Probing the Microbial-Mineral Interface by Neutron Reflectivity

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Outline

Introduction

- Motivation, Scientific Challenges

- Model System
 - Bacterial-Mineral Interface
- Neutron Reflectivity
 - Experimental Challenges
- Complementary Methods
 - Structure, Electrochemistry
- Future Research Perspectives



Mineral respiration by *Geobacter sulfurreducens*

Geobacter sulfurreducens

- gram-negative, anaerobic
- transfers electrons to soluble and insoluble electron acceptors
- Reductive precipitation of contaminant metals and radionuclides
 - − Uranium: U(VI) \rightarrow U(IV)
 - Technetium: $Tc(VII) \rightarrow Tc(IV)$
 - − Chromium: $Cr(VI) \rightarrow Cr(III)$
- Geobacter is predominating in many subsurface environments, also at DOE contamination sites





Microbial fuel cells

- Geobacter sulfurreducens is capable of direct electron transfer to insoluble electron acceptors (Fe₂O₃, MnO₂, Graphite)
- Design of microbial fuel cells to generate electricity



Lovley D.R., Nature Reviews, 4(7), 497-508 (2006).



Scientific Challenges

- Fundamental knowledge gaps exist in the electron transfer reaction mechanisms between microorganisms and sediment mineral surfaces
- Outer membrane cell protein (e.g. cytochrome) function and transport are poorly known

Objectives

- Proof-of-principle experiments to study the molecular interaction of microbial proteins with mineral surfaces with biomimetic lipid model membranes and neutron reflectivity
- Reveal molecular arrangement to understand mechanisms of electron transport and function of bio-macromolecules (proteins and lipids) on mineral surfaces

Hypothesis

 Transport of electrons by cytochromes across the cell membrane, and direct contact with minerals, are necessary for the reduction of minerals such as iron oxide



Cytochromes identified to be involved in Fe(III) reduction

The genome of *Geobacter sulfurreducens* contains **111** c-type cytochrome coding sequences (43 unique) with up to 27 heme moieties.

| MacA | 36 kDa | 2 hemes | associated to inner membrane |
|------|---------|----------|---|
| РрсА | 9.6 kDa | 3 hemes | shuttles electrons between inner and outer membrane |
| OmcB | 86 kDa | 12 hemes | most abundant cytochrome at outer membrane |
| OmcC | 89 kDa | 12 hemes | highly homologous to OmcB |
| OmcE | 30 kDa | 4 hemes | probably involved in transmembrane electron transfer |
| OmcS | 50 kDa | 6 hemes | extracellular surface (LPS/EPS layer) possibly terminal electron donor |



Plasma membrane

Periplasmatic space

Plasma membrane

Methé, B.A. et al., Science 302, 1967 (2003)



Proposed mechanism for extracellular electron transfer to insoluble electron acceptors

Simplified model of the electron transfer system in *Geobacter sulfurreducens*



Gram-negative bacterial cell

Lovley, Nature Reviews Microbiology, 4(7),497-508, 2006 Weber *et al.*, Nature Reviews Microbiology, 4(10), 752-764, 2006



Isolation and Purification of Geobacter Cytochromes

- **PpcA** from wild-type Geobacter sulfurreducens
- 9.6 kDa periplasmatic cytochrome
- 3 hemes with His-His coordination
- Hydrophobic loop
- Positively charged, pl 9.5



Crystal structure of recombinant PpcA



Isolation and Purification of Geobacter Cytochromes

- OmcB from a ∆OmcC mutant of Geobacter sulfurreducens
- 86 kDa outer membrane cytochrome
- 12 hemes with His-His coordination
- Flexible modules

Purified OmcB from *G. sulfurreducens*. Molecular masses at left. Each lane is a fraction from the final purification step (ion exchange on DEAE-Sepharose).



