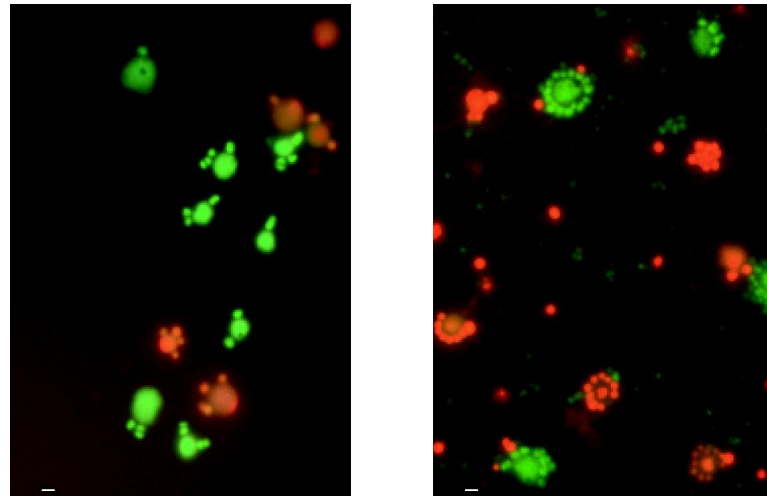


Analysing protein exchange between *Ignicoccus hospitalis* KIN4/1T and *Nanoarchaeum equitans*



Epifluorescence micrographs of *Ignicoccus/Nanoarchaeum* coculture stained with BacLight (Boulos *et al.*, 1999) adapted from (Jahn *et al.*, 2008)
(Scale bar: 1 μm)

Summary

- Objectives of research
- Relevance and importance of research
- Discovery of co-culture
- Biology of *I. hospitalis* and *N. equitans*
- Current understanding of association
- Experimental limitations of biological system
- Proposed methodologies to investigate protein exchange between *I. hospitalis* and *N. equitans*
- Conclusions

Objectives

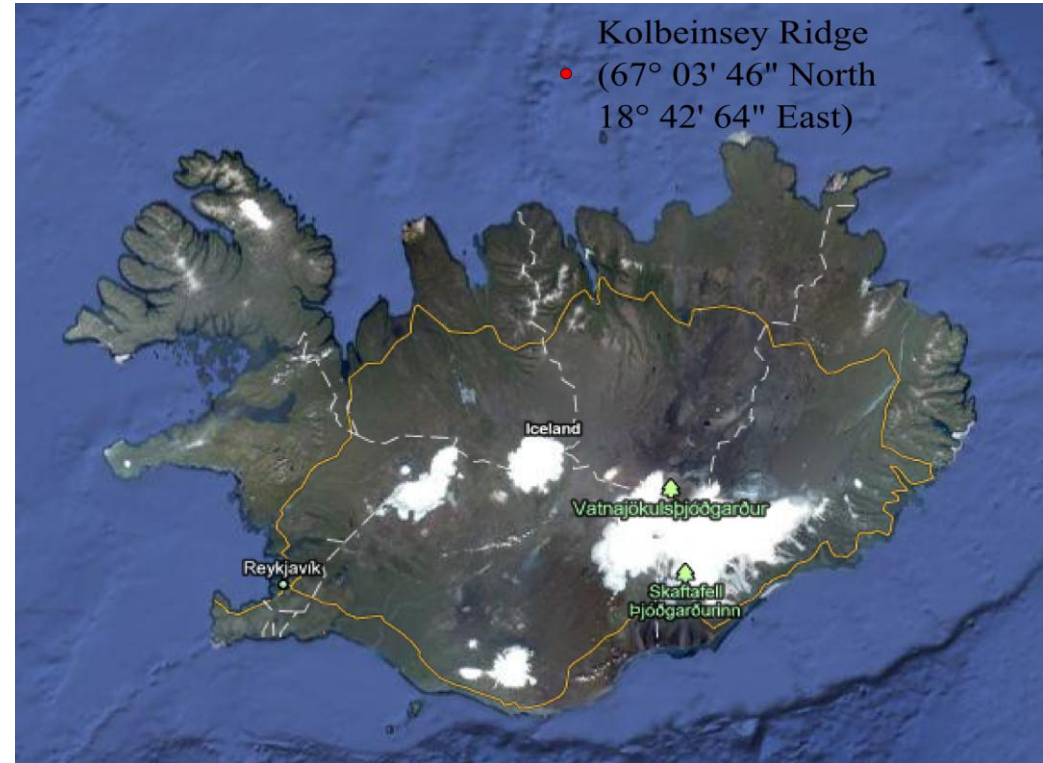
- Determine the role of Secretory (Sec) and Twin-Arginine-Transporters (TAT) in the exchange of proteins between *Nanoarchaeum equitans* and *Ignicoccus hospitalis*
 - Identify candidate proteins for Sec or TAT transport
 - Determine the localisation of TAT transporters in *I. Hospitalis*
 - Test competence of *I. hospitalis* Sec and TAT complexes for export of identified candidate proteins
 - Test competence of *N. equitans* SecDF complex for candidate protein uptake
 - Identify further avenues of research

