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12 Abstract

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Watersheds are the fundamental Earth surface functioning unit that connects the land to aquatic systems. Many watershed-scale models represent hydrological processes but often lack the representation of multi-component reactive transport processes that are relevant to soil and aquatic biogeochemical reactions. The lack of mechanism-based representation of reaction thermodynamics and kinetics at the watershed scale has limited our ability to understand and predict solute export and water quality, particularly under changing climate and anthropogenic conditions. Here we present a recently developed BioRT-Flux-PIHM (BioRT hereafter) v1.0, a watershed-scale biogeochemical reactive transport model. Augmenting the previously developed RT-Flux-PIHM that integrates land-surface interactions, surface hydrology, and abiotic geochemical reactions (Bao et al., 2017, WRR), the new development enables the simulation of 1) biotic processes including plant uptake, soil respiration, and microbially mediated reactions such as carbon decomposition and nutrient transformation; and 2) shallow and deep water partitioning to represent surface, shallow groundwater, and deep groundwater interactions. The reactive transport part of the code has been verified against the widely used reactive transport code CrunchTope. BioRT-Flux-PIHM v1.0 has recently been applied to understand reactive transport processes in multiple watersheds under diverse climate, vegetation, and geological conditions. This paper briefly introduces the governing equations and model structure with a focus on new model developments. It also showcases one hydrology example that simulates shallow and deep water interactions, and two biogeochemical examples relevant to nitrate and dissolved organic carbon (DOC). These examples are illustrated in two simulation modes of varying complexity. One is the spatially lumped mode (i.e., two land cells connected by one river segment) that focuses on processes and average behavior of a watershed. Another is the spatially distributed mode (i.e., hundreds of cells) that includes details of topography, land cover, and soil property conditions. The spatially distributed mode can be used to understand the impacts of spatial structure and identify hot spots of biogeochemical reactions.

1. Introduction

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Watersheds are the fundamental Earth surface units that receive and process water, mass, and energy (Li, 2019;Li et al., 2020;Ranalli and Macalady, 2010;Hubbard et al., 2018;Seyfried et al., 2018). Watershed processes include land surface interactions that regulate evapotranspiration and discharge, and water partitioning between shallow soil lateral flow going into streams versus downward flow and recharge into the deeper subsurface (Edwards et al., 2015) (Figure 1). Complex biogeochemical interactions also occur among soil, water, roots, and microbes, dictating gas effluxes (e.g., CO₂) via soil respiration, export of solute products derived from chemical weathering and biogeochemical transformation (Fatichi et al., 2019;van der Velde et al., 2010;Grathwohl et al., 2013).

These hydrological and biogeochemical processes determine how land surface responds to external forcings such as hydroclimatic drivers and human perturbations (van der Velde et al., 2014; Miller et al., 2020; Han et al., 2019; Steimke et al., 2018). Understanding these processes remains challenging due to the complex coupling of land surface processes, hydrology, and biogeochemical reactions (Kirchner, 2003). An example is the concentration-discharge (C-Q) relationships of solutes at stream and river outlets. These relationships encode the integrated signature of land surface responses to changes in hydrological conditions (Brooks et al., 2015; Zhi et al., 2020; Zhi and Li, 2020). Similar C-Q relationships have been observed for some solutes across watersheds under diverse geological and climatic conditions (Godsey et al., 2009; Basu et al., 2010; Moatar et al., 2017; Zarnetske et al., 2018; Godsey et al., 2019), whereas different solutes have shown contrasting patterns in the same watershed (Miller et al., 2017; Herndon et al., 2015; Zhi et al., 2019; Musolff et al., 2015). A general theory that can explain contrasting C-Q observations (e.g., flushing vs. dilution behaviors) under diverse watershed characteristics and forcing conditions remains elusive. The lack of understanding of mechanisms governing hydrological and biogeochemical interactions presents major roadblocks for forecasting water quality, including water issues such as eutrophication that persist worldwide.

One of the challenges to answering questions relevant to water quality and biogeochemical reactions is the lack of modeling tools that mechanistically link hydrological and biogeochemical processes at the watershed scale. Model development has been advancing separately within the disciplinary boundaries of hydrology and biogeochemistry (Li, 2019). Hydrologic models focus on solving for water storage and fluxes at the watershed scale and beyond (Fatichi et al., 2016). Reactive transport models (RTMs) have traditionally centered on transport and multi-component biogeochemical reactions typically in groundwater systems, which often have limited interactions with climate and other surficial watershed processes (Steefel et al., 2015;Li et al., 2017b;Mayer et al., 2002). Biogeochemical reactions in shallow soils that are often driven by environmental factors such as soil temperature and moisture cannot be well simulated in these models.

Previous modeling works has shown some integration across these two lines. For example, SWAT (Soil & Water Assessment Tool) (Gassman et al., 2007;Lam et al., 2010; Moriasi et al., 2013; Neitsch et al., 2011) includes a version that couples with the groundwater model MODFLOW and simulates surface water and groundwater quality in RT3D (Bailey et al., 2017; Ochoa et al., 2020). CATHY (Catchment Hydrology) includes processes of pesticide decay (Gatel et al., 2019; Scudeler et al., 2016). Some other hydrological models, including Hydrologiska Byråns Vattenbalansavdelning (HBV) and the Hydrological Predictions for the Environment (HYPE), also have modules that simulate processes relevant to nutrients and contaminants (Lindström et al., 2005;Lindström et al., 2010). While many of these models can simulate reaction processes such leaching of nutrients from agriculture lands (Lindström et al., 2005;Lindström et al., 2010;Bailey et al., 2017), most of them do not explicitly solve the multi-component reactive transport equations. In other words, they have relatively crude representations of solute leaching out of element bulk mass as part of the solute export but do not represent kinetics and thermodynamics of multi-component biogeochemical reactions typically included in reactive transport models (RTMs). They also do not simulate processes such as chemical weathering. As an example, nutrient leaching is often calculated based on empirical equations without explicitly solving reactive transport equations. Reaction rates are often represented using first-order decay (Gatel et al.,

2019), assuming reaction rate constants do not change with time and environmental conditions. However, biogeochemical processes including carbon decomposition and nutrient cycling are highly variable in space and time, depending on local environments such as substrate availability, soil temperature, and soil moisture (Li et al., 2017a;Suseela et al., 2012;HARTLEY et al., 2007). In filling in this model need, recently we augmented our watershed model RT-Flux-PIHM (Bao et al., 2017) with new developments of microbially mediated reactions, which allows us to model the interactions between biogeochemical reactions and environmental factors that are driven by land surface and hydrological processes.

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This manuscript introduces BioRT-Flux-PIHM (BioRT hereafter) v1.0, augmented based on RT-Flux-PIHM with two new additions. One is the capability of simulating biotic processes including plant uptake of nutrients, soil respiration, and microbially mediated reactions in the soil. Examples include the soil respiration that produces CO₂ and carbon decomposition that generates dissolved organic carbon (DOC), and other nutrient transformation processes such as nitrification and denitrification. The other is the introduction of an optional deeper layer below the shallow soil that enables the simulation of interactions of deep water and shallow soil water flow (Figure 1). Here the deep water is loosely defined as the water beyond the soil zone, typically in less weathered, fractured subsurface that harbors relatively old and slow-moving groundwater contributing to streams. This contrasts with shallow water in highly permeable soils. Mounting evidence in recent years has shown that deeper water beyond the shallow soil interacts with streams, introduces water with distinct chemistry, sustains base flow in dry times, and buffers climate variability (Gurdak, 2017; Green, 2016; Taylor et al., 2013; Condon et al., 2013; Anyah et al., 2008; Maxwell et al., 2011; Gleeson et al., 2015). Stream chemistry often reflects the distinct chemistry from the shallow soil and deeper groundwater zones, i.e., the so called Shallow and Deep Hypothesis (Zhi et al., 2019; Zhi and Li, 2020). Deeper groundwater is thus a fundamental component of the hydrologic cycle and water budget. The groundwater-surface water interactions can also modulate land-atmospheric energy exchanges and soil moisture dynamics (Keune et al., 2016; Martínez-de la Torre and Miguez-Macho, 2019). Including the deep water component thus enables the simulation of such interactions and the dynamics of water quality.

