



Multi-phase Biogeochemical Model for Microbially Induced Desaturation and Precipitation

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Abstract. A next-generation biogeochemical model was developed to explore the impact of the native water source on microbially induced desaturation and precipitation (MIDP) via denitrification. MIDP is a non-disruptive, nature-based ground improvement technique that offers the promise of cost-effective mitigation of earthquake-induced soil liquefaction under and adjacent to existing structures. MIDP leverages native soil bacteria to reduce the potential for liquefaction triggering in the short term through biogenic gas generation (treatment completed within hours to days) and over a longer term through calcium carbonate precipitation (treatment completed in weeks to months). This next-generation biogeochemical model expands earlier modeling to consider multi-phase speciation, bacterial competition, inhibition, and precipitation. This biogeochemical model was used to explore the impact of varying treatment recipes on MIDP products and by-products in a natural seawater environment. The case study presented herein demonstrates the importance of optimizing treatment recipes to minimize unwanted by-products (e.g., H₂S production) or incomplete denitrification (e.g., nitrate and nitrite accumulation).

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Keywords: biogeochemical modeling, liquefaction, denitrification, desaturation, precipitation



1 Introduction

30 Microbially induced desaturation and precipitation (MIDP) is a biogeotechnical technique that takes advantage of native subsurface denitrifying bacteria to mitigate earthquake-induced soil liquefaction (O'Donnell et al., 2017a, b; Pham et al., 2018). MIDP mitigates liquefaction in two ways: generation of nitrogen gas (N_2) that desaturates the soil and mineral precipitation (usually calcium carbonate, $CaCO_3$) that improves soil strength. A primary benefit of MIDP for liquefaction mitigation is, being non-disruptive, it can be used underneath existing structures (O'Donnell et al., 2017a; Hall, 2021).

35 For environmental and economic reasons, local source water is used to prepare the MIDP treatment solution (i.e., dissolve the substrates needed to induce denitrification). However, naturally occurring constituents in the source water may affect denitrification. For example, a competing electron acceptor (e.g., sulfate) may consume the electron donor (e.g., acetate), leading to incomplete denitrification and the formation of unwanted products (e.g., hydrogen sulfide). In addition, the biogeochemical reactions during MIDP result in changes to alkalinity and pH that may alter process kinetics and final MIDP
40 products.

To quantify the impact of source-water composition and to understand the complex biogeochemical interactions that occur during field application, we developed a next-generation biogeochemical model of MIDP. This mathematical model expands upon previous modeling of MIDP, which did not consider the impact of source water on MIDP or the impact of MIDP on the aqueous subsurface environment (Pham, 2017; O'Donnell et al., 2019). Our next-generation MIDP model includes all
45 essential biogeochemical processes based on the constituents commonly observed in the natural groundwater environments, substrates added to stimulate MIDP, and mechanisms that lead to desaturation and precipitation: e.g., N_2 -gas formation, acid-base speciation, and $CaCO_3$ precipitation. Since MIDP often is deployed in coastal areas (due to the prevalence of liquefiable soil deposits in this environment), we include conditions typical for coastal seawater in our model.

2 Model Foundation

50 The next-generation model builds upon previous MIDP models (Pham, 2017; O'Donnell et al., 2019), but broadens the range of processes considered by the model. Our next-generation model considers microbial growth and decay, alternative microbial metabolic processes, gas production, mineral-solids production, alkalinity and pH, microbial inhibition, and desaturation and precipitation in both fresh water and coastal environments. A comparison of the components and processes considered by the two earlier MIDP models and our next-generation model is provided in Table A1 in Appendix A.

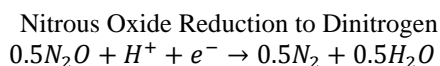
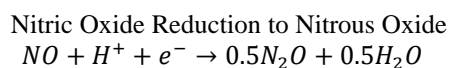
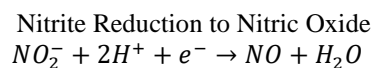
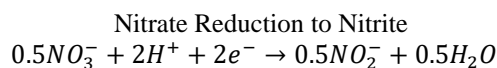
55 The next-generation model was constructed in Matlab (Little and Moler, 2017), and the code and necessary files are publicly available online at doi.org/10.5281/zenodo.7410676. The modeling equations (e.g., microbial growth, $CaCO_3$ precipitation, and biogenic gas evolution) were programmed within the original, publicly available van Turnhout Toolbox, a general-form mechanistic model for environmental systems (van Turnhout et al., 2016). The van Turnhout Toolbox simulates chemical speciation with ORCHESTRA (Meeussen, 2003), an extensive database of established geochemical equilibria. The
60 MIDP-specific biogeochemical model components (i.e., stoichiometry, type of inhibition and kinetics, potential chemical



species) were specified in an input spreadsheet that the program accesses. The degree of saturation and percent (by weight) of mineral precipitation were calculated outside of the van Turnhout Toolbox using model results, as discussed in Section 3.2 of this paper.

3 Model Principles

65 Denitrification is a multi-step process of nitrogen-species reduction. During each reduction step, energy and biomass are produced when paired with oxidation of an electron donor that produces a thermodynamically favorable reduction-oxidation (redox) reaction. The four steps of nitrogen reduction in denitrification conform to the following reduction half reactions, each consuming two or one electron equivalent (e^-) (Rittmann and McCarty, 2020):



70 In the model developed herein, the four steps were simplified to two steps, nitrate to nitrite and nitrite to dinitrogen gas. The reductions of nitric oxide and nitrous oxide are assumed to occur completely, as they are thermodynamically favorable (Appendix A). In contrast, the accumulation of NO_2^- must be considered explicitly because it is a denitrification inhibition ‘bottleneck’ and presents a risk to human health (Pham et al., 2018; Almeida et al., 1995). The variables for the equations used in the model described in the next several sections are detailed in Table 1.



