

A kinetic model for epidemic spread

M Pulvirenti, S Simonella

▶ To cite this version:

M Pulvirenti, S Simonella. A kinetic model for epidemic spread. Mathematics and Mechanics of Complex Systems, 2020, 8 (3). hal-02614994

HAL Id: hal-02614994

https://hal.science/hal-02614994

Submitted on 21 May 2020

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

A kinetic model for epidemic spread

M. Pulvirenti¹ and S. Simonella²

1. Dipartimento di Matematica, Università di Roma La Sapienza Piazzale Aldo Moro 5, 00185 Rome – Italy, and International Research Center M&MOCS, Università dell'Aquila, Palazzo Caetani, 04012 Cisterna di Latina – Italy.

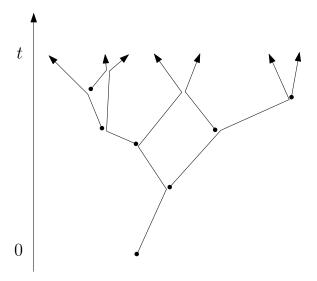
> 2. UMPA UMR 5669 CNRS, ENS DE LYON 46 ALLÉE D'ITALIE, 69364 LYON CEDEX 07 – FRANCE

Abstract. We present a Boltzmann equation for mixtures of three species of particles reducing to the Kermack-McKendrick (SIR) equations for the time-evolution of the density of infected agents in an isolated population. The kinetic model is potentially more detailed and might provide information on space mixing of the agents.

Keywords. Boltzmann equation; SIR model; low-density limit; stochastic particle system; forward cluster.

1 Boltzmann–SIR equations

Consider a population of identical individuals (particles) moving in the physical space and interacting upon contact. One (or several) of the individuals, say particle 1, has an infected status at time zero. As the dynamics runs, the infection can be transmitted, at the interaction times, to the individuals entering in contact with 1 or with the newly infected individuals. A cluster $\{i_1, i_2, \dots\}$ of infection grows in time, determined by the particle evolution: an individual is potentially infected at time t > 0 if it is involved, directly or indirectly, in the forward-in-time dynamics of 1. The "forward cluster of particle 1" (according to a terminology of [1, 22]) is represented symbolically in the picture below.



For concreteness, we may want to fix an idealized mechanical setting. Let us then proceed, as it is customary in kinetic theory, by looking at N hard spheres of unit mass and diameter $\varepsilon > 0$. The balls move in $\Lambda \subset \mathbb{R}^d$, d = 2, 3, and interact through elastic collisions. Each particle flies freely with constant velocity and, when two hard spheres collide with positions x, x_* at distance ε and incoming velocities v, v_* the latter are instantaneously transformed to outgoing velocities v', v'_* by the relations

$$\begin{cases} v' = v - \omega[\omega \cdot (v - v_*)] \\ v'_* = v_* + \omega[\omega \cdot (v - v_*)] \end{cases}, \tag{1.1}$$

where ω is the normalized relative distance $\omega = (x - x_*)/|x - x_*| = (x - x_*)/\varepsilon \in \mathbb{S}^{d-1}$.

We shall mimic the basic model in the mathematical theory of epidemics [14], by means of several assumptions. There are three different species of particles: S, I and R which stay for susceptible, infected and recovered, respectively. Upon collision between a particle of type S and a particle of type I, the reaction

$$S + I \rightarrow I + I$$

occurs instantaneously with rate $\beta \in [0,1]$. All the other collisions do not change the particle type but, in addition, a decay

$$I \to R$$

occurs with rate $\gamma \in [0, 1]$. Note that the population size is fixed (no deaths), and that the infection implies complete immunity. Finally for simplicity, we shall assume that β and γ are constants (they do not depend on time).

We are relying on the idea that the details of the interactions should not be of crucial importance (see [27] for a recent popular article simulating a similar system of particles). The main features are instead the following.

- The interactions are binary, and localized.
- The number of interactions per unit time is expected to be finite.
- The qualitative behaviour is independent of the number of particles N, provided that this is large in a suitable scaling limit.
- A statistical description is appropriate.

Under these assumptions, the Boltzmann equation for rarefied gases provides a tool of investigation.

