

Numerical gradient methods for flux identification in a system of conservation laws

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Abstract The identification of the flux for a system of conservation laws is studied from a numerical point of view, on the specific example of chromatography. Different strategies to compute the exact gradient of the discretized optimization problem are developed and compared. Numerical evidence of the convergence of the method is also given in the scalar and binary case. Finally a ternary mixture with real experimental data is studied and the identified isotherm is compared with results obtained by chemical engineers.

Keywords Chromatography · Discrete gradient method · Flux identification · Hyperbolic systems of conservation laws · Measure-valued solutions

1 Introduction

The accuracy of any kind of mathematical model relies on the precise knowledge of all the involved parameters, in the widest sense. Among these one must think of the initial data, which are only partially known in several applications (meteorology, for instance), and state laws, appearing as nonlinearities in partial differential equations. The inverse problem consists in recovering such data or parameters from experimental observations, in order to improve the current model. Specific problems arise when the governing equations are systems of nonlinear hyperbolic conservation laws, which are involved in numerous examples in physics and chemistry. We focus in this paper on the problem of identifying the flux in a system of conservation laws, motivated by the specific example of chromatography.

The chromatography process is a powerful tool to separate or analyze mixtures. It is widely used in chemical industry (pharmaceutical, perfume, and oil industry, ...) to produce relatively high quantities of very pure components. In these conditions, diffusive effects can be neglected, and the behavior can be reasonably modeled by a system involving a mass-balance law (see [1,2] for more aspects concerning models). The process is therefore

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mainly governed by a nonlinear function of the mixture concentrations, the so-called isotherm function, which appears as the flux of the system. Thermodynamical properties of the isotherm ensure that the resulting system is hyperbolic.

The precise knowledge of the isotherm is crucial, from the theoretical viewpoint of physico-chemical modeling, as well as the more practical consideration of accurately governing the experiment to improve separation. Chromatography can be used to identify isotherms, but its application is limited because it requires a rather heavy experimental apparatus. We describe another approach here, which consists in using numerical simulations of the process in order to compare the solutions with experimental outputs.

The problem consists in finding the parameters such that the solution of the model is “as close as possible” to some experimental observation at a given position, which means that a suitable cost function is chosen, typically a least-square estimate of the difference between the solution and the observation. We are thus confronted with an optimization problem, for which descent-type methods are appropriate, so that computing the gradient of the cost function is now the problem we focus on. Two strategies can be followed. The first one consists in directly computing the directional derivative of the cost function. It leads to a formula involving the solution to the linearized version of the original system of conservation laws. The other one is a reformulation of the problem in the spirit of control theory, which introduces an adjoint state, solution to a backward system of linear transport equations. A major problem for both formulations is that the coefficients in the linear equations are discontinuous as soon as shocks arise in the nonlinear solution. Thus, a correct formulation for the gradient is presently out of reach.

Therefore we turn to some discrete formulation of the problem, and we follow a strategy which consists in computing the exact gradient of the discretized problem rather than some arbitrary discretization of the continuous formulation. For the above reasons, we do not try to prove any convergence results, but we give numerical evidence that the schemes we obtain behave nicely when refining the discretization. We also give a few ways of comparing the two strategies.

The paper is organized as follows. In Sect. 2 we recall the physical context, sketch the main properties of the model, and precisely state the identification problem. Section 3 is devoted to the computation of the gradient, from both the formal and numerical point of view. Next we give numerical results for scalar equations and 2×2 systems (in Sect. 4) and show an application on a real set of experimental data for a 3×3 system (in Sect. 5). Finally, some technical computations are collected in the Appendix.

2 Description of the physical problem and model

We recall here the physics and chemistry principles underlying the chromatography process. The separation results from the interaction between two phases in relative motion. The experimental—or industrial—apparatus consists of a column filled with a porous medium in which a neutral solvent circulates at a fixed velocity. A given concentration of the mixture is introduced at the head of the column for a limited time. As the mixture travels down the column, part of it is adsorbed at the grain surface and forms what is called the stationary phase. The separation between the different components of the mixture results from the competition between two phenomena: on the one hand, the mobile phase propagates while obeying the fluid-dynamics laws, on the other hand, the balance between the two phases relies on thermodynamical laws from which the notion of diphasic equilibrium is defined.

2.1 Equations for the direct problem

It turns out that the actual experimental or industrial context allows to simplify the hydrodynamical model greatly. Indeed, since the length of the column is large compared to its diameter, we can safely neglect all radial effects, thus obtaining a one-dimensional model. Next, since the involved concentrations are high, precisely because we wish to observe nonlinear effects, all diffusive phenomena can also be removed. We do not need the energy-conservation equation, because the experiment usually takes place at constant temperature (more generally heat-transfer effects

can be neglected). Finally, the velocity of the mobile phase is assumed to be constant and equal to the vector solvent. This last hypothesis is quite reasonable in the case of liquid incompressible mixtures, and once again this is relevant for many chromatography manipulations (HPLC, High Performance Liquid Chromatography). All these hypotheses allow us to simplify the physics and leave only mass conservation to be considered.

The thermodynamical model invokes complicated mechanisms in order to simulate adsorption of the chemical components. Without going into details (see [1,2]), we merely recall the essential point which is to suppose the existence and uniqueness of a stable equilibrium state for the thermodynamical system of the two phases.

More precisely, for a mixture of p components, we denote by $\mathbf{c}^1, \mathbf{c}^2 \in \mathbb{R}^p$ with $c_i^j \geq 0$ for $1 \leq i \leq p$ and $j = 1, 2$, the concentrations in phases 1 and 2, with respect to the total volume in the column, of the p chemical components. The equilibrium is modeled by a smooth function $\mathbf{h}: \mathbb{R}^p \rightarrow \mathbb{R}^p$, such that $\mathbf{c}^2 = \mathbf{h}(\mathbf{c}^1)$. Furthermore \mathbf{h} has the following properties

$$\mathbf{h}(\mathbf{0}) = \mathbf{0}, \tag{1}$$

$$\mathbf{h}'(\mathbf{c}^1) \text{ is diagonalizable with eigenvalues } \mu_i(\mathbf{c}^1) > 0. \tag{2}$$

The function \mathbf{h} is called an *isotherm*, which comes from the fact that the local equilibrium is reached at constant temperature. Chemistry literature on isotherms is plentiful (see [1,2] and the bibliographies therein). A very classical example of such an isotherm is the *Langmuir isotherm* [3,4]

$$\mathbf{h}_i(\mathbf{c}) = N^* \frac{K_i c_i}{1 + \sum_{i=1}^p K_i c_i}. \tag{3}$$

Here the model is completely determined by $p + 1$ parameters, the so-called *Langmuir coefficients* K_i , which are homogeneous to the inverse of a concentration, and the *saturation coefficient* N^* , which corresponds to some limit concentration when the stationary phase is saturated. We shall mainly use this function, or variants of it, for our numerical simulations throughout the paper.

In order to write the mass-conservation equation we consider the following two phases: the mobile one—phase 1—moves downward with a speed $u > 0$; the adsorbed phase—phase 2—has null speed $v = 0$. Therefore we have

$$\partial_t(\mathbf{c}^1 + \mathbf{c}^2) + \partial_x(u\mathbf{c}^1) = \mathbf{0}. \tag{4}$$

In this experimental setting, we can assume that the equilibrium between the two phases is instantaneous (quasi-static process). At all times and everywhere in the column we can write the closing relation between the concentrations in the two phases, using the isotherm. Equation (4) can then be rewritten using only the mobile-phase concentrations $\mathbf{c} \equiv \mathbf{c}^1$

$$\begin{cases} \partial_x \mathbf{c} + \partial_t \mathbf{F}(\mathbf{c}) = \mathbf{0}, & t \in]0, T[, \quad x \in]0, L[, \\ \mathbf{c}(0, t) = \mathbf{c}_{inj}(t), \\ \mathbf{c}(x, 0) = \mathbf{0}, \end{cases} \tag{5}$$

where the function \mathbf{F} is given by

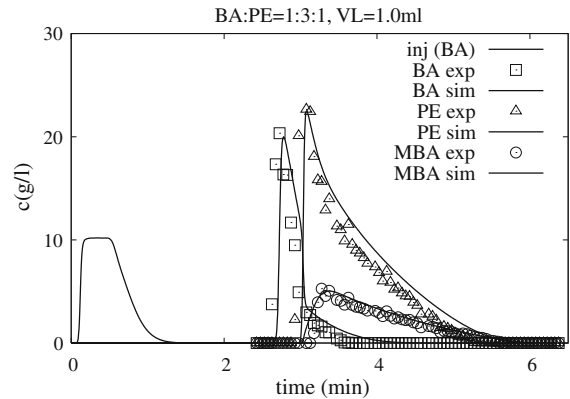
$$\mathbf{F}(\mathbf{c}) = \frac{1}{u} \left(\mathbf{c} + \frac{1 - \varepsilon}{\varepsilon} \mathbf{h}(\mathbf{c}) \right), \tag{6}$$

and $0 < \varepsilon < 1$ is the void fraction of the column.

Because of the properties of the isotherm (2), the system (5) is hyperbolic. Notice that, in (5), the time and space variables exchange their usual roles: the evolution variable is x here. This trick avoids the inversion of the function \mathbf{F} during simulations, and is made possible because the eigenvalues λ_i of \mathbf{F}' turn out to be positive (this is a direct computation using the positivity of the μ_i -s in (2)). This ensures that characteristics always enter the quadrant $\{t > 0, x > 0\}$. To be more specific, notice that one can check that $0 < \lambda_i < u$ for all i , so that the “concentration waves” propagate with a smaller velocity than the inert tracer: we actually model a retention phenomenon.

The injection of the mixture takes place at the head of the column during a limited time: this boundary condition is given along with experimental data and has more or less the shape of a notch. As an example Fig. 1 shows the concentrations measured at the output of the column as a function of time: these curves are called chromatograms.

Fig. 1 Concentrations in BA, PE, and MBA components at the column output



The mixture under consideration has three different components (BA, PE, and MBA) injected in proportion to 1:3:1 with the time injection profile denoted by ‘inj (BA)’ in the figure.

For each component the experimental concentrations are displayed with symbols and the concentrations computed with the model (5) and a special isotherm function provided by the chemical engineers are displayed with lines (see (23), Table 5 and [5]). In this experiment the component BA (\square) has been clearly separated from the other two (\triangle and \circ) with a peak in concentration reaching the bottom of the column half a minute ahead of the other two. Another remarkable feature is the strong hyperbolic behavior of the chromatograms with clearly identifiable shocks. This is due to the high concentration level which makes this dataset a very interesting benchmark that will be further studied in Sect. 5.

2.2 Identification

A problem of major practical interest for chemical engineers is the identification of the physical parameters of the isotherm. Chromatography can be used as a tool to obtain experimental values of the pairs $(\mathbf{c}, \mathbf{h}(\mathbf{c}))$. Several methods can be used for this; for a synthetic review, see [6], and for a more complete discussion see [1, 2]. For competitive isotherms, that is $\mathbf{c} \in \mathbb{R}^p$ with $p > 1$, the so-called *Frontal Analysis* (FA) methods are the only ones available. However, they are very slow and require significant amounts of usually expensive pure chemicals. Furthermore, to obtain relevant information for a function from \mathbb{R}^p to \mathbb{R}^p , one needs a significant number of such pairs $(\mathbf{c}, \mathbf{h}(\mathbf{c}))$, and each of these results from an involved experiment. For instance, the isotherm in Sect. 5 below has been identified by a FA method from a set of 30 experimental points. The PDE model is not used here; the parameters are obtained by direct fitting on the experimental measurements of isotherms.

The alternative approach we use here consists in using as observation the concentration profiles with respect to time at the output of the column (chromatograms). The underlying principle is quite simple: one measures the concentrations in the mixture at the exit of the column. The fit between these experimental data and the model is measured through a cost function $J(\mathbf{F})$, which we choose here as the classical least-squares observation:

$$J(\mathbf{F}) = \frac{1}{2} \int_0^T \|\mathbf{c}_{\mathbf{F}}(L, t) - \mathbf{c}_{\text{obs}}(t)\|^2 dt, \quad (7)$$

where $\mathbf{c}_{\mathbf{F}}$ denotes the solution to (5), and $\|\cdot\|$ is some norm on \mathbb{R}^p . This formulation presumes that we have a complete knowledge of each component of the mixture, as in Fig. 1. If the experimental apparatus does not provide such a measure, the observation is then the total concentration of the mixture, say $\mu_{\text{obs}}(t)$, that is the sum of the different components. In that case, the function J can be chosen as

$$J(\mathbf{F}) = \frac{1}{2} \int_0^T \left| \sum_{i=1}^p \mathbf{c}_{\mathbf{F}_i}(L, t) - \mu_{\text{obs}}(t) \right|^2 dt.$$

For instance, had the dataset in Fig. 1 been recorded as the sum of all three component concentrations, the separation of the last two components would have been hardly recognizable.

