An emulation-based approach for interrogating reactive transport models

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8	Abstract
9	We present an emulation-based approach to understand the interactions among different chemical
10	and biological processes modelled in environmental reactive transport models (RTMs) and explore
11	how the parameterisation of these processes influences the results of multi-component RTMs. We
12	utilize a previously published RTM consisting of 20 primary species, 20 secondary complexes, 17
13	mineral reactions and 2 biologically-mediated reactions which describes bio-stimulation using
14	sediment from a contaminated aquifer. We choose a subset of the input parameters to vary over a
15	range of values. The result is the construction of a new dataset that describes the model behaviour
16	over a range of environmental conditions. Using this dataset to train a statistical model creates an
17	emulator of the underlying RTM. This is a condensed representation of the original RTM that
18	facilitates rapid exploration of a broad range of environmental conditions and sensitivities. As an

19 illustration of this approach, we use the emulator to explore how varying the boundary conditions in 20 the RTM describing the aquifer impacts the rates and volumes of mineral precipitation. A key result 21 of this work is the recognition of an unanticipated dependency of pyrite precipitation on pCO₂ in the 22 injection fluid due to the stoichiometry of the microbially-mediated sulphatesulfate reduction 23 reaction. This complex relationship was made apparent by the emulator, while the underlying RTM 24 was not specifically constructed to create such a feedback. We argue that this emulation approach to 25 sensitivity analysis for RTMs may be useful in discovering such new coupled sensitives in 26 geochemical systems and for designing experiments to optimise environmental remediation. Finally, 27 we demonstrate that this approach can maximise specific mineral precipitation or dissolution 28 reactions by using the emulator to find local maxima, which can be widely applied in environmental 29 systems.

30 Synopsis

This study explores key factors regulating mineralization reactions in near surface environments
revealed by a machine learning approach to reactive transport modelling.

33 1 Introduction

34 Reactive transport modelling has been extensively applied across a wide variety of environmental

35 systems, providing a powerful means of quantifying, and even predicting, processes across Earth's

36 (near) surface environments (Richter and DePaolo, 1987; Bain et al., 2000; Johnson et al., 2004; van

- 37 Breukelen et al., 2004; Gaus et al., 2005; Torres et al., 2015; Li et al., 2017; Arora et al., 2020;
- 38 Molins and Knabner, 2020; Rolle and Borgne, 2020; Druhan et al., 2020; Cama et al., 2020).

39 Reactive transport models (RTMs) are constructed by combining multiple physical, chemical, and

40 biological processes to simulate the behaviour of environmental systems. As applications and

41 software have concurrently expanded (Steefel et al., 2015; Li et al. 2017; Maher & Mayer, 2019;

Druhan & Tournassatt, 2019), it is becoming increasingly common to explicitly calculate the rates of production and consumption for a variety of coexisting chemical species, as well as their equilibria with mineral phases, and their transport as they evolve in time and space. This type of multi-phase, multi-component RTM is a type of forward modelling where the results of the simulation emerge from a complex suite of interacting pathways, and hence the causes of observed behaviour are not always obvious.

- 48 <u>RTMs are often designed to describe the behaviour of specific field sites and systems. Due to their</u>
- 49 process-based nature, designing RTMs requires selection of a suite of chemical reactions and

50 transport mechanisms which are thought to dominate the geochemistry of the system over the scales

51 of interest. However, the parameterisation of various selected processes is often not unique and can

52 impact system behaviour (Williams et al., 2011; Martinez et al., 2014; Seigneur et al., 2021; Steefel

53 et al., 2005a). To assess the impact of the choice of parameterisation and the values chosen for

54 different parameters on model predictions, sensitivity analyses are generally performed (Malaguerra

55 <u>et al., 2013; Gatel et al., 2019). However, as RTMs become increasingly sophisticated, they</u>

56 incorporate disparate processes that can interact with each other in complex ways (Dwivedi et al.,

57 <u>2018; Hubbard et al., 2018, 2019; Maavara et al., 2021a, b; Dwivedi et al., 2017).</u>

58 The sensitivity analysis of an RTM in application to a specific environmental system can elucidate

59 the relative importance of specific interactions; for example, testing the solubility of mineral phases

60 relative to changes in the solution chemistry. However, results might emerge that were not

61 anticipated. These results might represent a real, but unexpected, interactions in which case the

62 sensitivity analysis has yielded new insight into the system being modelled. Equally, the result might

63 represent an incorrect interaction between two different processes that are known to act

64 independently of each other, in which case the RTM can be improved. Unfortunately, due to the

65 computational expense of many modern multi-component RTMs (e.g. Abd and Abushaikha, 2021;

66	Seigneur et al., 2021; Gharasoo et al., 2022), it is normally impractical to perform sensitivity analyses
67	in more than a few dimensions and it is up to the investigator to use their knowledge of the system to
68	choose which sensitivity analyses are necessary to explore (Steefel et al., 2005b). Ideally, we would
69	be able to systematically perform sensitivity analyses over many model parameters, considering how
70	model outputs vary as a function of multiple input parameters simultaneously (i.e. in a multivariate
71	way), while also lightening the computational burden that commonly occurs when using inverse
72	modelling approaches implemented by codes like PEST and iTOUGH2 (Doherty, 2004; Finsterle et
73	al., 2017). Such a capacity could direct future laboratory-based investigations to test whether these
74	model results are real-world phenomena, ultimately offering improved parameterisation of critical
75	components within the reaction network.
76	Here, we demonstrate a method for exploring a wide variety of potential model parameters by
77	adopting an emulator-based approach. Ours is not the first work to apply emulators to RTM
78	simulations. Notably, a rich vein of research based around replacing the geochemical solver in RTMs
79	with an emulator has emerged over the past few years (see Laloy and Jacques (2021) and Kyas et al.
80	(2022), among others). However, the work presented here is less concerned with speeding up
81	individual RTM simulations as it is with developing new methods to explore geochemical parameter
82	spaces. We also investigate the effect of changing geochemical parameters on the overall outcome of
83	RTM simulations, with an eye towards predicting system outcomes in real world scenarios. This is
84	similar in nature to recent work conducted by Ahmmed et al. (2021), which explores the ability of
85	different machine learning methods to predict the degree of mixing and the progression of a
86	simplified, generic reaction $(A + B \rightarrow C)$ in a finite element simulation, and we extend the idea of
87	predicting the final state of a simulation to published RTMs describing real world systems.
88	Such emulation approaches in predicting the outcomes of physical systems have a long history
80	including applications in physics-based animation (Grzeszczuk et al. 1008) complex multiphysics
07	menumg appreations in physics-based anniation (Orzeszezuk et al., 1996),complex multi-physics