Let us perform the so called Boltzmann-Grad limit [9] on the hard sphere system under consideration. Denoting the one-particle distribution functions by

$$f_S = f_S(t, x, v)$$
$$f_I = f_I(t, x, v)$$
$$f_R = f_R(t, x, v)$$

for the three species of particles, we obtain the following set of equations:

$$\begin{cases}
(\partial_t + v \cdot \nabla_x) f_S = Q(f_S, f_S) + Q(f_S, f_R) + (1 - \beta)Q(f_S, f_I) - \beta Q_-(f_S, f_I) \\
(\partial_t + v \cdot \nabla_x) f_I = Q(f_I, f) + \beta Q_+(f_S, f_I) - \gamma f_I \\
(\partial_t + v \cdot \nabla_x) f_R = Q(f_R, f) + \gamma f_I
\end{cases} , (1.2)$$

where

$$f = f_S + f_I + f_R$$

and Q is Boltzmann's operator (expressed in asymmetric form)

$$Q = Q_{+} - Q_{-}$$

$$Q_{+}(f,g)(v) := \int_{\mathbb{R}^{d}} \int_{\mathbb{S}^{d-1}} B(\omega; v - v_{*}) f(v') g(v'_{*}) d\omega dv_{*} .$$

$$Q_{-}(f,g)(v) := f(v) \int_{\mathbb{R}^{d}} \int_{\mathbb{S}^{d-1}} B(\omega; v - v_{*}) g(v_{*}) d\omega dv_{*} .$$

Note that the sum $f = f_S + f_I + f_R$ satisfies the classical Boltzmann equation

$$(\partial_t + v \cdot \nabla_x) f = Q(f, f) .$$

Here we chose $B(\omega; v - v_*) = (\omega \cdot (v - v_*)) \mathbbm{1} (\omega \cdot (v - v_*) \ge 0)$, corresponding to the hard-sphere cross section. However as said above, conclusions drawn from the kinetic model should not be very sensitive on the interaction rule; e.g. we shall consider as well different kernels $B(\omega; v - v_*) \ge 0$ such that

$$\int_{\mathbb{S}^{d-1}} B(\omega; v - v_*) d\omega = |v - v_*|^b,$$

for some $b \geq 0$.

In the second part of this exposition we will give more details on the passage from the particle dynamics to (1.2). Before that, we make a few elementary remarks on the equations themselves.

1.1 Maxwell collisions: Kermack-McKendrick equations

Averaging (1.2) over velocities, the Q operators vanish (because $\int Q_+ = \int Q_-$) and, in the spatially homogeneous case (no dependence on x), the expected fractions of individuals of the species $A \in \{S, I, R\}$, $A(t) = \int f_A(t, v) dv$, satisfy the equations

$$\begin{cases} \dot{S} = -\beta \int |v - v_*|^b f_S(v) f_I(v_*) dv dv_* \\ \dot{I} = \beta \int |v - v_*|^b f_S(v) f_I(v_*) dv dv_* - \gamma I \\ \dot{R} = \gamma I \end{cases}.$$

These equations are not closed, except when dealing with "Maxwellian molecules" (case b = 0 [4]) for which we get

$$\begin{cases} \dot{S} = -\beta I S \\ \dot{I} = \beta S I - \gamma I \\ \dot{R} = \gamma I \end{cases}$$
(1.3)

namely the epidemiology model of [14] in the case of time-independent rates. This model has been analysed and used extensively, and several generalizations have been conceived; see e.g. [2, 5, 17, 11], and the many references therein. The kinetic equation (1.2) stands as an extension accounting for dependence on space and velocity of the individuals.

