

The Biotic Ligand Model: Unresolved Scientific Issues and Site- and Species-specific Effects on Predicted Cu Toxicity

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EPA Hardrock Mining Conference 2012:
Advancing Solutions for a New Legacy
Denver, CO
April 4, 2012



Background

- The Biotic Ligand Model (BLM) is used to evaluate the site-specific toxicity of copper to aquatic organisms
 - Can be used to develop site-specific water quality criteria (EPA, 2007)
 - Ongoing investigations into different aspects of the Cu-BLM: geochemical, biological
- Current research: quantifying Cu-organic carbon complexation in low hardness waters and subsequent implications for predicting fish toxicity using the BLM



Presentation Outline

- Overview of BLM
- Site-specific Cu-binding studies and metal-DOM binding
- Cu toxicity in low-hardness waters
- Approaches to incorporating Cu binding constants of “biotic ligands” into BLM

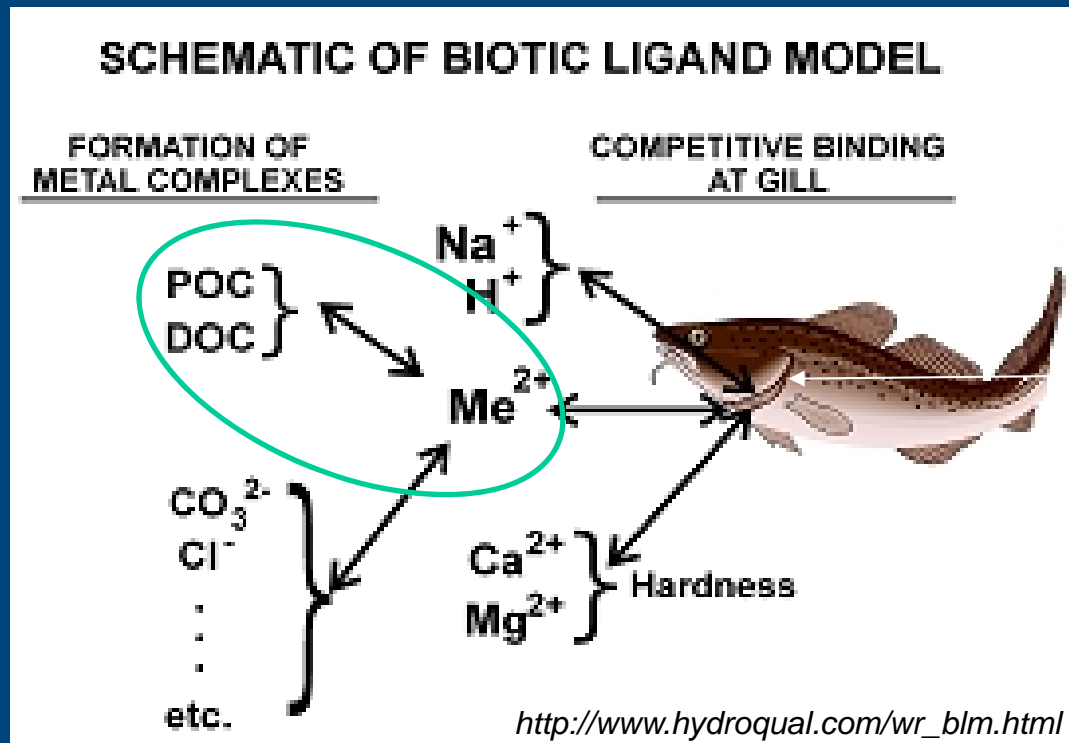


BLM: Background

- Water quality criteria for Cu (and many other metals) expressed as a function of hardness.
 - Increased hardness => decreased toxicity => higher WQC
 - Observed in many controlled experiments
- Well understood that Cu toxicity to aquatic biota is affected by other constituents in water
 - Dissolved organic carbon has been found to reduce Cu toxicity
- BLM developed to numerically address the influence of multiple chemical factors on Cu toxicity

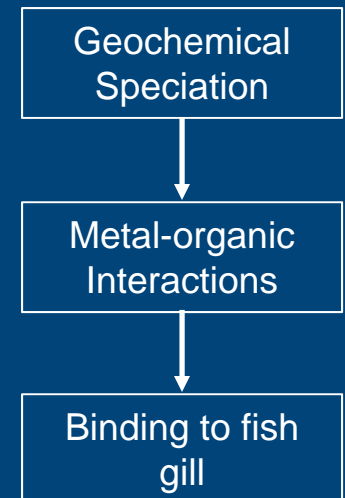
BLM: Conceptual Model

Cu speciation/sorption to gill binding sites (“biotic ligand”) affects bioavailability and toxicity



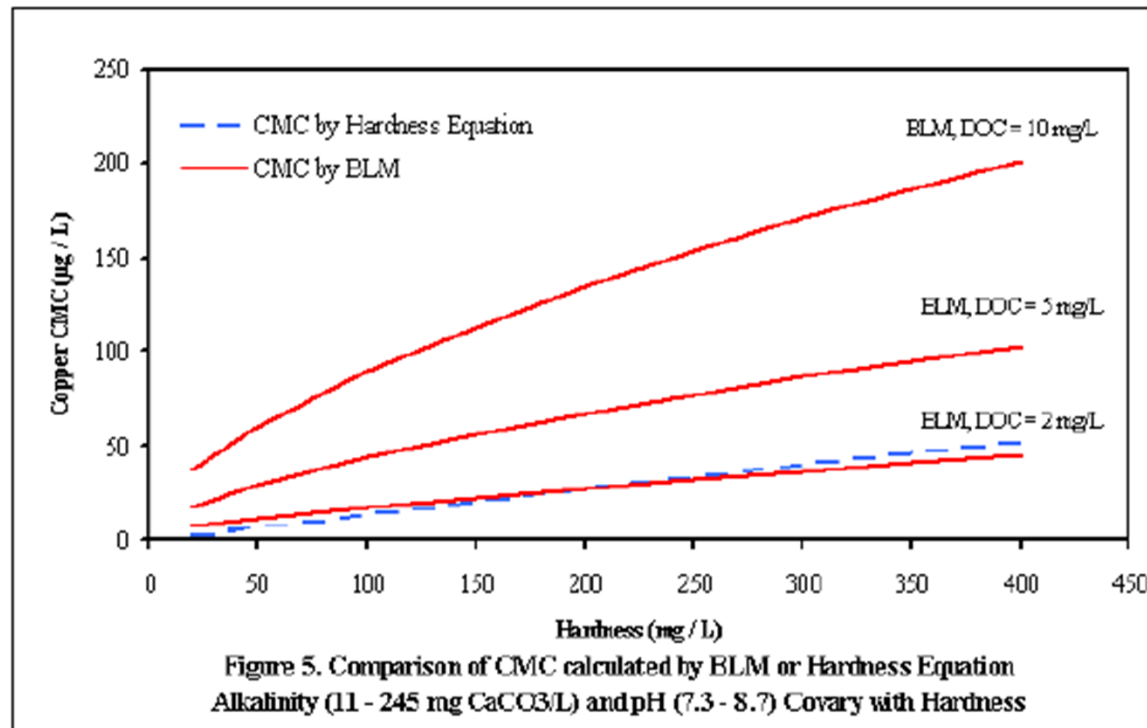
BLM: Conceptual Model (cont.)

- BLM: predict concentration of dissolved Cu that would cause toxicity to aquatic biota over a range of water quality conditions
 - BLM uses “lethal accumulation” on gill to estimate toxicity
- Three elements of model
 - Geochemical speciation code CHES (Santore and Driscoll, 1995)
 - Calculates inorganic metal speciation
 - WHAM V model (Tipping, 1994)
 - Calculates degree of metal-organic interaction
 - Biotic ligand (e.g., fish gill) binding constant (Di Toro et al., 2001)



BLM Illustration: Acute WQC in the Presence of DOC

Site-water chemistry parameters are needed to evaluate a criterion. This is analogous to the situation that previously existed for the hardness-based WQC, where a hardness concentration was necessary in order to derive a criterion. Examples of CMC calculations at various water chemistry conditions are presented in Figure 5 and Appendix G.



