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# Dilution of reactive plumes: evolution of concentration statistics under diffusion and nonlinear reaction

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#### Abstract

Concentration fields of solutes in porous media often exhibit large fluctuations, driven by physical and chemical heterogeneity from the pore to the Darcy scale. For many applications, ranging 8 from reactive transport modeling to toxicology, the knowledge of mean concentrations is not suf-9 ficient, and quantifying concentration variability is necessary. The probability density function 10 (PDF) of concentration quantifies the frequency of occurrence of concentration values throughout 11 a spatial domain. While evolution equations and analytical solutions for the concentration PDF 12 exist for conservative solutes, less is known about its evolution under the joint action of transport 13 and reaction. In this work, we investigate how dilution of a reactive plume by diffusion affects 14 the statistics of concentrations. While mixing has no effect on first-order reactions, its coupling 15 with nonlinear reactions leads to non-trivial effective kinetics relevant for a broad range of reac-16 tive transport problems. We study the evolution of the concentration PDF under diffusion and 17 nonlinear reaction in one spatial dimension, which represents a critical step towards further cou-18 pling with heterogeneous advection. We show that the dependence of the scalar dissipation rate 19 on concentration encodes the impact of diffusive transport on the concentration PDF and derive 20 a dynamical equation for its time evolution. Using a weak-coupling approximation for the reac-21 tion and diffusion dynamics, we derive analytical predictions for the concentration PDF and its 22 moments. Our results provide new insights into how diffusion and reaction control concentration 23 variability and open new opportunities for coupling mixing models with chemical reactions. 24

#### Highlights 25

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• We introduce a general framework quantifying the link between concentration PDFs and spatial 26 concentration profiles. 27

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• We derive a dynamical equation for the evolution of the concentration PDF under diffusion and nonlinear reaction in 1D.

• We derive analytical predictions for the concentration PDF and its moments using a weak-30 coupling approximation. 31

#### 1 Introduction 32

Quantifying the variability in solute concentrations in the subsurface is of primary importance to a 33 large range of applications, including reactive transport modeling (e.g., [1]), characterizing the natural 34 chemical signature of groundwater systems (e.g., [2]), and assessing human health risks associated with 35 contaminant transport (e.g., [3]). In particular, when reaction rates depend nonlinearly on available 36 concentrations, effective reaction rates can differ by orders of magnitude from reaction rates determined 37 under well-mixed conditions [4–7]. In such cases, knowledge of mean concentrations is not sufficient 38

<sup>39</sup> to predict overall reaction rates, which requires additional knowledge about the spatial variability of

40 concentration values and local reaction rates.

Concentration probability density function (PDF) methods aim to quantify transport and reaction 41 processes in terms of the spatial statistics (i.e., frequency of occurrence of given values) of transported 42 scalars [8-11]. These methods were originally developed in the context of turbulent flows, where they 43 have been extensively employed to model both conservative transport [12-17] and reactive transport 44 with an emphasis on combustion [18-23]. More recently, these methods have gained popularity in the 45 context of porous media to study conservative [2, 24-33] and reactive [34-40] mixing. The statistics of 46 concentrations have been studied in relation to different types of fluctuations. The first approach is to 47 study concentration fluctuations with respect to an average transport model, such as the advection-48 dispersion equation [24-26], in terms of a stochastic transport model such as the continuous time 49 random walk [28, 41]. Regarding, for instance, average concentrations at a given spatial position, 50 fluctuations typically arise from variability in particle transit times due to heterogeneity. A second 51 approach is to characterize uncertainty in concentrations due to uncertainty in model parameters, such 52 as flow velocities and reaction rates [29, 42-45]. These methods quantify the variability in concentra-53 tions due to uncertainty in physical and chemical parameters. A third point of view is to consider 54 the variability in concentrations over the spatial extension of a solute plume which arises from known 55 transport and reaction processes. Even with homogeneous physical and chemical parameters, mean 56 concentrations and associated fluctuations evolve under the combined effect of transport and reaction. 57 In this context, the concentration PDF provides a useful tool to quantify the spatial frequency of 58 occurrence of different solute concentration values, and to provide information about the mixing state 59 of solute plumes without explicitly resolving the corresponding spatial profiles [31]. 60 Here, we follow this last approach to study how the dilution of a reactive plume by diffusive mixing 61 affects concentration statistics (Fig. 1). In this context, evolution equations have been derived for the 62

conservative concentration PDF by quantifying the formation and interactions of lamellar structures 63 in concentration fields subject to advection and diffusion [30, 31, 46]. Recent developments have 64 extended these theories to mixing-limited bimolecular reactions [47]. Here, we consider a different 65 reactive transport problem in which mixing does not act to bring segregated reactants into contact 66 but rather to dilute a reactive plume. This system is relevant to a broad range of reactive transport 67 problems [6, 7], including situations where transported solute plumes react nonlinearly with a solid 68 phase as they get diluted by mixing. This leads to new dynamics that fundamentally differ from what 69 is known regarding mixing-limited reactions across diffusing or dispersing fronts [48-51]. 70

We investigate the coupling of dilution by diffusive mixing with the nonlinear degradation of a sin-71 gle species locally undergoing diffusion in one spatial dimensional and decaying at a rate proportional 72 to an arbitrary power of concentration (Fig. 1). We generalize the results of [7] for the evolution 73 of total mass and mean concentration to the evolution of the full concentration PDF, and we obtain 74 a dynamical equation for the latter in terms of the scalar dissipation rate [52-56], expressed as a 75 function of concentration. In addition to the mean concentration, the concentration PDF formulation 76 allows us to quantify higher-order concentration moments, such as the concentration variance, which 77 provides information about the variability in concentration values observed throughout a spatial do-78 main. Building on the weak-coupling approximation developed in [7], we relate the evolution of the 79 concentration PDF and its moments to the evolution of the maximum concentration, and quantify the 80 resulting anomalous kinetics. 81

In the interest of developing and illustrating the new method proposed here, the class of reaction-82 diffusion problems we analyze introduces significant simplifications. In particular, we consider a one-83 dimensional problem and neglect heterogeneous advection effects. In addition, the weak-coupling 84 approach involves a Gaussian approximation for the solute plumes at late times. Although reaction 85 under diffusion or dispersion in the absence of advective heterogeneity has been studied in its own 86 right in the context of porous media [4, 57-61], these conditions may be seen as fundamental building 87 blocks in the theory of mixing in heterogeneous flows. In the presence of heterogeneous flow fields 88 characteristic of porous media, solute plumes are deformed into a collection of elongated structures 89 termed lamellae (Fig. 1) [31]. Using the Ranz transform [62], the resulting compression-diffusion-90 reaction equation can be mapped onto a one-dimensional diffusion-reaction equation, which motivates 91



Figure 1: Illustration of the reactive transport problem under consideration (adapted from a simulation of transport at the Darcy scale in a heterogeneous permeability field [30]). We consider a plume of reactive solute (green) that undergoes nonlinear decay either by reaction with the solid, with another solute in excess everywhere, or with another solute that is well mixed inside the plume. The plume is diluted by mixing with the non-reactive resident fluid (black) while it reacts, leading to non-trivial effective reaction rates [7]. The latter are governed by the coupling of nonlinear reaction with the diffusive flux along concentration gradients. Here, we study the one-dimensional diffusion-reaction problem, which represents the first step towards solving the full advection-diffusion-reaction system.

<sup>92</sup> the present study.

The paper is structured as follows. Section 2 introduces a general definition for the concentration 93 PDF associated with a given spatial profile and explores its key properties. In Section 3, in light 94 of these concepts, we derive a governing equation for the dynamics of the concentration PDF of a 95 solute undergoing diffusion-induced dilution in one spatial dimension and nonlinear reaction described 96 locally by a power-law decay of concentration. In this context, we also discuss the relationship between 97 this simplified one-dimensional problem and reaction-diffusion in the presence of heterogeneous flows 98 typically found in porous media, based on the Ranz transform. We then study the one-dimensional 99 problem and the associated effective kinetics in terms of the weak-coupling approximation in Section 4. 100 Conclusions and an overall discussion of the results are presented in Section 5. Some additional 101 technical derivations are provided in the Appendices. 102

## <sup>103</sup> 2 The concentration PDF

This section is concerned with providing a general definition of the concentration PDF associated with 104 arbitrary spatial concentration profiles. We first present the general multispecies formulation, and then 105 discuss the single-species case. We provide a detailed discussion of the properties of the concentration 106 PDF, as well as of its numerical computation. The approach developed here, which forms the basis for 107 the subsequent sections, builds on that introduced in [63] regarding Eulerian velocity PDFs. The main 108 concepts behind the concentration PDF, which quantifies the spatial frequency of occurrence of different 109 concentration values, are illustrated in Fig. 2 for a one-dimensional, single-species concentration profile. 110 Before proceeding, we introduce some notation that will be employed throughout. The concentra-111 tion  $\underline{C}(\boldsymbol{x};t)$  at a given location  $\boldsymbol{x}$  and time t, with  $\underline{C} = (C_1, \ldots, C_{n_s})$ , is defined so that  $\underline{C}(\boldsymbol{x};t) d\boldsymbol{x}$  is 112 the mass in an infinitesimal volume  $d\mathbf{x}$  centered at  $\mathbf{x} = (x_1, \ldots, x_d)$  of each chemical species  $1, \ldots, n_s$ . 113 We use an underscore to denote quantities whose components relate to concentration species (of which 114 there are a number  $n_s$ ), and boldface to denote spatial vectors (which have components according to 115



Figure 2: Illustration of the key concepts underlying the concentration PDF. (a): Example spatial concentration profile, a Gaussian with unit variance and unit height, restricted to the spatial domain  $\Omega = [-3, 3]$  of size  $|\Omega| = 6$ . (b): Corresponding concentration PDF. Given a spatial profile, the concentration PDF quantifies the spatial frequency of occurrence of different concentration values c = C(x) occurring at some position(s) x. When concentration values are discretized into bins of width  $\Delta c$ , the PDF value p(c) is approximated by the fraction of the domain of the domain where the concentration c falls in this range, divided by  $\Delta c$  to obtain a probability density (bars in panel b). The continuous PDF (solid line in panel b) is obtained in the limit of infinitesimal bin widths dc. Each infinitesimal region dx around a point where C(x) = c contributes a probability density  $(dx/dc)/|\Omega| = 1/(|\Omega||dC(x)/dx|)$ . This causes a divergence of the PDF at the maximum concentration  $c_m = C(3) = C(-3)$ , where the gradient is small. These concepts are quantified and explored in detail throughout the text.

the spatial dimension d). We denote by |A| the d-volume of a spatial domain A (volume for d = 3-116 dimensional A, area for d = 2, length for d = 1, and number of points for d = 0). As usual, |a| for a 117 vector a denotes its Euclidean norm. We consider a spatial domain  $\Omega(t)$  where the concentrations are 118 defined for each time t, and denote by  $\mathbf{X}(t)$  a random variable with a uniform distribution over  $\Omega(t)$ . 119 The average with respect to X(t), i.e., the spatial average, is denoted by  $\overline{\cdot}$ . We also introduce, for 120 each time t, a random variable C(t) taking concentration values according to their spatial probability 121 of occurrence, and denote the average with respect to it (i.e., over concentrations) by  $\langle \cdot \rangle$ . As will be 122 discussed in more detail below, the two averages, while conceptually different, yield equivalent results. 123 That is, for an arbitrary function  $f(\underline{c}, t)$  of concentrations  $\underline{c}$  and times t, 124

$$\overline{f(\underline{C}[\boldsymbol{X}(t),t])} = \frac{1}{|\Omega(t)|} \int_{\Omega(t)} d\boldsymbol{x} f(\underline{C}[\boldsymbol{X}(t),t]) = \int_{0}^{\infty} d\underline{c} f(\underline{c},t) p(\underline{c};t) = \langle f[\underline{C}(t),t] \rangle, \tag{1}$$

where, at each time t,  $p(\cdot; t)$  is the concentration PDF, to be formally defined momentarily.

#### <sup>126</sup> 2.1 General formulation for a multispecies system

127 At each time t, the concentration PDF  $p(\cdot; t)$  is defined so that  $p(\underline{c}; t) d\underline{c}$  is the probability, in the sense

of spatial frequency of occurrence, of finding a concentration in the infinitesimal vicinity  $d\underline{c} = \prod_{j=1}^{n_s} dc_j$ of  $\underline{c}$  in the domain  $\Omega(t)$ . According to this definition, we have

$$p(\underline{c};t) = \overline{\delta[\underline{c} - \underline{C}[\boldsymbol{X}(t);t]]} = \frac{1}{|\Omega(t)|} \int_{\Omega(t)} d\boldsymbol{x} \,\delta[\underline{c} - \underline{C}(\boldsymbol{x};t)], \tag{2}$$

where  $\delta(\cdot)$  is the Dirac delta. Intuitively, this expression encodes the fact that the contributions to the probability of having a value <u>c</u> of concentration at time t correspond to points  $\boldsymbol{x}$  where  $\underline{C}(\boldsymbol{x};t) = \underline{c}$ , which are "counted" by the Dirac delta in accordance with the continuous nature of concentrations (for the general theory of PDFs as averages of Dirac deltas, see, e.g., [64]). This corresponds to the continuum limit described in Fig. 2.

