Mathematical Modeling, Laboratory Experiments, and Sensitivity Analysis of Bioplug Technology at Darcy Scale

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Summary

In this paper we study a Darcy-scale mathematical model for biofilm formation in porous media. The pores in the core are divided into three phases: water, oil, and biofilm. The water and oil flow are modeled by an extended version of Darcy’s law and the substrate is transported by diffusion and convection in the water phase. Initially there is biofilm on the pore walls. The biofilm consumes substrate for production of biomass and modifies the pore space which changes the rock permeability. The model includes detachment of biomass due to water flux and death of bacteria, and is implemented in MRST. We discuss the capability of the numerical simulator to capture results from laboratory experiments. We perform a novel sensitivity analysis based on sparse-grid interpolation and multi-wavelet expansion to identify the critical model parameters. Numerical experiments using diverse injection strategies are performed to study the impact of different porosity-permeability relations in a core saturated with water and oil.

Introduction

After primary and secondary production, up to 85% of the oil remains in the reservoir (Patel et al. 2015). Microbial improved and enhanced oil recovery (MIEOR) is one of the secondary and tertiary methods to increase the oil production using microorganisms (Wood 2019). Bioplug technology is a MIEOR strategy that consists in plugging the most permeable zones in the reservoir, which provokes water to flow through new paths and recovering the oil in these new zones.

Experiments in microsystems allow us to observe processes in more detail, which leads to improvement of the experimental methods in core-scale experiments. For example, in Liu et al. (2019) the effects of flow velocity and substrate (also referred to as nutrients) concentration on biofilm in a microchannel was studied, finding values of substrate concentration and flow velocity for a strong plugging effect. Core samples from reservoirs can be used to study changes in permeability due to biofilm formation, e.g., in Suthar et al. (2009) two-phase flow experiments were performed to study the selective plugging strategy for MIEOR. In that study, the MIEOR effects increased the oil recovery around 25%.

Mathematical models of bioplug technology are important as they help to predict the applicability of this MIEOR strategy and to optimize the benefits. In Kim (2006) a mathematical model for single-phase flow was proposed which includes changes of rock porosity and permeability as a result of biofilm growth. Li et al. (2011) built a mathematical model for two-phase flow including the effects of bio-surfactants and biomass on improving the oil recovery. The authors also compare the numerical results for different porosity-permeability relations. These porosity-permeability relations can also include the permeability of biofilm and be derived as a result of upscaling pore-scale models (van Noorden et al. 2010; Hommel et al. 2018; Landa-Marbán et al. In press). In this work, we present a two-phase core-scale model of bioplug technology and study the oil production for different porosity-permeability relations and injection strategies.

Sensitivity studies of mathematical models are of great interest because they provide estimates of the influence of physical parameters on a quantity of interest, e.g., biofilm formation. In Brockmann et al. (2006) a regional steady-state sensitivity analysis was performed to identify parameters with the largest impact on a mathematical model for deammonification in biofilm systems. Sensitivity analysis by means of Sobol decomposition provides rigorous estimates of parameter dependencies, but are prohibitively expensive to compute if the number of parameters is large. This is remedied for smooth problems by first computing spectral (generalized polynomial chaos) expansions in the parameters, which then leads to efficient evaluation of the sensitivity indices via post-processing of spectral coefficients (Sudret 2008). The latter method was employed in Landa-Marbán et al. (2019), where a
global sensitivity analysis was performed using Sobol indices to identify the critical parameters of a pore-scale model for permeable biofilm. In this paper, we consider nonsmooth models in the dependent parameters, hence spectral expansions with global smooth basis functions are not a robust choice. Instead, we propose a two-stage method where we first use sparse grids to estimate a piecewise linear interpolant of the function of interest, which yields a surrogate that can be further sampled at negligible cost. Secondly, we compute a multi-wavelet representation of the interpolant, from which the sensitivities can be directly evaluated.

The mathematical model consists of coupled nonlinear partial differential equations. Two-point flux approximation (TPFA) and backward Euler (BE) are used for the space and time discretization respectively. To solve the resulting nonlinear algebraic system, Newton’s method is used. The scheme is implemented in the MATLAB reservoir simulation tool (MRST), a free open-source software for reservoir modeling and simulation (Lie 2019).

To summarize, the new contributions of this work are:

- To compare the experimental results to numerical simulations of a core-scale porous medium including biofilm formation.
- Performing a novel global sensitivity analysis of the model parameters to identify the critical parameters.
- Performing numerical simulations for different porosity-permeability relations to study their impact on the predicted oil recovery.
- The study of diverse substrate injection strategies for the bioplug technology.

The paper is structured as follows. Firstly, we introduce and describe the implementation of the core-scale mathematical model for two-phase flow including the effects of biofilm formation. Secondly, we present a comparison of numerical simulations to laboratory experiments of this core-scale model. Thirdly, we introduce a novel method for global sensitivity analysis and apply it to the mathematical core-scale model. Diverse numerical experiments for different porosity-permeability relations and injection strategies are also explained. Finally, we present the conclusions.

**Core-Scale Model**

We consider a core sample of radius $r$ and length $L$ initially filled with oil and water and a given biofilm distribution. Fig. 1 shows the schematic representation of this system. As water and substrate are injected, some biomass is detached due to erosion (shear forces caused by the water flow). The biofilm consumes substrate to produce metabolites (i.e., gases) and to grow which modifies the rock porosity and hence the rock permeability. Consequently, the flow pattern is modified in the sense that pores where oil was replaced by water, and which were forming a preferential path for water flow, become less permeable. Therefore water enters other pores, mobilizing the oil present there and leading to improved oil production. The mathematical model presented in this section aims to describe the following processes in a core after bacterial inoculation: two-phase flow, substrate transport, permeability changes due to porosity modification, and biofilm growth, detachment, and death.

![Fig. 1—Core sample for laboratory experiments after inoculation of bacteria.](image)

We assume that the fluids are immiscible and incompressible. We consider that the biofilm has water and biomass as the only two components, and that the biofilm porosity and biofilm permeability are constants. In order to determine the amount of fluid outside the biofilm in the representative element volume (REV), we introduce the saturation of a fluid $S_\alpha$ (for oil $\alpha = o$ and for water $\alpha = w$) given by the ratio of volume of fluid $\alpha$ outside the biofilm over the volume of voids outside the biofilm (in REV). For this two-phase flow model, we have that $S_o + S_w = 1$.

