

# The evolutionary limit for models of populations interacting competitively via several resources

Nicolas Champagnat, Pierre-Emmanuel Jabin

► **To cite this version:**

Nicolas Champagnat, Pierre-Emmanuel Jabin. The evolutionary limit for models of populations interacting competitively via several resources. *Journal of Differential Equations*, Elsevier, 2011, 251 (1), pp.179-195. <10.1016/j.jde.2011.03.007>. <inria-00488979v2>

**HAL Id: inria-00488979**

**<https://hal.inria.fr/inria-00488979v2>**

Submitted on 9 Mar 2016

**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.



# The evolutionary limit for models of populations interacting competitively via several resources

Nicolas Champagnat<sup>1</sup>, Pierre-Emmanuel Jabin<sup>1,2</sup>

## Abstract

We consider an integro-differential nonlinear model that describes the evolution of a population structured by a quantitative trait. The interactions between individuals occur by way of competition for resources whose concentrations depend on the current state of the population. Following the formalism of [16], we study a concentration phenomenon arising in the limit of strong selection and small mutations. We prove that the population density converges to a sum of Dirac masses characterized by the solution  $\varphi$  of a Hamilton-Jacobi equation which depends on resource concentrations that we fully characterize in terms of the function  $\varphi$ .

*MSC 2000 subject classifications:* 35B25, 35K55, 92D15.

*Key words and phrases:* adaptive dynamics, Hamilton-Jacobi equation with constraints, Dirac concentration, metastable equilibrium.

## 1 Introduction

We are interested in the dynamics of a population subject to mutation and selection driven by competition for resources. Each individual in the population is characterized by a quantitative phenotypic trait  $x \in \mathbb{R}$  (for example

---

<sup>1</sup>TOSCA project-team, INRIA Sophia Antipolis – Méditerranée, 2004 rte des Lucioles, BP. 93, 06902 Sophia Antipolis Cedex, France,  
E-mail: [Nicolas.Champagnat@sophia.inria.fr](mailto:Nicolas.Champagnat@sophia.inria.fr)

<sup>2</sup>Laboratoire J.-A. Dieudonné, Université de Nice – Sophia Antipolis, Parc Valrose, 06108 Nice Cedex 02, France, E-mail: [jabin@unice.fr](mailto:jabin@unice.fr)

the size of individuals, their age at maturity, or their rate of intake of nutrients).

We study the following equation

$$\partial_t u_\varepsilon(t, x) = \frac{1}{\varepsilon} \left( \sum_{i=1}^k I_i^\varepsilon(t) \eta_i(x) - 1 \right) u_\varepsilon(t, x) + M_\varepsilon(u_\varepsilon)(t, x), \quad (1.1)$$

where  $M_\varepsilon$  is the mutation kernel

$$M_\varepsilon(f)(x) = \frac{1}{\varepsilon} \int_{\mathbb{R}} K(z) (f(x + \varepsilon z) - f(x)) dz, \quad (1.2)$$

for a  $K \in C_c^\infty(\mathbb{R})$  such that  $\int_{\mathbb{R}} zK(z) dz = 0$ . Among many other ecological situations [15], this model is relevant for the evolution of bacteria in a chemostat [14, 16]. With this interpretation,  $u^\varepsilon(t, x)$  represents the concentration of bacteria with trait  $x$  at time  $t$ , the function  $\eta_i(x)$  represents the growth rate of the population of trait  $x$  due to the consumption of a resource whose concentration is  $I_i^\varepsilon$ , and the term  $-1$  corresponds to the decrease of the bacteria concentration due to the constant flow out of the chemostat. Note that the growth and mutation terms in (1.1) are of different nature, as the total mutation rate  $\int_{\mathbb{R}} K(z)$  of the population of trait  $x$ , which could either be produced by mutations during individuals' lives or occurring at births, does not depend on the resources concentrations  $I_i^\varepsilon$ . Note also that the resources consumption rates depend linearly on the resources concentrations. From a biological point of view, one might prefer a saturating functional response. Our method actually extends to all the models of [9, Thm. 1], including competitive Lotka-Volterra PDEs, but we chose here to keep the model simple to ease the presentation of the mathematical arguments. This model extends the one proposed in [16] to an arbitrary number of resources.

This equation has to be coupled with equations for the resources  $I_i$ , namely

$$I_i(t) = \frac{c_i}{1 + \int_{\mathbb{R}} \eta_i(x) u_\varepsilon(x) dx}, \quad (1.3)$$

where  $c_i > 0$ . This corresponds to an assumption of fast resources dynamics with respect to the evolutionary dynamics. The resources concentrations are assumed to be at a (quasi-)equilibrium at each time  $t$ , which depends on the current concentrations  $u^\varepsilon$ .

The limit  $\varepsilon \rightarrow 0$  corresponds to a simultaneous scaling of fast selection and small mutations. It was already considered in [16]. The following argument

explains what limit behaviour for  $u^\varepsilon$  can be expected when  $\varepsilon \rightarrow 0$ . Defining  $\varphi_\varepsilon$  as

$$u_\varepsilon = e^{\varphi_\varepsilon/\varepsilon}, \quad \text{or } \varphi_\varepsilon = \varepsilon \log u_\varepsilon, \quad (1.4)$$

one gets the equation

$$\partial_t \varphi_\varepsilon = \sum_{i=1}^k I_i^\varepsilon(t) \eta_i(x) - 1 + H_\varepsilon(\varphi_\varepsilon), \quad (1.5)$$

where

$$H_\varepsilon(f) = \int_{\mathbb{R}} K(z) (e^{(f(x+\varepsilon z)-f(x))/\varepsilon} - 1) dz. \quad (1.6)$$

At the limit  $\varepsilon \rightarrow 0$  the Hamiltonian  $H_\varepsilon$  simply becomes

$$H(p) = \int_{\mathbb{R}} K(z) (e^{pz} - 1) dz, \quad (1.7)$$

where  $p$  stands for  $\partial_x f(x)$ . So one expects Eq. (1.5) to lead to

$$\partial_t \varphi = \sum_{i=1}^k I_i(t) \eta_i(x) - 1 + H(\partial_x \varphi), \quad (1.8)$$

for some  $I_i$  which are unfortunately unknown since one cannot pass to the limit directly in (1.3). Therefore one needs to find a relation between the  $\varphi$  and the  $I_i$  at the limit. Under quite general assumptions on the parameters (see Lemma 3.1 below), the total population mass  $\int_{\mathbb{R}} u_\varepsilon$  is uniformly bounded over time. This suggests that  $\max_{x \in \mathbb{R}} \varphi(t, x) = 0$  should hold true for all  $t \geq 0$ . Together with (1.8), this gives a candidate for the limit dynamics as a solution to a Hamilton-Jacobi equation with Hamiltonian  $H$  and with unknown Lagrange multipliers  $I_i$ , subject to a maximum constraint. The limit population is then composed at time  $t$  of Dirac masses located at the maxima of  $\varphi(t, \cdot)$ .

This heuristics was justified in [16] in the case of a single resource (and when the resources evolve on the same time scale as the population), and the case of two resources was only partly solved. The mathematical study of the convergence to the Hamilton-Jacobi equation with maximum constraint and the study of the Hamilton-Jacobi equation itself have only been done in very specific cases [16, 2, 5, 26]. In fact the main problem in this proposed model is that the number of unknowns (the resources) may easily be larger than the

dimension of the constraint (formally equal to the number of points where  $\varphi = 0$ ).

Our goal in this paper is to prove the convergence of  $\varphi_\varepsilon$  to a solution of (1.8), where we give a full characterization of the functions  $I_i$ . Those are no more considered as Lagrange multipliers for a set of constraints but are given by the solution  $\varphi$  itself. The new resulting model describes the evolution of a population as Dirac masses and is formally well posed.

The general problem of characterizing evolutionary dynamics as sums of Dirac masses under biologically relevant parameter scalings is a key tool in *adaptive dynamics*—a branch of biology studying the interplay between ecology and evolution [20, 22, 23, 13, 7]. The phenomenon of *evolutionary branching*, where evolution drives an (essentially) monotype population to subdivide into two (or more) distinct coexisting subpopulations, is particularly relevant in this framework [23, 18, 19]. When the population state can be approximated by Dirac masses, this simply corresponds to the transition from a population composed of a single Dirac mass to a population composed of two Dirac masses.

Several mathematical approaches have been explored to study this phenomenon. One approach consists in studying the stationary behaviour of an evolutionary model involving a scaling parameter for mutations, and then letting this parameter converge to 0. The stationary state has been proved to be composed of one or several Dirac masses for various models (for deterministic PDE models, see [4, 5, 12, 21, 17], for Fokker-Planck PDEs corresponding to stochastic population genetics models, see [3], for stochastic models, see [27], for game-theoretic models, see [11]). Closely related to these works are the notions of ESS (evolutionarily stable strategies) and CSS (convergence stable strategies) [23, 14], which allow one in some cases to characterize stable stationary states [4, 12, 21, 11].

The other main approach consists in studying a simultaneous scaling of mutation and selection, in order to obtain a limit dynamics where transitions from a single Dirac mass to two Dirac masses could occur. Here again, deterministic and stochastic approaches have been explored. The deterministic approach consists in applying the scaling of (1.1). The first formal results have been obtained in [16]. This was followed by several works on other models and on the corresponding Hamilton-Jacobi PDE [5, 26]. For models of the type we consider here, rigorous results (especially for the well posedness of the Hamilton-Jacobi eq. at the limit) mainly only exist in the case with just one resource, see [2] and [1] (one resource but multidimensional traits).

