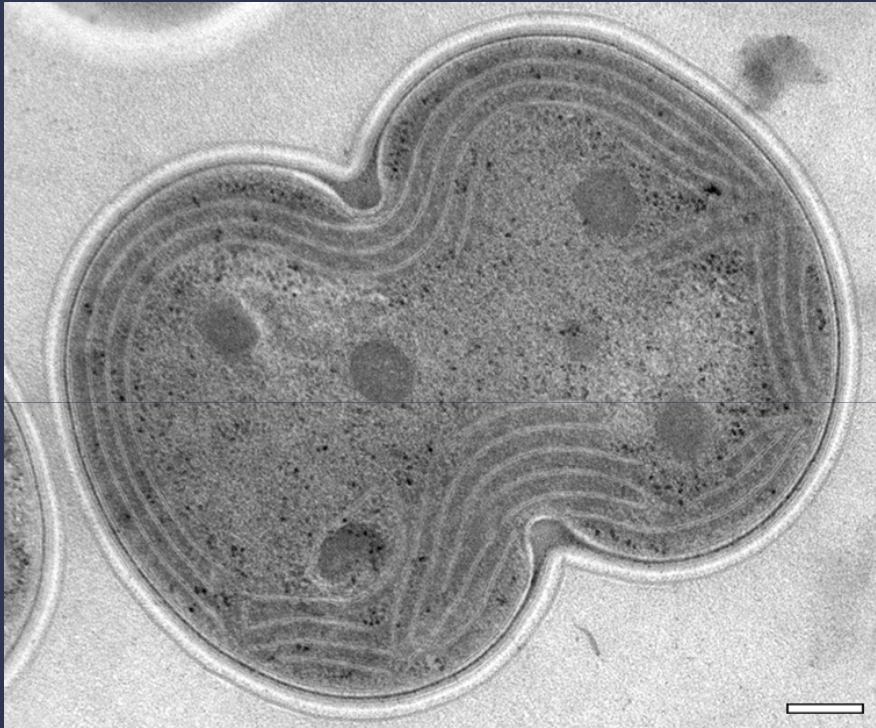
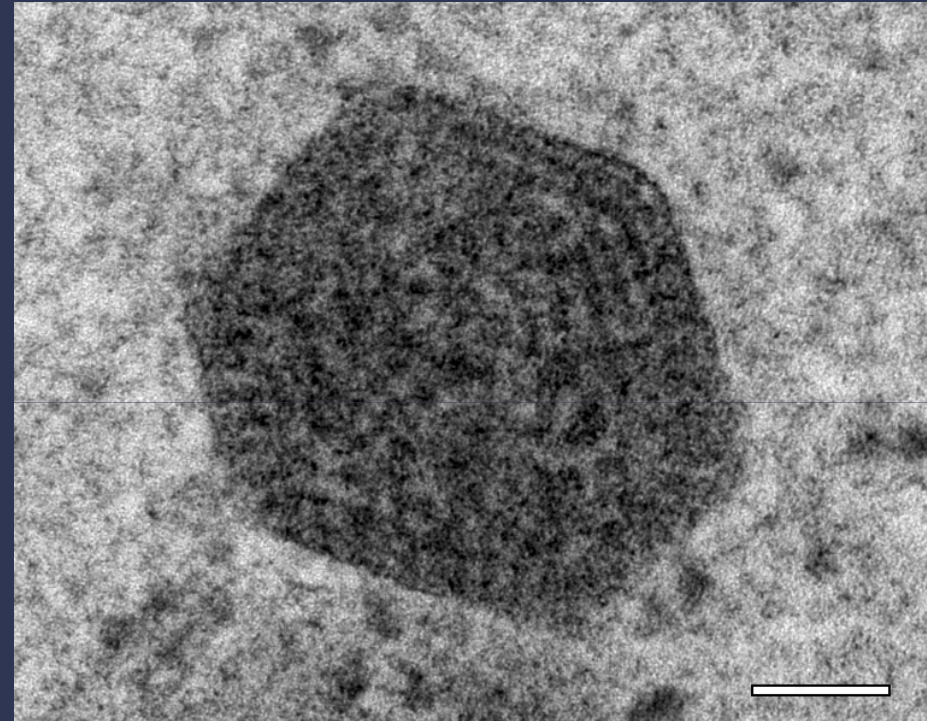


# Elucidating the Structural Basis of Carboxysome Function: A Progress Report

The **Carboxysome**, a Prokaryotic Organelle Composed Entirely of **Protein**, in Cyanobacteria and Chemoautotrophs



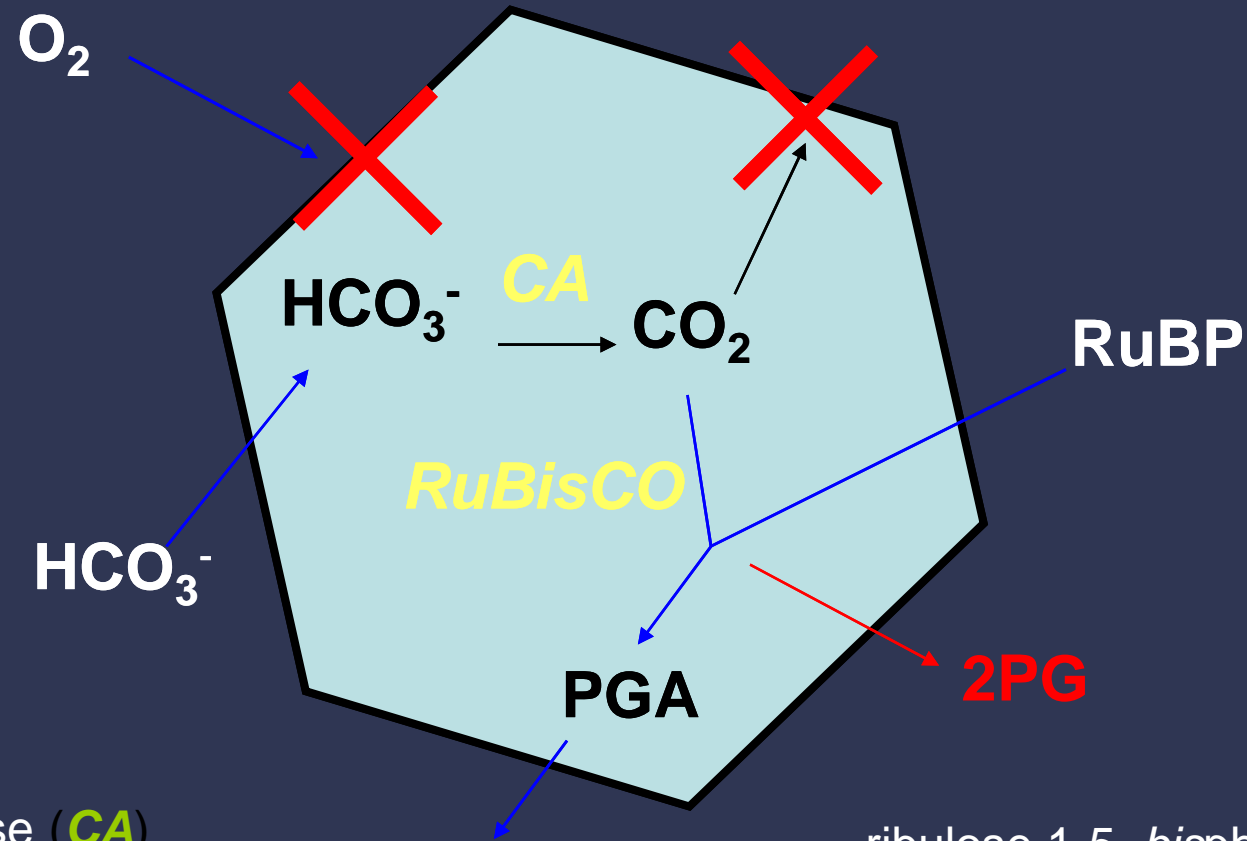
Scale bar = 200 nm



Scale bar = 50 nm

Transmission electron micrographs of *Syn 6803* cells courtesy of Robby Roberson, Wim Vermass and Allison van de Meene (Arizona State University).

# The carboxysome is a bacterial organelle for fixing CO<sub>2</sub>



carbonic anhydrase (**CA**)

Ribulose bis-phosphate (**RuBP**)

ribulose 1-5, bisphosphate  
carboxylase-oxygenase  
(**RuBisCO**).

3-phosphoglycerate (**PGA**)