The mass conservation and extended Darcy’s law for the oil phase are given by

$$\frac{\partial}{\partial t} (\rho_o \phi_f S_o) + \nabla \cdot (\rho_o \vec{v}_o) = Q_o, \quad \vec{v}_o = -\frac{k_{r,o}}{\mu_o}k(\nabla p_o - \rho_o \vec{g}), \quad \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cdots \cd -2
and for the water phase

\[
\frac{\partial}{\partial t}[\rho_w(\phi_f S_w + \phi_b \theta_w)] + \nabla \cdot (\rho_w \vec{v}_w) = Q_w, \quad \vec{v}_w = -\frac{k_{r,w}}{\mu_w} k (\nabla p_w - \rho_w \vec{g}),
\]

where \(\phi_f\) is the porosity outside the biofilm, \(\vec{v}_w\) the flow velocity, \(k_{r,w}\) the relative permeability, \(\rho_w\) the fluid density, \(k\) the absolute rock permeability, \(\vec{g}\) the gravity, \(\mu_w\) the viscosity, \(\theta_w\) the volume fraction of biofilm in the REV, \(\phi_b\) the biofilm water content, and \(Q_w\) source/sink terms. The minimum value of \(\phi_b\) is zero while the maximum value of \(\phi_b\) is equal to the initial porosity of the rock \(\phi_0\), corresponding to the case when the pores are filled with biofilm. In this work, we assume that there are no fluid sources/sinks, neglect the gravity effects, set to zero the residual phase saturations. Simplified expressions of these relations are given by

\[
k_{r,w}(S_w) = S_w^\gamma, \quad k_{r,o}(S_w) = (1 - S_w)^\beta, \quad .......................... (3)
\]

where \(\gamma\) and \(\beta\) are experimentally fitted factors. We assume that water can flow inside the biofilm even when there is only oil outside the biofilm (\(S_w = 0\)). Then, the previous relation is not a good model because for \(S_w = 0\) it results in \(k_{r,w}(0) = 0\) which leads to zero water velocity. Recalling that the relative permeabilities are used to extend Darcy’s law to multiphase flow, we look for relationships which account for the water saturation and volume fraction of biofilm. These relationships need to fulfill the following criteria: when there is only oil outside the biofilm, then \(k_{r,w}(S_w = 0, \phi_b) = 1\); when there is only biofilm in the REV, then \(k_{r,w}(S_w, \phi_b = \phi_0) = 1\). Then, we propose the following two relative permeability relations:

\[
k_{r,w}(S_w, \phi_b) = S_w^\gamma \left(1 - \frac{\phi_b}{\phi_0}\right) + \left(\frac{\phi_b}{\phi_0}\right), \quad k_{r,o}(S_w, \phi_b) = (1 - S_w)^\beta \left(1 - \frac{\phi_b}{\phi_0}\right). \quad .......................... (4)
\]

To describe the movement of substrate, we consider the following convection-diffusion-reaction transport equations:

\[
\frac{\partial}{\partial t}[\phi f (\phi f S_w + \phi_b \theta_w)] + \nabla \cdot \vec{j}_n = R_n, \quad \vec{j}_n = -D_n(\phi f S_w + \phi_b \theta_w) \nabla C_n + C_n \vec{v}_w. \quad .......................... (5)
\]

In the previous equations, \(C_n\) is the substrate concentration in the water, \(\vec{j}_n\) the substrate flux in water, and \(D_n\) the substrate dispersion coefficient which includes mechanical dispersion plus diffusion. The reaction term \(R_n\) is given by

\[
R_n = -\mu_n \frac{C_n}{K_n + C_n} \rho_b \phi_b, \quad .......................... (6)
\]

where \(\mu_n\) is the maximum rate of substrate utilization, \(\rho_b\) the biomass density, and \(K_n\) is the Monod half-velocity coefficient.

The following equation describes the biofilm evolution:

\[
\frac{\partial}{\partial t}(\rho_b \phi_b) = \mu_n \frac{C_n}{K_n + C_n} \rho_b \phi_b - K_d \rho_b \phi_b - K_{str} ||\nabla p|| \rho_b \phi_b. \quad .......................... (7)
\]

Here, we assumed that all consumed substrate is used to produce biomass, a linear death of bacteria given by \(K_d\), and erosion of bacteria given by the magnitude of the pressure gradient times a constant depending on the biofilm \(K_{str}\). After biomass has been detached, in this model we assume that the detached biomass flows out of the core and does not affect the rock properties; therefore, we do not include a transport equation for the detached biomass.

Diverse porosity-permeability relations have been proposed for the last decades. We refer to Hommel et al. (2018) for a recent review of these relations. Three porosity-permeability relations commonly used in modeling are the following:

\[
k_b = \left(\frac{\phi f}{\phi_0}\right)^\eta, \quad k_{vp} = \left(\frac{\phi f - \phi_{crit}}{\phi_0 - \phi_{crit}}\right)^\eta, \quad k_h = a \left(\frac{\phi f - \phi_{crit}}{\phi_0 - \phi_{crit}}\right)^3 + (1 - a) \left(\frac{\phi f - \phi_{crit}}{\phi_0 - \phi_{crit}}\right)^2, \quad .......................... (8)
\]
The first one is called the power law (Carman 1937; Ives and Pienvichitr 1965), where $\eta$ is a fitting factor calibrated either from experimental data or taken from a process-specific literature. The second relation is a variation of the power law known as the Verma-Pruess relation (Verma and Pruess 1988), where $\phi_{crit}$ is a critical porosity when the permeability becomes zero, which value is between 70 and 90% of the initial porosity. The third relation is proposed by Thullner et al. (2002), where $a$ is a weighting factor between -1.7 and -1.9. These three relations do not include the biofilm permeability. Vandevivere (1995) proposes the following relation of permeability and porosity for a plugging model

$$\frac{k_v}{k_0} = \exp \left[ \frac{1}{2} \left( \frac{B_r}{B_c} \right)^2 \left( \frac{\phi_f}{\phi_0} \right)^2 \right] \left[ 1 - \exp \left( -\frac{1}{2} \left( \frac{B_r}{B_c} \right)^2 \right) \right] \frac{k_b \phi_0}{k_0 \phi_0 - (k_0 - k_b) \phi_f}, \hspace{1cm} (9)$$

where $k_b$ is the biofilm permeability, $B_r$ is a relative porosity given by $B_r = 1 - \phi_f/\phi_0$, and $B_c$ is the critical point where biofilm begins to detach and form plugs. Thullner et al. (2002) presents the following relation which includes the biofilm permeability:

$$\frac{k_{th}}{k_0} = \left[ \left( \frac{\phi_f - \phi_{crit}}{\phi_0 - \phi_{crit}} \right)^\eta \right] + \frac{k_b}{k_0 + k_b}. \hspace{1cm} (10)$$

All these relationships are postulated at the Darcy-scale, based on experimental observations. A different approach is followed in van Noorden et al. (2010), Schulz and Knabner (2016), and Landa-Marbán et al. (In press), based on homogenization. Starting with models at the pore scale, one applies expansion methods to derive the mathematical models valid at the Darcy scale. In Landa-Marbán et al. (In press) effective porosity-permeability relations are derived for two different pore geometries: thin channels $k_c$ and tubes $k_t$. In this work we compare the following four porosity-permeability relations: Vandevivere $k_v$ (9), Thullner $k_{th}$ (10), channel $k_c$, and tube $k_t$. We refer to Appendix A for the mathematical expressions of $k_c$ and $k_t$.

The porosity in the porous medium changes in time as a function of the biofilm volume fraction $\phi_b(t)$. When there is no biofilm, the porosity $\phi_f(t)$ is equal to the initial porosity. In the case when the porous medium is filled with biofilm ($\phi_f(t) = 0$), the porosity in the REV is equal to the biofilm porosity $\theta_w$. The porosity-permeability relations (9-10) are given as a function of the rock porosity and do not include the biofilm porosity. The porosity-permeability relations $k_c$ and $k_t$ do include the biofilm porosity. Using the definitions of $\phi_f(t)$ and $\phi_b(t)$, we have the following relation for the void space outside the biofilm

$$\phi_f(t) = \phi_0 - \phi_b(t). \hspace{1cm} (11)$$

This mathematical model is implemented in a 3D domain with a uniform cell-centered grid. The TPFA is used for the spatial discretization, while BE for the time discretization. The Newton’s method is used as linearization scheme. The implementation of this core-scale mathematical model is done in MRST, based on the modification of the polymer example in the module ad-eor (see Bao et al. 2007).