90	simulators (Lu et al., 2021	; Bianchi et al., 2016), climate models ((Beucler et al., 2019	; Krasno	<u>polsk</u>	y
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- 91 et al., 2005; Castruccio et al., 2014; Kashinath et al., 2021), and emulating fluid flow through
- 92 Dolomite using a neural network (Li et al., 2022). In an emulator approach, the underlying physical
- 93 system is approximated by a statistical model (the emulator) which can be evaluated more quickly
- 94 than a conventional forward model. How this emulator is constructed varies by implementation and
- 95 may encode assumptions about the underlying system to be modelled (e.g. conservation of energy
- 96 (Beucler et al., 2019)). In this study, we are primarily interested in exploring and emulating the
- 97 geochemical behaviour of RTMs; therefore we focus less on transport effects and restrict ourselves to
- 98 <u>emulating one dimensional RTMs.</u>
- 99 In our implementation the emulator is built by training a Gradient Boosted Trees (GBT) regressor
- 100 (Chen and He, 2015) on a synthetic dataset generated from the original RTM. By training such a
- 101 <u>GBT model on the synthetic dataset generated by the original RTM, we create an emulator of the</u>
- 102 original system. This emulation approach is general and can be applied to a wide range of RTMs,
- 103 <u>using "off the shelf" statistical libraries, requiring no special construction of the statistical model</u>
- 104 beyond the choice of some training parameters. This approach can identify the critical processes and
- 105 parameters within RTMs and address the requirement for comprehensive, multivariate sensitivity
- 106 <u>analyses.</u>
- 107 We first present a tool that automates creation of synthetic datasets: a Python wrapper for the RTM
- 108 software CrunchTope (Druhan et al., 2013; Steefel et al., 2015), which we have named Omphalos.
- 109 Omphalos edits and runs CrunchTope input files in an automated fashion, systematically changing
- 110 model parameters according to user specification. It then records the output data, along with the
- 111 corresponding model input parameters for later analysis. We then apply a machine learning method
- 112 (Gradient Boosted Trees) to these recorded inputs and outputs to create a predicative model that can

113 reproduce RTM outputs based on the input variables, which we term a Reactivate Transport
114 Emulator (RTE).

115 We suggest that our contribution to the development of Reactive Transport Emulators could be used 116 to direct new experimental investigation to identify and corroborate predicted dependencies; 117 providing much-needed multivariate analysis of RTMs and helping to identify effects that can, in the 118 future, be considered explicitly when developing new RTMs. In pursuit of this goal, we demonstrate 119 our emulator approach in application to an RTM built for biostimulation of a contaminated aquifer. 120 We also show an additional application of this approach to efficiently predict the condition which 121 maximises an RTM-predicted time-integrated rate over the set of chosen parameters. We also 122 present, in the Supporting InformationSupplement, another example in application to a deep-sea

123 sediment column.

124 **2** Description of the Case Study

125 2.1 Old Rifle Site, Colorado

126 The Old Rifle site is located near Rifle, Colorado, USA. The location historically hosted a vanadium 127 and uranium ore processing facility, and the groundwater at the site remains high in aqueous 128 uranium. Oxidised uranium (U(VI)) is fluid-mobile and highly toxic, while reduced uranium (U(IV)) 129 is much less soluble and forms stable precipitates such as uraninite (UO₂) (Anderson et al., 2003; Wu 130 et al., 2006; Dullies et al., 2010; Williams et al., 2011; Long et al., 2015)(Anderson et al., 2003; Wu et al., 2006; Dullies et al., 2010; Williams et al., 2011; Long et al., 2015). Thus, uranium reduction 131 132 has been suggested as a means for remediating uranium contamination in groundwater. It has been 133 shown that iron sulfide minerals (FeS_{2(s)}) aid the reduction of soluble U(VI) to insoluble U(IV) 134 precipitates even after active remediation has ceased (Komlos et al., 2008; Moon et al., 2010; Bargar 135 et al., 2013; Long et al., 2015; Bone et al., 2017).

136 The RTM published for Old Rifle, upon which the RTE is based, was originally created as a 137 comprehensive model of microbial sulfate reduction and sulfide precipitation in Old Rifle sediment 138 during stimulation of microbial activity by amendment with $C_2H_3O_2^-$ (Druhan et al., 2014) (for a 139 schematic illustration of this RTM, see Fig. $\frac{8283}{10}$. In this context, we choose to vary the influent 140 boundary condition chemistry, representing changes to the chemical composition of the artificial 141 groundwater injectate. The original experiment was designed to model microbial sulfate and iron reduction in the sediment; therefore, we use net amorphous iron (II) sulfide (FeS_(am)), and pyrite 142 (FeS_{2(s)}) precipitation (both hereafter referred to simply as 'pyrite') as an observable that will record 143 144 the sensitivities of the model predictions to changes in the injection fluid. We also demonstrate the 145 utility of the emulator in predicting the chemical composition of the injection fluid that will 146 maximise the volume of pyrite precipitated in the sediments when amended with a labile organic 147 carbon source via injection wells.

148 **3** Methodology

149 **3.1 General Strategy**

150 To explore the dependence of the RTM on the chosen environmental variables, we begin with a 151 Monte Carlo approach; we draw random values for each parameter and record the model output 152 under that randomised condition. We then fit a model to this Monte-Carlo-generated dataset using a 153 GBT regressor. This fitting results in a model (our emulator—RTE) that reproduces the complex 154 interdependencies of chemical species that are encoded in the original, underlying, RTM. This 155 emulator can be interrogated to examine the dependence of the RTM outputs on the originally chosen 156 environmental variables in an efficient, multivariate way. This new way of performing sensitivity 157 analyses has the potential to give insight into trends and relationships that would not be apparent 158 otherwise and ultimately allows us to investigate the sensitivity of the model outputs with respect to

- 159 the RTM's original parameterisation. First, we will describe how we use the Monte Carlo approach
- 160 to generate data and then how we fit a model to this data. The overall workflow is shown in Figure
- 161 <u>1</u>Fig. 1.