The concentration PDF contains information about the full structure of point statistical variability of concentrations. According to the definition, Eq. (2), moments of concentration, such as the component-wise mean concentrations  $\langle \underline{C}(t) \rangle$  and variances  $\underline{\sigma_c}^2 = \langle \underline{C}^2(t) \rangle - \langle \underline{C}(t) \rangle^2$  (representing not a spatial variance but variability with respect to the mean concentration), may be computed given knowledge of the PDF, and agree with those obtained from the spatial profile. In particular,

$$\langle \underline{C}(t) \rangle = \int_{0}^{\infty} d\underline{c} \, \underline{c} p(\underline{c}; t) = \frac{\underline{M}(t)}{|\Omega(t)|} = \overline{\underline{C}[\mathbf{X}(t); t]},\tag{3}$$

<sup>140</sup> where the total component masses are given by

$$\underline{M}(t) = \int_{\Omega(t)} d\boldsymbol{x} \, \underline{C}(\boldsymbol{x}; t). \tag{4}$$

It is important to note that  $\Omega(t)$  may be time-dependent, but its size  $|\Omega(t)|$  must be finite at all 141 times, because a uniformly-distributed random variable X(t) does not exist in an infinite domain. 142 Intuitively, this is related to the fact that the concentration PDF is defined in terms of the spatial 143 frequency of occurrence of concentration values. Thus, if the total component masses are finite, most 144 regions in an infinite domain  $\Omega(t)$  must have a vanishing concentration, and  $p(\underline{c};t) = 0$  for all  $\underline{c} \neq 0$ , 145 so that we must have trivially  $p(\underline{c};t) = \delta(\underline{c})$  by normalization. In practice, as will be discussed in more 146 detail below, this means that in order to define a non-trivial concentration PDF we must consider 147 either a finite, fixed reference domain  $\Omega$ , or a minimum concentration threshold  $c_m$  such that  $\Omega(t)$  is 148 the region where concentrations are above this threshold. Finally, we note that, in an infinite periodic 149 system, the concentration PDF may be naturally defined over a single period. The previous argument 150 can be circumvented in this case because the total mass in such an infinite system is infinite, although 151 the mass per period is finite. 152

#### <sup>153</sup> 2.2 Single-species formulation

Writing the concentration PDF in a more useful analytical form requires changing variables in the Dirac delta in Eq. (2), in order to identify the spatial points  $\boldsymbol{x}$  that contribute to the integral for a given value of concentration  $\underline{c}$ . In other words, we must invert the relations  $\underline{C}(\boldsymbol{x};t) = \underline{c}$ , which form a set of  $n_s$  constraints in d dimensions. These constraints are not necessarily independent, and this is a difficult problem in general from an analytical perspective. We now derive a simple analytical form for the single-species case. We write  $C = C_1$  and  $c = c_1$  for the single species component. A numerical computation procedure, also valid for the general multispecies PDF, is discussed in Appendix A.

Assuming  $C(\cdot; t)$  to be smooth, the Dirac delta can be expressed as a simple-layer integral [65],

$$\delta[c - C(\boldsymbol{x}; t)] = \int_{\Lambda(c;t)} dS(\boldsymbol{y}) \, \frac{\delta(\boldsymbol{x} - \boldsymbol{y})}{|\nabla C(\boldsymbol{y}; t)|},\tag{5}$$

where  $dS(\boldsymbol{x})$  is the (d-1)-area element at point  $\boldsymbol{x}$  on the (d-1)-surface consisting of the points in the domain at time t at which the concentration has value c, and which do not constitute a local extremum (or zero-gradient inflection point). Formally, this surface is defined as  $\Lambda(c;t) = \{\boldsymbol{x} \in \Omega(t) \mid C(\boldsymbol{x};t) = c \land |\nabla C(\boldsymbol{x};t)| \neq 0\}$ . Note that, for the one-dimensional example in Fig. 2,  $\Lambda(c) = 2$  for all concentrations strictly between the minimum and maximum values. This is discussed in more detail in Section 2.4. Substituting Eq. (5) in Eq. (2), we find

$$p(c;t) = \int_{\Lambda(c;t)} \frac{dS(\boldsymbol{x})}{|\Omega(t)||\nabla C(\boldsymbol{x};t)|}.$$
(6)

This form corresponds to the intuitive notion of a change of variables mentioned above: it expresses the contribution to the probability density of a given value c of concentration as a spatial integral over the spatial region where the concentration takes the required value (see also Fig. 2). This spatial region has dimensionality one less than the full space due to the constraint  $C(\mathbf{x};t) = c$ . This result holds for smooth, non-constant  $C(\cdot;t)$ . The contributions of concentrations found near spatial concentration extrema and in extended regions of constant concentration are discussed below.

Equation (6) can also be expressed so as to highlight the role of concentration gradients associated with a given concentration in determining the value of the PDF. To this end, consider the harmonic average  $g_h(c;t)$  of the concentration gradient magnitudes in  $\Lambda(c;t)$ . By definition, it is given by

$$g_h(c;t) = \left[\frac{1}{|\Lambda(c;t)|} \int_{\Lambda(c;t)} \frac{dS(\boldsymbol{x})}{|\nabla C(\boldsymbol{x};t)|}\right]^{-1}.$$
(7)

<sup>177</sup> Substituting this definition in Eq. (6) immediately yields the simple form

$$p(c;t) = \frac{|\Lambda(c;t)|}{|\Omega(t)|g_h(c;t)}.$$
(8)

The proportionality to the inverse of a concentration gradient magnitude is a consequence of the 178 fact that p(c;t) is a density, i.e., a probability per unit concentration. Spatial regions where the 179 concentration gradient is low correspond to a higher density around the corresponding concentration 180 value due to the slow variation. To understand this, consider a surface  $\Lambda(c;t)$  over which the gradient 181 magnitude  $q_h(c;t) = |\nabla C|$  is homogeneous and therefore equal to its harmonic average. A small 182 variation dc of concentration occurs over a distance  $dz = dc/q_h(c;t)$  along the concentration gradient. 183 The gradient is locally perpendicular to the surface of constant concentration, so that the spatial 184 volume associated with an occurrence of concentration in [c, c + dc] is  $|\Lambda(c; t)|dz$ . Correspondingly, 185 the contribution to the probability p(c;t) dc of concentration in this range is  $|\Lambda(c;t)| dz/|\Omega(t)|$ . Thus, 186 its contribution to the concentration PDF p(c;t) is  $|\Lambda(c;t)|/[|\Omega(t)|q_h(c;t)]$ , in agreement with Eq. (8). 187 This is illustrated for a one-dimensional profile in Fig. 2. More generally, if the gradient is not constant 188 over the surface, the same argument can be applied locally, using the area element dS introduced above 189 and the volume element dSdz. The inverse proportionality to the harmonic average then arises due to 190 the inverse proportionality of the local contributions to the inverse gradient magnitude, which must 191 be integrated over. This provides an intuitive explanation for Eq. (6) and recovers Eq. (8). 192

#### <sup>193</sup> 2.3 Extrema and constant-concentration regions

We now study the impact on the concentration PDF of local extrema of the spatial concentration profile, at which  $|\nabla C(\boldsymbol{x};t)| = 0$ . Near an extremum  $c_0$  at position  $\boldsymbol{x}_0$  and time t, we have  $C(\boldsymbol{x}_0 + \Delta \boldsymbol{x};t) \approx c_0 + \Delta \boldsymbol{x} \cdot H(\boldsymbol{x}_0;t)\Delta \boldsymbol{x}/2$  and  $\nabla C(\boldsymbol{x};t) \approx H(\boldsymbol{x}_0;t)\Delta \boldsymbol{x}$ , where  $H(\boldsymbol{x}_0;t)$  is the Hessian matrix of concentration at position  $\boldsymbol{x}_0$  and time t, with components  $H_{ij}(\boldsymbol{x}_0;t) = \nabla_i \nabla_j C(\boldsymbol{x};t)|_{\boldsymbol{x}=\boldsymbol{x}_0}$ . We find that in d = 1 the PDF of concentration behaves in a vicinity  $\Delta c$  of  $c_0$  as

$$p(c_0 + \Delta c; t) \approx \left( |\Omega(t)| \sqrt{\Delta c \frac{\partial^2 C(x; t)}{\partial x^2}} \Big|_{x=x_0} \right)^{-1},$$
(9)

to first order in  $|\Delta c| \ll c_0$ , for  $\Delta c < 0$  near a maximum and  $\Delta c > 0$  near a minimum. This type of divergent behavior due to a spatial maximum can be observed in Fig. 2. We note that this is an integrable divergence, that is, the integral of  $p(\cdot;t)$  converges near  $c_0$  as necessary for a PDF. This result is a direct consequence of the fact that for a smooth function near an extremum the behavior of the function is quadratic, and the behavior of its derivative is linear. We have assumed the extremum is non-degenerate, so that the second derivative is nonzero; otherwise, higher order terms are important and the behavior is different.

In arbitrary dimension, we provide an argument based on dimensional analysis. First, note that the qualitative behavior near an extremum is determined by the Hessian determinant |H| and the distance  $|\Delta \boldsymbol{x}|$  to the maximum; thus, the gradient near an extremum obeys  $|\nabla C| \propto |H|^{1/d} |\Delta \boldsymbol{x}|$ , and the concentration changes as  $|\Delta c| \propto |\nabla c| |\Delta \boldsymbol{x}| \propto |H|^{1/d} \Delta x^2$ . The second equation gives  $|\Delta \boldsymbol{x}| \propto |H|^{-1/(2d)} |\Delta c|^{1/2}$ , and substituting in the first leads to  $p(c_0 + \Delta c) \approx |\Lambda(c_0 + \Delta c; t)|/[|\Omega(t)|| |H|^{1/(2d)} |\Delta c|^{1/2}]$ . Substituting  $|\Lambda(c_0 + \Delta c; t)| \propto |\Delta \boldsymbol{x}|^{d-1}$  for the (d-1)-area of the surface of concentration  $c_0 + \Delta c$  yields

$$p(c_0 + \Delta c; t) \propto \frac{|\Delta c|^{d/2 - 1}}{|\Omega(t)||H|^{1/2}}.$$
 (10)

Thus, there is no divergence in  $d \ge 2$ . In d = 2, extrema of the spatial concentration correspond to extrema of the concentration PDF, and in d = 3 they correspond to zeros. As before, we have assumed that the minimum is non-degenerate, i.e.,  $|H| \ne 0$ . As a simple example of degeneracy, consider a concentration field which is constant along the second and third dimensions, and exhibits a regular extremum along the first dimension. As expected, the divergence as the extremum is neared then depends only on the distance  $\Delta x$  along the first dimension, and the square-root divergence of the concentration PDF characteristic of d = 1 is recovered.

If  $C(\cdot;t)$  is constant in space (i.e., the concentration is homogeneous), or if we relax the smoothness assumption so that  $|\nabla C(\cdot;t)|$  may be zero in open sets by letting  $C(\cdot;t)$  be non-differentiable at their boundary (i.e, piecewise-smooth), these sets may contribute additional terms to the concentration PDF. Consider first that the concentration is piecewise-constant. Each set  $\Omega_i(t)$  where the concentration takes a given position-independent value  $c_i(t)$  at time t is associated with a probability  $|\Omega_i(t)|/|\Omega(t)|$ . Note that there must be a finite number N(t) of such constant-concentration sets at each time, and thus a finite number of corresponding concentration values, because open sets have nonzero measure and sets corresponding to different concentrations are disjoint, so that  $|\Omega(t)| = \sum_{i=1}^{N(t)} |\Omega_i(t)|$ . Therefore,

$$p(c;t) = \sum_{i=1}^{N(t)} \frac{|\Omega_i(t)|}{|\Omega(t)|} \delta[c - c_i(t)].$$
(11)

The Dirac deltas express the fact that concentration takes a single specific value within spatiallyhomogeneous regions. If the concentration is piecewise-smooth and non-constant over certain parts of the domain, Eq. (8) may be applied separately to each such region, and the contributions of all constant and non-constant regions summed over to obtain the full concentration PDF.