**Model Test**

Core-scale experiments under controlled conditions are performed in the laboratory for studying the effect on biofilm growth in porous media. These experiments aim to provide a better understanding of different features, i.e., the relation between biofilm composition and growth conditions, the plugging potential of different bacteria, and the adaptability of biofilm at diverse substrate flux. Fig. 2 shows a photograph of a typical core sample to study these effects.

In this section, we describe an experiment performed by NORCE for Equinor. The aim of this experiment is to test different brine qualities on the same system and the same biofilm. The core has a length of 29.50 cm and a radius of 1.90 cm. Initially, the core has an approximately homogeneous porosity of $\phi_0=0.23$ and permeability $k_0=1528$ md. The core sample is introduced inside a core holder, fully saturated with brine. Heating cables wrapped around the exterior of the core holder are used to conduct the experiments at a constant temperature (30°C). A backpressure regulator (BPR) is used to control the outlet pressure. Bacteria are injected into the core and the core is left standstill overnight to allow the bacteria to attach themselves to the pore walls. Fig. 3 shows a diagram of the experimental set-up.

The core is injected with two different brines with different substrate concentrations, where glucose is added as a source of carbon. We refer to these brines as $B_1=30.81$ kg/m³ and $B_2=42.63$ kg/m³. The two different brines are injected at a constant flow rate of $8.33 \times 10^{-10}$ m³/s (Darcy velocity of 6.4 cm/day). The injection strategy is the following: $B_1$ is injected during 53 days, after $B_2$ is injected during 82 days and finally $B_1$ is injected again during 61 days. The core permeability is estimated at different times, using the measurement of the pressure difference along the core divided by the pressure drop on the clean core. This ratio is known as resistance factor.
In the previous section we introduced the model equations for biofilm growth and two-phase flow in porous media. We simplify the mathematical model to compare to the experiment. This experiment can be modeled as a 1D single-phase flow system, where in addition we consider that the biofilm is impermeable and only diffusion for the substrate to reduce the number of parameters. Table 1 presents this simplified version of the mathematical model which includes only six variables: the water velocity $v_w$, water pressure $p_w$, permeability $k$, porosity $\phi$, substrate concentration $C_n$, and volume fraction of biomass $\phi_b$.

$$R_f$$ and it is given by $R_f(t) = \Delta p_w(t)/\Delta p_0$. Assuming that there is not change in the fluid rate, viscosity, and density, the resistance factor gives an estimate of the current rock permeability $k(t) = R_f(t)k_0$.

In the previous section we introduced the model equations for biofilm growth and two-phase flow in porous media. We simplify the mathematical model to compare to the experiment. This experiment can be modeled as a 1D single-phase flow system, where in addition we consider that the biofilm is impermeable and only diffusion for the substrate to reduce the number of parameters. Table 1 presents this simplified version of the mathematical model which includes only six variables: the water velocity $v_w$, water pressure $p_w$, permeability $k$, porosity $\phi$, substrate concentration $C_n$, and volume fraction of biomass $\phi_b$.

<table>
<thead>
<tr>
<th>Name</th>
<th>Equation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Darcy</td>
<td>$v_w = -(k/\mu_w)\partial_x p_w$, $\partial_t \phi_f + \partial_x v_w = 0$</td>
</tr>
<tr>
<td>Permeability</td>
<td>$k = k_0(\phi_f/\phi_0)\eta$</td>
</tr>
<tr>
<td>Porosity</td>
<td>$\phi_f = \phi_0 - \phi_b$</td>
</tr>
<tr>
<td>Substrate</td>
<td>$\partial_t(C_n\phi_f) + \partial_x (v_w C_n - D_n \phi_f \partial_x C_n) = -\mu_n p_b \phi_b C_n / (K_n + C_n)$</td>
</tr>
<tr>
<td>Biofilm</td>
<td>$\partial_t \phi_b = \mu_n \phi_b C_n / (K_n + C_n) - K_d \phi_b - K_{str}</td>
</tr>
</tbody>
</table>

Table 1—Core-scale single-phase equations.

Nine parameters are needed to solve the mathematical model in Table 1. To obtain a better estimate of the model parameters, it is necessary to perform various experiments under controlled input quantities. However, these experiments are expensive and time consuming. In Landa-Marbán et al. (2019) a pore-scale mathematical model is calibrated based on the experiments performed by Liu et al. (2019), where measurements of the biofilm
amount over time are taken for different flux velocities. For the core-scale system described in this section, only one experiment is performed. Then, we select parameter values from the literature in order to run numerical simulations and compare qualitatively the results to the experimental observations. Table 2 shows the selected values of these parameters.

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Notation</th>
<th>Value</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>Bacterial death rate</td>
<td>$K_d$</td>
<td>$3.2 \times 10^{-4}$ s$^{-1}$</td>
<td>Kundu et al. (1983)</td>
</tr>
<tr>
<td>Maximum growth rate</td>
<td>$\mu_n$</td>
<td>$5 \times 10^{-6}$ s$^{-1}$</td>
<td>Li et al. (2011)</td>
</tr>
<tr>
<td>Monod-half velocity</td>
<td>$K_n'$</td>
<td>$0.915$ kg/m$^3$</td>
<td>Linville et al. (2013)</td>
</tr>
<tr>
<td>Substrate diffusion coefficient</td>
<td>$D_n$</td>
<td>$5 \times 10^{-10}$ m$^2$/s</td>
<td>Hardy and Sarko (1993)</td>
</tr>
<tr>
<td>Biomass density</td>
<td>$\rho_b$</td>
<td>$20$ kg/m$^3$</td>
<td>Ro and Neethling (1991)</td>
</tr>
<tr>
<td>Power law constant</td>
<td>$\eta$</td>
<td>$2.5$</td>
<td>Hommel et al. (2018)</td>
</tr>
<tr>
<td>Water dynamic viscosity</td>
<td>$\mu_w$</td>
<td>$10^{-3}$ Pa·s</td>
<td>Standard</td>
</tr>
<tr>
<td>Water density</td>
<td>$\rho_w$</td>
<td>$10^3$ kg/m$^3$</td>
<td>Standard</td>
</tr>
</tbody>
</table>

Table 2—Model parameters for the verification study.