However, at the cost of more expensive experimentation, one can obtain such experimental datasets where the concentrations could be measured separately which, of course, enhances the sensitivity of the cost function (7) with respect to the isotherm parameters considerably. We will always use this type of dataset, artificially generated by numerical simulation in Sect. 4, or experimentally obtained in Sect. 5.

At this point, we emphasize the fact that, for practical applications, one cannot expect to identify the isotherm directly as a function from \mathbb{R}^p to \mathbb{R}^p . Indeed, the problem is severely ill-posed as soon as shocks are present in the observation, and no uniqueness is ensured. Instead we prefer to identify the parameters of some given analytical model, which enjoy a convenient physical interpretation, and sometimes can be roughly predicted from experimental data. For instance, consider the Langmuir isotherm (3). It is completely determined by the knowledge of the coefficients K_i and N^* . On the one hand, N^* is a concentration at saturation for the porous medium, on the other hand, the products N^*K_i can be more or less accurately estimated from the data by selecting on the chromatograms the second crossing of each component of the concentration with the time axis. These times correspond to the end of the rarefaction wave and are related to the gradient of the flux function as

$$T_i = \frac{L}{u} \left(1 + \frac{1 - \varepsilon}{\varepsilon} \frac{\partial \mathbf{h}}{\partial \mathbf{c}_i}(0) \right).$$

If we denote by $\alpha = (\alpha_1, \dots, \alpha_q)$ the parameters of the isotherm to be identified, the functional $J(\mathbf{F})$ becomes a function $\tilde{J}(\alpha)$ from \mathbb{R}^q to \mathbb{R}_+ , which has to be minimized over some subset of \mathbb{R}^q . For the Langmuir isotherm, we have $q = p + 1$, but for more realistic (and complex) models q can be much larger than p .

Notice finally that the minimization problem as it has been stated definitely does not fall within classical settings. The function J has no reason to be convex; actually numerical results show that there are local minima. Differentiability is an open problem (see below Sect. 3). Thus one can think of using nonlocal optimization methods to address this problem. There have indeed been some attempts for chromatography, using evolutionary algorithms (see [7,8]). These methods have the advantage of requiring little a priori knowledge on \mathbf{F} (in particular they do not use any kind of derivative), but their main drawback is that they require a large number of direct simulations of the numerical model.

2.3 Discrete formulation

We now turn to the discrete version of the optimization problem, since the experimental data will be provided at a discrete sampling rate. Furthermore the direct model cannot be solved in its continuous version (5). We choose to obtain an approximate solution by discretizing and solving it numerically using a standard finite-volume method that is well-adapted to this type of hyperbolic system.

We define a uniform grid in time and space

$$\begin{aligned} x_0 = 0 < x_1 = \Delta x < \dots < x_k = k\Delta x < \dots < x_K = K\Delta x = L, \\ t_0 = 0 < t_1 = \Delta t < \dots < t_n = n\Delta t < \dots < t_N = N\Delta t = T, \end{aligned}$$

where we will compute the solution using a Godunov scheme:

$$\mathbf{c}_{k+1}^n = \mathbf{c}_k^n - \lambda \left(\mathbf{F}(\mathbf{c}_k^n) - \mathbf{F}(\mathbf{c}_k^{n-1}) \right). \tag{8}$$

Here \mathbf{c}_k^n is an approximation of the mean value of the solution \mathbf{c} at points $x_k = k\Delta x$ and times $t_n = n\Delta t$. In particular, \mathbf{c}_0^n is an approximation of the injection condition in $x_0 = 0$. Similarly, the initial condition $\mathbf{c}(x, t = 0) = 0$ is discretized as $\mathbf{c}_k^0 = 0$ for $k = 0, \dots, K$.

There is no difficulty with this scheme which is known to be of order one in time and space (see [9]) given the nice monotonicity properties of the flux function \mathbf{F} : since all eigenvalues of \mathbf{F}' are positive, we are left with a simple upwind scheme. Notice, however, that contrary to standard usage, it is here the space variable which is the

abscissa along the column that plays the role of evolution variable in the numerical scheme. The cost function to be minimized can be obtained by discretizing (7) on this spatial grid:

$$\tilde{J}(\mathbf{F}) = \frac{1}{2} \Delta t \sum_{n=1}^N \|\mathbf{c}_K^n - \mathbf{c}_{\text{exp}}(t_n)\|^2. \tag{9}$$

One should note, however, that the definition of the discrete cost function is not as straightforward as it seems: first of all, the optimization relies on the assumption that the system (5) correctly models the experiment; hence there exists a function \mathbf{F} , completely determined by a set of parameters α , for which the experimental dataset is its solution—up to measurement precision. Therefore, in the ideal case where the measurements can be obtained with a sampling rate as small as possible and where the exact solution $\mathbf{c}(x, t; \alpha)$ of (5) can be computed for any set of parameters α , the cost function is

$$\tilde{J}(\alpha) = \frac{1}{2} \Delta t \sum_{n=1}^N \|\mathbf{c}(x_K, t_n; \alpha) - \mathbf{c}_{\text{exp}}(t_n)\|^2. \tag{10}$$

One should therefore question the convergence of the minimization algorithm when the sampling rate Δt goes to zero. Do the parameters α minimizing (10) tend, to the limit and in a sense yet to be defined, towards the parameters α which minimize (9)?

This is a very difficult problem, and in any case hardly useful from a practical point of view since experimental data are far from perfect. Furthermore, they are sampled on a time grid that is mostly very coarse with respect to a sampling rate needed to get a good numerical approximation of (5) using (8).

Therefore a good way of measuring the fit of the experiment and the model is to fix the discretization of the cost function J equal to the sampling rate of the experimental data, but to compute the model approximations of $\mathbf{c}(x_K, t_n; \alpha)$ with the numerical scheme, with a discretization fine enough to ensure good convergence. This implies a resampling on a coarser grid of the numerical solution and its gradient which has not been implemented in practice so far. The state of the art consists in interpolating the experimental data on a fine numerical grid using linear interpolation.

Given the high theoretical difficulty of the convergence of the optimization process, we will address in this study two intermediate problems which are interesting for understanding the minimization convergence which we pose directly at the discretized level. Both will be illustrated by numerical simulations.

First we define a discrete cost function

$$\hat{J}(\alpha) = \frac{1}{2} \Delta t \sum_{n=1}^N \|\mathbf{c}_K^n - \hat{\mathbf{c}}_K^n\|^2, \tag{11}$$

where $\hat{\mathbf{c}}_K^n$ is the discrete solution computed with scheme (8) with α set equal to the target value which we denote by $\hat{\alpha}$. This functional has a global minimum in $\hat{\alpha}$ and we will numerically illustrate that our minimization method correctly handles this simple case, and is numerically stable when we let Δt tend to 0.

In a second problem, we define the cost function as

$$\tilde{J}_\delta(\alpha) = \frac{1}{2} \Delta t \sum_{n=1}^N \|\mathbf{c}_K^n - \mathbf{c}_\delta(t_n)\|^2, \tag{12}$$

where the “experimental data” \mathbf{c}_δ is the numerical solution computed with the scheme (8) with α set equal to the target values $\hat{\alpha}$ and a very fine discretization in time and space δ , so that it correctly mimics the exact continuous solution of (5). The values $\mathbf{c}_\delta(t_n)$ are obtained by resampling this solution on the coarser grid of sampling Δt . This time, the global minimum of \tilde{J}_δ is not trivially obtained in $\hat{\alpha}$, except for the limiting case for which $\Delta t = \delta$, where we are back to the previous cost function (11) with $\tilde{J}_\delta(\hat{\alpha}) = 0$. Here again we will numerically illustrate the convergence of the minimization algorithm as Δt goes to δ .

3 Gradient computation

As soon as there is more than one parameter to identify, it is necessary to be able to estimate the gradient of the cost function with respect to these parameters to ensure a good behavior of the descent methods. Finite-difference estimation of the partial derivatives can be used but these introduce additional sampling rates—in the parameter directions—which have to be calibrated.

The gradient of the functional J with respect to \mathbf{F} (or with respect to the parameters α) is hard to study directly starting from the continuous formulation (7)–(5). However, in order to understand the kind of objects we have to deal with, we first perform some formal computations at the continuous level, and present two possible strategies to get a formulation of the gradient. As we shall see, the equivalence of the two formulations is an open problem, and this suggests that the functional is not differentiable in general. Notice by the way that, in the scalar case, it is in some sense Lipschitz continuous. Indeed, several authors, see for instance [10, 11] for conservation laws, [12] for degenerate parabolic equations, provide L^1 dependence results of the entropy solution with respect to the flux. Application of Hölder’s inequality then gives the result for J .

3.1 Continuous equations

First, we compute some kind of directional derivative of $J(\mathbf{F})$: let $\delta\mathbf{F}$ be some admissible direction, at this level one can think of any smooth function. For $\lambda > 0$, we denote by \mathbf{c} (resp. \mathbf{c}_λ) the solution to (5) corresponding to the flux \mathbf{F} (resp. $\mathbf{F} + \lambda\delta\mathbf{F}$), so that

$$\frac{J(\mathbf{F} + \lambda\delta\mathbf{F}) - J(\mathbf{F})}{\lambda} = \int_0^T \left\langle \frac{\mathbf{c}_\lambda(L, t) - \mathbf{c}(L, t)}{\lambda}, \frac{\mathbf{c}_\lambda(L, t) + \mathbf{c}(L, t)}{2} - \mathbf{c}_{\text{obs}}(t) \right\rangle dt. \tag{13}$$

Passing to the limit $\lambda \rightarrow 0$ in this relation is an open problem in general. Indeed on the one hand, we can expect that

$$\frac{\mathbf{c}_\lambda(L, \cdot) + \mathbf{c}(L, \cdot)}{2} - \mathbf{c}_{\text{obs}} \rightarrow \mathbf{c}(L, \cdot) - \mathbf{c}_{\text{obs}} \text{ in } L^1_{\text{loc}}(\mathbb{R}),$$

and this actually holds for scalar conservation laws; see [10, 11]. On the other hand, if $(\mathbf{c}_\lambda(L, t) - \mathbf{c}(L, t))/\lambda$ has a limit, say $\delta\mathbf{c}$, it turns out that $\delta\mathbf{c}$ has to solve the linearized equation

$$\begin{cases} \partial_x \delta\mathbf{c} + \partial_t (\partial_{\mathbf{c}}\mathbf{F}\delta\mathbf{c}) + \partial_t \delta\mathbf{F}(\mathbf{c}) = 0, & t \in]0, T[, \quad x \in]0, L[, \\ \delta\mathbf{c}(0, t) = 0, \quad \delta\mathbf{c}(x, 0) = 0, \end{cases} \tag{14}$$

where $\partial_{\mathbf{c}}\mathbf{F}$ is the matrix of the partial derivatives of \mathbf{F} with respect to the concentrations \mathbf{c}

$$(\partial_{\mathbf{c}}\mathbf{F})_{i,j} = \left(\frac{\partial \mathbf{F}_i}{\partial \mathbf{c}_j} \right). \tag{15}$$

The trouble here is that the solutions to (14) take values in the space of measures in t , so that the only convergence one can hope for is too weak to deal with the product in (13). Several notions of solutions have been developed in this context, in the scalar case. For multidimensional transport equations, see [13–15]. For conservation equations, which are involved here, see [16] for the one-dimensional case and [17] for multidimensional results. Notice that DiPerna–Lions and Ambrosio work with renormalized solutions, and assume (roughly speaking) the divergence of the velocity field to be integrable. This is not the case here, because the velocity involves the derivative of \mathbf{c} , which can be a nonpositive measure if shocks are present. One can find a justification for the weak-convergence property in the scalar case by performing a differentiation with respect to the initial data; see [18]: it makes use of the duality solutions developed in [16].