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Table 1. Notation, units, and description for variables used in the biogeochemical model

Symbol	Units	Description
\hat{q}	$\text{mol}_{\text{donor}} \text{mol}_{\text{biomass}}^{-1} \text{d}^{-1}$	maximum specific rate of electron-donor utilization
X_a	mol L^{-1}	active biomass concentration
C_d	mol L^{-1}	electron donor concentration
K_d	mol L^{-1}	electron donor half-maximum substrate concentration of the
C_a	mol L^{-1}	electron acceptor concentration
K_a	mol L^{-1}	electron acceptor half-maximum substrate concentration
I_i		inhibition factor ($0 < I_i < 1$)
Y	$\text{mol}_{\text{biomass}} \text{mol}_{\text{donor}}^{-1}$	biomass yield from consumed electron donor substrate
μ_{max}	d^{-1}	maximum specific microbial growth rate; product of Y and \hat{q}
b	d^{-1}	endogenous decay
K_i	mol L^{-1}	inhibition constant
C_i	mol L^{-1}	concentration of the inhibiting species
$v_{i[g]}$	$\text{mol L}^{-1} \text{d}^{-1}$	transfer rate from the aqueous phase to the gas phase
k_{La}	d^{-1}	mass transfer rate constant
$C_i[g]$	mol L^{-1}	gas phase concentration of the gas species i
$C_i[aq]$	mol L^{-1}	aqueous phase concentration of the biogenic gas species i
K_H	L atm mol^{-1}	Henry's Law constant
R	$\text{L atm mol}^{-1} \text{K}^{-1}$	universal gas constant
T	K	system's absolute temperature
$[N_2]_g$	$\text{mol L}_{\text{pore}}^{-1}$	produced N_2 gas during MIDP
$[CO_2]_g$	$\text{mol L}_{\text{pore}}^{-1}$	produced CO_2 gas
p	atm	pressure at treatment depth
S_g	$\text{L}_{\text{gas}} \text{L}_{\text{pore}}^{-1}$	gas saturation level
l	$\text{L}_{\text{aq}} \text{L}_{\text{pore}}^{-1}$	aqueous solution in the pore space
p_{N_2}	atm	partial pressure of N_2 gas
K_{H,N_2}	$\text{L}_{\text{aq}} \text{atm}_{N_2} \text{mol}_{N_2}^{-1}$	Henry's constant for N_2 at standard temperature
p_{CO_2}	atm	partial pressure of CO_2 gas
K_{H,CO_2}	$\text{L}_{\text{aq}} \text{atm}_{CO_2} \text{mol}_{N_2}^{-1}$	Henry's constant for CO_2 at standard temperature
$Y_{NO_3^-}$	$\text{mol}_{NO_3^-} \text{mol}_{\text{donor}}^{-1}$	stoichiometric coefficients of NO_3^-
Y_{N_2}	$\text{mol}_{N_2} \text{mol}_{\text{donor}}^{-1}$	stoichiometric coefficients of N_2
Y_{CO_2}	$\text{mol}_{CO_2} \text{mol}_{\text{donor}}^{-1}$	stoichiometric coefficients of CO_2
ϕ	$\text{L}_{\text{pore}} \text{L}_{\text{total}}^{-1}$	soil porosity
R_p	$\text{mol L}^{-1} \text{d}^{-1}$	net rate of precipitation ($R_p > 0$) or dissolution ($R_p < 0$) of minerals
ka	L d^{-1}	combined coefficient for constant mineral growth rate and the average crystal surface area
K_{sp}	$\text{mol}^2 \text{L}^{-2}$	constant solubility product.
Y_{CaCO_3}	$\text{mol CaCO}_3 \text{mol}_{\text{donor}}^{-1}$	$CaCO_3$ yield
$[NO_3^-]_C$	$\text{mol L}_{\text{pore}}^{-1}$	NO_3^- needed to achieve the target $CaCO_3$
e	$\text{L}_{\text{pore}} \text{L}_{\text{soil}}^{-1}$	void ratio
ρ_{soil}	$\text{kN L}_{\text{soil}}^{-1}$	soil density
u_{CaCO_3}	$\text{g CaCO}_3 \text{mol}^{-1} \text{CaCO}_3$	molarity to molecular weight conversion coefficient



3.1 Microbial Metabolism, Growth, and Decay

The processes within the model follow Monod kinetics, represented as multiplicative dual-substrate limitation
80 (O'Donnell et al., 2019; Bae and Rittmann, 1996).

$$\frac{dC_d}{dt} = -\hat{q}X_a \frac{C_d}{K_d+C_d} \cdot \frac{C_a}{K_a+C_a} I_i \quad (1)$$

Eq. 1 considers the electron-donor substrate (C_d) and three electron-acceptor substrates (C_a): NO_3^- and NO_2^- for denitrifying
bacteria and SO_4^{2-} for sulfate-reducing bacteria. For preliminary analysis, we assumed an initial denitrifier biomass
concentration of 0.5 mmol L^{-1} and sulfate-reducing biomass concentration of 0.25 mmol L^{-1} . The values of the constants (\hat{q} ,
85 K_d , and K_a) are in Appendix A. Derivations of important microbial kinetics parameters, found in Appendix A, are based on
Rittmann and McCarty (2020). These derivations were used to determine reaction stoichiometry, true yield (Y) and μ_{max}
(maximum specific growth rate) for all electron-donor and -acceptor pairs and the nitrogen source. Values of the kinetic and
stoichiometric parameters are detailed in Table 2. The inhibition factor I_i is described in a later section.

Values of half-maximum-rate concentrations (K_d and K_a) in the literature show variability for each electron donor and
90 acceptor pair due to the wide range of environments of the microorganisms (e.g., sediment, estuarine water, wastewater) and
the high degree of diversity of microorganisms able to carry out these reactions (Abdul-Talib et al., 2002; Papaspyrou et al.,
2014; Vavilin and Rytov, 2015). Table 3 details the constants we used as representative values for each K_d and K_a (for Eq. 1)
based on relevant electron-donor and -acceptor pairs and sources of those values. While, these values are not specific to a
coastal seawater environment, they have been experimentally validated.

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Table 2. Reaction stoichiometry, yield (Y), and maximum specific growth rates (μ_{max}) expected during MDP, considering acetate as the electron donor and natural electron acceptors. Units for all parameters are in Table 1.

Electron Acceptor	Nitrogen Source	\hat{q}	Y	μ_{max}	Reaction Stoichiometry
Nitrate	Nitrate	8.12	0.82	6.68	$0.222NO_3^- + 0.125C_2H_3O_2^- + 0.146H^+$ $\rightarrow 0.202NO_2^- + 0.147H_2CO_3$ $+ 0.103CH_{1.8}O_{0.5}N_{0.2} + 0.021H_2O$
Nitrite	Nitrate	11.69	0.99	11.6	$0.054NO_3^- + 0.202NO_2^- + 0.270C_2H_3O_2^- + 0.525H^+$ $\rightarrow 0.101N_2 + 0.272H_2CO_3$ $+ 0.268CH_{1.8}O_{0.5}N_{0.2} + 0.154H_2O$
Sulfate	Nitrate	3.74	0.58	2.18	$0.015NO_3^- + 0.072SO_4^- + 0.125C_2H_3O_2^- + 0.284H^+$ $\rightarrow 0.072H_2S + 0.177H_2CO_3$ $+ 0.073CH_{1.8}O_{0.5}N_{0.2} + 0.015H_2O$
Nitrate	Ammonium	6.95	1.01	6.99	$0.236NO_3^- + 0.125C_2H_3O_2^- + 0.025NH_4^+ + 0.10H^+$ $\rightarrow 0.236NO_2^- + 0.124H_2CO_3$ $+ 0.126CH_{1.8}O_{0.5}N_{0.2} + 0.050H_2O$
Nitrite	Ammonium	9.65	1.26	12.2	$0.235NO_2^- + 0.261C_2H_3O_2^- + 0.066NH_4^+ + 0.431H^+$ $\rightarrow 0.118N_2 + 0.193H_2CO_3$ $+ 0.328CH_{1.8}O_{0.5}N_{0.2} + 0.249H_2O$
Sulfate	Ammonium	3.63	0.18	0.66	$0.113SO_4^- + 0.125C_2H_3O_2^- + 0.005NH_4^+ + 0.346H^+$ $\rightarrow 0.113H_2S + 0.227H_2CO_3$ $+ 0.023CH_{1.8}O_{0.5}N_{0.2} + 0.009H_2O$

