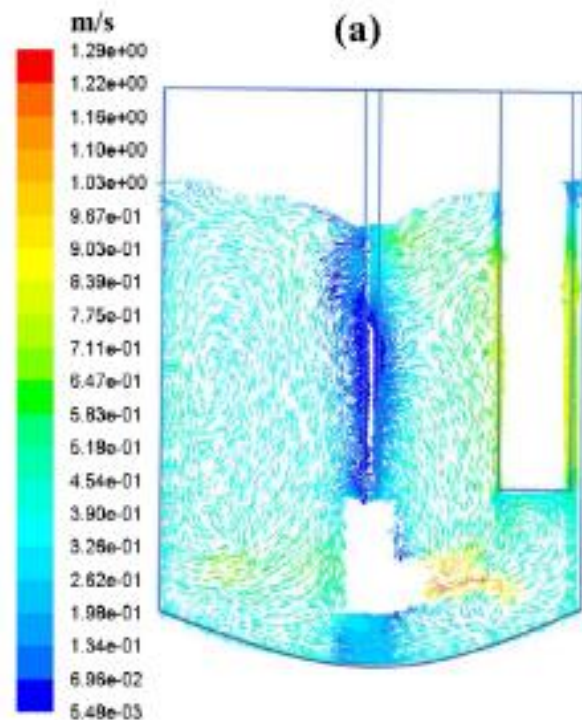


# The nucleation of molecules in solution using shear flow

# Flow conditions in industrial crystallization processes

## CSTR (also true for MSMPR) [1]

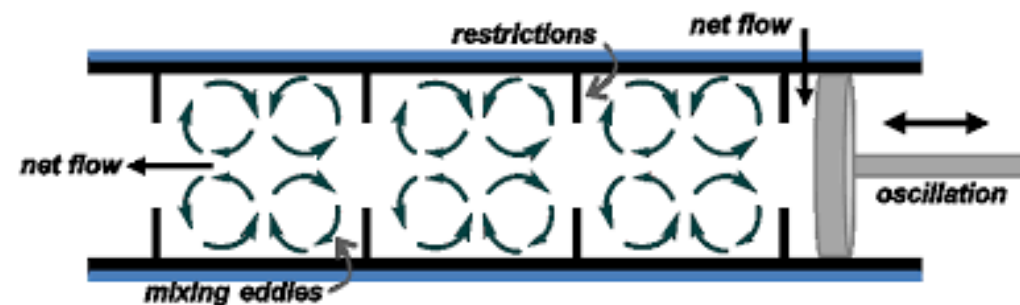


⇒ Inhomogeneous

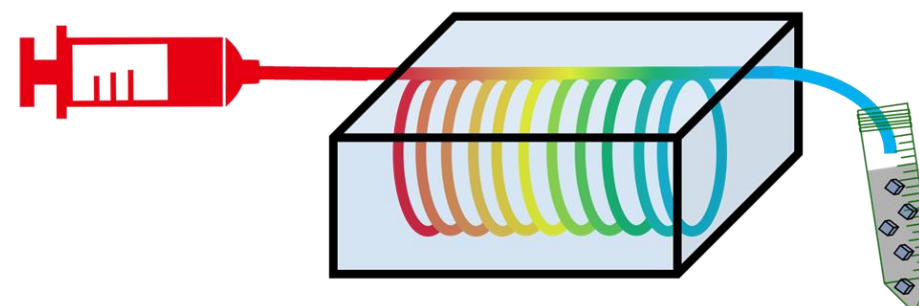
Homogenizing

Controlling

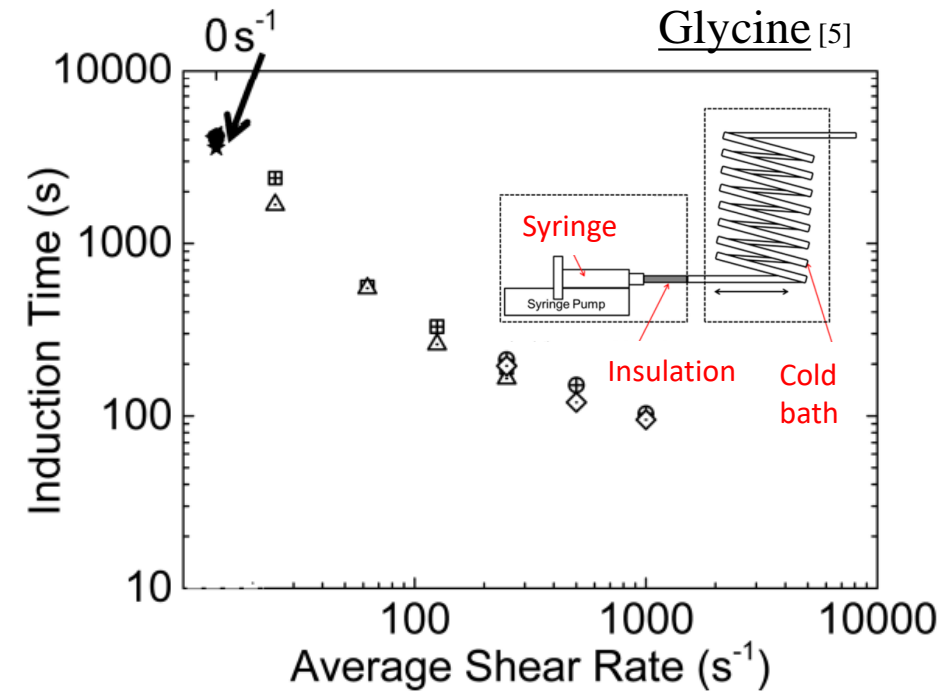