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Probing the Microbial-Mineral Interface by Neutron Reflectivity





Homology model of Geobacter omcB

- 12 heme moieties
- Hydrophobic membrane anchor
- Template structure: hexadecaheme cytochrome Hmc from *Desulfovibrio vulgaris*
- E-value of alignment: 6.5-10⁻¹²
- 3 domains
- Flexible linker

Model created using *Phyre and LOOPP* http://www.sbg.bio.ic.ac.uk/~3dpssm/ http://cbsuapps.tc.cornell.edu/loopp.aspx



Outer membrane cytochrome – OmcS

- 50 kDa
- 6 hemes
- OmcS may be the terminal electron donor transferring electrons to mineral surfaces
- OmcS is associated with lipopolysaccharides in the outer membrane and may also play an essential role in the conductivity of pili



Secondary structure prediction for OmcS



Model system

• Langmuir-Blodgett deposition of Fe₂O₃ nanoparticles



Deposition of hydrophobic iron oxide nanoparticles on silicon from solution by the Langmuir-Blodgett technique on a silicon wafer from the air/water interface.



SEM image of an iron oxide nanoparticle monolayer on silicon.

- Nanoparticle size: 5-60 nm
- Nanocrystalline hematite (Fe₂O₃) as determined by X-ray diffraction
- Surface coverage: ~ 60%



Biomimetic model membranes

- Biomimetic phospholipid model membranes
- Phosphatidylcholine (PC): zwitterionic, net charge: 0
- Phosphatidylglycerol (PG): anionic, net charge: -1





- Uncharged and negatively charged phospholipids to mimic the electrostatic environment of the bacterial membrane
- Interactions with lipid membranes may induce conformational changes in the proteins



Neutron Reflectivity









- Unique properties of neutrons to probe the complex biological-mineral interface
- Contrast variation to resolve model ambiguity
- Production and purification of deuterated proteins in the ORNL deuteration lab



Iron Oxide Nanoparticle Thin Films



 Reflectivity profiles from 25 and 50 µm iron oxide nanoparticle thin films deposited by the Langmuir-Blodgett technique. Data acquired at SNS BL-4B.



Phospholipid model membrane data



- Reflectivity profile of a phospholipid bilayer stack
- Fluid cell setup for studies in excess water



Simulated Neutron Reflectivity profiles



Neutron specular reflectivity from the protein *Geobacter sulfurreducens* OmcB deposited atop the fully deuterated DPPC bilayer from D_2O . A layer with 50% coverage is readily distinguishable from the bare phospholipid bilayer, while 10% coverage is more difficult to resolve and so represents a lower limit.



Complementary structural characterization

- Atomic Force Microscopy to investigate surface properties, roughness
- Spectroscopic methods: Fluorescence Spectroscopy (protein-lipid interactions) and Circular Dichroism (conformational changes)
- Small Angle X-ray and Neutron Scattering with contrast variation to obtain solution structures of single proteins and protein complexes, influence of pH and ions



Complementary methods



Thin layer electrode setup to study electrochemical properties of cytochrome proteins in lipid environments

Option to integrate thin layer electrochemical cell into sample holder for neutron reflectivity, simultaneous measurements

Control redox potential, electron transfer kinetics.



Redox potential of a small monoheme cytochrome c



UV/VIS absorption spectra of oxidized and reduced forms of PpcA



Future research perspectives

- Application of experimental system to study electron transport in Shewanella oneidenis or other microorganisms
- X-ray crystallographic data to obtain atomic resolution structural models
- Novel model systems incorporating Exopolysaccharides (EPS) and Lipopolysaccharides (LPS) from cell extracts





 Interaction of outer membrane cytochromes with oriented LPS layers
Recent progress in characterization of LPS on solid support*

*Abraham et al., JPhysChem B, 111, 2477-2483, 2007



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Single DMPC bilayer on Si



• T. Gutberlet, R. Steitz, G. Fragneto, B. Kloesgen, J. Phys.: Conden. Matter 16 (2004)



Contrast Variation

- Calculated scattering length density (SLD) for Fe₂O₃ in the hematite crystal modification is 7.2 · 10⁻⁶ A⁻². Surface coverage between 60% and 80%:
- SLD for the iron oxide layer between 4.32 • 10⁻⁶ A⁻² and 5.76 • 10⁻⁶ A⁻² corresponding to a D₂O/H₂O ratio of 70 -92%.

