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# Benchmarks for multicomponent diffusion and electrochemical migration

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#### 1 Abstract

2 In multicomponent electrolyte solutions, the tendency of ions to diffuse at different rates results 3 in a charge imbalance that is counteracted by the electrostatic coupling between charged species 4 leading to a process called "electrochemical migration" or "electromigration". Although not 5 commonly considered in solute transport problems, electromigration can strongly affect mass 6 transport processes. The number of reactive transport models that consider electromigration has 7 been growing in recent years, but a direct model inter-comparison that specifically focuses on the 8 role of electromigration has not been published to date. This contribution provides a set of three 9 benchmark problems that demonstrates the effect of electric coupling during multicomponent 10 diffusion and electrochemical migration and at the same time facilitates the inter-comparison of 11 solutions from existing reactive transport codes. The first benchmark (Lichtner, 1995) focuses on 12 the 1D-transient diffusion of  $HNO_3$  (pH = 4) in a NaCl solution into a fixed concentration 13 reservoir, also containing NaCl - but with lower HNO<sub>3</sub> concentrations (pH = 6). The second benchmark describes the 1D steady-state migration of the sodium isotope <sup>22</sup>Na triggered by 14 15 sodium chloride diffusion in neutral pH water. The third benchmark (Rolle et al., 2013) presents 16 a flow-through problem in which transverse dispersion is significantly affected by 17 electromigration. The system is described by 1D transient and 2D steady-state models. Very 18 good agreement on all of the benchmarks was obtained with the three reactive transport codes used: CrunchFlow, MIN3P and PHREEQC. 19

20

Keywords: Reactive transport modeling, multicomponent diffusion, electromigration, model
 intercomparison, benchmark

#### 23 **1. Introduction**

24 It is well known that diffusive transport in multicomponent electrolyte systems cannot be fully 25 described by Fickian diffusion alone, but is affected by a variety of processes including the electrostatic interactions between individual ions (Vinograd and McBain, 1941; Newman, 1973; 26 Ben-Yaakov, 1981; Cussler, 1997). Each dissolved species is subject to its own species-27 28 dependent diffusion coefficient, affected by parameters such as charge and size of the ion (Cussler, 1997) and ionic conductivity (Lasaga, 1979). As a result, dissolved species will tend to 29 30 diffuse at different rates, promoting the development of a charge imbalance in solution. 31 However, positively and negatively charged species are also affected by electric coupling, which 32 ensures that charge balance in solution is maintained. Generally speaking, "large" cations and 33 "small" anions are tied together electrostatically (Newman, 1973; Cussler, 1997) to enforce 34 electroneutrality at the macroscale - an essential condition in electrolyte solutions (Lichtner, 35 1996; Van Cappellen and Gaillard, 1996). This electric coupling leads to an additional mass 36 transport process called "electrochemical migration" or "electromigration" (Newman, 1991; 37 Ben-Yaakov, 1981). Fick's law neglects these interactions, describes ion migration solely based 38 on concentration gradients, and consequently does not consider the electric field generated by 39 electrostatic bonding (coulombic interactions) of charged species (Lasaga, 1979; McDuff and 40 Ellis, 1979; Newman, 1991; Lichtner, 1996; Van Cappellen and Gaillard, 1996). In a 41 multicomponent system that includes charged species, diffusive ion migration is therefore better 42 described by the Nernst-Plank equation, a formulation that explicitly considers the electric 43 coupling between species and ensures the conservation of charge (Lasaga, 1979; McDuff and 44 Ellis, 1979; Newman, 1991; Lichtner, 1996; Van Cappellen and Gaillard, 1996; Boudreau et al., 45 2004; Liu et al., 2011; Steefel et al., 2014).

46 In some cases, electrostatic interactions between diffusing species can have a strong 47 effect on ion mobility and can produce unexpected behavior such as uphill diffusion (e.g.: 48 Oelkers, 1996). In addition, apparent diffusion coefficients (i.e. diffusion coefficients derived 49 from Fick's law) may show a strong dependency on concentrations. Considering that the 50 quantification of diffusion coefficients is labor-intensive (Tyrell, 1961; Cussler, 1997), it is 51 impractical to determine apparent diffusion coefficients as a function of solution composition for 52 a range of conditions. Instead, it is advantageous to consider electrochemical interactions 53 affecting diffusion explicitly rather than lumping this effect into empirically measured apparent 54 diffusion coefficients.