#### <sup>232</sup> 2.4 One-dimensional Gaussian profile

As a relevant example which will play a central role in Section 3, we consider a single-species Gaussian concentration profile in d = 1 spatial dimension (Fig. 1), characterized by spatial variance  $\sigma^2(t)$  and total mass M(t):

$$C(x;t) = \frac{M(t)}{\sqrt{2\pi\sigma^2(t)}} \exp\left[-\frac{x^2}{2\sigma^2(t)}\right].$$
(12)

The corresponding regions of constant concentration c are zero-dimensional,  $\Lambda(c;t) = \{x | x = \pm x_c(c;t)\}.$ 

Inverting C(x;t) = c for x,

$$x_c(c;t) = \sqrt{2\sigma^2(t)\ln[c_M(t)/c]},$$
(13)

238 where

$$c_M(t) = \frac{M(t)}{\sqrt{2\pi\sigma^2(t)}} \tag{14}$$

is the maximum concentration at time t. The zero-dimensional area element on  $\Lambda(c;t)$  is a sum of point measures concentrated at the two points  $\pm x_c(c;t)$  where the concentration takes a given value,  $dS(x) = (\delta[x + x_c(c;t)] + \delta[x - x_c(c;t)]) dx$ . For all  $c < c_M(t)$ ,  $|\Lambda(c;t)| = 2$  because it consists of two points. The gradient of concentration magnitudes at  $\pm x_c(c;t)$  have the same value, so that their harmonic average is equal to this value and given by

$$g_h(c;t) = |\nabla C(x;t)|_{x=\pm x_c(c;t)} = \sqrt{\frac{2c^2 \ln[c_M(t)/c]}{\sigma^2(t)}}.$$
(15)

As discussed in Section 2, we must consider a finite domain in order for the concentration PDF 244 to be well-defined. This can be conveniently achieved through one of two possible approaches. The 245 first considers a fixed (i.e., time-independent) reference domain  $\Omega$  centered at x = 0. In this case, 246 a time-dependent minimum concentration  $c_m(t) = c_M(t) \exp\{-|\Omega|^2/[8\sigma^2(t)]\}$  is found at the fixed 247 domain boundaries. Alternatively, we fix a (time-independent) minimum concentration  $c_m$ , and con-248 sider  $\Omega(t) = \{x | C(x; t) > c_m\}$ , the time-dependent spatial domain where concentration are above the 249 minimum value. In this case,  $c_m$  can be thought of as a lower detection limit, and the concentration 250 PDF as being computed over the region of space where non-zero concentration is detected. In either 251 case, using Eq. (8) leads to the concentration PDF 252

$$p(c;t) = \frac{H[c_M(t) - c]H[c - c_m(t)]}{2c\sqrt{\ln[c_M(t)/c]\ln[c_M(t)/c_m(t)]}},$$
(16)

where under the second approach  $c_m(t) = c_m$  is constant. This PDF is illustrated in Fig. 2, and it agrees with the result reported, e.g., in [30]. By noting that  $|\Omega(t)| = 2x_c[c_m(t); t]$  and using Eqs. (13) and (14), we find

$$|\Omega(t)| = \frac{2M(t)}{c_M(t)} \sqrt{\frac{\ln[c_M(t)/c_m(t)]}{\pi}},$$
(17)

<sup>256</sup> and the concentration PDF can thus also be expressed as

$$p(c;t) = \frac{M(t)H[c_M(t) - c]H[c - c_m(t)]}{|\Omega(t)|cc_M(t)\sqrt{\pi \ln[c_M(t)/c]}}.$$
(18)

Note that, for fixed t, setting  $c = c_M(t) - \Delta c$  and expanding to first order in  $\Delta c$  recovers the expected  $\Delta c^{-1/2}$  divergence due to the presence of a spatial maximum discussed in Section 2.3. For small concentrations compared to the peak value, this PDF scales as 1/c with logarithmic corrections.

Using these results and Eq. (12), it can be verified by direct computation that the mean concentration is given by

$$\langle C(t) \rangle = \overline{C(X;t)} \approx \frac{M(t)}{|\Omega(t)|} = \frac{c_M(t)}{2} \sqrt{\frac{\pi}{\ln[c_M(t)/c_m(t)]}},\tag{19}$$

<sup>262</sup> and for the second moment of concentration we have

$$\langle C^2(t) \rangle = \overline{C^2(X;t)} \approx \frac{M(t)c_M(t)}{\sqrt{2}|\Omega(t)|} = \frac{c_M^2(t)}{2} \sqrt{\frac{\pi}{2\ln[c_M(t)/c_m(t)]}},$$
 (20)

where the approximation corresponds to approximating the total mass M(t) in the infinite domain by the mass within  $\Omega(t)$ , over which concentration is larger than  $c_m(t)$ . Note that, in particular, this approximation requires  $c_M(t) \gg c_m(t)$ , that is, the lower detection limit must be small compared to the peak concentration. Under the same approximation, the variance of concentration is thus

$$\sigma_c^2(t) \approx \frac{\pi c_M^2(t)}{4\ln[c_M(t)/c_m(t)]} \left[ \sqrt{\frac{2\ln[c_M(t)/c_m(t)]}{\pi}} - 1 \right],$$
(21)

267 or equivalently

$$\frac{\sigma_c^2(t)}{\langle C(t) \rangle^2} \approx \sqrt{\frac{2\ln[c_M(t)/c_m(t)]}{\pi}} - 1.$$
(22)

Note that the mean and variance of concentration represent the mean value and the associated variability in the frequency of occurrence of concentration values found throughout the domain. They differ from the more common metrics of mean plume position, which here is zero, and spatial or plume variance, which here is  $\sigma^2(t)$ .

# <sup>272</sup> 3 Concentration PDF evolution under nonlinear reaction and <sup>273</sup> diffusion

<sup>274</sup> In this section, we apply the concepts developed above to the single-species, one-dimensional reaction– <sup>275</sup> diffusion problem

$$\frac{\partial C(x;t)}{\partial t} = D \frac{\partial^2 C(x;t)}{\partial x^2} - r[C(x;t)], \qquad (23)$$

<sup>276</sup> where the reaction rate as a function of concentration has the form

$$r(c) = \kappa c^{\beta},\tag{24}$$

with  $\beta > 0$  the order of the reaction and  $\kappa$  the reaction rate constant in units of  $[C^{1-\beta}T^{-1}]$ . Under-277 standing the role and dynamics of concentration statistics under one-dimensional reaction-diffusion 278 represents a fundamental first step towards treating more complex reaction problems in heterogeneous 279 flow fields. In this connection, we discuss the Ranz transform, which was developed to quantify mixing 280 and reaction in the context of turbulence and combustion problems [62, 66] and has more recently been 281 employed to study these phenomena in porous media [30, 67, 68]. The goal of the Ranz transform is to 282 exploit the fact that material elements in an incompressible flow tend to orient themselves according 283 to the principal directions of strain-induced stretching (Fig. 1), forming thin lamellar structures (lines 284 in 2D or sheets in 3D) [48, 69–74, 74–76]. 285

<sup>286</sup> Due to incompressibility, the thickness of such a material element along the principal axis of com-<sup>287</sup> pression is given by s(t) = s(0)A(0)/A(t), where A(t) is its area (or length in 2D). Assuming that local <sup>288</sup> compression is well described by a first-order Taylor expansion of the flow, and neglecting concentra-<sup>289</sup> tion gradients on the surface in comparison to normal gradients along the compression direction  $\zeta$ , the <sup>290</sup> advection-diffusion-reaction equation becomes [30, 67, 74, 75]

$$\frac{\partial C}{\partial t} = -\frac{\zeta}{s} \frac{ds}{dt} \frac{\partial C}{\partial \zeta} + D \frac{\partial^2 C}{\partial \zeta^2} - r(C).$$
(25)

These approximations provide a very good description of transport at high Péclet number [30, 74, 75]. Applying the Ranz transform [62],

$$\tau = \int_0^t dt' \frac{s^2(0)}{s^2(t')}, \qquad \xi = \frac{s(0)}{s(t)}\zeta, \qquad (26)$$

reduces Eq. (25) to a diffusion-reaction equation in the new time  $\tau$  and rescaled transverse coordinate

 $\xi$ , which takes into account the history of advective deformation:

$$\frac{\partial C}{\partial t} = D \frac{\partial^2 C}{\partial \xi^2} - \frac{s^2}{s_0^2} r(C).$$
(27)

Thus, the full advection-diffusion-reaction problem reduces to a collection of one-dimensional diffusionreaction problems over lamellae. Solving this problem remains highly nontrivial, as it requires determining the statistics of thicknesses s(t) and those of lamella coalescence (overlapping) [30, 46, 67, 68, 77, 78], and the solution of the one-dimensional reaction-diffusion problem (27) for given  $s(\tau)$ , where the reaction rate becomes time-dependent according to a rescaling by  $s^2(\tau)/s_0^2$ . Treating the full problem is beyond the scope of this work. Here, we restrict ourselves to the purely-diffusive case, Eq. (23), which corresponds to setting  $s^2/s_0 = 1$  in Eq. (27), with the power-law reaction rate (24).

9

The dynamics remain nontrivial even under this simplified scenario. When the reaction is nonlinear 302 (order  $\beta$  nonzero and different from unity), the effective kinetics, representing the overall reaction rate 303 for a given amount of available mass, depend on the spatial variability of concentration. In other 304 words, the global reaction rate dM(t)/dt for the total mass is not given by  $\kappa M^{\beta}(t)$ , except when 305  $\beta = 1$  or when the system is well mixed, i.e., uniformly characterized by a single concentration value. 306 Similar considerations apply to the reaction rate in terms of the mean concentration instead of the total 307 mass. Nonetheless, knowledge of the point statistics of concentration as encoded in the concentration 308 PDF is sufficient to fully characterize the global reaction rate, which can be obtained by integrating 309 Eq. (23) over the domain  $\Omega(t)$ . Using integration by parts, the diffusive term can be seen to contribute 310  $2D|\partial C/\partial x|$  evaluated at the domain boundaries, which we neglect under the assumption of a small 311 lower detection limit  $c_m$  as above, we obtain 312

$$\frac{dM(t)}{dt} = -\int_{0}^{\infty} dc r(c)p(c;t) = -\kappa \int_{0}^{\infty} dc \, c^{\beta}p(c;t), \qquad (28)$$

where in the last equality we have expressed the integral of the reactive contribution in terms of the 313 concentration PDF using its definition, Eq. (2). 314

In what follows, we will first derive a dynamical equation for the evolution of the concentration 315 PDF, in terms of a concentration-dependent scalar dissipation rate. We will then discuss and employ 316 the Gaussian spatial profile arising from the weak-coupling approximation of [7] to directly compute 317 the evolution of the concentration PDF using the results of Section 2.4, and compare it to the PDF 318 obtained from numerical solutions of Eq. (23).

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#### Dynamical equation for the concentration PDF 3.1320

The concentration PDF evolves under the combined action of transport and reaction processes. The 321 structure of the reaction-diffusion equation (23), where transport and reaction terms locally contribute 322 independently to the evolution of the concentration field, ensures that the same holds for the evolution 323 of the concentration PDF for each value of concentration. We thus seek to express its evolution as 324

$$\frac{\partial p(c;t)}{\partial t} = \Delta p_D(c;t) + \Delta p_R(c;t), \qquad (29)$$

where the two contributions respectively represent the effect of diffusion and reaction. Note that, while 325 this independence holds locally, the global reaction dynamics (i.e., the evolution of the total mass and 326 other global quantities such as the concentration mean and variance) are affected by the dynamical 327 evolution of the plume through the combined effect of transport and reaction. These dynamics will 328 be discussed in detail in Section 4 using a weak-coupling approximation. We focus in what follows on 329 the case of a fixed minimum detection limit  $c_m$  and a time-varying domain  $\Omega(t)$  where concentrations 330 are above  $c_m$ . This choice avoids the introduction of an arbitrary domain devoid of physical meaning, 331 and corresponds more closely to real-world situations, where a lower detection limit always exists. 332 Nonetheless, note that the introduction of a fixed reference domain may be appropriate if one is in 333 fact interested in a specific spatial region, and the formalism can be easily adapted to that case. 334 As shown in Appendix B, the reactive contribution in Eq. (29) is given by 335

$$\Delta p_R(c;t) = \frac{\partial r(c)p(c;t)}{\partial c} + p(c;t)p(c_m;t)r(c_m).$$
(30)

Note that it is completely determined given knowledge of the concentration PDF and local reaction 336 rate as a function of concentration. The first term represents the net change in the probability density 337 of concentration c due to reaction, and the second term guarantees the PDF remains normalized by 338 accounting for reactive decay of concentrations below the threshold value  $c_m$ . 339

It should be clear that, in general, knowledge of the frequencies of occurrence of concentration 340 values, as expressed by p(c;t), is not sufficient to determine their evolution under transport. This is 341

because concentration fluxes due to advection and diffusion are sensitive to the spatial gradients in the concentration profiles. Thus, without additional information, an equation for the concentration PDF, which encodes only the point statistics of concentration, cannot be closed. However, it is possible to express the concentration PDF's evolution in terms of an additional mixing metric, which corresponds to the scalar dissipation rate (see, e.g., [53, 54]),

$$\chi_x(x;t) = (\nabla C) \cdot (D\nabla C), \tag{31}$$

expressed as a function of concentration. We make the simplifying assumption that the initial concentration profile C(x, 0) is symmetric about the origin and monotonically decreasing away from it (as happens, e.g., for a Gaussian profile or a square pulse). In that case, these properties hold for all times, so that a given concentration value is associated with a single concentration gradient magnitude, simplifying the description. Under these assumptions, the concentration-dependent scalar dissipation rate is given by

$$\chi(c;t) = \chi_x[\pm x_c(c;t);t] = Dg_h^2(c;t),$$
(32)

where the harmonic average gradient magnitude  $g_h$  is equal to the gradient magnitude at both points  $\pm x_c$  where the concentration is equal to c. As shown in Appendix B, we have for the transport contribution:

$$\Delta p_D(c;t) = p(c;t) \left[ \frac{1}{4} \frac{\partial \ln \chi(c;t)}{\partial c} \frac{\partial \chi(c;t)}{\partial c} - \frac{1}{2} \frac{\partial^2 \chi(c;t)}{\partial c^2} - \frac{p(c_m;t)}{2} \frac{\partial \chi(c;t)}{\partial c} \bigg|_{c=c_m} \right],$$
(33)

where the first and second terms arise from the impact of diffusion on the concentration gradients associated with a given concentration value, and the third term is due to the change in domain size due to concentrations dropping below  $c_m$  by diffusion.

