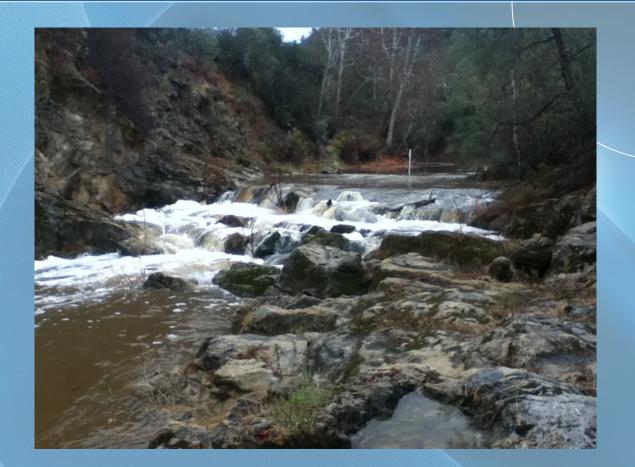
Unexpected Relationships between Methylmercury Enrichment in Fresh Waterbodies and Food-Web Uptake

Steve Dent PhD, Eric Blischke, Andy Greazel PG- CDM Smith; John Hillenbrand - USEPA





American Society of Mining and Reclamation Spokane WA

June 8, 2016





Overview

- Background: Mercury in the Environment
 - Mercury Advisories
 - Food-Web Interactions
 - Trace Mercury Assessments
- Site Introduction
- Small Reservoir Mercury Methylation Assessment
- Summary
- Implications for Future Mercury Assessments
- Potential Remedial Options







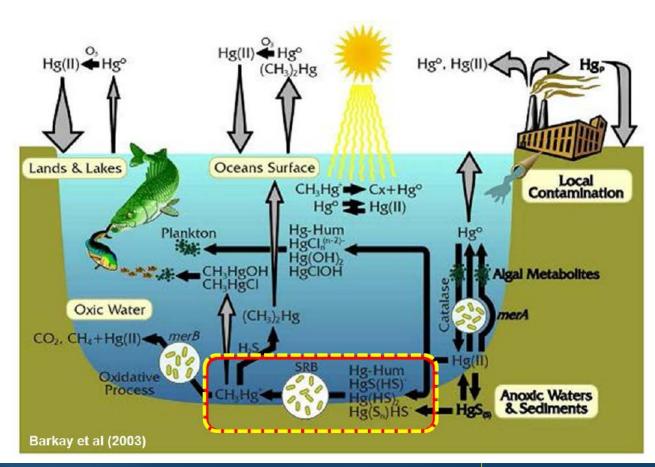
Background: Mercury Advisories in US



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Mercury Transformations in the Environment

Hg Biogeochemical Cycle



Key Methylation Ingredients

- Anoxia
- Organic Carbon
- Sulfate/Ferric Iron
- And last, but not least,
 <u>Labile</u>
 <u>Inorganic</u>
 <u>Mercury</u>





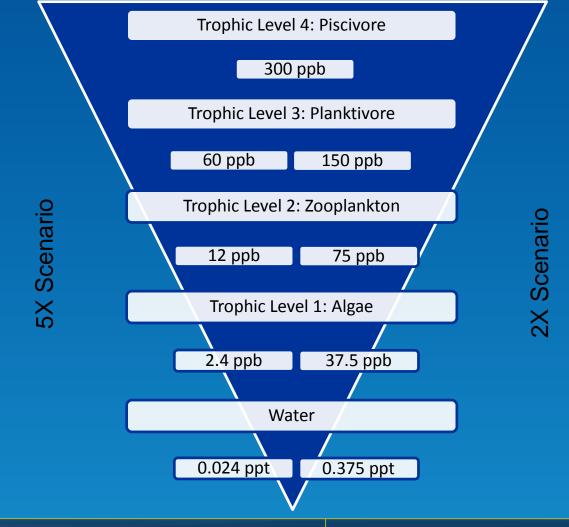
Food-Web Biomagnification



Background: Mercury in the Environment



Trace Mercury is Important – And the Numbers are Real





Smith

Background: Mercury in the Environment



Method 1630 Trace Methylmercury & 1631 Trace Total Mercury

Detection Limits (Brooks Rand Instruments)
THg 1631: <0.03 ng/L
MeHg 1630: <0.002 ng/L

