

Active spreading: Hydraulics for enhancing groundwater remediation

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ABSTRACT

During in situ groundwater remediation, reactions occur in a narrow reaction front, where the amendment and contaminant are close enough to mix. Active spreading, in which injection or extraction wells create spatially-variable velocity fields, can be used to increase the surface area of the reaction front, thereby enhancing reaction. This study uses four active spreading flow fields, that are building blocks to more complex remediation hydraulics, to evaluate how the flow field and the plume position control contaminant degradation in both homogeneous and heterogeneous aquifers. At the plume scale, reaction depends on mechanical dispersion across the reaction front, which is proportional to both the local velocity and the local contaminant concentration gradient. Mechanical dispersion, and consequently the amount of degradation, is highest when the reaction front is perpendicular to the local velocity, producing a high local dispersion coefficient. This effect is amplified where flow is diverging, due to sharpening of the concentration gradient.

INTRODUCTION

In-situ remediation is a common method for cleaning up contaminated groundwater. During in-situ remediation, a chemical or biological amendment is introduced into the aquifer to react with and degrade the contaminant. Reaction between the amendment and the contaminant take

24 place only within a relatively narrow region, called the reaction front, where the contaminant
25 and amendment are sufficiently close to react. Degradation reactions along the reaction front are
26 controlled by spreading and mixing. Mixing involves the smoothing of concentration gradients by
27 molecular diffusion and pore-scale dispersion (Bellin et al. 2011), which brings molecules together
28 to react. Spreading involves the reconfiguration of the contaminant and amendment plumes as
29 a result of spatial variations in velocity (Le Borgne et al. 2010). Spreading enhances mixing by
30 sharpening the concentration gradient and elongating the interface along which mixing can occur
31 (Le Borgne et al. 2013). It is helpful to distinguish passive spreading and active spreading. Passive
32 spreading results from velocity variations caused by aquifer heterogeneity, while active spreading
33 results from induced velocity variations (e.g., by injecting or withdrawing water through wells).

34 The key to successful in-situ remediation is to deliver the amendment in such a way that the
35 reaction front is spread throughout the contaminant plume. The most basic method of amendment
36 delivery is to install a well in the plume, typically where the contaminant concentrations are highest,
37 through which the amendment is injected. Then the contaminant and amendment plumes are left to
38 travel through the subsurface under natural hydraulic gradients. The reaction front is reconfigured
39 by passive spreading due to aquifer heterogeneity, which may elongate the reaction front, thereby
40 creating more contact between the contaminant and amendment (Le Borgne et al. 2013; Le Borgne
41 et al. 2014; Bandopadhyay et al. 2017). Passive spreading also increases the concentration gradients
42 at the reaction front, increasing the driving force for mass flux by molecular diffusion and dispersion
43 (Ou and Ranz 1983; Le Borgne et al. 2013). Although this passive spreading enhances reaction to
44 some degree, large portions of the amendment plume may remain isolated from the contaminant
45 plume; thus, the delivery of the amendment into the contaminant plume is incomplete.

46 Most in-situ remediation installations rely on the use of injection and extraction wells to enhance
47 delivery of the amendment and to direct the movement of the amendment through the contaminated
48 area. Injection and extraction wells can be used in pairs (dipoles) for smaller plumes, in multi-
49 well cells, or in multiple groups of multi-well cells for large plumes (Suthersan et al. 2009).
50 More advanced in-situ remediation designs reconfigure the active spreading flow fields during the

51 remediation process, which has been shown to further enhance remediation (Suthersan et al. 2015).
52 In these active spreading systems, the reaction front is reconfigured as imposed flow forcings from
53 the injection and extraction wells create spatially variable velocity fields (Zhang et al. 2009; Lester
54 et al. 2010; Trefry et al. 2012; Mays and Neupauer 2012; Suthersan et al. 2015). As with passive
55 spreading, active spreading elongates the reaction front and increases concentration gradients along
56 the reaction front, which has been shown to enhance mixing (Le Borgne et al. 2010) and reaction
57 (Piscopo et al. 2013; Bandopadhyay et al. 2017; Rodríguez-Escales et al. 2017; Cho et al. 2019).

58 While active spreading has been shown to enhance reaction during in-situ remediation, the
59 nature of the causal relationships between the active spreading flow fields, the geometry of the
60 reaction front, and the degree of enhanced mixing-controlled reaction in porous media are not
61 well understood. To design remediation systems that most effectively capitalize on the spatio-
62 temporal variations in spreading and mixing conditions, an investigation into the relationship
63 between active spreading and mixing-controlled reaction under flow fields typical of remediation
64 systems is necessary. Accordingly, the goal of this study is to evaluate how active spreading at the
65 plume scale impacts the overall degradation of a contaminant plume.

66 Several studies have investigated spreading, mixing, and reaction at scales much smaller than
67 the plume scale. A subset of studies consider situations in which one species is invading the region
68 occupied by another; thus, the reaction front is approximately perpendicular to the local velocity
69 vectors. Due to pore-scale velocity variations and incomplete mixing in the pore space, spreading
70 occurs as the invading species forms lamellae within the pore (Le Borgne et al. 2013), which
71 elongates the fluid interface and sharpens the concentration gradients that drive diffusive mixing
72 and reaction. The relationship between lamella formation and diffusive mixing and reaction has
73 been studied in uniform flow with passive spreading (Chiogna et al. 2012; de Anna et al. 2014;
74 Le Borgne et al. 2014; Perez et al. 2020), shear flow (Le Borgne et al. 2014; Bandopadhyay et al.
75 2017), stratified random flow (Le Borgne et al. 2014), and radial flow (Le Borgne et al. 2014).
76 While radial flow falls under the category of active spreading, it is a special case in which the bulk
77 velocity is everywhere perpendicular to the plume interface on the macro-scale. Where one species

78 invades another at the plume scale (as shown in the left callout of Figure 1), the lamella formation
79 at the pore scale underpin the up-scaled process of longitudinal dispersion, with dispersion length
80 scales that have been shown to increase with time at a rate that depends on the heterogeneity of the
81 porous medium (Le Borgne et al. 2013; Perez et al. 2020).

82 Several other studies have investigated spreading, mixing, and reaction where the reaction
83 front is approximately parallel to the local velocity vector, as shown in the right callout of Figure
84 1. This reaction front geometry forms from a continuous point release of a solute (Rolle et al.
85 2009; Chiogna et al. 2011; Chiogna et al. 2012), which grows into an elongated plume, with
86 high concentration gradients and a long interface parallel to flow, such that reactions depend on
87 transverse dispersion. A related study considered transverse dispersion between two adjacent
88 solute plumes with the interface aligned in the main flow direction (Cirpka et al. 2011). Mixing
89 and reaction were enhanced by the spatial variability in velocity caused by spatially segregated
90 high permeability zones (Rolle et al. 2009) or small-scale heterogeneity (Cirpka et al. 2011), while
91 temporal variations in uniform flow had little effect (Rolle et al. 2009).

92 These past studies of spreading, mixing, and reaction have considered flow fields and plume
93 configurations in which the orientation of the plume boundary is either perpendicular to or parallel
94 to the direction of the bulk flow. During in-situ remediation, an amendment plume (species B in
95 Figure 2) is emplaced within a contaminant plume (species A in Figure 2), with the reaction front
96 represented by the interface between the two plumes. Since the amendment plume and reaction
97 front are finite in size, the orientation of the reaction front relative to the local velocity varies along
98 the reaction front between the two previously investigated end-member orientations (perpendicular
99 and parallel to flow). Furthermore, under active spreading flow fields, the reaction front experiences
100 different local velocities over time as it travels through the spatially varying velocity field. Thus, an
101 individual segment of the reaction front will experience different orientations relative to the local
102 velocity in time, which was not considered in past studies of spreading, mixing, and reaction. In
103 these more complicated active spreading flows, the local orientation of the reaction front relative
104 to the local flow direction varies both spatially and temporally, leading to spreading and mixing

105 conditions that evolve in time. Thus, local details drive plume-scale reaction.

106 In this paper, we investigate the relative importance of the two orientations of the reaction
107 front relative to the local velocity and the spatio-temporal variation of these relative orientations
108 in promoting reaction during in-situ remediation of contaminated groundwater. Understanding
109 how the local geometries of the reaction front and the flow field control the amount of reaction
110 will enable design of active spreading protocols that exploit the local efficiencies to increase the
111 overall amount of degradation during in-situ remediation. We consider the plume scale and focus
112 on the continuum behavior over the Darcy and larger scales, because that is the scale of interest
113 in groundwater remediation activities. While several studies have shown that incomplete mixing
114 in pores limits reaction (Gramling et al. 2002; Raje and Kapoor 2000), the errors associated with
115 neglecting incomplete mixing at the Darcy and larger scales may be negligible for fast reactions
116 (Porta et al. 2013), which we consider herein.

117 In this work, we derive an explicit relationship between the global reaction rate, quantified
118 for the plume as a whole, and local characteristics of the plumes and velocity field along the
119 reaction front. We apply this relationship to four different active spreading protocols that represent
120 components of flow fields that may be used to drive flow during in situ remediation, and we compare
121 the result to uniform flow (i.e., no active spreading). Since this study focuses on active spreading,
122 we initially remove the effects of passive spreading by considering homogeneous aquifers only.
123 This simplification allows us to evaluate the features of the flow field alone that promote reaction
124 on the plume scale, and to identify patterns of flow and reaction front geometry that produce
125 the most degradation. We evaluate how the topology of the flow field enhances or inhibits the
126 overall degradation of the contaminant. Later we extend the investigation to heterogeneous aquifers
127 and we demonstrate that the same patterns of flow and reaction front geometry that produce
128 the most degradation in homogeneous aquifers also generally produce the most degradation in
129 the heterogeneous aquifers we tested. Insights gained from this research will provide crucial
130 information for the optimal design of active spreading groundwater remediation systems in the
131 field.

132 REACTIVE TRANSPORT THEORY

133 In the first part of this study, we consider a two-dimensional, rectangular, confined, homoge-
 134 neous, isotropic aquifer, centered at the origin. We assume the aquifer contains a circular plume of
 135 species A (contaminant) surrounding a circular plume of species B (amendment) as shown in Figure
 136 2. The chemical reaction between the species follows an instantaneous, irreversible bimolecular
 137 reaction, given by $A + B \rightarrow C$, which can represent, for example, oxidation of chlorinated solvents
 138 by potassium permanganate (Yoon and Schwartz 1999). We assume that reactive transport is
 139 governed by the advection-dispersion-reaction equation, given by

$$\frac{\partial C_i}{\partial t} = -\nabla \cdot (\mathbf{v}C_i) + \nabla \cdot \mathbf{D}\nabla C_i + R_i, \quad (1)$$

140 where C_i is the dimensionless concentration of the i^{th} species ($i = A, B, C$), t is dimensionless time,
 141 \mathbf{v} is the dimensionless groundwater velocity vector, R_i is the dimensionless reaction rate of species
 142 i , with $R_A = R_B = -R_C$, and \mathbf{D} is the dimensionless dispersion tensor, with components given by

$$\begin{aligned} D_{xx} &= \alpha_L \frac{v_x^2}{|\mathbf{v}|} + \alpha_T \frac{v_y^2}{|\mathbf{v}|} + D_m, \\ D_{xy} = D_{yx} &= (\alpha_L - \alpha_T) \frac{v_x v_y}{|\mathbf{v}|}, \\ D_{yy} &= \alpha_L \frac{v_y^2}{|\mathbf{v}|} + \alpha_T \frac{v_x^2}{|\mathbf{v}|} + D_m, \end{aligned} \quad (2)$$

143 where α_L and α_T are the dimensionless longitudinal and transverse dispersivities, respectively,
 144 and D_m is the dimensionless molecular diffusion coefficient. The dimensionless velocity comes
 145 from the dimensionless form of Darcy's law, given by $\mathbf{v} = -(K/n)\nabla h$, where K is dimensionless
 146 hydraulic conductivity, n is porosity, and h is dimensionless head obtained from

$$\nabla \cdot Kb\nabla h + \sum_{j=1}^3 Q_j \delta(\mathbf{x} - \mathbf{x}_{w_j}) = 0, \quad (3)$$

147 where b is the dimensionless thickness of the aquifer, and Q_j in the dimensionless injection rate at
 148 well j ($Q_j < 0$ indicates extraction), located at \mathbf{x}_{w_j} . Boundary conditions on (1) are $\nabla C_i \cdot \mathbf{n} = 0$

149 on all boundaries, where \mathbf{n} is the outward unit normal vector. This initial condition is shown
150 graphically in Figure 2. The boundary conditions on (3) are

$$\frac{\partial h}{\partial y} = 0 \text{ at } y = \pm L/2 \quad (4)$$

$$h = h_L \text{ at } x = -L/2 \quad (5)$$

$$h = 0 \text{ at } x = L/2, \quad (6)$$

151 where L is the dimensionless length of the domain and h_L is the dimensionless head at the boundary
152 at $x = -L/2$. For the active spreading scenarios, we assume background flow is negligible, so
153 $h_L = 0$; for comparison, we also consider uniform flow (i.e., no active spreading), with $h_L \neq 0$.