Substituting Eqs. (30) and (33) in Eq. (29), we conclude that the combined effect of transport and reaction leads to

$$\frac{\partial p(c;t)}{\partial t} = \frac{\partial r(c)p(c;t)}{\partial c} + \left[ \frac{1}{4} \frac{\partial \ln \chi(c;t)}{\partial c} \frac{\partial \chi(c;t)}{\partial c} - \frac{1}{2} \frac{\partial^2 \chi(c;t)}{\partial c^2} + \left( r(c_m) - \frac{1}{2} \frac{\partial \chi(c;t)}{\partial c} \Big|_{c=c_m} \right) p(c_m;t) \right] p(c;t).$$
(34)

This equation shows that the dynamical evolution of the concentration PDF for this problem is fully 361 determined given knowledge of the local reaction rates r(c) and the concentration-dependent scalar 362 dissipation rate  $\chi(c;t)$ . The first (reactive) term on the right-hand side has the form of an advective 363 contribution: the reaction rate r(c) functions as an advective "velocity" (units of inverse concentration 364 per time) generating a net flux of probability towards lower concentrations. The remaining contri-365 butions have the form of concentration-dependent sink/source terms. The first two terms in square 366 brackets, which are governed by the scalar dissipation rate, encode the impact on the PDF of changes 367 in concentration values and the associated gradients due to diffusive transport. The remaining terms 368 concern the reactive and diffusive fluxes of probability towards concentration values below the de-369 tection limit. These affect the concentration PDF through the normalization across the remaining 370 concentration values, since  $\int_{c_m}^{c_M(t)} dc p(c;t) = 1$  by definition. As discussed in the introduction, the scalar dissipation rate has been studied in connection with 371

As discussed in the introduction, the scalar dissipation rate has been studied in connection with mixing in porous media. By clarifying its role in the evolution of the concentration PDF, this approach provides a promising departure point for upscaling of reaction dynamics under transport.

## <sup>375</sup> 4 Concentration dynamics under the weak-coupling approxi-<sup>376</sup> mation

The weak-coupling approximation developed in [7] simplifies the full reaction-diffusion dynamics of Eq. (23) by assuming that, after a time on the order of the diffusion time, the concentration profile

remains Gaussian as for the conservative problem, but with a spatial variance and mass that are affected by reaction. We briefly review this approach before applying the theory and concepts developed here

to analyze the evolution of concentration statistics.

#### <sup>382</sup> 4.1 Problem setup and nondimensionalization

We will focus here, as in [7], on a rectangular pulse initial condition, corresponding to constant concentration  $c_0 = M_0/s_0$  over a domain  $\Omega(t)$  centered at the origin and of initial width  $s_0 = |\Omega(0)|$ . This setup correspond to a Dirac delta initial condition for the PDF,  $p(c; 0) = \delta(c-c_0)$ . The initial condition allows for meaningful direct comparison with the well-mixed batch problem in the domain  $\Omega(0)$ , which corresponds to the purely-reactive scenario in the absence of diffusion. In that case, D = 0, so that the scalar dissipation rate  $\chi \equiv 0$ , and we denote the associated uniform solution for concentration as a function of time as  $c_B(t)$ .

In addition, in order to meaningfully compare the behavior of the system across different parameter 390 values, it is convenient to nondimensionalize concentration according to the initial concentration  $c_0$ , 391 position according to the initial width  $s_0$ , and time according to the characteristic reaction time [7] 392  $\tau_R = \kappa^{-1} c_0^{1-\beta}$ . Thus, denoting nondimensionalized quantities by an asterisk, nondimensional time 393 is given by  $t_*/\tau_R$ , positions by  $x_* = x/s_0$ , and concentrations by  $C_* = C/c_0$ , seen as a function of 394 nondimensional times and positions. Functions of these quantities, such as the concentration PDF, are 395 nondimensionalized accordingly; further details may be found in Appendix D. In what follows, we work 396 exclusively in nondimensional units and drop the asterisks for notational brevity. The nondimensional 397 transport equation is given by 398

$$\frac{\partial C(x;t)}{\partial t} = \frac{1}{2 \operatorname{Da}} \frac{\partial^2 C(x;t)}{\partial x^2} - r[C(x,t)], \qquad (35)$$

<sup>399</sup> where we have introduced the Damköhler number

$$Da = \frac{\tau_D}{\tau_R}, \qquad \qquad \tau_D = \frac{s_0^2}{2D}. \tag{36}$$

<sup>400</sup> A large Damköhler number corresponds to fast reaction compared to the diffusion time  $\tau_D$  associated <sup>401</sup> with the nondimensionalization lengthscale  $s_0$ , and a low Damköhler number to slow reaction. Note <sup>402</sup> that Eq. (34) for the evolution of the concentration PDF remains unchanged when all intervening <sup>403</sup> quantities are nondimensionalized.

#### 404 4.2 Early times – Batch dynamics

<sup>405</sup> Under the weak-coupling approximation, up to times on the order of the diffusion time, the concen-<sup>406</sup> tration distribution is approximated as not having deformed substantially with respect to the initial <sup>407</sup> condition, here a square pulse of unit width in nondimensional coordinates. Thus, up to nondimen-<sup>408</sup> sional time t = Da, the system is taken to evolve according to reaction in a batch reactor. Since <sup>409</sup> variations in the spatial distribution up to this time are considered negligible and not resolved, under <sup>410</sup> this approximation the concentration PDF remains a Dirac delta for  $t \leq Da$ ,  $p(c; t) \approx \delta[c - c_B(t)]$ . We <sup>411</sup> have (see also Appendix E):

$$\frac{dc_B}{dt} = -c_B^\beta. \tag{37}$$

<sup>412</sup> The special case  $\beta = 1$  corresponds to linear reactions. Decay is then exponential,  $c_B(t) = \exp(-t)$ . <sup>413</sup> In this classical case, reaction and transport dynamics are fully decoupled: the exact solution of the <sup>414</sup> full reactive transport problem is simply the conservative transport solution, multiplied by  $\exp(-t)$  to <sup>415</sup> account for mass loss due to reaction. For nonlinear reactions,  $\beta \neq 1$ , integrating Eq. (37) yields

$$c_B(t) = \left[1 - (1 - \beta)t\right]^{\frac{1}{1 - \beta}}.$$
(38)

In nondimensional units, the total mass and concentration coincide for batch dynamics, because the width of the pulse is approximated as constant and equal to the initial unit width in this time window. Thus, for t < Da we approximate  $M(t) \approx c_B(t)$ . Note that for  $\beta < 1$  the batch reaction is fully depleted in a finite time

$$t_f^B = \frac{1}{1 - \beta}.\tag{39}$$

If  $t_f^B < \text{Da}$ , the full dynamics are thus simply treated as a well-mixed batch reactor.

#### 421 4.3 Late times – Gaussian profile

For times t < Da, [7] have verified that the dynamics are reasonably well approximated by a well-mixed 422 batch reactor, disregarding transport and spatial variability in concentration. We thus focus now on 423 times  $t \ge Da$ , for which spatial concentration variability cannot be neglected. Starting at t = Da, when 424 diffusion has had sufficient time to deform the concentration pulse, the weak-coupling approximation 425 consists in approximating the spatial profile as a Gaussian, whose properties are governed by the 426 combined effect of reaction and diffusion as detailed below. Consider first Eq. (34) for the dynamical 427 evolution of the concentration PDF. Under the Gaussian approximation, the scalar dissipation rate 428 can be computed analytically. From Eqs. (12) and (13) for the Gaussian profile and (77b) for the 429 scalar dissipation rate, we find 430

$$\chi(c;t) = \frac{2\pi c_M^2(t)c^2 \ln[c_M(t)/c]}{\mathrm{Da}\,M^2(t)} H[c_M(t) - c]H[c - c_m],\tag{40}$$

where we have used Eq. (14) to express the spatial variance in terms of the mass and peak concentration. Note that, expanding the logarithm in Taylor series for small  $\Delta c$  near the peak value,  $c = c_M(t) - \Delta c$ , we find that the scalar dissipation rate approaches zero linearly with  $\Delta c$ . At low concentrations compared to the peak value, it scales as  $c^2$  with logarithmic corrections. Using Eq. (40), Eq. (34) for the evolution of the concentration PDF becomes

$$\frac{\partial p(c;t)}{\partial t} = \frac{\partial c^{\beta} p(c;t)}{\partial c} + \left\{ \frac{1 + 2\ln[c_M(t)/c]}{2\ln[c_M(t)/c]} + \left(1 - 2\ln\frac{c_M(t)}{c_m} + c_m^{\beta-1}\right) c_m p(c_m;t) \right\} \frac{\pi c_M^2(t) p(c;t)}{\mathrm{Da} \, M^2(t)}.$$
(41)

Consider first the conservative case, corresponding to omitting the reactive terms involving  $c^{\beta}$  and 436  $c_m^{\beta}$ . As explained above, we consider a Gaussian initial condition at t = Da, and take the detection 437 limit to be sufficiently low for the mass corresponding to concentrations above  $c_m$  to be approximately 438 equal to the total mass for all times of interest. Then, M(t) = 1, and the peak concentration is given 439 by Eq. (14) with a spatial variance  $\sigma^2(t) = \sigma^2(0) + (t - Da)/Da$ . Using Eq. (16) for the Gaussian 440 concentration PDF, it can be verified by direct calculation that Eq. (34) holds identically for all c and 441 t. As a direct verification of the combined effect of reaction and diffusion on the concentration PDF. 442 consider the particular case of a linear decay reaction, r(c) = c, for which analytical solutions are 443 available. In this case, the concentration profile remains exactly Gaussian as before, but with a total 444 mass that evolves in time as  $M(t) = \exp(-t)$ . The spatial variance remains unchanged with respect to 445 the conservative case, and Eq. (16) remains valid with  $c_M(t)$  given by Eq. (14) as before. Once again, 446 direct computation of both sides of Eq. (41) shows that they are identical for all values of c and t. 447

Equation (41) quantifies the full evolution of the concentration PDF and sheds light on the fun-448 449 damental governing processes. However, solving it directly is challenging. We will thus focus instead on extending the weak-coupling approach to directly compute the concentration PDF and associated 450 quantities based on the evolution of the total mass M(t) and the peak concentration  $c_M(t)$ . This will 451 allow us to obtain additional analytical insights into the dynamics. [7] focused on the evolution of 452 the total mass and spatial variance. For the purpose of determining the concentration PDF, and in 453 particular the mean concentration above a detection threshold and the associated variability, it is more 454 convenient to quantify the dynamics in terms of the evolution of total mass and peak concentration, 455

as the latter plays a central role (see Eqs. (12), (19), and (21)). At time t = Da, when we begin approximating the concentration profile as Gaussian, the mass and peak are approximated as

$$M(Da) = c_M(Da) = c_B(Da) = [1 - (1 - \beta) Da]^{\frac{1}{1 - \beta}},$$
(42)

where  $c_B(Da)$  is computed according to the batch solution (38). We note that in [7], instead of matching the peak value to the batch dynamics at t = Da, the spatial variance was set to match that of the initial square pulse,  $\sigma^2(Da) = 1/12$ . This leads to qualitatively similar results and makes no difference at late times. The present choice enforces continuity of  $c_M(t)$  at t = Da, while the previous choice enforces continuity of  $\sigma^2(t)$  instead. Due to Eq. (14), the approximation of a sharp transition between batch dynamics and a Gaussian profile cannot enforce the continuity of both, and for the current choice we have  $\sigma^2(Da) = 1/(2\pi)$ , roughly double the initial spatial variance.