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Table 3. Half-maximum-rate concentrations, K_d and K_a , used for each electron-donor and -acceptor pair

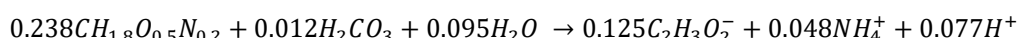
Electron Donor	K_d	Reference	Electron Acceptor	K_a	Reference
Acetate ($C_2H_3O_2^-$)	$1.0 \cdot 10^{-5}$	(Jia et al., 2020)	Nitrate (NO_3^-)	$5.4 \cdot 10^{-5}$	(Abdul-Talib et al., 2002)
Acetate ($C_2H_3O_2^-$)	$1.0 \cdot 10^{-5}$	(Jia et al., 2020)	Nitrite (NO_2^-)	$2.4 \cdot 10^{-5}$	(Abdul-Talib et al., 2002)
Acetate ($C_2H_3O_2^-$)	$7.1 \cdot 10^{-5}$	(Ingvorsen et al., 1984)	Sulfate (SO_4^-)	$2.00 \cdot 10^{-4}$	(Ingvorsen et al., 1984)



115 Microbial growth within the model is represented via reaction kinetics and stoichiometry expressed in Eq. 2:

$$\frac{dX_a}{dt} = X_a Y \hat{q} - b \quad (2)$$

Biomass yields (Y) are listed in Table 2. For sulfate-reducing bacteria, b was set to 0.03 d^{-1} , whereas it was set to 0.05 d^{-1} for denitrifiers (Rittmann and McCarty, 2020). As a result of decay, NH_4^+ is released and can serve as a nitrogen source for denitrification. Since NH_4^+ is thermodynamically favorable over NO_3^- as a nitrogen source, it is used first before NO_3^- during denitrification. Decay involves endogenous respiration, and we assumed that 80% of decayed biomass is available as an acetate for metabolism, while 20% becomes inert biomass (Rittmann and McCarty, 2020). The stoichiometry for decay is:



3.2 Inhibition

Denitrification inhibition, which slows nitrate and nitrite reduction rates (Glass et al., 1997), was included for the reduction of nitrate to nitrite and nitrite to N_2 gas. I_i is a general term for inhibition of either step, with i indicating which reaction. The form of I_i , shown in Eq. 3 is for non-competitive inhibition, and the inhibition coefficients for each inhibitor are found in Table 4:

$$I_i = \frac{K_i}{K_i + C_i} \quad (3)$$

Table 4. Non-competitive inhibition coefficients (K_i)

Inhibiting Compound	Reduction Process Inhibited	K_i (mol L^{-1})	Source
HNO_2	Nitrate	$2 \cdot 10^{-6}$	(Ma et al., 2010)
HNO_2	Nitrite	$8 \cdot 10^{-8}$	(Glass et al., 1997)
Salinity (as NaCl)	Nitrate, nitrite	0.51^a ; 0.78^b	^a (Panswad and Anan, 1999) ^b (Mariangel et al., 2008)
H_2S	Nitrate, nitrite	$6 \cdot 10^{-5}$	(Pan et al., 2019)
NO_3^-	Sulfide	$1 \cdot 10^{-3}$	(Veshareh et al., 2021)
NO_2^-	Sulfate	$1 \cdot 10^{-3}$	(Veshareh et al., 2021)

^aUnacclimated environments were DI and drinking water, ^bAcclimated environments were groundwater and sea water

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Although several inhibitors could affect MIDP, HNO_2 is the most important inhibitor of the MIDP process (Lilja and Johnson, 2016). Significant inhibition to overall denitrification has been reported at $0.04 \text{ mg HNO}_2 \text{ L}^{-1}$ during NO_2^- reduction (approximately 95% rate reduction) (Glass et al., 1997; Abeling and Seyfried, 1992), and a 60% decrease in NO_3^- reduction at $0.08 \text{ mg HNO}_2 \text{ L}^{-1}$ also was reported (Ma et al., 2010). Within the model, HNO_2 inhibits NO_3^- and NO_2^- reductions using the same inhibition coefficient (Table 4). The inhibition by HNO_2 is driven by pH speciation because NO_2^- is dominant at a pH of 3.4 and higher and HNO_2 is negligible for $\text{pH} \geq 7.6$. However, only a small concentration of HNO_2 can have a significant impact on denitrification, which underscores the importance accumulation of the intermediate NO_2^- and pH.

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Competitive inhibition between nitrate and nitrite reductions has been identified, with the presence of nitrate having a larger effect on nitrite reduction than nitrite on nitrate reduction (Lilja and Johnson, 2016; Glass et al., 1997; Almeida et al., 1995; Soto et al., 2007). Nitrite accumulation increases in the presence of nitrate until nitrate is depleted, such that nitrite reduction becomes the dominant process (Glass and Silverstein, 1998). When only nitrite remains, the rate of nitrite reduction increases. However, others have described that, as long as the electrons are adequately provided by the electron donor, competitive inhibition between nitrate and nitrite reductions is not significant (Soto et al., 2007; Ma et al., 2010; van den Berg et al., 2017). Therefore, the model does not include competitive inhibition, although it naturally includes competition for the electron donor between nitrate and nitrite reductions.

The model applies different non-competitive inhibition constants for salinity (as NaCl) for nitrate and nitrite reduction because nitrite reduction is more sensitive to salinity than nitrate. Because the magnitude of inhibition depends on experimental conditions and adaptation of the microorganisms, K_i value may differ for local conditions (Krishna Rao and Gnanam, 1990).