154 All dimensionless lengths are relative to the diameter, d , of the initially circular reaction front.
155 Because we are considering steady flow, the situation does not have a natural characteristic time.
156 Instead, we define the characteristic time in terms of a characteristic pumping rate, defined as the sum
157 of the magnitudes of all active pumping rates (per unit aquifer thickness). With this characteristic
158 length and pumping rate, the characteristic time is the time required for the characteristic pumping
159 rate to fill a cuboid whose top and bottom surfaces are squares of length d and whose height is equal
160 to the aquifer thickness. Dimensionless concentrations are relative to the maximum concentration
161 of species A in the aquifer at $t = 0$. See Section 1 of the Supplemental Material for the development
162 of the dimensionless forms of the equations.

163 **NUMERICAL MODEL AND ACTIVE SPREADING PROTOCOLS**

164 We consider four different active spreading protocols (Figure 3). The two basic active spreading
165 protocols that we consider are Diverging (injection in Well 1) and Converging (extracting from
166 Well 3) protocols (Figure 3a,b). These two protocols can be superimposed in space and time
167 to create more complex active spreading flow fields. The third active spreading protocol is the
168 Dipole protocol (Figure 3c), which is the superposition of the Diverging and Converging protocols.
169 The Stagnation protocol (Figure 3d) is similar the the Dipole protocol, except that the injected
170 water is split between Wells 1 and 2, leading to significant elongation of the reaction front. This

171 protocol is used to evaluate the importance of spreading on reaction. For all four active spreading
172 protocols, ambient flow is assumed negligible, and the amendment and contaminant plumes are
173 initially between Wells 1 and 2 (see plumes in Figure 2 and reaction front in Figure 3); thus the
174 amendment plume is assumed to have been emplaced with a well at (x_p, y_p) that is not shown and
175 not used in the active spreading protocols.

176 For comparison, we also evaluate a uniform flow field (Uniform protocol), which, by definition,
177 exhibits no plume spreading because spreading is caused by velocity variations which are absent in
178 uniform flow in a homogeneous aquifer. To make equivalent comparisons across protocols, all pro-
179 tocols, except the Stagnation protocol, are designed to have the same advective travel time between
180 the center of the initial plume $(x_p, y_p) = (-1.33, 0)$ and $(x, y) = (1.33, 0)$, allowing comparison of
181 protocols across time. For the Uniform protocol, this design resulted in a dimensionless velocity
182 higher than typical groundwater flow velocities.

183 The flow fields for the four protocols and the uniform flow condition are generated by solving
184 (3) numerically using MODFLOW-2000 (Harbaugh et al. 2000). Parameter values for the flow
185 simulations are given in Table 1, and dimensionless pumping rates for each protocol are shown in
186 Table 2.

187 The reactive transport equation (1) is solved numerically using RW3D (Salamon et al. 2006),
188 which uses random walk particle tracking. Particle tracking is a common method for modeling
189 solute transport in aquifers known for its computational efficiency and absence of numerical
190 dispersion (Berkowitz et al. 2006; Le Borgne et al. 2008a; Le Borgne et al. 2008b). Parameter
191 values are given in Table 1. Instead of solving (1) directly for species A, B, and C, we simulated
192 the transport of two conservative species A+C and B+C, eliminating the reaction term in (1)
193 (Gramling et al. 2002). Assuming a 1:1 mass ratio and 1:1 stoichiometric ratio of A and B, the

194 mass concentrations of A, B, and C are obtained from the conservative components as

$$C_C(x, y, t) = \min [2C_{A+C}(x, y, t), 2C_{B+C}(x, y, t)], \quad (7)$$

$$C_A(x, y, t) = C_{A+C}(x, y, t) - 0.5C_C(x, y, t), \quad (8)$$

$$C_B(x, y, t) = C_{B+C}(x, y, t) - 0.5C_C(x, y, t). \quad (9)$$

195 The initial distributions of concentrations of species A and B were obtained using analytical
196 expressions of transport in radial flow presented in [Neupauer et al. \(2020\)](#). The initial condition
197 is created by injecting fluid at a unit rate at location $(x, y) = (-1.33, 0)$. The injection period was
198 divided into two steps. In the first step of duration of $t_A = 0.466$, the injected fluid contained
199 species A at a dimensionless concentration of 1.02. In the second step of duration $t_B = 0.155$,
200 the injected fluid contained species B at a dimensionless concentration of 4.08. Advection (due to
201 injection), dispersion, and reaction were all simulated during the injection steps. The distributions
202 of the species A and B at the end of the injection period are shown in Figure 2, which corresponds
203 to the initial condition (at $t = 0$) for all protocols, while the concentration of species C was set to
204 zero everywhere. The maximum dimensionless concentrations of species A and B in the initial
205 plume distribution are 1.00 and 4.08, respectively; and the dimensionless masses of species A and
206 B in the aquifer at time $t = 0$ are 0.39 and 0.54, respectively. During emplacement of the initial
207 plumes, no flow other than the advection caused by injection of the plumes was present. At time
208 $t = 0$, the flow for each protocol was initiated and was assumed to immediately attain steady state
209 conditions.

210 In RW3D, species A+C and B+C are represented as collections of 6×10^6 and 2×10^6 randomly
211 placed particles, respectively, each of which has dimensionless mass of 6.53×10^{-8} and 2.73×10^{-7} ,
212 respectively. The smoothness of the concentration distribution depends on the number of particles
213 per unit area. Since the plume of species B covers a smaller area than the plume of species A, fewer
214 particles of species B were used. Relative to the plume centroid at (x_p, y_p) , particle positions in the
215 radial direction were randomly placed to match the distribution in Figure 2b, and particle position

216 in the angular direction were drawn from a uniform distribution.

217 At each output time, the particle positions and masses provided by RW3D were converted to
218 a concentration field by binning the particles into square bins of size 0.0089. Subsequently, the
219 concentration field was smoothed by convolution with a two-dimensional Gaussian with mean
220 zero. The standard deviation of the Gaussian ranged from 0.0089 at early times to 0.3560 at late
221 times. A smaller standard deviation was necessary for early time to avoid over smoothing the sharp
222 concentration gradients along the reaction front while a larger standard deviation was necessary
223 for later time because the area of the plumes increases with time. All analyses and presentation of
224 results use the smoothed concentration distributions.

225 GLOBAL AND LOCAL MEASURES OF SPREADING, MIXING, AND REACTION

226 The metrics used in this paper to quantify spreading, mixing, and reaction include global
227 measures that characterize the behavior of the entire plume, and local measures that characterize
228 behavior on scales much smaller than the plume scale. We consider three global measures. Since
229 the purpose of groundwater remediation is to remove the contaminant from the aquifer, the first
230 global measure is the cumulative amount of species A (contaminant) that has reacted, M_{rxn} at or
231 before time t , given by

$$M_{\text{rxn}}(t) = M_{A_0} - M_A(t), \quad (10)$$

232 where $M_A(t)$ is the dimensionless mass of species A in the domain at time t , calculated as

$$M_A(t) = \int \int_{x,y} nC_A(x, y, t) dx dy \quad (11)$$

233 and $M_{A_0} = M_A(t = 0)$ is the initial dimensionless mass of species A. Note that (10) is valid as long
234 as the mass of species A extracted from an active spreading extraction well is minimal. For the
235 scenarios in this study, less than 2.4% of the mass of species A was extracted in any simulation,
236 which we consider minimal.

237 The second global measure is the global reaction rate dM_{rxn}/dt , which is a measure of the
238 required duration of the remediation. The global reaction rate can be obtained by evaluating the

239 change in M_{rxn} between subsequent time steps of the numerical simulation. We also note that it is
 240 related to the total mass flow rate of species A (which is equivalent to the mass flow rate of species
 241 B) across the reaction front Γ , given by

$$\frac{dM_{\text{rxn}}}{dt} = b \oint_{\Gamma} J_{A\beta} d\Gamma, \quad (12)$$

242 where $J_{A\beta}$ represents the mass flux of species A across the reaction front and β is the direction
 243 perpendicular to the reaction front (see Fig. 2c). For very fast, irreversible reactions, the reaction
 244 front reduces to a surface along which the concentrations of both species vanish; therefore, the
 245 advective mass fluxes of both species across the reaction front vanish. Furthermore, if molecular
 246 diffusion is negligible (see justification in Figure S1 of the supplemental material), $J_{A\beta}$ represents
 247 the dispersive mass flux in the β direction,

248 The third global measure is the reaction front length, L_{Γ} , which is a measure of spreading. This
 249 measure allows us to evaluate the overall importance of spreading in in situ remediation.

250 We use local measures to quantify how local characteristics of the plume, reaction front geometry
 251 and the velocity field affect contaminant degradation. As shown in (12), the global reaction rate is
 252 controlled by the local dispersive mass flux, $J_{A\beta}$, of species A across the reaction front, given by

$$J_{A\beta} = -nD_{\beta} \frac{\partial C_A}{\partial \beta}, \quad (13)$$

253 where D_{β} is the dispersion coefficient in the direction β and $\partial C/\partial \beta$ is the concentration gradient
 254 of species A in the direction β . Thus we identify $J_{A\beta}$, D_{β} , and $\partial C/\partial \beta$ are three local measures of
 255 mixing.