However, one can say that, at least formally, derivation in the direction $\delta\mathbf{F}$ leads to

$$J'(\mathbf{F}) \delta\mathbf{F} = \int_0^T \langle \mathbf{c}(L, t) - \mathbf{c}_{\text{obs}}(t), \delta\mathbf{c}(L, dt) \rangle,$$

where $\delta \mathbf{c}$ is some solution to the system (14). The first strategy to compute the gradient of J consists therefore in finding some numerical evaluation of (16) and (14). Discretizing (14) is not straightforward; we refer to [19] for some results in the scalar case.

Notice that, if the function \mathbf{F} depends explicitly on a number q of parameters $(\alpha_1, \dots, \alpha_q)$, then, instead of computing the derivative of J in some direction $\delta \mathbf{F}$ which has not been clearly defined, we have to compute the gradient of J with respect to the α_j -s. Therefore, we consider $p \times q$ admissible directions, given by

$$\begin{aligned} \delta \mathbf{F} &= \partial_\alpha \mathbf{F} \text{ (in matrix form),} \\ (\delta \mathbf{F})_{i,j} &= \left(\frac{\partial \mathbf{F}_i}{\partial \alpha_j} \right), \quad i = 1, \dots, p, \quad j = 1, \dots, q, \end{aligned} \tag{16}$$

and the system (14) has to be interpreted now as a matrix-valued equation, where the components of the unknown $\delta \mathbf{c}$ are

$$(\delta \mathbf{c})_{ij} = (\partial_\alpha \mathbf{c})_{ij} = \frac{\partial \mathbf{c}_i}{\partial \alpha_j}, \quad i = 1, \dots, p, \quad j = 1, \dots, q.$$

Explicit expressions for $\delta \mathbf{F}$ and $\partial_\alpha \mathbf{F}$ are given in Appendix C below, for different isotherms.

Another possible expression for $J'(\mathbf{F})$ is obtained by reinterpreting the minimization problem

$$\min_{\mathbf{F}} \frac{1}{2} \int_0^T \|\mathbf{c}_{\mathbf{F}}(L, t) - \mathbf{c}_{\text{obs}}(t)\|^2 dt$$

as a constrained minimization problem:

$$\min \left\{ \tilde{J}(\mathbf{v}) = \frac{1}{2} \int_0^T \|\mathbf{v}(t) - \mathbf{c}_{\text{obs}}(t)\|^2 dt, \quad \mathbf{v}(t) = \mathbf{c}_{\mathbf{F}}(L, t) \text{ solution to (5)} \right\}. \tag{17}$$

This formulation is rather classical in control theory and parameter identification, and previous results on chromatography are based on it (see [6,20,21]). For the sake of completeness, we propose a detailed computation in Appendix A, and merely recall the results here.

The Lagrange multiplier corresponding to the constraint turns out to be a function $\mathbf{p}(x, t)$, namely the solution to the backward linear transport equation

$$\begin{cases} \partial_x \mathbf{p} + \partial_\alpha \mathbf{F}(\mathbf{c})^T \partial_t \mathbf{p} = 0, & t \in]0, T[, \quad x \in]0, L[, \\ \mathbf{p}(L, t) = \mathbf{c}(L, t) - \mathbf{c}_{\text{obs}}(t), \\ \mathbf{p}(x, T) = 0. \end{cases} \tag{18}$$

Using the fact $\tilde{J}(\mathbf{c}_{\mathbf{F}}(L, t)) = J(\mathbf{F})$, and the Lagrangian corresponding to the constraint, we get another formula for the gradient of $J(\mathbf{F})$:

$$J'(\mathbf{F})\delta \mathbf{F} = \int_0^L \delta \mathbf{F}(\mathbf{c}_{\text{ini}}(x))\mathbf{p}(x, 0) dx + \int_0^L \int_0^T \delta \mathbf{F}(\mathbf{c})\partial_t \mathbf{p} dt dx \tag{19}$$

for any \mathbf{p} solution to the adjoint equation (18).

The equivalence between (19) and (16) is justified only for smooth solutions, or under specific assumptions in the scalar case; see [22]. See also [23] for the case of differentiation with respect to the initial condition. A specific problem, when discontinuities occur in the solution \mathbf{c} to (5), is that uniqueness is not ensured for the backward problem (18). Therefore stability problems can arise when discretizing the equations. Also, as mentioned in [23], a crucial point to prove equivalence is a convenient definition of the nonconservative product $\partial_\alpha \mathbf{F} \times \delta \mathbf{c}$, when $\delta \mathbf{c}$ is measure-valued. This may indicate that the functional J is not differentiable in general. The nonuniqueness of the adjoint equation might be related to the notion of subdifferential. For a partial result in this direction, see [22], where the convergence of a viscous regularized problem is studied. This remains a very interesting and difficult open question.

3.2 Discrete version

We propose now a strategy to obtain a numerical approximation for both formulations of the gradient, and also give some ways of comparing them. The key strategy is here to compute the exact gradient of the discretized problem, instead of applying arbitrary schemes to the above systems of PDEs. Actually, we start from a given scheme for the direct problem (5), and then mimic the derivation of the continuous formulæ for the gradient. This actually provides numerical schemes both for (14) and (18), and it turns out that the numerical behavior is good. A reason for this could be that such discretizations implicitly define a nonconservative product which is in some sense consistent with the equations. The detailed understanding of this phenomenon, as well as the convergence of the discrete objects remain unresolved problems up to now. Concerning the adjoint formulation, a few steps in this direction can be found in [22]: we have convergence of the sequence of discrete gradients, in the scalar case, provided the discontinuities of \mathbf{c}_{obs} are exactly observed. The situation is worse for the direct formulation, since we only have some consistency and convergence results for numerical schemes for (14), once again in the scalar case; see [19].

We proceed now to the derivation of the schemes, following the same strategy as in the continuous case. Therefore we compute some exact gradient of the discretized problem. We have the following lemma, which will be proved in Appendix B:

Lemma 1 *The gradient of \tilde{J} is given by*

$$\tilde{J}'\delta\mathbf{F} \equiv \lim_{\alpha \rightarrow 0} \frac{\tilde{J}(\mathbf{F} + \alpha\delta\mathbf{F}) - \tilde{J}(\mathbf{F})}{\alpha} = \Delta t \sum_{n=0}^N (\mathbf{c}_K^n - \mathbf{c}_{\text{exp}}(t_n))^T \delta\mathbf{c}_K^n, \tag{20}$$

where $\delta\mathbf{c}_k^n$ is computed with the scheme

$$\delta\mathbf{c}_{k+1}^n = \delta\mathbf{c}_k^n - \lambda \left((\partial_{\mathbf{c}}\mathbf{F}(\mathbf{c}_k^n)) \delta\mathbf{c}_k^n - (\partial_{\mathbf{c}}\mathbf{F}(\mathbf{c}_k^{n-1})) \delta\mathbf{c}_k^{n-1} \right) - \lambda \left(\delta\mathbf{F}(\mathbf{c}_k^n) - \delta\mathbf{F}(\mathbf{c}_k^{n-1}) \right), \tag{21}$$

with $\delta\mathbf{c}_0^n = 0$ and $\delta\mathbf{c}_k^0 = 0$ as initial and boundary conditions.

By construction, the values obtained for the gradient using this method are rigorously equal to the values obtained using the constrained formulation used until now for this problem (see [6,20,21]):

$$\tilde{J}'(\mathbf{F})\delta\mathbf{F} = -\lambda \sum_{n=1}^{N-1} \sum_{k=0}^{K-1} \delta\mathbf{F}(\mathbf{c}_k^n) \left(\mathbf{a}_{k+1}^n - \mathbf{a}_{k+1}^{n+1} \right) + \lambda \sum_{k=0}^{K-1} \delta\mathbf{F}(\mathbf{c}_k^0) \mathbf{a}_{k+1}^0 - \lambda \sum_{k=0}^{K-1} \delta\mathbf{F}(\mathbf{c}_k^N) \mathbf{a}_{k+1}^{N-1},$$

where $\mathbf{a} = (\mathbf{a}_k^n)_{k,n}$ is given by the so-called adjoint scheme of (8) (see Appendix A)

$$\begin{cases} \mathbf{a}_k^n = \mathbf{a}_{k+1}^n - \lambda \left(\partial_{\mathbf{c}}\mathbf{F}(\mathbf{c}_k^{n+1}) \right)^T \left(\mathbf{a}_{k+1}^n - \mathbf{a}_{k+1}^{n+1} \right), & 0 \leq k < K, 0 \leq n \leq N, \\ \mathbf{a}_K^n = \Delta t \left(\mathbf{c}_K^{n+1} - \mathbf{c}_{\text{exp}}(t_{n+1}) \right), & 0 < n < N, \\ \mathbf{a}_k^N = 0, & 0 \leq k \leq K. \end{cases} \tag{22}$$

Although it is difficult to justify theoretically the computation of the gradient using the measure equation (14) when the discretization goes to 0, its numerical behavior is very stable and can be interpreted in terms of a delta-function approximation as will be seen in the numerical simulations in the following section.

Since they are equivalent in terms of accuracy, it is interesting to compare the two schemes in terms of numerical complexity. In the “direct derivation” method, the unknown $\delta\mathbf{c}$ in the problem (21) is a $p \times q$ matrix which can be computed along with the direct-problem unknown \mathbf{c} which is a p vector. On the other hand, the unknown \mathbf{a} in the adjoint method is only a p vector—instead of a $p \times q$ matrix—but its computation requires storing the direct-problem solution \mathbf{c} for the N times and K abscissa, since the scheme (22) has final boundary conditions in time and space and its coefficients $\partial_{\mathbf{c}}\mathbf{F}(\mathbf{c}_k^n)$ depend on the direct-scheme solution. The memory requirement is then $N \times p \times q$ for the *direct derivation* method and $N \times K \times p$ for the *adjoint* one. For one evaluation of the gradient, both methods require $NK(p + q)$ calls to the isotherm-dependent functions $\delta\mathbf{F}$ and $\partial_{\mathbf{c}}\mathbf{F}$. In terms of

elementary operations they also require of the order of $N \times K \times p \times (q + p)$ multiplications in the *adjoint* case against $N \times K \times q \times p^2 + N \times q \times p$ in the *direct derivation* case. The leading-order term is $\mathcal{O}(p \times N \times K)$ in both cases with a factor $q + p$ in the *adjoint* case and a factor $q \times p$ in the *direct derivation* case. We estimate that, for reasonably small values of the number of parameters q , the huge memory requirement of the *adjoint* method makes it prohibitive compared to the *direct derivation*. Another argument in favor of the direct computation is the possibility of computing both the solution \mathbf{c} and its derivatives $\delta\mathbf{c}$ on the same adaptive grid, while in the adjoint method the solutions of the direct and adjoint problems can have discontinuities in different places. The time varying adaptive grid devised to compute the direct problem cannot be easily used to compute the adjoint solution.

On the other hand, if the goal of the identification problem was the injection profile, instead of the isotherm parameters, the number of unknowns would become large enough to make the adjoint method more efficient than that of the direct derivation (see [23–25]).

4 Numerical verification of the convergence

4.1 Scalar case

For this set of experiments we have simulated a chromatogram using a Langmuir isotherm with the parameters in Table 1.

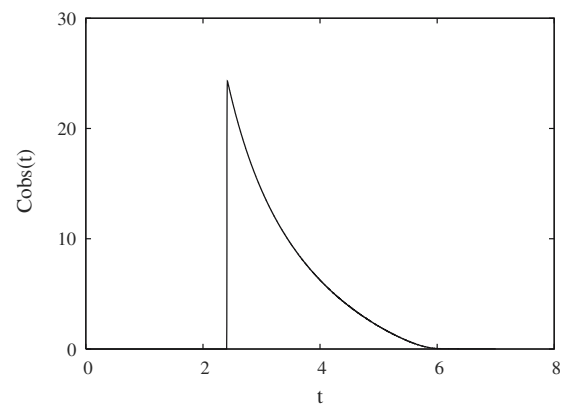
The chromatogram is displayed in Fig. 2. Using this as experimental data, we performed several identifications of the isotherm parameters using different space and time discretizations, from 200 to 4,000 time steps. When we use the same discretization as the one used to generate the dataset, we should and actually do recover the exact parameters of the isotherm. For coarser discretizations the gradient identification converges to a target slightly away from the true parameters but converges towards it with the discretization. This behavior is displayed in Fig. 3 which represents the distance between the result of the identification and the true target in the parameter space (K' and N^*) as a function of the time step on a logarithmic scale.