Evaluating Cu-Organic Complexation in a Low-hardness Stream



Site-specific Cu Binding Studies

- Purpose: Evaluate Cu binding properties of ambient DOM
- Performed laboratory studies of site-specific Cu binding in low-hardness waters
 - Finding: Stream DOM had less ability to complex Cu than calculated by the BLM



Methods

- Isolated DOM from three low hardness headwater streams in AK
- Cu-ISE titration
 - Fit to a 2-ligand model
- CLE-SPE (competitive ligand exchange-solid phase extraction)
 - Environmentally relevant [Cu]
- Used MINTEQA2 and empirically derived “effective log K” to estimate free Cu^{2+}
- Compared to BLM free Cu

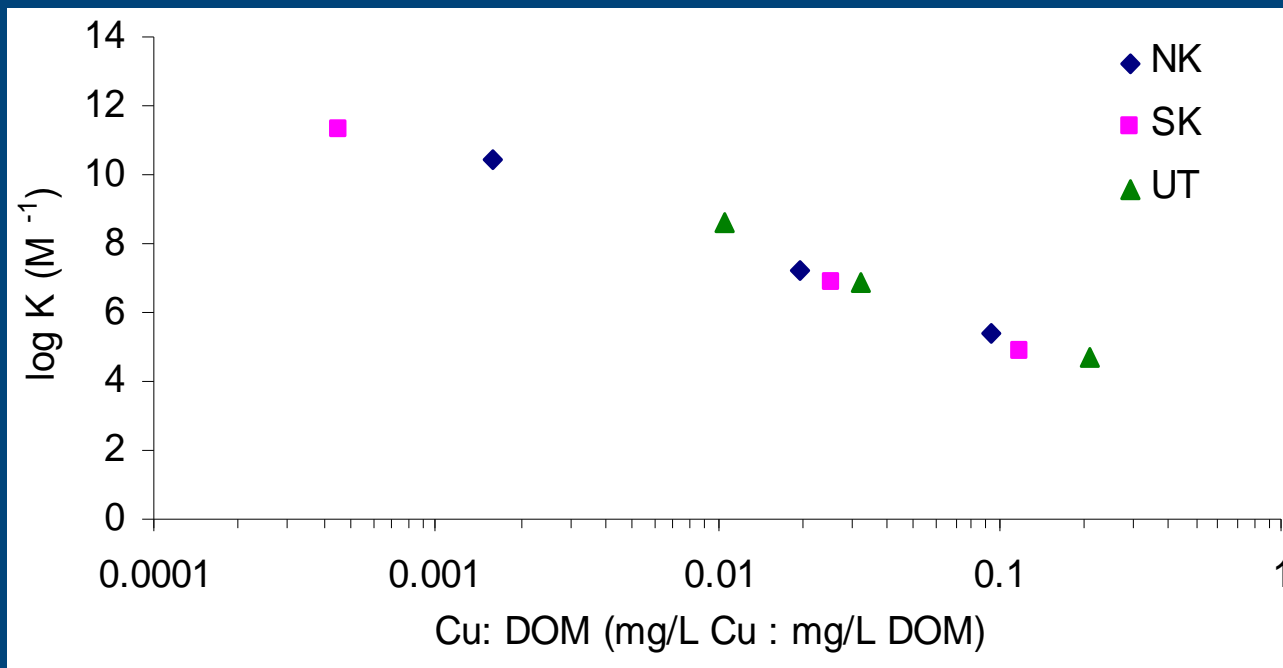


Ambient Water Quality

- pH: 7.1–7.6
- Alkalinity: 13.5–33.9 mg/L as CaCO₃
- Hardness: 13.4–28.4 mg/L as CaCO₃
- Dissolved Cu: 0.2–1.3 µg/L
- DOC: 1.3–2.2 mg/L

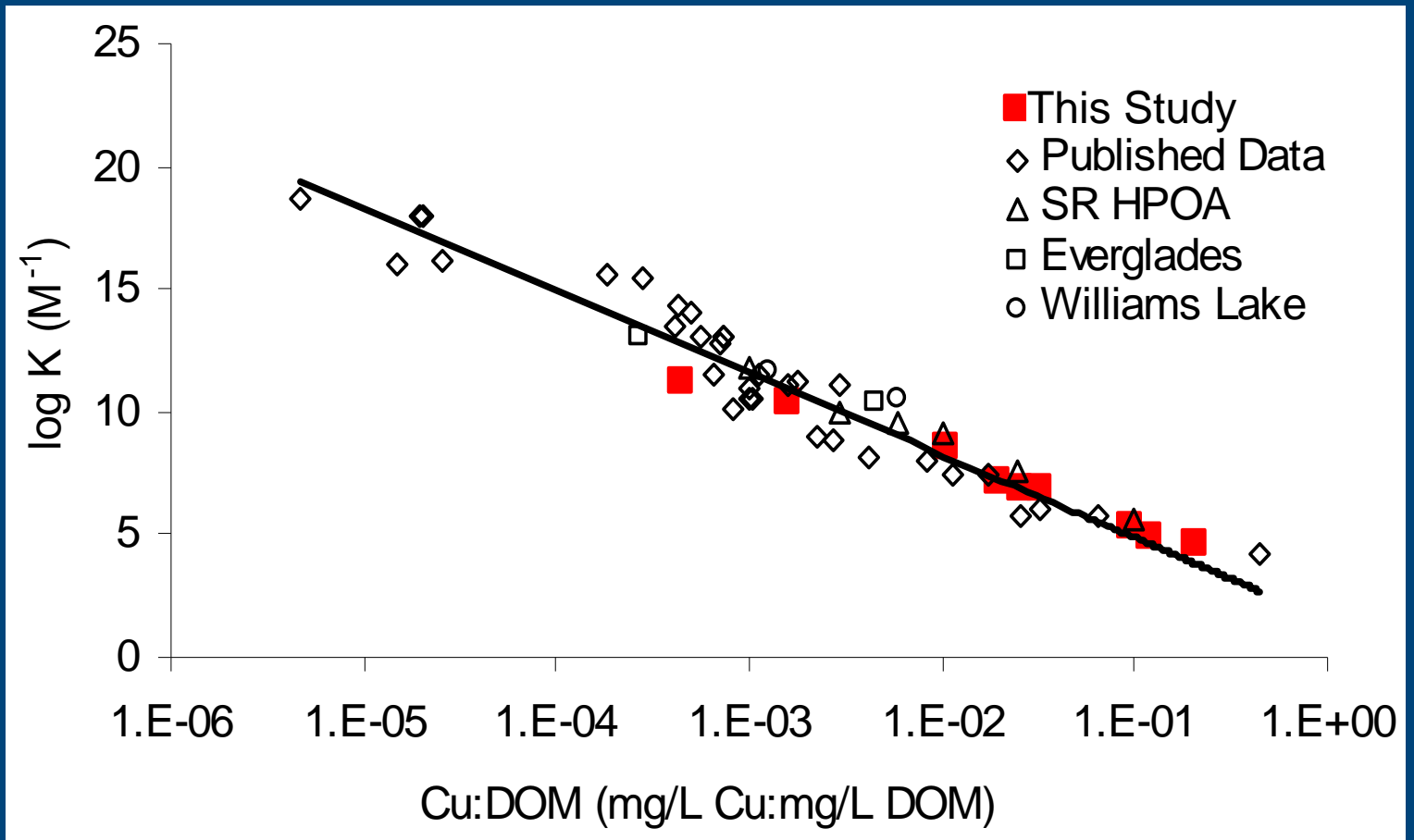
Results: Titration and CLE-SPE

“Effective log K” (net Cu complexation) of site waters
a function of Cu:DOM ratio



