

1 **BioRT-Flux-PIHM v1.0: a watershed biogeochemical reactive transport**
2 **model**

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12

Abstract

13 Watersheds are the fundamental Earth surface functioning units that connect the land to
14 aquatic systems. Many watershed-scale models represent hydrological processes but
15 lack the representation of multi-component reactive transport processes. This has limited
16 our capability to understand and predict solute export, water chemistry, and earth system
17 response to changing climate and anthropogenic conditions. Here we present a recently
18 developed BioRT-Flux-PIHM (BioRT hereafter) v1.0, a watershed-scale biogeochemical
19 reactive transport model. The model augments the previously developed RT-Flux-PIHM
20 that integrates land-surface interactions, surface hydrology, and abiotic geochemical
21 reactions. It enables the simulation of 1) shallow and deep water partitioning to represent
22 surface water, shallow soil water, and deeper groundwater; 2) biotic processes including
23 plant uptake, soil respiration, and microbially mediated reactions such as nutrient
24 transformation. The reactive transport part of the code has been verified against the
25 widely used reactive transport code CrunchTope. BioRT-Flux-PIHM v1.0 has recently
26 been applied in multiple watersheds under diverse climate, vegetation, and geological
27 conditions. This paper briefly introduces the governing equations and model structure
28 with a focus on new model developments. It also showcases one hydrology example that
29 simulates shallow and deep water interactions, and two biogeochemical examples
30 relevant to nitrate and dissolved organic carbon (DOC). These examples are illustrated
31 in two simulation modes of varying complexity. One is the spatially lumped mode (i.e.,
32 two land cells connected by one river segment) that focuses on processes and average
33 behavior of a watershed. Another is the spatially distributed mode (i.e., hundreds of cells)
34 that includes details of topography, land cover, and soil properties. Whereas the spatially
35 lumped mode represents more of averaged properties and processes and temporal
36 variations, the spatially distributed mode can be used to understand the impacts of spatial
37 structure and identify hot spots of biogeochemical reactions.

38 **1. Introduction**

39 Watersheds are the fundamental Earth surface units that receive and process
40 water, mass, and energy (Li, 2019;Li et al., 2020;Hubbard et al., 2018). Watershed
41 processes include land surface interactions that regulate evapotranspiration and
42 discharge, and water partitioning between shallow soil lateral flow going into streams
43 versus downward flow and recharge into the deeper subsurface (Figure 1). Complex
44 biogeochemical interactions occur among soil, water, roots, and microbes along water's
45 flow paths, regulating gas effluxes (e.g., CO₂) and solute export (Fatichi et al., 2019;van
46 der Velde et al., 2010;Grathwohl et al., 2013).

47 These hydrological and biogeochemical processes determine how Earth surface
48 systems respond to hydroclimatic forcing and human perturbations (van der Velde et al.,
49 2014;Miller et al., 2020;Han et al., 2019). Understanding these processes remains
50 challenging due to the complex process interactions. An example is the concentration-
51 discharge (C-Q) relationships of solutes at stream and river outlets. Similar C-Q
52 relationships have been observed for some solutes across watersheds under diverse
53 geological and climatic conditions (Godsey et al., 2009;Basu et al., 2010;Moatar et al.,
54 2017;Zarnetske et al., 2018;Godsey et al., 2019), whereas different solutes have shown
55 contrasting patterns in the same watershed (Miller et al., 2017;Herndon et al.,
56 2015;Musolff et al., 2015). A general theory that can explain contrasting C-Q observations
57 (e.g., flushing vs. dilution behaviors) under diverse watershed characteristics and external
58 conditions remains elusive. The lack of mechanistic understanding presents major
59 roadblocks to forecasting water quality and earth system dynamics in the future.

60 One of the outstanding challenges is the lack of modeling tools that mechanistically
61 link hydrological and biogeochemical processes at the watershed scale. Model
62 development has advanced mostly within the disciplinary boundaries of hydrology and
63 biogeochemistry (Li, 2019). Watershed hydrologic models focus on solving for water
64 storage and fluxes (Fatichi et al., 2016). Reactive transport models (RTMs) have
65 traditionally focused on transport and multi-component biogeochemical reactions mostly
66 in groundwater systems with limited interactions with climate and other surficial conditions
67 (Steeffel et al., 2015;Li et al., 2017b). Some integration crossing disciplinary boundaries

68 did occur in recent years. For example, SWAT (Soil & Water Assessment Tool) has a
69 version that couples with the groundwater model MODFLOW and the surface water and
70 groundwater quality model in RT3D (Bailey et al., 2017; Ochoa et al., 2020). CATHY
71 (Catchment Hydrology) includes processes of pesticide decay (Gatel et al.,
72 2019; Scudeler et al., 2016). Hydrologiska Byråns Vattenbalansavdelning (HBV) and the
73 Hydrological Predictions for the Environment (HYPE) have modules that simulate
74 processes relevant to nutrients and contaminants (Lindström et al., 2005; Lindström et al.,
75 2010). While these models can simulate processes such as leaching of nutrients from
76 agriculture lands (Lindström et al., 2005; Lindström et al., 2010; Bailey et al., 2017), they
77 do not explicitly solve the multi-component reactive transport equations. In other words,
78 reactions are often represented rudimentarily without honoring kinetics and
79 thermodynamics theories in soil biogeochemistry and geochemistry. For example,
80 nutrient leaching is calculated based on empirical equations without explicitly solving
81 reactive transport equations. Reaction rates are represented using first-order decay
82 (Gatel et al., 2019), assuming constant reaction rates that do not change with
83 environmental conditions. Biogeochemical processes however are highly variable with
84 seasonal dynamics and depend on local environments such as substrate availability, soil
85 temperature, and soil moisture (Li et al., 2017a; HARTLEY et al., 2007). These models
86 therefore cannot capture the temporal variations in environmental factors that regulate
87 soil biogeochemical reactions and stream and water chemistry.

88 To fill this model capability need, we augmented the watershed model RT-Flux-
89 PIHM (Bao et al., 2017) into BioRT-Flux-PIHM (BioRT hereafter). Compared to RT-Flux-
90 PIHM, BioRT has two additions. One is the capability of simulating biotic processes
91 including plant uptake of nutrients, soil respiration, and other microbe-mediated redox
92 reactions. Examples include soil respiration that produces CO₂ and dissolved organic
93 carbon (DOC), and nutrient transformation reactions such as nitrification and
94 denitrification. The other is the addition of an optional deeper layer below shallow soil to
95 enable the simulation of interacting deep water and shallow soil water flow (Figure 1).
96 Here the deep water is loosely defined as the water below the soil zone, typically in less
97 weathered, fractured subsurface that harbors relatively old and slow-moving groundwater
98 contributing to streams. It is a fundamental component of the hydrologic cycle and water

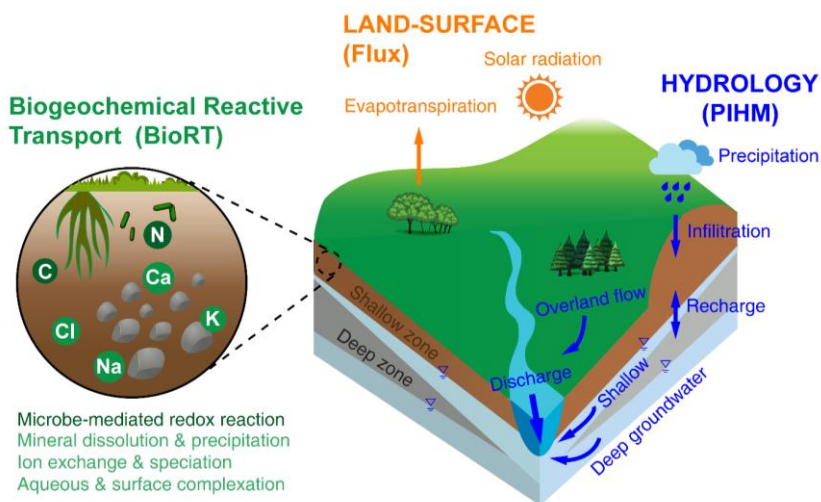
99 budget. The groundwater-surface water interactions also modulate land-atmospheric
100 energy exchanges and soil moisture dynamics (Keune et al., 2016). Evidence has been
101 mounting in recent years that deeper water below the shallow soil interacts with streams,
102 introduces water with distinct chemistry, sustains base flow in dry times, and buffers
103 climate variability (Gurdak, 2017;Green, 2016;Taylor et al., 2013). Stream chemistry often
104 reflects distinct chemistry from shallow soil water and deeper groundwater at different
105 time, i.e., the so called Shallow and Deep Hypothesis (Zhi et al., 2019;Zhi and Li,
106 2020;Botter et al., 2020). Including the deep water component thus is essential for
107 understanding mechanisms and predicting dynamics of water quality under changing
108 climate and human conditions.