Relevance

- Why are *Ignicoccus hospitalis* and *Nanoarchaeum equitans* of interest?
 - Hyperthermophiles (Leigh *et al.*, 2011)
 - Novel proteins (Podar *et al.*, 2008a)
 - Very ancient lineages? (Podar *et al.*, 2008a)
 - Novel phyla in case of *Nanoarchaeum equitans*? (Huber *et al.*, 2003)
 - Evolution of the eukaryotic cell? (Kuper *et al.*, 2010)
 - Evolution of a vesicle trafficking system (Podar *et al.*, 2008b)
 - Evolution of species co-associations (Mevarech and Allers, 2007)

Discovery of organisms

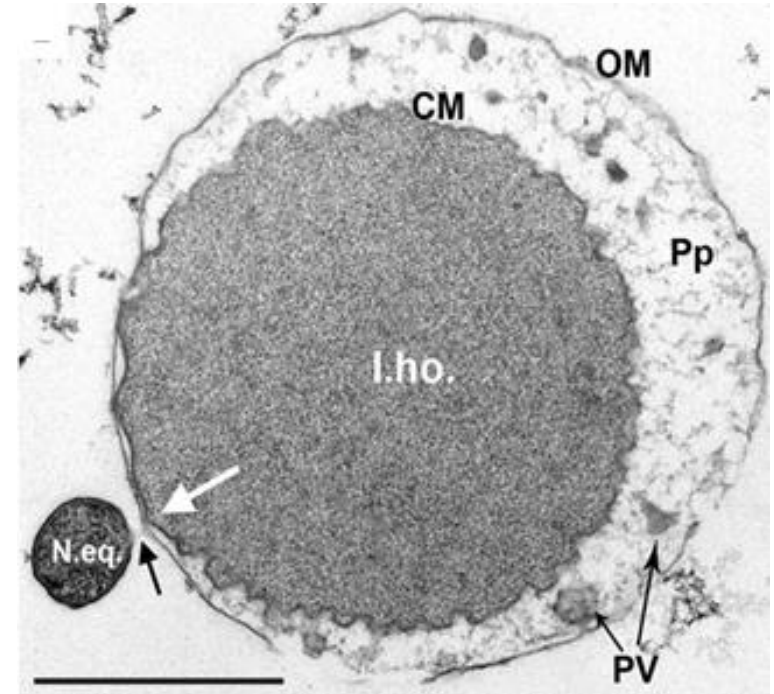
- Hydrothermal system at Kolbeinsey Ridge from depth of 106m (Fricke *et al.*, 1989)
- *Ignicoccus hospitalis* KIN4/I isolate
- Discovery of *Nanoarchaeum equitans* by Karl Stetter in 2002
- Unique relationship (Burghardt *et al.*, 2009)
- Stable co-culture established at University of Regensburg