256 These local mixing measures can be evaluated by equating $J_{A\beta}$ with its components in the
 257 direction of the local velocity vector, s , and in the direction perpendicular to the local velocity
 258 vector η (see Fig. 2c). The local dispersive mass fluxes in the s and η directions are, respectively,

$$J_{As} = -n\alpha_L |\mathbf{v}| \frac{\partial C_A}{\partial s} \quad (14)$$

259 and

$$J_{A\eta} = -n\alpha_T|\mathbf{v}|\frac{\partial C_A}{\partial \eta} . \quad (15)$$

260 Let θ be the angle between the local velocity vector and the direction locally perpendicular to
 261 the reaction front (Fig. 2c). Then $J_{A\beta}$ is given by

$$\begin{aligned} J_{A\beta} &= J_{A\eta} \sin \theta + J_{As} \cos \theta \\ &= -n\alpha_T|\mathbf{v}|\frac{\partial C_A}{\partial \eta} \sin \theta - n\alpha_L|\mathbf{v}|\frac{\partial C_A}{\partial s} \cos \theta \\ &= -n\alpha_T|\mathbf{v}|\frac{\partial C_A}{\partial \beta} \sin^2 \theta - n\alpha_L|\mathbf{v}|\frac{\partial C_A}{\partial \beta} \cos^2 \theta \\ &= -n \left(\alpha_L \cos^2 \theta + \alpha_T \sin^2 \theta \right) |\mathbf{v}|\frac{\partial C_A}{\partial \beta} \\ &= -n\alpha_\beta|\mathbf{v}|\frac{\partial C_A}{\partial \beta} \end{aligned} \quad (16)$$

262 where $\alpha_\beta = \alpha_L \cos^2 \theta + \alpha_T \sin^2 \theta$ is the dispersivity in the β direction and D_β is the dispersion
 263 coefficient in the β direction, given by

$$D_\beta = \alpha_\beta|\mathbf{v}|. \quad (17)$$

264 The final expression in (16) is obtained by substituting (14) and (15) into the first line to obtain the
 265 second line, and the third line follows from the second line through the use of the chain rule on the
 266 partial derivatives of C_A .

267 Integrating (12) over dimensionless time after substituting (16) and (17) demonstrates the
 268 relationship between the global measure $M_{\text{rxn}}(t)$ and the three local mixing measures:

$$\begin{aligned} M_{\text{rxn}}(t) &= b \int_0^t \oint_{\Gamma} J_{A\beta} d\Gamma dt' \\ &= nb \int_0^t \oint_{\Gamma} \frac{\partial C_A}{\partial \beta} D_\beta d\Gamma dt' \\ &= nb \int_0^t \oint_{\Gamma} \frac{\partial C_A}{\partial \beta} \alpha_\beta |\mathbf{v}| d\Gamma dt' . \end{aligned} \quad (18)$$

269 Finally, we consider one local measure of spreading called instantaneous strain (Zhang et al.
 270 2009), which is given by

$$\zeta = \frac{d\rho}{dt} \frac{\Delta t}{\rho} \approx \frac{\Delta\rho}{\rho}, \quad (19)$$

271 where ρ is the length of a small arc of the reaction front Γ , Δt is the time interval over which
 272 the strain is calculated, and $\Delta\rho$ is the change in the arc length over time Δt . In this work, strain
 273 was estimated by calculating $\Delta\rho$ over a time interval of $\Delta t = 0.016$. The concentration gradient,
 274 $\partial C_A / \partial \beta$, was approximated through linear interpolation of $C_A(\beta)$ at each point along the reaction
 275 front. We applied a Savitzky-Golay filter (Orfanidis 1995) to the concentration gradient along
 276 the reaction front to remove noise resulting from the numerical solution. We used a second order
 277 polynomial, with a frame length (normalized by the number of points along the curve) of between
 278 0.0382 and 0.4437 to ensure maximum smoothing while still preserving the spatial character. M_{rxn}
 279 was calculated from (18) using this smoothed curve and is plotted as dots in Figure 4a. The results
 280 are visually indistinguishable from the those calculated using (10) and (11), verifying the accuracy
 281 of the smoothed curves.

282 RESULTS FOR A HOMOGENEOUS AQUIFER

283 Figure 4a shows that $M_{\text{rxn}}(t)$ (normalized by M_{A0}) grows with time for each protocol. Compared
 284 to the Uniform protocol (no active spreading, Figure 4d), the Diverging and Dipole protocols
 285 produce more reaction, with the Dipole protocol producing only slightly more; while the Converging
 286 and Stagnation protocols produce less reaction. Thus, having active spreading is not sufficient to
 287 enhance reaction relative to uniform flow.

288 For each protocol, the temporal evolution of the dispersive mass flow rate of species A across
 289 the reaction front, dM_{rxn}/dt , is shown in Figure 4b (normalized by M_{A0}). For the Diverging,
 290 Dipole, and Uniform protocols, dM_{rxn}/dt decreases monotonically at a decreasing rate. Both the
 291 Converging and Stagnation protocols, however, show brief periods when dM_{rxn}/dt increases. At
 292 early times, dM_{rxn}/dt for the Diverging and Dipole protocols is higher than for the Uniform protocol
 293 and is lower for late times. The opposite behavior is observed for the Converging and Stagnation

294 protocols (Figure 4e).

295 The premise of active spreading is to increase the length, L_Γ , of the reaction front, which, in
296 turn, increases the length along which mixing, and therefore reaction, can occur. The temporal
297 evolution of L_Γ for each protocol is shown in Figure 4c (normalized by the initial length of the
298 reaction front, L_{Γ_0}). The position of the reaction front is obtained by finding the zero contour of
299 $C_{A+C} - C_{B+C}$ in Matlab, to which we applied a Savitzky-Golay filter (Orfanidis 1995) to remove
300 noise resulting from the numerical solution. We used a second order polynomial, with a frame
301 length (normalized by the number of points along the curve) of between 0.0382 and 0.1656 to
302 ensure maximum smoothing while still preserving the spatial character.

303 With uniform flow, as the plumes move down-gradient, the reaction front propagates into the
304 plume of species A, from a Lagrangian perspective (see Figure 5a), because the concentration
305 of species B is higher than the concentration of species A. Thus, the length of the reaction front
306 increases over time, even without active spreading. With active spreading, the reaction front length
307 increases even more (Figure 4f). However, the reaction front length does not by itself explain the
308 relative amount of reaction shown in Fig. 4a. For example, the Stagnation protocol has the longest
309 reaction front (dashed line in Fig. 4c), but it produces the least amount of reaction (dashed line in
310 Fig. 4a). Thus, global stretching alone is not sufficient to explain the global amount of reaction.

311 To investigate the causes of the relative amount of contaminant degradation across the four
312 active spreading protocols, we evaluate the local measures along the reaction fronts. Figures 5–9
313 show the plumes of species A and B and the position of the reaction fronts at four different times
314 for the Uniform, Diverging, Converging, Dipole, and Stagnation protocols, respectively, along with
315 the time evolution of the dispersion coefficient D_β , instantaneous strain ζ , concentration gradient
316 $\partial C_A / \partial \beta$, and dispersive mass flux, $J_{A\beta}$. Note that the color scale for C_A in the plume plot is
317 different than in Figure 2 to allow for more resolution.

318 The overall mass reacted depends, in part, on the reaction front length. Comparison of Fig. 4c
319 and Figs. 5b–9b show that the reaction front length is correlated with the amount of positive strain.
320 Instantaneous strain is positive where the reaction front is stretched locally. Local stretching occurs

321 where flow is diverging and the reaction front is perpendicular to flow, e.g., at $\ell/L_\Gamma = 0$ and 0.5 for
 322 the Diverging and Stagnation protocols for all times (Fig. 6, Fig. 9); at $\ell/L_\Gamma = 0$ for Dipole protocol
 323 at early times (Fig. 8). Local stretching also occurs where flow is converging and the reaction front
 324 is parallel to flow, e.g., at $\ell/L_\Gamma = 0.25$ for the Converging protocol at all times Fig. 7; at $\ell/L_\Gamma \approx 0.3$
 325 for the Dipole protocol at later times (Fig. 8). These relationships are summarized in Table 3. The
 326 Stagnation protocol has the longest reaction front, consistent with its having the highest positive
 327 strain (see Figures 5-9b), followed by the Converging, Diverging, Dipole, and Uniform protocols,
 328 in decreasing order. For the Converging protocol, the amount of positive strain is low at early times
 329 and increases at later times (white shaded region in Figure 7b,c), consistent with the reaction front
 330 length being shorter at early times and higher at later times relative to the other protocols (Fig. 4c).
 331 These results are summarized in Table 3.

332 The overall mass reacted also depends on the dispersive mass flux. Figures 5–9 show that in
 333 all cases the spatial average of the dispersive mass flux, $J_{A\beta}$, across the reaction front (subplot c)
 334 decreases over time. Since the instantaneous reaction rate is directly related to the dispersive mass
 335 flux, the decrease in $J_{A\beta}$ is consistent with the decreasing trend of dM_{rxn}/dt in Figure 4b. The
 336 differences in the global reaction rate and other global measures across the different protocols can
 337 be explained by the variability of D_β and $\partial C_A/\partial \beta$, which are components of $J_{A\beta}$, along the front.

338 Figures 5b–9b show the variation of $D_\beta = \alpha_\beta |\mathbf{v}|$ along the reaction front for all protocols. The
 339 value of local dispersivity, α_β , depends on the direction of the local velocity vector relative to the
 340 orientation of the reaction front through (16), as summarized in Table 3. Since $\alpha_L > \alpha_T$, α_β is
 341 higher where the reaction front is perpendicular to the local flow. Therefore, D_β tends to be higher
 342 where the reaction front is perpendicular to local flow.

343 The magnitude of velocity also affects D_β . All protocols have approximately the same mean
 344 velocity but different spatial distributions of velocity, which impacts the amount of reaction. The
 345 Diverging protocol exhibits high velocities for $x < 0$ (see Fig. 3), which corresponds to the locations
 346 of the reaction front at early times, when the concentration gradient remains high (Figs. 5c–9c).
 347 Thus, the Diverging protocol exhibits high dispersive mass flux and high amounts of reaction,

348 especially at early times (Fig. 4d). On the other hand, the Converging protocol exhibits high
 349 velocities for $x > 0$ (see Fig. 3), which corresponds to locations of the reaction front at late times,
 350 when the concentration gradient has diminished (Fig. 5c–9c). Thus, the Converging protocol
 351 exhibits less dispersive mass flux and less overall reaction, with the rate of reaction increasing over
 352 time (Fig. 4d). The Dipole protocol exhibits moderate velocities near the upstream and downstream
 353 wells and lower velocities near $x = 0$. resulting in less reaction than the Diverging protocol at
 354 early times (Fig. 4d), but the rate of reaction decreases more slowly (Fig. 4e). The Stagnation
 355 protocol also exhibits moderate velocities at the upstream and downstream wells, but it has very
 356 low velocities near the stagnation point, so D_β is low near the stagnation point even though the
 357 reaction front is perpendicular to local flow, leading to less reaction overall than the other protocols.

358 The concentration gradient, $\partial C_A / \partial \beta$ (Figures 5c–9c), is smoothed by dispersion and negative
 359 strain, and is sharpened by positive strain. For a given protocol, the concentration gradient decreases
 360 more quickly over time where D_β is high and more slowly where it is low. For example, in all
 361 protocols except Stagnation, $\partial C_A / \partial \beta$ decreases rapidly near $\ell / L_\Gamma = 0$ and 0.5 and slowly near
 362 $\ell / L_\Gamma = 0.25$, corresponding to high and low values, respectively, of D_β . The concentration gradient
 363 decreases most rapidly where strain is negative and the reaction front is perpendicular to flow (i.e.,
 364 $\alpha_\beta \approx \alpha_L$, so D_β is high), such as near $\ell / L_\Gamma = 0$ or 0.5 in the Converging protocol. On the other hand,
 365 $\partial C_A / \partial \beta$ decreases most slowly where strain is positive and the reaction front is parallel to flow (i.e.,
 366 $\alpha_\beta \approx \alpha_T$, so D_β is low, such as near $\ell / L_\Gamma = 0.25$ in the Converging protocol. The concentration
 367 gradient decreases at a moderate rate where $\zeta > 0$ and D_β is high, and where $\zeta < 0$ and D_β is
 368 low. These behaviors are summarized in Table 3. In uniform flow, $\zeta = 0$ everywhere at all times,
 369 which neither sharpens nor smooths the concentration gradient. Thus, in the Uniform protocol,
 370 $\partial C_A / \partial \beta$ decreases in time as a result of dispersion only. As a result, the rate of degradation in the
 371 Converging protocol increases over time relative to the Uniform protocol because the concentration
 372 gradient remains high (due to positive strain along most of the reaction front, see Table 3). On the
 373 other hand, the rate of degradation in the Diverging protocol decreases over time relative to the
 374 Uniform protocol because the concentration gradient decreases more rapidly (high α_β where strain

375 is positive, and low α_β where strain is negative).