Trace Mercury Sampling

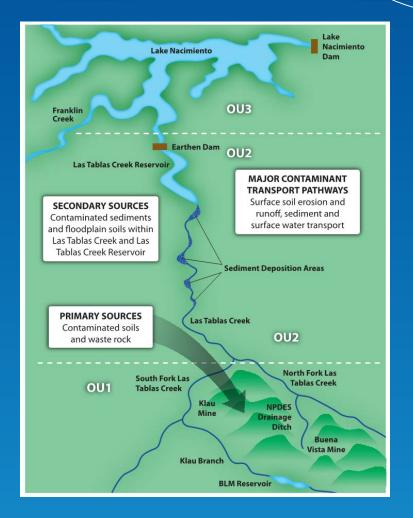
EPA Method 1669
Clean Hands/Dirty Hands
Trace Clean Bottles and

Preservative



Site Introduction - Conceptual Site Model

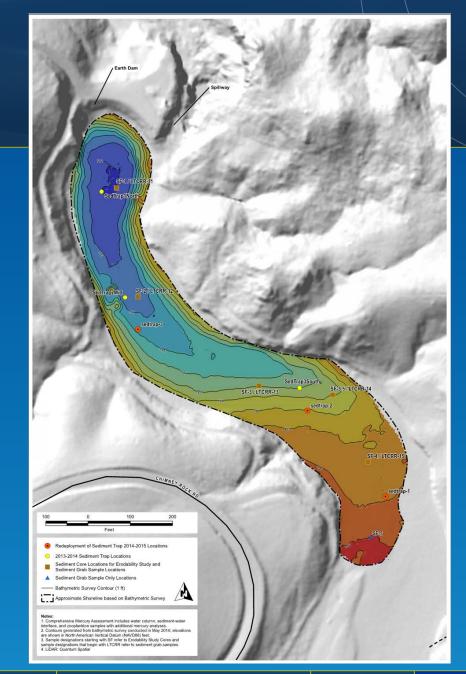
- Mercury mining and processing activities at the Klau and Buena Vista Mines Superfund site has resulted in mercury contamination throughout the watershed
- Contaminant transport is dominated by particulate transport during winter precipitation events
- Mercury is methylated in Las Tablas Creek Ranch Reservoir (LTCRR) sediments and enters food-web
- Loading assessment revealed LTCRR a net source to Lake Nacimiento





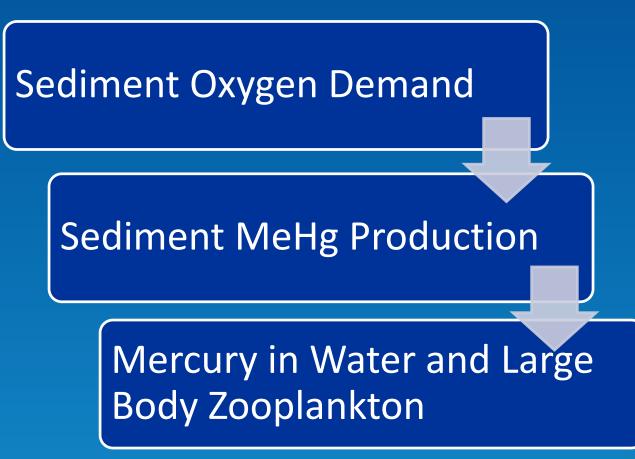
Las Tablas Creek Ranch Reservoir

- Small Reservoir
 - ~52 acre feet
- Fed by Intermittent Stream
 - Storm Flows >10X Reservoir
 Volume Observed
- Shallow and Warm
 - 8 to 13 ft @ 50 to 70 °F
- Net Mercury Source
 - Annual THg Loading Increased
 3x to 4x
 - Fish Body Burden: 2.3 PPM