To remind the reader of the original motivations for such SIR models ([24, 14]), we recall that an epidemic is not necessarily terminated by the exhaustion of the susceptible individuals, nor by the extinction of the virulence. This is apparent from (1.3), over a threshold value of the density. Setting indeed $A_{\infty} := \lim_{t\to\infty} A(t)$, $A_0 = A(0)$ and

 $R(t) = R_0 + \gamma \int_0^t I(\tau) d\tau$ (showing that $I(t) \to 0$ as $t \to \infty$), one has that $\frac{dS}{dR} = -\frac{\beta}{\gamma} S$ and hence (by $R_\infty + S_\infty = 1$ and the assumption $R_0 = 0$) $S_\infty = S_0 e^{-\frac{\beta}{\gamma}(1 - S_\infty)}$, or

$$e^{-\frac{\beta}{\gamma}S_{\infty}}\frac{\beta}{\gamma}S_{\infty} = S_0 \frac{\beta}{\gamma}e^{-\frac{\beta}{\gamma}}.$$
 (1.4)

Since $\max y e^{-y} = \frac{1}{e}$, given a value of β/γ one can find non vanishing solutions for S_{∞} , provided that $I_0 = 1 - S_0$ is sufficiently large.

1.2 Confinement

The model can be easily adapted to investigate several different situations. Examples might be boundary conditions or external potentials, imposing internal spatial constraints or local enhancing of density. There has been recent intense interest in the effects of isolation of individuals, and of the reduction of social mixing, by means of physical distancing measures ([16, 18]). At the level of (1.2), the energy can be used as simple parameter regulating the interaction rate.

We just make an example of one adaptation of (1.2), intended to model a confinement effect. Following [27] we assume that, for each species, there are two types of particles: wandering and confined. We denote by g_A , $A \in \{S, I, R\}$, the distribution of confined particles, while we maintain the notation f_A for the wandering particles. The distribution of the species A is $h_A := f_A + g_A$ and $f = \sum_A h_A$. Wandering particles have mass $m_w = 1$, while confined particles have mass $m_c = +\infty$ and zero velocity. The distribution g_A is proportional to a Dirac delta in velocity. Confined particles are frozen and their total distribution is stationary:

$$g_S(t,x) + g_I(t,x) + g_R(t,x) = \text{const.} \quad \forall t .$$

The collision law becomes

$$\begin{cases} v' = v - \frac{2m_*}{m + m_*} \omega[\omega \cdot (v - v_*)] \\ v'_* = v_* + \frac{2m}{m + m_*} \omega[\omega \cdot (v - v_*)] \end{cases}$$

where m, m_* are the masses of the incoming particles, and Eq. (1.2) is replaced by

$$\begin{cases} (\partial_t + v \cdot \nabla_x) f_S = Q(f_S, h_S) + Q(f_S, h_R) + (1 - \beta)Q(f_S, h_I) - \beta Q_-(f_S, h_I) \\ (\partial_t + v \cdot \nabla_x) f_I = Q(f_I, f) + \beta Q_+(f_S, h_I) - \gamma f_I \\ (\partial_t + v \cdot \nabla_x) f_R = Q(f_R, f) + \gamma f_I. \\ \dot{g}_S = -\beta Q_-(g_S, f_I) \\ \dot{g}_I = \beta Q_+(g_S, f_I) - \gamma g_I \\ \dot{g}_R = \gamma g_I \end{cases}$$
(1.5)

In the spatially homogeneous case, integrating Eq.s (1.5) in v, calling $A_w = \int f_A dv$ and $A_c = \int g_A dv$, A = S, I, R, we obtain

$$\begin{cases} \dot{S}_w = -\beta \int |v - v_*|^b f_S(v) h_I(v_*) dv dv_* \\ \dot{I}_w = \beta \int |v - v_*|^b f_S(v) h_I(v_*) dv dv_* - \gamma I_w \\ \dot{R}_w = \gamma I_w \\ \dot{S}_c = -S_c \int |v_*|^b f_I(v_*) dv_* \\ \dot{I}_c = S_c \int |v_*|^b f_I(v_*) dv_* - \gamma I_c \\ \dot{R}_c = \gamma I_c \end{cases}$$

Again, Eq.s (1.6) reduce to a standard SIR model in the case of Maxwellian molecules:

$$\begin{cases}
\dot{S}_{w} = -\beta S_{w} \left(I_{w} + I_{c} \right) \\
\dot{I}_{w} = \beta S_{w} \left(I_{w} + I_{c} \right) - \gamma I_{w} \\
\dot{R}_{w} = \gamma I_{w} \\
\dot{S}_{c} = -S_{c} I_{w} \\
\dot{I}_{c} = S_{c} I_{w} - \gamma I_{c} \\
\dot{R}_{c} = \gamma I_{c}.
\end{cases} (1.6)$$

1.3 Related problems

The kinetic model presented above should be interpreted as a remark of mathematical physics character: we do not pretend that it can be of use in epidemiology. It is more detailed than the classical SIR, insofar it includes space and velocities of the agents. Presumably, its main potential interest in applications is the identification of spatial patterns having an impact on the history of epidemics. Moreover, a dynamical representation in

terms of forward (or backward) clusters would provide information on the tracing of the infection. We comment next on a few other problems arising naturally.

