

Numerical Modeling of Flocking Dynamics with Topological Interactions

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Abstract. In this paper, we propose a numerical investigation of topological interactions in flocking dynamics. Starting from a microscopic description of the phenomena, mesoscopic and macroscopic models have been previously derived under specific assumptions. We explore the role of topological interactions by describing the convergence speed to consensus in both microscopic and macroscopic dynamics, considering different forms of topological interactions. Additionally, we compare mesoscopic and macroscopic dynamics for monokinetic and non-monokinetic initial data. Finally, we illustrate with some simulations in one- and two-dimensional domains the sensitive dependence of solutions on initial conditions, including the case where the system exhibits two solutions starting with the same initial data.

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1 Introduction

Collective dynamics have been widely investigated over the last years in different fields of research, due to the great variety of living and non-living systems exhibiting such complex behaviors. From flocks of birds, schools of fish to bacterial colonies and robotic swarms, coordinated movements of individuals interacting within a group give rise to the so called *emergent behaviors*. Understanding the underlying principles of self-organized behaviors has emerged as a fundamental challenge in applied mathematics,

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aiming at providing the framework to describe, predict, and analyze the basic mechanisms determining the observed collective dynamics. A crucial aspect of mathematical modeling is selecting the appropriate scale for the description for the phenomena under investigation. In traditional microscopic, agent-based approaches, the dynamics of each individual in the group are tracked. While this approach allows to highly represent the interactions between agents, it also leads to increasing computational cost as the number of agents grows. For large flocks, mesoscopic and macroscopic models are generally more efficient. When the number of agents increases, mesoscopic and macroscopic models can be considered as efficient alternatives to pure microscopic ones. This is mainly due to the fact that microscopic models, focusing on the dynamics of each individual, become computationally demanding for large groups. Conversely, mesoscopic and macroscopic approaches allow to describe collective behaviors using aggregated variables such as probabilistic distribution of agents, macroscopic density and average velocity field (see [25] for an updated reference paper on the derivation from microscopic to macroscopic models). Several mathematical models have been proposed in the literature to model *flocking* behaviors, see [33] for an introduction to the field. Typically displayed by flocks of birds, flocking is used to define the coordinated motion of agents reaching the same velocity when moving towards the same direction. The majority of the models, starting from the seminal papers [12, 13], assumes interactions involving metric quantities (*metric interactions*), where individuals adjust their movement in response to the other members of the group, and the interaction depends on the relative Euclidean distance. However, increasing empirical evidence suggests that in many biological systems, interactions are *topological*, meaning that individuals interact with a fixed number of neighbors, regardless of their absolute distances [3, 6]. As a consequence, several mathematical models adopting this novel paradigm have appeared, ranging from microscopic to macroscopic ones, see e.g. [2, 7, 10, 17, 18, 28]. This physically non-metric nature of interactions plays a crucial role for the stability, robustness and efficiency of the observed motion, affecting not only the local organization of individuals but also the emergent macroscopic patterns.

1.1 Topological interactions models: From microscopic to macroscopic scale

In this paper we focus on the topological model of the Cucker-Smale type firstly proposed in [17]. Starting from the microscopic scale, the model reads

$$\begin{cases} \dot{X}_i(t) = V_i(t), \\ \dot{V}_i(t) = \frac{1}{N} \sum_{j=1}^N p_{i,j} (V_j(t) - V_i(t)), \end{cases} \quad i \in \{1, \dots, N\}. \quad (1.1)$$

Here $X_i \in \mathbb{R}^d$, $V_i \in \mathbb{R}^d$ denote positions and velocities of N indistinguishable interacting agents in a general $d \geq 1$ dimensional space. A key factor of the model is the so-called communication weights $p_{i,j}$, modeling the kind of considered interactions.

In the original version of the Cucker-Smale model [12,13], the interaction $p_{i,j}$ between two agents $i, j \in \{1, \dots, N\}$ depends on the metric distance between them, i.e. $p_{i,j} = p_{i,j}(|X_i - X_j|)$. In the topological framework, $p_{i,j}$ are chosen as

$$p_{i,j} := K\left(M_{i,j}^N\right), \tag{1.2}$$

where $K: [0,1] \rightarrow \mathbb{R}^+$ is a non increasing function. The quantity $M_{i,j}^N$ is defined as

$$M_{i,j}^N := \frac{1}{N} \sum_{h=1}^N \chi_{\bar{B}(X_i, |X_i - X_j|)}(X_h), \tag{1.3}$$

where $\chi_{\bar{B}(X_i, |X_i - X_j|)}$ denotes the characteristic function of the closed ball centered in X_i with radius $|X_i - X_j|$. The quantity $M_{i,j}^N$ hence counts the number of agents inside the ball $\bar{B}(X_i, |X_i - X_j|)$, and it is known as *topological rank* between two agents $i, j \in \{1, \dots, N\}$. Note that non-decreasing property of function K reflects the fact that the farther apart two agents are, also in a topological sense, the less influence one has on the other. In the next section, specific choices of non-increasing function K are performed, see the following (3.1a), (3.1b).

Starting from the microscopic model (1.1), the kinetic and hydrodynamic equations have been first formally derived in [17]. At kinetic level, the Vlasov-type equation related to (1.1) reads as

$$\begin{aligned} & \partial_t f_t(x, v) + v \cdot \nabla_x f_t(x, v) \\ &= \nabla_v \cdot \left(f_t(x, v) \int K(M[v_t^0](x, |x - y|)) (v - w) df_t(y, w) \right), \end{aligned} \tag{1.4}$$

where f_t denotes a distribution function of agents at time t depending on $(x, v) \in \mathbb{R}^d \times \mathbb{R}^d$. We have introduced the following notation for the zero-th and first order moments of $f_t = f_t(x, v)$, respectively

$$v_t^0(x) := \int f_t(x, v) dv, \tag{1.5}$$

$$v_t^1(x) := \int v f_t(x, v) dv \tag{1.6}$$

for any $t \geq 0$ and

$$M[v_t^0](x, |x - y|) := \int_{\bar{B}(x, |x - y|)} dv_t^0 \tag{1.7}$$

the mass inside the ball $\bar{B}(x, |x - y|)$.

At the macroscopic scale, the following pressureless Euler-type system is formally obtained in [17]:

$$\begin{cases} \partial_t \rho_t(x) + \nabla_x \cdot (\rho_t(x) u_t(x)) = 0, \\ \partial_t u_t(x) + (u_t(x) \cdot \nabla_x) u_t(x) = \int_{\mathbb{R}^d} K\left(M[\rho_t](x, |x - y|)\right) (u_t(y) - u_t(x)) \rho_t(y) dy, \end{cases} \tag{1.8}$$

under a monokinetic assumption on the kinetic distribution function, namely

$$f_t(x, v) = \rho_t(x) \delta(v - u_t(x)). \quad (1.9)$$

Here $(\rho_t(x), u_t(x)) : [0, T] \times \mathbb{R}^d \rightarrow \mathbb{R} \times \mathbb{R}^d$, $T > 0$, denote density and velocity field, respectively.

Remark 1.1. The quantities v_t^0 and v_t^1 denote mass and momentum of the distribution function f_t at the kinetic level. Under a monokinetic assumption, the kinetic moments and the macroscopic moments coincide as

$$v_t^0(x) = \rho_t(x), \quad v_t^1(x) = \rho_t(x) u_t(x). \quad (1.10)$$

We remark that the derivation in [17] is only formal. Indeed the rigorous proof of the mean-field limit and propagation of chaos results for the topological kinetic equation (1.4), under the assumption of Lipschitz continuity for the interaction function K in (1.2), has been presented in [4, 5]. The results on propagation of chaos have been further extended to the case of general $K : [0, 1] \rightarrow [0, 1]$ non-increasing functions in [24]. From the analytical point of view, local and global well-posedness results for the multi-dimensional case of the Euler system can be found in [21, 27], and rigorous results have been also obtained for stochastic models of topological type [14, 15].

From a numerical perspective, the literature is still quite poor and several issues need to be tackled. In this work we present a numerical investigation of the topological models in (1.1)-(1.8) at different scales. In particular, we focus on different choices of function K in (1.2), which rules topological interactions, studying how this affects the convergence to consensus. Moreover, we investigate the similarities and differences between the mesoscopic and macroscopic dynamics both within and outside the monokinetic regime. Finally, we illustrate with examples in one- and two-dimensional scenarios the sensitive dependence of the solutions on the initial data.

