



HAL
open science

Allee effect promotes diversity in traveling waves of colonization

Lionel Roques, Jimmy Garnier, François Hamel, Etienne K. Klein

► **To cite this version:**

Lionel Roques, Jimmy Garnier, François Hamel, Etienne K. Klein. Allee effect promotes diversity in traveling waves of colonization. *Proceedings of the National Academy of Sciences of the United States of America*, 2012, 109 (23), pp.8828-8833. 10.1073/pnas.1201695109 . hal-01257401

HAL Id: hal-01257401

<https://hal.science/hal-01257401>

Submitted on 16 Jan 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.

Allee effect promotes diversity in traveling waves of colonization

Lionel Roques^{a,1}, Jimmy Garnier^{a,b}, François Hamel^{b,c}, and Etienne K. Klein^a

^aUR 546 Biostatistique et Processus Spatiaux, Institut National de la Recherche Agronomique, F-84000 Avignon, France; ^bAix-Marseille Université, Laboratoire d'Analyse, Topologie, Probabilités (Unité Mixte de Recherche 7353), Faculté des Sciences et Techniques, Avenue Escadrille Normandie-Niemen, F-13397 Marseille Cedex 20, France; and ^cInstitut Universitaire de France

Edited by* Haim Brezis, Rutgers, The State University of New Jersey, Piscataway, NJ, and approved March 28, 2012 (received for review January 31, 2012)

Most mathematical studies on expanding populations have focused on the rate of range expansion of a population. However, the genetic consequences of population expansion remain an understudied body of theory. Describing an expanding population as a traveling wave solution derived from a classical reaction-diffusion model, we analyze the spatio-temporal evolution of its genetic structure. We show that the presence of an Allee effect (i.e., a lower per capita growth rate at low densities) drastically modifies genetic diversity, both in the colonization front and behind it. With an Allee effect (i.e., pushed colonization waves), all of the genetic diversity of a population is conserved in the colonization front. In the absence of an Allee effect (i.e., pulled waves), only the furthest forward members of the initial population persist in the colonization front, indicating a strong erosion of the diversity in this population. These results counteract commonly held notions that the Allee effect generally has adverse consequences. Our study contributes new knowledge to the surfing phenomenon in continuous models without random genetic drift. It also provides insight into the dynamics of traveling wave solutions and leads to a new interpretation of the mathematical notions of pulled and pushed waves.

Rapid increases in the number of biological invasions by alien organisms (1) and the movement of species in response to their climatic niches shifting as a result of climate change have caused a growing number of empirical and observational studies to address the phenomenon of range expansion. Numerous mathematical approaches and simulations have been developed to analyze the processes of these expansions (2, 3). Most results focus on the rate of range expansion (4), and the genetic consequences of range expansion have received little attention from mathematicians and modelers (5). However, range expansions are known to have an important effect on genetic diversity (6, 7) and generally lead to a loss of genetic diversity along the expansion axis due to successive founder effects (8). Simulation studies have already investigated the role of the geometry of the invaded territory (9–11), the importance of long-distance dispersal and the shape of the dispersal kernel (12–14), the effects of local demography (15), or existence of a juvenile stage (13). Further research is needed to obtain mathematical results supporting these empirical and simulation studies, as such results could determine the causes of diversity loss and the factors capable of increasing or reducing it.

In a simulation study using a stepping-stone model with a lattice structure, Edmonds et al. (16) analyzed the fate of a neutral mutation present in the leading edge of an expanding population. Although in most cases the mutation remains at a low frequency in its original position, in some cases the mutation increases in frequency and propagates among the leading edge. This phenomenon is described as “surfing” (15). Surfing is caused by the strong genetic drift taking place on the edge of the population wave (5, 17) because the local growth rate of the low density individuals on the edge of the expanding wave is typically higher than the growth rate for the bulk of the population (15). The existence of surfing events has a significant impact on the subsequent genetic patterns of the population after expansion (18) and their occurrence can

be influenced by the existence of long-distance dispersal events (14), local demography (15), or selection (11).

The existence of an Allee effect is another critical factor affecting the dynamics of the leading edge of a population expansion. The Allee effect is characterized by a decrease in the per capita growth rate at low densities, and this can be due to increased damage from bioaggressors, increased mortality due to interspecific competition or reduced fitness due to suboptimal mating opportunities (19). This dynamic has been observed in many populations (20–22). The Allee effect is known to affect the rate of spread of a population (23, 24) and is expected to modify genetic drift on the edge of that population. Using simulation models with stochastic demography, Hallatschek and Nelson (25) provided a numerical analysis of the surfing phenomenon in the presence of an Allee effect. More precisely, using a backward-time approach, they analyzed the initial position of successful surfers in the wave. Significant differences in the probability distributions of the successful surfers were found between populations that experience an Allee effect and populations that do not. Using the framework of reaction-diffusion equations, they were able to connect their numerical findings to analytical formulas. Among other things, they concluded that surfing is not possible in deterministic reaction-diffusion, Kolmogorov–Petrovsky–Piskunov (KPP) type models (26, 27) (i.e., without an Allee effect). The goal of our study is to investigate how the Allee effect determines genetic diversity in a colonization front and to do so in a context broader than that considered in the surfing phenomenon.