The typical question concerning SIR equations is determining the long-time behaviour in relation with the parameters β, γ , and its dependence on local characteristics of the initial data. We are interested in masses, but also in local densities in the presence of spatial inhomogeneities. From the mathematical side, little can be done, but the problem is suited to numerical investigation. In analogy with gas dynamics, it is natural to use stochastic methods, as we will discuss in the next section.

At the theoretical level, it would be interesting to detect large scale limits and derive, starting from (1.2), equations for locally conserved quantities. Eq. (1.2) can be useful in fact for limited amounts of time. Preliminarily, one should characterize the equilibria. Let $F_A = \lim_{t\to 0} f_A$ be the asymptotic distributions. Then we expect $F_I = 0$, and the other two distributions should satisfy

$$\begin{cases} Q(F_S, F_S) + Q(F_S, F_R) = 0 \\ Q(F_R, F_R) + Q(F_R, F_S) = 0 \end{cases}$$

The latter equation is satisfied if both F_S and F_R are Maxwellians

$$F_A = A_\infty \frac{e^{-\frac{(v-u)^2}{2\sigma^2}}}{(2\pi\sigma^2)^{d/2}}$$

for some constants S_{∞} and R_{∞} , with σ and u determined by the initial conditions. A_{∞} would be obtained as in (1.4). Notice that, when $f = f_S + f_I + f_R$ is a global equilibrium, a solution (f_S, f_I, f_R) of Eq. (1.2) for b = 0 is given by the same global equilibrium with densities S(t), I(t), R(t) driven by (1.3).

2 Particle systems

2.1 Stochastic particle system

In this section we introduce a particle system yielding, in a suitable scaling limit, kinetic equations of type (1.2). The interest of this dynamics is twofold. First, it can be considered as a microscopic model to be accepted as the phenomenology, covering a large variety of kernels B. It would be somewhat funny to believe that the laws of Newton can be used to describe efficiently the interaction among individuals. On the other hand, we do not know so much concerning the details of such interactions, thus a stochastic collision appears

to be more robust than a deterministic one. Secondly, the particle scheme corresponds numerically to the direct simulation Monte Carlo method, widely used to approximate rarefied gas dynamics. There are several variants of such methods ([3, 25]). Below, we will deal with an inhomogeneous Kac model for three species with reactions [13].

We start by regularizing the collision operator (1.2). The strictly local interaction is smeared as follows:

$$\begin{split} Q^h &= Q^h_+ - Q^h_- \\ Q^h_+(f,g)(x,v) &:= \int_{\mathbb{R}^d} \int_{\mathbb{R}^d} \int_{\mathbb{S}^{d-1}} B(\omega;v-v_*) \, h(|x-y|) \, f(x,v') g(y,v_*') \, d\omega \, dv_* dy \\ Q^h_-(f,g)(x,v) &:= \int_{\mathbb{R}^d} \int_{\mathbb{R}^d} \int_{\mathbb{S}^{d-1}} B(\omega;v-v_*) \, h(|x-y|) \, f(x,v) g(y,v_*) \, d\omega \, dv_* dy \end{split}$$

where $h: \mathbb{R}^+ \to \mathbb{R}^+$ is a smooth approximation of the delta function.

To simplify the notation we limit ourselves to the case of (1.2) with $\beta = 1$, being the more general cases a trivial extension. We therefore consider the following equations:

$$\begin{cases} (\partial_t + v \cdot \nabla_x) f_S = Q^h(f_S, f_S) + Q^h(f_S, f_R) - Q_-^h(f_S, f_I) \\ (\partial_t + v \cdot \nabla_x) f_I = Q^h(f_I, f) + Q_+^h(f_S, f_I) - \gamma f_I \\ (\partial_t + v \cdot \nabla_x) f_R = Q^h(f_R, f) + \gamma f_I. \end{cases}$$
(2.1)

We can pass to the limit $Q^h \to Q$ inside (2.1), whenever we have a smooth solution of the initial value problem.