The two curves correspond to identifications starting from different initial guesses (\square for $K' = 1.1, N^* = 90$) and (\circ for $K' = 0.9, N^* = 150$). This figure shows that the descent algorithm converges towards a minimum

Table 1 Parameters of simulation in the scalar case

N^*	100	U	3
K'	5	Inj	25
FlowRate	1	Porosity	0.59
CFL	0.8	Number of time steps	4000
Diameter	0.39	Length	15.

Fig. 2 Chromatogram simulated with Table 1



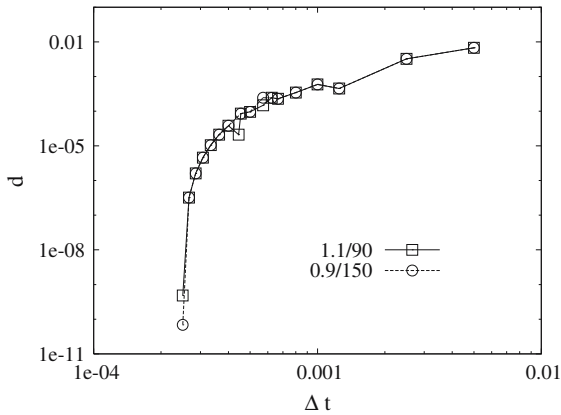


Fig. 3 Distance of the optimization result to the target as a function of discretization parameter Δt

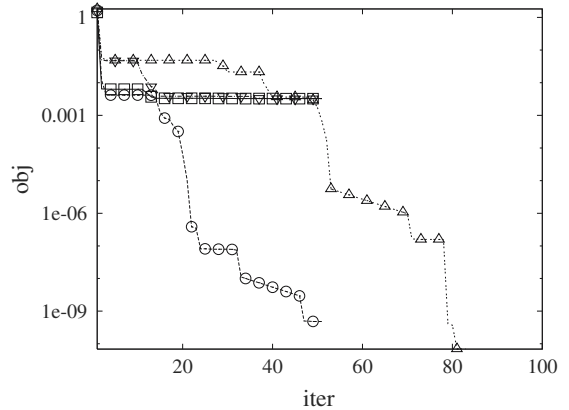


Fig. 4 Convergence paths for $Nt = 400, 4,000$ and the two initial guesses

Table 2 Best parameters for different discretization in the scalar case

Nt	K'	N^*
200	4.91223	97.5226
1000	4.9804	99.5889
2000	4.99402	99.8801
4000	5.	100.00

independent of the starting point; it gets closer to the theoretical target as the discretization is refined. Figure 4 displays the evolution of the objective function with the iterations for two different discretizations and the two initial guesses. When starting from $(K' = 1.1, N^* = 90)$, \square indicate the behavior of the minimization done for 400 time steps and \circ correspond to the 4,000 time-step computation. The minimization starting from the other initial guess $(K' = 0.9, N^* = 150)$ is displayed with ∇ for the 400 time steps computation and Δ for the 4,000 one. The two coarse-grid computations (\square and ∇) converge to roughly the same level of 0.003 for the objective function, while the fine-grid computations (\circ and Δ) reach very small values below 10^{-9} . For both discretizations the first initial guess $(K' = 1.1, N^* = 90)$ indicated by \square and \circ) leads to the convergence state more rapidly than the other initial guess indicated by the triangles.

Using the results of this convergence study, we can also perform a closer analysis of the numerical behavior of the measure equation (21). For four different discretizations ($Nt = 200, 1,000, 2,000,$ and $4,000$) we compute the solution to (21), using the best isotherm parameters for the corresponding discretization, as recorded in Table 2. It is a vector of two components corresponding to the derivatives of the concentration with respect to the two parameters of the isotherms K' and N^* . In Fig. 5 we display the first coordinate at the output of the column $\delta c_{N^*}(L, t)$, in Fig. 6 the second one $\delta c_{K'}(L, t)$ and in Fig. 7 the difference between the simulated and experimental chromatograms $c(L, t) - c_{\text{obs}}(t)$. The left-hand-side graphs display a zoom in the time range of interest. When the convergence isotherm is used, the derivatives converge theoretically towards a delta function located at the shock position ($t = 2.41$) and this behavior is well reproduced numerically in Figs. 5 and 6. Figure 7 allows one to verify that the difference between the experimental and simulated chromatograms decreases in amplitude as the discretization is refined.

We also display the solution of the measure equation computed with parameters of the isotherm slightly away from the best fit, for two discretizations $Nt = 1,000$ in Fig. 8 and 4,000 in Fig. 9. This last case corresponds to the discretization used to generate the “experimental” chromatogram, and the measure is a delta function when the target parameters are used (middle graphs (b) for $N^* = 100$ and $K' = 5$). In fact, derivatives computed for

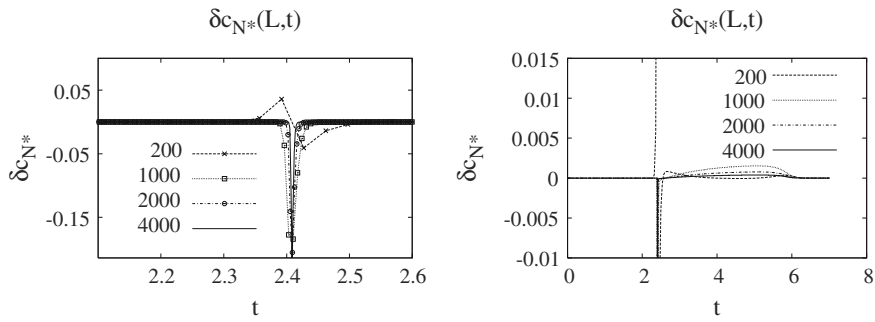


Fig. 5 First component of the measure solution $\delta c_1(L, t)$ for $Nt = 200, 1,000, 2,000,$ and $4,000$ time steps, convergence isotherm used in each case

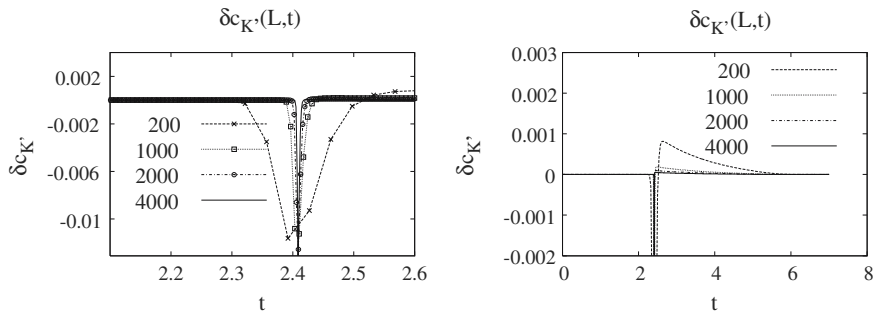


Fig. 6 Second component of the measure solution $\delta c_1(L, t)$ for $Nt = 200, 1,000, 2,000,$ and $4,000$ time steps, convergence isotherm used in each case

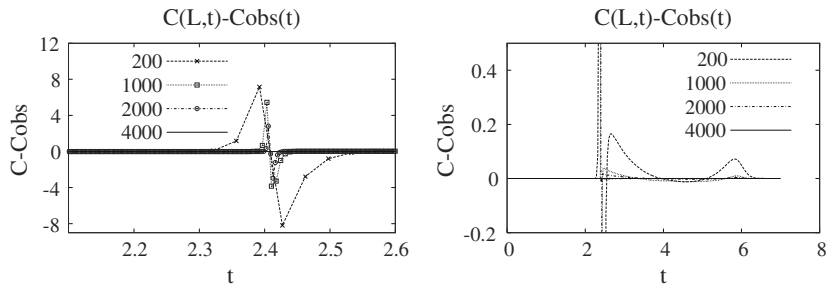


Fig. 7 Difference between the simulated and experimental chromatograms for $Nt = 200, 1,000, 2,000,$ and $4,000$ time steps, convergence isotherm used in each case

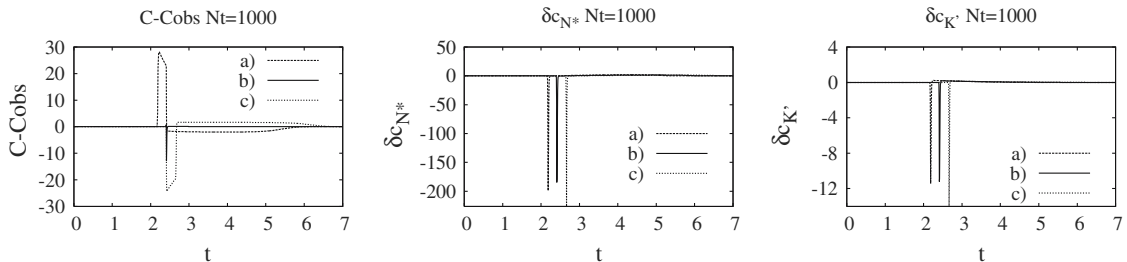


Fig. 8 $c(L, t) - C_{obs}(t)$ and $\delta c(L, t)$ for $Nt = 1,000$ time steps, and three parameter sets (N^*, K') (a) $(90, 4)$, (b) $(100, 5)$, and (c) $(110, 6)$

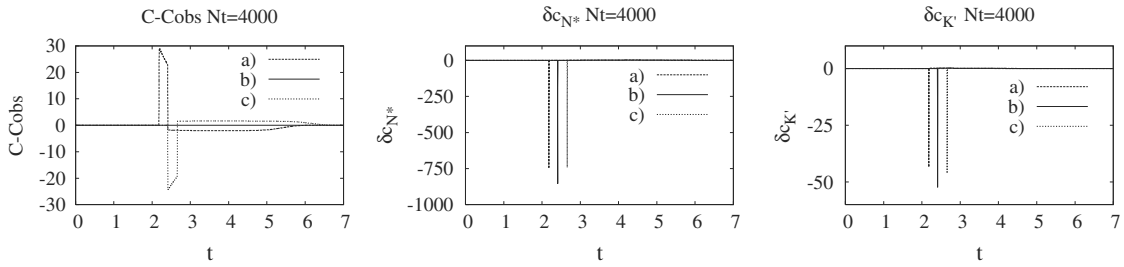


Fig. 9 $c(L, t) - C_{\text{obs}}(t)$ and $\delta c(L, t)$ for $Nt = 4,000$ time steps, and three parameter sets (N^*, K') (a) (90, 4), (b) (100, 5), and (c) (110, 6)

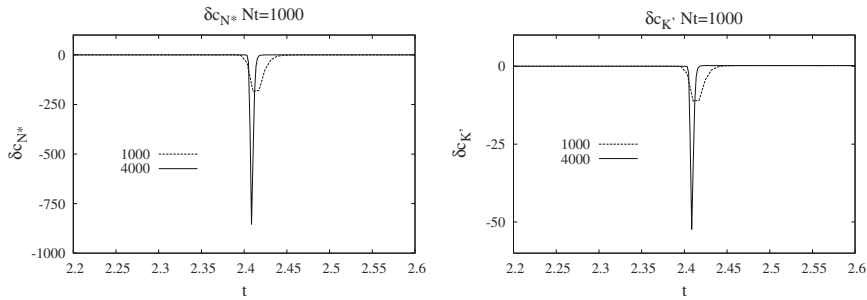


Fig. 10 Zoom of $\delta c(L, t)$ for $Nt = 1,000$ and $4,000$ time steps, and target parameters (100, 5)

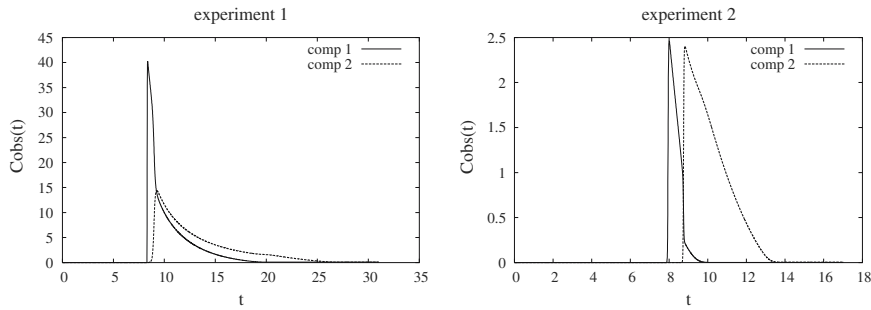


Fig. 11 Chromatograms simulated with Table 3

isotherm parameters away from the target value all look like delta functions but positioned at the wrong time. Figure 10 displays a zoom of the derivatives obtained using the target parameters for the two discretizations. It corroborates clearly the convergence towards a delta function.