The stochastic approach is based on individual-based models, which are related to evolutionary PDE models as those in [12, 21] through a scaling of large population [8]. Using a simultaneous scaling of large population and rare mutations, a stochastic limit process was obtained in [6] in the case of a monotype population (i.e. when the limit process can only be composed of a single Dirac mass), and in [10] when the limit population can be composed of finitely many Dirac masses.

Finally note that the total population of individuals is typically very high, for bacteria for example. This is why even stochastic models will usually take some limit with infinite populations. Of course, this has some drawbacks. In particular the population of individuals around a precise trait may turn out to be low (even though the total population is large). As in the scaling under consideration, one has growth or decay of order  $\exp(C/\varepsilon)$ , it is in fact quite common that the population density of large sets of traits becomes much smaller than the density corresponding to a single individual, making the notion of density meaningless. One of the most important open problem would be to derive models that are both able of dealing with very large populations and still treat correctly the small subpopulations (keeping the stochastic effects or at least truncating the population with less than 1 individual).

There are already some attempts in this direction, mainly proposing models with truncation, see [25] and very recently [24]. In these works however the truncation is only made at some level  $\exp(-C/\varepsilon)$ .

In order to state our main result, we need some regularity and decay assumptions on the  $\eta_i$ , namely

$$\eta_i > 0, \quad \exists \bar{\eta} \in C_0(\mathbb{R}), \quad \forall x, \quad \sum_{i=1}^k \left( |\eta_i(x)| + |\eta_i'(x)| + |\eta_i''(x)| \right) \leq \bar{\eta}(x), \quad (1.9)$$

where  $C_0(\mathbb{R})$  is the set of continuous function, tending to 0 as  $x \rightarrow \pm\infty$ .

Our method of characterization of the environmental variables  $I_i$  in (1.8) is based on the general principle that there exists a unique environment  $I_1, \dots, I_k$  corresponding to any given population state. Here, the population state at time  $t$  is given by the set  $\{x \in \mathbb{R} : \varphi(t, x) = 0\}$ . This principle was the basis of the work of Barles and Perthame [2] in the case of a single resource, where the previous correspondance is automatically satisfied under very general assumptions. In our case, we characterize  $I_i$  from  $\omega := \{\varphi(t, \cdot) =$

0} by first constructing a *population measure* corresponding to the set  $\omega$ , which has support in this set and which is a sort of metastable population equilibrium when there is no mutations (more specifically, we require it to be an ESS in the set of population measures with support in  $\omega$ , see Prop. 1.1 below and [21]). Next, we construct the  $I_i$  from this population measure. The following assumptions characterize the situations where our approach can be applied, that is where uniqueness holds in the two steps previously described. The first assumption deals with the number of possible roots of the reproduction rate:

$$\exists 1 \leq \bar{k} \leq k, \forall I_1 \in [0, c_1], \dots, I_k \in [0, c_k],$$

$$\text{the function } x \mapsto \sum_{i=1}^k I_i \eta_i(x) - 1 \text{ has at most } \bar{k} \text{ roots.} \quad (1.10)$$

We also require an invertibility condition on the matrix  $\eta_i(x_j)$

$$\forall x_1 \dots x_{\bar{k}} \text{ distinct, the } \bar{k} \text{ vectors of } \mathbb{R}^k$$

$$(\eta_i(x_1))_{i=1 \dots k}, \dots, (\eta_i(x_{\bar{k}}))_{i=1 \dots k} \text{ are linearly independent} \quad (1.11)$$

Assumption (1.11) holds for generical choices of the functions  $\eta_i$ . Assumption (1.10) is more restrictive, specifically if  $k = 1$ , since it basically means that  $\eta_1$  is monotone.

Note however that sometimes, one may have uniqueness of the environmental variables whereas the population measure is not unique (and (1.11) is violated). Most of our method would remain valid in such a case. This is the situation of a single resource, where almost nothing is required [2]. Nevertheless in more general situations, the conditions for which this kind of property holds are not currently identified. Even though (1.10) and (1.11) are likely more restrictive than would be strictly necessary, they have the advantage of being explicit and of enabling us to reconstruct the limit of the original  $u_\varepsilon$ . Moreover from the point of view of the biological interpretation, it is interesting to have a population equilibrium.

It may be hard to check (1.10) in specific models, but it is at least satisfied in large classes of parameters, for example if the derivatives  $\eta_i^{(k)}$  are positive (or negative) for all  $i = 1, \dots, k$ , in which case  $\bar{k} = k$ . For instance in the case  $k = 2$ , (1.10) and (1.11) are ensured by the convexity (or concavity) of  $\eta_1$  and  $\eta_2$ .

This issue of uniqueness of the environmental variables is of course easier if the environment is larger. In particular, it would be considerably simplified by considering a model with (implicitly) infinitely many resources, like the Lotka-Volterra model

$$\partial_t u_\varepsilon(t, x) = \frac{1}{\varepsilon} \left( r(x) - \int_{\mathbb{R}} b(x-y) u_\varepsilon(t, y) dy \right) u_\varepsilon(t, x) + M_\varepsilon(u_\varepsilon)(t, x),$$

with an even function  $b$ . In this case, assumptions (1.10) and (1.11) should be replaced by the much simpler assumption that  $b$  as an operator is positive, i.e. that the Fourier transform of  $b$  is positive (see [9]). However, we chose to study here a model with explicit resources, as these are easier to interpret in terms of biological modeling.

Now, we may uniquely define the metastable measure associated with a set  $\omega$  by

**Proposition 1.1** *For any closed  $\omega \subset \mathbb{R}$ , there exists a unique finite nonnegative measure  $\mu(\omega)$  satisfying*

*i)  $\text{supp } \mu \subset \omega$*

*ii) denoting  $\bar{I}_i(\mu) = c_i / (1 + \int \eta_i(x) d\mu(x))$ ,*

$$\sum_{i=1}^k \bar{I}_i(\mu) \eta_i(x) - 1 \leq 0 \text{ in } \omega, \quad \sum_{i=1}^k \bar{I}_i(\mu) \eta_i(x) - 1 = 0 \text{ on } \text{supp } \mu.$$

Now the limiting  $I_i$  in (1.8) are directly obtained by

$$I_i(t) = \bar{I}_i(\mu(\{x \in \mathbb{R} : \varphi(t, x) = 0\})), \quad (1.12)$$

i.e.  $\varphi$  is solution to the (closed) Hamilton-Jacobi equation

$$\partial_t \varphi = \sum_{i=1}^k \bar{I}_i(\mu(\{\varphi(t, \cdot) = 0\})) \eta_i(x) - 1 + H(\partial_x \varphi). \quad (1.13)$$

We prove

**Theorem 1.1** *Assume  $K \in C_c^\infty(\mathbb{R})$ ,  $\int_{\mathbb{R}} zK(z) dz = 0$ , (1.9), (1.10), (1.11), that the initial data  $u_\varepsilon(t=0) > 0$  or  $\varphi_\varepsilon(t=0)$  are  $C^2$ , satisfy*

$$\sup_{\varepsilon} \int_{\mathbb{R}} u_\varepsilon(t=0, x) dx < \infty, \quad \sup_{\varepsilon} \|\partial_x \varphi_\varepsilon(t=0, \cdot)\|_{L^\infty(\mathbb{R})} < \infty, \quad (1.14)$$

$$\inf_{\varepsilon} \inf_{x \in \mathbb{R}} \partial_{xx} \varphi(t=0, x) > -\infty, \quad (1.15)$$



and that  $\varphi_\varepsilon(t = 0, \cdot)$  converges to a function  $\varphi^0$  for the norm  $\|\cdot\|_{W^{1,\infty}(\mathbb{R})}$ . Then up to the extraction of a subsequence in  $\varepsilon$ ,  $\varphi_\varepsilon$  converges to some  $\varphi$  uniformly on any compact subset of  $[0, T] \times \mathbb{R}$  and in  $W^{1,p}([0, T] \times K)$  for any  $T > 0$ ,  $p < \infty$  and any compact  $K$ . In particular,  $\varphi$  is continuous. The function  $I_i^\varepsilon$  converges to  $I_i$  in  $L^p([0, T])$  for any  $T > 0$ ,  $p < \infty$ , where  $I_i$  is defined from  $\varphi$  as in (1.12), and  $I_i$  is approximately right-continuous for **all**  $t \geq 0$ . The function  $\varphi$  is a solution to (1.8) almost everywhere in  $t, x$  with initial condition  $\varphi(t = 0, \cdot) = \varphi^0$ . Moreover if one defines  $\psi = \varphi - \sum_{i=1}^k \int_0^t I_i(s) ds \eta_i(x)$ , then  $\psi$  is a viscosity solution to

$$\partial_t \psi(t, x) = H \left( \partial_x \psi + \sum_{i=1}^k \int_0^t I_i(s) ds \eta_i(x) \right). \quad (1.16)$$

We recall that a function  $f$  on  $[0, +\infty)$  is approximately right-continuous at  $t \geq 0$  if  $t$  is a point of Lebesgue right-continuity of  $f$ , i.e.

$$\lim_{s \rightarrow 0} \frac{1}{s} \int_t^{t+s} |f(\theta) - f(t)| d\theta = 0.$$

Notice that, under the assumptions of Theorem 1.1,  $\varphi_\varepsilon(t = 0, x) \rightarrow -\infty$  when  $x \rightarrow \pm\infty$  since  $\int_{\mathbb{R}} u_\varepsilon < \infty$  and  $\varphi_\varepsilon$  is uniformly Lipschitz. Be also careful that we assume  $\|\varphi_\varepsilon(t = 0, \cdot) - \varphi^0\|_{W^{1,\infty}(\mathbb{R})} \rightarrow 0$  even though  $\varphi_\varepsilon(t = 0)$  (and thus  $\varphi^0$ ) is not bounded.

