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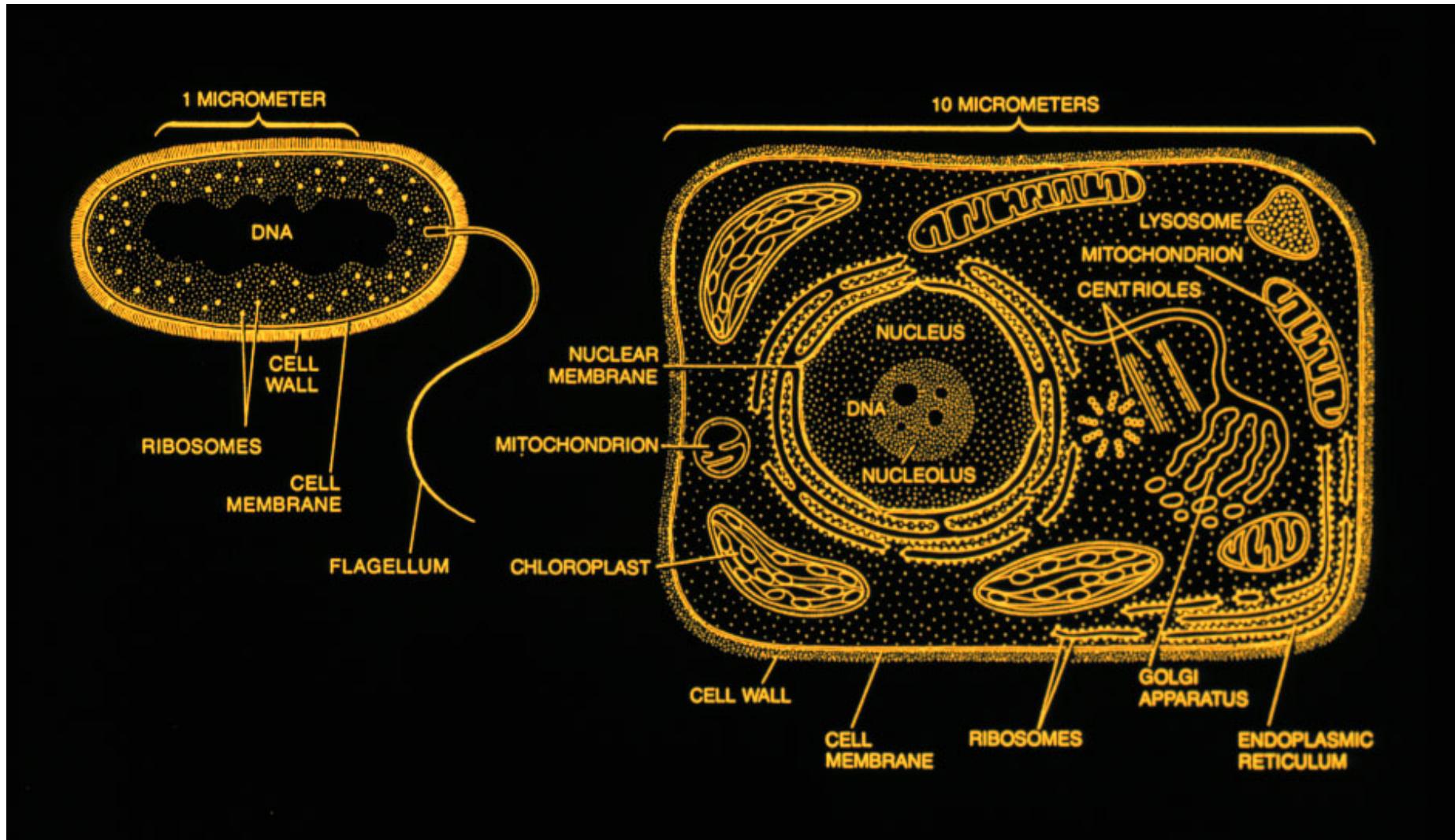
# **Planctomycetes and PVC superphylum bacteria and their similarities in structure and function to eukaryotes**

John A. Fuerst

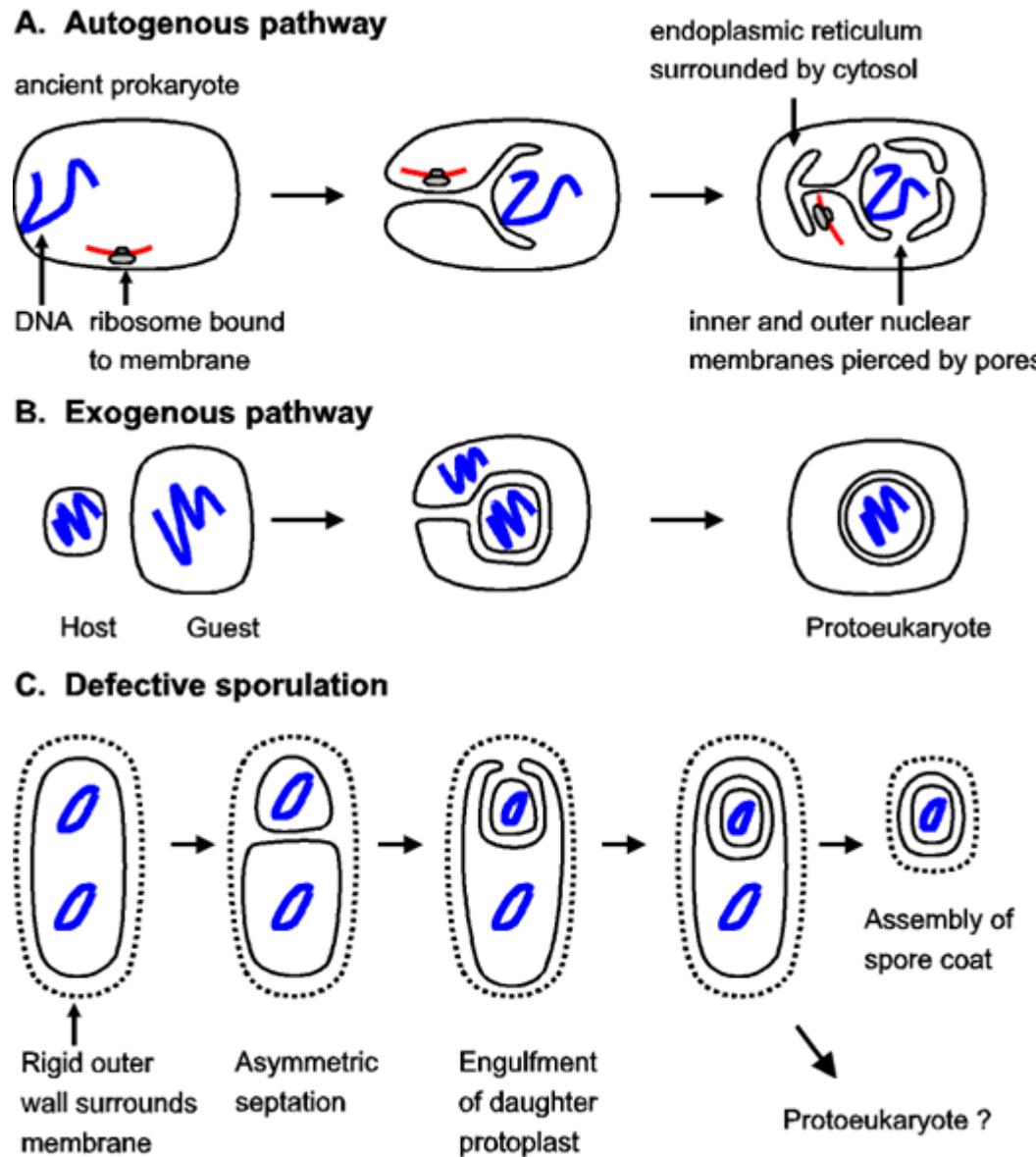
**School of Chemistry and Molecular Biosciences,  
The University of Queensland,  
St.Lucia 4072, Australia**

Tutorial for ICTS EvoCell 2012, NCBS Bangalore

# Prokaryote vs. Eukaryote



## Major ways proposed for evolving endomembranes and the eukaryotic nucleus



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From: Peter Cook, Principles of Nuclear Structure and Function Wiley, 2001

# Evolutionary advantages of the nucleus?

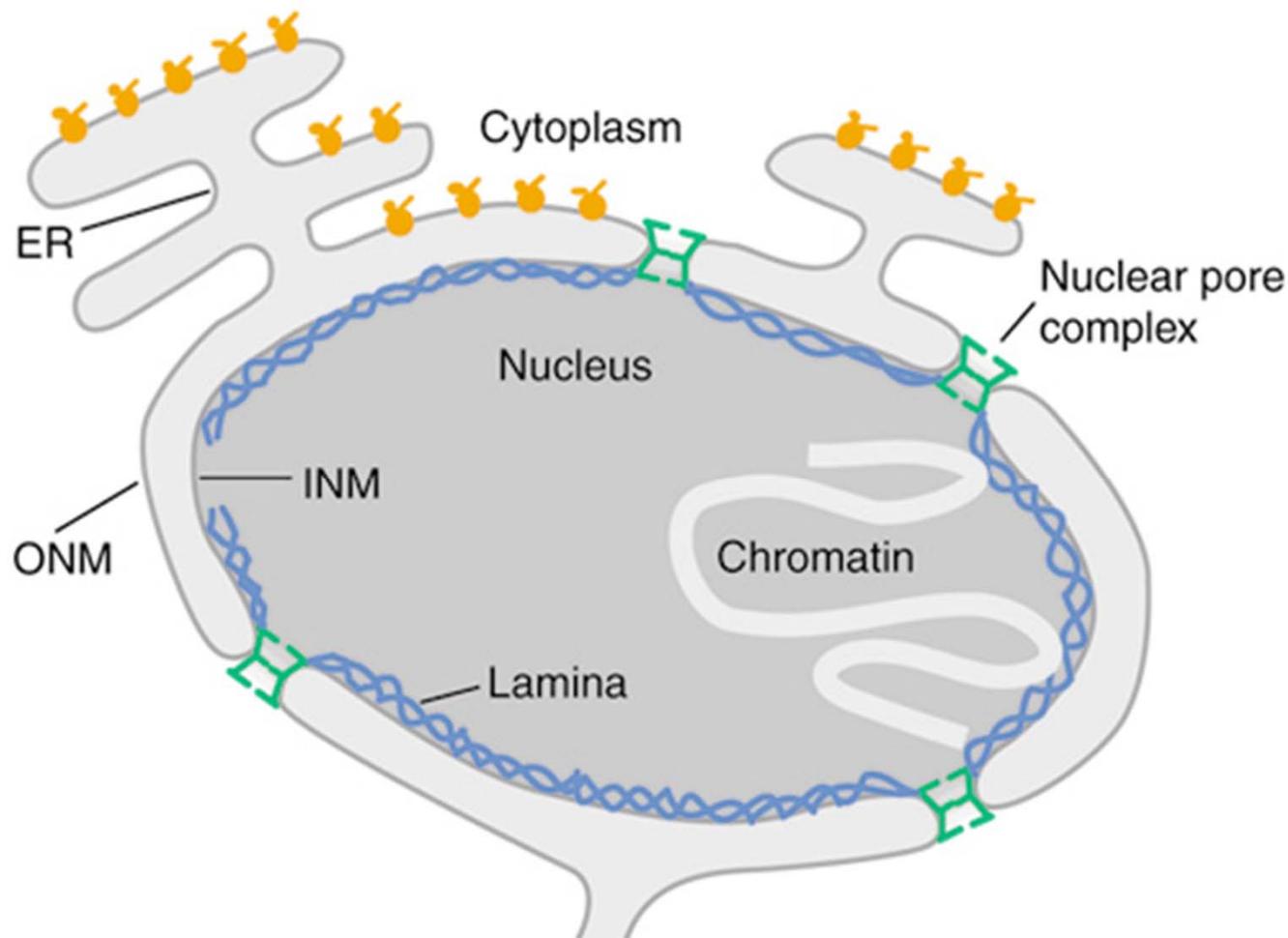
# Evolutionary advantages of the nucleus?

- 1) overcomes diffusion limits in large cells - advantages of 'molecular crowding'
- 2) Protection of DNA from shearing forces on mitosis? Protection from oxidation products after the mitochondrial invasion?
- 3) Separation of slow transcription & RNA processing from fast translation - prevent translation of unprocessed nonsense pre-mRNA
- 4) Corollary of endomembranes accompanying endocytosis/phagocytosis as a new eukaryotic modes of nutrition

OR

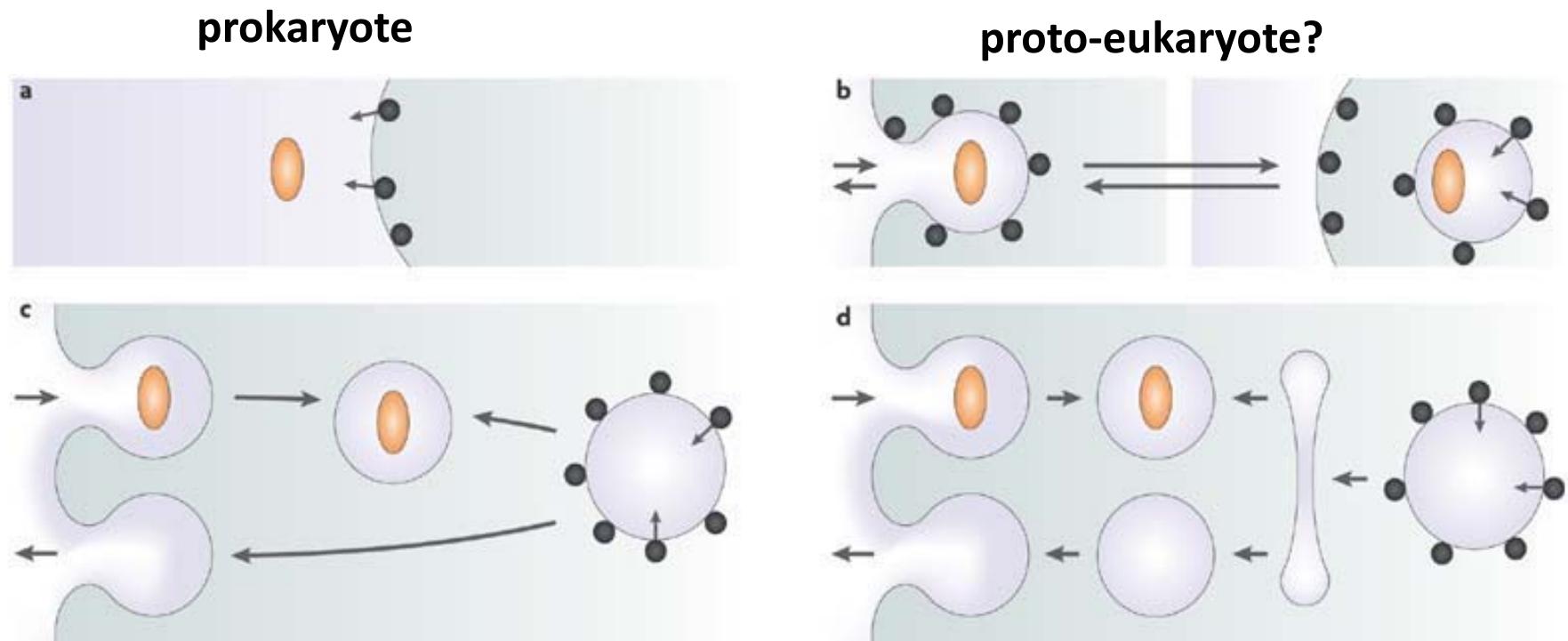
a neutral non-adaptive evolutionary accident?

## Eukaryote nuclear envelope – ER connection – origins of endomembranes and nucleus connected?



From: Schirmer and Gerace *Genome Biology* 2002 3:reviews1008.1

# Is the origin of endocytosis and vesicle formation a key to endomembrane evolution & thus cell compartments?



Nature Reviews | Genetics

almost eukaryote (LECA?)

From: De Duve, C 2007. The origin of eukaryotes: a reappraisal. Nature Reviews Genetics 8, 395 - 403

Do any organisms living today resemble such proto-eukaryotes?

i.e.

are there any 'living fossils' preserving features of the proto-eukaryote?

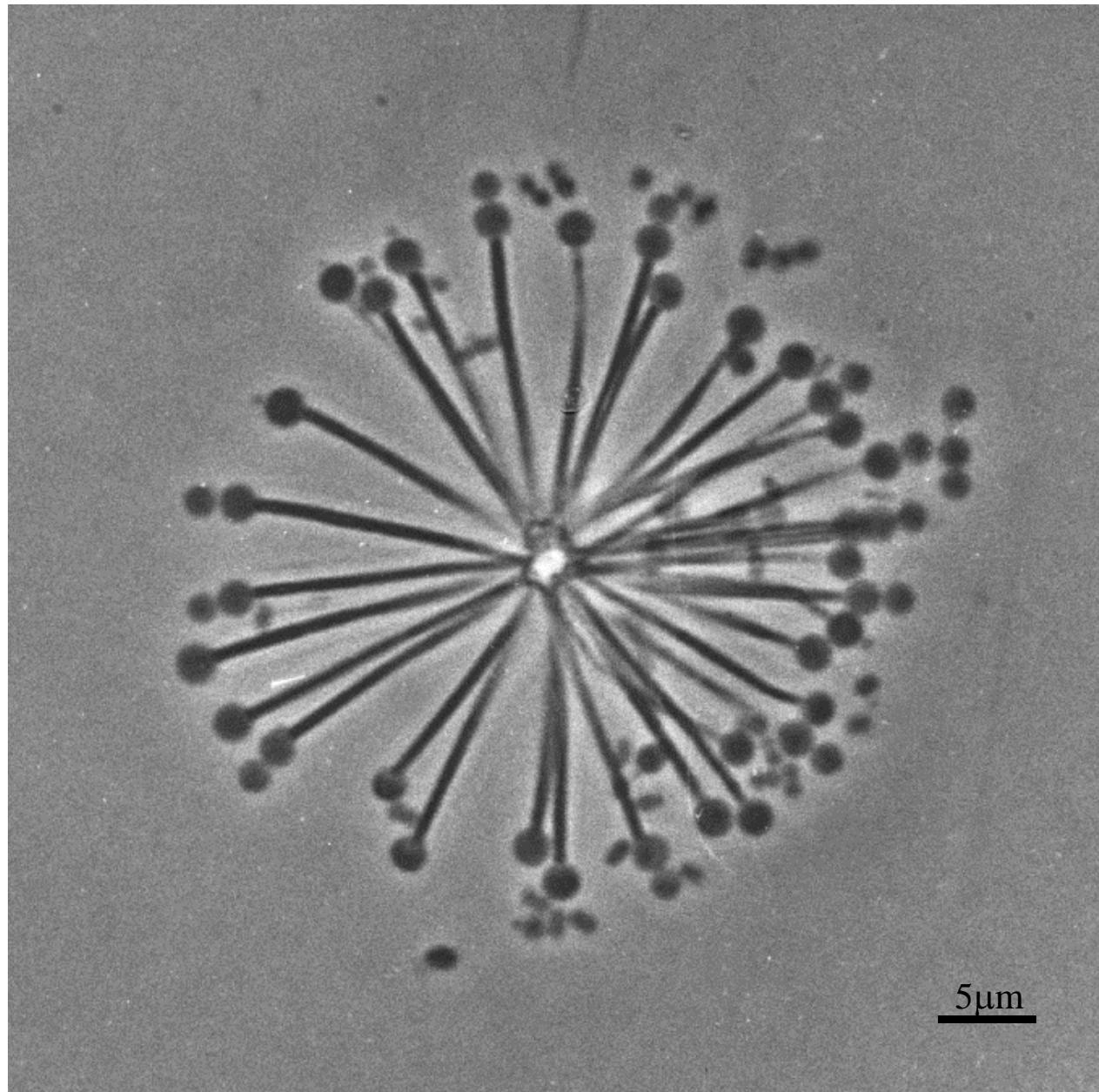
if so, what do they imply about evolutionary models for compartmentalization origins?

# A Bacterial model for origin of eukaryote compartmentalization? - The Planctomycetes

**Planctomycetes** are:

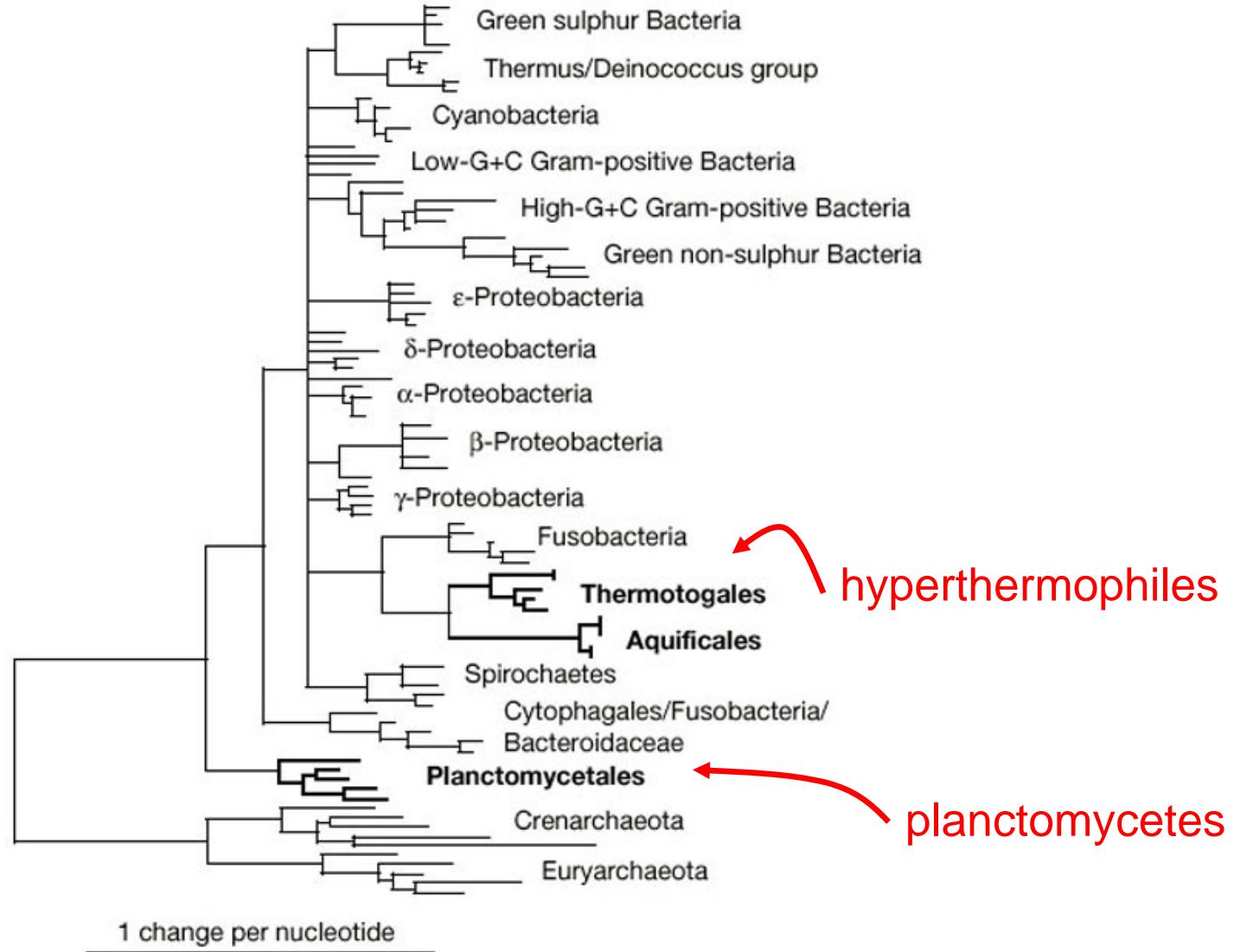
- Members of **Domain Bacteria** (e.g. by 16S rRNA sequence trees)
- Form distinct **separate Phylum** within Domain –but may be member of the '**PVC' superphylum**
- Budding, aquatic/soil bacteria
- Cell walls mostly protein - possess **no peptidoglycan** (unlike most other domain Bacteria members)
- All possess **compartments with DNA enveloped by membrane**

Uncultured *Planctomyces bekefii* rosette from lake water – stalks are non-cellular

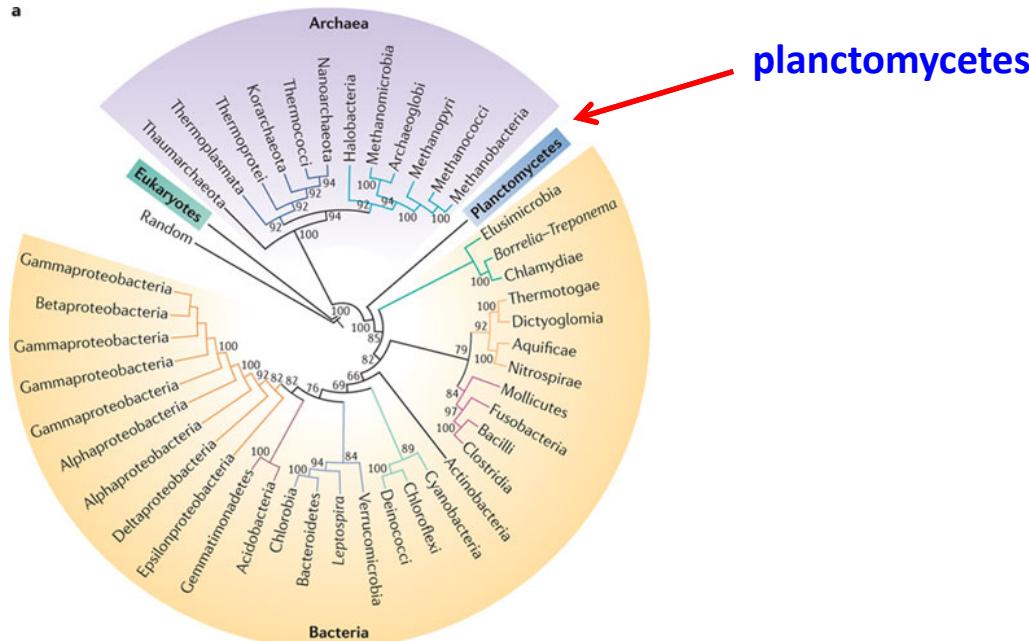


# Are planctomycetes the deepest branching Bacterial phylum?

From: Brochier, C.  
and Philippe , H.  
2002. Phylogeny:  
A non-  
hyperthermophilic  
ancestor for  
Bacteria  
*Nature* 417, 244

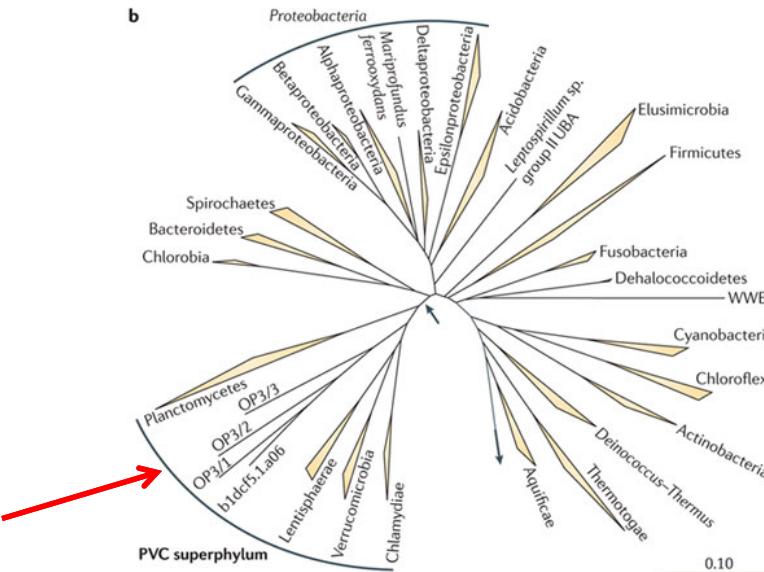


## Proteome-derived tree



## 23S rRNA-derived tree

PVC superphylum



# Markers of the unusual evolutionary significance of planctomycetes

- Sterols & simple sterol synthesis pathway in *Gemmata obscuriglobus*
- Mixed ether and ester ladderane lipids in anammox planctomycetes
- Unique form of anaerobic chemolithoautotrophy in anammox planctomycetes
- C<sub>1</sub> - transfer enzymes in *Gemmata* and *Rhodopirellula* with phylogenetic position between the Archaea and Bacteria
- Deep-branching position in 16S rRNA tree of Bacteria when slowly-evolving positions used
- Membrane-bounded compartments & condensed nucleoids

## Similarities of planctomycetes to eukaryotes

- Cells compartmentalized via internal membranes (nucleoid in a compartment with at least one surrounding membrane)
- In *Gemmata obscuriglobus*, nuclear compartment surrounded by 2-membrane envelope (within planctomycete major membrane-bounded compartment)
- Condensed chromosomes
- Endocytosis – uptake of external proteins via membrane infolding
- MC (membrane-coat) proteins homologous with eukaryotic clathrins
- Sterols (*G. obscuriglobus*)

# Compartmentalized cells of planctomycetes

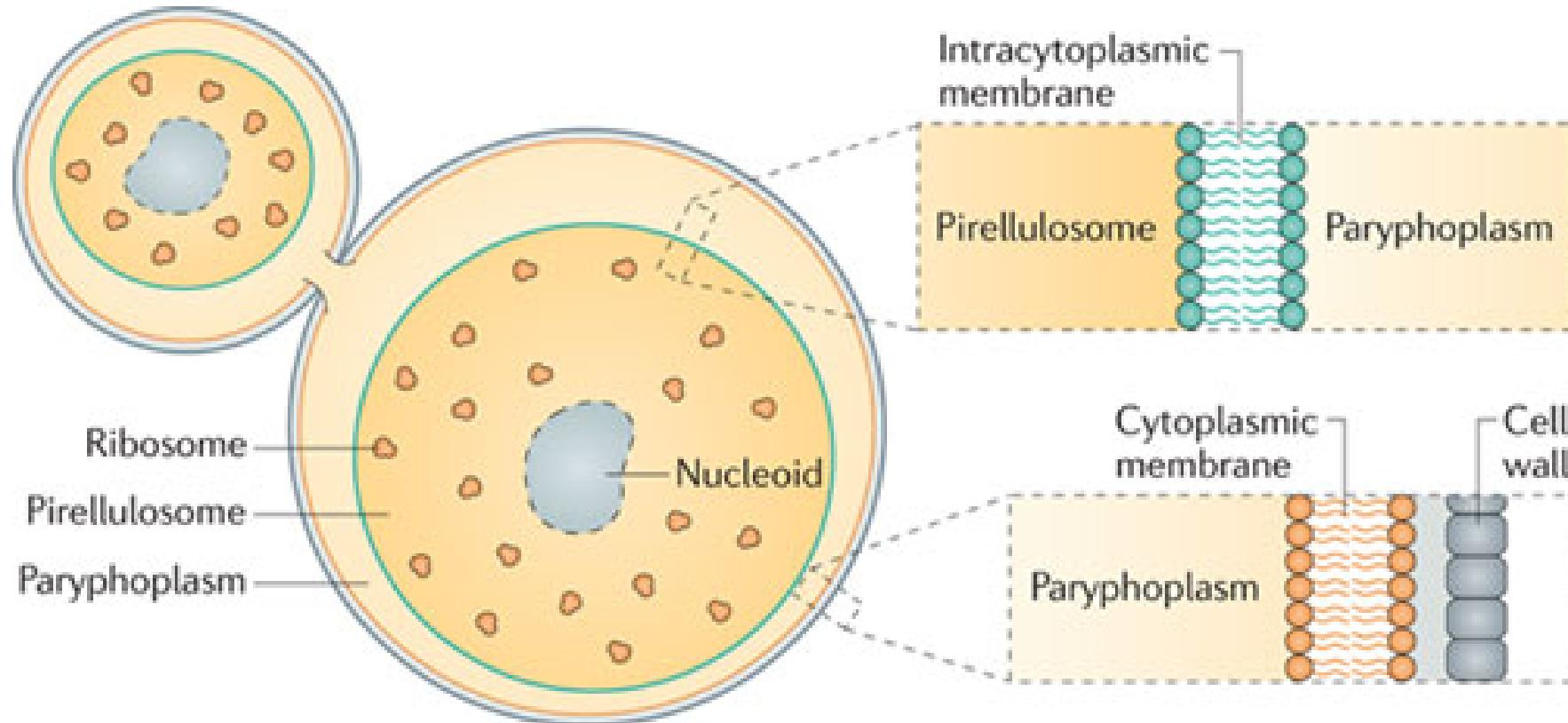
- All planctomycetes examined share a common **compartmentalized cell plan** including an **intracytoplasmic membrane** separating 2 major internal cell regions
  - - ribosome-free paryphoplasm
  - - ribosome-rich pirellulosome
- Some planctomycetes have further compartments – **Gemmata** with a **double-membrane-bounded nuclear compartment** within the pirellulosome

All known planctomycetes share a new type of bacterial cell compartment: the pirellulosome

A cell compartment :

- bounded by a single membrane
- contains nucleoid and electron-dense ribosome-like particles, all cell DNA + most (but not all) cell RNA

# The shared planctomycete plan

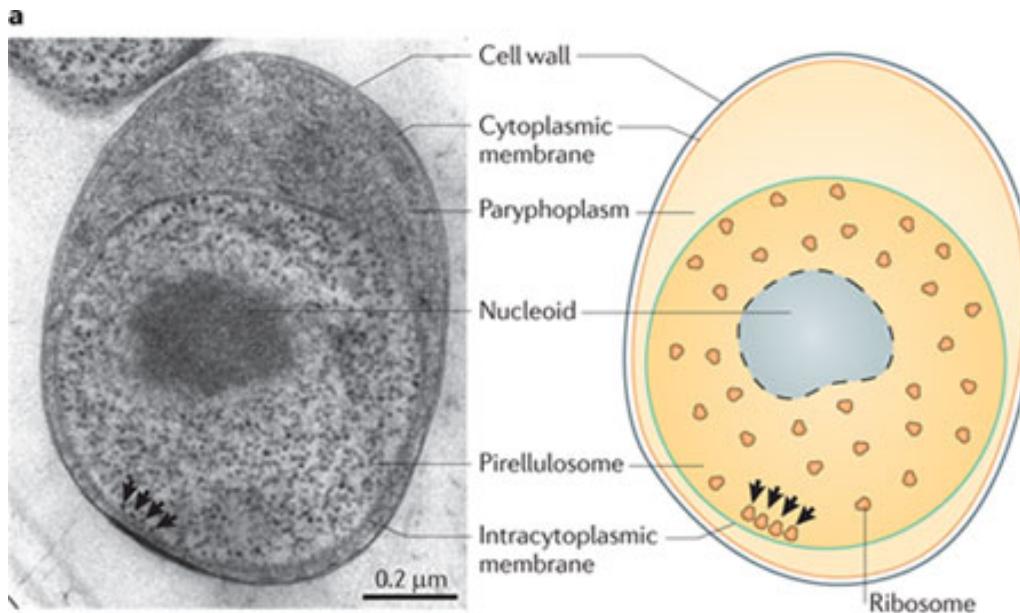


Nature Reviews | Microbiology

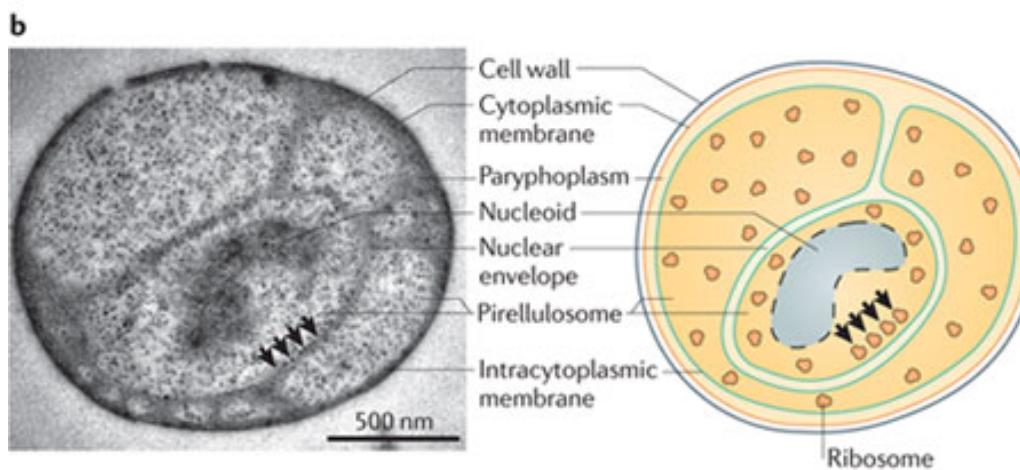
From: Fuerst JA, Sagulenko E. 2011. *Nat Rev Microbiol*. 9(6):403-13.