Hydrogen sulfide (H_2S) also can be inhibitory to denitrification (Pan et al., 2019). Nitrate, nitrite, and N_2O reductions have been inhibited by H_2S , though the extent and sensitivity of reduction in the presence of H_2S was experiment-dependent (Senga et al., 2006; Pan et al., 2013; Tugtas and Pavlostathis, 2007; Liang et al., 2020; Cardoso et al., 2006). Within the model, one aqueous-phase H_2S -inhibition constant was used for both NO_3^- and NO_2^- reduction steps.

A $\text{pH} < 6$ can significantly slow denitrification (Glass and Silverstein, 1998) by inhibiting enzyme activity (Šimek and Cooper, 2002) and microbial growth (Estuardo et al., 2008). When the pH goes higher than 8, enzyme activity also can be impeded, leading to reduced denitrification rates or incomplete denitrification. Incidents of a high pH often are temporary, as CaCO_3 precipitation in MIDP buffers the pH (Salek et al., 2015). The benefit of including a pH -inhibition function when predicting denitrification has been demonstrated, but the values of their governing parameters are environment-specific and require fitting (Estuardo et al., 2008). Within the model, we considered the indirect net effect of pH only through HNO_2 inhibition, which does not require environment-specific parameters because the concentration of HNO_2 is automatically calculated within the model structure.

3.3 Biogenic Gas Production

O'Donnell et al. (2019) considered the production of N_2 and CO_2 during denitrification but did not consider the varying subsurface stresses that would influence phase transfer. The relative concentrations of the produced biogenic gas can affect the distribution of gas at depth, since the gases have different solubilities, as well as different stoichiometries for electron-donor consumption.

Our next-generation MIDP model includes mass-transfer kinetics for transfers of N_2 , CO_2 , and H_2S from the aqueous phase to the gas phase (or from the gas phase). N_2 , CO_2 , and H_2S concentrations were modeled in the aqueous and gas phases. The rate of transfer of a gaseous compound from the aqueous phase to (or from) the gas phase, $v_{i(g)}$, depends on the gas's degree of super-saturation and a mass-transfer-rate coefficient (Salek et al., 2015):



$$170 \quad v_{i[g]} = k_1 a (C_{i[g]} - \frac{C_{i[aq]} RT}{K_H}) \quad (4)$$

We assigned $k_1 a$ values for N_2 , CO_2 , and H_2S of 5 d^{-1} (Yongsiri et al., 2004), though the values can vary widely based on porous medium conditions and temperature. We did not include pore-scale kinetics. The aqueous concentrations of CO_2 and H_2S depend on the pH, as described below.

The biogenic gas volume needed to achieve a target level of desaturation (S_g) by N_2 ($[N_2]_g$) and CO_2 ($[CO_2]_g$) was
 175 determined by:

$$[N_2]_g + [CO_2]_g = \frac{p S_g}{RT} \quad (5)$$

in which p was assumed to be equal to the sum of the hydraulic pressure at the treatment depth (7.6 m in an upcoming example) and the atmospheric pressure. Gas-phase H_2S was not included in the desaturation calculations because its solubility is much higher than N_2 and CO_2 .

Eq. 6 describes the amount of input NO_3^- required for desaturation by N_2 and CO_2 (NO_{3^-d} , $\text{mol}_{NO_3} \text{ L}_{\text{pore}}^{-1}$) at the deepest
 180 target treatment depth, which is the lowest depth of the treated zone. The depth increases the pressure (p_{N_2} and p_{CO_2}) and the needed amount of gas production to exceed the solubility threshold (K_{H,N_2} and K_{H,CO_2}) and enter the gas phase, according to Henry's Law. The equation considers the amount of gas needed to overcome the solubility threshold to achieve the target level of desaturation (Hall et al., 2018; Pham, 2017):

$$185 \quad NO_{3^-d} = \frac{\left(\frac{[N_2]_g + \frac{p_{N_2}}{K_{H,N_2}}}{l}\right) Y_{NO_3^-}}{Y_{N_2}} + \frac{\left(\frac{[CO_2]_g + \frac{p_{CO_2}}{K_{H,CO_2}}}{l}\right) Y_{NO_3^-}}{Y_{CO_2}} \quad (6)$$

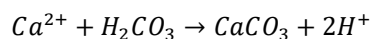
K_{H,N_2} was set to 1600 and K_{H,CO_2} to 29 (L atm mol^{-1}). We determine the amount of nitrate needed to achieve target desaturation levels at the lowest depth, because greater depth requires a higher concentration of gas to achieve target desaturation levels, as the pressures are at their respective maxima.

Eq. 7 was used to determine the biogenic gas volume (V_g , $\text{L}_{\text{gas}} \text{ L}_{\text{tot}}^{-1}$),

$$190 \quad V_g = \frac{S_g RT \varphi}{p} \quad (7)$$

3.4 Solids Precipitation and Dissolution

Precipitation occurs when dissolved inorganic carbon (DIC), produced from microbial substrate conversion of the electron donor, exceeds the solubility of $CaCO_3$ for the concentration of Ca^{2+} present. The stoichiometry for $CaCO_3$ precipitation is:





195 The van Turnhout Toolbox considers precipitation based on equilibrium calculations from the ORCHESTRA module
(Meeussen, 2003). This assumption is valid when the rates of precipitation and dissolution of minerals are much faster than
the phase transfer between the aqueous and solid phases (Salek et al., 2015). Previous MIDP modeling did not consider
precipitation kinetics, but assumed instantaneous equilibrium (Pham, 2017; O'Donnell et al., 2019). Instantaneous equilibrium
may be an over-simplification for environmental conditions (Singurindy et al., 2004) in which the mechanisms of crystal
200 nucleation, crystal growth, and mass transfer of reactants to the contact point of crystal growth are important (Rittmann et al.,
2002). Therefore, we included precipitation and dissolution kinetics in the next-generation model.

The model considers first-order precipitation and dissolution kinetics with respect to the Ca^{2+} concentration (Rittmann
et al., 2002):

$$R_p = ka \left(1 - \frac{K_{sp}}{[\text{Ca}^{2+}][\text{CO}_3^{2-}]} \right) [\text{Ca}^{2+}] \quad (8)$$

205 K_{sp} was set to $1.83 \cdot 10^{-8} \text{ mol}^2 \text{ L}^{-2}$ at 25°C for CaCO_3 , ka is a combined kinetic coefficient, because it is difficult to separate mass
transfer kinetics, crystal growth rate, and solid surface area (Rittmann et al., 2002; Spanos and Koutsoukos, 1998; Rittmann et
al., 2003). ka can have a large range depending on the environment and the ease of establishing precipitation nucleation points.
We assumed ka was 100 L d^{-1} , though this value should be used as a fitting parameter subject to experimental validation.
Precipitation was implemented using the van Turnhout Toolbox's method for biochemical reactions; ka was specified as a
210 reaction rate, and K_{sp} was among the governing input parameters.

