



Assembly Principles in Two Dimensional Ordered Virus Arrays

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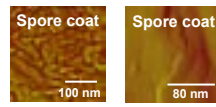
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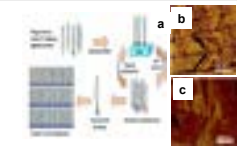
Introduction

Motivation: Scientific & technological challenges associated with macromolecular assembly

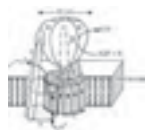
Physics of protein aggregation



Bio-inspired synthesis of materials



Formation of protein machines



Protein structure determination



Cowpea Mosaic Virus (CPMV) is an ideal system to investigate the assembly with nanoscale materials



Unique characteristics

- Robust – stable in 60 °C water
- Icosahedron shape + well-studied crystal structure
- Multiple-charged sites to coordinate crystallization
- Can be produced in grams per harvest

Objectives

- Study the 2-D assembly of ordered CPMV arrays induced by evaporation of electrolytes
- Identify the effect of surfactants in controlling the interaction strength and configurations between virions
- Determine assembly principles: Predict packing configurations

Experimental Method

Drop-dry experiments to determine appropriate sets of parameters for ordered packing

1. Deposit virus solution on mica

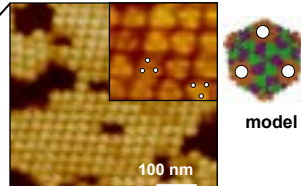
Virus solution:
30 μ l volume, 0.50 mg/ml virus,
2 mM phosphate buffer pH = 7

Vary solution composition:
- Virus concentration
- Surfactant composition
(6kDa polyethylene glycol, PEG)

2. Dry under a controlled environment

Petri dish

3. Image films by atomic force microscopy



Results

2D arrays on mica show effect of interaction potential on degree of order

Dependence of packing on % PEG

Evolution of packing

PEG: 0 %

Low order:
ranging from square to rhombic

PEG: 0.0002 %

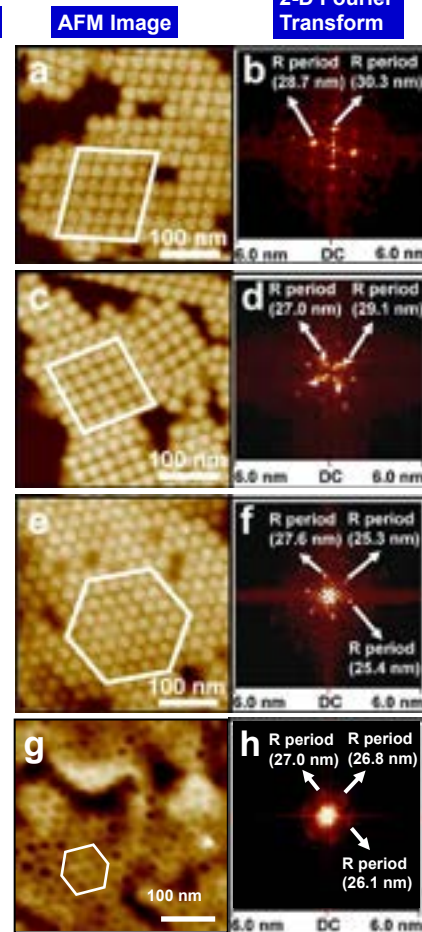
Better order:
mostly rhombic,
some square

PEG: 0.001 %

Well ordered:
hexagonal

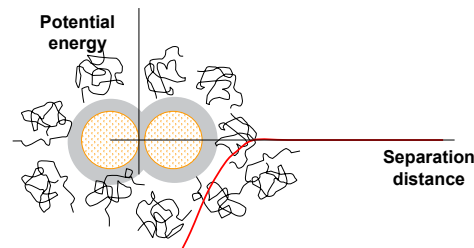
PEG: 0.04 %

PEG co-deposited
prevents imaging



Discussions

Addition of polymers can dramatically change the interaction potentials on macromolecular assembly

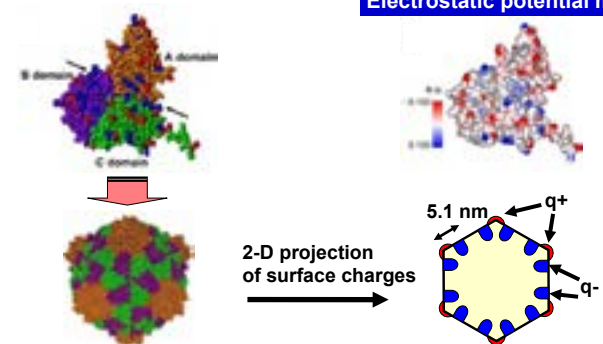


Addition of surfactant polymers (typically PEG):