109 This paper introduces the new developments in BioRT. The paper starts with a
110 brief overview of water and energy related processes. It then introduces governing
111 equations and reaction kinetics used in BioRT, followed by three examples that illustrate
112 the new capabilities. The examples include the surface water and groundwater
113 interactions, nitrate transformation and transport, and the production and export of DOC.
114 The model can be set up in both spatially lumped or spatially explicit modes. The source
115 code and the examples shown here are ~~hosted on~~archived on the Zenodo website
116 (<https://doi.org/10.5281/zenodo.3936073>) and the GitHub website ([https://github.com/Li-](https://github.com/Li-Reactive-Water-Group/BioRT-Flux-PIHM)
117 [Reactive-Water-Group/BioRT-Flux-PIHM](https://github.com/Li-Reactive-Water-Group/BioRT-Flux-PIHM)).

118

119 **2. Model overview**

120 BioRT-Flux-PIHM integrates three modules (Figure 1). The Flux module is for land-
121 surface processes including surface energy balance, solar radiation, and ET (Shi et al.,
122 2013). The hydrology module PIHM simulates water processes including precipitation,
123 interception, infiltration, recharge, surface runoff, subsurface lateral flow, and deep water
124 flow (Qu and Duffy, 2007). The BioRT module simulates solute transport reactions. The
125 abiotic reactions included in RT-Flux-PIHM (Bao et al., 2017) are mineral dissolution and
126 precipitation, aqueous and surface complexation, and ion exchange reactions. The newly
127 added reactions include plant uptake of nutrients, soil respiration, microbe-mediated
128 redox reactions (e.g., carbon decomposition and nutrient transformation).



129
 130 **Figure 1.** A conceptual diagram for processes at the watershed scale. Land surface interactions
 131 include processes such as energy balance, solar radiation, and evapotranspiration; hydrological
 132 processes partition water between surface runoff, shallow soil water, and deeper water entering
 133 the stream. Soil biogeochemical reactions include abiotic reactions such as weathering (e.g.,
 134 mineral dissolution and precipitation), ion exchange, surface complexations), and biotic
 135 processes such as plant uptake of nutrients, soil respiration, and other microbe-mediated
 136 reactions. These processes are represented in three modules: The Flux module for land-surface
 137 interactions, the PIHM module for catchment hydrology, and the BioRT module for
 138 biogeochemical reactions. Conceptually the shallow zone is the shallow soil and weathered zone
 139 that are more conducive to water flow (e.g., soil lateral flow or interflow). The deep zone
 140 refers to the less weathered zone that often harbors the old and slow flowing groundwater.
 141 Reactions can occur in both shallow and deep zones.

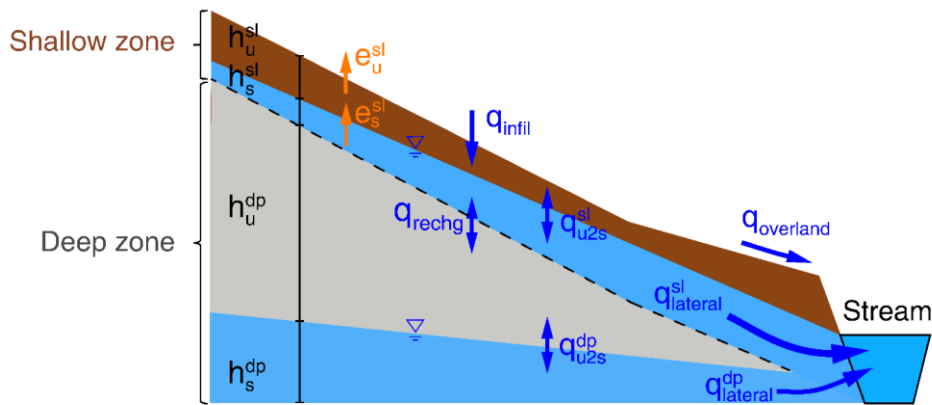
142
 143 The land surface and hydrology modules solve for soil temperature and water
 144 storage, from which water fluxes are calculated for surface runoff, shallow and deep water
 145 fluxes. The BioRT module uses the calculated soil temperature, water storage, and water
 146 fluxes to simulate solute transport (advective and diffusive/dispersive) and
 147 biogeochemical reactions in both shallow and deep zones (see governing equations in
 148 later sections). The reactions include kinetically controlled (e.g., microbe-mediated redox
 149 reaction, mineral dissolution and precipitation) or equilibrium-controlled (e.g., ion
 150 exchange, surface complexation (sorption), and aqueous complexation). Users can

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

154 The model can be set up running in either spatially lumped or spatially explicit
155 modes. When running in spatially explicit mode, the simulation domain can be structured
156 as prismatic grids based on topography. Each grid is partitioned into surface and shallow
157 and deep subsurface layers. The surface layer calculates water flow above ground
158 (surface runoff). The shallow zone is loosely defined as the highly permeable subsurface
159 that contrasts the deep zone that is broadly defined as the lower permeability zone
160 beyond the shallow zone. In many places, the shallow zone is the soil zone that is most
161 conductive to water flow (e.g., lateral flow) and is responsive to hydroclimatic forcing. The
162 deep subsurface zone is the less weathered layer that harbors the old ground water that
163 contributes to stream flow. Note that these definitions differ from those in the hydrology
164 community, which often refers to the shallow soil lateral flow as groundwater, in a way
165 that distinguishes it from the surface runoff (Dingman, 2015). These source waters from
166 different depths of the subsurface often have distinct solid and water chemistry, and are
167 dominant at different hydrological conditions in different time of the year, as have been
168 observed in many catchments and watersheds (Brantley et al., 2018;Zhi et al., 2019;Zhi
169 and Li, 2020;Sullivan et al., 2016). The model is flexible for taking inputs from online data
170 portals or local measurements and it can accommodate low data availability (see the
171 following section of 5 for data need and domain setup).

172

173 **3. Governing equations and processes**



174
 175 **Figure 2.** Hillslope view of the shallow and deep zones and relevant water flows. Streams
 176 received water primarily from three water flows: the surface runoff (q_{overland}), and lateral flow from
 177 shallow zone ($q_{\text{lateral}}^{\text{sl}}$), and the lateral flow that has been recharged and eventually come out from
 178 deeper zone ($q_{\text{lateral}}^{\text{dp}}$). The symbol of “h”, “e”, and “q” denotes water storage, evapotranspiration,
 179 and water flow, respectively. The superscript letter “sl” and “dp” refer to shallow and deep zone,
 180 respectively. The subscript letters “u” and “s” refer to unsaturated and saturated layers,
 181 respectively. Detailed equations are in section 3.1 – 3.2. The terms “infil”, “u2s”, and “recharge”
 182 refer to infiltration, unsaturated to saturated zones, and recharge.

183

184 3.1 Water equations

185 Flux-PIHM simulates surface runoff and a lumped subsurface flux into streams
 186 without distinguishing shallow soil water and deeper groundwater flow paths. Mounting
 187 evidence has shown that the shallow soil water and deeper groundwater have distinct
 188 chemistry and are dominant at different times of the year (Xiao et al., 2021; Zhi and Li,
 189 2020; Zhi et al., 2019). This means that a lumped subsurface flow cannot describe the
 190 dynamics of stream chemistry. We therefore added deeper groundwater zone to simulate
 191 deeper water flows that interact with streams. Each prismatic element now has three
 192 zones in the vertical direction: surface (or above ground), shallow and deep zones in the
 193 subsurface.