Eq. 9 was used to determine the amount of substrate needed to achieve a target precipitation level, which is determined
by the ratio between mass of precipitated CaCO_3 and mass of the soil solids ($[\text{CaCO}_3]$, $\text{kg CaCO}_3 \text{ kg soil}^{-1}$).

$$[\text{NO}_3^-]_c = \frac{[\text{CaCO}_3] \rho_{\text{soil}} Y_{\text{NO}_3^-}}{e u_{\text{CaCO}_3} Y_{\text{CaCO}_3}} \quad (9)$$

The stoichiometric coefficients considered the total amount of input NO_3^- and produced H_2CO_3 for the total assumed two-step
215 denitrification process. The DIC available for precipitation to provide Y_{CaCO_3} is estimated based on pH-driven speciation at
each time step.

3.5 Determining pH

Because pH governs the concentration of important aqueous species based on acid/base speciation, the pH influences
many of the geochemical reactions involved in MIDP. The pH was determined using the geochemical equilibrium software
220 ORCHESTRA, which is part of the van Turnhout Toolbox. ORCHESTRA uses a mass balance on all species within the system
and the products of rate-dependent processes as a function of time (i.e., kinetic, biogeochemical, and phase transfer processes).
At each time step, the program performs a mass balance on all complexed species and their fate (e.g., transformed through
microbial processes, precipitation, gas phase transfer) (van Turnhout et al., 2016; Meeussen, 2003). The program's logic flow
and calculation sequence are found in Appendix A.



225 4 Case Study MIDP Behavior Seawater Conditions: Model Results and Discussion

To demonstrate the capabilities of the model, we illustrate MIDP behavior when targeting desaturation for liquefaction mitigation in a coastal geochemical environment. In this case study, we demonstrate the impact of precipitation on biochemical reactions and the resulting multi-phase products and by-products resulting from MIDP and other environmental biogeochemical processes (e.g., sulfate reduction). However, we only consider desaturation as a target treatment mechanism and do not model an MIDP treatment recipe optimized for precipitation as a liquefaction-mitigation mechanism.

Table 5 details the chemical characteristics used to simulate coastal groundwater conditions, which were assumed to have the same characteristic of seawater due to intrusion (hereafter, referred to as “seawater”). The treatment substrate was added to the baseline level of these components. We based the target treatment zone’s soil properties on a case study of microbial desaturation via denitrification in Portland, Oregon presented by Moug et al. (2022). The deepest target treatment depth was 7.6 m. We assumed typical values of total unit weight, dry density, and porosity for uniform clean sand for the soil (Christopher et al., 2006): total unit weight of 19.5 kN m^{-3} (dry unit weight of 15.6 kN m^{-3} ; bulk density of 1950 kg m^{-3}) and porosity of 0.39.

Table 5. Chemical Characteristics Assumed for a Coastal Seawater Environment

Compound	Coastal Seawater
Nitrate	$20.3^{[1]} \mu\text{mol L}^{-1}$
Nitrite	$0.14^{[1]} \mu\text{mol L}^{-1}$
Sulfate	$28.2^{[2]} \text{mmol L}^{-1}$
DIC	$2.13^{[1]} \text{mmol L}^{-1}$
pH	$7.61^{[1]}$
Ammonium	$0.25^{[1]} \mu\text{mol L}^{-1}$
Iron	$0.60^{[3]} \text{nmol L}^{-1}$
Sodium	$0.47^{[2]} \text{mol L}^{-1}$
Calcium	$10.3^{[2]} \text{mmol L}^{-1}$
Chloride	$0.55^{[2]} \text{mol L}^{-1}$

^[1]Average of measured values (Alin et al., 2017)

240 ^[2]Reference composition of “standard seawater” from and calculated for pH = 7.61 for acid-base species (Millero et al., 2008; European Commission. Directorate General for Research., 2011)

^[3](Bruland et al., 2001)

The reported desaturation levels required to increase the cyclic shear resistance for liquefaction mitigation range between 2 to 10% (He and Chu, 2014; O’Donnell et al., 2017a). We chose 10%, which is at the high end of the mitigation range, but well below the desaturation level at which gas starts to migrate upward or spread laterally, reported to occur at 20% for poorly graded (i.e., uniform) fine sands (Pham, 2017).

Following Eq. 5, 7.10 mmol L^{-1} of total N_2 gas is required to meet a minimum target desaturation level of 10% throughout the entire treatment zone (assuming only desaturation via N_2 gas). To meet the target desaturation, we estimated the treatment recipe to be 22.4 mmol L^{-1} of nitrate ($1.84 \text{ g calcium nitrate L}^{-1}$) and 32.1 mmol L^{-1} of acetate ($2.54 \text{ g calcium acetate L}^{-1}$) using Eq. 6. However, based on background levels of nitrate and nitrite and the use of released ammonium as a nitrogen source, these levels were adjusted to establish the treatment recipe detailed in Table 6. The adjusted values were set



to result in complete denitrification (i.e., no residual nitrate or nitrite that relies on bacterial decay as the electron donor) and to not exceed 0.1 mmol L⁻¹ of acetate after complete denitrification. We compared the impact of varying the input levels of acetate (as calcium acetate) on the MIDP treatment to the matched treatment recipe. We tested the impact of addition of an extra 25% of acetate over our original estimations, referred to as the ‘Excess Acetate’ treatment recipe, and 25% less acetate from our original estimation, referred to as the ‘Reduced Acetate’ treatment recipe. For the excess- and reduced-acetate comparisons, we did not adjust the input levels of nitrate (as calcium nitrate) from our original estimations of 22.4 mmol L⁻¹ (1.84 g calcium nitrate L⁻¹).

260 **Table 6. MIDP treatment recipes for each modeled condition.**

	Matched	Excess Acetate	Reduced Acetate
Nitrate (mmol L ⁻¹)	19.0 (1.56 g calcium nitrate L ⁻¹)	22.4 (1.84 g calcium nitrate L ⁻¹)	22.4 (1.84 g calcium nitrate L ⁻¹)
Acetate (mmol L ⁻¹)	22.4 (1.77 g calcium acetate L ⁻¹)	40.1 (3.17 g calcium acetate L ⁻¹)	24.0 (1.90 g calcium acetate L ⁻¹)

The results of the matched treatment recipe on the subsurface gas volume and saturation profile are shown in the top two panels of Figure 1. For the coastal seawater conditions, the target desaturation level of 10% at 7.6 m (or a degree of saturation of 90%) was achieved by N₂ generation in approximately 2.1 days. The amount of CO₂ produced did not reach its saturation threshold, and CO₂ did not contribute to desaturation at any of the modeled depths. The difference in volume of gas at the different levels is due to the increase in pressure with depth.