Increasing
Cu relative
to ambient
DOM
results in
lower log K

Comparison with Other Studies

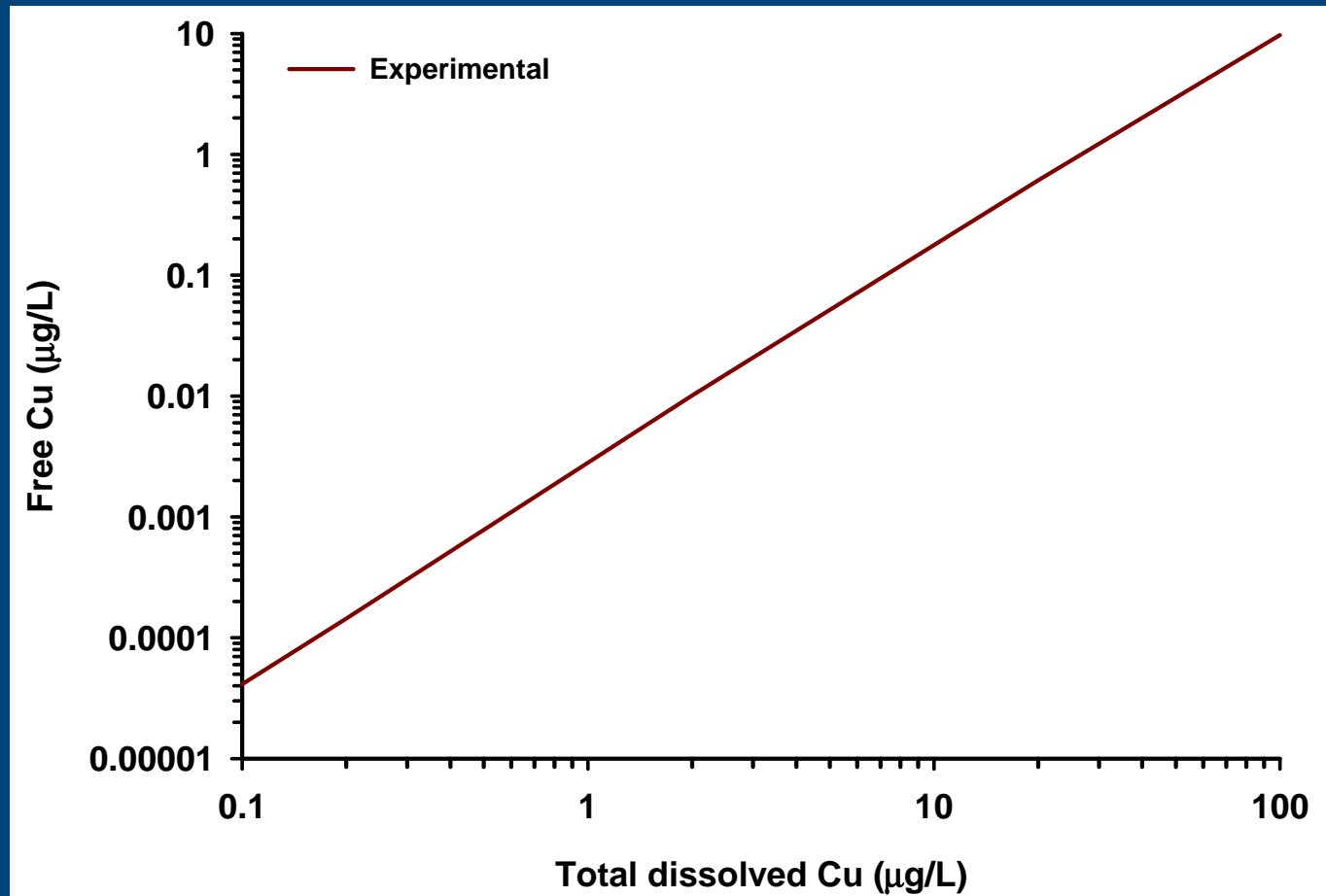




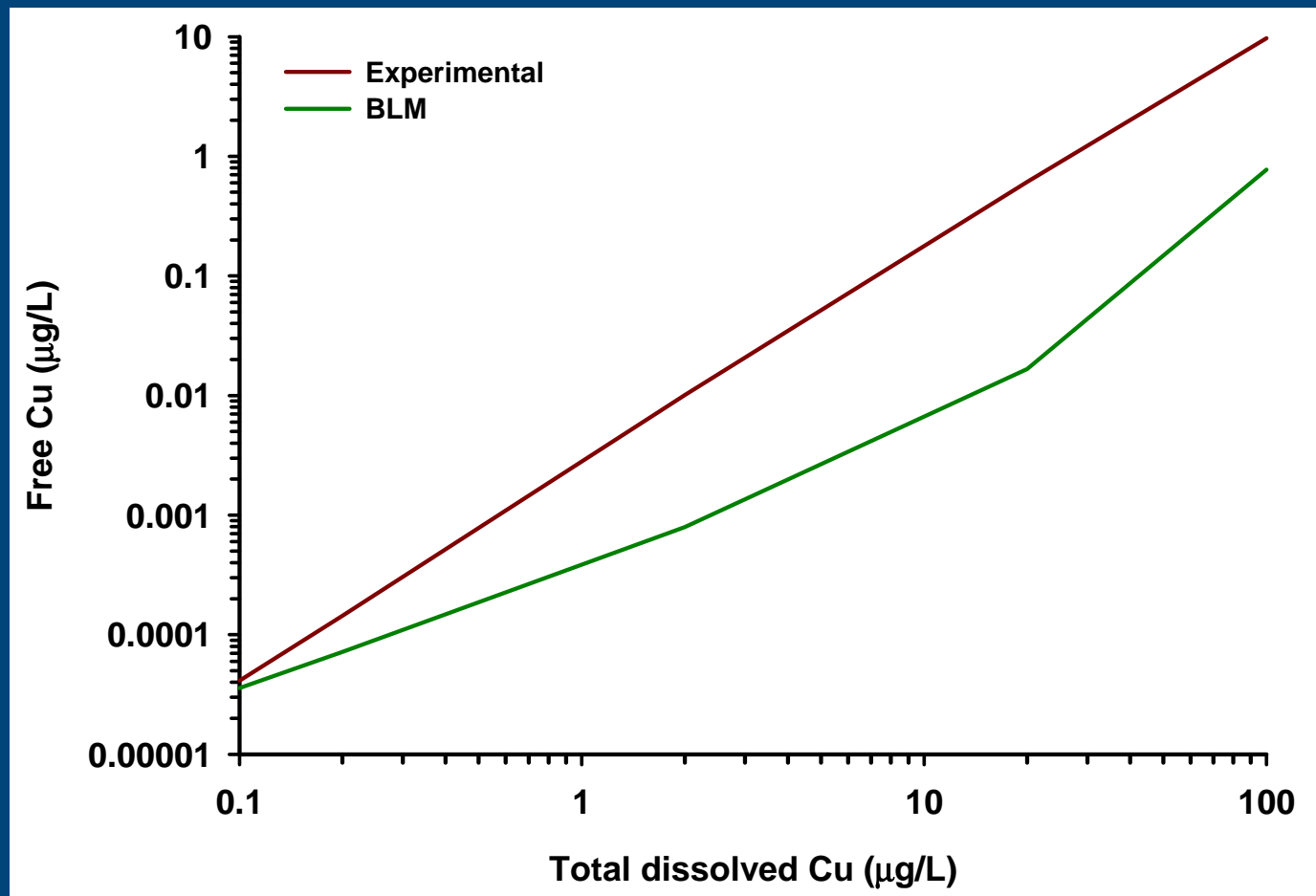
Site-Specific Cu Binding Summary

- Cu-organic binding a function of relative amounts of Cu and DOM present – net affinity changes as more Cu is added
 - Distribution of binding sites in DOM
 - High affinity (high log K) sites less abundant than lower affinity sites
 - As Cu concentrations increase, progressive shift to binding with lower affinity sites
- Cu:DOM ratio is important in predicting complexation

