1	Supporting Information		
2	to		
3	Dissolved organic matter enhances microbial mercury		
4	methylation under sulfidic conditions		
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Table S1. Thermodynamic data for equilibrium speciation modeling.

Reaction	log K	Reference	
Hg-S Aqueous Speciation			
$Hg^{2+} + 2HS^{-} = Hg(SH)_{2}^{0}$	37.7	Benoit <i>et al.</i> ¹	
$Hg^{2+} + 2HS^{-} = HgS_2H^{-} + H^{+}$	31.5	Benoit <i>et al.</i> ¹	
$Hg^{2+} + 2HS^{-} = HgS_{2}^{2-} + 2H^{+}$	23.5	Benoit <i>et al.</i> ¹	
$Hg^{2+} + HS^{-} = HgSH^{+}$	30.2	Benoit <i>et al.</i> ¹	
Metacinnabar Precipitation			
$Hg^{2+} + HS^{-} = HgS(s) + H^{+}$	38±2	NIST Critical Database ²	
Hg-DOM Complexation			
$Hg^{2+} + 2RS^{-} = Hg(SR)_2$	42.0	Skyllberg ³	
$RS^- + H^+ = RSH$	10.0	Skyllberg ³	
Hg-CYS Complexation			
$Hg^{2+} + CYS^{2-} = HgCYS^{0}$	38.5	Basinger et al. ⁴	
$Hg^{2+} + 2H^{+} + 2CYS^{2-} = Hg(HCYS)_{2}^{0}$	39.8	Basinger <i>et al.</i> ⁴	
$Hg^{2+} + 2CYS^{2-} = Hg(CYS)_2^{2-}$	45.3	Basinger et al. ⁴	





Figure S1. Initial and final **a**) pH, **b**) optical density, **c**) total cell protein, and **d**) sulfide in washed cell Hg-methylation assays of *D. desulfuricans* ND132 in the presence of Suwannee River humic acid (SRHA) or L-cysteine (CYS). Washed cells were incubated with 5.0 nM ²⁰¹HgCl₂ for 3 hours at 31 °C. Initial measurements were taken at the beginning of the incubation, and final measurements after the 3 h incubation period. Error bars are standard deviations of triplicate methylation assays. These data correspond to the THg and MeHg data shown in Figure 1 in the main text.



Figure S2. Initial and final **a**) pH, **b**) optical density, **c**) total cell protein, and **d**) sulfide in washed cell Hg-methylation assays of *D. desulfuricans* ND132 in the presence of Suwannee River humic acid (SRHA) or L-cysteine (CYS). Washed cells were incubated with 0.5 nM ²⁰¹HgCl₂ for 3 hours at 31 °C. Initial measurements were taken at the beginning of the incubation, and final measurements after the 3 h incubation period. Error bars are standard deviations of triplicate methylation assays. These data correspond to the MeHg and THg data shown in Figure 1 in the main text.





Figure S3. Initial and final a) pH, b) optical density, c) total cell protein, and d)
sulfide in washed cell Hg-methylation assays of *D. desulfuricans* ND132 in the
presence of Williams Lake hydrophobic acid (WLHPoA) or L-cysteine (CYS).
Washed cells were incubated with 0.5 nM ²⁰¹HgCl₂ for 3 hours at 31 °C. Initial
measurements were taken at the beginning of the incubation, and final
measurements after the 3 h incubation period. Error bars are standard deviations
of triplicate methylation assays. These data correspond to the MeHg and THg data

- shown in Figure 2.

- DOM enhances Hg methylation



Figure S4. Relationship between filterable total Hg (THg) and total methylmercury115(MeHg) production in washed cell assays with (**a**) dissolved organic matter or (**b**) or116L-cysteine. SRHA = Suwannee River humic acid. WLHPoA = Williams Lake117hydrophobic acid. Filterable 201 THg and total MeHg production were not strongly118correlated in experiments with DOM isolates ($r^2 = 0.15$, p = 0.011). For experiments119with L-cysteine addition, total MeHg production was strongly correlated with120filterable THg ($r^2 > 0.99$, p < 10-7).</td>





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Figure S5. Methylmercury production is linearly dependent upon dissolved organic
matter concentration in experiments with either (a) Suwannee River humic acid
(SRHA) or (b) Williams Lake hydrophobic acid (WLHPoA).

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132 **References**

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