55 Reactive transport models are commonly used for the quantitative investigation of flow, transport and reaction processes in porous media. These models aid with the verification of 56 conceptual models, are used to design and evaluate experiments, and assist with the 57 58 interpretation of field data in the fields of geology, engineering and environmental research 59 (Boudreau, 1997; Kang et al., 2006; Steefel et al., 2003; Wang and Van Cappellen, 1996; 60 MacQuarrie and Mayer 2005). Traditionally, diffusion has been implemented into reactive transport models based on Fick's law and diffusion coefficients are often treated as adjustable 61 62 parameters (Cussler, 1997). However, the number of reactive transport models that include 63 electromigration and consider the chemical potential gradient as the driving force of diffusion 64 has been growing in recent years (Parkhurst and Appelo, 1999; Giambalvo et al., 2002; Shiba et 65 al., 2005; Johannesson et al 2007; Paz-Garcia et al., 2011; Muniruzzaman et al., 2014). Although some aspects of electromigration on solute transport have been investigated (Oelkers, 1996; 66 67 Giambalvo et al., 2002; Steefel and Maher, 2009), a direct model inter-comparison that

specifically focuses on the role of electromigration and electrostatic effects on ion transport hasnot been published to date.

70 This contribution was motivated by the need for benchmark problems suited to evaluate 71 the effect of electric coupling during multicomponent diffusion and electrochemical migration 72 and to facilitate an inter-comparison of existing reactive transport codes. The following 73 benchmark problems are specifically designed to highlight effects of electromigration. The first 74 two benchmarks are one-dimensional and the third benchmark includes two parts, involving one-75 and two-dimensional scenarios. Three reactive transport codes were used independently for the 76 inter-comparison, namely CrunchFlow (Steefel et al., 2014), MIN3P (Mayer et al., 2002) and 77 PHREEQC (Parkhurst and Appelo, 1999).

# 78 **2.** Governing Equations

#### 79 Mass Transfer in Electrolytic Systems

Species-specific diffusion is necessary to describe the behavior of electrolyte systems (Steefel and Maher, 2009) where diffusive transport is the dominant mass transport process. The most important feature that distinguishes the electrolyte systems from non-electrolyte systems is the electric coupling of the ionic fluxes (Helfferich, 1962; Newman, 1973). In the electrolyte systems, electric interaction of ion-ion, ion-solvent and ion-interface induces an electric field. The treatment of electrolytic diffusion follows naturally from the generalized treatment of diffusion (Taylor and Krishna, 1993).

#### 87 Nernst-Planck Equation for Multicomponent Systems

The migration of interacting species is described by the Nernst-Planck equation, which can be derived from expressions for the diffusive flux written in terms of the chemical potential (Steefel

90 et al, 2014). Written in terms of the flux of an arbitrary species *i*, the Nernst-Planck equation is91 given by:

92 
$$\mathbf{J}_{i} = -\mathbf{D}_{i} \left( \nabla c_{i} + c_{i} \nabla \ln \gamma_{i} + \frac{c_{i} F}{RT} z_{i} \nabla \psi \right)$$
(1)

where  $D_i$  is the species-dependent diffusion coefficient (m<sup>2</sup> s<sup>-1</sup>),  $c_i$  is the concentration (mol L<sup>-1</sup> H<sub>2</sub>O),  $\gamma_i$  is the activity coefficient (-), F is the Faraday constant (96485 C mol<sup>-1</sup>), R is the gas constant (8.341 J K<sup>-1</sup> mol<sup>-1</sup>), T is the absolute temperature (K),  $z_i$  is the charge number (-) and  $\psi$ is the electric potential (V or J C<sup>-1</sup>). In the presence of advection with a Darcy's velocity **q** (m s<sup>-1</sup>), the modified flux term is:

98 
$$\mathbf{J}_{i} = -\mathbf{D}_{i} \left( \nabla c_{i} + c_{i} \nabla \ln \gamma_{i} + \frac{c_{i} F}{RT} z_{i} \nabla \psi \right) + c_{i} \mathbf{q}$$
(2)

99 This expression is known as the extended Nernst-Planck equation and holds, in ideal systems, for 100 all mobile species. It describes the movement of ions in a solution with or without external 101 electric field (Helfferich, 1962, Bard, 1980 and Bagotsky, 2006). In a multicomponent system, 102 the set of Nernst-Planck equations, one for each species, must be solved simultaneously.

