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# Benthic biolayer structure controls whole-stream reactive transport

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## Key Points:

- Storage in non-reactive sublayer causes long contaminant survival times
- Reach-scale reaction kinetics cannot be captured by constant rate model
- Upscaled model quantifies all aspects of reaction and mass transfer

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## Abstract

Hyporheic zone reaction rates are highest just below the sediment-water interface, in a shallow region called the benthic biolayer. Vertical variability of hyporheic reaction rates leads to unexpected reaction kinetics for stream-borne solutes, compared to classical model predictions. We show that deeper, low-reactivity locations within the hyporheic zone retain solutes for extended periods, which delays reactions and causes solutes to persist at higher concentrations in the stream reach than would be predicted by classical approaches. These behaviors are captured by an upscaled model that reveals the fundamental physical and chemical processes in the hyporheic zone. We show how time scales of transport and reaction within the biolayer control solute retention and transformation at the stream scale, and we demonstrate that accurate assessment of stream-scale reactivity requires methods that integrate over all travel times.

## Plain Language Summary

Dissolved materials such as carbon, nutrients, and contaminants react as they move through the river network. Some locations in the river are far more reactive than others, and it is challenging to predict how this spatial variability of reaction rates controls the reactivity of the entire stream. One hotspot of high reactivity is the benthic biolayer, a thin region below the sediment-water interface with an abundance of microbial activity, and below which reactivity decreases to very low values. We use a mathematical model to quantify the benthic biolayer's contribution to whole river material transformation, based on the biolayer's thickness and reactivity. We show that thin or less reactive biolayers allow dissolved mass to become sequestered for long periods deep in the streambed, leading to low but persistent concentrations long after the mass is introduced to the river. These theoretical advances improve our understanding of how measurable features of the river – namely, the depth-dependent reaction rates within the streambed – are directly related to biogeochemical transformations and contaminant retention timescales in rivers.

## 1 Introduction

A defining feature of rivers is the transition in physical and chemical characteristics across the sediment-water interface (SWI). Downstream velocities, mixing rates, and light availability decrease rapidly at the SWI to viscous flows and light limited conditions (Jones & Mulholland, 1999). This transition zone, called the benthic biolayer, contributes disproportionately to the biologically-mediated transformation of reactive solutes and fine particulate matter in the river corridor, including heterotrophic carbon respiration, nutrient cycling, and trace contaminant degradation (Battin et al., 2008; Kunkel & Radke, 2008; Marzadri et al., 2017). Fluvial ecosystems are highly sensitive to physical perturbations and elevated contaminant concentrations in the biolayer, since this region supplies refugia and energy (as microbial biomass) for freshwater vertebrates (Jones & Mulholland, 1999; Cardenas et al., 2016; Moran et al., 2017). Structural features of the biolayer, such as its depth and reactivity, are therefore important predictors of whole-river reactivity and ecosystem health.

Dissolved oxygen depletes as aerated surface water propagates deeper into the hyporheic zone (HZ), leading to conditions that sustain microbial communities with higher tolerance for anoxia and slower metabolism. The stratification of chemical conditions and microbial biomass below the sediment-water interface (SWI) creates sharp gradients in reaction rates, as well as regions of low reactivity below the biolayer (Kunkel & Radke, 2008; Harvey et al., 2013; Knapp et al., 2017). The vertical profile of reaction rates is challenging to determine not only because steep concentration gradients are difficult to measure in pore waters, but also because various transport processes are simultaneously active (e.g., advective pumping, molecular and turbulent diffusion, mechanical dispersion). These processes are often grouped using scaling laws to estimate vertical solute

fluxes across the SWI. Scaling predictions are related to subsurface concentrations by assuming that solutes diffuse vertically (O'Connor & Harvey, 2008; Grant et al., 2012), which allows the reaction profile to be inferred from a diffusion-reaction or similar 1-D transport model (Harvey et al., 2013; Knapp et al., 2017; Schaper et al., 2019).

Recent modeling efforts strongly suggest that spatial variability of reaction rates in the biolayer controls the fate of reactive solutes at the stream scale. Numerical simulations show that whole-stream transformation is 5-25 $\times$  greater when HZ reaction rates are highest near the SWI, compared to a stream with the same average reactivity uniformly distributed in the HZ (Li et al., 2017). These differences arise because solutes entering the HZ typically propagate through shallow, high reactivity flowpaths before returning to the water column. Process-based models must therefore account for the correlation between residence time in the stream and residence time in reactive regions of the river bed. Traveltime based approaches for advection-dominated hyporheic flows account for the variability of reaction rates between flow paths by assuming that fluid parcels move unmixed through the HZ (Azizian et al., 2015; Reeder et al., 2018). However, we currently lack a physics-based upscaling framework that accounts for the joint impact of spatially varying reaction rates and diffusive mass transfer in the HZ. This knowledge gap limits our understanding of how measurable features of the HZ contribute to river corridor biogeochemistry, as well as how long streamborne contaminants are retained in the benthic biolayer and the less reactive sublayer that can act as a secondary source.