We shall indicate by $\mathcal{A} = \mathcal{S}, \mathcal{I}, \mathcal{R} \subset \{1, 2, \dots, N\}$ the (random) disjoint sets of particles of type A = S, I, R respectively. They form a partition of $\{1, 2, \dots, N\}$, so that the process $Z_N : \mathbb{R}^+ \to \mathcal{X}, Z_N = Z_N(t) = (z_1(t), \dots, z_N(t)), z_i = (x_i, v_i)$, takes values in

$$\mathcal{X} = \bigcup_{\mathcal{S},\mathcal{I},\mathcal{R}} \mathcal{X}(\mathcal{S},\mathcal{I},\mathcal{R}), \quad \mathcal{X}(\mathcal{S},\mathcal{I},\mathcal{R}) = \{ (Z_{\mathcal{S}}, Z_{\mathcal{I}}, Z_{\mathcal{R}}) \}$$

with

$$|\mathcal{S}| + |\mathcal{I}| + |\mathcal{R}| = N ,$$

and $z_i \in \Lambda \times \mathbb{R}^d$. Here $|\mathcal{A}|$ denotes the cardinality of the set \mathcal{A} . The configuration of particles in the three species are $Z_{\mathcal{S}} = (z_{s_1}, z_{s_2}, \cdots), \ Z_{\mathcal{I}} = (z_{i_1}, z_{i_2}, \cdots)$ and $Z_{\mathcal{R}} = (z_{r_1}, z_{r_2}, \cdots),$ respectively.

Le us define the time evolution. Particles move freely for a random time, exponentially distributed with intensity scaling like N. Then two particles are randomly chosen, say particles j and k, according to $\int B(\omega; v_j - v_k) h(|x_j - x_k|) d\omega$ and their velocities are updated

as in (1.1) with $\omega \sim B(\cdot; v_j - v_k)$. If the pair of colliding particles is of type (A, A) or (S, R) or (I, R), the particles do not change their species. If the pair is of type (S, I), then the outgoing pair is of type (I, I). We abbreviate from now on $h_{j,k} = h(|x_j - x_k|)$, and we denote by J_{jk} the linear operator transforming the velocities j and k to a postcollisional pair with scattering vector ω . The generator of the process reads

$$\mathcal{L} = \mathcal{L}_0 + \mathcal{L}_i + \mathcal{L}_d$$

where $\mathcal{L}_0 = \sum v_i \cdot \nabla_{x_i}$ is the generator of the free motion,

$$\mathcal{L}_{i}\phi(Z_{N}) = \frac{1}{N} \sum_{j \in \mathcal{S}} \sum_{k \in \mathcal{I}} \int B(\omega; v_{j} - v_{k}) h_{j,k}$$

$$\times \left(J_{jk}\phi \left(Z_{\mathcal{S} \setminus \{j\}}, Z_{\mathcal{I} \cup \{j\}}, Z_{\mathcal{R}} \right) - \phi(Z_{N}) \right) d\omega$$

$$+ \frac{1}{N} \left(\sum_{j \in \mathcal{S}} \sum_{k \in \mathcal{R}} + \sum_{j \in \mathcal{I}} \sum_{k \in \mathcal{R}} \right) \int B(\omega; v_{j} - v_{k}) h_{j,k} \left(J_{jk}\phi \left(Z_{N} \right) - \phi(Z_{N}) \right) d\omega$$

$$+ \frac{1}{2N} \sum_{\mathcal{A} = \mathcal{S}, \mathcal{I}, \mathcal{R}} \sum_{\substack{j,k \in \mathcal{A} \\ j \neq k}} \int B(\omega; v_{j} - v_{k}) h_{j,k} \left(J_{jk}\phi \left(Z_{N} \right) - \phi(Z_{N}) \right) d\omega , \qquad (2.2)$$

and

$$\mathcal{L}_d \phi(Z_N) = \gamma \sum_{i \in \mathcal{I}} \left(\phi(Z_{\mathcal{S}}, Z_{\mathcal{I} \setminus \{i\}}, Z_{\mathcal{R} \cup \{i\}}) - \phi(Z_N) \right) . \tag{2.3}$$

