



Article Mass Transport in Membrane Systems: Flow Regime Identification by Fourier Analysis

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Abstract: The numerical calculation of local mass distributions in membrane systems by computational fluid dynamics (CFD) offers indispensable benefits. However, the concept to calculate such distributions in response to separate variations of operation conditions (OCs) makes it difficult to address overall, flow-physics-related questions, which require the consideration of the collective interaction of OCs. It is shown that such understanding-related relationships can be obtained by the analytical solution of the advection–diffusion equation considered. A Fourier series model (FSM) is presented, which provides exact solutions of an advection–diffusion equation for a wide range of OCs. On this basis, a new zeroth-order model is developed, which is very simple and as accurate as the complete FSM for all conditions of practical relevance. Advection-dominated blocked and diffusion-dominated unblocked flow regimes are identified (depending on a Péclet number which compares the flow geometry with a length scale imposed by the flow), which implies relevant requirements for the use of lab results for pilot- and full-scale applications. Analyses reveal the equivalence of variations of OCs, which offers a variety of options to accomplish desired flow regime changes.

Keywords: membrane; mass transport; flow regimes

1. Introduction

Computational studies of membrane systems as illustrated in Figure 1 enable very valuable insight into the effectiveness of performance measures, e.g., with respect to concentration polarization and fouling [1–11], as well as evaluating flow unsteadiness induced by different spacer configurations [12–28]. Such computational insights are invaluable given the experimental challenges associated with directly quantifying such phenomena. In one recent study, Liang et al. [29] used two-dimensional (2D) computational fluid dynamics (CFD; the use of this term refers here to the numerical solution of partial differential equations) simulations to resolve the roles played by bulk flow and slip velocities at a membrane surface for generating shear forces at the membrane surface. From this work, it was determined that the origin of shear induction was less important when compared with the resonant frequency of the perturbations themselves for increasing flux through the disruption of foulant and concentration polarization boundary layers. Similarly valuable insights into improving the hydrodynamic conditions within spiral wound membrane elements were made by Foo et al. [30]. Using CFD, these researchers identified optimum spacer geometries for enhancing water flux (up to a 40% flux enhancement) through a membrane through maximizing unsteady-state shear forces at a membrane surface. Reaching this outcome, as is the case in other similar studies, required computationally intensive evaluations of the hydrodynamic environments within the membrane flow channels. Re-



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Copyright: © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). gardless of the complexities involved, the impacts that such studies have on improving the performance and energy efficiency of membrane processes is obvious.

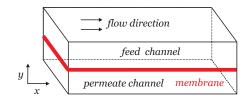


Figure 1. Membrane system illustration.

Despite the indispensable benefits of CFD, the use of such numerical investigation methods is non-trivial, time-consuming, dependent on the availability of numerical codes, and affected by the numerical schemes applied [31–39]. For example, the influence of Péclet numbers (see Equation (6)) has consequences for the equation type considered and the numerical solution of these equations. The equation shows a parabolic (hyperbolic) behavior if the Péclet number is small (large). The numerical solution of such equations becomes increasingly difficult as the Péclet number increases due to the onset of spurious oscillations or excessive numerical damping if standard finite difference or finite element formulations are used [40]. A particular problem is the usual application of Reynoldsaveraged Navier-Stokes (RANS) equations. If spacers are involved, separated flow will appear (recirculation zones), and RANS equations are known to not have the capability to deal with such flow features (the latter requires the use of rather sophisticated, partially resolving flow simulation methods [41-48]). On top of knowledge about detailed mass distributions in membrane systems for specific operation conditions OCs provided by CFD, knowledge of specific flow regimes (which may have desired or undesired characteristics) is very helpful for the membrane system design process. Diffusion-advection processes are characterized by diffusion-dominated and advection-dominated flow regimes, which are separated by corresponding Péclet numbers [40,49]: when this number is small (large), then diffusion (advection) dominates. However, there is a large variety of Péclet numbers that can be considered [50], there are often fuzzy criteria for characterizing different flow regimes (such as Péclet numbers larger or smaller than unity), and the practical consequences of such regime separations are not always obvious. So, there are questions regarding the membrane system design (see also Figure 2):

- Q1. What determines the existence of qualitatively different mass transport regimes? Which interplay of geometry and OCs is implied, which matters to upscaling?
- Q2. How is it possible to accomplish desired flow regime changes by equivalent variations of OCs?
- Q3. For different flow regimes, how is it possible to understand the overall mass transport through membranes and characteristic mass distribution features?

The CFD approach often provides an inappropriate basis to address these questions: separate parameter variations hardly allow conclusions about collective parameter effects.

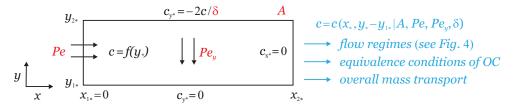


Figure 2. An illustration of the flow configuration considered and flow physics questions addressed (in blue). Operation condition parameters are given in red. Partial derivatives (c_{x*} and c_{y*}) are indicated here by subscripts.

In addition to CFD, the use of analytical simulation methods is beneficial because of several reasons. Their use hardly depends on the availability of codes. They are by orders of magnitude more efficient than the numerical integration of partial differential equations in time because of the independence of discretizations and the need to honor numerical stability criteria under significant model parameter variations. Such analytical simulation methods can be used for the verification of numerical solvers because they provide exact results (i.e., independent evidence is not needed to support them). However, first of all, the availability of such models offers the chance to address questions about basic mass transport mechanisms, e.g., regarding the questions Q1–Q3 presented above.

One question related to the use of analytical simulation methods is about their required complexity and simplicity of using them. For the case considered, analytical simulation methods are given by Fourier series representations, there is the question of whether the complexity of such Fourier series representations can be significantly reduced to enhance the clarity of conclusions. Another question related to analytical simulation methods is that their development requires the neglect of effects of spacers, which modify basic flow characteristics implied by geometry, inflow and boundary conditions. The specification of spacer effects on the basic flow configuration varies in the literature [30,51,52]. There are indications that spacers can produce higher fluxes of about 30–40% compared with cases without spacers [30,51,52]. This means analytical simulation methods can be expected to provide valuable guidelines for the characterization of basic flow features, even under conditions where spacers and other effects (flow unsteadiness) are involved. Thus, there are questions in addition to Q1–Q3 considered above:

- Q4. What is the most simple analytical model which still enables accurate calculations equivalent to complete Fourier series solutions?
- Q5. Given the required approximate representation of the flow field, which arguments support the use of analytical simulation methods under more complex flow conditions?

Analytical solutions obtained by Laplace transforms were presented for a variety of problems, for example, in regard to turbulent dispersion of air pollution [53] and diffusion of oxygen into the blood [54], but no corresponding solutions were presented in regard to membrane systems. The corresponding value of analytical simulation methods is demonstrated here by addressing the questions Q1–Q5 presented above. Fourier analysis is applied for the derivation of analytical results, which offers (via the analysis of eigenvalue regimes) essential new insight. The model development is presented in Section 2 in conjunction with the consideration of questions about the model evaluation and computational aspects. Model applications are reported in Section 3 by focusing on the influence of OCs on species concentration distributions. Section 4 deals with conclusions about the questions considered.