- Figure 1: Flowchart describing the overall reactive transport emulation workflow developed in 163
- 164 this study. It is divided into two key sections: preparation of the input reactive transport model for submission to Omphalos, and the analysis and emulation of the resultant data. 165

166 3.2 **Generating Data**

167 We use the open-source software CrunchTope as the reactive transport framework for the models in

168 this study. To generate the synthetic datasets necessary for our approach, and given the time169 consuming nature of generating a single point (requiring a complete run of the RTM, along with 170 modified boundary conditions), we developed a software package in Python to automate this process. 171 This software package can manage the automatic generation and submission of unique input files to 172 CrunchTope, as well as recording the output of each run, storing it in a manageable data structure for 173 future use. Use of the software package is straight-forward, requiring the configuration of a single file 174 listing which species/parameters are to be varied, and how they should be varied.

175 We have named this software package Omphalos (available for download—Sect. <u>6.1</u>5.1). Omphalos

176 can be run on clusters using Simple Linux Utility for Resource Management (Yoo et al., 2003)(Yoo

177 <u>et al., 2003</u>) to execute input files in parallel, <u>or run locally with CrunchTope simulations on</u>

178 <u>individual CPUs</u>, which considerably reduces the time required to generate large datasets. Omphalos

179 works by taking random values which are drawn from uniform distributions (other statistical

180 distributions are possible) of the chosen variables, sampling the space evenly. This provides a

181 complete dataset for training the emulator.

182 While the underlying principle of training emulators on synthetic data can be applied to any reactive

183 transport code, currently the software used to implement the approach is only compatible with

184 CrunchTope, because the input file reading and writing must be in a specific format. The approach is

readily generalized, however, and the methodology could be applied to any RTM software (e.g.

186 Geochemist's Workbench, <u>TOUGHREACT</u>), provided that the string input/output code is adapted

187 for compatibility. To use other RTMs with Omphalos, two key factors need to be addressed:

188 compatibility with Omphalos, and the computational expense of a single RTM run.

189 **3.3** Application to Contaminated Aquifer Case Study

190 We begin by applying the emulation methodology to our case study. To create the dataset for training

191 the emulator, we collected the results of 10,927 unique CrunchTope simulations based on the original

192 RTM describing Old Rifle using Omphalos, drawing random concentrations for each species five chosen species (NH_4^+ , SO_4^{2-} , Ca^{2+} , $C_2H_3O_2^-$, and pCO₂) in the boundary condition. Of these 10,927 193 194 runs, 9416 provide useable data because some runs fail to converge within the specified timeframe, 195 or the geochemical condition generated cannot be charge balanced. Excluding these runs helps ensure 196 that our dataset is kept realistic, because our RTM is built on a mechanistic understanding and 197 implementation of the physical processes at work in the system that have been validated in some 198 way. Therefore, runs that take excessively long to run are failing to converge in the simulation 199 scheme of CrunchTope, and hence likely to be unphysical in some way. Similarly, runs that fail to 200 speciate or charge balance indicate some kind of extreme physical condition that is unlikely to be realistic and so are excluded. The concentrations for NH₄⁺, SO₄²⁻, Ca²⁺, and C₂H₃O₂⁻ are varied 201 202 between 0–30 mM. The pCO₂ is varied between 0–10 bar. We acknowledge that these ranges of 203 concentrations are somewhat higher than those that occur in natural systems, but we extend the range 204 to observe RTM behaviours at limiting concentrations. Related to this, it is possible for the dominant 205 reaction mechanism in a system to change under differing conditions (e.g. the change in calcite 206 dissolution mechanism as a function of pH (Dolgaleva et al., 2005)) and any such behaviour should 207 be explicitly encoded into the RTM, otherwise the emulator may give invalid predictions under 208 conditions that are far from the original model run. We have assumed in this study that the 209 mechanisms governing the precipitation of pyrite do not change under very low or very high 210 concentrations of these species.

The injection fluid was constrained at pH 7.2. This constraint, in conjunction with the concentration of various species iterated in Omphalos, speciates according to CrunchTope's internal speciation calculation. Therefore, for example, although the total amount of SO_4^{2-} in the injection will be iterated in, and dictated by, Omphalos, the amount that speciates into other aqueous complexes (i.e.

10

secondary species) like HSO_4^- or $H_2SO_{4(aq)}$ is controlled by CrunchTope. For the sake of simplicity, we will report the input concentration, not the concentration after speciation.

217 The RTM describing Old Rifle has 100 grid cells with a size of 1 cm. Each run of the RTM took 218 approximately 90 seconds, so the total time to generate the dataset was roughly four hours when run 219 on a remote machine with 200 CPUs. The number of runs was chosen as a balance between what was 220 computationally tractable and the ability of the emulator to achieve a good fit. We have intentionally 221 chosen to vary some chemical species in the influent boundary condition that do not play an obvious 222 role in the mineral precipitation process we are particularly interested in, namely, the precipitation of pyrite in Old Rifle sediments (e.g., NH₄⁺ or Ca²⁺, respectively). We did this to see if we can use the 223 224 emulator to detect behaviour in the RTM beyond what we might initially hypothesise.

225 **3.4** Fitting the emulator

We implement the GBT regressor using XGBoost (Chen and He, 2015) in Python. The code for
fitting the models is available in the Supplement. For a precision GBT models, see the supplement
Sect. S1.2.

229 **3.4.1 Data Strategy**

230 Data generated by Omphalos was imported into a Jupyter notebook environment from the .pkl output 231 file. There are 9416 different input file runs in this data file, having excluded 1511 runs on the grounds of them being unrealistic, as discussed previously. The relevant data was indexed out of the 232 data structure; in our case this meant the concentrations of NH_4^+ , SO_4^{2-} , Ca^{2+} , Ca^{2+} , and $C_2H_3O_2^-$ in 233 234 the boundary condition, as well as value of pCO₂. This results in a 5×9416 array of floating-point 235 numbers for the features. Each feature was then normalised to be in the range 0 to 1 for training. For example, values of SO_4^{2-} concentration in the simulations were drawn randomly between 0 and 30 236 mM, so all SO_4^{2-} concentrations were divided through by 30 to have values in the range 0–1. We did 237

this to improve the training performance of the GBT model over different datasets (i.e. so that the
same GBT model can be applied to both the Old Rifle case study, and our supplementary case study
of ODP Site 1086 (see Supplement, Sect. 3).

Similarly, the relevant data was also extracted from the data file: for each cell in the gridded RTM, we calculated the net pyrite precipitation over the course of the simulation, and then summed this value over the column to get the net pyrite precipitated across the domain. This results in a 1×9416 array of floating-point labels to be predicted from the feature array. We scale this feature array by a factor of 1×10^4 to avoid issues with small floating-point numbers in XGBoost.

We prepared these data for training the GBT regressor with a hold-out strategy using the scikitlearn.train_test_split method, keeping 10% of the dataset back for validating the model. Data was split randomly within the dataset. This means that 8474 randomly selected data points were used to train the model and 942 randomly selected data points were used to test it by using the model to predict a value based on the held back data and comparing the prediction to the true value.