By assuming small gradients in ionic strength, a dilute solution with low ionic strength and isothermal conditions, the contribution of the flux from the gradients in the logarithms of the activity coefficients can be neglected (Giambalvo et al., 2002; Steefel and Maher, 2009). With this approximation, the flux of an individual species becomes:

107 
$$\mathbf{J}_{i} = -\mathbf{D}_{i} \left( \nabla \mathbf{c}_{i} + \frac{\mathbf{c}_{i} \mathbf{F}}{\mathbf{RT}} \mathbf{z}_{i} \nabla \psi \right) + \mathbf{c}_{i} \mathbf{q}$$
(3)

108 This equation represents the contributions of diffusion, electromigration and advection to the 109 total mass transfer. Assuming there is no externally induced current (null current assumption), a 110 simplified version of the mass flux can be derived (Giambalvo et al., 2002):

111 
$$\mathbf{J}_{i} = -\mathbf{D}_{i} \left( \nabla \mathbf{c}_{i} + \mathbf{c}_{i} \mathbf{z}_{i} \frac{\mathbf{q} \sum_{j}^{N_{aq}} \mathbf{z}_{j} \mathbf{c}_{j} - \sum_{k}^{N_{aq}} \mathbf{D}_{k} \mathbf{z}_{k} \nabla \mathbf{c}_{k}}{\sum_{i}^{N_{aq}} \mathbf{D}_{i} \mathbf{z}_{i}^{2} \mathbf{c}_{i}} \right) + \mathbf{c}_{i} \mathbf{q}$$
(4)

112 This formulation has the advantage that the electric potential does not appear as a primary 113 unknown and is therefore well suited for implementation in standard reactive transport codes.

# 114 **3.** Participating Codes

115 The three reactive transport codes participating in this benchmarking exercise are CrunchFlow 116 (Steefel et al., 2014), MIN3P (Mayer et al., 2002) and PHREEQC (Parkhurst and Appelo, 1999). 117 CrunchFlow and MIN3P are 3D block-centered finite difference (finite volume) models using 118 the global implicit approach (GIA) to solve the fully coupled transport and reaction equations. 119 PHREEQC solves the multicomponent diffusion problem with a 1D finite difference method 120 using the sequential non-iterative approach (SNIA). A detailed description of the formulation 121 and capabilities of the codes is discussed elsewhere (Steefel et al., 2014 and references therein). 122 PHREEQC considers the gradients of the activity coefficients in its implementation (Appelo and 123 Wersin, 2007) whereas CrunchFlow and MIN3P neglect this contribution.

124

#### 4. Benchmark Descriptions

The three benchmark problems are summarized in **Table 1**. The first benchmark (Benchmark 1) focuses on the role of electromigration in driving the flux of the various charged species to maintain local charge balance and was first presented by Lichtner (1995). This problem considers diffusion of HNO<sub>3</sub> from a low pH solution (pH = 4) into a circum-neutral reservoir (pH = 6) with low HNO<sub>3</sub> concentrations, both with the same elevated NaCl background concentrations.

Benchmark	Description	Processes	Dimension
1	HNO <sub>3</sub> (pH 4) diffusion into a circum-neutral pH reservoir	Diffusion/Electromigration	1D
2	Sodium isotope fractionation induced by sodium chloride diffusion in neutral pH water	Diffusion/Electromigration	1D
3	Transverse dispersion affected by electromigration	Advection/Diffusion/ Electromigration	1D/2D

132 **Table 1.** Summary of multicomponent diffusion benchmarks

131

134 The second benchmark (2) shows the electromigration and subsequent fractionation of the sodium isotope <sup>22</sup>Na<sup>+</sup> due to diffusion of NaCl under neutral pH conditions. The problem is 135 136 loosely based on Glaus et al (2013); however, the benchmark case presented here is set up for a 137 uniform relatively coarse-grained uncharged porous medium and does not include diffusion 138 through charged micropores as would be the case if porous clay were considered. In the first two 139 benchmarks, diffusion and electromigration are the only transport processes and models are set 140 up in one dimension. The third benchmark (3) investigates the effect of electromigration on 141 transverse dispersion and is based on experiments and modeling carried out by Rolle et al. 142 (2013). This benchmark also includes advection and is characterized by a higher level of 143 complexity; it is simulated in one- and two-dimensions.