2. Model Development

2.1. Equation Considered

Incompressible flow and a 2D domain are considered, see the illustration in Figure 2. The velocity field is approximated by assuming constant velocities U and V in the x and y directions, which is a requirement to obtain analytical solutions. The suitability of this assumption is considered in Section 4, conclusion #5, in conjunction with mass transport properties. A mixture (e.g., an oil-in-water emulsion) of a continuous phase (e.g., water) and a dispersed phase (e.g., oil at low concentrations) is considered. The solutions presented below are not specific to oil-in-water emulsions, they can be applied to many other systems. According to refs. [7,10,55], the dispersed phase transport equations in the feed/permeate region and membrane region (see Figure 1), respectively, read

$$\frac{\partial c}{\partial t} + U\frac{\partial c}{\partial x} + V\frac{\partial c}{\partial y} = D_M \Big(\frac{\partial^2 c}{\partial x^2} + \frac{\partial^2 c}{\partial y^2}\Big), \qquad \frac{\partial \epsilon_p c}{\partial t} + U\frac{\partial c}{\partial x} + V\frac{\partial c}{\partial y} = D_c \Big(\frac{\partial^2 c}{\partial x^2} + \frac{\partial^2 c}{\partial y^2}\Big).$$
(1)

Here, *c* refers to the dispersed phase concentration, *t* is time, D_M refers to the molecular diffusion (of oil in water), D_c refers to the capillary diffusion, and ϵ_p is the membrane

porosity. The constant diffusivities D_M and D_c are considered in consistency with the consideration of constant velocities U and V. Diffusion in x direction (which is sometimes neglected) is included. No attempt is made to differentiate between the velocities U and V involved in Equation (1) because these equations are covered by one equation in the following, in particular, the advection–diffusion equation of the membrane region,

$$\frac{\partial c}{\partial t} + \hat{U}\frac{\partial c}{\partial x} + \hat{V}\frac{\partial c}{\partial y} = \hat{D}_c \left(\frac{\partial^2 c}{\partial x^2} + \frac{\partial^2 c}{\partial y^2}\right).$$
(2)

Here, $\hat{U} = U/\epsilon_p$, $\hat{V} = V/\epsilon_p$, and $\hat{D}_c = D_c/\epsilon_p$ are introduced. The solution obtained can be easily applied as a solution to the feed/permeate region equation by a corresponding adjustment of model parameters (by setting $\epsilon_p = 1$ and replacing D_c by D_M). The domain considered is $0 \le x \le x_2$ and $y_1 \le y \le y_2$; see the illustration in Figure 2. By involving imposed functions g(x, y) and f(y), the initial distribution and x boundary conditions (BCs) at $x = x_1 = 0$ and $x = x_2$ are given by

$$c(x,y,0) = g(x,y), \quad c(0,y,t) = f(y), \quad \frac{\partial c}{\partial x}(x_2,y,t) = 0.$$
 (3)

The latter condition applies to the case that diffusion in *x* direction is included. The *y* BCs at $y = y_1$ and $y = y_2$ are given by

$$\frac{\partial c}{\partial y}(x, y_1, t) = 0, \qquad c(x, y_2, t)\hat{V}/\delta + \hat{D}_c \frac{\partial c}{\partial y}(x, y_2, t) = 0.$$
(4)

The *y* BCs modify a constant mass transfer (zero gradient) along *y* by the condition at y_2 , which leads to an accumulation of the dispersed phase in the membrane region; see the discussion in the beginning of Section 3. The parameter δ introduced here enables variations of the mass transfer at y_2 ; Equation (A2) in the Appendix A shows that $2/\delta$ represents the standardized membrane permeability. The introduction of δ is relevant to the identification of the critical Péclet number $Pe_c = 4/(2 + \delta)$ in Section 3.1. The measurement of δ is addressed in Section 3.2. Implications of these BCs may be well seen in figures below (although the effects may be small, see the distributions along x_*).

An important property of the solution c(x, y, t) is its boundedness: if the initial and boundary values of c(x, y, t) lie within a minimum and maximum, $c_{min} \le c \le c_{max}$, then all c(x, y, t) values are bounded by this range [56]. The consideration of an advection–diffusion equation for the dispersed phase is equivalent to the consideration of a stochastic particle model for this phase (a random model for the positions of oil droplets, a Brownian motion model originally applied to describe the motion of pollen grains in water) where the mass concentration represents a non-normalized probability density function (PDF) [57]: the concentration has all the properties of a PDF except that it is not normalized.

It is helpful to introduce normalized variables: $x_* = \hat{U}x/(2\hat{D}_c)$, $y_* = \hat{V}y/(2\hat{D}_c)$, and $t_* = \hat{V}^2 t/(4\hat{D}_c)$. By using these normalized variables, Equation (2) reads

$$\frac{\partial c}{\partial t_*} + 2\frac{U^2}{V^2} \frac{\partial c}{\partial x_*} + 2\frac{\partial c}{\partial y_*} = \frac{U^2}{V^2} \frac{\partial^2 c}{\partial x_*^2} + \frac{\partial^2 c}{\partial y_*^2}.$$
(5)

The equations derived in the following reveal that $c = c(x_*, y_* - y_{1*}, t_* | x_{2*}, \Delta, \delta, U/V)$, where $\Delta = y_{2*} - y_{1*}$. It is convenient to introduce usually applied non-dimensional numbers, the aspect ratio *A*, the Péclet number *Pe*, and the Péclet number *Pe*_y in *y* direction,

$$A = \frac{x_2}{y_2 - y_1}, \quad Pe = \frac{\hat{U}x_2}{\hat{D}_c}, \quad Pe_y = \frac{\hat{V}(y_2 - y_1)}{\hat{D}_c}.$$
 (6)

Pe and *Pe*_y represent the product of corresponding Reynolds numbers $(\hat{U}x_2/\nu \text{ and } \hat{V}(y_2 - y_1)/\nu$, respectively) with the Schmidt number ν/\hat{D}_c , where ν is the kinematic viscosity. By applying the relationships

$$x_{2*} = Pe/2, \qquad \Delta = Pe_y/2, \qquad U/V = Pe/(APe_y), \tag{7}$$

the solution $c = c(x_*, y_* - y_{1*}, t_* | x_{2*}, \Delta, \delta, U/V)$ can be written in the following way:

$$c = c(x_*, y_* - y_{1*}, t_* | A, Pe, Pe_y, \delta).$$
(8)

The solution $c(x_*, y_*, t_*)$ to Equation (5) involves two ingredients, the stationary solution $s(x_*, y_*)$ for infinitely large t_* and the transitional solution $w(x_*, y_*, t_*)$,

$$C(x_*, y_*, t_*) = s(x_*, y_*) + w(x_*, y_*, t_*).$$
(9)

The stationary and transitional solutions $s(x_*, y_*)$ and $w(x_*, y_*, t_*)$, which are referred to as stationary Fourier series model (FSM) and transitional Fourier series model (FSM), are given by Equation (A3) and Equation (A17), respectively; see the explanations in the Appendix A.