## Cell structure of Planctomycetes – variation within a shared plan

*Pirellula*



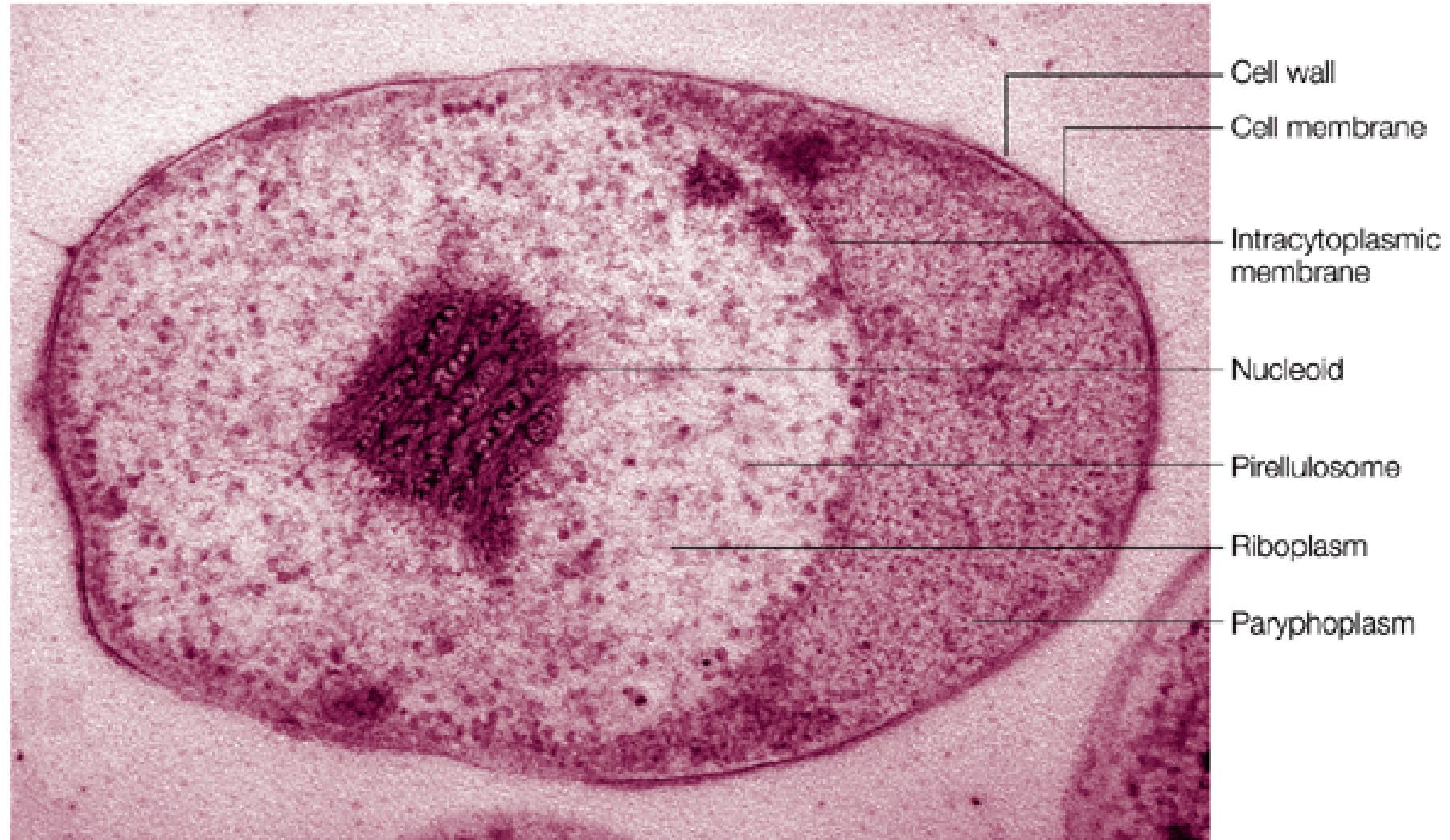
*Gemmata*



Nature Reviews | Microbiology

From: Fuerst JA, Sagulenko E. 2011. *Nat Rev Microbiol.* 9(6):403-13.

**Cell plan of *Pirellula* group planctomycetes- TEM of sectioned cryosubstituted cell of *Blastopirellula marina***



From: Nature Reviews Microbiology 1:11-12 (2003)

Nature Reviews | Microbiology

# Planctomycetes may belong to a postulated wider PVC Superphylum

- At least 3 phyla:

*Planctomycetes*

*Verrucomicrobia*

*Chlamydiae*

- + phyla such as marine *Lentisphaerae* and *Poribacteria*

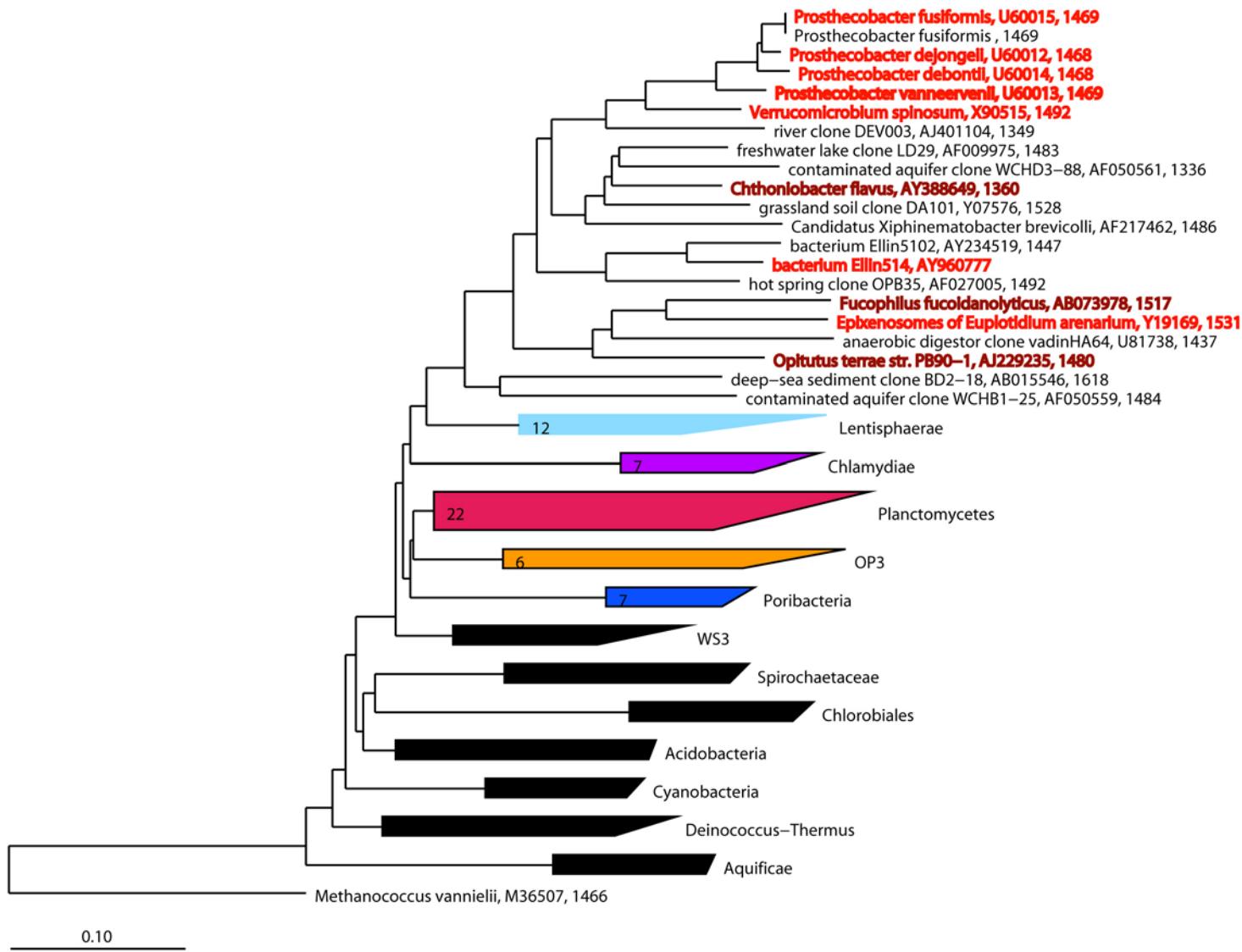
(Wagner M, Horn M. 2006. Curr Opin Biotechnol. 17:241-249).

# Hypothesis

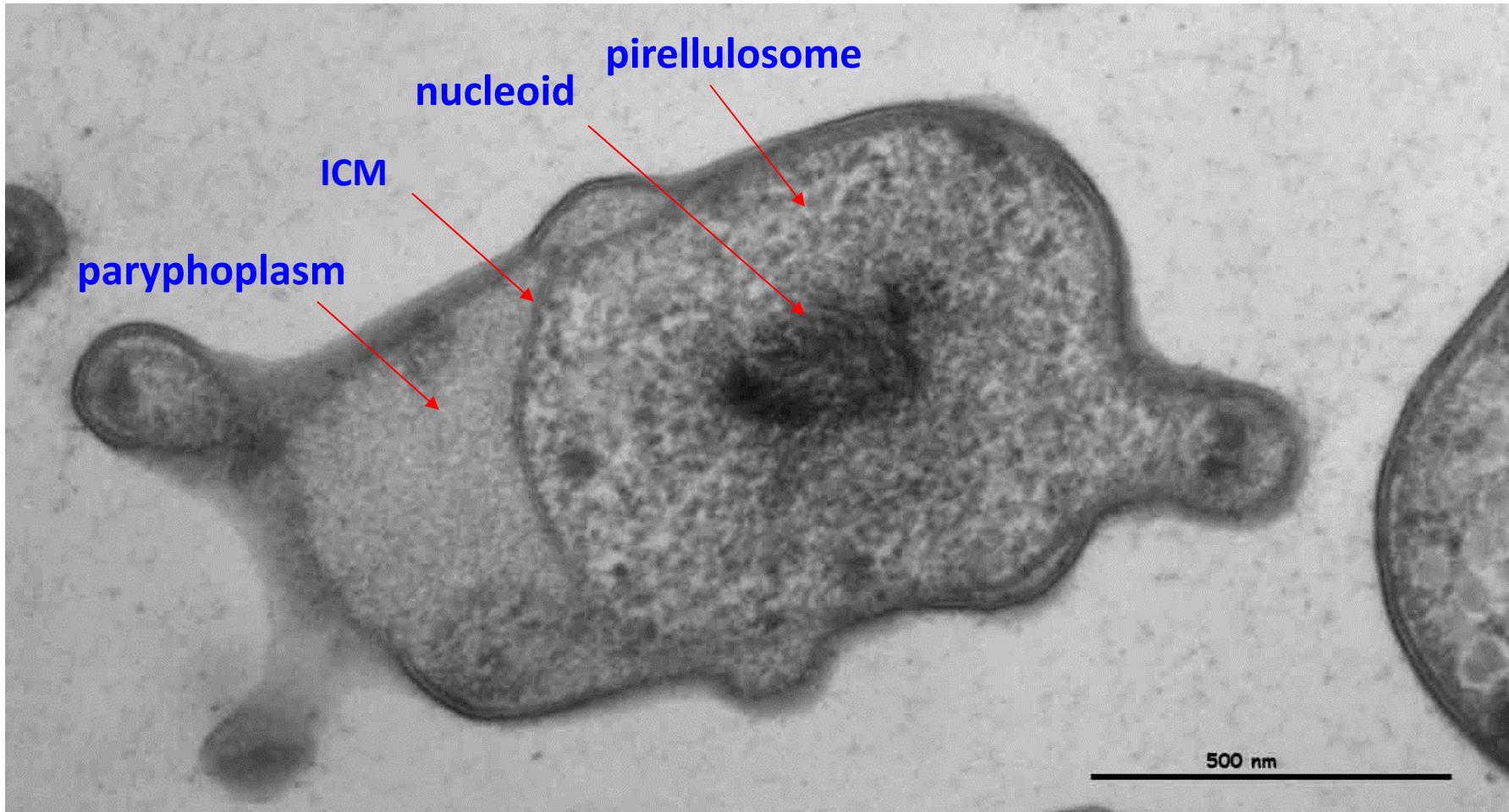
**If** PVC superphylum reflects valid evolutionary relationship

**then** phylum *Verrucomicrobia* and/or *Chlamydiae* or other members of the superphylum might display **cell compartments** homologous to those in Planctomycetes

## Phylum *Verrucomicrobia* : compartmentalized cells in all species examined

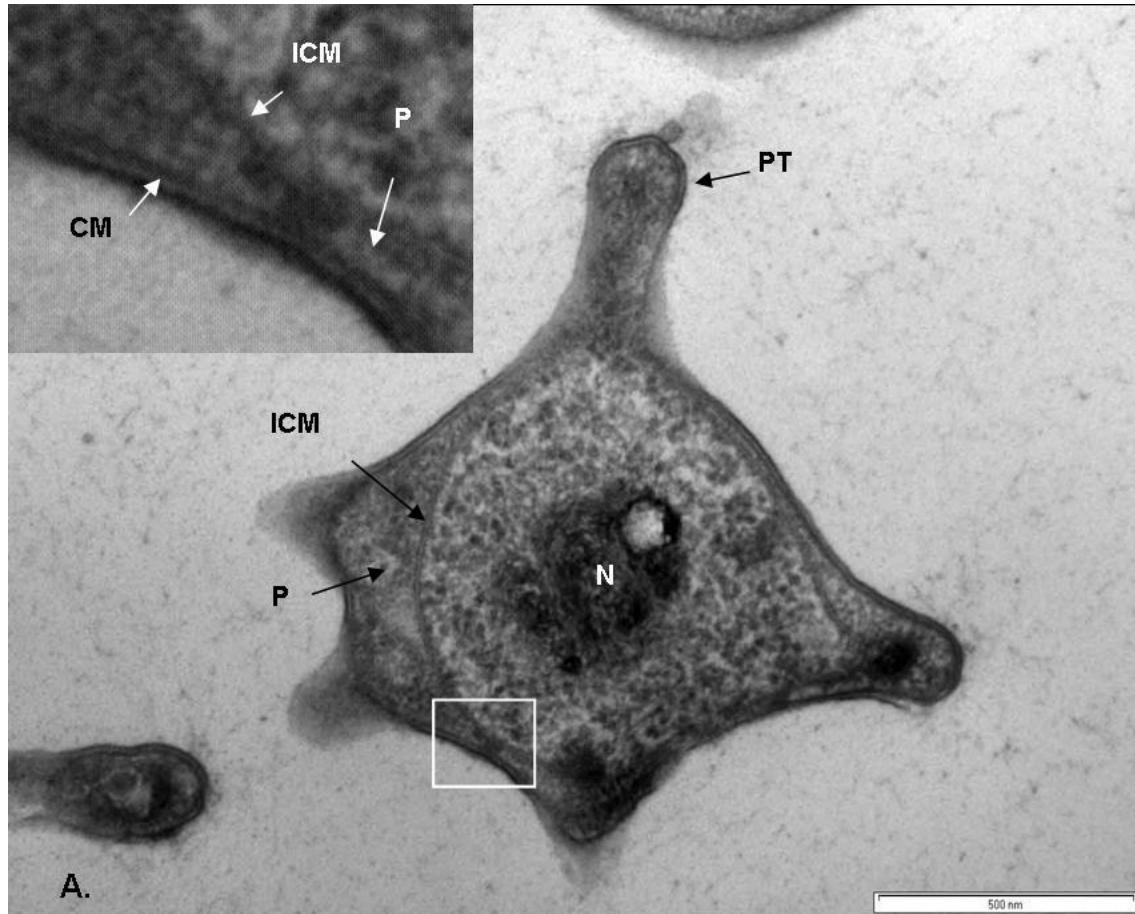


Major cell compartment of *Verrucomicrobium spinosum* is topologically equivalent to pirellulosome of planctomycete shared cell plan

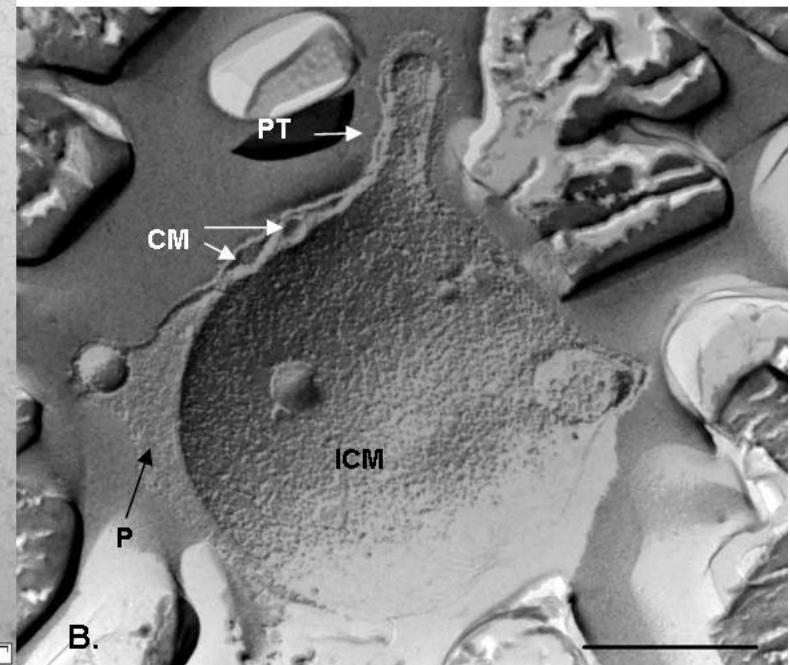


# *Verrucomicrobium spinosum* compartmentalization

Sectioned HPF-frozen

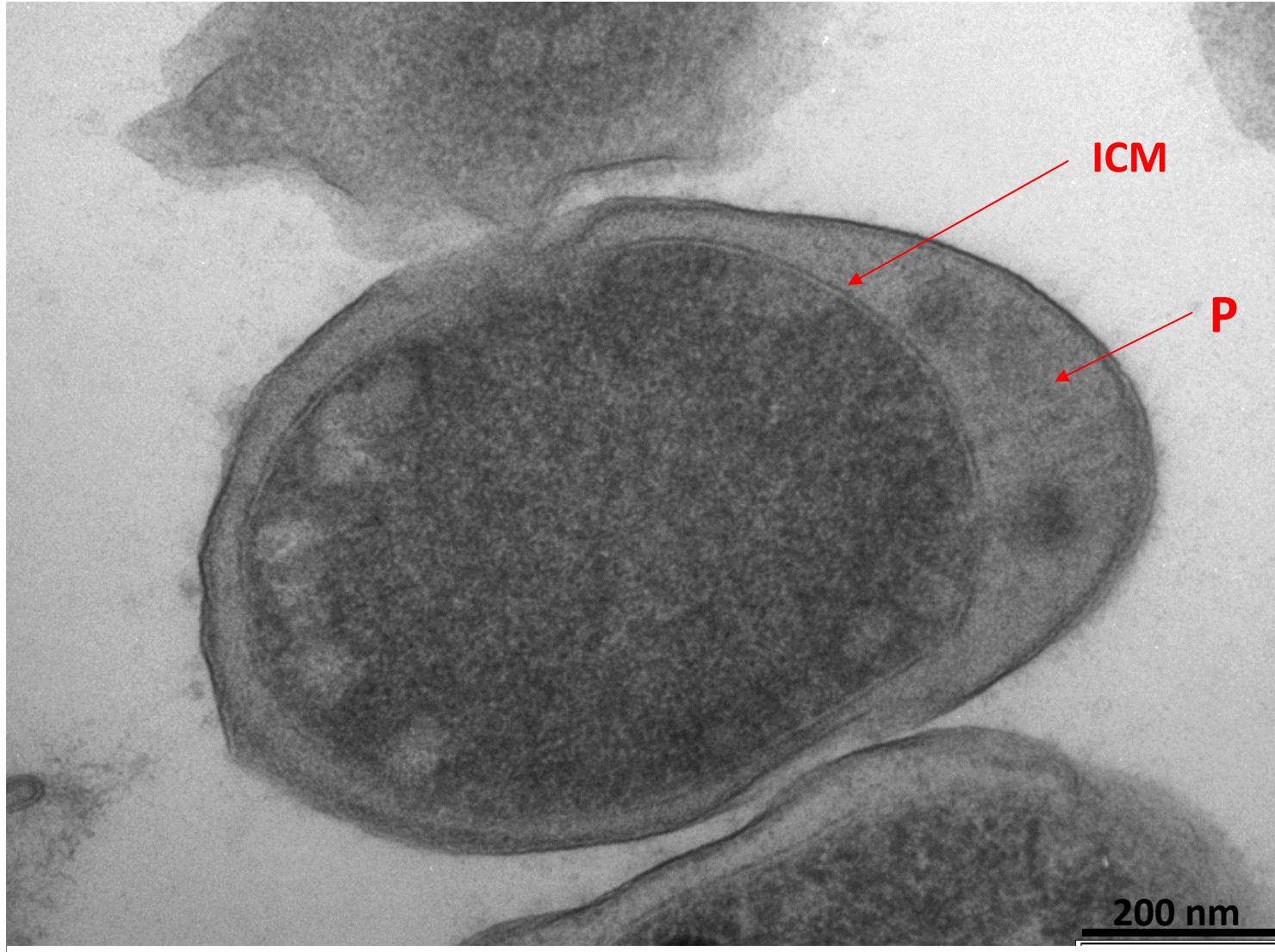


Freeze-fractured

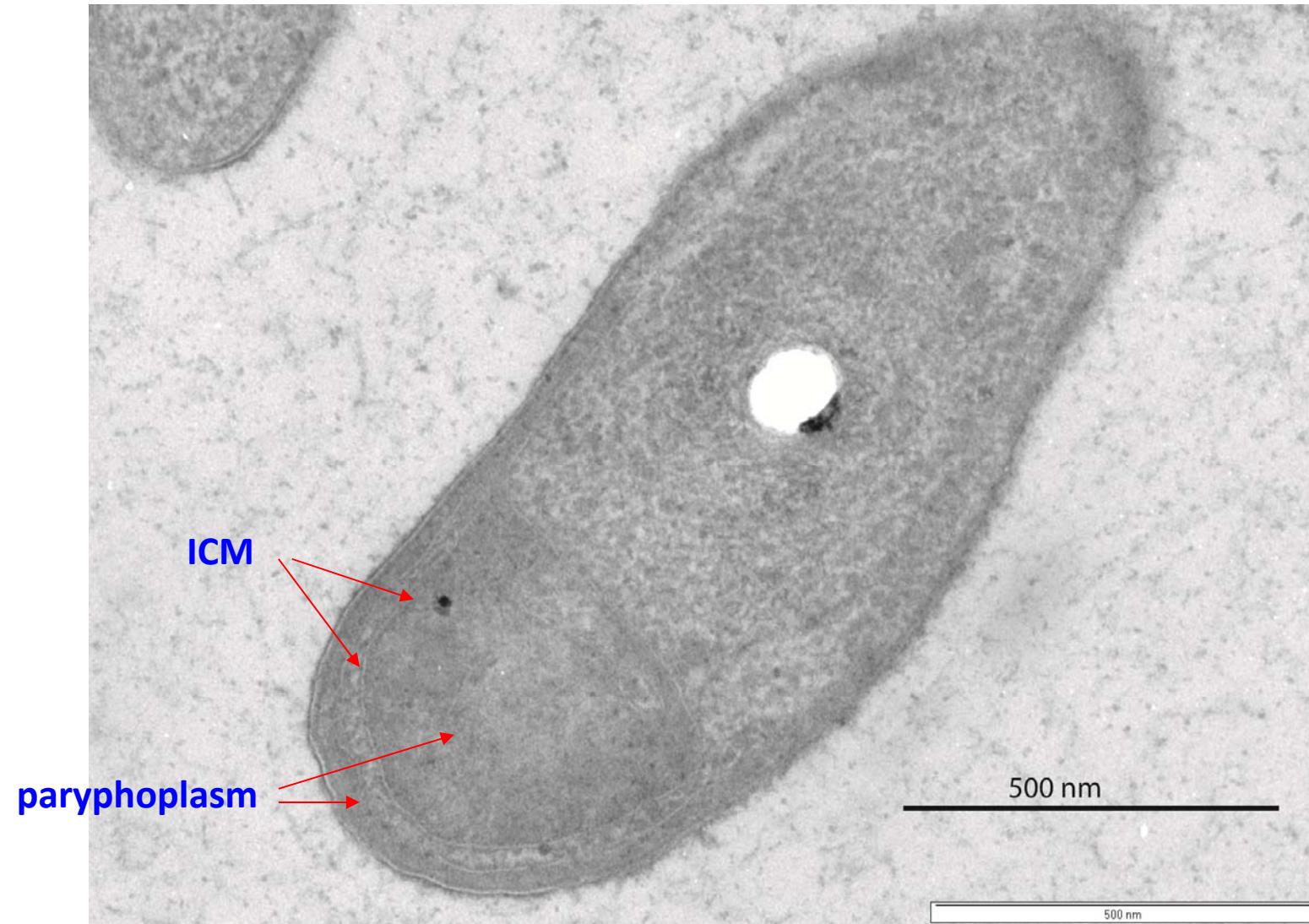


From: Kuo-Chang Lee, Richard I Webb, Peter H Janssen, Parveen Sangwan, Tony Romeo, James T Staley, John A Fuerst  
Phylum Verrucomicrobia representatives share a compartmentalized cell plan with members of bacterial phylum Planctomycetes  
BMC Microbiology 2009, 9:5 (8 January 2009)

Compartmented cells of marine *Lentisphaera araneosa* -  
another example of the planctomycete cell plan in a  
separate phylum *Lentisphaerae* of PVC

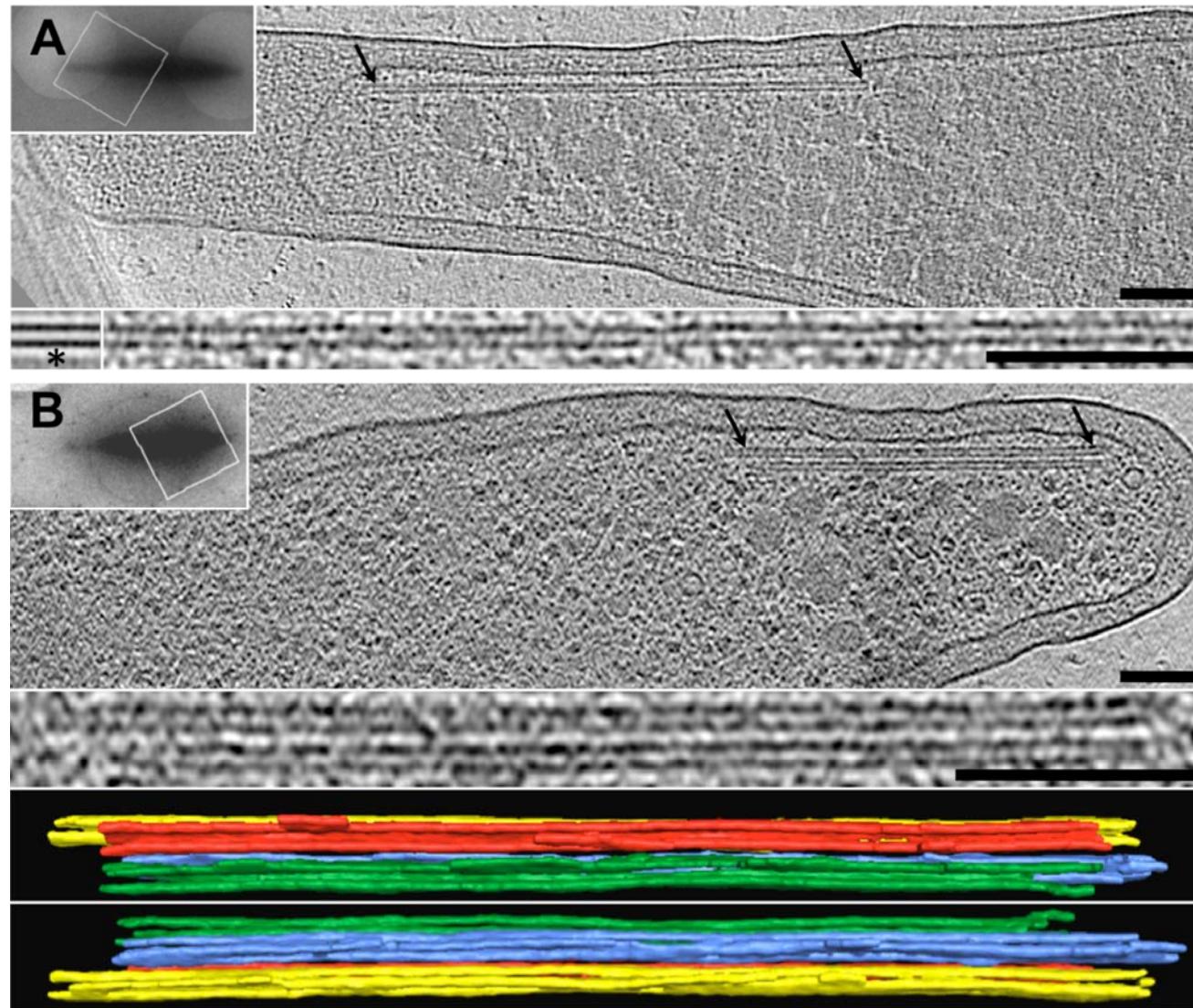


*Prosthecobacter dejongeii*, the first tubulin-syntheizing verrucomicrobia, has compartmentalized cells also! (when prepared by high-pressure freezing)