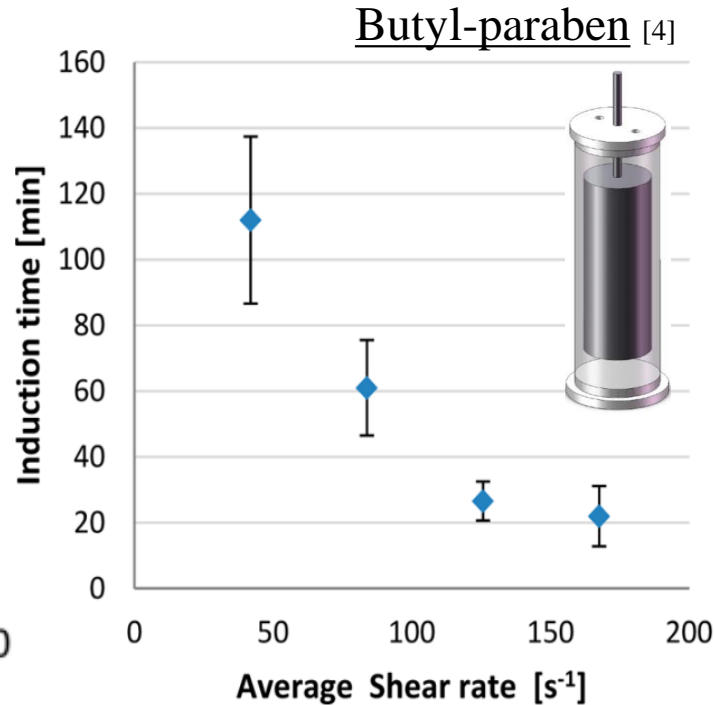
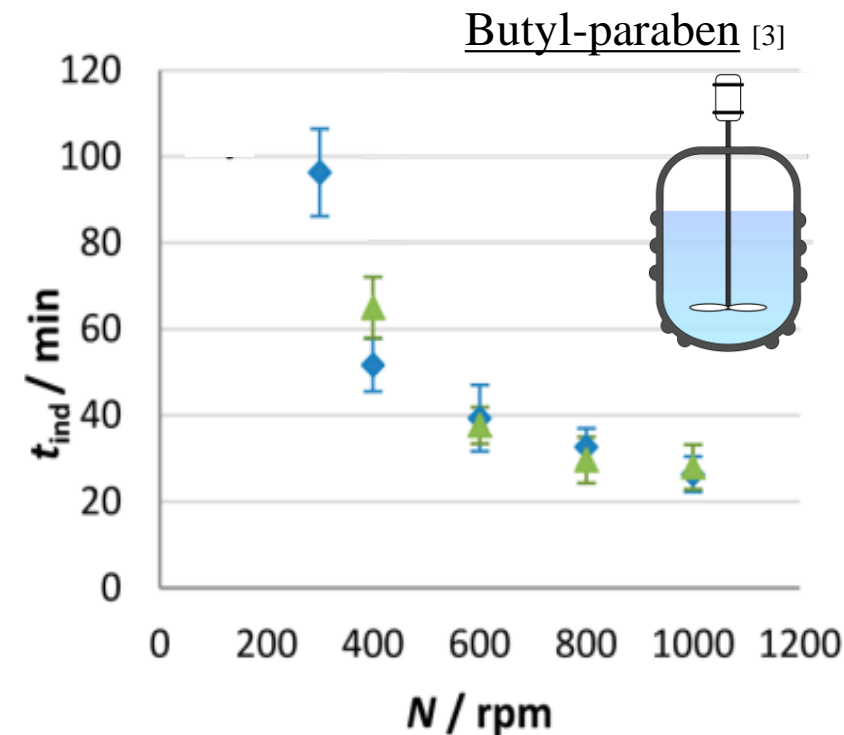
## COBRC [2]



## LFC



# Flow behaviour and crystallization

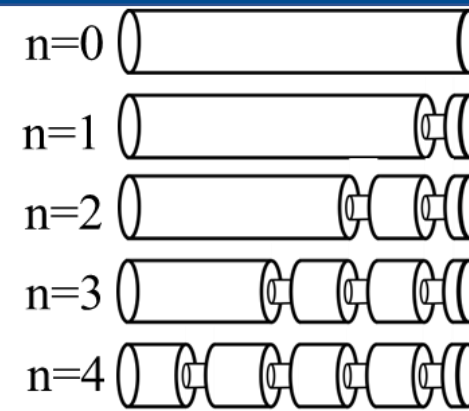
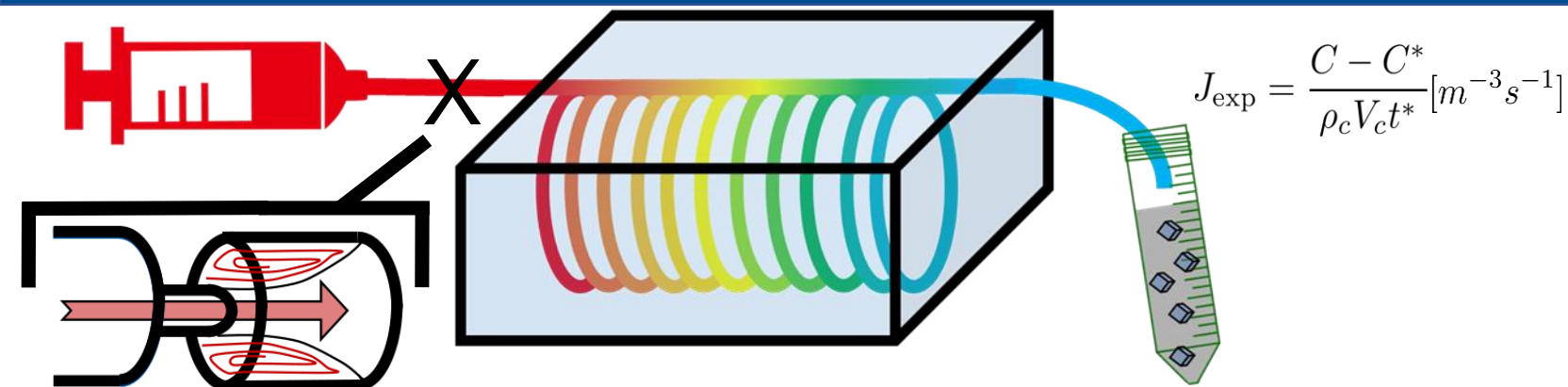


[3] Liu, J., Svard, M., & Rasmuson, Å. C. (2015). *Crystal Growth & Design*, 15(9), 4177-4184.