### 144 Benchmark 1: Transient Electromigration

This problem was initially presented by Lichtner (1995) and has previously been used as an example to illustrate the multicomponent capabilities of PHREEQC (Appelo, 2007). It is a 1D transient problem with a fixed concentration (Dirichlet) boundary condition on the left (at x = 0),

148 representing the reservoir, and a no-flux (Neumann) boundary condition on the right (x = 0.01149 m). The chemical system is composed of four primary (component) species ( $H^+$ ,  $NO_3^-$ ,  $Na^+$  and 150 Cl<sup>-</sup>) and one secondary species (OH<sup>-</sup>). The porosity is set to 1.0 and the domain is discretized 151 into 100 equally spaced cells of 100 microns each. The temperature is 25°C and there is no flow, 152 the only transport process is multicomponent diffusion according to the Nernst-Planck equation. 153 Activity coefficients are calculated with the extended Debye-Hückel equation. Species-154 dependent diffusion coefficients, as well as the initial and boundary conditions defining the 155 chemical system are given in Table 2.

156

**Table 2.** Boundary conditions, initial conditions, and species dependent diffusion coefficients for
 Benchmark 1 (transient electromigration problem).

Species	Boundary Condition	Initial Condition	<b>Diffusion Coefficient</b> (m <sup>2</sup> s <sup>-1</sup> )
	( <b>mM</b> )	( <b>mM</b> )	
pН	6.001	4.007	$9.31 \times 10^{-9}$
Na <sup>+</sup>	0.1	0.1	$1.33 \times 10^{-9}$
Cl	0.1	0.1	$2.03  imes 10^{-9}$
NO <sub>3</sub> <sup>-</sup>	0.001	0.1	$1.90 \times 10^{-9}$
OH	$^{a}1.03 \times 10^{-5}$	$^{a}1.06 \times 10^{-7}$	$5.27 \times 10^{-9}$

<sup>a</sup>OH- concentrations are only provided for completeness, calculated from H<sup>+</sup> and H<sub>2</sub>O ( $K_w = 10^{-14}$ )

160

161 The problem is run for 1 hour using a constant time step of 0.001 hour (corresponding to 1,000 162 time steps). Results are compared along the spatial profile after T = 1 hour for  $H^+$ ,  $Na^+$ ,  $NO_3^-$  and 163 Cl<sup>-</sup>.

#### 164 Benchmark 2: Tracer Isotope Diffusion

165 This 1D problem involves three primary (component) species,  $Na^+$ ,  $CI^-$ , and  $H^+$ , along with an 166 isotope of Na that is also treated as a distinct component, <sup>22</sup>Na<sup>+</sup>. In addition, a single secondary 167 species,  $OH^-$ , is considered. In this case, fixed concentration (Dirichlet) boundary conditions are 168 considered at either end of the domain. The initial condition in the domain is divided into two

169 regions; concentrations in half of the domain are equivalent to those at the left boundary, while 170 concentrations in the other domain half are defined by the right boundary condition. However, 171 the initial conditions are not significant since the simulation is run until steady state conditions 172 are achieved. The porosity is set to a constant and uniform value of 0.5 and the domain is 173 discretized into 100 equally spaced cells of 100 microns each. The diffusion coefficients of Na<sup>+</sup> and <sup>22</sup>Na<sup>+</sup> are assumed to be identical. A constant time step of 1 hour is used and the simulation 174 175 is run to 1,500 days to ensure that steady-state is achieved. Concentrations at the boundaries and 176 species-dependent diffusion coefficients are described in Table 3. The simulation also assumes 177 no flow.

178

179 **Table 3.** Boundary conditions and diffusion coefficients for Benchmark 2 (isotope tracer problem).