Map showing location of Kolbeinsey Ridge

Ignicoccus hospitalis

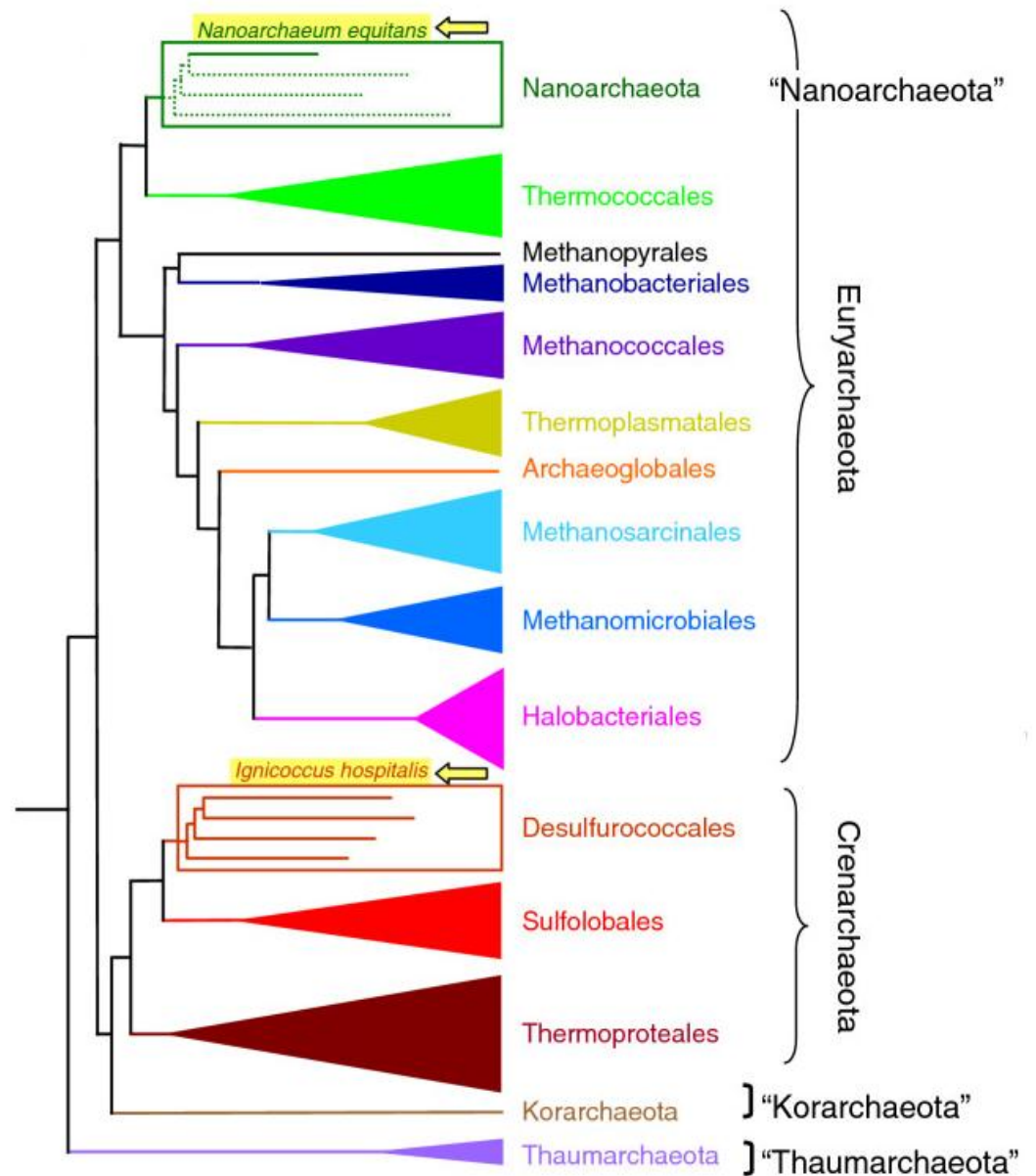
- Obligate anaerobe (Forterre *et al.*, 2009)
- Hyperthermophile (Forterre *et al.*, 2009)
- Ancient organism? (Podar *et al.*, 2008a)
- Unusual morphology (Paper *et al.*, 2007, Burghardt *et al.*, 2007)
- Unusual metabolism (Junglas *et al.*, 2008)
- Unique carbon assimilation (Junglas *et al.*, 2008)
- Smallest free-living genome (Podar *et al.*, 2008)



**Transmission electron micrographs
of ultrathin sections
of *I. hospitalis* and *N. equitans***
CM: Cytoplasmic membrane
OM: Outer membrane
Pp: Periplasm
Figure from (Jahn *et al.*, 2008)
(Scale Bar: 1µm)