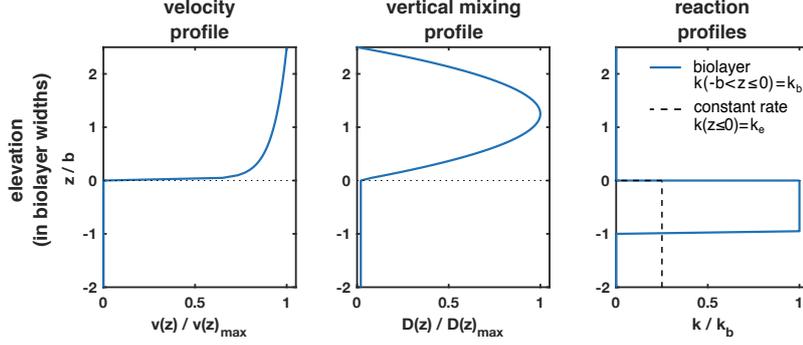
This work is motivated by the questions of how the size and reactivity of the benthic biolayer influence reach-scale mass fate, and how these properties manifest in up-scaled observations of reactive transport. We isolate the effects of depth-dependent HZ reaction rates on up-scaled predictions of solute fate in a stream. Solute transport in the HZ is considered to be dominated by vertical diffusion, which aligns our analysis with existing diffusion-based scaling laws that predict hyporheic exchange fluxes from measurable stream parameters. We present streambed-scale and reach-scale simulations designed to mimic a pulse tracer injection, which is a common method for assessing the processes controlling reactive transport in rivers and whose results are extendable to other boundary conditions (e.g., plateau injection experiments). We compare numerical results to predictions from a classical mobile-immobile model with uniform reactions in the subsurface, as well as predictions from a novel mobile-immobile model that explicitly represents the vertically-varying reaction profile in the biolayer.

## 2 Methods

### 2.1 Transport scenario

We consider the transport and reaction scenario illustrated in Figure 1 that comprises the water column and HZ. The SWI is located at  $z = 0$ , the water column extends from  $z = 0$  to  $z = d$ , and the HZ extends from  $z = -h$  to  $z = 0$ . We set the stream velocity to  $v(z) = v_s + v_0\kappa^{-1} \ln(z/z_0)$  for  $z > z_0$  and zero for  $z < z_0$  (Fischer et al., 1979). The length  $z_0$  is the width of a roughness layer at the SWI and represents the effect of the porous streambed on the stream velocity,  $\kappa \approx 0.41$  is the von Karman coefficient,  $v_0$  the shear velocity, and  $v_s$  the slip velocity at the SWI. We set  $v(z) = 0$  in the HZ, which assumes that streamwise velocity in the HZ is negligible compared to the stream. Vertical mixing is quantified by the dispersion coefficient  $D(z)$ , which is set equal to  $D(z) = \kappa v_0 z (1 - z/d)$  for  $z > z_0$  in the water column (Fischer et al., 1979) and equal to the constant effective diffusivity  $D_h$  in the HZ (Grant et al., 2012). We disregard streamwise dispersion and diffusion because advection in the water column dominates streamwise transport.

Solutes undergo first-order reactions in the HZ, which is a reasonable assumption when the modeled solute is limiting, that is, reactions are independent of the concen-



**Figure 1.** (Left to right) Spatial velocity, vertical mixing, and reaction profiles across the surface-subsurface continuum. The SWI is located at  $z = 0$ . The figures are truncated at  $z/b = -2$  since all values are constant at deeper locations in the HZ.