Species	Left Boundary Condition (mM)	<b>Right Boundary</b> <b>Condition (mM)</b>	<b>Diffusion Coefficient</b> (m <sup>2</sup> s <sup>-1</sup> )
pН	7.0	7.0	$9.31  imes 10^{-9}$
Na <sup>+</sup>	0.5	0.1	$1.33 \times 10^{-9}$
$^{22}Na^{+}$	10 <sup>-6</sup>	10 <sup>-6</sup>	$1.33 \times 10^{-9}$
Cl	0.5	0.1	$2.03  imes 10^{-9}$
OH	$^{a}1.05 \times 10^{-4}$	$^{a}1.03 \times 10^{-4}$	$5.27 \times 10^{-9}$

<sup>a</sup>OH- concentrations are only provided for completeness, calculated from H<sup>+</sup> and H<sub>2</sub>O ( $K_w = 10^{-14}$ )

181

#### 182 Benchmark 3: Transverse Dispersion

Rolle et al. (2013) investigate the effect of electromigration on transverse dispersion under steady state flow conditions. In the full 2D case, the problem involves unidirectional flow and transport of a multicomponent tracer plume down the length of a 2D flow-through chamber.



187

188 Figure 1. Schematic of the 2D flow and transverse disperison experiment conducted by Rolle et al.(2013).

191 Using PHREEQC, Rolle et al. (2013) solved the problem numerically by simulating transverse 192 dispersion and electromigration perpendicular to the flow path as a 1D problem. This approach 193 simplifies a 2D steady-state problem into a 1D transient problem by making use of the 194 transformation t = x/v, where x is the distance from the source for the 2D problem, v is the 195 uniform average linear groundwater velocity, and t defines the travel time to reach the location x. 196 At the same time, t defines the simulation time for the 1D transient transverse dispersion problem 197 (Rolle et al., 2013). Coinciding with experimental conditions, a 1 cm source in the middle of the 198 12 cm wide cross section at x = 0 describes the continuous release of the electrolyte solution. The simulation was run for the case of an average linear velocity of 1.5 m day<sup>-1</sup>. The results of the 199 200 1D transient simulations are compared among the three participating codes, whereas fully 2D 201 simulations with explicit treatment of flow were performed with CrunchFlow and MIN3P.

- 202
- 203 **Table 4.** Chemical conditions and transverse dispersion coefficients for Benchmark 3 (transverse dispersion methods)
- dispersion problem).

Species	Tracer	Initial Condition (1D)	Diffusion	Transverse
	Injection	and Remaining Injection	Coefficient	Dispersion
	Ports (mM)	Ports (2D) (mM)	$(\mathbf{m}^2 \mathbf{s}^{\cdot 1})$	Coefficient (m <sup>2</sup> s <sup>-1</sup> )
$\mathbf{K}^+$	0.29	10 <sup>-6</sup>	$1.77  imes 10^{-9}$	2. 405 $\times 10^{-9}$
$Mg^{2+}$	0.29	10-6	$6.26 \times 10^{-10}$	$1.745 \times 10^{-9}$
Cl	0.87	$3 \times 10^{-6}$	$1.81 \times 10^{-9}$	$2.425 \times 10^{-9}$

The dispersion coefficients used in these simulations require some discussion. In fact, the parameterization of the hydrodynamic transverse dispersion coefficient used in Rolle et al. (2013) differs from the classical linear model commonly adopted in subsurface applications of solute transport and reads as:

210 
$$D_i^T = D_i^P + D_i^{aq} \left(\frac{Pe^2}{Pe + 2 + 4\delta^2}\right)^{\beta}$$
 (5)

where  $D_i^{P}$  is the pore diffusion coefficient approximated as the product of the aqueous diffusion 211 coefficient of a species *i* and the porosity of the medium (0.41).  $Pe=vd/D_i^{aq}$  is the grain Péclet 212 213 number where d is the average grain size (1.25 mm).  $\delta$ =6.2, and  $\beta$ =0.47 are empirical parameters 214 determined in previous multitracer experiments and pore-scale simulations (Rolle et al., 2012). 215 Equation 7 explicitly retains a direct dependence of the mechanical dispersion term on the 216 aqueous diffusivity of the transported species; the non-linear dependence on the average flow 217 velocity arises from the incomplete mixing in the pore channels (e.g. Hochstetler et al., 2013; 218 Rolle and Kitanidis, 2014).

