



Supplement of

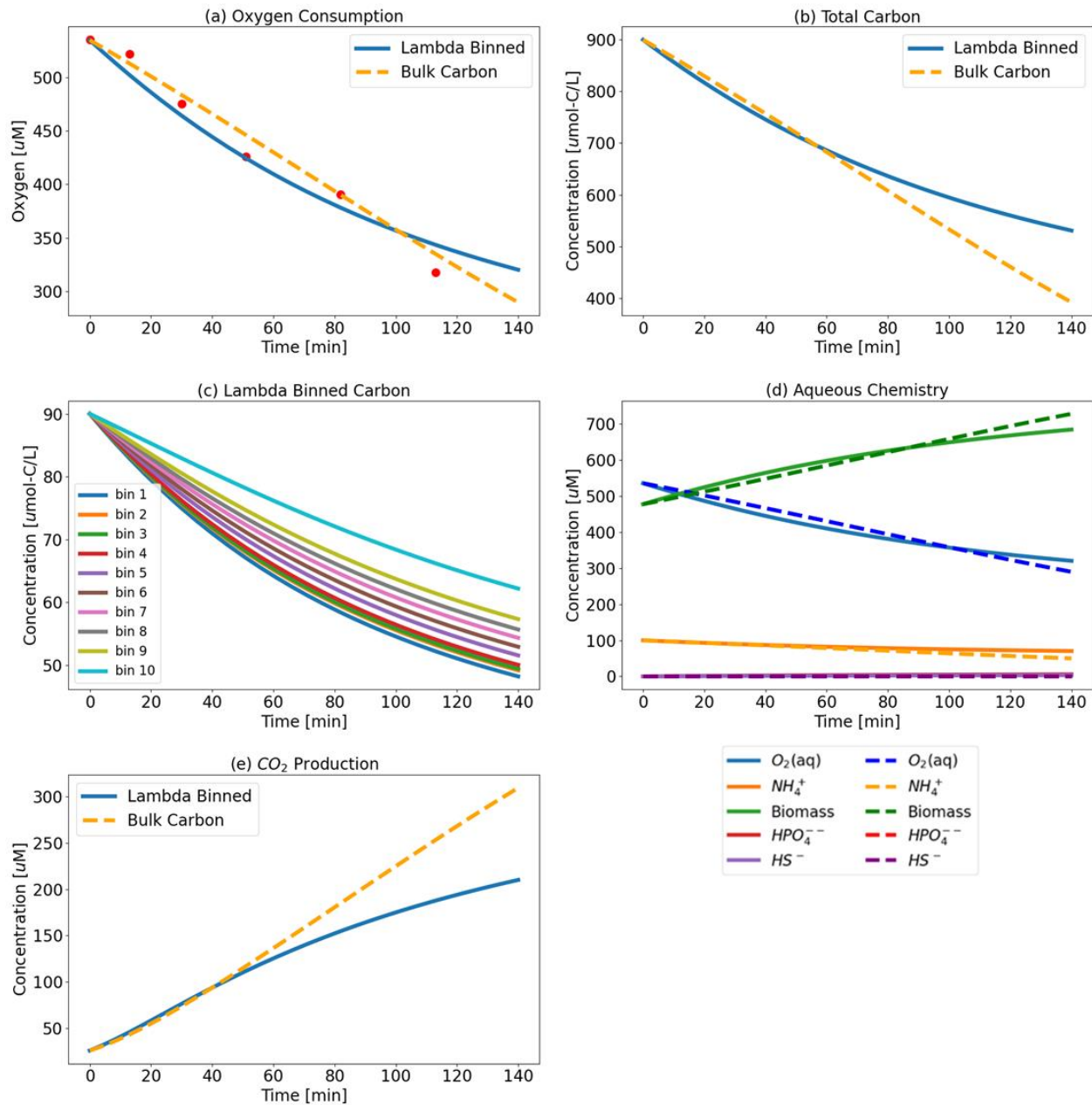
Lambda-PFLOTRAN 1.0: a workflow for incorporating organic matter chemistry informed by ultra high resolution mass spectrometry into biogeochemical modeling

Katherine A. Muller et al.