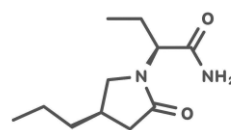
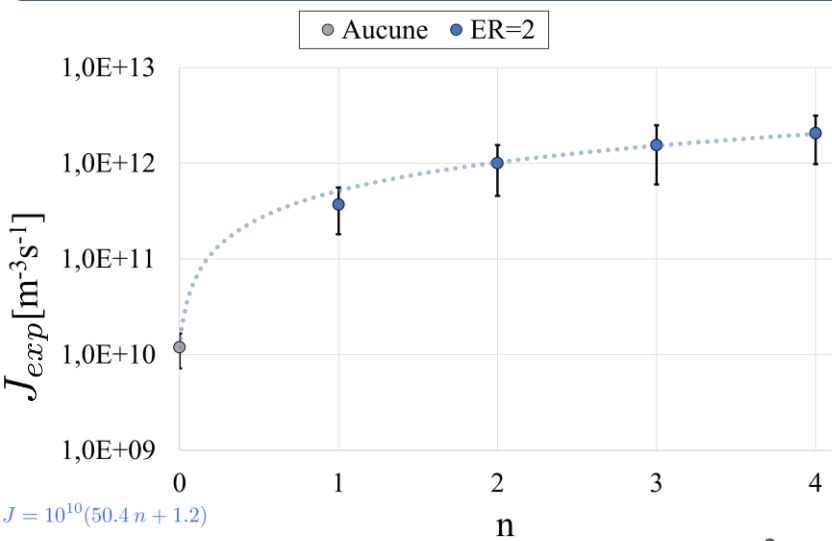
[4] Liu, J., & Rasmuson, Å. C. (2013). *Crystal growth & design*, 13(10), 4385-4394.

[5] Forsyth, C., Mulheran, P. A., Forsyth, C., Haw, M. D., Burns, I. S., & Sefcik, J. (2015). *Crystal Growth & Design*, 15(1), 94-102.

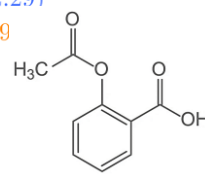
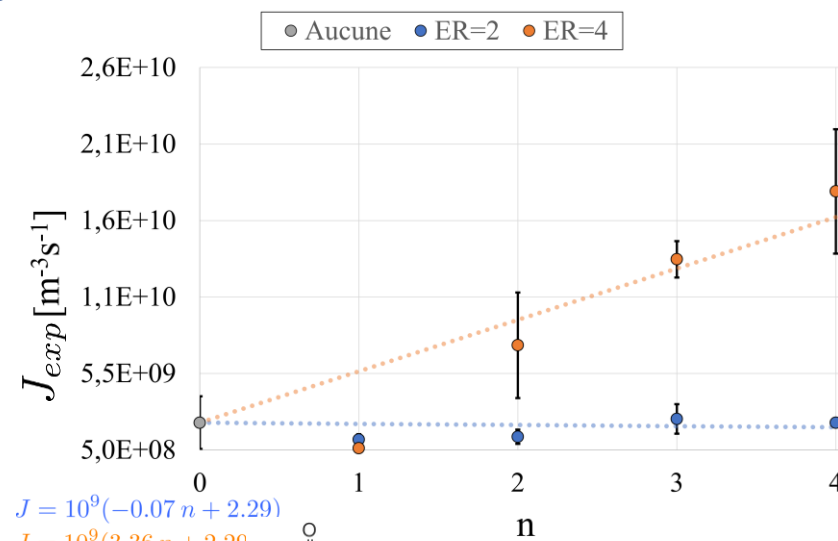
# Introduction



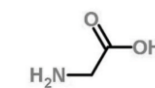
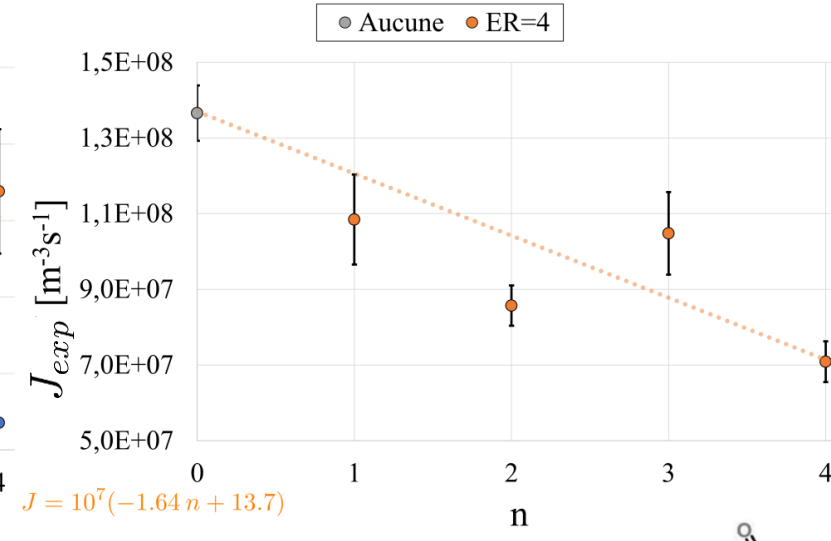
Brivaracetam [6]