The total mass dynamics are then obtained from the local reaction kinetics. That is, integrating Eq. (35) in space for fixed time and using Eq. (12) for the Gaussian profile, neglecting mass present at concentrations below the detection limit  $c_m$ , we obtain the mass evolution for  $\beta > 0$  as

$$\frac{dM(t)}{dt} = -\frac{M^{\beta}(t)}{\sqrt{\beta}} \left[2\pi\sigma^{2}(t)\right]^{\frac{1-\beta}{2}}.$$
(43)

<sup>468</sup> In order to obtain an equation for the peak concentration, we employ Eq. (35) at x = 0. This gives

$$\frac{dc_M(t)}{dt} = -\frac{c_M(t)}{2\,\mathrm{Da}\,\sigma^2(t)} - c_M^\beta(t).$$
(44)

<sup>469</sup> Using Eq. (14) to express the spatial variance in terms of the mass and peak concentration, we obtain

$$\frac{dM(t)}{dt} = -\frac{M(t)}{\sqrt{\beta}} c_M^{\beta-1}(t), \qquad (45a)$$

470

$$\frac{dc_M(t)}{dt} = -\frac{\pi c_M^3(t)}{\operatorname{Da} M^2(t)} - c_M^\beta(t).$$
(45b)

<sup>471</sup> Note that Eq. (45a) for the overall reaction rate can also be obtained directly from the concentration <sup>472</sup> PDF by using Eqs. (18) and (28). The result follows from setting  $c_m = 0$  in the lower integration <sup>473</sup> limit, which, for  $\beta > 0$ , is equivalent to neglecting the mass present at concentrations lower than  $c_m$ , <sup>474</sup> as was done for the spatial profile in Eq. (43). This confirms that, as expected, knowledge of the <sup>475</sup> concentration PDF directly determines overall reaction rates and mass evolution, without need for <sup>476</sup> additional detailed knowledge about the spatial profile.

Equations (45), with initial conditions at time t = Da according to Eq. (42), fully determine the 477 evolution of total mass and peak concentration in this regime. Given knowledge of the mass and peak 478 concentration, the concentration PDF for  $t \ge Da$ , based on the Gaussian assumption, is given by 479 Eq. (16). In the examples considered here, we fix the minimum concentration, corresponding to an 480 arbitrary lower detection limit, to  $c_m = 10^{-6}$ . Recall that, in nondimensional units, this refers to a 481 detection limit six orders of magnitude below the initial concentration characterizing the rectangular 482 pulse injection. The results are qualitatively similar for different  $c_m$ , so long as the peak concentration 483  $c_M(t)$  remains large compared to  $c_m$  for the times of interest. If  $c_M(t)$  approaches  $c_m$ , an appreciable 484 amount of mass loss will start occurring as it drops below the detection limit. The system will then 485 appear completely depleted, with all detectable mass and concentrations reaching zero. 486

In [7], good agreement was shown to hold between the total mass and spatial variance predicted by the weak-coupling approximation and numerical simulations of Eq. (35). Here, we verify the predictions of this approach, cast in terms of the total mass and peak concentration, for the full concentration PDF, as well as the time-evolution of the associated mean concentration and concentration variance. To this end, we first conduct resolved numerical simulations of the coupled diffusion and reaction problem, Eq. (35). To compute the quantities associated with model predictions, we employ Eqs. (45), which constitute ordinary rather than partial differential equations and can be efficiently solved numerically. The numerical methods used for both these computations are detailed in Appendix C. Furthermore, we obtain analytical predictions for late-time behaviors under the weak-coupling approximation.

In order to test the applicability of the Gaussian assumption for predicting concentration variability at times  $t \ge Da$ , we note that the concentration PDF associated with a Gaussian profile, Eq. (16) can be expressed in terms of concentrations rescaled by their (time-dependent) peak value,  $c' = c/c_M(t)$ , as

$$p'(c';t) = c_M(t)p[c_M(t)c';t] = f_p(c')\frac{H(1-c')H[c'-c'_m(t)]}{\sqrt{\ln[1/c'_m(t)]}},$$
(46)

500 where  $c'_m(t) = c_m/c_M(t)$ , and

$$f_p(c') = \frac{1}{2c'\sqrt{\ln(1/c')}}.$$
(47)

The concentration PDF p'(c';t) of rescaled concentration is independent of time except through the lower cutoff  $c'_m(t)$  (on which it depends in terms of both its range and normalization). The functional dependency  $f_p(c')$  on rescaled concentration under the Gaussian approximation is both timeindependent and independent of the system parameters  $\beta$  and Da. In order to compare this to the resolved numerical results, below we compute the functional dependency according to Eq. (46), i.e., as  $\sqrt{\ln[1/c'_m(t)]}c_M(t)p[c_M(t)c';t]$ .

Similarly, to assess the quality of the Gaussian approximation regarding the scalar dissipation rate, we consider the rescaled form (see Eq. (40))

$$\chi'(c';t) = \frac{\operatorname{Da} M^2(t)}{2\pi c_M^4(t)} \chi[c_M(t)c',t] = f_{\chi}(c')H(1-c')H[c'-c'_m(t)],$$
(48)

<sup>509</sup> where the functional dependency on rescaled concentration is given by

$$f_{\chi}(c') = c'^2 \ln(1/c'). \tag{49}$$

Below, we employ Eq. (48) to compute the functional dependency from numerical simulations, similarly to the case above for the concentration PDF. Comparison of the concentration PDF functional forms provides a metric for the performance of the Gaussian approximation regarding the relative frequency of occurrence of different concentration values, and comparison of the scalar dissipation rate functional forms provides complementary information about the approximation's performance regarding the gradient values associated with each concentration value.

As they correspond to qualitatively different regimes, we consider the cases  $\beta < 1$ ,  $1 < \beta < 3$ , and  $\beta > 3$  separately below. In what follows, we also discuss analytical approximations for late-time behaviors. Before proceeding, we note that the starting point for these results is the fact that, for  $t \ge Da$ , the concentration profile may be approximated as Gaussian with a spatial variance that increases diffusively as  $\sigma^2(t) \approx t/Da$  [7]. Substituting into Eq. (43) and solving the resulting equation, we obtain an approximation for the mass evolution for  $t \ge Da$ ,

$$M(t) \approx \left[ M(\mathrm{Da})^{1-\beta} - \frac{2}{\sqrt{\beta}} \frac{1-\beta}{3-\beta} \left(\frac{2\pi}{\mathrm{Da}}\right)^{\frac{1-\beta}{2}} \left[ t^{\frac{3-\beta}{2}} - \mathrm{Da}^{\frac{3-\beta}{2}} \right] \right]^{\frac{1}{1-\beta}},$$
 (50)

valid for  $\beta \neq 1, 3$ . Recall that the case  $\beta = 1$ , corresponding to a first-order decay reaction, can be fully solved analytically. The case  $\beta = 3$ , which we do not analyze in detail here for brevity, behaves qualitatively similarly to  $\beta > 3$ , although extremely slow (logarithmic) decay of mass occurs at late times [7].

#### <sup>526</sup> 4.4 Reaction order $\beta < 1$ — Full-depletion regime

For  $\beta < 1$ , reaction rates increase with decreasing concentration, leading to full depletion in finite time. Under fully-mixed conditions, the depletion time  $t_f^B$  is given by Eq. (39). Recall that for <sup>529</sup> sufficiently large Da full depletion occurs before transport must be taken into account. We thus focus <sup>530</sup> on Damköhler numbers such that  $Da < t_f^B$ .

<sup>531</sup> Under the combined effect of reaction and diffusion, mass is still fully depleted in a finite time  $t_f$ , <sup>532</sup> different from  $t_f^B$ . Setting  $M(t_f) = 0$  in Eq. (50) and solving for  $t_f$  yields [7]

$$t_f \approx \left[ \mathrm{Da}^{\frac{3-\beta}{2}} + \frac{\sqrt{\beta}}{2} \frac{3-\beta}{1-\beta} \left( \frac{\mathrm{Da}}{2\pi} \right)^{\frac{1-\beta}{2}} \left[ 1 - (1-\beta) \,\mathrm{Da} \right] \right]^{\frac{2}{3-\beta}}.$$
 (51)

For sufficiently small detection limit  $c_m$ , we expect this prediction to provide a good estimate of the time at which the peak, mean concentration, and concentration variance reach zero. Interestingly, the depletion time decreases with decreasing Da. This decrease is due to the fact that, because in this regime reaction is faster at lower concentrations, stronger diffusion (smaller Da) leads to faster depletion when compared to a system with the same reaction rate. For this reason, both mass and peak concentration decay faster than the batch solution. In particular, for small Da  $\ll$  1, we find a simple power-law scaling with Da,

$$t_f \approx \left(\frac{\sqrt{\beta}}{2}\frac{3-\beta}{1-\beta}\right)^{\frac{2}{3-\beta}} \left(\frac{\mathrm{Da}}{2\pi}\right)^{\frac{1-\beta}{3-\beta}}.$$
(52)

Note however that  $t_f$  / Da increases with decreasing Da, meaning that with increasing diffusion the depletion time occurs increasingly later than the time t = Da at which the Gaussian approximation kicks in.



Figure 3: Functional dependency of the concentration PDF in terms of concentration rescaled by peak value, for  $\beta = 1/2$  and (a): slow reaction,  $Da = 10^{-2}$ , (b): intermediate reaction,  $Da = 10^{-1}$ , and (c): fast reaction, Da = 1. Colors stand for different times, with markers showing the results of resolved numerical simulations of the fully-coupled reaction–diffusion problem (35). Solid lines represent the analytical solution (47) under the weak-coupling model. The dashed lines show a pure 1/c' scaling for reference at low concentrations and the scaling of the inverse-square-root divergence near the maximum concentration value (see Eqs. (9) and (47)).

To illustrate these results, we take  $\beta = 1/2$  as a representative example. First, we compare the functional dependency of the concentration PDF (see Eqs. (46) and (47)) under the weak-coupling approximation to that obtained from resolved numerical simulations of Eq. (35) (see Appendix A and Appendix C). We focus on times t such that  $Da \leq t < t_f$  (see Eq. (51)). Results for different Damköhler numbers and times are shown in Fig. 3. Overall, the Gaussian approximation provides good estimates of the relative frequency of concentration values. Because, for  $\beta < 1$ , reaction is faster at lower concentrations, lower PDF values compared to the Gaussian prediction are observed at low values



Figure 4: Functional dependency of the scalar dissipation rate in terms of concentration rescaled by peak value, for  $\beta = 1/2$  and (a): slow reaction,  $Da = 10^{-2}$ , (b): intermediate reaction,  $Da = 10^{-1}$ , and (c): fast reaction, Da = 1. Colors stand for different times, with markers showing the results of resolved numerical simulations of the fully-coupled reaction–diffusion problem (35). Solid lines represent the analytical solution (49) under the weak-coupling model. The dashed lines show a pure  $c'^2$  scaling for reference at low concentrations and the linear approach to zero near the peak concentration (see Eqs. (40) and (49)).

of concentration. This discrepancy at low concentrations also leads to a corresponding overprediction of higher concentrations, due to the normalization condition for the PDF. Although the bulk of the mass distribution is generally well captured, as discussed below, high Da leads to significant depletion at early times and results in a worse approximation.

Similar conclusions are reached from consideration of the functional dependency of the scalar dissipation rate (see Eqs. (48) and (49)), shown in Fig. 4. The Gaussian approximation again provides good estimates, but higher scalar dissipation rates, corresponding to higher gradients, are observed at low concentration values. The discrepancy is more pronounced in the same cases and for the same reasons as discussed regarding the concentration PDF, although in this case normalization does not play a role and thus no associated discrepancy at higher concentration values is observed.