Nanoarchaeum equitans

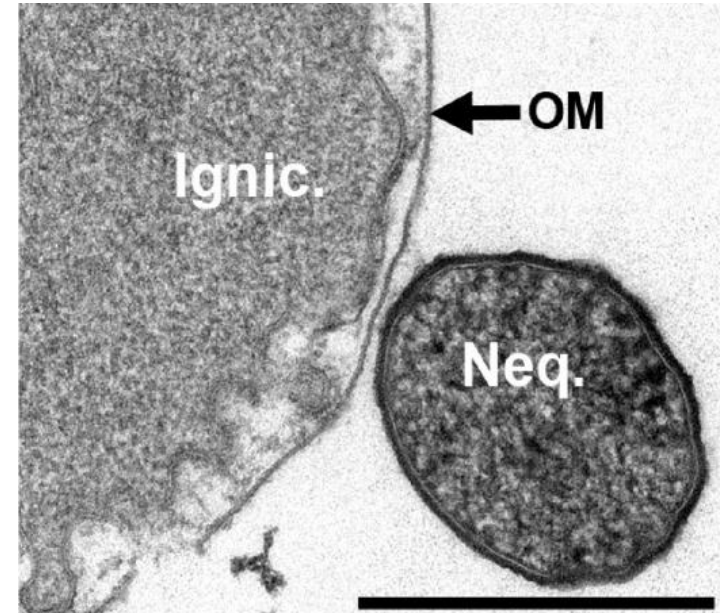
- Nanoarchaeota (Huber *et al.*, 2002)
- Smallest genome in archaea (Huber *et al.*, 2003)
- Obligate symbiont / parasite (Waters *et al.*, 2003)
- Lacks key genes (Podar *et al.*, 2008a)
- Unknown metabolism (Lewalter and Muller, 2006)



Archael Phylogeny from (Forterre *et al.*, 2009)

Physiological dependence

- Host-derived
 - Amino acids (Jahn *et al.*, 2008)
 - Lipids (Jahn *et al.*, 2004)
- *Ignicoccus* protein exporters:
 - SecYE/61 β complex (Burghardt *et al.*, 2009)
 - Twin-arginine translocation (TAT) system (Podar *et al.*, 2008a)
- *Nanoarchaeum* putative protein importer:
 - SecDF complex (Burghardt *et al.*, 2009)



**Electron micrograph showing
Nanoarchaeum equitans attached
to *Ignicoccus hospitalis*
OM: Outer membrane**

**Figure from (Forterre *et al.*, 2009)
(Scale bar: 100nm)**

Limitations of experimental system

- Genetic methods unavailable (Burghardt *et al.*, 2009)
- Key difficulties: (Mevarech and Allers, 2007)
 - Solid media cultivation
 - Transformation systems
 - Enrichment
 - RNAi unavailable
- Divergent from the standard genetic models (Leigh *et al.*, 2011)
- Enigmatic genes (Podar *et al.*, 2008a)
- Culture density (Huber *et al.*, 2003)

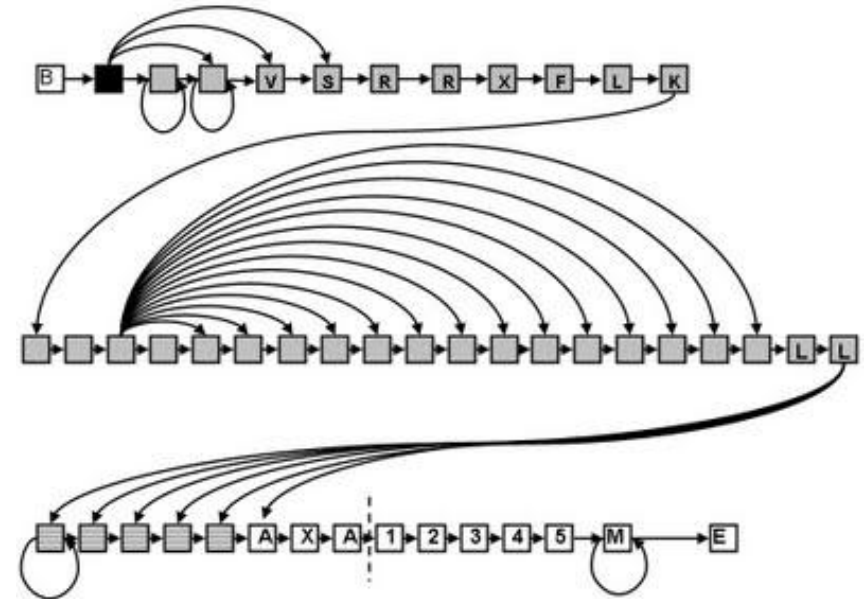


**BD BioSciences FACS Aria-II
cell sorter**

From (<http://www.bdbiosciences.com>)