Modeling Free Cu: Empirical Data



Modeling Free Cu: Comparison to BLM

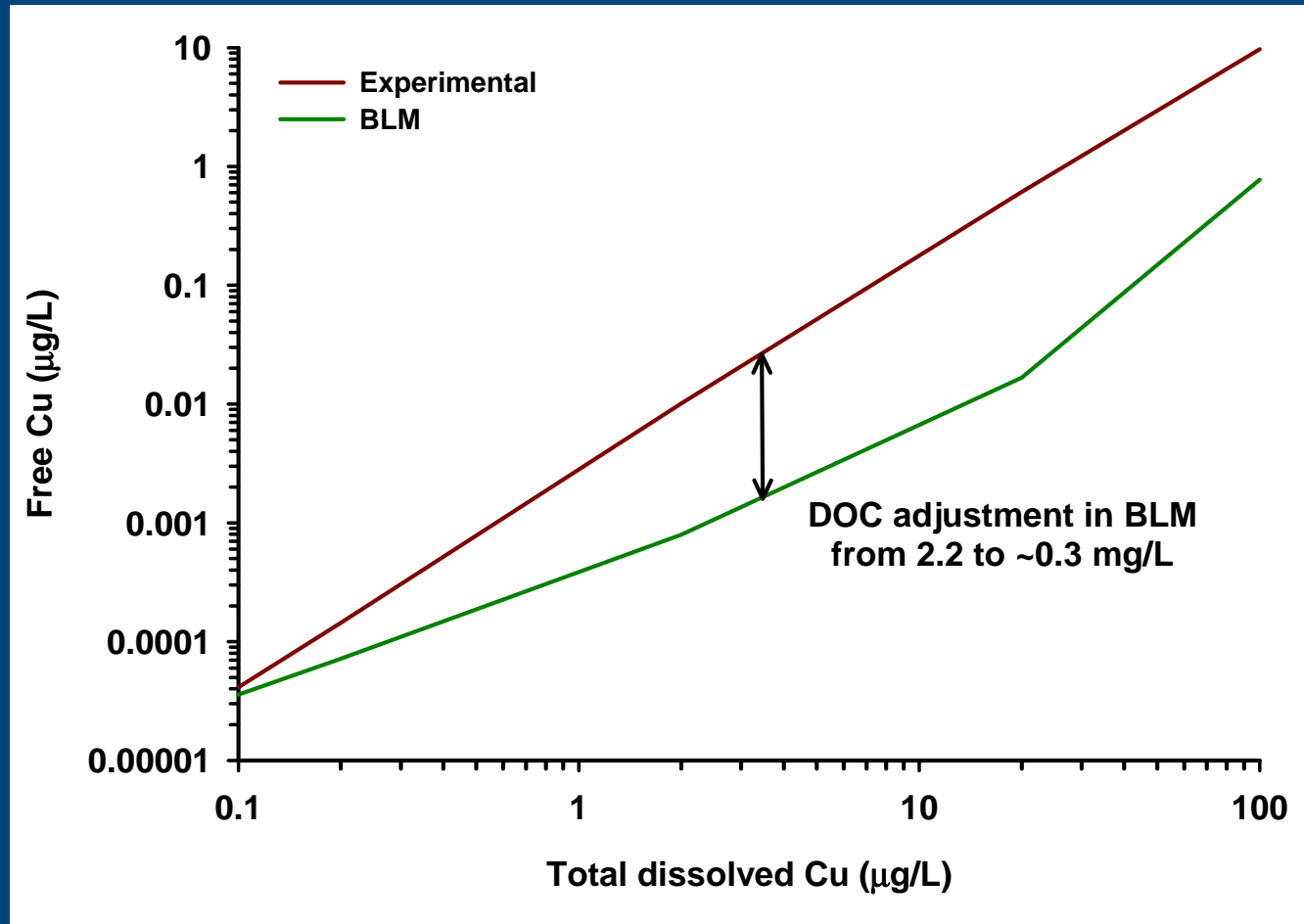




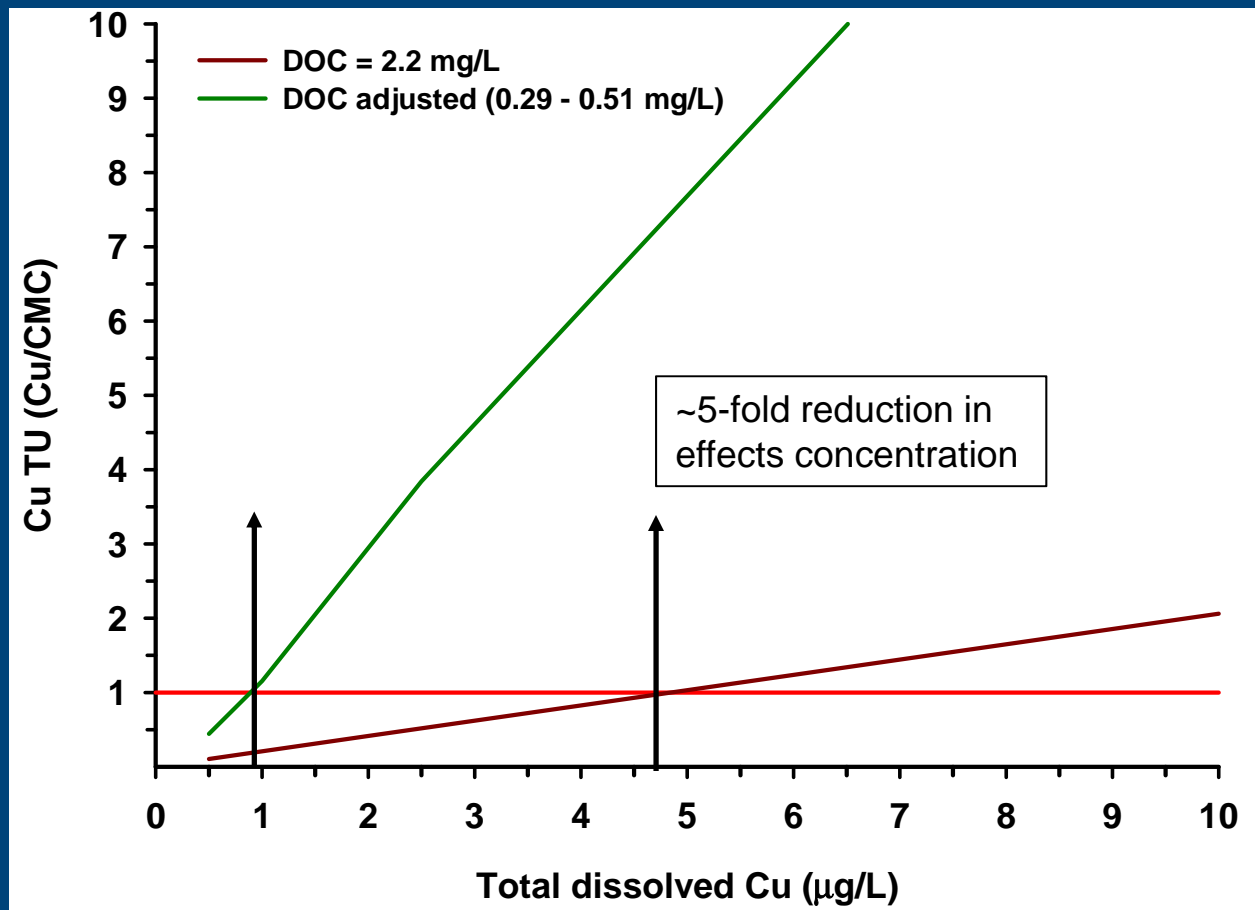
Adjusting DOC Concentrations in BLM to “Match” Empirical Data

- Previous authors (De Schamphelaere et al., 2004; Welsh et al., 2008) proposed adjusting DOC concentration (input to BLM) to match Cu-DOC complexation toxicity results
 - Adjustment factor of 2 used
- This study: adjust [DOC] from 2.2 mg/L to approx. 0.3 mg/L to match experimental data
 - Adjustment factor of approximately 8