2.2. Validation of the Model Implementation

The equations were implemented in an in-house code written for this application. In numerical model applications, the use of the stationary and transitional FSMs obtained ensures exact solutions to the advection-diffusion equation considered. Relatively simple analysis also shows that the use of the structures of $X_n(x_*)$ and $U_n(y_*)$ given in the Appendix A ensures the correct upper and lower y_* BCs and outflow BCs. However, the inflow BCs represented by the sum of Fourier series contributions in the model implementation require validation because they are affected by the numerical implementation, for example, the correct calculation of eigenvalues. This question is addressed in terms of Figure 3, which shows the concentration distribution along y_*/Δ for reference parameters $Pe_y = 1$, $Pe = 10^2$, $A = 10^3$, $\delta = 10$, and $\delta = 1$ at $x_* = 0$. The dashed lines represent the imposed profile, the colored lines arise from imitated implementation errors: the neglect of one eigenfunction contribution, and the incorrect calculation of one eigenvalue (β_0^m or $\beta_{p_{1}}^{p}$, respectively) given by a modification by a factor of 2 or 0.5. It may be seen that such incorrect implementations have serious consequences. Both the neglect of one eigenfunction contribution or an eigenvalue modification (by a factor of 2 or 0.5) implies that the model is incapable to correctly provide the imposed profile. Therefore, checks whether concentration calculations are in consistency with imposed profiles are sufficient to conclude that the model implementation is correct. A relatively simple way to provide additional support for the correct model implementation is to ask whether the solutions obtained satisfy the boundedness of mass transport requirement. This is, e.g., not the case for the cases with implementation errors presented in Figure 3: such model results disagree with the boundedness of mass transport because concentration values below zero and above one are obtained. For all the cases presented below, it was found that the boundedness requirements were satisfied by model solutions.

2.3. Computational Cost

The computational cost of the presented method depends very much on the convergence behavior of Fourier representations involved. The stationary solution is driven by the convergence of y_* eigenfunctions. A scaling analysis reveals that $U_n(y_{2*}) \sim (q_1 - q_2)Pe_y/n^2$ at sufficiently high n, where q_1 and q_2 are positive and negative numbers of order unity (which are related to the two contributions to a_n in Table A1). Correspondingly, transitional solution contributions scale with $w_2(y_{2*}, 0) \sim q_1Pe_y/n^2$, where q_0 is a positive number of order unity. The easiest way to see these scalings is to multiply series contributions with the inverse scaling to demonstrate that series contributions do not change with n and parameter variations. Let us suppose the convergence requirement that the related Fourier series contributions need to be smaller than 10^{-9} . This implies the condition $Pe_y/N^2 \leq 10^{-9}$, i.e., $N \geq 10^{4.5}Pe_y^{1/2} \approx 3.2 \times 10^4 Pe_y^{1/2}$. For usual values $Pe_y \leq 1$, this condition can be easily satisfied (in less than a minute). A corresponding scaling analysis of x_* eigenfunctions involved in the transitional solution part shows that $w_1(x_{2*}, 0) \sim (-1)^{n+1}qPe(1+2/Pe)/n^3$, where q is a positive number of order unity. By using again a convergence criterion that corresponding Fourier series contributions need to be smaller than 10^{-9} , the corresponding convergence condition reads $Pe/N^3 \leq 10^{-9}$ for sufficiently high Pe, this means $N \geq 10^3 Pe^{1/3}$. For $Pe = (10^2, 10^3, 10^4)$, for example, one finds $N \geq (4.6 \times 10^3, 10^4, 2.2 \times 10^4)$. Such simulations take much less than a minute on a personal computer. A very essential observation is the following. In contrast to the stationary solution $s(x_*, y_*)$, the transitional solution $w(x_*, y_*, t_*) = c_{in}w_1(x_*, t_*)w_2(y_*, t_*)$ is given by separated variations in x_* and y_* . This makes computations by orders of magnitude more efficient because a double loop over x_* and y_* eigenvalues does not appear.

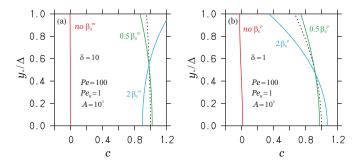


Figure 3. The concentration distribution along y_* from y_{1*} (lower bound) to y_{2*} (upper bound) for reference parameters $Pe_y = 1$, $Pe = 10^2$, $A = 10^3$, and $\delta = 10$ in (**a**) and $\delta = 1$ in (**b**) at $x_* = 0$ (dashed lines). The red lines refer to the neglect of β_0^m or β_0^p , respectively, eigenfunction contributions. The blue and green lines refer to a modification of β_0^m or $-\beta_0^p$, respectively, by the given factors (2 and 0.5).

The computational development of the simulation method leads to four observations. (i) The computational method is computationally highly efficient: exact solutions are obtained on a personal computer in less than a minute. (ii) A specific question related to the computational cost is about the scaling of cost with model parameters such as the Péclet numbers involved (Pe and Pe_y). It was shown that such model parameter variations have very minor effects on the solution convergence with respect to both the stationary and transitional solution. (iii) One key element of correct solutions is the exact iterative solution of eigenvalue equations for all parameter regimes, which represents a challenging task. An efficient algorithm for handling this question was presented here. (iv) Another key element is, in particular, the correct identification of the lowest order eigenvalues for all parameter regimes, otherwise the method simply does not work.

3. Model Application

Results using the model obtained are presented now by focusing for simplicity on the stationary solution (the transitional solution is discussed in Section 3.2)

$$c = c(x_*, y_* - y_{1*} | A, Pe, Pe_y, \delta).$$
(10)

The parameters involved are given by Equation (6). As shown in the Appendix A in the paragraph below Equation (A6), the critical parameter that separates different eigenvalue regimes is given by $p = 2/\delta - (1 + 2/\delta)Pe_y/2$ (see also the illustration in Figure 4). To restrict attention, it is assumed that $\delta \ge 0$. According to the BC, $s(x_*, y_{2*}) + [\delta/2][\partial s/\partial y_*](x_*, y_{2*}) = 0$, which implies a negative concentration gradient at y_{2*} ; this is equivalent to the calculation of a concentration (of oil) that has at the lower boundary y_{1*} a larger value than at the upper boundary y_{2*} . It is worth noting that the

assumption $\delta < 0$, which implies decreasing concentration values toward the lower boundary, may well be in conflict with the boundedness of mass transport for the conditions considered because concentration values larger than unity can be obtained.

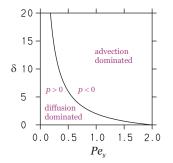


Figure 4. Parameter regimes: the black curve shows Pe_y values for which $p = 2/\delta - (1 + 2/\delta)Pe_y/2 = 0$.

3.1. FSM: Flow Regimes and Equivalent OC

The concentration distribution at $x_* = x_{2*}$ and $y_* = y_{1*}$ is shown in Figure 5 for varying parameters A, Pe, Pe_y , and δ . The most relevant observation is that $p = 2/\delta - (1+2/\delta)Pe_y/2$, which separates different eigenvalue regimes and also separates different concentration regimes; see Figure 5a. For p < 0, one finds high concentration values which do not change much with δ . We refer to this case as blocked flow below. For p > 0, one finds much lower concentration values which are, approximately, linearly controlled by δ . We refer to this case below as unblocked flow. The critical Pe_y for which p = 0 reads $Pe_c = 4/(2+\delta)$, which is bounded, $0 \le Pe_c \le 2$. Hence, p can be written $p = 2(1 - Pe_y/Pe_c)/\delta$. Here, Pe_y/Pe_c represents a normalized Péclet number. By introducing $L_f = Pe_c\hat{D}_c/\hat{V}$, it can be written $Pe_y/Pe_c = (y_2 - y_1)/L_f$. Correspondingly, the conditions for the unblocked (blocked) flow regimes are given by

$$y_2 - y_1 < L_f \quad (y_2 - y_1 > L_f),$$
 (11)

respectively. Here, L_f represents a characteristic length scale which depends only on flow properties. Equation (11) state, therefore, the requirement that the domain height has to be sufficiently small to enable unblocked flow. The relationship between unblocked/blocked flow and diffusion/advection-dominated flow regimes can be seen as follows. In unblocked flow, the diffusion dominates: molecular transport governs this regime basically unaffected by the directional influence of advection. It is, therefore, very natural to identify the unblocked (blocked) flow regimes as diffusion (advection)-dominated flow. According to $p = 2(1 - Pe_y/Pe_c)/\delta$, a Péclet number Pe_y/Pe_c smaller (larger) than unity is the condition to have diffusion (advection)-dominated flow.