251 **3.4.2 Training Strategy**

We use the test set of data points generated by Omphalos to train an XGBoost regressor using squared error as the loss function to predict the amount of pyrite precipitated in the column as a function of varied species concentrations in the boundary condition. Squared log loss, and pseudo-Huber error wewere also tried but squared loss performed best overall. Training curves showing the testing and training loss as training progressions are given the supplement, Fig. S2.

-Hyperparameter choices for the model are explained and given in the supplement, Sect. S1.3, Table
S1. The choice of hyperparameters is the same for each emulator model, and we are able to achieve
high quality fits using the default XGBoost regularisations, only changing a few settings relating to
tree growth policy. While it is a known problem in machine learning that the choice of optimal

261 hyperparameter is dependent on the data being modelled (Claesen and De Moor, 2015), it appears

that in the context of these RTEs, the hyperparameters chosen give a good fit for both Old Rifle and

263 our supplementary case study of ODP Site 1086: datasets describing very different natural

264 environments, with different length and time scales. This makes the workflow applicable across a

- 265 wide variety of reactive transport modelling domains.
- 266 It is possible that with more complex hyperparameter tuning, better emulator fits may be achieved,

but for the purposes outline in this paper, we suggest that this automated optimisation of a subset of

268 the available hyperparameters is sufficient, and represents a balance between emulator fit,

269 generalisability across differing RTMs, and time spent by the user.

270 3.4.3 Model metrics

271 We report our model goodness-of-fit to the underlying dataset as the R² value for the model, using

272 the built in XGBRegressor.score() method from XGBoosts Scikit-Learn API, on both the original

273 training dataset, as well as the 10% of the dataset held back for validation, show in Table 1. We also

274 report the root-mean-square error (RMSE) over both the training and validation datasets, calculated

275 <u>using the Booster.eval() method. Training curves are shown in the supplementary, Figure S2. We</u>

276 report the same metrics for our second model in the supplementary, Table S4.

Dataset	<u>R²</u>	<u>RMSE</u>
Training	<u>0.99996</u>	<u>1.95012</u>
Validation	<u>0.99964</u>	<u>5.65848</u>

277 <u>Table 1: Training and validation metrics for the XGBoost regressor model fit to the Old Rifle</u>

278 dataset. R² represent a normalized measure of the fit quality, with the best possible score being

279 **<u>1. RMSE is the root-mean-squared error in the predictions, where we recall that in data pre-</u>**

processing that the values to be predicted were multiplied by a factor of 10⁴, and so the RMSE
 should be divided by that factor when assessing the average error on data presented in Figures
 2, 3, and 4.

283 4 Results and discussion

284 **4.1** Application to the Old Rifle Site

285 The synthetic data generated using Omphalos to interrogate the underlying RTM are shown in Fig. 2, 286 colour mapped by the pCO₂ with which the injectate solution is in equilibrium. The colour mapping 287 helps visualise how variability in the precipitated volume of pyrite over the 43-day RTM simulation 288 might be considered in conjunction with other model parameters. Ultimately, pyrite forms because 289 aqueous hydrogen sulfide, produced through microbial sulfate reduction, reacts with reduced ferrous 290 iron (Fe(II)) to form pyrite. Thus, we aim to explore the interdependencies between the mechanisms 291 driving microbial sulfate reduction and the subsequent precipitation of pyrite, as they emerge due to 292 variations in injectate chemical composition.



Figure 2: Scatter plots of chemical concentrations in the fluid injectate (influent boundary condition) for an RTM adapted to Old Rifle sediments colour-mapped by the pCO₂ with which the inlet boundary condition is in equilibrium. The dataset comprises 9416 points generated by drawing concentrations for all five species independently from uniform random distributions, with the corresponding net increase in pyrite volume fraction precipitated (y-axis) calculated by running the Old Rifle RTM designed by Druhan et al. (2014) with the randomised influent

boundary condition.(2014) with the randomised influent boundary condition. The green diamond indicates the net pyrite volume fraction generated from the original boundary condition used in Druhan et al. (2014).

303 We then train the emulator on this synthetic dataset. Fitting a GBT regressor to the data in Fig. 2 304 means Fig. 3 can be generated by the emulator. This figure shows how the emulator predicts the 305 change in pyrite volume fraction as the concentration of each of the species in the injection fluid is 306 varied (other species in the RTM not defined as variables in this study are held constant at values reported by Druhan et al. (2014)). The convergence of the emulator is shown in Fig. S3.(2014)). We 807 308 stress that the RTM results shown in Fig. 3 are not part of the training dataset, and that the emulator 309 has not been exposed to these exact values. This demonstrates the capability of the emulator to 310 reproduce the underlying RTM itself. For example, Fig. 2A suggests visually that the concentration 311 of NH₄⁺ in the system is uncorrelated with net pyrite precipitation at the Old Rifle Site. Fig. 3A 812 confirms this lack of dependence on NH₄⁺, capturing the correct trend (with some noise) although 313 being slightly offset. This slight offset also applies to Fig. 3C in the fit of the Ca²⁺ dependence. We 314 suggest that these slight offsets to the fits in the cases of the weakly or uncorrelated variables is due 315 to the emulator preferentially capturing stronger dependencies and slightly drawing down the 316 predicated variable on average.

In contrast to the minimal impact that changing NH_4^+ concentration has on pyrite precipitation,

318 $C_2H_3O_2^-$ and SO_4^{2-} concentrations correlate strongly with net pyrite precipitation. This is as expected

319 in a system where $C_2H_3O_2^-$, which is the electron donor for microbial sulfate reduction, enables

- 320 sulfate to be reduced to sulfide and thus drive pyrite precipitation in the presence of Fe(II).
- 321 Approximately 20 days after $C_2H_3O_2^-$ amendment, microbial sulfate reduction takes over from
- 322 dissimilatory iron reduction as the dominant process consuming $C_2H_3O_2^-$. As microbial sulfate

323	reduction requires eight-times the number of electrons per mole of SO_4^{2-} reduced than dissimilatory
324	iron reduction requires (per mole of iron reduced), the electron donor $(C_2H_3O_2^-)$ begins to be rapidly
325	consumed, whereas during dissimilatory iron reduction it was effectively in excess. As a result of this
326	new scarcity of $C_2H_3O_2^-$, the rate of dissimilatory iron reduction drops and so does the concentration
327	of Fe(II). However, dissimilatory iron reduction is still active in the column, releasing a small-but
328	non-zero—flux of aqueous Fe(II) that allows for continued pyrite precipitation. The emulator
329	interprets this as Fe(II) being 'always' available in this system, and thus predicts that pyrite
330	precipitation can scale linearly with SO_4^{2-} and $C_2H_3O_2^{-}$, as shown in Fig, 4A. The sediment itself
331	would need to contain abundant ferrihydrite, goethite, or another bioavailable ferri(hydr)oxide for
332	this reduction to continue indefinitely; this may not be the case. This highlights the need for the range
333	of parameters sampled when training the emulator to be sufficiently wide to capture all the RTM
334	behaviour, otherwise it may extrapolate and "learn" incorrect assumptions about the system: in this
335	case that bioavailable iron never limits dissimilatory iron reduction. One solution would be to expand
336	the range over which concentrations are drawn to reach the limit where iron-bearing mineral volume
337	fraction becomes a limiting factor so that the model can learn what happens when this occurs.