Adjusting DOC Concentrations



Implications: Estimating Cu Toxicity with Adjusted DOC





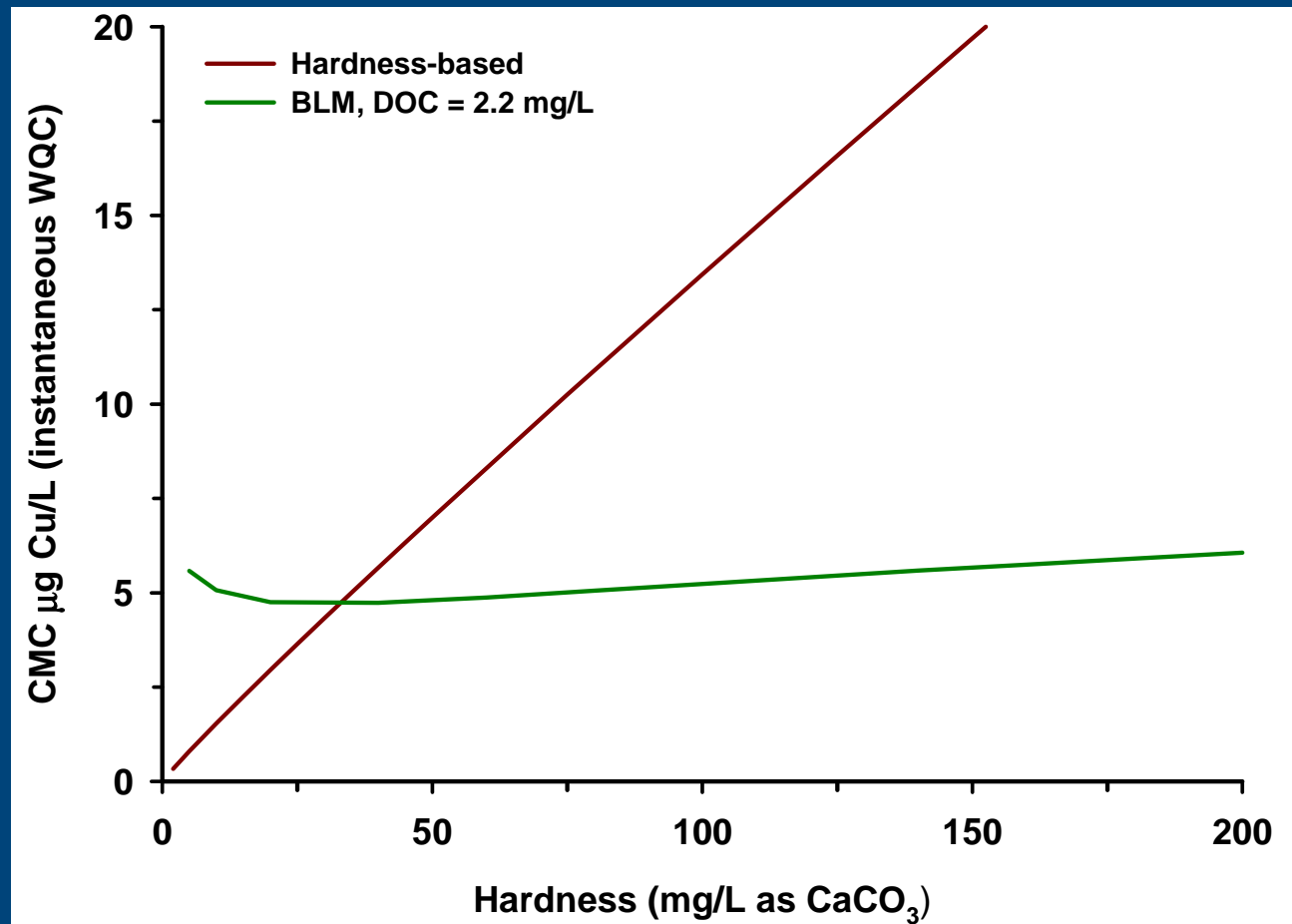
Summary of Cu Binding Results

- BLM under-predicted free Cu compared to site-specific estimates
- Needed to lower DOC in BLM to attain same free Cu results – similar findings to other researchers (e.g., De Schamphelaere et al., 2004; Welsh et al., 2008), but somewhat greater magnitude of adjustment
- Results in a ~ 5-fold decrease in instantaneous WQC compared to BLM

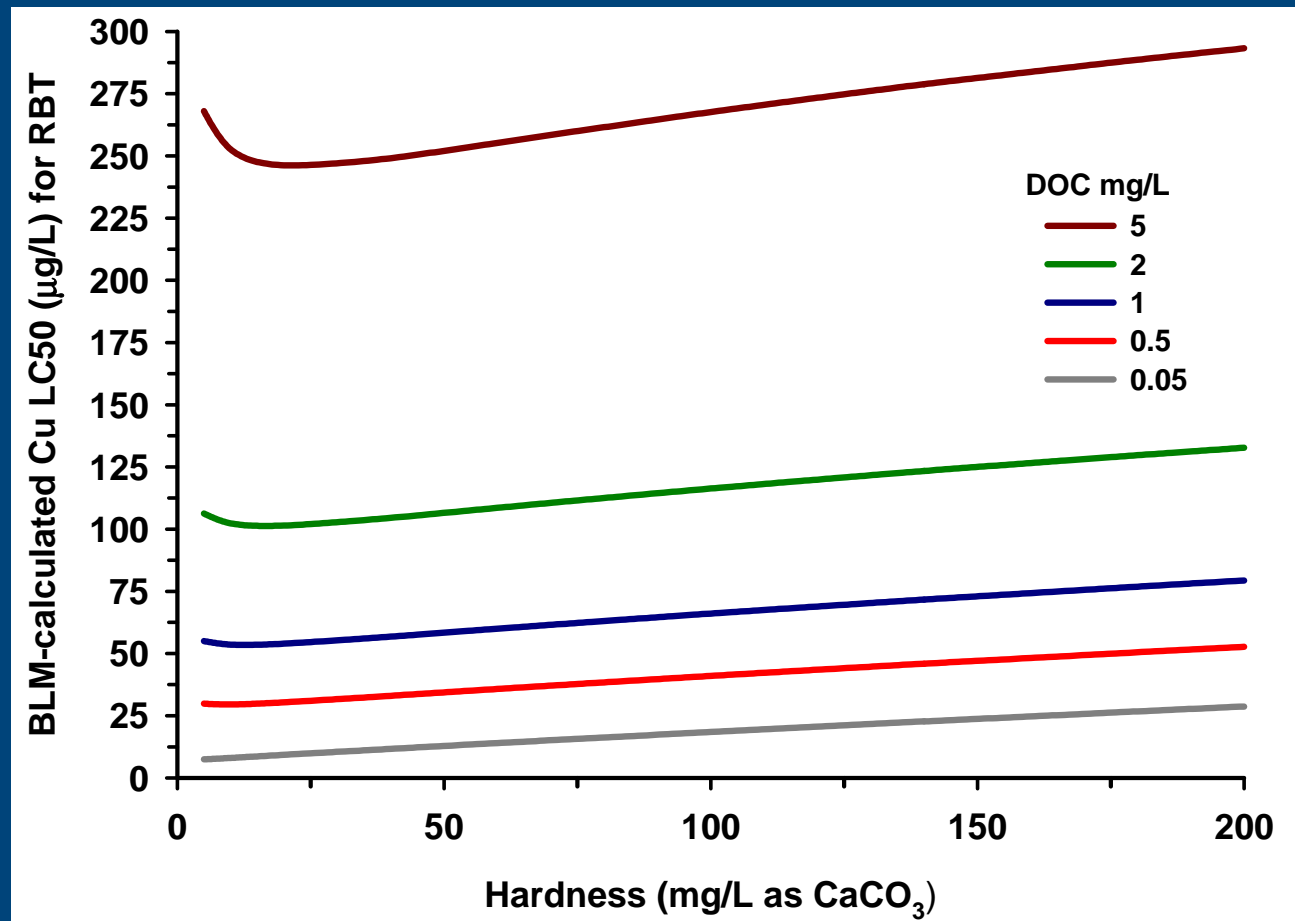
Other Issues: Modeling Cu in Low Hardness Waters?