376 Given that the dispersive mass flux $J_{A\beta}$ (16) depends on the product of D_β and $\partial C_A/\partial\beta$, and
377 $\partial C_A/\partial\beta$ also depends on D_β , $J_{A\beta}$ is more sensitive to the spatio-temporal pattern of D_β than
378 to that of $\partial C_A/\partial\beta$. Thus, $J_{A\beta}$ (Figs, 5c–9c) follows the same pattern as D_β , except where the
379 concentration of species A (and therefore its concentration gradient and mass flux) falls to zero
380 (e.g., near $\ell/L_\Gamma = 0$ and 0.5 at late times in the Diverging protocol, where $C_A \approx 0$, but D_β is
381 relatively high). Thus, the overall amount of reaction is higher in active spreading protocols in
382 which the reaction front tends to be more perpendicular to the local velocity vectors, i.e., where
383 $\alpha_\beta \approx \alpha_L$, such as in the Diverging and Dipole protocols. Likewise, the overall amount of reaction is
384 lower in active spreading protocols in which the reaction front tends to be more parallel to the local
385 velocity vectors, such as in the Converging and Stagnation protocols. For comparison, in uniform
386 flow, the reaction front has equal proportions perpendicular to and parallel to the local velocity
387 vectors, so the amount of reaction in the Uniform protocol falls in the middle of the five protocols
388 considered in this work.

389 RESULTS FOR HETEROGENEOUS AQUIFERS

390 In this section, we investigate how heterogeneity (passive spreading) impacts reaction during
391 active spreading. Reactive transport was simulated in flow fields generated from the four active
392 spreading protocols with nine different heterogeneous hydraulic conductivity (K) fields. Random
393 fields of $\ln K$ were generated using sequential Gaussian simulation (Deutsch and Journel 1992)
394 with a spherical variogram with correlation lengths, λ , of $\lambda = 0.125d$, $0.25d$, and $0.5d$, with mean
395 value of dimensionless K of 3.57 and $\sigma_{\ln K}$ of 0.25 , 0.5 , and 1 . Plots of the $\ln K$ fields in the
396 vicinity of the wells are shown in Figure S2 in the Supplemental Material. For all simulations
397 presented in this section, species A and B are represented as a collection of 6×10^5 and 2×10^5
398 randomly placed particles, respectively, each of which has a dimensionless mass of 6.53×10^{-7} and
399 2.73×10^{-6} , respectively. A smaller number of particles were required compared to the simulations
400 in the previous section because only the global behavior is analyzed here.

401 Figure 10 shows the difference of cumulative mass reacted in the heterogeneous aquifer (active

402 and passive spreading) and the cumulative mass reacted in the homogeneous aquifer, denoted as
403 $M_{\text{rxn-AH}}$, (active spreading only) for each protocol and each heterogeneity model. This difference
404 represents the contribution of passive spreading to reaction. For low $\sigma_{\ln K}$, passive spreading
405 contributes little extra reaction for all protocols, and the relative contribution increases as $\sigma_{\ln K}$
406 increases, with the Converging protocol having the largest contribution. At certain times, passive
407 spreading leads to a reduction in the overall amount of reaction, particularly when the combination
408 of the active spreading protocol and the heterogeneity pattern cause the plume to constrict (See
409 Figures S2-S11 in the Supplemental Material for plots of the plume evolution). The behavior
410 depends on the spatial distribution of $\ln K$, which is consistent with the findings of [de Barros et al.](#)
411 [\(2012\)](#).

412 Figure 11 shows ratio of cumulative mass reacted by each active spreading protocol in the
413 heterogeneous aquifer (active and passive spreading) relative to the cumulative mass reacted by
414 the uniform protocol in the heterogeneous aquifer, denoted as $M_{\text{rxn-UHet}}$ (passive spreading only).
415 This ratio represents the contribution of active spreading. The results show that as heterogeneity
416 increases, the contribution of active spreading to reaction also increases, demonstrating that the
417 amount of reaction is strongly controlled by the active spreading protocol, even in heterogeneous
418 aquifers. The results also show that the relative amounts of reaction across the four active spreading
419 protocols remain the same as in homogeneous aquifers, i.e., the Diverging protocol exhibits the
420 most reaction, followed, in decreasing order, by Dipole, Converging and Stagnation protocols. One
421 exception is that as heterogeneity increases, the Converging protocol produces more reaction than
422 the other protocols at later times. As heterogeneity increases, the reaction front becomes more
423 irregular, resulting in portions of the reaction front that are more perpendicular to the flow direction
424 near the extraction well where velocities are high. For example, the third and fourth panels of
425 Figure S8c in the Supplemental Material show that the reaction front near the leading edge of the
426 plume of species B (white shaded region between the blue and red shading) crosses the streamlines
427 (i.e., is perpendicular to the local velocity), while for the homogeneous aquifer (Fig. 7a) the reaction
428 front remains aligned with the streamlines. Thus, the combination of high dispersivity ($\alpha_B \approx \alpha_L$,

429 because portions of the reaction front are perpendicular to flow), high velocity (i.e., high D_β), and
430 positive strain enhances reaction. This enhancement also occurs in the Dipole protocol, but to a
431 lesser extent.

432 **DISCUSSION**

433 Spreading, mixing, and reaction in porous media are manifestations of spatial or spatio-temporal
434 variations in the velocity field. Spreading, by definition, is the reconfiguration of a plume due to
435 velocity variation; while mixing and reaction tend to be enhanced as a result of spreading. A
436 high degree of spreading has been accomplished under conditions of chaotic advection (Lester
437 et al. 2013; Aref et al. 2017; Speetjens et al. 2021), but spreading can also be accomplished to a
438 lesser degree through natural heterogeneity of the porous medium (passive spreading) and imposed
439 pumping schemes (active spreading).

440 A number of recent studies point to the intriguing result that stretching may be sufficient to
441 predict reactive transport. Specifically, de Barros et al. (2012) used the Okubo-Weiss parameter
442 to predict reaction based on stretching; Engdahl et al. (2014) found a similar result that stretching
443 predicts reaction. Both studies argue that reaction depends on flow, which is supported by the results
444 of the present study. However, both studies assumed a constant isotropic dispersion coefficient and
445 therefore did not account for the impact of the orientation of the reaction front within the flow
446 field nor on the spatial variation of velocity. In the present study, we used a dispersion coefficient
447 that varies with flow direction and magnitude of velocity, which is the observed behavior of solute
448 transport at the plume scale. We quantified stretching as the length of the reaction front, and we
449 showed (Fig. 4) that stretching alone is not sufficient to enhance reaction – the stagnation protocol
450 exhibits double the amount of stretching of any other method, yet it produces the least amount of
451 reaction. In fact, the results show that spatial variations in velocity (i.e., the required ingredient
452 for spreading) does not necessarily enhance reaction. Relative to uniform flow, two of the four
453 protocols we investigated (Converging and Stagnation) produce less reaction than uniform flow
454 (Fig. 4d), so in those cases, active spreading inhibited reaction. The key conclusion here is that
455 global measures are not sufficient to predict the overall amount of contaminant degradation.

456 Our analysis of local relationships between velocity field and plume geometry allowed us
457 to identify that instantaneous reactions in porous media proceed more quickly where the plume
458 interface, and therefore the reaction front, is perpendicular to local velocity, owing to the notion
459 that longitudinal dispersivity is greater than transverse dispersivity. These results are consistent
460 with and expand upon earlier works that considered simpler plume geometries.

461 In the context of flow and transport in porous media, those studying lamellae (Meunier and
462 Villiermaux 2010; Le Borgne et al. 2013; de Anna et al. 2014) have articulated the interplay between
463 stretching forming longer lamellae versus diffusion coalescing lamellae together. A similar interplay
464 has been explored by researchers studying spreading and mixing using the scalar dissipation rate
465 that, under certain assumptions described by de Dreuzy et al. (2012), is $(1/2)dM/dt$, where M is
466 the integral over space of the concentration squared (Le Borgne et al. 2010; de Dreuzy et al. 2012;
467 Engdahl et al. 2013). As a group, these papers explain the interplay between spreading that sharpens
468 gradients and elongates interfaces versus mixing that softens gradients and blurs interfaces. While
469 these earlier works often considered simpler plume geometries, the results of the present study are
470 consistent with their findings. For example, studies that considered one species invading another
471 species (Le Borgne et al. 2013; Perez et al. 2020) identified the formation of lamella within a pore
472 that sharpened concentration gradients and promoted mixing and reaction. At the plume scale,
473 which is the focus of the present study, the plumes form lamella-like features as they travel through
474 heterogeneous material (See Figures S3-S11 in the Supplemental Material), which locally sharpens
475 concentration gradients. Consistent with studies of lamella formation at the pore scale, we found
476 that increasing heterogeneity leads to high amounts of reaction (see Figs. 10 and 11).

477 As another example, our results are consistent with results of studies that considered a plume
478 emanating from a continuous source, in which the plume boundary is aligned with the flow direction
479 and mixing across the boundary is controlled by transverse dispersion. Cirpka et al. (2011) found
480 that small-scale heterogeneity enhanced mixing and reaction of such a plume. This result is
481 consistent with our finding that heterogeneity increases the amount of reaction in the Converging
482 protocol relative to the Uniform protocol. Without heterogeneity (Fig. 7), the reaction front is

483 aligned with the flow direction throughout most of its extent, especially where velocity is high (near
484 the extraction well), but with heterogeneity, portions of the reaction front are perpendicular to the
485 local flow direction, and longitudinal dispersion dominates, enhancing reaction. In another study
486 with a plume emanating from a continuous source, [Rolle et al. \(2009\)](#) found that temporal variation
487 in uniform flow had essentially no effect on reaction. This behavior is consistent with the results of
488 our study, i.e., the plume boundary remained aligned with the flow direction, even as the velocity
489 changes, but reaction proceeds most rapidly where the reaction front is perpendicular to the local
490 velocity, so the temporal change in uniform flow had little effect.

491 To further demonstrate that a reaction front aligned perpendicular to the local flow field enhances
492 reaction, we repeated the Dipole protocol; however, after $t=9.48$, we rotated the dipole by 90° so that
493 injection occurs at $(0, +4.44)$ and extraction occurs at $(0, -4.44)$. Figure 12 shows that cumulative
494 mass reacted increases when the dipole orientation is changed, leading to more cumulative mass
495 reacted for the Two-Step Dipole protocol as compared with the standard one-step Dipole protocol.
496 The cause of this behavior is apparent through comparison of Figures 8 and 13, which show the
497 plume evolution and local measures along the reaction front at various times for the one-step Dipole
498 and Two-Step Dipole protocols, respectively. In both figures, the left panels are showing results
499 for $t = 4.74$, and both figures are identical because both protocols have the same flow rate. The
500 second time that is plotted for the Two-Step Dipole protocol (Fig. 13) is just after the change in
501 the flow field, and very close in time to the second time plotted in Figure 8. At this time, the
502 highest concentration gradients along reaction front occur near $\ell = 0.25L_\Gamma$. In the one-step Dipole
503 protocol, D_β is low at that point because the reaction front is aligned with the local flow direction.
504 On the other hand, in the Two-Step Dipole protocol, the flow field is rotated 90° , so the reaction
505 front is perpendicular to the flow direction at that point, leading to high D_β , high $J_{A\beta}$, and therefore
506 a higher reaction rate. Over time, the concentration gradients are smoothed as the reaction front
507 propagates toward the operating extraction well (at $(x, y) = (0, -4.44)$), leading to lowering of $J_{A\beta}$
508 and of the reaction rate.

