Optimal design of experiment for model discrimination

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ABSTRACT

This study investigates the process of chamomile oil extraction from flowers. A parameter-distributed model consisting of a set of partial differential equations is used to describe the governing mass transfer phenomena in a solid-fluid environment under supercritical conditions using carbon dioxide as a solvent. The concept of quasi-one-dimensional flow is applied to reduce the number of spatial dimensions. The flow is assumed to be uniform across any cross-section, although the area available for the fluid phase can vary along the extractor. The physical properties of the solvent are estimated from the Peng-Robinson equation of state. The empirical correlations used in the model are based on the set of laboratory experiments performed under multiple constant operating conditions: $30-40^{\circ}C$, 100-200 bar, and $3.33-6.67 \cdot 10^{-5}$ kg/s. The model-based design of the experiments technique is applied in this work to design an experiment to discriminate between two competing process models.

1. Introduction

Supercritical CO_2 is defined as carbon dioxide that is pressurized and heated above its critical point (31.1 °C, 74 bar). Depending on the operating conditions, the fluid properties such as viscosity and density can vary, which leads to multiple industrial applications of CO_2 .

The supercritical carbon dioxide is commonly used for impregnation as described by Weidner [1], Machado et al. [2] or Fathi et al. [3]. Impregnation is defined as modifying the properties of bulk substances by physically or chemically binding/adsorbing impregnates to a bulk material or surface, such as the hydrophobization of surfaces. The main advantage of using supercritical CO₂ is that after depressurization, it desorbs from the surface and evaporates, leaving a solvent-free product. On the other hand, the main disadvantage of using carbon dioxide for impregnation is the low solubility of many drugs of interest.

Another application of supercritical CO₂ is nanoparticles formation as investigated by Padrela et al. [4], Franco and De Marco [5], Saadati Ardestani et al. [6] or Sodeifian et al. [7]. Supercritical carbon-dioxide-assisted technologies enable the production of different morphologies of different sizes, including nanoparticles and nanocrystals, by modulating operating conditions. Supercritical fluid-based processes have advantages over techniques conventionally employed to produce nanosized particles or crystals, such as reduced use of toxic solvents. Moreover, the CO₂ is completely removed from the final product by simple depressurization.

One of the most popular applications of supercritical CO_2 is the extraction of essential oils, as described by many researchers, for example, by Sodeifian and Sajadian [8], Reverchon et al. [9] or Sovova [10]. Traditional methods, such as distillation and organic solvent extraction, are commonly employed but have drawbacks. Distillation, involves high temperatures that can lead to the thermal degradation

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of heat-sensitive compounds. This limitation has increased the popularity of alternative techniques, such as supercritical fluid extraction. Supercritical CO₂ is appealing due to its distinctive properties: it is inflammable, non-toxic and non-corrosive. Supercritical fluids can exhibit both gasand liquid-like properties, allowing for adjustable dissolving power through changes in operating conditions.

This study investigates the extraction of essential oil from chamomile flowers (Matricaria chamomilla L.) via supercritical fluid extraction techniques and the modelling of this process. Chamomile is a medicinal herb widely cultivated in southern and eastern Europe — in countries such as Germany, Hungary, France and Russia. It can be found outside Europe, for instance in Brazil as discussed by Singh et al. [11]. This plant is distinguished by its hollow, bright gold cones, housing disc or tubular florets and surrounded by about fifteen white ray or ligulate florets. Chamomile has been used for its medicinal benefits, serving as an anti-inflammatory, antioxidant, mild astringent, and healing remedy. Extracts of chamomile are widely used to calm nerves and mitigate anxiety, hysteria, nightmares, insomnia and other sleep-related conditions, according to Srivastava [12]. Orav et al. [13] reported that oil yields from dried chamomile samples ranged from 0.7 to 6.7 mL/kg. The highest yields of essential oil, between 6.1 and 6.7 mL/kg, were derived from chamomile sourced from Latvia and Ukraine. In comparison, chamomile from Armenia exhibited a lower oil content of 0.7 mL/kg.

The literature offers various mathematical models to describe the extraction of valuable compounds from biomass. Selecting a process model is case-to-case dependent and requires analysis of each model's specific assumptions about mass transfer and thermodynamic equilibrium.