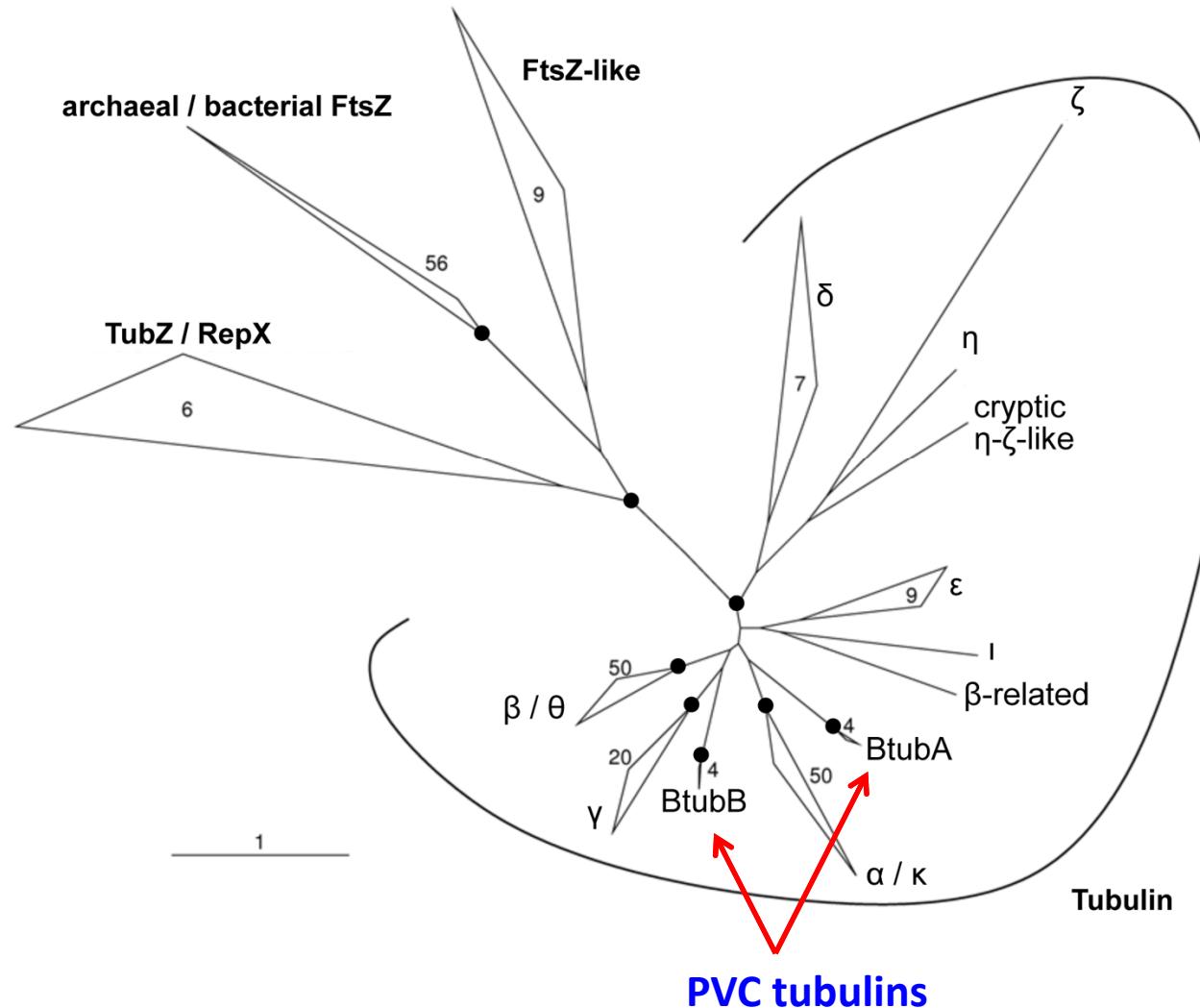


From: Kuo-Chang Lee, Richard I Webb, Peter H Janssen, Parveen Sangwan, Tony Romeo, James T Staley, John A Fuerst  
BMC Microbiology 2009, 9:5 (8 January 2009)

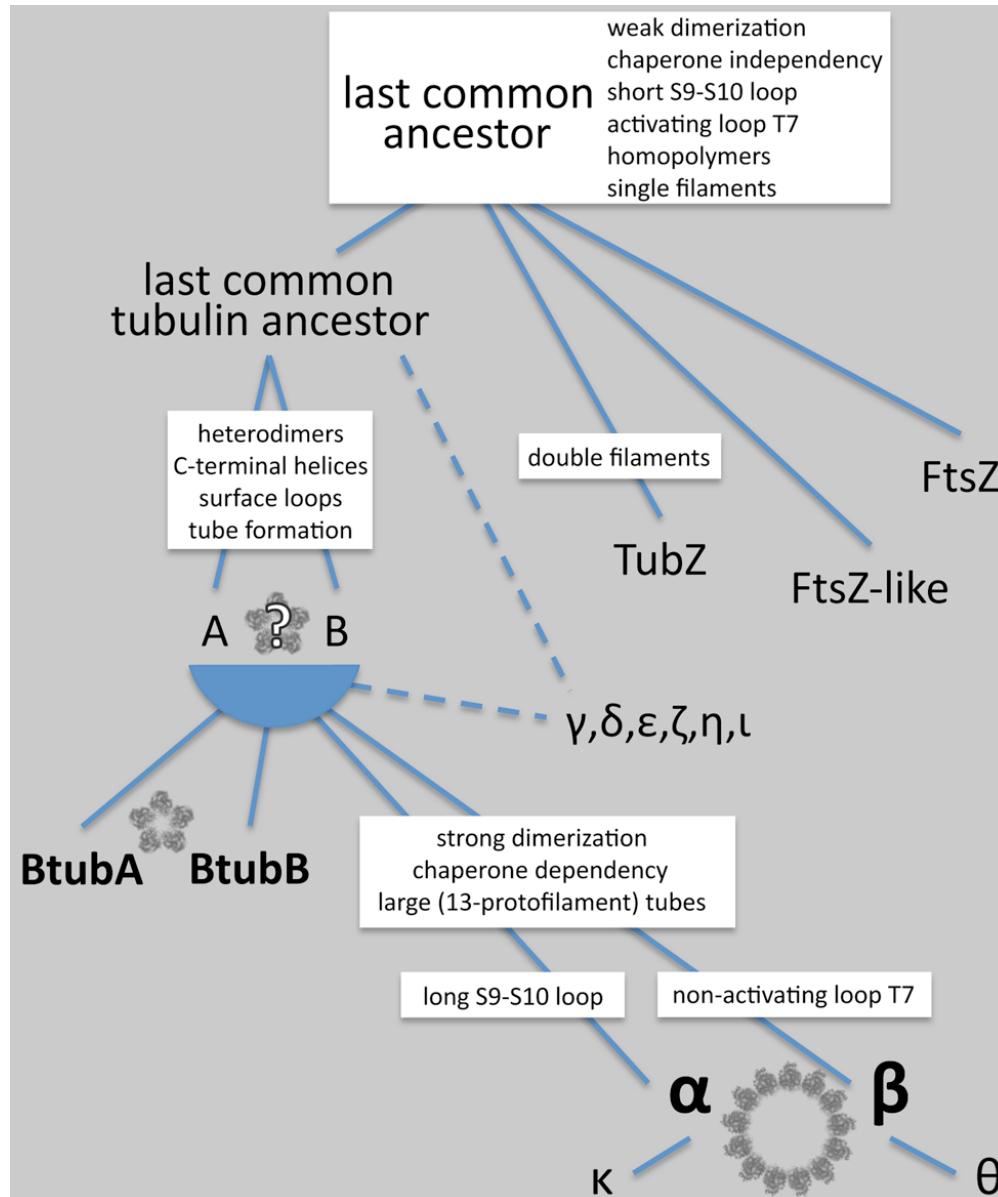
# *Prosthecobacter* (PVC member) has tubulin-based 5-protofilament microtubules



# PVC tubulins are eukaryote-like



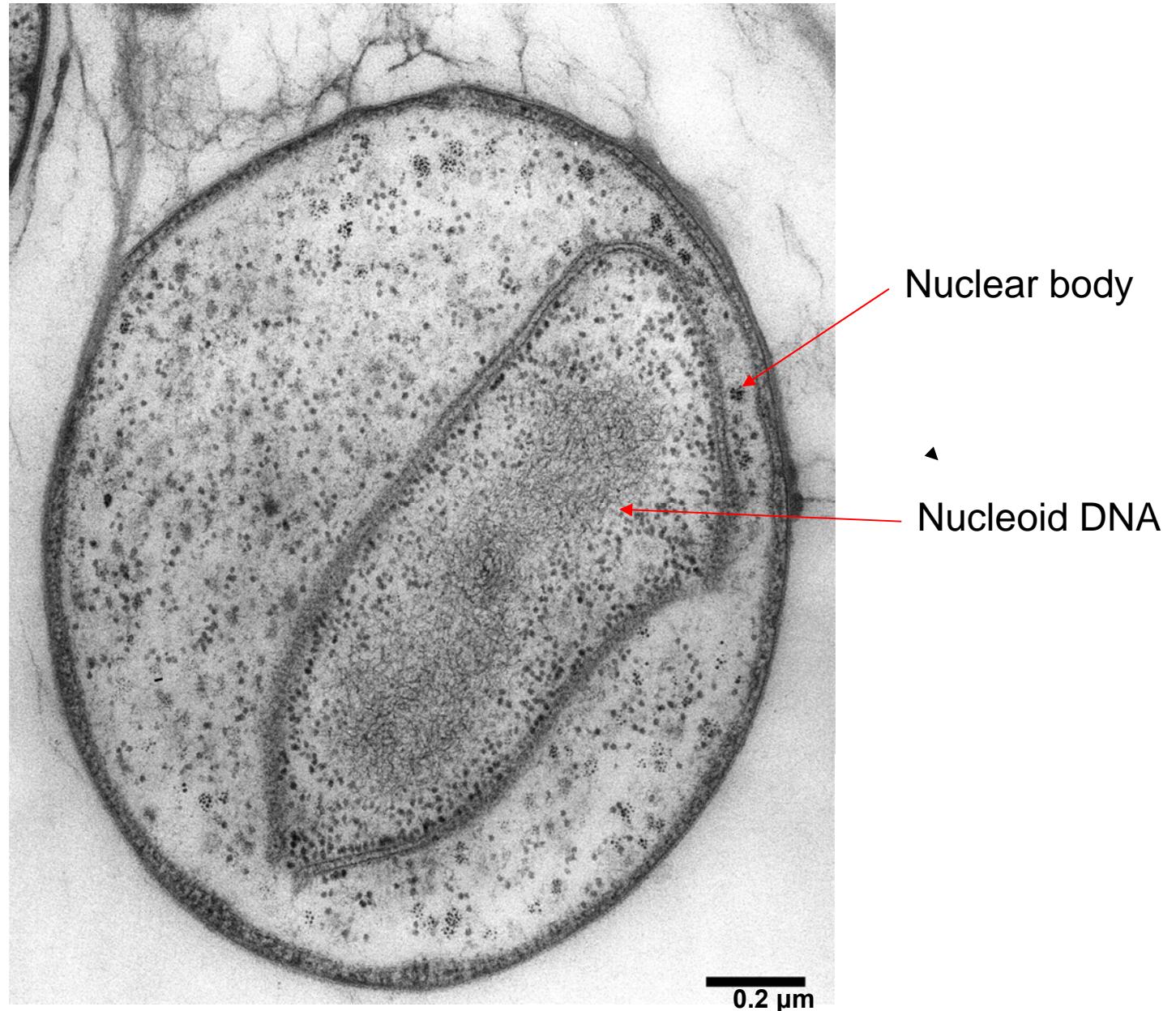
# Tubulin evolution model - PVC tubulins ancient, not result of HGT from modern eukaryote clade



# *Gemmata obscuriglobus*- a 'nucleated' bacterium?

- Freshwater aerobic heterotroph from Maroon Dam, Queensland
- Possesses 'double'-membrane envelope surrounding their nucleoid DNA (but envelope appears to be a folded single membrane topologically)
  - Lindsay et al. 2001 Archives of Microbiology 175: 413-429
- Possesses **sterols** and the simplest pathway for sterol synthesis known
- **radiation resistant** (UV and gamma rays) - 40X more resistant to UV radiation than *E. coli* (Lieber, A et al., J Bacteriol. Dec 2008):
- Draft 9 Mb (?) genome available

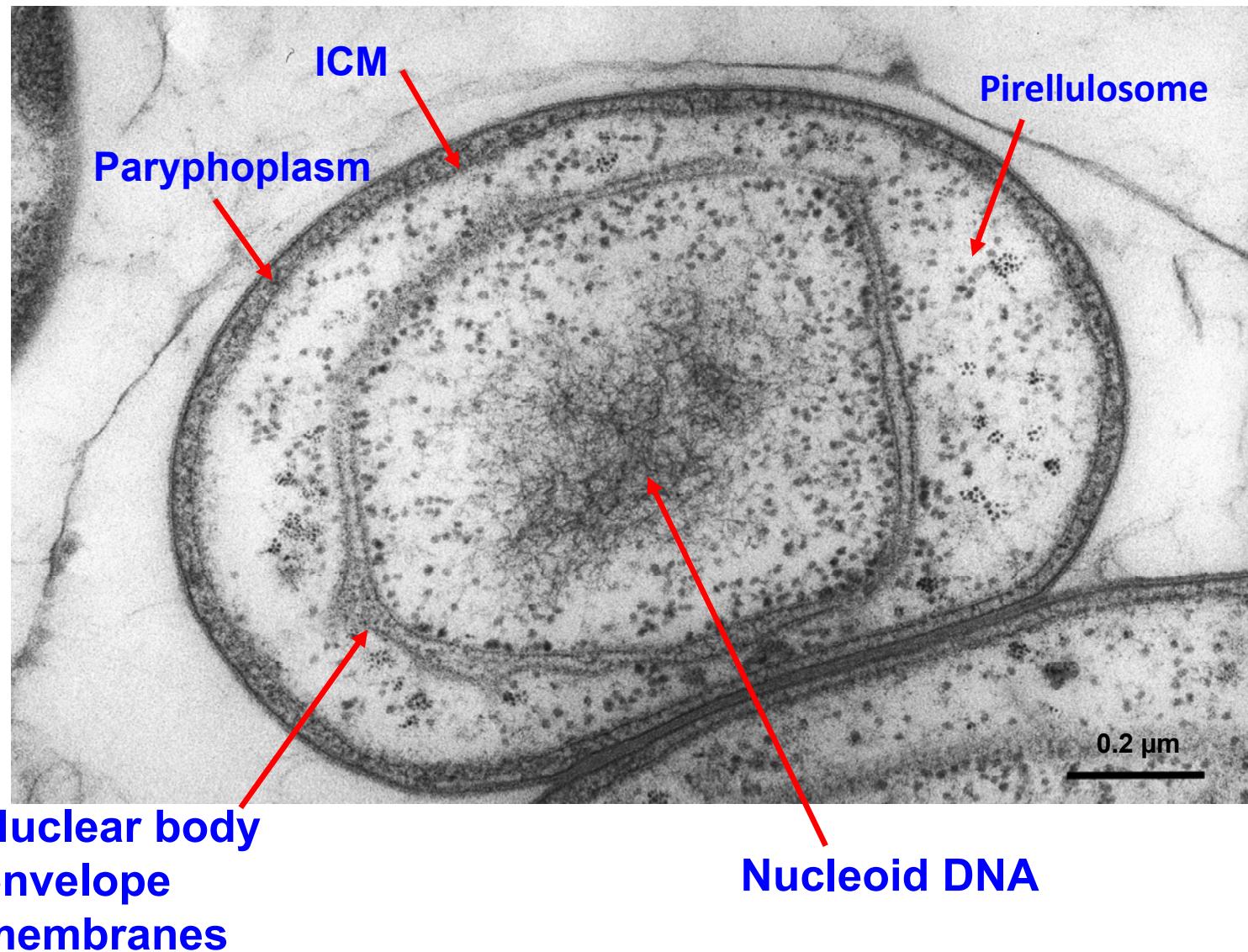
*Gemmata obscuriglobus* – section of cryosubstituted cell in TEM



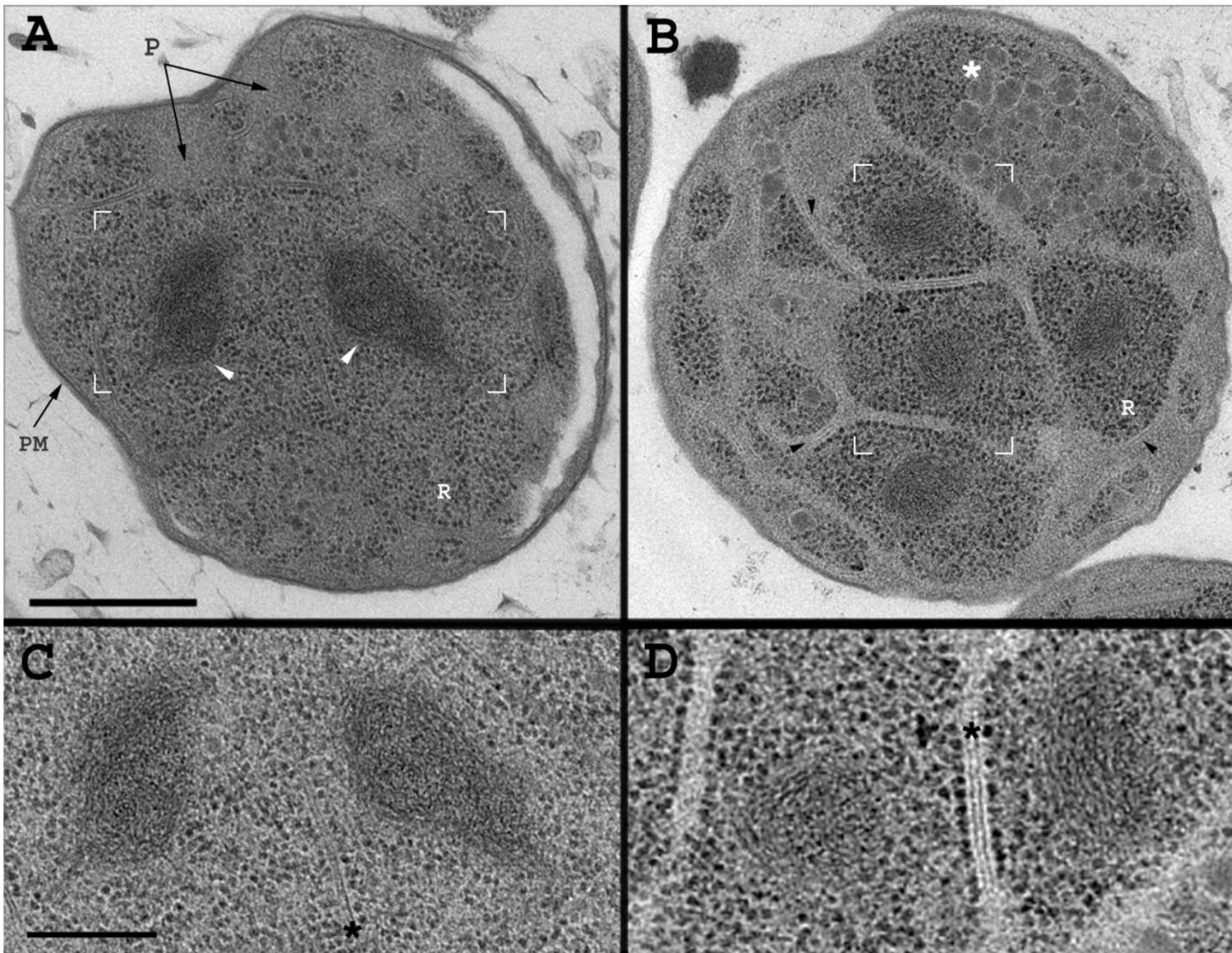
Lindsay MR, Webb RI, Strous M, Jetten MS, Butler MK, Forde RJ, Fuerst JA.  
Arch Microbiol. 2001 Jun;175(6):413-29.

*Gemmata obscuriglobus*: a planctomycete with a double membrane-bounded nucleoid within the pirellulosome

Thin section of cryosubstituted *Gemmata obscuriglobus* bud



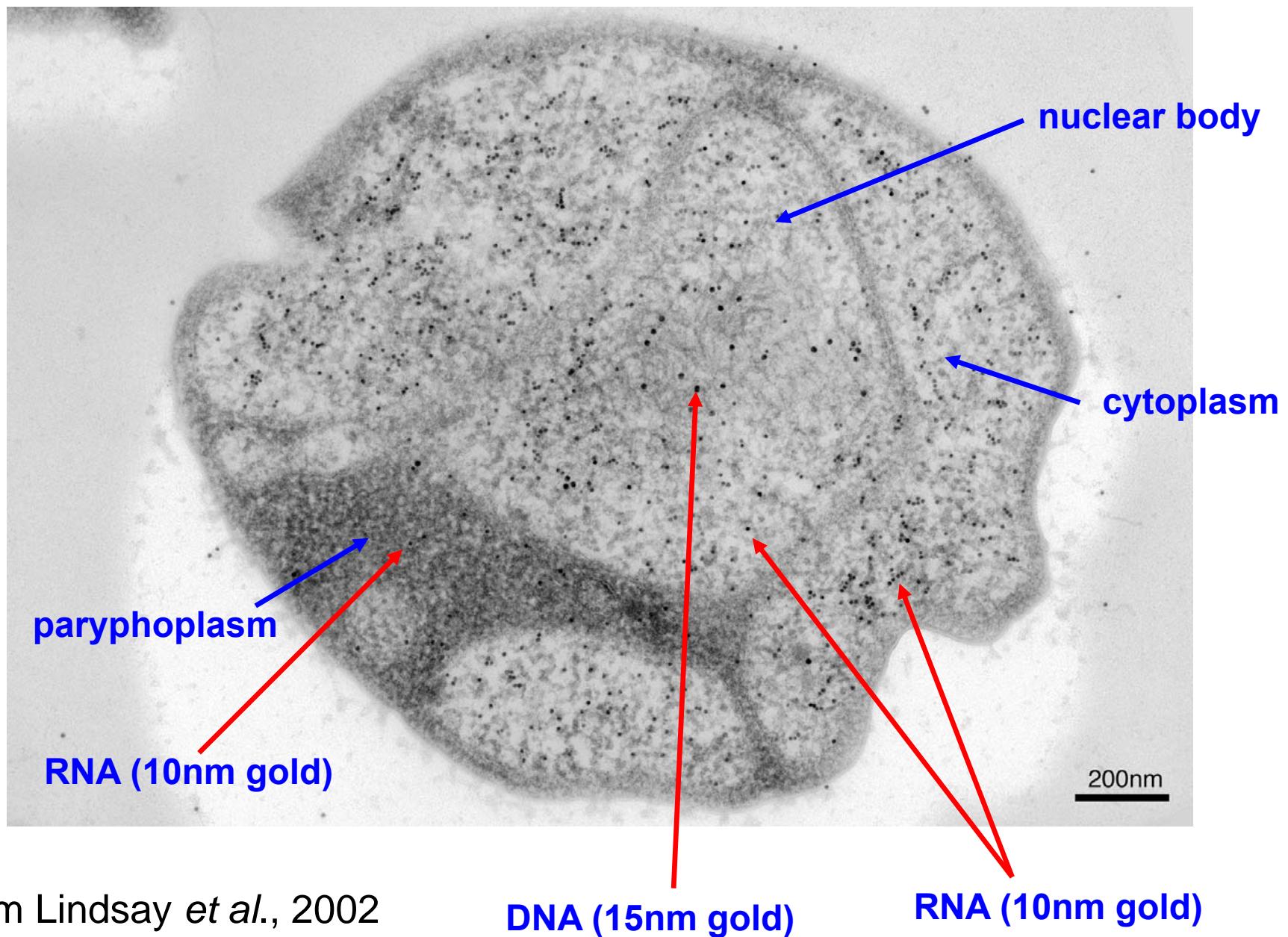
# Condensed chromatin correlates with radiation resistance in *Gemmata*



Arnon Lieber, Andrew Leis, Ariel Kushmaro, Abraham Minsky, and Ohad Medalia

**Chromatin organization and radio-resistance in the bacterium *Gemmata obscuriglobus*** J. Bacteriol.  
published ahead of print on 12 December 2008, doi:10.1128/JB.01513-08

**Location of DNA and RNA within *Gemmata obscuriglobus*  
(via anti-DNA antibody immunogold and RNase-gold localization)**

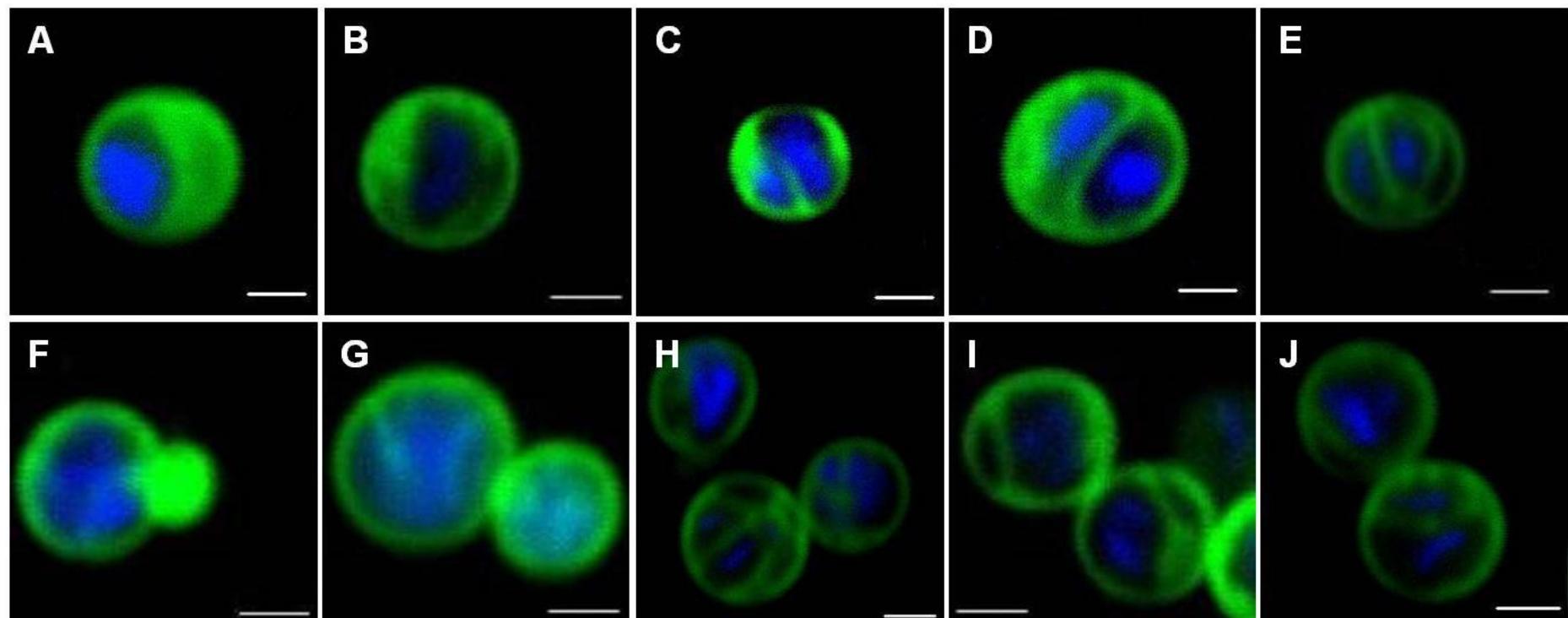


# **Cell division in *Gemmata obscuriglobus***

- No FtsZ involved
- of divisome proteins found in other Bacteria only FtsK appears (and this could operate in chromosome segregation independently)
- Budding involving asymmetric formation of daughter bud and eventual separation of mother and daughter – can be repeated
- Nuclear body and envelope reforms in bud, but nucleoid may be transferred through bud neck
- New envelope formed from ICMs of mother and daughter (if ICM= ER membrane then similar to eukaryote nuclear envelope formation)

## *Gemmata* cell cycle stages – Fate of nuclear DNA relative to nuclear envelope during division

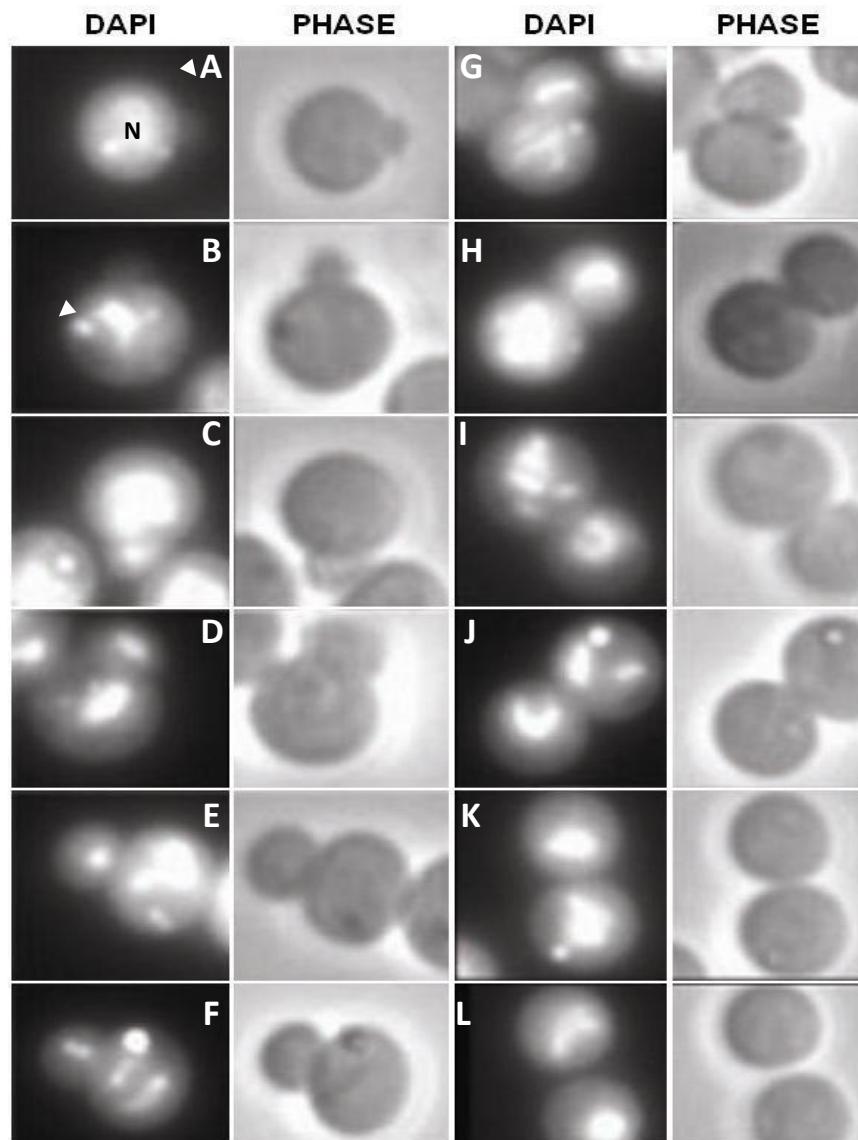
Lee, K-C , Webb, RI and Fuerst, JA, *BMC Cell Biology* January 2009



