

BIOSORPTION OF ACTINIDES TOWARDS WIPP MICROORGANISMS

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Biosorption of Actinides in the WIPP

The Waste Isolation Pilot Plant (WIPP) transuranic repository remains a cornerstone of the U.S. Department of Energy's (DOE) nuclear waste management effort. Waste disposal operations began at WIPP on March 26, 1999 but a requirement of the repository license is that the WIPP needs to be recertified every five years for its disposal operations. The WIPP is now pursuing its third recertification (submitted in March 2014) and there are many ongoing discussions about the possibility of expanded missions and additional nuclear repository concepts in a salt geology.

Microbial processes have been shown to influence the long-term migration of actinides in a wide variety of subsurface environments [1]. These studies however tend to focus on the near-surface contaminant problems in low ionic-strength groundwater that is present at many DOE sites. The ongoing recertification of the WIPP TRU repository and the broader consideration of salt repository concepts for the permanent disposal of HLW/SF waste has focused attention on the effects of the halophilic microorganisms that are indigenous to salt and high ionic-strength brine systems.

A key and important actinide microbial interaction is the bioassociation of actinide species with microorganisms to form biocolloids. This interaction can potentially contribute to the "mobile" actinide source when physical transport due to intrusion scenarios is the predominant release pathway. If transport pathways are present, bioassociation can lead to reduced mobilization (e.g., if the microbe itself is not mobile) or enhanced mobilization (if the microbe or the degradation "fragments" are mobile). In the self-sealing geology of a salt repository with no interconnected groundwater, only biocolloidal contributions to the source term are considered in PA..

Herein, we report progress in investigating the bioassociation of actinides toward halophilic microorganisms to extend what has been observed in soil bacteria [2-3] to the microorganisms that are typically found in the high ionic-strength brine systems predicted for the WIPP. A number of isolates have been found and both halophilic archaea and bacteria have been identified [4]. The biosorption of actinides towards these microorganisms is being investigated using redox-invariant analogs (e.g., thorium and neodymium) or actinides that can be stabilized in specific oxidation states (e.g., Am³⁺ or NpO₂⁺) since the biosorption observed is specific to the oxidation state of the actinide and the its aqueous speciation. These data are used to define the biocolloid enhancement factors which are used in WIPP PA.

Actinide Biocolloids in Brine

The biocolloidal contribution to dissolved actinide concentrations was re-evaluated using WIPP-indigenous microorganisms as a function of pC_{H+} and brine composition. In almost all cases investigated corption was observed although these were often coupled with hio precipitation at high pH. Estimated biosorption effects were used to develop enhancement factors to determine the biocolloidal contribution to the mobile actinide concentration source term.



Biosorption of Nd³⁺ and Th⁴⁺

Thorium and neodymium were used as the redox-invariant analogs for the An(IV) and An(III) oxidation state to avoid surface moderated redox changes in the biosorption experiments.





Experimental Approach: Biosorption Experiments

GWB Brine

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Overall Approach

- · Simulated WIPP brine, with simplified brine (sodium perchlorate or chloride) with some variability in ionic strengths
- Same-pH mixing of actinide/metal and microorganism (OD₆₀₀ check) · Pre-filtration of actinide/analog to differentiate colloidal contribution (if
- possible) - 1-2 hour experiment duration under constant mixing
- ICP-MS analysis of metal after 100K Da (~ 20 nm) filtration (Amicon
- ultrafilters)
- Followup analysis to confirm oxidation state and status of microbes
 Reversibility, kinetics and viability/toxicity checks

Bacteria Chrom



Model/Developmental Study: Biosorption of Np(V)

Np(V) was used as a model/developmental system since it does not readily hydrolyze and micro-molar concentrations are stable towards precipitation across the pH range of interest.





• For Chromohalobacter sp., titration data at 2 and 4 M have the same shape and are similar what is seen with other organisms at low ionic strength - best-fit model invokes 4 surface complexation sites (as with soil bacteria) $_{\odot}$ Differences observed in sorption at 2 and 4M over $pC_{H^{+}}$ range

- \circ Model invoking carbonate complexation reactions fits best at higher pC_{H-} For *Halobacterium noricense*, much lower sorption was noted for the same biomass that likely reflects a different functional group distribution than observed for the bacteria

Summary of Observations: Nd and Th Results

- Np(V) at 5 g/L
- Archaea: ~ 10% sorption
- Bacteria: 70-85% sorption

Nd(III) - at 10 g/L

- Archaea: ~ 25% sorption
- Bacteria: 50-65% sorption
- · EDTA reduced sorption in both cases by ~ 40%
- Th(IV) at 10 g/L
 - Archaea: ~ 10-20% (no EDTA) and ~0% (EDTA) sorption
- Bacteria: ~ 80% (no EDTA) and 57% (EDTA) sorption
- At pC_{H+} > 9, cannot differentiate between sorption and precipitation, due to colloids and "induced" precipitation

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