339 Figure 3: Plots of the GBT model fit (blue line) plotted over the results from the underlying 340 RTM (black + symbols) when interrogated with the same input parameters (which are taken as 341 ground truth). Each plot shows the net volume fraction due to pyrite precipitation as a 342 percentage of the initial volume fraction of the sediment as each parameter is varied while all 343 other parameters are held at the values used in the original experiment by Druhan et al. (2014). 344 The emulator (blue line) captures the overall trends in the data. The lack of smoothness in the 345 emulator predications arises from the inability to encode this as a condition in XGBoost and 346 the discreet nature of the decision tree algorithm. The RTM results compared to here are not 847 part of the training dataset, and so the emulator has not been exposed to those exact values. 348 We also note that our emulator suggests that increasing pCO₂ leads to decreased pyrite precipitation

349 (<u>Figure 4Fig. 3</u>E), a relationship that may not have been apparent in a single run of the RTM. Three-

dimensional visualisation of the data confirms that the pyrite-volume-fraction-change varies as a

function of pCO₂ net pyrite precipitated decreasing as pCO₂ increases (Fig. 4B and Fig. 4C). This

352 three-dimensional visualisation allows us to see that the gradient of the pyrite-volume-fraction-

change with respect to SO_4^{2-} and $C_2H_3O_2^{-}$ is itself a function of pCO₂ and flattens as pCO₂ increases.

To understand why the gradient changes, we must first understand why pCO_2 affects the amount of pyrite precipitated in the first place.

356 Sediment samples from Old Rifle are initially poised for dissimilatory iron reduction and there is a 357 sizeable community of iron-reducing bacteria naturally present in the system. The background 358 sulfate-reducing microbial community is initially relatively small and thus, for microbial sulfate 359 reduction to proceed at significant rates, the mass of sulfate-reducing bacteria must first increase. In 360 the original experiment by Druhan et al. (2014), the sulfate-reducing biomass begins reaching a size 361 where it can start consuming large quantities of $C_2H_3O_2^-$ around day 20 of the experiment. This 362 biomass growth is modelled in CrunchTope using a Monod-biomass rate law (Jin and Bethke, 2005), 363 which has both an anabolic and catabolic component. In the formulation of this Monod-Biomass rate 364 law as implemented in CrunchTope, the thermodynamic term (Gibbs free energy of the reaction) is 365 calculated exclusively using the catabolic pathway. The catabolic pathway for this reaction (in terms 366 of the exchange of one electron) is given below in Equation (4.1), and the form of the Gibbs free 367 energy is this context is given in Equation (4.2) (we take the phosphorylation potential to be 0, and 368 the average stoichiometric number to be 1, see derivation in Jin and Bethke (2005) for further 369 details).

370
$$\frac{1}{8}SO_4^{2-} + \frac{1}{8}C_2H_3O_2^{-} + \frac{3}{8}H^+ \rightarrow \frac{1}{8}H_2S_{(aq)} + \frac{1}{4}CO_{2(aq)} + \frac{1}{4}H_2O$$
 (4.1)

371
$$\Delta G = \mathcal{RT} \ln \left(\frac{\left[CO_{2(aq)} \right]^{\frac{1}{4}} \left[H_2 S_{(aq)} \right]^{\frac{1}{8}}}{\left[SO_4^{2-1} \right]^{\frac{1}{8}} \left[C_2 H_3 O_2^{-1} \right]^{\frac{1}{8}} \left[H^+ \right]^{\frac{3}{8}}} \right)$$
(4.2)

Taking this form for the Gibbs free energy of the reaction and substituting it into the thermodynamic
term of the reaction rate calculation as implemented in CrunchTope (Steefel et al., 2015) gives

Equation (4.3) below describing the rate of microbial sulfate reduction in the Rifle RTM.

375
$$R_{MB} = k_{max} B \frac{[C_2H_3O_2^-]}{[C_2H_3O_2^-] + K_{half[Ace]}} \frac{[SO_4^{2-}]}{[SO_4^{2-}] + K_{half[SO_4^{2-}]}} F_T$$
(4.3)

376 where

377
$$\boldsymbol{F_T} = \left(\mathbf{1} - \frac{\left[\text{CO}_{2(\text{aq})} \right]^{\frac{1}{4}} \left[\text{H}_2 \text{S}_{(\text{aq})} \right]^{\frac{1}{8}}}{\left[\text{SO}_4^{2-1} \right]^{\frac{1}{8}} \left[\text{C}_2 \text{H}_3 \text{O}_2^{-1} \right]^{\frac{1}{8}} \left[\text{H}^+ \right]^{\frac{3}{8}}} \right)$$
(4.4)

 R_{MB} is the overall rate of microbial sulfate reduction, k_{max} the rate constant for microbial sulfate 378 reduction, B is the biomass concentration, and $K_{half[X]}$ is a half-saturation constant. The two Monod 379 kinetic factors for the electron donor ($C_2H_3O_2^-$) and the electron acceptor (SO_4^{2-}) are referred to as F_D 380 381 and F_A respectively (Jin and Bethke, 2003, 2005, 2007). Equation (4.4) illustrates the underlying 382 relationship between pCO₂ in the injectate solution and the resulting accumulation of pyrite. As the 383 pCO₂ of in the injectate increases, the F_T term becomes smaller, inhibiting the overall rate of 384 microbial sulfate reduction (Fig. S5). Consequently, biomass growth is also inhibited, and the rate of microbial sulfate reduction is never high enough to produce the concentration of $H_2S_{(aq)}$ required for 385 significant pyrite precipitation. This explains why the model suggests that the gradient of the pyrite 386 volume precipitated with respect to both $C_2H_3O_2^-$ and SO_4^{2-} varies as a function of pCO₂ in the 387

injectate. When pCO₂ is low and both SO_4^{2-} and $C_2H_3O_2^{-}$ are large with respect to their half



389 saturation constants (Equation (4.4)), the overall Monod-biomass rate law will approach Bk_{max} .