112 tration of co-reactants, abundance of catalysts such as enzymes, or thermodynamic con-  
 113 straints (Dodds et al., 2002; Garayburu-Caruso et al., 2020). The depth-dependent re-  
 114 action rate  $k(z)$  can be an arbitrary function of streambed elevation, but typically de-  
 115 creases sharply with depth due to the presence of a reactive biolayer. We follow Li et  
 116 al. (2017) and consider a biolayer structure consistent with field observations (Knapp  
 117 et al., 2017; Schaper et al., 2019; Inwood et al., 2007; O’Connor & Harvey, 2008). The  
 118 reaction rate is set to  $k(z) = k_b$  within a layer of thickness  $b$  just below the SWI. It is  
 119 set to  $k(z) = 0$  in the non-reactive sublayer of thickness  $\ell = h - b$ . Other profiles that  
 120 decay on a characteristic length scale  $b$  are expected to show a qualitatively similar be-  
 121 havior. The characteristic time scales in the HZ are given by the characteristic reaction  
 122 time  $\tau_r = k_b^{-1}$ , as well as the characteristic diffusion times  $\tau_h = h^2/D_h$  across the HZ,  
 123  $\tau_b = b^2/D_h$  across the biolayer, and  $\tau_\ell = \ell^2/D_h$  across the non-reactive sublayer. The  
 124 Damköhler number  $Da = \tau_b k_b$  compares the diffusion and reaction times in the bio-  
 125 layer. We consider the order of time scales  $\tau_r \leq \tau_b < \tau_\ell$ , which means that  $Da \geq 1$ .  
 126 This implies that reactions can occur before solute is transmitted to the sublayer. If,  
 127 on the contrary  $\tau_r > \tau_b$  (i.e.,  $Da < 1$ ), only a small amount of solute can react before it  
 128 reaches the sublayer. In this case, the behaviors of reactive and conservative solutes are  
 129 very similar, and thus are not presented. The evolution of solute concentration  $C(x, z, t)$   
 130 in the combined stream-HZ system is expressed by the advection-dispersion equation

$$131 \theta(z) \frac{\partial C}{\partial t} + v(z) \frac{\partial C}{\partial x} - \frac{\partial}{\partial z} \left[ \theta(z) D(z) \frac{\partial C}{\partial z} \right] = -\theta(z) k(z) C, \quad (1)$$

133 where the porosity  $\theta(z)$  equals 1 in the water column and a constant value  $\theta_h$  in the HZ.  
 134 The horizontal boundaries  $z = -h$  and  $z = d$  are impermeable. In line with the ex-  
 135 perimental design of field tracer studies, we assume that the HZ is initially free of reac-  
 136 tive mass, and solute is introduced as a line injection in the water column. The direct  
 137 advection-dispersion problem (1) is solved numerically using a reactive time-domain ran-  
 138 dom walk approach (TDRW), based on the implementation of Russian et al. (2016) for  
 139 conservative solutes. The TDRW method is computationally efficient for media with spa-  
 140 tially heterogeneous advection, diffusion and reaction properties. Details on the imple-  
 141 mentation, discretization, and parameterization of the TDRW simulations are given in  
 142 supporting information Section SI-VI. In the following, we present the mobile-immobile  
 143 model that upscales this reactive transport scenario. In order to analyze the impact of  
 144 the benthic biolayer on whole stream reactive transport, we consider solute breakthrough  
 145 curves at a downstream control plane for different biolayer scenarios.

## 2.2 Mobile-immobile biolayer (MIM-B) model

We employ a mobile-immobile approach (Villermanx, 1974; Haggerty & Gorelick, 1995) to upscale the reactive transport problem. Eq. (1) is decomposed into an advection-dispersion equation for transport in the stream, a diffusion-reaction equation for the biolayer, and an equation for vertical diffusion in the sublayer. These equations are coupled through concentration and flux continuity at their respective interfaces. By vertical averaging we obtain a temporally non-local evolution equation for the average stream concentration  $\bar{C}_s(x, t)$ :

$$\begin{aligned} \frac{\partial \bar{C}_s}{\partial t} + \frac{\theta_h}{d} \frac{\partial}{\partial t} \int_0^t dt' \varphi_h(t-t') \bar{C}_s(x, t') \\ + \bar{v} \frac{\partial \bar{C}_s}{\partial x} - D^* \frac{\partial^2 \bar{C}_s}{\partial x^2} = -\frac{\theta_h k_b}{d} \int_0^t dt' \varphi_b(t-t') \bar{C}_s(x, t'). \end{aligned} \quad (2)$$

Details are provided in Section SI-II of the supporting information. The mean velocity in the stream is denoted by  $\bar{v}$ , and the shear dispersion coefficient  $D^* = 5.93v_0d$  (Fischer et al., 1979) quantifies the impact of vertical velocity variability on longitudinal dispersion in the stream. The non-local term (second term on left side) denotes the time derivative of the concentration in the HZ. It quantifies solute trapping, release, and degradation in the hyporheic zone. The term on the right side demonstrates that the upscaled reaction kinetics are temporally non-local, but nonetheless linear. The non-locality stems from the fact that solute first diffuses into the biolayer before it reacts. Thus, the mass degraded at a given time  $t$  is proportional to the stream concentrations  $\bar{C}_s(x, t')$  at earlier times  $t'$ . The memory kernel  $\varphi_h(t)$  describes the evolution of mass in the HZ in response to an instantaneous solute pulse at the SWI. It is decomposed into  $\varphi_h(t) = \varphi_b(t) + \varphi_0(t)$ , wherein  $\varphi_b(t)$  and  $\varphi_0(t)$  encode the diffusive and reactive mass transfer mechanisms across the biolayer and the non-reactive sublayer, respectively. Explicit Laplace space expressions for  $\varphi_b(t)$  and  $\varphi_0(t)$  are given in Section SI-II C. While this formulation can be generalized to account for water column reactions (sensu Roche et al., 2019), we exclusively model reactions in the HZ to elucidate the biolayer's influence on stream-scale reactivity.