Methylation and Food-Web Connectivity Evaluation



Small Reservoir Mercury Methylation Assessment

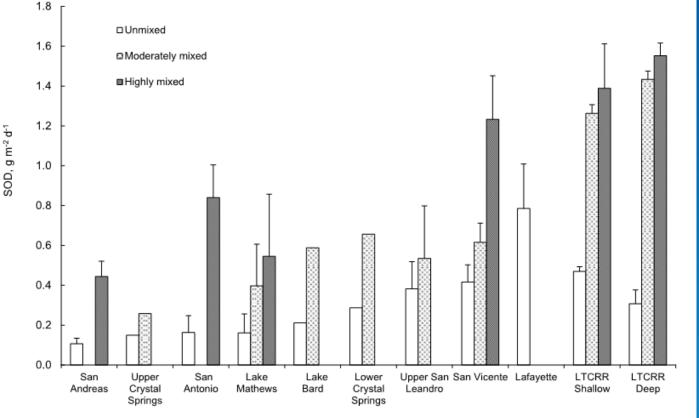


CDM Smith

Las Tablas Creek Ranch Reservoir Sediment Oxygen Demand







Beutel, 2003

Small Reservoir Mercury Methylation Assessment

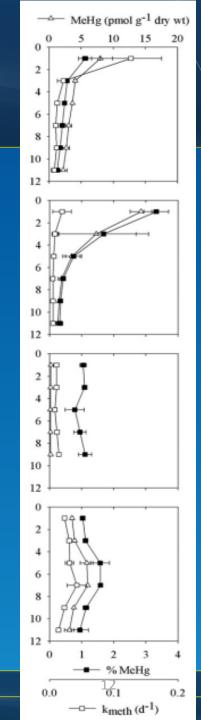


Methylmercury Production

- Surrogate for MeHg production
 Windham-Myers et al., 2009
 - %MeHg of THg in sediment
- Sites with highest surface methylation also have highest fish concentrations Benoit et al., 2003
- Growing database in literature to use for comparison



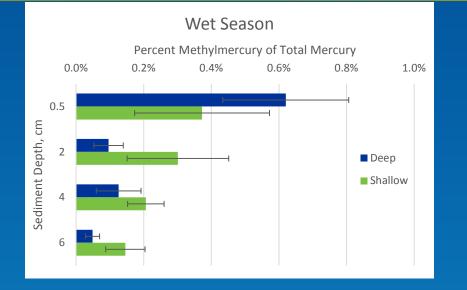
Small Reservoir Mercury Methylation Assessment

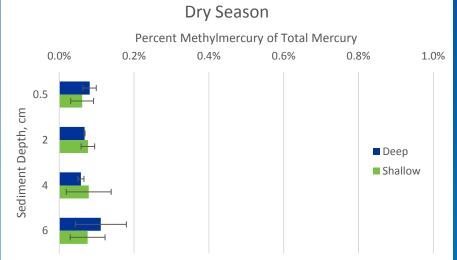


Hollweg, et al. 2009



Sediment Methylation and Bottom Water Enrichment





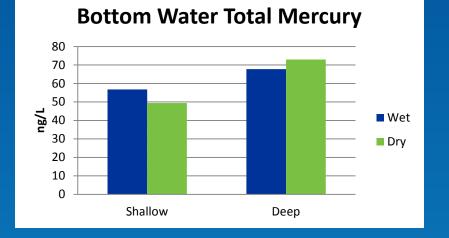
Highest Surface Methylation in Deep Site in Wet Season

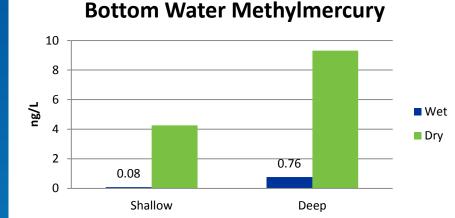
- Highest Sub-Surface Methylation in Shallow Site in Wet Season
- Methylmercury Essentially Shut Down in Sediments During Dry Season