Goto et al. [14] presented the Shrinking Core (SC) model, which describes a process of irreversible desorption that is followed by diffusion through the pores of a porous solid. When the mass transfer rate of the solute in the non-extracted inner region is significantly slower than in the outer region, where most of the solute has already been extracted, or when the solute concentration exceeds its solubility in

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the solvent, a distinct boundary may form between the inner and outer regions. As extraction progresses, the core of the inner region shrinks. The model envisions supercritical ${\rm CO}_2$ extraction as a sharp, inward-moving front, with a completely non-extracted core ahead of the front and a fully extracted shell behind it.

Sovova [10] proposed The Broken-and-Intact Cell (BIC) model, which assumes that a portion of the solute, initially stored within plant structures and protected by cell walls, is released during the mechanical breakdown of the material. The solute located in the region of broken cells near the particle surface is directly exposed to the solvent, while the core of the particle contains intact cells with undamaged walls. This model describes three extraction phases: a fast extraction phase for accessible oil, a transient phase, and a slow phase controlled by diffusion. The model has been successfully applied to the extraction of grape oil (Sovová et al. [15]) and caraway oil (Sovova et al. [16]).

The Supercritical Fluid Extraction (SFE) process can be be treated similarly to heat transfer, considering solid particles like hot balls cooling down in a uniform environment. Bartle et al. [17] introduced the hot ball diffusion (HBD) model, where spherical particles with uniformly distributed solute diffuse similarly to heat diffusion. Unlike the BIC model, where solute is readily available on the particle surface, the HBD model is suited for systems with small quantities of extractable materials and is not limited by solubility. The model is particularly relevant when internal diffusion controls mass transfer, allowing results from single particles to be extended to the entire bed under uniform conditions. Reverchon et al. [9] have further elaborated on the HBD model and used it to simulate extraction processes for natural materials.

Reverchon [18] proposed a model for extraction of essential oils, which are mainly located inside the vegetable cells in organules called vacuoles. Only a small fraction of essential oil might be near the particle surface due to the breaking up of cells during grinding or in epidermal hairs located on the leaf surface. The fraction of oil freely available on the particle surface should not be significant in the case of SFE from leaves. Consequently, SFE of essential oil from leaves should be mainly controlled by the internal mass-transfer resistance. Therefore, the external mass-transfer coefficient was neglected in the development of the model of Reverchon [18]. The mass balances were developed in the additional hypotheses that the axial dispersion can be neglected and that the solvent density and flow rate are constant along the bed.

This work builds upon the linear kinetic model suggested by Reverchon [18], deriving fundamental governing equations to develop a comprehensive model for the chamomile oil extraction process. This model aims for control-oriented simplicity, assuming a semi-continuous operation within a cylindrical vessel. The process involves a supercritical solvent being pumped through a fixed bed of finely chopped biomass to extract the solute, followed by separation of the solvent and solute in a flush drum to

collect the extract. Parameters such as pressure (P), feed flow rate (F) and inlet temperature (T^{in}) are adjustable and measurable, while the outlet temperature (T^{out}) and the amount of product at the outlet can only be monitored. Figure 1 presents a simplified process flow diagram.

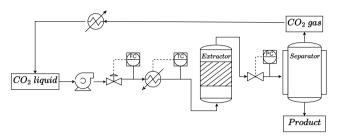


Figure 1: Process flow diagram

2. Materials and methods

2.1. Governing equations

Following the work of Anderson [19], the governing equations for a quasi-one-dimensional flow were derived. A quasi-one-dimensional flow refers to a fluid flow scenario assuming that the flow properties are uniformly distributed across any cross-section. This simplification is typically applied when the flow channel's cross-sectional area changes, such as through irregular shapes or partial filling of an extractor. According to this assumption, velocity and other flow properties change solely in the flow direction.

As discussed by Anderson [20], all flows are compressible, but some of them can be treated as incompressible when the Mach number is smaller than 0.3. This assumption leads to the incompressible condition: $\nabla \cdot u = 0$, which is valid for constant density (strict incompressible) or varying density flow. The assumption allows for removing acoustic waves and large perturbations in density and/or temperature. In the 1-D case, the incompressibility condition becomes $\frac{du}{dz} = 0$, so the fluid velocity is constant.