Following the framework provided in (25) and (28), we focus on one-dimensional reaction-diffusion equations of the form:

$$\partial_t u = \partial_{xx} u + f(u), \quad t > 0, \quad x \in (-\infty, +\infty), \quad [1]$$

where $u = u(t, x)$ is the density of the population (of genes or haploid individuals) at time t and space position x . It evolves in time under the joint effects of local dispersal accounted for by the diffusion term and local reproduction described by the growth function f . Since Skellam's work (29), these models have commonly been used to explore population range expansions. However, little is known regarding the evolution of the inside structure of these models' solutions, that is, the dynamics of the components through which the structure of a population is determined.

Under some assumptions on the function f and the initial population u_0 , the solutions of Eq. 1 converge to traveling wave solutions (27, 30, 31). These solutions describe the invasion of the unoccupied region with a constant speed c and a constant density profile U , and the population density can be written as $u(t, x) = U(x - ct)$. In this study, we focus on such traveling wave solutions,

Author contributions: L.R., J.G., F.H., and E.K.K. designed research, performed research, contributed new reagents/analytic tools, and wrote the paper.

The authors declare no conflict of interest.

*This Direct Submission article had a prearranged editor.

Freely available online through the PNAS open access option.

¹To whom correspondence should be addressed. E-mail: lionel.roques@avignon.inra.fr.

Results: How the Fractions Propagate

We consider an arbitrarily chosen fraction v^k that satisfies Eq. 2—we call it v in the sequel—and we study the evolution of the density $v(t, x)$. From our assumptions, the total population $u(t, x)$ satisfies that $u(t, x) = U(x - ct)$. Thus, the density of the fraction satisfies

$$\partial_t v = \partial_{xx} v + v g(U(x - ct)), \quad t > 0, \quad x \in \mathbb{R}. \quad [7]$$

Moreover, when $t = 0$, $u_0(x) = U(x)$ and $v_0(x) = v(0, x)$ corresponds to a fraction of the quantity $U(x)$, thus $0 \leq v_0(x) \leq U(x)$ for all $x \in \mathbb{R}$. From the stability properties of the traveling waves, considering the initial condition $u_0(x) = U(x)$ is equivalent to defining the fractions inside a population that has already reached its stationary profile. Biologically, this means that we consider the spatio-temporal evolution of the diversity distributed at a given time in an ongoing wave of colonization.

Using the properties of the growth function f and the subsequent properties of the profile U and of the speed c , we can describe the evolution of v in a moving interval with speed c , as well as behind the waves.

Evolution of the Density v of the Fraction in a Moving Frame. In this section, we study the evolution of the density v of the fraction in an interval moving at the same speed c as the total population. Two situations may occur herein. Either the fraction is able to follow the total population and spreads with the same speed c , or the fraction is not able to follow the total population.

Our first result is concerned with the KPP case and is valid under general assumptions on the initial density v_0 of the fraction. These assumptions include the particular case of compactly supported initial fractions [i.e., $v_0(x) = 0$ outside a bounded set] and are satisfied by all the fractions depicted in Fig. 1, with the exception of the rightmost (light gray) one.

Result 1a (KPP case; see proof in Appendix A). *If the initial density v_0 of the fraction converges to 0 faster than U as $x \rightarrow +\infty$,* then, for any $A \in \mathbb{R}$, the density $v(t, x)$ of the fraction converges (as $t \rightarrow \infty$) to 0 uniformly in the moving half-line $[A + ct, \infty)$.*

This result shows that under the KPP assumption any fraction v whose initial density $v_0(x)$ is 0 for large x cannot expand with the total population.

It was stated in ref. 25 that gene surfing was not possible for the logistic growth function $f(u) = u(1 - u)$. If by gene surfing we mean that the wave tends to be made of a single fraction, then our result also shows that surfing is not possible for fractions with compactly supported initial densities. However, the surfing of fractions that are not initially compactly supported can occur, even with logistic growth functions. Consider the rightmost fraction v^r (the light gray fraction in Fig. 1): At $t = 0$, $v_0^r = U \cdot \mathbf{1}_{[\alpha, \infty)}$, where $\mathbf{1}_{[\alpha, \infty)}$ denotes the indicator function of the interval $[\alpha, \infty)$ for some $\alpha \in \mathbb{R}$. The fraction corresponding to the remaining part of the population satisfies $v_0^l = U \cdot \mathbf{1}_{(-\infty, \alpha)}$ as well as the assumption of Result 1a. Because $u(t, x) = U(x - ct) = v^l(t, x) + v^r(t, x)$, this result shows that $v^r(t, x)$ converges to $U(x - ct)$ in any moving half-line $[A + ct, \infty)$. This means that the fraction v^r manages to “surf” on the wave.

Result 2a (Allee case; see proof in Appendix B). *For any $A \in \mathbb{R}$ the density v of the fraction converges (as $t \rightarrow \infty$) to a proportion $p[v_0]$ of the total population $u(t, x)$ in the moving half-line $[A + ct, \infty)$ that is $v(t, x) - p[v_0]u(t, x) = v(t, x) - p[v_0]U(x - ct) \rightarrow 0$ as*

$t \rightarrow \infty$, uniformly in $[A + ct, \infty)$. The proportion $p[v_0]$ can be computed explicitly:

$$p[v_0] = \frac{\int_{-\infty}^{+\infty} v_0(x) U(x) e^{cx} dx}{\int_{-\infty}^{+\infty} U^2(x) e^{cx} dx} \in [0, 1]. \quad [8]$$

This result shows that any fraction v with a nonzero initial density v_0 follows the total population. Moreover, in the interval moving with speed c , the profile $v(t, ct + \cdot)$ of the fraction tends to resemble the profile U of the total population, with a scaling factor $p[v_0]$ dependent on the initial density v_0 . Note that the integral terms in the expression of p are well-defined, and this can be easily checked using the formulas in Eq. 5. This would not be true under the KPP assumption (30).