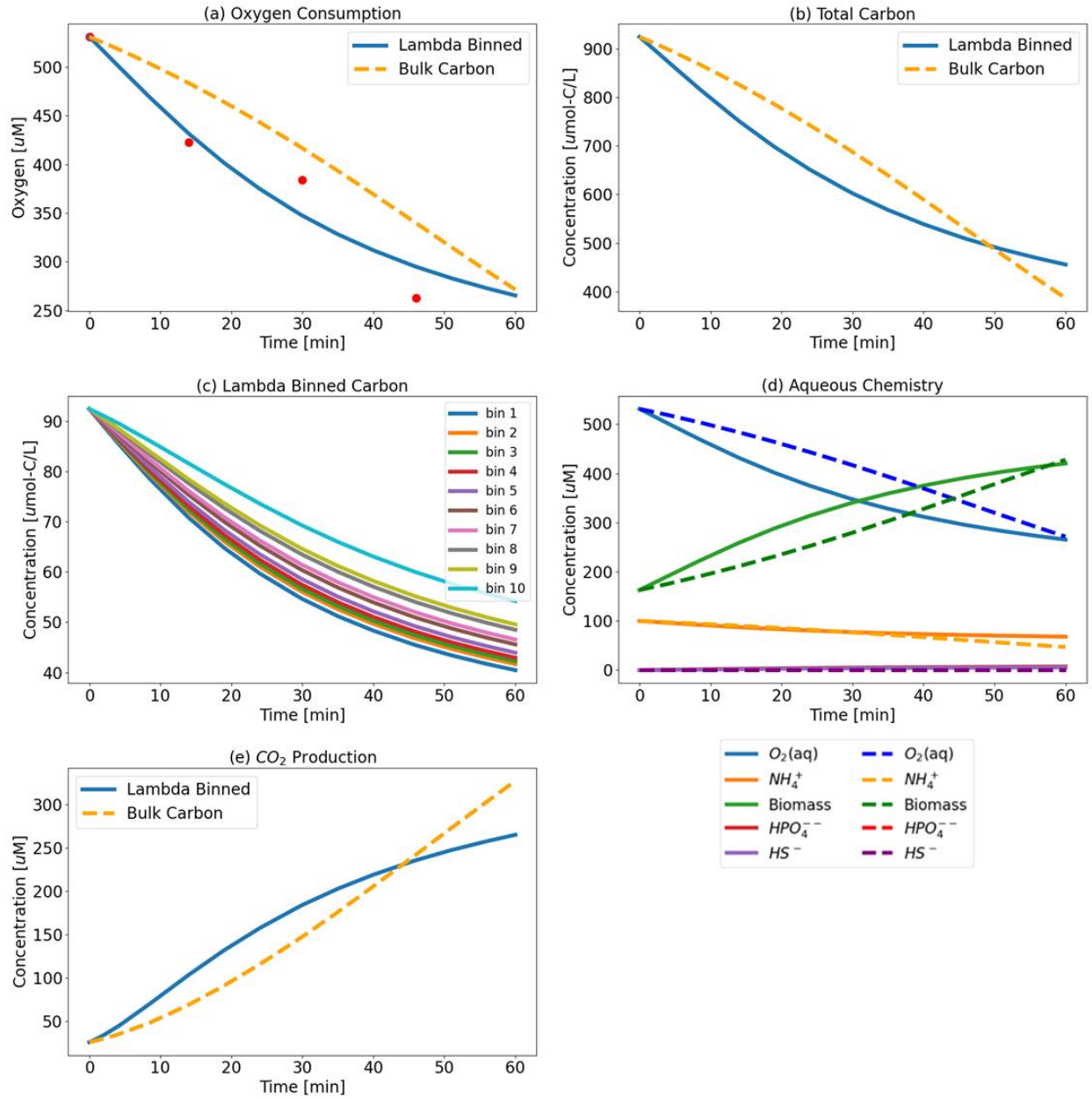
Correspondence to: Katherine A. Muller (katherine.muller@pnnl.gov)

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14 **SI.1 Test Case 1b and 1c Results**



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 16 **Figure S1:** Test Case 1b Results – (a) Oxygen Consumption where Lambda-PFLOTRAN workflow was used to fit (blue line) to
 17 experimental respiration data (red dots) and (b) the corresponding Total Carbon Consumption; (c) Individual Organic Matter
 18 Consumption by λ bin; and (d) biogeochemistry including O_2 (aq) (blue); Biomass (green); NH_4^+ (orange); HS^- (purple); and HPO_4^-
 19 $^-$ (red); and (e) CO_2 production for the upstream incubation. The dashed orange lines in (a, b and e) show simulation results assuming
 20 a generic OM species of CH_2O for comparison. Fitted μ_{max} values for the lambda binned model is 0.055 min^{-1} ($R^2 = 0.95$) and fitted
 21 μ_{max} to the bulk OM CH_2O model is 0.0065 min^{-1} ($R^2 = 0.97$). V_h and CC are fixed at assumed values of 10 m^3 and 1 M , respectively
 22 in both simulations.



23
 24 **Figure S2:** Test Case 1c Results – (a) Oxygen Consumption where Lambda-PFLOTTRAN workflow was used to fit (blue line) to
 25 experimental respiration data (red dots) and (b) the corresponding Total Carbon Consumption; (c) Individual Organic Matter
 26 Consumption by λ bin; and (d) biogeochemistry including O_2 (aq) (blue); Biomass (green); NH_4^+ (orange); HS^- (purple); and
 27 HPO_4^- (red); and (e) CO_2 production for the upstream incubation. The dashed orange lines in (a, b, and e) show simulation results
 28 assuming a generic OM species of CH_2O for comparison. Fitted μ_{max} values for the lambda binned model is 0.35 min^{-1} ($R^2 = 0.92$)
 29 and fitted μ_{max} to the bulk OM CH_2O model is 0.13 min^{-1} ($R^2 = 0.48$). V_h and CC are fixed at assumed values of 10 m^3 and 1 M ,
 30 respectively in both simulations.