4.2 Binary mixture

We simulate in this case the two chromatograms displayed in Fig. 11 using a Bi-Langmuir isotherm for a two-component mixture, namely

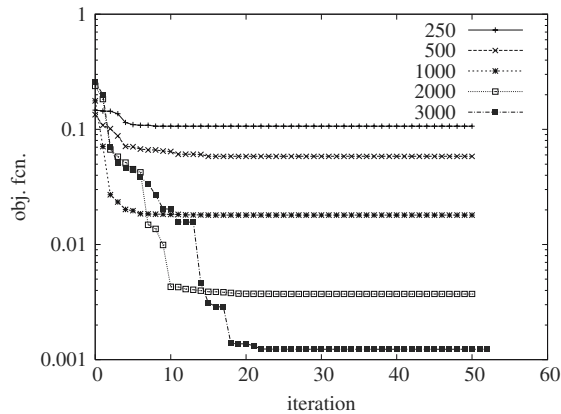
$$h_i(\mathbf{c}) = N_1^* \frac{K_i^1 c_i}{1 + K_1^1 c_1 + K_2^1 c_2} + N_2^* \frac{K_i^2 c_i}{1 + K_1^2 c_1 + K_2^2 c_2}.$$

This model takes into account the fact that there are two possible kinds of adsorption sites; it is determined by six coefficients, whose values are listed in Table 3. We used a very refined discretization of 4,000 time steps for each experiment, the second one until $T = 17$, with a flow rate of 1.2 ml/min, the first and third one until $T = 31$, with a flow rate of 0.6 ml/min. The CFL condition is conservatively ensured by imposing

Table 3 Isotherm and injection parameters for the binary mixture simulation

	Component 1		Component 2			
K_1	0.203564		0.283886		N_1^*	14.30
K_2	0.0325631		0.0407128		N_2^*	120.55
Experiment	1		2		3	
Component	1	2	1	2	1	2
Injection	30.72	30.72	1.49	4.74	3.72	3.72

Fig. 12 Convergence history for different discretizations, starting from $N^* = (10, 130)$



$$\frac{\Delta x}{\Delta t} \sup_{\mathbf{c}} \max_i |\lambda_i(\mathbf{c})| < 0.8,$$

where λ_i denote the eigenvalues of \mathbf{F}' (see (6)).

We then perform identifications of the coefficients N_1^* and N_2^* , the coefficients K_1 and K_2 being kept equal to their theoretical values, which were used to generate the experimental data. Different discretizations and initial guesses are used. All the results tend to confirm the robustness of the identification algorithm. The behavior of the objective function with the iterations of the minimization algorithm is displayed in Fig. 12 for different discretizations. As expected, the value of the objective function reached at convergence diminishes for finer discretization. Less predictably, this simulation shows that convergence is reached faster for coarser discretization. Figure 13 shows the distance of the parameters found by the minimization algorithm with the target parameters (used to generate the experimental chromatogram). There are two curves depicted in this figure, corresponding to two different initial guesses $(N_1^*, N_2^*) = (130, 10)$ and $(110, 20)$. As expected, in both cases, the distance goes to zero when the discretization is refined. The parameters reached by the minimization algorithm are displayed for different discretizations and the two initial guesses in Fig. 14. Even for coarse discretizations the dependence on the initial guess is very small.

5 Application on real datasets

5.1 Experimental identification of isotherms

We test the identification method on real datasets which were extensively studied by Quiñones et al. in [5]. This paper is remarkable because it provides a very important set of isotherms measurements for a three-components mixture; see Fig. 15 below, where each symbol corresponds to a pair $(\mathbf{c}, \mathbf{h}(\mathbf{c}))$, and a different experimental setting. Let us explain briefly how to read this kind of figure. There are three components, 1 = BA = benzylalcohol,

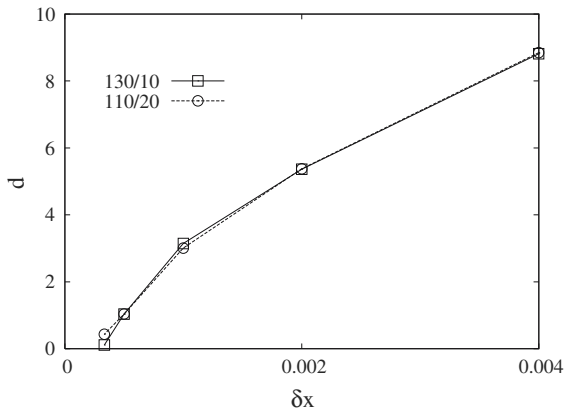


Fig. 13 Distance to target at convergence as a function of the discretization

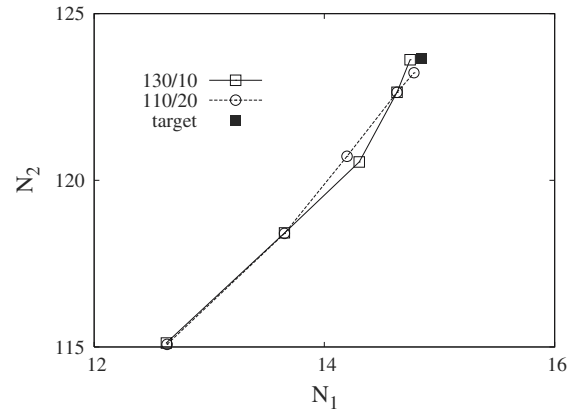


Fig. 14 Target at convergence as a function of the discretization

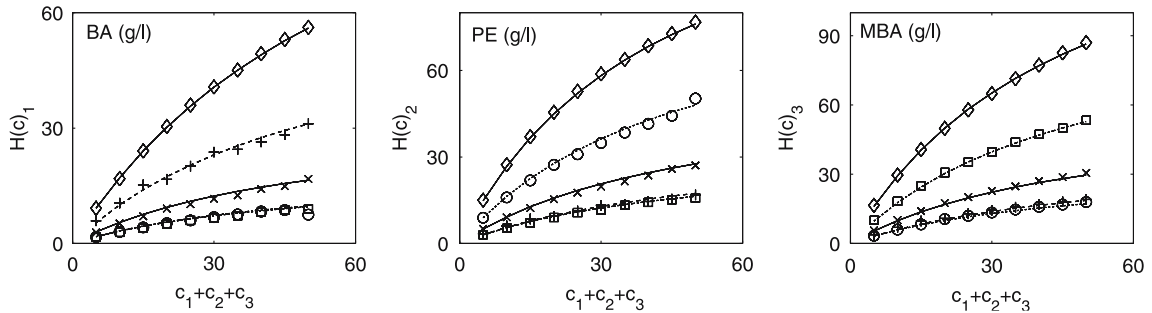


Fig. 15 Isotherms experimental values and model values with the parameters of Table 4

2 = PE = 2-phenylethanol, 3 = MBA = 2-methylbenzylalcohol. For each component, $i = 1, 2, 3$, the adsorbed quantity $h(c)_i$ is displayed as a function of the total amount of mixture ($c_1 + c_2 + c_3$), for five different compositions of the mixture, namely \diamond for single-component, $+$ for 3:1:1 mixture,¹ \times for 1:1:1, \circ for 1:3:1 and \square for 1:1:3 mixture.

In this paper the authors make use of an isotherm function that is slightly different from the Langmuir isotherm, with six independent parameters

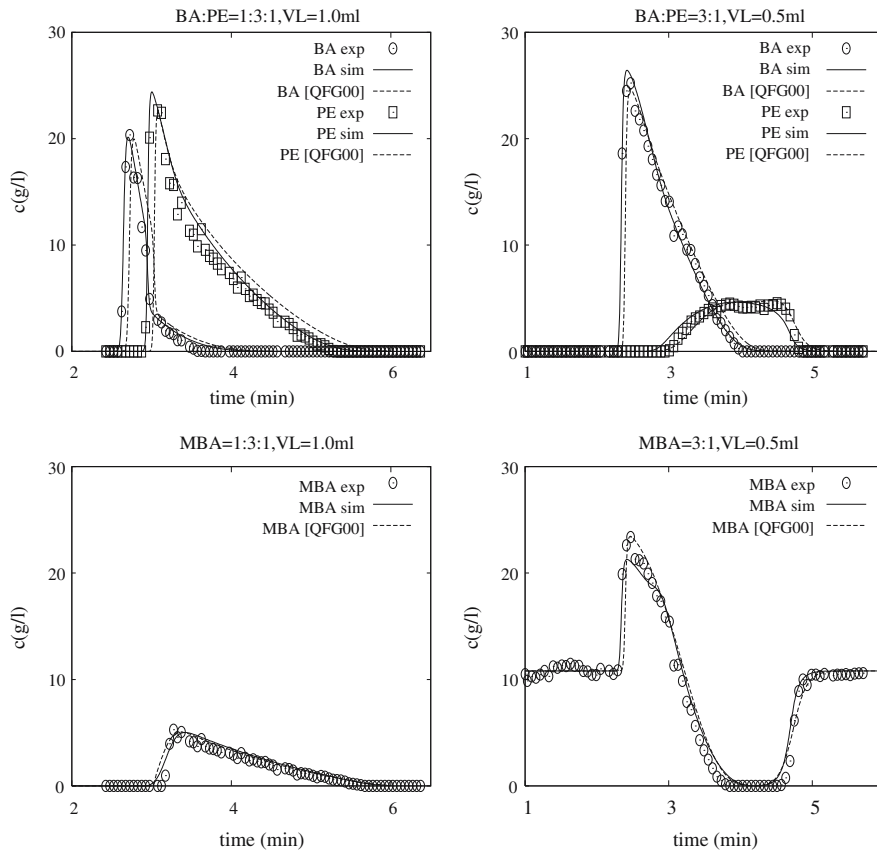
$$h_i(c) = N_i^* \frac{K_i c_i}{1 + \sum_{i=1}^p K_i c_i} \tag{23}$$

They identify the coefficients K_i and N_i^* for $i = 1, \dots, 3$ by fitting the curves obtained with (23) with the experimental points for single-component mixtures (\diamond in Fig. 15), that is, for a total of 30 experiments. The identified parameters are given in Table 4, and all the lines displayed in Fig. 15 represent the isotherm computed for these values. Then simulated chromatograms are computed using (5), the modified isotherm law (23) with the values obtained for the six parameters, and visually compared with a set of seven experimental chromatograms, corresponding to different proportions of the three components and different injection profiles. Two of them are displayed in Fig. 16. The left-hand side corresponds to a mixture of BA, PE, and MPA with proportions 1:3:1, while the right-hand side corresponds to a mixture of BA, PE, and MPA with proportions 3:1. Experimental profiles are represented with symbols. The chromatograms computed with (5), and the parameter values in Table 4 are represented by dashed lines (---), while solid lines (—) represent the chromatograms computed with the values of Table 5.

¹ Three parts of BA, one part of PE and MBA

Table 4 Isotherm parameters for the ternary mixture [5]

Name / #	BA / 1	PE / 2	MBA / 3
K_i	0.01516	0.02341	0.02107
$K'_i = N_i^* K_i$	1.97	3.303	3.55
N_i^*	129.99	141.09	168.50

**Fig. 16** Experimental chromatograms (symbols) and simulated chromatograms using parameters from Table 4 (—) and Table 5 (---). BA and PE components (*top*), MBA component (*bottom*)**Table 5** Isotherm parameters after optimization starting from Table 4

Name/#	BA / 1	PE / 2	MBA / 3
K_i	0.0137	0.0214	0.0206
$K'_i = N_i^* K_i$	1.78046	3.00974	3.47049
N_i^*	129.986	141.07	168.495

Notice that these computations are performed with an injection condition recorded from experimental data as well, see Fig. 17, which is not a step function. The influence of this realistic boundary condition on the shape of the chromatograms is important; see [5] for a more detailed discussion. The left-hand injection profile corresponds to the 1:3:1 experiment, the right-hand one to 3:1.