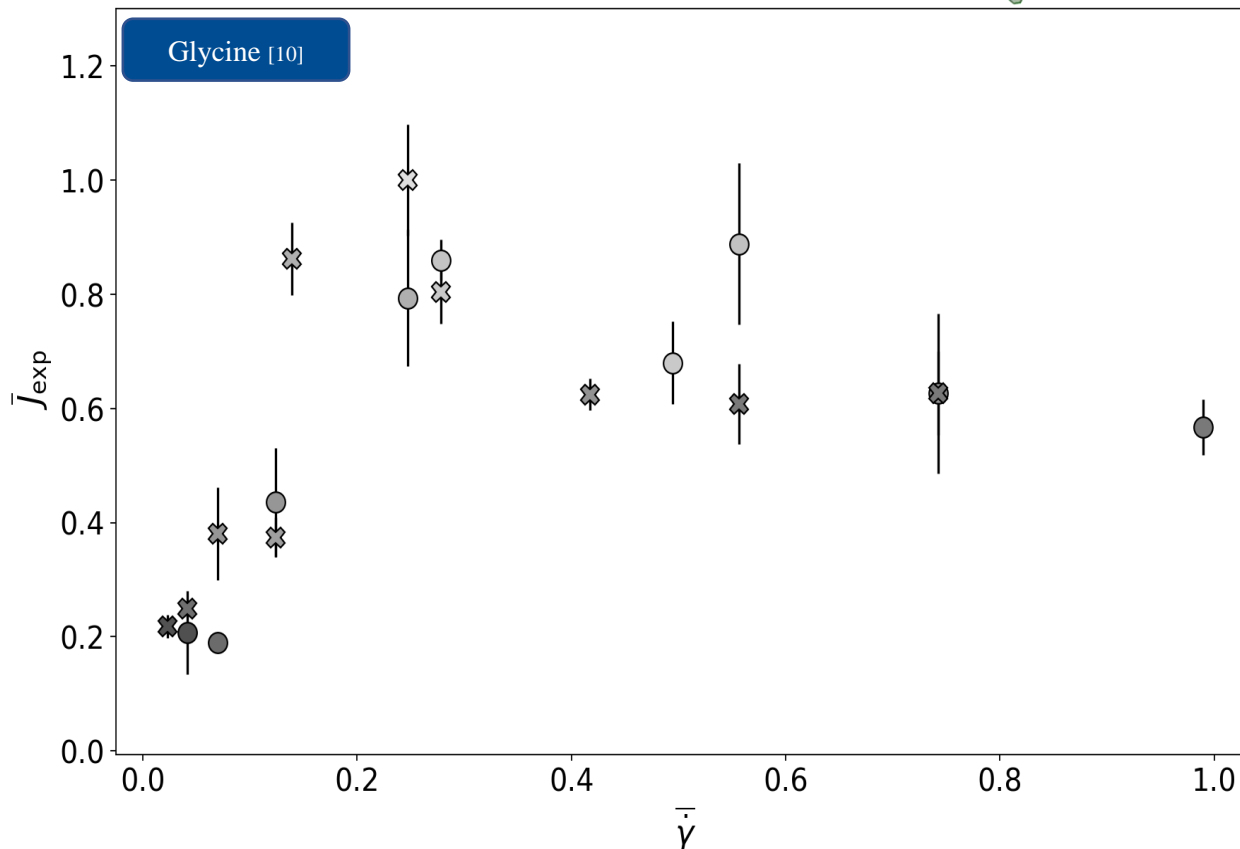
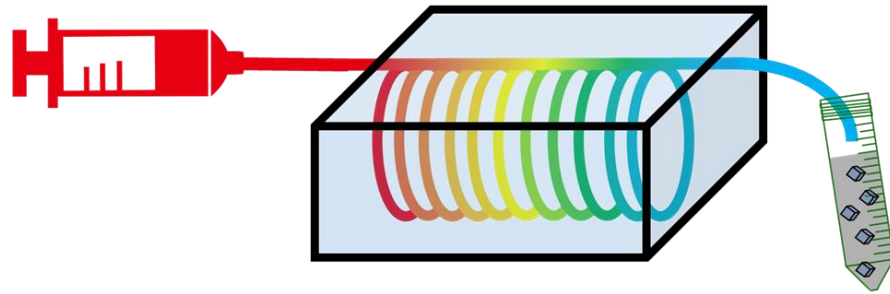
Aspirin [7]



Glycine [8]



# Experimental shear-rate analysis



In crystallizer **WITHOUT** diameter constriction, changing

- $Q$  : flow rate
- $r$  : internal radius
- $L$  the length to keep the residence time constant

Shear rate :

$$\dot{\gamma} = \frac{4Q}{\pi r^3}$$

CNT with shear [9]

$$J = N_t j Z e^{-\frac{F(R^*)}{k_b T}} [m^{-3} s^{-1}]$$

$$\Rightarrow J_{th} = \frac{J}{N_t} = j Z e^{-\frac{F(R^*)}{k_b T}} [s^{-1}]$$

# Theoretical shear-rate analysis

$$J_{\text{th}} = \frac{J}{N_t} = jZe^{-\frac{F(R^*)}{k_b T}} [s^{-1}]$$

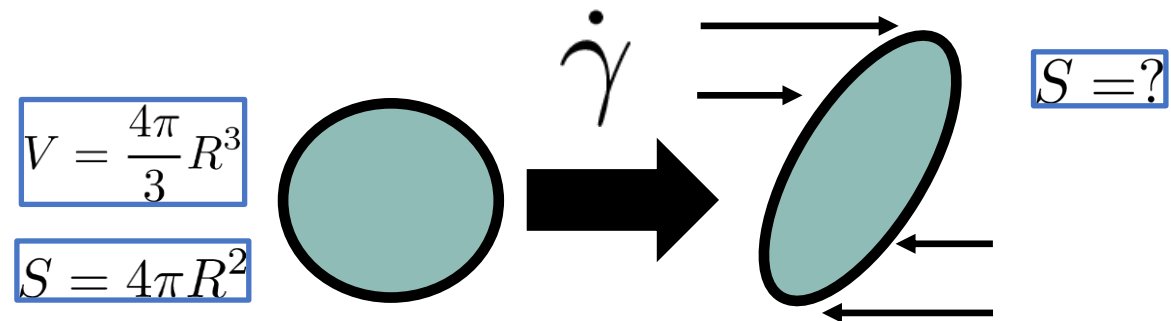
Free energy [9]

From the classical expression as spherical nucleus :

$$F(R) = -VF_v + S\nu$$

1) Shear induces an increase of the internal energy due to stretching

2) Volume is conserved but not the surface !