This paper introduces new developments in the BioRT model. The code has been verified against the widely used reactive transport code CrunchTope (Supporting Information, SI). This paper briefly overviews water and energy related processes incorporated in the model. Readers are referred to previous publications for more details of processes such as evapotranspiration (ET), hydrological flow, and abiotic reactions (Shi et al., 2013;Bao et al., 2017;Li et al., 2017a;Qu and Duffy, 2007). We showcase three examples that illustrate the new model capabilities and the simulation of biogeochemical reactions. The first hydrological example shows the surface water and groundwater interactions. This second example focuses on nitrate transformation and transport in a spatially lumped mode. The third example examines the production and export of DOC with the representation of spatial details. The source code and the examples shown here are hosted on the GitHub website (https://github.com/PSUmodeling/BioRT-Flux-PIHM).

2. Model overview

BioRT-Flux-PIHM integrates three modules (Figure 1). The Flux module is for land-surface processes including surface energy balance, solar radiation, and ET (Shi et al., 2013). The hydrology module PIHM simulates water processes including precipitation, interception, infiltration, recharge, surface runoff, subsurface lateral flow, and deep water flow (Qu and Duffy, 2007). The BioRT module simulates solute transport, bio-relevant processes such as plant uptake of nutrients from water, and multi-component reactions. The reaction processes can include soil respiration, microbially mediated redox reactions (e.g., soil respiration, carbon decomposition and nutrient transformation), ion exchange, aqueous and surface complexation, and mineral dissolution and precipitation. Note that geochemical reactions in our previous RT-Flux-PIHM are abiotic (Bao et al., 2017), including mineral dissolution and precipitation, aqueous and surface complexation, and ion exchange reactions.

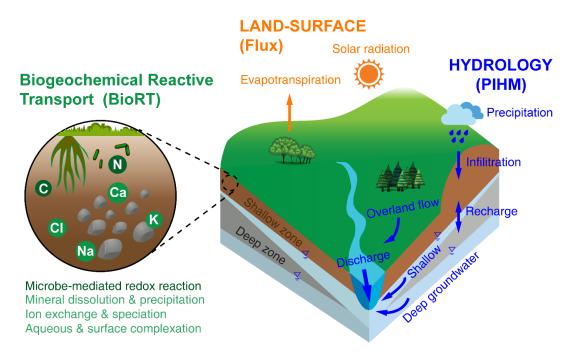


Figure 1. A conceptual diagram for processes at the watershed scale. This includes land surface interactions such as energy balance, solar radiation, evapotranspiration (e.g., evaporation, transpiration, and snow sublimation); hydrological processes partitioning water between surface runoff, shallow soil water, and deeper water entering the stream. Soil biogeochemical reactions include abiotic reactions (e.g., mineral dissolution and precipitation, ion exchange, surface complexations), and biotic processes such as soil respiration, plant uptake of nutrients, and other microbe-mediated reactions such as transformation of carbon and nitrogen. These processes are represented in three modules: The Flux module for land-surface interactions, the PIHM module for catchment hydrology, and the recently augmented BioRT module for biogeochemical reactions. Conceptually the shallow zone is the shallow soil and weathered zone that are more conductive to water flow (e.g., soil lateral flow or interflow). The deep zone refers to the less weathered zone that often harbors the relatively old and slow flowing groundwater. Reactions can occur in both shallow and deep zones. For the BioRT, the light and dark greens refer to abiotic and biological reactions, respectively.

The land surface and hydrology modules solve for soil temperature and water storage, from which water fluxes can be quantified for surface runoff, shallow and deep water fluxes. The BioRT module uses the calculated soil temperature, water storage, and water fluxes to simulate advection, diffusion/dispersion, and biogeochemical reactions in both shallow and deep zones (see governing equations in later sections). The reactions can be kinetically controlled (e.g., microbial redox reaction) or equilibrium-controlled (e.g., ion exchange, surface complexation (sorption), and aqueous complexation). Users can

define the types of reactions to be included and the form of reaction kinetics in input files. The output of BioRT includes the spatial distribution and time series of aqueous and solid concentrations, from which we can also infer reaction rates.

The simulation domain is structured as prismatic grids based on topography. Each grid is partitioned into surface and shallow and deep subsurface layers. The surface layer calculates water flow above ground (surface runoff). The shallow zone is loosely defined as the highly permeable subsurface that is contrasting to the deep zone that is broadly defined as the lower permeability zone beyond the shallow zone. In many places, this shallow zone is the soil zone that is most conductive to water flow (e.g., soil lateral flow) and is responsive to hydroclimatic forcings. The deep subsurface zone is the less weathered layer that harbors the relatively older and slower flowing ground water that contributes to stream flow. Note that these definitions differ from those in the hydrology community, which often refers to the shallow soil water flow or lumped shallow soil as groundwater, in a way that distinguishes it from the surface runoff (Winter et al., 1998; Dingman, 2015; Todd and Mays, 2005). These shallow and deep waters often have distinct solid and water chemistry, and are dominant at different time of the year, as have been observed and inferred in many catchments and watersheds (Brantley et al., 2018;Zhi et al., 2019;Zhi and Li, 2020;Li et al., 2020;Sullivan et al., 2016). Despite the model complexity, the model is flexible for taking inputs from online data portals or local measurements and it can accommodate low data availability (see the following section of 5 for data need and domain setup). The model is developed as a research tool to understand coupled watershed processes rather than as a policy model to guide management.

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3. Governing equations and processes

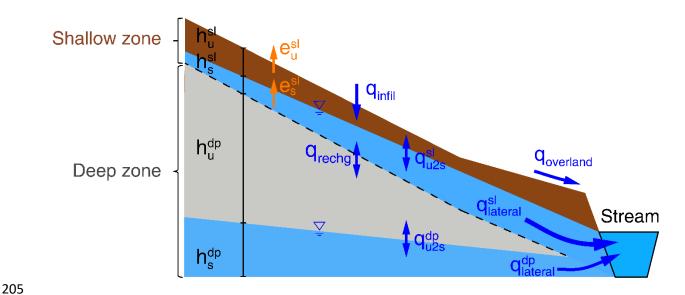


Figure 2. Hillslope view of the shallow and deep zones and relevant water flows. The symbol of "h", "e", and "q" denotes water storage [m], evapotranspiration [m/s], and water flow [m/s], respectively. The superscript letter "sl" and "dp" refer to shallow and deep zone, respectively. The subscript letters "u" and "s" refer to unsaturated and saturated layer, respectively. Detailed equations are listed in the following sections. The terms "infil", "u2s", and "recharge' refer to infiltration, unsaturated to saturated zones, and recharge.

3.1 Water equations

The original Flux-PIHM simulates surface runoff and a lumped subsurface flux into streams without distinguishing shallow soil water flow and deeper groundwater flow. In working with stream water chemistry data, we realized that a lumped subsurface flow cannot describe the dynamics of stream chemistry, as the shallow soil water and deeper groundwater have distinct chemistry and are dominant at different times of the year (Zhi et al., 2019; Zhi and Li, 2020). We therefore added an optional deeper groundwater zone in the code to simulate the deeper water that interacts with streams. Each prismatic element now has three zones in the vertical direction: surface (or above ground), shallow and deep zones in the subsurface.