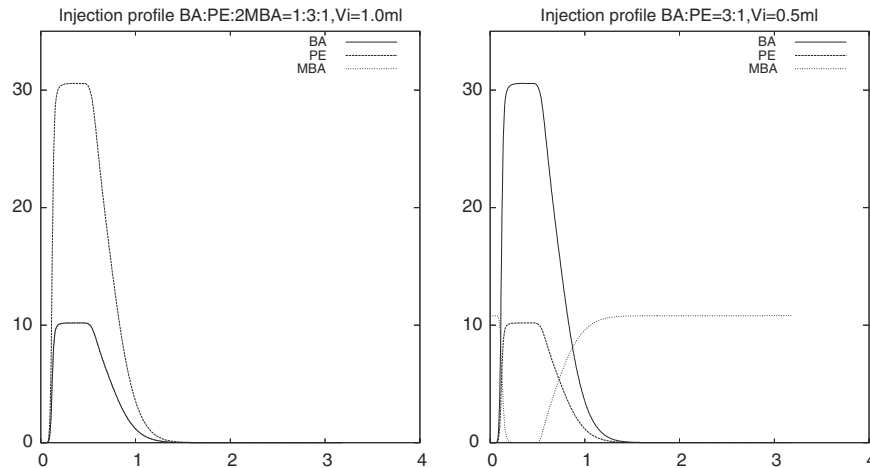


Fig. 17 Injection profiles used for the two experiments

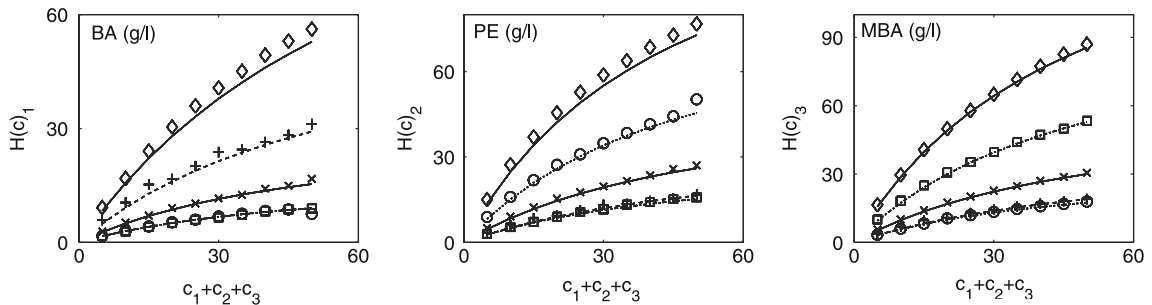


Fig. 18 Isotherms experimental values and model values with the optimized parameters of Table 5

Table 6 Experimental values of K'_i , from [5]

Name / #	BA / 1	PE / 2	MBA / 3
$K'_i = N_i^* K_i$	1.833	3.108	3.511

Remark From now on, all the numerical simulations we use are performed using the Godunov scheme, with 1,000 space points, 1,391 time steps, and the gradient is computed with the direct strategy described in Sect. 3.2.

5.2 Numerical identification of the isotherm

Now we bring our method into operation. We emphasize again that this is a completely different identification strategy from the previous one, which makes use of direct measurements of the isotherm. It is of course very reliable, but unfortunately such experimental estimates for isotherms are seldom available. On the contrary, our method indeed makes use of indirect measurements, such as the two sets of chromatograms in Fig. 16 above as an observation, for which it is much easier to obtain experimental data.

In order to recover the coefficients with the same accuracy on each component and use the information from both datasets, we consider the following weighted cost function, where $\gamma_j = 1/\max_n |\mathbf{c}_{\text{exp}}(t_n)_j|$:

$$\tilde{J}(\alpha) = \frac{1}{2} \Delta t \sum_{\text{exp}=1,2} \sum_{j=1}^3 \gamma_j \sum_{n=1}^N \left| \mathbf{c}_{K_j}^n - \mathbf{c}_{\text{exp}}(t_n)_j \right|^2. \tag{24}$$

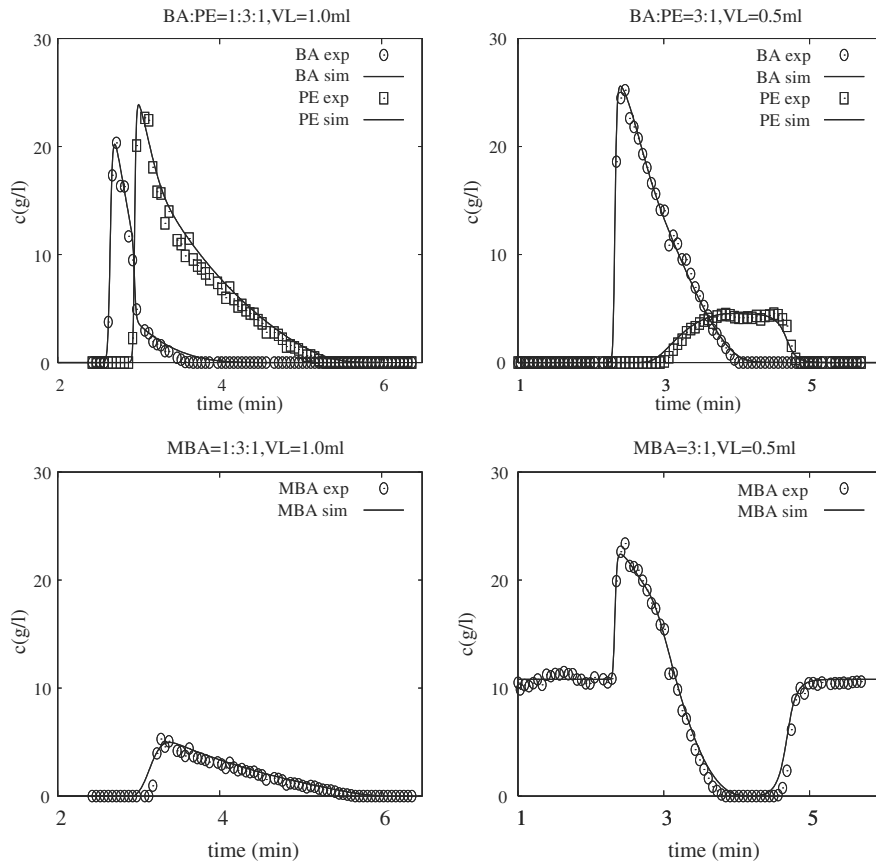


Fig. 19 Experimental chromatograms and simulated chromatograms using optimized parameters from Table 8, BA and PE components (*top*) and MBA component (*bottom*)

Table 7 Isotherm parameters N^* identified from several starting points

Starting point	$N_{BA/1}^*$	$N_{PE/2}^*$	$N_{MBA/3}^*$	Initial cost value	Final cost value
200. 200. 200.	120.937	134.14	169.401	0.206	0.00892
100. 100. 100.	123.724	135.768	158.81	0.411	0.00901
200. 200. 200.	124.059	135.537	159.29	0.432	0.00902
200. 100. 100.	123.342	135.968	157.67	0.292	0.00909
200. 100. 200.	123.373	135.704	159.637	0.174	0.00896
100. 200. 100.	120.003	133.503	173.244	0.122	0.00912
200. 200. 100.	124.167	135.949	157.592	0.237	0.00909
100. 100. 200.	118.536	133.099	178.772	0.210	0.00945
100. 150. 200.	116.171	133.053	182.999	0.046	0.00980
200. 150. 100.	124.46	136.054	156.623	0.071	0.00917
Mean value	122.614	135.441	165.685		
Relative variance	0.09	0.030	0.45		

Table 8 Isotherm parameters for the fifth experiment in Table 7

Name / #	BA / 1	PE / 2	MBA / 3
K_i	0.01486	0.0229	0.02199
$K'_i = N_i^* K_i$	1.833	3.108	3.511
N_i^*	123.373	135.704	159.637

As a first test, we start the descent with the parameters of Table 4 as an initial guess; the value of the cost function is then 0.115, and we optimize on the whole set of six parameters. After convergence of the conjugate-gradient optimization, its value is 0.0107, it has been divided by ten and the parameters of the isotherm are displayed in Table 5. The fit with the experimental isotherm is displayed in Fig. 15, the fit with experimental chromatograms in Fig. 16 with solid lines (—), together with the simulations of [5] (dashed lines - -). Several comments are in order here.

First, it is clear that the fit on the isotherms is worse for simulated parameters. This emphasizes the fact that the kind of data we used to identify are completely different from those of [5]. The concentration range of the chromatograms is not the same as in the isotherm data, and in particular we have no information at all on the single-component adsorption (\diamond in Figs. 15 and 18), for which the simulated parameters are quite poor.

On the other hand, the value of the cost function has been divided by ten, and it is clear from Fig. 16 that shocks are much better identified than from the experimental parameters. This is not very surprising, since the numerical identification process is fully based upon the hyperbolic model, which is very sensitive to the shock position.

Next, we tried another identification series, taking into account some experimental values for the parameters $K'_i = N_i^* K_i$, which are given in [5], and recalled in Table 6 below (in the original paper, the displayed values are $K'_i \times (1 - \epsilon) / \epsilon$, where $\epsilon = 0.59$ is the porosity of the medium). These values, corresponding to “analytical conditions”, for very small injected concentrations, are obtained with good precision, and correspond to the propagation at concentration zero, which is given by $\mathbf{F}'(0)$. Therefore, we performed ten optimizations, keeping the values K'_i constant, and with different starting values for the three remaining parameters N_i^* . The results are collected in Table 7. We notice that the cost function has been divided at least by a factor 10. The third coefficient seems to be more difficult to identify (relative variance 0.45), possibly because its concentration is rather small in one of the experiments. To give an illustration of this result, we display the chromatograms (Fig. 19), for the fifth computation (the values are extracted from the italic line of Table 7, and recalled in Table 8). Concerning isotherms, the results are comparable to those in Fig. 18. The results on the chromatograms are good; in particular, we still have the correct position of shocks, which was not so good in [5].

6 Conclusion

The computation of the gradient required to use a descent-optimization method in the flux identification for a system of conservation laws has now been thoroughly studied numerically. The direct computation of the gradient from the original problem, formulated with partial differential equations is still an open problem, even in the scalar case. However, convergence with respect to the discretization step is indicated by all the numerical tests. Moreover, the application of the identification method proved to be reliable for the case of a 3×3 system fully documented in the chemical engineering literature. We emphasize that additional experimental information on the parameters can drastically improve the results.

On the other hand, it is clear that the complexity of the objective function makes it necessary to combine some global optimization method with the gradient to avoid, for instance, local minima. A first idea is to use the global method to find a good starting point for the gradient (or any descent method), but more intricate couplings are currently under study, using evolutionary algorithms. Finally, the strategies for the formulation of the discrete gra-

dient have to be applied to more complete models, taking into account, for instance, diffusive effects which are also widely used in chemical engineering.

Acknowledgements This work was partially supported by the MESR-CNRS grant ACI-NIM #2003-84 “CHROMALGEMA”

Appendix A: Adjoint formulation for the gradient estimation

In this Appendix we describe a method to compute the gradient of the cost function (7) with a weak formulation. This method has been described and used in [20] and [21], but we give here the full computation, including the case of a non-zero initial state, which is used in Sect. 5.

The gradient with respect to the model parameters will be derived through a Lagrangian formulation. We start with the development in the continuous case which gives a good feeling for the method. We will next present the discrete Lagrangian model which is actually used in the numerical simulations.

A.1 Gradient computation in the continuous case

The constraint for our optimization problem is that $\ll \mathbf{c}_\alpha$ is solution to (5) \gg .