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

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

$$199 \quad \theta_i^{sl} \frac{dh_{i,u}^{sl}}{dt} = q_{i,inf} - q_{i,u2s}^{sl} - e_{i,u}^{sl} \quad (1)$$

$$200 \quad \theta_i^{sl} \frac{dh_{i,s}^{sl}}{dt} = q_{i,u2s}^{sl} - q_{i,rechg} - e_{i,s}^{sl} + \sum_1^{N_{ij}} q_{ij}^{sl} \quad (2)$$

201 Where θ_i^{sl} [m^3 pore space/ m^3 total volume] is the shallow zone porosity in element i ; $h_{i,u}^{sl}$
 202 and $h_{i,s}^{sl}$ [m] are the unsaturated and saturated water storage in the shallow zone,
 203 respectively. The storages h here are the height of soil column with equivalent saturated
 204 water, not the height of the pure water (100% volume) column. That is why porosity is in
 205 the equation. For saturation zones, this height is needed to quantify the depths of water
 206 tables and determines the direction of water flow between neighboring grids. The
 207 $q_{i,inf}$ [m/s] is the infiltration rate from the surface to the shallow zone; $q_{i,u2s}^{sl}$ [m/s] is the
 208 vertical flow from the unsaturated layer to the saturated layer in the shallow zone; $q_{i,rechg}$
 209 [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
 210 evapotranspiration (ET) from the unsaturated and saturated layer in the shallow zone,
 211 respectively; q_{ij}^{sl} [m/s] are the lateral flows in the shallow saturated layer between the
 212 element i and its neighbor element j ; N_{ij} (≤ 3) is the number of neighbor elements j . For
 213 a prismatic element i , a boundary cell has one or two neighbors; a non-boundary cell has
 214 three neighbors. ET is calculated by the Penman potential evaporation scheme (detailed
 215 equations in Shi (2012)). A similar set of water equations for the deep zone are in the SI
 216 (Eqn. S1 and S2).

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

$$221 \quad q_{i,inf} = K_{i,inf} \frac{H_{i,sur} - H_{i,u}^{sl}}{d_{i,inf}} \quad (3)$$

222
$$q_{i,u2s}^{sl} = K_{i,V}^{sl} \frac{H_{i,u}^{sl} - H_{i,s}^{sl}}{0.5d_i^{sl}} \quad (4)$$

223 Where $d_{i,inf}$ and d_i^{sl} [m] are the thickness of infiltration layer and shallow zone depth,
 224 respectively; $K_{i,inf}$ [m/s] is the hydraulic conductivity of the infiltration layer, the top 0.1 m
 225 of the subsurface that has different conductivity from the rest of subsurface; $K_{i,V}^{sl}$ [m/s] is
 226 the hydraulic conductivity in the vertical direction (i.e., weighted average of macropore
 227 $K_{i,macV}$ 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
 228 and saturated layer, respectively. The lateral flow in the shallow saturated layer is
 229 calculated using Darcy's law:
 230

231
$$q_{ij}^{sl} = K_{ij}^{sl} \frac{H_{i,s}^{sl} - H_{j,s}^{sl}}{d_{ij}} \quad (5)$$

232 Where d_{ij} [m] is the distance between the centers of elements i and j ; K_{ij}^{sl} [m/s] is the
 233 harmonic mean of shallow hydraulic conductivity in the horizontal direction between
 234 elements i ($K_{i,H}^{sl}$) and j ($K_{j,H}^{sl}$). The interaction between the shallow saturated zone and
 235 stream channel also follows Eqn. 5, except that the adjacent head is replaced by the level
 236 of stream water. Similar to the shallow zone, hydrological equations in the deep zone are
 237 detailed in the SI (Eqn. S1 – S8).
 238

239 3.2 Reactive transport equations

240 The governing advection dispersion reaction (ADR) equation for an arbitrary solute
 241 m in an element i is as follows (Bao et al., 2017):

242
$$V_i \frac{d(S_{w,i} \theta_i C_{m,i})}{dt} = \sum_1^{N_{ij}} \left(A_{ij} D_{ij} \frac{C_{m,j} - C_{m,i}}{d_{ij}} - q_{ij} A_{ij} C_{m,j} \right) + R_{m,i}, \quad m = 1, \dots, nm \quad (6)$$

243 Where V_i [m³ total volume] is the total volume of element i ; $S_{w,i}$ [m³ water/m³ pore space]
 244 is soil water saturation; θ_i [m³ pore space/m³ total volume] is the porosity; $C_{m,i}$ [mol/m³
 245 water] is the aqueous concentration of species m ; N_{ij} is the number of fluxes from

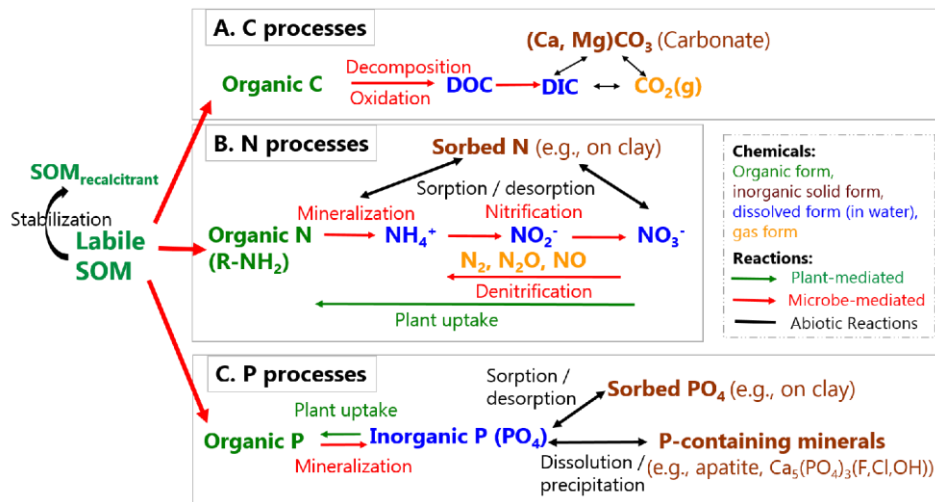
246 neighbor element j for element i , N_{ij} is 2 for the unsaturated zone (infiltration, recharge)
247 with only vertical flows and 5 for saturated zone with flux from (or to) the unsaturated
248 zone, from (or to) the deeper zone, and fluxes between i and three neighbor elements j
249 in lateral flow directions for non-boundary grids; A_{ij} [m^2] is the grid area shared by i and
250 its neighbor grid j ; D_{ij} [m^2/s] is the hydrodynamic dispersion coefficient (i.e., sum of
251 mechanical dispersion and effective diffusion coefficient) normal to the shared surface
252 A_{ij} ; d_{ij} [m] is the distance between the center of i and its neighbor elements j ; q_{ij} [m/s]
253 is the flow rate across A_{ij} ; $R_{m,i}$ [mol/s] is the total rate of kinetically controlled reactions in
254 element i that involve species m ; nm is the total number of independent primary species
255 to be solved for reactive transport equations. Equation (6) states that the change of solute
256 mass (the left term) is driven by dispersive transport, advective transport, and reactions
257 (i.e., the 1st, 2nd, and 3rd right-hand side terms, respectively).

258

259 **3.3 Biogeochemical processes and reaction kinetics**

260 **3.3.1 Biogeochemical processes**

261 Here we discuss representative biogeochemical processes that involve plants and
262 microbes that can be included in BioRT. BioRT differs from general water quality models
263 that primarily target a few contaminants (e.g., N, P, metals). The framework of BioRT is
264 flexible and the users can define reactions and solutes of interests in the input files. For
265 abiotic reactions such as mineral dissolution and surface complexation or ion exchange,
266 readers are referred to (Bao et al. (2017)). Generally speaking, shallow soils contain more
267 weathered materials and soil organic matters (SOM) including roots, leaves, and
268 microbes. SOM can be decomposed partially into organic molecules that dissolve in
269 water, i.e., Dissolved Organic Carbon (DOC). It can also become oxidized completely
270 into CO_2 , which can emit back to the atmosphere in gas form (Davidson, 2006) or
271 transport and enter streams in the form of dissolved inorganic carbon (DIC). With
272 coexisting cations (e.g., Ca, Mg), DIC can precipitate out and become carbonate minerals
273 (e.g., CaCO_3).



274
 275 **Figure 3.** Biotic and abiotic reactions relevant to the transformation of soil organic matter (SOM).
 276 SOM can become stabilized (recalcitrant) through sorption on clay and separation from reactants.
 277 Labile OM can decompose into inorganic forms, releasing C, N, and P that further transform into
 278 various forms (adopted from Li (2019), permission with Mineralogical Society of America).