From a practical point of view, computing the solution  $u_\varepsilon$  of Eq. (1.1) is often too costly for small  $\varepsilon$ . This result allows to approximate the population density  $u_\varepsilon$  for small  $\varepsilon$  by the simpler  $\mu(\{\varphi(t, \cdot) = 0\})$ , where  $\varphi$  may be obtained by a discretization of (1.13), in the fashion of those done in [16]. Rigorous numerical analysis of this kind of Hamilton-Jacobi equations is however still very preliminary.

On a more theoretical level, this theorem justifies in simple cases the classical view of evolution (population at local equilibrium at every time). It also indicates that for small  $\varepsilon$ , the behaviour of the population is determined by a closed equation which means that evolution remains deterministic and stable in the parameter  $\varepsilon$ .

In the proofs below,  $C$  denotes a numerical constant which may change from line to line.

## 2 Proof of Prop. 1.1

### 2.1 Uniqueness

Assume that two measures  $\mu_1$  and  $\mu_2$  satisfy both points of Prop. 1.1. We first prove that they induce the same resources  $\bar{I}_i$  and then conclude that they are equal.

*1st step: Uniqueness of the  $\bar{I}_i$ .* The argument here is essentially an adaptation of [21]. First note that

$$\int_{\mathbb{R}} \left( \sum_{i=1}^k \bar{I}_i(\mu_1) \eta_i(x) - 1 \right) d\mu_2 + \int_{\mathbb{R}} \left( \sum_{i=1}^k \bar{I}_i(\mu_2) \eta_i(x) - 1 \right) d\mu_1 \leq 0, \quad (2.1)$$

since  $\mu_1$  and  $\mu_2$  are non negative and by the point *ii*,  $\sum_{i=1}^k \bar{I}_i(\mu_j) \eta_i(x) - 1$  is non positive on  $\omega$  for  $j = 1, 2$ .

On the other hand since  $\sum_{i=1}^k \bar{I}_i(\mu_j) \eta_i - 1$  vanishes on the support of  $\mu_j$ , one has for instance

$$\begin{aligned} \int_{\mathbb{R}} \left( \sum_{i=1}^k \bar{I}_i(\mu_1) \eta_i - 1 \right) d\mu_2 &= \int_{\mathbb{R}} \left( \sum_{i=1}^k (\bar{I}_i(\mu_1) - \bar{I}_i(\mu_2)) \eta_i \right) d\mu_2 \\ &= \sum_{i=1}^k (\bar{I}_i(\mu_1) - \bar{I}_i(\mu_2)) \int_{\mathbb{R}} \eta_i d\mu_2 \\ &= \sum_{i=1}^k (\bar{I}_i(\mu_1) - \bar{I}_i(\mu_2)) (c_i / \bar{I}_i(\mu_2) - 1), \end{aligned}$$

by the definition of  $\bar{I}_i(\mu_2)$ .

Since one has

$$\sum_{i=1}^k c_i (\bar{I}_i(\mu_1) - \bar{I}_i(\mu_2)) (1/\bar{I}_i(\mu_2) - 1/\bar{I}_i(\mu_1)) = \sum_{i=1}^k c_i \frac{(\bar{I}_i(\mu_1) - \bar{I}_i(\mu_2))^2}{\bar{I}_i(\mu_1) \bar{I}_i(\mu_2)} \geq 0.$$

one deduces from (2.1) that

$$\bar{I}_i(\mu_1) = \bar{I}_i(\mu_2), \quad i = 1 \dots k. \quad (2.2)$$

*2nd step: Uniqueness of  $\mu$ .* It is not possible to deduce that  $\mu_1 = \mu_2$  directly from (2.2). This degeneracy (the possibility of having several equilibrium

measures, all corresponding to the same environment) is the reason why we require additional assumptions on the  $\eta_i$ .

First of all by Assumption (1.10), point  $i$  and thanks to (2.2), we know that  $\mu_1$  and  $\mu_2$  are both supported on a set consisting of at most  $\bar{k}$  points  $\{x_1, \dots, x_{\bar{k}}\}$ , which are the roots of  $\sum_i \bar{I}_i(\mu_1)\eta_i(x) - 1 = \sum_i \bar{I}_i(\mu_2)\eta_i(x) - 1$  (possibly completed by arbitrary distinct points if this function has less than  $\bar{k}$  roots). Therefore one may write

$$\mu_j = \sum_{l=1}^{\bar{k}} \alpha_l^j \delta_{x_l}.$$

Now (2.2) tells that  $\int \eta_i d\mu_1 = \int \eta_i d\mu_2$  which means that

$$\sum_{l=1}^{\bar{k}} \alpha_l^1 \eta_i(x_l) = \sum_{l=1}^{\bar{k}} \alpha_l^2 \eta_i(x_l), \quad \forall i = 1 \dots k.$$

To conclude it remains to use condition (1.11) and get that  $\alpha_l^1 = \alpha_l^2$ .

### 2.1.1 Existence

The basic idea to get existence is quite simple: Solve the equation<sup>1</sup>

$$\partial_t \nu = \left( \sum_{i=1}^k \bar{I}_i(\nu) \eta_i(\cdot) - 1 \right) \nu, \quad (2.3)$$

and obtain the equilibrium measure  $\mu$  as the limit of  $\nu(t)$  as  $t \rightarrow +\infty$ .

This is done by considering the entropy

$$\begin{aligned} L(\nu) &= \sum_{i=1}^k c_i \log \bar{I}_i(\nu) + \int d\nu \\ &= \sum_{i=1}^k c_i \log c_i - \sum_{i=1}^k c_i \log \left( 1 + \int \eta_i d\nu \right) + \int d\nu. \end{aligned} \quad (2.4)$$

---

<sup>1</sup>Existence and uniqueness are trivial for (2.3), for example by Cauchy-Lipschitz theorem in the set of finite positive measures equipped with the total variation norm.

As  $-\log$  is convex and  $\eta_i \geq 0$ , then  $L$  itself is a convex function of  $\nu$ . Moreover if  $\nu(t)$  solves (2.3), one has

$$\frac{d}{dt}L(\nu(t)) = - \int \left( \sum_{i=1}^k \bar{I}_i(\nu) \eta_i(x) - 1 \right)^2 d\nu. \quad (2.5)$$

Therefore one expects the limit of  $\nu$  and the equilibrium measure we are looking for to be the minimum of  $L$ .

Since the  $\eta_i$  are bounded, one finds

$$L(\nu) \geq -C + c \int d\nu,$$

for two numerical constants  $C$  and  $c$ . Consequently  $L$  is bounded from below on  $M_+^1(\omega)$  the set of nonnegative Radon measures on  $\omega$ . In addition, one may restrict to a bounded subset of  $M_+^1(\omega)$  to compute the infimum of  $L$ . As any ball of  $M_+^1(\omega)$  is compact for the weak-\* topology (dual of continuous functions with compact support),  $L$  attains its infimum, or

$$M_0 = \{ \nu \in M_+^1(\omega), L(\nu) \leq L(\nu') \forall \nu' \in M_+^1(\omega) \} \neq \emptyset.$$

Now take any  $\mu \in M_0$  then take  $\nu$  the solution to (2.3) with  $\nu(t=0) = \mu$ .  $L(\nu)$  is non increasing and since it is already at a minimum initially, it is necessarily constant. By (2.5), this means that

$$\sum_{i=1}^k \bar{I}_i(\mu) \eta_i - 1 = 0 \text{ on } \text{supp } \mu.$$

Hence  $\mu$  is in fact a stationary solution to (2.3) and it satisfies point  $i$  and the second part of point  $ii$  of Prop. 1.1. Note by the way that the uniqueness argument in fact tells that there is a unique element in  $M_0$ .

It only remains to check the first part of point  $ii$ . By contradiction assume that there exists a point  $x_0 \in \omega$  s.t.

$$\sum_{i=1}^k \bar{I}_i(\mu) \eta_i(x_0) - 1 > 0.$$

Let  $\alpha > 0$  and define  $\nu_\alpha = \mu + \alpha\delta_{x_0} \in M_+^1(\omega)$ . Now compute

$$\begin{aligned}
L(\nu_\alpha) &= \int d\mu + \alpha - \sum_{i=1}^k c_i \log \left( 1 + \int \eta_i d\mu + \alpha\eta_i(x_0) \right) + \sum_{i=1}^k c_i \log c_i \\
&= \int d\mu + \alpha - \sum_{i=1}^k c_i \left( \log \left( 1 + \int \eta_i d\mu \right) + \frac{\alpha \eta_i(x_0)}{1 + \int \eta_i d\mu} + O(\alpha^2) \right) \\
&\quad + \sum_{i=1}^k c_i \log c_i \\
&= L(\mu) - \alpha \left( \sum_{i=1}^k \bar{I}_i(\mu) \eta_i(x_0) - 1 \right) + O(\alpha^2).
\end{aligned}$$

Thus  $L(\nu_\alpha) < L(\mu)$  for  $\alpha$  small enough which is impossible as  $\mu$  is an absolute minimum of  $L$ .

Consequently the first part of *ii* is satisfied and the proof of Prop. 1.1 complete.