509 At the pore scale, soothing of concentration gradients is driven by molecular diffusion, thus,

510 the Peclet number, which reflects the relative contributions of advection and molecular diffusion,
511 is a fundamental parameter. At the plume scale, smoothing of the concentration gradients occurs
512 through molecular diffusion, but at this larger scale, it is also driven by velocity variations that
513 bring together water parcels that take different paths around solid grains and around small-scale
514 low permeability features. In groundwater remediation applications, the Peclet number is typically
515 large, particularly near the active spreading wells, so much of the reaction is driven by advection.
516 A sensitivity analysis was conducted on the molecular diffusion coefficient in our simulations and
517 showed that the amount of reaction is not sensitive to molecular diffusion (See Figure S1 of the
518 Supplemental Material).

519 An important caveat is that the simulations presented here, like many others in the reactive
520 transport literature, assume instantaneous bimolecular reaction of $A + B \rightarrow C$, where all species
521 are aqueous and non-sorbing. Not all of these assumptions will hold in practical situations, for
522 example, it is known that the pumping schemes that optimize degradation of sorbing contaminants
523 are quite different from those that optimize degradation of non-sorbing contaminants (Neupauer
524 and Mays 2015). Future work will be required to generalize the results of this study beyond the
525 simplified chemistry assumed here. The goal of this study is to demonstrate that it may be possible
526 to improve practical applications of reactive transport in porous media by taking into account the
527 geometry of plume spreading.

528 CONCLUSION

529 Several recent studies have shown that velocity variations within a pore (Meunier and Villermaux
530 2010; Le Borgne et al. 2013) or in a porous medium at larger scales (de Barros et al. 2012; Engdahl
531 et al. 2014) lead to elongation of fluid interfaces and sharpening of concentration gradients at
532 the interface, both of which can enhance mixing and therefore reaction. In this work, we found
533 elongation of the interface and sharpening of the gradients was not sufficient to enhance reaction.
534 Furthermore, some patterns of spatially varying velocity actually inhibit reaction. In other words, a
535 global enhancement of reaction is not necessarily the result of global stretching of a plume interface,
536 but rather is a result of local characteristics of flow and plume geometry along the reaction front.

537 Thus, the objective of the present study was to identify the local mechanisms that lead to a global
538 enhancement of reaction.

539 In this study, we evaluated spreading, mixing, and reaction of two contiguous solute plumes in
540 porous media under uniform flow conditions and under four different active spreading protocols:
541 Diverging, Converging, Dipole, and Stagnation (Figure 3). We considered irreversible, instanta-
542 neous bimolecular reaction, in which reaction takes place along a narrow moving reaction front
543 between the two plumes. We demonstrated that the amount of reacted mass is equivalent to the
544 cumulative mass of the species that disperses across the reaction front.

545 Reaction occurs because the two reactant species come together by dispersion across the reaction
546 front. By definition, the local dispersive mass flux across the reaction front depends on the
547 local concentration gradient and the local dispersion coefficient, both defined in the direction
548 perpendicular to the reaction front. To discern the contributions of each of these to reaction, we
549 evaluated the temporal evolution of characteristics of the local flow field and plume along the
550 moving reaction front. We found that the spatial variability in the flow field, and the position of
551 the reaction front within it, controls the amount of reaction according to the orientation of the
552 reaction front relative to the local velocity, and whether flow is diverging or converging. The
553 highest amount of reaction occurs where flow is diverging and the reaction front is perpendicular to
554 the local velocity. In these locations, dispersion is dominated by longitudinal dispersivity, which is
555 greater than transverse dispersivity. In addition, strain is positive, which sharpens the concentration
556 gradient; thus although dispersion smooths the concentration gradient, the positive strain partially
557 counteracts the effects of smoothing. The least amount of reaction occurs where the reaction front
558 is aligned parallel to the local velocity, especially where flow is not converging. In these locations,
559 transverse dispersion dominates, and negative strain rapidly reduces the concentration gradient.

560 The active spreading protocols in this study are representative of components of in situ ground-
561 water remediation systems in which an amendment is emplaced in the contaminant plume to react
562 with and degrade the contaminant. These protocols can be used individually or superimposed in
563 space and time, to create more robust flow fields that can be designed to address the specific plume

564 geometry, aquifer characteristics, and remediation goals at a particular site. The combination of the
565 protocols in space and time can lead to faster and more complete degradation of the contaminant,
566 as shown with the Two-Step Dipole protocol which is a superposition in space and time of the
567 Diverging and Converging protocols.

568 We note that the flow fields, aquifer heterogeneity models, and reaction chemistry model consid-
569 ered here are not exhaustive, and additional complexities may be encountered in practice that have
570 not been addressed here. For example, contaminants may be trapped in low permeability materials
571 which cannot be accessed by the active spreading protocols described here; or contaminants may
572 sorb and desorb kinetically onto the porous material, so the reaction front may not be a discrete
573 linear feature as it was in this study. Nevertheless, the main results of this study are still applicable,
574 i.e., that the global amount of reaction is controlled by the local mechanisms that bring species
575 together to react, which, in our scenario, are the concentration gradient and dispersion coefficient
576 in the direction locally transverse to the reaction front. Additional work is needed to identify active
577 spreading protocols that can address these more challenging problems.

578 **DATA AVAILABILITY STATEMENT**

579 All data, models, and code generated or used during the study appear or are cited herein.

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TABLE 1. Parameter values in flow and reactive transport simulations. All values are dimensionless.

Parameter	Value
Hydraulic conductivity, K	3.57
Aquifer thickness, b	1.78
Porosity, n	0.25
Length of side of square aquifer, L	53.4
Finite difference grid discretization	0.0222
Head at $x = -L/2$, h_L	
Active spreading protocols	0
Uniform flow protocol	0.534
Coordinates of well 1, \mathbf{x}_{w1}	(-4.44, 0)
Coordinates of well 2, \mathbf{x}_{w2}	(0, 0)
Coordinates of well 3, \mathbf{x}_{w3}	(4.44, 0)
Longitudinal dispersivity, α_L	0.0178
Transverse dispersivity, α_T	0.00178
Molecular diffusion coefficient, D_m	0

TABLE 2. Dimensionless injection (positive) and extraction (negative) rates for the protocols.

Protocol Name	Injection/Extraction Pattern		
	Well 1	Well 2	Well 3
Diverging	1.0	0	0
Converging	0	0	-1.0
Dipole	0.5	0	-0.5
Stagnation	0.4	0.1	-0.5
Uniform	0	0	0

TABLE 3. Summary of the relationship between the reaction front and flow conditions on stretching and mixing parameters.

Character of flow	Orientation of Reaction Front	
	\perp to flow	\parallel to flow
diverging	$\zeta > 0$	$\zeta < 0$
	$\alpha_\beta \approx \alpha_L$	$\alpha_\beta \approx \alpha_T$
	$\frac{\partial C_A}{\partial \beta}$ decreases moderately	$\frac{\partial C_A}{\partial \beta}$ decreases moderately
converging	$\zeta < 0$	$\zeta > 0$
	$\alpha_\beta \approx \alpha_L$	$\alpha_\beta \approx \alpha_T$
	$\frac{\partial C_A}{\partial \beta}$ decreases rapidly	$\frac{\partial C_A}{\partial \beta}$ decreases slowly

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709 (b) Slice through $y = 0$. (c) Enlargement showing vectors representing the direction

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711 locally tangential to the reaction front ξ , and the direction transverse to local flow

712 η from an arbitrary point (yellow dot), assuming uniform flow in the x direction.

713 The green circle in subplot a represents the reaction front, which initially is circular

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716 well locations. In subplot b, the horizontal dashed lines represent the maximum

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724 that flows are simulated over the domain $-26.7 \leq x \leq 26.7$ and $-26.7 \leq y \leq 26.7$,

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753		black circles represent $\ell/L_\Gamma = 0, 0.25, \text{ and } 0.5$, respectively. L_Γ is the reaction	
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771		black circles represent $\ell/L_\Gamma = 0, 0.25, \text{ and } 0.5$, respectively. L_Γ is the reaction	
772		front length. Gray shading indicates negative strain. Note that the limits on the	
773		plume plot and on the plot of dispersive mass flux are different in this figure than	
774		in Figure 5–9.	48

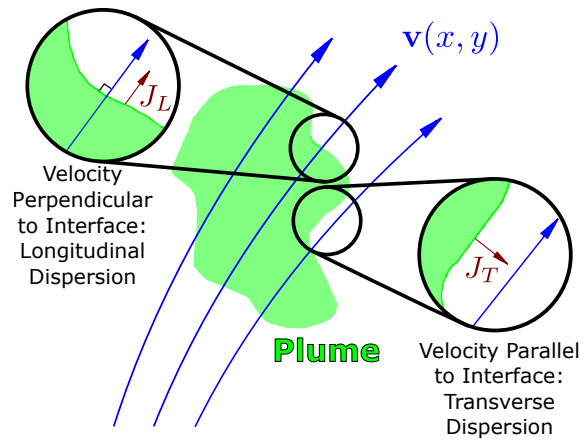


Fig. 1. Schematic of plume and location velocity field. Left and right callouts show positions where, respectively, purely longitudinal dispersive mass flux, J_L , and purely transverse dispersive mass flux, J_T , occur.

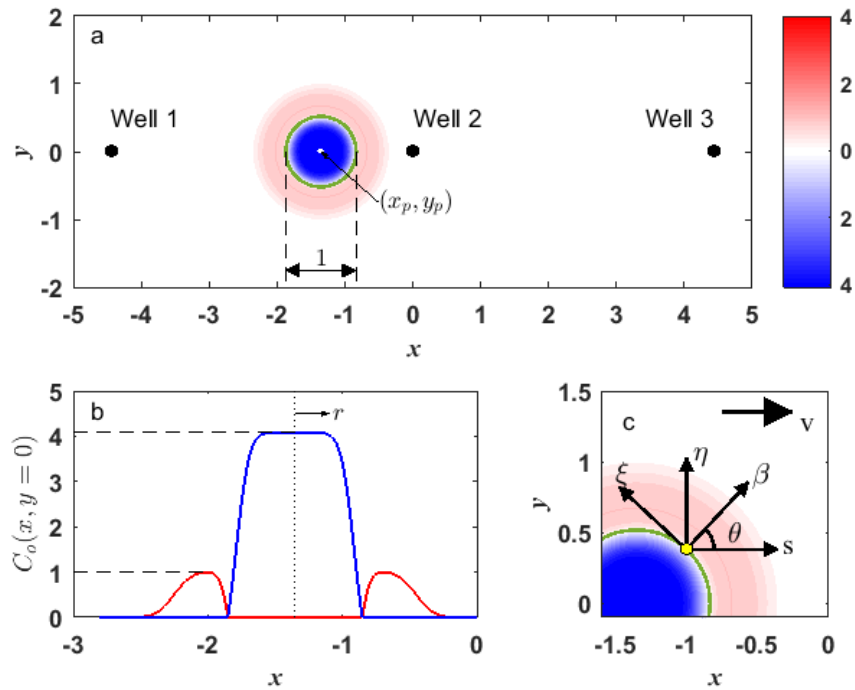


Fig. 2. Initial concentration distributions of species A (red) and B (blue). (a) Plan view. (b) Slice through $y = 0$. (c) Enlargement showing vectors representing the direction locally perpendicular to the reaction front β , the local flow direction s , the direction locally tangential to the reaction front ξ , and the direction transverse to local flow η from an arbitrary point (yellow dot), assuming uniform flow in the x direction. The green circle in subplot a represents the reaction front, which initially is circular with a unit dimensionless diameter. Dimensionless lengths are normalized by this diameter. The plumes are centered at (x_p, y_p) . Small black circles represent the well locations. In subplot b, the horizontal dashed lines represent the maximum dimensionless concentrations of species A and species B at $t = 0$, both relative to the initial maximum concentration of species A, and the vertical dashed line represents $r = 0$, where r is the radial distance from the center of Plume B.