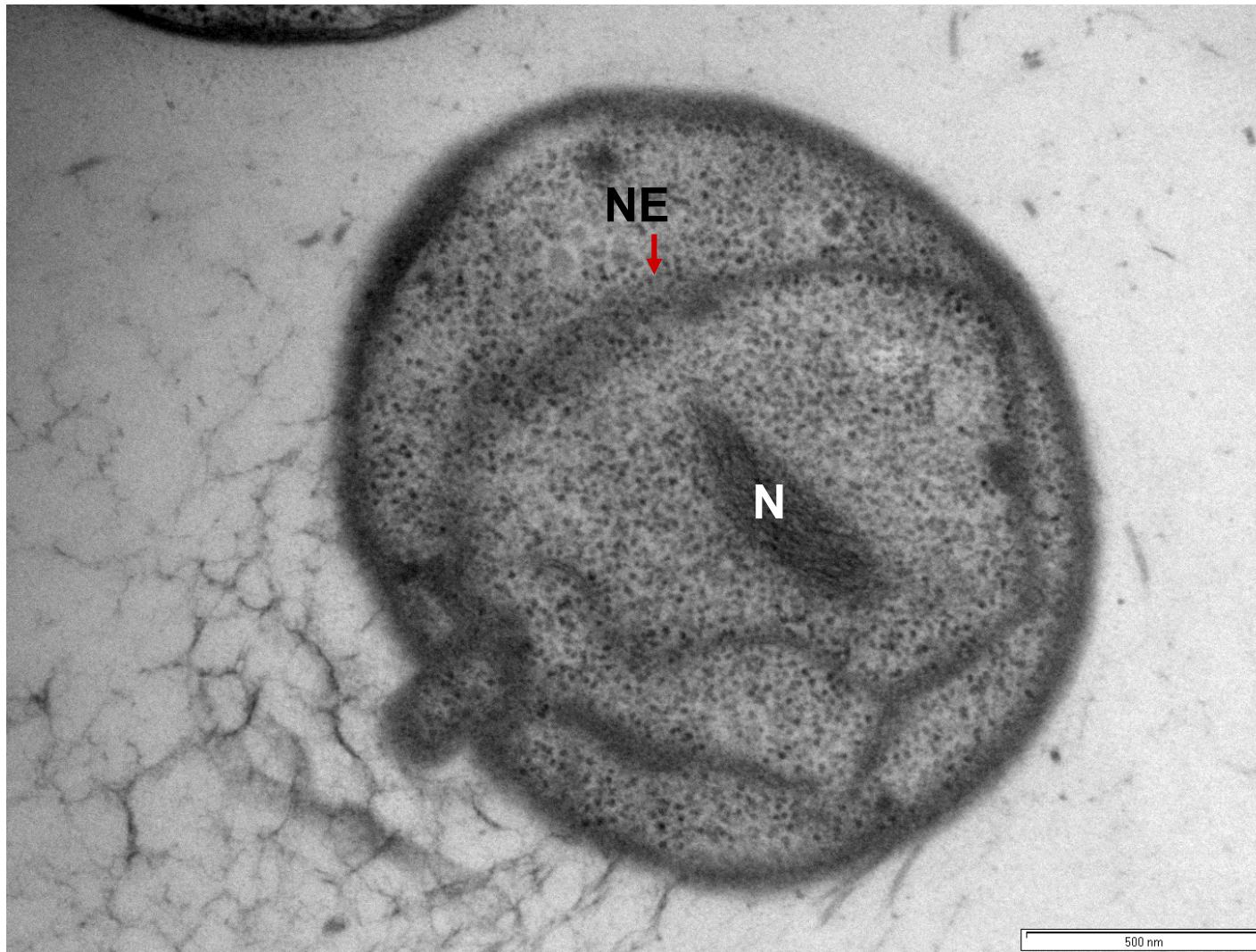
Confocal laser scanning micrographs Blue – DAPI, Green – DiOC<sub>6</sub>(3), Bar - 1 μm

## Fate of the nucleoid DNA during budding

- Formation of bud and translocation of nucleoid **do not occur** at the same time
- Bud initiates **without** nucleoid
- Nucleoid appears in bud later, before bud matures to full size

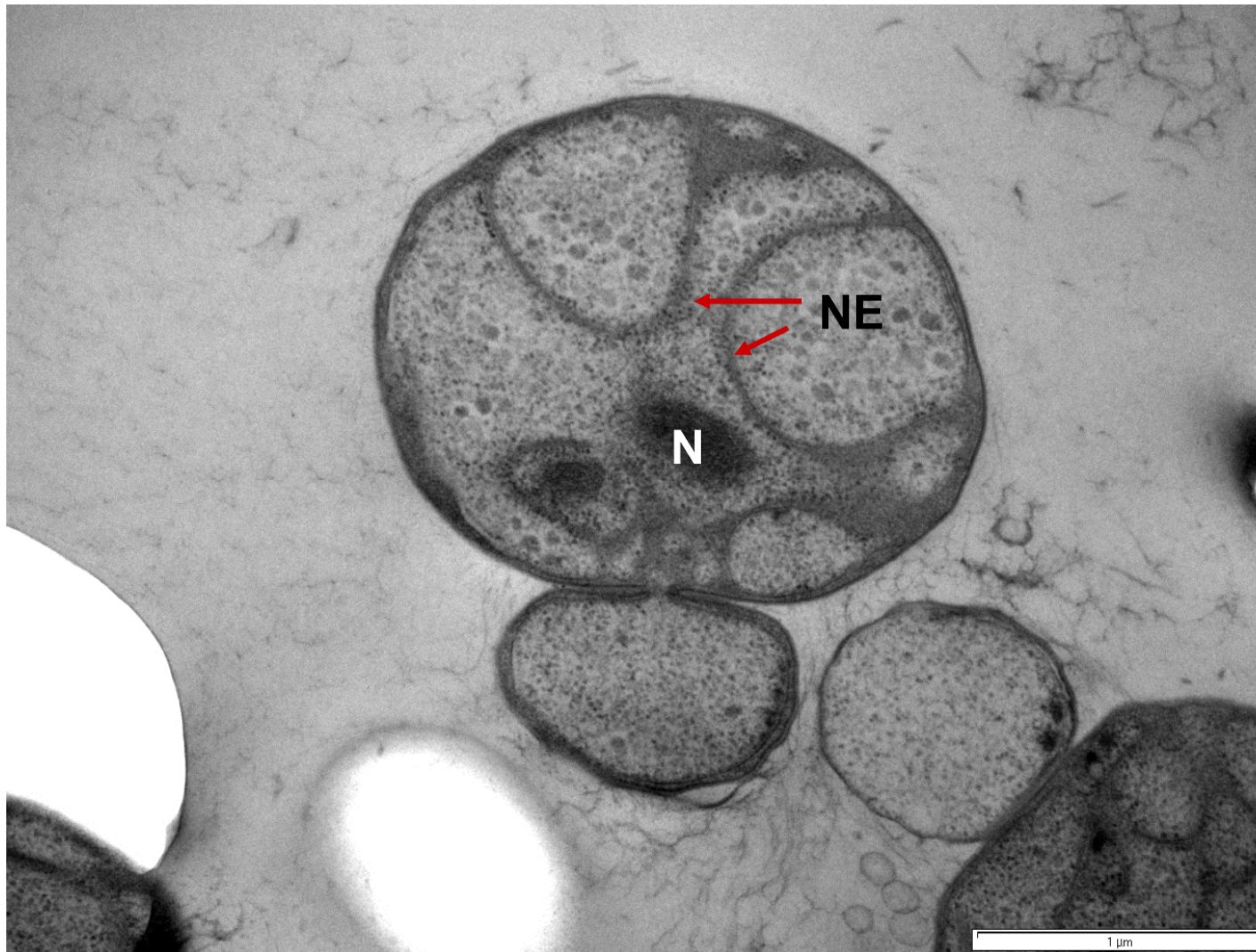


# *G. obscuriglobus* budding cell - 1



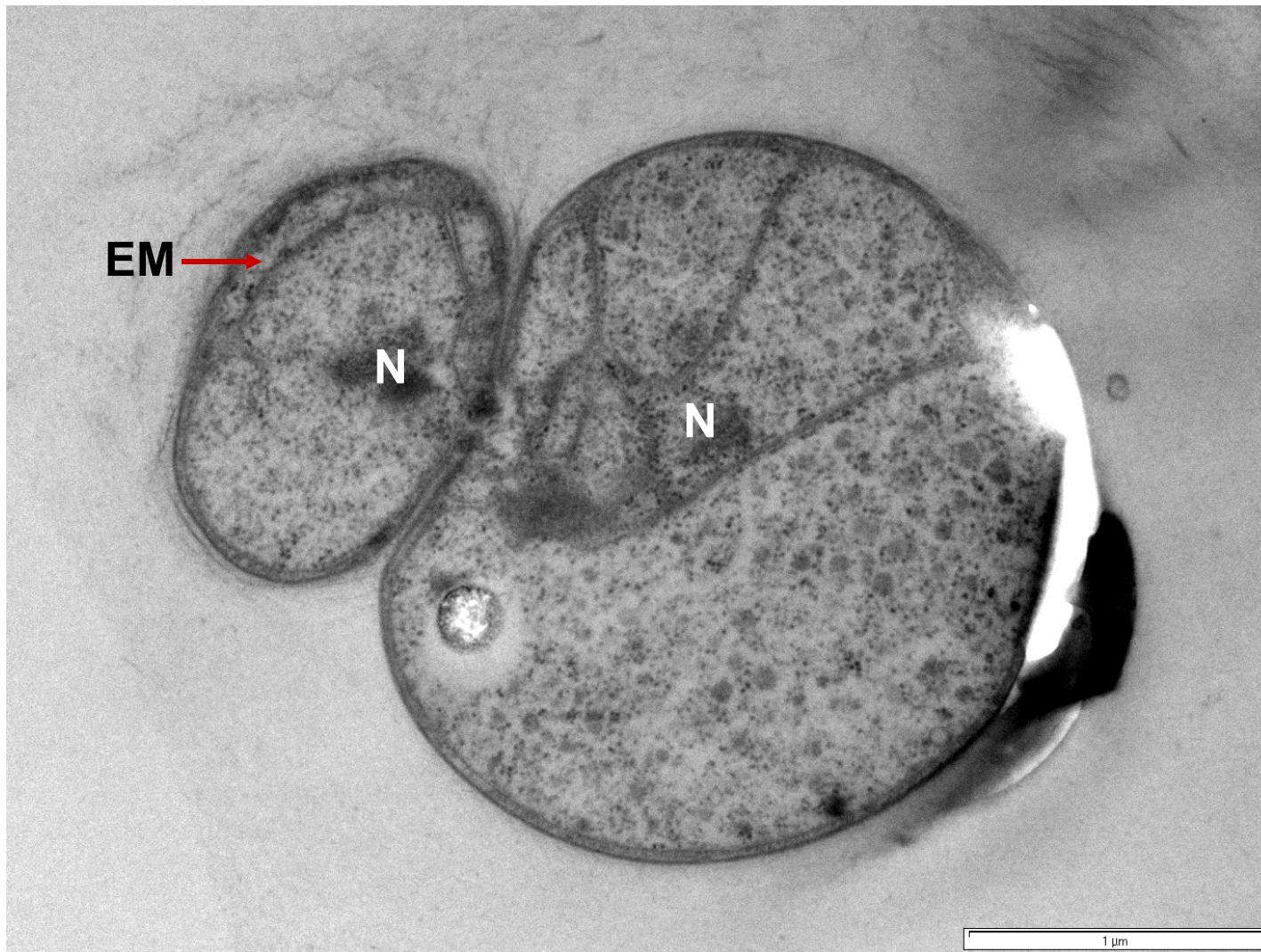
TEM, Thin section of high pressure-frozen cells grown on M1 agar

# *G. obscuriglobus* budding cell - 2



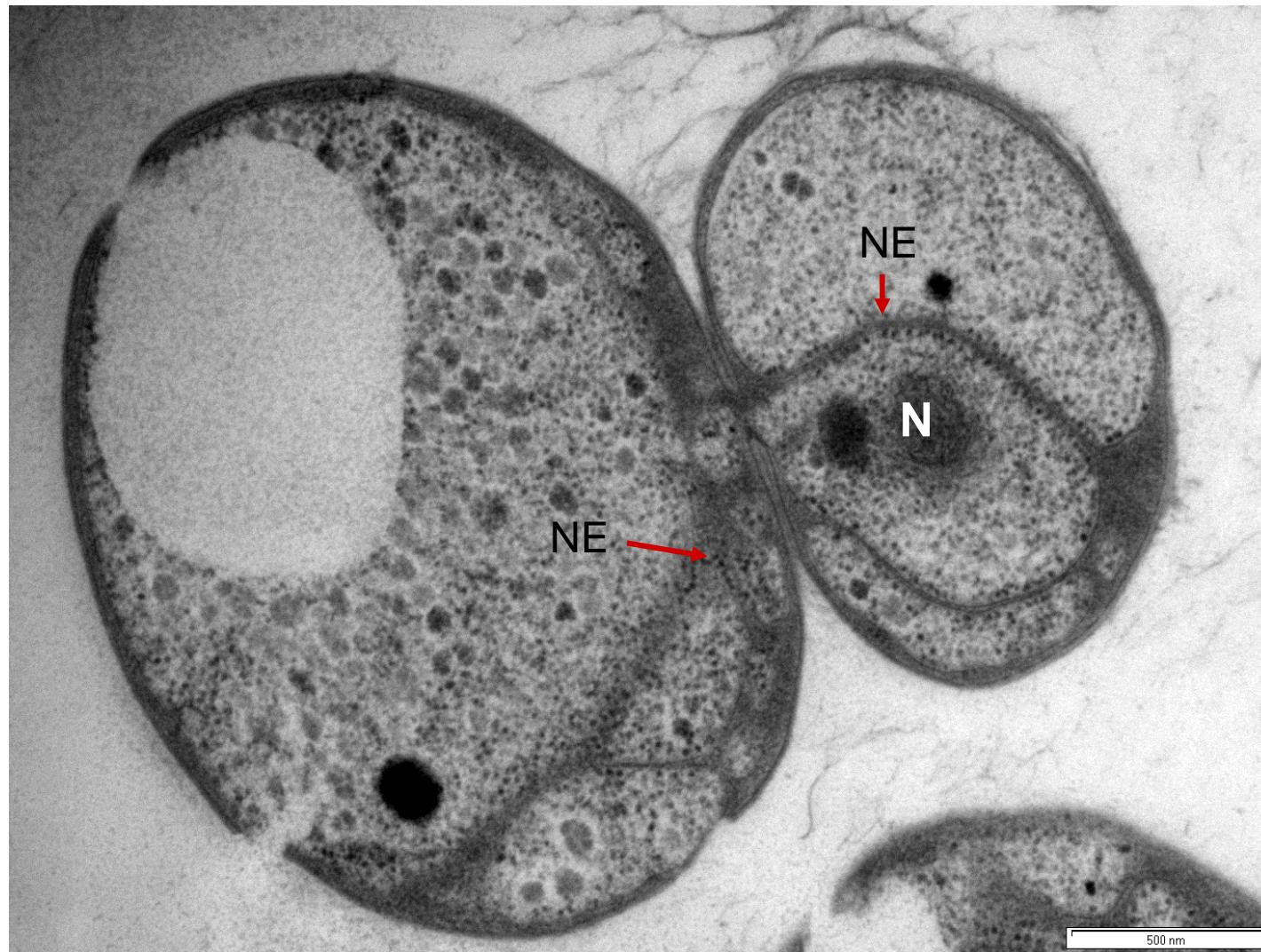
TEM, Thin section of high pressure-frozen cells grown on M1 agar

# *G. obscuriglobus* budding cell - 3



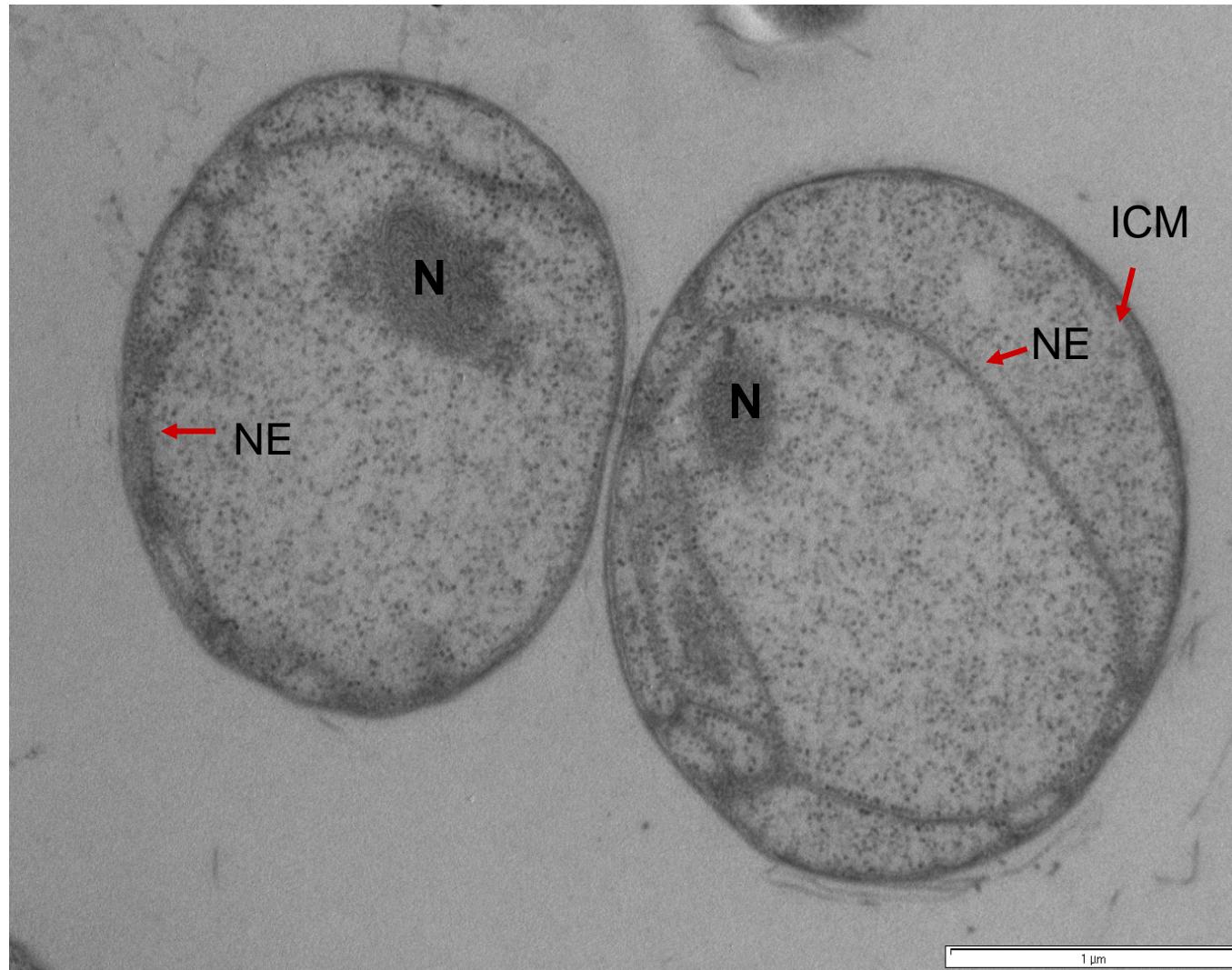
TEM, Thin section of high pressure-frozen cells grown on M1 agar

# *G. obscuriglobus* budding cell - 4



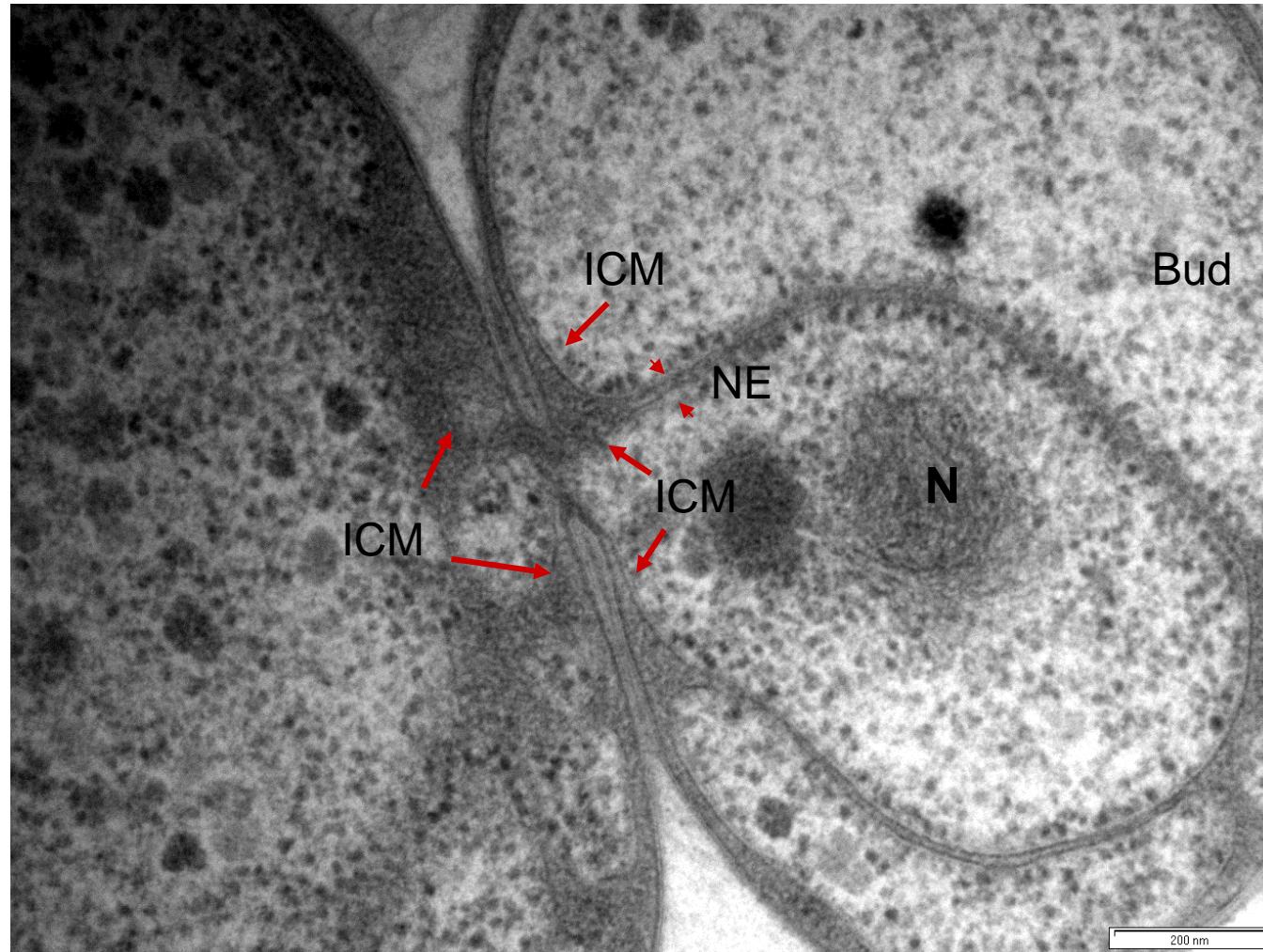
TEM, Thin section of high pressure frozen cells grown on M1 agar

# *G. obscuriglobus* budding cell - 5



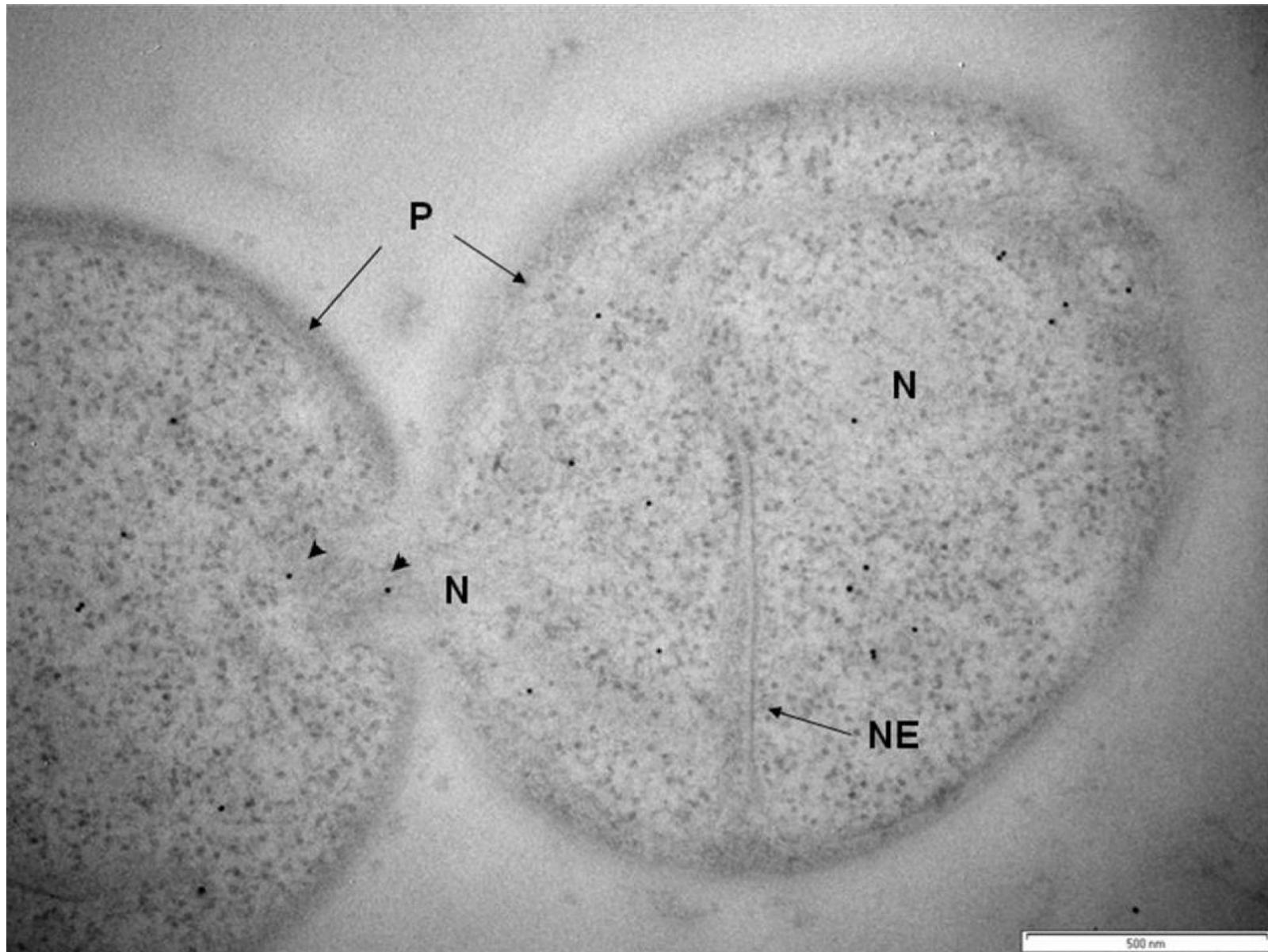
TEM, Thin section of high pressure frozen cells grown on M1 agar

**ICM membranes of mother cell and the bud form the 2-membraned nuclear envelope in the bud**



TEM, Thin section of high pressure frozen cells grown on M1 agar

*A Gemmata* mother cell nucleoid associated with FtsK passes through bud neck

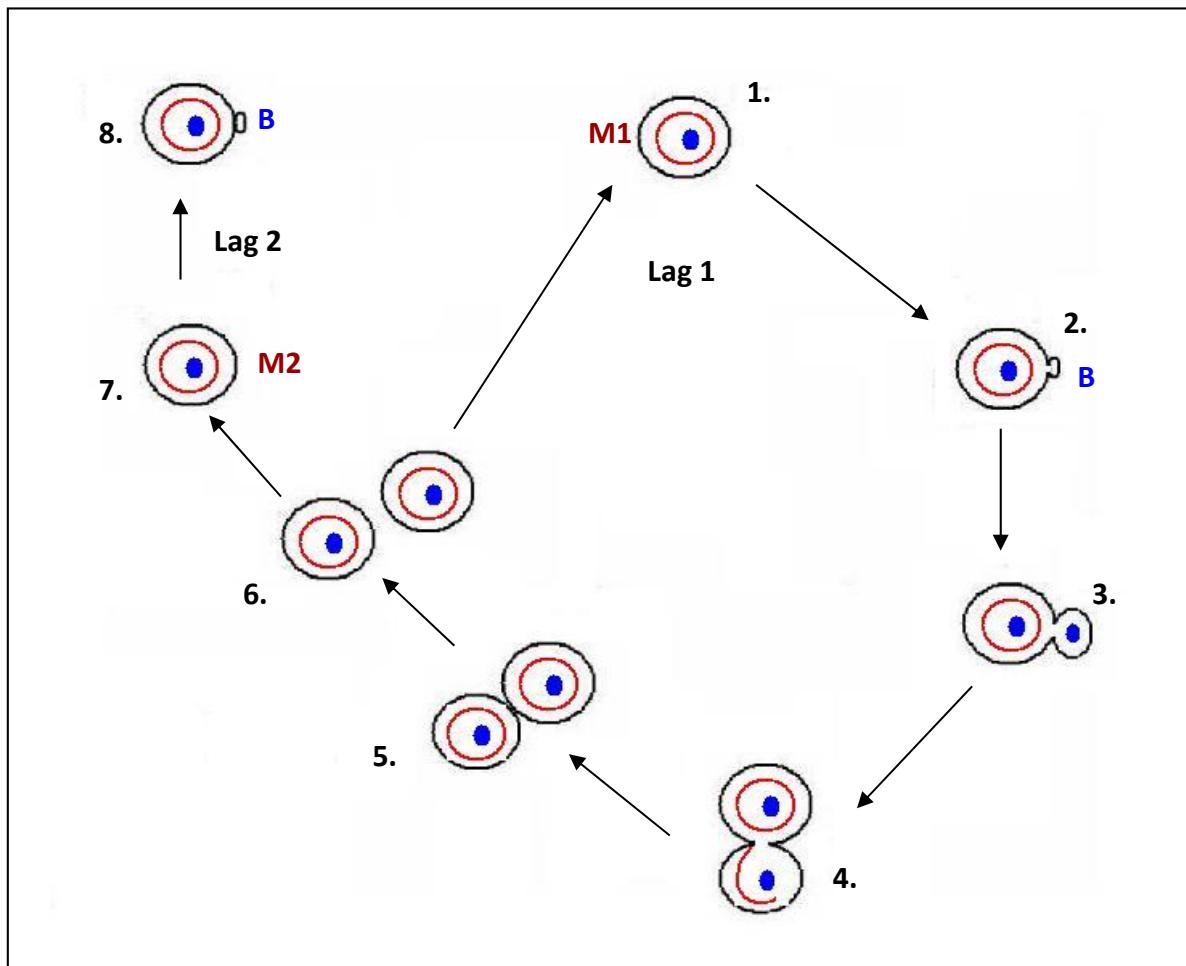


Jeffery Lee PhD thesis, Fuerst lab

# Nuclear segregation during division in *Gemmata obscuriglobus*

- ✿ Bud initiates **without** nucleoid, then nucleoid appears in bud before bud matures to full size
- ✿ Naked nucleoid appears in bud **before** its **envelopment in nuclear membranes**
- ✿ **Nuclear envelope** of bud derives from **intracytoplasmic membrane**, but from different cells - **inner membrane** is from ICM of **mother cell**, **outer membrane** from ICM of **bud**
- ✿ **Ribosomes** associated with the new nuclear envelope

# *Gemmata obscuriglobus* cell budding cycle



**M1** – mother cell giving rise to the first bud

**M2** – mother cell derived from the first bud

**B** – bud

**Blue region** – nucleoid

**Red circle** - nuclear envelope

**Lag 1** – 2~4 hours

**Lag 2** – 3~5.5 hours

'Eukaryality' implies nuclei AND endomembranes forming vesicles with the help of coatomer proteins

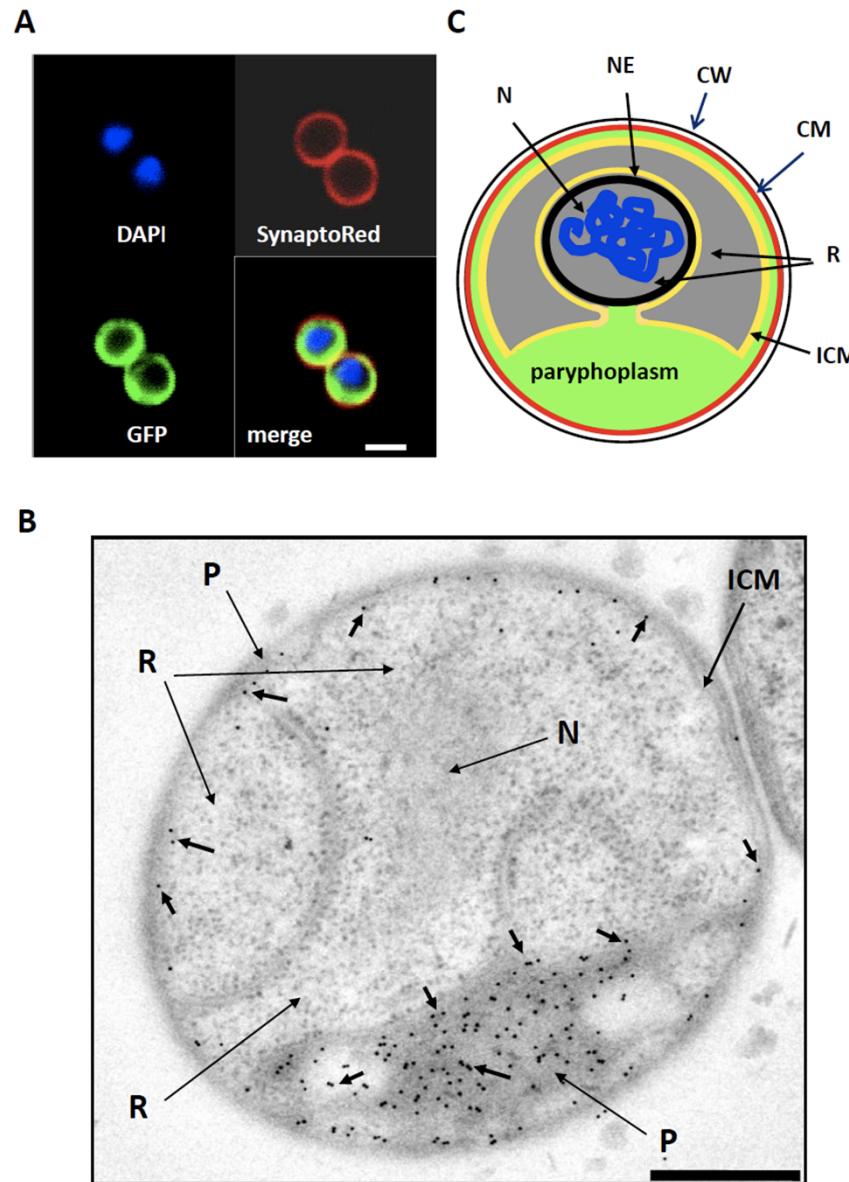
- *Gemmata* should be ideal system for - Experimental test for endocytosis and membrane trafficking in a bacterium!