279
 280 OM decomposition releases organic nitrogen (R-NH₂), which can further react to
 281 become NH₄⁺ and other nitrogen forms (N₂, N₂O, NO, NO₂⁻, NO₂) (Figure 3). The gases
 282 can emit back to the atmosphere. Denitrification requires anoxic conditions and occurs
 283 less commonly in shallow soils owing to the pervasive presence of O₂ (Sebestyen et al.,
 284 2019); denitrification can become important under wet conditions and in O₂-depleted
 285 groundwater aquifers. Phosphorous (P) can be in organic forms in organic matter, sorbed
 286 on fine soil particles, dissolved in water, or in solid forms as P-containing minerals.
 287 Transformation of nutrients occurs through various bio-mediated or abiotic reactions. A
 288 representative P-containing mineral in the Earth's crust is apatite Ca₅(PO₄)₃(F, Cl, OH).
 289 Once liberated via rock dissolution, P is biologically assimilated and locked in organic
 290 forms. These organic forms have very low solubility, allowing them to bind on and be
 291 transported together with soil particles in the form of orthophosphate or pyro-diphosphate.

292
 293 **3.3.2 Reaction kinetics in natural soils**

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

$$299 \quad r = kA f(T) f(S_w) f(Z_w) \quad (7)$$

300 Where the reaction rate r [mol/s] depends on rate constant k [mol/m²/s], the surface area
 301 A [m²] is a lumped parameter that quantitatively represents SOM content and biomass
 302 abundance, $f(T)$ and $f(S_w)$ describe the temperature and soil moisture dependence,
 303 respectively, $f(Z_w)$ can be included to account for the depth distribution of SOM (Seibert
 304 et al., 2009), and Z_w [m] is the water table depth. An example for the depth distribution is
 305 $f(Z_w) = \exp\left(-\frac{Z_w}{b_m}\right)$ (Weiler and McDonnell, 2006; Bai et al., 2016), with b_m as the depth
 306 coefficient describing the gradient of SOM content over depth. Users can choose to
 307 include either one or all of these dependencies in input or database files.

308 The temperature dependence follows a Q₁₀-based form (Lloyd and Taylor,
 309 1994; Friedlingstein et al., 2006; Hararuk et al., 2015) as follows:

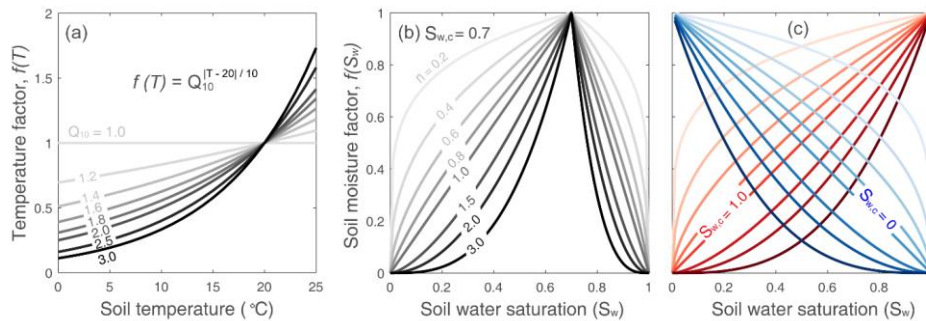
$$310 \quad f(T) = Q_{10}^{(T-20)/10} \quad (8),$$

311 where Q_{10} is the relative increase in reaction rates when temperature increases by 10 °C
 312 (Davidson and Janssens, 2006). Values of Q_{10} (Figure 4a) can vary from 1.0 to 3.0,
 313 depending on climatic conditions, substrate availability, and ecosystem type (e.g.,
 314 grassland, forest) (Davidson et al., 2006; Liu et al., 2017). The mean values are in the
 315 range of 1.4 to 2.5 (Zhou et al., 2009; Bracho et al., 2016). The Q_{10} value can be specified
 316 in the input file.

317 The soil moisture dependence function $f(S_w)$ is coded in the following form:

$$318 \quad \left\{ \begin{array}{l} \left(\frac{S_w}{S_{w,c}}\right)^n, S_w \leq S_{w,c} \\ \left(\frac{1-S_w}{1-S_{w,c}}\right)^n, S_w > S_{w,c} \end{array} \right. \quad (9)$$

319 Here $S_{w,c}$ [0 to 1] is the critical soil moisture at which rates are highest, and n is the
 320 exponent reflecting the dependence of rates on soil moisture. A typical n value is 2 (Yan
 321 et al., 2018) with a range between 1.2 and 3.0 (Hamamoto et al., 2010), depending on
 322 soil structure and texture. As shown in Figure 4b, the form indicates an intermediate
 323 critical soil moisture $S_{w,c}$ at which $f(S_w)$ reaches its maximum. When $S_w \leq S_{w,c}$, $f(S_w)$
 324 increases with S_w ; when $S_w > S_{w,c}$, $f(S_w)$ decreases with S_w (Figure 4b) (Yan et al.,
 325 2018). Under the extreme conditions of $S_{w,c}$ equals to 0 or 1, $f(S_w)$ monotonically
 326 increase or decrease (Figure 4c). The two parameters, $S_{w,c}$ and n , determines the shape
 327 of the curve. They can be specified in input or database files. One can also choose not to
 328 have temperature or soil moisture dependence by choosing parameters that would lead
 329 to the value of exponent being zero.



330 **Figure 4.** Reaction rate dependence. (a) Function form of soil temperature dependence and (b,
 331 c) soil moisture dependence for reaction rates. The $f(T)$ takes Q_{10} form (Equation 8). The soil
 332 moisture factor $f(S_w)$ depends on $S_{w,c}$ and n and soil water saturation S_w (Equation 9). The soil
 333 moisture function can represent three types of behaviors: the threshold behavior (b, $0 < S_{w,c} < 1$),
 334 increase behavior (red in (c), $S_{w,c} = 1$), and decrease behavior (blue in (c), $S_{w,c} = 0$). Values of n
 335 = 1 leads to a linear threshold dependence of S_w while $n < 1$ and $n > 1$ lead to concave and
 336 convex dependences, respectively.
 337

338

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

344 on concentrations of electron donors and non-oxygen electron acceptors (e.g., nitrate,
345 iron oxides, sulfate) that are often limited under anoxic conditions (Bao et al., 2014;Li,
346 2019;Benettin et al., 2020). The order of redox reactions typically follows the
347 biogeochemical redox ladder, which is based on how much microbe can harvest energy
348 by reducing different types of electron acceptors. Monod reaction rate laws are often used
349 for quantifying rates of these redox conditions. These rate laws are detailed in the section
350 S2 of Supporting Information. Users can combine these Monod rate laws and the
351 temperature and soil moisture dependence described above, if needed.

352

353 **3.4 Plant related processes: root uptake of nitrate as an example**

354 Nutrient uptake by plants is complex and remains poorly understood. A variety of
355 plant uptake models exists with varying degrees of complexity (Neitsch et al., 2011;Fisher
356 et al., 2010;Cai et al., 2016). These models are mostly based on plant growth module or
357 supply and demand approach that often requires detailed phenological and plant
358 attributes including growth cycle, root age and biomass, nutrient availability, and carbon
359 allocation, in addition to local temperature and soil moisture (Neitsch et al.,
360 2011;Porporato et al., 2003;Dunbabin et al., 2002;Buysse et al., 1996;Fisher et al., 2010).
361 Without detailed mechanistic understanding, we assume a simple and operational
362 approach. In the Example 2 that we show later, for example, nitrate uptake was modelled
363 with dependence on NO_3^- concentration, soil temperature and moisture, and rooting
364 density (McMurtrie et al., 2012;Yan et al., 2012;Buljovic and Engels, 2001).

$$365 \quad r_{\text{uptake}} = k_{\text{uptake}} C_{\text{NO}_3^-} f(T) f(S_w) f_{\text{root}}(d_w) \quad (13)$$

$$366 \quad f_{\text{root}}(d_w) = \exp((-d_w + \delta) / \lambda) \quad (14)$$

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

373

374 **4. Numerical scheme and model verification**

375 The system of differential equations for water storages (e.g., Eqn. 1 and 2, and Eqn.
376 S1 and S2) are assembled into a global system of ordinary differential equations (ODEs).
377 It is solved in CVODE (short for C-language Variable-coefficients ODE solver,
378 <https://computing.llnl.gov/projects/sundials/cvode>), a numerical ODE solver in the SUite
379 of Nonlinear and Differential / ALgebraic equation Solvers (SUNDIALS) (Hindmarsh et
380 al., 2005). In BioRT, the transport step is first solved with water by the preconditioned
381 Krylov (iterative) method and the Generalized Minimal Residual Method (Saad and
382 Schultz, 1986). All primary species in element I are then assembled in a local matrix
383 and solved iteratively using the Crank-Nicolson and Newton-Raphson methods in
384 CVODE (Bao et al., 2017).