In each prismatic element *i*, the shallow zone includes unsaturated and saturated water storages. The unsaturated zone receives water from the surface via infiltration and only flows vertically to the saturated zone. The saturated zone flows both vertically to the

deep zone (recharge) and laterally to neighboring grids j or the stream (lateral). The code solves the following equations for the shallow zone:

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$$\theta_i^{sl} \frac{dh_{i,u}^{sl}}{dt} = q_{i,inf} - q_{i,u2s}^{sl} - e_{i,u}^{sl} \quad (1)$$

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$$\theta_i^{sl} \frac{dh_{i,s}^{sl}}{dt} = q_{i,u2s}^{sl} - q_{i,rechg} - e_{i,s}^{sl} + \sum_{1}^{N_{ij}} q_{ij}^{sl} \quad (2)$$

Where θ_i^{sl} [m³ pore space/m³ total volume] is the shallow zone porosity in the element i; $h_{i,u}^{sl}$ and $h_{i,s}^{sl}$ [m] are the unsaturated and saturated water storage in the shallow zone, respectively. Note that the storages h here are essentially the height of soil column with equivalent saturated water, not the height of the pure water (100% volume) column. That is why porosity is in the equation. For saturation zones, this height is needed to quantify the depths of water tables and determines the direction of water flow between neighboring grids. The $q_{i,inf}$ [m/s] is the infiltration rate from the surface to the shallow zone; $q_{i,u2s}^{sl}$ [m/s] is the vertical flow from the unsaturated layer to the saturated layer in the shallow zone; $q_{i,rechg}$ [m/s] is the recharge rate from the shallow zone to the deep zone; $e_{i,u}^{sl}$ and $e_{i,s}^{sl}$ [m/s] are evapotranspiration (ET) from the unsaturated and saturated layer in the shallow zone, respectively; q_{ij}^{sl} [m/s] are the lateral flows in the shallow saturated layer between the element i and its neighbor element j; N_{ij} (≤ 3) is the number of neighbor elements i. For a prismatic element i, a boundary cell could have one or two neighbors; a non-boundary cell has three neighbors. The ET is calculated by the Penman potential evaporation scheme and detailed equations can be found in Shi (2012). A similar set of water equations for the deep zone are in the SI (Eqn. S1 and S2).

Infiltration and vertical fluxes from the unsaturated to the saturated layer in the shallow zone are based on the Richards equation, in which hydraulic water head H (i.e., the summation of water storage h and elevation head z) and hydraulic conductivity K are used to determine the fluxes in each element i:

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$$q_{i,inf} = K_{i,inf} \frac{H_{i,sur} - H_{i,u}^{sl}}{d_{i,inf}}$$
 (3)

$$q_{i,u2s}^{sl} = K_{i,v}^{sl} \frac{H_{i,u}^{sl} - H_{i,s}^{sl}}{0.5d_i^{sl}}$$
 (4)

Where $d_{i,inf}$ and d_i^{sl} [m] are the thickness of infiltration layer and shallow zone depth for the elements i, respectively; $K_{i,inf}$ [m/s] is the hydraulic conductivity of the infiltration layer, the top 0.1 m of the subsurface and is considered to have different conductivity from the rest of subsurface; $K_{i,V}^{sl}$ [m/s] is the hydraulic conductivity in the vertical direction (i.e., weighted average of macropore $K_{i,macV}$ and soil matrix $K_{i,satV}$, Eqn. S7); $H_{i,sur}$ [m] is the surface hydraulic water head (= $h_{i,sur} + z_{i,sur}$); $H_{i,u}^{sl}$ and $H_{i,s}^{sl}$ [m] are the shallow hydraulic water heads in the unsaturated and saturated layer, respectively. The lateral flow in the shallow saturated layer is calculated using Darcy's law:

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$$q_{ij}^{sl} = K_{ij}^{sl} \frac{H_{i,s}^{sl} - H_{j,s}^{sl}}{d_{ij}}$$
 (5)

Where d_{ij} [m] is the distance between the centers of elements i and j; K_{ij}^{sl} [m/s] is the harmonic mean of shallow hydraulic conductivity in the horizontal direction between elements i ($K_{i,H}^{sl}$) and j ($K_{j,H}^{sl}$). The interaction between the shallow saturated zone and stream channel also follows Eqn. 5, except that the adjacent head is replaced by the level of the channel water. Similar to the shallow zone, hydrological equations in the deep zone are detailed in the SI (Eqn. S1 – S8).

3.2 Reactive transport equations

The governing advection dispersion reaction (ADR) equation for an arbitrary solute m in grid i is as follows (Bao et al., 2017), i.e., the change of solute mass (i.e., the left term in Eqn. 6) is driven by dispersive transport, advective transport, and reactions (i.e., the 1st, 2nd, and 3rd right-hand side terms, respectively):

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$$V_{i}\frac{d(S_{w,i}\theta_{i}C_{m,i})}{dt} = \sum_{1}^{N_{ij}} \left(A_{ij}D_{ij}\frac{C_{m,j} - C_{m,i}}{d_{ij}} - q_{ij}A_{ij}C_{m,j}\right) + R_{m,i}, \qquad m = 1, ..., nm \quad (6)$$

Where V_i [m³ total volume] is the total volume of element i; $S_{w,i}$ [m³ water/m³ pore space]

is soil water saturation; θ_i [m³ pore space/m³ total volume] is porosity; $C_{m,i}$ [mol/m³ water] is the aqueous concentration of species m; N_{ij} is the number of fluxes from neighbor element j for element i, N_{ij} is 2 for the unsaturated zone (infiltration, recharge) with only vertical flows and 5 for saturated zone with flux from (or to) the unsaturated zone, from (or to) the deeper zone, and fluxes between i and three neighbor elements j in lateral flow directions for non-boundary grids; A_{ij} [m²] is the grid area shared by i and its neighbor grid j; D_{ij} [m²/s] is the hydrodynamic dispersion coefficient (i.e., sum of mechanical dispersion and effective diffusion coefficient) normal to the shared surface A_{ij} ; d_{ij} [m] is the distance between the center of i and its neighbor elements j; q_{ij} [m/s] is the flow rate across A_{ij} ; $R_{m,i}$ [mol/s] is the total rate of kinetically controlled reactions in element i that involve species m; nm is the total number of independent primary species to be solved for reactive transport equations.

3.3 Biogeochemical processes and reaction kinetics

3.3.1 Biogeochemical processes

Here we discuss some representative biogeochemical processes that involve plants and microbes that can be included in BioRT. BioRT differs from general water quality models that often primarily target a few contaminants (e.g., N, P, metals). The framework of the code is flexible and the users can define their reactions and solutes of interests in the input files. For abiotic reactions such as mineral dissolution and surface complexation or ion exchange, readers are referred to an earlier paper (Bao et al. (2017). Generally speaking, shallow soils compared to deep soils contain more weathered materials and soil organic matters (SOM) including roots, leaves, and microbes. SOM can be decomposed partially into organic molecules that dissolve in water (Wieder et al., 2015), i.e., DOC, or it can be oxidized completely into CO₂ that is released back to the atmosphere as a gas (Davidson, 2006) or surface water in the form of dissolved inorganic carbon (DIC). With coexisting cations (e.g., Ca, Mg), DIC can often precipitate and become carbonate minerals (e.g., CaCO₃).

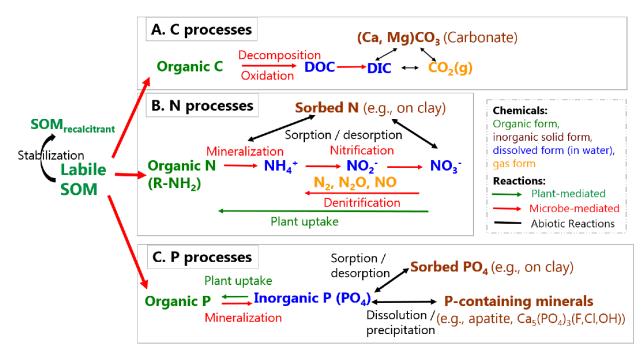


Figure 3. Biotic and abiotic reactions relevant to the transformation of soil organic matter (SOM). It can become stabilized through sorption on clay and separation from reactants. Labile OM can decompose into inorganic forms, releasing C, N, and P that further transform between different forms (adopted from Li (2019), permission with Mineralogical Society of America).