The formula in Eq. 8 provides precise information regarding the origin of the individuals that compose the wave at large times. Let us again consider the “leftmost” fraction defined by $v_0^l = U \cdot \mathbf{1}_{(-\infty, \alpha)}$ for $\alpha \in \mathbb{R}$. The asymptotic proportion of this fraction in any moving half-line with speed c is $p(\alpha) := p[v_0^l]$. Differentiating $p(\alpha)$ with respect to α we obtain a quantity $p'(\alpha)$ that can be interpreted as the relative contribution to the wave of the individuals with an initial position α :

$$p'(\alpha) = U^2(\alpha) e^{c\alpha} / \left(\int_{-\infty}^{+\infty} U^2(x) e^{cx} dx \right).$$

Using a similar formula, and replacing U with the solution of a stochastic simulation model incorporating an Allee effect, ref. 25 obtained a good fit of the probability of gene surfing in their model.

Here, the density profile U is known explicitly. Using the formula in Eq. 5, we observe that $p'(\pm\infty) = 0$ and that p' reaches a unique maximum at

$$\alpha_{\max} = \sqrt{2} \ln \left(\frac{1 - 2\rho}{1 + 2\rho} \right). \quad [9]$$

Interestingly, α_{\max} is a decreasing function of ρ , with $\alpha_{\max}(0) = 0$, corresponding to the position of the inflexion point of the profile U , and $\alpha_{\max}(1/2) = -\infty$. This formula emphasizes the advantageous role of the Allee effect for the fractions situated deep in the core of the population; the stronger the Allee effect, the more these individuals will contribute to the wave.

Evolution of the Density v of the Fraction Behind the Wave. The aim of the previous section was to analyze the behavior of an arbitrarily chosen fraction in a moving interval $[A + ct, \infty)$ with speed c equal to the spreading speed of the total population. Here, we analyze the evolution of the density of the fraction in the remaining part of the space: $(-\infty, A + ct)$. We assume that the initial density $v_0(x)$ of the fraction converges to 0 as $x \rightarrow -\infty$.

Result 1b (KPP case; see proof in Appendix C). *If the initial density v_0 of the fraction converges to 0 faster than U as $x \rightarrow +\infty$,* then, for any $A \in \mathbb{R}$, the density of the fraction converges (as $t \rightarrow \infty$) to 0 uniformly in the moving half-line $(-\infty, A + ct)$.*

This result, together with Result 1a, implies that the density $v(t, x)$ of the fraction converges to 0 uniformly in \mathbb{R} as $t \rightarrow \infty$. Thus, under the KPP assumption, any compactly supported fraction will vanish in the sense that, at large times, its density becomes negligible at any point of the space, as a result of dilution.

Result 2b (Allee case, see proof in Appendix D). *For any speed $c' \in (0, c)$ and any $A \in \mathbb{R}$, the density $v(t, x)$ of the fraction converges (as $t \rightarrow \infty$) to the proportion $p[v_0]u(t, x)$ in a set of the form $(c't, A + ct)$. Besides, for any $\varepsilon > 0$ and $B > 0$, $p[v_0]/2 - \varepsilon < v(t, x) < p[v_0] + \varepsilon$ in the set $[-B, A + ct)$ for $t > 0$ and $-A > 0$ large enough.*

Result 2a shows that, in any moving half-line $[A + ct, \infty)$, the density v of the fraction tends to resemble a proportion $p[v_0]$ of the total population u when the Allee effect is present. Result 2b

*This means that $\int_0^{+\infty} v_0^2(y) dy < \infty$. Note that if v_0 is compactly supported, then $v(t, x)$ converges to 0 faster than $U(x)$ for each $t \geq 0$, see the formula in Eq. 10 in Appendix A.

shows that this is actually true in $(c't, +\infty)$, for any $c' > 0$. This result also indicates that, in the reference frame, the fraction v propagates to the right with the same speed as the total population, but also diffuses in the opposite direction, given that B can be chosen arbitrarily large. For any initially compactly supported fraction v , the fraction centroid can be defined for $t > 0$ as the unique point $\bar{x}(t)$ satisfying $\int_{-\infty}^{\bar{x}(t)} v(t, y) dy = \int_{\bar{x}(t)}^{+\infty} v(t, y) dy$. Result 2b shows that $\bar{x}(t)$ cannot spread faster than $(c' + c)/2$, for all $c' > 0$. This means that the fraction centroid moves to the right at a speed less or equal to $c/2$.