## 2.3 Surrogate models

To illustrate the benthic biolayer's impact on whole stream reactive transport, we contrast the MIM-B with two surrogate models, termed *S1* and *S2*. Model *S1* assumes the hyporheic zone is uniformly reactive over all depths. Model *S2* assumes that water column and HZ are in equilibrium.

### 2.3.1 Fully reactive hyporheic zone (*S1*)

In agreement with classical assumptions (e.g., Runkel, 2007; Haggerty et al., 2009; Aubeneau et al., 2015), *S1* assumes that the HZ is fully reactive and characterized by an equivalent reaction rate  $k_e$ . The evolution equation for  $\bar{C}_s$  is obtained from (2) by substituting  $k_b$  with  $k_e$  and setting  $\varphi_h(t) = \varphi_b(t) \equiv \varphi_e(t)$ . The latter can be written in terms of the memory kernel  $\phi(t)$  for a non-reactive solute as  $\varphi_e(t) = \phi(t) \exp(-k_e t)$  (Dentz et al., 2011). We define  $k_e$  such that the total reacted mass in the HZ, in response to an instantaneous solute pulse at the SWI, is equal to the total reacted mass in the MIM-B. Using this definition, we derive the following transcendental equation for  $k_e$  (see, SI-IV)

$$\sqrt{\frac{k_e}{k_b}} \tanh\left(\sqrt{k_e \tau_h}\right) = \tanh\left(\sqrt{Da}\right). \quad (3)$$

191 The solution of (3) can be approximated by  $k_e = k_b \tanh(\sqrt{Da})^2$  for  $k_e \tau_h > 10$ . This  
 192 implies that the equivalent streambed reactivity increases monotonically with  $Da$  and  
 193 asymptotes towards  $k_b$  as  $Da \rightarrow \infty$ .

### 194 2.3.2 Equilibrium model for the water column (S2)

195 Model S2 assumes that the water column and the HZ are in equilibrium. The evo-  
 196 lution equation for  $\bar{C}_s$  in this limit is obtained from Eq. (2) by localization of the mem-  
 197 ory kernels on the left and right sides. This gives (Section SI-V)

$$198 \quad R_a \frac{\partial \bar{C}_s}{\partial t} + \bar{v} \frac{\partial \bar{C}_s}{\partial x} - D^* \frac{\partial^2 \bar{C}_s}{\partial x^2} = -k_a \bar{C}_s(x, t'), \quad (4)$$

200 where the apparent retardation coefficient  $R_a$  and the apparent reaction rate  $k_a$  are de-  
 201 fined by

$$202 \quad R_a = 1 + \frac{\theta_h}{d} \int_0^\infty dt' \varphi_h(t'), \quad k_a = \frac{\theta_h k_b}{d} \int_0^\infty dt' \varphi_b(t'). \quad (5)$$

### 204 2.4 Reach-scale reactivity

205 The two surrogate models S1 and S2 are defined such that they have the same down-  
 206 stream mass recovery as the MIM-B model. The fraction of mass recovered  $M_R$  at a down-  
 207 stream location is obtained by integration of the solute breakthrough curves over time  
 208 from zero to infinity. Thus, we obtain from S2 the explicit expression

$$209 \quad M_R(x) = \exp \left[ -\frac{x\bar{v}}{2D^*} \left( \sqrt{1 + \frac{4k_a D^*}{\bar{v}^2}} - 1 \right) \right]. \quad (6)$$

211 This predicted exponential decrease is commonly observed in field experiments. The reach  
 212 scale reactivity  $K_r$  [ $L^{-1}$ ] has been defined in the literature as the slope of the logarithm  
 213 of  $M_R(x)$ , that is,  $K_r \equiv -x^{-1} \ln M_R(x)$  (Tank et al., 2017). Reach scale reactivity  $K_r$   
 214 reads in terms of the apparent reaction rate  $k_a$  as

$$215 \quad K_r = \frac{\bar{v}}{2D^*} \left( \sqrt{1 + \frac{4k_a D^*}{\bar{v}^2}} - 1 \right). \quad (7)$$