These parameter values have the same order of magnitude as the ones used in mathematical modeling (Alpkvist and Klapper 2007; Duddu et al. 2009; Hommel et al. 2018; Landa-Marbán et al. 2019). The stress coefficient $K_{str}$ is neglected before the simulations and its value is chosen to better fit the experimental measures. In addition to the model parameters and core dimensions, initial, and boundary conditions are needed to complete the model. The initial porosity and permeability are $\phi_0=0.23$ and $k_0=1528$ md respectively. We assume that initially there is no substrate in the brine. The pressure is set to zero initially (in this system, we are only interested in pressure differences that affects the detachment and transport of substrate). Recalling that the time after inoculation and before starting the substrate injection is roughly half day, we assume that the initial volume fraction of biofilm is $0.04$. From subplot (a) that after a day of substrate injection the constant flow rate $B_1$ was returned to $B_2$ and the resistance factor does not increase significantly. When increasing the substrate concentration to $B_2$ from $B_1$, both simulations and experimental measures show that the resistance factor does not increase significantly. When increasing the substrate concentration to $B_2$, we observe that the resistance factor increases due to major biofilm activity. When the substrate injection is set back to $B_1$, we observe a steady state where the resistance factor has increased by a factor of three. From the simulation, it is possible to observe how the resistance factor decreases from changing the substrate injection $B_2$ to $B_1$, which means a sensible response of the biofilm to substrate input. One of the outcomes of this experiment is that the plugging can be controlled by modifying the substrate flux. In addition, the growth conditions of the microorganism are affected by the different physical environments. From the experimental data we observe fluctuations on the measurements. This behavior is attributed to the dynamical attachment and detachment of biomass, a complex process not included in the mathematical model. However, this mathematical model is simple enough to have few parameters and complex enough to capture processes such as dynamical changes on the resistance factor and a steady state where the biofilm growth is balanced with the detachment and death of bacteria.

Fig. 5 shows values of the simulations for pressure, substrate, and biomass along the core. We observe from subplot (a) that after a day of substrate injection the constant flow rate $B_1$ requires an inlet pressure of approximately 200 Pa. From subplot (b) we observe that the inlet pressure has increased more than twice in order to keep the flow rate, as a result of permeability reduction due to the biofilm formation. From the substrate profile (c), this decreases from the inlet to the outlet as bacteria consume them. Given the initial homogeneous volume fraction of biofilm, this has increased more than three order of magnitude in comparison with the initial volume fraction, but it has stopped growing due to the shear forces.

**Global Sensitivity Analysis**

We keep the exposition on sensitivity analysis general to emphasize that the methodology is not restricted to the problems presented in this paper. Let $g(y)$ be a scalar multidimensional function with $y = (y_1, \ldots, y_n) \in \Omega \subset \mathbb{R}^n$, defined by a range of independent parameters, i.e. $\Omega = (a_1, b_1) \otimes \ldots \otimes (a_n, b_n)$. We associate a nonnegative weight $w_j(y_j) = 1/(b_j - a_j)$ with each parameter $y_j \in [a_j, b_j]$ for $j = 1, \ldots, n$, and use the product measure $w = \prod_{j=1}^n w_j$. More general weight functions are possible, including non-product measures corresponding to inter-dependence between the parameters (Rahman 2014). In this paper we do not aim to quantify uncertainty by stochastic models, but sensitivities in outputs given bounds on the input parameters.
To determine the relative effect of each of the \( n \) input parameters on the output quantity of interest \( q \), we perform a global sensitivity analysis in terms of a Sobol decomposition (Sobol 2001). The function \( q \) is decomposed as a series expansion in all subsets of variables; the sum of contributions from the individual variables in isolation, all
combinations of pairs of variables and so on, leading to the expression:

\[ q(\vec{y}) = q^{[0]} + \sum_{i=1}^{n} q^{[i]}(y_i) + \sum_{i=1, j>i}^{n} q^{[i,j]}(y_i, y_j) + \ldots + q^{[1,\ldots,n]}(\vec{y}), \]

(12)

where the terms are defined recursively by

\[ q^{[0]} = \int_{\Omega} q(\vec{y}) w(\vec{y}) d\vec{y}, \]

(13)

\[ q^{[i]}(y_i) = \int_{\Omega_{+i}} q(\vec{y}) w_{\sim i}(\vec{y}) d\vec{y}_{\sim i} - q^{[0]}, \quad 1 \leq i \leq n, \]

(14)

\[ q^{[i,j]}(y_i, y_j) = \int_{\Omega_{+i,j}} q(\vec{y}) w_{\sim i,j}(\vec{y}) d\vec{y}_{\sim i,j} - q^{[i]}(y_i) - q^{[j]}(y_j) - q^{[0]}, \quad 1 \leq i < j \leq n, \]

(15)

and so on for higher-order terms. The subscript \( \sim i \) means that the \( i \)th index is omitted, e.g., \( w_{\sim i} = \prod_{j \neq i} w_j \).

The Sobol index for the \( s \)-parameter combination \( \{y_1, y_2, \ldots, y_s\} \) is given by

\[ S_{(1,\ldots,i_s)} = \frac{1}{\text{Var}(q)} \int_{\Omega_{Y_{s-1},\ldots,i_s}} [q^{\{1,\ldots,i_s\}}(y_1, \ldots, y_s)]^2 w_i(y_1) \ldots w_s(y_s) dy_1 \ldots dy_s. \]

(16)

The total variability in \( q \) due to variable \( i \) is obtained by summing over all subsets of parameters including parameter \( i \), denoted \( I_i \), which yields the total Sobol index for parameter \( i \),

\[ S_{(i)} = \sum_{s=1}^{n} \sum_{i \in \{1,\ldots,i_s\}} S_{(1,\ldots,i_s)}. \]

(17)

Directly computing global sensitivities of a computationally complex function, e.g., a computer model, over a domain of only moderately high dimensionality is often infeasible due to the computational cost. As an additional challenge, we are interested in nonsmooth \( q \), excluding the use of Sobol decomposition based on generalized polynomial chaos for smooth problems (Sudret 2008). As a remedy, we approximate the function of interest with an interpolant on a Clenshaw-Curtis type sparse grid, using the software in Klimke and Wohlmuth (2005). Subsequently, a multi-wavelet decomposition of the interpolant is performed, and the global sensitivity indices are evaluated directly from the multi-wavelet coefficients. The accuracy in the sensitivity indices as determined by the computer model is thus largely determined by two kinds of errors: interpolation error and basis truncation error introduced when replacing the interpolant by a series expansion in a finite set of multi-wavelets.

**Multi-Linear Collocation.** The outline of multi-linear collocation closely follows the exposition in Klimke and Wohlmuth (2005), which also provides the source code for the numerical implementation. We perform interpolation of possibly nonsmooth (but continuous) functions and therefore rely on localized basis functions. For discontinuous functions, we refer to the adaptive sparse grid methods introduced in Ma and Zabaras (2009). Multidimensional collocation on sparse grids is built from tensor products of low-order single dimensional collocation rules. Temporarily assume that \( n = 1 \), denote \( y = \vec{y} \) and let \( Y_\ell \) be a set of interpolation nodes in the parameter space, for which the function \( q \) is evaluated. The index \( \ell \) refers to a refinement level: the higher \( \ell \), the bigger the set of interpolation nodes. An interpolant of \( q \) is given by

\[ I_\ell(q) = \sum_{y_\ell \in Y_\ell} q(y_\ell) \psi_{y_\ell}(y), \]

(18)

where \( \psi_{y_\ell}(y) \) are nodal basis functions with the property

\[ \psi_{y_\ell}(y) = \begin{cases} 1 & \text{if } \ell = \ell', \\ 0 & \text{otherwise}. \end{cases} \]

(19)

In this work we employ the piecewise linear ‘hat’ functions for level \( \ell > 1 \)

\[ \psi_{y_\ell}(y) = \begin{cases} 1 - 2^{1-i}|y - y_\ell| & \text{if } |y - y_\ell| < 2^{1-i}, \\ 0 & \text{otherwise}, \end{cases} \]

(20)

supplemented with the unit function (level \( \ell = 1 \)). The collocation nodes are nested, i.e., \( Y_\ell \subseteq Y_{\ell+1} \). Rather than expressing the interpolant as a function of the nodes in some sufficiently refined set \( Y_\ell \), it can be expressed as a sum of hierarchical surpluses, i.e., differences between the interpolants at successive levels,

\[ \Delta_\ell(q) = I_\ell(q) - I_{\ell-1}(q), \quad I_0(q) \equiv 0. \]

(21)
Then, Eq. (18) can be expressed
\[
I_\ell(q) = \sum_{j=1}^\ell \Delta_j(q) = \sum_{j=1}^\ell \sum_{y_j \in Y_j \sim Y_{j-1}} q(y_j) \psi^{(j)}(y_{j,Y}),
\]
(22)
where the superscript \((j)\) is used to emphasize that the basis functions are level specific. Using this construction, one may use the local difference between the interpolant and the true function as a measure to determine where further refinement in terms of more nodes are needed.