Identification of candidate transferred proteins

- Combination survey using existing bioinformatic tools and heuristic approaches:
 - PRED-TAT (Bagos *et al.*, 2010)
 - TatP (Bendtsen *et al.*, 2005)
 - TATFIND (Rose *et al.*, 2002)
 - SignalP 3.0 (Bendtsen *et al.*, 2004)
 - Phobius (Kall *et al.*, 2004)
- Preliminary survey of *I. hospitalis* protein database:
 - 8 Sec signal peptide-containing proteins
 - 3 TAT signal peptide-containing proteins



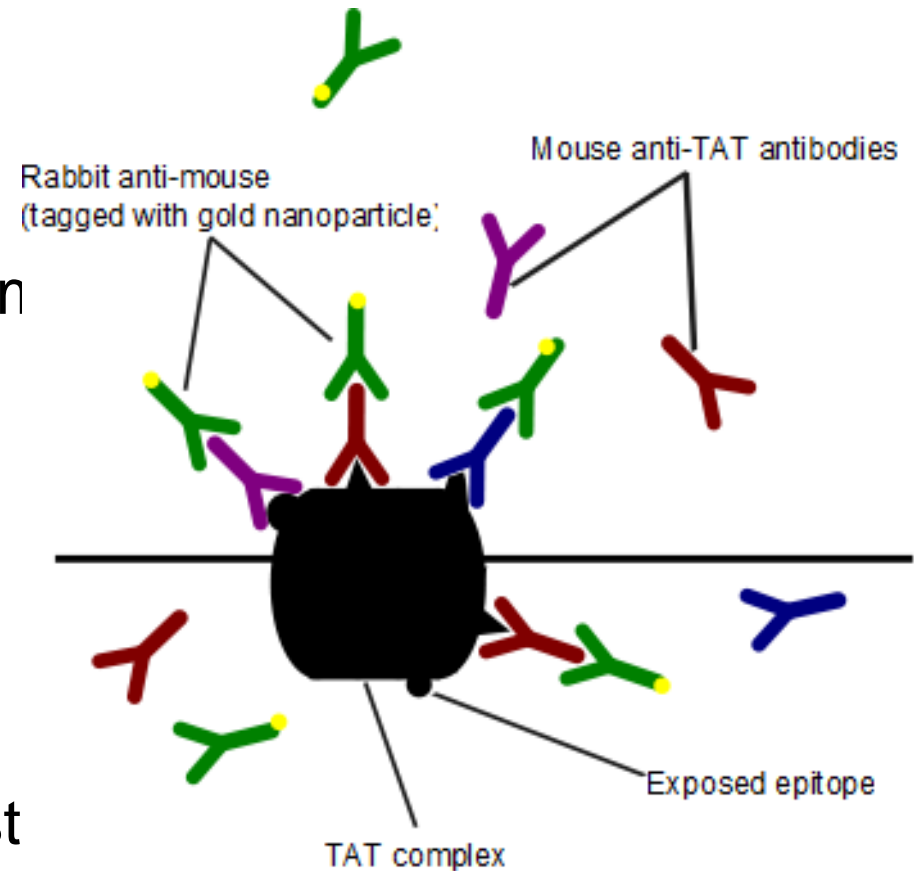
PRED-TAT Hidden Markov Model diagram
Figure from (Bagos *et al.*, 2010)

Culturing organisms

- Basic growth conditions:
 - Seawater medium (Huber *et al.*, 2000)
 - Anoxic: Gas phase of H₂-CO₂ (80/20 vol/vol) at 300kPa (Paper *et al.*, 2007)
 - pH 5.5-6.0 (Paper *et al.*, 2007)
 - Temperature: 90°C (Mevarech and Allers, 2007)
- Final cell densities: 2x10⁷ cells ml⁻¹ (Huber *et al.*, 2003)
- Modifications to increase cell density:
 - Cellulose capillaries (increase to 3x10⁷ cells ml⁻¹) (Paper *et al.*, 2007, Kuper *et al.*, 2009)
 - H₂S stripping (increase of *Nanoarchaeum* density to 3x10⁸ cells ml⁻¹) (Mevarech and Allers, 2007)