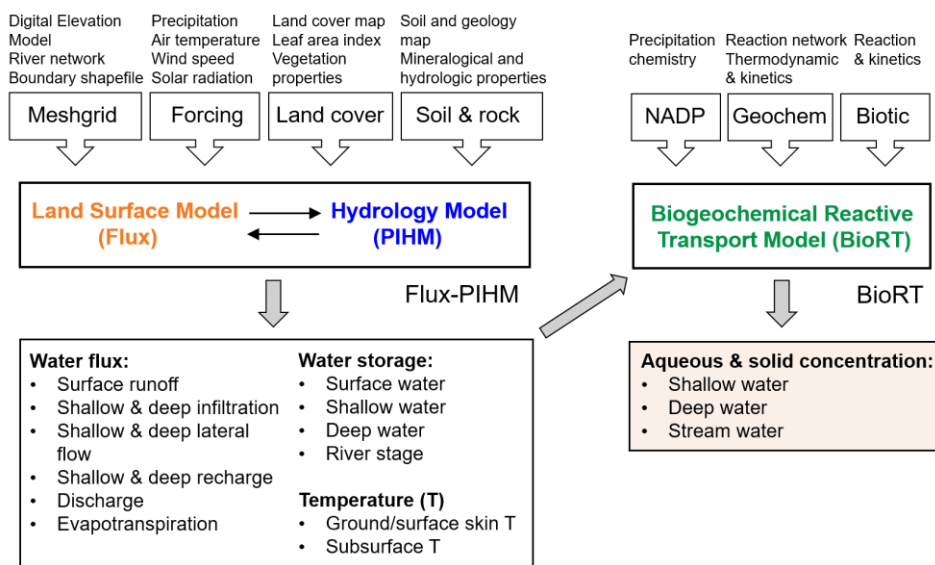
385 **Model verification.** The BioRT module had been verified against CrunchTope
386 under different transport and reaction conditions (Figures S1 – S7 in SI). CrunchTope is
387 a widely used subsurface reactive transport model (Steefel and Lasaga, 1994; Steefel et
388 al., 2015), and is often used as a benchmark to verify other reactive transport models.
389 Verification was performed under simplified hydrological conditions with 1-D column and
390 constant flow rates such that it focuses on advection, diffusion, dispersion, and
391 biogeochemical reactions. Specifically, three cases were verified. The phosphorus case
392 that involves kinetics-controlled apatite dissolution and thermodynamics-controlled
393 phosphorous speciation was first tested for solution accuracy of the bulk code that was
394 inherited from the original RT-Flux-PIHM. Soil carbon and nitrogen processes were
395 further verified for solution accuracy of the augmented BioRT module. Table S7 shows
396 an average percent bias and Nash Sutcliffe efficiency (NSE) of 1.1% and 0.98, indicating
397 a robust performance for a variety of solutes under different transport and reaction
398 conditions. Note NSE ranges from $-\infty$ to 1, with NSE = 1 being the perfect fit (Moriassi et
399 al., 2007).

401 **5. Model structure, data needs, and domain setup**

402 **Model structure.** The model takes meteorological forcing time series as input and
403 solves for water storages and soil temperature, along with other hydrologic and land

404 surface states and fluxes (Figure 5). BioRT reads in the model output of water and
405 temperature from Flux-PIHM, and solves the biogeochemical reactive transport
406 equations. At the time scale of months to years that are typical for BioRT simulations,
407 alterations in solid phase properties, including, porosity, permeability, and reactive
408 surface area, are considered negligible such that hydrological parameters remain
409 constant with time.

410 **Data needs.** The code sets up the model domain based on watershed
411 characteristics including topography, land cover, and shallow and deep zone properties
412 (Figure 5). When the model is used in a spatially distributed form, the model domain is
413 set up using elevation, land cover, soil and geology maps supplied by the user. A useful
414 data portal is the Geospatial Data Gateway (<https://datagateway.nrcs.usda.gov>). Another
415 geospatial data source is the HydroTerre (<http://www.hydroterre.psu.edu/>), where users
416 can obtain data on elevation, land cover, geology, and soil (Leonard and Duffy, 2013).
417 Meteorological forcing data can be downloaded from the North American Land Data
418 Assimilation Systems Phase 2 (NLDAS-2, <https://ldas.gsfc.nasa.gov/nldas/v2/forcing>).
419 The vegetation forcing, i.e., Leaf Area Index (LAI), can be obtained from MODIS
420 (Moderate Resolution Imaging Spectroradiometer, <https://modis.gsfc.nasa.gov/data>).
421 Other vegetation properties (e.g., shading fraction, rooting depth) can be adopted from,
422 for example, the Noah vegetation parameter table embedded in the Weather Research
423 and Forecasting model (WRF; Skamarock and Klemp (2019)). Local measurements from
424 meteorological stations and field campaigns (e.g., land cover, soil, geology) can be used
425 in the model. Initial water and solid phase chemistry can be based on measurements or
426 general knowledge of the simulated sites. The form of reaction rate laws can be defined
427 in the input files and calibrated to reproduce field data. Reaction thermodynamics, mostly
428 equilibrium constants, are from the geochemical database EQ3/6 by default (Wolery,
429 1992). These reaction parameters can be modified when necessary. The model outputs
430 include aqueous and solid concentrations of shallow and deep zone and stream water.



431
 432 **Figure 5.** Model structure, input, and output of BioRT-Flux-PIHM. The Flux-PIHM takes in
 433 watershed characteristics including topography (digital elevation model, DEM), land cover,
 434 shallow and deep zone properties, and meteorological forcing time series and solves for water
 435 storage, and ground and soil temperature. BioRT takes in water- and temperature-related output
 436 from Flux-PIHM and additional inputs such as precipitation chemistry and shallow and deep water
 437 chemistry and biogeochemical kinetics parameters, and solve for aqueous and solid
 438 concentrations in the shallow and deep zone, and stream water. NADP stands for the National
 439 Atmospheric Deposition Program.

440
 441 **Domain set up: from simple, spatially lumped to complex, spatially distributed**
 442 **domains.** The domain can be set up at different spatial resolutions with different numbers
 443 of grids. A simple domain can be set up with two land grids representing two sides of a
 444 watershed connected by one river cell. This setup uses averaged properties without
 445 needs for larger spatial data. Alternatively, a complex domain can be set up to track “hot
 446 spots” of biogeochemical reactions using many grids with explicit representation of spatial
 447 details (e.g., topographic map, river network, land use map, soil and geology map, mineral
 448 distribution). The model domain can be set up using PIHM-GIS
 449 (http://www.pihm.psu.edu/pihmgis_home.html), a standalone GIS interface for watershed
 450 delineation, domain decomposition, and parameter assignment (Bhatt et al., 2014). The

451 same processes (e.g., hydrology, reaction network) can be setup in both types of spatial
 452 configurations. Auto-calibration is not built into the model, but a global calibration
 453 coefficient approach is used to reduce parameter dimension and facilitate manual
 454 calibration. A typical model application requires 20 to 30 hydrological parameters to be
 455 calibrated. These parameters include land surface parameters (e.g., canopy resistance,
 456 surface albedo), soil and geology parameters (e.g., hydraulic conductivity, porosity, Van
 457 Genuchten, macropore properties) (Shi et al., 2013). Reaction-related parameters (e.g.,
 458 reaction rate constant, mineral surface area, Q_{10} , $S_{w,c}$, and n) are additionally needed for
 459 calibration, the number of which depends on the numbers of reactions involved in a
 460 particular system.

461

462 **6. Model applications**

463 The original RT-Flux-PIHM has been applied to understand processes related to
 464 the geogenic solutes of Cl and Mg at the Shale Hills watershed and for Na in a watershed
 465 on Volcán Chimborazo in the Ecuadorian Andes (Table 1). The new BioRT-Flux-PIHM
 466 has been demonstrated for understanding the dynamics of DOC and nitrate at Shale Hills
 467 and Coal Creek. This section will present one hydrology and two biogeochemical
 468 examples in the Susquehanna Shale Hills Critical Zone Observatory (SSHCZO), a small
 469 headwater watershed in central Pennsylvania, USA. The mean annual precipitation is
 470 approximately 1,070 mm and the mean annual temperature is 10°C (Brantley et al.,
 471 2018). Soil carbon storage and respiration and nitrogen budget and fluxes have been
 472 studied in detail (Andrews et al., 2011; Hasenmueller et al., 2015; Weitzman and Kaye,
 473 2018). Modeling work has been conducted to understand hydrological dynamics (Shi et
 474 al., 2013; Xiao et al., 2019), transport of the non-reactive tracer Cl, and the weathering-
 475 derived solute Mg (Bao et al., 2017; Li et al., 2017a).