In multiple dimensions \((n > 1)\), a tensor-product formula would lead to very large sets of nodes even for moderate \(n\). As a remedy, a sparse grid based on Smolyak’s construction (Smolyak 1963) is used, adding only a subset of the tensor product Clenshaw-Curtis nodes at each new level. The hierarchical multidimensional interpolant of \(q = q(y_1, \ldots, q_n)\) on total level \(p \geq n\), is given by
\[
I_{p,n}(q) = I_{p-1,n}(q) + \Delta I_{p,n}(q),
\]
(23)
with \(I_{n-1,n} = 0\) and
\[
\Delta I_{p,n}(q) = \sum_{|i|=p} (\Delta_{i_1} \otimes \cdots \otimes \Delta_{i_n})(q) = \sum_{|i|=p} \sum_j \left( \psi^{(i_1)}_{j_1} \otimes \cdots \otimes \psi^{(i_n)}_{j_n} \right) \cdot q(y_{j_1}) - I_{p-1,n}(q)(y_{j_2}^{(i)}),
\]
(24)
For more details on the multi-linear collocation method we refer to Klimke and Wohlmuth (2005). The hierarchical multi-linear interpolant (23) is a surrogate from which we can obtain approximations of \(q\). The Smolyak algorithm yields the multidimensional interpolant at low computational cost, but the global sensitivity indices are not directly available. Next, we introduce a series expansion of \(I_{p,n}\) in multi-wavelets to obtain the Sobol decomposition of \(I_{p,n}\).

**Sobol Indices via Multi-Wavelet Expansions.** An alternative representation of \(q\) is via a spectral expansion in a set of multidimensional orthogonal basis functions \(\{\varphi_j\}\),
\[
q(Y) = \sum_{j=1}^{\infty} c_j \varphi_j(Y),
\]
(25)
where the infinite sum is truncated to some finite set in practical application. This is known as generalized polynomial chaos expansion if the basis functions are orthogonal polynomials (Xiu and Karniadakis 2002), and it has been demonstrated that the Sobol indices can be directly identified from the expansion coefficients, once these have been computed (Sudret 2008). Nonsmooth functions can be represented by means of multi-resolution analysis where the basis functions are piecewise polynomial multi-wavelets (Le Maitre et al. 2004), and the identification of the Sobol indices is straightforward.

A piecewise linear multi-wavelet expansion of sufficient resolution can exactly represent the piecewise linear interpolant, provided that the support nodes are aligned with the support of the multi-wavelets. The advantage of using the multi-wavelet expansions rather than trying to directly evaluate the Sobol component functions, is that we may rely on a single interpolation of the function of interest itself (and not its square, conditional on some subset of the variables, and so on).

Let \(\{\varphi_{\ell}(y)\}_{\ell \in \text{MW}}\) be a finite-dimensional basis of multi-wavelets indexed by some set of nonnegative integers \(I_{\text{MW}} \subset \mathbb{N}_0^n\). Ideally, the number of basis functions should be limited but chosen in order to be a good approximation of the interpolant \(I_{p,n}(q)\) of the function of interest. The approximation properties of the multi-wavelet basis are determined by the order of the piecewise polynomial multi-wavelets, and the resolution level that governs the localization in parameter space. Analogous to the hierarchical surplus defined for the multi-linear interpolant, the hierarchical surplus of the multi-wavelet expansion is given by the contribution captured by the finest resolution level. The multi-wavelet basis is hierarchical, and can be enriched by adding multi-wavelets to regions labeled important due to large surpluses.

We may now either project the interpolant \(I_{p,n}(q)\) onto the basis \(\{\varphi_{\ell}(y)\}_{\ell \in \text{MW}}\), or interpolate the product \(q \varphi_{\ell}\), and then perform the projection by computing the expected value. The latter is simpler, as it only involves integration of a piecewise linear function to compute each multi-wavelet coefficient. However, we do not know a priori what multi-wavelets should be included in the truncated basis, and will therefore settle for a simple strategy that also admits adaptivity. Unlike \(q\) itself, the interpolant \(I_{p,n}(q)\) can be sampled repeatedly at moderate computational cost. Given \(N\) random samples of the parameters \(y\), and a tentative multi-wavelet basis of size \(P\), we form the linear system for the \(P\)-vector of multi-wavelet coefficients \(\varphi\)
\[
\Phi \tilde{c} = \tilde{q},
\]
(26)
where the matrix $\Phi \in \mathbb{R}^{N \times P}$ is defined by $[\Phi]_{k,j} = \varphi_j(\vec{y}(k))$, and $\vec{q} \in \mathbb{R}^N$ contains the evaluations of $I_{p,n}(q)$ in the samples $y_j$, $j = 1, \ldots, N$. The ordinary least squares solution to (26) gives the multi-wavelet coefficients of the tentative bases. If the hierarchical surplus (defined as the difference between the interpolants of two successive grid levels) is above a user specified threshold, the approximation of $I_{p,n}(q)$ is not sufficiently resolved, and we may increase the tentative multi-wavelet basis functions in unresolved regions of parameter space. This is done by solving (26) with an extended basis. As a further measure of how well we represent $I_{p,n}$, we may compare the variance predicted by the multi-wavelet expansion with the sample variance of $I_{p,n}(q)$.

**Numerical Experiments**

For the numerical experiments, we consider a core-sample saturated with water and oil. We impose a more permeable zone ($10^2 k_0$) in the middle of the core, as shown in Fig. 6a. This more permeable zone is set to study the effects of biofilm on oil recovery when it modifies the rock properties in a thief zone. We consider that initially there is only biofilm in the middle part of the thief zone with a volume fraction of $2.5 \times 10^{-3}$, as shown in Fig. 6b.

![Permeability at t=0](image1)

![Biomass at t=0](image2)

**Fig. 6**—Initial rock permeability (a) and initial volume fraction of biofilm (b).

The core has an initial water saturation of 0.2 and an oil saturation of 0.8. The substrate is injected on the left side of the core at a velocity of 1 m/day. The initial water pressure is set to $10^6$ Pa. Table 3 lists the parameters for the numerical simulations. These boundary conditions, initial conditions, and parameters are selected to perform the diverse numerical studies in a computational time of order of minutes.