The middle panels of Fig. 1 indicate that adding excess acetate increased the degree of saturation at 7.6 m, which is shown by less than 90% saturation at 7.6 m. In contrast, the bottom two panels show that adding less acetate slowed N₂ generation so that 90% saturation was not reached at 7.6 m in 60 days.



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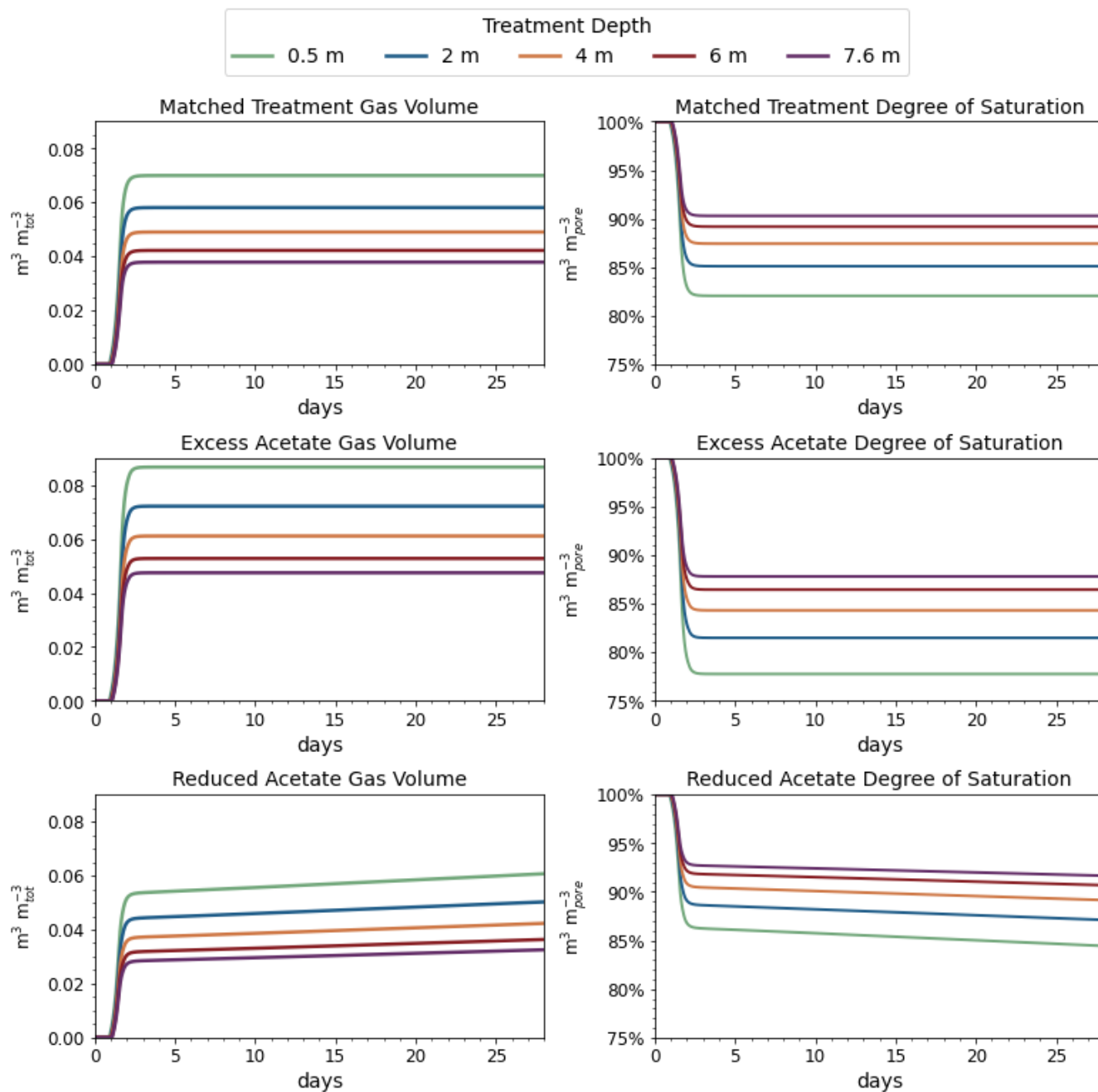


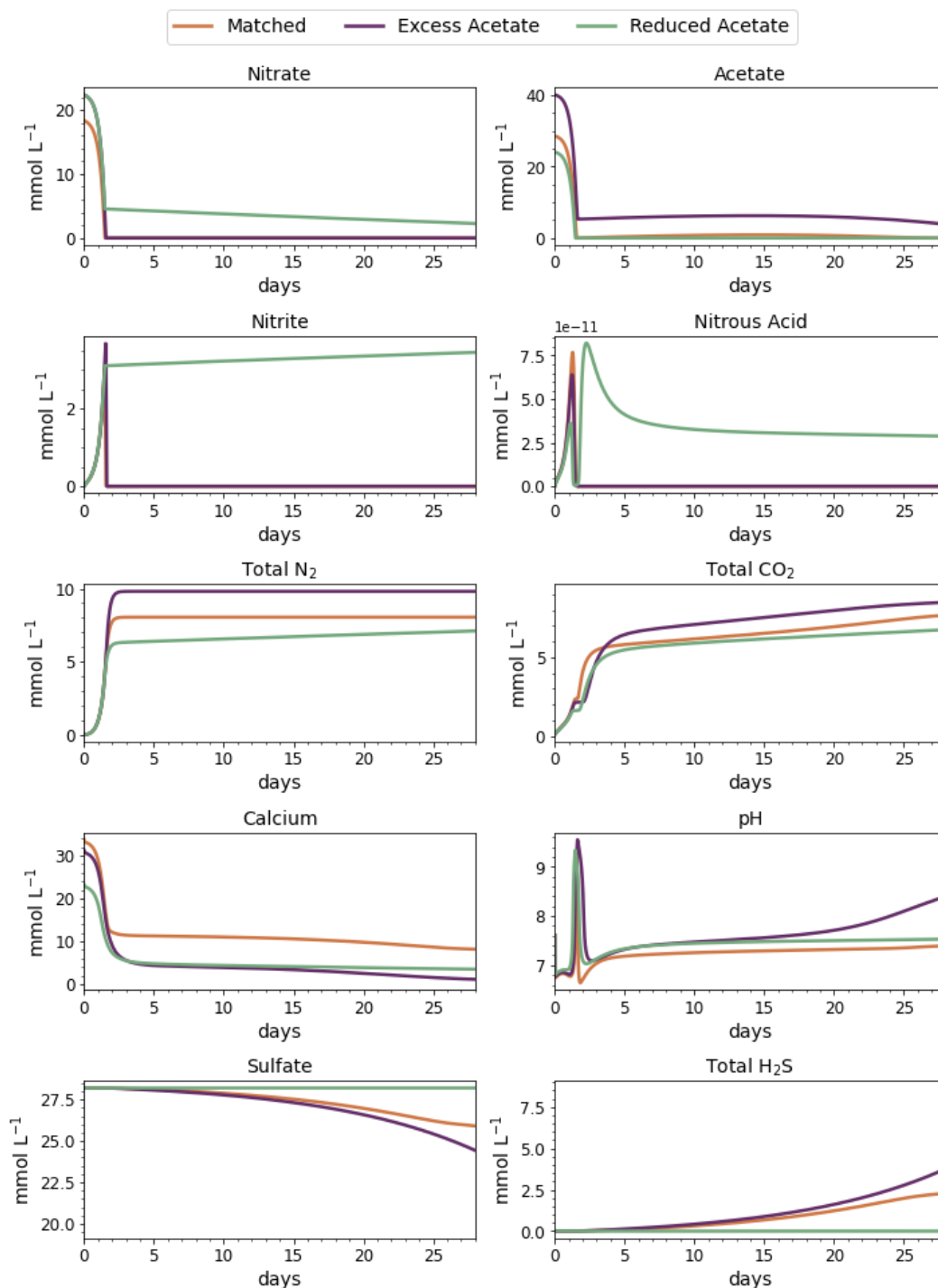
Figure 1. Gas volumes normalized to the total soil volume (left) and degree of saturation by depth for the simulated Coastal seawater conditions. The desaturation target was 10%, or a saturation ratio of 90%.



