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**Supporting information for article:**

**On some aspects of crystallization process energetics, logistic new-phase nucleation kinetics, crystal size distribution, Ostwald ripening**

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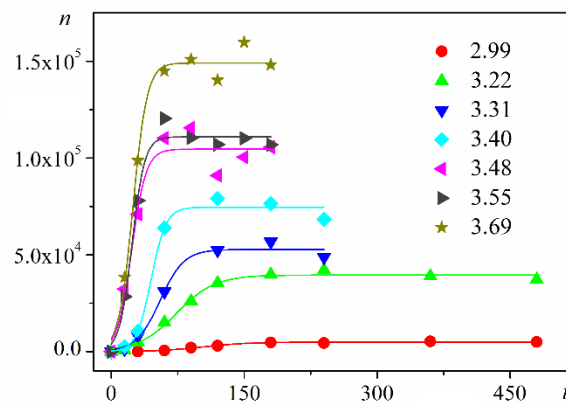
## SUPPORTING INFORMATION

### I. Original experimental data fitted to logistic dependence:

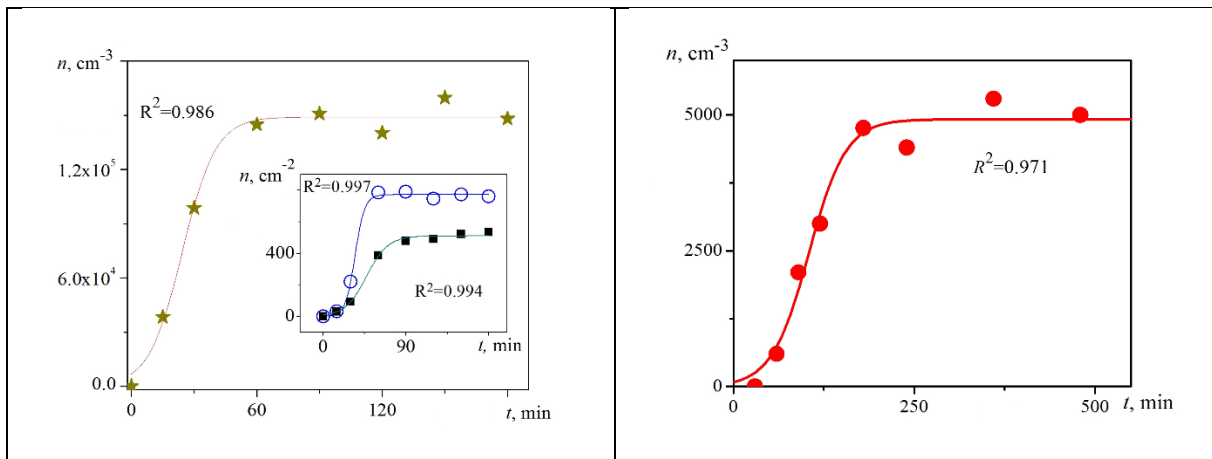
$$n = \frac{n_s}{1 + \exp[-k(t - t_c)]}$$

where  $n$  [ $\text{cm}^{-3}$ ] is nuclei number density,  $n_s$  is saturated nuclei number density,  $k$  [ $\text{s}^{-1}$ ] is adjusting coefficient,  $t$  is time, and  $t_c$  being the point when  $n = n_s/2$ .