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Notation</th>
<th>Values</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>Bacterial death rate</td>
<td>$K_d$</td>
<td>$3.2 \times 10^{-6}$ s$^{-1}$</td>
<td>Kundu et al. (1983)</td>
</tr>
<tr>
<td>Maximum growth rate</td>
<td>$\mu_n$</td>
<td>$1.59 \times 10^{-4}$ s$^{-1}$</td>
<td>Linville et al. (2013)</td>
</tr>
<tr>
<td>Monod-half velocity</td>
<td>$K_n$</td>
<td>$0.915$ kg/m$^3$</td>
<td>Linville et al. (2013)</td>
</tr>
<tr>
<td>Substrate diffusion</td>
<td>$D_n$</td>
<td>$5 \times 10^{-10}$ m$^2$/s</td>
<td>Hardy and Sarko (1993)</td>
</tr>
<tr>
<td>Biomass density</td>
<td>$\rho_b$</td>
<td>$20$ kg/m$^3$</td>
<td>Ro and Neethling (1991)</td>
</tr>
<tr>
<td>Factor Brooks-Corey</td>
<td>$\gamma, \beta$</td>
<td>$2$</td>
<td>Lacerda et al. (2012)</td>
</tr>
<tr>
<td>Critical point</td>
<td>$B_c$</td>
<td>$0.35$</td>
<td>Vandevivere (1995)</td>
</tr>
<tr>
<td>Fitting factor Power law</td>
<td>$\eta$</td>
<td>$2.5$</td>
<td>Hommel et al. (2018)</td>
</tr>
<tr>
<td>Stress coefficient</td>
<td>$K_{str}$</td>
<td>$10^{-10}$ m/Pa·s</td>
<td>Landa-Marbán et al. (2019)</td>
</tr>
<tr>
<td>Critical porosity</td>
<td>$\phi_{crit}$</td>
<td>$0$</td>
<td>Assumed</td>
</tr>
<tr>
<td>Initial permeability</td>
<td>$k_0$</td>
<td>$2.45 \times 10^{-12}$ m$^2$</td>
<td>Assumed</td>
</tr>
<tr>
<td>Biofilm permeability</td>
<td>$k_b$</td>
<td>$2.45 \times 10^{-13}$ m$^2$</td>
<td>Assumed</td>
</tr>
<tr>
<td>Initial porosity</td>
<td>$\phi_0$</td>
<td>$0.21$</td>
<td>Assumed</td>
</tr>
<tr>
<td>Core length</td>
<td>$L$</td>
<td>$0.30$ m</td>
<td>Assumed</td>
</tr>
<tr>
<td>Core radius</td>
<td>$r$</td>
<td>$0.19$ m</td>
<td>Assumed</td>
</tr>
<tr>
<td>Injected substrate concentration</td>
<td>$C_i$</td>
<td>$5$ kg/m$^3$</td>
<td>Assumed</td>
</tr>
<tr>
<td>Injected water velocity</td>
<td>$v_i$</td>
<td>$1.16 \times 10^{-5}$ m/s</td>
<td>Assumed</td>
</tr>
<tr>
<td>Biofilm water content</td>
<td>$\theta_w$</td>
<td>$0.9$</td>
<td>Assumed</td>
</tr>
<tr>
<td>Water viscosity</td>
<td>$\mu_w$</td>
<td>$10^{-3}$ Pa·s</td>
<td>Standard</td>
</tr>
<tr>
<td>Oil viscosity</td>
<td>$\mu_o$</td>
<td>$3.92 \times 10^{-3}$ Pa·s</td>
<td>Standard</td>
</tr>
<tr>
<td>Water density</td>
<td>$\rho_w$</td>
<td>$10^3$ kg/m$^3$</td>
<td>Standard</td>
</tr>
<tr>
<td>Oil density</td>
<td>$\rho_o$</td>
<td>$800$ kg/m$^3$</td>
<td>Standard</td>
</tr>
</tbody>
</table>

Table 3—Table of input variables and model parameters for the verification study.
Impact of Porosity-Permeability Relations. We compare the oil recovery for four porosity-permeability relations: Vandevivere $k_v$ (9), Thullner $k_{th}$ (10), channel $k_c$ (Appendix A), and tube $k_t$ (Appendix A). These four relations include the biofilm permeability. Fig. 7 shows the profiles of these four porosity permeability relations for a highly permeable biofilm and a less permeable biofilm respectively.

From Fig. 7 we can observe that these permeability-porosity relationships give different values of permeability for the same values of rock porosity. Thus, we expect to observe different oil recovery predictions for each of these relationships. Fig. 8 shows the comparison of percentage of oil recovery in comparison with the initial oil in the core for the four different porosity-permeability relations. The biofilm permeability is set to $k_b = 0.1k_0$.

We observe that after one day of water injection, the Vandevivere porosity-permeability relation predicts larger oil recovery than the tube relation. In addition, the channel relation predicts slightly more oil recovery than the Thullner relation. These results are expected from Fig. 7b, where the Vandevivere relationship gives the fastest permeability reduction as a function of the porosity reduction in contrast to the tube relationship which gives the slowest permeability reduction. From Fig. 8, we conclude that the predicted oil recovery differs for the different porosity-permeability curves. The oil recovery could be over- or underestimated depending on the assumed porosity-permeability relation.

Injection Strategies. The study of injection strategies is important for the optimization of the bioplug technology. Comparison of oil recovery for different injection techniques is possible through numerical experiments. In this work, we compare the oil recovery for different substrate injection strategies for the same initial conditions. For this study, we use the $k_c$ porosity-permeability relation. Fig. 9 shows a reference profile where only water is injected and the five different strategies tested.

The reference strategy consists in injecting only water at a constant flux rate. Strategy A consists in injecting water at a higher flux rate after half the time of the experiment. The second injection strategy is found commonly in the literature. Water and substrate are injected continuously at a fixed flux rate and after increasing the water flow to diverge the water flow. Strategy C consists in inverting the direction of the high flux to study the effect on the oil recovery in comparison with Strategy B. For Strategy E, first substrate is injected at a fixed flux rate and double concentration. The system is then closed in order to let the bacteria consume the suspended substrate. Afterwards, the water flux is reactivated at the high rate. We observe that Strategy B and Strategy E use the same amount of substrate. Fig. 10 shows the three different oil predictions for the different injection strategies.

From the previous plot we observe that Strategy E predicts the largest oil recovery. Changing the flow direction at half of the injection (Strategy C) does not result in an improvement on the oil recovery in comparison to Strategy B. Strategies C and D give similar predictions of oil recovery. Fig. 11 shows the simulation results for pressure,
Fig. 8—Percentage of oil recoveries for the different porosity-permeability relationships.

Reference Strategy A Strategy B

| Substrate, and biomass along the core for Strategy D. The cells with water saturation above 0.5 are shown in subplot (a). We observe that after three hours of injection the water has displaced most of the oil in the thief zone. In subplot (b) the cells where oil saturation is above 0.5 after 6 hours of injection are shown. We observe that most of the oil recovered is from the thief zone, but there is still significant amount of oil in the core. Subplot |
Fig. 10—Comparison of the oil recoveries for the different injection strategies over time.