**2PG, 2-phosphoglycolate (Eisenhut et al., 2006)**

# Carboxysome Genes

*Halothiobacillus neapolitanus*\*

Alpha

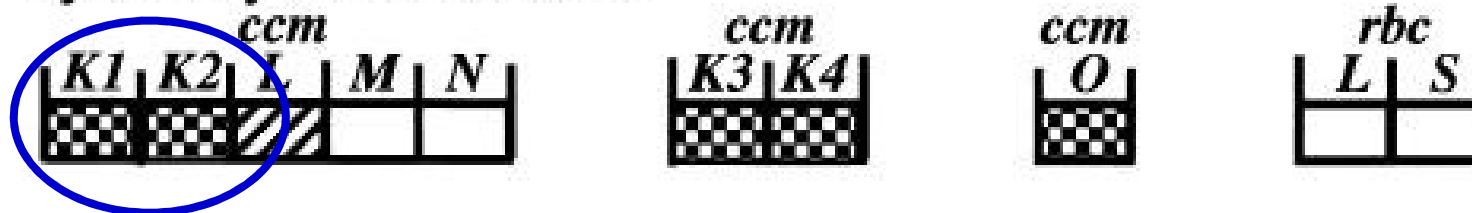


*Synechococcus* PCC7942\*\*



Beta

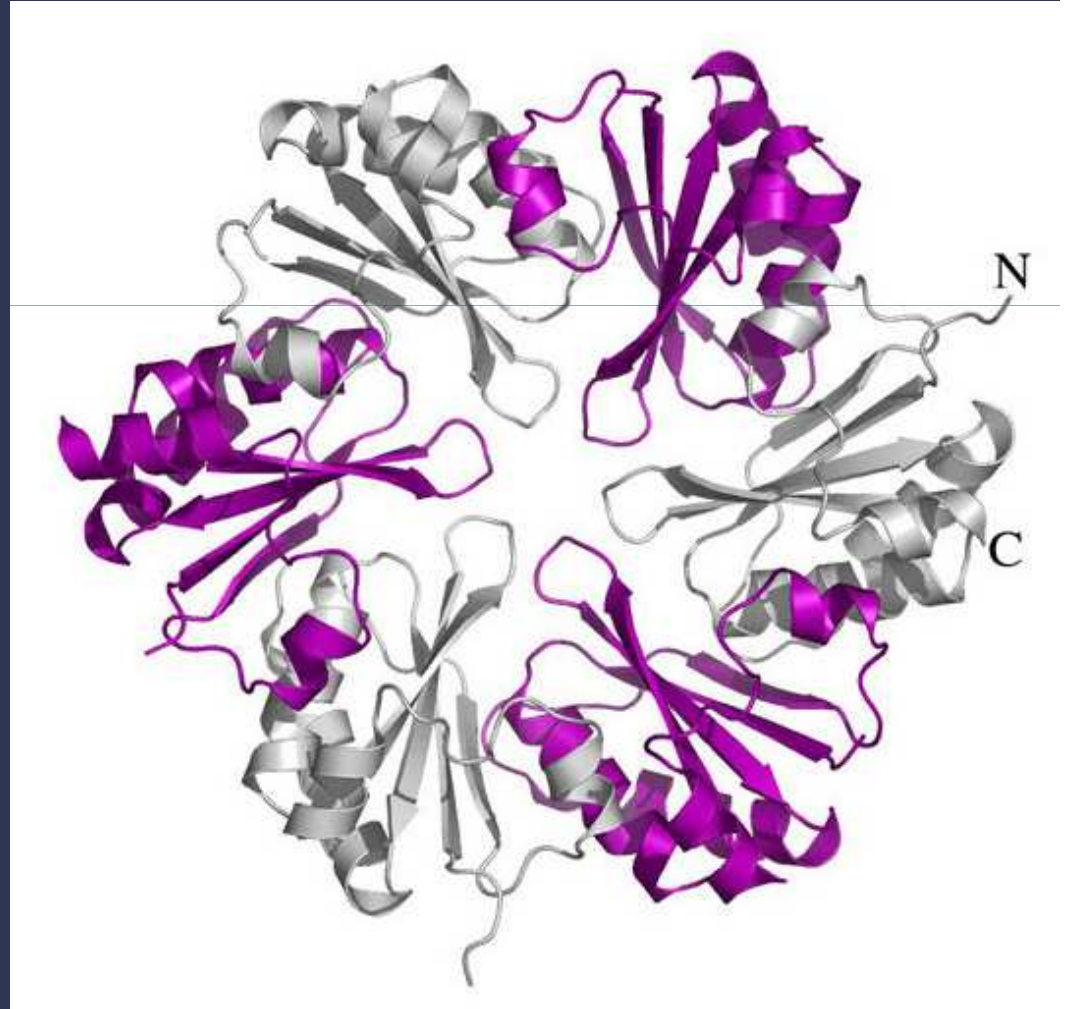
*Synechocystis* PCC6803\*\*\*

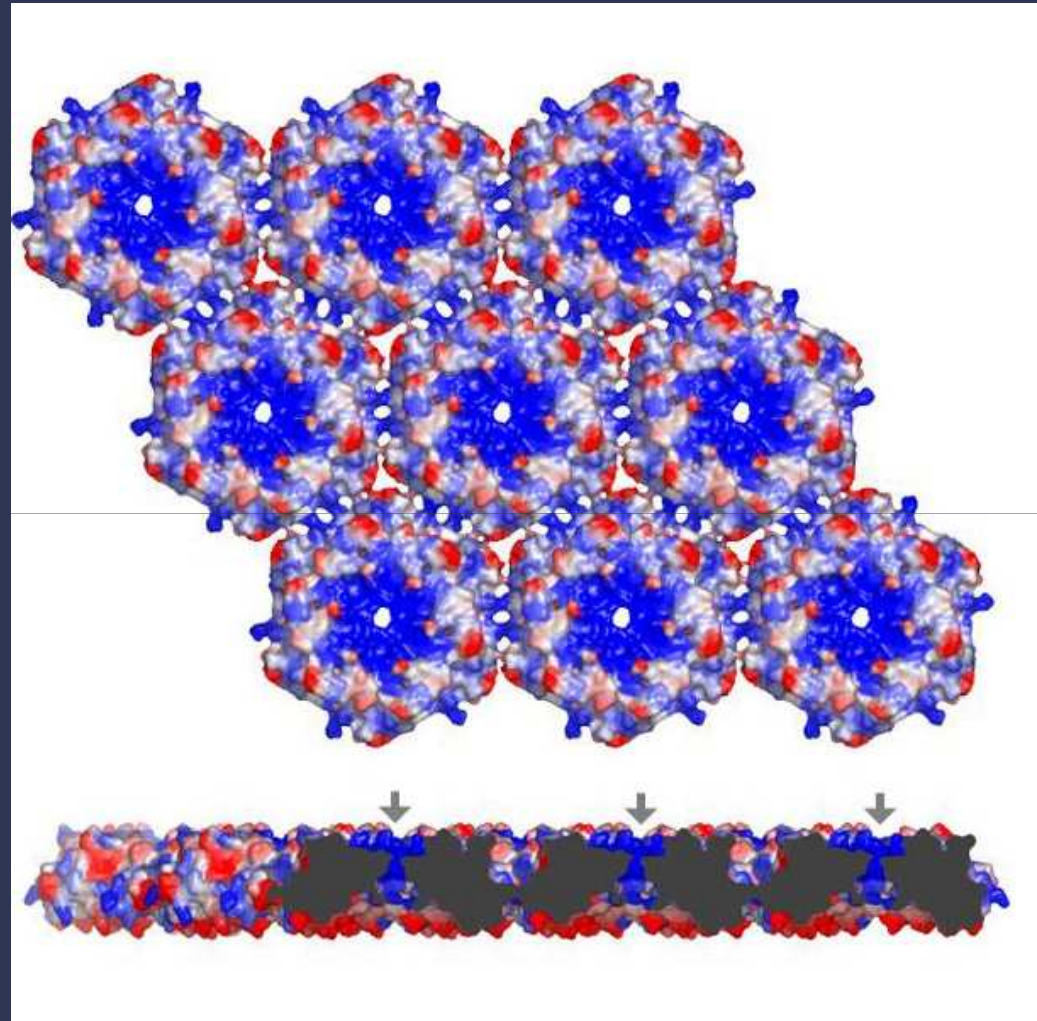
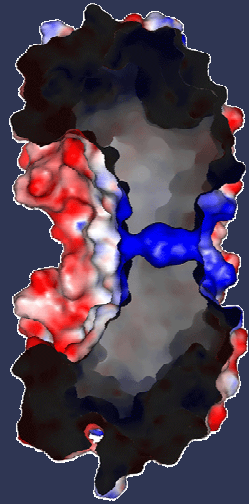
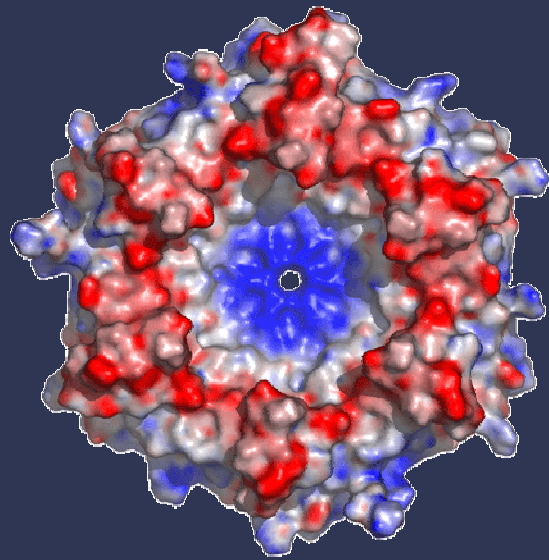


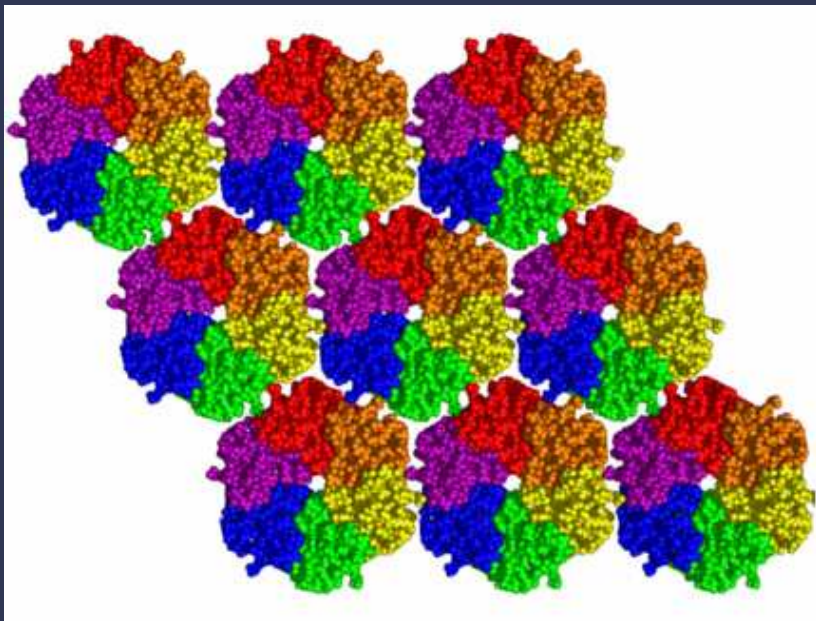
*Prochlorococcus marinus* MIT9313\*\*\*\*

Alpha

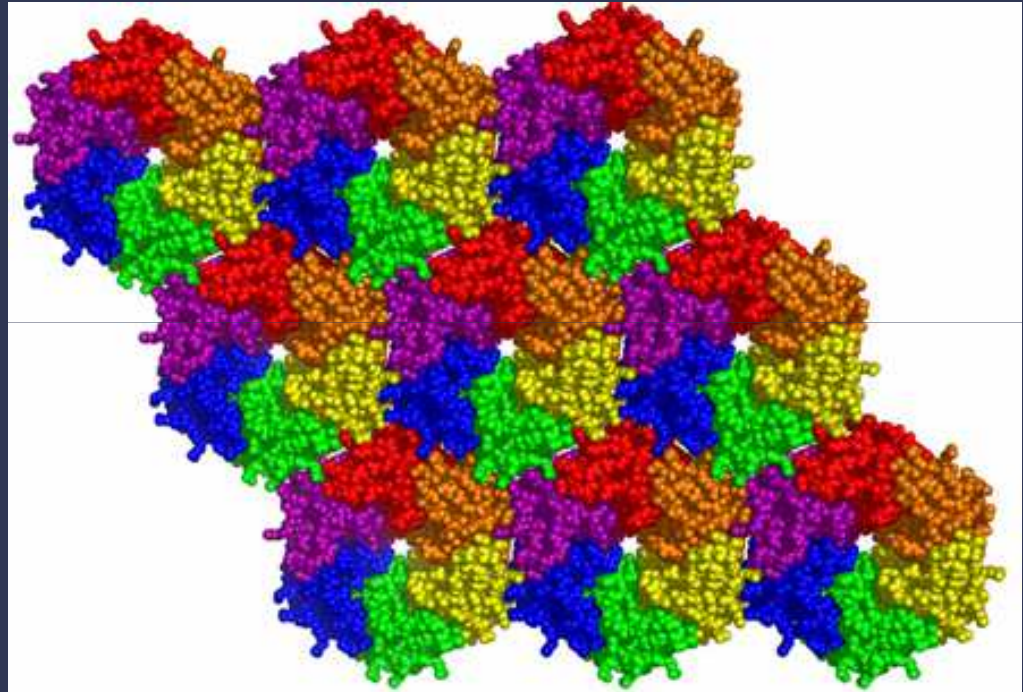








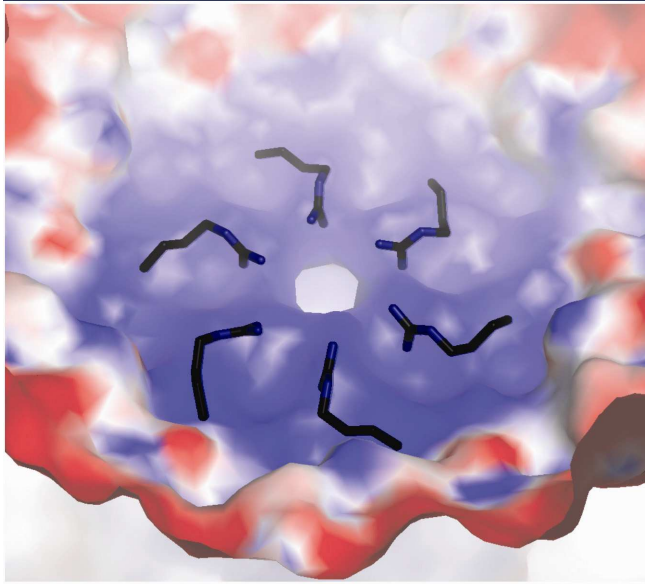
CcmK2 *Syn PCC6803* (Beta)



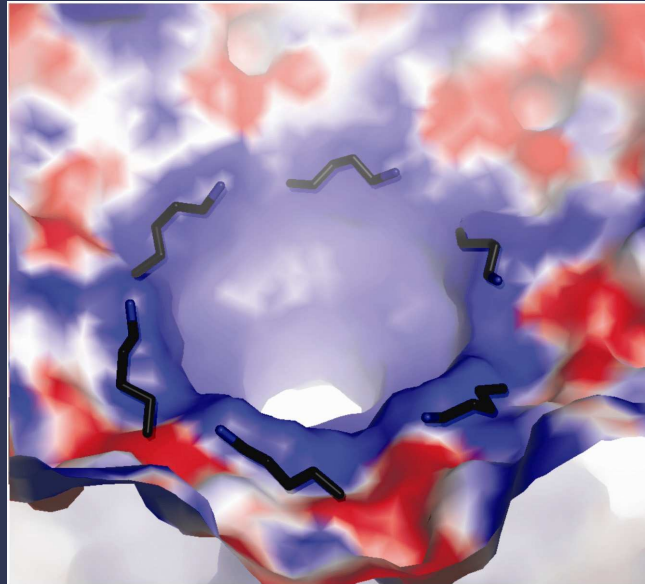
CsoS1A *H. neopolitanus* (Alpha)