The paper is organized as follows: in Section 2 we detail the numerical schemes implemented to approximate the solutions at the kinetic and macroscopic level. The main results of our study are reported in Section 3, where different scenarios, at different scales, in 1D and 2D setting are discussed, including the case with non uniqueness of solutions. Conclusions and future perspective on the topic conclude the paper in Section 4.

2 Numerical approximation

In this section we describe the numerical schemes implemented to approximate the solutions of the models at different scales. The microscopic model is numerically approximated using classical implicit Euler scheme both in the one and two dimensional case, avoiding restrictions on the time step due to explicit integration. The kinetic and the macroscopic models are approximated using finite volume schemes, as detailed in the following. The time discretization on $[0, T]$ is performed introducing a discretization time step Δt , which is chosen according to a CFL stability condition for the kinetic and the macroscopic equation.

2.1 Numerical scheme for the kinetic model

We detail the construction of the scheme in the one dimensional case, using a classical upwind scheme both in the spatial and in the velocity direction. Let us consider uniform spatial cells $\mathcal{C}_i = (x_{i-1/2}, x_{i+1/2})$, where x_i are equally distributed points and $x_{i+1/2} = x_i + \Delta x/2$. Analogously, we define uniform velocity cells $\mathcal{V}_k = (v_{k-1/2}, v_{k+1/2})$, where v_k are equally distributed points and $v_{k+1/2} = v_k + \Delta v/2$. Let us consider the kinetic equation (1.4) in the one-dimensional case rewritten as

$$\partial_t f + \partial_x F(f) + \partial_v G(f) = 0, \tag{2.1}$$

where

$$F(f) = v f, \quad G(f) = -f \int K(M[v_t^0](x, |x-y|)) (v-w) df(y,w). \tag{2.2}$$

The fully discrete upwind scheme reads as

$$f_{i,k}^{n+1} = f_{i,k}^n - \frac{\Delta t}{\Delta x} (F_{i+1/2,k}^n - F_{i-1/2,k}^n) - \frac{\Delta t}{\Delta v} (G_{i,k+1/2}^n - G_{i,k-1/2}^n), \tag{2.3}$$

where $F_{i+1/2,k}^n$ and $G_{i,k+1/2}^n$ are first-order upwind numerical fluxes. In particular, we can explicitly define

$$F_{i+1/2,k}^n = v_k f_{i+1/2,k}^n = \begin{cases} v_k f_{i,k}^n, & \text{if } v_k > 0, \\ v_k f_{i+1,k}^n, & \text{if } v_k \leq 0, \end{cases} \tag{2.4}$$

and by introducing $\xi_{i,k+1/2}$ as a quadrature approximation of the integral term in (2.2) at $(x_i, v_{k+1/2})$, we can also define

$$G_{i,k+1/2}^n = \xi_{i,k+1/2} f_{i,k+1/2}^n = \begin{cases} \xi_{i,k+1/2} f_{i,k}^n, & \text{if } \xi_{i,k+1/2} > 0, \\ \xi_{i,k+1/2} f_{i,k+1}^n, & \text{if } \xi_{i,k+1/2} \leq 0. \end{cases} \tag{2.5}$$

The choice of a first order scheme relies in its positivity-preserving property, which is fundamental for a probability distribution function. We briefly describe the derivation of the CFL condition to guarantee the positivity-preserving of the scheme, inspired by [31].

2.1.1 Positivity preserving property

It is possible to rewrite (2.3) as

$$f_{i,k}^{n+1} = \frac{1}{2} \left(f_{i,k}^n - \frac{2\Delta t}{\Delta x} v_k f_{i+1/2,k}^n - \frac{2\Delta t}{\Delta v} \xi_{i,k+1/2} f_{i,k+1/2}^n \right) + \frac{1}{2} \left(f_{i,k}^n + \frac{2\Delta t}{\Delta x} v_k f_{i-1/2,k}^n + \frac{2\Delta t}{\Delta v} \xi_{i,k-1/2} f_{i,k-1/2}^n \right). \tag{2.6}$$

Then, if $\xi_{i,k+1/2} \leq 0$, we have

$$\begin{aligned} & f_{i,k}^n - \frac{2\Delta t}{\Delta x} v_k f_{i+1/2,k}^n - \frac{2\Delta t}{\Delta v} \xi_{i,k+1/2} f_{i,k+1/2}^n \\ &= f_{i,k}^n - \frac{2\Delta t}{\Delta x} v_k f_{i+1,k}^n - \frac{2\Delta t}{\Delta v} \xi_{i,k+1/2} f_{i,k+1}^n > 0, \quad \text{if } v_k \leq 0, \end{aligned} \tag{2.7}$$

and

$$\begin{aligned}
 & f_{i,k}^n - \frac{2\Delta t}{\Delta x} v_k f_{i+1/2,k}^n - \frac{2\Delta t}{\Delta v} \zeta_{i,k+1/2} f_{i,k+1/2}^n \\
 &= \left(1 - \frac{2\Delta t}{\Delta x} v_k \right) f_{i,k}^n - \frac{2\Delta t}{\Delta v} \zeta_{i,k+1/2} f_{i,k+1}^n > 0, \quad \text{if } v_k > 0,
 \end{aligned} \tag{2.8}$$

under the CFL condition $\Delta t < \frac{\Delta x}{2 \max_k |v_k|}$. In the same spirit, if $\zeta_{i,k+1/2} > 0$

$$\begin{aligned}
 & f_{i,k}^n - \frac{2\Delta t}{\Delta x} v_k f_{i+1/2,k}^n - \frac{2\Delta t}{\Delta v} \zeta_{i,k+1/2} f_{i,k+1/2}^n \\
 &= \left(1 - \frac{2\Delta t}{\Delta v} \zeta_{i,k+1/2} \right) f_{i,k}^n - \frac{2\Delta t}{\Delta x} v_k f_{i,k+1}^n > 0, \quad \text{if } v_k \leq 0,
 \end{aligned} \tag{2.9}$$

under the CFL condition $\Delta t < \frac{\Delta v}{2 \max_{i,k} |\zeta_{i,k+1/2}|}$. Again we have

$$\begin{aligned}
 & f_{i,k}^n - \frac{2\Delta t}{\Delta x} v_k f_{i+1/2,k}^n - \frac{2\Delta t}{\Delta v} \zeta_{i,k+1/2} f_{i,k+1/2}^n \\
 &= \left(1 - \frac{2\Delta t}{\Delta x} v_k - \frac{2\Delta t}{\Delta v} \zeta_{i,k+1/2} \right) f_{i,k}^n > 0, \quad \text{if } v_k > 0,
 \end{aligned} \tag{2.10}$$

under the CFL condition

$$\Delta t < \frac{\Delta v \Delta x}{2 \max_{i,k} (|\Delta v v_k + \Delta x \zeta_{i,k+1/2}|)}.$$

In the same way, the second term in (2.6) is positive under the same condition. We conclude that the first order scheme under the CFL condition

$$\Delta t < \frac{\Delta v \Delta x}{2 \max_{i,k} (|\Delta v v_k + \Delta x \zeta_{i,k+1/2}|)} \tag{2.11}$$

is positivity preserving, namely $f_{i,k}^{n+1} > 0$ if $f_{i,k}^n > 0$ for all i and k .