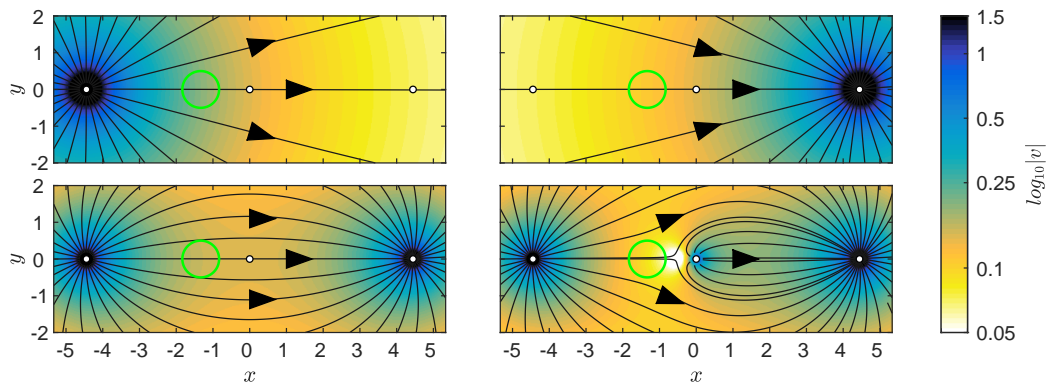


Fig. 3. Velocity field for (a) Diverging, (b) Converging, (c) Dipole and (d) Stagnation active spreading protocols. Color shading represents the magnitude of velocity. Thin black lines are streamlines; arrows show the direction of flow. White circles are locations of the wells. The green circles represent the reaction front at $t = 0$. Note that flows are simulated over the domain $-26.7 \leq x \leq 26.7$ and $-26.7 \leq y \leq 26.7$, but only the portion of the domain in the vicinity of the wells is shown.

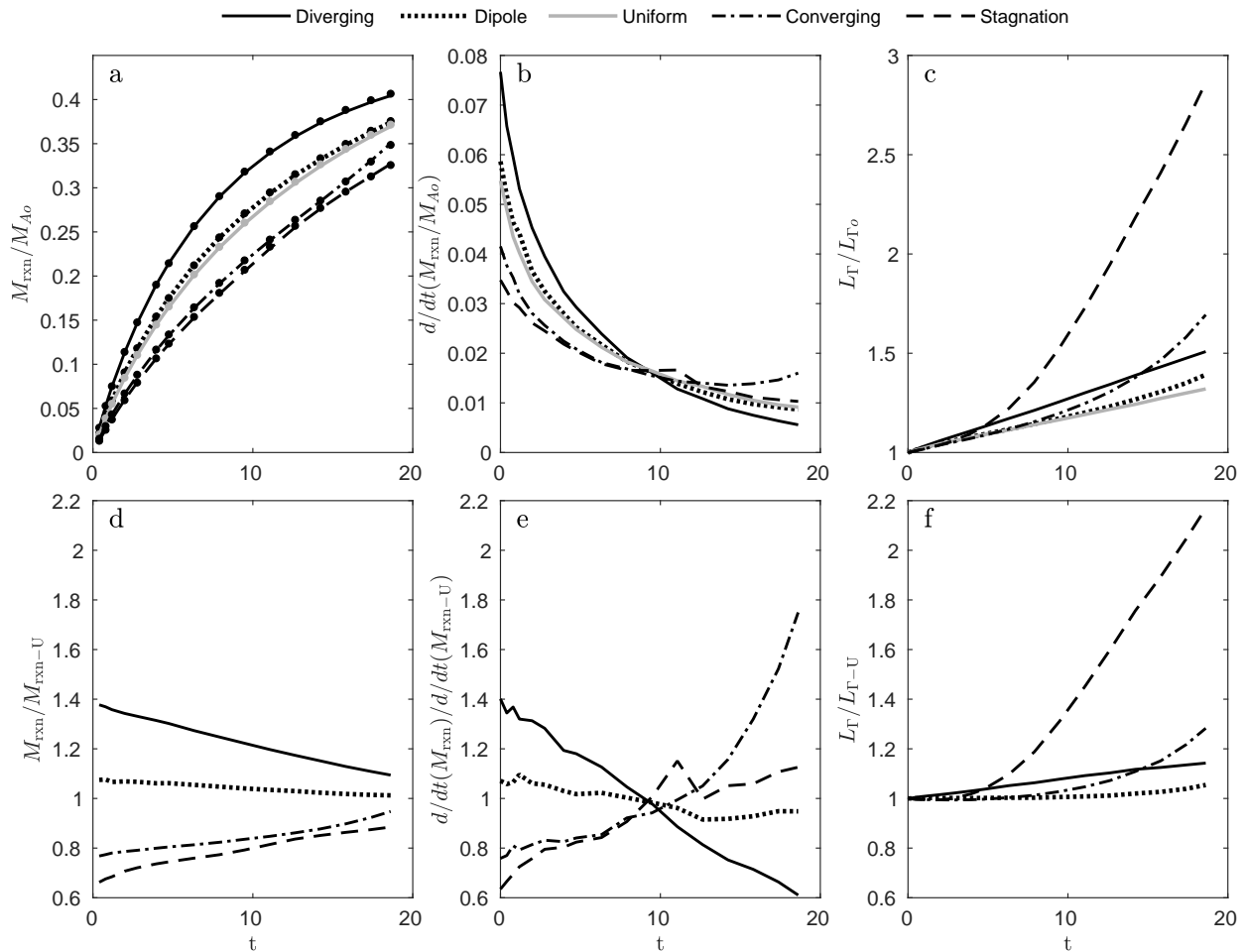


Fig. 4. Temporal evolution of measures of global spreading, mixing, and reaction. (a) Temporal evolution of the cumulative mass reacted of species A, M_{rxn} , calculated using (10) (lines) and (18) (points) and normalized by M_{A_0} . (b) Temporal evolution of the dispersive mass flow rate of species A across the reaction front, $d/dt(M_{\text{rxn}}/M_{A_0})$. (c) Temporal evolution of the length of the reaction front, L_{Γ} , normalized by the initial length, L_{Γ_0} . (d) M_{rxn} normalized by $M_{\text{rxn-U}}$. (e) $d/dt(M_{\text{rxn}})$ normalized by $d/dt(M_{\text{rxn-U}})$. (f) L_{Γ} normalized by $L_{\Gamma-U}$. The subscript U refers to the Uniform protocol.

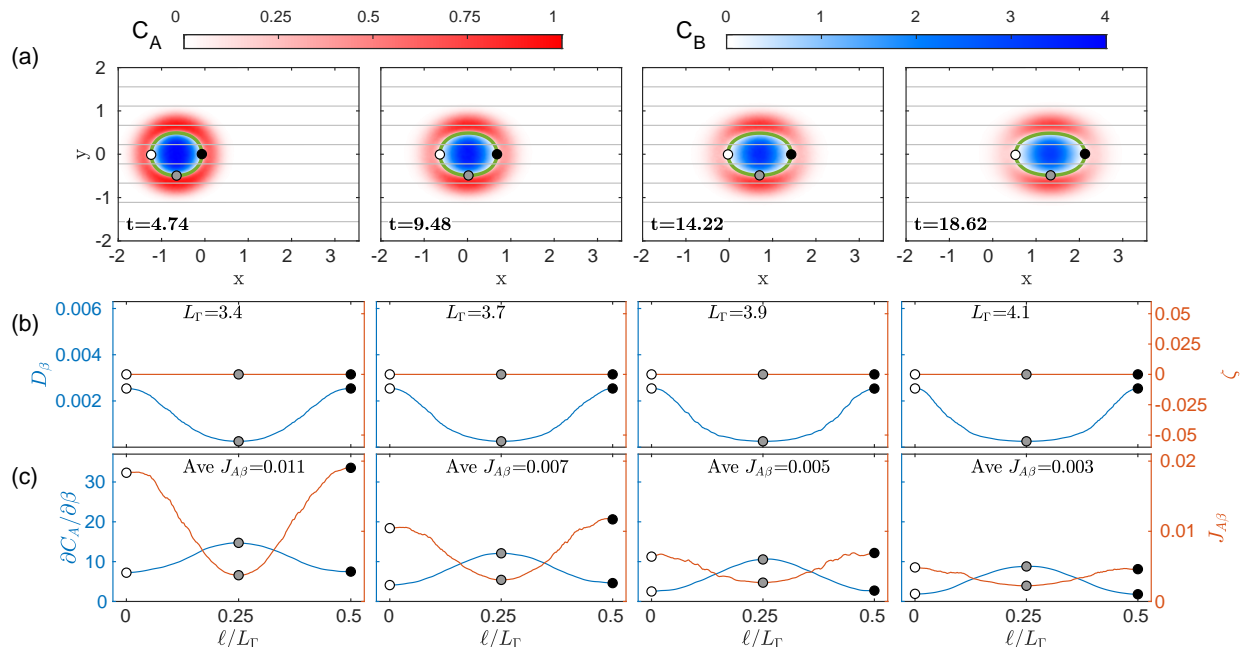


Fig. 5. Local analysis of results of Uniform protocol at four times. (a) Plumes of Species A (red) and B (blue) and reaction front (green line). Gray lines show streamlines. D_β (b, left axis), strain (b, right axis), $\partial C_A/\partial \beta$ (c, left axis) and dispersive mass flux, $J_{A\beta}$ (c, right axis) along the reaction front. White, gray, and black circles represent $\ell/L_\Gamma = 0, 0.25,$ and $0.5,$ respectively. L_Γ is the reaction front length.