476 **Table 1.** Example Model applications of BioRT-Flux-PIHM

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

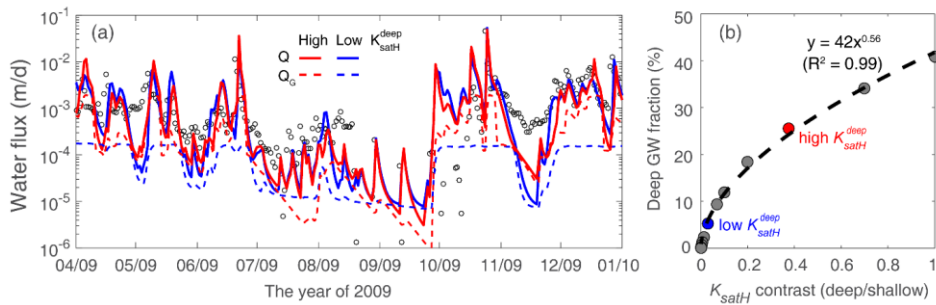
| | | | | | |
|-----------------------------|------|-----------------------|----------------------------------|--|-----------------------------------|
| | | Spatially distributed | Cl, Mg | <ul style="list-style-type: none"> • Chlorite dissolution¹ • Illite dissolution¹ • Cation exchange | Bao et al., 2017; Li et al., 2017 |
| Shale Hills (PA, USA) | 0.08 | Spatially distributed | DOC | <ul style="list-style-type: none"> • SOC decomposition² • DOC sorption | Wen et al., 2020 |
| | | Spatially lumped | NO ₃ ⁻ | <ul style="list-style-type: none"> • Soil N leaching² • Denitrification² • Plant uptake³ | This work |
| Coal Creek (CO, USA) | 53 | Spatially lumped | DOC, Na | <ul style="list-style-type: none"> • SOC decomposition² • DOC sorption • Albite dissolution¹ | Zhi et al., 2019 |
| Volcán Chimborazo (Ecuador) | | Spatially distributed | Cl, Na, Ca, Mg, SiO ₂ | <ul style="list-style-type: none"> • Albite dissolution¹ • Diopside dissolution¹ | Saberi et al. (2021) |

477 Note: Transition State Theory (TST) is a classic kinetic rate law for mineral dissolution and
478 precipitation (Brantley et al., 2008) (Eqn. S15); . SOC stands for soil organic carbon.

479

480 6.1 Example 1: Shallow and deep water interactions

481 The model was set up using the spatially lumped mode with two grids and one
482 river grid characterized by average land cover, soil and rock properties based on previous
483 work. The model assumed the dominant soil type (Weikert soil) at Shale Hills. The
484 porosity of the deep zone was set to a tenth of the shallow soil porosity based on
485 measurements of the groundwater aquifer (Brantley et al., 2018; Kuntz et al., 2011). In a
486 headwater catchment like Shale Hills where the deep groundwater is most likely sourced
487 from recharge, the deep groundwater contribution to the stream can be primarily
488 controlled by the hydraulic conductivity (K_{satH}) contrast between the deep and shallow
489 zones (i.e., $K_{satH}^{dp} / K_{satH}^{sl}$). This is because the K_{satH} contrast determines the partitioning
490 of infiltrating water between the shallow lateral flow and the downward recharge to the
491 deep zone and then deep groundwater flow. Two cases of high (red) and low (blue) K_{satH}^{dp}
492 were set up to showcase the control of K_{satH} contrast on deep groundwater (Figure 6a).
493 By changing the deep zone K_{satH}^{dp} from 2.6 to 0.22 (m/d), the annual deep groundwater
494 (Q_G) contribution to discharge (Q) decreased from 26% to 5.2%, although the total stream
495 discharge remains the same. This indicates that the changing K_{satH}^{dp} mostly changes the
496 flow partitioning between the shallow soil flow and deeper groundwater flow into streams.



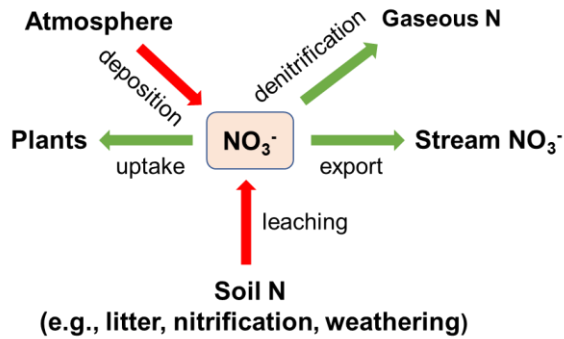
497
 498 **Figure 6.** (a) Hydraulic conductivity (K_{satH}) contrast controls the proportion of deep groundwater
 499 (Q_G). The cases of high ($K_{satH}^{dp} = 2.6 \text{ m/d}$, red) and low conductivity ($K_{satH}^{dp} = 0.22 \text{ m/d}$, blue) led
 500 to 26% and 5.2% of annual Q_G contribution to discharge (Q), respectively. (b) Deep groundwater
 501 fraction as a function of K_{satH} contrast between the deep and shallow zone. The upper limit of the
 502 deep / shallow K_{satH} contrast was set to 1 as most watersheds have smaller K_{satH} in the deep
 503 zone than in the shallow zone. The two red and blue dots correspond to the two cases in left
 504 panel.

505
 506 Several additional cases were further tested to examine the relationship between
 507 deep groundwater fraction (%) of discharge and K_{satH} contrast. Figure 6b shows that the
 508 deep groundwater fraction rapidly increases with the increasing ratio of $K_{satH}^{dp} / K_{satH}^{sl}$,
 509 reaching a limit when K_{satH} contrast is sufficiently high. The deep groundwater
 510 contribution to the stream reaches ~ 40% when K_{satH}^{dp} and K_{satH}^{sl} are equal. In natural
 511 systems, we do see places, for example, karst formations, where groundwater contributes
 512 to more than 40% (Hartmann et al., 2014; Husic, 2018). These places may have higher
 513 deeper conductivity than shallow soils due to the development of highly conductive
 514 conduits.

516 **6.2 Example 2: Nitrate dynamics in a spatially implicit domain**

517 This example focuses on nitrate (NO_3^-), a dominant dissolved N form in water
 518 (<https://criticalzone.org/shale-hills/data/datasets/>) (Weitzman and Kaye, 2018). The N
 519 processes at Shale Hills include atmospheric N deposition, soil N leaching, stream export,
 520 denitrification, and plant uptake (Figure 7). Based on field measurements, the
 521 atmospheric deposition at the site is the dominant N input; N export via discharge is only

522 a small fraction (2.5%) of atmospheric N input. Most deposited N is tightly cycled by plants
 523 or lost to the atmosphere via denitrification.



524
 525 **Figure 7.** Modeled nitrogen processes in Example 2. Atmospheric N deposition is the major N
 526 input; denitrification and plant uptake are the major N loss and sink. Export via discharge only
 527 occupies a small fraction.
 528

529 The soil N leaching process was represented using a lumped reaction that
 530 generates NO_3^- . Conceptually this could represent the total rates of reactions including
 531 the decomposition of soil organic matter (SOM), nitrification, and rock weathering that
 532 generates NO_3^- . Its rate was assumed to depend on soil temperature and moisture and
 533 follows the equation $r_{leach} = kA f(T) f(S_w)$, where r_{leach} [mol/s] is the leaching rate, $k =$
 534 $10^{-9.7}$ [mol/m²/s] is the leaching rate constant (Regnier and Steefel, 1999), and A [m²] is
 535 the surface area that represents the contact area between substrates and N transforming
 536 microbe, and $f(T)$ and $f(S_w)$ are soil temperature (Eqn. 8) and soil moisture (Eqn. 9)
 537 functions, respectively. The surface area was calculated based on SOM volume fraction
 538 [m³/m³], specific surface area (SSA, [m²/g]), substrate density [g/cm³], and element
 539 volume [m³].