In order to write a Lagrangian for this constraint, we first define a weak formulation: $E(\mathbf{c}, \mathbf{p}, \alpha) = 0$, $\forall \mathbf{p}$, where the functional E is obtained by multiplying the PDE in (5) by a test function \mathbf{p} , as smooth as \mathbf{c} , and integrating by parts on the domain $[0, L] \times [0, T]$:

$$\begin{aligned} E(\mathbf{c}, \mathbf{p}, \alpha) &= \int_0^L \int_0^T \langle (\partial_x \mathbf{c} + \partial_t \mathbf{F}(\mathbf{c})), \mathbf{p} \rangle dt \, dx \\ &= \int_0^T \left([\langle \mathbf{c}, \mathbf{p} \rangle]_0^L - \int_0^L \langle \mathbf{c}, \partial_x \mathbf{p} \rangle dx \right) dt + \int_0^L \left([\langle \mathbf{F}(\mathbf{c}), \mathbf{p} \rangle]_0^T - \int_0^T \langle \mathbf{F}(\mathbf{c}), \partial_t \mathbf{p} \rangle dt \right) dx \\ &= \int_0^T \langle \mathbf{c}(L, t), \mathbf{p}(L, t) \rangle dt - \int_0^T \langle \mathbf{c}_{inj}(t), \mathbf{p}(0, t) \rangle dt + \int_0^L \langle \mathbf{F}(\mathbf{c}(x, T)), \mathbf{p}(x, T) \rangle dx \\ &\quad - \int_0^L \langle \mathbf{F}(\mathbf{c}_{ini}(x)), \mathbf{p}(x, 0) \rangle dx - \int_0^L \int_0^T \langle (\mathbf{c}, \partial_x \mathbf{p}) + (\mathbf{F}(\mathbf{c}), \partial_t \mathbf{p}) \rangle dt \, dx. \end{aligned}$$

The Lagrangian for the constrained minimization problem is

$$L(\mathbf{c}, \mathbf{p}, \alpha) = J(\mathbf{c}) - E(\mathbf{c}, \mathbf{p}, \alpha). \quad (25)$$

We notice that

$$\tilde{J}(\alpha) = J(\mathbf{c}_\alpha) = L(\mathbf{c}_\alpha, \mathbf{p}, \alpha),$$

so that when we apply formally the chain rule, we obtain

$$J'(\alpha) \delta \alpha = \frac{\partial L}{\partial \mathbf{c}}(\mathbf{c}_\alpha, \mathbf{p}, \alpha) \frac{\partial \mathbf{c}}{\partial \alpha} \delta \alpha + \frac{\partial L}{\partial \alpha}(\mathbf{c}_\alpha, \mathbf{p}, \alpha) \delta \alpha.$$

Since it is hard to compute $\frac{\partial \mathbf{c}}{\partial \alpha} \delta \alpha$ (for reasons mentioned in Sect. 3) we will choose \mathbf{p} such that $\frac{\partial L}{\partial \mathbf{c}}(\mathbf{c}_\alpha, \mathbf{p}, \alpha) = 0$ and will next compute $\tilde{J}'(\alpha) \delta \alpha = \frac{\partial L}{\partial \alpha}(\mathbf{c}_\alpha, \mathbf{p}, \alpha) \delta \alpha$ for this special \mathbf{p} .

When we differentiate the Lagrangian with respect to \mathbf{c} , the terms involving \mathbf{c}_{inj} and \mathbf{c}_{ini} disappear, because they are fixed data of the problem, the term $\langle \mathbf{F}(\mathbf{c}), \partial_t \mathbf{p} \rangle$ leads to $\langle \partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}) \delta \mathbf{c}, \partial_t \mathbf{p} \rangle = \langle \delta \mathbf{c}, (\partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}))^T \partial_t \mathbf{p} \rangle$, so that, for the cost function (7), we have

$$\begin{aligned} \frac{\partial L}{\partial \mathbf{c}} &= \frac{\partial J}{\partial \mathbf{c}} - \frac{\partial E}{\partial \mathbf{c}} = \int_0^T \langle (\mathbf{c}(L, t) - \mathbf{c}_{\text{obs}}(t)), \delta \mathbf{c} \rangle dt - \int_0^T \langle \delta \mathbf{c}, \mathbf{p}(L, t) \rangle dt \\ &\quad - \int_0^L \langle \delta \mathbf{c}, \partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}(x, T))^T \mathbf{p}(x, T) \rangle dx + \int_0^L \int_0^T \langle \delta \mathbf{c}, (\partial_x \mathbf{p} + \partial_{\mathbf{c}} \mathbf{p}(\mathbf{c})^T \partial_t \mathbf{p}) \rangle dt dx. \end{aligned}$$

Putting this equal to 0 for all possible $\delta \mathbf{c}$ can be interpreted as a weak formulation of a linear transport equation for \mathbf{p} , with a boundary condition on $x = L$, readily given by the first two terms in this formula, and a final datum on $t = T$, which reads $\partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}(x, T))^T \mathbf{p}(x, T) = 0$. But since $\partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}(x, T))$ is invertible, we can finally choose the adjoint \mathbf{p} solution to (18).

A formula for the gradient of $\tilde{J}(\alpha)$ is then obtained by picking any \mathbf{p} solution to (18) and computing

$$\begin{aligned} \tilde{J}(\alpha) \delta \alpha &= \frac{\partial L}{\partial \alpha} \delta \alpha = - \frac{\partial E}{\partial \alpha} \delta \alpha \\ &= \int_0^L \delta \mathbf{F}(\mathbf{c}_{\text{ini}}(x)) \mathbf{p}(x, 0) dx - \int_0^L \delta \mathbf{F}(\mathbf{c}(x, T)) \mathbf{p}(x, T) dx + \int_0^L \int_0^T \delta \mathbf{F}(\mathbf{c}) \partial_t \mathbf{p} dt dx. \end{aligned}$$

Here $\delta \mathbf{F}$ is obtained by differentiating \mathbf{F} with respect to the parameters we want to identify (see (16) and Appendix C).

A.2 Discrete formulation

We first write the discrete analogue of formula (25) for the numerical scheme (8)

$$\begin{aligned} \tilde{E}(\mathbf{c}, \mathbf{a}, \alpha) &= \sum_{n=1}^N \sum_{k=0}^{K-1} \langle \mathbf{c}_{k+1}^n - \mathbf{c}_k^n + \lambda (\mathbf{F}(\mathbf{c}_k^n) - \mathbf{F}(\mathbf{c}_k^{n-1})), \mathbf{a}_{k+1}^{n-1} \rangle \\ &= \sum_{n=1}^N \sum_{k=0}^{K-1} \langle \mathbf{c}_{k+1}^n, \mathbf{a}_{k+1}^{n-1} \rangle - \sum_{n=1}^N \sum_{k=0}^{K-1} \langle \mathbf{c}_k^n, \mathbf{a}_{k+1}^{n-1} \rangle + \lambda \sum_{n=1}^N \sum_{k=0}^{K-1} \langle \mathbf{F}(\mathbf{c}_k^n), \mathbf{a}_{k+1}^{n-1} \rangle - \lambda \sum_{n=1}^N \sum_{k=0}^{K-1} \langle \mathbf{F}(\mathbf{c}_k^{n-1}), \mathbf{a}_{k+1}^{n-1} \rangle. \end{aligned} \tag{26}$$

Since we will not differentiate with respect to the terms \mathbf{c}_0^n which are fixed by the injection condition, we set them apart. Furthermore, we move up the k indices in the first sum and the n indices in the fourth sum:

$$\begin{aligned} \tilde{E}(\mathbf{c}, \mathbf{a}, \alpha) &= \sum_{n=1}^N \sum_{k=1}^K \langle \mathbf{c}_k^n, \mathbf{a}_k^{n-1} \rangle - \sum_{n=1}^N \sum_{k=0}^{K-1} \langle \mathbf{c}_k^n, \mathbf{a}_{k+1}^{n-1} \rangle + \lambda \sum_{n=1}^N \sum_{k=0}^{K-1} \langle \mathbf{F}(\mathbf{c}_k^n), \mathbf{a}_{k+1}^{n-1} \rangle - \lambda \sum_{n=0}^{N-1} \sum_{k=0}^{K-1} \langle \mathbf{F}(\mathbf{c}_k^n), \mathbf{a}_{k+1}^n \rangle \\ &= \sum_{n=1}^{N-1} \sum_{k=1}^{K-1} \left\{ \langle \mathbf{c}_k^n, \mathbf{a}_k^{n-1} \rangle - \langle \mathbf{c}_k^n, \mathbf{a}_{k+1}^{n-1} \rangle + \lambda (\langle \mathbf{F}(\mathbf{c}_k^n), \mathbf{a}_{k+1}^{n-1} \rangle - \langle \mathbf{F}(\mathbf{c}_k^n), \mathbf{a}_{k+1}^n \rangle) \right\} \\ &\quad + \sum_{k=1}^K \langle \mathbf{c}_k^N, \mathbf{a}_k^{N-1} \rangle + \sum_{n=1}^{N-1} \langle \mathbf{c}_K^n, \mathbf{a}_K^{n-1} \rangle - \sum_{k=0}^{K-1} \langle \mathbf{c}_k^N, \mathbf{a}_{k+1}^{N-1} \rangle - \sum_{n=1}^{N-1} \langle \mathbf{c}_0^n, \mathbf{a}_1^n \rangle \\ &\quad + \lambda \left(\sum_{k=0}^{K-1} \langle \mathbf{F}(\mathbf{c}_k^N), \mathbf{a}_{k+1}^{N-1} \rangle + \sum_{n=1}^{N-1} \langle \mathbf{F}(\mathbf{c}_0^n), \mathbf{a}_1^{n-1} \rangle - \sum_{k=0}^{K-1} \langle \mathbf{F}(\mathbf{c}_k^0), \mathbf{a}_{k+1}^0 \rangle - \sum_{n=1}^{N-1} \langle \mathbf{F}(\mathbf{c}_0^n), \mathbf{a}_1^n \rangle \right). \end{aligned}$$

As in the continuous case, we write the discrete Lagrangian where the constraint $\ll (\mathbf{c}_k^n) \gg$ is taken into account by $\tilde{E}(\mathbf{c}, \mathbf{a}, \alpha) = 0$. We are led to differentiate $\tilde{L}(\mathbf{c}, \mathbf{a}, \alpha) = \tilde{J}(\mathbf{c}) - \tilde{E}(\mathbf{c}, \mathbf{a}, \alpha)$ with respect to all components of (\mathbf{c}_k^n) . On the one hand, we get for the cost function

$$\frac{\partial \tilde{J}}{\partial \mathbf{c}_k^n} = \begin{cases} \Delta t \langle (\mathbf{c}_K^n - \mathbf{c}_{\text{exp}}(t_n)), \delta \mathbf{c}_k^n \rangle, & 0 \leq n \leq N, \\ 0, & 0 \leq k < K, \quad 0 \leq n \leq N. \end{cases}$$

On the other hand, the derivatives of the weak formulation are given by

$$\begin{aligned} \frac{\partial \tilde{E}}{\partial \mathbf{c}_k^n} &= \langle \delta \mathbf{c}_k^n, \mathbf{a}_k^{n-1} - \mathbf{a}_{k+1}^{n-1} \rangle + \lambda \langle \partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}_k^n) \delta \mathbf{c}_k^n, (\mathbf{a}_{k+1}^{n-1} - \mathbf{a}_{k+1}^n) \rangle \\ &= \langle \delta \mathbf{c}_k^n, \mathbf{a}_k^{n-1} - \mathbf{a}_{k+1}^{n-1} + \lambda (\partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}_k^n))^T (\mathbf{a}_{k+1}^{n-1} - \mathbf{a}_{k+1}^n) \rangle \end{aligned}$$

for $k = 1, \dots, K - 1$ and $n = 1, \dots, N - 1$,

$$\frac{\partial \tilde{E}}{\partial \mathbf{c}_K^n} = \langle \delta \mathbf{c}_K^n, \mathbf{a}_K^{n-1} \rangle, \quad 1 \leq n \leq N - 1,$$

$$\frac{\partial \tilde{E}}{\partial \mathbf{c}_k^N} = \langle \delta \mathbf{c}_k^N, \mathbf{a}_k^{N-1} - \mathbf{a}_{k+1}^{N-1} + \lambda (\partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}_k^N))^T \mathbf{a}_{k+1}^{N-1} \rangle, \quad 1 \leq k \leq K - 1.$$

Imposing that all partial derivatives of $\tilde{L} = \tilde{J} - \tilde{E}$ with respect to \mathbf{c}_k^n must be zero, we obtain the following formulae for \mathbf{a}_k^n :

$$\begin{cases} \mathbf{a}_k^{n-1} = \mathbf{a}_{k+1}^{n-1} - \lambda (\partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}_k^n))^T (\mathbf{a}_{k+1}^{n-1} - \mathbf{a}_{k+1}^n), & 0 \leq k < K, \quad 0 < n \leq N - 1, \\ \mathbf{a}_K^n = \Delta t (\mathbf{c}_K^{n+1} - \mathbf{c}_{exp}(t_{n+1})), & 0 < n < N, \\ \mathbf{a}_k^{N-1} = \mathbf{a}_{k+1}^{N-1} - \lambda (\partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}_k^N))^T \mathbf{a}_{k+1}^{N-1}, & 1 \leq k \leq K - 1. \end{cases} \tag{27}$$

In order to recover a discretization scheme for the continuous backward equation (18) that is compatible with the final condition at $t = T$, we are led to impose $\mathbf{a}_k^N = 0$ for all $0 < k \leq K$, so that the third relation in (27) rewrites exactly as the first one, for $0 < n \leq N$ and $0 \leq k < K$. Thus we obtain (22).