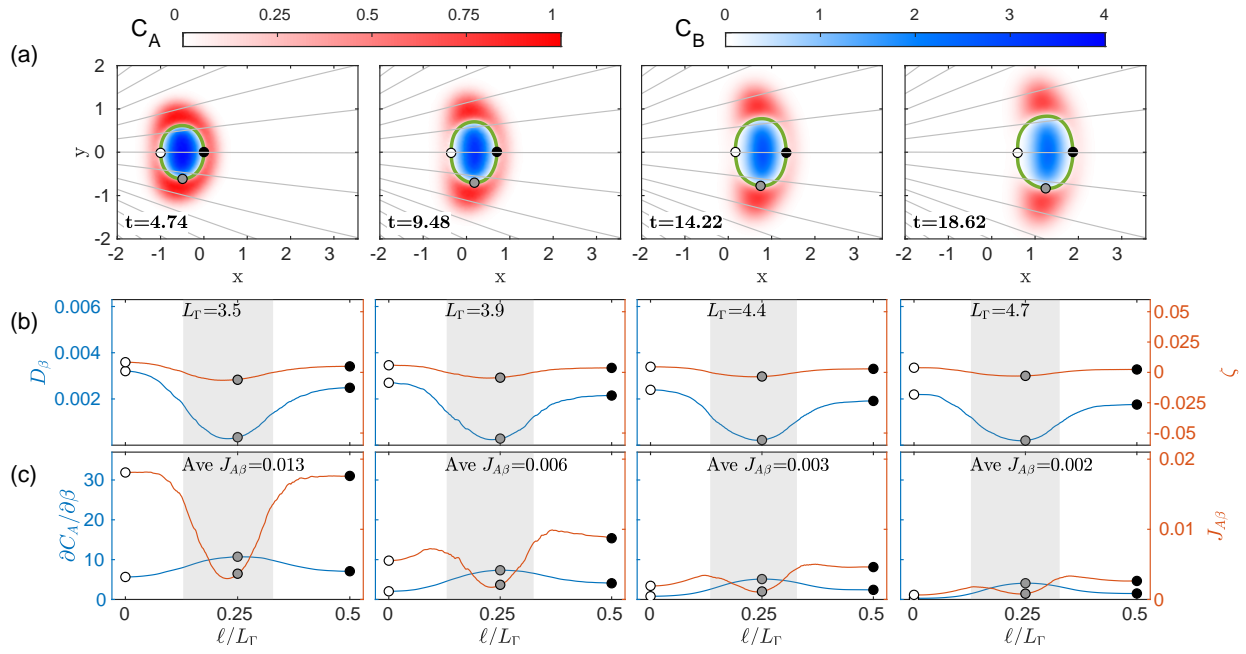


Fig. 6. Local analysis of the Diverging protocol at four different times. (a) Plumes of Species A (red) and B (blue) and reaction front (green line). Gray lines show streamlines. D_β (b, left axis), strain (b, right axis), $\partial C_A/\partial \beta$ (c, left axis) and dispersive mass flux, $J_{A\beta}$ (c, right axis) along the reaction front. White, gray, and black circles represent $\ell/L_\Gamma = 0, 0.25,$ and 0.5 , respectively. L_Γ is the reaction front length. Gray shading indicates negative strain.

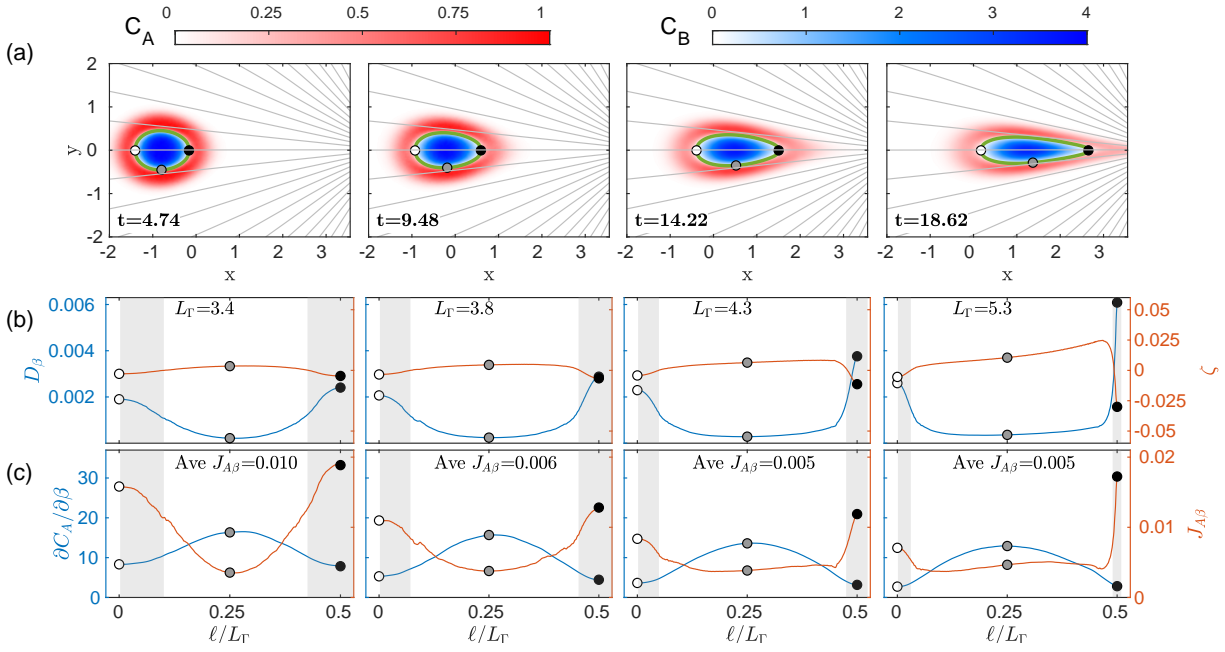


Fig. 7. Local analysis of results of Converging protocol at four times. (a) Plumes of Species A (red) and B (blue) and reaction front (green line). Gray lines show streamlines. D_β (b, left axis), strain (b, right axis), $\partial C_A / \partial \beta$ (c, left axis) and dispersive mass flux, $J_{A\beta}$ (c, right axis) along the reaction front. White, gray, and black circles represent $\ell/L_\Gamma = 0, 0.25$, and 0.5 , respectively. L_Γ is the reaction front length. Gray shading indicates negative strain.

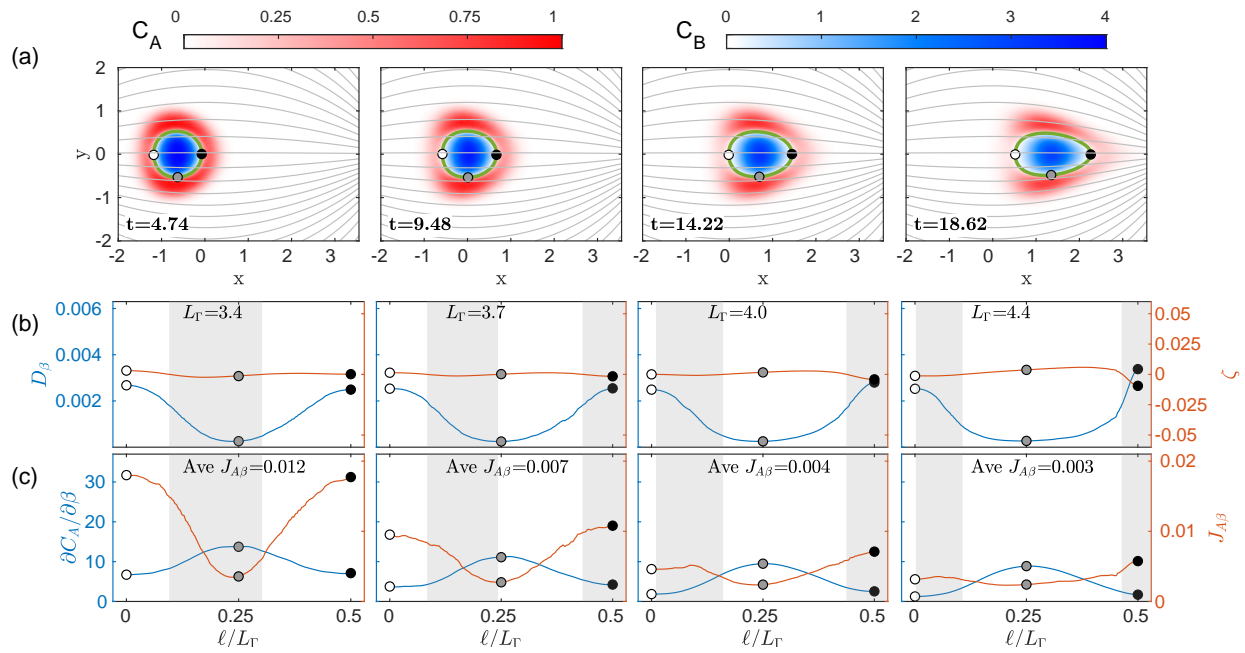


Fig. 8. Local analysis of the Dipole protocol at four different times. (a) Plumes of Species A (red) and B (blue) and reaction front (green line). Gray lines show streamlines. D_β (b, left axis), strain (b, right axis), $\partial C_A / \partial \beta$ (c, left axis) and dispersive mass flux, $J_{A\beta}$ (c, right axis) along the reaction front. White, gray, and black circles represent $\ell/L_\Gamma = 0, 0.25,$ and $0.5,$ respectively. L_Γ is the reaction front length. Gray shading indicates negative strain.

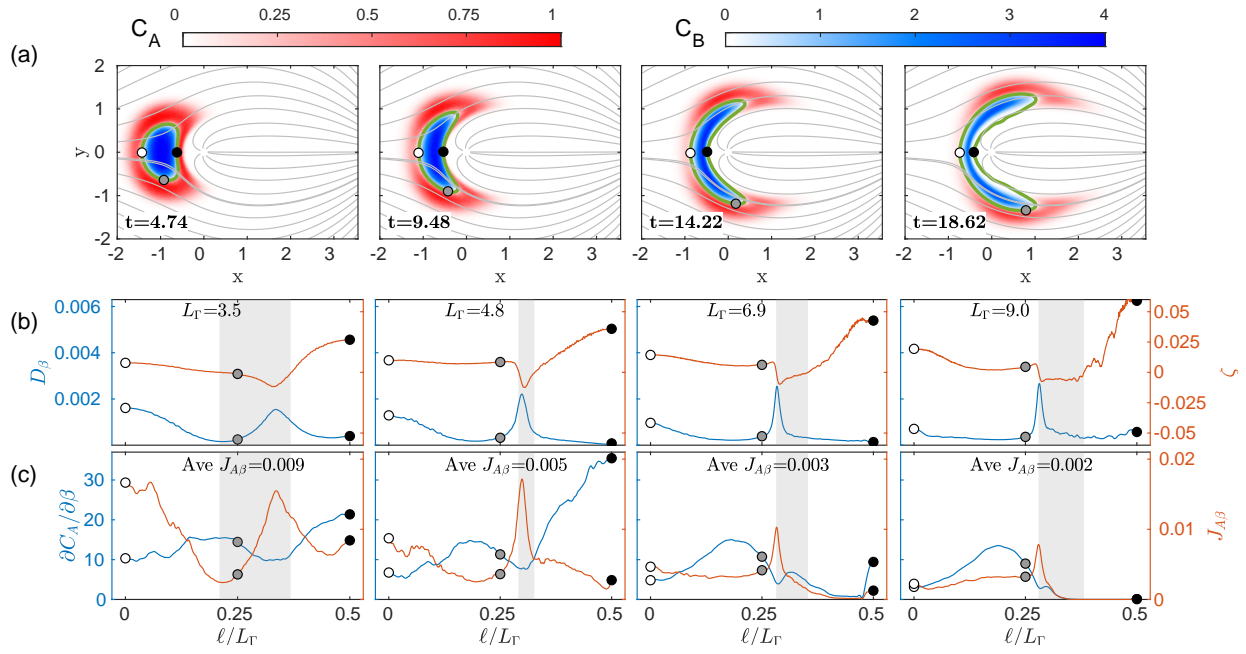


Fig. 9. Local analysis of the Stagnation protocol at four different times. (a) Plumes of Species A (red) and B (blue) and reaction front (green line). Gray lines show streamlines. D_β (b, left axis), strain (b, right axis), $\partial C_A/\partial \beta$ (c, left axis) and dispersive mass flux, $J_{A\beta}$ (c, right axis) along the reaction front. White, gray, and black circles represent $\ell/L_\Gamma = 0, 0.25,$ and 0.5 , respectively. L_Γ is the reaction front length. Gray shading indicates negative strain.