Next, we turn to the evolution of the mass and peak concentration, shown in Fig. 5. As predicted 560 by the weak-coupling approximation, both quantities reach zero in finite time as complete depletion 561 occurs. However, due to the nonlinear nature of the reaction and the presence of diffusion, the peak 562 dynamics differ from the total mass dynamics, as captured by Eqs. (45). Note the onset of a diffusive 563 scaling of peak decay for low Damköhler, showing that the dynamics are controlled by diffusion at 564 times  $t \sim Da$  (corresponding to dimensional times on the order of the diffusion time). For late times, 565 as concentration becomes smaller, reaction becomes more efficient. The dynamics depart from the 566 diffusive scaling and full depletion eventually occurs. The slight underprediction of the depletion time  $t_f$ 567 by the analytical approximation compared to the numerically-computed weak-coupling approximation 568 results is due mostly to the breakdown of the linear variance growth approximation just before complete 569 depletion, as observed in [7]. When comparing the predictions of the weak-coupling approximation to 570 the resolved numerical simulations, we find overall good agreement. Regarding peak concentrations, 571 the approximation of a sharp transition between batch and Gaussian-profile dynamics at t = Da572 leads to some discrepancy at times  $t \sim Da$ , which becomes more pronounced with increasing Da. As 573 the Damköhler number increases and the depletion time approaches Da from above  $(t_f/\text{Da} \sim 1)$ , see 574 Eq. (51), stronger discrepancies are observed for both the mean concentration and the total mass, 575 due to significant depletion before an approximately Gaussian profile is achieved. Nonetheless, the 576 predictions for  $t_f$  still provide a reasonable approximation. Recall that, for higher Da >  $t_f^B = 2$ 577 (Eq. (39)), diffusion does not play a role before depletion the dynamics become well described by 578 simple batch kinetics. 579



Figure 5: Temporal evolution of (a): total mass and (b): peak concentration for  $\beta = 1/2$ . Colors stand for different Damköhler numbers, markers show the results of resolved numerical simulations of the fully-coupled reaction-diffusion problem (35), solid lines are numerical results based on the weakcoupling model (Eqs. (45)), and dash-dotted lines show the purely-diffusive behavior. The vertical dashed lines show approximate analytical results for the depletion time (Eq. (51)).

Note that defining a mean concentration in terms of a fixed, reference spatial region, as done 580 in [7], leads to a mean concentration scaling identical to that of the total mass, since in that case 581  $\langle C(t) \rangle = M(t)/|\Omega|$  with constant domain size  $|\Omega|$ . Interestingly, when the mean concentration is 582 instead defined, as here, in terms of the temporally-variable region where concentration is above a 583 detection limit, the concentration mean and variance are instead fully controlled by the peak dynamics, 584 see Eqs. (19) and (21). This leads also to a similar quality of the weak-coupling approximation when 585 compared to the resolved simulations, as shown in Fig. 6. In particular, for low Damköhler numbers, 586 our approach accurately predicts a transition from diffusion- to reaction-controlled dynamics, governed 587 by the peak behavior described above, for both the mean concentration and the concentration variance. 588

#### $_{599}$ 4.5 Reaction order $1 < \beta < 3$ — Power-law-decay regime

In this regime, the interplay between reaction and diffusion leads to power-law mass decay at late times. Indeed, for  $t \gg \text{Da}$ , Eq. (50) reduces to [7]

$$M(t) \approx \sqrt{\frac{2\pi}{\mathrm{Da}}} \left(\frac{\sqrt{\beta}}{2} \frac{3-\beta}{\beta-1}\right)^{\frac{1}{\beta-1}} t^{-\frac{3-\beta}{2(\beta-1)}}.$$
(53)

<sup>592</sup> Substituting in Eq. (45b), and noting that, for  $1 < \beta < 3$ , as the peak concentration decreases the <sup>593</sup> reactive term eventually dominates over the diffusive term, we have for late times

$$\frac{dc_M(t)}{dt} \approx -c_M^\beta(t). \tag{54}$$

<sup>594</sup> Solving this equation, we obtain the late time peak decay

$$c_M(t) \approx [(\beta - 1)t]^{-\frac{1}{\beta - 1}}.$$
 (55)

This behavior is characteristic of reaction-dominated conditions, and identical to the late-time batch dynamics, see Eq. (38). However, note that the total mass, Eq. (53), exhibits a different temporal scaling, as it results from the reactive decay of Gaussian-distributed concentration values undergoing diffusive spreading. This results, in contrast to the  $\beta < 1$  regime, in slower asymptotic mass decay than predicted by batch dynamics, because in this case reaction is slower at lower concentrations.



Figure 6: Temporal evolution of (a): mean concentration and (b): concentration variance for  $\beta = 1/2$ . Colors stand for different Damköhler numbers, markers show the results of resolved numerical simulations of the fully-coupled reaction-diffusion problem (35), solid lines are numerical results based on the weak-coupling model (Eqs. (45)), and dash-dotted lines show the purely-diffusive behavior. The vertical dashed lines show approximate analytical results for the depletion time (Eq. (51)).

We now take  $\beta = 2$  as a representative example. Again, we start by assessing the performance 600 of the Gaussian assumption for the functional dependency of the concentration PDF (Fig. 7). The 601 approximation provides very good estimates of the frequency of occurrence of different concentrations. 602 A slight underestimation of low concentrations, and corresponding overestimation of intermediate con-603 centrations, occurs at late times for large Damköhler number. When considering the scalar dissipation 604 rate (Fig. 8), we again observe very good agreement, with a corresponding slight but more discernible 605 overestimation of concentration gradients at low concentrations. In this case, reaction is less efficient 606 at low concentrations, so these results are somewhat counter-intuitive. Indeed, reaction is stronger 607 and therefore has a more pronounced effect compared to diffusion at high concentrations. However, 608 when the corresponding spatial profile is compared to a Gaussian of the same variance and maximum 609 concentration, the latter underestimates the intermediate values but overestimates the tails. This is 610 illustrated in Fig. 9 for the case of  $Da = 10^2$  at time t = 10 Da. This effect is slight, but it is more 611 pronounced at the level of concentration gradients. 612

Next, we consider the evolution of total mass and peak concentration, see Fig. 10. The analytical 613 approximation captures the asymptotic scalings correctly, although it slightly underpredicts the mass 614 and overpredicts the peak concentration when compared to the numerically-computed weak-coupling 615 approximation results. These discrepancies result in this case from small corrections to the  $\sigma^2(t) \approx$ 616 t/ Da diffusive spatial variance growth approximation of [7], implicit in Eq. (50) for the approximate 617 late-time mass evolution. Again, the onset of a purely-diffusive scaling of peak decay is visible for low 618 Damköhler before the asymptotic regime is reached. The weak-coupling approximation provides very 619 good estimates of the resolved dynamics, even for  $Da \sim 1$ . As expected, the largest discrepancies occur 620 for times  $t \sim Da$ , around which the approximation of a sharp transition between batch and Gaussian-621 profile dynamics has the most impact. In addition, a sharp depletion of mass can be observed at 622 late times for the high-Da cases due to the detection limit. As expected, the transition from the 623 dynamics predicted by the weak-coupling approximation to this sharp depletion occurs when the peak 624 concentration becomes comparable to  $c_m$ . 625

As for the  $\beta < 1$  case, the concentration mean and variance (Fig. 11) are controlled by the peak dynamics, see Eqs. (19) and (21). However, the changes in the size of the domain where concentration is larger than the detection limit lead to logarithmic corrections to the purely-reactive power-law scaling observed for the peak concentration. As for the mass and peak, the weak-coupling approximation



Figure 7: Functional dependency of the concentration PDF in terms of concentration rescaled by peak value, for  $\beta = 2$  and (a): slow reaction, Da =  $10^{-2}$ , (b): intermediate reaction, Da = 1, and (c): fast reaction, Da =  $10^2$ . Colors stand for different times, with markers showing the results of resolved numerical simulations of the fully-coupled reaction–diffusion problem (35). Solid lines represent the analytical solution (47) under the weak-coupling model. The dashed lines show a pure 1/c' scaling for reference at low concentrations and the scaling of the inverse-square-root divergence near the maximum concentration value (see Eqs. (9) and (47)).

<sup>630</sup> provides very good predictions of the full dynamics, with the quality of the approximation improving <sup>631</sup> for late times  $t \gtrsim 10$  Da. The discrepancies in the analytical vs numerical results for the weak-coupling <sup>632</sup> approximation result from the discrepancies in the computation of the peak concentration discussed <sup>633</sup> above.

#### <sup>634</sup> 4.6 Reaction order $\beta > 3$ — Incomplete-depletion regime

In this case, the substantial reaction slowdown associated with decreasing concentrations, enhanced by diffusion-induced dilution, prevents the complete depletion of mass by reaction, even for arbitrarily large times. Taking the limit  $t \to \infty$  in Eq. (50) yields the leftover mass [7]

$$M_{\infty} \approx \left[1 + \left(1 + \frac{2}{(\beta - 3)\sqrt{(2\pi)^{\beta - 1}\beta}}\right)(\beta - 1)\operatorname{Da}\right]^{-\frac{1}{\beta - 1}}.$$
(56)

<sup>638</sup> Substituting in Eq. (45b) as before, we now conclude that the late-time dominant contribution is <sup>639</sup> diffusive, so that

$$\frac{dc_M(t)}{dt} \approx -\frac{\pi c_M^3(t)}{\operatorname{Da} M_\infty^2}.$$
(57)

<sup>640</sup> Solving this equation leads, for large times, to

$$c_M(t) \approx M_\infty \sqrt{\frac{\mathrm{Da}}{2\pi t}}.$$
 (58)

As the mass becomes constant at late times, the decay of peak concentration exhibits the signature of diffusive plume spreading. The mass in a well-mixed batch decreases asymptotically to zero, meaning the diffusing system leads to less efficient reaction, as expected since the reaction slows down strongly with decreasing concentration in this case. In contrast, however, the diffusion-governed peak decay is asymptotically faster than the corresponding batch prediction, Eq. (38).

To illustrate this regime, we take  $\beta = 4$  as a representative example. As predicted by the weakcoupling approximation, reaction is sufficiently inefficient at low concentrations for diffusive spreading



Figure 8: Functional dependency of the scalar dissipation rate in terms of concentration rescaled by peak value, for  $\beta = 2$  and (a): slow reaction,  $Da = 10^{-2}$ , (b): intermediate reaction, Da = 1, and (c): fast reaction,  $Da = 10^2$ . Colors stand for different times, with markers showing the results of resolved numerical simulations of the fully-coupled reaction–diffusion problem (35). Solid lines represent the analytical solution (49) under the weak-coupling model. The dashed lines show a pure  $c'^2$  scaling for reference at low concentrations and the linear approach to zero near the peak concentration (see Eqs. (40) and (49)).

to completely arrest mass decay. This means that the system dynamics are essentially dominated by diffusion. Correspondingly, both the concentration PDF and the scalar dissipation rate show excellent agreement with the Gaussian prediction, as shown in Figs. 12 and 13, respectively.

Next, we consider the evolution of total mass and peak concentration, see Fig. 14. In this case, the 651 analytical asymptotic predictions provide very accurate estimates of the weak-coupling dynamics. As 652 expected from the analytical discussion, the results become indistinguishable from a purely diffusive 653 system for sufficiently low Da. However, when comparing to the resolved numerical simulations, we 654 find that the weak-coupling approximation underpredicts the asymptotic mass at higher Da. The main 655 reason for the discrepancy is that, because the reaction is particularly inefficient at low concentrations, 656 the system transitions to the Gaussian regime earlier in this case. Indeed, as shown by the dash-dotted 657 lines in Fig. 14, assuming the transition occurs at t = Da / 10 rather than at t = Da leads to improved 658 estimates of the asymptotic mass. To obtain this prediction, we proceed as for Eq. (56) but set the 659 initial condition of the weak-coupling Gaussian regime at t = Da/10 (see Eq. (42)), leading to 660

$$M_{\infty} \approx \left[ 1 + \left( 1 + \frac{2 \cdot 10^{\frac{\beta - 1}{2}}}{(\beta - 3)\sqrt{(2\pi)^{\beta - 1}\beta}} \right) (\beta - 1) \frac{\text{Da}}{10} \right]^{-\frac{1}{\beta - 1}}.$$
 (59)

<sup>661</sup> Note that the scaling of asymptotic mass with Da, and the relationship between asymptotic peak <sup>662</sup> dynamics and asymptotic mass, Eq. (58), remain unchanged.

The concentration mean and variance for this case are shown in Fig. 15. The considerations 663 pertaining to the  $1 < \beta < 3$  case regarding the scaling of concentration mean and variance in terms of 664 the peak concentration remain valid here, but in this case we observe logarithmically-corrected diffusive 665 rather than reactive scalings. Once again, the low-Da case is indistinguishable from purely-diffusive 666 dynamics. In addition, due to the low efficiency of reaction at low concentrations, the scaling behavior 667 of both mean concentration and concentration variance remains purely diffusive, even for large Da. 668 Overall, as for the total mass and peak concentration, the analytical approximation provides very good 669 asymptotic estimates of the weak-coupling approximation, but better predictions of the fully-resolved 670 dynamics are obtained by assuming the transition to the Gaussian regime occurs at time t = Da/10671 672 rather than at t = Da.



Figure 9: Concentration profile for  $\beta = 2$  and  $\text{Da} = 10^2$ , at time t = 10 Da. The solid line shows the profile computed from resolved numerical simulations of Eq. (35), and the dashed colored line shows a Gaussian profile with the same variance  $\sigma^2(t)$  and maximum concentration  $c_M(t)$ . The horizontal dashed line shows the lower detection limit.