217 Eq. (7) simplifies to  $K_r = k_a \bar{v}^{-1}$  in the limit  $D^* \rightarrow 0$ , meaning that  $M_r$  decays as  $\exp(-\tau_a k_a)$   
 218 in this limit, where  $\tau_a = x\bar{v}^{-1}$  is the advective travel time. Note that  $K_r$  is not a re-  
 219 action rate. It facilitates the estimation of reach-scale mass removal integrated over all  
 220 times. This is important to note because the time scales for reaction can be very large  
 221 due to mass transfer limitation in the HZ, which has a dramatic impact on contaminant  
 222 removal and secondary release as discussed below. The evaluation of  $K_r$  is one of sev-  
 223 eral methods that are often inter-compared to determine reach-scale reaction kinetics (e.g.,  
 224 Finkler et al., 2021). For ease of interpretation, we present reach-scale reactivity as up-  
 225 take velocities, which are commonly used for comparison across rivers. The inverse of  
 226  $K_r$  denotes the nutrient spiraling length  $S_w = K_r^{-1}$  [L], which describes the character-  
 227 istic distance a reactant travels downstream before reacting. The uptake velocity  $v_f =$   
 228  $K_r \bar{v} d$  [ $L T^{-1}$ ] measures demand for reactants relative to in-stream concentration (Tank  
 229 et al., 2017).

## 230 3 Results and discussion

### 231 3.1 Interplay between biolayer structure and solute fate in the HZ

232 Memory functions, which quantify the mass in the HZ resulting from an instan-  
 233 taneous solute pulse at the SWI, are shown in Figure 2 for (a) the sublayer, (b) the bi-

234 olayer, and (c) the entire HZ, obtained from the direct numerical simulations and the  
 235 analytical MIM-B.

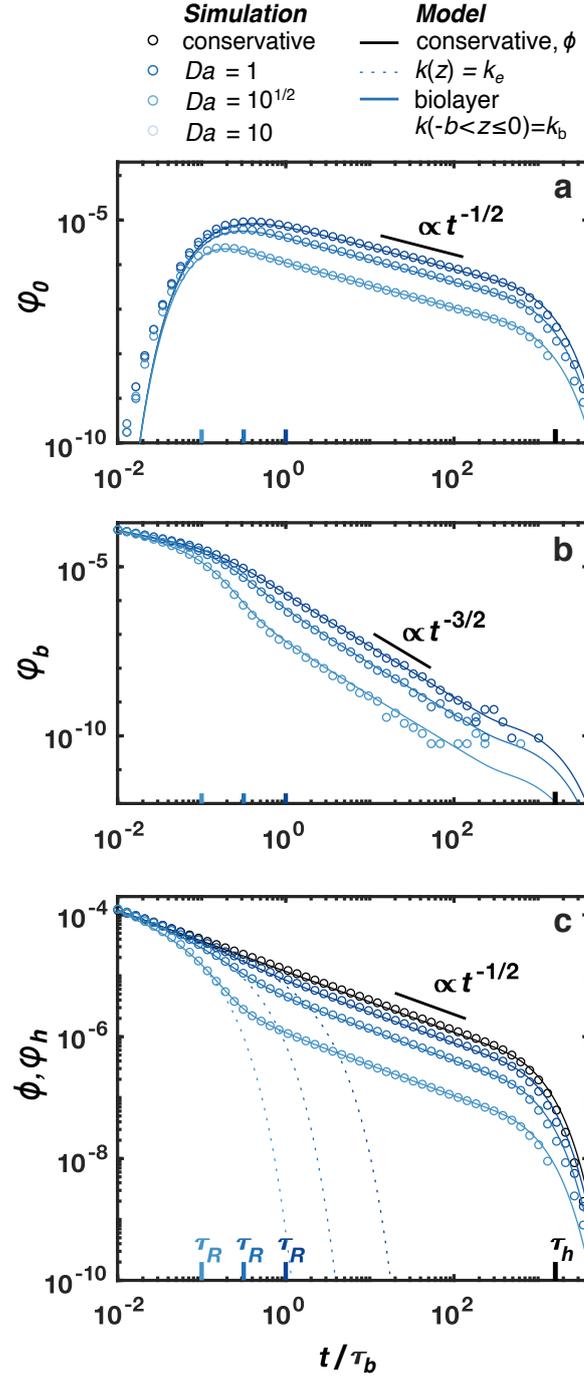
236 The sublayer memory function  $\varphi_0$  increases from 0 to a maximum on the time scale  
 237  $\tau_b$ , which is the time for solute transmission across the biolayer. It then decreases as  $t^{-1/2}$ ,  
 238 as for a conservative solute, due to diffusion back to the biolayer. Last, it tempers ex-  
 239 ponentially on the time scale  $\tau_\ell$  as the sublayer depletes by diffusion.