Evidence for endocytosis in a  
bacterium - a functional  
'eukaryotic' process in the  
planctomycete *Gemmata*  
*obscuriglobus*

# Endocytosis-like processes in planctomycetes

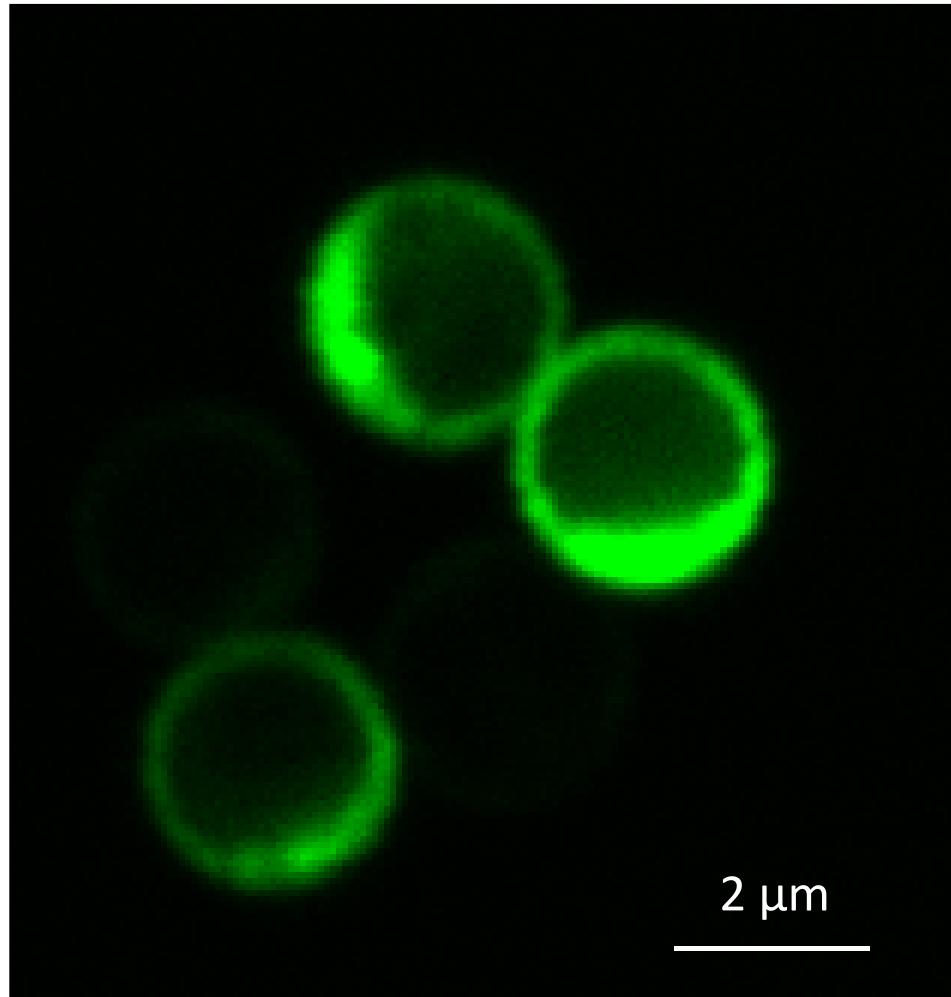
- Protein uptake – receptor-mediated, energy-dependent & vesicle-associated
- Polysaccharide uptake – vesicle-associated but probably via fluid-phase pinocytosis, energy-dependent

# Protein uptake by *Gemmata*

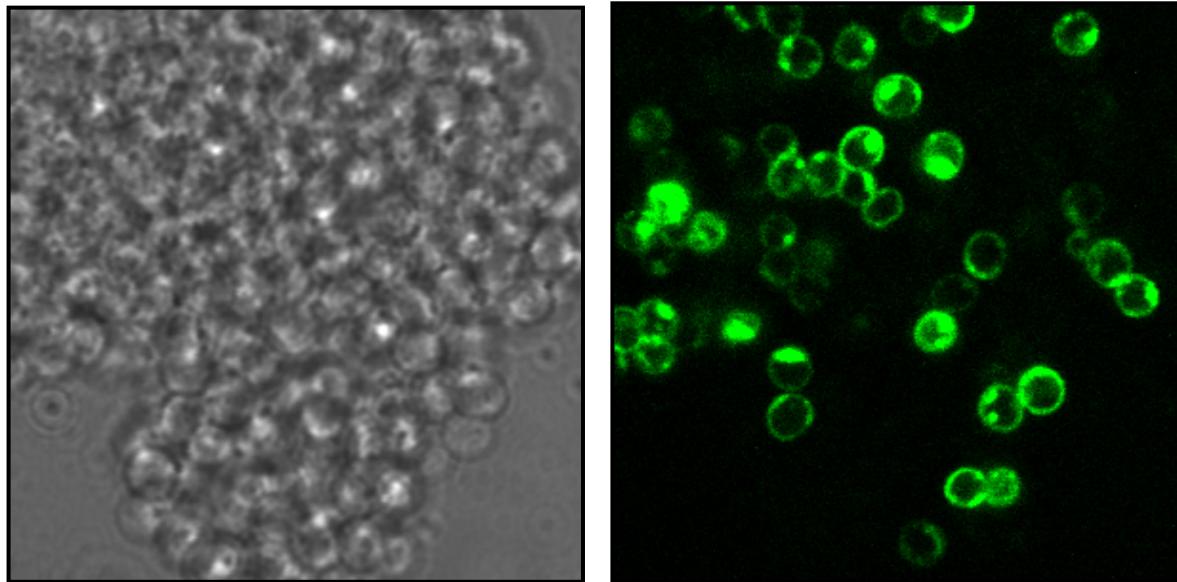


From: Lonhienne TG, Sagulenko E, Webb RI, Lee KC, Franke J, Devos DP, Nouwens A, Carroll BJ, Fuerst JA. Endocytosis-like protein uptake in the bacterium *Gemmata obscuriglobus*. Proc Natl Acad Sci U S A. 2010 Jul 20; 107(29):12883-8.

Uptake of GFP by cells of *Gemmata obscuriglobus*  
seen via CLSM

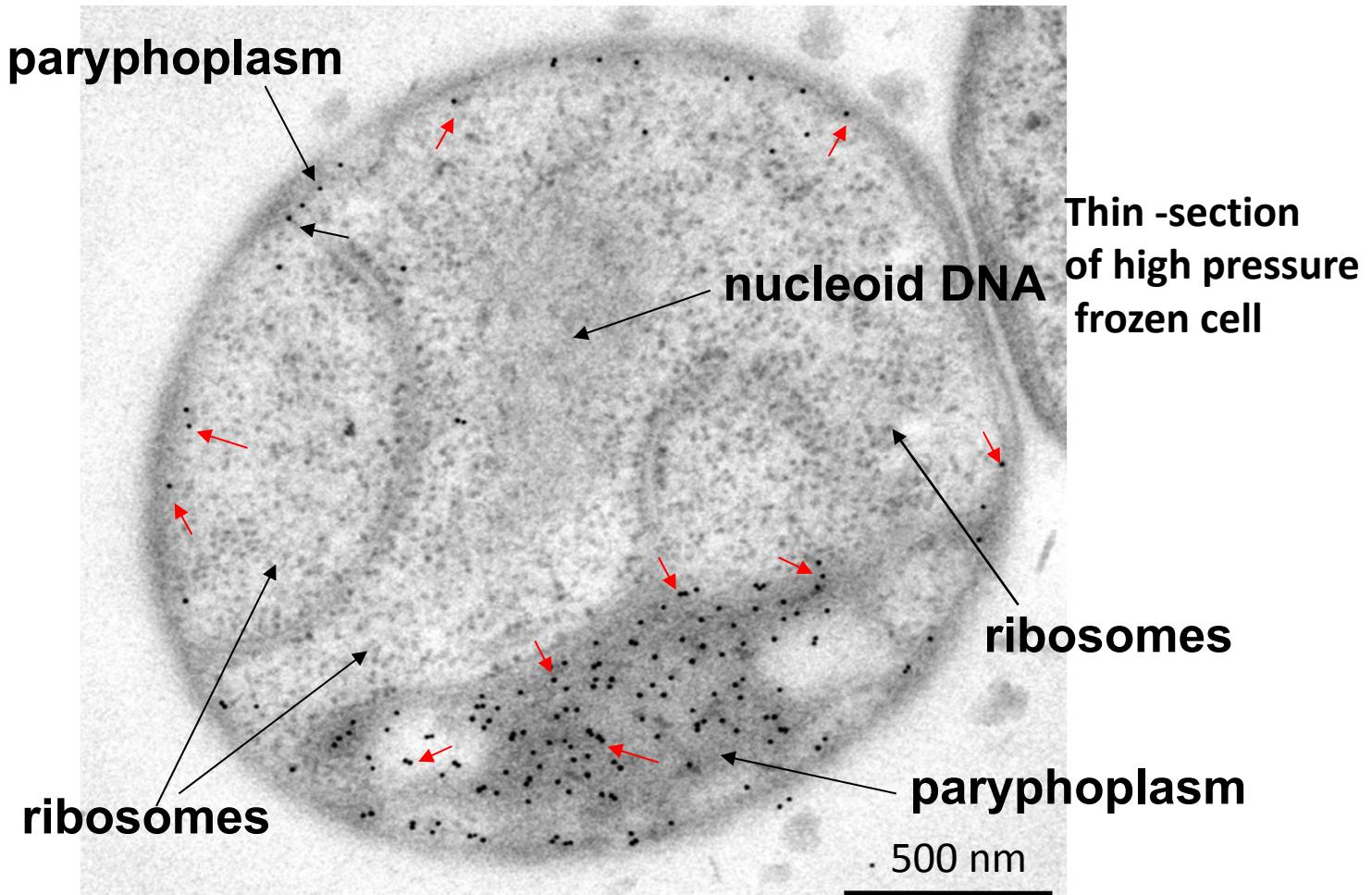


# GFP is taken up by Gemmata cells



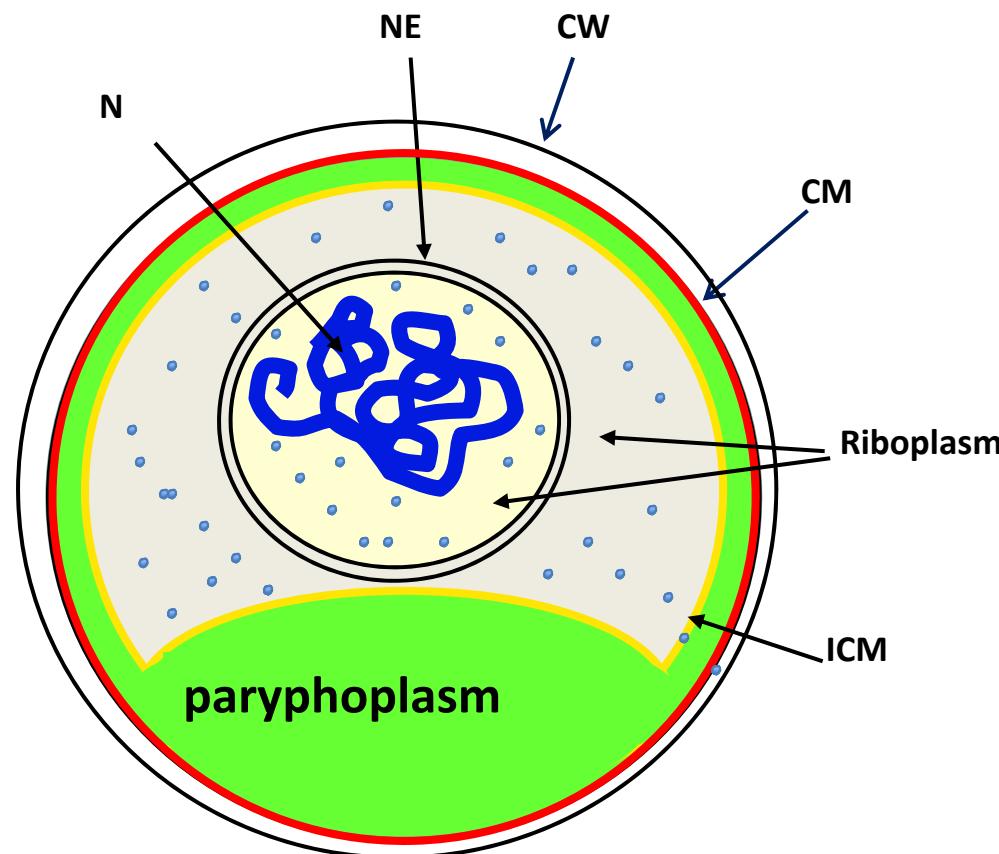
From: Lonhienne TG, Sagulenko E, Webb RI, Lee KC, Franke J, Devos DP, Nouwens A, Carroll BJ, Fuerst JA. Endocytosis-like protein uptake in the bacterium *Gemmata obscuriglobus*. Proc Natl Acad Sci U S A. 2010 Jul 20; 107(29):12883-8.

# Immunogold labelling of GFP taken up by *Gemmata obscuriglobus* cells

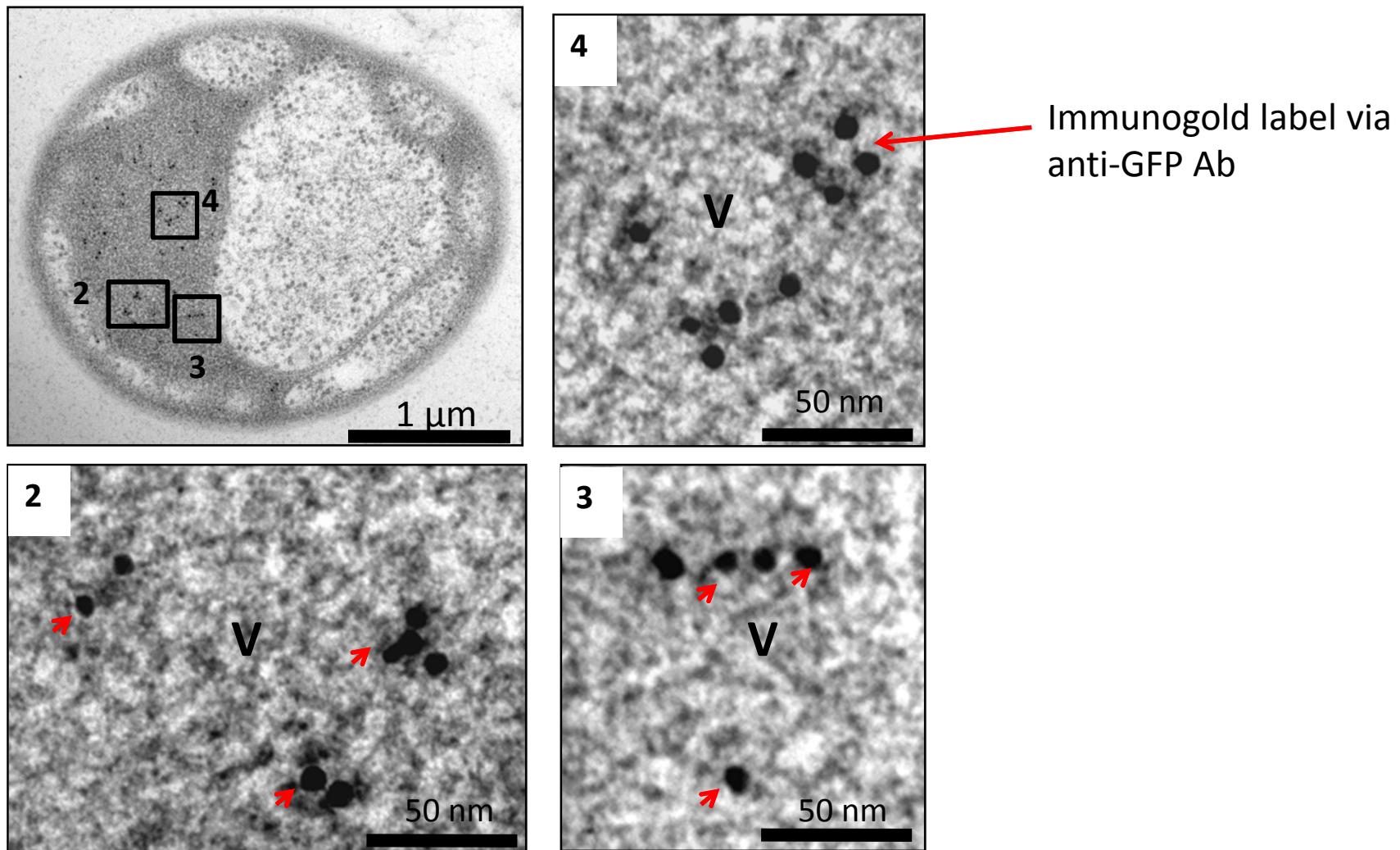


From: Lonhienne TG, Sagulenko E, Webb RI, Lee KC, Franke J, Devos DP, Nouwens A, Carroll BJ, Fuerst JA. Endocytosis-like protein uptake in the bacterium *Gemmata obscuriglobus*. Proc Natl Acad Sci U SA. 2010 Jul 20;107(29):12883-8.

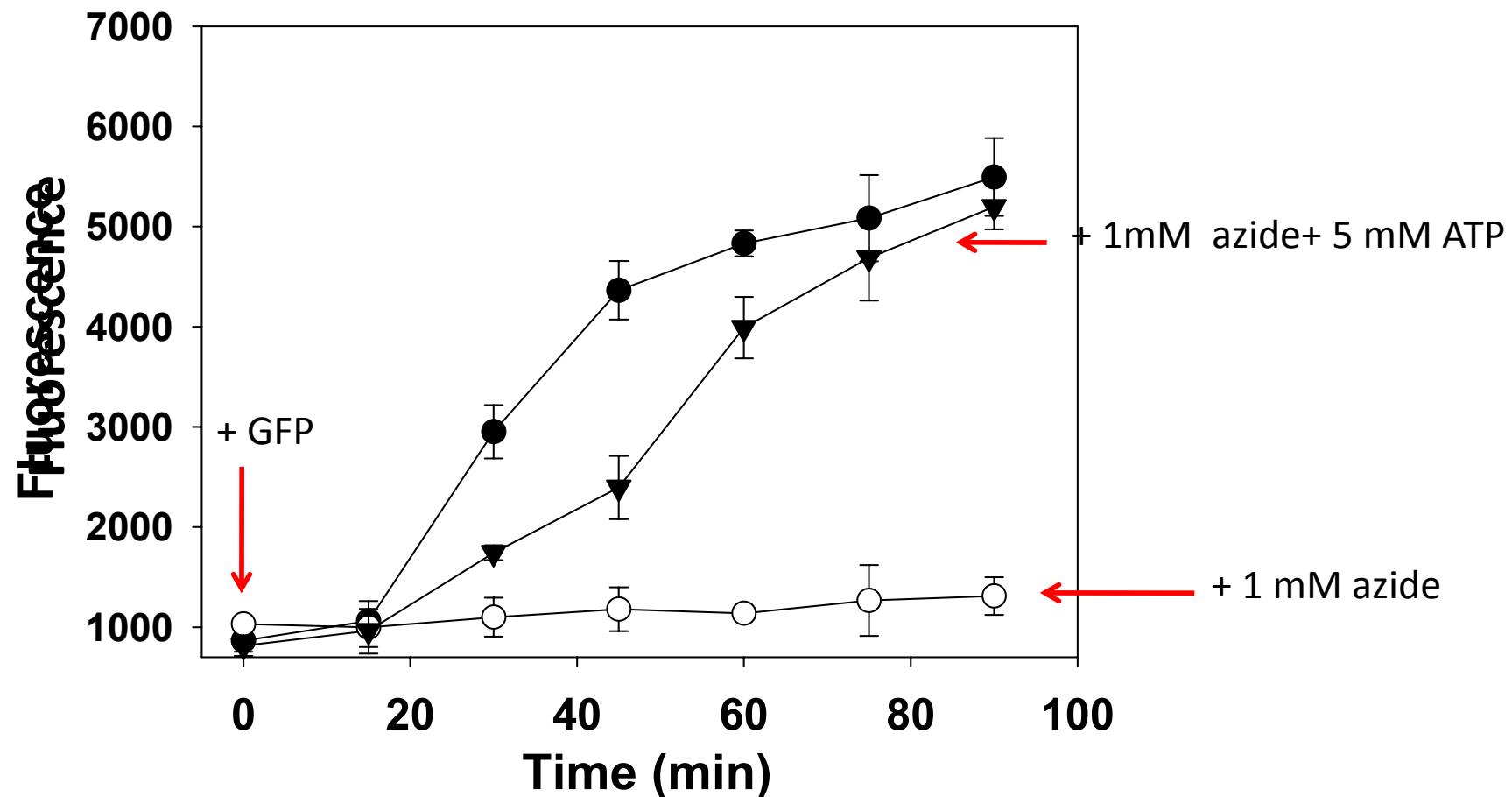
# *Gemmata obscuriglobus* cell plan



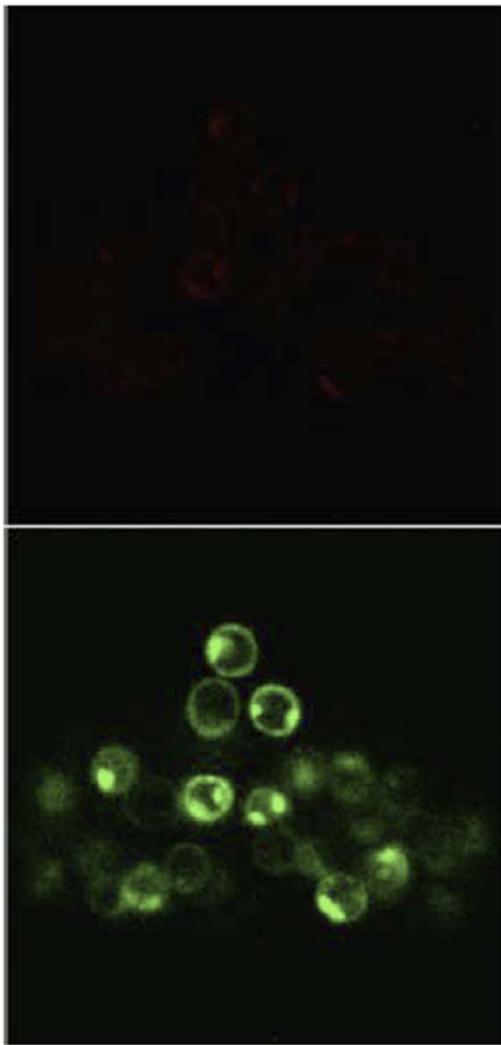
# GFP is incorporated into membrane-bounded vesicles within one compartment (the paryphoplasm)



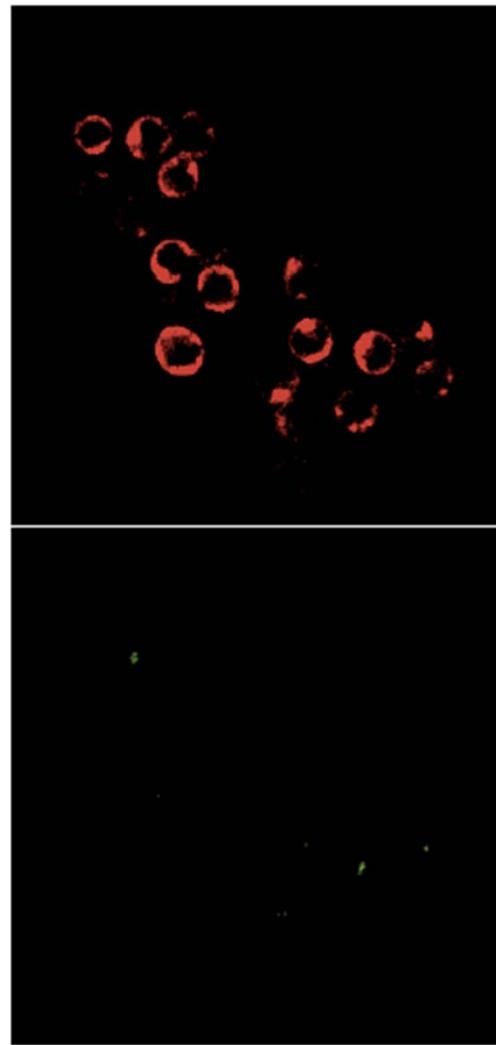
**Fluorescence plot for GFP incubation with *Gemmata obscuriglobus* : GFP uptake is an energy-dependent process.**