2.2 Numerical scheme for the macroscopic model

Let us describe the numerical scheme for the approximation of the macroscopic system (1.8) in the one- and two-dimensional cases. We present the details of the approximation in the two-dimensional case, since the one-dimensional case can be easily derived. In the two-dimensional case denoting $u = (u_1, u_2)$, $\nabla = (\partial_x, \partial_y)$ and omitting the subscript t for simplicity of notation, system (1.8) reads as

$$\begin{cases}
 \partial_t \rho + \partial_x(\rho u_1) + \partial_y(\rho u_2) = 0, \\
 \partial_t u_1 + u_1 \partial_x u_1 + u_2 \partial_y u_1 = \int_{\mathbb{R}^2} K(M[\rho](\mathbf{z}, |\mathbf{z} - \mathbf{z}'|)) (u_1(\mathbf{z}') - u_1(\mathbf{z})) \rho(\mathbf{z}') d\mathbf{z}', \\
 \partial_t u_2 + u_1 \partial_x u_2 + u_2 \partial_y u_2 = \int_{\mathbb{R}^2} K(M[\rho](\mathbf{z}, |\mathbf{z} - \mathbf{z}'|)) (u_2(\mathbf{z}') - u_2(\mathbf{z})) \rho(\mathbf{z}') d\mathbf{z}'
 \end{cases} \tag{2.12}$$

where $\mathbf{z} = (x, y) \in \mathbb{R}^2$. The numerical scheme for the approximation of the solution combines the finite volume second-order central scheme proposed by Kurganov and Tadmor [20]. A variant of this scheme has been implemented in [1] for the numerical approximation of a pressureless Euler alignment system without topological interactions. We here extend the approach proposed in [20] with the global flux technique [9,16,19] to treat non-conservative terms. In this spirit, we rewrite system (2.12) in a quasi-conservative form as

$$\partial_t w + \partial_x \tilde{A}(w) + \partial_y \tilde{B}(w) = S(w), \tag{2.13}$$

where

$$w = \begin{pmatrix} \rho \\ u_1 \\ u_2 \end{pmatrix}, \quad S(w) = \begin{pmatrix} 0 \\ \int_{\mathbb{R}^2} K(M[\rho](\mathbf{z}, |\mathbf{z} - \mathbf{z}'|)) (u_1(\mathbf{z}') - u_1(\mathbf{z})) \rho(\mathbf{z}') d\mathbf{z}' \\ \int_{\mathbb{R}^2} K(M[\rho](\mathbf{z}, |\mathbf{z} - \mathbf{z}'|)) (u_2(\mathbf{z}') - u_2(\mathbf{z})) \rho(\mathbf{z}') d\mathbf{z}' \end{pmatrix}. \tag{2.14}$$

The fluxes are defined by

$$\tilde{A}(w) = \begin{pmatrix} \rho u_1 \\ \frac{1}{2} u_1^2 \\ u_1 u_2 + \int_{\hat{x}} N_1(\zeta, y) d\zeta \end{pmatrix}, \quad \tilde{B}(w) = \begin{pmatrix} \rho u_2 \\ u_1 u_2 + \int_{\hat{y}} N_2(x, \eta) d\eta \\ \frac{1}{2} u_2^2 \end{pmatrix} \tag{2.15}$$

with

$$N_1(x, y) = -u_2(x, y) \partial_x u_1(x, y), \quad N_2(x, y) = -u_1(x, y) \partial_y u_2(x, y) \tag{2.16}$$

and \hat{x}, \hat{y} arbitrary values.

Let us consider uniform discretization in space, with cells $C_{i,j} = (x_{i-1/2}, x_{i+1/2}) \times (y_{j-1/2}, y_{j+1/2})$ of volume $\Delta x \times \Delta y$, whose centers are given by (x_i, y_j) . The semi-discrete version of our scheme in the two-dimensional case reads as

$$\frac{d}{dt} \bar{w}_{i,j} = -\frac{1}{\Delta x} \left(\mathcal{F}_{i+\frac{1}{2},j}^x(t) - \mathcal{F}_{i-\frac{1}{2},j}^x(t) \right) - \frac{1}{\Delta y} \left(\mathcal{F}_{i,j+\frac{1}{2}}^y(t) - \mathcal{F}_{i,j-\frac{1}{2}}^y(t) \right) + S_{i,j}(t), \tag{2.17}$$

where $\bar{w}_{i,j}$ represents the cell average of w in $C_{i,j}$ and \mathcal{F}^x (or \mathcal{F}^y) is the numerical flux corresponding to \tilde{A} (or \tilde{B} respectively). To obtain a second-order scheme we introduce a piecewise linear reconstruction for w of the form

$$w_{i,j}(x, y) = \bar{w}_{i,j} + (w_x)_{i,j} (x - x_i) + (w_y)_{i,j} (y - y_j). \tag{2.18}$$

The derivatives w_x and w_y are approximated by using a minmod limiter [30] to avoid oscillations in the reconstruction procedure, namely

$$(w_x)_{i,j} = \text{minmod} \left(\theta \frac{\bar{w}_{i+1,j} - \bar{w}_{i,j}}{\Delta x}, \theta \frac{\bar{w}_{i+1,j} - \bar{w}_{i-1,j}}{2\Delta x}, \theta \frac{\bar{w}_{i,j} - \bar{w}_{i-1,j}}{\Delta x} \right),$$

$$(w_y)_{i,j} = \text{minmod} \left(\theta \frac{\bar{w}_{i,j+1} - \bar{w}_{i,j}}{\Delta y}, \theta \frac{\bar{w}_{i,j+1} - \bar{w}_{i,j-1}}{2\Delta y}, \theta \frac{\bar{w}_{i,j} - \bar{w}_{i,j-1}}{\Delta y} \right),$$

where $\theta \in [1, 2]$. We can now easily define the reconstructed values at the interfaces $w_{i,j \pm \frac{1}{2}}$ and $w_{i \pm \frac{1}{2},j}$ through the expression (2.18) and the characteristic speeds at the interfaces. We define the following quantities

$$w_{i,j}^N = w_{i,j}(x_i, y_{j+1/2}), \quad w_{i,j}^S := w_{i,j}(x_i, y_{j-1/2}), \quad w_{i,j}^E := w_{i,j}(x_{i+1/2}, y_j), \quad w_{i,j}^W := w_{i,j}(x_{i-1/2}, y_j).$$

In particular $a_{i+1/2,j}^\pm$ will denote the largest (lowest) characteristic speed of \tilde{A} along the interface $i+1/2$ and $b_{i,j+1/2}^\pm$ the largest (lowest) characteristic speed of \tilde{B} along the interface $j+1/2$.

The numerical fluxes read as

$$\mathcal{F}_{i+\frac{1}{2},j}^x = \frac{a_{i+\frac{1}{2},j}^+ \tilde{A}(w_{i,j}^E) - a_{i+\frac{1}{2},j}^- \tilde{A}(w_{i+1,j}^W)}{a_{i+\frac{1}{2},j}^+ - a_{i+\frac{1}{2},j}^-} + \frac{a_{i+\frac{1}{2},j}^+ a_{i+\frac{1}{2},j}^-}{a_{i+\frac{1}{2},j}^+ - a_{i+\frac{1}{2},j}^-} (w_{i+1,j}^W - w_{i,j}^E)$$

and

$$\mathcal{F}_{i,j+\frac{1}{2}}^y = \frac{b_{i,j+\frac{1}{2}}^+ \tilde{B}(w_{i,j}^N) - b_{i,j+\frac{1}{2}}^- \tilde{B}(w_{i,j+1}^S)}{b_{i,j+\frac{1}{2}}^+ - b_{i,j+\frac{1}{2}}^-} + \frac{b_{i,j+\frac{1}{2}}^+ b_{i,j+\frac{1}{2}}^-}{b_{i,j+\frac{1}{2}}^+ - b_{i,j+\frac{1}{2}}^-} (w_{i,j+1}^S - w_{i,j}^N).$$

The integral term $S_{i,j}$ is approximated by using a nested quadrature rule (the first one for the computation of the quantity M and the second one for the full integral).

Specifically, the approximation of the second component of the source term in (2.13), here denoted as $S_{i,j}^{(2)}$, is computed as

$$S_{i,j}^{(2)} := \rho_{i,j} \sum_{\mathbf{z}'_{p,q} \in \mathbb{Z}} \omega_{\mathbf{z}'} K(M[\rho_\Delta](\mathbf{z}_{i,j}, \|\mathbf{z}_{i,j} - \mathbf{z}'_{p,q}\|)) (u_{p,q} - u_{i,j}) \rho_{p,q}, \tag{2.19}$$

where $\mathbf{z}_{i,j} := (x_i, y_j)$, $u_{i,j} \approx u_1(x_i, y_j)$ and the quantity M is approximated by

$$M[\rho_\Delta](\mathbf{z}_{i,j}, \|\mathbf{z}_{i,j} - \mathbf{z}'\|) \approx \sum_{\mathbf{h}_{l,m} \in \mathbb{Z}} \tilde{\omega}_{\mathbf{h}} \rho_{l,m} \mathbb{1}_{\{\|\mathbf{h}_{l,m} - \mathbf{z}_{i,j}\| \leq \|\mathbf{z}_{i,j} - \mathbf{z}'\|\}}. \tag{2.20}$$

The quantities $\omega_{\mathbf{z}'}$ and $\tilde{\omega}_{\mathbf{h}}$ are the corresponding quadrature weights. The third component $S_{i,j}^{(3)}$ is computed analogously, by defining $u_{i,j} \approx u_2(x_i, y_j)$.