# A comparison of pores

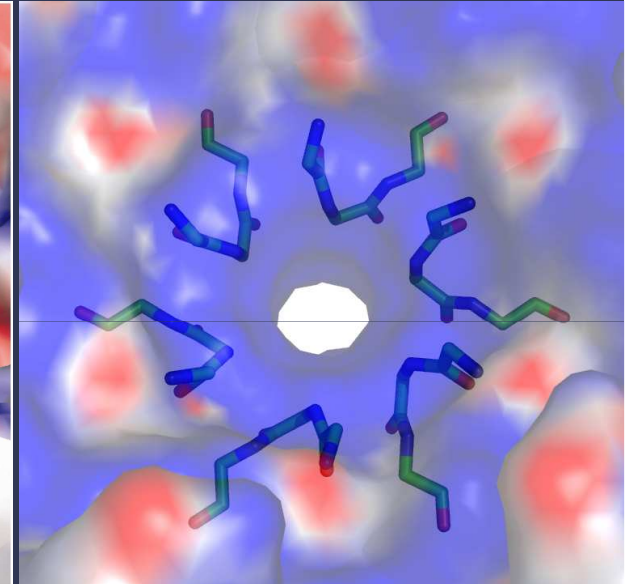
CcmK4



CcmK2/CcmK1



CsoS1A



Arg 38

Lys 36

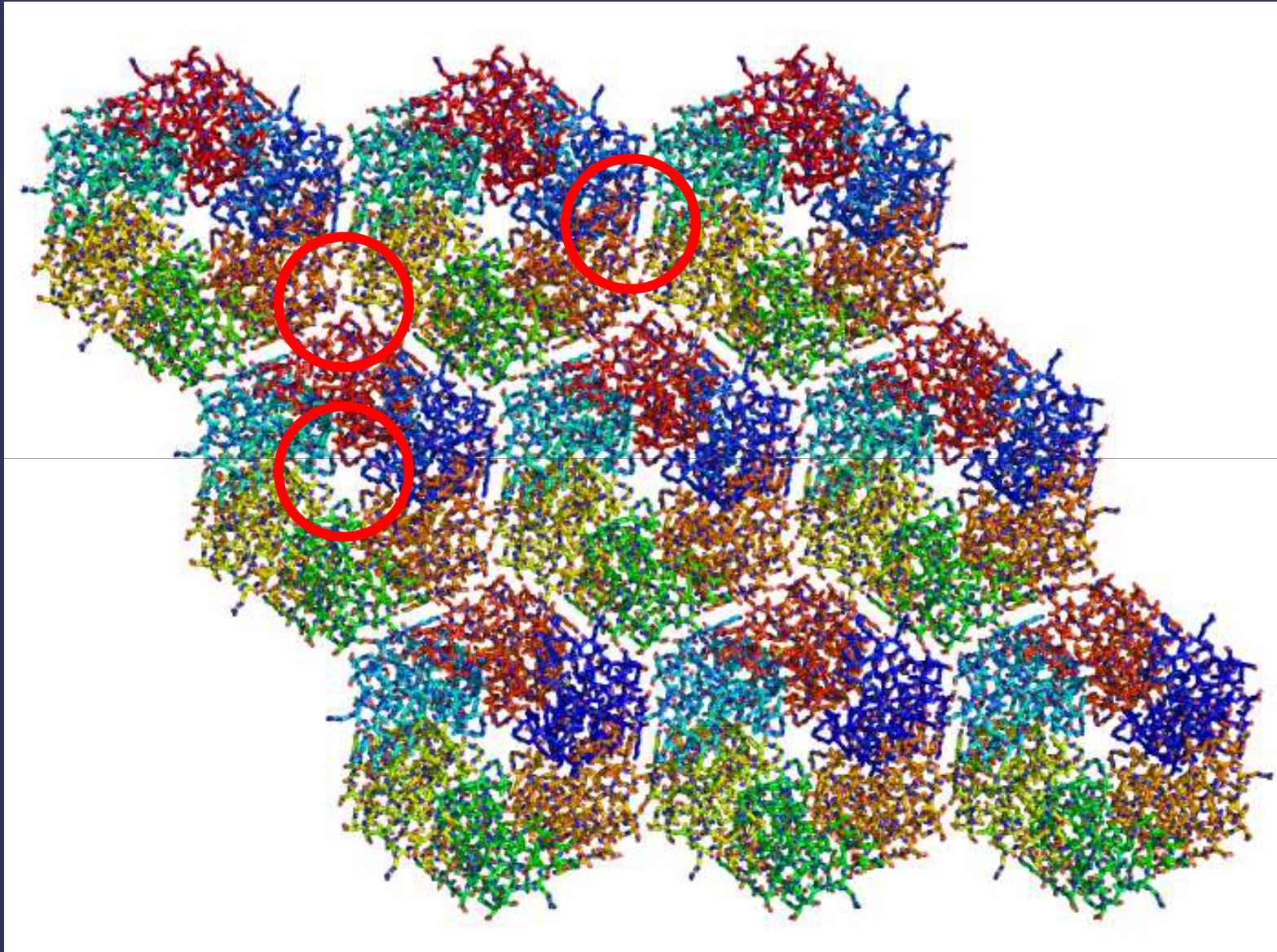
Loss of charge in pore: K→ F  
G triplet, inward facing amides

CcmK2    ----LVGYEKIGSGRVTVI----

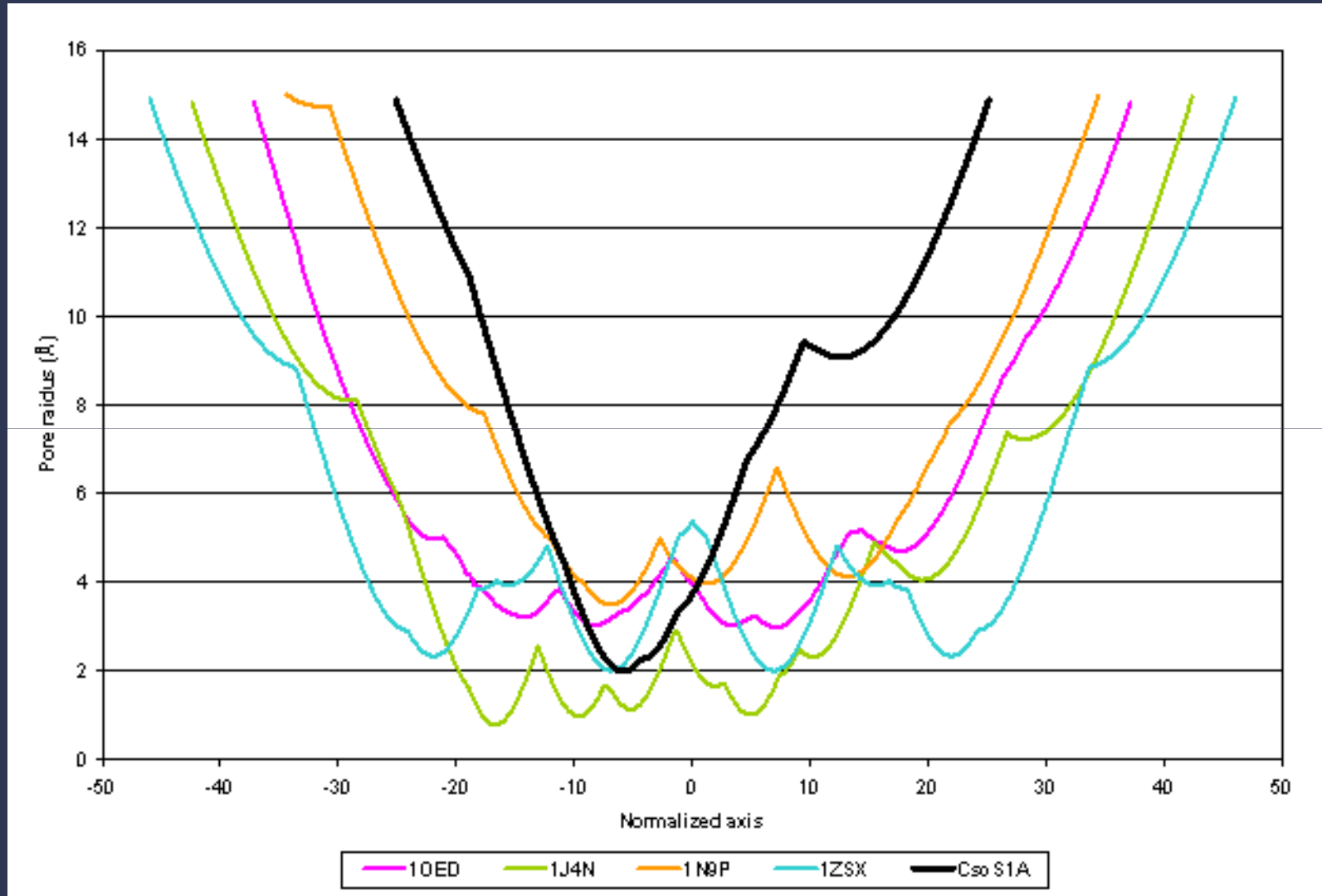
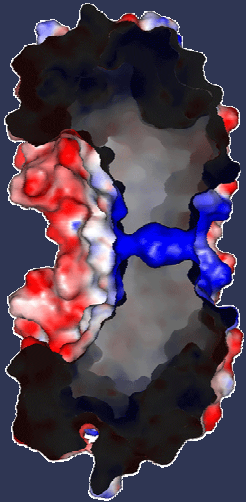
CsoS1A    ----LVGRQFVGGGYVTVL----



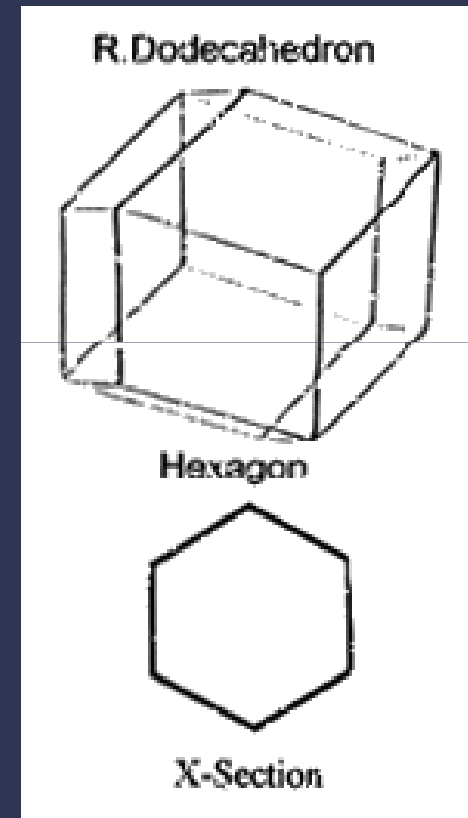
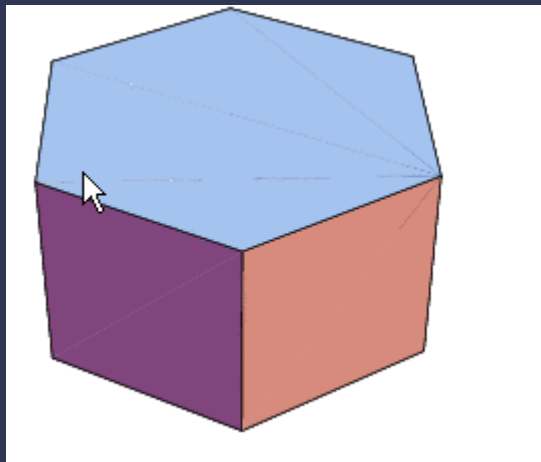
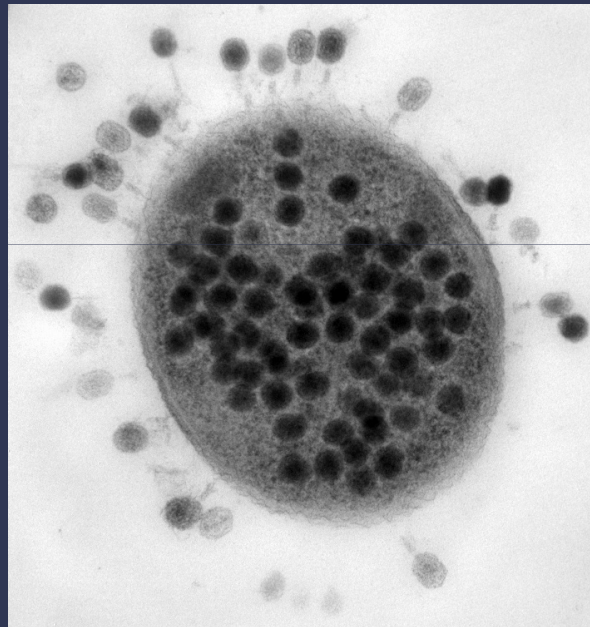
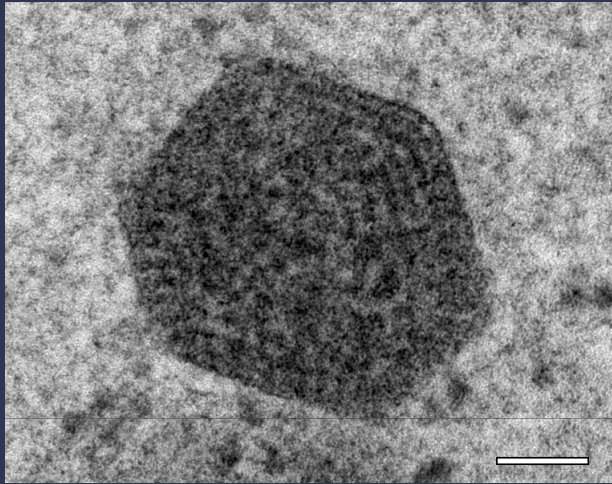
# Sulfate Soaks

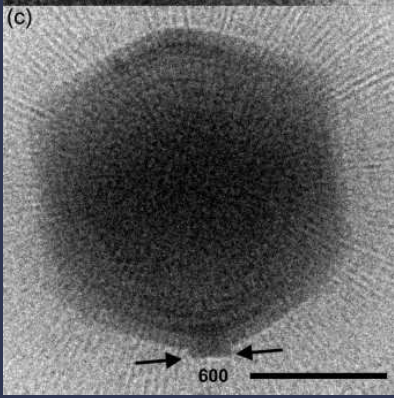
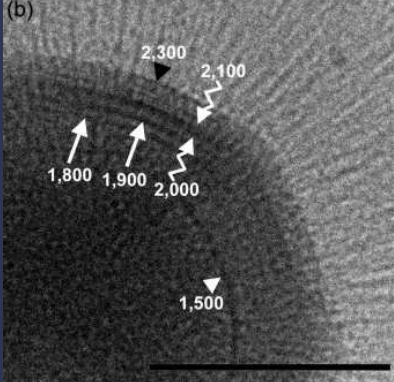
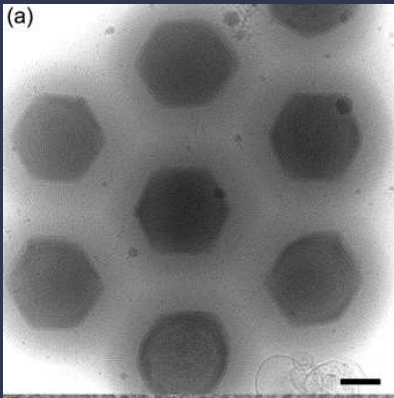


# Analogy to membrane channel proteins



# Possible Geometries?





And two tomography studies.....