We choose now test functions of the form

$$\phi_A(Z_N) = \frac{1}{N} \sum_{\ell \in \mathcal{A}} \varphi(z_\ell)$$

and focus, for instance, on the case $\mathcal{A} = \mathcal{S}$. We have that $\mathcal{L}_d \phi_S = 0$. Evaluating Eq. (2.2) in ϕ_S we notice that, given j and k, all the terms with $\ell \neq j, k$ cancel out. In the second line of (2.2) we find

$$\sum_{\substack{\ell \in \mathcal{S} \\ \ell \neq j}} J_{jk} \, \varphi(z_{\ell}) - \sum_{\ell \in \mathcal{S}} \varphi(z_{\ell}) = -\varphi(z_{j}) \; .$$

Therefore

$$\mathcal{L}_{i}\phi_{S}(Z_{N}) = -\frac{1}{N^{2}} \sum_{j \in \mathcal{S}} \sum_{k \in \mathcal{I}} \int B(\omega; v_{j} - v_{k}) h_{j,k} \varphi(z_{j}) d\omega$$

$$+ \frac{1}{N^{2}} \sum_{j \in \mathcal{S}} \sum_{k \in \mathcal{R}} \int B(\omega; v_{j} - v_{k}) h_{j,k} \left(\varphi(x_{j}, v'_{j}) - \varphi(z_{j}) \right) d\omega$$

$$+ \frac{1}{2N^{2}} \sum_{\substack{j,k \in \mathcal{S} \\ j \neq k}} \int B(\omega; v_{j} - v_{k}) h_{j,k} \left(\varphi(x_{j}, v'_{j}) + \varphi(x_{k}, v'_{k}) - \varphi(z_{j}) - \varphi(z_{k}) \right) d\omega .$$

(2.4)

Next, we introduce a probability measure with density $W^N: \mathcal{X} \to \mathbb{R}^+$, assumed to be symmetric in the exchange of the particle labels within each one of the species. An example is provided by the fully factorized (chaotic) state, which we shall assume, to fix ideas, as initial distribution of the particle process: $W^N(0) = f_0^{\otimes N}$ with $f^0 = \sum_A f_A^0$, A = (S, I, R), where f_A^0 are the initial data for (2.1). We further denote by $f_A^N = f_A^N(z)$ the one-particle marginals of W^N , defined as

$$\int f_A^N(z)\varphi(z)dz = \int W^N(Z_N)\phi_A(Z_N)dZ_N .$$

It is the probability density of finding a particle of type A in z. Similarly, $f_{A_1,A_2}^N = f_{A_1,A_2}^N(z_1,z_2)$ denotes the two-particle marginal, namely the probability density of finding two particles of type A_1 and A_2 in z_1 and z_2 :

$$\int f_{A_1,A_2}^N(z_1,z_2)\varphi(z_1,z_2)dz_1dz_2 = \int W^N(Z_N)\phi_{A_1,A_2}(Z_N)dZ_N$$

for $\phi_{A_1,A_2}(Z_N) = \frac{1}{N(N-1)} \sum_{j \in A_1} \sum_{\substack{k \in A_2 \\ k \neq j}} \varphi(z_j, z_k)$. Even though the initial measure is factorized, the time-evolved density $W^N(t)$ is not, due to correlations generated by the dynamics. The factorization is however recovered in the limit $N \to \infty$ and

$$f_{A_1,A_2}^N(z_1,z_2) \approx f_{A_1}^N(z_1) f_{A_2}^N(z_2)$$
 (2.5)

We are ready to compute

$$\frac{d}{dt} \int W^N(t) \, \phi_S = \int W^N(t) \mathcal{L} \phi_S \, .$$

Using (2.4), the definition of marginal and (2.5), we deduce that, as $N \to \infty$,

$$\frac{d}{dt} \int f_S^N(t) \varphi \approx \int f_S^N(v \cdot \nabla_x \varphi) + \int Q^h(f_S^N, f_S^N) \varphi + \int Q^h(f_S^N, f_R^N) \varphi - \int Q_-^h(f_S^N, f_I^N) \varphi ,$$
that is the first equation of (2.1) in weak formulation.