- Ran series of BLM simulations to further evaluate implications of Cu-DOC complexation in low hardness waters
- Used site-water data as base water quality
 - Temperature = 19°C
 - pH = 7.13
 - DOC = 2.17 mg/L (HA = 10%)
 - Ca, Mg = 4.09, 1.1 mg/L (hardness = 14.7 mg/L CaCO₃)
 - K = 0.1 mg/L
 - SO₄ = 1.7 mg/L
 - Cl = 0.5 mg/L
 - Alkalinity = 22.3 mg/L CaCO₃
 - S = 0.001 mg/L (default, non-functional)

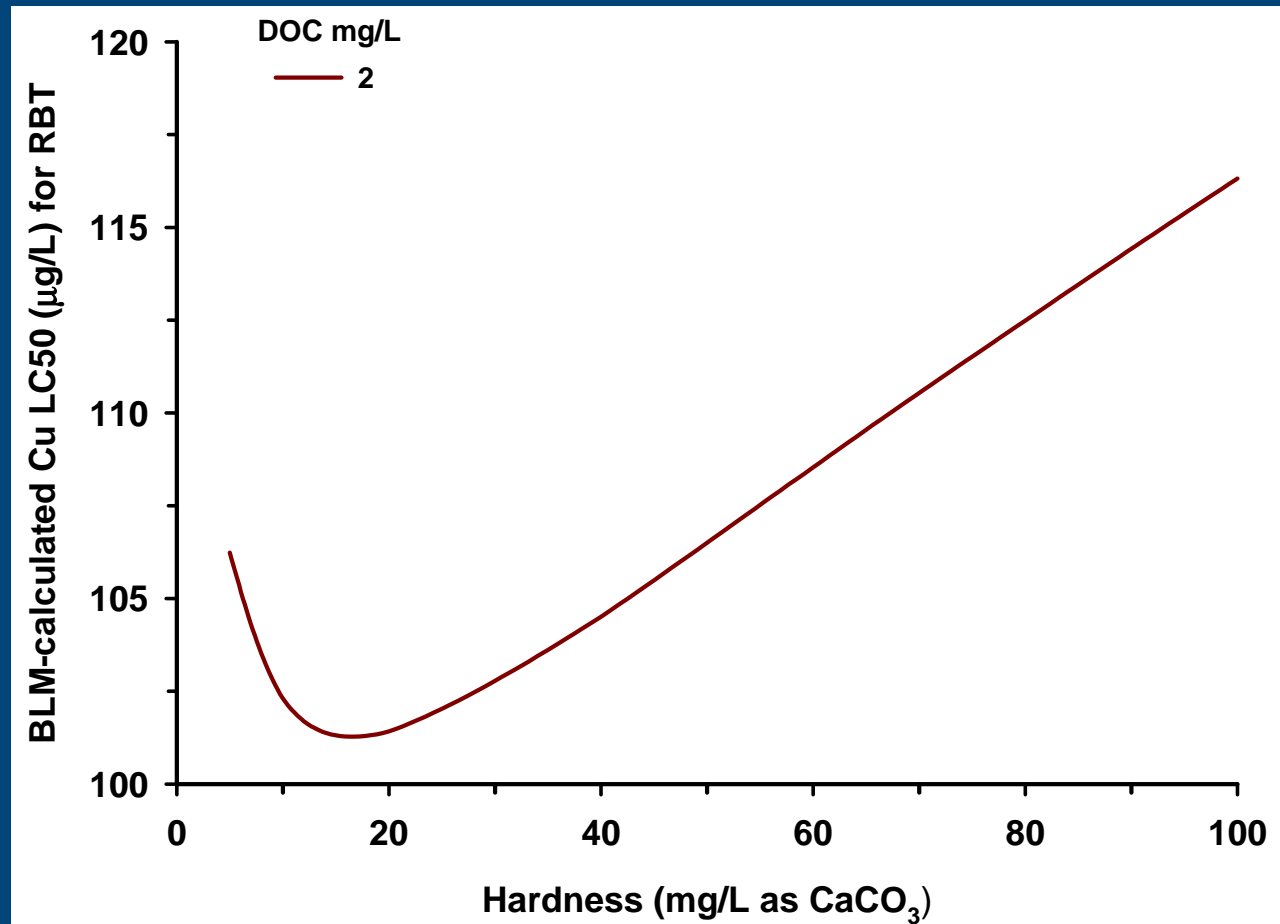
Simulation Results: Varying Hardness; Unadjusted DOC



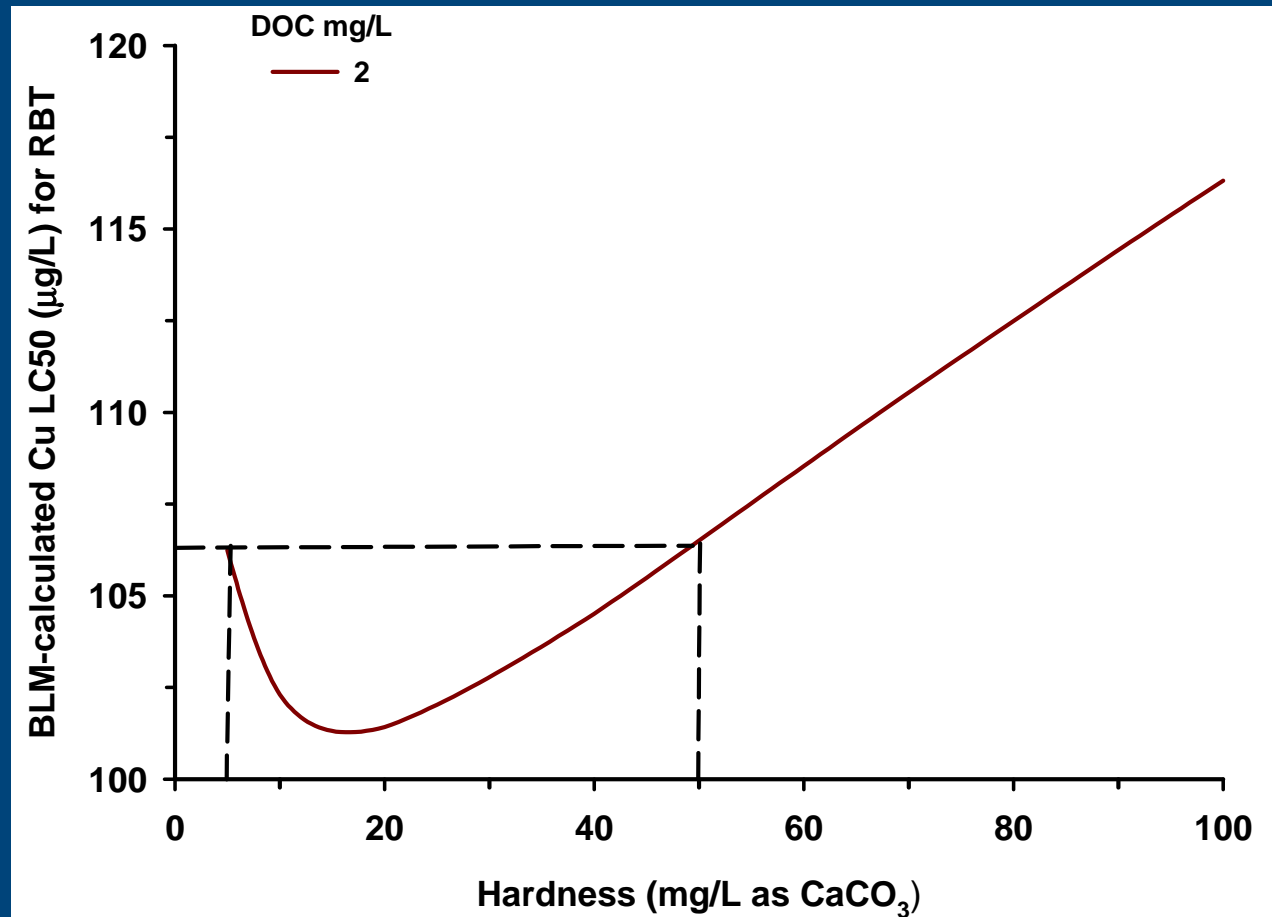
Simulation Results: Rainbow Trout LC50 Varying Hardness and DOC



Hardness Simulation: Artifact of DOC Complexation?