Figure 4: A selection of the GBT model predictions of the percentage volume fraction increase
 due to pyrite precipitation as a result of varying two parameters simultaneously for selected

- 393 pairs of variables. Other model parameters are held at the values used in Druhan et al. (2014).
- 394 The remaining variable-pair plots are provided in Fig. S4.
- 395 This dependence emerged somewhat unexpectedly from the emulator, as one would not inherently
- 396 expect a relationship between injectate pCO_2 and SO_4^{2-} reduction rates, yet it agrees with results
- 397 previously reported by Jin and Kirk (2016, 2018) as well as Paper et al. (2021). These studies related
- 398 the influence of pCO₂ and pH to the rate of microbial reactions, both *in vitro*, *in situ*, and *in silico*.
- 399 We suggest that our type of analysis could be used to direct future lab and field work to test
- 400 <u>hypotheses suggested by the results generated by running the emulator.</u>
- 401 This analysis also explains some of the features observed in Fig. 4A: the gradients of $C_2H_3O_2^-$ and
- 402 SO_4^{2-} are coupled in such a way as to indicate that if one is in excess, then the other becomes limiting
- 403 in the production of $H_2S_{(aq)}$ and hence the precipitation of pyrite. However, the limiting behaviour
- 404 when both are in excess seems to indicate that given enough SO_4^{2-} and $C_2H_3O_2^{-}$, pyrite precipitation
- 405 can continue indefinitely assuming suitably low pCO₂. Given this prediction, it is sensible to check

406 whether, at such high levels of SO_4^{2-} and $C_2H_3O_2^{-}$ as the model suggests for maximum pyrite 407 precipitation, there is indeed enough Fe(II) available in the system to precipitate pyrite: this is a 408 second potential dependence as mentioned above.

409 Lastly, the model can be interrogated in all 5 dimensions and the amendment fluid composition that 410 corresponds to the largest net pyrite precipitation over the modelled interval can be determined. We 411 do this simple optimisation by evaluating the emulator at regular intervals across all 5 dimensions at 412 intervals of $\sim 2 \text{ mM}$ (intervals of $\sim 0.67 \text{ mM}$ for pCO₂) in the range that the emulator was trained on 413 (0-30 mM, except for pCO₂ which has a range of 0-10 mM). This corresponds to checking 759375 414 different boundary conditions to find which boundary condition results in the most net pyrite 415 precipitation and takes ~7 minutes. This amendment composition is shown in Table S1. The total 416 change in volume fraction due to pyrite precipitation predicted by the emulator is 0.143 and the 417 actual RTM modelled precipitation when this boundary condition is used is 0.150. There is a 4.7% 418 absolute error on the net pyrite volume fractions change predicted by the emulator when compared to 419 the actual net pyrite precipitation calculated by the RTM. This error is inherent in statistical learning 420 techniques but can be further mitigated with larger training datasets, in conjunction with different 421 emulator training hyperparameterisations: an area for future improvement to the methodology. These 422 optimised conditions represent an almost four-fold increase in the amount of pyrite precipitated in the 423 original RTM for Old Rifle (Druhan et al., 2014).

424 **4.2** Advantages and drawbacks of the emulation approach

In this study, 9416 individual RTM simulations were used to train a GBT regression model to predict a specific model output, in this case net pyrite precipitation. This emulator is a reduced representation of the complex system of equations in the underlying RTM, having a faster computational time but introducing some prediction errors. We now discuss the key advantages and drawbacks of this emulation approach.

430 **4.3** Advantages of the emulation approach

431 9416 RTM runs were used to train the emulator (the data shown in Fig. 2). This number of runs could 432 instead be used to perform a sensitivity analysis in all five variables at a spacing of ~4.8 mM between 433 points by directly interrogating the simulator. What then, is the advantage of the emulation approach, 434 if the same information can be visualised from discreet runs of the original RTM without having to 435 go to the extra effort to train the model, which introduces prediction errors? The key advantages are 436 outlined below.

437 **4.3.1** Advantages over directly interrogating the simulator

438 The first and most obvious advantage is the lack of a need for an explicit interpolation scheme. 439 Correlations generated by directly plotting simulator results in both test cases lead to data points 440 lying on a grid of finite resolution. If intermediate values on this grid were to be determined, an 441 explicit interpolation scheme would have to be applied, which would introduce errors of its own that 442 would then need to be quantified. Furthermore, an improvement in the interpolation scheme would 443 come at the expense of adding one extra point to the grid in each dimension: in the context of Old Rifle this is an extra 9031 data points $(7^5 - 6^5 = 9031 \text{ going from a 5D grid of 6 points in all})$ 444 445 directions to 7) roughly doubling the dataset size. In contrast, since any number of points can be 446 submitted to the emulator for inference, concerns relating dataset size to sampling resolution are 447 assuaged. Beyond that, the errors in the model fit are already quantified during training.

448 More broadly, to explore the dataspace, emulators are extremely fast compared to simulators. The 449 time for a single query of the emulator is on the order of milliseconds rather than the 450 seconds/minutes/hours for a single forward RTM simulation. This allows the emulator to be used as a 451 tool for efficiently exploring the simulator by rapidly developing intuition for the space itself and 452 how the simulator behaves in different circumstances. Furthermore, emulator models are easy to distribute and share with collaborators. Model weights can be published directly or distributed as
standalone files. This means that a well-trained emulator can be made once and then the encoded data
shared.

456 Lastly, performing a direct interrogation of the simulator requires choices of parameters and ranges, 457 and results in a grid of points over the region of interest at limited resolution. A similar procedure 458 must be undertaken when creating a dataset to train the emulator, in so far as ranges and parameters 459 of interest must be chosen. However, the dataset can always be further added to in a straightforward 460 manner, further drawing from the random distribution to increase the size of the dataset and thus 461 improve model performance. With both approaches, using Omphalos means that the data generation 462 process can be parallelised and using high-performance computing facilities can reduce the 463 computational expense of interrogating the simulator. This means that all the computational expense 464 is upfront in both cases since the emulator need only be fit once.