### 3 Proof of Theorem 1.1

#### 3.1 A priori estimates for Eq. (1.5)

We denote by  $BV_{\text{loc}}(\mathbb{R})$  the set of functions on  $\mathbb{R}$  with bounded variation on any compact subset of  $\mathbb{R}$ , by  $M^1(\omega)$  the set of signed Radon measures on the subset  $\omega$  of  $\mathbb{R}$  equipped with the total variation norm.

We show the following estimates on the solution to (1.5)

**Lemma 3.1** *Let  $\varphi_\varepsilon$  be a solution to (1.5) with initial data  $\varphi_\varepsilon^0$  such that  $\int_{\mathbb{R}} e^{\varphi_\varepsilon^0(x)/\varepsilon} dx < \infty$ ,  $\partial_x \varphi_\varepsilon^0 \in L^\infty(\mathbb{R})$  and  $\partial_{xx} \varphi_\varepsilon^0$  uniformly bounded from below. Then for any  $T > 0$*

$$\begin{aligned}
&\|\partial_t \varphi_\varepsilon\|_{L^\infty([0,T] \times \mathbb{R})} + \|\partial_x \varphi_\varepsilon\|_{L^\infty([0,T], BV_{\text{loc}}(\mathbb{R}) \cap L^\infty(\mathbb{R}))} + \|\partial_{tx} \varphi_\varepsilon\|_{L^\infty([0,T], M^1)} \leq C_T, \\
&\forall t \leq T, x \in \mathbb{R}, \quad \partial_{xx} \varphi_\varepsilon(t, x) \geq -C_T, \quad H_\varepsilon(\varphi_\varepsilon) \geq -C_T \varepsilon,
\end{aligned}$$

$$\forall t \leq T, \quad \int_{\mathbb{R}} u_\varepsilon(t, x) dx \leq C_T, \quad \varphi_\varepsilon(t, x) \leq C_T \varepsilon \log 1/\varepsilon.$$

where  $C_T$  only depends on the time  $T$ ,  $\int_{\mathbb{R}} e^{\varphi_\varepsilon^0(x)/\varepsilon} dx$ ,  $\|\partial_x \varphi_\varepsilon^0\|_{L^\infty(\mathbb{R})}$  and the infimum of  $\partial_{xx} \varphi_\varepsilon^0(x)$ .

**Proof.** We start with the easy bound on the total mass.

*Step 0: Bound on the total mass.* First notice that because of (1.9), there exists  $R > 0$  s.t.

$$\forall |x| > R, \quad \sum_{i=1}^k \eta_i(x) \leq 1/2.$$

Let  $\psi$  be a regular test function with support in  $|x| > R$ , taking values in  $[0, 1]$  and equal to 1 on  $|x| > R+1$ . Using the fact that  $I_i^\varepsilon(t) \leq 1$ , we compute

$$\begin{aligned} \frac{d}{dt} \int_{\mathbb{R}} \psi(x) u_\varepsilon(t, x) dx &\leq -\frac{1}{2\varepsilon} \int_{\mathbb{R}} \psi(x) u_\varepsilon(t, x) dx \\ &\quad + \frac{1}{\varepsilon} \int_{\mathbb{R}^2} K(z) (\psi(x - \varepsilon z) - \psi(x)) u_\varepsilon(t, x) dz dx \\ &\leq -\frac{1}{2\varepsilon} \int_{\mathbb{R}} \psi(x) u_\varepsilon(t, x) dx + C \int_{\mathbb{R}} u_\varepsilon(t, x) dx. \end{aligned}$$

On the other hand as each  $\eta_i > 0$ , one has for some constant  $C$

$$I_i^\varepsilon(t) = \frac{c_i}{1 + \int \eta_i u_\varepsilon dx} \leq \frac{C}{1 + \int (1 - \psi) u_\varepsilon dx}.$$

Therefore with the same kind of estimate

$$\begin{aligned} \frac{d}{dt} \int_{\mathbb{R}} (1 - \psi(x)) u_\varepsilon(t, x) dx &\leq C \int_{\mathbb{R}} u_\varepsilon(t, x) dx \\ &\quad + \frac{1}{\varepsilon} \left( \frac{C}{1 + \int (1 - \psi) u_\varepsilon dx} - 1 \right) \int_{\mathbb{R}} (1 - \psi(x)) u_\varepsilon(t, x) dx. \end{aligned}$$

Summing the two

$$\begin{aligned} \frac{d}{dt} \int_{\mathbb{R}} u_\varepsilon(t, x) dx &\leq \frac{1}{\varepsilon} \left( \frac{C}{1 + \int (1 - \psi) u_\varepsilon dx} - 1 \right) \int_{\mathbb{R}} (1 - \psi(x)) u_\varepsilon(t, x) dx \\ &\quad - \frac{1}{2\varepsilon} \int_{\mathbb{R}} \psi(x) u_\varepsilon(t, x) dx + C \int_{\mathbb{R}} u_\varepsilon(t, x) dx. \end{aligned}$$

Since the sum of the first two terms of the r.h.s. is negative if  $\int u_\varepsilon$  is larger than a constant independent of  $\varepsilon$ , this shows that  $\int u_\varepsilon(t, x) dx$  remains uniformly bounded on any finite time interval.

*Step 1: Bound on  $\partial_x \varphi_\varepsilon$ .* This is a classical bound for solutions to Hamilton-Jacobi equations. Here we still have to check that it remains true uniformly at the  $\varepsilon$  level. Compute

$$\begin{aligned} \partial_t \partial_x \varphi_\varepsilon &= \sum_{i=1}^k I_i^\varepsilon(t) \eta'_i(x) \\ &+ \int K(z) e^{\frac{\varphi_\varepsilon(t, x + \varepsilon z) - \varphi_\varepsilon(t, x)}{\varepsilon}} \frac{\partial_x \varphi_\varepsilon(t, x + \varepsilon z) - \partial_x \varphi_\varepsilon(t, x)}{\varepsilon} dz. \end{aligned} \quad (3.1)$$

We first observe that, as  $I_i^\varepsilon \in [0, \max_i c_i]$  and  $\sum_i |\eta'_i(x)| \leq \bar{\eta}(x)$

$$\begin{aligned} |\partial_t \partial_x \varphi_\varepsilon| &\leq \max_i c_i \bar{\eta}(x) + \frac{2}{\varepsilon} \int_{\mathbb{R}} K(z) e^{|z| \|\partial_x \varphi_\varepsilon(t, \cdot)\|_{L^\infty(\mathbb{R})}} \|\partial_x \varphi_\varepsilon(t, \cdot)\|_{L^\infty(\mathbb{R})} dz \\ &\leq \frac{C}{\varepsilon} e^{C \|\partial_x \varphi_\varepsilon(t, \cdot)\|_{L^\infty(\rho)}}, \end{aligned}$$

since  $K$  has compact support. This entails

$$\|\partial_x \varphi_\varepsilon(t, \cdot)\|_{L^\infty(\mathbb{R})} \leq \|\partial_x \varphi_\varepsilon^0\|_{L^\infty(\mathbb{R})} + \frac{C}{\varepsilon} \int_0^t e^{C \|\partial_x \varphi_\varepsilon(s, \cdot)\|_{L^\infty(\mathbb{R})}} ds,$$

from which easily follows that  $\partial_x \varphi_\varepsilon \in L^\infty([0, t_\varepsilon], \mathbb{R})$  for some  $t_\varepsilon > 0$ , which may (for the moment) depend on  $\varepsilon$ .

Now we use the classical maximum principle. Fix  $t \in [0, T]$  such that  $C_{\varepsilon, t} := \|\partial_x \varphi_\varepsilon(t, \cdot)\|_{L^\infty(\mathbb{R})} < \infty$ . For any  $x \in \mathbb{R}$  such that  $\partial_x \varphi_\varepsilon(t, x) > \sup_y \partial_x \varphi_\varepsilon(t, y) - \alpha$ , where the constant  $\alpha > 0$  will be specified later, we have

$$\partial_t \partial_x \varphi_\varepsilon(t, x) \leq \max_i c_i \bar{\eta}(x) + \int_{\mathbb{R}} K(z) e^{|z| C_{t, \varepsilon}} \frac{\alpha}{\varepsilon} dz \leq C \left(1 + \frac{\alpha}{\varepsilon} e^{C C_{t, \varepsilon}}\right).$$

Therefore, choosing  $\alpha = \varepsilon e^{-C C_{t, \varepsilon}}$ , we obtain

$$\frac{d}{dt} \sup_x \partial_x \varphi_\varepsilon(t, x) \leq C,$$

for a constant  $C$  independent of  $t < t_\varepsilon$  and of  $\varepsilon$ . Using a similar argument for the minimum, we deduce that  $t_\varepsilon > T$  and that  $\partial_x \varphi_\varepsilon$  is bounded on  $[0, T] \times \mathbb{R}$  by a constant depending only on  $T$  and  $\|\partial_x \varphi_\varepsilon^0\|_{L^\infty(\mathbb{R})}$ .

*Step 2: First bound on  $H_\varepsilon(\varphi_\varepsilon)$  and bounds on  $\partial_t \varphi_\varepsilon$  and  $\varphi_\varepsilon$ .* Simply note that

$$\begin{aligned} - \int_{\mathbb{R}} K(z) dz &\leq H_\varepsilon(\varphi_\varepsilon(t))(x) = \int_{\mathbb{R}} K(z) e^{\frac{\varphi_\varepsilon(t, x + \varepsilon z) - \varphi_\varepsilon(t, x)}{\varepsilon}} dz - \int_{\mathbb{R}} K(z) dz \\ &\leq \int K(z) e^{|z| \|\partial_x \varphi_\varepsilon\|_{L^\infty([0, T], \mathbb{R})}} dz \leq C. \end{aligned}$$

Consequently, directly from Eq. (1.5),

$$|\partial_t \varphi_\varepsilon| \leq \max_i c_i \bar{\eta}(x) + C,$$

hence ending the proof of the full Lipschitz bound on  $\varphi_\varepsilon$ .