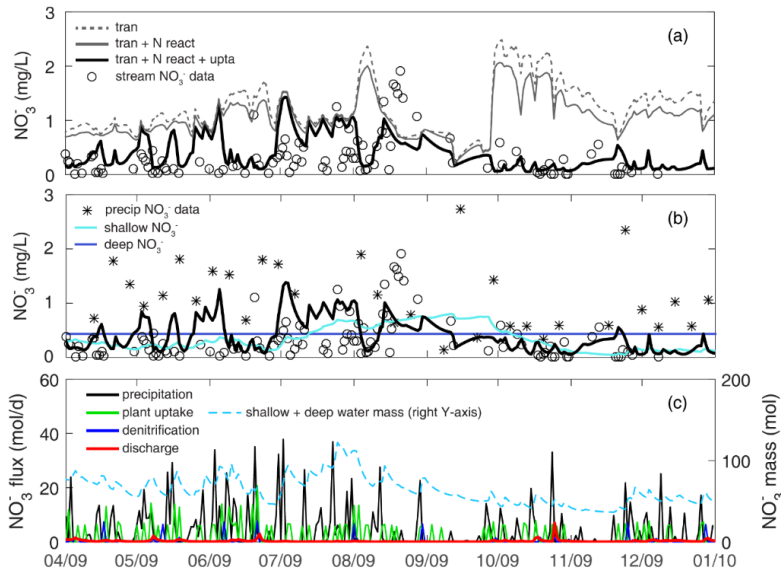
540 Denitrification converts NO_3^- to N_2 gas under anaerobic conditions. Here this
 541 process was modeled by the Monod rate law with DOC as the electron donor (Di Capua
 542 et al., 2019), NO_3^- as the electron acceptor, and with an inhibition term $f(O_2)$ (Eqn. S13).

543 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 $k =$
 544 10^{-10} [mol/m²/s] is the denitrification rate constant (Regnier and Steefel, 1999), half-

545 saturation constants $K_{m,DOC} = 15 [uM]$ and $K_{m,NO_3^-} = 45 [uM]$ (Regnier and Steefel,
546 1999). For soil N leaching and denitrification, the SSA were respectively tuned as $1.6 \times$
547 10^{-6} and $7.5 \times 10^{-5} [m^2/g]$ to reproduce observed stream nitrate dynamics. The calibrated
548 values were orders of magnitude lower than the lab measured SSA of natural materials
549 (e.g., SOM, $0.6 \sim 2 m^2/g$) (Rutherford et al., 1992). Such discrepancies between
550 calibrated effective reactive surface area (i.e., solid-water contact area) and lab measured
551 absolute surface area are consistent with other observations in literature (Li et al.,
552 2014; Heidari et al., 2017; Wen and Li, 2017, 2018). The uptake rate constant was
553 calibrated by constraining the partitioning of N transformation flux between denitrification
554 and plant uptake by the ratio of 1:5, a value estimated from field measurements of
555 gaseous N outputs (3.53 kg-N/ha/yr) and plant N uptake (18.3 kg-N/ha/yr) (Weitzman and
556 Kaye, 2018). The uptake rate constant in the deep zone (> 2 m in depth) was considered
557 negligible (Hasenmueller et al., 2017). Groundwater nitrate was initialized as 0.43 mg/L,
558 the average of measured groundwater concentration during 2009 - 2010.

559
560 **Temporal nitrate dynamics.** Three cases were set up to understand and quantify the
561 effects of different processes in determining nitrate dynamics (Figure 8a). The *transport-*
562 *only* case (dashed line, *tran*) simulates nitrate input from precipitation (at 1.4 ± 0.96 mg/L,
563 based on the 2009 data of NADP PA42 site) and N transport but without any reactions. It
564 overestimated stream nitrate data (0.33 ± 0.39 mg/L) throughout the year. The *transport*
565 *+ N reactions* case (gray line, *tran + N react*) has denitrification and soil N leaching
566 processes but not plant uptake. These two reactions lowered the nitrate concentration
567 slightly, as these two processes compensate each other in adding and removing nitrate
568 from water. The *transport + N reactions + uptake* case (thick black line, *tran + N react +*
569 *upta*) have all processes. It significantly lowered the nitrate concentration, especially in
570 April-May and October-December. Nitrate peaks from May to July, exhibiting comparable
571 levels of high nitrate concentration (Figure 8b). It is noticeable that the three cases almost
572 overlapped at these overestimated short nitrate peaks, suggesting nitrate-rich
573 precipitation may not be routed into the subsurface where denitrification and plant uptake
574 could occur.

575 Although precipitation from April to August accounted for 70% of the total
 576 simulation period, larger storm events in October contributed more to nitrate export.
 577 Deeper groundwater had higher nitrate concentration than shallow water, because most
 578 plant uptake occurred in the shallow zone. The nitrate fluxes into the deeper zone
 579 however only contributed 26% of stream nitrate export at the annual scale, due to the
 580 relatively small groundwater contribution (9.5%) to the stream. Denitrification and plant
 581 uptake largely occurred during wet spring with leaves growing. Denitrification peaks often
 582 appeared after major storm events because wet conditions facilitate denitrification.
 583 Comparing the three outfluxes (Figure 8c), nitrate export via discharge (red) was
 584 negligible compared to denitrification (blue) and plant uptake (green). At the annual scale,
 585 stream export accounted for 9.5%, whereas denitrification and plant uptake took up 15%
 586 and 75% of deposited NO_3^- , respectively. In other words, as Nitrate enters this system via
 587 precipitation, plant uptake can play a significant role in reducing nitrate level, indicating
 588 precipitated nitrate is tightly cycled in the system.



589
 590 **Figure 8.** Stream nitrate dynamics and fluxes at Shale Hills in Example 2. (a) Three simulation
 591 scenarios with different processes are demonstrated here: *transport-only* (dashed line, *tran*),
 592 *transport + N reaction* (gray line, *tran + N react*), *transport + N reaction + plant uptake* (thick black

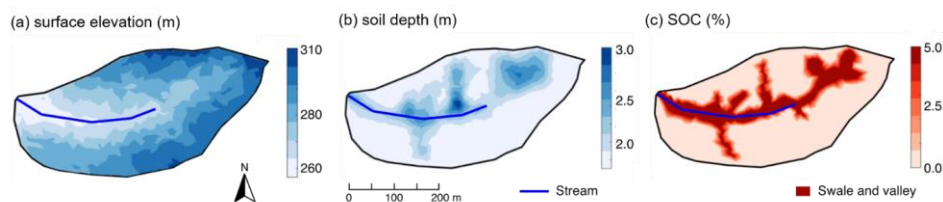
593 line, $tran + N_{react} + upta$), where N reactions include both nitrate leaching and denitrification (see
594 Figure 7); (b) nitrate concentration in precipitation, shallow and deep water; (c) nitrate fluxes and
595 budget. Note that nitrate leaching was ignored in (b) due to its minimal flux as N deposition from
596 rainfall was the dominant input (Weitzman and Kaye, 2018).

597

598 **6.3 Example 3: DOC production and export in a spatially distributed domain**

599 This example showcases the application of BioRT-Flux-PIHM in a spatially
600 distributed mode. This work has been documented with full details in Wen et al. (2020).
601 Here we only introduce some key features and capabilities in the spatially distributed
602 mode. The Shale Hills catchment was discretized into 535 prismatic land elements and
603 20 stream segments through PIHMgis based on the topography (Figure 9a). The
604 heterogeneous distributions of soil depth and solid organic carbon within the domain
605 (Figure 9b-c) were interpolated through ordinary kriging based on field surveys (Andrews
606 et al., 2011; Lin, 2006). Other soil and mineralogy properties such as hydraulic
607 conductivity, van Genuchten parameters, and ion exchange capacity were spatially
608 distributed following intensive field measurements (Jin and Brantley, 2011; Jin et al., 2010)
609 (criticalzone.org/shale-hills/data/).