Figure 5b–d shows the same huge discrepancies between blocked and unblocked flow regimes seen in Figure 5a. Figure 5c,d indicates an equivalence between Pe and inverse A effects. The structure of A curves can be seen as follows. An increased $A^{-1} = (y_2 - y_1)/x_2$ is equivalent to a decreased x_2 , which implies a higher \hat{U} to keep $Pe = \hat{U}x_2/\hat{D}_c$ unchanged. The higher the \hat{U} , the higher is the concentration reduction. The same mechanism implies Pe effects: an increased Pe is equivalent to a higher \hat{U} . With respect to Figure 5b, it is worth noting the following. According to Equation (A4), the x_*/x_{2*} dependence of the concentration distribution is basically controlled by U/V: x_*/x_{2*} variations will disappear if U/V vanishes. This case is almost irrelevant to applications, because usually U >> V. For $Pe_y > 1$ in Figure 5b, one finds $Pe/(APe_y) = U/V < 0.1$, i.e., this range of variations is unlikely to be seen in applications. For $Pe_y \leq 1$, the $Pe_y = \hat{V}(y_2 - y_1)/\hat{D}_c$ effects result from the following. An increased Pe_y implies a larger $y_2 - y_1$ and x_2 to keep A unchanged. This implies a smaller \hat{U} to keep Pe constant. Hence, an increased Pe_y increases the concentration.

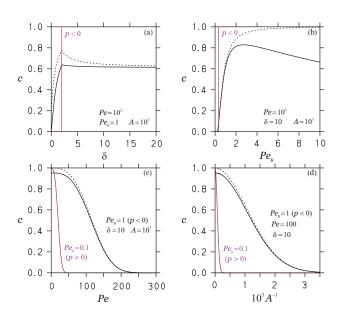


Figure 5. The concentration distribution at $x_* = x_{2*}$, $y_* = y_{1*}$ (dashed lines) and $y_* = y_{2*}$ (solid lines) for reference parameters $\delta = 10$, $Pe_y = 1$, $Pe = 10^2$, and $A = 10^3$. Variations of δ , Pe_y , Pe, A are shown in (**a**), (**b**), (**c**), and (**d**), respectively. The reference parameters kept constant are also given. In (**a**,**b**), the vertical line separates p > 0 and p < 0 cases. In (**c**,**d**), both p > 0 and p < 0 cases are shown.

Spatial concentration distributions are shown in Figures 6 and 7. The figures confirm the close relation between Pe, A^{-1} and Pe_y effects. As discussed above, an increase in Pe and A^{-1} reflects an increase in \hat{U} , i.e., a change in Pe can be accomplished by a corresponding change in A^{-1} . An analysis of such changes in the blocked flow regime (by looking at requirements for equivalent variations) reveals that increasing Pe by a factor k is approximately equivalent to dividing A by $k^{1.5}$. The relation between Pe_y and A variations is slightly different; see the discussion related to Figure 5. Different patterns can be seen for U/V < 0.1. However, according to the discussion related to Figure 5, this is the regime of little relevance to applications. Apart from this regime, an analysis shows that multiplying A by a factor k is approximately equivalent to multiplying Pe_y with a factor slightly larger that k (1.2k or 1.3k), i.e., there is an equivalence of Pe_y and A variations.

The huge discrepancy between the high-concentration blocked flow and much lowerconcentration unblocked flow regimes can be seen again in Figures 6 and 7. In addition to this difference, there is another factor coming into play: $U/V = Pe/(APe_y)$, which controls the x_*/x_{2*} variation according to $R = [1 + (1 + \beta_n^2)U^2/V^2]^{1/2}$. In the small U/V regime, which is of little relevance to applications (see above), one finds the following features. For U/V = 0.05 - 0.1, there are almost linear variations; see Figure 7b,c. For U/V < 0.05, one finds relatively minor concentration variations along x_*/x_{2*} , including almost constant concentration values if U/V << 0.05; see Figure 7b. This is the region where U/V has almost no influence on the concentration distribution. Otherwise, for U/V > 0.1, one observes a significant (exponentially) spatial concentration reduction, even for the blocked flow regime; see Figure 7c. Regarding the y_*/Δ variations shown in Figure 6, there are almost homogeneous concentration distributions. The latter is a consequence of two facts: the inflow distribution at $y_* = y_{1*}$, and the boundedness of scalars which excludes the appearance of minimum and maximum values in between boundary values.

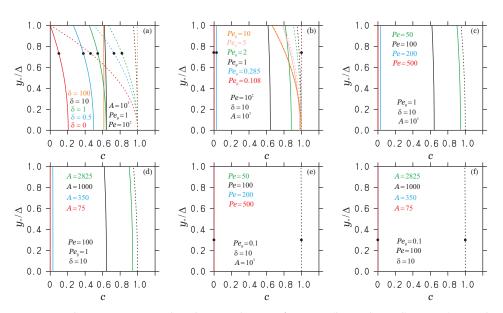


Figure 6. The concentration distribution along y_* from y_{1*} (lower bound) to y_{2*} (upper bound) at $x_* = 0$ (dashed lines) and $x_* = x_{2*}$ (solid lines). Reference values $\delta = 10$, $Pe_y = 1$, $Pe = 10^2$, and $A = 10^3$ are considered in (**a**–**d**) in conjunction with variations of δ , Pe_y , Pe, A. In (**e**,**f**), complementary cases to (**c**,**d**) are considered to cover the p > 0 case, where $Pe_y = 0.1$ in contrast to $Pe_y = 1$ in (**c**,**d**). The reference parameters kept constant are also given. The black dots on curves indicate p > 0 cases. The $x_* = 0$ profiles are unaffected by Pe, A variations. In (**e**,**f**), all solid curves coincide.

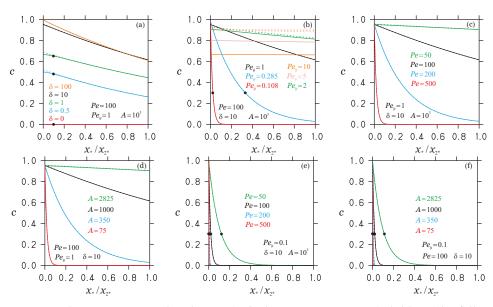


Figure 7. The concentration distribution (**a**–**f**) along x_* at $y_* = y_{2*}$ (solid lines) by following the notation applied in Figure 6. The dashed lines show the results of model Equation (12).