To get the upper bound on  $\varphi_\varepsilon$ , simply note that because of the uniform Lipschitz bound on  $\varphi_\varepsilon$

$$\varphi_\varepsilon(t, y) \geq \varphi_\varepsilon(t, x) - C_T |y - x|,$$

so

$$\int_{\mathbb{R}} u_\varepsilon(t, y) dy \geq \int_{\mathbb{R}} e^{\varphi_\varepsilon(t, x)/\varepsilon} e^{-C_T |y-x|/\varepsilon} dy \geq 2C_T^{-1} \varepsilon e^{\varphi_\varepsilon(t, x)/\varepsilon}.$$

Hence the bound on the total mass yields that  $\varphi_\varepsilon \leq C_T \varepsilon \log 1/\varepsilon$ .

*Step 3: BV bound on  $\partial_x \varphi_\varepsilon$ .* As for  $\partial_x \varphi_\varepsilon$ , we begin with a maximum (actually, minimum) principle. First from (1.5)

$$\begin{aligned} \partial_t \partial_{xx} \varphi_\varepsilon &\geq -\bar{\eta}(x) + \int_{\mathbb{R}} K(z) e^{\frac{\varphi_\varepsilon(t, x+\varepsilon z) - \varphi_\varepsilon(t, x)}{\varepsilon}} \frac{\partial_{xx} \varphi_\varepsilon(t, x + \varepsilon z) - \partial_{xx} \varphi_\varepsilon(t, x)}{\varepsilon} dz \\ &\quad + \int_{\mathbb{R}} K(z) e^{\frac{\varphi_\varepsilon(t, x+\varepsilon z) - \varphi_\varepsilon(t, x)}{\varepsilon}} \frac{(\partial_x \varphi_\varepsilon(t, x + \varepsilon z) - \partial_x \varphi_\varepsilon(t, x))^2}{\varepsilon} dz. \end{aligned}$$

The last term is of course non negative and so with the same argument as before, we get

$$\frac{d}{dt} \inf_x \partial_{xx} \varphi_\varepsilon(t, x) \geq -C,$$

where  $C$  does not depend on  $\varepsilon$ . This proves the uniform lower bound on  $\partial_{xx} \varphi_\varepsilon$ . On the other hand, for any measurable subset  $A$  of  $[x_1, x_2]$ ,

$$\begin{aligned} \int_{x_1}^{x_2} (\mathbb{I}_{x \in A} - \mathbb{I}_{x \notin A}) \partial_{xx} \varphi_\varepsilon(t, x) dx &= \int_{x_1}^{x_2} \partial_{xx} \varphi_\varepsilon(t, x) dx \\ &\quad - 2 \int_{x_1}^{x_2} \partial_{xx} \varphi_\varepsilon(t, x) \mathbb{I}_{x \notin A} dx \\ &\leq \partial_x \varphi_\varepsilon(t, x_2) - \partial_x \varphi_\varepsilon(t, x_1) + C |x_2 - x_1| \leq 2 \|\partial_x \varphi_\varepsilon\|_{L^\infty([0, T], \mathbb{R})} + C |x_2 - x_1|. \end{aligned}$$

This indeed shows that  $\partial_{xx} \varphi_\varepsilon(t, \cdot)$  belongs to  $M^1([x_1, x_2])$  with total variation norm less than  $2 \|\partial_x \varphi_\varepsilon\|_{L^\infty([0, T], \mathbb{R})} + C |x_2 - x_1|$ . Thus,  $\partial_{xx} \varphi_\varepsilon$  belongs to the space  $L^\infty([0, T], M^1(\mathbb{R}))$ , which entails  $\partial_x \varphi_\varepsilon \in L^\infty([0, T], BV_{\text{loc}}(\mathbb{R}))$ .



Finally, it follows from (3.1) that

$$\begin{aligned} |\partial_{tx}\varphi_\varepsilon(t, x)| &\leq \bar{\eta}(x) \max_i c_i + \int K(z) e^{|z| \|\partial_x \varphi_\varepsilon\|_{L^\infty}} \frac{|\partial_x \varphi_\varepsilon(t, x + \varepsilon z) - \partial_x \varphi_\varepsilon(t, x)|}{\varepsilon} dz \\ &\leq c \left( \bar{\eta}(x) + \int K(z) \int_0^z |\partial_{xx}\varphi_\varepsilon(t, x + \varepsilon\theta)| d\theta dz \right). \end{aligned}$$

Integrating, by Fubini

$$\int_{x_1}^{x_2} |\partial_{tx}\varphi_\varepsilon| dx \leq C \int_{x_1}^{x_2} \bar{\eta}(x) dx + C \int_{x_1 - \varepsilon\rho}^{x_2 + \varepsilon\rho} |\partial_{xx}\varphi_\varepsilon| dx,$$

where  $\rho$  is such that the support of  $K$  is included in the ball centered at 0 of radius  $\rho$ . This ends the proof of all the bounds on the derivatives of  $\varphi_\varepsilon$ .

*Conclusion.* It only remains to show the sharp lower bound on  $H_\varepsilon(\varphi_\varepsilon)$ . Let us write

$$H_\varepsilon(\varphi_\varepsilon) \geq \int_{\mathbb{R}} K(z) \exp\left(\int_0^1 z \partial_x \varphi_\varepsilon(t, x + \theta z \varepsilon) d\theta\right) dz - \int_{\mathbb{R}} K(z) dz.$$

The *BV* bound on  $\partial_x \varphi_\varepsilon$  shows that this function admits right and left limits at all  $x \in \mathbb{R}$ . Let us denote  $\partial_x \varphi_\varepsilon(t, x^+)$  the limit on the right and  $\partial_x \varphi_\varepsilon(t, x^-)$  the limit on the left. As  $\partial_{xx}\varphi_\varepsilon$  is bounded from below, we know in addition that

$$\forall x, \quad \partial_x \varphi_\varepsilon(t, x^+) \geq \partial_x \varphi_\varepsilon(t, x^-).$$

By differentiating once more

$$\begin{aligned} \int_0^1 z \partial_x \varphi_\varepsilon(t, x + \theta z \varepsilon) d\theta &\geq z \partial_x \varphi_\varepsilon(t, x^+) \\ &\quad + \int_0^1 z \int_0^1 \theta z \varepsilon \partial_{xx}\varphi_\varepsilon(t, x + \theta' \theta z \varepsilon) d\theta' d\theta \\ &\geq z \partial_x \varphi_\varepsilon(t, x^+) - C \varepsilon z^2, \end{aligned}$$

again as  $\partial_{xx}\varphi_\varepsilon$  is bounded from below. Finally

$$\begin{aligned} H_\varepsilon(\varphi_\varepsilon) &\geq \int_{\mathbb{R}} K(z) \exp(z \partial_x \varphi_\varepsilon(t, x^+) - C \varepsilon z^2) dz - \int_{\mathbb{R}} K(z) dz \\ &\geq H(\partial_x \varphi_\varepsilon(t, x^+)) - C \varepsilon, \end{aligned}$$

where  $H$  is defined as in (1.7) and since  $K$  is compactly supported. Because we assumed that  $\int_{\mathbb{R}} z K(z) dz = 0$ , we have  $H(p) \geq 0$  for any  $p$ , which ends the proof of Lemma 3.1.

### 3.2 Passing to the limit in $\varphi_\varepsilon$

From the assumptions in Theorem 1.1, Lemma 3.1 gives uniform bounds on  $\varphi_\varepsilon$ .

Therefore up to an extraction in  $\varepsilon$  (still denoted with  $\varepsilon$ ), there exists a function  $\varphi$  on  $[0, T] \times \mathbb{R}$  such that  $\partial_t \varphi \in L^\infty([0, T] \times \mathbb{R})$ ,  $\partial_x \varphi \in L^\infty([0, T], BV_{\text{loc}} \cap L^\infty(\mathbb{R}))$ ,  $\partial_{tx} \varphi \in L^\infty([0, T], M_{\text{loc}}^1(\mathbb{R}))$  and  $\partial_{xx} \varphi$  uniformly lower bounded on  $[0, T] \times \mathbb{R}$ , satisfying

$$\begin{aligned} \varphi_\varepsilon &\longrightarrow \varphi \quad \text{uniformly in } C(K) \text{ for any compact } K \text{ of } [0, T] \times \mathbb{R}, \\ \partial_x \varphi_\varepsilon &\longrightarrow \partial_x \varphi \quad \text{in any } L_{\text{loc}}^p([0, T], \mathbb{R}), \quad p < \infty. \end{aligned} \quad (3.2)$$

The first convergence follows from Arzéla-Ascoli theorem. For the second convergence, observe that  $\|\partial_x \varphi_\varepsilon\|_{L^\infty([0, T], BV_{\text{loc}}(\mathbb{R}))} + \|\partial_{tx} \varphi_\varepsilon\|_{L^\infty([0, T], M^1(\mathbb{R}))} \leq C_T$  implies that  $\partial_x \varphi_\varepsilon$  is uniformly bounded in  $L^\infty([0, T] \times \mathbb{R}) \cap BV_{\text{loc}}([0, T] \times \mathbb{R})$ . The convergence in  $L_{\text{loc}}^p$  follows by compact embedding. We also have  $\varphi \leq 0$  since otherwise the uniform bound on  $\int_{\mathbb{R}} u_\varepsilon(t, x) dx$  would be contradicted. As the  $I_i^\varepsilon$  are bounded, it is possible to extract weak-\* converging subsequences (still denoted with  $\varepsilon$ ) to some  $I_i(t)$ .