The semi-discrete scheme obtained in (2.17) is discretized in time using classical explicit Euler method.

3 Numerical tests

We present several numerical tests simulating microscopic, kinetic and macroscopic dynamics. We investigate the role of topological interactions on the overall dynamics, comparing the results obtained at different scales. Further comparison will involve kinetic and macroscopic dynamics. We conclude our analysis exploiting sensitive dependence on initial conditions in a 2D setting. All the tests performed have been run a laptop equipped with an Intel Core i7-1060NG7 processor and 16 GB RAM.

3.1 Effects of topological interactions

We investigate the role of topological interactions on the overall dynamics, comparing the results obtained for different choices of function K in (1.2). We consider a reference microscopic scenario in 1D, with $N=100$ agents. At initial time, a smaller group of $N_1=20$ agents is randomly located in $[-3,-1]$ with initial velocity randomly assigned in $[-2,0]$, whereas the remaining $N_2=80$ agents are located in $[1,3]$ with initial random velocity in $[0,2]$. Fig. 1 shows the initial condition plotted in the position-velocity space.

In the following Test1–Test3 we run system (1.1) over the time interval $[0,500]$, with time discretization step $\Delta t = 0.02$.

Test 1: Study of consensus velocities for different values of \bar{M} . We here consider a non Lipschitz function, choosing $K = \mathbb{1}_{[0, \bar{M}/N]}$. Varying the value of $\bar{M} > 0$, we show the effect of variations in the number of neighbors considered for interactions. Fig. 2 shows the velocity evolution of the agents for four different values of \bar{M} , which are representative for the simulated scenario. Indeed, choosing $\bar{M} < N_1$ implies that agents in the smaller group do not interact with the remaining ones, and different consensus velocities are reached within the groups (Fig. 2(a)). In order to let the two groups communicating, we consider $\bar{M} > N_1$ (Fig. 2(b)). We observe that the number of consensus velocities decreases as \bar{M} increases, since interacting with a higher number of neighbors reduces the formation of cluster dynamics. Assuming $\bar{M} = N$ we recover an all-to-all type of interaction. Comparing the results obtained for $\bar{M}=60$, we observe that, keeping N fixed, the value of \bar{M} influences the time at which flocking is reached (see Fig. 2(c)-(d)). This is due to the fact, keeping N fixed, the process of achieving consensus is faster increasing the number of neighbors each agent considers for interactions.

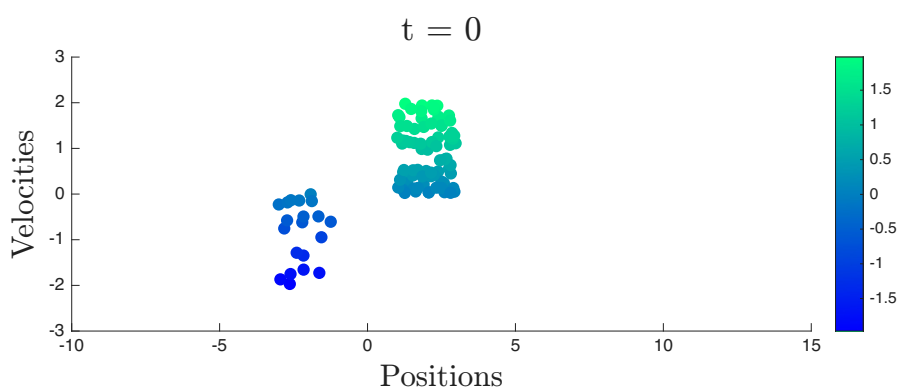


Figure 1: Test 1: initial condition $N = 100$ agents. Each position is marked with colors based on its initial velocity: the velocities of the agents located in $[-3,-1]$ are randomly chosen in $[-2,0]$, whereas the velocities of the agents located in $[1,3]$ are randomly chosen in $[0,2]$.

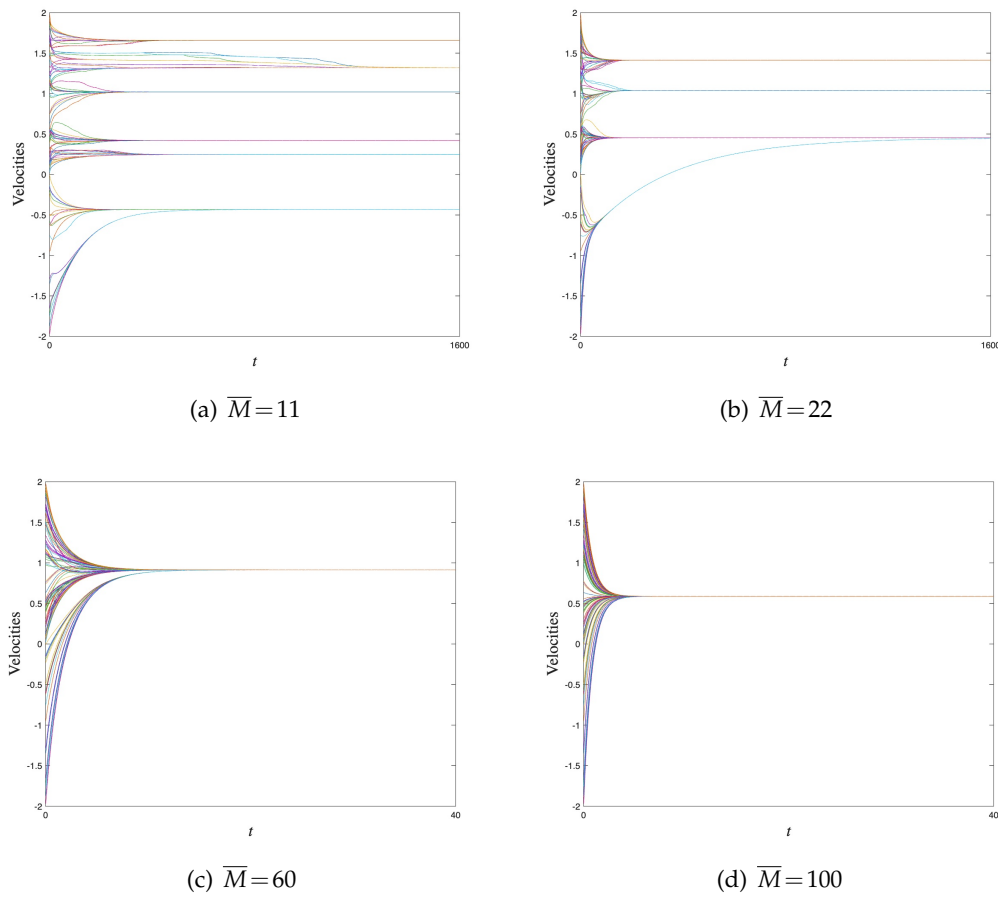


Figure 2: Test 1: plot of $V_i(t)$, $i=1, \dots, N$, solution to (1.1) with $K = \mathbb{1}_{[0, \bar{M}/N]}$ for different values of \bar{M} .

Test 2: Study of consensus velocities for different choices of K . In Test 2 we compare the velocity of convergence to a flocking state for two different $K : [0,1] \rightarrow [0,1]$, both Lipschitz and decreasing functions:

$$K_1(x) = 1 - x, \tag{3.1a}$$

$$K_2(x) = (1 - x)^2. \tag{3.1b}$$

Fig. 3(a)-(b) shows the results obtained for K_1 and K_2 , respectively. Function K_2 decays to zero faster with respect to K_1 , reflecting in a weaker interactions among agents. This also implies a slower convergence to the flocking state. The obtained results can be better visualized in Fig. 3(c), showing the evolution in time of the microscopic mean velocity \bar{V} , for the two different choices of K .