For this benchmark analysis we considered the mixed electrolyte case described in Rolle et al., 2013, where a dilute solution of KCl and MgCl<sub>2</sub> was continuously injected in ambient deionized water. The free aqueous diffusion coefficients of the ions at T=20 °C are  $D_{K+} = 1.77 \times 10^{-9} \text{ m}^2 \text{ s}^ 1, D_{Mg2+} = 6.26 \times 10^{-10} \text{ m}^2 \text{ s}^{-1}$ , and  $D_{Cl-} = 1.81 \times 10^{-9} \text{ m}^2 \text{ s}^{-1}$ . These values used in Eq. 7 yield the transverse dispersion coefficients given in the last column of **Table 4**.

1D Benchmark: The 1D benchmark consists of a pure transverse diffusion problem discretized into 48 grid cells of 2.5 mm. In the 1D system, the injection ports constitute initial conditions used at grid cells 23-26, corresponding to a 10 mm wide region in the center of the symmetrical system. The initial condition is used everywhere else in the domain and is intended to represent deionized water. The transverse dispersion coefficients given in **Table 4** are used. The boundaries at either end of the system are treated as no-flux, but they do not influence the system behavior for the 16 hour simulation time (corresponding to x = 1 m, i.e. the outflow boundary of the domain). The simulation was run with a constant 0.001 hour time step.

232 **2D Benchmark:** For the full 2D problem solved with CrunchFlow and MIN3P, the transverse 233 discretization is 50 grid cells with a spacing of 2.4 mm (corresponding to a total width of 0.12 234 m). At the inlet boundary, grid cells 24, 25, 26 and 27 in the transverse direction are set at the 235 tracer injection port concentrations of 0.29 mM K<sup>+</sup>, 0.29 mM Mg<sup>2+</sup>, and 0.87 mM Cl<sup>-</sup> (see

236 Table 4), while the remaining injection ports carry deionized water. The longitudinal 237 discretization is 500 grid cells with a spacing of 2.4 mm thus a total length of 1.2 m; the 238 concentrations are reported at x = 1.0 m, corresponding to the outflow boundary of the 239 experimental setup. The additional length of 0.2 m is considered in the models to avoid any 240 possible boundary effects. In this case, lateral flow can be calculated, or simply prescribed at 1.5 m day<sup>-1</sup>. A maximum time step of 1 hour is used with an initial minimum time step of  $10^{-6}$  hours. 241 242 The simulation time is 32 hours to ensure that the final results correspond to steady state 243 conditions representative of the experiment.

244 **5. Results and Discussion** 

#### 245 Benchmark 1

Simulation results for the Benchmark 1 (Lichtner, 1995) depict the diffusion of  $HNO_3$  (pH = 4.007) from the solution domain towards the boundary where  $NO_3^-$  concentration are 100 times lower and pH = 6.001. Results for  $NO_3^-$  and H<sup>+</sup> reveal that both ions continue to diffuse towards

249 the left boundary after 1 hour simulation time (Figure 2). Because the diffusion coefficient for 250  $H^+$  is much larger than the corresponding value for NO<sub>3</sub>,  $H^+$  has become substantially more depleted in the domain than  $NO_3^-$ . The discrepancy in diffusion rates of H<sup>+</sup> and  $NO_3^-$  triggered 251 electromigration of Na<sup>+</sup> and Cl<sup>-</sup> to maintain local charge balance; Na<sup>+</sup> is entering the domain to 252 253 offset the preferential loss of  $H^+$ , while  $Cl^-$  is leaving the system to counterbalance  $NO_3^-$ , which is preferentially retained. Migration of Na<sup>+</sup> and Cl<sup>-</sup> occurs despite the fact that there was no 254 255 initial concentration gradient of either species (Table 2) and takes place even against the 256 developing concentration gradients of Na<sup>+</sup> and Cl<sup>-</sup>. If Fick's Law were used to describe this multispecies diffusion problem, there would be no change in Na<sup>+</sup> and Cl<sup>-</sup> concentration and 257 258 consequently electroneutrality would be violated.

There is very good agreement between the simulation results of all three codes and they demonstrate near identical outputs. Simulations were executed on a desktop computer equipped with an Intel Core 2 Quad CPU with two 2.4 GHz processors, 8 GB RAM and a 64-bit operating system.



Figure 2: Species concentrations after 1 hour simulation time for HNO<sub>3</sub> diffusion (Benchmark 1). The left boundary is a fixed concentration (Dirichlet) boundary, while the right boundary is no-flux.