Now, we write again

$$H_\varepsilon(\varphi_\varepsilon) = \int_{\mathbb{R}} K(z) \left( \exp \left( \int_0^1 z \partial_x \varphi_\varepsilon(t, x + \varepsilon z \theta) d\theta \right) - 1 \right) dz.$$

From the  $L^\infty$  bound on  $\partial_x \varphi_\varepsilon$  and its strong convergence, one deduces that

$$H_\varepsilon(\varphi_\varepsilon) \longrightarrow H(\partial_x \varphi) \text{ in } L_{\text{loc}}^1. \quad (3.3)$$

Therefore one may pass to the limit in (1.5) and obtain (1.8) (for the moment in the sense of distribution; the equality *a.e.* will follow from the convergence of  $I_i^\varepsilon$  in  $L^p([0, T])$ , proved below).

In addition by following [16] or [2], one may easily show that  $\psi(t, x) = \varphi(t, x) - \sum_{i=1}^k \int_0^t I_i(s) ds \eta_i(x)$  is a viscosity solution to (1.16). We refer the reader to these references for this technical part.

It remains to obtain (1.12), the approximate right-continuity of  $I_i$  for *all* time  $t$  and the convergence of  $I_i^\varepsilon$  to  $I_i$  in  $L^p([0, T])$  for  $p < \infty$ . This requires some sort of uniform continuity on the  $I_i^\varepsilon$  which is the object of the rest of the proof.

### 3.3 Continuity in time for the $I_i^\varepsilon$

First of all note that, as suggested by the simulations of [16], there are examples where the  $I_i$  have jumps in time at the limit. So we will only be able to prove their right-continuity.

This regularity in time comes from the stability of the equilibrium defined through (1.12) and Prop. 1.1. Therefore let us define

$$\bar{I}_i(t) = \bar{I}_i(\mu(\{\varphi(t, \cdot) = 0\})),$$

where  $\bar{I}_i$  and  $\mu$  are given by Prop. 1.1 and  $\varphi$  is the uniform limit of  $\varphi_\varepsilon$  as taken in the previous subsection.

Our first goal is the following result.

**Lemma 3.2** *For any fixed  $s$ , there exist functions  $\sigma_s, \tilde{\sigma} \in C(\mathbb{R}_+)$  with  $\sigma_s(0) = \tilde{\sigma}(0) = 0$  s.t.*

$$\int_s^t |I_i^\varepsilon(r) - \bar{I}_i(s)|^2 dr \leq (t - s) \sigma_s(t - s) + \tilde{\sigma}(\varepsilon).$$

**Remark.** Of course the whole point is that  $\sigma_s$  is uniform in  $\varepsilon$ . It is also crucial for the following that  $\tilde{\sigma}$  does not depend on  $s$ .

#### 3.3.1 Proof of Lemma 3.2

*Step 0:  $\varphi$  has compact level sets.*

Observe that  $\varphi_\varepsilon(t = 0, x) \rightarrow -\infty$  when  $x \rightarrow \pm\infty$  since  $\int_{\mathbb{R}} u_\varepsilon(t = 0, x) dx < \infty$  and  $\partial_x \varphi(t = 0)$  is bounded. Because of the uniform convergence of  $\varphi_\varepsilon(t = 0)$  to  $\varphi^0$  on  $\mathbb{R}$ , one deduces that  $\varphi^0(x) \rightarrow -\infty$  when  $x \rightarrow \pm\infty$ .

Since  $\partial_x \varphi \in L^\infty([0, T], \mathbb{R})$  and  $I_i(t) \in [0, \max_i c_i]$ , it follows from (1.8) that  $\partial_t \varphi \in L^\infty([0, T], \mathbb{R})$  and thus  $\varphi(t, x) \rightarrow -\infty$  when  $x \rightarrow \pm\infty$  for all  $t \geq 0$ .

Therefore, the set

$$\Omega := \{(t, x) \in [0, T] \times \mathbb{R} : \varphi(t, x) \geq -1\}$$

is compact.

*Step 1: One basic property of  $\{\varphi = 0\}$ .*

Let us start with the following crucial observation

$$\begin{aligned} \forall s, \exists \tau_s \in C(\mathbb{R}_+) \text{ with } \tau_s(0) = 0, \text{ s.t. } \forall t \geq s, \\ \forall x \in \{\varphi(t, \cdot) = 0\}, \exists y \in \{\varphi(s, \cdot) = 0\} \text{ with } |y - x| \leq \tau_s(t - s). \end{aligned} \quad (3.4)$$

This is a sort of semi-continuity for  $\{\varphi = 0\}$ . It is proved very simply by contradiction. If it were not true, then

$$\begin{aligned} & \exists s, \exists \tau_0 > 0, \exists t_n \rightarrow s, t_n \geq s, \exists y_n \in \{\varphi(t_n, \cdot) = 0\}, \\ & d(y_n, \{\varphi(s, \cdot) = 0\}) \geq \tau_0, \end{aligned}$$

where  $d(y, \omega) = \inf_{x \in \omega} |x - y|$  is the usual distance.

Since all the  $y_n$  belong to the compact set  $\Omega$  of Step 0, we can extract a converging subsequence  $y_n \rightarrow y$ . As  $\varphi$  is continuous,  $\varphi(s, y) = 0$  or  $y \in \{\varphi(s, \cdot) = 0\}$ . On the other hand one would also have  $d(y, \{\varphi(s, \cdot) = 0\}) \geq \tau_0$  which is contradictory.

*Step 2: The functional.*

Denote

$$\mu_s = \mu(\{\varphi(s, \cdot) = 0\}),$$

as given by Prop. 1.1. We look at the evolution of

$$F_\varepsilon(t) = \int_{\mathbb{R}} \log u_\varepsilon(t, x) d\mu_s(x) = \frac{1}{\varepsilon} \int_{\mathbb{R}} \varphi_\varepsilon(t, x) d\mu_s(x),$$

for  $t \geq s$ . Compute

$$\frac{d}{dt} F_\varepsilon(t) = \frac{1}{\varepsilon} \int_{\mathbb{R}} \left( \sum_{i=1}^k I_i^\varepsilon(t) \eta_i(x) - 1 \right) d\mu_s(x) + \frac{1}{\varepsilon} \int_{\mathbb{R}} H_\varepsilon(\varphi_\varepsilon(t)) d\mu_s.$$

Now write

$$\begin{aligned} \frac{1}{\varepsilon} \int_{\mathbb{R}} \left( \sum_{i=1}^k I_i^\varepsilon(t) \eta_i(x) - 1 \right) d\mu_s(x) &= \frac{d}{dt} \int_{\mathbb{R}} u_\varepsilon(t, x) dx \\ &\quad - \frac{1}{\varepsilon} \int_{\mathbb{R}} \left( \sum_{i=1}^k I_i^\varepsilon(t) \eta_i(x) - 1 \right) (u_\varepsilon(t, x) dx - d\mu_s(x)). \end{aligned}$$

As  $\sum_{i=1}^k \bar{I}_i(s) \eta_i(x) - 1$  vanishes on the support of  $\mu_s$ ,

$$\begin{aligned} \frac{1}{\varepsilon} \int_{\mathbb{R}} \left( \sum_{i=1}^k I_i^\varepsilon(t) \eta_i(x) - 1 \right) d\mu_s(x) &= \frac{d}{dt} \int_{\mathbb{R}} u_\varepsilon(t, x) dx - \frac{A(t)}{\varepsilon} \\ &\quad - \frac{1}{\varepsilon} \int_{\mathbb{R}} \left( \sum_{i=1}^k (I_i^\varepsilon(t) - \bar{I}_i(s)) \eta_i(x) \right) (u_\varepsilon(t, x) dx - d\mu_s(x)), \end{aligned}$$

with

$$A(t) = \int_{\mathbb{R}} \left( \sum_{i=1}^k \bar{I}_i(s) \eta_i(x) - 1 \right) u_\varepsilon(t, x) dx.$$

Notice that

$$\int_{\mathbb{R}} \left( \sum_{i=1}^k (I_i^\varepsilon(t) - \bar{I}_i(s)) \eta_i(x) \right) (u_\varepsilon(t, x) dx - d\mu_s(x)) = - \sum_{i=1}^k c_i \frac{(I_i^\varepsilon(t) - \bar{I}_i(s))^2}{I_i^\varepsilon(t) \bar{I}_i(s)}.$$

So we deduce

$$\begin{aligned} \frac{1}{\varepsilon} \int_s^t \sum_{i=1}^k c_i \frac{(I_i^\varepsilon(r) - \bar{I}_i(s))^2}{I_i^\varepsilon(r) \bar{I}_i(s)} dr &= \int_{\mathbb{R}} \log \frac{u_\varepsilon(t, x)}{u_\varepsilon(s, x)} d\mu_s - \int_{\mathbb{R}} (u_\varepsilon(t, x) - u_\varepsilon(s, x)) dx \\ &\quad + \int_s^t \frac{A(r)}{\varepsilon} dr - \frac{1}{\varepsilon} \int_s^t \int_{\mathbb{R}} H_\varepsilon(\varphi_\varepsilon(r)) d\mu_s. \end{aligned} \tag{3.5}$$