- Introduces a non-specific, depletion mediated attraction
 - When particles are close, absence of PEG between the particles creates an osmotic pressure gradient to promote aggregation of particles
- Increase the nucleation rate of proteins
- Size and concentration of polymer controls range of attraction

Analysis of amino acids groups on virus capsids provides links to their charge potential landscape

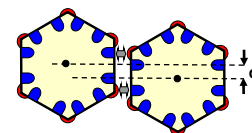
Distribution of the negatively (Asp/Glu) and positively (Arg/Lys/His) charged amino acid residues at pH 7



Steric & electrostatic complementarity of two neighboring CPMV capsids

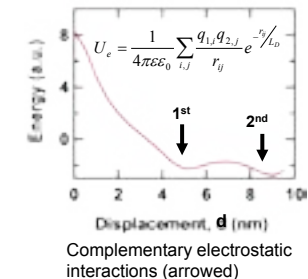
Interaction Model:
Two neighboring CPMVs

Hexagons with -ve charged protrusions & +ve charged patches

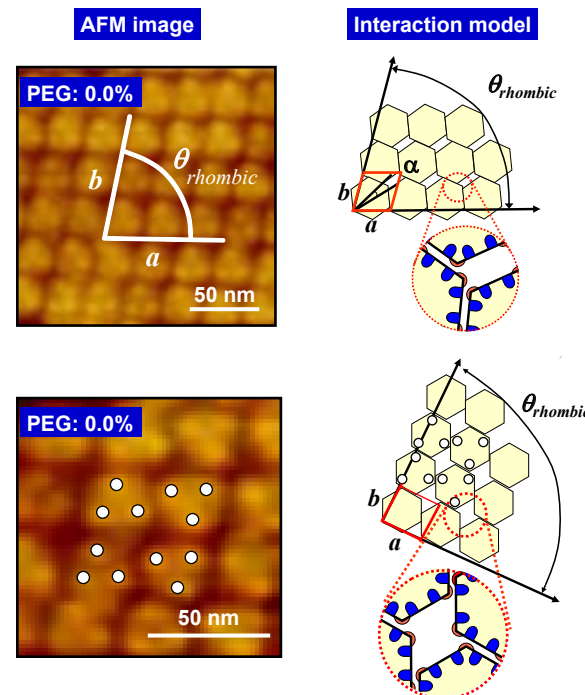


d: transverse shift of the position of the symmetry axes through the center of the projected images

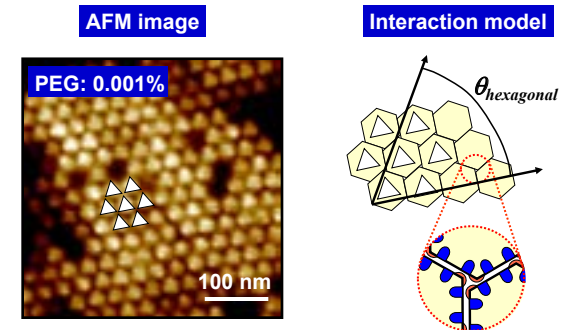
Electrostatic potential energy landscape of neighboring CPMVs



Steric & electrostatic model for 2-D CPMV assembly

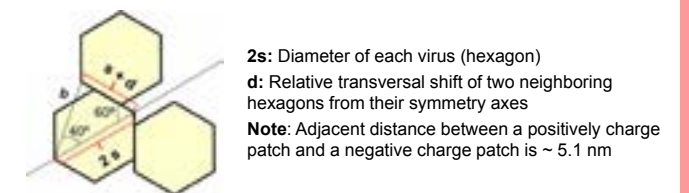


Steric & electrostatic model for 2-D CPMV assembly



Deduction of the average size of a CPMV virion from the 2D array AFM data

Hexagon representation of a virus array



Average size of a CPMV virion and the transversal shift (d) between two virions obtained from the 2D CPMV arrays data

PEG concentration in the virus solution for the array formation (wt. %)	0	0.0002	0.001
Average size of a CPMV virion in the array, 2s (nm)	30.2	30.2	28.6
Average transversal shift, d, between two adjacent virions (nm)	5.88	5.88	5.15

Conclusions

- Electrostatic potential and geometric arrangement of charge patches on CPMV capsids can be used to deduce interaction configurations between neighboring CPMVs.
- Postulated electrostatic and steric complementarity principles explain three different observed 2-D CPMV assembly configurations.
- Steric and electrostatic complementarity principles can be applied to design and assemble other complex biological building blocks.

Acknowledgements

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