OM decomposition releases organic nitrogen (R-NH₂), which can further react to become NH₄+, and other nitrogen forms (N₂, N₂O, NO, NO₂-, NO₂) (Figure 3). The gases can be emitted back to the atmosphere (Saha et al., 2017;Maavara et al., 2018). Denitrification requires anoxic conditions and occurs less commonly in shallow soils owing to the pervasive presence of O₂ (Sebestyen et al., 2019); it can become important under wet conditions and in O₂-depleted groundwater systems. Phosphorous (P) can be in an organic form (e.g., leaves), sorbed on fine soil particles, dissolved in water, or in solid forms as P-containing minerals. Transformation of nutrients occurs through various bio-mediated or abiotic reactions. A representative P-containing mineral in the Earth's crust is apatite Ca₅(PO₄)₃(F, CI, OH). Once liberated via rock dissolution, P is mostly biologically assimilated and locked in organic forms. These organic forms have very low solubility, allowing them to bind on and be transported together with soil particles in the form of orthophosphate or pyro-diphosphate. Overall, these reactions are a combination of biotic and abiotic reactions.

3.3.2 Reaction kinetics in natural soils

Rate dependence on temperature and soil moisture. Reactions such as soil respiration and plant uptake typically depend on environmental conditions (temperature or soil moisture). For example, in shallow oxic soils where organic carbon and O₂ are often abundant, the rate law for carbon decomposition can be simplified to the following form assuming microorganism concentrations are relatively constant.

$$r = kAf(T)f(S_w)f(Z_w)$$
 (7)

Where the reaction rate r [mol/s] depends on rate constant k [mol/m²/s], the surface area A [m²] is a lumped parameter that quantitatively represents SOM content and biomass abundance, f(T) and $f(S_w)$ describe the temperature and soil moisture dependence, respectively, $f(Z_w)$ can be included to account for the depth distribution of SOM (Seibert et al., 2009), and Z_w [m] is the water table depth. An example for the depth distribution is $f(Z_w) = \exp\left(-\frac{Z_w}{b_m}\right)$ (Weiler and McDonnell, 2006;Ottoy et al., 2016;Bai et al., 2016), with b_m as the depth coefficient describing the gradient of SOM content over depth. Users can choose to include either one or all of these dependences in input or database files.

The temperature dependence follows a Q_{10} -based form (Friedlingstein et al., 2006;Hararuk et al., 2015) as follows:

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$$f(T) = Q_{10}^{|T-20|/10}$$
 (8),

where Q_{10} is the relative increase in reaction rates when temperature increases by 10 °C (Davidson and Janssens, 2006). Values of Q_{10} (Figure 4a) can vary from 1.0 to 3.0, depending on climatic conditions, substrate availability, and ecosystem type (e.g., grassland, forest) (Davidson et al., 2006;Liu et al., 2017). The mean values are in the range of 1.4 to 2.5 (Zhou et al., 2009;Bracho et al., 2016). The Q_{10} value can be specified in the input file. The soil moisture dependence function $f(S_w)$ is coded based on the following form:

$$\begin{cases}
 \left(\frac{S_w}{S_{w,c}}\right)^n, S_w \leq S_{w,c} \\
 \left(\frac{1-S_w}{1-S_{w,c}}\right)^n, S_w > S_{w,c}
\end{cases} \tag{9}$$

Here $S_{w,c}$ [0 to 1] is the critical soil moisture at which rates are highest, and n is the exponent reflecting the dependence of rates on soil moisture. A typical n value is 2 (Yan et al., 2018) with a range between 1.2 and 3.0 (Hamamoto et al., 2010), depending on soil structure and texture. As shown in Figure 4b, the form indicates an intermediate critical soil moisture $S_{w,c}$ at which $f(S_w)$ reaches its maximum. When S_w is below this value, $f(S_w)$ increases with S_w ; when S_w is above this value, $f(S_w)$ decreases with S_w behavior (Figure 4b) (Yan et al., 2018). Under the extreme conditions of $S_{w,c}$ equals to 0 or 1, $f(S_w)$ monotonically increase or decrease (Figure 4c). The two parameters, $S_{w,c}$ and n, determines the shape of the curve. They can be specified in input or database files. One can also choose not to have temperature or soil moisture dependence by choosing parameters that would lead to the value of exponent being zero.

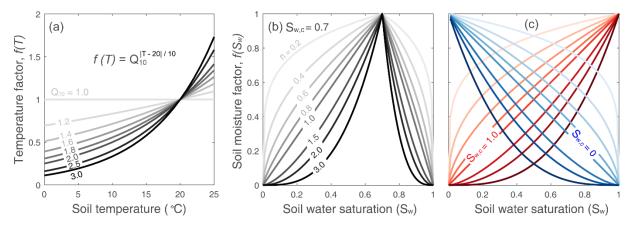


Figure 4. (a) Function form of soil temperature dependence and (b, c) soil moisture dependence for reaction rates. The temperature factor f(T) is a function of the Q_{10} (defined by users) and soil temperature. The soil moisture factor $f(S_w)$ is a function of two user-defined parameters $S_{w,c}$ and n and soil water saturation S_w . The soil moisture function can represent three types of behaviors: the threshold behavior (b, $0 < S_{w,c} < 1$), increase behavior (red in (c), $S_{w,c} = 1$), and decrease behavior (blue in (c), $S_{w,c} = 0$). Values of n = 1 leads to a linear threshold dependence of S_w while n < 1 and n > 1 lead to concave and convex dependences, respectively.

Rate dependence on substrates: Monod kinetics and biogeochemical redox ladder. Deeper groundwater aquifers often experience anoxic conditions that lead to processes such as denitrification or methanogenesis. This can also happen in wetlands or wet soils. The rates of microbe-mediated redox reactions depend not only on temperature and soil moisture as discussed above, they also depend on concentrations

of electron donors and non-oxygen electron acceptors (e.g., nitrate, iron oxides, sulfate) that are often limited under anoxic conditions (Bao et al., 2014;Li, 2019). The order of the redox reactions often follows the biogeochemical redox ladder, which is based on how much microbe can harvest energy by reducing different types of electron acceptors. Monod reaction rate laws are often used for quantifying rates of these redox conditions. These rate laws are detailed in the section S2 of Supporting Information and also in Li (2019). Users can combine these Monod rate laws and the temperature and soil moisture dependence described above, if needed.

3.4 Plant related processes: root uptake of nitrate as an example

Nitrate uptake by plants is intrinsically complex and remains poorly understood (Devienne-Barret et al., 2000;Crawford and Glass, 1998;Hachiya and Sakakibara, 2016). A variety of plant uptake models exists with varying degrees of complexity (Neitsch et al., 2011;Fisher et al., 2010;Cai et al., 2016). These models are mostly based on plant growth module or supply and demand approach that often requires detailed phenological and plant attributes such as growth cycle, root age and biomass, nitrate availability, phosphorous stress, and carbon allocation, in addition to local climate conditions such as temperature and soil moisture (Neitsch et al., 2011;Porporato et al., 2003;Dunbabin et al., 2002;Buysse et al., 1996;Fisher et al., 2010). Without detailed information, we can assume a simple and operational approach (Eqn. 13 and 14). In the Example 2 that we show later, for example, we modeled nitrate uptake with dependence on NO₃-concentration, soil temperature and moisture, and rooting density (McMurtrie et al., 2012;Yan et al., 2012;Buljovcic and Engels, 2001).

$$r_{uptake} = k_{uptake} C_{NO_3}^- f(T) f(S_w) f_{root}(d_w)$$
 (13)
$$f_{root}(d_w) = \exp((-d_w + \delta) / \lambda)$$
 (14)

Where k_{uptake} [L/s] is the nitrate uptake rate, $f_{root}(d_w)$ is the normalized rooting density term in the range of 0 to 1 as a function of water depth to the groundwater (d_w) . The rooting term (Eqn. 14) was exponentially fitted ($\delta = 0.013, \lambda = 0.20$) based on field measurements of root distribution along depth (Hasenmueller et al., 2017). It is common to observe root density decrease exponentially in forests (López et al., 2001). Other form

of user-tailored plant uptake rate law can be added if needed.