Moreover, we compare the results obtained at the microscopic scale with the macro-

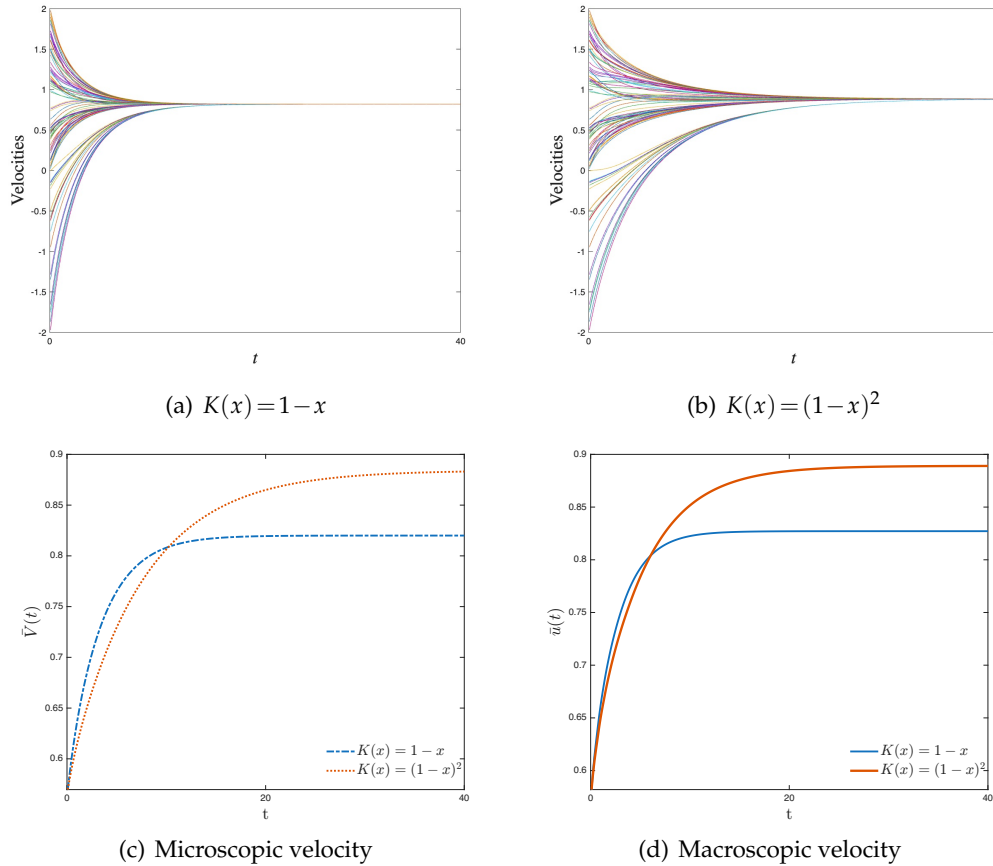


Figure 3: Test 2: plot of the microscopic velocity $V_i(t)$, $i = 1, \dots, N$, solution to (1.1) with a) K linear b) K convex function and the comparison between c) $\bar{V}(t)$ and d) $\bar{u}(t)$, for different K functions.

scopic one. In order to link microscopic initial data to macroscopic ones, different approaches can be adopted, see [29, 32] for a complete introduction to seminal methods in the field. Our choice allows us to introduce the initial data for the kinetic scale, that will be used in Test 5, creating a full path, from microscopic to macroscopic scale.

Starting from the initial data for the microscopic scale, we build initial data ρ_0, u_0 to run system (1.8). Denoting $X^0 \in \mathbb{R}^N$, $V^0 \in \mathbb{R}^N$ initial positions and velocities of the agents, we define an initial distribution probability considering the sum of N Gaussian functions

$$f^0(x, v) = \sum_{i=1}^N \frac{1}{2\pi\sigma_x\sigma_v} e^{-\frac{(x-X_i^0)^2}{2\sigma_x^2} - \frac{(v-V_i^0)^2}{2\sigma_v^2}} \tag{3.2}$$

with $\sigma_x = 0.1$, $\sigma_v = 0.1$. Here $X_i^0, V_i^0 \in \mathbb{R}$ denotes the position and the velocity of each agent i , respectively. Recalling the definition of v_t^0 and v_t^1 in (1.5)-(1.6), initial data at the

macroscopic scale are given as

$$\begin{cases} \rho_0 = v_0^0, \\ Q_0 = v_0^1, \end{cases} \tag{3.3}$$

where $Q_t := \rho_t u_t$ denotes the momentum, for any $t \geq 0$.

We run (1.8) for both choices of the function K in (3.1a)-(3.1b), aiming at comparing the influence of the performed choice on the convergence time to the consensus state also at the macroscopic scale. We set $T = 40$, $\Delta x = 0.4$, $\Delta t = 0.02$. In particular, we evaluate the macroscopic average velocity $\bar{u}(t)$, computed as the integral of the velocity field over the spatial domain. Fig. 3(d) shows that also at this scale the evolution depends on the choice of K , with faster convergence to the consensus equilibrium corresponding to the linear one.

Test 3: Evolution of the microscopic model for a linear K . We conclude our analysis on the microscopic scale in 1D showing some screenshots of the dynamics of agents assuming $K = K_1$ in (3.1a), see Fig. 4. We remark that initial condition in Test 1 and its evolution in Test 3 represents the reference microscopic dynamics for the comparison with different scales performed in Section 3.2.

3.2 Comparison of the microscopic, kinetic and macroscopic models

Starting from the results obtained at the microscopic scale, we here focus on the kinetic and macroscopic scales. We recall that the derivation of the hydrodynamic limit (1.8) is obtained under a monokinetic assumption on the initial data of (1.4). Indeed, far from the monokinetic case, no rigorous proofs of convergence have been established.

We approximate the solution to (1.4) in the space-velocity domain $[-15, 20] \times [-5, 5]$. The simulation runs over the time interval $[0, T] = [0, 3]$ with Δt chosen according to the CFL condition (2.11).

Test 4: Monokinetic initial condition. We recall that analytical results obtained in [5] ensure a complete correspondence between Vlasov moments and Euler solutions, in case of monokinetic initial data, namely

$$f^0 = \rho^0(x) \delta(v - u^0(x)). \tag{3.4}$$

This is a strong *ansatz*, since the solution is completely described by its hydrodynamic fields and no further closure assumptions are required. In Test 4 we simulate a monokinetic-like initial data scenario. We approximate the initial distribution choosing a Gaussian function with a small value of σ_v , namely:

$$f^0(x, v) = \frac{1}{2\pi\sigma_x\sigma_v} e^{-\frac{(x-x_0)^2}{2\sigma_x^2} - \frac{(v-v_0)^2}{2\sigma_v^2}} \tag{3.5}$$

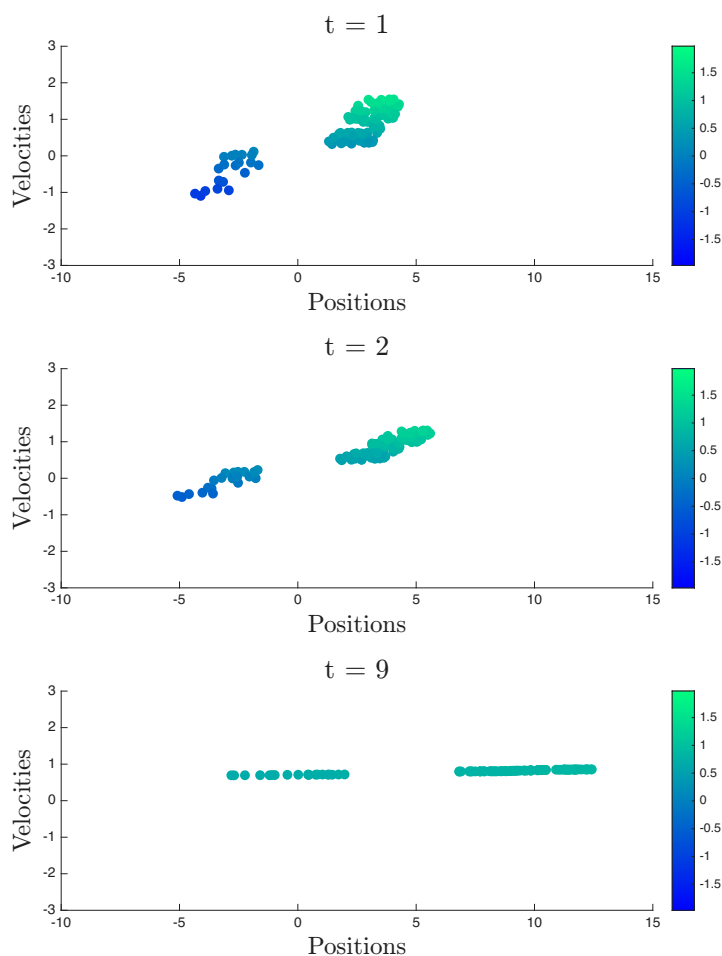


Figure 4: Test 3: screenshots of numerical simulation of (1.1) with $K=K_1$ as in (3.1a).