A. Insulin crystal nucleation (insulin from BioChemika,  $\geq 85\%$  (GE),  $\sim 24$  IU/mg).



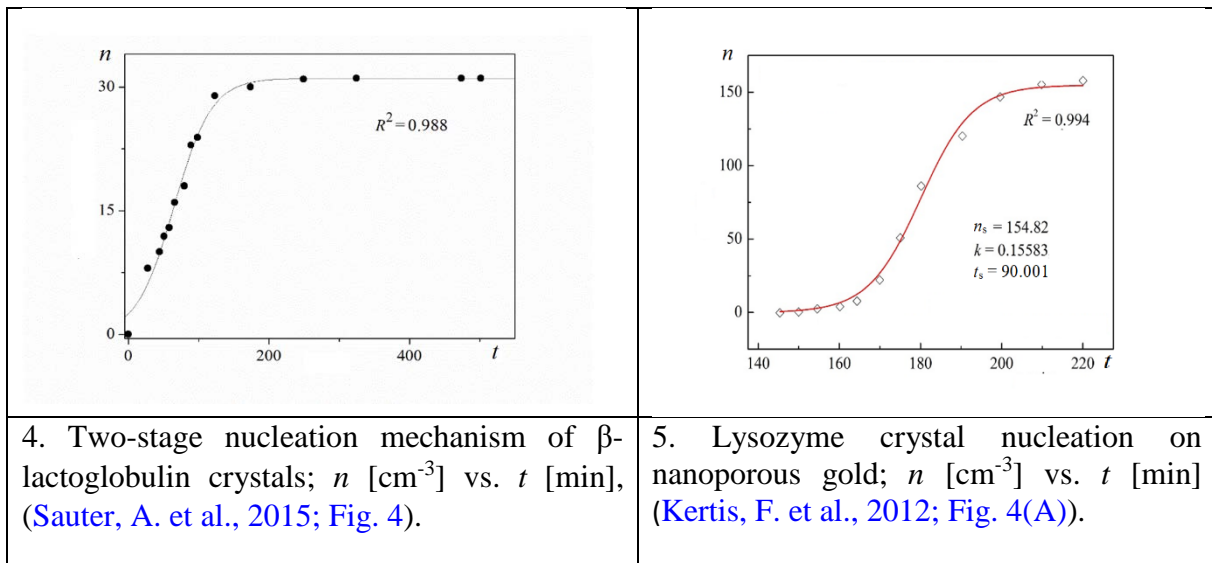
1. Plots of experimental  $n$  [ $\text{cm}^{-3}$ ] vs.  $t$  [min] data for insulin crystal nucleation in bulk solution under series of supersaturations. Color symbols denote dimensionless supersaturations,  $\ln(c/c_e)$ , where  $c$  is the actual insulin concentration,  $c_e$  is the equilibrium one (Nanev, C. N. et al., 2011).



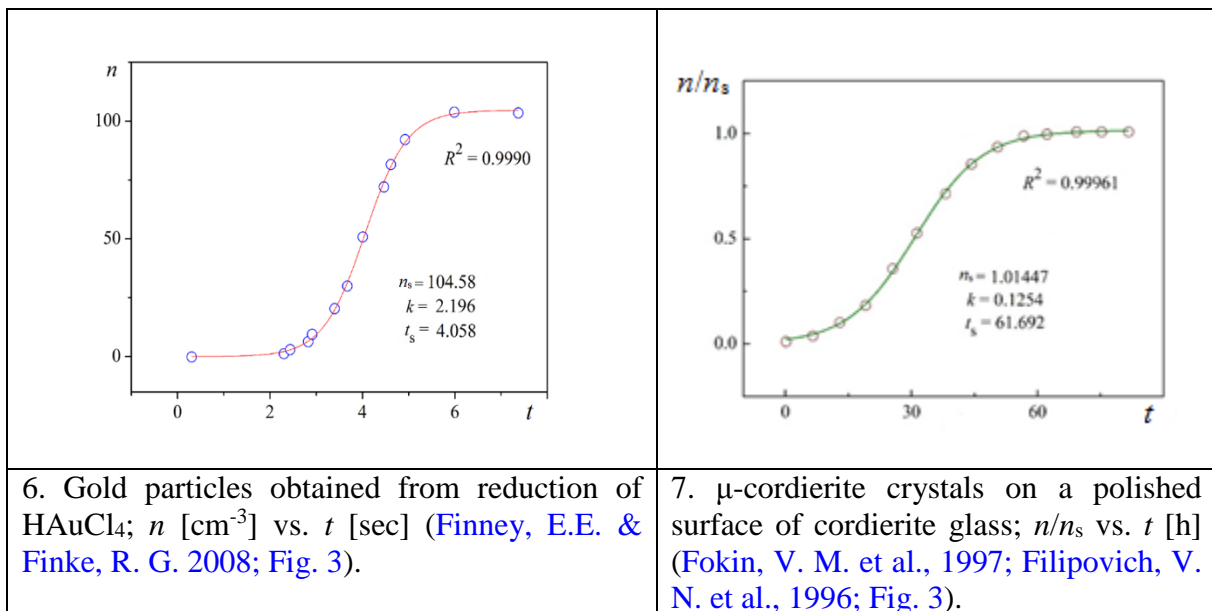
2. Plots of  $n$  vs.  $t$ , measured for insulin crystal nucleation at supersaturation 3.69: stars represent data for nucleation in bulk solution; the inserted graph shows nucleation on glass surface (open circles) and nucleation on the air/solution interface, black squares (Nanev, C. N. & Tonchev, V. D., 2015; Fig. 4).