*Step 3: Easy bounds.*

Lemma 3.1 tells that

$$-H_\varepsilon(\varphi_\varepsilon) \leq C_T \varepsilon.$$

The total mass stays bounded in time so

$$- \int_{\mathbb{R}} (u_\varepsilon(t, x) - u_\varepsilon(s, x)) dx \leq \int_{\mathbb{R}} (u_\varepsilon(t, x) + u_\varepsilon(s, x)) dx \leq C.$$

And furthermore

$$\begin{aligned} \int_{\mathbb{R}} \log \frac{u_\varepsilon(t, x)}{u_\varepsilon(s, x)} d\mu_s &= \frac{1}{\varepsilon} \int_{\mathbb{R}} (\varphi_\varepsilon(t, x) - \varphi_\varepsilon(s, x)) d\mu_s \\ &\leq \frac{1}{\varepsilon} \int_{\mathbb{R}} (\varphi(t, x) - \varphi(s, x)) d\mu_s + \frac{2}{\varepsilon} \|\varphi_\varepsilon - \varphi\|_{L^\infty(\Omega)}, \end{aligned}$$

where the last bound comes from the fact that, by Prop. 1.1,  $\mu_s$  is supported on  $\{\varphi(s, \cdot) = 0\} \subset \Omega$ , where  $\Omega$  is defined in Step 0. Since in addition we know that  $\varphi \leq 0$ ,

$$\int_{\mathbb{R}} \log \frac{u_\varepsilon(t, x)}{u_\varepsilon(s, x)} d\mu_s \leq \frac{2}{\varepsilon} \|\varphi_\varepsilon - \varphi\|_{L^\infty(\Omega)}.$$

Consequently we deduce from (3.5) the bound

$$\frac{1}{\varepsilon} \int_s^t \sum_{i=1}^k c_i \frac{(I_i^\varepsilon(r) - \bar{I}_i(s))^2}{I_i^\varepsilon(r) \bar{I}_i(s)} dr \leq C + \frac{2}{\varepsilon} \|\varphi_\varepsilon - \varphi\|_{L^\infty(\Omega)} + \int_s^t \frac{A(r)}{\varepsilon} dr. \quad (3.6)$$

*Step 4: Control on A and the measure of  $\{x, \varphi_\varepsilon \sim 0\}$ .*

For some  $\alpha_\varepsilon$  to be chosen later, decompose

$$\begin{aligned} \int_s^t A(r) dr &= \int_s^t \int_{\mathbb{R}} \left( \sum_{i=1}^k \bar{I}_i(s) \eta_i(x) - 1 \right) u_\varepsilon(r, x) \mathbb{I}_{\varphi_\varepsilon(r, x) \leq -\alpha_\varepsilon} dx dr \\ &\quad + \int_s^t \int_{\mathbb{R}} \left( \sum_{i=1}^k \bar{I}_i(s) \eta_i(x) - 1 \right) u_\varepsilon(r, x) \mathbb{I}_{\varphi_\varepsilon(r, x) \geq -\alpha_\varepsilon} dx dr. \end{aligned}$$

For the first part, note again that by (1.9), there exists  $R$  s.t.

$$\forall |x| > R, \quad \sum_{i=1}^k \bar{I}_i(s) \eta_i(x) \leq 1/2.$$

Therefore we may simply dominate

$$\int_s^t \int_{\mathbb{R}} \left( \sum_{i=1}^k \bar{I}_i(s) \eta_i(x) - 1 \right) u_\varepsilon(r, x) \mathbb{I}_{\varphi_\varepsilon(r, x) \leq -\alpha_\varepsilon} dx dr \leq C (t - s) e^{-\alpha_\varepsilon/\varepsilon}.$$

Concerning the second part, we constrain  $1/2 \geq \alpha_\varepsilon \geq \|\varphi - \varphi_\varepsilon\|_{L^\infty(\Omega)}$  and may therefore bound

$$\begin{aligned} &\int_s^t \left( \sum_{i=1}^k \bar{I}_i(s) \eta_i(x) - 1 \right) u_\varepsilon(r, x) \mathbb{I}_{\varphi_\varepsilon(r, x) \geq -\alpha_\varepsilon} dr \\ &\leq \int_s^t \left( \sum_{i=1}^k \bar{I}_i(s) \eta_i(x) - 1 \right)^+ u_\varepsilon(r, x) \mathbb{I}_{\varphi(r, x) \geq -2\alpha_\varepsilon} dr, \end{aligned}$$

where  $(a)^+$  denotes the positive part of  $a \in \mathbb{R}$ . Now  $\sum_{i=1}^k \bar{I}_i(s) \eta_i(x) - 1$  is nonpositive on  $\{\varphi(s, \cdot) = 0\}$  and so

$$\left( \sum_{i=1}^k \bar{I}_i(s) \eta_i - 1 \right)^+ \mathbb{I}_{\varphi(r, \cdot) = 0} \leq C \sup_{x \in \{\varphi(r, \cdot) = 0\}} \inf_{y \in \{\varphi(s, \cdot) = 0\}} |y - x| \leq C \tau_s(t - s),$$

by Step 1 as the  $\eta_i$  are uniformly Lipschitz. For two sets  $O_1$  and  $O_2$ , define in general

$$\delta(O_1, O_2) = \sup_{x \in O_1} \inf_{y \in O_2} |x - y|.$$

By the same argument, one gets

$$\begin{aligned} \left( \sum_{i=1}^k \bar{I}_i(s) \eta_i - 1 \right)^+ \mathbb{I}_{\varphi(r, \cdot) \geq -2\alpha_\varepsilon} &\leq C \tau_s(t - s) \\ &+ C \delta(\{\varphi(r, \cdot) \geq -2\alpha_\varepsilon\}, \{\varphi(r, \cdot) = 0\}). \end{aligned}$$

Inequality (3.5) now becomes

$$\begin{aligned} \int_s^t \sum_{i=1}^k c_i \frac{(I_i^\varepsilon(r) - \bar{I}_i(s))^2}{I_i^\varepsilon(r) \bar{I}_i(s)} dr &\leq C \varepsilon + 2 \|\varphi_\varepsilon - \varphi\|_{L^\infty(\Omega)} + C(t - s) e^{-\alpha_\varepsilon/\varepsilon} \\ &+ C \int_s^t \tau_s(r - s) dr + C \int_s^t \delta(\{\varphi(r, \cdot) \geq -2\alpha_\varepsilon\}, \{\varphi(r, \cdot) = 0\}) dr. \end{aligned} \tag{3.7}$$

*Conclusion.* Eq. (3.5) indeed gives Lemma 3.2 if one defines

$$\begin{aligned} \sigma_s(t - s) &= \frac{1}{t - s} \int_s^t \tau_s(r - s) dr, \\ \tilde{\sigma}(\varepsilon) &= C \varepsilon + 2 \|\varphi_\varepsilon - \varphi\|_{L^\infty(\Omega)} + C T e^{-\alpha_\varepsilon/\varepsilon} \\ &+ C \int_0^T \delta(\{\varphi(r, \cdot) \geq -2\alpha_\varepsilon\}, \{\varphi(r, \cdot) = 0\}) dr. \end{aligned}$$

Of course  $\sigma_s$  is continuous and, as  $\tau_s$  is continuous and  $\tau_s(0) = 0$ , then trivially  $\sigma_s(0) = 0$ . Since  $\{\varphi(r, \cdot) \geq -2\alpha_\varepsilon\}$  and  $\{\varphi(r, \cdot) = 0\}$  are subsets of  $\Omega$ ,  $\tilde{\sigma}(\varepsilon)$  is bounded for  $\varepsilon \leq 1$ , and thus, in order to complete the proof of Lemma 3.2, we only have to check that  $\tilde{\sigma}(\varepsilon) \rightarrow 0$  when  $\varepsilon \rightarrow 0$  for a convenient choice of  $\alpha_\varepsilon$ . If we take  $\alpha_\varepsilon \geq \|\varphi_\varepsilon - \varphi\|_{L^\infty(\Omega)}$  converging to 0 slowly enough to have  $\alpha_\varepsilon/\varepsilon \rightarrow +\infty$ , we only have to prove that

$$C \int_0^T \delta(\{\varphi(r, \cdot) \geq -2\alpha_\varepsilon\}, \{\varphi(r, \cdot) = 0\}) dr \longrightarrow 0 \quad \text{as } \varepsilon \rightarrow 0.$$

By dominated convergence it is enough that for any  $r$

$$\delta(\{\varphi(r, \cdot) \geq -2\alpha_\varepsilon\}, \{\varphi(r, \cdot) = 0\}) \longrightarrow 0.$$

Just as in Step 1 this is a direct consequence of the continuity of  $\varphi$ .

### 3.4 Compactness of the $I_i^\varepsilon$ and the proof of (1.12)

First notice that, simply passing to the limit in Lemma 3.2,

**Lemma 3.3** *There exists  $\sigma_s \in C(\mathbb{R}_+)$  with  $\sigma_s(0) = 0$  s.t.  $\forall i$*

$$\int_s^t |I_i(r) - \bar{I}_i(s)|^2 dr \leq (t-s) \sigma_s(t-s).$$

This means that at any point of Lebesgue continuity of  $I_i$ , one has  $I_i = \bar{I}_i$ . We recall that *a.e.* point is a Lebesgue point for  $I_i$ . As the  $I_i$  were defined only almost everywhere anyhow (they are weak-\* limits), we may identify  $I_i$  and  $\bar{I}_i$ . This proves (1.12) and that  $\bar{I}_i$  is approximately continuous on the right for any time  $t$  (and not only *a.e.*  $t$ ).