465 The advantages we outline make the case for the emulator as a tool to be used in conjunction with the 466 RTM, rather than a replacement for it. The alacrity with which the emulator can be interrogated 467 means that it is an invaluable tool for investigating RTM behaviour in multiple dimensions. Further 468 to this, the ability to evaluate the state of a system after a fixed period of time makes the emulator 469 approach ideally suited for modelling more complex time-series models with time varying boundary 470 conditions: instead of having to run the RTM forward each time the system changes boundary 471 conditions, the emulator can be interrogated for the expected result given the system's current state 472 from the previous regime.

Emulation makes sensitivity analysis for RTMs simple and allows us to identify correlations and
 interactions among parameters that would otherwise be difficult to anticipate, for example the CO₂
 dependency of microbially mediated reactions (Bethke et al., 2011; Jin and Kirk, 2016, 2018; Paper

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- 476 et al., 2021). This ability to elucidate unexpected but key model dependencies and sensitivities could
- 477 prove invaluable in helping direct RTM development.
- 478 4.3.2 Application to Bayesian optimization
- 479 <u>4.3.2 A critical advantage of the technique proposed here is that working emulators are</u>
- 480 essential to Using emulators to identify new feedbacks
- 481 As modern RTMs grow in sophistication and complexity, they increasingly draw on large suites of
- 482 chemical and mineralogical information from vast databases, which constitute large sets of non-linear
- 483 equations all coupled through transport and fluid chemistry. While it is true that for a sufficiently
- 484 simple model, coupled geochemical behavior could be deduced by reasoning about the governing
- 485 equation of the systems, for a large, modern RTM it is inevitable that during development some
- 486 <u>feedbacks will be overlooked.</u>
- 487 <u>Emulation makes sensitivity analysis for RTMs simple and allows us to identify correlations and</u>
- 488 interactions among parameters that would otherwise be difficult to anticipate by allowing an
- 489 investigator to quickly test a wide variety of hypotheses. We demonstrate this in the case of Old
- 490 Rifle by identifying the CO₂ dependency of microbially mediated reactions (Bethke et al., 2011; Jin
- 491 and Kirk, 2016, 2018; Paper et al., 2021). This ability to elucidate unexpected but key model
- 492 <u>dependencies and sensitivities could prove invaluable in helping direct RTM development.</u>

493 <u>4.3.3 Application to Bayesian optimization</u>

- 494 <u>A critical advantage of the technique proposed here is that emulation is an essential part of Bayesian</u>
- 495 optimization. Bayesian optimisation is an approach for finding global maxima and minima in systems
- 496 whose objective function is expensive to evaluate and does not return the gradients of that function
- 497 (of which RTMs are an example) (Frazier, 2018).(Frazier, 2018). Bayesian optimisation works by
- 498 applying an acquisition function that calculates the point that will give the most information about

the function that requires optimisation. An emulator is then fit using these data points selected by the acquisition function and the emulator is updated with a new point each iteration. In this way, the optimiser balances exploitation of known optima, and exploration of unevaluated regions of the function. Such an approach can find the global maximum with relatively few evaluations of the RTM.

504 This study lays the groundwork for future application of Bayesian optimization to highly 505 dimensioned RTMs, potentially allowing for effective optimization over many different (twenty or 506 more) parameters at once. By demonstrating that broad (but local) fits to the RTM with an emulator 507 are possible, we have demonstrated that a GBT regressor can be used as an emulator informing a 508 Bayesian optimization algorithm in this context. This allows for a constellation of local fits in a 509 highly dimensioned space as the algorithm searches for the global optimum in problems that would 510 otherwise be computationally intractable. Bayesian optimisation could even be applied, with a 511 suitable loss function, to optimise for multiple objectives at once (subject to trade-offs among 512 objectives).

513 4.4 Disadvantages of the emulation approach

514 This emulation approach relies on the relative computational inexpensiveness of the RTM. In 515 situations where the underlying model is expensive or time-consuming to evaluate, and 516 computational resources are limited, then this modelling approach becomes unfeasible. One way to 517 overcome this limitation is to reduce the resolution of the RTM (as was done in this work), both in 518 time and space, to lower computation time but this comes at the expense of RTM accuracy. In the 519 context of analysing the interaction of underlying modelled processes in an RTM, this loss of 520 resolution may be less of a problem, as we would be primarily concerned with the relationship 521 among parameters and their impact on outputs, rather than their magnitudes. However, this issue of

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522 computational expense is primarilycan be allayed by the parallelised generation of data alluded to 523 earlier and only the most expensive RTMs would be intractable for a full emulator fit if this 524 technique was deployed correctly, and even in this extreme case, Bayesian optimisation would still 525 be possible.

Additionally, caution is needed when choosing the ranges over which the parameters will be drawn from the uniform random distributions. Key considerations include the number of points being generated relative to the size of the space being covered—a denser cluster of training data will result in a tighter fit, at the expense of range. Conversely, too small of a range and the emulator will not capture key behaviour, or be unable to learn about simulator edge cases, as discussed above with respect to the bioavailable iron in the Old Rifle RTM.

532 4.5 Choice of learning algorithm

Gradient boosted trees outperformed other machine-learning methods that we tested while building the emulators, such as Gaussian process regression. The downsides of GBT include the lack of ability to encode smoothness to preclude sharp discontinuities in the concentration-precipitation space or other such prior assumptions. Furthermore, a low root mean squared error over the entire model fit region does not necessarily imply a good fit globally; it may be that there are some regions of good fit and other regions of poor fit which make up an acceptable root mean square error over the whole space.

540 **4.6** The effect of scale on emulator predictions

541 Our case study relies on the capacity of CrunchTope to predict changes in mineral volume fraction. 542 Therefore, the errors in the predictions, and hence the utility of the approach, ultimately depend on 543 the scale of the system being modelled and thus the sensitivity to what could be very small changes 544 in mineral volume fraction. 545 When analysing the emulator to investigate how different processes in the underlying RTM affect 546 each other, we are primarily considering an issue of whether the emulator can correctly learn the 547 underlying model behaviour. We are also considering whether the emulator can capture the 548 behaviour in the output variables with respect to a changing subset of RTM parameters (some of 549 which we may not have expected at the outset). In this use-case, the emulator is largely concerned 550 with trends and gradients; Figs. 2, 3, 3, 4, S4, S8, and S9 show that this is accurately reported in all 551 case studies. Comparing the case study considered in this paper to the additional case study presented 552 in the Supplement we see that they are discretised at different scales (2 m and 1 cm for the deep-sea 553 sediment column and Old Rifle respectively). However, the emulator for each RTM has root mean 554 squared error over the dataset (and hence absolute error in prediction) of the same order of 555 magnitude. This implies that the error in absolute volume precipitated that each model predicts is 556 different. However, the analysis of the trends and interactions emerging from both RTMs is equally 557 valid in both cases.