Numerical Computations

Our analytical results were derived for a front-like initial condition $u_0(x) = U(x)$, corresponding to an already established traveling wave. We investigate numerically whether these results remain qualitatively true when u_0 is a compactly supported step function (Fig. 2A) that has not yet reached a traveling wave profile. We assume that the population is made of $N = 8$ fractions v^k , which satisfy, at $t = 0$, $v_0^1 = \mathbf{1}_{(-40, -21]}$, $v_0^k = \mathbf{1}_{(x_{k-1}, x_k]}$ for $k = 2, \dots, N$ for the sequence $x_1 < x_2 < \dots < x_N = 0$ of evenly spaced points. We numerically solved Eq. 2 with KPP and Allee growth terms. In both cases, we observed that the solution $u(t, x)$ rapidly converges to a traveling wave profile (dashed curves in Fig. 2).

KPP Case. Fig. 2B shows the evolution of the spatial structure of the solution $u(t, x)$ of Eq. 1 with a KPP growth term. As predicted by Results 1a and 1b, only the rightmost fraction follows the propagation to the right of the total population. The mass of the rightmost fraction, which was initially small, increases linearly with time. This could be interpreted as a form of surfing. We can observe that this fraction slowly diffuses into the bulk of the population, but with a null speed (i.e., sublinearly).

We observe that the spatial structure of the population has a “vertical pattern,” meaning that the population is highly spatially structured. Fig. 2B shows that the evolution of this pattern is slow compared to the rate at which the population propagates. Actually, because the growth term $v^k g(u)$ in Eq. 2 is always positive, the density of the fractions 2, ..., $N - 1$ cannot decrease

to 0 faster than the solution of the heat equation $\partial_t v = \partial_{xx} v$ (i.e., cannot decrease faster than the order $1/\sqrt{t}$).

Allee Case. With the Allee growth function in Eq. 4, the numerical results of Fig. 2C and Fig. 2D show that the theoretical predictions of Results 2a and 2b remain qualitatively true when u_0 is compactly supported. In particular, the stronger the Allee effect is, the more the fractions situated deep in the core of the population contribute to the wave.

The rightmost fraction remains the most represented in the colonization front. However, contrary to the KPP case, all the other fractions are conserved in the colonization front, leading to a spatial structure of the population with a “horizontal pattern.”

Discussion

Using a mathematical model commonly recognized in the literature as a robust descriptor of a population colonizing an empty space (2–4), we showed that the presence of an Allee effect drastically modifies genetic diversity in the colonization front. When an Allee effect is present, all of the fractions of a population are conserved in the colonization front, even if their proportions differ according to their initial distribution. In the absence of an Allee effect, only the furthest forward fraction in the initial population eventually remains in the colonization front, indicating a strong erosion of diversity due to the demographic advantage of isolated individuals ahead of the colonization front. Under this classical KPP model, any fraction except that located at the head of the front vanishes progressively; this shows that the “enhanced transport” of neutral fractions described by Vlad et al (28) for similar equations is only a transient phenomenon.

These results diversify the commonly held perspective that the Allee effect possesses net adverse consequences. This perspective is inherited from demographic studies of range expansion, which demonstrate that an Allee effect reduces the speed of colonization (23, 35) and can even stop it in heterogeneous environments (24). Reducing the growth rate of the individuals ahead of the colonization front simultaneously reduces the speed of colonization and enables a diversity of genes coming from the core of the population to remain on the front, as demonstrated by Result 2a and the formula in Eq. 9. Other mechanisms that reduce the growth rate of the individuals ahead of the front should also result in greater conservation of the genetic diversity of a population. For instance, Pluess (36) demonstrated how a retreating glacier limited the spread of a population of European larch, thereby functioning as an extreme Allee effect where all of the seeds falling on the icecap die. Given our results, this should lead to high genetic diversity in the colonization front, which was actually observed (36). The Allee effect could be a partial or alternative explanation to the argument of long-distance mixing of genes advanced in ref. 36. Our results are also consistent with the findings in ref. 37, which showed that the existence of a juvenile phase (i.e., a nonreproductive life-stage) in the life-cycle of an organism can lead to higher levels of genetic diversity. Although the juvenile phase does not generate an Allee effect in the strictest sense, it slows down the colonization process in a similar way, thus enabling an accumulation of genetic diversity in the colonization front.

The Allee effect also affects the spatial distribution of diversity. As Fig. 2C and D illustrate, this effect leads to a “horizontal pattern” of genetic diversity (i.e., an absence of genetic differentiation in space), and Result 2b shows that, after the population has reached its maximum capacity, this pattern diffuses in the opposite direction within the core of the population. Conversely, we observed a “vertical pattern” of genetic diversity in the absence of an Allee effect (i.e., a strongly structured spatial distribution of population fractions; see Fig. 2B). This genetic structure eventually attenuates due to diffusion in the saturated population, and each fraction becomes negligible. However, in both cases the diffusion in the saturated population occurs at