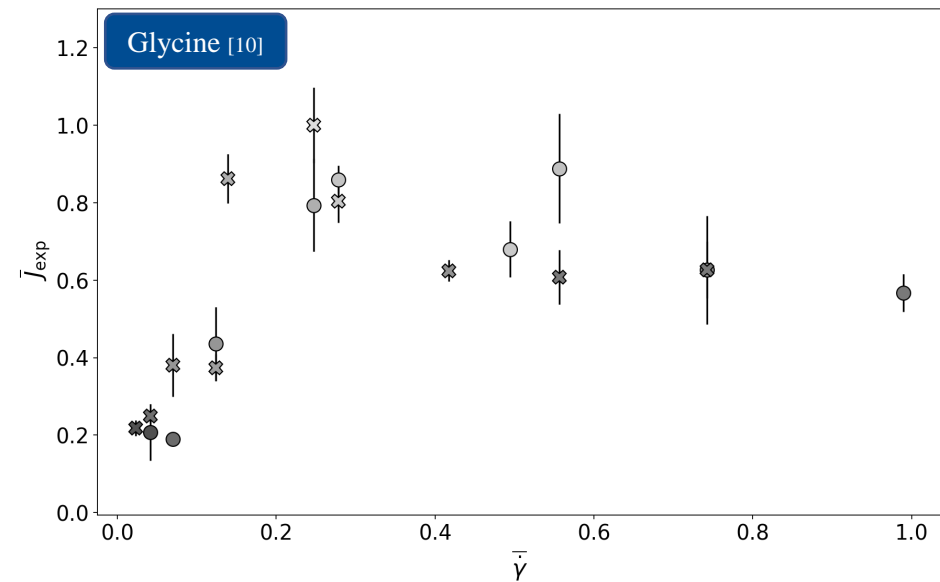


$F(R^*) \propto \dot{\gamma}$  : Increase of the energy barrier with shear rate (unfavorable)

Successful aggregation rate [9]

$Zj \propto \dot{\gamma}$  : Mass transfer enhancement with shear rate (favorable)

# Theoretical shear-rate analysis

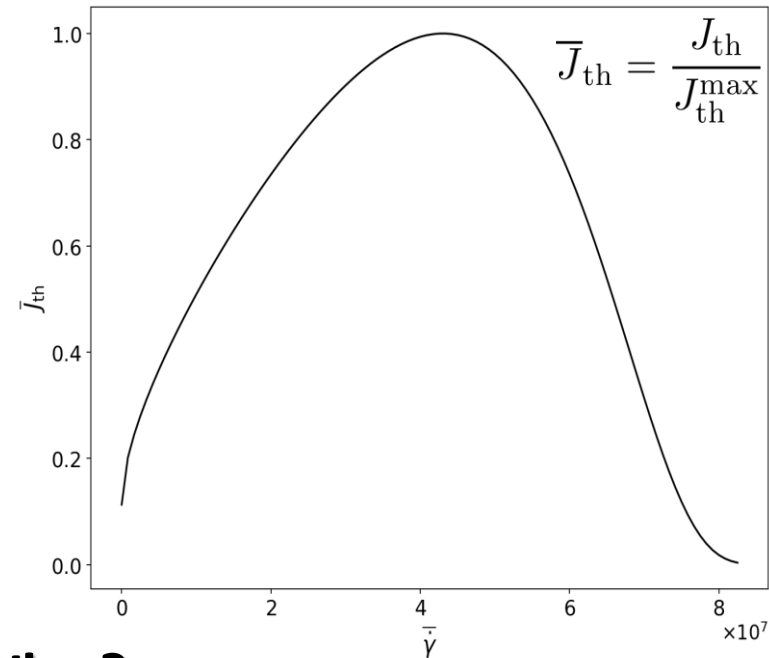
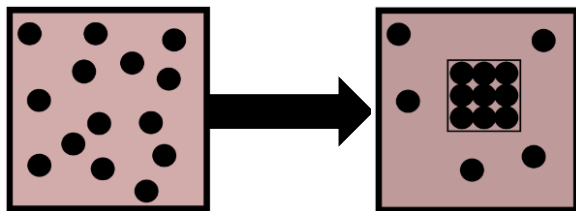


With :