Equivalent LC50, 10-fold Difference in Hardness





BLM Simulations: Summary

- Outputs at low hardness in BLM suggests Cu preferentially bound to DOC rather than the biotic ligand (gill)
- BLM may under-predict toxicity of Cu because of DOC complexation (log K data)
- Degree of under-predicted toxicity of Cu may be exacerbated in soft water



Predicting Cu Toxicity: Implications of Biotic Ligand Component

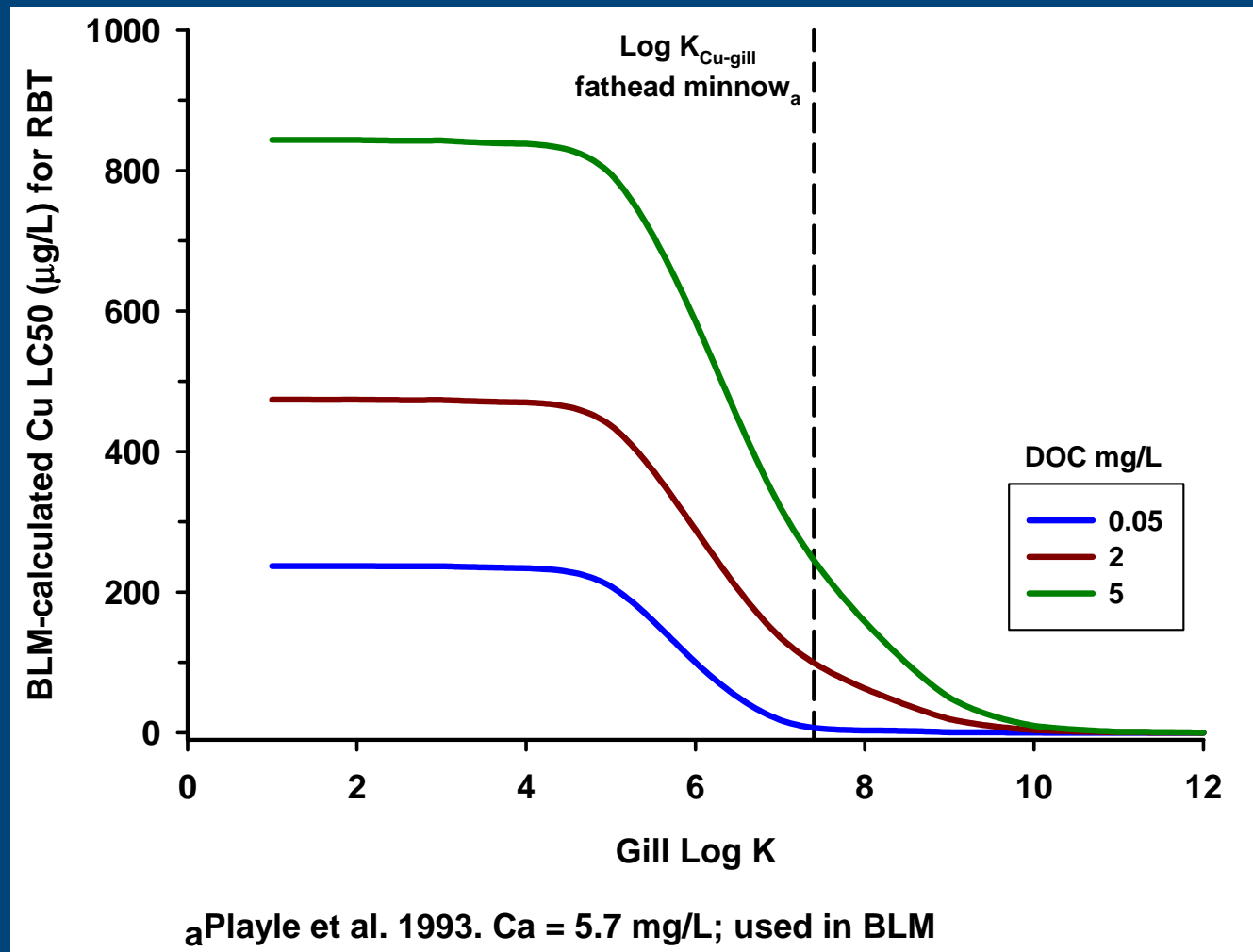
- Cu toxicity a function of relative complexation: $\log K$ of DOC v. $\log K$ of biotic ligand
- Biotic ligand not as refined as other two BLM components
- Current BLM uses a constant $\log K$ value for the biotic ligand
 - Shifts in relative $\log K$ of DOC in water v. constant $\log K$ in biotic ligand alter predicted toxicity

Biotic Ligand (gill) Log K in the BLM

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Cu_Rainbow_Trout_06-10-07 - Notepad
File Edit Format View Help
Column model parameter file, ver 3.00
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Number of Components, Species, Phases, Linked Lists
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Component Charge Type Activity site Den
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Cu, 2, 2, 2, 1
DOC, -1, 2, 1, .0006
Ca, 2, 2, 2, 1
Mg, 2, 2, 2, 1
Na, 1, 2, 2, 1
K, 1, 2, 2, 1
SO4, -2, 2, 2, 1
Cl, -1, 2, 2, 1
CO3, -2, 2, 2, 1
S, -2, 2, 2, 1
BL, -1, 11, 1, 3E-05
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BL-CuOH, 11, 1, 0, 0, -1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, -1.300 0.000 0.000
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BL-Mg, 11, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 1, 3.600 0.000 0.000
BL-H, 11, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 5.400 0.000 0.000
BL-Na, 11, 1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 1, 3.000 0.000 0.000
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Linked Lists
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Phase iTemp iMonte H Cu DOC Ca Mg Na K SO4 Cl CO3 S BL Log Ks Var Log K Delta H Temp
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File I/O and System Description
D:\MODELS\CHESS315\CU\LC50.WK1 | File with sequence of input chemistry (next line)
2 | Type of file (1 = ASCII, 2 = LOTUS)
  
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Log K in the BLM



Shifts in Apparent Gill Log K with Hardness?