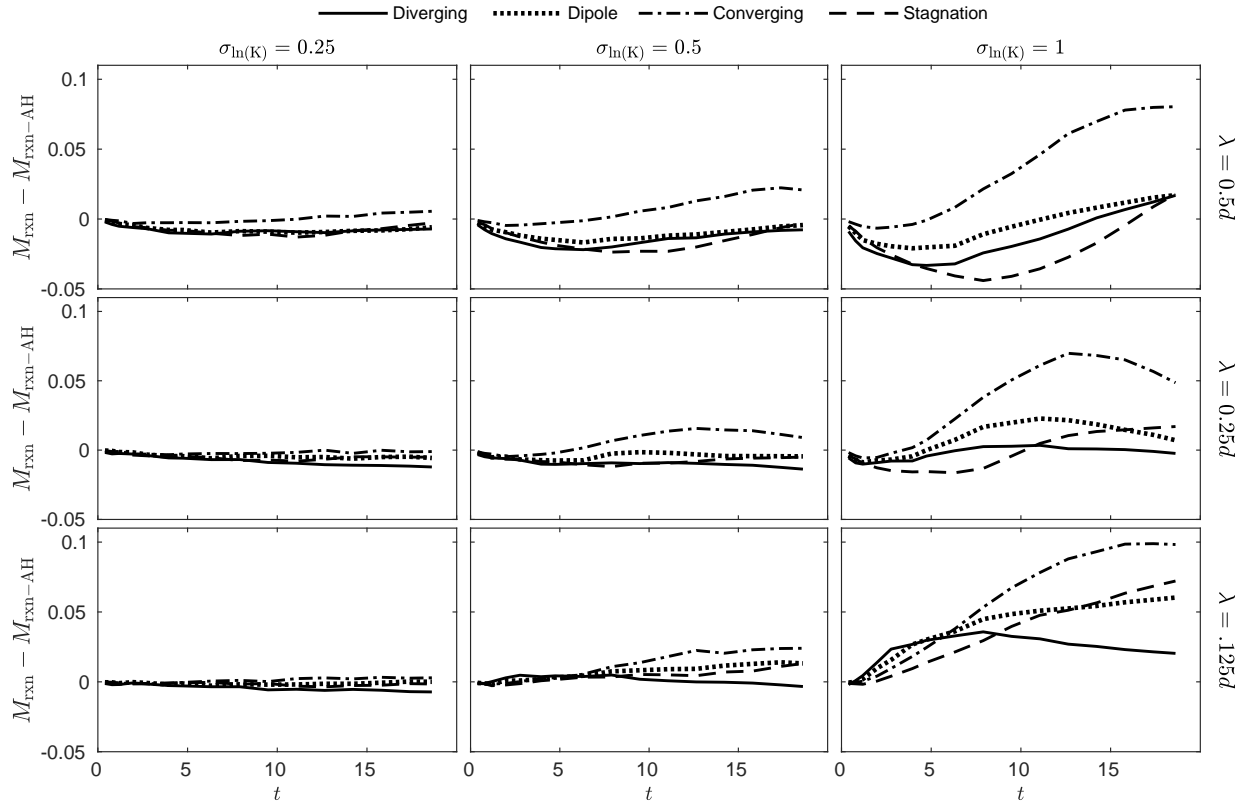


Fig. 10. $M_{\text{rxn}} - M_{\text{rxn-AH}}$ for the four active spreading protocols in nine different models of heterogeneous aquifers, where $M_{\text{rxn-AH}}$ is M_{rxn} for the respective protocol in the homogeneous aquifer. Heterogeneity increases from left to right across the rows, with increasing $\sigma_{\ln K}$ (labeled above each column), and from top to bottom along the columns, with decreasing correlation length λ (labeled to the right of each row).

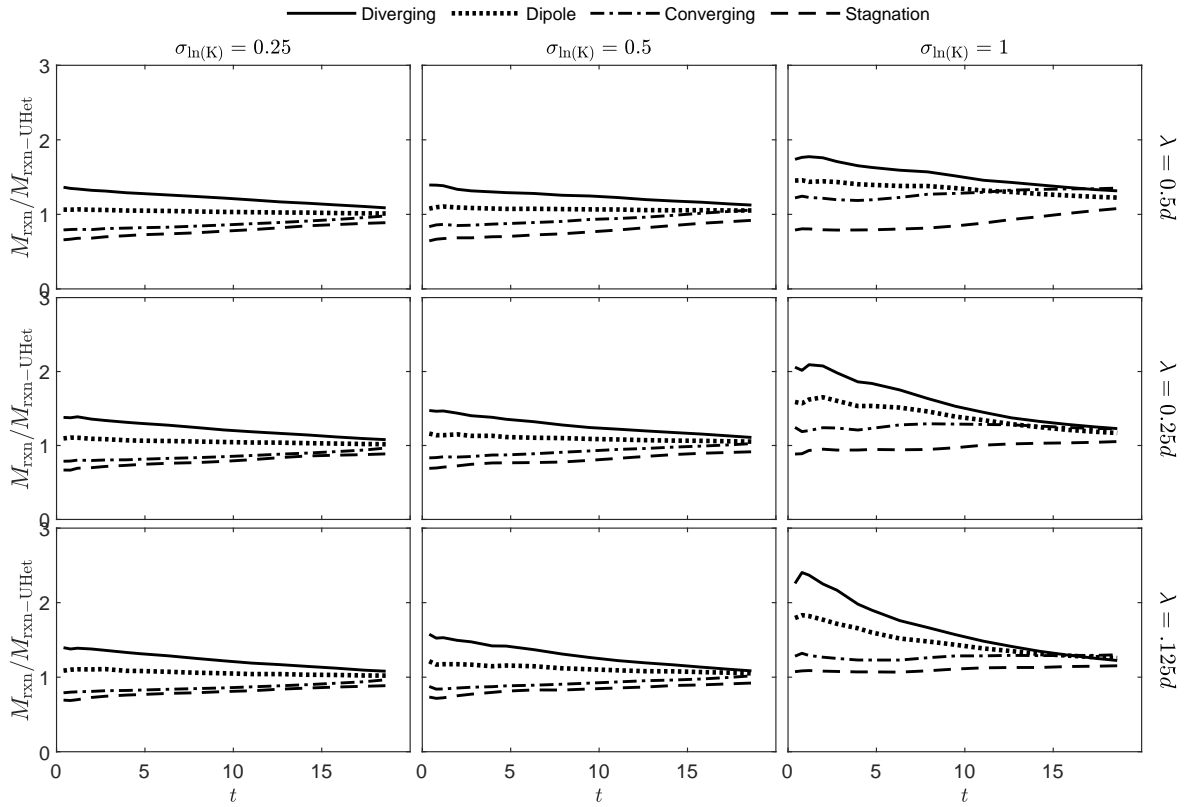


Fig. 11. $M_{\text{rxn}}/M_{\text{rxn-UHet}}$ for the four active spreading protocols in nine different models of heterogeneous aquifers, where $M_{\text{rxn-UHet}}$ is M_{rxn} for the Uniform protocol in each heterogeneous aquifer. Heterogeneity increases from left to right across the rows, with increasing $\sigma \ln K$ (labeled above each column), and from top to bottom along the columns, with decreasing correlation length λ (labeled to the right of each row).

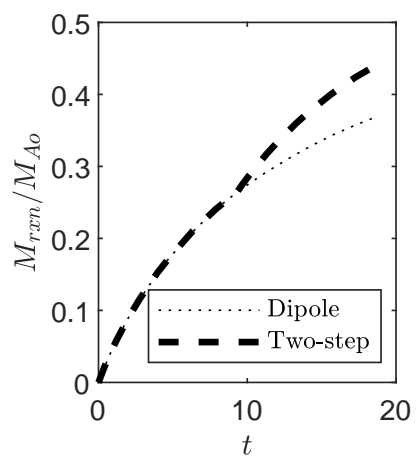


Fig. 12. Temporal evolution of cumulative mass reacted of species A, M_{rxn} , normalized by M_{Ao} for the Dipole and Two-Step Dipole protocols.

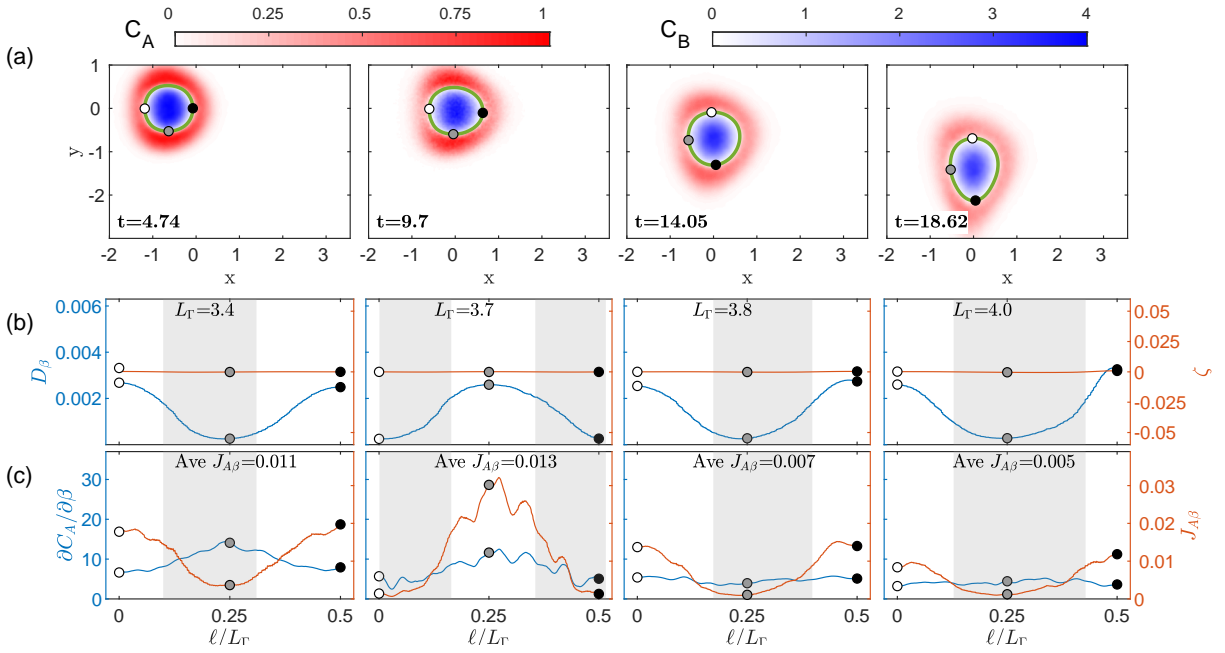


Fig. 13. Local analysis of the Two-Step Dipole protocol at four different times. (a) Plumes of Species A (red) and B (blue) and reaction front (green line). Gray lines show streamlines. D_β (b, left axis), strain (b, right axis), $\partial C_A / \partial \beta$ (c, left axis) and dispersive mass flux, $J_{A\beta}$ (c, right axis) along the reaction front. White, gray, and black circles represent $\ell/L_\Gamma = 0, 0.25,$ and $0.5,$ respectively. L_Γ is the reaction front length. Gray shading indicates negative strain. Note that the limits on the plume plot and on the plot of dispersive mass flux are different in this figure than in Figure 5–9.