with $x_0 = -2$, $v_0 = 1.5$, $\sigma_x = 2$, $\sigma_v = \sqrt{0.001}$. The solution is approximated running the scheme detailed in Section 2.1 setting $\Delta x = 0.05$, $\Delta v = 0.005$. Following the approach in [23], we compare the moments of the solution to Vlasov with the solution of the Euler system. We run the macroscopic system (1.8) starting with

$$\begin{cases} \rho_0 = v_0^0, \\ Q_0 = v_0^1, \end{cases} \quad (3.6)$$

with v_0^0, v_0^1 defined as in (1.5)-(1.6).

Fig. 5 shows the comparison between the scales at two different time instants. As expected, we observe a good agreement between zeroth and first order moments of the solution to (1.4), and density and momentum solutions to (1.8). The small discrepancy between the profiles is due to the approximation of monokineticity assumption.

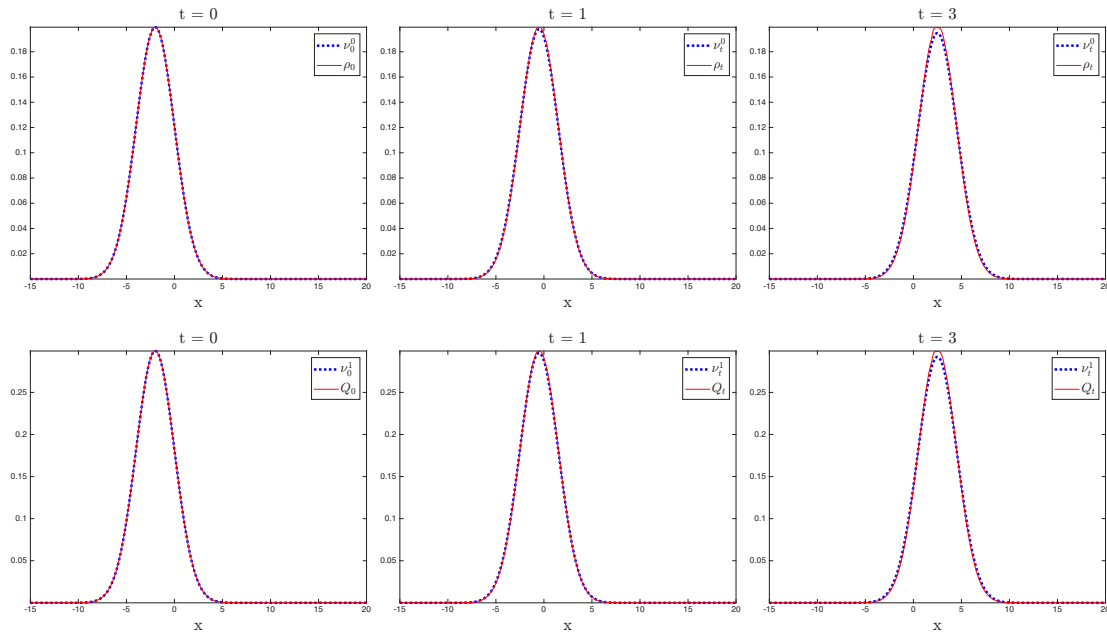


Figure 5: Test 4: approximation of monokinetic initial data. Comparison of v_t^0 and ρ_t (first line) and v_t^1 and Q_t (second line) at different time steps.

Test 5: Non-monokinetic initial condition. In Test 5 we consider a general initial data, to perform a comparison far from the monokinetic assumption. We run (1.4) setting $\Delta x = 0.2$ $\Delta v = 0.025$, starting with the initial distribution defined in (3.2). Screenshots of the distribution at different times are reported in the phase space, see Fig. 6. Hence we run the macroscopic model (1.8) with initial condition given by (3.6). Fig. 7 shows the comparison between kinetic and macroscopic scale, at different time instant. The non-monokinetic structure of the initial data seems to play a crucial role. Indeed the agreement between kinetic moments and macroscopic solution is no longer observed. Numerical results seem to suggest that the comparison is still valid only for short time, getting worse for longer time.

3.3 Sensitive dependence on initial data: numerical results in 1D and 2D

We conclude our multiscale analysis focusing on the sensitive dependence of the model on initial conditions. Indeed, as shown in [4], there exist pathological initial configurations for which the dynamics is not well-defined. Specifically, this occurs when two agents are equidistant from a third agent, and this equidistance is preserved in the past or future.

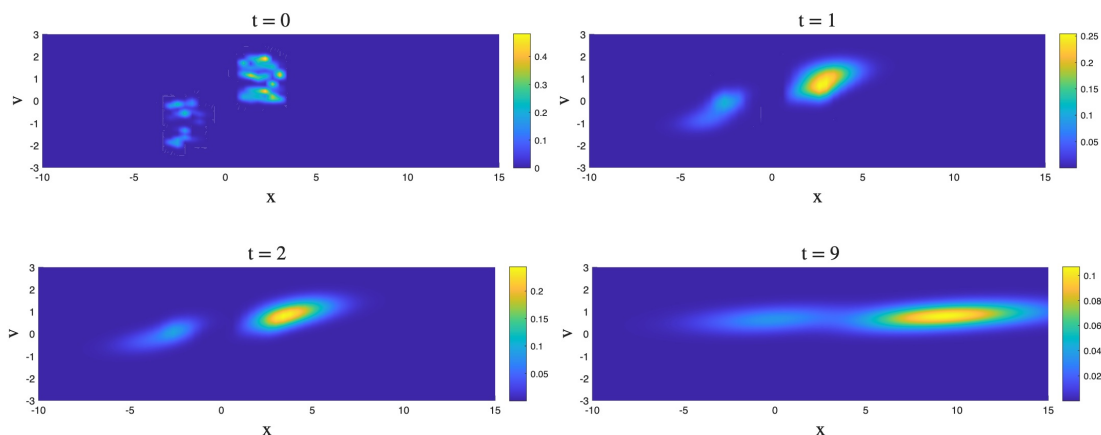


Figure 6: Test 5: screenshots of a numerical simulations of (1.4) in the phase space.