240 The memory function  $\varphi_b$  for the biolayer decays as  $t^{-1/2}$  for times smaller than the  
 241 reaction time,  $t < \tau_R$ , due to diffusion across the SWI. For  $\tau_R < t < \tau_b$  mass is de-  
 242 pleted from the biolayer by chemical reaction, which manifests in an exponential decrease  
 243 of  $\varphi_b$  (Figure 2b). For times  $t \gg \tau_b$ , the biolayer can be considered well-mixed, and  $\varphi_b$   
 244 transitions to a  $t^{-3/2}$  decay because mass in the biolayer changes in a quasi-static fash-  
 245 ion due to the mass flux from the sublayer (see SI-II D),

$$246 \varphi_b(t) = -\frac{\tau_b}{1 + Da} \frac{d\varphi_0}{dt} \propto t^{-3/2}. \quad (8)$$

248 The memory function  $\varphi_h$  integrates the diffusion-reaction process in the biolayer  
 249 and retention in the sublayer (Figure 2c). For times  $t \ll \tau_R$ , mass removal in the streambed  
 250 is primarily caused by diffusion upward across the SWI, and we observe the character-  
 251 istic  $t^{-1/2}$  decay of a conservative solute. As discussed above, solute is depleted by re-  
 252 action in the biolayer for  $\tau_R < t < \tau_b$ , giving rise to an exponential decay of  $\varphi_h$ . For  
 253  $t < \tau_b$  all remaining mass resides at shallow depth in the benthic biolayer, and the sys-  
 254 tem behaves as a scenario of constant streambed reactivity. For  $t > \tau_b$ , however, so-  
 255 lute diffuses into the inert sublayer. Eventually, most mass remaining in the streambed  
 256 is sequestered below the biolayer. The upward diffusion of mass from the inert sublayer  
 257 into the biolayer results in a second regime of  $\varphi_h(t) \sim t^{-1/2}$  (Figure 2c) because dif-  
 258 fusion from the sublayer through the biolayer and to the stream is the dominant deple-  
 259 tion process. Exponential tempering of  $\varphi_h(t)$  then occurs on the time scale  $\tau_\ell$  (Figure  
 260 2c). For comparison, we show memory functions for the corresponding surrogate model  
 261 *SI* parameterized with  $k_e$  (Figure 2c dotted line). It decays as  $t^{-1/2}$  for times smaller  
 262 than the reaction time  $\tau_e = k_e^{-1}$  and exponentially fast for  $t > \tau_e$  as solute degrades  
 263 throughout the HZ. Thus, *SI* predicts much faster depletion of reactant than the MIM-  
 264 B because it does not account for long survival in the sublayer.

265 In summary, the interaction of reaction and diffusion processes in the HZ is gov-  
 266 erned by three distinct timescales: the characteristic reaction time  $\tau_R = k_b^{-1}$ , which sets  
 267 the time for solute depletion from the biolayer by reaction; the diffusion time  $\tau_b$ , which  
 268 sets the time for solute transmission through the biolayer to the inert sublayer; and  $\tau_\ell$ ,  
 269 which sets the time for diffusive depletion of solute from the sublayer. The match be-  
 270 tween simulated and modeled memory functions shows that the MIM-B correctly cap-  
 271 tures the long survival times in the HZ and the spatial segregation of reactants between  
 272 the biolayer and the non-reactive sublayer.



**Figure 2.** Modeled and simulated memory functions of varying biolayer  $Da$ . a) Memory functions for the inert sublayer show all mass in  $-h \leq z < b$ . b) Memory functions for the benthic biolayer show all mass in  $-b \leq z < 0$ . c) Full memory functions for conservative (black) and reactive (colored) solutes. Model and simulations transition to  $t^{-1/2}$  tailing for  $t \gg \tau_b(1 + Da)^{-1}$ . For all experiments,  $b = 0.05$  m,  $D_h = 1.042 \times 10^{-6}$  m<sup>2</sup>s<sup>-1</sup>,  $h = -2$  m, and  $k_b$  is varied.

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## 3.2 Reach-scale observations and model predictions

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### 3.2.1 Breakthrough curves