The set of quasi-one-dimensional governing equations in Cartesian coordinates is described by Equations 1 - 3:

$$\frac{\partial \left(\rho_f A_f\right)}{\partial t} + \frac{\partial \left(\rho_f A_f v\right)}{\partial z} = 0 \tag{1}$$

$$\frac{\partial \left(\rho_f v A_f\right)}{\partial t} + \frac{\partial \left(\rho_f A_f v^2\right)}{\partial z} = -A_f \frac{\partial P}{\partial z} \tag{2}$$

$$\frac{\partial \left(\rho_{f} e A_{f}\right)}{\partial t} + \frac{\partial \left(\rho_{f} A_{f} v e\right)}{\partial z} = -P \frac{\left(A_{f} v\right)}{\partial z} + \frac{\partial}{\partial z} \left(k \frac{\partial T}{\partial z}\right) \tag{3}$$

where ρ_f is the density of the fluid, A_f is the function which describes a change in the cross-section, v is the velocity, P is the total pressure, e is the internal energy of the fluid, t is time and z is the spatial direction.

2.2. Extraction model

2.2.1. Continuity equation

The previously derived quasi-one-dimensional continuity equation (Equation 1) is redefined by incorporating the function $A_f = A\phi$. This modification distinguishes

constant and varying terms, where the varying term accounts for changes in the cross-sectional area available for the fluid. Equation 4 shows the modified continuity equation:

$$\frac{\partial(\rho_f\phi)}{\partial t} + \frac{\partial(\rho_f v A\phi)}{\partial z} = 0 \tag{4}$$

where A is the total cross-section of the extractor and ϕ describes porosity along the extractor.

Assuming that the mass flow rate is constant in time, the temporal derivative becomes the mass flux F, and the spatial derivative can be integrated along z as

$$\int \frac{\partial(\rho_f v A \phi)}{\partial z} dz = F \to F = \rho_f v A \phi \tag{5}$$

To simplify the system dynamics, it is assumed that F is a control variable and affects the whole system instantaneously (due to $\nabla \cdot u = 0$), which allows finding the velocity profile that satisfies mass continuity based on F, ϕ and ρ_f :

$$v = \frac{F}{\rho_f A \phi} \tag{6}$$

Similarly, superficial velocity may be introduced:

$$u = v\phi = \frac{F}{\rho_f A} \tag{7}$$

The fluid density ρ_f can be obtained from the Peng-Robinson equation of state if the temperature and thermodynamic pressure are known along z. Variation in fluid density may occur due to pressure or inlet temperature changes. In a non-isothermal case, in Equations 6 and 7 ρ_f is considered the average fluid density along the extraction column.

2.2.2. Mass balance for the fluid phase

Equation 8 describes the movement of the solute in the system, which is constrained to the axial direction due to the quasi-one-dimensional assumption. Given that the solute concentration in the solvent is negligible, the fluid phase is described as pseudo-homogeneous, with properties identical to those of the solvent itself. It is also assumed that the thermodynamic pressure remains constant throughout the device. The analysis further simplifies the flow dynamics by disregarding the boundary layer near the extractor's inner wall. This leads to a uniform velocity profile across any cross-section perpendicular to the axial direction. Thus, the mass balance equation includes convection, diffusion and kinetic terms representing the fluid phase behaviour:

$$\frac{\partial c_f}{\partial t} + \frac{1}{\phi} \frac{\partial \left(c_f u\right)}{\partial z} = \frac{1 - \phi}{\phi} r_e + \frac{1}{\phi} \frac{\partial}{\partial z} \left(D_e^M \frac{\partial c_f}{\partial z}\right) \tag{8}$$

where c_f represents the solute concentration in the fluid phase, r_e is the mass transfer kinetic term and D_e^M is the axial diffusion coefficient.

2.2.3. Mass balance for the solid phase

As given by Equation 9, the solid phase is considered stationary, without convection and diffusion terms in the mass balance equation. Therefore, the only significant term in this equation is the kinetic term of Equation 10, which connects the solid and fluid phases. For simplicity, the extract is represented by a single pseudo-component:

$$\frac{\partial c_s}{\partial t} = \underbrace{r_e}_{\text{Kinetics}} \tag{9}$$

2.2.4. Kinetic term

As the solvent flows through the bed, CO₂ molecules diffuse into the pores and adsorb on the particle surface to form an external fluid film around the solid particles due to the solvent-solid matrix interactions. The dissolved solute diffuses from the particle's core through the solid-fluid interface, the pore and the film into the bulk. Figure 2 shows the mass transfer mechanism, where the mean solute concentration in the solid phase is denoted as c_s , and the equilibrium concentrations at the solid-fluid interface are denoted as c_s^* and c_n^* for the solid and fluid phases, respectively. The concentration of the solutes in the fluid phase in the centre of the pore is denoted as c_p . As the solute diffuses through the pore, its concentration changes and reaches c_{pf} at the pore opening. Then, the solute diffuses through the film around the particle and reaches bulk concentration c_f . The two-film theory describes the solid-fluid interface inside the pore. The overall mass transfer coefficient can be determined from the relationship between the solute concentration in one phase and its equilibrium concentration.