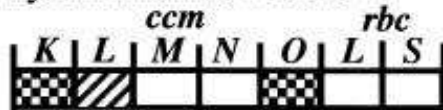
# Pentameric Defects?

*Halothiobacillus neapolitanus*\*



**Pfam 03319**

*Synechococcus* PCC7942\*\*



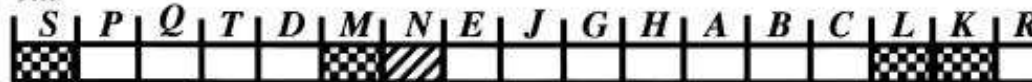
*Synechocystis* PCC6803\*\*\*



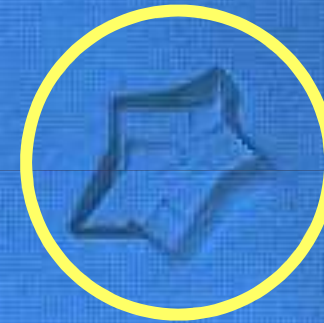
*Prochlorococcus marinus* MIT9313\*\*\*\*



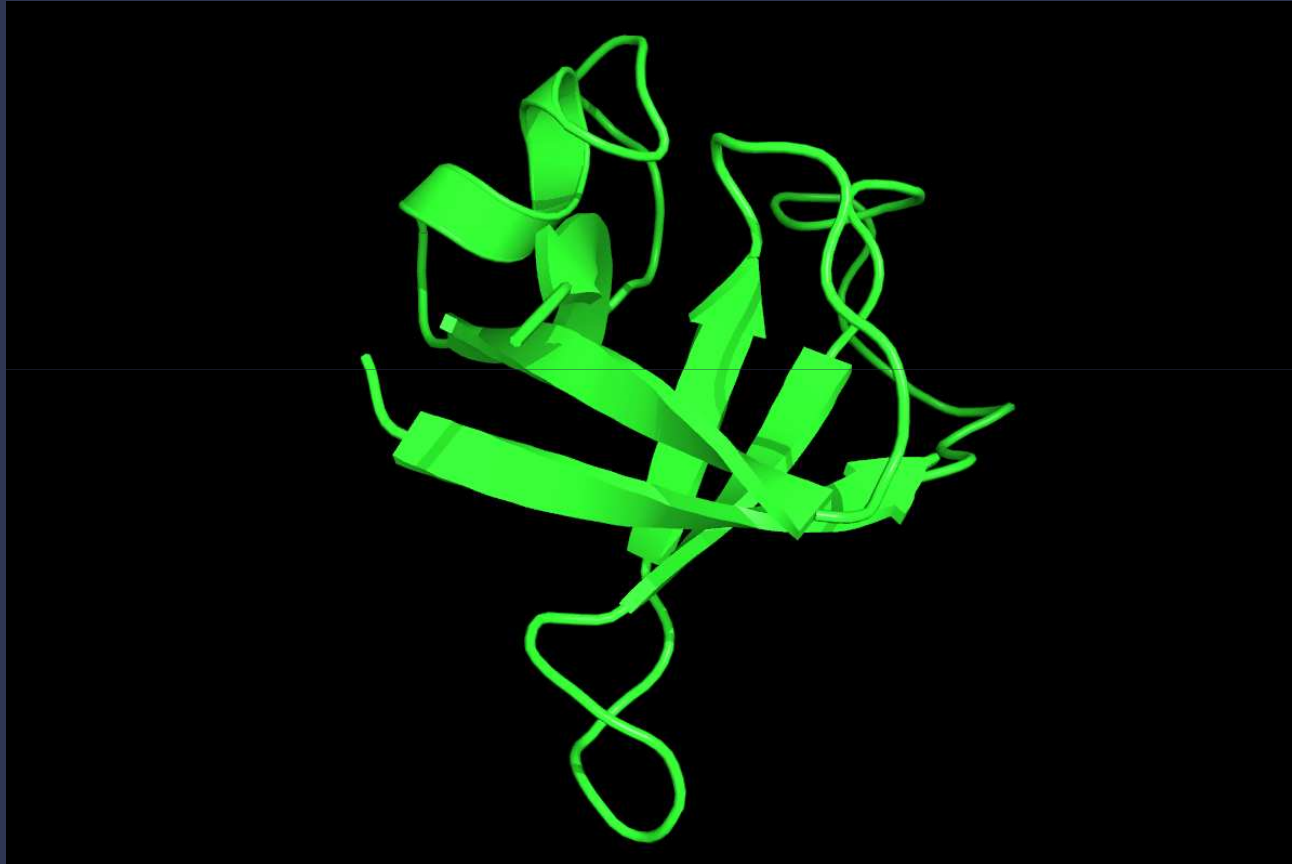
*Salmonella typhimurium*/*Escherichia coli*  
*eut*



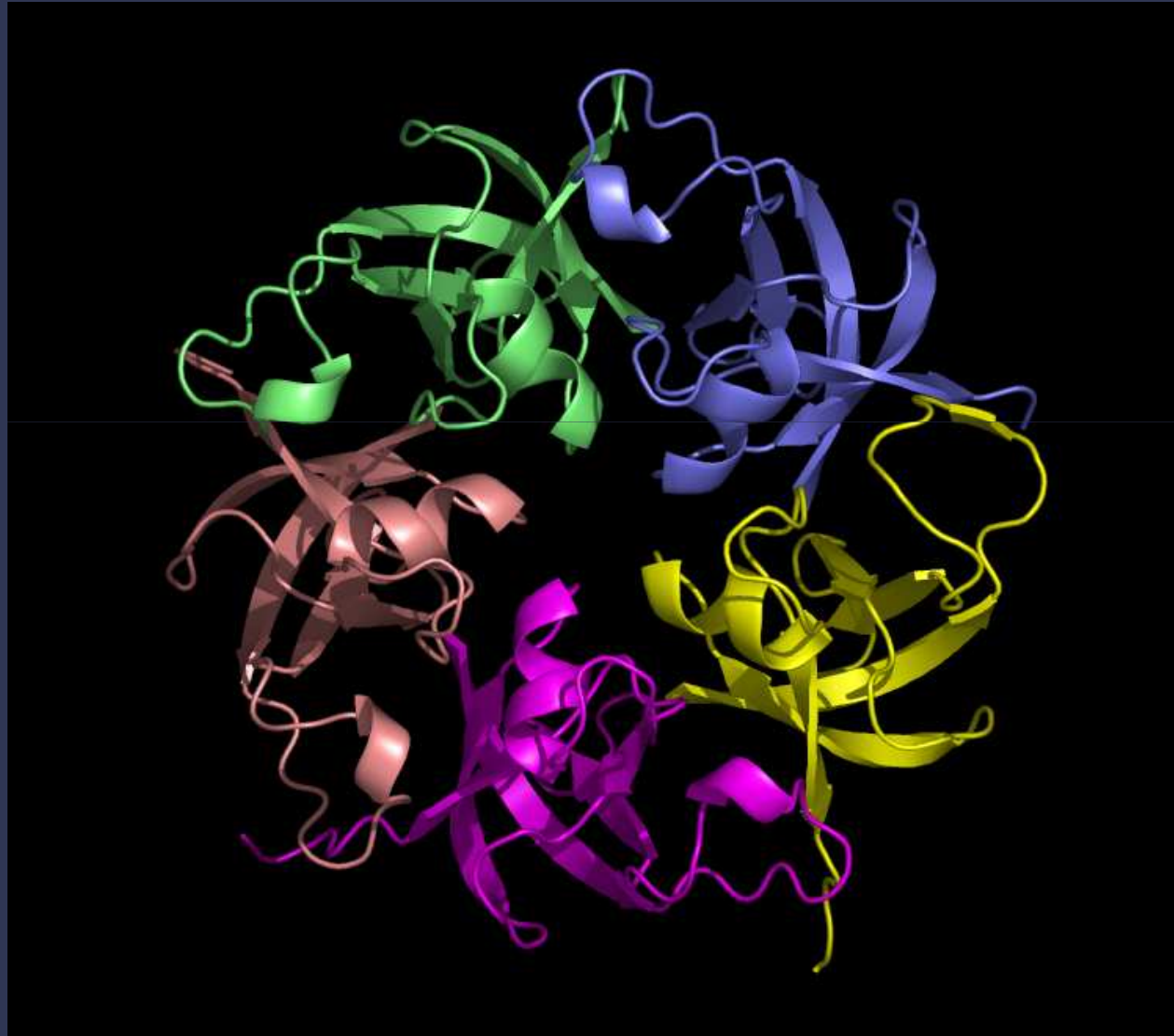
*Salmonella enterica*

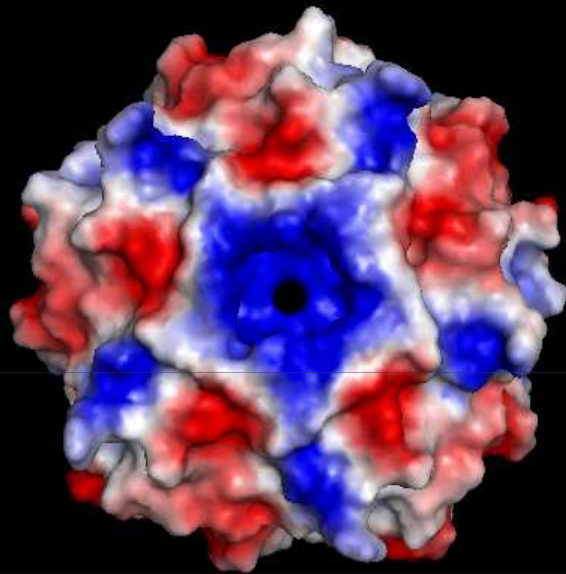


# OrfA structure



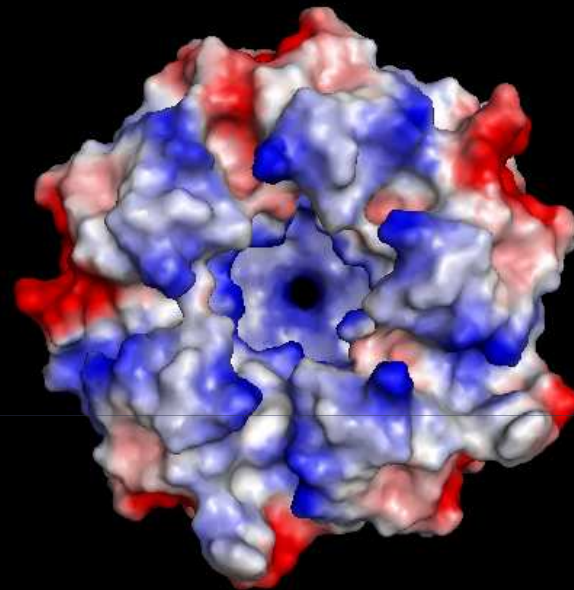
# OrfA structure





-64.978

64.978

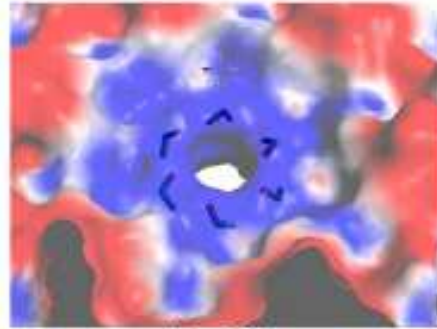


-64.978

64.978

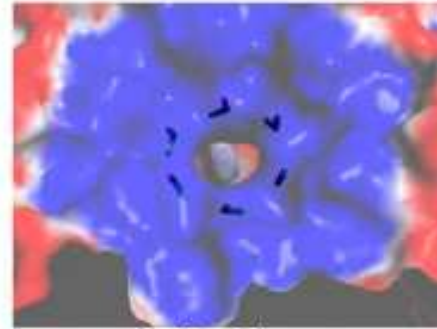


**CcmK1**



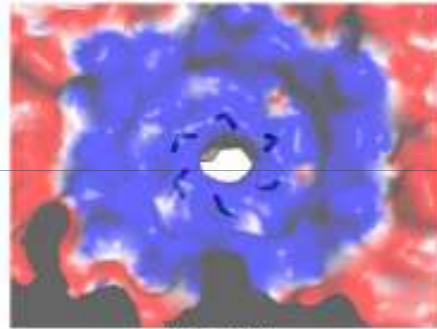
(Ser 39)

**CcmK2**



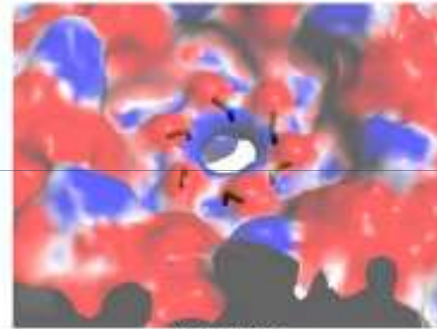
(Ser 39)