Localisation of complexes

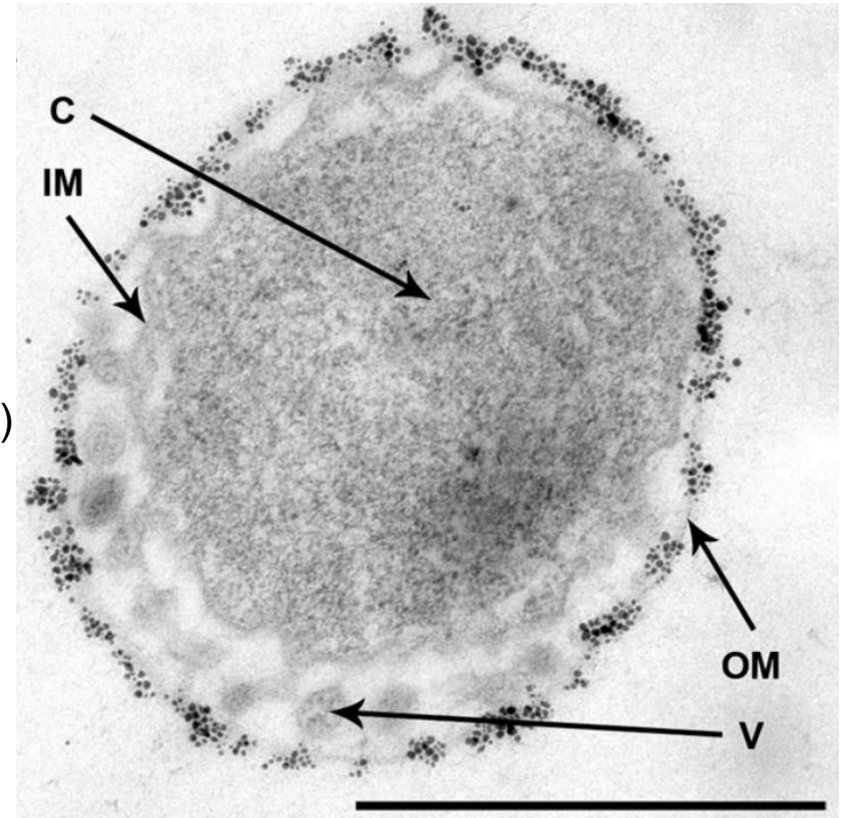
- Sec complexes previously isolated at interaction site (Burghardt *et al.*, 2009)
- Isolate and purify TAT complex from *I. hospitalis* via procedure used in (Porcelli *et al.*, 2002)
 - Membrane solubilisation
 - Ultracentrifugation
 - SDS-PAGE
- Raise polyclonal antibodies against purified TAT protein using mouse system



Immunolocalisation using polyclonal antibodies and secondary antibody markers

Sectioning and labelling

- Cryoimmobilisation via high-pressure freezing (Kuper *et al.*, 2009)
- Freeze-substitution dehydration (Walther and Ziegler, 2002)
- Embed in Epon resin (Junglas *et al.*, 2008)
- Serial ultrathin sections (70nm) (Junglas *et al.*, 2008)
- Incubate with primary rabbit anti-TAT antibody
- Incubate with secondary anti-rabbit antibody with gold nanoparticles
- Transmission electron micrography (Kuper *et al.*, 2009)