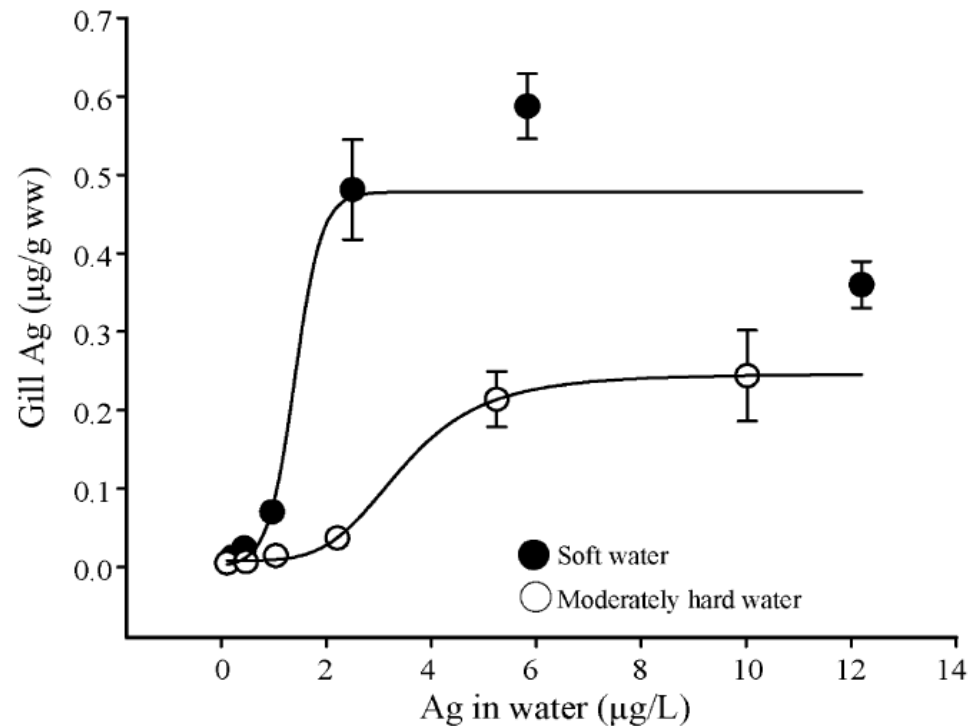
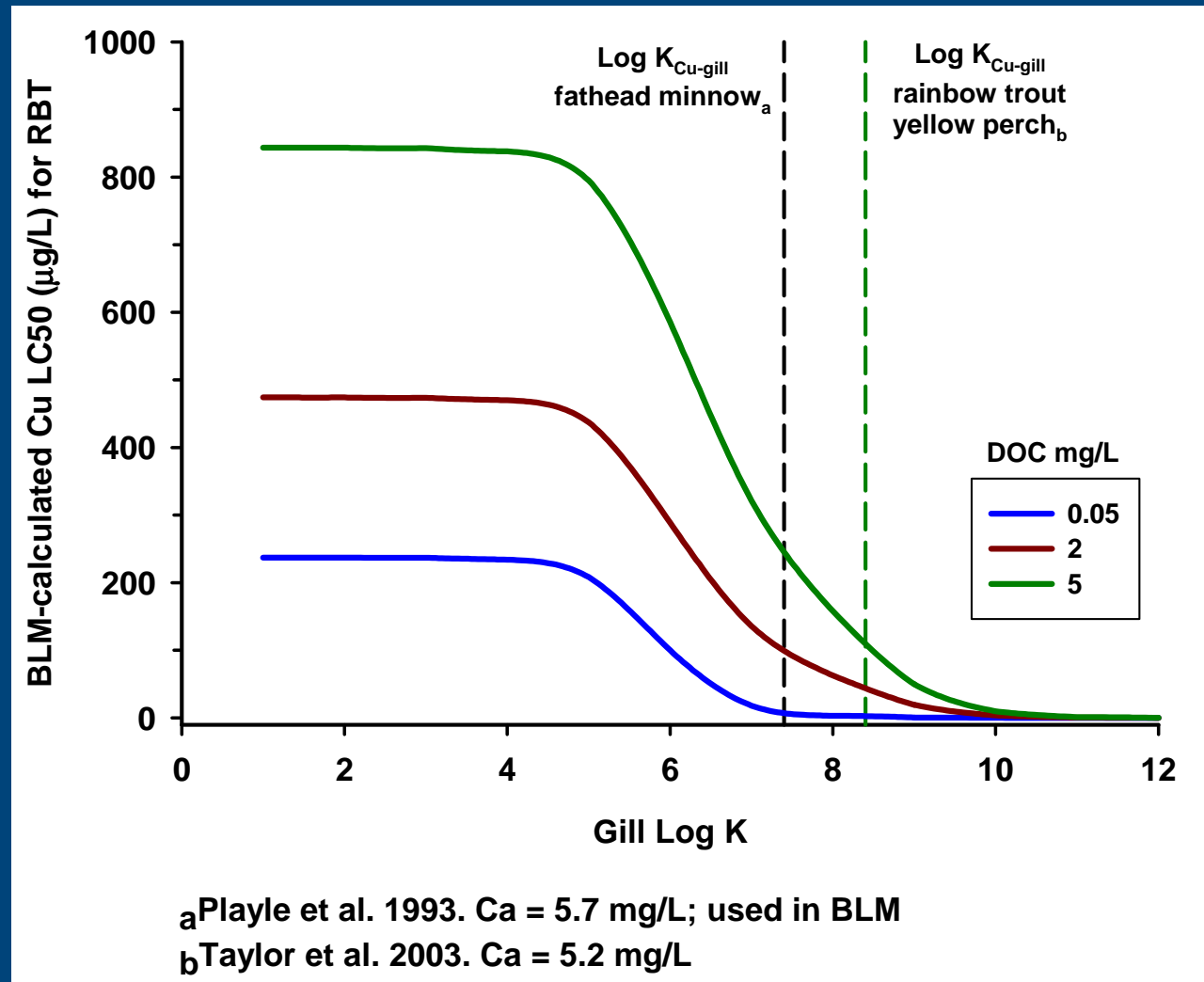


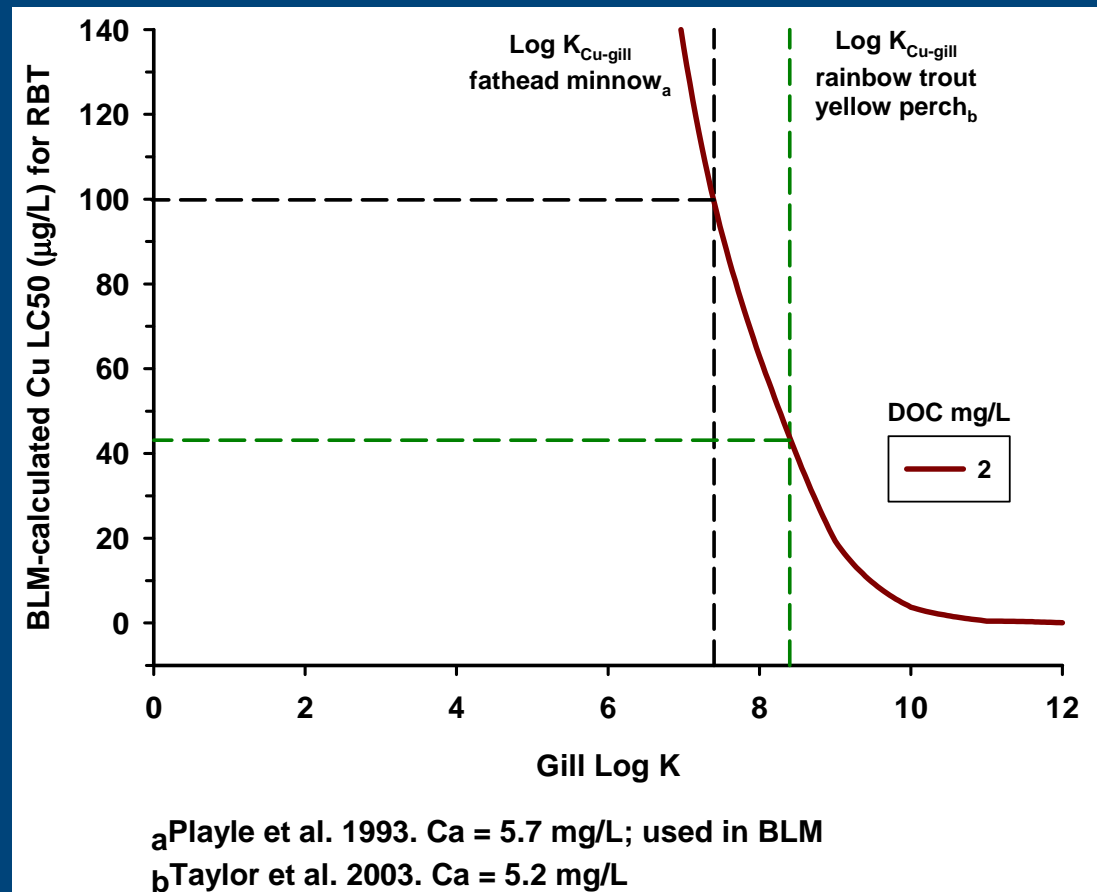
Fig. 4. Silver (Ag) accumulation on the gill (wet weight; ww) of *Pimephales promelas* (previously acclimated to soft water or moderately hard water) during 1-h Ag exposure in soft water. Error bars represent standard error.

Bielmyer et al., 2008.

Measured Gill Log Ks in Different Species












Effects of Varying Log K on Predicted Toxicity





Biotic Ligand Log K Summary

-  □
 -  □
 -  □
 -  □
 -  □
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 -  □
 -  □
 -  □
- Gill Log K known to change with water chemistry – dynamic
 - Using Log Ks developed for different species may result in ~ 2-fold change in LC50 at DOC = 2 mg/L
 - Variable log K in gill + variable log K in site water = variable predicted toxicity



Conclusions

- BLM under-predicted free Cu compared to site-specific estimates
- Needed to lower DOC in BLM to attain same free Cu results – similar findings to other researchers (e.g., De Schamphelaere et al., 2004; Welsh et al., 2008), but somewhat greater magnitude of adjustment
 - ~ 5-fold decrease in instantaneous WQC



Conclusions (cont.)

- Simulation modeling with BLM suggests Cu preferentially bound to DOC rather than the biotic ligand (gill) at low hardness
- Degree of under-predicted Cu toxicity
- Variable log K in gill + variable log K in site water = variable predicted toxicity
- Uncertainty in Cu toxicity can be reduced with supplemental site-specific data
 - Cu-DOC complexation
 - Species-specific toxicity testing