The gradient of \tilde{J} for this adjoint is next computed by plugging it into (26) and differentiating \tilde{E} with respect to α . Only the two terms depending on \mathbf{F} plays a role. We obtain

$$\tilde{J}'(\alpha) \delta \alpha = - \frac{\partial \tilde{E}}{\partial \alpha} \delta \alpha = - \lambda \sum_{n=1}^N \sum_{k=0}^{K-1} \delta \mathbf{F}(\mathbf{c}_k^n) (\mathbf{a}_{k+1}^{n-1} - \mathbf{a}_{k+1}^n) + \lambda \sum_{k=0}^{K-1} \delta \mathbf{F}(\mathbf{c}_k^0) \mathbf{a}_{k+1}^0,$$

where $\delta \mathbf{F} = \partial_{\alpha} \mathbf{F} \delta \alpha$, (see (16) and Appendix C below).

Appendix B: Proof of Lemma 1

We study the limit of the Newton ratio

$$\frac{\tilde{J}(\mathbf{F} + \alpha \delta \mathbf{F}) - \tilde{J}(\mathbf{F})}{\alpha} = \Delta t \sum_{n=0}^N \left\langle \frac{\mathbf{d}_K^n - \mathbf{c}_K^n}{\alpha}, \frac{\mathbf{d}_K^n + \mathbf{c}_K^n}{2} - \mathbf{c}_{exp}(t_n) \right\rangle.$$

where \mathbf{d}_k^n is the solution of the Godunov scheme (8) associated to the perturbed flux $\mathbf{F} + \alpha \delta \mathbf{F}$

$$\mathbf{d}_{k+1}^n = \mathbf{d}_k^n - \lambda \left((\mathbf{F} + \alpha \delta \mathbf{F})(\mathbf{d}_k^n) - (\mathbf{F} + \alpha \delta \mathbf{F})(\mathbf{d}_k^{n-1}) \right).$$

We actually prove that uniformly in $k = 0, \dots, K$

$$\lim_{\alpha \rightarrow 0} \|\mathbf{d}_k - \mathbf{c}_k\| = 0 \quad \text{and} \quad \lim_{\alpha \rightarrow 0} \left\| \frac{\mathbf{d}_k - \mathbf{c}_k}{\alpha} - \delta \mathbf{c}_k \right\| = 0, \tag{28}$$

where $\delta \mathbf{c}_k^n$ is given by the scheme (21) and

$$\|\mathbf{c}_k\| = \max_{n=0, \dots, N} |\mathbf{c}_k^n|.$$

We set $\mathbf{r}_k^n = \mathbf{d}_k^n - \mathbf{c}_k^n$, which verifies

$$\begin{cases} \mathbf{r}_{k+1}^n - \mathbf{r}_k^n + \lambda \left((\mathbf{F} + \alpha \delta \mathbf{F})(\mathbf{d}_k^n) - (\mathbf{F} + \alpha \delta \mathbf{F})(\mathbf{d}_k^{n-1}) \right) - \lambda \left(\mathbf{F}(\mathbf{c}_k^n) - \mathbf{F}(\mathbf{c}_k^{n-1}) \right) = 0, \\ \mathbf{r}_0^n = 0, \\ \mathbf{r}_k^0 = 0. \end{cases}$$

After some algebra, and provided that \mathbf{F} is sufficiently smooth, we obtain that the quantity $\mathbf{z}_k^n = \mathbf{r}_k^n / \alpha$ satisfies

$$\begin{cases} \mathbf{z}_{k+1}^n - (1 - \lambda \partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}_k^n)) \mathbf{z}_k^n - \partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}_k^{n-1}) \mathbf{z}_k^{n-1} + \lambda \left(\delta \mathbf{F}(\mathbf{d}_k^n) - \delta \mathbf{F}(\mathbf{d}_k^{n-1}) \right) = \mathcal{O}(\alpha, \|\mathbf{z}_k\|^2), \\ \mathbf{z}_0^n = 0, \\ \mathbf{z}_k^0 = 0. \end{cases} \tag{29}$$

We prove now that $\|\mathbf{z}^n\| \leq M$. Provided that λ verifies a CFL condition $\lambda < 1 / \|\partial_{\mathbf{c}} \mathbf{F}\|$, we have from (29)

$$\|\mathbf{z}_{k+1}\| \leq \|\mathbf{z}_k\| + a \|\mathbf{z}_k\|^2 + b,$$

with $a = C\lambda\alpha$, where C is a bound of $\|\partial_{\mathbf{c}\mathbf{c}} \mathbf{F}\|$ and $b = 2\lambda\|\delta \mathbf{F}\|$. Using $\|\mathbf{z}_0\| = 0$ we obtain by summation

$$\|\mathbf{z}_K\| \leq a \sum_{k=0}^{K-1} \|\mathbf{z}_k\|^2 + Kb.$$

In order to bound $\|\mathbf{z}_k\|$ by M , we therefore need to have

$$aKM^2 - M + Kb \leq 0 \tag{30}$$

which can be true if $4abK^2 < 1$, in other words if

$$\alpha < \frac{1}{8\lambda^2 C \|\delta \mathbf{F}\| K^2}.$$

In that case the smallest root of (30) provides a bound for $\|\mathbf{z}_k\|$

$$M = \frac{1 - \sqrt{1 - 4abK^2}}{2aK} = \lambda \|\delta \mathbf{F}\| K + \mathcal{O}(\alpha) < \frac{\lambda}{2} \|\delta \mathbf{F}\| K,$$

provided that α is sufficiently small. Since $\mathbf{r}_k^n = \alpha \mathbf{z}_k^n$, this proves the first limit in (28).

Now, for the difference $\Delta_k^n = \mathbf{z}_k^n - \delta \mathbf{c}_k^n$, we obtain in the same way, using the bound on $\|\mathbf{z}_k\|$, and if $\delta \mathbf{F}$ is smooth enough,

$$\Delta_{k+1}^n - \Delta_k^n (1 - \lambda \partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}_k^n)) - \partial_{\mathbf{c}} \mathbf{F}(\mathbf{c}_k^{n-1}) \Delta_k^{n-1} = \mathcal{O}(\alpha).$$

We use again $\lambda < 1 / \|\partial_{\mathbf{c}} \mathbf{F}\|$ to obtain that, for α small enough,

$$\|\Delta_{k+1}\| \leq \|\Delta_k\| + \mathcal{O}(\alpha).$$

Since $\|\Delta_0\| = 0$, this gives in turn

$$\|\Delta_k\| \leq \mathcal{O}(\alpha).$$

Therefore, for a given discretization, $\|\Delta_k\| \rightarrow 0$ when $\alpha \rightarrow 0$. □

Appendix C: Gradient of isotherms

In this Appendix we have put together the explicit formulæ to compute the $\delta \mathbf{F}$ involved in (28) or (20–21), for several isotherm functions \mathbf{h} (see (16) and (6)).

In the case of Langmuir isotherm (3), one can optimize with respect to the parameters N^* and $(K_i)_{i=1}^p$. In that case the partial derivatives are

$$\left(\frac{\partial \mathbf{h}(\mathbf{c})}{\partial N^*} \right)_i = \frac{K_i \mathbf{c}_i}{1 + \sum_{j=1}^p K_j \mathbf{c}_j}, \quad \left(\frac{\partial \mathbf{h}(\mathbf{c})}{\partial K_l} \right)_i = N^* \frac{\delta_{li} \mathbf{c}_i}{1 + \sum_{j=1}^p K_j \mathbf{c}_j} - N^* \frac{K_i \mathbf{c}_i \mathbf{c}_l}{\left(1 + \sum_{j=1}^p K_j \mathbf{c}_j \right)^2}. \tag{31}$$

One can also write this isotherm as a function of N^* and $(K'_i = N^*K_i)_{i=1}^p$.

$$\mathbf{h}_i(\mathbf{c}) = N^* \frac{K'_i \mathbf{c}_i}{N^* + \sum_{j=1}^p K'_j \mathbf{c}_j}.$$

This last option is often more interesting because experimental chromatograms provide us with a reliable and direct estimation of the K'_i values. Partial derivatives with respect to N^* and K'_i are

$$\left(\frac{\partial \mathbf{h}(\mathbf{c})}{\partial N^*} \right)_i = \frac{K'_i \mathbf{c}_i \sum_{j=1}^p K'_j \mathbf{c}_j}{\left(N^* + \sum_{j=1}^p K'_j \mathbf{c}_j \right)^2}, \quad \left(\frac{\partial \mathbf{h}(\mathbf{c})}{\partial K'_l} \right)_i = N^* \frac{\delta_{li} \mathbf{c}_i}{N^* + \sum_{j=1}^p K'_j \mathbf{c}_j} - N^* \frac{K'_i \mathbf{c}_i \mathbf{c}_l}{\left(N^* + \sum_{j=1}^p K'_j \mathbf{c}_j \right)^2}. \quad (32)$$

The Bi-Langmuir isotherm (23) used to validate the binary mixture identification has the following partial derivatives

$$\left(\frac{\partial \mathbf{h}(\mathbf{c})}{\partial N_l^*} \right)_i = \frac{K_l^i \mathbf{c}_i}{1 + \sum_{j=1}^p K_j^m \mathbf{c}_j}, \quad \left(\frac{\partial \mathbf{h}(\mathbf{c})}{\partial K_l^m} \right)_i = N_l^* \frac{\delta_{li} \mathbf{c}_i}{1 + \sum_{j=1}^p K_j^m \mathbf{c}_j} - N_l^* \frac{K_l^m \mathbf{c}_i \mathbf{c}_l}{\left(1 + \sum_{j=1}^p K_j^m \mathbf{c}_j \right)^2}. \quad (33)$$

In the case of the isotherm (23) used to model the experimental data there are $2p$ parameters $(N_i^*)_{i=1, \dots, p}$ and $(K_i)_{i=1, \dots, p}$ versus which the derivatives are the following

$$\left(\frac{\partial \mathbf{h}(\mathbf{c})}{\partial N_l^*} \right)_i = \delta_{li} \frac{K_i \mathbf{c}_i}{1 + \sum_{j=1}^p K_j \mathbf{c}_j}, \quad \left(\frac{\partial \mathbf{h}(\mathbf{c})}{\partial K_l} \right)_i = N_i^* \frac{\delta_{li} \mathbf{c}_i}{1 + \sum_{j=1}^p K_j \mathbf{c}_j} - N_i^* \frac{K_i \mathbf{c}_i \mathbf{c}_l}{\left(1 + \sum_{j=1}^p K_j \mathbf{c}_j \right)^2}. \quad (34)$$

As done in the Langmuir case, this isotherm can be rewritten as a function of the $(N_i^*)_{i=1}^p$ and $(K'_i = N_i^* K_i)_{i=1}^p$

$$\mathbf{h}_i(\mathbf{c}) = \frac{K'_i \mathbf{c}_i}{1 + \sum_{j=1}^p \frac{K'_j \mathbf{c}_j}{N_j^*}}.$$

Partial derivatives with respect to N_i^* and K'_i are

$$\left(\frac{\partial \mathbf{h}(\mathbf{c})}{\partial N_l^*} \right)_i = \frac{K'_i K'_i \mathbf{c}_i \mathbf{c}_l}{N_l^{*2} \left(1 + \sum_{j=1}^p \frac{K'_j \mathbf{c}_j}{N_j^*} \right)^2}, \quad \left(\frac{\partial \mathbf{h}(\mathbf{c})}{\partial K'_l} \right)_i = \frac{\delta_{li} \mathbf{c}_i}{1 + \sum_{j=1}^p \frac{K'_j \mathbf{c}_j}{N_j^*}} - \frac{K'_i \mathbf{c}_i \mathbf{c}_l}{N_l^* \left(1 + \sum_{j=1}^p \frac{K'_j \mathbf{c}_j}{N_j^*} \right)^2}. \quad (35)$$

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