3.2. Zeroth-Order Model

The relevance of the lowest eigenvalue contributions is certainly of interest. To address this question, the stationary FSM is reduced to only these contributions by considering $s_0(x_*, y_*) = e^{y_*}a_0X_0(x_*)U_0(y_*)$. By involving the BC Equation (A13), a more appropriate writing reads $s_0(x_*, y_*) = f(y_*)X_0(x_*)$, i.e.

$$s_0(x_*, y_*) = c_1 \left(1 - \frac{(y_* - y_{1*})^2}{\Delta(\Delta + \delta)}\right) e^{(1 - R_0)x_*} \frac{1 + R_0 - (1 - R_0)e^{-2R_0(x_{2*} - x_*)}}{1 + R_0 - (1 - R_0)e^{-2R_0x_{2*}}},$$
(12)

where $R_0 = [1 + (1 + \beta_0^2)U^2/V^2]^{1/2}$, $x_{2*} = Pe/2$, $\Delta = Pe_y/2$, and $U/V = Pe/(APe_y)$. Depending on negative or positive *p* values, one needs to use either β_0^m or β_0^p , respectively, in R_0 . Here, β_0^p and β_0^m have to be found iteratively by solving Equation (A8) and Equation (A10), respectively; this means they are found via

$$\beta_0^p \Delta = \arctan\left\{\frac{\beta_0^p}{1 + \delta(1 + [\beta_0^p]^2)/2}\right\}, \quad \beta_0^m \Delta = \arctan\left\{\frac{\beta_0^m}{1 + \delta(1 - [\beta_0^m]^2)/2}\right\}.$$
(13)

Figure 8 demonstrates that simple approximations for these eigenvalues (simple functions of Pe_y or δ) are unavailable, meaning that they need to be determined iteratively via Equation (13). A simple alternative is to use an online calculator [58] to accurately solve the equations considered. According to Figure 8, zero eigenvalues may be found if $Pe_y = Pe_c = 4/(2 + \delta)$. The model Equation (12) is referred to below as zeroth-order model (ZOM). For given model parameters $\Delta(\Delta + \delta) = Pe_y(Pe_y + 2\delta)/4$, R_0 , and Pe, this model completely specifies the structure of the concentration distribution via the imposed BC $f(y_*)$ and $X_0(x_*)$, which represents exponential decay along x_* .

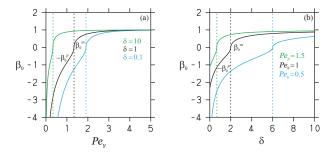


Figure 8. Eigenvalues β_0^m and $-\beta_0^p$ as function of (**a**) Pe_y (depending on δ) and (**b**) δ (depending on Pe_y). The vertical lines separate β_0^m from $-\beta_0^p$.

The performance of Equation (12) is also shown in Figure 7. It may be seen that the performance of this model is excellent with one minor exception: the U/V < 0.05 cases in Figure 7b. However, as discussed in relation to Figure 5, this is a case that is almost irrelevant to applications. On the one hand, the performance of Equation (12) reveals the fundamental relevance of including the eigenvalues of lowest order. On the other hand, this performance enables it to perform accurate simulations based on the simple analytical model Equation (12).

Figure 9, where the same cases are considered as in Figure 7, confirms the assumption that the same applies to the transitional solution (a constant initial value $c(x_*, y_*, 0) = c_{in} =$ 1 is considered): the reduction of Equations (A17)–(A19) to only considering the lowest eigenvalue contributions results in transitional solutions that do not show any visible difference from the complete solutions (A17)–(A19). As given in regard to the stationary solution, this fact enables the use of a simple analytical formula for performing highly accurate simulations.

First, the ZOM is very beneficial regarding the characterization of the concentration distribution by global maximum and minimum values, which exist because of the boundedness of mass transport. The spatial concentration distributions shown in Figures 6 and 7 reveal a global concentration maximum and minimum at $(x_*, y_*) = (0, y_{1*})$ and $(x_*, y_*) = (x_{2*}, y_{2*})$, respectively. The ZOM then provides the maximum and minimum values

$$s_{min} = s_0(x_{2*}, y_{2*}) = c_1 \Big[1 - \Delta/(\Delta + \delta) \Big] X_0(x_{2*}), \qquad s_{max} = s_0(0, y_{1*}) = c_1.$$
(14)

Thus, by involving $Pe_y = 2\Delta$, the stationary ZOM solution is found to be bounded by

$$c_1 \Big[1 - \frac{Pe_y}{Pe_y + 2\delta} \Big] X_0(x_{2*}) \le s_0(x_*, y_*) \le c_1.$$
(15)

Second, the ZOM is very beneficial regarding the estimation of model parameters, which are required to apply the ZOM and FSM and are relevant to scaling questions (upscaling). The aspect ratio $A = x_2/(y_2 - y_1)$ is known, but a relevant question is about how δ , $Pe = \hat{U}x_2/\hat{D_c}$, and $Pe_y = \hat{V}(y_2 - y_1)/\hat{D_c}$ can be determined. This question is not trivial because $Pe = \hat{U}x_2/\hat{D_c}$ and $Pe_y = \hat{V}(y_2 - y_1)/\hat{D_c}$ involve the membrane diffusivity $\hat{D_c}$, which is difficult to measure, and δ , too, is difficult to determine on the basis of $s(x_*, y_{2*}) + [\delta/2][\partial s/\partial y_*](x_*, y_{2*}) = 0$. This question is considered by assuming the measured values of the stationary solution provided by the ZOM. It is possible to address this question in a more general set up, but for simplicity, it is assumed here that U/V is known in addition to A. Because of $X_0(0) = 1$, it is found that $s_0(0, y_*) = f(y_*) = c_1(1 - (y_* - y_{1*})^2/[\Delta(\Delta + \delta)])$. Hence, concentration values taken at two appropriate positions enable us to determine $\Delta = Pe_y/2$ and δ . Because U/V is known, δ , Pe_y , and $Pe_y = APe_yU/V$ are found.

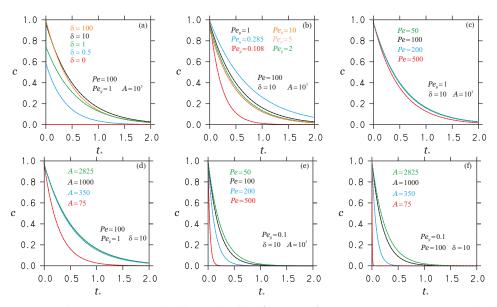


Figure 9. The concentration distribution (**a**–**f**) as function of time t_* at $x_* = x_{2*}$, $y_* = y_{2*}$ (solid lines) by considering the same cases as in Figure 7 (solid lines). The dashed lines show the results of using only the lowest eigenvalue contributions in Equations (A17)–(A19): there is no visible difference to solid curves.

Third, the ZOM can be used to obtain exact conclusions about the equivalence conditions for model parameter variations discussed in Section 3.1. As a first example, let us consider $s_0(x_*, y_{1*}) = c_1 X_0(x_*)$ by excluding variations of Pe and δ . In this case, $X_0(x_*)$ is affected by variations of Pe_y and A only via $(1 + \beta_0^2)/[Pe/(APe_y)]^2$ in R_0 . Then, variations of A can be compensated by variations of Pe_y which ensure an unchanged $X_0(x_*)$. As a second example, let us consider $s_0(x_*, y_{1*}) = c_1 X_0(x_*)$ by excluding variations of Pe_y and δ . Then, $X_0(x_*)$ is unchanged in response to A variations if Pe is calculated by the condition to have an unchanged $X_0(x_*)$. As a third example, let us consider the parameter $\Delta/(\Delta + \delta)$, which determines the minimal concentration along y_* according to Equation (15). This parameter can be used to balance membrane fouling reflected by a change in δ from δ_1 to δ_2 . The latter requires to determine a change Δ_2 of Δ_1 such that $\Delta_1/(\Delta_1 + \delta_1) = \Delta_2/(\Delta_2 + \delta_2)$. The latter requirement can also be written $\delta_2/\delta_1 = Pe_{y,2}/Pe_{y,1}$. This means an increase in δ_2 (a reduced flux) can be compensated by a corresponding higher Pe_y .