**CcmK4**



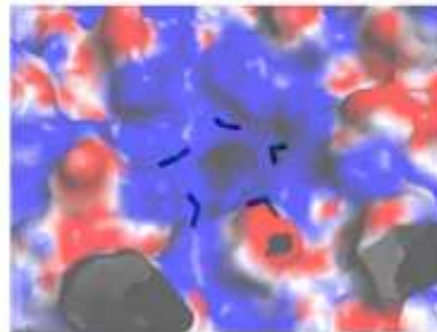
(Ser 41)

**CsoS1A**



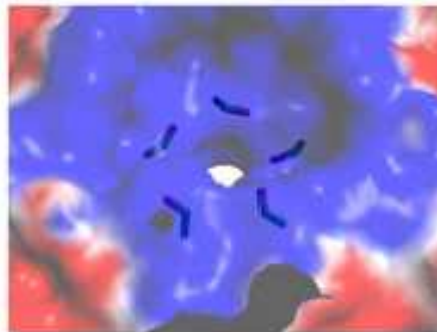
(Gly 43)

**CcmL**



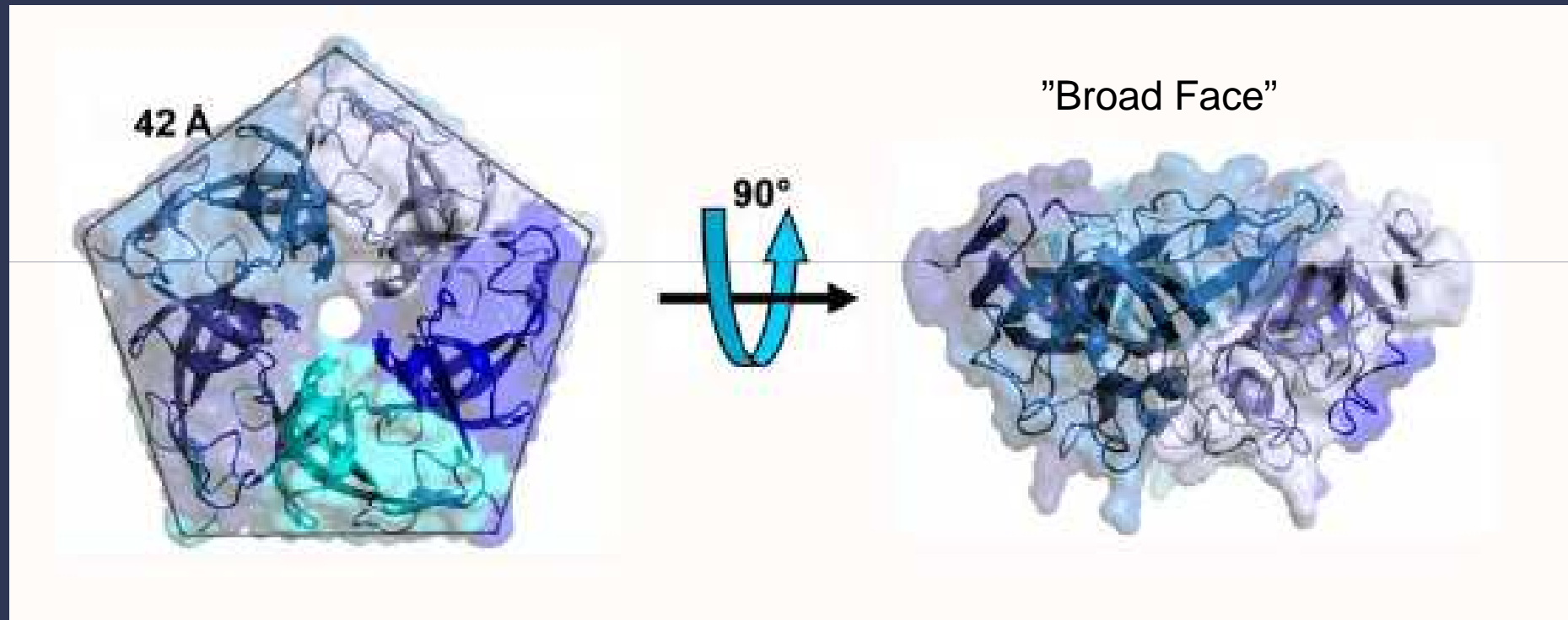
(Ser 61)

**OrfA**

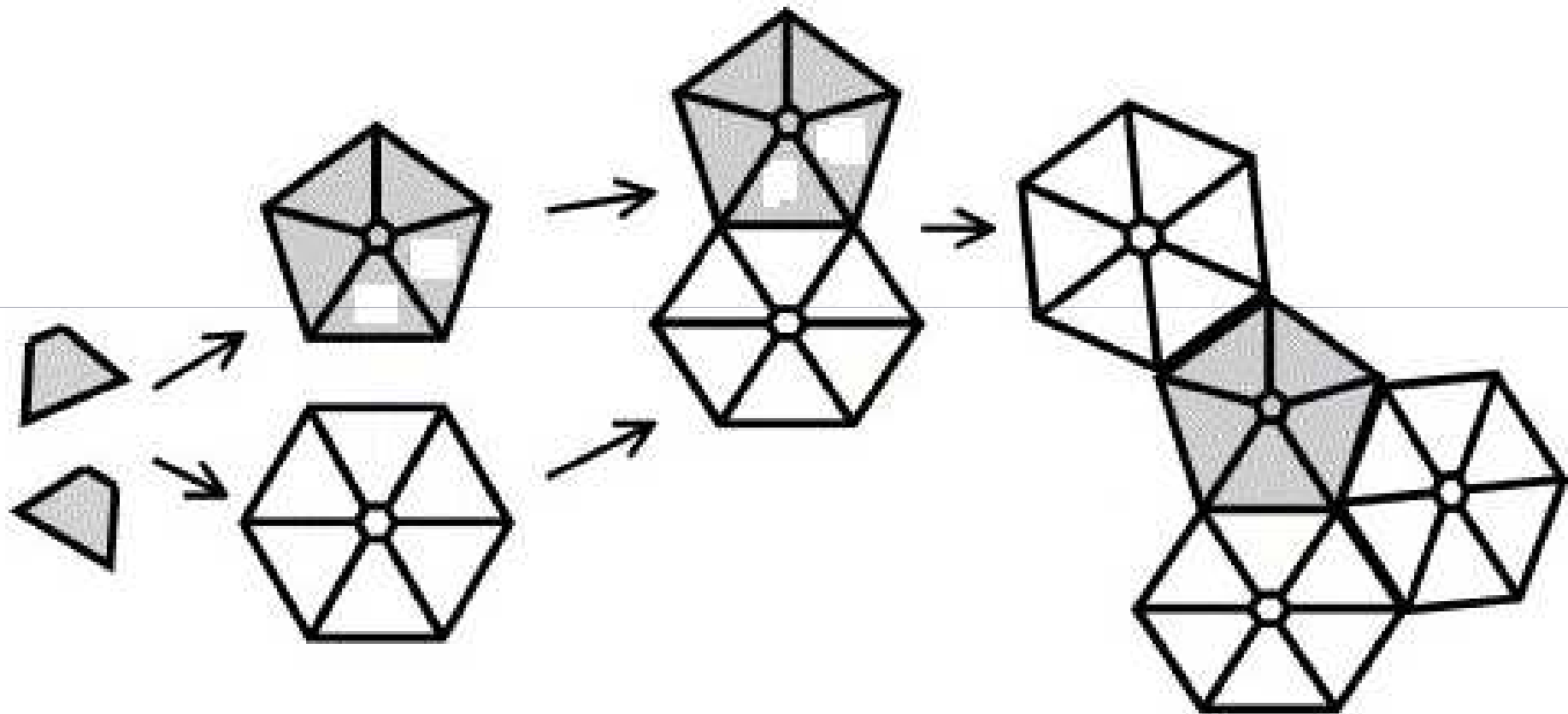


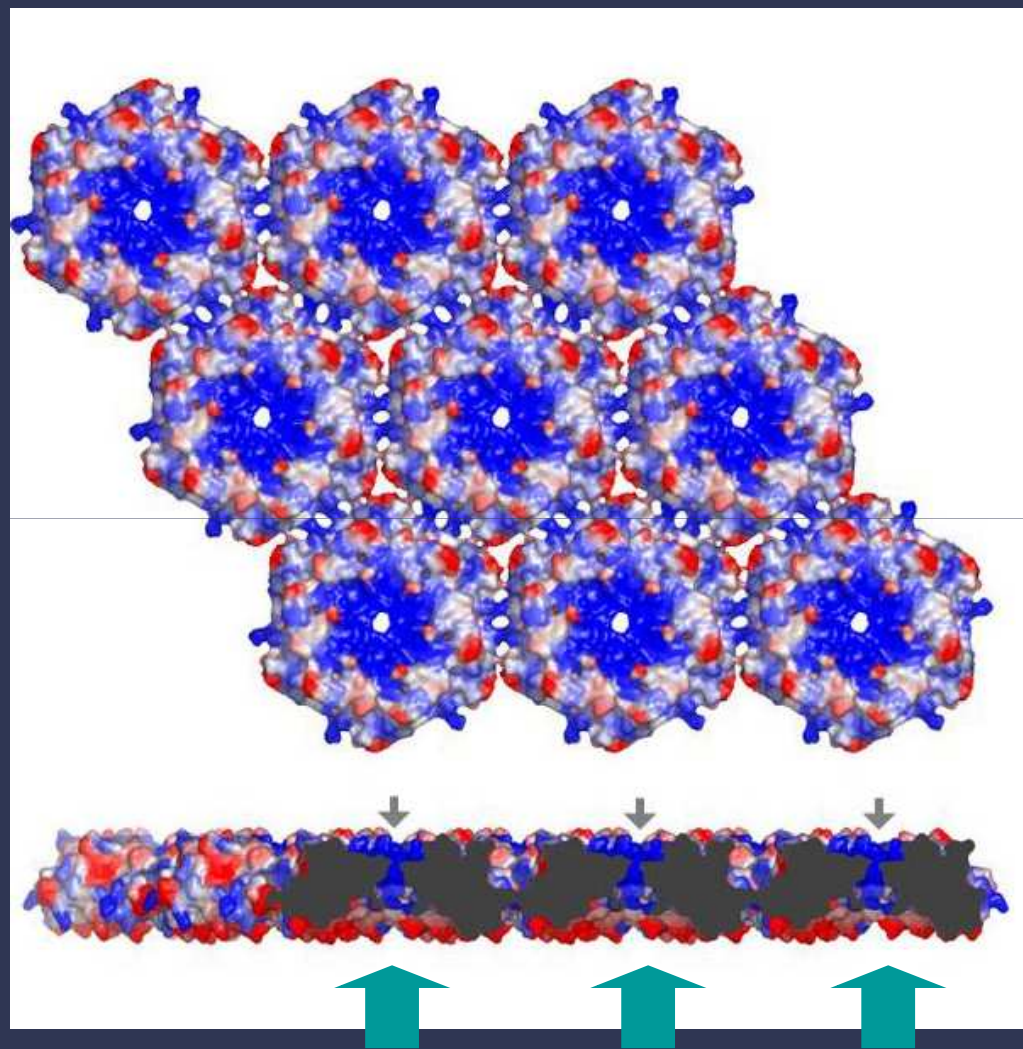
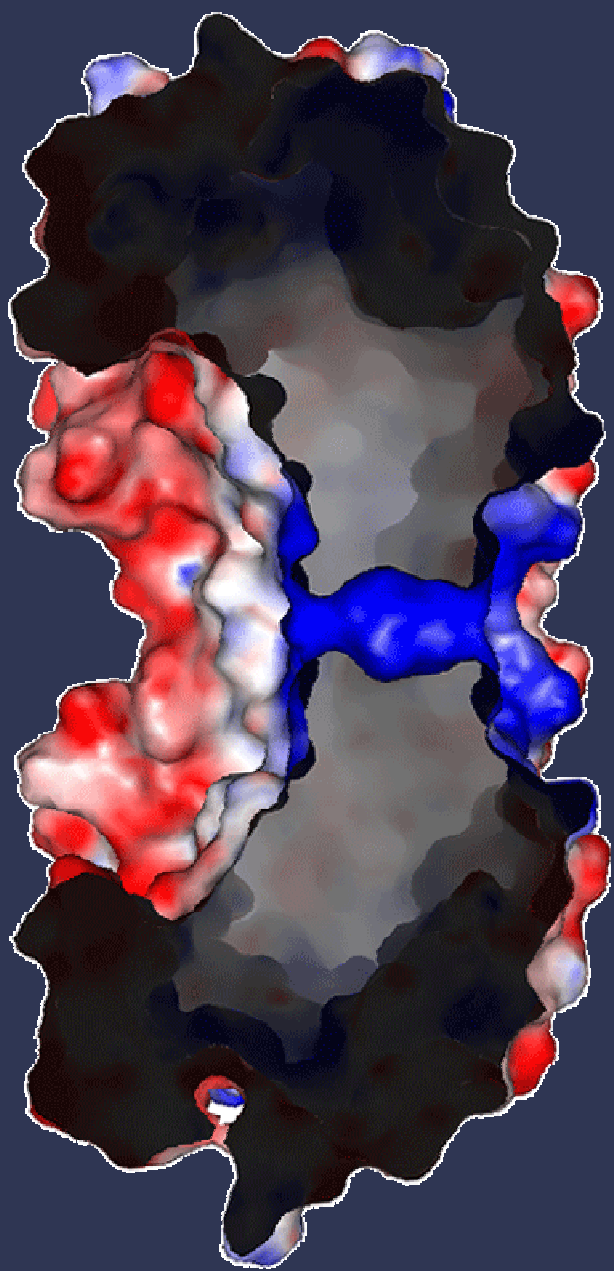
(Ser 55)