$$\bar{J}_{\text{exp}} = \frac{J_{\text{exp}}}{J_{\text{exp}}^{\text{max}}}$$

and

$$\bar{\dot{\gamma}} = \frac{\dot{\gamma}}{\dot{\gamma}_{\text{max}}}$$



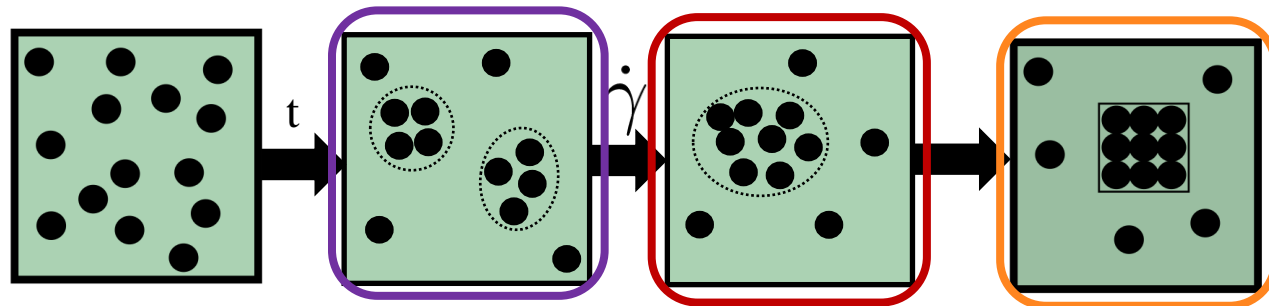
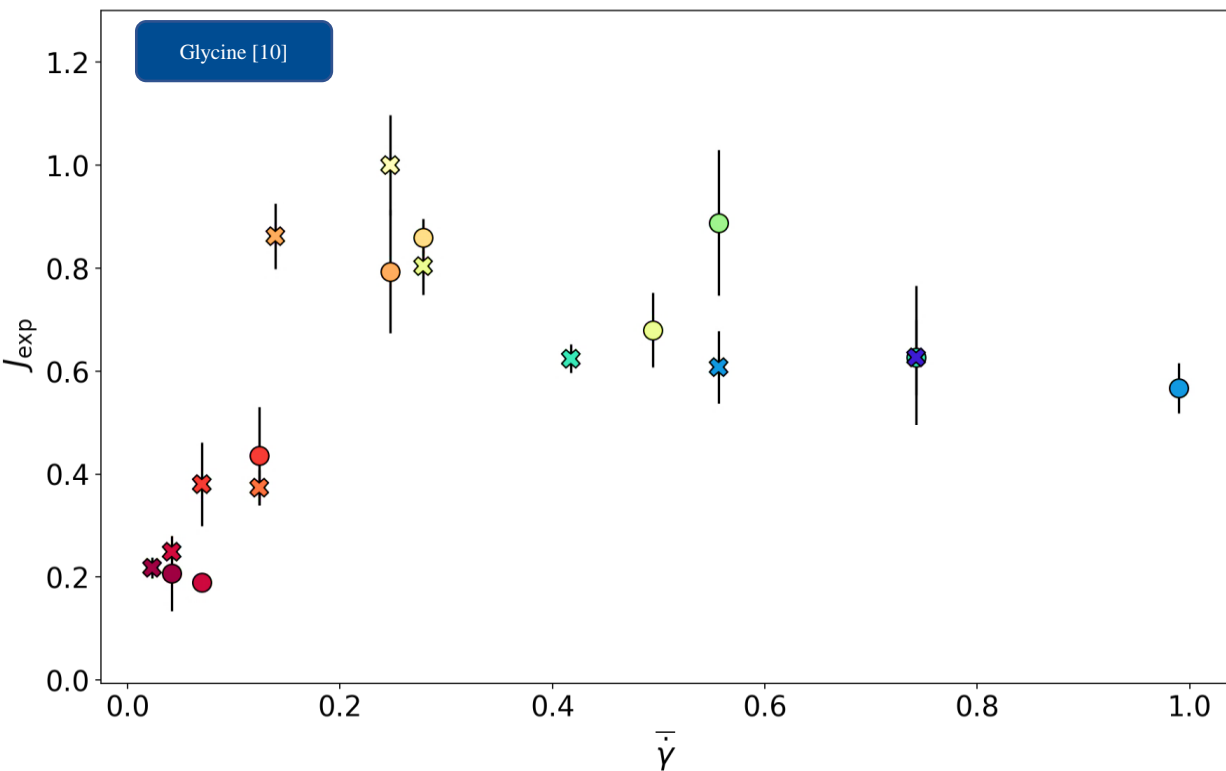
For molecular aggregation  
 $\Rightarrow$  Not working [10]

## Why ?

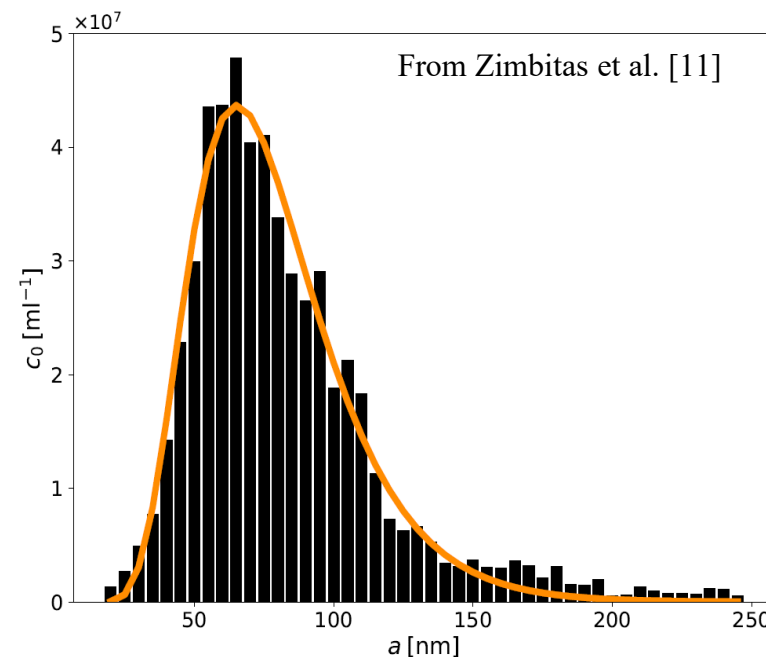
- Low nucleus scale (  $\sim 1$  nm [10])
- Rigid glycine nucleus ( $G \sim 10^{10}$  Pa [11])

$\Rightarrow$  Deformation effect  $\ll$  mass transfer effect

# Theoretical shear-rate analysis



⇒ Molecular assemblies (clusters) exist in undersaturated conditions ! [11]



Distribution population nearly independent of the supersaturation :