275 The impacts of the different MIDP treatment recipes on water quality are shown in Figure 2, and the initial five days
of treatment are highlighted in Figure 3. With the matched-treatment recipe, almost all nitrate and nitrite were consumed by
1.6 days, and only 10^{-4} mol L⁻¹ of acetate remained (Figure 3). Nitrite accumulation was transient and modest (3 mM at its
peak); thus, complete denitrification was achieved with this treatment recipe for coastal seawater conditions (Figure 3). After
1.6 days, sulfate reduction began and continued to occur, driven by microbial endogenous respiration, at a small rate that
280 resulted in the production of more total CO₂ and total H₂S than the matched treatment over time (Figure 2).

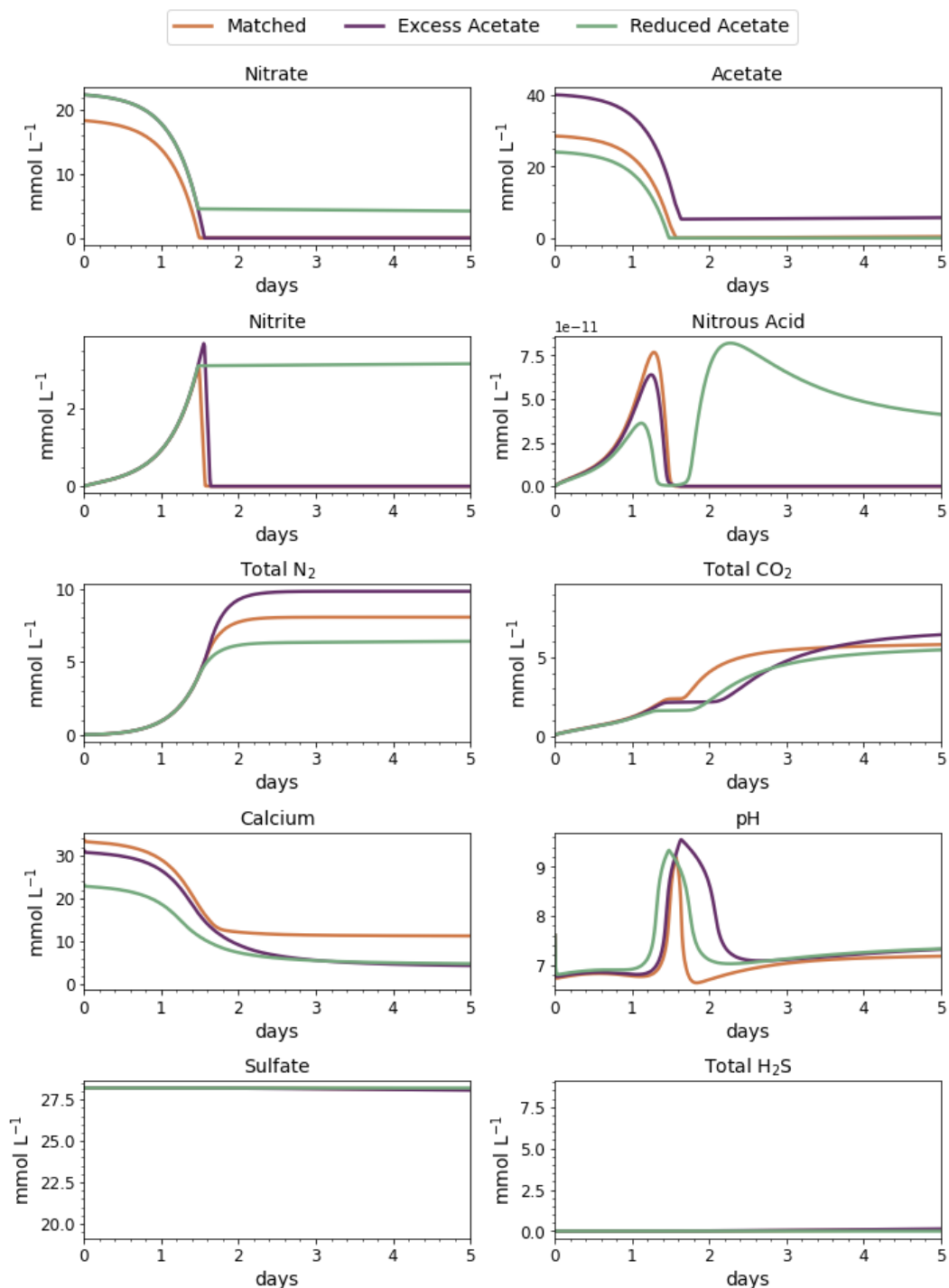
 With the excess-acetate recipe, all the nitrate was completely reduced, although small, transient accumulations of
nitrite and nitrous acid occurred (Figure 3). As expected, not all the acetate was consumed with the excess-acetate recipe, and
the remaining acetate led to sulfate reduction and the highest amount of produced H₂S of the three modeled scenarios (Figure
2). Additional N₂ was produced because of the higher amounts of nitrate in the treatment recipe, leading to a level of
285 desaturation at 7.6 m that exceeded the target 10% value.