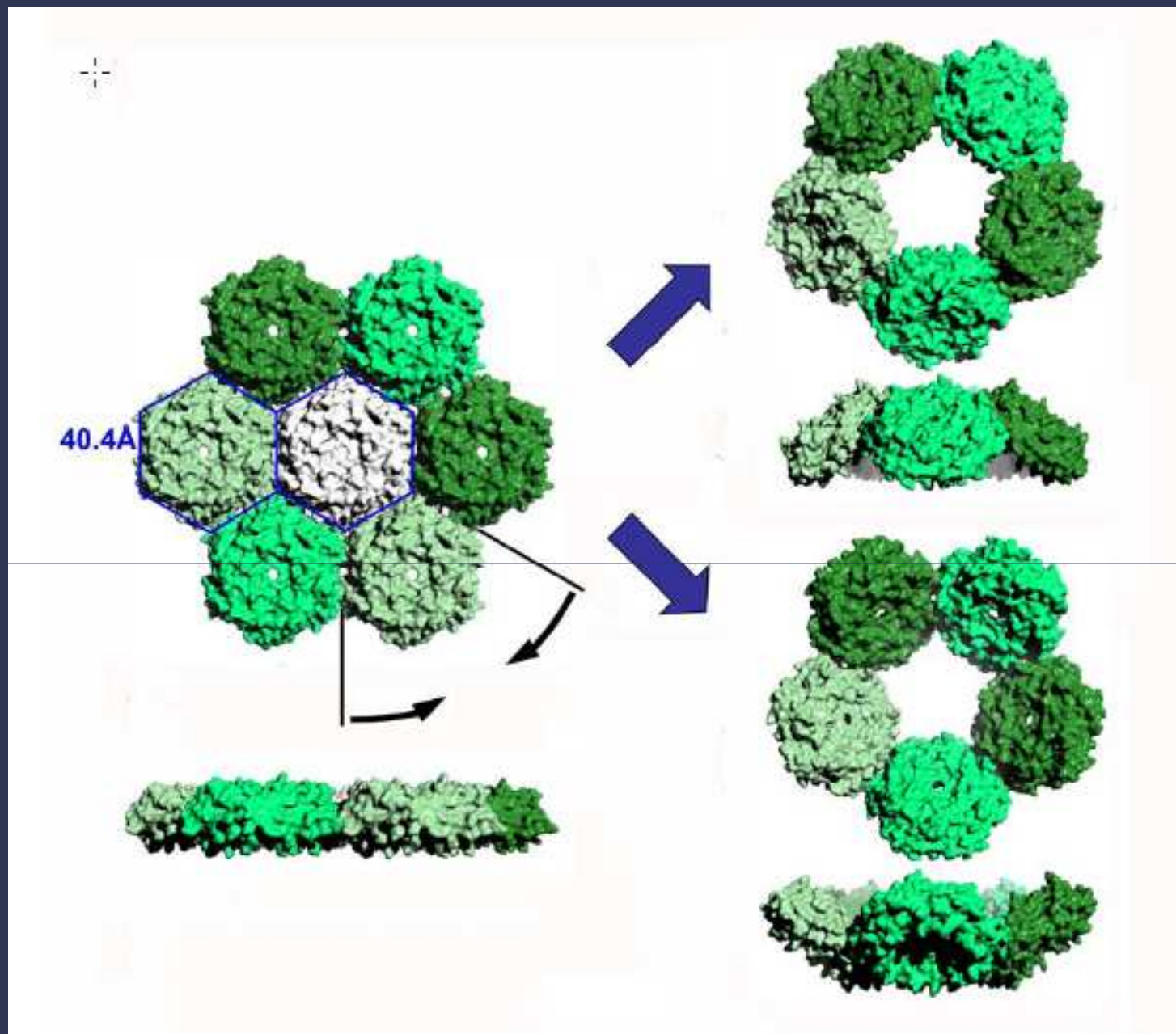
# The Pentamer is a Truncated Pyramid

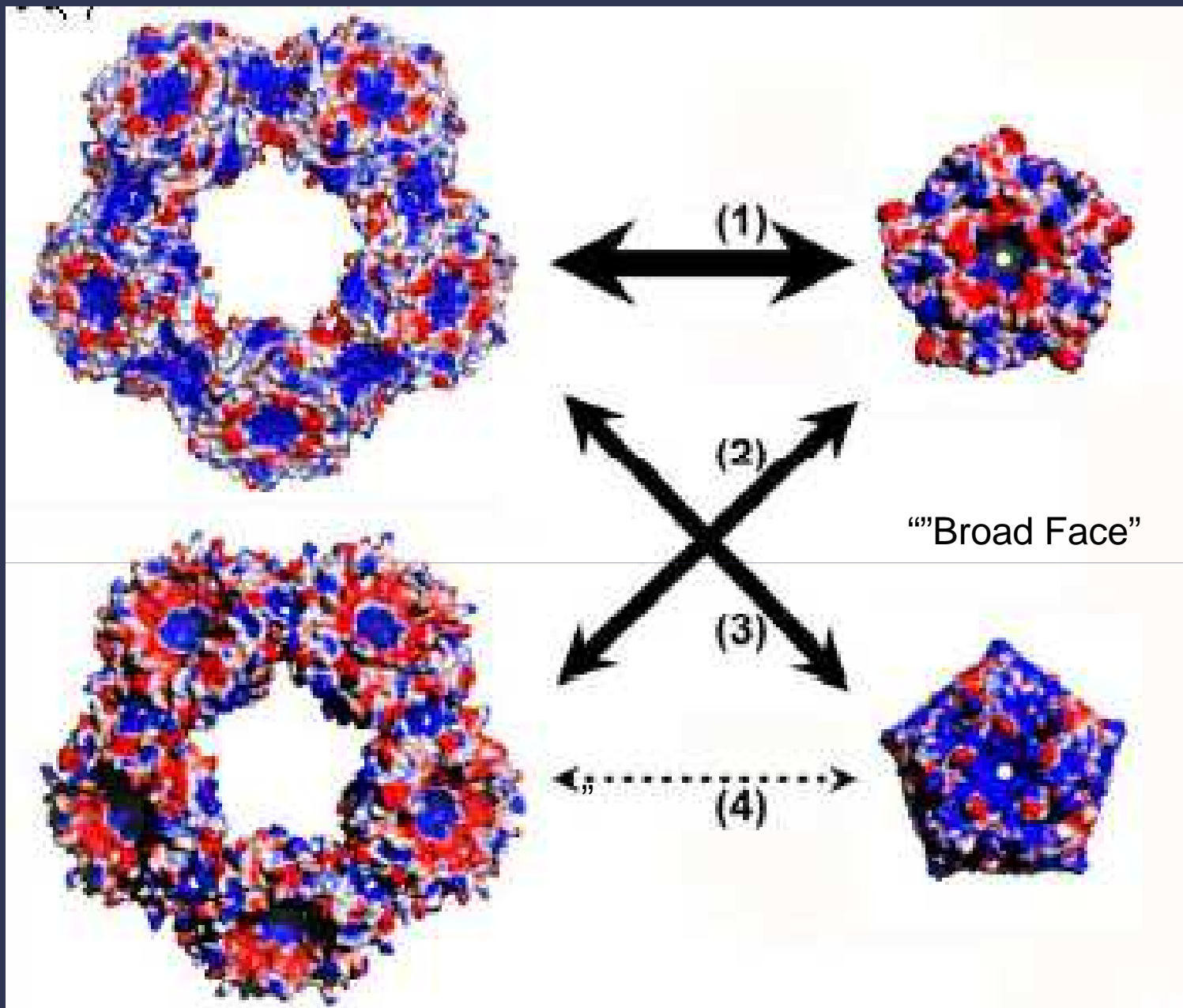


# Assembly of an Icosahedron

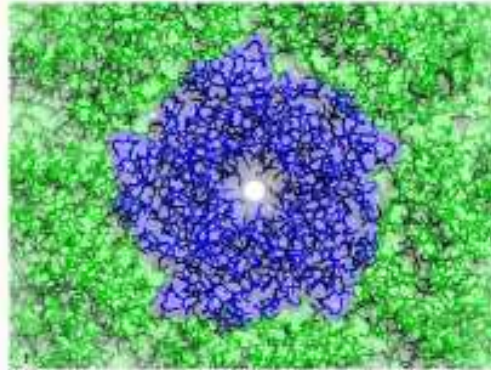




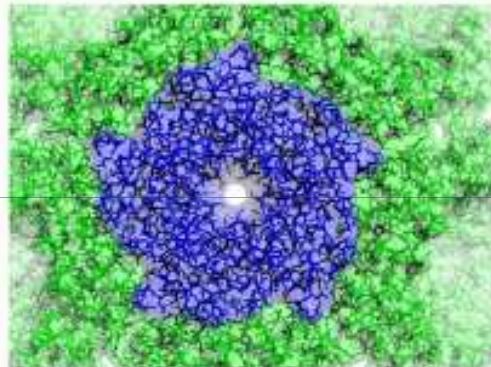




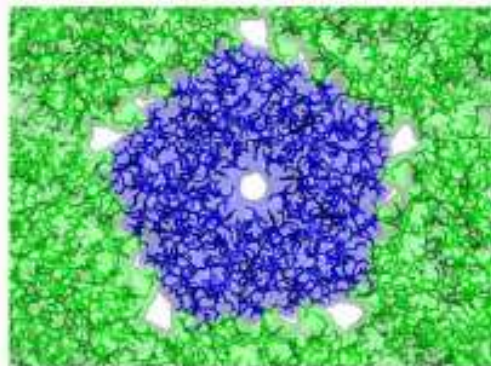
**B**



(1) RD = -35.62, SC = 0.496, SA = 996

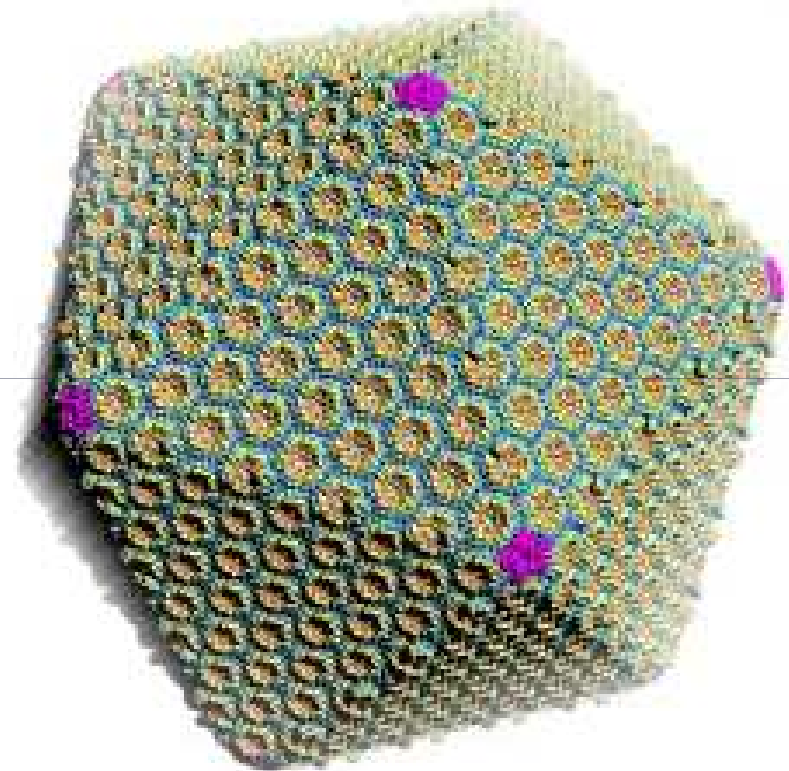
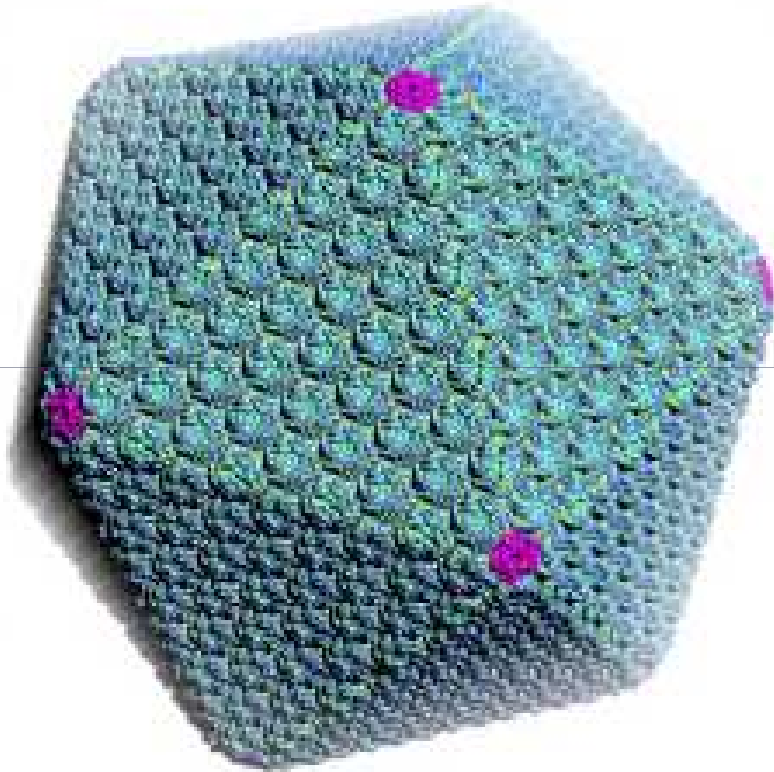