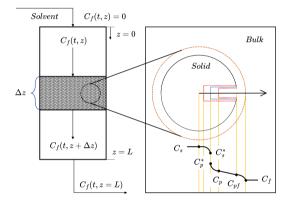


Figure 2: Mass transfer mechanism.

Bulley et al. [21] suggest a process where the driving force for extraction is given by the difference between the concentration of the solute in the bulk, c_f , and in the centre of the pore, c_p^* . The concentration c_p^* is in equilibrium with c_s according to the equilibrium relationship. The rate of extraction is thus $r_e\left(c_f-c_p^*(c_s)\right)$. In contrast, Reverchon [18] proposes a driving force given by the difference between c_s and c_p^* . Concentration c_p^* is determined by the equilibrium relationship with c_f and the extraction rate given by Equation 10:

$$r_e = \frac{D_i}{\mu l^2} \left(c_s - c_p^* \right) \tag{10}$$

where μ is sphericity, l a characteristic dimension of particles that can be defined as l = r/3, r is the mean particle radius, ρ_s is the solid density, D_i corresponds to the overall diffusion coefficient and c_P^* is the concentration at the solid-fluid interface (which according to the internal resistance model is supposed to be at equilibrium with the fluid phase).

According to Bulley et al. [21], a linear equilibrium relationship (Equation 11) can be used to find the equilibrium

concentration of the solute in the fluid phase c_f^* based on the concentration of the solute in the solid phase c_s :

$$c_f^* = k_p c_s \tag{11}$$

The volumetric partition coefficient k_p acts as an equilibrium constant between the solute concentration in one phase and the corresponding equilibrium concentration at the solid-fluid interphase. According to Spiro and Kandiah [22], k_p can be expressed through the mass partition coefficient k_m :

$$k_m = \frac{k_p \rho_s}{\rho_f} \tag{12}$$

According to Reverchon [18], the kinetic term becomes

$$r_e = -\frac{D_i}{\mu l^2} \left(c_s - \frac{\rho_s c_f}{k_m \rho_f} \right) \tag{13}$$

2.2.5. Uneven solute's distribution in the solid phase

Following the idea of the Broken-and-Intact Cell (BIC) model (Sovova [23]), the internal diffusion coefficient D_i is considered to be a product of the reference value of D_i^R and the exponential decay function γ , as given by Equation 14:

$$D_i = D_i^R \gamma(c_s) = D_i^R \exp\left(\Upsilon\left(1 - \frac{c_s}{c_{s0}}\right)\right)$$
 (14)

where Υ describes the curvature of the decay function. Equation 15 describes the final form of the kinetic term:

$$r_e = -\frac{D_i^R \gamma}{\mu l^2} \left(c_s - \frac{\rho_s c_f}{k_m \rho_f} \right) \tag{15}$$

The γ function limits the solute's availability in the solid phase. Similarly to the BIC model, the solute is assumed to be contained in the cells, some of which are open because the cell walls were broken by grinding, with the rest remaining intact. The diffusion of the solute from a particle's core takes more time than the diffusion of the solute close to the outer surface. The same idea can be represented by the decaying internal diffusion coefficient, where the decreasing term is a function of the solute concentration in the solid.

Alternatively, the decay function γ can be interpreted by referring to the Shrinking Core model presented by Goto et al. [14], where the particle radius changes as the amount of solute in the solid phase decreases. As the particle size decreases due to dissolution, the diffusion path increases, which makes the diffusion slower and reduces the value of the diffusion coefficient. This analogy can be applied to Equation 14 to justify the application of a varying diffusion coefficient.

2.2.6. Empirical correlations

The empirical correlations for D_i and Υ were derived by article 1 and validated for temperatures between $30-40^{\circ}C$, pressures between 100-200 bar, and mass flow rates between $3.33-6.67\cdot10^{-5}$ kg/s. Figures 3 and 4 show the results of multiple linear regression applied to solutions of parameter estimation and selected independent variables.