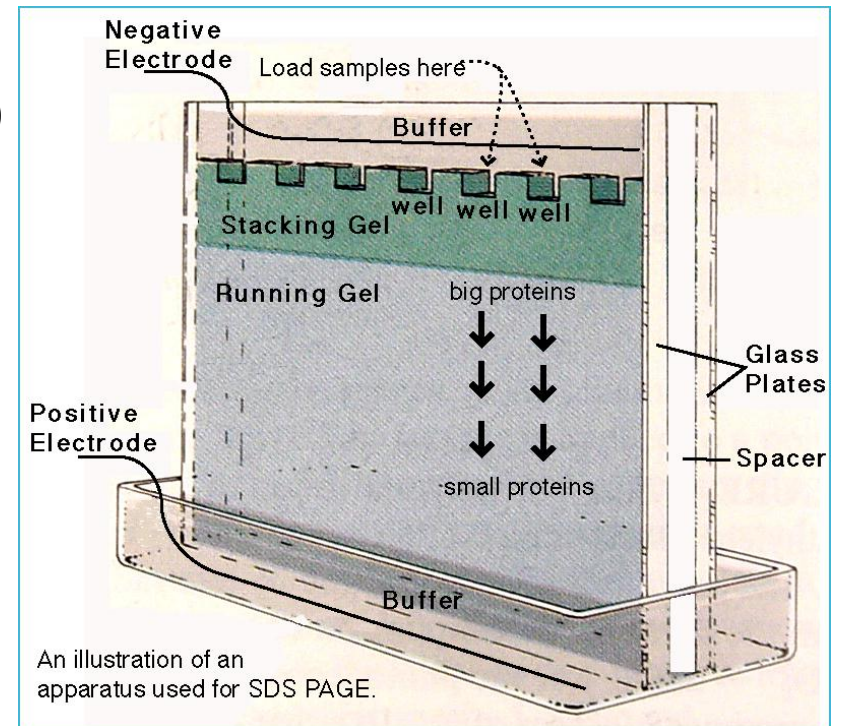


Immunolabelled A_1A_0 ATP-synthase

Figure from (Kuper *et al.*, 2009)
(Scale Bar 1 μ m)

Ignicoccus hospitalis Sec and TAT export competence

- Generation of *Ignicoccus* inverted membrane vesicles (Ring and Eichler, 2001)
 - French Press
 - Centrifugation and resuspension
- Isolation and purification of candidate proteins
 - Size-exclusion chromatography
 - Centrifugation
 - SDS-PAGE
 - Protein-specific biophysical separation

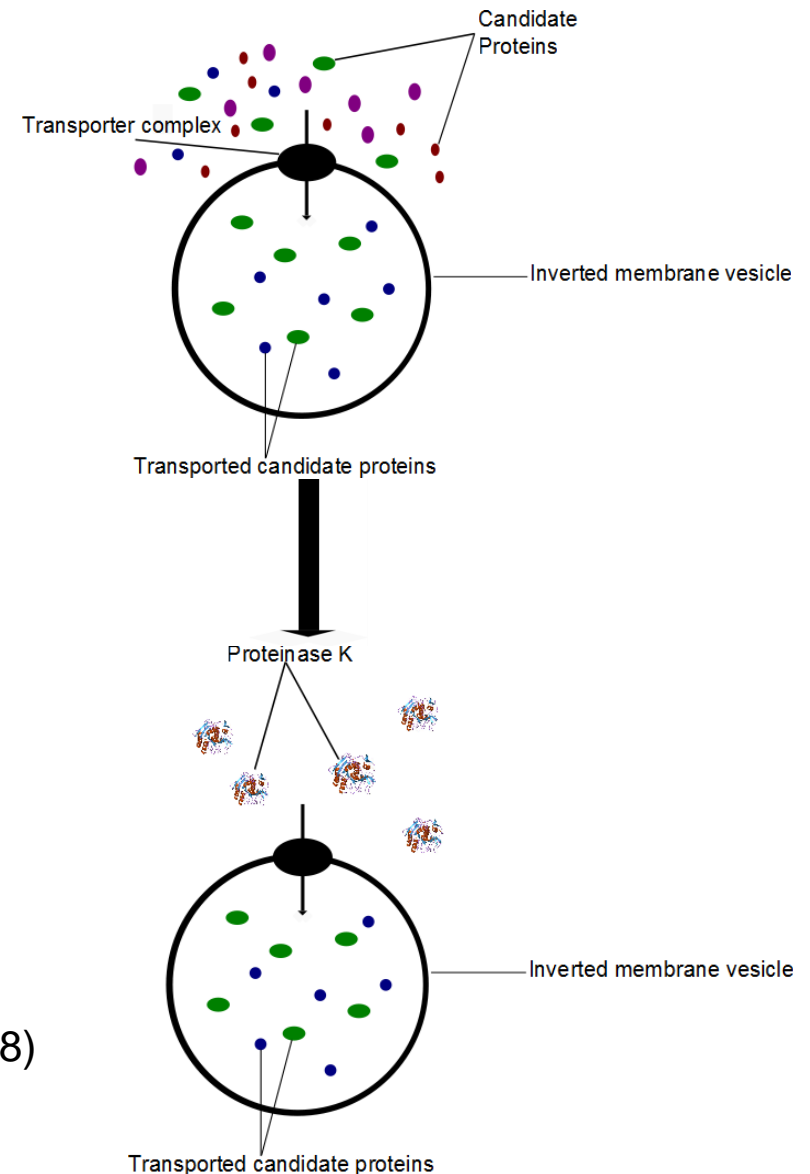


SDS-PAGE diagram

From (Georgia Institute of Technology)