(2) RD = -35.90, SC = 0.387, SA = 937



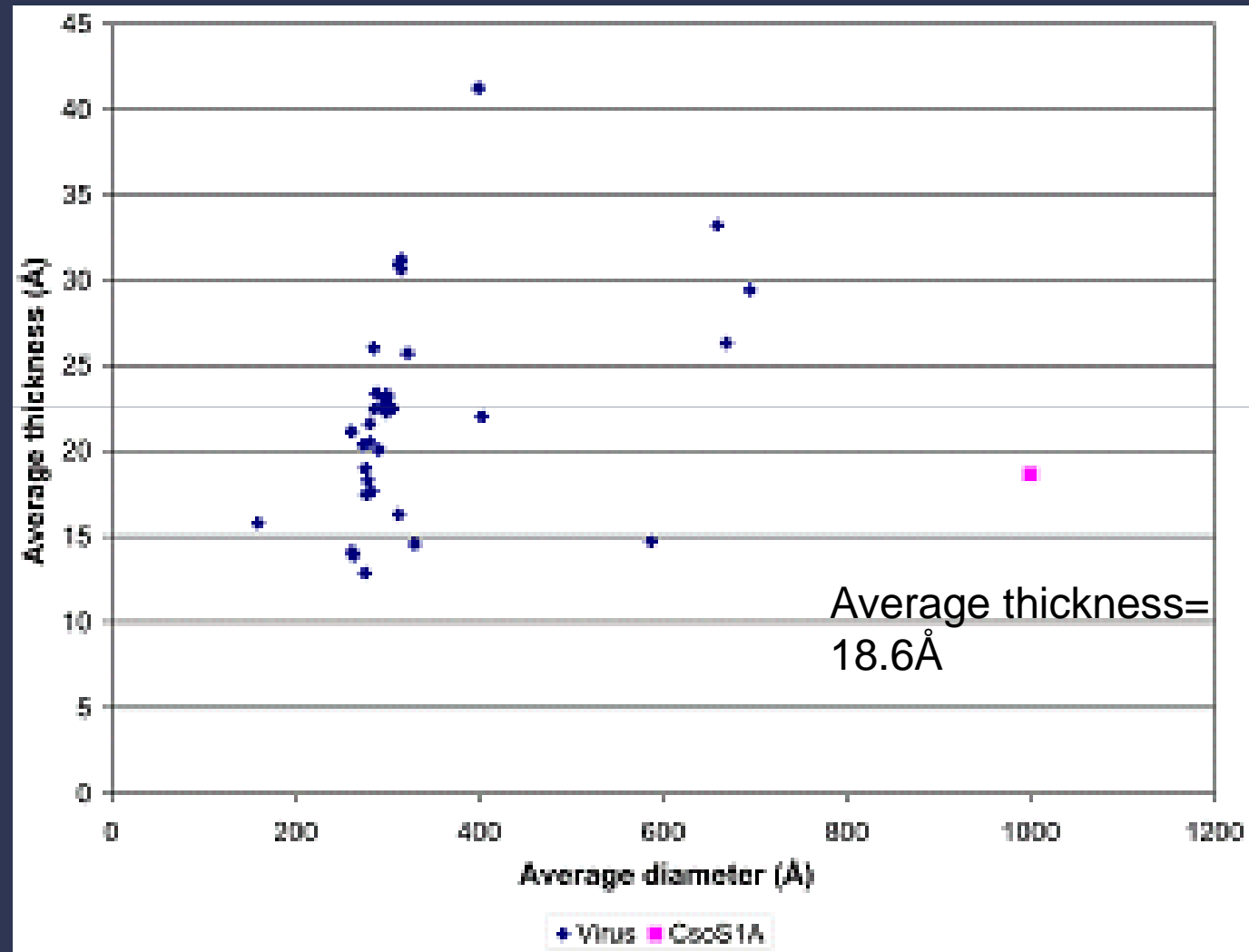
(3) RD = -26.35, SC = 0.376, SA = 816

# T=75 Models of the Carboxysome

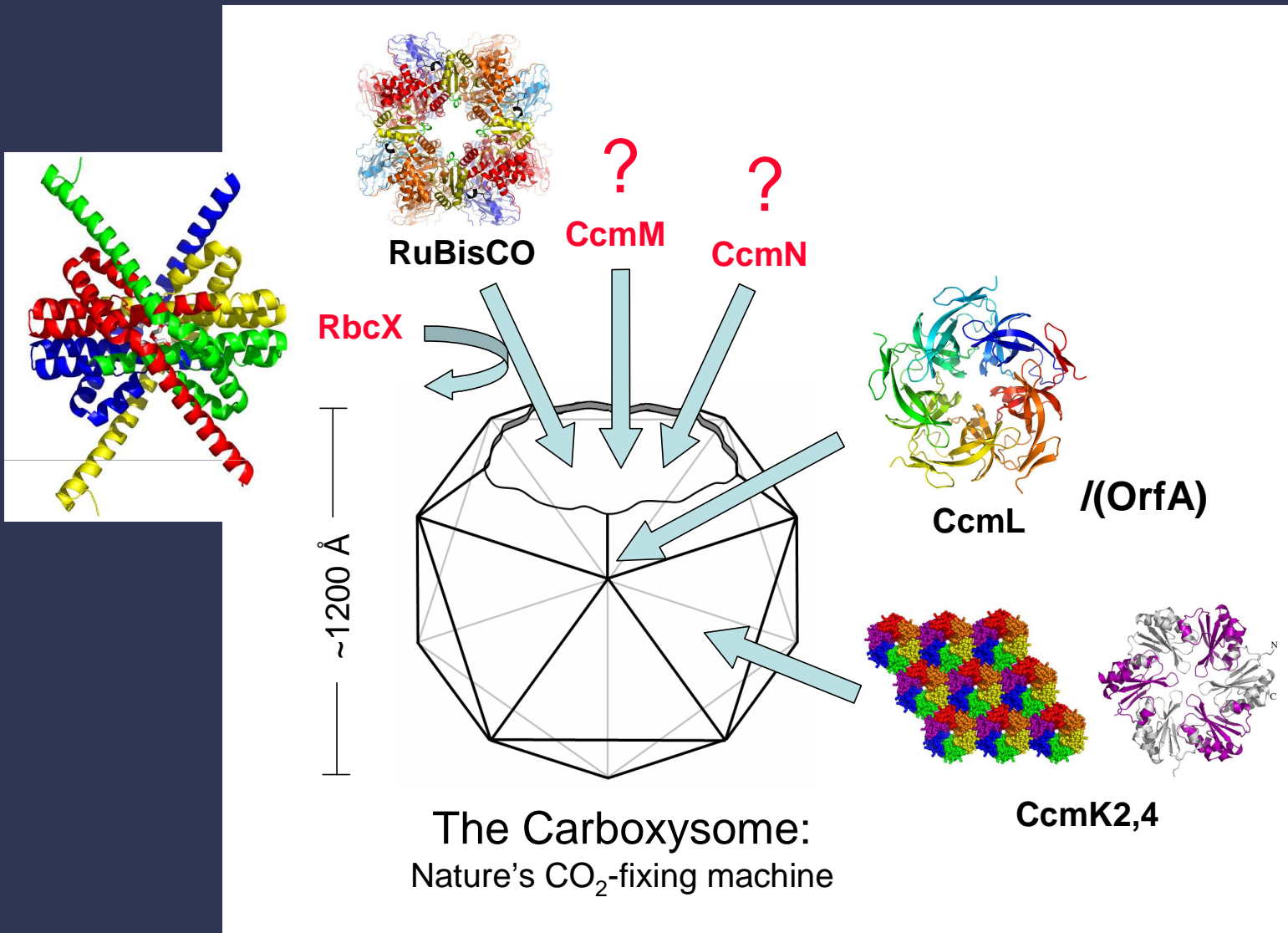


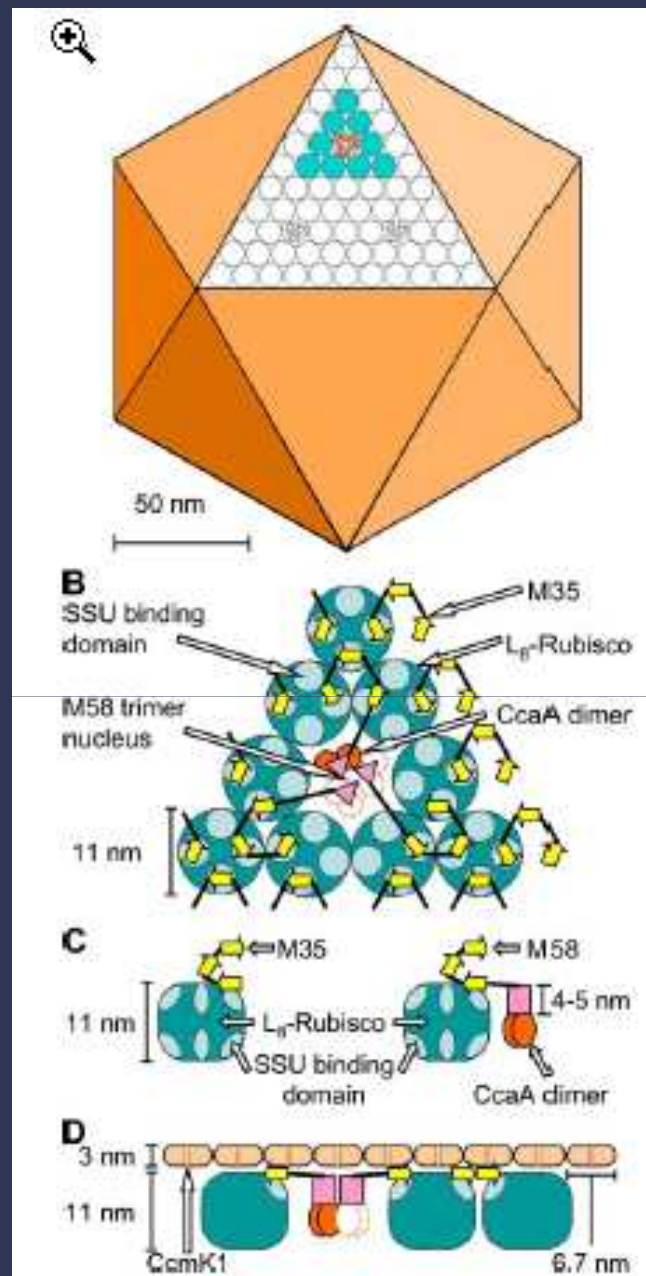


# The carboxysome shell is relatively thin compared to the shells of viruses

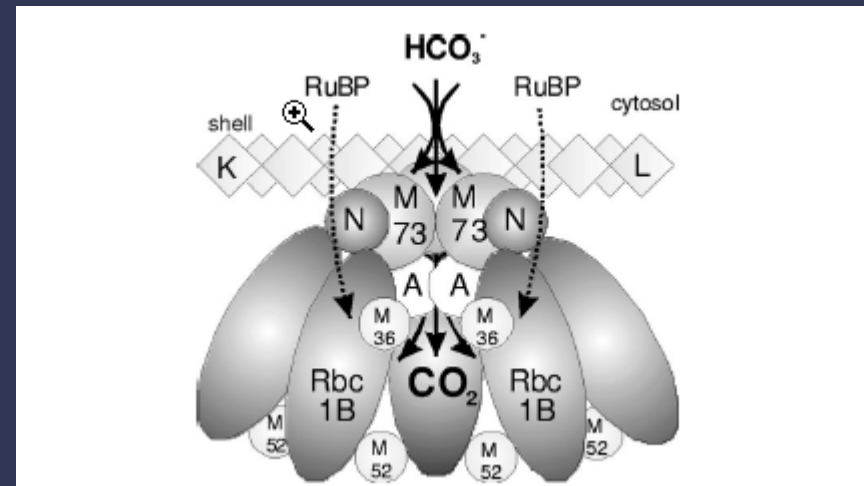


# Next Steps.....





Long et al., JBC 2007



Cot et al., J. Bact 2007

JBC Papers in Press. Published on March 10, 2008 as Manuscript M709214200  
The latest version is at <http://www.jbc.org/cgi/doi/10.1074/jbc.M709214200>

Recombinant organelle biogenesis

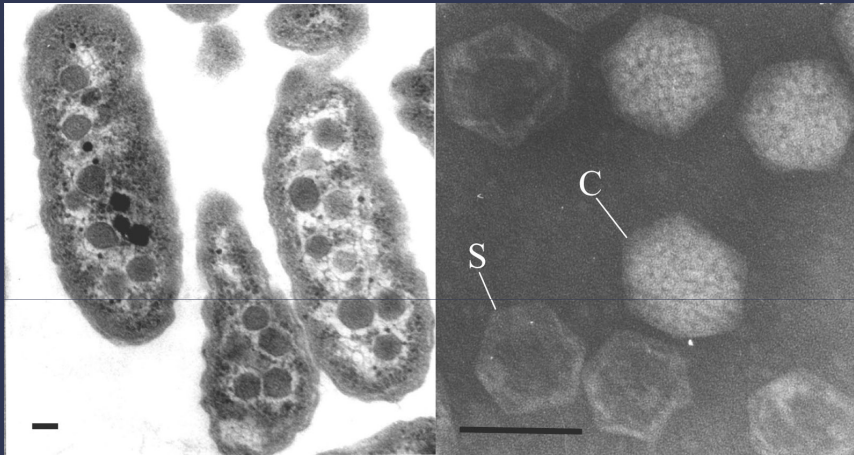
#### BIOCHEMICAL AND STRUCTURAL INSIGHTS INTO BACTERIAL ORGANELLE FORM AND BIOGENESIS.