 For the reduced-acetate test, approximately 20% of the input nitrate remained after all the acetate was consumed, and
this residual nitrate was slowly utilized beyond 1.5 days through biomass endogenous decay (Figure 2). The peak amount of
accumulated nitrite was not as high as the other conditions because of the overall limited nitrate reduction, but some nitrite
accumulation remained throughout the modeled 28 days due to the lack of acetate. The dip and quick increase in nitrous acid
290 around 1.5 to 1.7 days (Figure 3) was due to the shift in electron donor from input acetate to bacterial decay. Not enough
electron donor was available to reduce all of the input nitrate and the accumulated nitrite after 28 days (Figure 2), even though
bacterial decay caused added denitrification. N₂ produced in the reduced-acetate condition did not meet the 10% desaturation
threshold at 7.6 m at the end of the 28-day modeled period.



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Figure 2. Water-quality results for 28 days of MIDP in coastal seawater conditions targeting a desaturation level of 10% in three different treatment recipes: empirically matched, 25% excess acetate, and 25% reduced acetate.

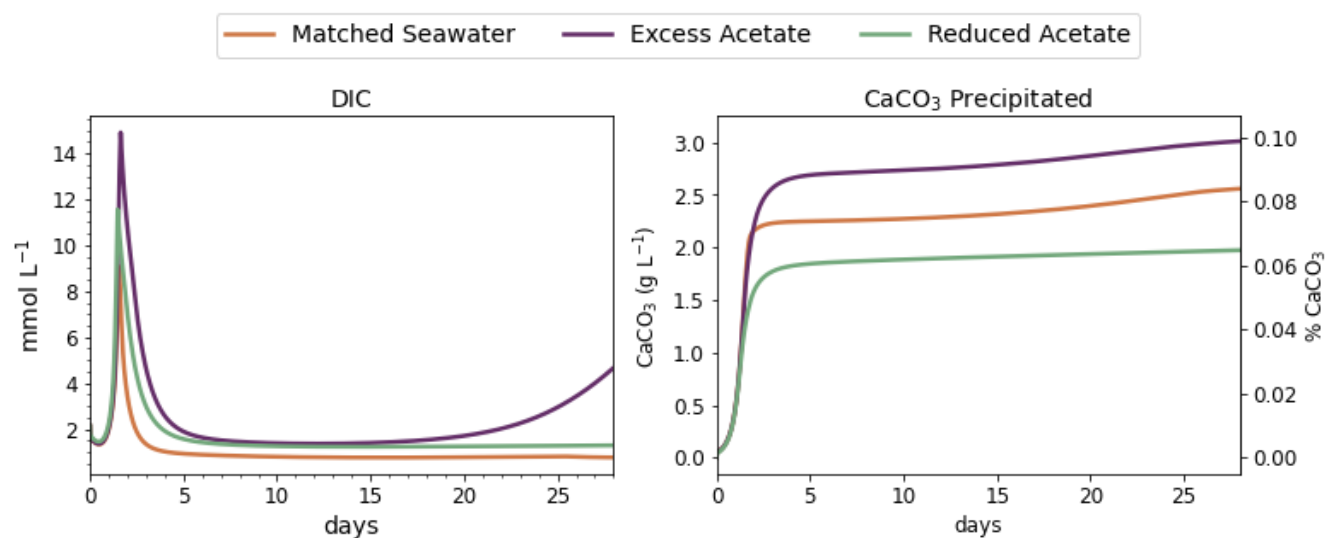


300 **Figure 3. Water-quality results for the first 3 days of MIDP in coastal seawater conditions targeting a desaturation level of 10% in three different treatment recipes: empirically matched, 25% excess acetate, and 25% reduced acetate.**



Nitrite reduction produced most of the base, which is demonstrated by the spike in pH between 1.3 and 1.8 days (Figure 3), when the rate of nitrite reduction was at its maximum in each treatment. In the matched case, the pH returned to circumneutral after 1.6 days due to the precipitation of CaCO_3 , which consumes base. This trend is reinforced by rapid Ca^{2+} consumption in the early treatment time period (Figure 3), along with production of CaCO_3 in the first ~ 2 days, shown in the right panel of Figure 4. DIC production in the excess-acetate treatment lagged the matched recipe and was slightly quicker in the reduced-acetate case, which also is seen with the pH trends in Figure 2 and Figure 3. However, the overall consumption of DIC and subsequent precipitation were slower in the excess- and reduced-acetate treatments, which correspond to the longer time for the pH to reach approximately neutral levels. The additional CaCO_3 precipitated with the excess acetate resulted from the excess of input calcium, since acetate was added as calcium acetate.

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Figure 4. DIC concentration and CaCO_3 precipitated during the first 3 days of MIDP in coastal seawater conditions targeting a desaturation level of 10%.

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Microbial decay coupled to sulfate reduction also produced more DIC (left panel of Fig. 4), promoting additional CaCO_3 precipitation after the completion of denitrification (after day 2 in Figure 4). In the excess-acetate treatment, an increase in DIC after 20 days was due to the increased rate of sulfate reduction because of the additional electron donor (Figure 2).

In summary, the simulations show that the matched-acetate recipe optimized MIDP treatment targeted at desaturation for coastal seawater conditions by maximizing the desired outcome (i.e., N_2 production for desaturation) while minimizing undesired by-products (e.g., nitrite and nitrous-oxide accumulation from incomplete denitrification, residual acetate, and H_2S from sulfate reduction).



320 **5 Conclusion**

The next-generation biogeochemical model expanded previous biogeochemical models for MIDP by considering microbial stoichiometry and kinetics for two steps of denitrification and for sulfate reduction. The model also includes gas-liquid mass-transfer kinetics for N_2 and CO_2 , $CaCO_3$ precipitation kinetics, microbial competition, and inhibition by HNO_2 , salinity, and sulfide. Model simulations demonstrated that adding nitrate and acetate using a properly matched recipe led to rapid desaturation without causing unwanted outcomes: incomplete desaturation and accumulations of nitrite and nitrous oxide with too-little acetate, or residual acetate and accelerated H_2S generation with excess acetate. The model can be used to optimize treatment recipes for maximizing desaturation or precipitation in most subsurface groundwater environments for liquefaction mitigation. However, field data describing the environmental biogeochemical characteristics (e.g., pH, background chemical concentrations) for the most optimized results is necessary to understand the potential biogeochemical reactions and processes that may impact MIDP, and subsequently, liquefaction mitigation.

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335 **Author Contribution**

CH and AvT co-developed the model. CH drafted the first version of this paper with input from AvT, LvP, EK, and BR.

Code Availability

The next-generation model was constructed in Matlab and the code and necessary files are publicly available online at doi.org/10.5281/zenodo.7410676 (CC 4.0).

340 **Competing Interests**

The authors have no competing interests to declare.



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