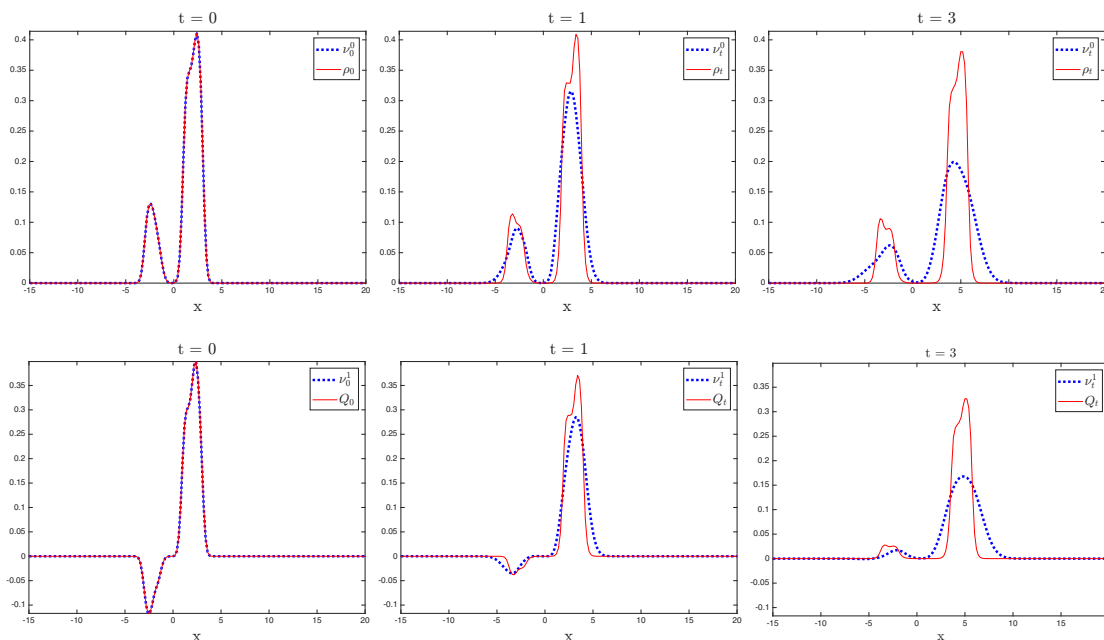


Figure 7: Test 5: Comparison of v_t^0 and ρ_t (first line) and v_t^1 and Q_t (second line) at different time steps.

Test 6: Sensitive dependence on initial data in 1D. In [4], authors prove well-posedness of (1.1) for almost all initial data, giving an example of critical initial datum for the system in 1D. Initial positions and velocities with $N = 3$ agents are given as $X^0 = (-1, \varepsilon, 1)$, $V^0 = (-1, 0, 1)$, with $\varepsilon \in (-1, 1), \varepsilon \neq 0$. The model coefficients p_{ij} take values $\{1/3, 2/3, 1\}$. Assuming $K(2/3) = 1$ and $K(1) = 0$, the dynamics varies depending on ε ,

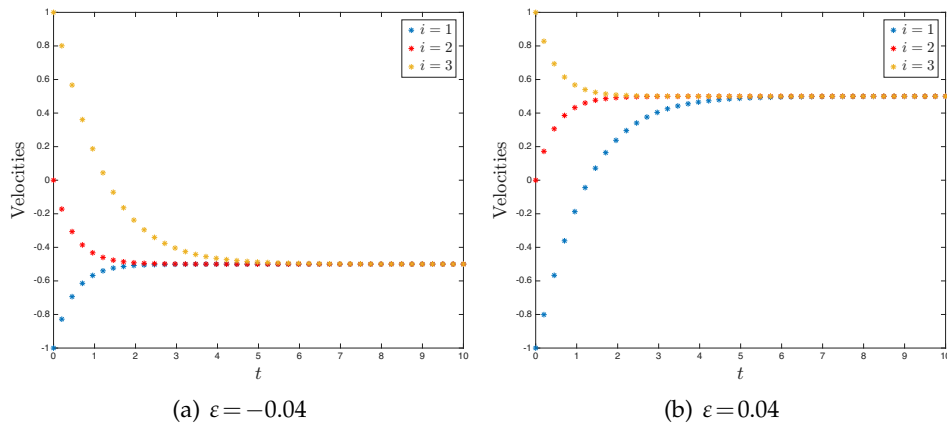


Figure 8: Test 6: numerical simulation in 1D of (1.1). Here $N=3$, $X^0 = (-1, \epsilon, 1)$, $V^0 = (-1, 0, 1)$.

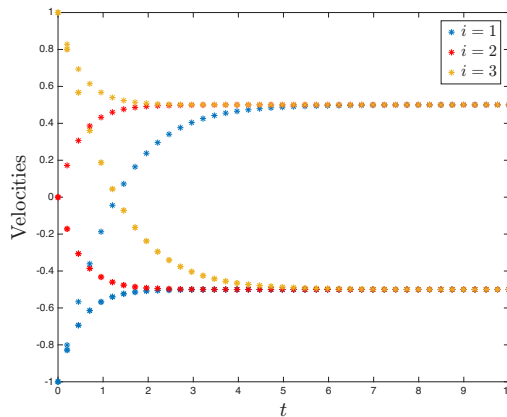


Figure 9: Test 6: numerical simulation in 1D of (1.1). Here $N=3$, $X^0 = (-1, 0, 1)$, $V^0 = (-1, 0, 1)$, depending on the two *by hand* selections of the first neighbor.

leading to a discontinuity in $\epsilon = 0$. Fig. 8 shows the evolution of V_i , for $i = 1, 2, 3$, for two different values of the initial position of agent labeled with $i = 2$. We observe that a small variation on the value of ϵ reflects in a fairly different dynamics. In particular, for $\epsilon = 0$, the solution to system (1.1) is not unique, since there exists at least two solutions, each being approached by the solution obtained for $\epsilon \rightarrow 0^+$ and $\epsilon \rightarrow 0^-$. Fig. 9 shows the two solutions obtained running the system assuming $\epsilon = 0$. Since in this case the first nearest neighbor is not unique, we obtain two different solutions, forcing the algorithm to select the particle located at X_1^0 or the one at X_3^0 , leading to two different consensus velocities. This is the same phenomena observed in standard one particle dynamics with low regularity (less than Cauchy-Lipschitz), see e.g. [22, Remark IV.3]. It is important to note that — in the one dimensional example considered — the ambiguity in selecting the nearest agent

persists over time. Indeed, in this case, two agents are equidistant from a third one, and the equidistance is preserved in the past or future. This situation corresponds to what are defined as irregular points of the *iso-rank manifold* in [4]. Therefore, any selection criterion, such as choosing the agent with the lowest (or highest) index, can only be arbitrary. It is worth noting that in [4] and [5], the derivation of the kinetic equation is performed outside this set of initial conditions which are pathological for the well-posedness of the dynamics and have Lebesgue measure zero.

Tests 7-8: Sensitive dependence on initial data in 2D. We then present a 2D scenario, generalizing the initial condition to N agents, showing the sensitive dependence of the solutions on the initial data in higher dimension, both at microscopic and macroscopic scales. First we run the microscopic model (1.1) with $N = 100$ agents. Test 7 and Test 8 differ on the initial condition. At initial time, two groups of 45 agents are randomly located in balls, centered in $(-0.3, -0.3)$ and $(0.3, 0.3)$, and radius $R = 0.1$. Each agent i belonging to the group starts moving with initial velocity $V_i^0 = (V_{ix}^0, V_{iy}^0)$ randomly chosen in $[-1, -0.2] \times [-1, -0.2]$, whereas initial velocities of agents belonging to the groups on the right are randomly chosen in $[0.2, 1] \times [0.2, 1]$. A small group of 10 agents with null velocities, is located in the ball centered in $(-0.08, -0.08)$ in Test 7, and in $(0.08, 0.08)$ in Test 8. In the following $(X_i^0, Y_i^0) \in \mathbb{R}^2$ denotes the initial position of each agent i . At the macroscopic scale, we run the system (1.8) considering the spatial domain $[-1, 1] \times [-1, 1]$ in the time interval $[0, 1]$, with a linear function K . We approximate the solution implementing the scheme detailed in Section 2.2, with $\Delta_x = \Delta_y = 0.04$. We consider an initial density given as the sum of Gaussian functions,

$$\rho_0(x, y) = \sum_{i=1}^N \frac{1}{2\pi\sigma_x\sigma_y} e^{-\frac{(x-X_i^0)^2}{2\sigma_x^2} - \frac{(y-Y_i^0)^2}{2\sigma_y^2}} \quad (3.7)$$

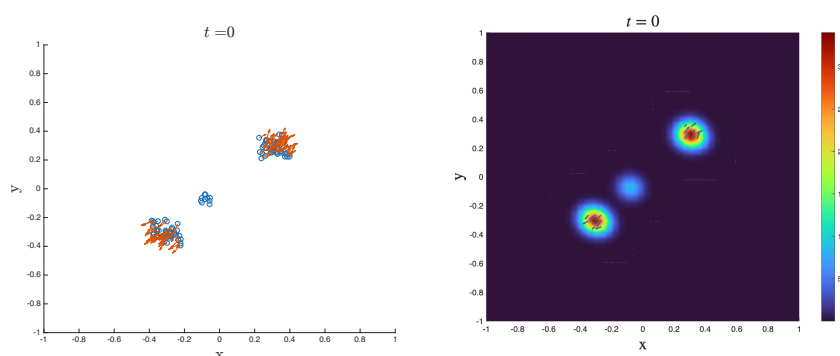


Figure 10: Test 7: initial conditions for the microscopic scale and the macroscopic one. The small group is located in a circle centered in $(-0.08, -0.08)$.