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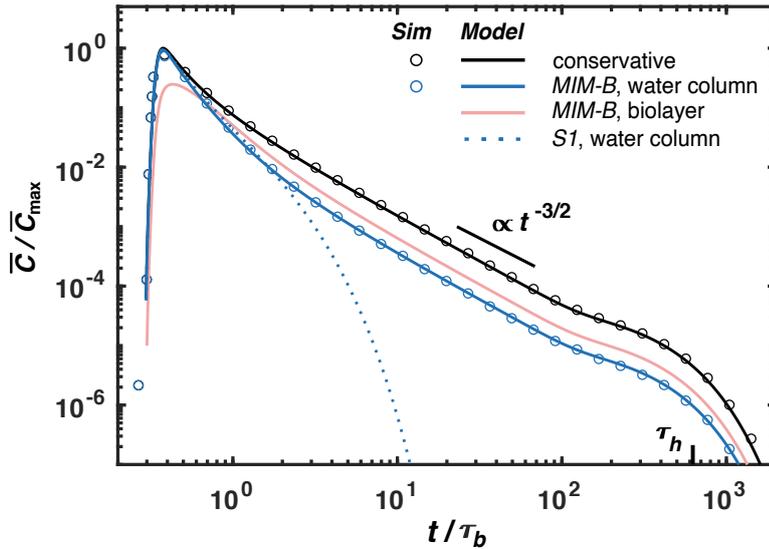
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Figure 3 shows BTCs for conservative and reactive solutes from numerical simulations and MIM-B model predictions, as well as MIM-B prediction for the biolayer concentration at a control plane 100 m downstream from the injection point. These results are compared to the prediction of surrogate model *S1* for a fully reactive HZ. The conservative BTC decays as a power law with  $t^{-3/2}$  and is cut off at the characteristic diffusion time across the HZ. This behavior is characteristic of diffusive mass transfer and secondary release from the HZ. The BTC for the reactive solute shows the same tailing features as the conservative BTC, albeit at lower concentrations due to degradation in the biolayer. The strong tailing of the contaminant concentration is caused by transmission of unreacted solute to the sublayer and release back into the stream through the biolayer. These behaviors are correctly quantified by the MIM-B, which predicts similar behavior for the contaminant concentration in the biolayer. This shows that both the stream and the biolayer are sourced by upward diffusion of solute sequestered in the sublayer. Results mirror results from memory function simulations, demonstrating that biolayer structure has a similar influence on degradation timescales at both the local scale and the whole-stream scale. On the other hand, *S1* predicts exponential decay of the BTC on the reaction timescale and thus severely under-predicts late time contaminant levels.



**Figure 3.** Simulated and MIM-B predicted BTCs for a pulse tracer injection with  $x = 100$  m and  $b = 0.08$  m, corresponding to  $Da = 1.2$ . See Section SI-VI for other parameter values.  $C_{\max}$  equals maximum concentration of the analytical solution for the conservative BTC.

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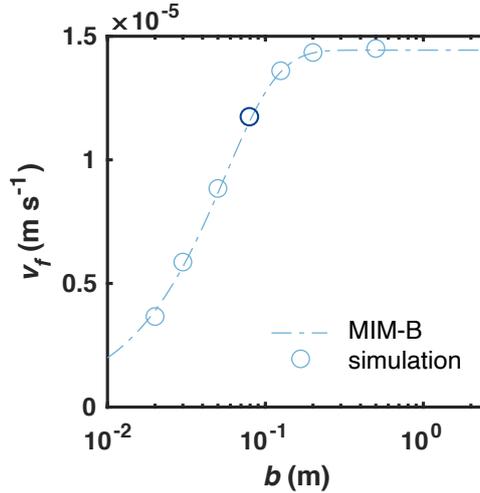
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The exact match between simulated BTCs and MIM-B predictions demonstrates that the MIM-B fully captures the impact of long survival times in the HZ, as well as the spatial segregation of reactants in the HZ, on reach scale transport and degradation. Notably, the model predicts a power law decay of survival times for all  $Da$ . This indicates the potential of MIM-B to provide correct estimates of trace contaminants in benthic sediments and the stream over a range of different physical and chemical conditions in the HZ. Trace organic contaminants (TOCs) are now detected in most rivers (Bernhardt

300 [et al., 2017](#)) and impair stream ecosystems at low levels, for example, endocrine disrupt-  
 301 ing compounds that alter fish physiology at nanomolar concentrations ([Adeel et al., 2017](#);  
 302 [Khanal et al., 2006](#)). Degradation rates of TOCs decrease rapidly with depth in the HZ  
 303 and span a range of half lives ([Kunkel & Radke, 2008](#); [Schaper et al., 2019](#)). These char-  
 304 acteristics allow TOCs to persist in sediments long after they enter the river network and  
 305 act as a secondary source ([Ciparis et al., 2012](#); [Cozzarelli et al., 2017](#)). This suggests that  
 306 their degradation timescales must be estimated by explicitly accounting for the verti-  
 307 cally varying reaction rates in the HZ (see [SI-VII](#)).

### 3.2.2 Reach-scale reactivity

309 Calculated  $v_f$  resulting from integration of the simulated BTC, sensu [Tank et al.](#)  
 310 ([2008](#)), agrees well with the analytical prediction from the MIM-B model (Eq. 7, Fig-  
 311 ure 4). The plot also shows that  $v_f$  approaches an asymptotic value for values  $b$  larger  
 312 than the characteristic survival depth  $s = (D_h/k_b)^{1/2}$ , which denotes the diffusion length  
 313 during the reaction times  $\tau_R$ . This implies that the spatial extension of the the biolayer  
 314 has little bearing on whole-stream reactivity for  $b > s$ . When  $b \gg s$ , nearly all solute  
 315 reacts before propagating below the biolayer. In contrast, a substantial amount of mass  
 316 propagates through the biolayer unreacted when  $b < s$ , resulting in a lower effective  
 reactivity of the HZ (see [SI-VII](#)) and a lower reach-scale reactivity.