- 267
- 268 Benchmark 2
- 269 The results of the Benchmark 2 simulation visually show steady state diffusion with same
- 270 concentration profiles for  $Na^+$  and  $Cl^-$  from left to right (
- Figure 3). However, it has to be kept in mind that the diffusion coefficient for Cl<sup>-</sup> is considerably
- 272 larger than the one for Na<sup>+</sup>. In fact, considering that the equations are based on the null current
- assumption, this holds back Cl<sup>-</sup> migration and accelerates Na<sup>+</sup> migration. Although there are no
- 274 initial concentration gradients for <sup>22</sup>Na<sup>+</sup>, H<sup>+</sup> and OH<sup>-</sup>, these species, present at much lower
- 275 concentrations, also become affected by the electrostatic coupling.
- 276





**Figure 3:** Na<sup>+</sup>, Cl<sup>-</sup>, H<sup>+</sup>, OH<sup>-</sup> and <sup>22</sup>Na<sup>+</sup> concentrations after 1500 days for system summarized in Table 2 (Benchmark 2). The left boundary is a fixed concentration (Dirichlet) boundary at 0.5 mM, while the right boundary is a fixed concentration boundary at 0.1 mM for Na<sup>+</sup>, Cl<sup>-</sup>. The fixed gradient in NaCl results in a flux of H<sup>+</sup>, OH<sup>-</sup> and <sup>22</sup>Na<sup>+</sup>, despite the fact that their concentrations are the same at either end of the column.

A closer look at the results reveals that  $H^+$  migrates from the left to the right to enhance the 285 286 positive charge flux, while OH<sup>-</sup> migrates from the right to the left to counteract the negative charge flux from the left to the right dominated by Cl<sup>-</sup>. Primarily, one would expect that <sup>22</sup>Na<sup>+</sup> 287 288 should also be subjected to a net flux from the left to the right; however, the sodium isotope is 289 present at very low concentrations and is more strongly affected by migration dynamics of H<sup>+</sup> 290 and OH, resulting in a net migration form the right to the left inducing an unexpected isotope fractionation. Solving this problem with Fick's law would not predict <sup>22</sup>Na<sup>+</sup> isotope fractionation, 291 292  $H^+$  and  $OH^-$  migration, and would result in a net negative charge flux across the domain. These 293 results suggest that multicomponent diffusion can introduce isotope fractionation, even in the 294 absence of fractionating reactions.

295 Overall, there is very good agreement between the three codes with better agreement between CrunchFlow and MIN3P. Slight differences are observed for the PHREEQC results. It is difficult 296 297 to decisively determine the reasons for these differences, but it is likely that the discrepancies are 298 due to slight variations in model formation (i.e. consideration of activity gradients in the 299 PHREEQC formulation, absent in the other two codes) and/or the use of different coupling 300 schemes (GIM vs. SNIA). However, all codes show identical trends and concentration 301 differences are small, implying that the residual discrepancies will not affect the interpretation of 302 the results.

303

#### 304 Benchmark 3: 1D Transverse Dispersion

The transverse concentration profiles for Cl<sup>-</sup>, K<sup>+</sup> and Mg<sup>2+</sup> are plotted at the outlet (x = 1.0 m) corresponding to a residence time of 16 hours in the 2D domain. The separation of the three tracer profiles (

308 Figure 4) demonstrates the effect of species-dependent dispersion coefficients and 309 electrochemical migration on transverse displacement. The Cl<sup>-</sup> concentration profile is located between  $K^+$  and  $Mg^{2+}$  despite having the largest diffusion coefficient. In fact,  $D_{Cl-}$  in liberated 310 311 state is considerably larger than  $D_{Mg2+}$  and also slightly larger than  $D_{K+}$  (Table 4). These results 312 show that Cl<sup>-</sup> migration is retarded due to electrostatic coupling with the cations and in particular with Mg<sup>2+</sup>, which diffuses more slowly. The outcomes reported in Fig. 4 demonstrate the 313 314 positive contribution of electromigration to transverse displacement of the two cations and the 315 negative contribution of electromigration to transverse displacement of chloride (Rolle et al., 316 2013).