Fig. 11—Saturations, substrate, and biomass profiles along the core.
(c) shows the substrate concentration on the lower half part of the core after 12 hours. As described in Strategy D, the flux direction has changed, so now substrate is being injected on the right side of the core, as shown in the figure. From this subplot we observe that the substrate concentration decreases along the core, where the lower values are located in the thief zone. Given the initial homogeneous volume fraction of biofilm, this has increased more than two orders of magnitude in comparison with the initial volume fraction, showing a greater value on the left side. This result shows that changing the substrate flux direction will not lead to a symmetric biofilm formation.

**Sensitivity Analysis: Numerical Results.** Sensitivity analysis is performed for two test cases based on the methodology presented in a previous section.

**Case I: Mean Permeability with Variability in Seven Parameters.** We consider the single-phase core-scale mathematical model in Table 1 and the same boundary and initial conditions as in Model Test. The quantity of interest is the percentage of oil extraction compared to the initial oil in the core (0-100%). We now consider the case of two-phase flow (oil and water). The injected water velocity changes after 8 hours from \(v_i=1\) m/day to a higher velocity of \(v_i=50\) m/day. Along the length of the core, we discretize with 30 elements, while on the transversal section we discretize with 9 times 9 elements. Due to the computational complexity, only three parameters are considered in the sensitivity study: initial volume fraction, position, and length of the biofilm, as presented in Table 6. The remaining parameters, boundary, and initial conditions are the same as described at the beginning of this section. The quantity of interest is the percentage of oil extraction compared to the initial oil in the core (0-100%). We expect nonsmooth parameter dependence due to sharp changes in velocity.

A sparse grid interpolant on 7 levels yields an estimated error of 0.014 between the two finest levels of resolution, where the relative error has again been estimated as the \(L^2\) norm of the hierarchical surplus divided by the norm 14.

**Parameter** | **Notation** | **Range** | **Total Sobol Index**
---|---|---|---
Maximum growth rate | \(\mu_p\) | \([4.5, 5.5]\times10^{-6}\) s\(^{-1}\) | 0.76
Bacterial death rate | \(K_b\) | \([2.88, 3.52]\times10^{-6}\) s\(^{-1}\) | 0.33
Stress coefficient | \(K_{str}\) | \([1.35, 1.65]\times10^{-9}\) m/Pa·s | 0.017
Power law constant | \(\eta\) | \([2.25, 2.75]\) | 0.0015
Monod-half velocity | \(K_n\) | \([0.82, 1.01]\) kg/m\(^3\) | 0.0017
Biomass density | \(\rho_b\) | \([18, 22]\) kg/m\(^3\) | 0.0012
Nutrient diffusion coefficient | \(D_n\) | \([4.5, 5.5]\times10^{-10}\) m\(^2\)/s | 0.0012

Table 4—Total variability contribution of input parameters to percentage of oil extraction.

The total contribution of each material parameter, isolated, and in combination with the others, are shown in Table 4. Only maximum growth rate and bacterial death rate exhibit significant effect on the variability in \(q\) for the parameter ranges considered here.

**Case II: Two-Phase Flow with Three Variable Parameters.** We now consider the case of two-phase flow (oil and water). The injected water velocity changes after 8 hours from \(v_i=1\) m/day to a higher velocity of \(v_i=50\) m/day. Along the length of the core, we discretize with 30 elements, while on the transversal section we discretize with 9 times 9 elements. Due to the computational complexity, only three parameters are considered in the sensitivity study: initial volume fraction, position, and length of the biofilm, as presented in Table 6. The remaining parameters, boundary, and initial conditions are the same as described at the beginning of this section. The quantity of interest is the percentage of oil extraction compared to the initial oil in the core (0-100%). We expect nonsmooth parameter dependence due to sharp changes in velocity.

A sparse grid interpolant on 7 levels yields an estimated error of 0.014 between the two finest levels of resolution, where the relative error has again been estimated as the \(L^2\) norm of the hierarchical surplus divided by the norm.

<table>
<thead>
<tr>
<th>(\mathcal{L})</th>
<th>(o)</th>
<th>0</th>
<th>1</th>
<th>2</th>
<th>3</th>
<th>4</th>
</tr>
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<tbody>
<tr>
<td>0</td>
<td>-</td>
<td>0.883</td>
<td>0.931</td>
<td>0.974</td>
<td>0.994</td>
<td></td>
</tr>
<tr>
<td>1</td>
<td>0.695</td>
<td>0.901</td>
<td>0.949</td>
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<td>0.997</td>
<td></td>
</tr>
<tr>
<td>2</td>
<td>0.883</td>
<td>0.917</td>
<td>0.952</td>
<td>0.982</td>
<td>0.999</td>
<td></td>
</tr>
</tbody>
</table>

Table 5—Fraction of sample variance in test Case I represented by multi-wavelet expansion of polynomial order \(o\) and resolution level \(\mathcal{L}\).
of the solution itself. A set of \( N = 10^6 \) samples are drawn from the interpolant surrogate model to fit a multi-wavelet model through ordinary least squares. As shown in Table 7, essentially all variance is captured by a multi-wavelet expansion in total level 2 and with piecewise quadratic basis functions. The total Sobol indices for this multi-wavelet representation are shown in Table 6. For the parameter ranges investigated, the variability in oil extraction is dominated by the initial volume fraction of the biofilm. The length and position of the biofilm should however not be entirely ignored.

Table 6—Total contribution of each initial parameter.

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Notation</th>
<th>Range</th>
<th>Total Sobol Index</th>
</tr>
</thead>
<tbody>
<tr>
<td>Volume fraction of the biofilm</td>
<td>( \phi_b )</td>
<td>[2.5( \times )10(^{-4}), 2.5( \times )10(^{-3})]</td>
<td>0.86</td>
</tr>
<tr>
<td>Length of the biofilm</td>
<td>( L_b )</td>
<td>[2, 20] cm</td>
<td>0.18</td>
</tr>
<tr>
<td>Position of the biofilm centre along the core</td>
<td>( X_b )</td>
<td>[11, 19] cm</td>
<td>0.05</td>
</tr>
</tbody>
</table>

Table 7—Fraction of sample variance in test Case II represented by multi-wavelet expansion of polynomial order \( o \) and resolution level \( L \).

<table>
<thead>
<tr>
<th>( L )</th>
<th>0</th>
<th>1</th>
<th>2</th>
<th>3</th>
</tr>
</thead>
<tbody>
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<td>0.887</td>
<td>0.964</td>
<td>0.991</td>
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<tr>
<td>1</td>
<td>0.662</td>
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<td>0.970</td>
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<td>2</td>
<td>0.872</td>
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<td>0.991</td>
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<tr>
<td>3</td>
<td>0.950</td>
<td>0.907</td>
<td>0.971</td>
<td>0.993</td>
</tr>
</tbody>
</table>

Conclusions

In this work we discuss a core-scale mathematical model for single- and two-phase flow including the transport of substrate and changes on the permeability due to formation of biomass. The single-phase laboratory experiment shows that the substrate input changes the plug-potential. The single-phase mathematical model captured the observed response of permeability to changes in the substrate flux. For the two-phase mathematical model, we investigated the effects on the simulated oil recovery for different empirical and upscaled porosity-permeability relations. These results show that the predicted oil recovery could be over- or underestimated depending on the assumed porosity-permeability relation in the mathematical model. Numerical simulations are performed for different injection strategies to study the oil recovery. After simulations, injecting substrate and stopping the water flow to let the bacteria consume the substrate and after reactivating the flow at a higher rate results in the largest oil recovery prediction. The sensitivity analysis for the single-phase core-scale model shows that two parameters are responsible for almost all variability in the mean permeability: maximum growth rate and bacterial death rate. Both parameters need to be estimated in the laboratory with sufficient accuracy to lead to a reliable estimate of the permeability changes. The sensitivity analysis for the two-phase core-scale model demonstrates less impact on the total variability in oil extraction from the initial position of biofilm as compared to the initial volume fraction and length of the biofilm. Thus, the amount of initial biofilm has a higher impact on the oil recovery in comparison to the initial position of biofilm in the thief zone.