610



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

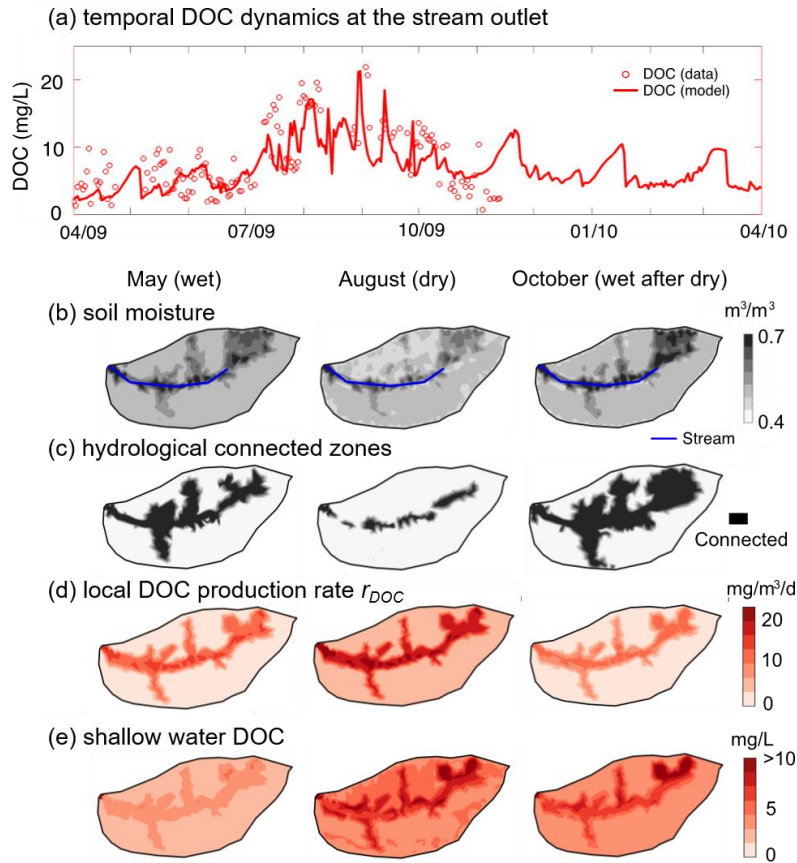
619

620 **Temporal and spatial patterns of DOC production and export.** The model outputs
621 followed the general trend of stream DOC measurements with the model evaluation index

622 NSE of 0.55 for monthly DOC concentration (Figure 10a). NSE ~~ranges from $-\infty$ to 1.0~~
623 ~~(i.e., perfect fit) with~~ values greater than 0.5 are considered good performance for monthly
624 water quality model (Moriassi et al., 2015). The model reproduced high DOC values (~15
625 mg/L) in the dry periods (July-September). The model enabled the identification of
626 reaction hot spots. In May when soil water is relatively abundant, the valley and swales
627 with deeper soils (Figure 10b) are generally wetter compared to the hillslope and ridgetop,
628 and are hydrologically connected to the stream (Figure 10b, c). The distribution of local
629 DOC production rate r_{DOC} and DOC concentration followed that of SOC (Figure 10c) and
630 water content (Figure 10b). Low r_{DOC} in relatively dry planar hillslopes and uplands
631 resulted in low soil water DOC. The average stream DOC (~5 mg/L) reflected soil water
632 DOC in the valley and swales.

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

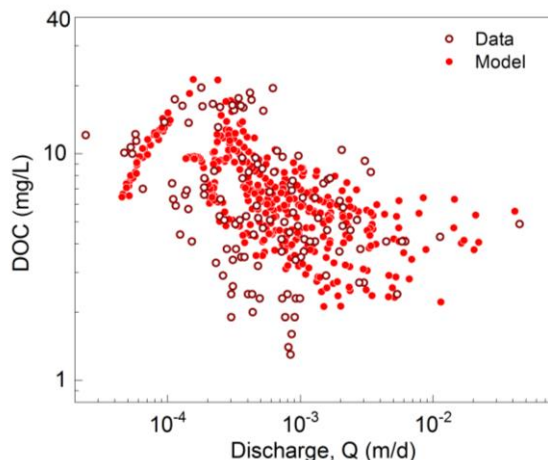
642 In October, precipitation wetted the catchment again. The hydrologically
643 connected zones expanded beyond swales and the valley to the upland hillslopes (Figure
644 10c). The increase in hydrological connectivity zones favored the mixing of shallow water
645 DOC sourced from upland hillslopes (low DOC), swales, and valley (high DOC) into
646 stream rather than only from the stream vicinity with high DOC in the dry August, leading
647 to a drop in stream DOC.



648
 649 **Figure 10.** (a) Temporal dynamics of stream DOC concentration; spatial profiles of (b) shallow
 650 soil moisture, (c) hydrologically connected zones, (d) local DOC production rates r_{DOC} and (e)
 651 shallow water DOC concentration in May (wet), August (dry), and October (wet after dry) of 2009.
 652 The soil DOC and r_{DOC} were high in swales and valley with relatively high shallow water and SOC
 653 content. August had the highest shallow water DOC concentration compared to May and October,
 654 because most DOC accumulated in zones that are disconnected to the stream.

655
 656 **C-Q patterns.** The DOC C-Q relationship showed a non-typical pattern with flushing first
 657 and transitioning into a dilution pattern, with an overall C-Q slope $b = -0.23$ (Figure 11).
 658 At low discharges ($< 1.8 \times 10^{-4}$ m/d) in the summer dry period, the stream DOC mainly
 659 came from the organic-rich swales and valley floor zones with high soil water DOC (Figure

660 10e). With discharge increasing in wetter period (i.e., spring and fall), the contribution
661 from planar hillslopes and uplands with lower DOC concentration increased (Figure 10e),
662 leading to the dilution of stream DOC.



663
664 **Figure 11.** Relationships between daily discharge (Q) and stream DOC concentration. With the
665 increasing Q, the stream water first shifted from the dominance of groundwater with low DOC at
666 very low discharge to the predominance of organic-rich soil water from swales and valley at
667 intermediate discharge. As the discharge increases further, the stream water switches to the
668 dominance of high flow with lower DOC water from planar hillslopes and uplands, resulting in a
669 dilution C-Q pattern (modified from Wen et al., 2020).

670

671 7. Discussion

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

682 This can be particularly useful for watersheds of higher stream orders, where a large
683 proportion of deep water may come from nearby regional aquifers.

684 The advantage and disadvantages of simple versus complex models have long
685 been debated in the modeling community (Fatichi et al., 2016; Li et al., 2020; Wen et al.,
686 2021). The computational cost of solving a spatially distributed, nonlinear, multi-
687 component reactive transport model is high, posing challenges for the application of
688 ensemble-based analysis. With additional reactions and transport processes, the model
689 includes more functions (such as reaction kinetic rate laws) and parameters (e.g., reaction
690 rate constants, surface area) than hydrological models. The complexity brings in issues
691 of equifinality, uncertainty, and data demands (Beven, 2001, 2006; Kirchner et al., 1996).
692 These issues will persist even though reactive transport models will be constrained by
693 additional chemical data.

694 It is in this spirit of balancing the cost and gain that we present both spatial
695 distributed and lumped modes for the BioRT model (Li et al., 2020). Compared to the
696 distributed version, the spatially implicit model requires less spatial data and is
697 computationally inexpensive. It can assess the average dynamics of water and solute
698 dynamics and focus on the interacting processes without resolving spatial details. The
699 lumped approach can accommodate basins with low data availability, and it can be easier
700 for students to learn. In contrast, spatially explicit representations enable the exploration
701 of the “hot spots” (e.g., swales and riparian zones with high soil water DOC concentrations
702 in Figure 10e) and their contribution to stream chemistry at different times. Spatial
703 heterogeneities in watershed properties (e.g., soil types and depth, lithology, vegetation,
704 biomass, and mineralogy) are ubiquitous in natural systems and are challenging to
705 resolve. A general understanding of the linkage between local catchment features and
706 catchment-scale dynamics (e.g., stream concentration dynamics and solute export
707 pattern) is often lacking. The spatially distributed model provides a tool to explore these
708 questions. Ultimately, the choice of the model complexity level depends on research
709 questions that the model is set to answer and the available data. At the end, we all need
710 to balance cost and gain when deciding to use a simple or complex model, striving to be
711 “simple but not simplistic” (Beven and Lane, 2019).

712 **8. Summary and conclusion**

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

735
736 **Data availability.** Field data (e.g., discharge, stream chemistry) is archived at Shale Hills
737 data portal: <http://criticalzone.org/shale-hills/data/datasets/> or maintained at HydroShare:
738 <https://www.hydroshare.org/group/147>.

739
740 **Code availability.** The current model release (BioRT-Flux-PIHM v1.0) is archived at:
741 <https://doi.org/10.5281/zenodo.3936073>. Documentation, source code, and examples
742 are available at GitHub repository: <https://github.com/Li-Reactive-Water-Group/BioRT-Flux-PIHM>.
743

744
745 **Competing interests.** The authors declare that they have no conflict of interest.
746
747 **Author contributions.** LL conceived the model idea and oversaw the model
748 development. WZ coded the BioRT module, verified the code against the benchmark
749 reactive transport model CrunchTope, and applied and tested the model at Shale Hills
750 watershed. YS developed the deep groundwater component and integrated the BioRT-
751 Flux-PIHM v1.0 into the MM-PIHM family. WH, LS, KS, DK, BS, and GHCH tested the
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