317

318

**Figure 4.** 1D simulation results of transverse profiles for  $Cl^-$ ,  $K^+$  and  $Mg^{2+}$  at the outlet (corresponding to a residence time of 16 hours) demonstrate the effect of species-dependent dispersion and electromigration on the transverse displacement of charged species (1D Benchmark 3 solved with MIN3P).

323 There is a very good agreement between the three codes and an excellent match between 324 CrunchFlow and MIN3P (

Figure 5). Similar to the two previous benchmarks, there are slight differences between the results of CrunchFlow and MIN3P on the one hand and PHREEQC on the other hand. Peak chloride concentrations predicted by PHREEQC are slightly higher than those calculated by CrunchFlow and MIN3P (~ 0.6%). Magnesium and potassium concentration profiles are in very good agreement for all codes.



332

**Figure 5.** 1D simulation of transverse multicomponent diffusion for the case of transport of mixed

electrolytes (KCl and MgCl<sub>2</sub> solution) in pure water described by Rolle et al. (2013), comparing
 CrunchFlow, MIN3P and PHREEQC results.

# 337 Benchmark 3: 2D Flow and Transverse Dispersion

Using CrunchFlow and MIN3P it was possible to carry out a full two-dimensional flow and multicomponent transport simulation of the flow-through system. The simulation was run for two pore volumes (32 hours) to ensure that steady state conditions at the outflow were reached. To illustrate the 2D concentration distributions and to provide a means for visual comparison of the CrunchFlow and MIN3P results, 2D contour plots are provided for K<sup>+</sup>, Mg<sup>2+</sup> and Cl<sup>-</sup> (**Figure** 

**343 6**).



**Figure 6.** Simulation results for Benchmark 3 considering flow (uni-directional) and multicomponent transverse dispersion for steady-state conditions, from top to bottom are shown:  $K^+$ ,  $Mg^{2+}$  and  $Cl^-$  for CrunchFlow and  $K^+$ ,  $Mg^{2+}$  and  $Cl^-$  for MIN3P.

Cross-sections extracted from two-dimensional steady state CrunchFlow and MIN3P results are compared at the outflow to one-dimensional transient PHREEQC results, corresponding to a residence time of 16 hours. Overall, there is an excellent agreement between MIN3P and CrunchFlow results (**Figure 7**) and results are also very close to the concentrations computed with the 1D PHREEQC approach. PHREEQC concentration profiles are slightly higher than CrunchFlow

and MIN3P (the differences of the peak concentrations are ~ 0.6% for Cl<sup>-</sup>, ~ 0.7% for Mg<sup>2+</sup> and 355 356 ~ 0.4% for  $K^+$ ).



359 Figure 7. Comparison of 1D PHREEQC results (no explicit consideration of flow, only following the 360 plume as it moves down the flow path) and transverse profiles derived from 2D CrunchFlow and MIN3P 361 runs for the transverse dispersion problem. The CrunchFlow 2D runs are based on GIMRT and use a first 362 order upwind formulation, along with a backwards Euler time stepping approach, the same numerical 363 methods are used in the MIN3P simulations.

### 365 6. Concluding Remarks

366 Three benchmark problems were presented, each with significant effects of multicomponent 367 diffusion and electromigration on transport of solutes in saturated porous media. The 368 benchmarks were specifically designed to be sensitive to the effect of electromigration on 369 diffusion and lateral concentration displacement. Benchmarks 1 and 2 are hypothetical problems 370 that provide opportunities to verify the implementation of multicomponent diffusion and 371 electromigration formulations in reactive transport codes. Benchmark 3 is based on the outcomes 372 of laboratory experiments (Rolle et al., 2013) and provides the opportunity to verify and validate 373 multicomponent diffusion and species-dependent transverse dispersion formulations under flow-374 through conditions. Three reactive transport codes with the capability of simulating 375 multicomponent diffusion and electrochemical migration participated in this study (CrunchFlow, 376 MIN3P and PHREEQC). For all benchmark problems considered in this work an overall very 377 good agreement between the simulation results obtained with the different codes. Despite some 378 residual discrepancies between the simulation results, all three codes were able to consistently 379 reproduce the same trends and evolution in concentration patterns induced by multicomponent 380 diffusion and by the electrostatic interactions between the charged species. Small discrepancies 381 between the results indicate that different approaches in implementing the governing equations 382 are not a significant source of uncertainties for model applications; uncertainties will rather be 383 dominated by the underlying conceptual model.

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