#### 5 Discussion and conclusions

We have presented a detailed discussion of the theoretical and numerical properties of the concentration PDF of a reactive plume evolving under the coupled action of dilution and nonlinear reaction. The concentration PDF encodes the full point statistical variability of concentration values found throughout a spatial domain, and is therefore of central interest to both fundamental and applied reactive transport and risk assessment problems.

As a first step towards solving the full advection-diffusion-reaction system, we have conducted a 679 detailed analysis of nonlinear decay of a single solute undergoing one-dimensional diffusion. We have 680 shown that the weak-coupling approximation developed in [7] to quantify the evolution of total mass 681 and mean concentration in a fixed reference domain can be extended to quantify the evolution of 682 mean concentrations above a lower detection limit, along with the associated variability. In particular, 683 we have found that the late-time Gaussian profile approximation leads to good predictions of the 684 concentration PDF. Even for this simple one-dimensional diffusive system, the effective reaction kinetics 685 governing these quantities exhibit rich dynamics. In contrast to well-mixed batch kinetics, the decay 686 of concentrations at different Damköhler numbers can be dominated by diffusive or reactive effects 687 at different times. Late-time concentrations can exhibit complete extinction, power-law decay, or 688 complete arrest of reaction depending on the reaction order. The Gaussian approximation underlying 689 the weak-coupling approach allows us to predict the associated concentration variance. Remarkably, 690 the dynamics of the maximum concentration value, which provide the key control on the mean and 691 variance of concentrations above a given detection threshold, differ qualitatively from the dynamics of 692 total mass. In particular, for low Damköhler numbers, both the mean concentration and associated 693 variability exhibit diffusion-controlled decay for times on the order of the diffusion time, for any reaction 694 order. This diffusive control persists across all times and Damköhler numbers for strongly superlinear 695 decay reactions (of order  $\beta > 3$ ), whereas reaction is the fundamental mechanism controlling late-time 696 and/or large-Da dynamics for lower reaction orders. 697

We have also derived a dynamical equation for the evolution of the concentration PDF in the nonlinear reaction-diffusion problem, showing that the impact of transport is encoded in the scalar dissipation rate seen as a function of concentration. It is important to note that the generalization of this equation to multiple dimensions, variable advection and/or diffusion tensor, and arbitrary reaction rates, for which concentration is no longer necessarily monotonically decreasing, is not straightforward.



Figure 10: Temporal evolution of (a): total mass and (b): peak concentration for  $\beta = 2$ . Colors stand for different Damköhler numbers, markers show the results of resolved numerical simulations of the fully-coupled reaction-diffusion problem (35), solid lines are numerical results based on the weakcoupling model (Eqs. (45)), dashed lines show approximate asymptotic analytical results (Eqs. (53) and (55)), and dash-dotted lines show the purely-diffusive behavior.

Nonetheless, we believe the approach developed here, highlighting the role of the scalar dissipation rate, has the potential to form the basis for upscaling procedures in more complex systems. In particular, as discussed in Section 3, the dynamics of reaction under advection-diffusion in heterogeneous flows, in both two- and three-dimensional porous media, can be reduced to one-dimensional reaction-diffusion equations along the principal direction of compression on a collection of lamellar structures. With this in mind, further work will focus on the generalization of the approach developed here to the dynamics of Gaussian profiles subject to stretching-enhanced diffusion over an ensemble of lamellae.

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717 Availability of data and material: This work did not make use of or generate any datasets.

718 Code availability Numerical simulations used the py-pde open-source Python package for solving

<sup>719</sup> partial differential equations [79]. Please contact the authors for any additional information or requests.

## $_{^{720}}$ A Discretization

<sup>721</sup> When computing the concentration PDF numerically, discretizations are typically employed both spa-<sup>722</sup> tially and for the concentration values. As a simple example that highlights the central concepts, <sup>723</sup> consider a regular spatial discretization into a grid with constant cell volume  $V_g$ , with each grid cell <sup>724</sup> associated with the average concentration within it. According to the argument for constant concen-<sup>725</sup> tration regions developed in the previous section, for each fixed time t, the concentration PDF is then



Figure 11: Temporal evolution of (a): mean concentration and (b): concentration variance for  $\beta = 2$ . Colors stand for different Damköhler numbers, markers show the results of resolved numerical simulations of the fully-coupled reaction-diffusion problem (35), solid lines are numerical results based on the weak-coupling model (Eqs. (45)), dashed lines show approximate asymptotic analytical results (Eqs. (19) and (21), using Eq. (55) for the peak concentration), and dash-dotted lines show the purely-diffusive behavior.

726 estimated as

$$p(c;t;N_g) = \frac{1}{N_g} \sum_{i=1}^{N_g} \delta[c - c_i(t)],$$
(60)

where  $N_g = |\Omega|/V_g$  is the number of grid points and  $c_i$  is the average concentration in cell *i*. Note that the  $c_i$  associated with different cells are not necessarily all different.

Further discretizing concentration into bins leads to a probability mass function with values for the probability of concentration in each bin  $B_k = [c^{(k)}, c^{(k+1)}], k \ge 0$ :

$$p_k(t; N_g) = \frac{1}{N_g} \sum_{i=1}^{N_g} H[c_i(t) - c^{(k)}] H[c^{(k+1)} - c_i(t)],$$
(61)

where  $H(\cdot)$  is the Heaviside step function. This means that the probability of finding a concentration value in bin k is the fraction of cells where the concentration falls within bin k. If we take concentration bin widths  $\Delta c$  to be constant,  $c^{(k)} = k\Delta c$  for  $k \ge 0$ , and approximate the PDF of concentration by dividing probabilities by  $\Delta c$  (as in Fig. 2), we obtain

$$p(c;t;N_g,\Delta c) = \frac{1}{N_g \Delta c} \sum_{i=1}^{N_g} H[c_i(t) - k\Delta c] H[(k+1)\Delta c - c_i(t)].$$
(62)

These discretization procedures generalize directly to a multispecies system. In that case, concentration bins refer to the simultaneous attainment of concentration values of each species. The associated probabilities are computed as above by counting spatial cells where these values occur simultaneously. The procedure also generalizes directly to non-uniform and/or time-dependent spatial cell sizes and/or concentration bins. More involved techniques employing kernel reconstructions of the concentration field may be formalized in a similar manner [80–82].

The maximum resolution of the discretized concentration PDF described above is given by  $1/(N_g\Delta c)$ , and the maximum PDF value is  $1/\Delta c$ . We now discuss the impact of local spatial extrema, which can be associated with divergences of the continuous PDF as discussed in Section 2.3, on the discretized computation. It is also important to note that sources of error typically come into play in the



Figure 12: Functional dependency of the concentration PDF in terms of concentration rescaled by peak value, for  $\beta = 2$  and (a): slow reaction,  $Da = 10^{-2}$ , (b): intermediate reaction, Da = 1, and (c): fast reaction,  $Da = 10^2$ . Colors stand for different times, with markers showing the results of resolved numerical simulations of the fully-coupled reaction-diffusion problem (35). Solid lines represent the analytical solution (47) under the weak-coupling model. The dashed lines show a pure 1/c' scaling for reference at low concentrations and the scaling of the inverse-square-root divergence near the maximum concentration value (see Eqs. (9) and (47)).

determination of the spatial concentration field. For example, if the latter is computed based on a particle tracking simulation of some transport and reaction dynamics, fluctuations arise due to the finite number of particles, whereas in a standard Eulerian simulation the spatial discretization impacts the determination of the true concentration field. We focus here on the error resulting from concentration discretization  $\Delta c$ , which is dominant given a sufficiently-resolved spatial concentration field.

Consider first a region of constant concentration. A delta peak  $\delta(c-c_0)|\Omega_0|/|\Omega|$  in the concentration 750 PDF, associated as discussed above to a subdomain  $\Omega_0$  of volume  $|\Omega_0|$  where  $C(\boldsymbol{x};t) = c_0$ , corresponds 751 under sufficiently fine spatial discretization to a discretized contribution  $|\Omega_0|/(|\Omega|\Delta c)$ . Thus, halving 752 the concentration discretization  $\Delta c$  leads to a doubling of the numerically-computed peak. More 753 generally, refining the discretization as  $\Delta c \rightarrow a\Delta c$ , a < 1, yields  $p(c_0; t; N_g, a\Delta c) = ap(c_0; t; \Delta c)$ . 754 On the other hand, smooth extrema correspond to divergences only for spatial dimension d = 1, for 755 which they lead to an inverse-square-root divergence, as shown in Section 2.3. In this case, averaging 756 the concentration over a range  $\Delta c$  near the spatial extreme value  $c_0$  yields  $p(c_0;t;\Delta c) \propto 1/\sqrt{\Delta c}$ , 757 so that  $p(c_0; t; a\Delta c) \approx \sqrt{ap(c_0; t; \Delta c)}$ . Thus, observing these behaviors in a numerical computation 758 is a signature of the presence of a spatial extremum, and the scaling behavior with concentration 759 discretization refinement indicates its type. 760

## <sup>761</sup> B Dynamical equation for the concentration PDF

In this appendix, we provide a detailed derivation of the dynamical equation for the concentration PDF 762 discussed in the main text, Eq. (34), by explicitly determining the diffusive transport and reaction 763 contributions in Eq. (29). We first consider the effect of diffusion,  $\Delta p_D(c;t)$ . For one-dimensional 764 diffusion and nonlinear concentration decay, assuming a symmetric initial condition about the origin 765 and monotonically decreasing with distance from the latter, the spatial concentration profile retains 766 these properties for all times. Therefore, we have  $|\Lambda(c;t)| = 2$  for all concentrations within the range 767 observed at time t, corresponding to the two points  $x = \pm x_c(c;t)$  where C(x;t) = c. Furthermore, the 768 concentration gradient magnitude is the same at  $\pm x_c$ , so that its harmonic average is simply  $g_h(c;t) =$ 769  $\nabla C[-x_c(c;t),t] = -\nabla C[x_c(c;t),t]$ . Using Eq. (8) and the fact that  $|\Lambda(c;t)|$  is time-independent within 770



Figure 13: Functional dependency of the scalar dissipation rate in terms of concentration rescaled by peak value, for  $\beta = 2$  and (a): slow reaction,  $Da = 10^{-2}$ , (b): intermediate reaction, Da = 1, and (c): fast reaction,  $Da = 10^2$ . Colors stand for different times, with markers showing the results of resolved numerical simulations of the fully-coupled reaction–diffusion problem (35). Solid lines represent the analytical solution (49) under the weak-coupling model. The dashed lines show a pure  $c'^2$  scaling for reference at low concentrations and the linear approach to zero near the peak concentration (see Eqs. (40) and (49)).

 $\tau_{11}$  the concentration range observed, we can write the change in the concentration PDF as

$$\frac{\partial p(c;t)}{\partial t} = -p(c;t) \left[ \frac{\partial \ln |\Omega(t)|}{\partial t} + \frac{\partial \ln g_h(c;t)}{\partial t} \right].$$
(63)

<sup>772</sup> If a fixed reference volume is considered, the term corresponding to the change of  $|\Omega(t)|$  in time is <sup>773</sup> zero. As discussed in the main text, we focus here on the case of a minimum detection limit  $c_m$  and a <sup>774</sup> time-varying domain  $\Omega(t)$  where  $c > c_m$ .

In order to compute the terms in square brackets, we consider the time evolution of quantities on a given concentration surface (in one dimension, at the points  $\pm x_c(c;t)$ ). By definition, the change in time of concentration over such a surface is zero, so that

$$\frac{\partial C[x_c(c;t);t]}{\partial t} = \left[\frac{\partial x_c(c;t)}{\partial t}\frac{\partial C(x;t)}{\partial x} + \frac{\partial C(x;t)}{\partial t}\right]_{x=x_c(c;t)} = 0$$
(64)

Taking into account that, at  $x = x_c$ ,  $\partial/\partial x = -g_h(c;t)\partial/\partial c$ , the changes associated with transport lead to

$$\frac{\partial x_c(c;t)}{\partial t} = \frac{1}{2g_h(c;t)} \frac{\partial \chi(c;t)}{\partial c},\tag{65}$$

where  $\chi(c;t) = \chi_x[\pm x_c(c;t);t] = Dg_h^2(c;t)$  is the concentration-dependent scalar dissipation rate. The same approach for the variation of the concentration gradient leads to

$$\frac{\partial g_h(c;t)}{\partial t} = -\left[\frac{\partial x_c(c;t)}{\partial t}\frac{\partial^2 C(x;t)}{\partial x^2} + \frac{\partial}{\partial t}\frac{\partial C(x;t)}{\partial x}\right]_{x=x_c(c;t)},\tag{66}$$

782 and we find

$$\frac{\partial \ln g_h(c;t)}{\partial t} = \frac{1}{2} \frac{\partial^2 \chi(c;t)}{\partial c^2} - \frac{1}{4} \frac{\partial \ln \chi(c;t)}{\partial c} \frac{\partial \chi(c;t)}{\partial c}.$$
(67)

The change in domain volume  $\Omega(t) = 2x_c(c_m; t)$  due to dilution of concentration below the detection limit  $c_m$  obeys

$$\frac{\partial |\Omega(t)|}{\partial t} = 2 \frac{\partial x_c(c_m; t)}{\partial t},\tag{68}$$



Figure 14: Temporal evolution of (a): total mass and (b): peak concentration for  $\beta = 4$ . Colors stand for different Damköhler numbers, markers show the results of resolved numerical simulations of the fully-coupled reaction-diffusion problem (35), solid lines are numerical results based on the weakcoupling model (Eqs. (45)), and dashed lines show approximate asymptotic analytical results (Eqs. (56) and (58)). Dash-dotted lines show analytical results assuming the transition to the Gaussian regime occurs at time t = Da/10 rather than at t = Da (Eq. (59)).