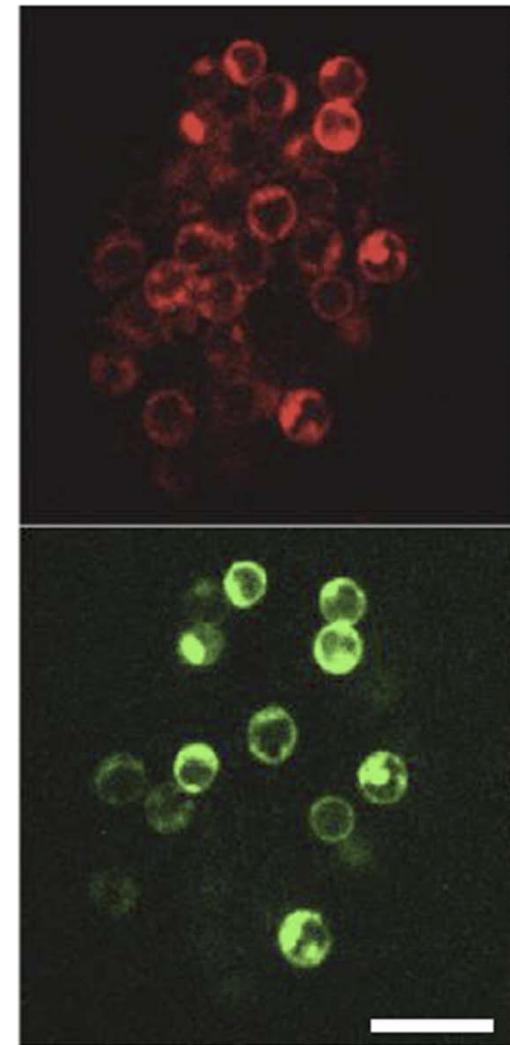
IgG-Cy3:GFP  
1:20



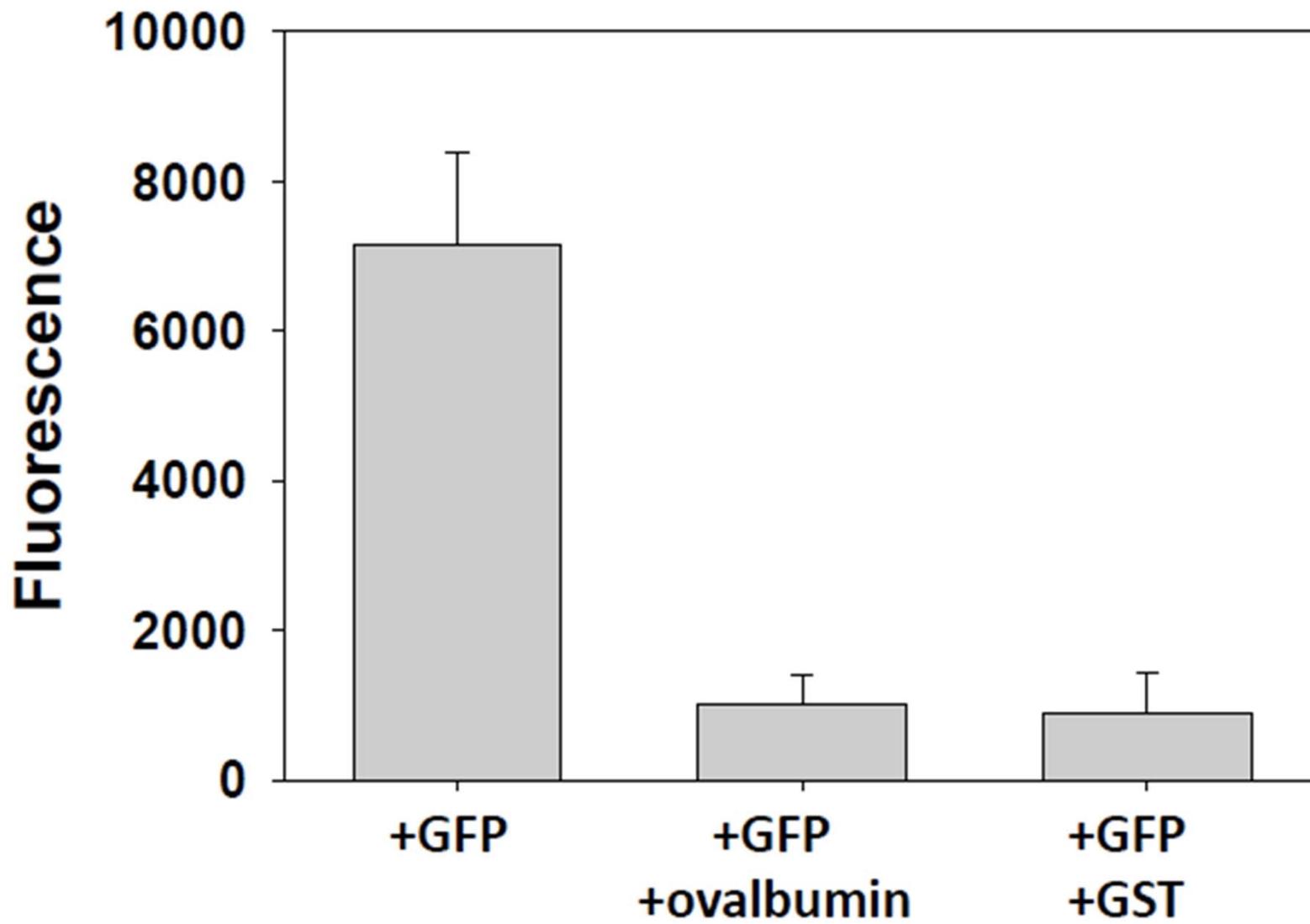
IgG-Cy3:GFP  
20:1



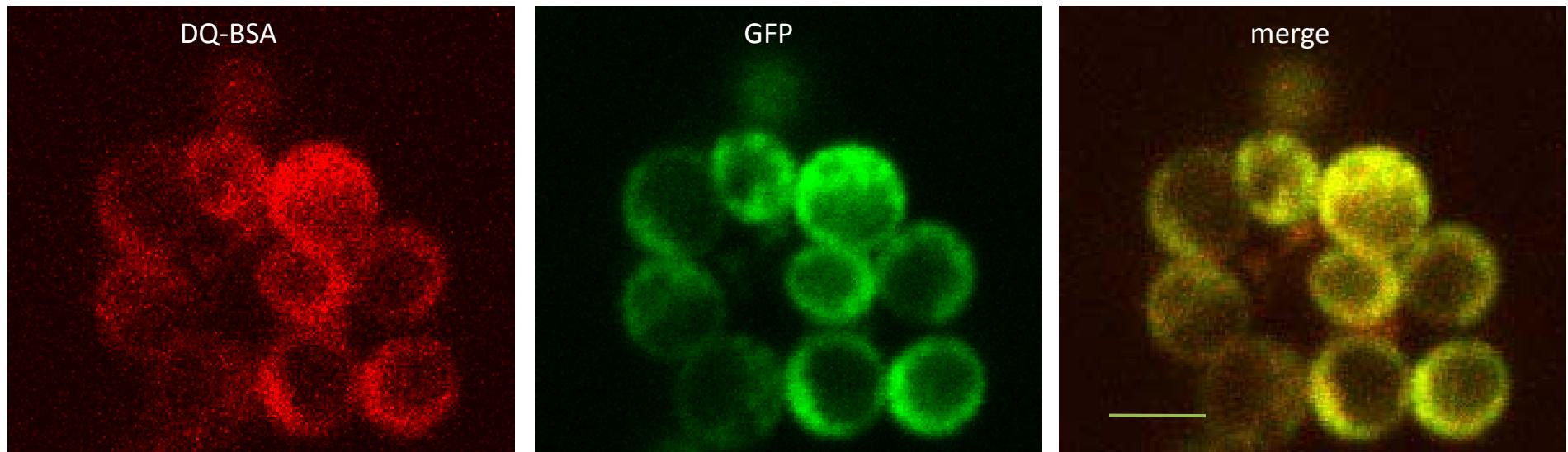
IgG-Cy3:GFP  
1:1



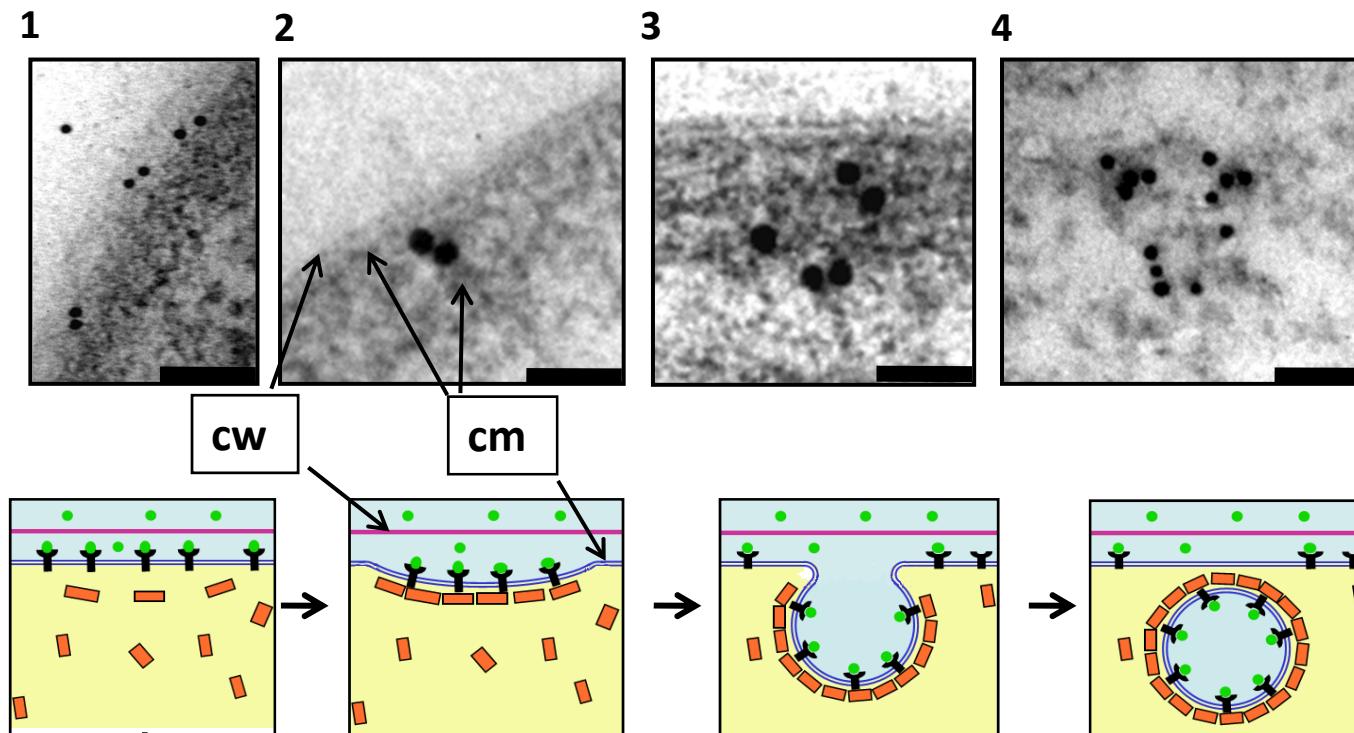
## Competition experiment



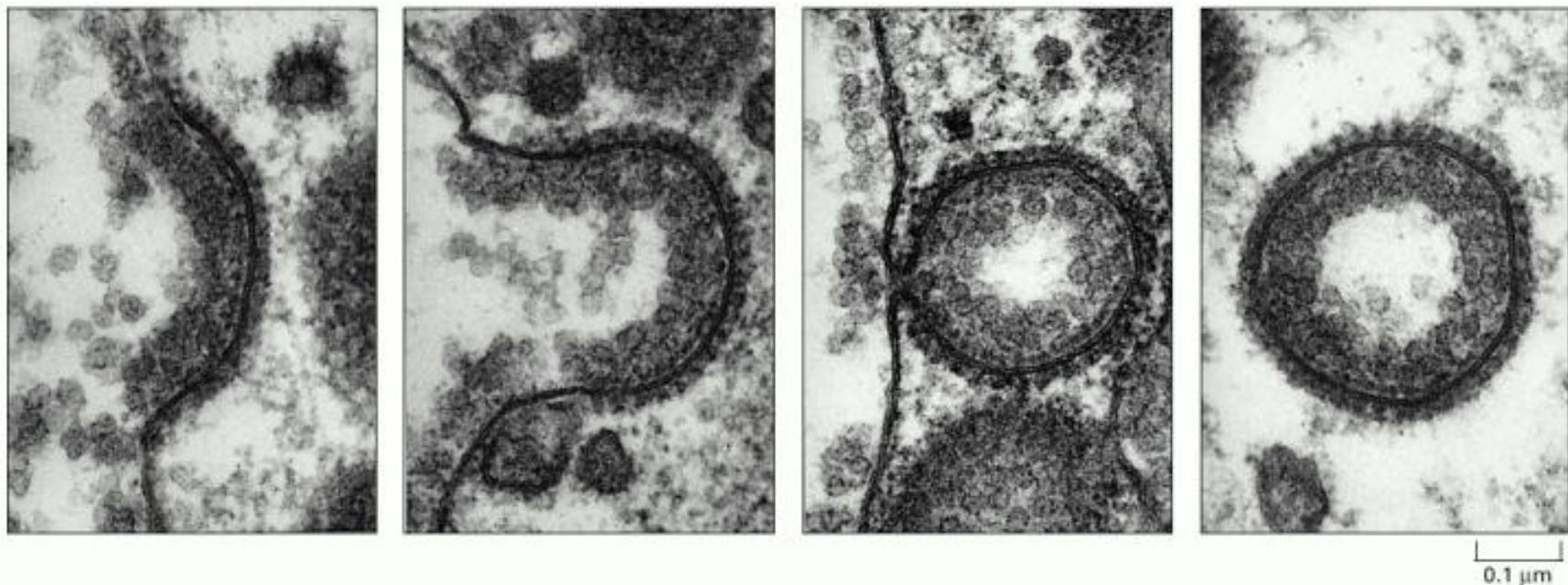
Protein DQ Green BSA (fluorescent only after degradation) is degraded after uptake into *Gemmata* cells (in same major compartment as one taking up GFP) - can there be a bacterial lysosome?



# Proposed mechanism of receptor-mediated endocytosis in *Gemmata obscuriglobus* cells



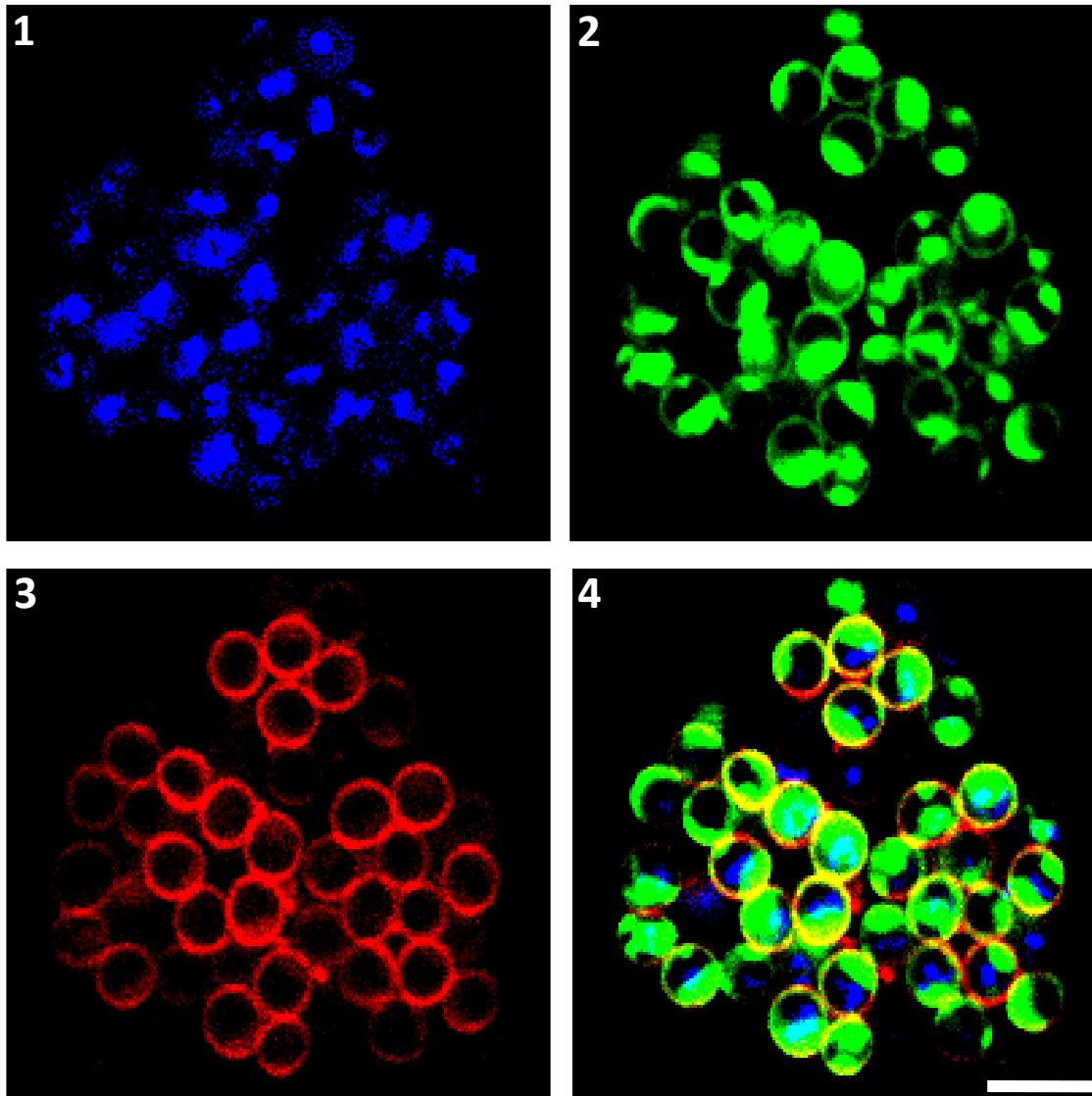
# Endocytosis in a eukaryote: TEM section showing endocytosis of lipoprotein by hen oocyte



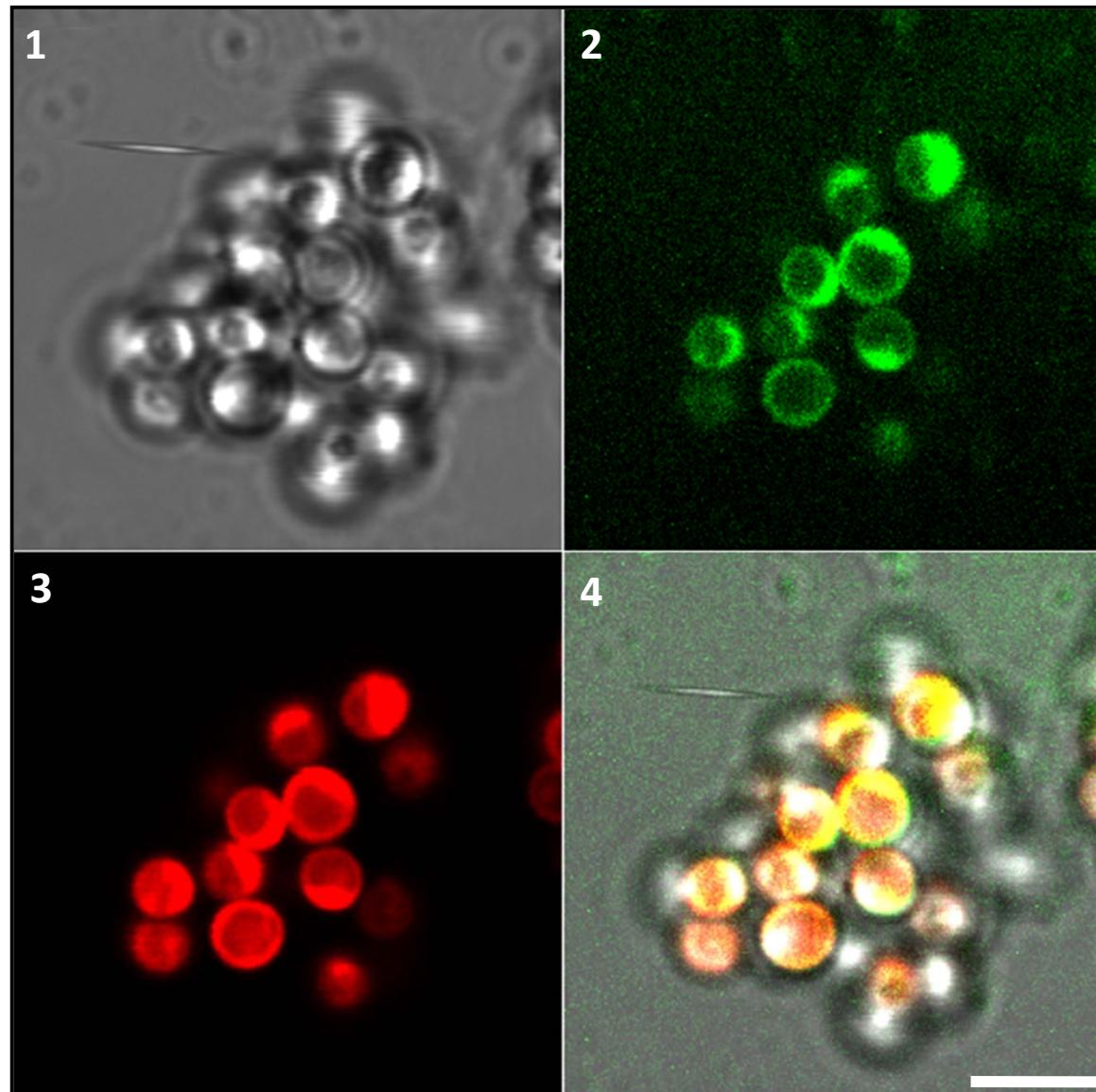
From:: Alberts, Molecular Biology of the Cell 2002 ed. Fig. 13-41  
modified figure from:

MM Perry and AB Gilbert 1979. Yolk transport in the ovarian follicle of the hen (*Gallus domesticus*): lipoprotein-like particles at the periphery of the oocyte in the rapid growth phase. J Cell Sci 39: 257-272

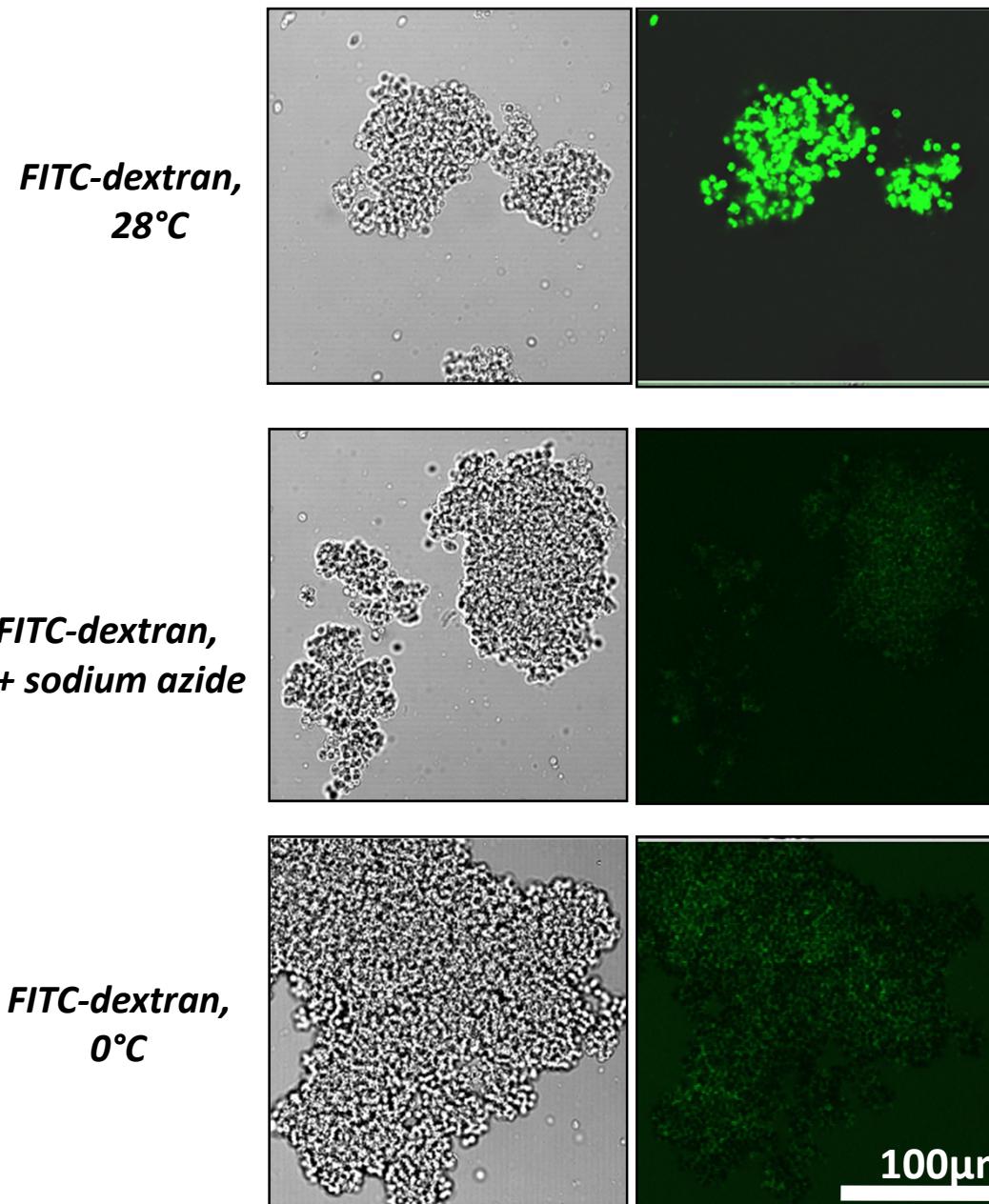
*Gemmata* the omnivore - uptake of FITC-dextran by *Gemmata obscuriglobus* cells



Rhodamine-dextran co-localizes with GFP in the paryphoplasm of *Gemmata obscuriglobus* cells



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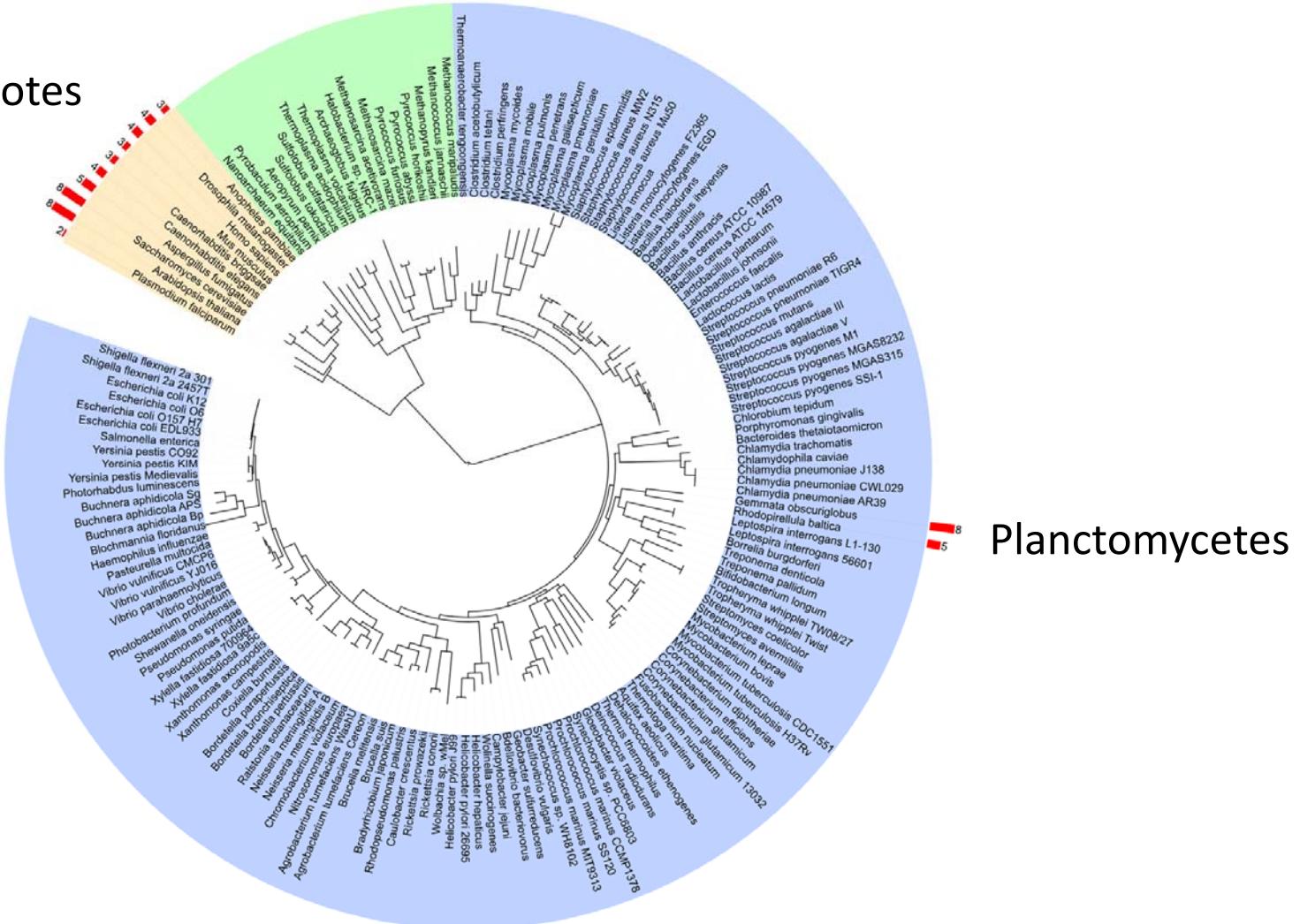


Sagulenko & Fuerst

# Protocoatomers & planctomycetes

- Within domain Bacteria, PVC superphylum members including planctomycetes share MC protein clathrin-like homologs
- Implies possibility of both clathrin-based endocytosis and MC-containing nuclear pores in PVC members, e.g. *Gemmata*

# Eukaryotes



Santarella-Mellwig R et al. PLoS Biol. 2010 Jan 19;8(1):e1000281.

# MC proteins in PVC members

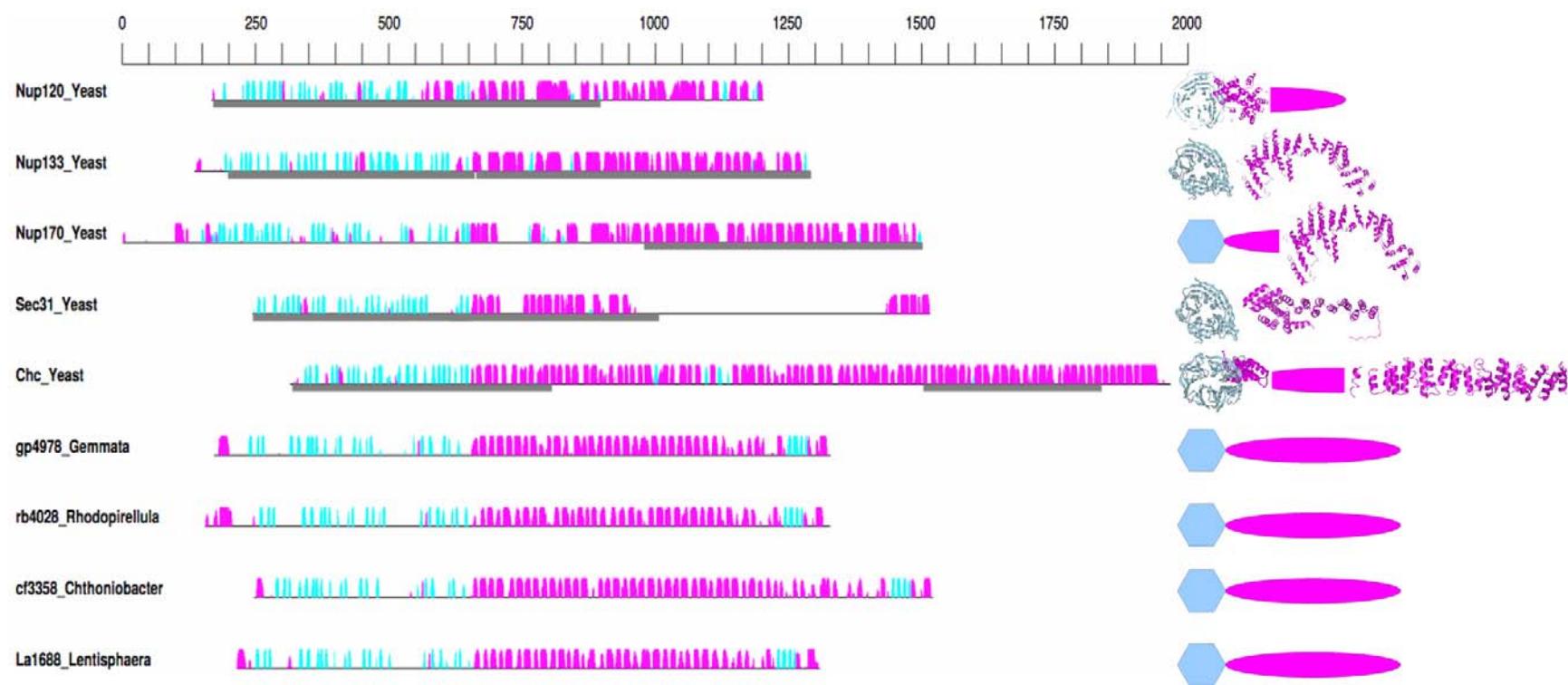
Species	Phylum	Genome Status	Total Number of Proteins	Number of MCs
<i>Chlamydophila felis</i> Fe/C-56	C	F	1,013	0
<i>Candidatus Protochlamydia amoebophila</i> UWE25	C	F	2,031	0
<i>Chlamydia muridarum</i> Nigg	C	F	911	0
<i>Victivallis vadensis</i> BAA-548	L	D	3,541	0
<i>Lentisphaera araneosa</i> HTCC2155	L	D	5,104	9
<i>Candidatus Kuenenia stuttgartiensis</i>	P	F	4,663	0
<i>Blastopirellula marina</i> DSM 3645	P	D	6,025	11
<i>Planctomyces maris</i> DSM 8797	P	D	6,480	11
<i>Rhodopirellula baltica</i> SH 1	P	F	7,325	5
<i>Gemmata obscuriglobus</i> UQM 2246	P	D	7,989	8
<i>Akkermansia muciniphila</i> BAA-835	V	F	2,176	0
<i>Methylacidiphilum infernorum</i> V4	V	F	2,462	0
<i>Opitutaceae bacterium</i> TAV2	V	D	4,036	0
<i>Opitutus terrae</i> PB90-1	V	F	4,632	0
<i>Pedosphaera parvula</i> Ellin514	V	D	6,402	9
<i>Verrucomicrobium spinosum</i>	V	D	6,509	16
<i>Chthoniobacter flavus</i> Ellin428	V	D	6,716	14

V, Verrucomicrobia; L, Lentisphaerae; P, Planctomycetes; C, Chlamydiae; D, draft; F, finished.

doi:10.1371/journal.pbio.1000281.t001

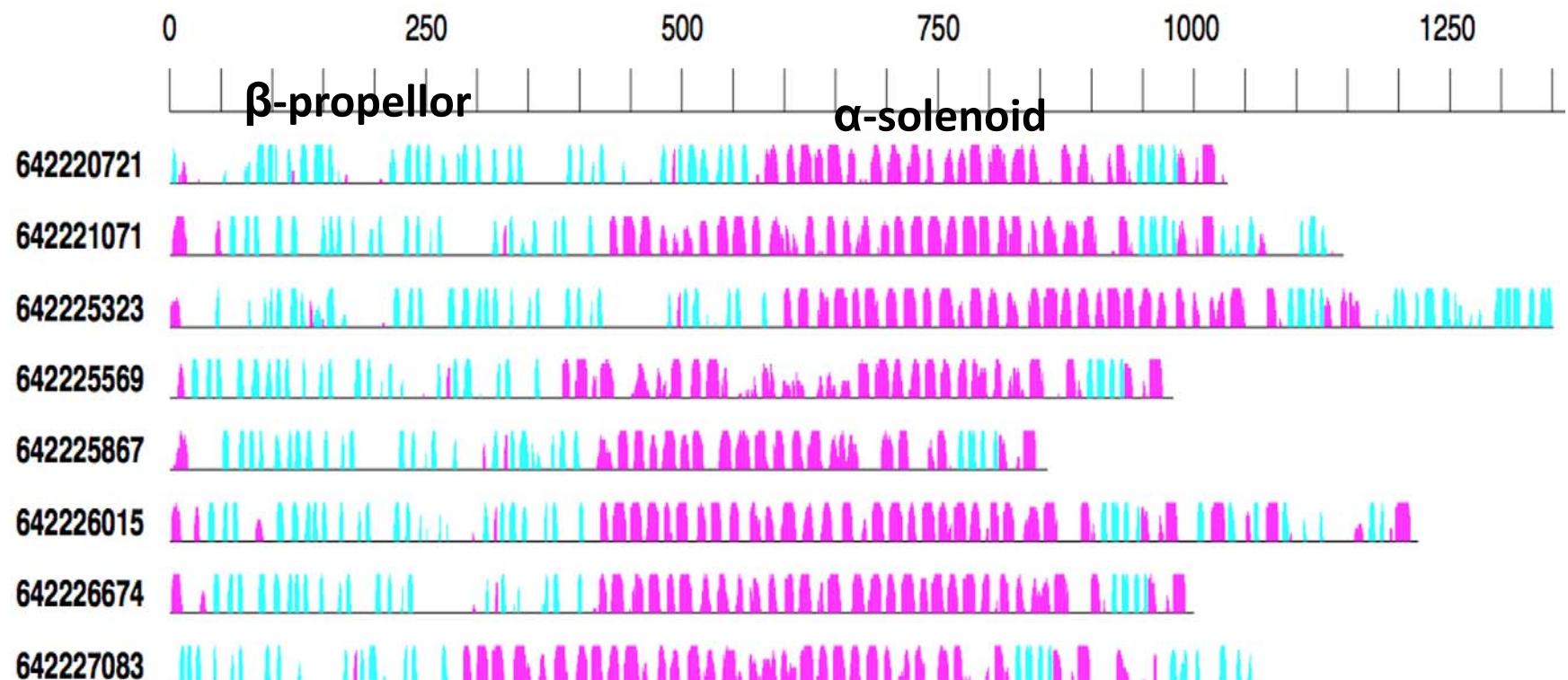
Santarella-Mellwig R et al. PLoS Biol. 2010 Jan 19;8(1):e1000281.