3. Plot of  $n$  vs.  $t$  data for insulin crystal nucleation in bulk solution at dimensionless supersaturation 2.99 (Nanev, C. N. & Tonchev, V. D., 2015; Fig. 3a).

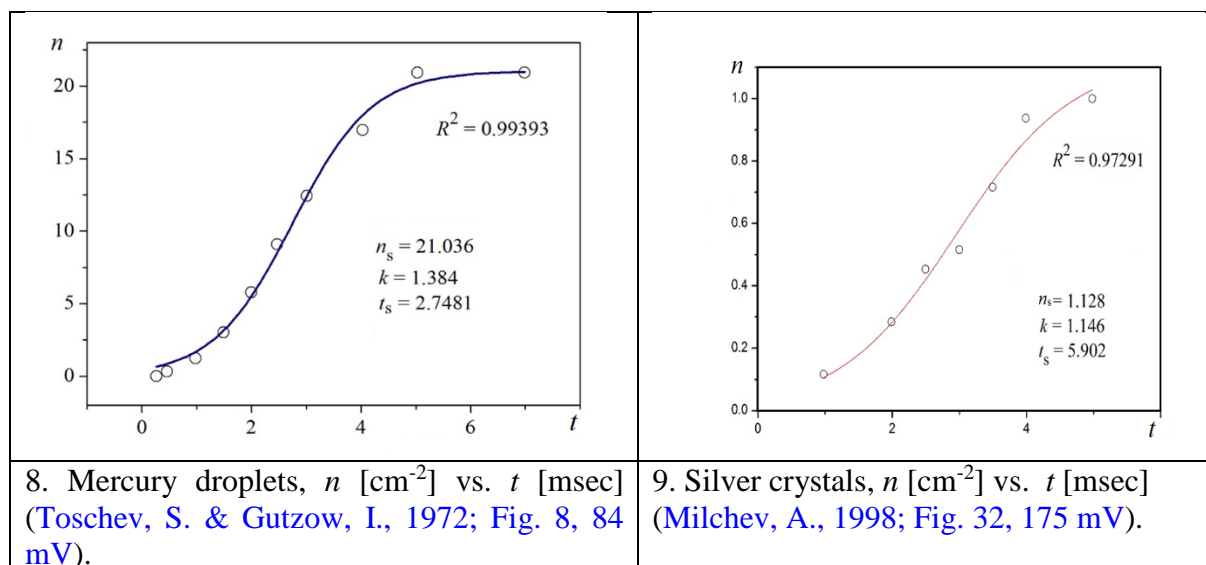
## B. Crystal nucleation of other proteins



## C. Crystal nucleation in other systems



## D. Electrochemical new-phase nucleation



**Conclusion:** Four nucleation kinetic time-scales are observed. The fastest is the electrochemical new-phase nucleation, [msec]. Crystal nucleation resulting from chemical reaction is a little bit slower, [sec]; the protein crystal nucleation proceeds on time-scale of minutes, while the most slower is the glass crystal nucleation, [h].

## Literature

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