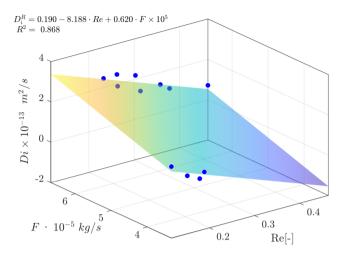


Figure 3: Multiple linear regression $D_i^R = f(Re, F)$

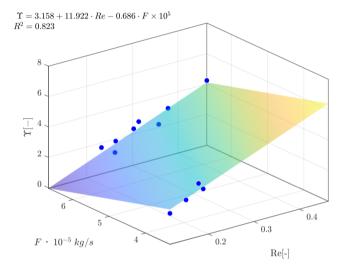


Figure 4: Multiple linear regression $\Upsilon = f(Re, F)$

2.2.7. Heat balance

The heat balance equation describe the evolution of the enthalpy in the system and it is given by Equation 16

$$\frac{\partial \left(\rho_f h A_f\right)}{\partial t} = -\frac{\partial \left(\rho_f h A_f v\right)}{\partial z} + \frac{\partial \left(P A_f\right)}{\partial t} + \frac{\partial}{\partial z} \left(k \frac{\partial T}{\partial z}\right) \tag{16}$$

If the value of enthalpy h is known from the time evolution of the energy equation, and pressure P is known from measurement, then the temperature T can be reconstructed based on the departure function. The departure function is a mathematical function that characterizes the deviation of a thermodynamic property (enthalpy, entropy, and internal energy) of a real substance from that of an ideal gas at the same temperature and pressure. As presented by Gmehling et al. [24], for the Peng-Robinson equation of state, the enthalpy departure function is defined by Equation 17.

$$h - h^{id} = RT \left[T_r(Z - 1) - 2.078(1 + \kappa) \sqrt{\alpha(T)} \ln \left(\frac{Z + \left(1 + \sqrt{2} \right) B}{Z + \left(1 - \sqrt{2} \right) B} \right) \right]$$
(17)

where α is defined as $\left(1 + \kappa \left(1 - \sqrt{T_r}\right)\right)^2$, T_r is the reduced temperature, P_r is the reduced pressure, Z is the compressibility factor, κ is a quadratic function of the acentric factor and B is calculated as $0.07780 \frac{P_r}{T}$.

Equation 17 requires an reference sate, which is assumed to be $T_{ref} = 298.15$ K and $P_{ref} = 1.01325$ bar.

A root-finder can be used to find a value of temperature, which minimizes the difference between the value of enthalpy coming from the heat balance and the departure functions. The root fining procedure to repeated at every time step to find a temperature profile along spatial direction z.

$$\min_{T} \left(\underbrace{h(t,x)}_{\text{Heat balance}} - \underbrace{h(T,P,\rho_f(T,P))}_{\text{Departure function}} \right)^2 \tag{18}$$

2.2.8. Pressure term

As explained in Chapters 2.1, at Low-Mach number conditions, the thermodynamic pressure is nearly constant in space due to the small pressure wave propagation that occurs at the speed of sound. Under such conditions, the term $\partial P/\partial t$ can be approximated by a difference equation, which describes the pressure change in the whole system. The pressure P in the system is considered a state variable, while the pressure in the new time-step P_{in} is considered a control variable.

$$\frac{\partial P}{\partial t} \approx \frac{P_{in} - P}{\Delta t} \tag{19}$$

Such a simplified equation allows for instantaneous pressure change in the system but does not consider a gradual pressure build-up and the effects of pressure losses. In a real system, the dynamics of pressure change would depend on a pump and a back-pressure regulator.