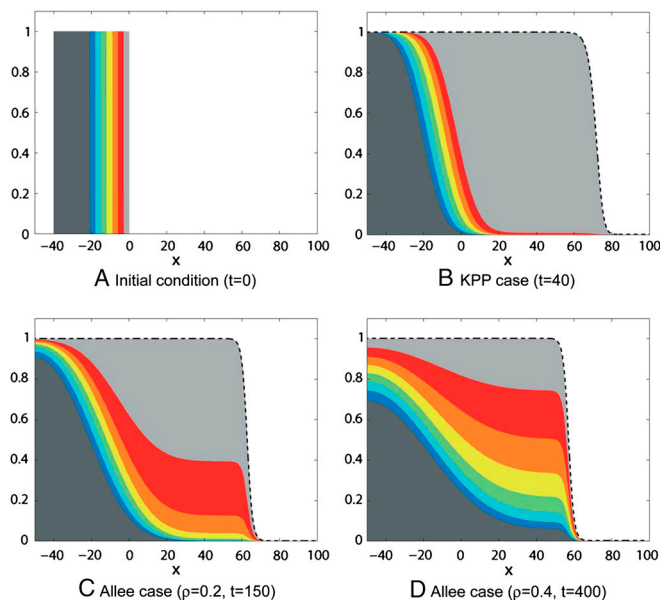


Fig. 2. Evolution of the spatial structure of the solution $u(t, x)$ of Eq. 1. (A) Initial distribution of the fractions; (B) spatial structure of the colonization wave $u(t, x)$ with the KPP growth term $f(u) = u(1 - u)$; (C and D) spatial structure of the colonization wave $u(t, x)$ with the growth term $f(u) = u(1 - u)(u - \rho)$. In each case, the dashed black curve corresponds to the profile U of the stable traveling wave solution of Eq. 1.

a much lower rate than the rate of colonization (sublinear diffusion vs. linear propagation). These two time-scales are consistent with the results obtained in ref. 13 with a stepping-stone model.

Genetic drift is not taken into account in our forward-time approach, and this is an important difference between this study and (25). In (25), the dominant role of genetic drift eventually leads to the fixation of a single gene in the colonization front, leading to a total loss of diversity. However, consistent results are obtained from their backward-time approach. In the absence of stochastic genetic drift, the deterministic evolution of the allelic densities does not depend on the distribution of the alleles within and among diploid individuals (38). Our results can therefore be applied to the dynamics of genetic diversity in haploid as well as diploid populations.

Our forward-time approach also underlines the ambiguity in the definition of the surfing phenomenon. Surfing can be associated with either (i) a rare gene becoming drastically dominant in the front or, (ii) with a gene initially present on the front being propagated alongside others present in the front. These two definitions lead to contradictory results, as with definition (i), gene surfing is only possible without an Allee effect and for the furthest forward fraction of the front, and definition (ii) dictates that gene surfing is promoted by the Allee effect. In the presence of an Allee effect, the centroid of any fraction of the front is propagated at speed $c/2$, which is consistent with the fact that any fraction spreads between its initial location and the leading point of the front (5, 16).

From a mathematical standpoint, our study contributes a unique perspective to the extensively studied topic of reaction-diffusion equations. One of the main features of these equations is their exhibition of traveling wave solutions that keep a constant profile. Earlier approaches were concerned with the dynamics of the total waves, making our study's mathematical analysis of the dynamics of the inside structure of these waves unique. Our results show that these dynamics are strongly dependent on the type of growth function f , and seem to be determined by the pulled-pushed nature of the waves. Conversely, our observations show that the notions of pulled and pushed solutions can be defined based on the inside structure of the solutions rather than on their speed of propagation. This conceptualization of pushed and pulled solutions, whose mathematical definitions will be given in a future work, has the advantage of being intuitive and adaptable to more complex models that do not necessarily admit traveling wave solutions. For instance, we should now be able to determine the pushed-pulled nature of the solutions of (i) integro-differential equations including long-distance dispersal events and resulting in accelerating waves (39, 40), (ii) reaction-diffusion equations with spatially heterogeneous coefficients that lead to pulsating or generalized transition waves (41, 42), (iii) reaction-diffusion equations with forced speed, which have been used in ref. 43 to study the effects of a shifting climate on the dynamics of a biological species. These processes are known in ecology to be determinants of the colonization patterns of many organisms (2, 7) and their effects on genetic diversity require further investigation (5).

Appendix A: Proof of Result 1a

The main idea of the proof is to compare the equation satisfied by v with a homogeneous linear equation, and to compute explicitly the solution of the linear equation.

Recall that v satisfies Eq. 7. From the KPP assumption, we know that $g(U(x-ct)) \leq f'(0)$. Thus, a comparison argument implies that v is smaller than the solution w of the equation: $\partial_t w = \partial_{xx} w + wf'(0)$, with the same initial condition $w(0, x) = v_0(x)$. This function w can be computed explicitly. Fix $A \in \mathbb{R}$ and consider an element $x_0 + ct$ in the moving half-line $[A + ct, \infty)$. Because $c = 2\sqrt{f'(0)}$, we have:

$$w(t, x_0 + ct) = \frac{e^{-\frac{x_0 c}{2}}}{\sqrt{4\pi t}} \int_{-\infty}^{+\infty} e^{\frac{y}{2}} v_0(y) e^{-\frac{(x_0 - y)^2}{4t}} dy. \quad [10]$$

Finally, because $x_0 \geq A$ and $\int_{-\infty}^{+\infty} e^{\frac{y}{2}} v_0(y) dy < \infty$, we get:

$$w(t, x_0 + ct) \leq \frac{e^{-\frac{A c}{2}}}{\sqrt{4\pi t}} \int_{-\infty}^{+\infty} e^{\frac{y}{2}} v_0(y) dy \rightarrow 0 \quad \text{as } t \rightarrow \infty,$$

and the above convergence is uniform in $x_0 \in [A, +\infty)$. Because $0 \leq v(t, x) \leq w(t, x)$, this implies the assertion in Result 1a.