Small Reservoir Mercury Methylation Assessment



Bottom Water Enrichment

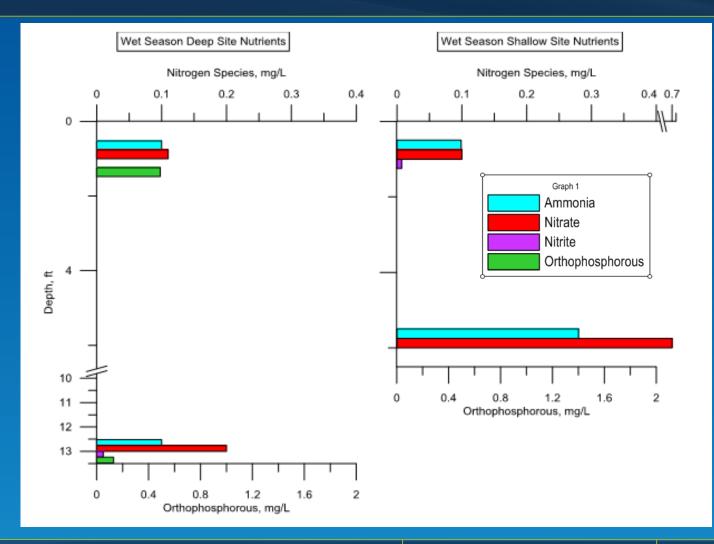




- Small Variability in Total Mercury Between Seasons
- Methylmercury Highest in Dry Season
 - Inverse to surface methylation observation
- Contributing Factors: In Water Methylation, No Flow Through, Incomplete Picture of Wet Season Dynamics



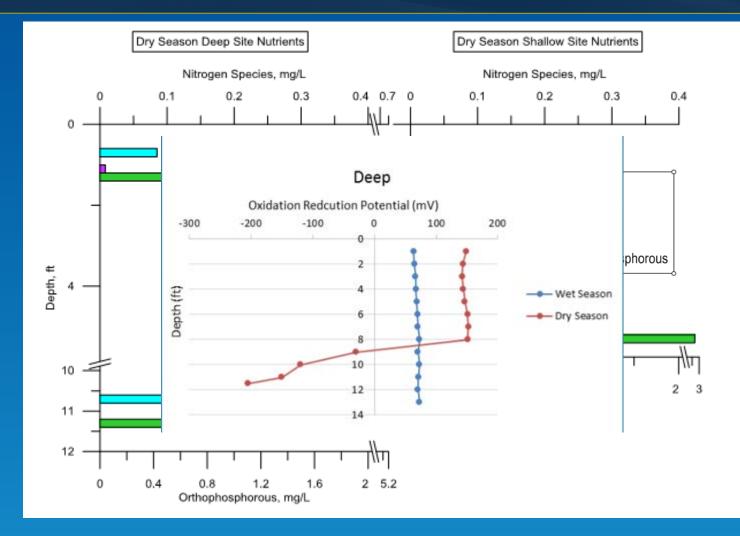
Reservoir Nutrient Dynamics – Wet Season



Small Reservoir Mercury Methylation Assessment



Reservoir Nutrient Dynamics – Dry Season



Small Reservoir Mercury Methylation Assessment



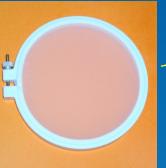
Zooplankton Body Burden/Enumeration

Large body > 243 um

Enumeration/ Speciation







DI Rinse

THg Analysis: EPA Method 7473 MeHg Analysis: EPA Method 1630





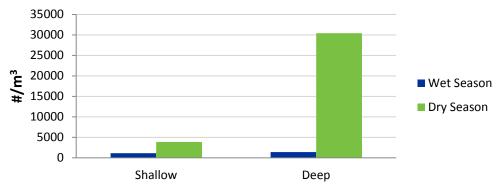




Zooplankton Body Burden: Unexpected Patterns

Wet Season Body Burden: **Dry Season Body Burden: MeHg ~20%** 30% 1800 1600 1400 1200 ng/g 1000 METHYL MERCURY 800 MERCURY 600 400 200 0 Shallow Shallow Deep Deep





