

# Modeling and numerical study on a collection of strongly coupled chemotaxis-fluid systems

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In this work, we study chemotaxis-fluid systems governed by the following general form

$$\begin{aligned} n_t + \mathbf{u} \cdot \nabla n &= \Delta n - \nabla \cdot (n \mathcal{S}(x, n, c) \nabla c) + f(n), & (x, t) \in \Omega \times (0, T), \\ c_t + \mathbf{u} \cdot \nabla c &= \Delta c - h(n, c), & (x, t) \in \Omega \times (0, T), \\ \mathbf{u}_t + \kappa(\mathbf{u} \cdot \nabla) \mathbf{u} &= \Delta \mathbf{u} + \nabla P + n \nabla \Phi + \mathbf{f}_u(x, t), & (x, t) \in \Omega \times (0, T), \\ \nabla \cdot \mathbf{u} &= 0, & (x, t) \in \Omega \times (0, T). \end{aligned}$$

which models the interaction between motile biological organisms and their chemical signaling in fluid environment. It is motivated by recent analytical studies on chemotaxis-Stokes/Navier-Stokes systems, particularly those involving singular sensitivities, logistic-type source terms, chemical signaling, and general fluid coupling.

We aim to numerically investigate various structurally distinct forms of this system by incorporating logistic source functions  $f(n)$  and sensitivity functions  $\mathcal{S}(x, n, c)$  in the  $n$ -equation, both consumption and production signaling mechanisms  $h(n, c)$  in the  $c$ -equation, and modeling the fluid dynamics through either the Stokes or Navier-Stokes in the  $u$ -equation depending on the variation considered. To capture the complexity of the underlying dynamics, we develop a high-resolution finite element method that incorporates stabilization techniques suited for resolving steep gradients and nonlinear interactions in convection- and chemotaxis-dominated regimes. Numerical simulations are carried out in both two- and three-dimensional domains, supporting analytical and numerical findings reported in the literature.

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