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4. Numerical scheme and model verification

The system of differential equations for the water storages (e.g., Eqn. 1 and 2, and Eqn. S1 and S2) are assembled into a global system of ordinary differential equations (ODEs) and solved in CVODE (short for C-language Variable-coefficients ODE solver, https://computing.llnl.gov/projects/sundials/cvode), a numerical ODE solver in the SUite of Nonlinear and Differential / ALgebraic equation Solvers (SUNDIALS) (Hindmarsh et al., 2005). In BioRT, the transport step is first solved with water by the preconditioned Krylov (iterative) method and the Generalized Minimal Residual Method (Saad and Schultz, 1986). In the following reaction step, all primary species in each finite volume are assembled in a local matrix and then solved iteratively by the Crank-Nicolson and Newton-Raphson method in CVODE (Bao et al., 2017).

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Model verification. The BioRT module had been verified against CrunchTope under different transport and reaction conditions (Figures S1 – S7 in SI). CrunchTope is a widely used subsurface reactive transport model (Steefel and Lasaga, 1994; Steefel et al., 2015), and is often used as a benchmark to verify other reactive transport models. Verification was performed under simplified hydrological conditions with 1-D column and constant flow rates such that it focuses on advection, diffusion, dispersion, and biogeochemical reactions. Specifically, three cases of soil phosphorus, carbon, and nitrogen were verified for temporal evolution and spatial pattern of relevant solute concentrations. The phosphorus that involves kinetics-controlled case apatite dissolution and thermodynamics-controlled phosphorous speciation was first tested for solution accuracy of the bulk code that was inherited from the original RT-Flux-PIHM. Soil carbon and nitrogen processes that involve microbe-driven processes, such as soil carbon decomposition and mineralization, nitrification and denitrification, were further verified for solution accuracy of the augmented BioRT module.

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5. Model structure, data needs, and domain setup

Model structure. The model takes meteorological forcing time series as input and solves for water storages and soil temperature, along with other hydrologic and land surface states and fluxes (Figure 5). BioRT reads in the model output of water and temperature from Flux-PIHM, and solves the biogeochemical reactive transport equations. At the time scale of months to years that are typical for BioRT-Flux-PIHM simulations, alterations in solid phase properties, including, porosity, permeability, and reactive surface area, are considered negligible such that hydrological parameters remain constant with time.

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Data needs. The code sets up the model domain based on watershed characteristics including topography, land cover, and shallow and deep zone properties (Figure 5). When the model is used in a spatially distributed form, the model domain can be set up using elevation, land cover, soil and geology maps supplied by the user or from the data portal of Geospatial Data Gateway (https://datagateway.nrcs.usda.gov). The meteorological forcing data can be downloaded from the North American Land Data Assimilation Systems Phase 2 (NLDAS-2, https://ldas.gsfc.nasa.gov/nldas/v2/forcing). The vegetation forcing, i.e., Leaf Area Index (LAI), can be obtained from MODIS (Moderate Resolution Imaging Spectroradiometer, https://modis.gsfc.nasa.gov/data). Other vegetation properties associated with land cover (e.g., shading fraction, rooting depth) can be adopted from, for example, the Noah vegetation parameter table embedded in the Weather Research and Forecasting model (WRF; Skamarock and Klemp (2019)). Local measurements from meteorological stations and field campaigns (e.g., land cover, soil, geology) can also be used in the model. Another data source is the HydroTerre (http://www.hydroterre.psu.edu/), where users can obtain geospatial data (e.g., elevation, land cover, geology, soil) (Leonard and Duffy, 2013). Initial water and solid phase chemistry can be based on measurements or general knowledge of the simulated sites. The form of reaction rate laws, including the Monod form, temperature and soil moisture dependence, can be defined in the input files and calibrated to reproduce field data. Reaction thermodynamics, mostly equilibrium constants, are from the geochemical database EQ3/6 by default (Wolery, 1992). These reaction parameters can be modified when necessary. The model outputs include aqueous and solid concentrations of shallow and deep zone and stream water.

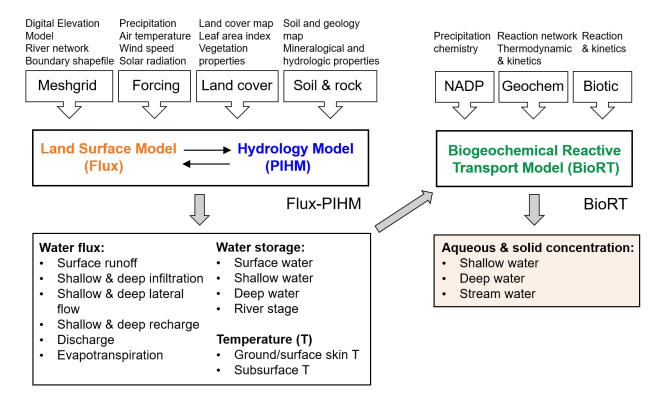


Figure 5. Model structure, input, and output of BioRT-Flux-PIHM. The Flux-PIHM takes in watershed characteristics including topography (digital elevation model, DEM), land cover, shallow and deep zone properties, and meteorological forcing time series and solves for water storage, and ground and soil temperature. BioRT takes in water- and temperature-related information from Flux-PIHM with additional inputs such as precipitation chemistry and shallow and deep water chemistry and biogeochemical kinetics parameters, and solve for aqueous and solid concentrations in the shallow and deep zone, and stream water. NADP stands for the National Atmospheric Deposition Program. This paper focuses on the BioRT component. The land-surface, hydrological processes, and abiotic reactive transport processes have been described in previous papers (Bao et al., 2017;Shi et al., 2013). Discussions on how air temperature and ET influence stream chemistry can be found in Li (2019).

Domain set up: from simple, spatially lumped to complex, spatially distributed domains. The domain can be set up at different spatial resolutions with different numbers of grids. A simple domain can be set up with only two land grids representing two sides of a watershed connected by one river cell. This setup uses averaged properties without needs for larger spatial data. Alternatively, a complex domain can be set up using many grids with explicit representation of spatial details. The model domain can be set up using PIHM-GIS (http://www.pihm.psu.edu/pihmgis_home.html), a standalone GIS interface for watershed delineation, domain decomposition, and parameter assignment (Bhatt et al.,

2014). It requires much more data and can be computationally expensive but can be used to identify "hot spots" of biogeochemical reactions within a watershed. The same model processes (e.g., hydrology, reaction network) can be setup in both spatial configurations. Auto-calibration is not built into the model, but a global calibration coefficient approach is used to reduce parameter dimension and facilitate manual calibration. A typical model application requires 20 to 30 hydrological parameters to be calibrated. These parameters include land surface parameters (e.g., canopy resistance, surface albedo), soil and geology parameters (e.g., hydraulic conductivity, porosity, Van Genuchten, macropore properties) (Shi et al., 2013). Reaction-related parameters (e.g., reaction rate constant, mineral surface area, Q₁₀, S_{w,c}, and n) are additionally needed for calibration, the number of which depends on the numbers of reactions involved in a particular system.

6. Model applications

The original RT-Flux-PIHM has been applied to understand processes related to the geogenic solutes of CI and Mg at the Shale Hills watershed and for Na in a watershed on Volcán Chimborazo in the Ecuadorian Andes (Table 1). The new BioRT-Flux-PIHM has been demonstrated for understanding the dynamics of DOC and nitrate at Shale Hills and Coal Creek. This section will present one hydrology and two biogeochemical examples in the Susquehanna Shale Hills Critical Zone Observatory (SSHCZO), a small headwater watershed in central Pennsylvania, USA. The mean annual precipitation is approximately 1,070 mm and the mean annual temperature is 10 °C (Brantley et al., 2018). Soil carbon storage and respiration and nitrogen budget and fluxes have been studied in detail (Andrews et al., 2011;Hasenmueller et al., 2015;Shi et al., 2018;Hodges et al., 2019;Weitzman and Kaye, 2018). Modeling work has been conducted to understand hydrological dynamics (Shi et al., 2013;Xiao et al., 2019), transport of the non-reactive tracer CI, and the weathering-derived solute Mg (Bao et al., 2017;Li et al., 2017a).