# MC proteins - Eukaryotes & PVC bacteria



Santarella-Mellwig R et al. PLoS Biol. 2010 Jan 19;8(1):e1000281

# Gemmata MC proteins & shared domains

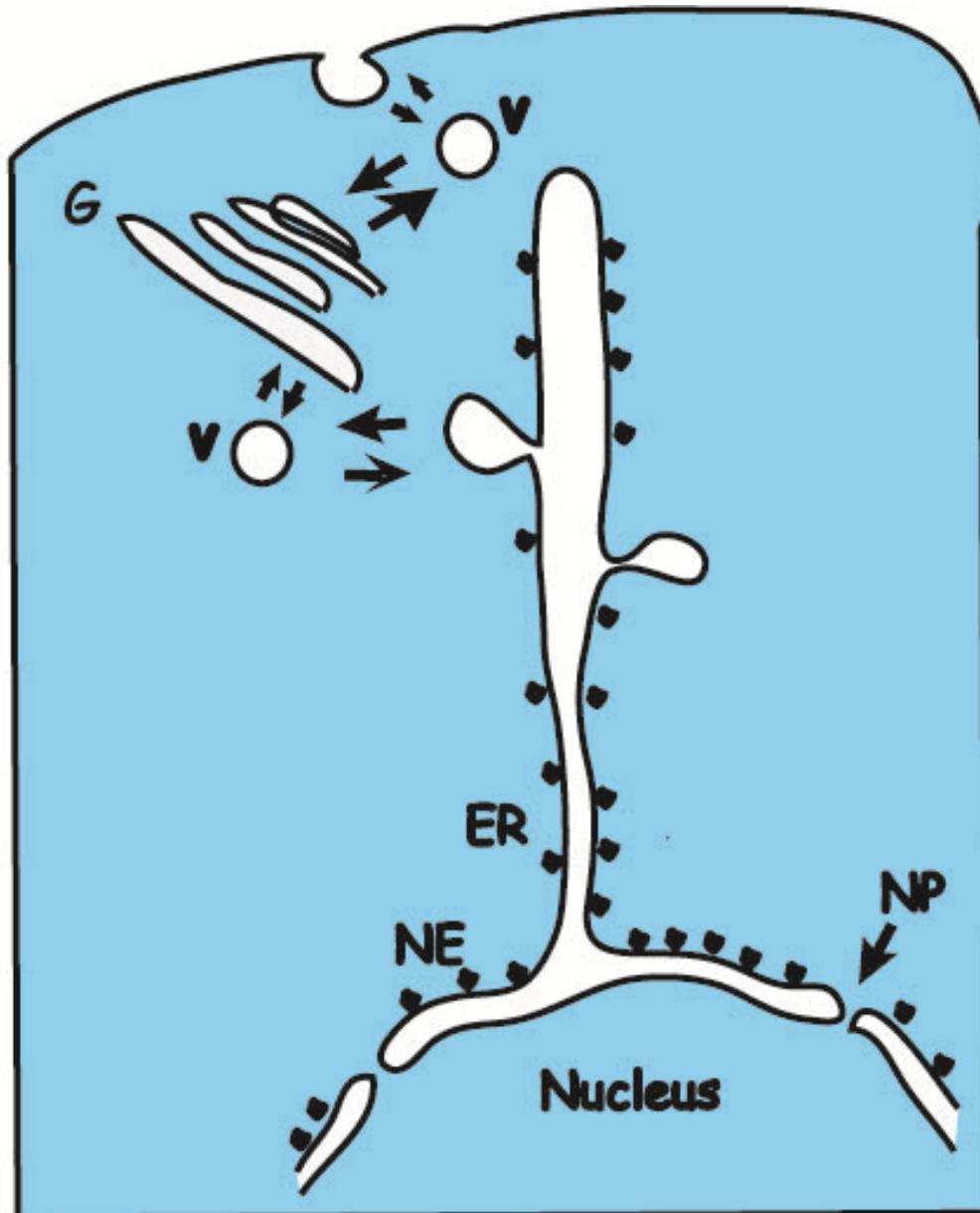


Santarella-Mellwig R et al. PLoS Biol. 2010 Jan 19;8(1):e1000281.

## Towards the first eukaryote ancestral nutrition

- First planctomycete may have taken up proteins and polysaccharides as a mode of nutrition
- This may be a clue to the origin of compartmentalization and the eukaryote endomembrane system re selective pressure for internal vesicles & trafficking

The **Evolved** Eukaryotic Endomembrane System - Essential Elements?  
Several features are already present in *Gemmata obscuriglobus*

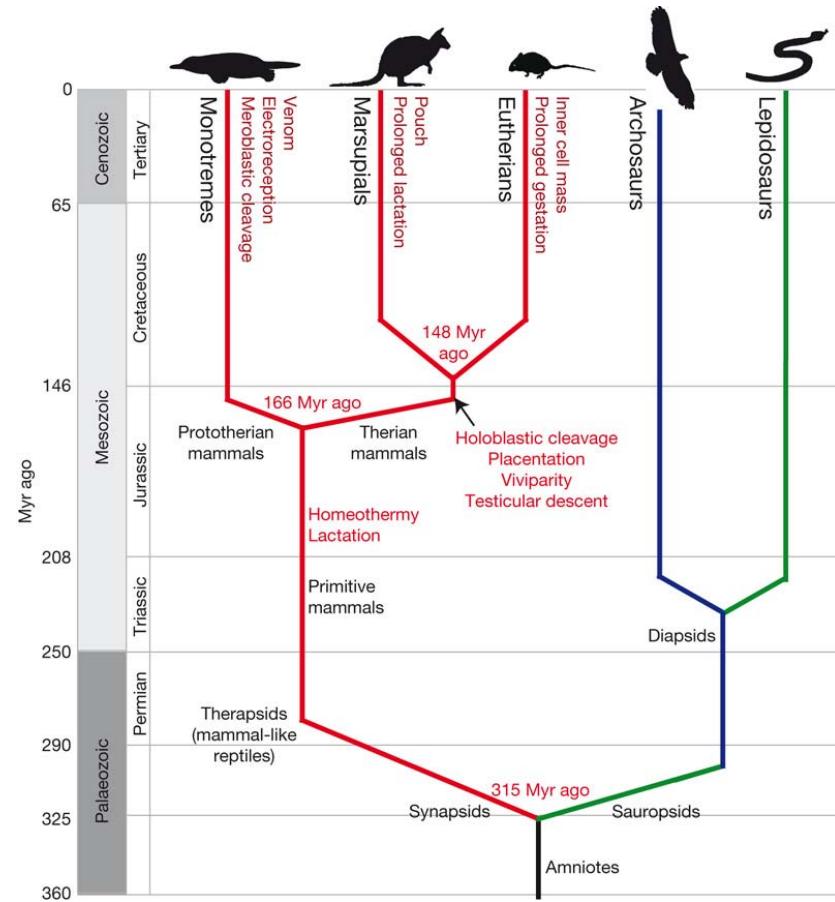


Modified from J Maynard Smith and Szathmary

Planctomycetes have both structural and functional features analogous to eukaryotes with some actual homology with eukaryote molecular mechanisms

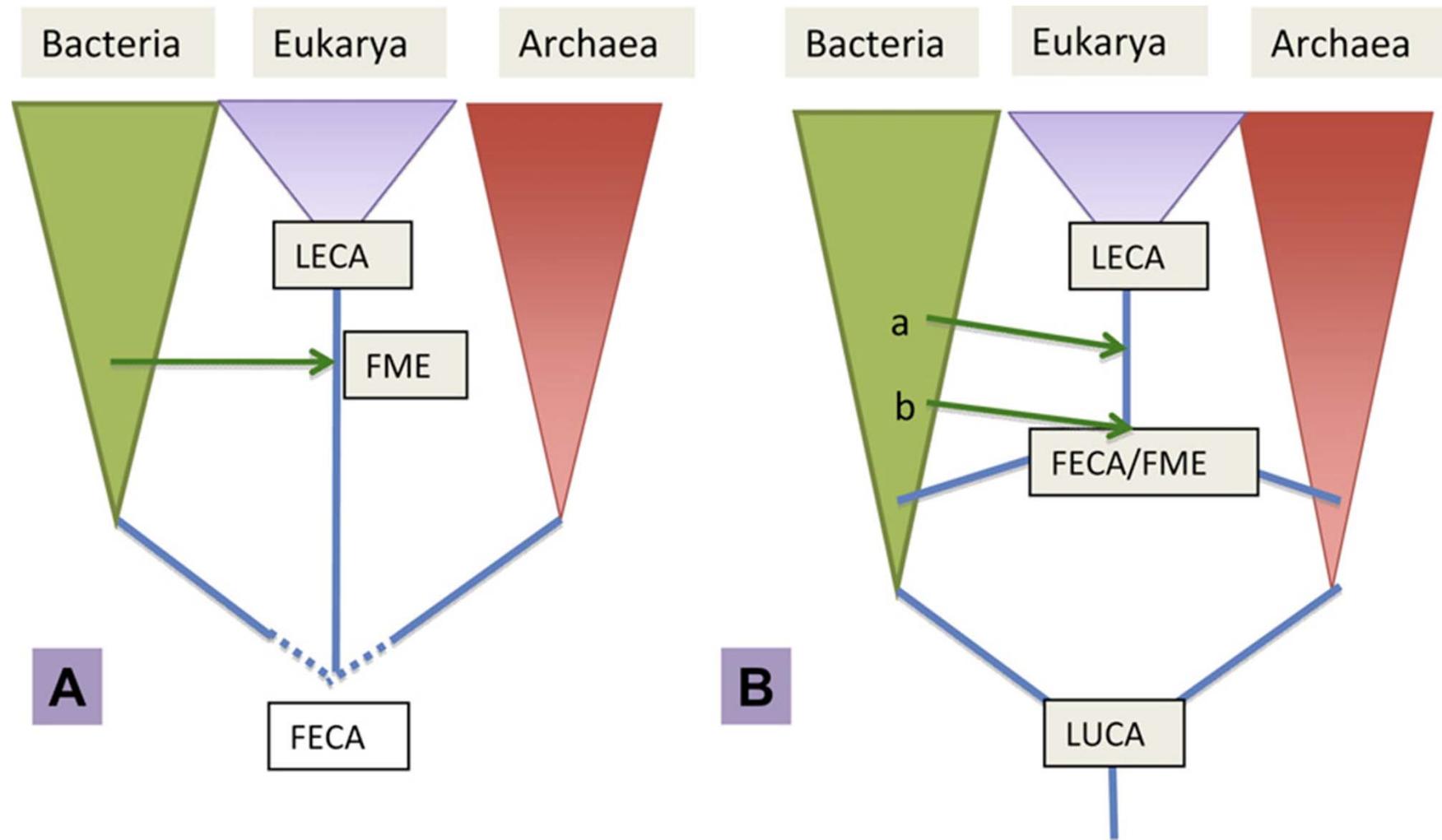
What's the evolutionary origin of such features?

# Is the planctomycete a Bacterial Platypus? - with a mix of features, some from deep ancestor of separate lineages?



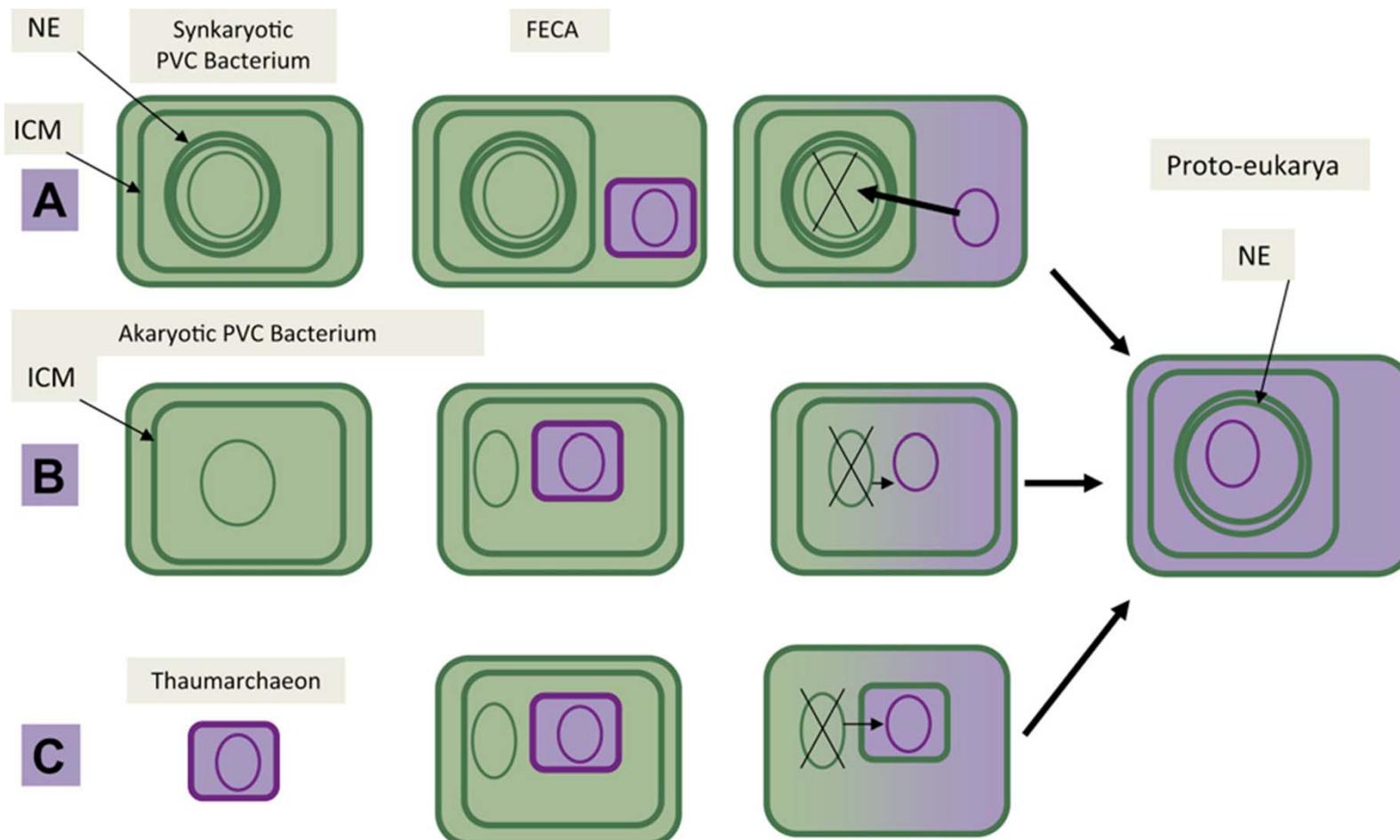
e.g. genomics reveals sex chromosome homology to birds but convergent evolution of venom peptides

# Domain tree models - eukarya ancestral or derived?



Forterre P. A new fusion hypothesis for the origin of Eukarya: better previous ones, but probably also wrong. Res Microbiol. 2011 Jan;162(1):77-91

# New improved but probably wrong fusion hypotheses for Euk origins

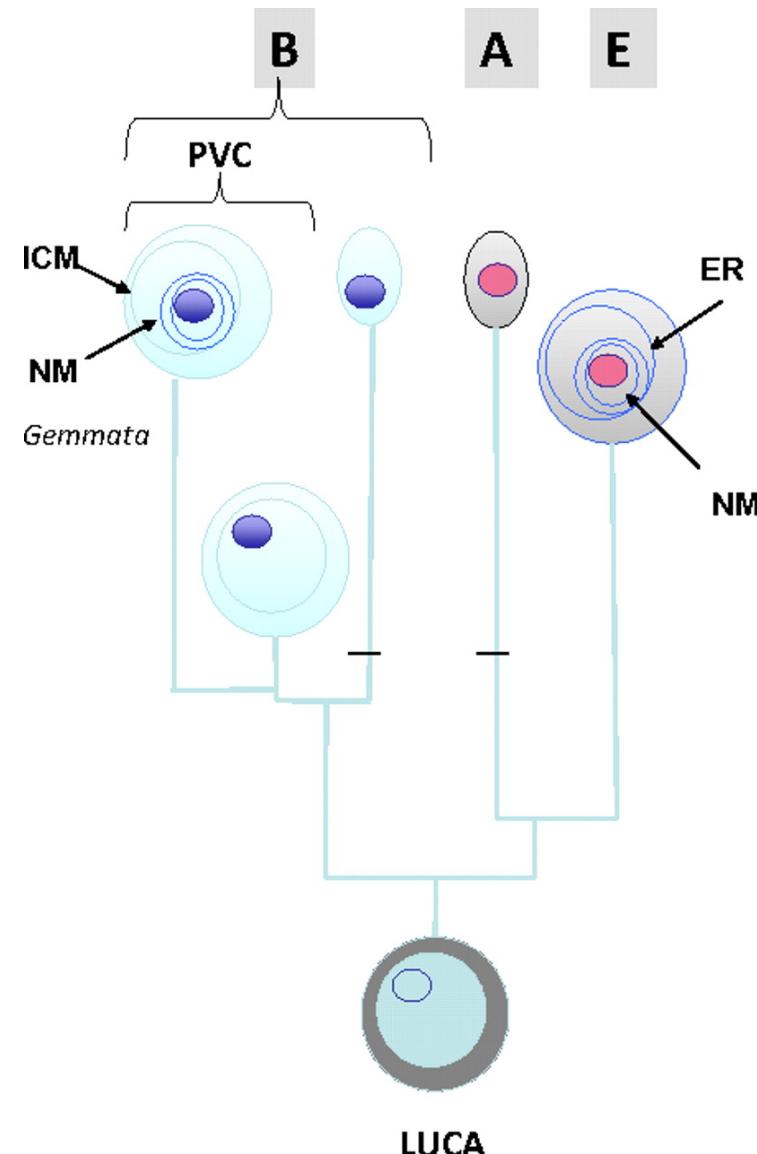


Forterre P. A new fusion hypothesis for the origin of Eukarya: better previous ones, but probably also wrong. Res Microbiol. 2011 Jan;162(1):77-91

# Lessons from *Gemmata*

- Fusion hypotheses are not necessary to explain origin of eukaryote nucleus, and some form of autogenous internal development of a nucleus is more parsimonious
- Uncouples origin of nucleus from origin of eukaryote chimeric genome
- Suggests retention from compartmentalized FECA or LUCA should again be considered

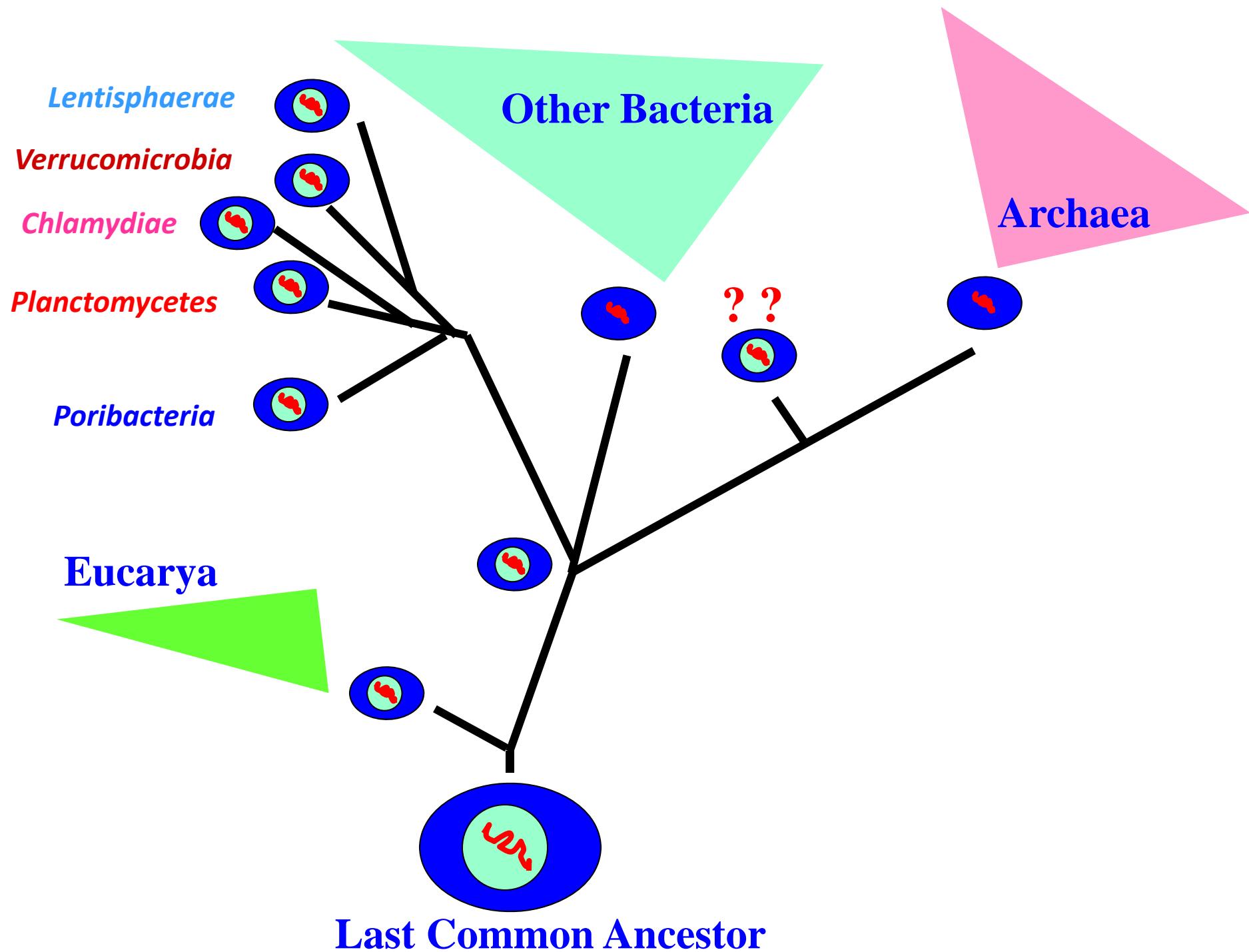
A ‘retention’ scenario for the origin of modern compartmentalized cells (Eukarya and PVC bacteria) from a compartmentalized LUCA. A, Archaea; B, Bacteria; E Eukarya.



Forterre P , Gribaldo S PNAS 2010;107:12739-12740

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PNAS





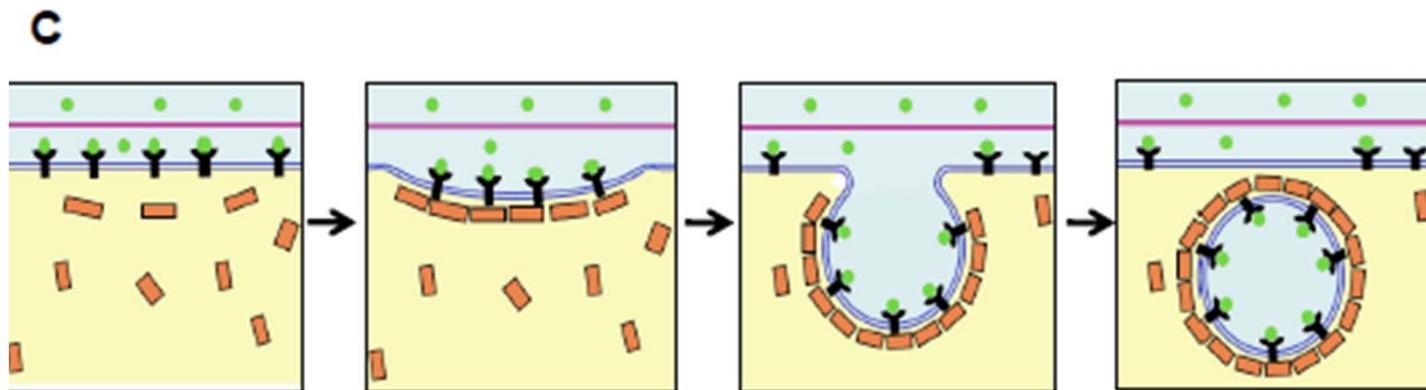
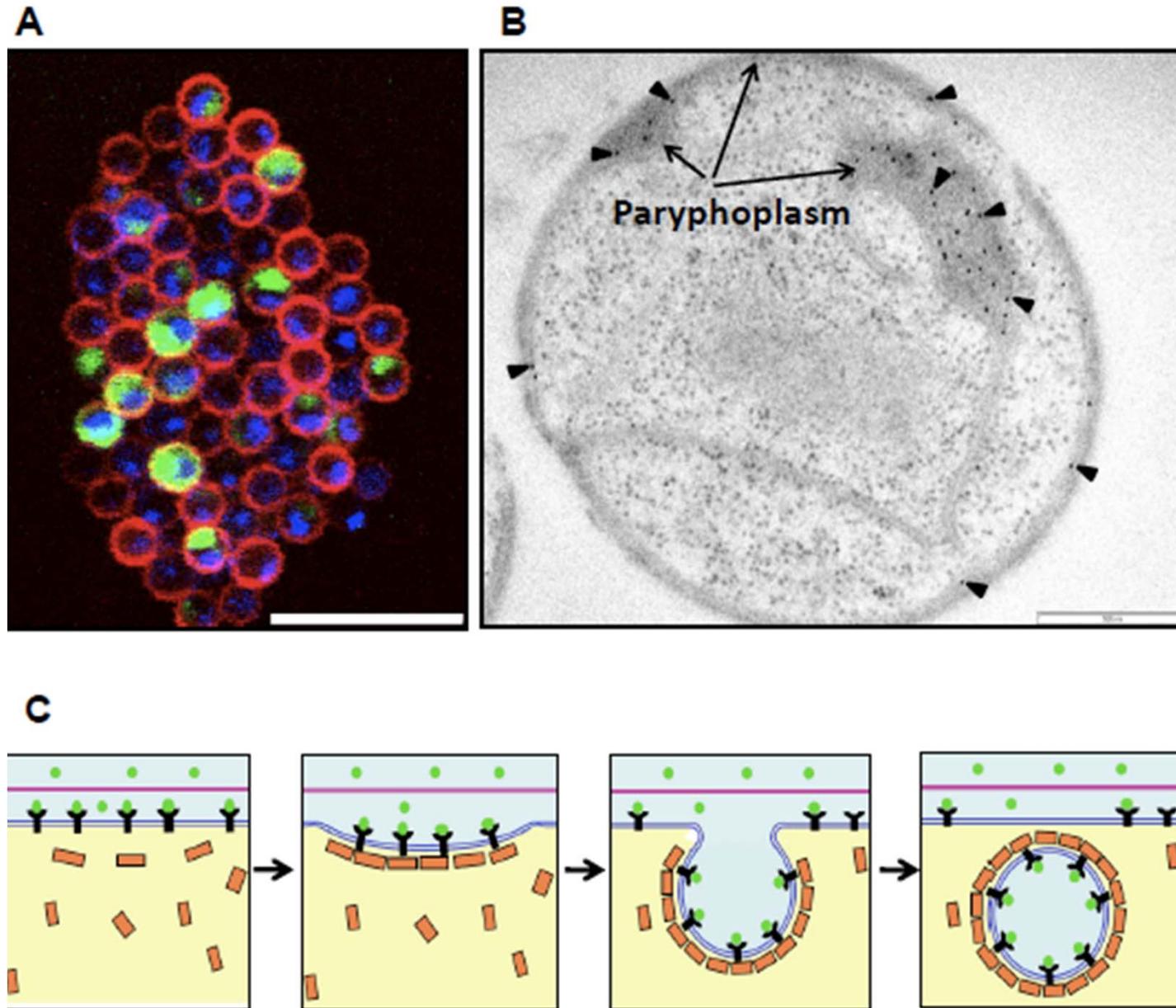
## Acknowledgements

- Australian Research Council
- Evgeny Sagulenko, Benjamin Yee, Jeffery Lee, Margaret Butler
- Collaborators on protein uptake research:  
Jody Franke, Mike Rout (Rockefeller University);  
Damien Devos (EMBL Heidelberg)  
Rick Webb (Centre for Microscopy and Microanalysis, The University of Queensland)



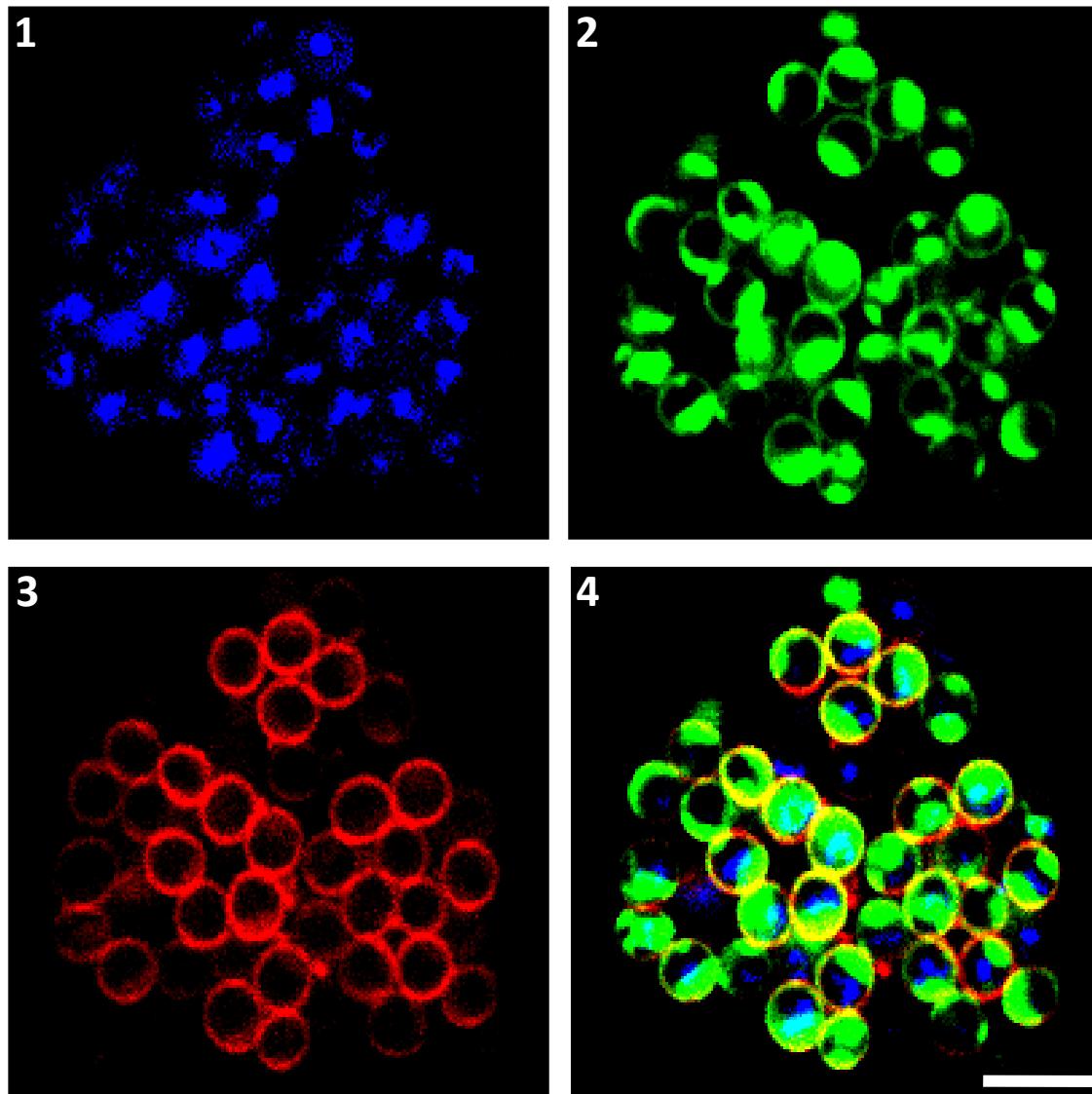


## Protein uptake by the planctomycete *Gemmata obscuriglobus*

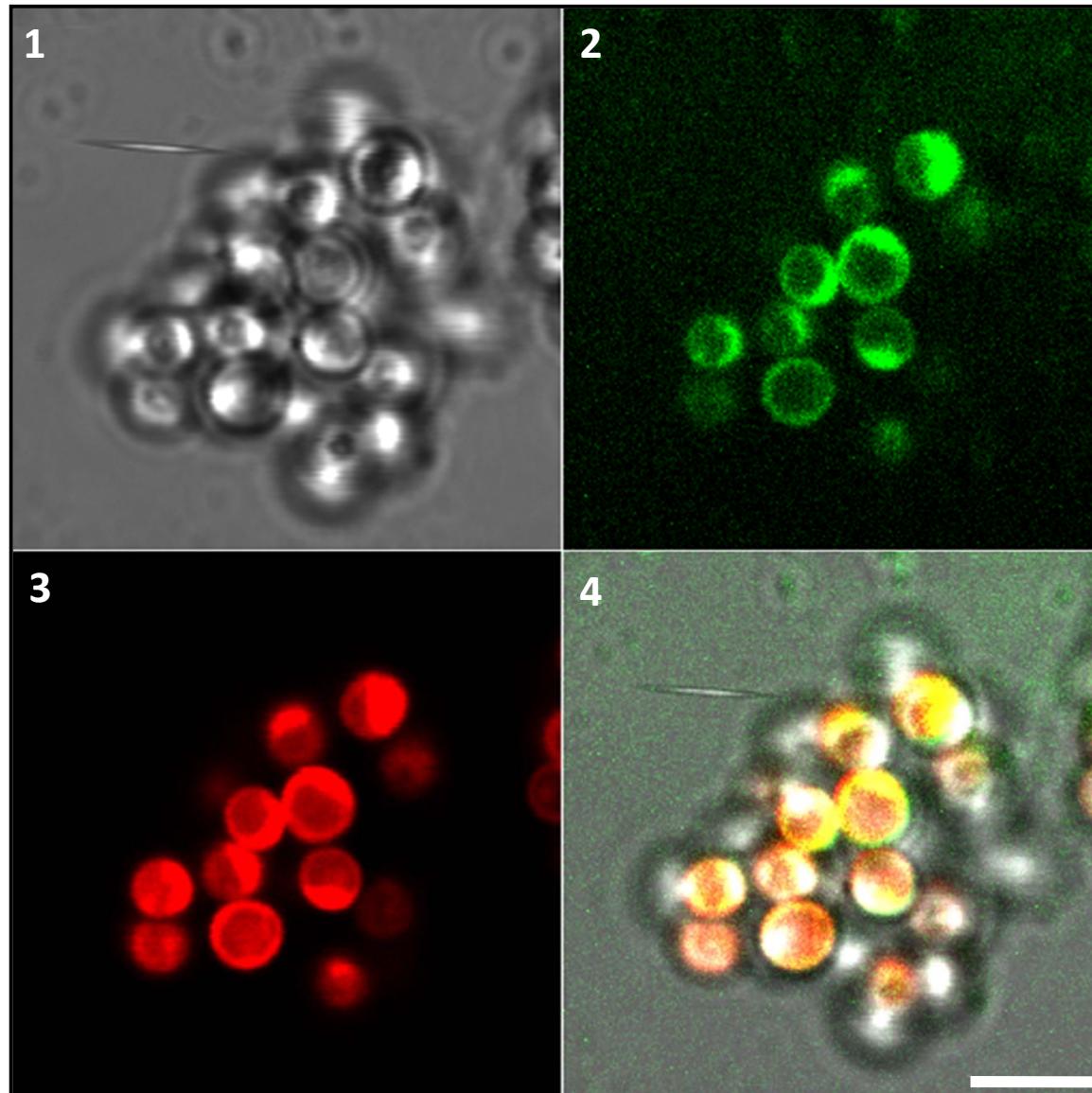


For B&C See: Fuerst, JA and Sagulenko, E. Communicative & Integrative Biology 3:6, 1-4, Nov/Dec 2010