4. Summary

An analytical FSM for the mass transport in membrane systems is presented here. Our method is computationally highly efficient and exact, i.e., no independent evidence is needed for model results, only the implementation needs justification. Evidence for the validity of results and computational features is described in Sections 2.2 and 2.3. Our conclusions were obtained by Fourier analysis, which provides analytical results that cannot be obtained by experimental or numerical studies.

Let us summarize the answers obtained in regard to the questions Q1–Q5 presented in Section 1. For doing so, the OCs (A, Pe, Pe_y , δ) are assumed to be known; questions in this regard can be addressed by following the discussion at the end of Section 3.2. Based on our core results, the identification of flow separation conditions $y_2 - y_1 \leq L_f$ (see Equation (11)), the discussions of equivalent OC variations (see Section 3), and the ZOM $s_0(x, y) = f(y)X_0(x)$ (see Equation (12)), there are the following conclusions:

- 1. Arguably, our most relevant observation is that $p = 2(1 - Pe_y/Pe_c)/\delta$, which separates different eigenvalue regimes and also separates different mass transport regimes, in particular diffusion ($p > 0, y_2 - y_1 < L_f$) and advection ($p < 0, y_2 - y_1 > L_f$)dominated regimes. These regime separation conditions compare geometric conditions (the domain size) with the characteristic length scale L_f imposed by the flow. Given a membrane size considered, knowledge of the regime separation conditions is beneficial for the understanding of upscaling requirements, i.e., the use of lab results for pilot- and full-scale applications; with respect to the same flow properties, upscaling can imply transitions from very efficient to very inefficient flow regimes. A very relevant observation is that diffusion-dominated and advection-dominated flow regimes correspond to unblocked (low concentration values) and blocked (high concentration values) flow. Hence, the mathematical characterization of the dominance of one process has relevant physical consequences. Advection-dominated flow implies blocked flow because the dominance of advection inhibits molecular diffusion, i.e., the reduction in concentration gradients.
- 2. Knowledge of analytical equivalence conditions for *A*, *Pe* and *Pe_y* parameter variations for cases of practical relevance enables the use of various parameter variations to realize desired effects (under conditions where certain parameter variations are inappropriate). The understanding of several ways to accomplish regime changes enables transitions to preferred flow regimes (see the discussion related to Figure 7). The ZOM can provide exact conclusions about equivalent variations of OCs.
- 3. The FSM, but in particular the ZOM, provide an answer to question Q3 about the understanding of the overall mass transport and characteristic mass distribution features: for both flow regimes, the ZOM explains the difference between (input and output) boundary values implied by OCs and characteristic concentration variations in between these bounds. In particular, the ZOM enables the explicit calculation of global maximum/minimum concentration values $c_1(1 Pe_y/[Pe_y + 2\delta])X_0(x_{2*}) \le s_0(x_*, y_*) \le c_1$, which is helpful for the understanding of concentration variations.
- 4. Based on the FSM, the ZOM $s_0(x, y) = f(y)X_0(x)$ was presented, which be can be easily applied. The ZOM performance was found to be excellent for all regimes of practical relevance; see above. The significant advantages offered by the ZOM are described above (see second and third points).
- 5. According to Equation (2), the mass transport is affected by mass transport properties (diffusivity \hat{D}_c), mass transport initial and BCs, and the structure of the velocity field. The transport properties are known, and there is no problem to exactly satisfy mass transport initial and BCs. Although the velocity field is only approximately represented, the boundedness property of mass transport ensures then proper transitions between the imposed exact BCs, i.e., more complex flow conditions can be covered by the method considered.

Our conclusions support the membrane system design via providing relevant guidelines for applications; see the preceding paragraph, in particular the conclusions regarding questions Q1–Q3. It can be expected that the results obtained provide valuable guidelines for CFD and experimental studies, even under conditions where spacers and other effects are present (see the conclusion to question Q5). Direct applications of our method to membrane system design are not the scope of our paper, the latter requires extensive comparisons with corresponding CFD results for a wide range of parameter variations.

The results presented may be seen as modeling of the solute concentration of oil-in an oil-in-water emulsion (see the reference to a corresponding stochastic particle system in Section 2.1). However, in fact, the same equations can be used to model a variety of other systems, as, for example, saltwater systems for brackish water and seawater reverse osmosis. It is worth noting that the method presented actually represents a design strategy that can be applied to other (radial) geometries; it can be extended to three-dimensional analyses, and it can be applied with other initial and boundary conditions. Results obtained on this basis can be expected to provide valuable additional information and conclusions in support of numerical and experimental studies of membrane systems.

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Nomenclature

Α	aspect ratio, $x_2/(y_2 - y_1)$
С	dispersed phase concentration
<i>c</i> _{in}	initial value in $c(x_*, y_*, 0) = c_{in}$
c_1	model parameter, see Equation (A13)
D_M	molecular diffusion coefficient
D_c	capillary diffusion coefficient
$\hat{D_c}$	D_c/ϵ_p
f(y)	imposed boundary condition
g(x,y)	imposed initial condition
L_f	characteristic length, $Pe_c\hat{D}_c/\hat{V}$
Pe	Péclet number, $\hat{U}x_2/\hat{D_c}$
Pe_y	Péclet number, $\hat{V}(y_2 - y_1) / \hat{D_c}$
Pe_c	critical Péclet number, $4/(2+\delta)$
р	parameter, $2/\delta - (1+2/\delta)Pe_y/2$
R	parameter, $[1 + (1 + \beta_n^2)U^2/V^2]^{1/2}$
s,w	stationary, transitional solutions
t	time
t_*	non-dim., $\hat{V}^2 t/(4\hat{D_c})$

U _n	shifted <i>y</i> eigenfunction
U, V	velocities in x, y directions
U/V	non-dim., $Pe/(APe_y)$
\hat{U},\hat{V}	$U/\epsilon_p, V/\epsilon_p$
X_n	x eigenfunction
<i>x</i> , <i>y</i>	positions in space
x_1, x_2	x domain bounds
<i>y</i> ₁ , <i>y</i> ₂	y domain bounds
x_{*}, y_{*}	non-dim., $\hat{U}x/(2\hat{D}_c)$, $\hat{V}y/(2\hat{D}_c)$
<i>x</i> _{2*}	non-dim., $\hat{U}x_2/(2\hat{D}_c) = Pe/2$
y_{1*}, y_{2*}	non-dim., $\hat{V}y_1/(2\hat{D_c})$, $\hat{V}y_2/(2\hat{D_c})$
eta_0,eta_0^m,eta_0^p	eigenvalues, see Equations (12) and (13)
Δ	non-dim., $y_{2*} - y_{1*} = Pe_y/2$
δ^{-1}	membrane permeability in Equation (4)
ϵ_p	membrane porosity
ν	kinematic viscosity
()min,max	minimum, maximum values
()0	zeroth order contributions

Appendix A. Stationary and Transitional Solutions

Appendix A.1. Stationary Solution

The stationary solution satisfies the stationary partial differential equation

$$2\frac{U^2}{V^2}\frac{\partial s}{\partial x_*} + 2\frac{\partial s}{\partial y_*} = \frac{U^2}{V^2}\frac{\partial^2 s}{\partial x_*^2} + \frac{\partial^2 s}{\partial y_*^2},\tag{A1}$$

combined with non-homogeneous BCs,

$$s(0, y_*) = f(y_*), \quad \frac{\partial s}{\partial x_*}(x_{2*}, y_*) = 0, \quad \frac{\partial s}{\partial y_*}(x_*, y_{1*}) = 0, \\ s(x_*, y_{2*}) + \frac{\delta}{2} \frac{\partial s}{\partial y_*}(x_*, y_{2*}) = 0.$$
(A2)