Nomenclature

\( a \) = weighting factor, dimensionless
\( B_c, B_r \) = critical and relative porosity, dimensionless
\( B_1, B_2 \) = injected brine concentrations, m/L\(^3\), kg/m\(^3\)
\( C_i, C_n \) = injected substrate concentration and substrate concentration, m/L\(^3\), kg/m\(^3\)
\( d \) = core diameter, L, m
\( D_n \) = substrate diffusion coefficient, L\(^2\)/s
\( E, F, G, V, W, X \) = integration coefficients, dimensionless
\( g \) = gravity, L/t\(^2\), m/s\(^2\)
\( h_b \) = biofilm thickness, dimensionless
\( I_i \) = all subsets of parameters including parameter \( i \) for global sensitivity analysis, dimensionless
\( I_{p,n}(q) \) = hierarchical multidimensional interpolant of \( q \) on total level \( p \geq n \), dimension dep. on \( q \)
\[ I_{MW} = \text{multi-wavelet index set of nonnegative integers} \]

\[ j_n = \text{substrate flux, m/t, L}^2, \text{kg/s m}^2 \]

\[ J_\nu = \text{Bessel function of order } \nu \text{ of first kind, dimensionless} \]

\[ k, k_0, k_b = \text{rock permeability, initial rock permeability, and biofilm permeability, L}^2, \text{mdarcy [m}^2] \]

\[ k_p, k_t, k_{vp} = \text{power law, weighted, and Verma-Pruess permeability relationships, L}^2, \text{mdarcy [m}^2] \]

\[ k_{r,o}, k_{r,w} = \text{oil and water relative permeabilities, dimensionless} \]

\[ \tilde{k}_b = \text{biofilm permeability, dimensionless} \]

\[ K_d = \text{bacterial death rate, t}^{-1}, \text{s}^{-1} \]

\[ K_{str} = \text{stress coefficient, L}^2 \text{t/m, m/s Pa} \]

\[ K_n = \text{Monod half-velocity coefficient, m/L}^3, \text{kg/m}^3 \]

\[ \ell = \text{refinement level, dimensionless} \]

\[ L = \text{number of resolution levels, dimensionless} \]

\[ L, L_b = \text{core and biofilm length, L, cm} \]

\[ n = \text{number of stochastic dimensions, dimensionless} \]

\[ N_0^n = \text{set of n-tuples of nonnegative integers, dimensionless} \]

\[ o = \text{polynomial order of the multi-wavelet expansion, dimensionless} \]

\[ p_o, p_w = \text{oil and water pressure, m/L t}^2, \text{Pa} \]

\[ P = \text{size of multi-wavelet basis} \]

\[ q = \text{quantity of interest for global sensitivity analysis, varying dimension} \]

\[ r = \text{core radius, L, cm} \]

\[ R_f = \text{resistance factor, dimensionless} \]

\[ R_n = \text{substrate reaction term, m/t L}^3, \text{kg/s m}^3 \]

\[ S_o, S_w = \text{saturation of oil and water, dimensionless} \]

\[ S(i) = \text{total Sobol index for parameter } i, \text{dimensionless} \]

\[ t = \text{time, t, days [hours]} \]

\[ T = \text{temperature, T, } °\text{C} \]

\[ v_i, v_o, v_w = \text{injected water velocity, oil, and water velocity, L/t, m/s} \]

\[ w = \text{variable depending on the biofilm thickness, dimensionless} \]

\[ X_b = \text{position of the biofilm centre along the core, L, cm} \]

\[ y_i = \text{general parameter for global sensitivity analysis, dimensionless} \]

\[ Y = \text{interpolation nodes in the parameter space} \]

\[ Y_\nu = \text{Bessel function of order } \nu \text{ of second kind, dimensionless} \]

\[ Y_r = \text{set of interpolation nodes in the parameter space, dimensionless} \]

\[ \beta = \text{fitting factor Brooks-Corey relationship, dimensionless} \]

\[ \gamma = \text{factor Brooks-Corey relationship, dimensionless} \]

\[ \eta = \text{fitting factor Power law, dimensionless} \]

\[ \theta_w = \text{biofilm water content, dimensionless} \]

\[ \mu_o = \text{maximum specific biomass production rate, t}^{-1}, \text{s}^{-1} \]

\[ \mu_o, \mu_w = \text{water and oil viscosity, m/L t, Pa s} \]

\[ \xi = \text{variable dependent on biofilm permeability and porosity, dimensionless} \]

\[ \rho_o, \rho_w = \text{density of biomass, oil, and water, m/L}^3, \text{kg/m}^3 \]

\[ \phi = \text{rock porosity, dimensionless} \]

\[ \phi_b, \phi_f = \text{volume fraction of biofilm and void space outside the biofilm, dimensionless} \]

\[ \varphi_j = j\text{th orthogonal basis function, dimensionless} \]

\[ \psi_{yt} = \text{piecewise linear interpolation functions, dimensionless} \]

\[ \Omega = \text{range of independent parameters for global sensitivity analysis, dimensionless} \]

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where

\[ E = \frac{2w\theta_w [J_0(\xi) Y_0(-w\xi) - J_0(w\xi) Y_0(-\xi)] + \xi\tilde{k}_b [J_0(w\xi) Y_1(-w\xi) + Y_0(-w\xi) J_1(w\xi)]}{4[J_0(\xi) Y_1(-w\xi) + \xi Y_0(-\xi) J_1(w\xi)]} \]

\[ = \frac{\xi \left( 4 \tilde{k}_b + w^2 \right) [J_0(\xi) Y_1(-w\xi) + Y_0(-\xi) J_1(w\xi)]}{4[J_0(\xi) Y_1(-w\xi) + \xi Y_0(-\xi) J_1(w\xi)]}, \quad \text{................................. (A-6)} \]

\[ F = \frac{2\tilde{k}_b \xi Y_1(-w\xi) + w\theta_w Y_0(-\xi)}{2[J_0(\xi) Y_1(-w\xi) + \xi Y_0(-\xi) J_1(w\xi)]}, \quad \text{................................. (A-7)} \]

\[ G = \frac{2\tilde{k}_b \xi J_1(w\xi) + w\theta_w J_0(\xi)}{2[J_0(\xi) Y_1(-w\xi) + \xi Y_0(-\xi) J_1(w\xi)]}, \quad \text{................................. (A-8)} \]

where \( \xi = i\sqrt{\theta_w/\tilde{k}_b} \) and \( i \) is the imaginary number. Here, \( J_\nu(z) \) and \( Y_\nu(z) \) are the Bessel function of order \( \nu \) of first and second kind respectively.
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