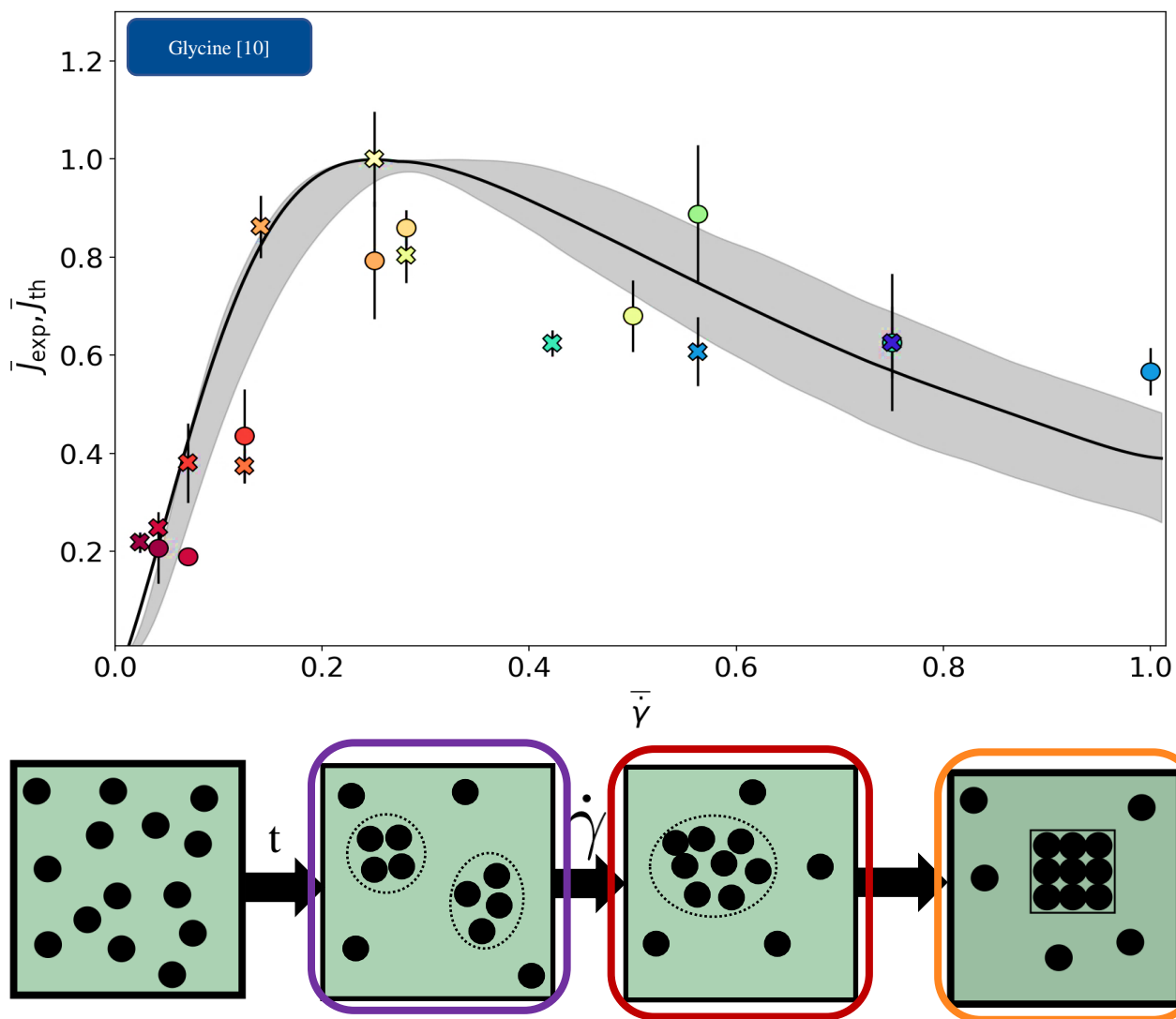
- Mean radius ~ 80 nm

⇒ Under agitation conditions bigger cluster are created : radius > 250 nm [12]

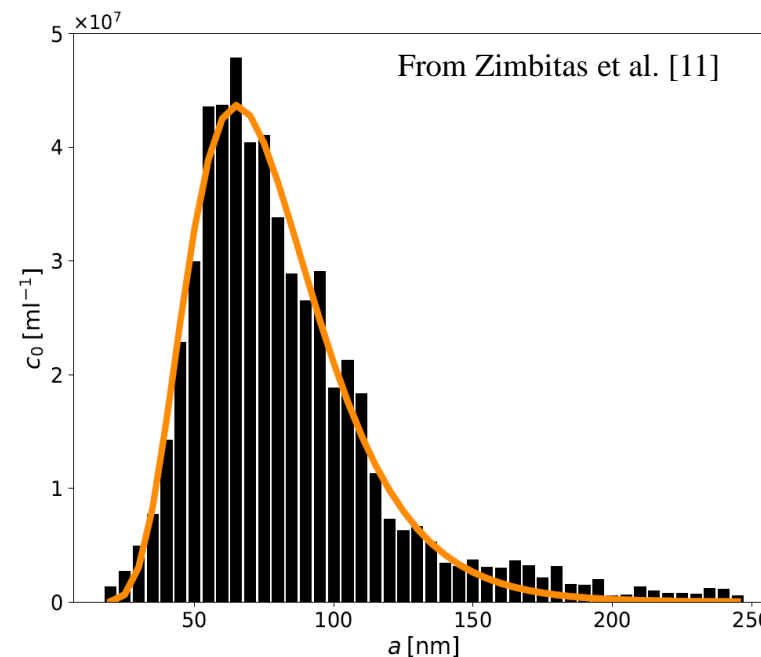
⇒ When big clusters are created, a huge decrease of the induction time is observed [12]



# Theoretical shear-rate analysis



⇒ Molecular assemblies (clusters) exist in undersaturated conditions ! [11]



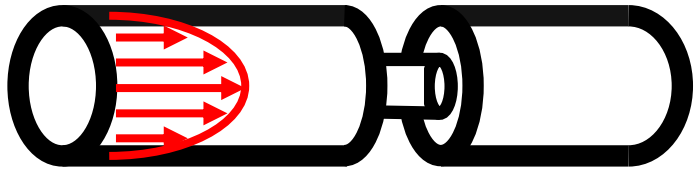
Distribution population nearly independent of the supersaturation :

- Mean radius ~ 80 nm

⇒ Under agitation conditions bigger cluster are created : radius > 250 nm [12]