Potential Factors

0

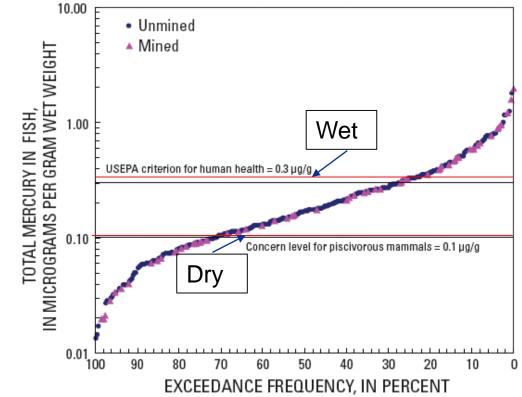
- Dissolved Organic Carbon Competition
- Algea Density
 - Secchi Wet: ~4 feet
 - Secchi Dry: ~3.5 feet





How Does Our Site Compare to Others: Fish Body Burden in US Basins

- 367 stream sites sampled across United States
- Sites with fish greater than 0.3 μg/g
 - 25% Exceed Target
- Sites with fish greater than 0.6 μg/g
 - 10% Exceed Target



USGS, 2009

Small Reservoir Mercury Methylation Assessment

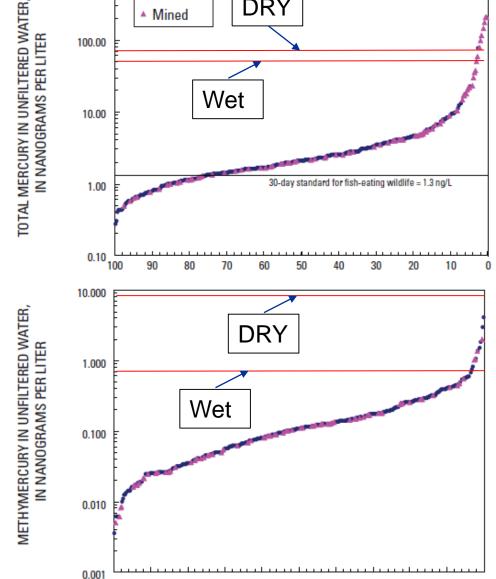




Total Mercury and Methylmercury Surface Water In US Basins

- LTCRR on High End of **Mercury Impacted** Systems in US
- Variability Between Seasons
 - Small THg Change
 - Large MeHg Change
- **Remember This???** •

0.024 ppt



DRY

1,000.00

100.00

Unmined

Mined

90

Small Reservoir Mercury Methylation Assessment

0.375 ppt



50

EXCEEDANCE FREQUENCY, IN PERCENT

60

70



10

Summary and Conclusions – Reservoir Assessment Summary

- LTCRR sediments consume oxygen rapidly and maintains conditions conducive to reduction in both wet and dry seasons
 - Elevated external nitrate loading in wet season
 - Elevated internal phosphorous and ammonia loading in dry season
- Inverse relationship between mercury methylation in sediment and methylmercury bottom water enrichment
- Inverse relationship between methylmercury enrichment water column relative to food-web uptake



Future Assessments and Technology Screening

- **Recommended Additions to Future LTCRR Assessments**
- DOC Analysis
 - UV 254: Aromatic Fraction of DOC
- Algae Enumeration
- Wet Season Sample Timing
 - Target a post storm event with longer antecedent dry condition ~mininum 2 to 4 weeks
- Dry Season Sample Timing
 - Target a sampling event at the end of the dry season prior to the first flush of the wet season.



Potential Remedial Options

Source Control:

- Reduce load of mercury from both the mine site and watershed.
- Reduce load of nutrients from watershed.
- Select In-Situ Remedial Options:
 - Dredging
 - Capping
 - Redox Controls
 - Aeration/Oxygenation
 - Nitrate Addition
 - Coagulation/Precipitation
 - Biomanipulation



Questions?



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Questions