The other two equations can be recovered similarly. For A = I, Eq. (2.3) yields

$$\mathcal{L}_d \phi_I(Z_N) = \frac{\gamma}{N} \sum_{i \in \mathcal{I}} \left(\sum_{\ell \in \mathcal{I} \setminus \{i\}} \varphi(z_\ell) - \sum_{\ell \in \mathcal{I}} \varphi(z_\ell) \right) = -\frac{\gamma}{N} \sum_{i \in \mathcal{I}} \varphi(z_i) ,$$

while in the second line of (2.2) we find

$$\sum_{\ell \in \mathcal{I} \cup \{j\}} J_{jk} \, \varphi(z_{\ell}) - \sum_{\ell \in \mathcal{I}} \varphi(z_{\ell}) = J_{j,k} \varphi(z_{j}) + (J_{j,k} \varphi(z_{k}) - \varphi(z_{k}))$$

so that

$$\frac{d}{dt} \int f_I^N(t) \approx \int f_I^N \left(v \cdot \nabla_x \varphi \right) + \int Q^h \left(f_I^N, \sum_A f_A^N \right) \varphi + \int Q_+^h (f_S^N, f_I^N) \varphi - \gamma \int f_I^N \varphi ,$$

which is the second equation of (2.1).

2.2 Mechanical system

We briefly come back to the deterministic particle model, which was our starting point. That is, N hard spheres of diameter ε moving in the physical space and colliding elastically, with reactions simulating infection and recovery. We call this system "mechanical" as the interaction is deterministic. Clealry there is still stochasticity in the reactions and, strictly speaking, we are dealing again with a stochastic process.

We can easily adapt to this case the formal arguments of the previous section. The process Z_N takes still values in \mathcal{X} , but in addition the hard core exclusion is imposed $\min_{i\neq j} |x_i - x_j| > \varepsilon$. In the generator (2.2), 1/N is replaced by ε^{d-1} , B is the hard-sphere kernel $(\omega \cdot (v_j - v_k)) \mathbbm{1}(\omega \cdot (v_j - v_k) \geq 0)$, $h_{j,k}$ is absent and the operator $(J_{j,k} - 1)$ is replaced by $(\delta(x_k - x_j - \omega \varepsilon)J_{j,k} - \delta(x_k - x_j + \omega \varepsilon))$. Following [21], Section 2.1, and assuming the chaos property (2.5), Eq. (1.2) is obtained in the limit $N \to \infty$, $\varepsilon \to 0$ with $\varepsilon^{d-1}N = 1$.

2.3 Rigorous results

We have derived formally the kinetic equations under proper scaling limits, presenting only the basic ideas. A rigorous approach is possible, based on existing literature. In the case of the stochastic system, one can apply martingale techniques as in [28], or the hierarchy of equations for the family of the marginals [23], or coupling techniques [10]. In the case of the mechanical model, one can resort to the validity techniques for the Boltzmann equation, leading to a short time result; see [15] and subsequent works [12, 26, 6, 8, 19, 20, 7].

Acknowledgments. We are indebted to Grégory Miermont for enlightening comments inspiring this work. We thank Nicola Cotugno for advice on bibliography, and Robert Patterson for sharing numerical simulations and for useful comments on the manuscript.

References

- [1] K. Aoki, M. Pulvirenti, S. Simonella and T. Tsuji. Backward clusters, hierarchy and wild sums for a hard sphere system in a low-density regime. *Math. Models Methods Appl. Sci.* **25**(05), 995-1010 (2015).
- [2] R.M. Anderson and R.M. May. Population Biology of Infectious Diseases: Part I. Nature 280, 361-367 (1979).

