:** for two positive sequences (λ_k) , (μ_k) and $\mu > 0$,

$$\kappa_+^k = \lambda_k, \quad \kappa_-^k = \begin{cases} N \mu_k, & \text{if } k < n_c \\ \mu, & \text{if } k \geq n_c \end{cases}$$

- ▶ $U_k^N(t) :=$ number of polymers of size k at time t ;
- ▶ Markov process $(U_k^N(t), k \in \mathbb{N})$ with generator

$$\begin{aligned} \Omega_N(f)(u) &= \sum_{k=1}^{+\infty} \lambda_k u_k \frac{u_1}{N} [f(u + e_{k+1} - e_k - e_1) - f(u)] \\ &+ \sum_{k=2}^{+\infty} (N \mu_k \mathbb{1}_{\{k < n_c\}} + \mu \mathbb{1}_{\{k \geq n_c\}}) u_k \int_{\mathcal{S}_k} [f(u + y - e_k) - f(u)] \nu_k(dy) \end{aligned}$$

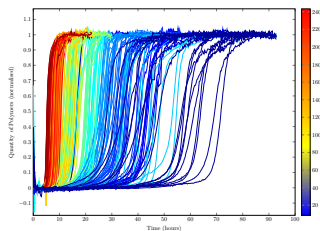
where (ν_k) are fragmentation measures and (\mathcal{S}_k) are the set of all possible fragmentations.

Mathematical Interpretation

- ▶ **Lag time:** for any fraction $\delta \in (0, 1)$,

$$L_\delta^N := \inf\{t \geq 0 : \sum_{k \geq n_c} k U_k^N(t) \geq \delta N\}.$$

- ▶ Observations in terms of lag time:



- ▶ sharp phase transition:
for any $\delta_1, \delta_2 \in (0, 1)$,

$$L_{\delta_1}^N \sim L_{\delta_2}^N$$

- ▶ high variance:

$$\mathcal{O}\left(\sqrt{\text{Var}(L_\delta^N)}\right) \sim \mathcal{O}\left(\mathbb{E}(L_\delta^N)\right)$$

Figure: Xue et al.(08' PNAS).

Main Results (for $n_c \geq 3$)

- ▶ The moment of the first nucleus:

$$T^N := \inf\{t \geq 0 : U_{n_c}^N(t) = 1\}.$$

With high probability, for any $\delta \in (0, \delta_0)$,

$$L_\delta^N \sim \mathcal{O}(T^N + \log(N)).$$

- ▶ For the convergence in probability

$$\lim_{N \rightarrow \infty} \left(\frac{T^N}{N^{n_c-3}} \right) = E_\rho,$$

where E_ρ is an **exponential random variable** with parameter ρ only depends on $(\lambda_k, \mu_k, k \leq n_c - 1)$.

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Therefore, $L_\delta^N \sim \mathcal{O}\left(E_\rho N^{n_c-3} + \log(N)\right).$

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If $n_c > 3$, $\boxed{\frac{L_\delta^N}{N^{n_c-3}} \sim \mathcal{O}(E_\rho)}$ ← Not depends on δ & Large var

- ▶ For the convergence in probability

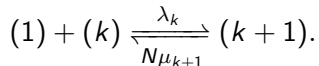
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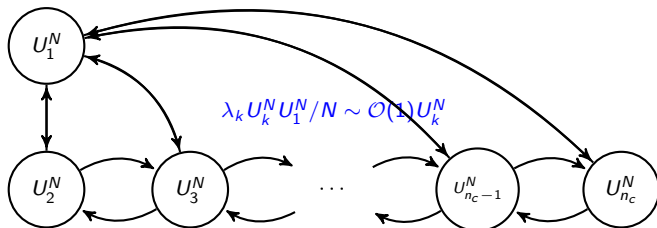
Sketch of proofs (Step I)

Before T^N , $U_k^N(t) \equiv 0$, for all $k > n_c + 1$.

A simple example, Becker-Döring reactions:



fast process



slow process

$$N\mu_{k+1} U_{k+1}^N \sim \mathcal{O}(N) U_{k+1}^N$$

Sketch of proofs (Step I)

- ▶ Distribution of T^N only depends on the fast-slow system $(U_1^N(t), \dots, U_{n_c}^N(t))$.
- ▶ Study the dynamic on the very large time interval $[0, N^{n_c-3}t]$ by using marked Poisson point processes;
- ▶ Main Difficulties:
 - ▶ very large fluctuations
(Time scale N^{n_c-3} v.s. Space scale N).
 - ▶ multi-dimensional stochastic averaging system:
hard to identify the limit of occupation measures
- ▶ Techniques: coupling, flow balance equations...
- ▶ Proofs work for general fragmentation measures under reasonable conditions.

Sketch of proofs (Step II)

After time T^N , by coupling, number of stable polymers

$$(U_{n_c}^N(t), U_{n_c+1}^N(t), \dots)$$

could be lower bounded by a supercritical branching process.

- ▶ The lag time of the branching process is less than $K \log N$ with probability $p_0 > 0$.
- ▶ Therefore, stochastically

$$T^N \leq L_\delta^N \leq \sum_{i=1}^{G_{p_0}} (T_i^N + K \log N)$$

where G_{p_0} is a geometric random variable.

Future work

- ▶ Experiments: fragmentation rates are more likely sublinear for smaller polymers, *i.e.*, for k small,

$$\kappa_-^k = \mathcal{O}(N^\alpha), \text{ for an } 0 < \alpha < 1.$$

See the biology review Morris et al. (09').

- ▶ In the general case $\kappa_-/\kappa_+ \sim \phi(N)$, the nucleation time should be $\mathcal{O}(\phi(N)^{n_c-2}/N)$.
- ▶ Nucleation in a multi-type polymers environment.

References

- ▶ (Eugène, Xue, Robert & Doumic, 16') Insights into the variability of nucleated amyloid polymerization by a minimalistic model of stochastic protein assembly. ([Journal of Chemical Physics](#))
- ▶ (Domic, Eugène & Robert, 16') Asymptotics of stochastic protein assembly models. ([SIAM Journal on Applied Mathematics](#))
- ▶ (Robert & Sun, 17') On the Asymptotic Distribution of Nucleation Times of Polymerization Processes. ([arXiv:1712.08347](#))

Thank you!