2.2.9. Extraction yield

The process yield is calculated according to Equation 20 as presented by Sovova et al. [16]. The measurement equation evaluates the solute's mass at the extraction unit outlet and sums it up. The integral form of the measurement (Equation 20) can be transformed into the differential form (Equation 21) and augmented with the process model.

$$y = \int_{t_0}^{t_f} \frac{F}{\rho_f} c_f \Big|_{z=L} dt \tag{20}$$

$$\frac{dy}{dt} = \frac{F}{\rho_f} c_f \bigg|_{z=L}$$
(21)

2.2.10. Initial and boundary conditions

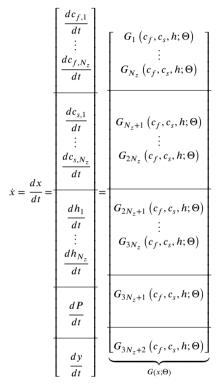
It is assumed that the solvent is free of solute at the beginning of the process $c_{f0}=0$, that all the solid particles have the same initial solute content c_{s0} , and that the system is isothermal, hence the initial state is h_0 . The fluid at the inlet is considered not to contain any solute. The initial and boundary conditions are defined as follows:

$$\begin{split} c_f(t=0,z) &= 0 \qquad c_s(t=0,z) = c_{s0} \qquad h(t=0,z) = h_0 \\ c_f(t,z=0) &= 0 \qquad h(t,z=0) = h_{in} \qquad \frac{\partial c_f(t,z=L)}{\partial x} = 0 \end{split}$$

$$\frac{\partial h(t, z = L)}{\partial x} = 0 \quad c_s(t, z = \{0, L\}) = 0 \quad y(0) = 0 \quad P(0) = P_0$$

2.2.11. Discretization methods

The method of lines is used to transform the process model equations into a set of ODEs denoted as $G(x;\Theta)$. The backward finite difference is used to approximate the first-order derivative, while the central difference scheme approximates the second-order derivative z direction. The length of the fixed bed is divided into N_z , i.e. equally distributed points in the z direction. The state-space model after discretization is denoted as x and defined as follows:



where $x \in \mathbb{R}^{N_x=3N_z+2}$ and $\Theta \in \mathbb{R}^{N_\Theta=N_\theta+N_u}$, N_θ is the number of parameters, N_u is the number of control variables.

For a derivative to be conservative, it must form a telescoping series. In other words, only the boundary terms should remain after adding all terms coming from the discretization over a grid, and the artificial interior points should be cancelled out. Discretization is applied to the conservative form of the model to ensure mass conservation.

3. General function approximators

3.1. Radial Basis Function

An alternative approach to the first principle modelling of physical process is to apply general function approximators and train them based on a dataset. Such an approach has an advantage of not pre-assuming a structure of a model, which results in higher flexibility of a model. On the other hand, this come with a cost of higher number of parameters to be fitted and choosing appropriate function approximator. In this work, a Radial Basis Function (RBF) is used to define $\frac{dc_s}{dt}$ based on a dataset. RBF is a sum real-valued

i	1	2	3	4	5	6	
$\tilde{c}^{c}_{si} Re^{c}_{i}$	-0.294	1.465	-0.303	1.124	-0.025	-0.424	
Re_i^c	0.410	0.281	0.057	0.913	1.185	0.269	
w_{i}	2.001	0.541	6.569	-1.208	-0.663	6.123	
$w_i = \sigma_1^2$	0.433						
σ_1^2	0.051						
$b^{'}$	-0.131						

Table 1Parameters of the RBF

functions (so called kernels) δ whose value depends only on the distance between the input and some fixed point, called a center c, so that $\delta(x) = \delta(||x-c||)$. The distance is usually Euclidean distance, although other metrics are sometimes used. Sums of radial basis functions are typically used to approximate given functions $y(x) = \sum_{i=1}^{N} w_i \delta(||x - c_i||) + b$. Where N corresponds to the number of kernels, w to weight in the summation and b is a bias. The kernel can be defined as a Gaussian, Inverse quadratic, Inverse multi-quadratic, Polyharmonic spline etc. In this work, the two-dimensional Gaussian is used. All the kernels are assumed to have the same shape, which means the all have the same widths in the same direction. Following observation from article 1, the two independent variables are normalized concentration of the solute in the solid phase, defined as $\tilde{c}_s(t) = 1 - \left(\frac{c_s(t)}{c_s}\right)$, and the Reynolds number.

$$\frac{dc_s}{dt} = \sum_{i=1}^{N} w_i \delta(||x - c_i||) + b$$

$$= \sum_{i=1}^{N} w_i \exp\left(-\frac{\left(\tilde{c}_s(t) - \tilde{c}_{si}^c\right)^2}{2\sigma_1^2} - \frac{\left(Re(t) - Re_i^c\right)^2}{2\sigma_2^2}\right) + b$$
(22)

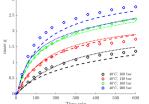
where \tilde{c}_{si}^c and Re_i^c correspond to centrers of each Gaussian kernels as defined above. σ_1 na σ_2 corresponds to the width of each kernel in direction of \bar{c}_{si} and Re_i , respectively. The unknowns of this equation are N, w_i , \tilde{c}_{si}^c and Re_i^c . If N is pre-selected, then the total number of unknowns can calculated as 3N+3. The new process model is defined by substituting Equation 9 with ??. The parameter estimation procedure follows article 1. The obtained parameters are presented in Table 1.