- [3] G.A. Bird. Molecular Gas Dynamics and the Direct Simulation of Gas Flows. Oxford Engineering Science Series **42**, Clarendon Press (1994).
- [4] A.V. Bobylev. The theory of the nonlinear spatially uniform Boltzmann equation for Maxwell molecules. In: *Math. Phys. Rev.*, Soviet Sci. Rev. Sec. C Math. Phys. Rev. 7 (1988).
- [5] F. Brauer and C. Castillo-Chávez. Mathematical Models in Population Biology and Epidemiology. Springer (2001).
- [6] C. Cercignani, R. Illner and M. Pulvirenti. The mathematical theory of dilute gases. Springer-Verlag, New York (1994).
- [7] R. Denlinger. The propagation of chaos for a rarefied gas of hard spheres in the whole space. Arch. Rat. Mech. Anal. 229, 885-952 (2018).
- [8] I. Gallagher, L. Saint Raymond and B. Texier. From Newton to Boltzmann: hard spheres and short-range potentials. Zürich Adv. Lect. in Math. Ser. 18, EMS (2014).
- [9] H. Grad. On the kinetic theory of rarefied gases. Comm. on Pure and App. Math. **2**(4), 331–407 (1949).
- [10] C. Graham and S. Méléard. Stochastic particle approximations for generalized Boltzmann models and convergence estimates. *Annals of Probab.* **25** (1997).
- [11] T. Harko, F. Lobo and M.K. Mak. Exact analytical solutions of the Susceptible-Infected-Recovered (SIR) epidemic model and of the SIR model with equal death and birth rates. Appl. Math. and Comp. 236, 184-194 (2014).
- [12] R. Illner and M. Pulvirenti. Global Validity of the Boltzmann equation for a Twoand Three-Dimensional Rare Gas in Vacuum: Erratum and Improved Result. Comm. Math. Phys. 121, 143-146 (1989).
- [13] M. Kac. Foundations of kinetic theory. *Proc. Third Berkeley Symp. on Math. Stat.* and *Prob.*, University of California Press, Berkeley and Los Angeles (1956).
- [14] W. O. Kermack and A. G. McKendrick. A contribution to the mathematical theory of epidemics. Proc. Roy. Soc. Lond. A 115, 772 (1927).

- [15] O.E. Lanford. Time evolution of large classical systems. Lect. Notes Phys. 38, 1–111 (1975).
- [16] Q. Li, X. Guan, P. Wu et al. Early transmission dynamics in Wuhan, China, of novel coronavirus-infected pneumonia. New Engl. J. Med. 382, 13 (2020).
- [17] J.D. Murray. Mathematical Biology: I. An Introduction. Springer-Verlag (2002).
- [18] K. Prem, Y. Liu et al. The effect of control strategies to reduce social mixing on outcomes of the COVID-19 epidemic in Wuhan, China: a modelling study. *Lancet Public Health* 5, 5 (2020).
- [19] M. Pulvirenti, C. Saffirio and S. Simonella. On the validity of the Boltzmann equation for short-range potentials. *Rev. Math. Phys.* **26**, 1–64 (2014).
- [20] M. Pulvirenti and S. Simonella. The Boltzmann-Grad limit of a hard sphere system: analysis of the correlation error. *Inventiones* **207**(3), 1135-1237, (2017).
- [21] M. Pulvirenti and S. Simonella. A brief introduction to the scaling limits and effective equations in kinetic theory. hal-02406490 (2019).
- [22] M. Pulvirenti and S. Simonella. On the cardinality of collisional clusters for hard spheres at low density (2020).
- [23] M. Pulvirenti, W. Wagner and M.B. Zavelani Rossi. Convergence of particle schemes for the Boltzmann equation. *Eur. J. Mech. B/Fluids* **13** 3, 339–351 (1994).
- [24] R. Ross. An Application of the Theory of Probabilities to the Study of a priori Pathometry. Part I. Proc. Roy. Soc. Lond. A 92, 638 (1916).
- [25] S. Rjasanow and W. Wagner. Stochastic Numerics for the Boltzmann Equation. Springer, Berlin (2005).
- [26] H. Spohn. Large-Scale Dynamics of Interacting Particles. Springer, Berlin (1991).
- [27] H. Stevens. Why outbreaks like coronavirus spread exponentially, and how to "flatten the curve". *The Washington Post*, March 14 (2020), https://www.washingtonpost.com/graphics/2020/world/corona-simulator/.
- [28] W. Wagner. A convergence proof for Bird's direct simulation Monte Carlo method for the Boltzmann equation. J. Stat. Phys. **66** (3/4), 1011-1044 (1992).