558 When concerned with the optimisation capabilities of the emulator, the absolute value of the 559 optimised quantity and hence the model scale must be considered. In large-scale systems, such as 560 weathering of the critical zone, the error in the volume fraction change $(5.5 \times 10^{-5} \text{ for pyrite})$ is below 561 the resolution of measurement techniques for mineral abundance (e.g. XRD and SEM - (Gu et al., 562 2020)(Gu et al., 2020)). However, in smaller-scale systems where the microscale environment 563 becomes increasingly important, these errors in volume fraction become much harder to ignore. For 564 example, in the RTM experiments exploring the effects of scale on simulating mineral dissolution in 565 porous media described by Jung and Navarre-Sitchler (2018), significant errors in changes in 566 predicted volume fraction would propagate into calculated dissolution/precipitation rates, losing 567 sensitivity in the results.

568 4.7 Extension to multiple outputs

569 Multiple output regression (the prediction of a vector of outputs, rather than a single label) is 570 experimentally available in active development for XGBoost and is currently available for supported 571 by other machine learning implementations that we explored, including GPFlow for Gaussian 572 process regression. Given that our approach is currently limited to the prediction of one label-per-573 emulator trained, the availability of regressors that can predict more than one label 'off the shelf' will 574 greatly improve the utility of reactive transport emulation. The prediction of multiple outputs 575 simultaneously will expand the scope of analysis to investigate the interaction of modelled processes 576 in multiple outputs at once. In the context of optimisation problems, one possible application of the 577 emulator like this could be to maximise mineral precipitation in one region of a system while trying 578 to maximise dissolution in another region.

579 **4.8** Improvements to the model

This proof-of-concept model demonstrates the fitting of an emulator over a relatively small range of environmental parameters. Future work will involve expanding the scope of the emulators both in terms of the number of parameters being varied, but also the range over which they are varied, so the whole behaviour of the underlying model can be captured with more accuracy. There is also scope for adding time dependency to the GBT modelling approach, to predict a time series <u>of</u> intermediate RTM states during the evolution of geochemical systems.

586 4.9 Potential applications

587 Our emulator approach is flexible; any quantity recorded by an RTM can be used as a target variable, 588 and so the behaviour of any RTM output can be explored in detail to evaluate the model formulation. 589 The behaviour of the system in response to the variation of any parameter under any other set of 590 conditions can be projected out of the model and plotted in a straight-forward manner. This approach 591 can be extended to two or even three dimensions and time series thereof and ultimately the emulator

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can be interrogated for local maxima and minima to solve optimisation problems. This-new approach
has potential applications in industry and in environmental remediation where the chemical
composition of amendments can be predicted using an underlying reactive transport simulation,
provided that that system is well understood.

596 Omphalos also has utility outside of generating datasets for emulation; its automated submission of 597 CrunchTope input files means it can be used to systematically explore sets of input variables in an 598 easy way, simply by editing the Omphalos configuration file.

599 **5** Conclusions

600 We have presented a newan emulator based approach for interrogating and understanding multi-601 component RTMs. By building an emulator of an RTM that captures the multidimensional nature of 602 the underlying model we have created a newdemonstrated that such an approach can be used as a tool 603 for performing global sensitivity analyses on RTMs. This allows us to investigate behaviour arising 604 from the interaction among the many disparate processes that comprise RTMs. For example, we 605 investigated how the Monod-biomass parameterisation of microbial sulfate reduction interacted with 606 the mechanism of pyrite precipitation. In this example, pyrite precipitation was inhibited when there 607 was an excess of CO₂ in the column because the catabolic pathway was partially dependent on CO₂ 608 concentration. This prevented the growth of sulfate reducing biomass, ultimately curtailing the 609 production of hydrogen sulfide required for pyrite precipitation. This behaviour reproduced results 610 previously reported by Jin and Kirk (2016, 2018) (2016, 2018), and suggest that our emulation 611 approach has approaches have utility in discovering unexpected, but nonetheless real, model 612 behaviours, potentially directing future lab and field work.

613 The approach<u>methodology we have laid out</u> is flexible; any quantity recorded by an RTM can be 614 used as a target variable, and so the behaviour of any RTM output can be explored in detail to

30

615	evaluate the model formulation. The behaviour of the system in response to the variation of any
616	parameter under any other set of conditions can be projected out of the model and plot in a straight-
617	forward manner. This approach Emulator approaches can be extended to two or even three
618	dimensions and ultimately the emulator can be interrogated for local maxima and minima to solve
619	optimisation problems. This new approach has We suggest that emulator based approaches to
620	exploring RTMs have potential applications in industry and in environmental remediation where the
621	chemical composition of amendments can be predicted using an underlying reactive transport
622	simulation, provided that that system is well understood. The presentation of this optimisation
623	process to Old Rifle (and to ODP Site 1086, see supplementary) represents a proof of concept.
624	6 Code availability
625	6.1 Omphalos
626	Omphalos is available on GitHub and Zenodo. Please note you must provide your own CrunchTope
627	executable.
628	https://github.com/a-fotherby/Omphalos
629	https://doi.org/10.5281/zenodo.7113298
630	6.2 GBT Models
631	Jupyter notebooks for fitting the GBT models and plotting the figures are available on GitHub, and a
632	permanent record is available on Zenodo.
633	https://github.com/a-fotherby/dissertation_xgboost.
634	https://doi.org/10.5281/zenodo.7113323
635	7 Data availability

- 636 The data used is available on GitHub and Zenodo.
- 637 <u>https://github.com/a-fotherby/GMD 2022</u>
- 638 <u>https://doi.org/10.5281/zenodo.7113379</u>

639 8 Supplement

- 640 Codebase for Omphalos. Model fitting code. Schematic figures of decision tree and the Old Rifle
- 641 RTM. Table of predicted optimal values for precipitating pyrite at Old Rifle. Convergence behaviour
- of the GBT regressors. Additional co-dependency plots for Old Rifle. Figure showing the effect of
- rate law choice on CO₂ dependency in the Old Rifle RTM. Supplementary Case Study detailing
- 644 application to a deep-sea sediment column. Description of XGBoost implementation.

645 9 Author contribution

- 646 AF and HJB conceived of the study. AF wrote the codebase and conducted the experiments. AF
- 647 prepared the manuscript with contributions from all co-authors.

648 10 Competing interests

649 The authors declare that they have no conflict of interest.

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