Using Eq. (65), this leads to

$$\frac{\partial |\Omega(t)|}{\partial t} = \frac{1}{g_h(c_m;t)} \frac{\partial \chi(c;t)}{\partial c} \bigg|_{c=c_m},\tag{69}$$

so that, dividing through by  $|\Omega(t)|$  and using Eq. (8),

$$\frac{\partial \ln |\Omega(t)|}{\partial t} = \frac{p(c_m; t)}{2} \frac{\partial \chi(c; t)}{\partial c} \bigg|_{c=c_m}.$$
(70)

<sup>787</sup> Using Eq. (63) for the transport contribution, these results lead to

$$\Delta p_D(c;t) = p(c;t) \left[ \frac{1}{4} \frac{\partial \ln \chi(c;t)}{\partial c} \frac{\partial \chi(c;t)}{\partial c} - \frac{1}{2} \frac{\partial^2 \chi(c;t)}{\partial c^2} - \frac{p(c_m;t)}{2} \frac{\partial \chi(c;t)}{\partial c} \Big|_{c=c_m} \right].$$
(71)

We now turn to the reaction term,  $\Delta p_R$ . Consider a small change dc in the concentrations due to reaction only, over a small time interval dt. The probability p(c; t + dt) dc of finding concentrations in the infinitesimal vicinity dc of c decreases due to reaction at rate r(c) away from c, and increases due to decrease in nearby concentrations towards c. Thus,

$$[p(c;t+dt) - p(c;t)] dc = \sum_{j=1}^{n_s} [r(c+dc)p(c+dc;t) - r(c)p(c;t)] dt.$$
(72)

<sup>792</sup> Expanding the first term on each side in Taylor series, and dividing through by dt and dc, we obtain

$$\frac{\partial p(c;t)}{\partial t} = \frac{\partial r(c)p(c;t)}{\partial c}.$$
(73)

<sup>793</sup> When a fixed minimum concentration detection limit is considered, as discussed above, it is necessary <sup>794</sup> to take the change in volume (in one dimension, length)  $|\Omega(t)|$  due to reactive decay of the minimum



Figure 15: Temporal evolution of (a): mean concentration and (b): concentration variance for  $\beta = 4$ . Colors stand for different Damköhler numbers, markers show the results of resolved numerical simulations of the fully-coupled reaction-diffusion problem (35), solid lines are numerical results based on the weak-coupling model (Eqs. (45)), and dashed lines show approximate asymptotic analytical results (Eqs. (19) and (21), using Eqs. (56) and (58) for the peak concentration). Dash-dotted lines show analytical results assuming the transition to the Gaussian regime occurs at time t = Da/10 rather than at t = Da (Eqs. (19) and (21), using Eqs. (58) and (59) for the peak concentration).

<sup>795</sup> concentration into account. Using the same techniques as before, we obtain for the change in the <sup>796</sup> domain size due to reaction:

$$\frac{\partial \ln |\Omega(t)|}{\partial t} = -p(c_m; t)r(c_m).$$
(74)

<sup>797</sup> Thus, the complete effect of reaction is

$$\Delta p_R(c;t) = \frac{\partial r(c)p(c;t)}{\partial c} + p(c;t)p(c_m;t)r(c_m).$$
(75)

<sup>798</sup> Note that integration of the right-hand side from  $c = c_m$  to  $c = \infty$  yields zero, which ensures the <sup>799</sup> reactive contribution conserves probability for arbitrary r(c). Substituting the effects of transport <sup>800</sup> and reaction, Eqs. (33) and (33), in Eq. (29) leads to the dynamical Eq. (34) for the evolution of the <sup>801</sup> concentration PDF under one-dimensional diffusion and nonlinear decay.

#### <sup>802</sup> C Numerical methods

In this appendix, we provide details on the numerical methods used to integrate the weak-coupling equations (45) and the reaction-diffusion equation (35). Regarding Eqs. (45), which are ordinary differential equations, we implemented a standard fourth-order Runge-Kutta method in the C++ language. This method was chosen for its simplicity of implementation and high accuracy, and also because, as an explicit method, it provides a convenient approach to integrate these nonlinear equations without requiring numerical root-finding methods. We employed a time step  $\Delta t = 10^{-2} \min{\{\text{Da}, 1/\text{Da}\}}$  for the temporal discretization, which we verified led to consistently converged results.

For the fully-coupled reaction-diffusion problem, Eq. (35), we employed the **py-pde** open-source Python package for solving partial differential equations [79]. We used a regular finite difference discretization of a one-dimensional domain of half-width L and second-order centered differences for the spatial derivative approximations. For the time integration, we employed an explicit Forward Euler scheme. We set reflecting boundary conditions at the edges of the computational domain, but we verified that the latter was sufficiently large that no appreciable mass reached the edges, rendering

Table 1: Discretization parameters used in computing solutions of Eq. (35) to determine the mass and concentration peak, mean, and variance. For these computations, we employed a temporal discretization  $\Delta t = a \min\{\text{Da}, 1/\text{Da}\}$  and a spatial discretization  $\Delta x = b\sqrt{\Delta t/\text{Da}}$ . The corresponding values of (a, b) for different reaction orders  $\beta$  and Damköhler numbers Da are given in the table.

$\beta \setminus Da$	$10^{-2}$	$10^{-1}$	$10^{0}$	$10^{1}$	$10^{2}$
1/2	$(10^{-4}, 1)$	$(10^{-4}, 1)$	$(5\cdot 10^{-7},1)$	_	_
2	$(10^{-2}, 1)$	$(10^{-2}, 1)$	$(10^{-2}, 1)$	$(10^{-1}, 2)$	(10, 4)
4	$(10^{-3}, 1)$	$(10^{-2}, 1)$	$(10^{-2}, 1)$	$(10^{-1}, 1)$	$(10^{-1}, 1)$

Table 2: Discretization parameters used in computing solutions of Eq. (35) to determine the concentration PDF and scalar dissipation rate. For these computations, we employed a temporal discretization  $\Delta t = a \min\{\text{Da}, 1/\text{Da}\}$  and a spatial discretization  $\Delta x = \sqrt{\Delta t/\text{Da}}$ . The corresponding values of a for different reaction orders  $\beta$  and Damköhler numbers Da are given in the table.

$\beta \setminus Da$	$10^{-2}$	$10^{-1}$	$10^{0}$	$10^{1}$	$10^{2}$
1/2	$10^{-4}$	$10^{-4}$	$5\cdot 10^{-6}$	_	_
2	$10^{-3}$	_	$10^{-4}$	_	$10^{-2}$
4	$10^{-4}$	_	$10^{-4}$	_	$10^{-1}$

the choice of boundary conditions irrelevant. Since the late-time variance growth is approximately 816 diffusive,  $L = 10\sqrt{t_m}/Da$ , where  $t_m$  is the maximum simulation time, may be used as a simple 817 estimate of necessary domain size. However, because of the lower detection limit  $c_m = 10^{-6}$  used in 818 the computation of the quantities of interest, we found that in practice it was never necessary to use 819 L > 1500 for the simulations conducted here. We note that, for  $\beta < 1$ , where complete depletion of 820 concentrations can happen in finite time, the increase of reaction rates with decreasing concentration 821 values can lead to numerical issues, because very low concentrations can drop below zero within a time 822 step. We avoid this issue by setting negative concentrations to zero before computing reaction rates. 823 We chose the spatial and temporal discretizations so as to ensure good accuracy while maintaining 824 reasonable simulation times. The discretization parameters for different system parameters  $\beta$  and Da 825 are summarized in Table 1 for the mass and concentration peak, mean, and variance calculations, and 826 in Table 2 for the concentration PDF and scalar dissipation rate. 827

The concentration PDF was obtained by counting discretized spatial locations where the concen-828 tration value fell within prescribed bins (see also Section A). In order to accurately resolve both low 829 and high concentrations, we employed  $n_{\ell}$  logarithmically-spaced concentration bins for concentrations 830 between the lower detection limit  $c_m$  and  $2c_M(t)/3$ , where the time-dependent peak value  $c_M(t)$  was 831 determined from the numerical concentration profiles, and  $n_h$  linearly-space bins for the remaining 832 concentrations between  $2c_M(t)/3$  and  $c_M(t)$ . For the lower, intermediate, and higher time examined 833 in each case, we employed  $(n_{\ell}, n_h) = (20, 10), (15, 8), \text{ and } (10, 6),$  respectively. The scalar dissipation 834 rate was calculated according to Eq. (77b) by numerically computing the spatial derivative at each dis-835 cretized spatial location (in the rising limb of the symmetric concentration profile), using second-order 836 central differences. The corresponding concentration values at each spatial location were recorded and 837 used to obtain the scalar dissipation rate as a function of concentration. 838

#### <sup>339</sup> D Problem setup and nondimensionalization

This appendix provides additional details on the nondimensionalization used in Section 4. Denoting
 nondimensionalized quantities by an asterisk, we have

$$C_*(x_*;t_*) = \frac{C(s_0x_*,\tau_R t_*)}{c_0}, \qquad r_*(c_*) = \frac{\tau_R}{c_0}r(c_0c_*) = c_*^\beta, \tag{76}$$

where the characteristic reaction time  $\tau_R = \kappa^{-1} c_0^{1-\beta}$ . The minimum and maximum concentrations  $c_M(t)$  and  $c_m(t)$  are normalized in the same manner. Similarly, the nondimensional spatial variance  $\sigma_*^{2}(t_*) = \sigma^2(\tau_R t_*)/s_0^2$ . The concentration PDF and scalar dissipation are then nondimensionalized accordingly as

$$p_*(c_*;t_*) = c_0 p(c_0 c_*;\tau_R t_*), \tag{77a}$$

$$\chi_*(c_*;t_*) = \frac{\tau_R}{c_0^2} \chi(c_0 c_*;\tau_R t_*) = \frac{1}{2 \operatorname{Da}} \left( \frac{\partial C_*(x_*,t_*)}{\partial x_*} \right)_{x_*=x_c(c)/s_0}^2,$$
(77b)

with the Damköhler number  $Da = \tau_D / \tau_R$ .

Note that, in nondimensional units, the pulse initial condition is given by a unit-width rectangle centered at the origin,

$$C_*(x_*;0) = H\left(1/2 - x_*\right) H\left(1/2 + x_*\right),\tag{78}$$

which implies  $p_*(c_*;0) = \delta(c_*-1)$ . The batch concentration  $c_B(t)$  for the well-mixed problem is nondimensionalized as above, and  $c_{B*}(0) = 1$ .

#### **E** Batch dynamics

Here, we provide some details on the equations governing the well-mixed batch problem discussed in Section 4 and its relation to the concentration PDF. Noting that  $p(c;t) = \delta[c - c_B(t)]$  for the batch problem, where  $c_B(t) = C(x;t)$  is the homogeneous concentration over the domain, and assuming  $c_B(t) > c_m$ , Eq. (34) becomes

$$\frac{\partial p(c;t)}{\partial t} = \frac{\partial r(c)p(c;t)}{\partial c}.$$
(79)

Multiplying through by c and integrating over c (using integration by parts on the right hand side), we recover the standard well-mixed rate law for the batch concentration as a function of time,

$$\frac{dc_B(t)}{dt} = -r[c_B(t)]. \tag{80}$$

Substituting Eq. (76) for the rate yields Eq. (37).

Once  $c_B(t)$  drops below  $c_m$ , at some time  $t_m$ , the domain  $\Omega(t)$  where concentrations are above this detection limit becomes empty, and the concentration PDF becomes ill-defined. By convention, we can set  $c_B(t > t_m) = 0$  and  $p(c; t > t_m) = \delta(c)$ , which conveys the meaning that concentrations are zero everywhere (below the detection limit).

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