The solution $s(x_*, y_*)$ of Equation (A1) can be obtained by separation of variables [56]. It reads

$$s(x_*, y_*) = \sum_{n=0}^{N} s_n = e^{y_*} \sum_{n=0}^{N} a_n X_n(x_*) U_n(y_*),$$
(A3)

where $N \rightarrow \infty$ is supposed. The x_* eigenfunctions, which read the eigenvalues β_n of y_* eigenfunctions, are given by

$$X_n(x_*) = e^{(1-R)x_*} \frac{1+R-(1-R)e^{-2R(x_{2*}-x_*)}}{1+R-(1-R)e^{-2Rx_{2*}}},$$
(A4)

where $R = [1 + (1 + \beta_n^2)U^2/V^2]^{1/2}$. Equation (A3) applies the modification $Y_n(y_*) = e^{y_*}U_n(y_*)$ of y_* eigenfunctions, which simplifies the ordinary differential equation (ODE) for y_* eigenfunctions because of a vanishing first-order derivative. For n = 1, 2, ..., the modified eigenfunctions are found to be given by

$$U_n(y_*) = \beta_n \cos[\beta_n(y_* - y_{1*})] - \sin[\beta_n(y_* - y_{1*})].$$
(A5)

For the case n = 1, 2, ... considered, the y_* eigenvalues β_n satisfy the equation

$$\pi^{-1}\beta_n(y_{2*} - y_{1*}) - n = \pi^{-1} \arctan\left\{\frac{\beta_n}{1 + \delta(1 + \beta_n^2)/2}\right\}.$$
 (A6)

This equation can be solved iteratively starting with $\beta_n = n\pi/(y_{2*} - y_{1*})$ on the right-hand side (RHS). The solution is obtained after fewer than 20 iterations. The converged solution

can be written $\beta_n = \nu \pi / (y_{2*} - y_{1*})$, where ν refers to the shifted eigenvalue (in contrast to n). Depending on the BCs (the setting of δ), it turns out that ν is bounded, $n \le \nu \le n + 1/2$. Here, the lower and upper limits correspond to $\delta \to \infty$ and $\delta = 0$, respectively.

However, there are also contributions for n = 0. In particular, these contributions are determined by the sign of $p = 2/\delta - (1 + 2/\delta)\Delta$, where $\Delta = y_{2*} - y_{1*}$ is applied [56]. Here, p arises from the BCs of U [56]. One finds $0 = U'(y_{1*}) - p_1U(y_{1*})$ and $0 = U'(y_{2*}) + p_2U(y_{2*})$, where $p_1 = -1$ and $p_2 = 1 + 2/\delta$. The latter implies $p = p_1 + p_2 + p_1p_2\Delta$. Let us consider these two cases in the following two paragraphs (the consideration of p = 0 has little practical value because it implies a specific setting for Δ which can be avoided).

(a) Case p > 0: In this case, there is one additional positive eigenvalue (denoted by β_0^p , p refers to a positive eigenvalue) below $\beta_1 = \pi/(y_{2*} - y_{1*})$. The eigenfunction follows Equation (A5),

$$U_0^p(y_*) = \beta_0^p \cos\left[\beta_0^p(y_* - y_{1*})\right] - \sin\left[\beta_0^p(y_* - y_{1*})\right].$$
 (A7)

The eigenvalue β_0^p is determined by the solution of Equation (A6) with n = 0,

$$\beta_0^p(y_{2*} - y_{1*}) = \arctan\left\{\frac{\beta_0^p}{1 + \delta(1 + [\beta_0^p]^2)/2}\right\}.$$
(A8)

This equation was solved iteratively in the same way as Equation (A6) by applying, however, 1000 iterations starting with $\beta_0^m = 10^{-8}$.

(b) Case p < 0: In this case, there is one additional negative eigenvalue (denoted by β_0^m , *m* refers to a negative eigenvalue). The corresponding eigenfunction reads

$$U_0^m(y_*) = \beta_0^m cosh[\beta_0^m(y_* - y_{1*})] - sinh[\beta_0^m(y_* - y_{1*})] = = \frac{\gamma - 1}{2} e^{\beta_0^m(y_* - y_{1*})} + \frac{\gamma + 1}{2} e^{-\beta_0^m(y_* - y_{1*})},$$
(A9)

where β_0^m is given by the solution of

$$\beta_0^m(y_{2*} - y_{1*}) = \operatorname{arctanh}\left\{\frac{\beta_0^m}{1 + \delta(1 - [\beta_0^m]^2)/2}\right\}.$$
(A10)

An alternative formulation of this equation, which is based on the solution of the quadratic equation for β_0^m , reads

$$\beta_0^m = -\frac{1}{\delta \tanh(\beta_0^m \Delta)} + \sqrt{\frac{1}{\delta^2 \tanh^2(\beta_0^m \Delta)} + 1 + \frac{2}{\delta}}.$$
 (A11)

The latter equation was found to deal correctly with the limit $\beta_0^m \to 1$. It was numerically solved in the same way as Equation (A8) by applying 1000 iterations starting with $\beta_0^m = 10^{-6}$.

The coefficients a_n in Equation (A3) are chosen such that the stationary solution s matches an imposed boundary function $f(y_*)$ at $x_* = 0$; this means $s(0, y_*) = f(y_*)$. The key for deriving the coefficient formula is to make use of the orthogonality of $U_n(y_*)$ eigenfunction, $\int_{y_{1*}}^{y_{2*}} U_n(y_*) U_m(y_*) dy_* = \delta_{nm} w_n^2$, where the squared norm w_n^2 of U_n is given in Table A1.

Positive eigenvalues
$a_{n} = c_{1} \frac{\sin(\beta_{n}\Delta)}{w_{n}^{2}} e^{-y_{2*}} - c_{1} \frac{e^{-y_{2*}}[k_{1n}\cos(\beta_{n}\Delta) + k_{2n}\sin(\beta_{n}\Delta)] - 4\beta_{n}e^{-y_{1*}}}{w_{n}^{2}\Delta(\Delta + \delta)(1 + \beta_{n}^{2})^{2}},$
$k_{1n} = 2\beta_n \Big[2 + (1 + \beta_n^2) \Delta \Big], \qquad k_{2n} = 1 - 2\beta_n^2 + \Big[1 + (1 + \beta_n^2) \Delta \Big]^2,$
$w_n^2 = \frac{1+\beta_n^2}{2}\Delta - \frac{1-\beta_n^2}{4\beta_n}sin(2\beta_n\Delta) + \frac{1}{2}cos(2\beta_n\Delta) - \frac{1}{2}$
Negative eigenvalue
$a_0^m = c_1 \frac{\sinh(\beta_0^m \Delta)}{w_m^2} e^{-y_{2*}} - c_1 \frac{e^{-y_{2*}} \left[k_{1m} \cosh(\beta_0^m \Delta) + k_{2m} \sinh(\beta_0^m \Delta) \right] - 4\beta_0^m e^{-y_{1*}}}{w_m^2 \Delta (\Delta + \delta) (1 - [\beta_0^m]^2)^2},$
$k_{1m} = 2eta_0^m \Big[2 + (1 - [eta_0^m]^2) \Delta \Big], \qquad k_{2m} = 1 + 2[eta_0^m]^2 + \Big[1 + (1 - [eta_0^m]^2) \Delta \Big]^2,$
$w_m^2 = rac{1}{2} - rac{1 - [eta_0^m]^2}{2} \Delta + rac{(1 - eta_0^m)^2}{8eta_0^m} e^{2eta_0^m\Delta} - rac{(1 + eta_0^m)^2}{8eta_0^m} e^{-2eta_0^m\Delta}$

Table A1. Fourier coefficients a_n and a_0^m of the stationary solution, where $a_0^p = a_0$.