Protection Assay

- Proteinase K treatment
- Lyse liposomes
- Re-isolate and purify candidate proteins
- Controls:
 - Treat candidate proteins with archaeal signal peptidases: Igni153 and Neq432 (Podar *et al.*, 2008a)
 - Trimethylene *N*-oxide reductase (TorA) TAT inhibitor (Chanal *et al.*, 2003)
 - Sec small peptide inhibitors (Li *et al.*, 2008)



Proteinase K protection assay

Nanoarchaea equitans SecDF import

- Problematic S-layer (Ring and Eichler, 2001)
- Isolate and purify SecDF complex (Nouwen *et al.*, 2005)
- Formation of liposomes (Cladera *et al.*, 1997)
- Reconstitution of SecDF complex into liposomes (Nouwen *et al.*, 2005)
- TEM validation
- Proteinase K protection assay
- Controls:
 - Treat candidate proteins with Igni153 and Neq432 (Podar *et al.*, 2008a)
 - Sec small peptide inhibitors (Li *et al.*, 2008)

Conclusions

- Enigmatic relationship
- Genetically intractable organisms
- Potentially important and interesting
- Investigation of TAT and Sec mediated protein exchange between *Nanoarchaeum equitans* and *Ignicoccus* requires:
 - Identification of potential transported proteins
 - Demonstration of transporter localisation to interaction site
 - Demonstration of transporter competence for candidate proteins
- Further work

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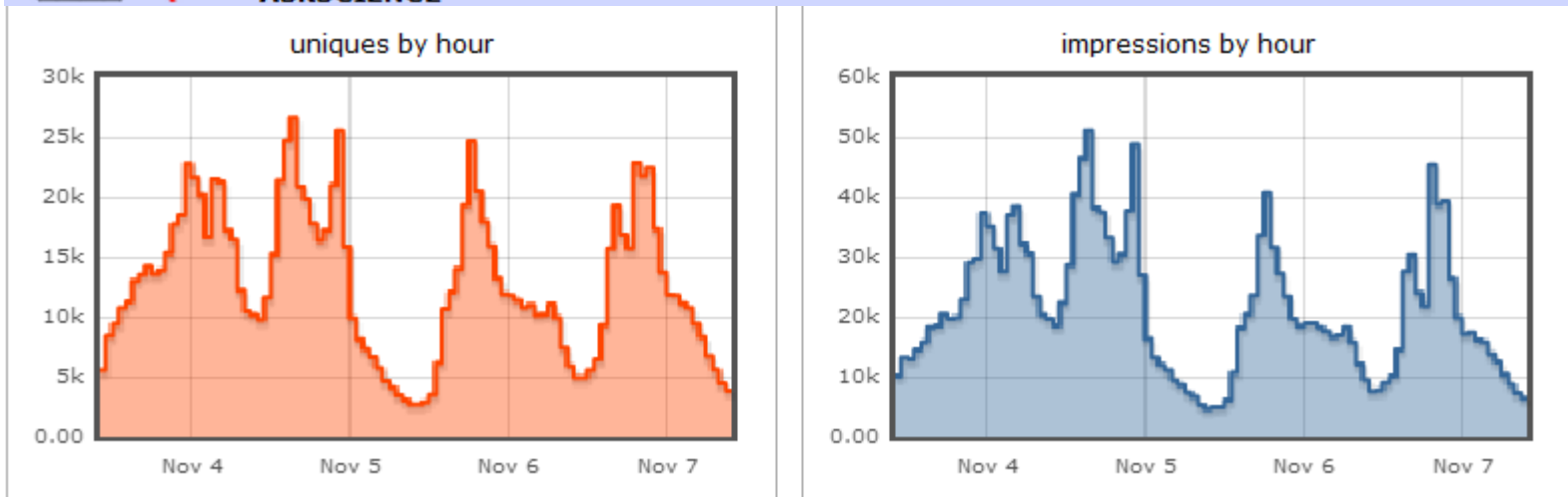
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