Joshua P Parsons<sup>1\*</sup>, Sriramulu D Dinesh<sup>2\*</sup>, Evelyne Deery<sup>1</sup>, Helen K Leech<sup>1</sup>, Amanda A Brindley<sup>1</sup>, Dana Heldt<sup>1</sup>, Steffi Frank<sup>1</sup>, C. Mark Smales<sup>1</sup>, Heinrich Lünsdorf<sup>3</sup>, Alain Rambach<sup>4</sup>, Mhairi H Gass<sup>5</sup>, Andrew Bleloch<sup>5</sup>, Kirsty J McClean<sup>6</sup>, Andrew W Munro<sup>6</sup>, Stephen E J Rigby<sup>7</sup>, Martin J Warren<sup>1</sup> and Michael B Prentice<sup>2</sup>

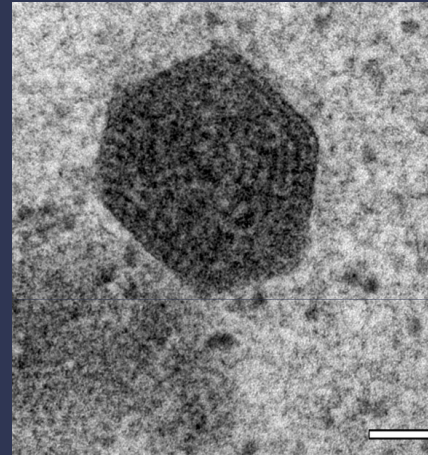
1. Protein Science Group, Department of Biochemistry, University of Kent, Canterbury, Kent CT2 7NJ, UK
2. Departments of Pathology and Microbiology, University College Cork, Ireland
3. Department of Vaccinology, Helmholtz Center of Infection Research, Braunschweig, Germany
4. CHROMagar, 4 place du 18 Juin 1940, F-75006 Paris, France
5. SuperSTEM facility, Daresbury Laboratories, Daresbury, UK
6. Faculty of Life Sciences, Manchester Interdisciplinary Biocentre, University of Manchester, 131 Princess Street, Manchester M1 7DN
7. School of Biological and Chemical Sciences, Queen Mary, University of London, Mile End Road, London E1 4NS, UK

# Other Bacterial Microcompartments

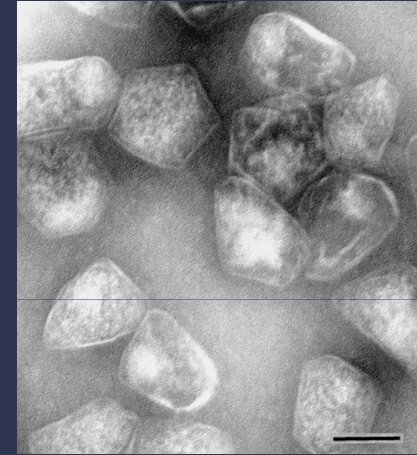
A



B



C

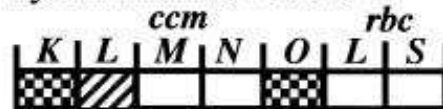


# Other Bacterial Microcompartments

*Halothiobacillus neapolitanus*\*



*Synechococcus* PCC7942\*\*



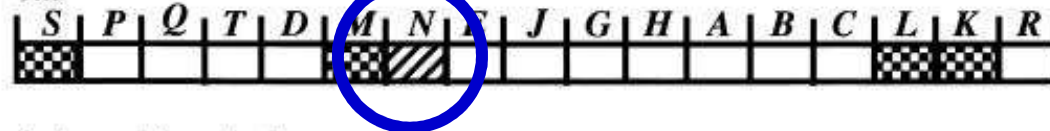
*Synechocystis* PCC6803\*\*\*



*Prochlorococcus marinus* MIT9313\*\*\*\*

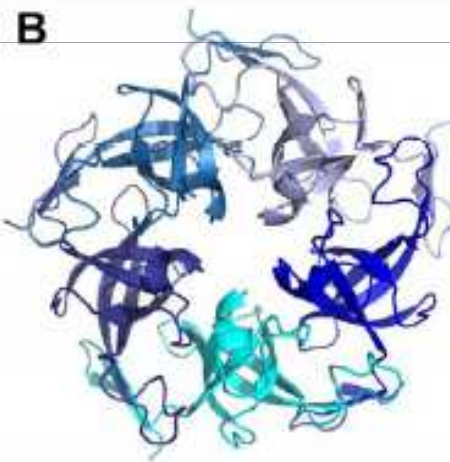
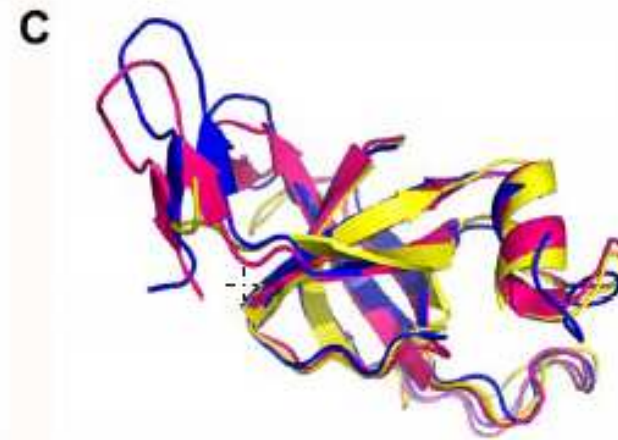
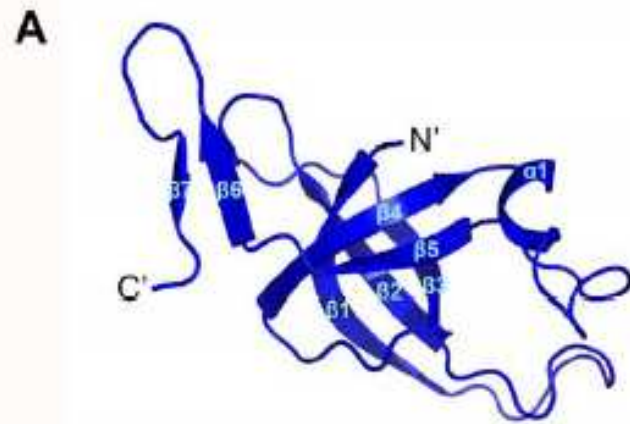


*Salmonella typhimurium*/*Escherichia coli*  
*eut*



*Salmonella enterica*  
*pdu*

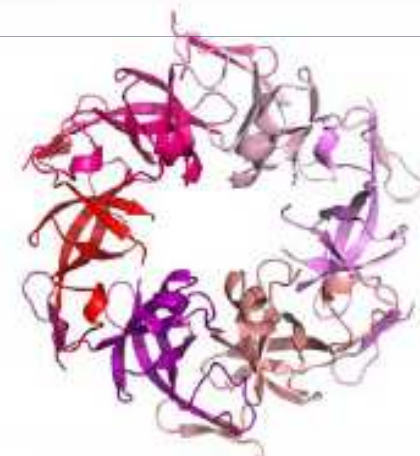




**Ccml**

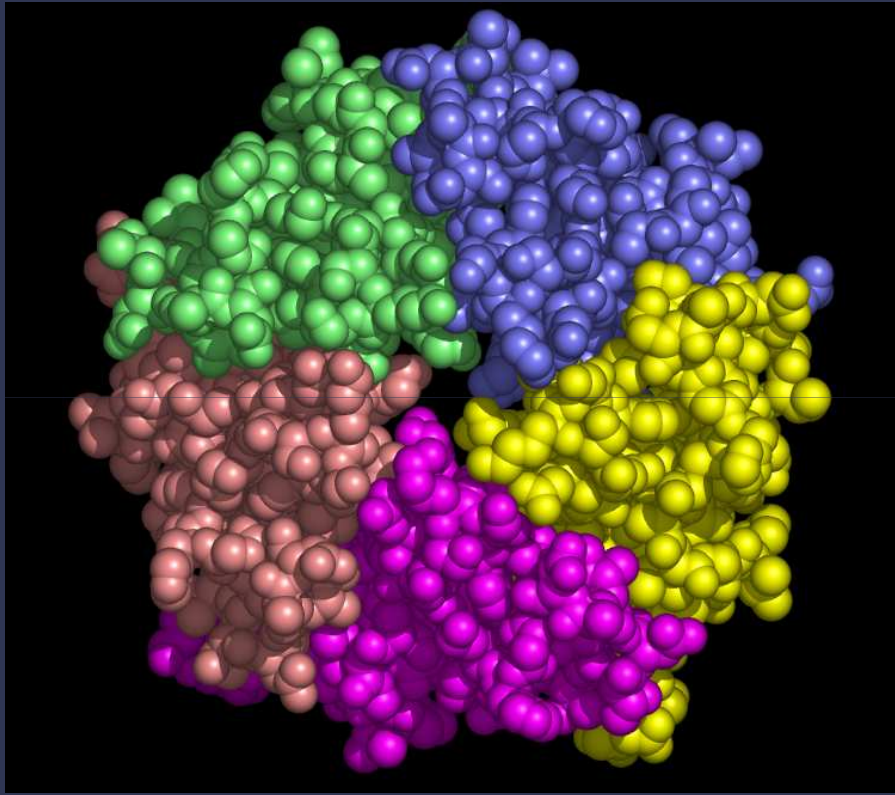


**OrfA**

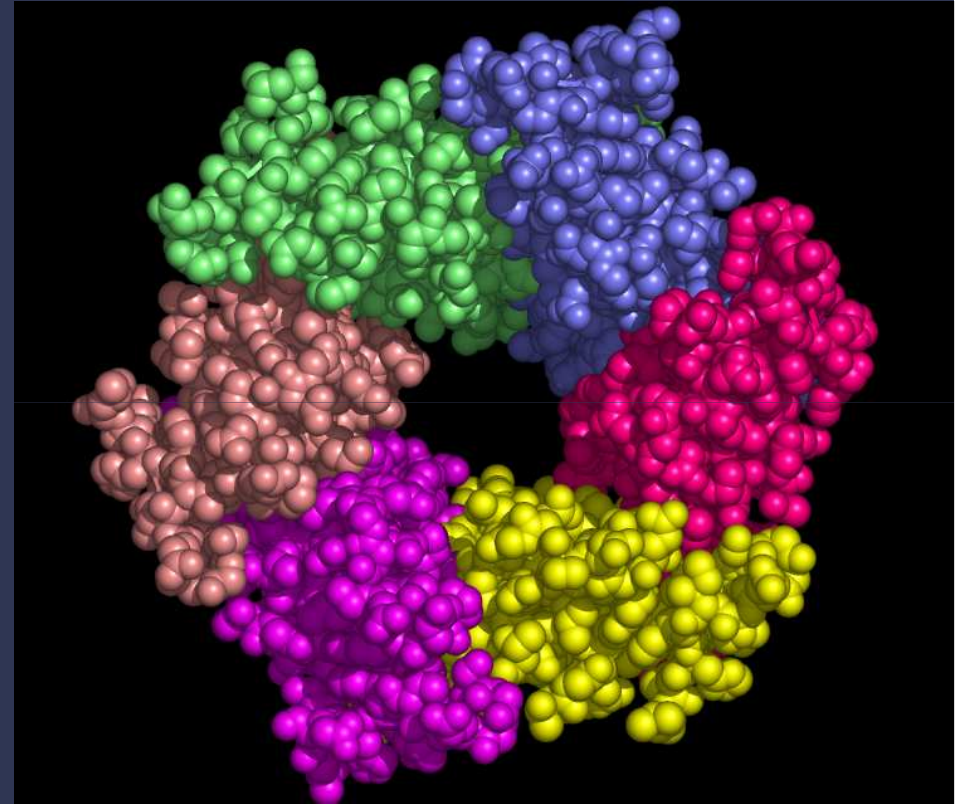


**EutN**

# OrfA/CcmL



# EutN



And other unusual shell protein building blocks forthcoming....

# Summary

- We have sufficient structural data to model the carboxysome shell
- Bacterial microcompartments are surprisingly widespread and diverse in function
- And likely to be architecturally diverse in their building blocks



# Opportunities.....

- Cyanobacterial Annotation Tools for Undergraduates (DOE)/Database needs
- Cyanobacterial Molecular Biology Workshop, June 2010 Lake Delevan, Wisconsin

# Acknowledgements

- **UCLA**

Todd Yeates' group

- **USM**

Sabine Heinhorst &  
Gordon Cannon's group

- **JGI and UC Berkeley**

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Kostas Mavramommatis

Edwin Kim

Michael Klein

Jay Kinney

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**USDA and DOE**