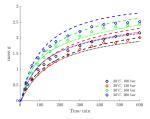
Good agreement between the simulation results and the dataset can be observed, as presented by calculated mean square error and standard deviation as presented in Table 2.

4. Optimal design of experiment for model discrimination

Let's consider two probability distributions $p(y_1)$ and $p(y_2)$ where each represents the Gaussian probability density function for a model.

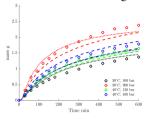
$$p(Y|y_1) = \prod_{i=1}^{n_t} \frac{1}{\sqrt{2\pi\sigma_1^2}} \exp\left(\frac{\sum (Y - y_1(t, x, p))^2}{2\sigma_1^2}\right)$$
(23)





(a) Simulation results at $6.67 \cdot 10^{-5}$ kg/s

(b) Simulation results at $6.67 \cdot 10^{-5}$ kg/s



(c) Simulation results at $3.33 \cdot 10^{-5}$ kg/s

Figure 5: Comparison of models against the dataset

If the ratio of two probability distributions is considered to indicate the measure of similarity, then $\ln\left(\frac{p(Y|y_1)}{p(Y|y_2)}\right)$ becomes a measure of the odds in favour of choosing hypothesis H_1 ($p(Y|y_1)$ is a true model) over hypothesis H_2 ($p(Y|y_2)$ is a true model). Alternatively, the ratio can be interpreted as the information in favour of hypothesis H_1 as opposed to the hypothesis H_2 . The so-called 'weight of evidence' or expected information in favour of choosing H_1 over H_2 can be defined through the Kullback–Leibler divergence and is represented by:

$$I(1:2) = \int_{-\infty}^{\infty} p(Y|y_1) \ln\left(\frac{p(Y|y_1)}{p(Y|y_2)}\right) dY$$
 (24)

The above equation can be written more explicitly as

$$I(1:2) = \int_{-\infty}^{\infty} p(Y|y_1) \left[\sum_{i=1}^{n_t} \left(\ln \left(\frac{\sigma_2}{\sigma_1} \right) - \frac{(Y_i - y_{1i})^2}{2\sigma_1^2} + \frac{(Y_i - y_{2i})^2}{2\sigma_2^2} \right) \right] dY$$

$$= \sum_{i=1}^{n_t} \int_{-\infty}^{\infty} \left(p(Y|y_1) \ln \left(\frac{\sigma_2}{\sigma_1} \right) \right) dY - \sum_{i=1}^{n_t} \int_{-\infty}^{\infty} \left(p(Y|y_1) \frac{(Y_i - y_{1i})^2}{2\sigma_1^2} \right) dY$$

$$+ \sum_{i=1}^{n_t} \int_{-\infty}^{\infty} \left(p(Y|y_1) \frac{(Y_i - y_{2i})^2}{2\sigma_2^2} \right) dY$$
(25)

The equation can be simplified if the expected error is constant for all the measurements: $\mathbb{E}[(Y_i - y_{1i})^2] = \mathbb{E}[\sigma_1^2]$:

$$I(1:2) = \sum_{i=1}^{n_t} \int_{-\infty}^{\infty} \left(p(Y|y_1) \ln \left(\frac{\sigma_2}{\sigma_1} \right) \right) dY - \sum_{i=1}^{n_t} \int_{-\infty}^{\infty} \left(\frac{1}{2} p(Y|y_1) \right) dY + \sum_{i=1}^{n_t} \int_{-\infty}^{\infty} \left(p(Y|y_1) \frac{(Y_i - y_{2i})^2}{2\sigma_2^2} \right) dY$$
 (26)

The first two terms can be simplified by taking a constant in front of integrals and by noticing that $\int p(x)dx = 1$.