 Table 1. Existing Model applications of BioRT-Flux-PIHM

| Watershed | Size | Model | Modeled | Reactions (rate laws: 1, TST; | Reference |
|------------|----------|--------|---------|-------------------------------|-----------|
| (location) | (km^2) | domain | solutes | 2, Monod based; 3, plant | |
| | | | | uptake) | |

| | | Spatially distributed | CI, Mg | Chlorite dissolution¹ Bao et al., Illite dissolution¹ 2017; Li et |
|-------------------------|------|-----------------------|-----------------------------|--|
| | | | | Carbonate dissolution & al., 2017 precipitation¹ |
| Shale Hills | 0.00 | | | Cation exchange |
| (PA, USA) | 0.08 | Spatially | DOC | SOC decomposition² Wen et al., |
| | | distributed | | • DOC sorption 2020 |
| | | Spatially | NO_3^- | Soil N leaching² This work |
| | | lumped | | Denitrification² |
| | | | | Plant uptake³ |
| Coal Creek | 53 | Spatially | DOC, Na | SOC decomposition² Zhi et al., |
| (CO, USA) | | lumped | | DOC sorption 2019 |
| | | | | Albite dissolution¹ |
| Volcán | | Spatially | Cl, Na, | Albite dissolution¹ Saberi et al. |
| Chimborazo (Ecuador) | | distributed | Ca, Mg, SiO ₂ | Diopside dissolution¹ (under review) |

Note: Transition State Theory (TST) is a classic kinetic rate law for mineral dissolution and precipitation (Brantley et al., 2008) (Eqn. S15); Monod rate law with environmental dependency (i.e., soil temperature and soil moisture) is widely used for microbially driven reactions. Monod-based and plant nitrate uptake rate law are detailed in the following section of 6.2. SOC stands for soil organic carbon.

6.1 Example 1: Shallow and deep water interactions

The model was set up using the spatially lumped mode with two land grids and one river grid characterized by average land cover, soil and rock properties based on previous work (Shi et al., 2013;Kuntz et al., 2011). The model assumed a Weikert soil, the dominant soil type at Shale Hills (Shi et al., 2013). The porosity of the deep zone was set to a tenth of the shallow soil porosity based on measurements of the groundwater aquifer (Brantley et al., 2018;Kuntz et al., 2011). In a headwater catchment like Shale Hills where the deep groundwater is most likely sourced from recharge, the deep groundwater contribution to the stream can be primarily controlled by the hydraulic conductivity (K_{satH}) contrast between the deep and shallow zones (i.e., $K_{satH}^{dp} / K_{satH}^{sl}$). This is because the K_{satH} contrast determines the partitioning of infiltrating water between the shallow lateral flow and the downward recharge to the deep zone and then deep groundwater flow. Two cases of high (red) and low (blue) K_{satH}^{dp} were set up to showcase

the control of K_{satH} contrast on deep groundwater (Figure 6a). By changing the deep zone K_{satH}^{dp} from 2.6 to 0.22 (m/d), the annual deep groundwater (Q_G) contribution to discharge (Q) decreased from 26% to 5.2%, although the total discharge is negligible. This indicates that the changing K_{satH}^{dp} mostly changes the portioning between the shallow soil lateral flow and recharge, whereas total infiltration and discharge remain very similar.

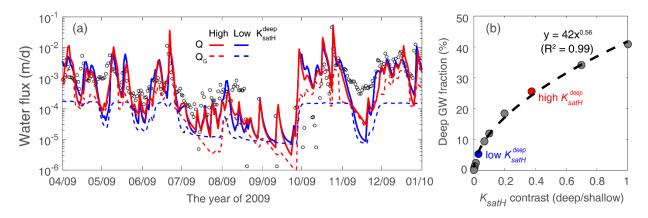


Figure 6. (a) Hydraulic conductivity (K_{satH}) contrast control on deep groundwater (Q_G). The cases of high ($K_{satH}^{dp} = 2.6 \ m/d$, red) and low conductivity ($K_{satH}^{dp} = 0.22 \ m/d$, blue) led to 26% and 5.2% of annual Q_G contribution to discharge (Q_G), respectively. (b) Deep groundwater fraction as a function of K_{satH} contrast between the deep and shallow zone. The K_{satH} contrast was limited to 1 in the figure as most watersheds exhibit a smaller K_{satH} in the deep zone than in the shallow zone. The two red and blue dots correspond to the two cases in left panel.

Several additional cases were further tested to examine the relationship between deep groundwater fraction (%) of discharge and K_{satH} contrast. Figure 6b shows that the deep groundwater fraction rapidly increases with the increasing ratio of K_{satH}^{dp} / K_{satH}^{sl} , reaching a limit when K_{satH} contrast is sufficiently high. The deep groundwater contribution to the stream reaches ~ 40% when the K_{satH}^{dp} and K_{satH}^{sl} are equal. In natural systems, we do see places, for example, karst formations, where groundwater contributes to more than 40% (Hartmann et al., 2014;Husic, 2018). These places may have higher deeper conductivity than shallow soils due to the development of highly conductive conduits.

6.2 Example 2: Nitrate dynamics in a spatially implicit domain

This example focuses on nitrate (NO₃-), a dominant dissolved N form in water with abundant measurements (https://criticalzone.org/shale-hills/data/datasets/) (Weitzman and Kaye, 2018). The N processes at Shale Hills include atmospheric N deposition, soil N leaching, stream export, denitrification, and plant uptake (Figure 7). Based on field measurements, the atmospheric deposition at the site is the dominant N input and N export via discharge is only a small fraction (2.5%) of atmospheric N input, indicating most deposited N is tightly cycled by plants or lost to the atmosphere via denitrification.

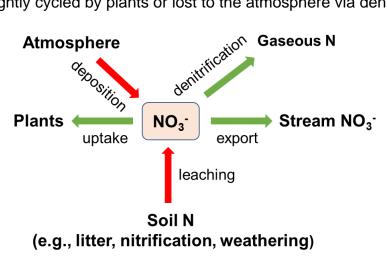


Figure 7. Modeled nitrogen processes in Example 2. Atmospheric N deposition is the major N input; denitrification and plant uptake are the major N loss and sink. Export via discharge only occupies a small fraction.

The soil N leaching process was represented using a lumped reaction that generates NO₃-. Conceptually this could represent the total rates of reactions including the decomposition of soil organic matter (SOM), nitrification, and rock weathering that generates NO₃-. Its rate was assumed to depend on soil temperature and moisture and follows the equation $r_{leach} = kAf(T)f(S_w)$, where r_{leach} [mol/s] is the leaching rate, $k = 10^{-9.7}$ [mol/m²/s] is the leaching rate constant (Regnier and Steefel, 1999), and A [m²] is the surface area that represents the contact area between substrates and N transforming microbe, and f(T) and $f(S_w)$ are soil temperature (Eqn. 8) and soil moisture (Eqn. 9) functions, respectively. The surface area was calculated based on SOM volume fraction [m³/m³], specific surface area (SSA, [m²/g]), substrate density [g/cm³], and element volume [m³]. Denitrification converts NO₃- to N₂ gas under anaerobic conditions. Here this process was modeled by the Monod rate law with DOC as the electron donor (Di Capua

et al., 2019), NO_3^- as the electron acceptor, and with an inhibition term $f(O_2)$ (Eqn. S13). The reaction rate: $r_{denitrification} = kA \frac{c_{DOC}}{K_{m,DOC} + C_{DOC}} \frac{c_{NO_3^-}}{K_{m,NO_3^-} + C_{NO_3^-}} f(O_2) f(T) f(S_w)$, where k = 110⁻¹⁰ [mol/m²/s] is the denitrification rate constant (Regnier and Steefel, 1999), halfsaturation constants $K_{m,DOC} = 15 [uM]$ and $K_{m,NO_3} = 45 [uM]$ (Regnier and Steefel, 1999; Billen, 1977). For soil N leaching and denitrification, the SSA were respectively tuned as 1.6×10^{-6} and 7.5×10^{-5} [m²/g] to reproduce observed stream nitrate dynamics. The calibrated values were orders of magnitude lower than the lab measured SSA of natural materials (e.g., SOM, $0.6 \sim 2 \text{ m}^2/\text{g}$) (Rutherford et al., 1992; Chiou et al., 1990). Such discrepancies between calibrated effective reactive surface area (i.e., solid-water contact area) and lab measured absolute surface area are consistent with other observations in literature (Li et al., 2014; Heidari et al., 2017). The uptake rate constant was calibrated by constraining the partitioning of N transformation flux between denitrification and plant uptake by the ratio of 1:5, a value estimated from field measurements of gaseous N outputs (3.53 kg-N/ha/yr) and plant N uptake (18.3 kg-N/ha/yr) (Weitzman and Kaye, 2018). The uptake rate constant in the deep zone (> 2 m in depth) was considered negligible (Weitzman and Kaye, 2018; Hasenmueller et al., 2017). Groundwater nitrate was initialized as 0.43 mg/L, the average of measured groundwater concentration during 2009 - 2010.

