

## 5. Crystal morphology

### 5.1 Introduction

shape, form, habit, morphology

#### - “*Equilibrium morphology*”

Crystal shape equilibrated with surroundings to minimize the surface free energy of crystals.

#### - “*Growth morphology*”

Crystal shape developed in the course of crystal growth.

### 5.2 Predicting crystal morphology

- **Attachment energy  $E_{att}$**  energy released per mole of layer added to a crystal surface, is directly proportional to the growth rate of a given face.

#### - **Bravais-Friedel-Donnay-Harker (prediction for growth morphology)**

The binding energy between crystal planes is inversely proportional to the interplanar spacing. As such, the growth rate  $v_{hkl}$  of (hkl) face is proportional to  $1/d_{hkl}$ .

#### - **Specific force field (molecular mechanics) (prediction for growth morphology)**

The interaction energy  $E_{ij}$  between the individual  $i$ -th and  $j$ -th non-bonded atoms of the molecules that constitute the crystal calculated in terms of van der Waals and electrostatic contributions.

$$E_{ij} = -\frac{A}{r_{ij}^6} + \frac{B}{r_{ij}^{12}} + \frac{q_i q_j}{\epsilon r_{ij}}$$

#### - **Atomistic lattice simulation (prediction for equilibrium morphology)**

$$\gamma_{hkl} = 0.5(E_{hkl}^B - 2E_{hkl}^S)$$

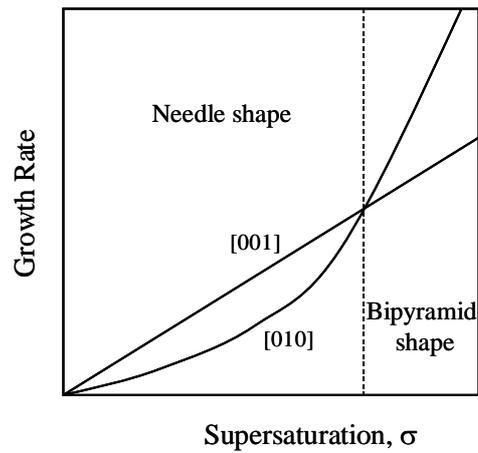
where  $E_{hkl}^B$  and  $E_{hkl}^S$  are energy of crystal plane (hkl) in bulk and surface, respectively.

### 5.3 Influence of supersaturation, temperature, solvents, additives, impurities.

#### - Supersaturation

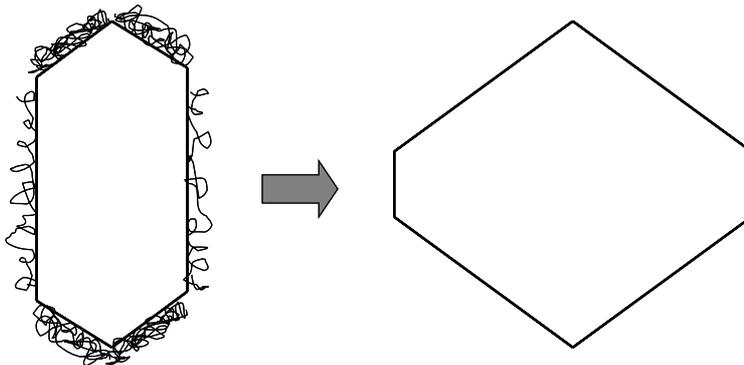
Growth rate of each crystal face depending on the supersaturation brings about the morphological change with supersaturation level.

Ex) paracetamol crystal



#### - Additives & Impurities

Selective adsorption of impurities on the crystal surface will modify the crystal morphology.



Langmuir isotherm (*Chem. Eng. Comm.*, Vol. 120, 119 (1993), *Chem. Eng. Sci.*, 55, 733-747 (2000, *Crystal Research & Technology*, Vol. 40(6), 586-592 (2005)))

$$\frac{L - L_0}{L_{eq} - L_0} = \frac{K_{ad} C_{add}}{1 + K_{ad} C_{Add}}$$

$$v_i = v_\infty (1 - 2r_c / l)^{1/2}$$

where  $r_c$  is the critical radius of two-dimensional nucleus and  $l$  is the distance separated between impurities adsorbed on crystal surface.

- **Some additive** is integrated into growth site of crystal, disrupting further growth.

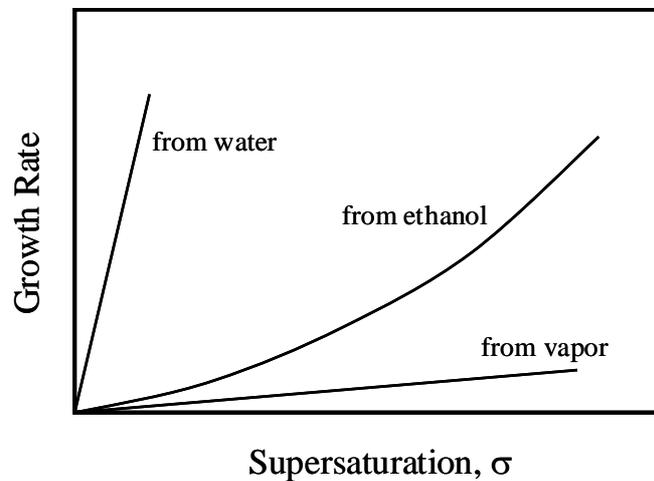
To perform this function effectively, the additive molecule must resemble the crystallizing molecules while containing small difference in stereochemistry or functionality.

→ “tailor-made additives”.

- **Solvent**

The solvent will modify  $\alpha$  in the solution crystallization.

$$\alpha = \xi \left[ \frac{\Delta H_f}{RT} - \ln x_{eq} \right]$$



## 5.5 Crystal purity

agrochemicals, pharmaceuticals, polymer intermediates, electronic materials.

- **Degree of purification ( $D_i$ )**

$$D_i = \frac{\text{mass of impurity per mass of solute in crystal}}{\text{mass of impurity per mass of solute in solution}}$$

If  $D_i > 1$ ,

no purification by crystallization.

If  $D_i < 1$ ,

purification by crystallization.

- In general, the extent of incorporation of an impurity or additive will depend on

- the relative size
- the stereochemistry
- the functionality

of additive molecules when compared with crystallizing solute.

- The more dissimilar the components, the purer the crystals are likely to be!!!