$$I(1:2) = n_t \ln\left(\frac{\sigma_2}{\sigma_1}\right) - \frac{n_t}{2} + \sum_{i=1}^{n_t} \frac{1}{2\sigma_2^2} \int_{-\infty}^{\infty} \left(p(Y|y_1) \left(Y_i^2 - 2Y_i y_{2i} + y_{2i}^2\right)\right) dY$$

$$= n_t \ln\left(\frac{\sigma_2}{\sigma_1}\right) - \frac{n_t}{2} + \sum_{i=1}^{n_t} \frac{1}{2\sigma_2^2} \int_{-\infty}^{\infty} \left(p(Y|y_1) \times Y_i^2\right) dY$$

Experiment	1	2	3	4	5	6	7	8	9	10	11	12
Mean squared error	0.0049	0.0100	0.0042	0.1255	0.0401	0.0368	0.0061	0.1292	0.0084	0.0109	0.0028	0.0130
Standard deviation of error	0.0705	0.0619	0.0334	0.1422	0.0791	0.0704	0.0779	0.1313	0.0466	0.1069	0.0287	0.0477

Table 2 Error between experimental data and model predictions

$$-\sum_{i=1}^{n_t} \frac{2y_{2i}}{2\sigma_2^2} \underbrace{\int_{-\infty}^{\infty} \left(p(Y|y_1) \times Y_i\right) dY}_{\text{expected value} = y_{1i}} + \sum_{i=1}^{n_t} \frac{y_{2i}^2}{2\sigma_2^2} \underbrace{\int_{-\infty}^{\infty} p(Y|y_1) dY}_{=1}$$

The remaining integral can be solved by recognizing that $\sigma^2 = \int_{-\infty}^{\infty} X^2 p(X) dX - \mathbb{E}[(X)]^2$, which leads to $\int_{-\infty}^{\infty} Y_i^2 p(Y|y_{1i}) dY = y_{1i}^2 + \sigma_1^2$. Finally the Kullback–Leibler divergence becomes:

$$I(1:2) = n_t \ln\left(\frac{\sigma_2}{\sigma_1}\right) - \frac{n_t}{2} + \sum_{i=1}^{n_t} \frac{y_{1i}^2 + \sigma_1^2}{2\sigma_2^2} \sum_{i=1}^{n_t} \frac{2y_{2i}y_{1i}}{2\sigma_2^2} + \sum_{i=1}^{n_t} \frac{y_{2i}^2}{2\sigma_2^2}$$

$$= n_t \ln\left(\frac{\sigma_2}{\sigma_1}\right) - \frac{n_t}{2} + \frac{n_t}{2\sigma_1^2} + \sum_{i=1}^{n_t} \frac{1}{\sigma_2^2} \left(y_{1i} - y_{2i}\right)^2$$

$$= n_t \ln\left(\frac{\sigma_1}{\sigma_2}\right) - \frac{n_t}{2} + \frac{n_t}{2\sigma_2^2} + \sum_{i=1}^{n_t} \frac{1}{2\sigma_1^2} \left(y_{1i} - y_{2i}\right)^2$$
(28)

While Kulback-Liebler divergence is a statistical distance, it is not a metric on the space of probability distributions. While metrics are symmetric and generalize linear distance, satisfying the triangle inequality, divergences are asymmetric in general and generalize squared distance. In general, $I(1:2) \neq I(2:1)$. By taking into account that the Kullbacl-Liebler divergence is additive for independent distribution, the function j for model discrimination can be defined as

$$j(1,2) = I(1:2) + I(2:1) = \int_{-\infty}^{\infty} [p(Y|y_2) - p(Y|y_1)] \ln \frac{p(Y|y_1)}{p(Y|y_2)} dy$$

$$= \frac{n_t(\sigma_1^2 - \sigma_2^2)}{2\sigma_1^2\sigma_2^2} + \frac{\sigma_1^2 + \sigma_2^2}{2\sigma_1\sigma_2} \times \sum_{i=1}^{n_t} (y_{1i} - y_{2i})^2$$
(29)

The first term of i is independent of changes in v_1 and y_2 , while the second term is equivalent to the sum of squared differences between two model outputs. By maximizing j, the y_1 and y_2 are spread apart. Although both models were fitted with the same dataset, they employ structurally different extraction kinetic terms. These structural differences lead to different outputs, particularly in regions not covered by the dataset.

5. Results

6. Conclusions

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