By using this property, one obtains

$$a_n = \frac{1}{w_n^2} \int_{y_{1*}}^{y_{2*}} e^{-y_*} f(y_*) U_n(y_*) dy_*.$$
(A12)

The requirement for the imposed function $f(y_*)$ is that it needs to satisfy the BCs. In order to do so, it is assumed that

$$f(y_*) = c_1 \left\{ 1 - \frac{(y_* - y_{1*})^2}{\Delta(\Delta + \delta)} \right\}.$$
 (A13)

The use of this expression in Equation (A12) provides a_n as given in Table A1. The abbreviations a_0^p and a_0^m are used to refer to the coefficients related to positive and negative eigenvalues, respectively. It is worth noting that w_n^2 and w_m^2 refer to the squared norm of U_n and U_0^m , respectively. In correspondence to the notation used before, w_0^2 refers to the squared norm of U_0^p . With respect to a_0^m , there arises a question about the limit $\beta_0^m \to 1$, which is relevant to some parameter regimes. In this limit case, one finds a_0^m to be given by

$$a_0^m = c_1 \frac{\sinh(\beta_0^m \Delta)}{w_m^2} e^{-y_{2*}} - c_1 \frac{1 - e^{-2\Delta}(1 + 2\Delta + 2\Delta^2)}{4w_m^2 \Delta(\Delta + \delta)} e^{-y_{1*}}.$$
 (A14)

The latter expression was applied for $|\beta_0^m - 1| \le 10^{-8}$. The model (A3) obtained in this way represents the stationary Fourier series model (FSM).

Appendix A.2. Transitional Solution

The transitional solution satisfies the non-stationary equation,

$$\frac{\partial w}{\partial t_*} + 2\frac{U^2}{V^2}\frac{\partial w}{\partial x_*} + 2\frac{\partial w}{\partial y_*} = \frac{U^2}{V^2}\frac{\partial^2 w}{\partial x_*^2} + \frac{\partial^2 w}{\partial y_*^2},\tag{A15}$$

combined with homogeneous BCs,

$$w(0, y_*, t_*) = \frac{\partial w}{\partial x_*}(x_{2*}, y_*, t_*) = \frac{\partial w}{\partial y_*}(x_*, y_{1*}, t_*) = 0, w(x_*, y_{2*}, t_*) + \frac{\delta}{2} \frac{\partial w}{\partial u_*}(x_*, y_{2*}, t_*) = 0,$$
(A16)

and the initial condition $w(x_*, y_*) = g(x_*, y_*)$. The solution $w(x_*, y_*, t_*)$ of Equation (A15) can be found via separation of variables. It is given by

$$w(x_*, y_*, t_*) = c_{in} w_1(x_*, t_*) w_2(y_*, t_*),$$
(A17)

where (by supposing $N \to \infty$ and $M \to \infty$)

$$w_1(x_*,t_*) = \sum_{m=1}^{M} c_m sin(\alpha_m x_*) exp \Big\{ x_* - (1+\alpha_m^2) [U/V]^2 t_* \Big\},$$
 (A18)

$$w_2(y_*,t_*) = \sum_{n=0}^N b_n \Big[\beta_n \cos[\beta_n(y_*-y_{1*})] - \sin[\beta_n(y_*-y_{1*})] \Big] exp\{y_* - (1+\beta_n^2)t_*\}.$$
 (A19)

With respect to $w_2(y_*, t_*)$, the terms of zeroth order are provided in dependence on p as described above regarding the stationary solution; see cases (a) and (b). This means, for p > 0 Equation (A19) is applied, whereas sin[...] and cos[...] are replaced by the corresponding sinh[...] and cosh[...] functions for p < 0.

In consistency with Table A1, the y_* Fourier coefficients (which were determined by the condition to integrate to one, meaning no initial y_* variation was considered) are given in Table A2.

Table A2. y_* Fourier coefficients b_n and b_0^m of the transitional solution ($b_0^p = b_0$).

Positive eigenvalues	
$b_n=\frac{\sin(\beta_n\Delta)}{w_n^2}e^{-y_{2*}},$	$w_n^2=rac{1+eta_n^2}{2}\Delta-rac{1-eta_n^2}{4eta_n}sin(2eta_n\Delta)+rac{1}{2}cos(2eta_n\Delta)-rac{1}{2}$
Negative eigenvalue	
$b_0^m = \frac{\sinh(\beta_0^m \Delta)}{w_m^2} e^{-y_{2*}}$	$, w_m^2 = -\frac{1 - [\beta_0^m]^2}{2}\Delta + \frac{1 + [\beta_0^m]^2}{4\beta_0^m} sinh(2\beta_0^m\Delta) - \frac{1}{2}cosh(2\beta_0^m\Delta) + \frac{1}{2}$

The corresponding x_* Fourier coefficients can be calculated on the basis of the orthogonality property of x_* eigenfunctions, $\int_0^{x_{2*}} \sin(\alpha_n x_*) \sin(\alpha_m x_*) dx_* = \delta_{nm} w_x^2$, where the squared norm of x_* eigenfunctions is given by

$$w_x^2 = \frac{x_{2*}}{2} - \frac{\sin(2\alpha_m x_{2*})}{4\alpha_m}.$$
 (A20)

The x_* eigenvalues α_m in Equation (A18) satisfy the equation

$$\pi^{-1}\alpha_m x_{2*} - n = -\pi^{-1} \arctan(\alpha_m).$$
(A21)

Here, n = 1, 2, ... refers to non-disturbed eigenvalue numbers. The technique to solve this equation for n = 1, 2, ... is equivalent to the solution of Equation (A6) related to the stationary solution, where 20 iterations were applied. There is one positive eigenvalue below $\alpha_1 = \pi/x_{2*}$ which is included in the solution of Equation (A22) for n = 1, 2, ...

With respect to the x_* Fourier coefficients, the condition reads

$$c_m = \frac{1}{w_x^2} \int_0^{x_{2*}} e^{-x_*} g(x_*) \sin(\alpha_m x_*) dx_*,$$
(A22)

where $g(x_*)$ is an imposed x_* profile. Regarding the latter,

$$g(x_*) = c_{in}e^{x_* - x_{2*}} \left[1 + (x_{2*} - x_*)(1 + 1/x_{2*}) \right] x_*/x_{2*}$$
(A23)

is applied. This function follows the e^{x_*} variation of $w_1(x_*, t_*)$; the factor $e^{x_{2*}}$ is included for normalization. The function $g(x_*)$ is positive, bounded, and correctly satisfies the x_* BCs; this means g(0) = 0 and $g'(x_{2*}) = 0$. By using Equation (A22), the x_* Fourier coefficients obtained read

$$c_m = \frac{2(1+1/x_{2*})e^{-x_{2*}}}{x_{2*}w_x^2\alpha_m^3} \left[1 - \cos(\alpha_m x_{2*})\right].$$
(A24)

The model Equation (A17) obtained represents the transitional Fourier series model (FSM).

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