Appendix B: Proof of Result 2a

In the moving frame with speed c , the fraction density can be written $\tilde{v}(t, x) = v(t, x + ct)$. In order to remove the advection terms that appear in the equation satisfied by \tilde{v} , we set $v^*(t, x) = \tilde{v}(t, x) e^{cx/2}$. Then, we can check that the function v^* is a solution of a linear equation without advection term:

$$\partial_t v^* = \partial_{xx} v^* + v^* \left[g(U(x)) - \frac{c^2}{4} \right], \quad [11]$$

with the initial condition $v^*(0, x) = v_0(x) e^{cx/2}$. In the remaining part of the proof, we show that v^* can be written as the sum of a stationary function and of a function that converges to 0 exponentially fast as $t \rightarrow \infty$.

Note that $\varphi(x) = e^{cx/2} U(x)$ is a positive eigenfunction of the operator that appears in the right hand side of Eq. 11, and that the associated eigenvalue is 0. The Sturm-Liouville theory implies that 0 is the largest eigenvalue of this operator, the remaining part of the spectrum being located at the left of some negative constant $-\mu$. Thus, we can write:

$$v^*(t, x) = p\varphi(x) + z(t, x), \quad [12]$$

where $p \in \mathbb{R}$ and z is "orthogonal" to φ in the sense that $\int_{-\infty}^{+\infty} z(t, x) \varphi(x) dx = 0$, for each $t \geq 0$. Moreover, $|z(t, x)| \leq K e^{-\mu t}$, for some constant $K > 0$. Multiplying the expression in Eq. 12 at $t = 0$ by φ and integrating, we get the expression in Eq. 8 for p .

Finally, we have $|v^*(t, x) - p\varphi(x)| \leq K e^{-\mu t}$ and therefore $|\tilde{v}(t, x) - pU(x)| \leq K e^{-\mu t - cx/2}$. This shows that $\tilde{v}(t, x)$ converges to $pU(x)$ uniformly in any moving half-line $[A - c't, +\infty)$ with $c' \in [0, 2\mu/c)$. In particular, taking $c' = 0$ and using $v(t, x) = \tilde{v}(t, x - ct)$ we obtain that, for any $A \in \mathbb{R}$, the fraction density converges to a proportion p of the total population $u(t, x) = U(x - ct)$, uniformly in the moving half-line $[A + ct, +\infty)$.

Appendix C: Proof of Result 1b

Take any $\varepsilon > 0$. From Result 1a, we already know that for any $A < 0$ there exists a time $t_A > 0$ such that $0 < v(t, A + ct) < \varepsilon/2$ for all $t \geq t_A$. Again, we place ourselves in the moving frame with speed c : $\tilde{v}(t, x) = v(t, x + ct)$ satisfies $0 < \tilde{v}(t, A) < \varepsilon/2$ and satisfies the equation:

$$\partial_t \tilde{v} = \partial_{xx} \tilde{v} + c \partial_x \tilde{v} + g(U(x)) \tilde{v}, \quad t > 0, x \in \mathbb{R}. \quad [13]$$

By constructing a super-solution to Eq. 13 close to the solution of the heat equation, we are going to show that $\tilde{v} \rightarrow 0$ in $(-\infty, A]$ as $t \rightarrow \infty$.

The assumption $v_0(x) \rightarrow 0$ as $x \rightarrow -\infty$ implies that, for any $t > 0$, $\tilde{v}(t, x) \rightarrow 0$ as $x \rightarrow -\infty$. In particular, there exists a point $x_A < A$ such that $\tilde{v}(t_A, x) \leq \varepsilon/2$ for all $x \leq x_A$. Thus, in the interval $(-\infty, A]$, and for times larger than t_A , \tilde{v} is smaller than the solution \tilde{v} of:

$$\begin{cases} \partial_t \tilde{v} = \partial_{xx} \tilde{v} + c \partial_x \tilde{v} + f(U(x)), & t \geq t_A, x < A, \\ \tilde{v}(t, A) = \varepsilon/2, & t \geq t_A, \\ \tilde{v}(t_A, x) = \begin{cases} 1 & \text{if } x \in (x_A, A), \\ \varepsilon/2 & \text{if } x \leq x_A. \end{cases} \end{cases} \quad [14]$$

Consider the auxiliary problem

$$\begin{cases} \partial_t \bar{v}^1 = \partial_{xx} \bar{v}^1 + c \partial_x \bar{v}^1, t \geq t_A, x \in \mathbb{R}, \\ \bar{v}^1(t_A, x) = \begin{cases} 1 & \text{if } x \in (x_A, A), \\ \varepsilon/2 & \text{if } x \leq x_A \text{ or } x \geq A. \end{cases} \end{cases} \quad [15]$$