**Figure 4.** Reach-scale reactivity reported as uptake velocity, for streambeds containing biolayers with varying  $b$ . The reaction rate is  $k_b = 2.0 \times 10^{-4} \text{ s}^{-1}$  and corresponds to a characteristic survival depth of  $s = 0.07 \text{ m}$ . See Section [SI-VI](#) for all parameter values. Dark blue circle is result from the simulation shown in [Figure 3](#).

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318 The MIM-B gives exact estimates of reach-scale reactivity under the assumed condi-  
 319 tions of diffusion dominated transport and stratified reactivity in the HZ. It should be  
 320 emphasized that these estimates are only valid when made at asymptotic times. Trans-  
 321 ient storage in the HZ delays transport through the reach, causing mass to arrive sig-  
 322 nificantly later than the advective timescale  $\tau_a = x \bar{v}^{-1}$ , upon which metrics such as  
 323  $v_f$  are typically based. This delay can cause time-resolved methods for estimating reach-  
 324 scale reactivity to deviate systematically from the reactivity calculated from integrated  
 325 mass transformation. For example, results from pulse tracer injection experiments are  
 326 commonly used to calculate a different effective reaction rate for each data point of the  
 327 BTC, wherein the reaction time is set to the breakthrough time ([Covino et al., 2010](#)).

328 The effective rates calculated from this method depend nonlinearly on reactant concen-  
329 tration even when reactions within the reach are linear, which may lead to the conclu-  
330 sion that reaction kinetics are nonlinear (Li et al., 2021). Integrated methods, such as  
331 integration of the BTC or constant rate injection experiments, account for the transport  
332 and reaction delays associated with non-local transport to the HZ and mass sequestra-  
333 tion below the biolayer. Nevertheless, using these methods may require very long obser-  
334 vation times in order to account for the power law decay of reactive mass released from  
335 the sublayer, and they provide no information about when concentrations may exceed  
336 critical thresholds. Finally, it is important to note that neither method gives informat-  
337 ion on the actual reaction kinetics and time evolution of the downstream contaminant  
338 concentration. The derived upscaled MIM-B shows that reaction kinetics are in fact non-  
339 local as expressed by the right side of Eq. (2) and characterized by a power-law decay  
340 of contaminant survival times.

## 341 4 Conclusion

342 Two fundamental challenges for providing mechanistic predictions of river corri-  
343 dor reactivity are to explicitly link local heterogeneity of the controlling physical pro-  
344 cesses to upscaled observations within a consistent modeling framework, and to identify  
345 the relative importance of microscale processes and structural features of the river cor-  
346 ridor (Ward & Packman, 2019; Kelleher et al., 2019). We address these challenges by  
347 analyzing and upscaling reactive transport in a river-streambed system characterized by  
348 a benthic biolayer. Isolating the dominant small scale features, we derive a novel upscaled  
349 model (MIM-B) that captures the dominant physical and chemical processes in the ben-  
350 thic biolayer, the HZ, and reach scale. The model predictions closely agree with detailed  
351 numerical simulations of transport and reaction in the river-streambed system.

352 We find that the biolayer structure strongly controls solute degradation in the HZ  
353 and at the reach scale. Accumulation in the sublayer leads to long survival times for re-  
354 active solute, characterized by a power-law decay of concentration and by the spatial seg-  
355 regation of mass in the HZ. This is in stark contrast to model predictions based on the  
356 classical assumption of a fully reactive HZ, for which the contaminant concentration de-  
357 cays exponentially fast on the characteristic reaction time scale and thus strongly under-  
358 predicts contaminant levels in the tail. The novel MIM-B captures all aspects of con-  
359 taminant degradation on the HZ and reach scales. Specifically, it correctly predicts tail  
360 concentrations and reach scale reactivity. Reach scale reactivity quantifies the reaction  
361 potential of the system; however, this potential can in principle only be observed at very  
362 long experimental times due to the role of the sublayer as a secondary release.

363 Although we assume solutes diffuse vertically through the HZ in order to align our  
364 model with empirical scaling laws, we expect similar qualitative behavior in any streambed  
365 with vertically varying reaction rates and a multiscale residence time distribution (e.g.,  
366 Elliott & Brooks, 1997). The characteristics of reactant fate identified within our model  
367 framework are critical for assessing contamination levels in streams and in shallow sed-  
368 iments, which are dramatically underestimated at late times by classical models that as-  
369 sume uniform reaction rates in the hyporheic zone.

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