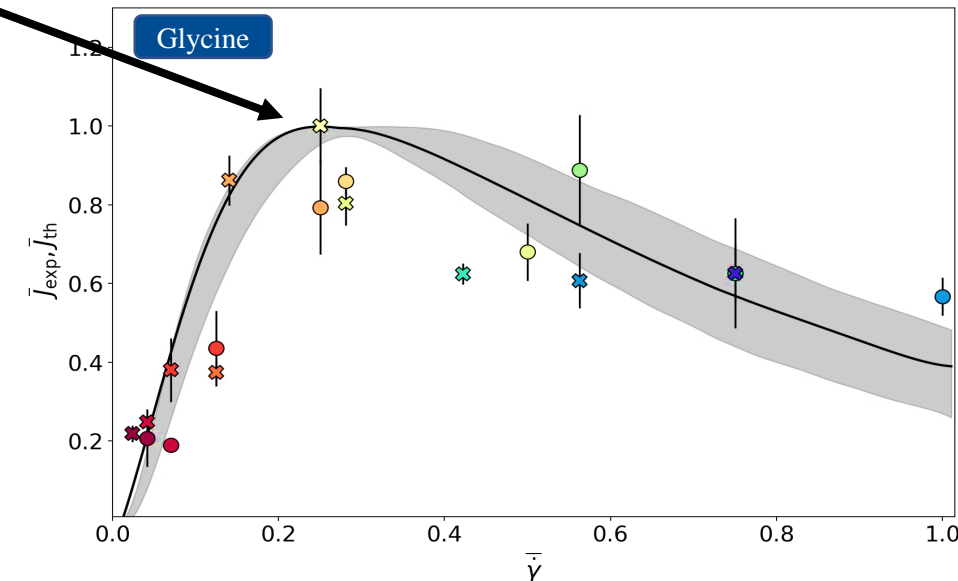
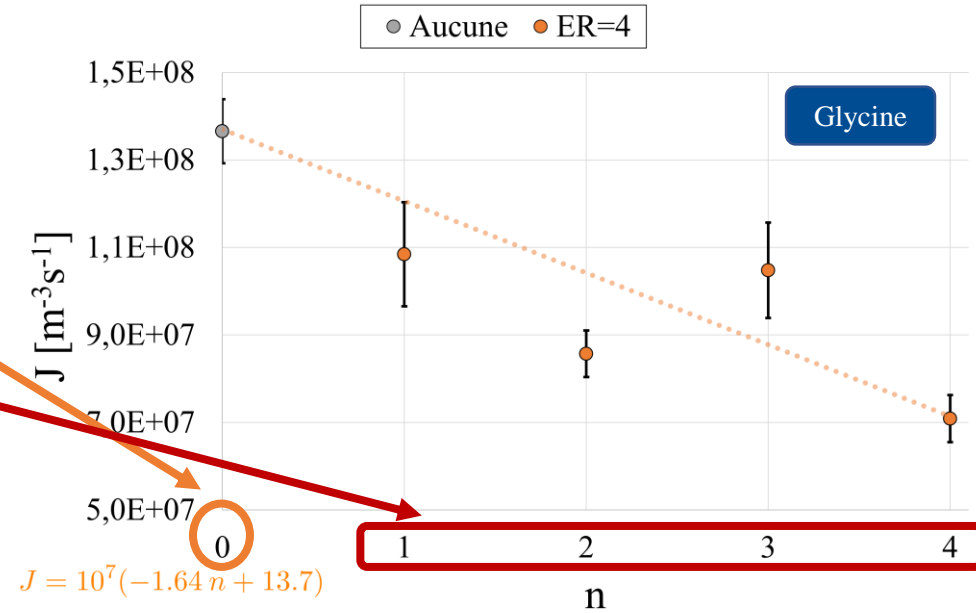
⇒ When big clusters are created, a huge decrease of the induction time is observed [12]

# Conclusion

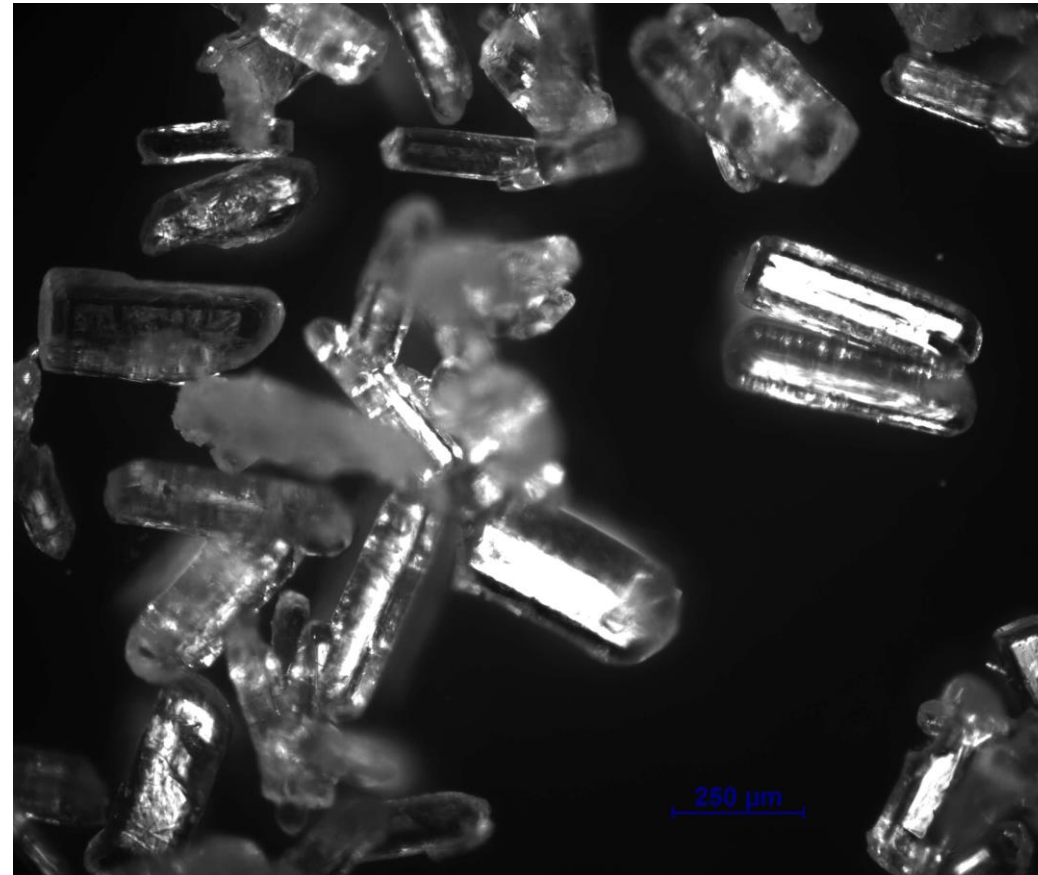


Initial point realized at  
~ 5 000 [s<sup>-1</sup>]

- Adding supplementary shear through the constrictions  
⇒ drop of nucleation after 3 000 [s<sup>-1</sup>]
- Brivaracetam, similar structures already observed [13].
- What about Aspirin ?
- Brivaracetam ≠ Glycine ≠ Aspirin :  
Different physico-chemical properties ( $G$ ,  $v$ ,  $\Delta\mu$ ) of the clusters  
⇒ change the position of the optimum ?



Thanks !



# Temperature verification