Temporal nitrate dynamics. Three cases were set up to understand and quantify the effects of different processes in determining nitrate dynamics (Figure 8a). The *transport-only* case (dashed line, *tran*) simulates nitrate input from precipitation (at 1.4 ± 0.96 mg/L, based on the 2009 data of NADP PA42 site) and N transport but without any reactions. It overestimated stream nitrate data (0.33 ± 0.39 mg/L) throughout the year. The *transport* + *N reactions* case (gray line, *tran* + *N react*) has denitrification and soil N leaching processes but not plant uptake. These two reactions lowered the nitrate concentration slightly, as these two processes compensate each other in adding and removing nitrate from water. The *transport* + *N reactions* + *uptake* case (thick black line, *tran* + *N react* + *upta*) have all processes. It significantly lowered the nitrate concentration, especially in April-May and October-December. Nitrate peaks from May to July, exhibiting comparable

levels of high precipitation nitrate concentration (Figure 8b). It is noticeable that the three cases almost overlapped at these overestimated short nitrate peaks, suggesting nitraterich precipitation may not be routed into the subsurface where denitrification and plant uptake could occur.

Although precipitation from April to August accounted for 70% of the total simulation period, larger storm events in October contributed more to the export. Deeper groundwater had higher nitrate concentration than shallow water, because most plant uptake occurred in the shallow zone. The nitrate fluxes into the deeper zone however only contributed 26% of stream nitrate export at the annual scale, due to the relatively small groundwater contribution (9.5%) to the stream. Denitrification and plant uptake largely occurred during the wet spring period, which is part of the leaf-on season. Denitrification peaks often appeared after major storm events. Comparing the three outfluxes (Figure 8c), nitrate export via discharge (red) was negligible compared to denitrification (blue) and plant uptake (green). At the annual scale, stream export only accounted for 9.5% outfluxes, whereas denitrification and plant uptake took up 15% and 75% of deposited NO₃-, respectively. In other words, as Nitrate enters this system via precipitation, plant uptake can play a significant role in reducing nitrate level, indicating precipitated nitrate is tightly cycled in the system.

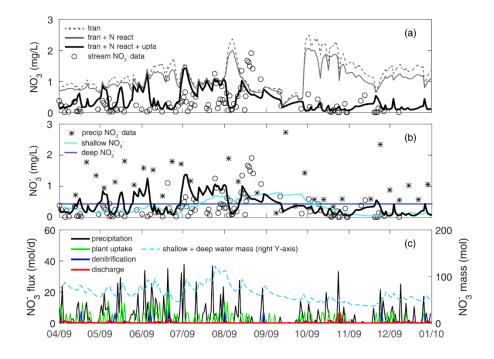


Figure 8. Stream nitrate dynamics and fluxes at Shale Hills in Example 2. (a) stream nitrate dynamics in three simulation conditions with different processes: *transport-only* (dashed line, *tran*), *transport + N reaction* (gray line, *tran + N react*), *transport + N reaction + plant uptake* (thick black line, *tran + N react + upta*), where N reactions include both nitrate leaching and denitrification (see Figure 7); (b) nitrate concentration in precipitation, shallow and deep water; (c) nitrate fluxes and budget. Note the nitrate leaching was ignored in (b) due to its minimal flux as precipitation N deposition was as the dominant input source (Weitzman and Kaye, 2018).

6.3 Example 3: DOC production and export in a spatially distributed domain

This example showcases the application of BioRT-Flux-PIHM in a spatially distributed mode. This work has been documented with full details in Wen et al. (2020). Here we only introduce some key features and capabilities in the spatially distributed mode. The Shale Hills catchment was discretized into 535 prismatic land elements and 20 stream segments through PIHMgis based on the topography (Figure 9a). The heterogeneous distributions of soil depth and solid organic carbon within the domain (Figure 9b-c) were interpolated through ordinary kriging based on field surveys (Andrews et al., 2011;Lin, 2006). Other soil and mineralogy properties such as hydraulic conductivity, van Genuchten parameters, and ion exchange capacity were spatially distributed following intensive field measurements (Jin and Brantley, 2011;Jin et al., 2010;Shi et al., 2013) (criticalzone.org/shale-hills/data/).

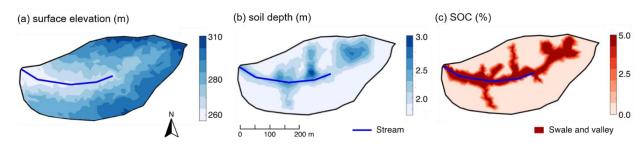


Figure 9. Attributes of Shale Hills in the spatially distributed mode in Example 3: (a) surface elevation, (b) soil depth, and (c) soil organic carbon (SOC). The surface elevation was generated from lidar topographic data (criticalzone.org/shale-hills/data); Soil depths and SOC were interpolated using ordinary kriging based on field surveys (Andrews et al., 2011;Lin, 2006). The SOC distribution in (c) was further simplified using the high, uniform SOC (5% v/v) in swales and valley soils based on field survey (Andrews et al., 2011). Swales and valley floor areas were defined based on surface elevation via field survey and a 10 m resolution digital elevation model (Lin, 2006).

Temporal and spatial patterns of DOC production and export. The model outputs followed the general trend of stream DOC measurements (NSE = 0.55 for monthly DOC concentration; Figure 10a), with high values (~15 mg/L) in the dry periods (July-September). The model enabled the identification of spatial patterns and hot spots of reactions. In May when soil water is relatively abundant, the valley and swales with deeper soils (Figure 10b) generally tended to be wetter compared to the hillslope and ridgetop, and were hydrologically connected to the stream (Figure 10b, c). The distribution of local DOC production rate r_{DOC} and DOC concentration followed that of SOC (Figure 10c) and water content (Figure 10b). Low r_{DOC} in relatively dry planar hillslopes and uplands resulted in low soil water DOC. The average stream DOC (~5 mg/L) reflected soil water DOC in the valley and swales.

In August, the hydrologically-connected zones with high water content shrank to the vicinity of the stream and river bed. With high temperature in summer, r_{DOC} increased by 2-fold from May across the whole catchment while still exhibited the highest values in the SOC-rich regions. Soil water DOC concentration increased by a factor of 2 because the produced DOC was trapped in low soil moisture areas that were not hydrologically connected to the stream. In the north side with low water content (Figure 10b), the soil water DOC (~7 mg/L in average) accumulated more than the south side (~5 mg/L in average). The high shallow water DOC (~10 mg/L) in the stream vicinity dominated the stream DOC in August.

In October, precipitation wetted the catchment again. The hydrologically connected zones expanded beyond swales and the valley to the upland hillslopes (Figure 10c). The increase in hydrological connectivity zones favored the mixing of shallow water DOC sourced from upland hillslopes (low DOC), swales, and valley (high DOC) into stream rather than only from the stream vicinity with high DOC in the dry August, leading to a drop in stream DOC.