Then, \bar{v}^1 is simply the solution of the heat equation with an advection term and can be computed explicitly. In particular, we have $\max_{x \in \mathbb{R}} \bar{v}^1(t, x) \leq \varepsilon/2 + C/\sqrt{t}$ for some constant $C > 0$. Setting $j(x) = -\varepsilon x/(1-x)$ for $x < 0$, we observe that $-j'' - cj'$ is positive and decreases like $c\varepsilon/x^2$ as $x \rightarrow -\infty$. Under the KPP assumption, it is known (44) that the profile $U(x)$ of the traveling wave converges exponentially to 1 as $x \rightarrow -\infty$. Using $f(1) = 0$ and $f'(1) < 0$, this implies that $f(U(x))$ converges exponentially to 0 as $x \rightarrow -\infty$. Thus, if A is chosen negative enough, $f(U(x)) \leq -j''(x) - cj'(x)$ in $(-\infty, A]$. The parabolic maximum principle then shows that $\bar{v}^1(t, x) + j(x) \geq \tilde{v}(t, x)$ in $(-\infty, A]$, for all $t \geq t_A$. We finally get:

$$\varepsilon + C/\sqrt{t} \geq \bar{v}^1(t, x) + j(x) \geq \tilde{v}(t, x) \geq \underline{v}(t, x), \quad [16]$$

for all $t \geq t_A, x \in (-\infty, A]$. Thus, $v(t, x) \leq \varepsilon + C/\sqrt{t}$ in $(-\infty, A + ct]$ for all $t \geq t_A$.

Appendix D: Proof of Result 2b

From Result 2a, we know that for any $A \in \mathbb{R}$ there exists a time t_A such that $|v(t, A + ct) - pU(A)| < \varepsilon/4$ for all $t \geq t_A$. Thus, if A is negative enough, because $U(-\infty) = 1$, we have $|v(t, A + ct) - p| < \varepsilon/3$ for $t \geq t_A$. Using the same arguments as in the proof of

Result 1b, we can show that $v(t, x)$ is smaller than $p + \varepsilon$ in the half-line $(-\infty, A + ct]$ for t large enough.

In order to construct a lower bound for v , we construct an appropriate subsolution. First, one can choose $A < 0$ such that $f(U(x)) \geq 0$ for all $x \leq A$. Then, setting

$$\begin{cases} \partial_t \underline{v} = \partial_{xx} \underline{v} + c \partial_x \underline{v}, t \geq t_A, x \in \mathbb{R}, \\ \underline{v}(t_A, x) = \begin{cases} 0 & \text{if } x \leq A, \\ p - \varepsilon/2 & \text{if } x \geq A, \end{cases} \end{cases} \quad [17]$$

a comparison argument implies that $\tilde{v}(t, x) = v(t, x + ct)$ is larger than $\underline{v}(t, x)$ for all $t \geq t_A$ and $x \leq A$. Again, Eq. 17 simply corresponds to the heat equation with an advection term, and its solution can be computed explicitly. For any $t > t_A$, the function $\underline{v}(t, \cdot)$ is increasing and therefore $\underline{v}(t + t_A, x) \geq \underline{v}(t + t_A, c'(t + t_A) - ct - B)$ for all $t > 0, B \geq 0$ and $x \geq c'(t + t_A) - ct - B$, which gives:

$$\underline{v}(t + t_A, x) \geq \frac{p - \varepsilon/2}{\sqrt{\pi}} \int_{\frac{A - B - c'(t + t_A)}{2\sqrt{t}}}^{\infty} e^{-z^2} dz.$$

As a consequence, if $0 < c' < c$, $v(t, x)$ is larger than $p - \varepsilon$ in $(c't, A + ct)$ for t large enough. We also observe (with $c' = 0$) that $v(t, x)$ is larger than $p/2 - \varepsilon$ in $[-B, A + ct)$ for t large enough.

ACKNOWLEDGMENTS. We thank Mickaël Chekroun, Bruno Fady, Claire Lavigne, Samuel Soubeyrand, and the anonymous reviewers for their insightful comments. The authors are supported by the French "Agence Nationale de la Recherche" within the projects ColonsGS, PREFERRED, and URTICLIM.