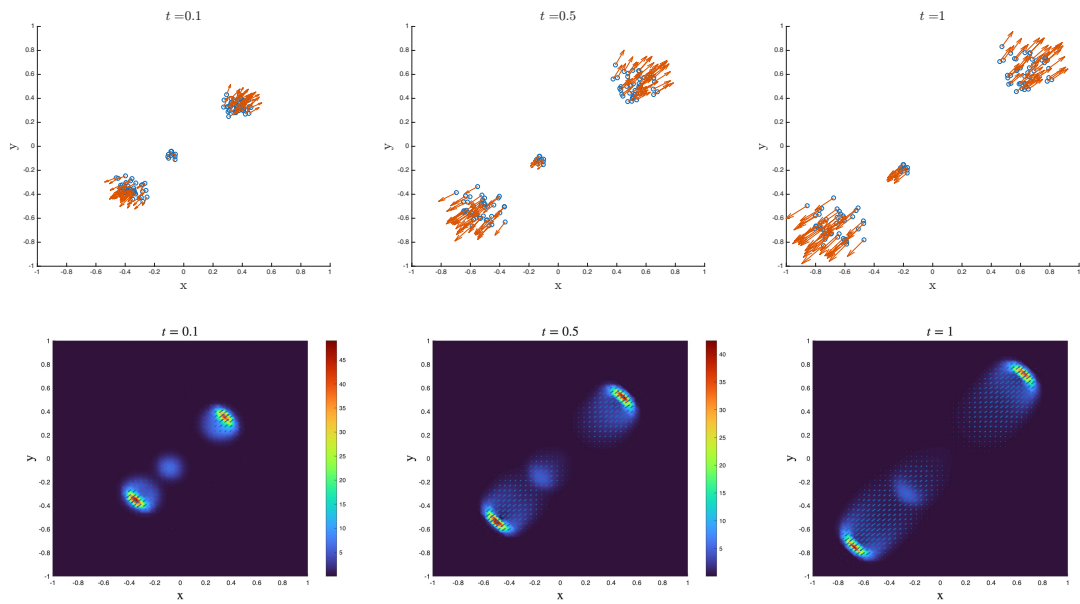


Figure 11: Test 7: screenshots of the numerical simulation at (first line) the microscopic scale and (second line) the macroscopic scale.

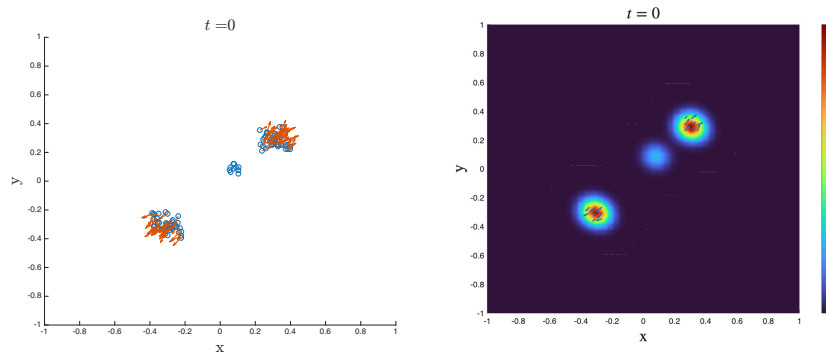


Figure 12: Test 8: initial conditions for the microscopic scale and the macroscopic one. The small group is located in a circle centered in $(0.08, 0.08)$.

with $\sigma_x = \sigma_y = 0.05$. A plot of initial conditions for Test 7 and Test 8 is shown in Fig. 10 and Fig. 12, respectively. Fig. 11 and Fig. 13 show several screenshots of the evolution of systems, both at microscopic and macroscopic scale. Depending on its initial position, the small group starts moving downward (Test 7) or upward (Test 8) reaching consensus with the closest group. A small variation on the initial data reflects in a different overall dynamics for the system, at both scales. Note that the well-posedness issue does not depend on the number of agents (in the microscopic case, $N = 3$ is anyway sufficient to

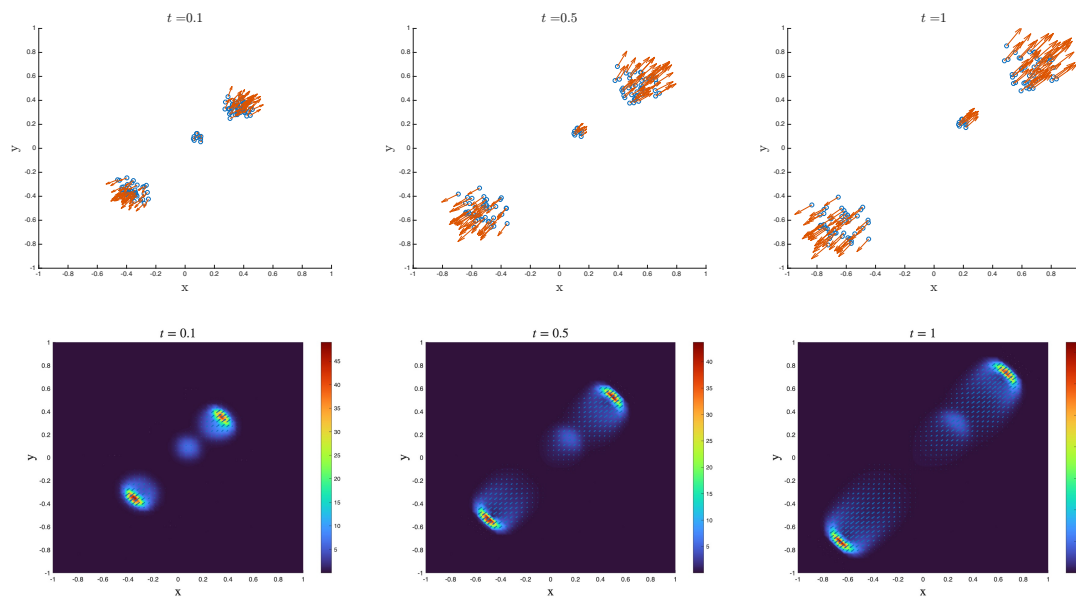


Figure 13: Test 8: screenshots of the numerical simulation at (first line) the microscopic scale and (second line) the macroscopic scale.

observe this phenomenon). The evolution considered in Test 7 and Test 8 —with two clouds of particles moving in opposite directions—represents perfectly valid initial conditions, close to the pathological one, and serve as an example illustrating the sensitive dependence on the initial data.

4 Conclusion

In this paper we investigated numerically flocking dynamics with topological interactions at different scales. The microscopic model represented the starting point of the analysis, for which several choices of interaction kernels were considered and analyzed. The corresponding Vlasov-type kinetic model was rigorously derived in [4] and its hydrodynamic limit to a pressureless Euler-type system was proved in [5], under monokinetic assumptions on the initial condition. In our work we studied the agreement between different scales in several scenarios, also far from monokinetic initial conditions. We finally focused on the sensitive dependence on the initial data in 1D and 2D scenarios, including the case of critical initial conditions leading to non uniqueness of solutions. Future perspectives include further investigations of this aspect, including analogy with the selection of the limit for solutions to ODE with rough vector fields outside the Cauchy-Lipschitz theory. Moreover, a complete investigation of the macroscopic model, extending the analysis to the case of different topological interactions (e.g. attraction and repulsion effects) will be performed. At kinetic and continuous scales, the role of topo-

logical interactions in continuous leader–follower transitions will be investigated, in the spirit of the work in [11] where metric interactions are considered. From a numerical point of view, we aim to extend the rescaling velocity method proposed in [26] to the case of kinetic flocking models with topological interactions.

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