Now let us prove the compactness in  $L_{\text{loc}}^1$  of each  $I_i^\varepsilon$ . We apply the usual criterion and hence wish to control

$$\int_0^T \frac{1}{h} \int_s^{s+h} |I_i^\varepsilon(t) - I_i^\varepsilon(s)| dt ds.$$

Decompose

$$\begin{aligned} \int_0^T \frac{1}{h} \int_s^{s+h} |I_i^\varepsilon(t) - I_i^\varepsilon(s)| dt ds &\leq \int_0^T \frac{1}{h} \int_s^{s+h} |I_i^\varepsilon(t) - I_i(s)| dt ds \\ &+ \int_0^T \frac{1}{h} \int_s^{s+h} |I_i(t) - I_i^\varepsilon(s)| dt ds + \int_0^T \frac{1}{h} \int_s^{s+h} |I_i(t) - I_i(s)| dt ds. \end{aligned}$$

The first and third terms are bounded directly from Lemmas 3.2 and 3.3: for example,

$$\begin{aligned} \int_0^T \frac{1}{h} \int_s^{s+h} |I_i^\varepsilon(t) - I_i(s)| dt ds &\leq \int_0^T \left( \frac{1}{h} \int_s^{s+h} |I_i^\varepsilon(t) - I_i(s)|^2 dt \right)^{1/2} ds \\ &\leq \int_0^T (\sigma_s(h) + \tilde{\sigma}(\varepsilon)/h)^{1/2} ds. \end{aligned}$$



The second term can be handled the same way after swapping the order of integration

$$\begin{aligned} \int_0^T \frac{1}{h} \int_s^{s+h} |I_i^\varepsilon(s) - I_i(t)| dt ds &= \int_0^{T+h} \frac{1}{h} \int_{\max(0, t-h)}^t |I_i^\varepsilon(s) - I_i(t)| ds dt \\ &\leq \int_0^{T+h} \left( \frac{1}{h} \int_{\max(0, t-h)}^t |I_i^\varepsilon(s) - I_i(t)|^2 ds \right)^{1/2} dt \\ &\leq \int_0^{T+h} (\sigma_t(h) + \tilde{\sigma}(\varepsilon)/h)^{1/2} dt. \end{aligned}$$

So finally we bound

$$\begin{aligned} \int_0^T \frac{1}{h} \int_s^{s+h} |I_i^\varepsilon(t) - I_i^\varepsilon(s)| dt ds &\leq 3 \int_0^{T+h} (\sigma_s(h) + \tilde{\sigma}(\varepsilon)/h)^{1/2} ds \\ &\leq 3 \int_0^{T+h} \sqrt{\sigma_s(h)} ds + 3(T+h)\sqrt{\tilde{\sigma}(\varepsilon)/h}. \end{aligned}$$

Since of course the functions  $\sigma_s(\cdot)$  can be chosen uniformly bounded in Lemma 3.2, again by dominated convergence, this shows that  $\forall \tau > 0, \exists h, \exists \varepsilon_0(h)$  s.t.  $\forall \varepsilon < \varepsilon_0(h)$

$$\int_0^T \frac{1}{h} \int_s^{s+h} |I_i^\varepsilon(t) - I_i^\varepsilon(s)| dt ds \leq \tau.$$

This is enough to get compactness of the  $I_i^\varepsilon$  in  $L_{\text{loc}}^1$  and then in any  $L_{\text{loc}}^p$  with  $p < \infty$ , which concludes the proof of Theorem 1.1.

## References

- [1] G. Barles, S. Mirrahimi, B. Perthame, Concentration in Lotka-Volterra parabolic or integral equations: a general convergence result. *Methods Appl. Anal.* **16**, No. 3, 321–340 (2009).
- [2] G. Barles, B. Perthame, Concentrations and constrained Hamilton-Jacobi equations arising in adaptive dynamics. *Recent developments in nonlinear partial differential equations*, 57–68, Contemp. Math., **439**, Amer. Math. Soc., Providence, RI, 2007.

- [3] Bürger, R., Bomze, I. M. Stationary distributions under mutation-selection balance: structure and properties. *Adv. Appl. Prob.*, **28**, 227–251 (1996).
- [4] Calsina, A., Cuadrado, S. Small mutation rate and evolutionarily stable strategies in infinite dimensional adaptive dynamics. *J. Math. Biol.*, **48**, 135–159 (2004).
- [5] Carrillo, J. A., Cuadrado, S., Perthame, B. Adaptive dynamics via Hamilton-Jacobi approach and entropy methods for a juvenile-adult model. *Math. Biosci.* **205**(1), 137–161 (2007).
- [6] N. Champagnat. A microscopic interpretation for adaptive dynamics trait substitution sequence models. *Stoch. Proc. Appl.*, **116**, 1127–1160 (2006).
- [7] N. Champagnat, R. Ferrière, G. Ben Arous. The canonical equation of adaptive dynamics: a mathematical view. *Selection*, **2**, 71–81 (2001).
- [8] N. Champagnat, R. Ferrière, S. Méléard. From individual stochastic processes to macroscopic models in adaptive evolution, *Stoch. Models*, **24** suppl. 1, 2–44 (2008).
- [9] N. Champagnat, P.-E. Jabin, G. Raoul. Convergence to equilibrium in competitive Lotka-Volterra and chemostat systems, *C. R. Math. Acad. Sci. Paris*, **348**(23–24), 1267–1272 (2010).
- [10] N. Champagnat, S. Méléard. Polymorphic evolution sequence and evolutionary branching. To appear in *Probab. Theory Relat. Fields* (published online, 2010).
- [11] R. Cressman, J. Hofbauer. Measure dynamics on a one-dimensional continuous trait space: theoretical foundations for adaptive dynamics. *Theor. Pop. Biol.*, **67**, 47–59 (2005).
- [12] L. Desvillettes, P.-E. Jabin, S. Mischler, G. Raoul. On selection dynamics for continuous structured populations. *Commun. Math. Sci.*, **6**(3), 729–747 (2008).
- [13] Dieckmann, U. and Law, R. The dynamical theory of coevolution: a derivation from stochastic ecological processes. *J. Math. Biol.*, **34**, 579–612 (1996).

- [14] O. Diekmann, A beginner's guide to adaptive dynamics. In Mathematical modelling of population dynamics, *Banach Center Publ.*, **63**, 47–86, Polish Acad. Sci., Warsaw (2004).
- [15] O. Diekmann, M. Gyllenberg, H. Huang, M. Kirkilionis, J.A.J. Metz, H.R. Thieme. On the formulation and analysis of general deterministic structured population models. II. Nonlinear theory. *J. Math. Biol.* **43**, 157–189 (2001).
- [16] O. Diekmann, P.E. Jabin, S. Mischler, B. Perthame, The dynamics of adaptation: An illuminating example and a Hamilton-Jacobi approach. *Theor. Popul. Biol.* **67**, 257–271 (2005).
- [17] Genieys, S., Bessonov, N., Volpert, V. Mathematical model of evolutionary branching. *Math. Comput. Modelling*, **49(11–12)**, 2109–2115 (2009).
- [18] Geritz, S. A. H., Metz, J. A. J., Kisdi, E., Meszéna, G. Dynamics of adaptation and evolutionary branching. *Phys. Rev. Lett.*, **78**, 2024–2027 (1997).
- [19] Geritz, S. A. H., Kisdi, E., Meszéna, G., Metz, J. A. J. Evolutionary singular strategies and the adaptive growth and branching of the evolutionary tree. *Evol. Ecol.*, **12**, 35–57 (1998).
- [20] J. Hofbauer, R. Sigmund. Adaptive dynamics and evolutionary stability. *Applied Math. Letters*, **3**, 75–79 (1990).
- [21] P.E. Jabin, G. Raoul, Selection dynamics with competition. To appear *J. Math Biol.*.
- [22] Metz, J. A. J., Nisbet, R. M. and Geritz, S. A. H. How should we define 'fitness' for general ecological scenarios? *Trends in Ecology and Evolution*, **7**, 198–202 (1992).
- [23] Metz, J. A. J., Geritz, S. A. H., Meszéna, G., Jacobs, F. A. J. and van Heerwaarden, J. S. Adaptive Dynamics, a geometrical study of the consequences of nearly faithful reproduction. In: van Strien, S. J. & Verduyn Lunel, S. M. (ed.), *Stochastic and Spatial Structures of Dynamical Systems*, North Holland, Amsterdam, pp. 183–231 (1996).

- [24] S. Mirrahimi, G. Barles, B. Perthame, P. E. Souganidis, Singular Hamilton-Jacobi equation for the tail problem. *Preprint*.
- [25] Perthame, B., Gauduchon, M. Survival thresholds and mortality rates in adaptive dynamics: conciliating deterministic and stochastic simulations. *IMA Journal of Mathematical Medicine and Biology*, to appear (published online, 2009).
- [26] Perthame, B., Génieys, S. Concentration in the nonlocal Fisher equation: the Hamilton-Jacobi limit. *Math. Model. Nat. Phenom.*, **2(4)**, 135–151 (2007).
- [27] F. Yu. Stationary distributions of a model of sympatric speciation, *Ann. Appl. Probab.*, **17**, 840–874 (2007).