- DAISIE (2009) *Handbook of Alien Species in Europe* (Springer, Dordrecht, The Netherlands).
- Shigesada N, Kawasaki K (1997) *Biological Invasions: Theory and Practice* (Oxford Univ Press, Oxford).
- Turchin P (1998) *Quantitative Analysis of Movement: Measuring and Modeling Population Redistribution in Animals and Plants* (Sinauer, Sunderland, MA).
- Hastings A, et al. (2005) The spatial spread of invasions: New developments in theory and evidence. *Ecol Lett* 8:91–101.
- Excoffier L, Foll M, Petit RJ (2009) Genetic consequences of range expansions. *Annu Rev Ecol Syst* 40:481–501.
- Hewitt GM (2000) The genetic legacy of the quaternary ice ages. *Nature* 405:907–913.
- Davis MB, Shaw RG (2001) Range shifts and adaptive responses to quaternary climate change. *Science* 292:673–9.
- Roussellet J, et al. (2010) The role of topography in structuring the demographic history of the pine processionary moth, *Thaumetopoea pityocampa* (Lepidoptera: Notodontidae). *J Biogeogr* 37:1478–1490.
- Bialozyt R, Ziegenhagen B, Petit RJ (2006) Contrasting effects of long distance seed dispersal on genetic diversity during range expansion. *J Ecol Biol* 19:12–20.
- Wegmann D, Currat M, Excoffier L (2006) Molecular diversity after a range expansion in heterogeneous environments. *Genetics* 174:2009–2020.
- Travis JMJ, et al. (2007) Deleterious mutations can surf to high densities on the wave front of an expanding population. *Mol Biol Evol* 24:2334–2343.
- Ibrahim KM, Nichols RA, Hewitt GM (1996) Spatial patterns of genetic variation generated by different forms of dispersal during range expansion. *Heredity* 77:282–291.
- Austerlitz F, Garnier-Géré PH (2003) Modelling the impact of colonisation on genetic diversity and differentiation of forest trees: Interaction of life cycle, pollen flow and seed long-distance dispersal. *Heredity* 90:282–290.
- Fayard J, Klein EK, Lefèvre F (2009) Long distance dispersal and the fate of a gene from the colonization front. *J Evol Biol* 22:2171–2182.
- Klopfstein S, Currat M, Excoffier L (2006) The fate of mutations surfing on the wave of a range expansion. *Mol Biol Evol* 23:482–490.
- Edmonds CA, Lillie AS, Cavalli-Sforza LL (2004) Mutations arising in the wave front of an expanding population. *Proc Natl Acad Sci USA* 101:975–979.
- Excoffier L, Ray N (2008) Surfing during population expansions promotes genetic revolutions and structuration. *Trends Ecol Evol* 23:347–351.
- Hallatschek O, Hersen P, Ramanathan S, Nelson DR (2007) Genetic drift at expanding frontiers promotes gene segregation. *Proc Natl Acad Sci USA* 104:19926–19930.
- Berec L, Angulo E, Courchamp F (2007) Multiple Allee effects and population management. *Trends Ecol Evol* 22:185–191.
- Dennis B (1989) Allee effects: Population growth, critical density, and the chance of extinction. *Nat Resour Model* 3:481–538.
- Veit RR, Lewis MA (1996) Dispersal, population growth, and the Allee effect: Dynamics of the house finch invasion of eastern North America. *Am Nat* 148:255–274.
- Kramer AM, Dennis B, Liebhold AM, Drake JM (2009) The evidence for Allee effects. *Population Ecology* 51:341–354.
- Lewis MA, Kareiva P (1993) Allee dynamics and the spread of invading organisms. *Theor Popul Biol* 43:141–158.
- Barton NH, Turelli M (2011) Spatial waves of advance with bistable dynamics: Cytoplasmic and genetic analogues of Allee effects. *Am Nat* 178:E48–E75.
- Hallatschek O, Nelson DR (2008) Gene surfing in expanding populations. *Theor Popul Biol* 73:158–170.
- Fisher RA (1937) The wave of advance of advantageous genes. *Ann Hum Genet* 7:355–369.
- Kolmogorov AN, Petrovsky IG, Piskunov NS (1937) Étude de l'équation de la diffusion avec croissance de la quantité de matière et son application à un problème biologique [Study of the diffusion equation with growth of the quantity of matter and its application to a biological problem]. *Bulletin de l'Université d'État de Moscou, Série Internationale A* 1:1–26 (in French).
- Vlad MO, Cavalli-Sforza LL, Ross J (2004) Enhanced (hydrodynamic) transport induced by population growth in reaction-diffusion systems with application to population genetics. *Proc Natl Acad Sci USA* 101:10249–10253.
- Skellam JG (1951) Random dispersal in theoretical populations. *Biometrika* 38:196–218.
- Aronson DG, Weinberger HG (1975) *Nonlinear diffusion in population genetics, combustion and nerve propagation Lectures Notes Math*, (Springer, New York), Vol. 446, pp 5–49.
- Fife PC, McLeod J (1977) The approach of solutions of nonlinear diffusion equations to traveling front solutions. *Arch Ration Mech Anal* 65:335–361.
- Keitt TH, Lewis MA, Holt RD (2001) Allee effects, invasion pinning, and species' borders. *Am Nat* 157:203–216.
- Stokes AN (1976) On two types of moving front in quasilinear diffusion. *Math Biosci* 31:307–315.
- Rothe F (1981) Convergence to pushed fronts. *Rocky Mountain J Math* 11:617–634.
- Lewis MA, Van Den Driessche P (1993) Waves of extinction from sterile insect release. *Math Biosci* 116:221–247.
- Pluess AR (2011) Pursuing glacier retreat: Genetic structure of a rapidly expanding larch decidua population. *Mol Ecol* 20:473–485.
- Austerlitz F, Mariette S, Machon N, Gouyon PH, Godelle B (2000) Effects of colonization processes on genetic diversity: Differences between annual plants and tree species. *Genetics* 154:1309–1321.
- Hartl DL, Clark AG (2006) *Principles of Population Genetics, Fourth edition* (Sinauer, Sunderland, MA).
- Kot M, Lewis M, van den Driessche P (1996) Dispersal data and the spread of invading organisms. *Ecology* 77:2027–2042.
- Garnier J (2011) Accelerating solutions in integro-differential equations. *SIAM J Math Anal* 43:1955–1974.
- Xin J (2000) Front propagation in heterogeneous media. *SIAM Rev* 42:161–230.
- Berestycki H, Hamel F (2012) Generalized transition waves and their properties. *Comm Pure Appl Math* 65:592–648.
- Berestycki H, Diekmann O, Nagelkerke CJ, Zegeling PA (2009) Can a species keep pace with a shifting climate? *Bull Math Biol* 71:399–429.
- Aronson DG, Weinberger HG (1978) Multidimensional non-linear diffusion arising in population-genetics. *Adv Math* 30:33–76.