# Uptake of FITC-dextran by *Gemmata obscuriglobus* cells

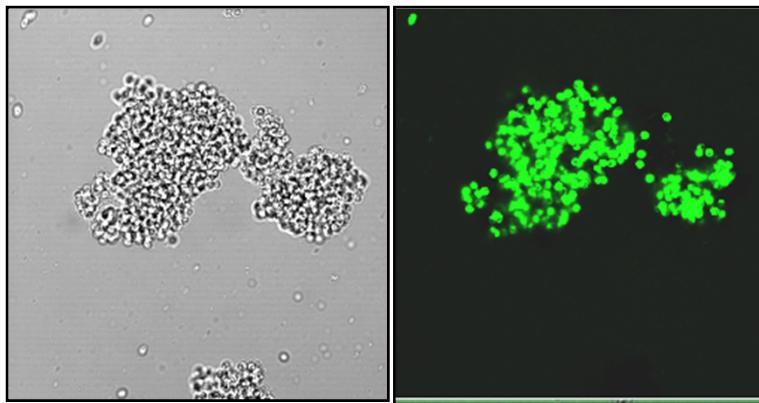


Rhodamine-dextran co-localizes with GFP in the paryphoplasm of *Gemmata obscuriglobus* cells

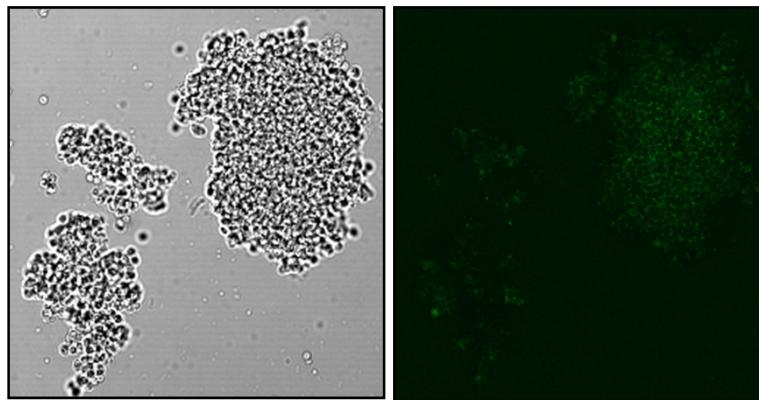


Energy dependent dextran uptake: Low temperature ( $0^{\circ}\text{C}$ ) and sodium azide inhibit uptake of FITC-dextran by *G. obscuriglobus*

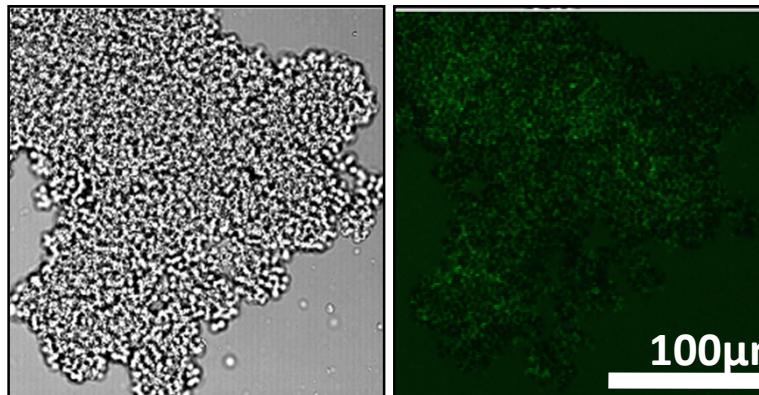
*FITC-dextran,*  
 $28^{\circ}\text{C}$



*FITC-dextran,*  
*+ sodium azide*

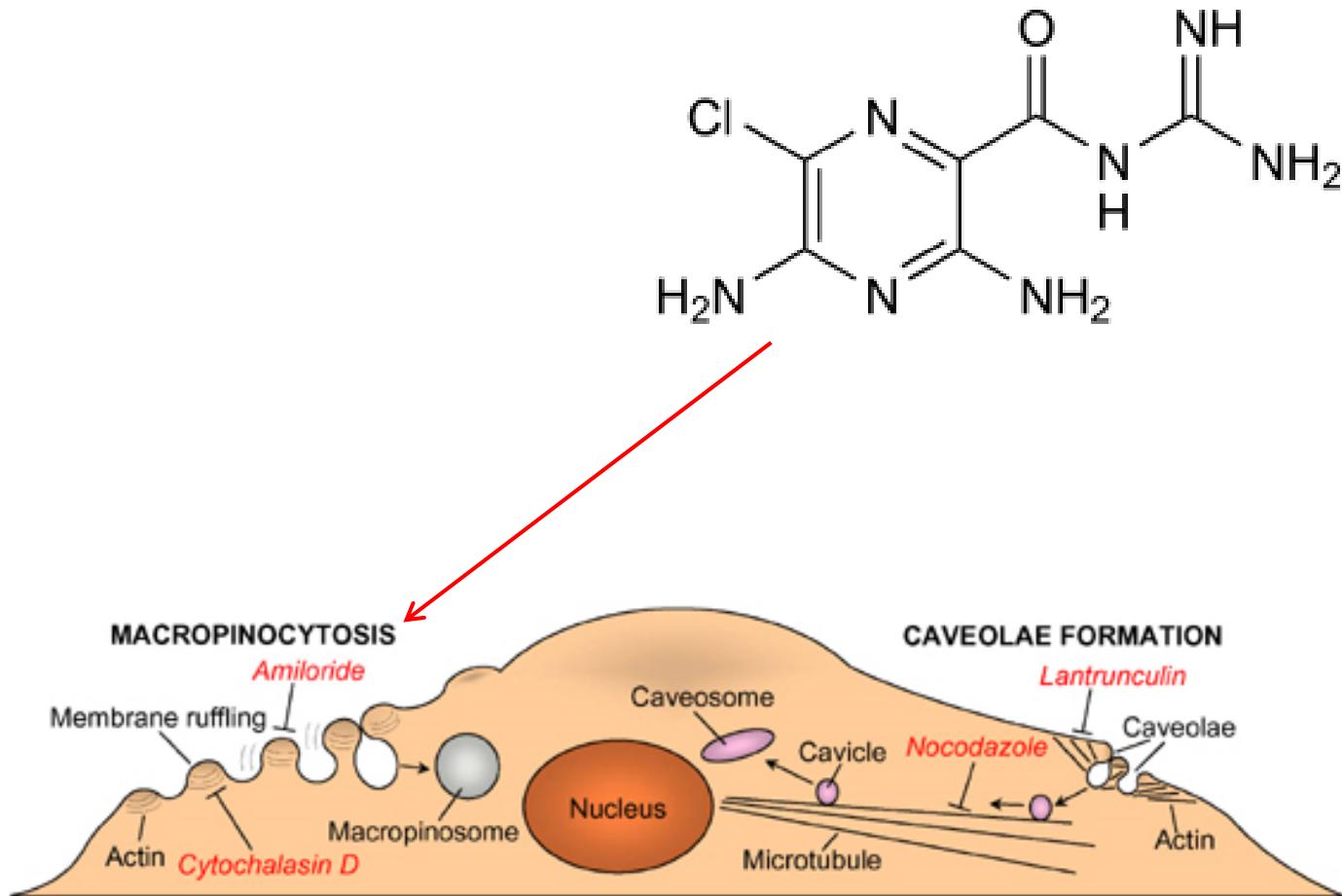


*FITC-dextran,*  
 $0^{\circ}\text{C}$



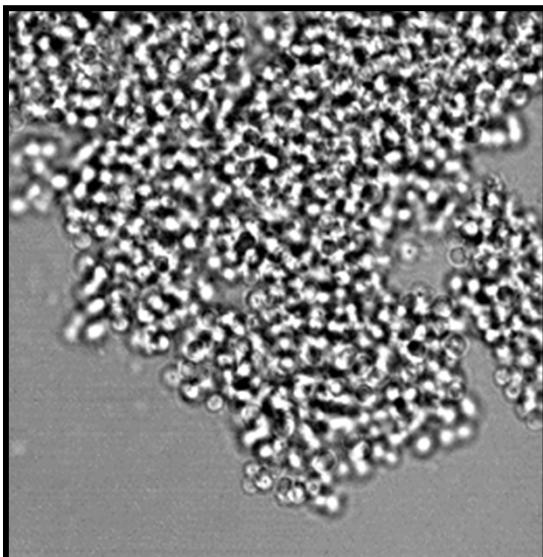
100 $\mu\text{m}$

# Macropinocytosis is inhibited by amiloride, which blocks the $\text{Na}^+/\text{H}^+$ exchanger and inhibits ruffling

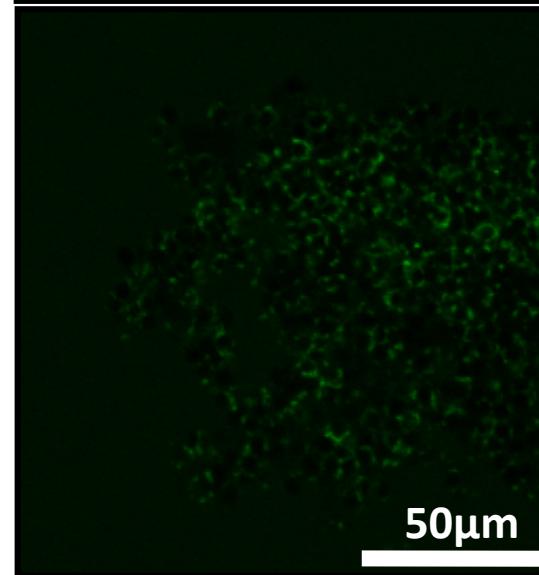
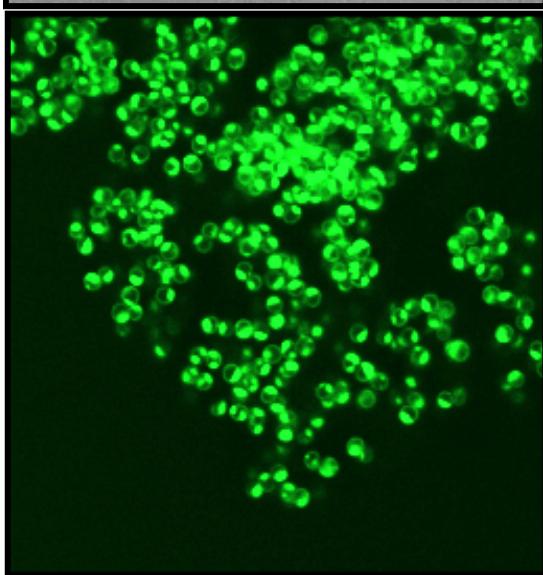
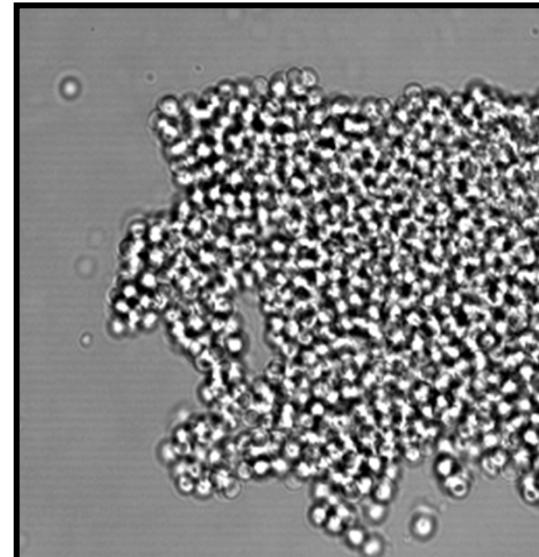


# Amiloride inhibits uptake of FITC-dextran by *Gemmata obscuriglobus*

FITC-dextran

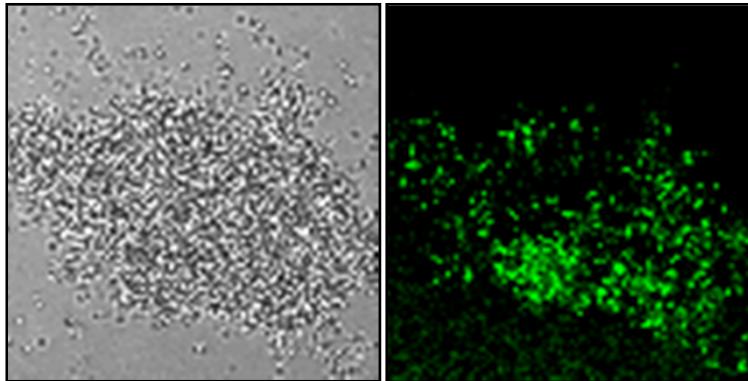


FITC-dextran + amiloride

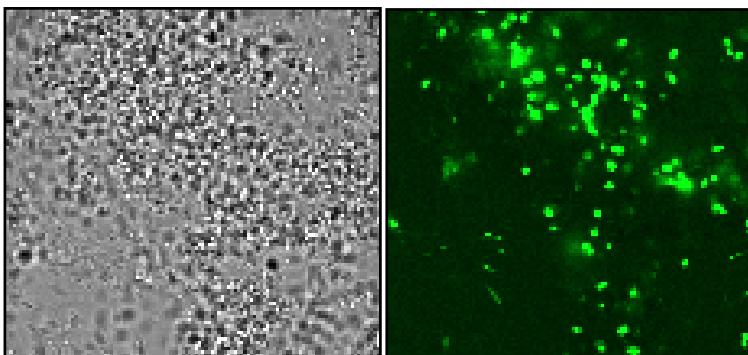


# Marine planctomycetes can uptake FITC-dextran

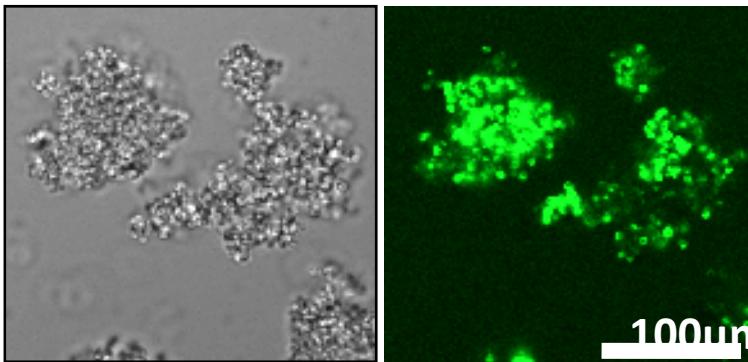
*Pirellula  
staleyi*



*Blastopirellula  
marina*

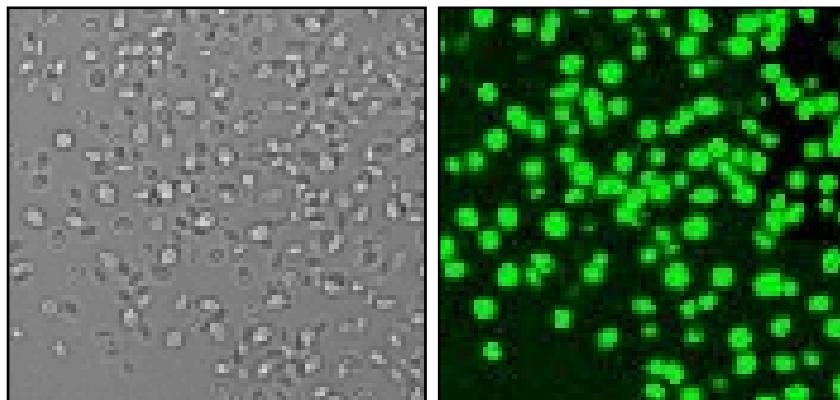


*DDSe3013*

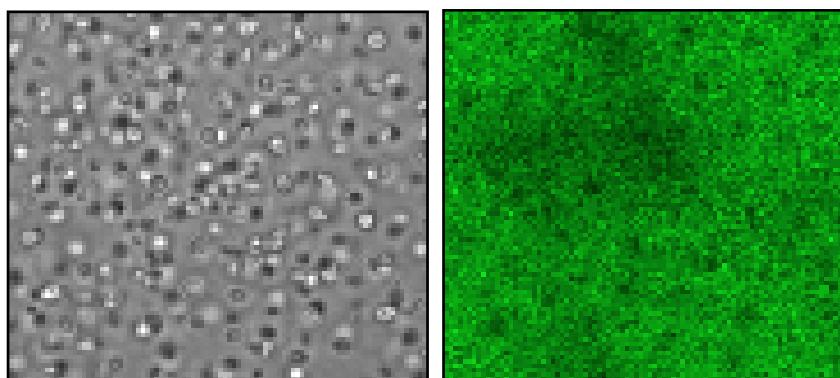


**Marine planctomycetes can uptake GFP-dextran only in salt water, freshwater bacteria - only in freshwater**

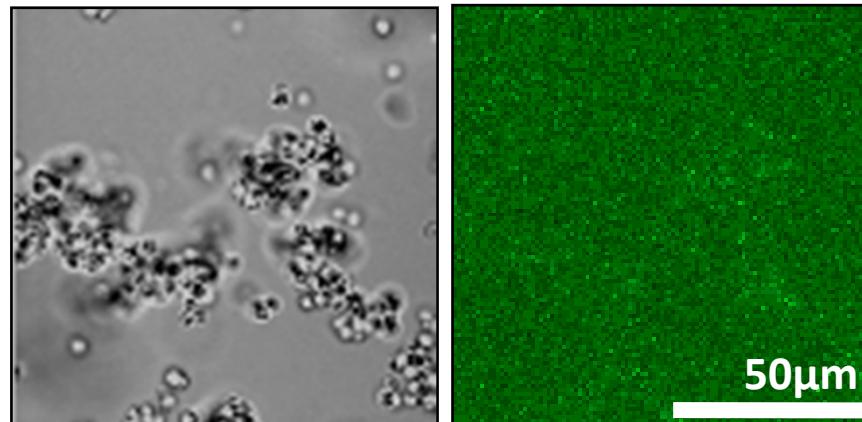
DDSe3017-  
artificial  
seawater



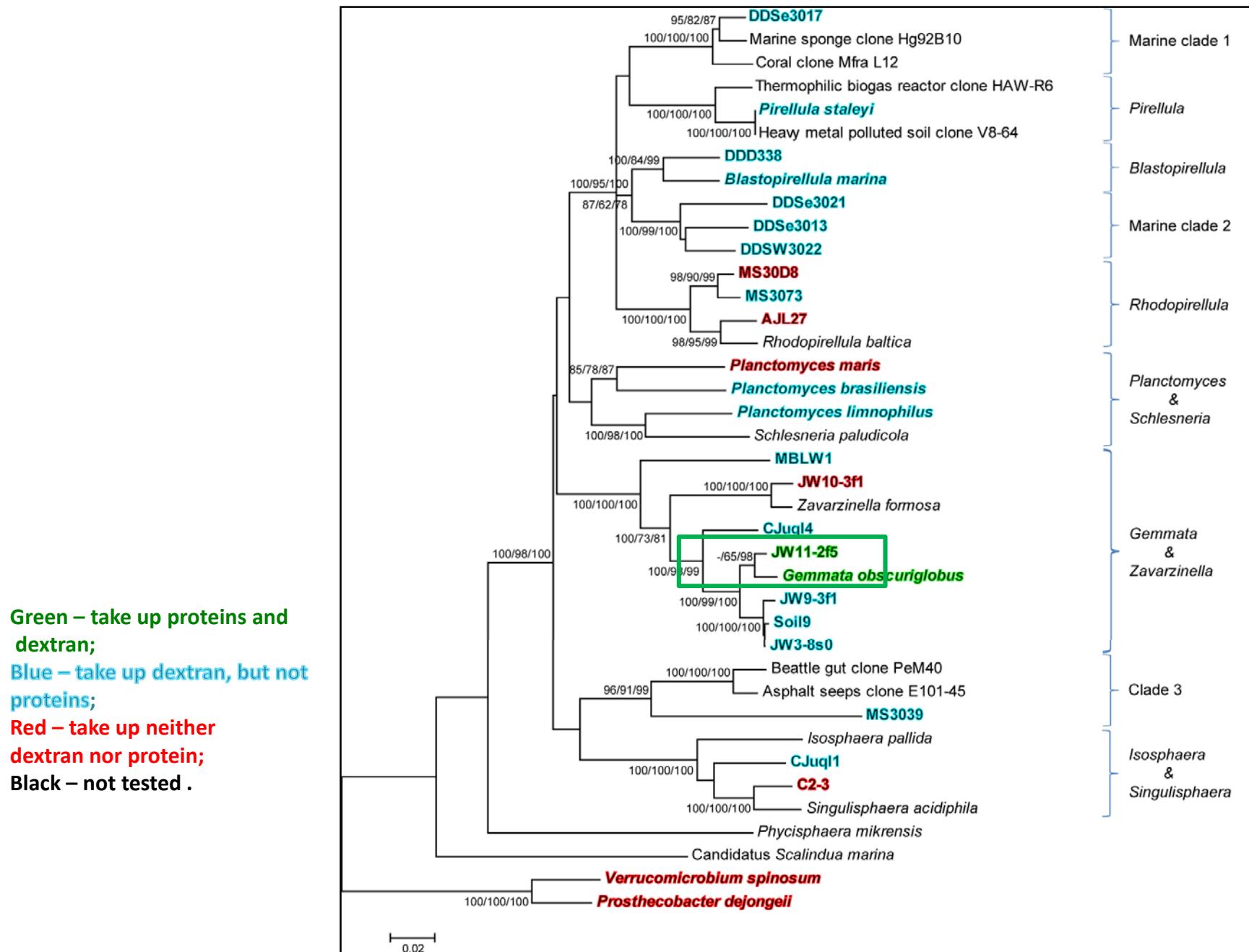
DDSe3017-  
freshwater



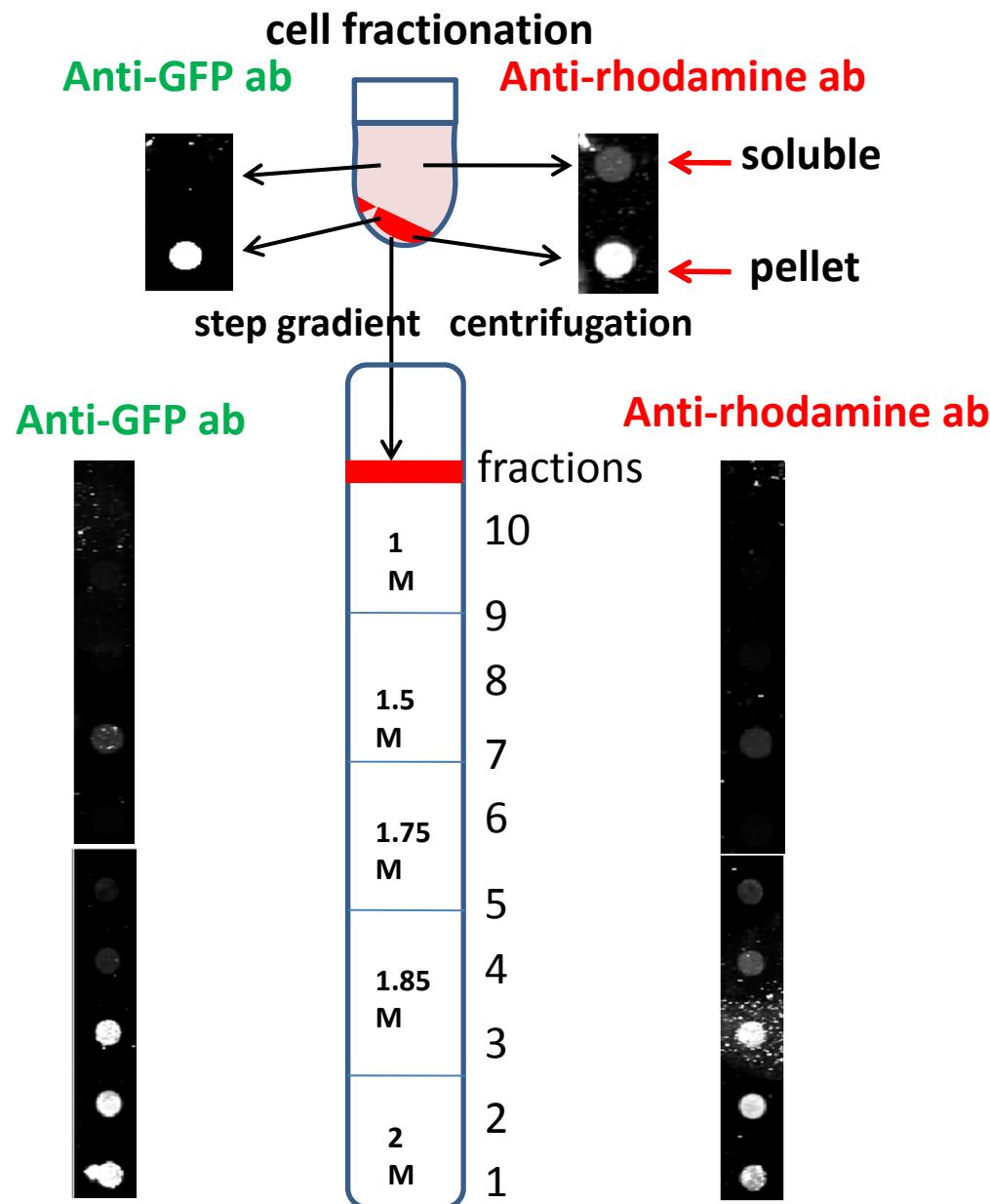
JW11-2F5-  
artificial  
seawater



# Phylogenetic tree based on 16S rRNA gene of planctomycetes and verrucomicrobia strains tested for macromolecule uptake



## GFP and rhodamine-dextran co-localize to the same lysed cell fraction



*Gemmata obscuriglobus* cell fractionation and step gradient centrifugation

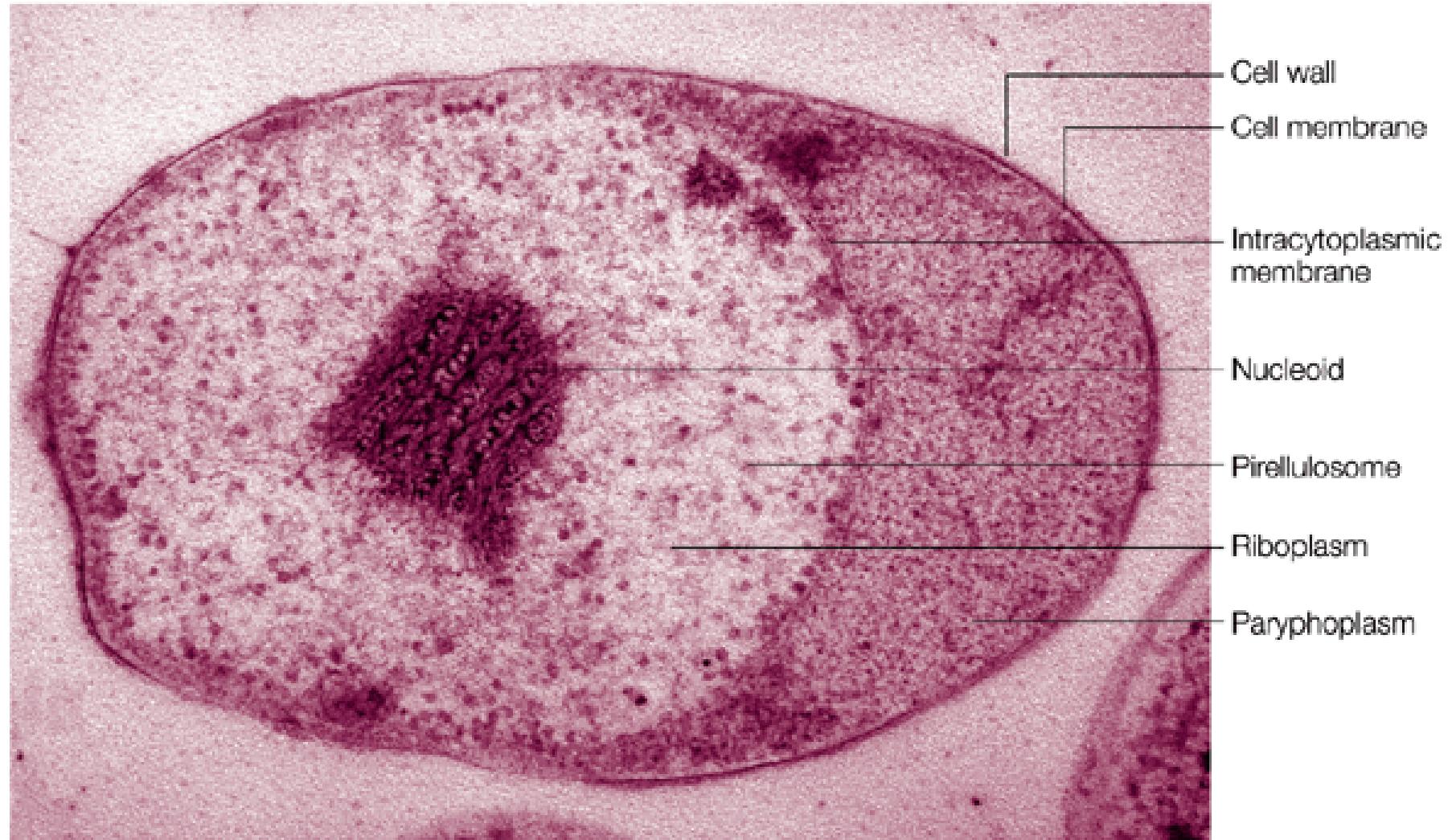
- Forterre's 'unlikely' PTV (planctomycete-thaumarchaea-virus) fusion models of PVC contributions to eukaryality -
- NE could be early & PVC-derived but FECA would still need archaeal contribution? (e.g. thaumarchaea supply actins, ESCRT genes, informational system, PVC supply tubulin, sterols and NE)
- Viral contributions needed to explain full eukaryal complexity

# A Bacterial model for origin of eukaryote compartmentalization? - The Planctomycetes

**Planctomycetes** are:

- Members of **Domain Bacteria** (e.g. by 16S rRNA sequence trees)
- Form distinct **separate Phylum** within Domain –but may be member of the '**PVC' superphylum**
- Budding, aquatic/soil bacteria
- Cell walls mostly protein - possess **no peptidoglycan** (unlike most other domain Bacteria members)
- All possess **compartments with DNA enveloped by membrane (single or single+double)**
- **Thus – may be analogous or homologous with an early eukaryote?**

**Cell plan of *Pirellula* group planctomycetes- TEM of sectioned cryosubstituted cell of *Blastopirellula marina***



From: Nature Reviews Microbiology 1:11-12 (2003)

Nature Reviews | Microbiology

## Planctomycete Cell Plans