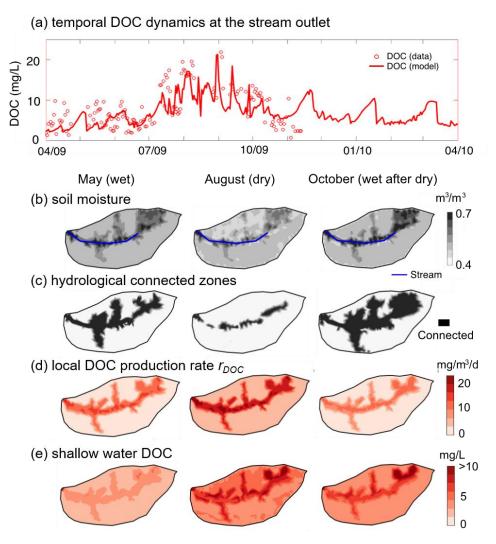


Figure 10. (a) Temporal dynamics of stream DOC concentration; spatial profiles of (b) shallow soil moisture, (c) hydrologically connected zones, (d) local DOC production rates r_{DOC} and (e) shallow water DOC concentration in May (wet), August (dry), and October (wet after dry) of 2009. The soil DOC and r_{DOC} were high in swales and valley with relatively high shallow water and SOC content. August had the highest shallow water DOC concentration compared to May and October, because most DOC accumulated in zones that are disconnected to the stream.

C-Q patterns. The DOC C-Q relationship showed a non-typical pattern with flushing first and transitioning into a dilution pattern, with an overall C-Q slope b = -0.23 (Figure 11). At low discharges (< 1.8×10^{-4} m/d) in the summer dry period, the stream DOC mainly came from the organic-rich swales and valley floor zones with high soil water DOC (Figure 10e). With discharge increasing in wetter period (i.e., spring and fall), the contribution

from planar hillslopes and uplands with lower DOC concentration increased (Figure 10e), leading to the dilution of stream DOC.

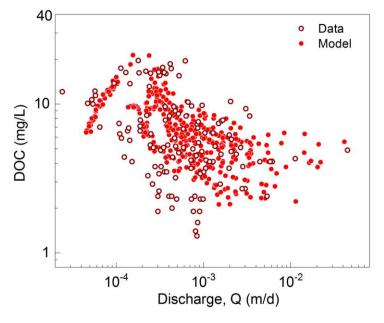


Figure 11. Relationships of daily discharge (*Q*) with stream DOC concentration. With the increase of *Q*, the stream water first shifted from the dominance of groundwater with low DOC at very low discharge to the predominance of organic-rich soil water from swales and valley at intermediate discharge. As the discharge increases further, the stream water switches to the dominance of high flow with lower DOC water from planar hillslopes and uplands, resulting in a dilution C-Q pattern (modified from Wen et al., 2020).

7. Discussion

The watershed biogeochemical model BioRT-Flux-PIHM brings the reactive transport modeling capabilities to the watershed scale, enabling the simulation of subsurface shallow and deep flow paths and biogeochemical reactions influenced by hydroclimatic conditions and land-surface interactions. The expanded model capability of simulating bio-mediated processes such as plant uptake, soil respiration, and microbe-mediated redox reactions enables the simulation of carbon and nutrient cycling in the shallow subsurface. The inclusion of the deep groundwater zone allows the exploration of the effects of subsurface structure on hydrological partitioning between shallow soil lateral flow and deep groundwater, and their relationships with stream discharge. Although not shown here, the model can also simulate deeper groundwater coming from regional aquifers across the outer boundary. This can be particularly useful for

watersheds of higher stream orders, where a large proportion of deep water may come from nearby regional aquifers.

The model presented here is complex and process-based. The computational cost of solving a spatially distributed, nonlinear, multi-component reactive transport model is high, posing challenges for the application of ensemble-based uncertainty analysis and model weighting/selection methods (Song et al., 2015). With additional reaction and transport processes, the model includes more functions (such as reaction kinetic rate laws) and parameters (e.g., reaction rate constants, surface area) than hydrological models, which have already been criticized for their complexity, equifinality, uncertainty, and data demands (Beven, 2001, 2006; Kirchner et al., 1996). These issues will persist even though reactive transport models will be constrained by additional chemical data. A major source of uncertainty in these models lies in epistemic uncertainties, i.e., the lack of specific knowledge in forcing data and details of reactivities (e.g., spatial distribution and abundance of reactive materials), on top of uncertainties related to hydrology (Beven, 2000; Beven and Freer, 2001). The model's conceptual foundations also represent a major source of uncertainty.

It is in this spirit of "balancing" the cost and gain that we present both spatial distributed and lumped modes for the BioRT model. Compared to the distributed version, the spatially implicit model requires less spatial data, is computationally inexpensive, and is relatively easy to set up. It can assess the average dynamics of the water and solute dynamics and focus on the interactions among processes without resolving spatial details. The lumped approach can also accommodate basins with low data availability, and it can be easier for students to learn to use the model. In contrast, spatially explicit representations enable the exploration of the "hot spots" (e.g., swales and riparian zones with high soil water DOC concentrations in Figure 10e) and their contribution to stream chemistry at different times. Spatial heterogeneities in watershed properties (e.g., soil types and depth, lithology, vegetation, biomass, and mineralogy) are ubiquitous in natural systems. However, a general understanding of the linkage between local catchment features and catchment-scale dynamics (e.g., stream concentration dynamics and solute export pattern) is often lacking. We generally do not understand how spatial heterogeneity affects water flow paths, stream water chemistry, and biogeochemical reaction rates. The

spatially distributed model provides a tool to further explore these questions. Ultimately, the choice of the model complexity level depends on the research questions that the model is set to answer. At the end, we all need to balance cost and gain when deciding to use a simple or complex model, striving to be "simple but not simplistic" (Beven and Lane, 2019).

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8. Summary and conclusion

This paper introduces the watershed-scale biogeochemical reactive transport code BioRT (short for BioRT-Flux-PIHM). The code integrates processes of land-surface interactions, surface hydrology, and multi-component biogeochemical reactive transport. The new development enables the simulation of 1) biotic reactions including plant uptake, soil respiration, and microbe-mediated redox reactions, and 2) surface water interactions with groundwater from deeper subsurface that still interacts with streams. BioRT has been verified against the widely used reactive transport code CrunchTope for soil carbon, nitrogen, and phosphorus processes. The BioRT module has been applied to understand carbon, nitrogen, and weathering processes in Shale Hills in Pennsylvania, Coal Creek in Colorado, and Volcán Chimborazo watershed in Andes in Ecuador. Here we showcase the modeling capability of surface-groundwater interactions and reactive transport processes relevant to nitrate and DOC in Shale Hills in two simulation modes. One is in a spatially lumped mode using averaged properties and another is in a spatially distributed mode with consideration of spatial heterogeneity. Results show that the deep groundwater flow that interacts with the stream is primarily controlled by the hydraulic conductivity contrast between shallow and deep zone. biogeochemical reactions in shallow soil primarily determine the stream water chemistry under high flow conditions. The spatially lumped method with two lumped grids can capture the temporal dynamics of average behavior and mass balance; the spatially distributed running mode can be used to understand the spatial dynamics and to identify "hot spots" of reactions. The code can be used for biogeochemical reactive transport simulations in watersheds under diverse climate, land cover, and geology conditions.

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Data availability. Field data (e.g., discharge, stream chemistry) is archived at Shale Hills 782

data portal: http://criticalzone.org/shale-hills/data/datasets/ or maintained at HydroShare: 783

784 https://www.hydroshare.org/group/147.

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Code availability. The current model release (BioRT-Flux-PIHM v1.0), including 786

documentation, source code, example data, is available at GitHub repository: 787

https://github.com/PSUmodeling/BioRT-Flux-PIHM. 788

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Competing interests. The authors declare that they have no conflict of interest.

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Author contributions. LL conceived the model idea and oversaw the model development. WZ coded the BioRT module, verified the code against the benchmark reactive transport model CrunchTope, and applied and tested the model at Shale Hills watershed. YS developed the deep groundwater component and integrated the BioRT-Flux-PIHM v1.0 into the MM-PIHM family. WH, LS, KS, DK, BS, and GHCN tested the

code during its development and contributed study cases.

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