

http://[www.aimspress.com](http://http://www.aimspress.com/journal/MBE)/journal/MBE

Mathematical Biosciences and Engineering, 16(4): 3047–3070. [DOI: 10.3934](http://dx.doi.org/10.3934/mbe.2019151)/mbe.2019151 Received: 05 December 2018 Accepted: 11 March 2019 Published: 10 April 2019

Research article

Dynamics of a general model of host-symbiont interaction

Eric Foxall[∗]

Mathematical and Statistical Sciences, 632 Central Academic Building, University of Alberta, Edmonton AB, T6G 2G1, Canada

* Correspondence: Email: efoxall@ualberta.ca

Abstract: We consider a model of host-symbiont interactions, in which symbionts can only live in association with their host and are transmitted both vertically from associated hosts to their offspring and horizontally from associated hosts to nearby unassociated hosts. The effect of the symbiont is modelled by a change in the birth rate of associated hosts. We analyze the two-dimensional dynamics in the resulting four-dimensional parameter space, and determine the qualitative behaviour for all parameter values. We find that for all but one choice of parameter values, solutions in the feasible region, apart from a 0- or 1-dimensional set of initial conditions, tend either to a unique equilibrium, or to one of two distinct equilibria. Moreover, the bistable case occurs only when the symbiont is a mutualist whose horizontal spread rate through the host population exceeds the positive change in the birth rate of associated hosts.

Keywords: host-symbiont; ordinary differential equation; pathogen; mutualist

1. Introduction

Symbiosis is an important aspect of ecology [\[1\]](#page-21-0). In fact, an entire journal, titled Symbiosis, is devoted to the study of symbiotic interactions at all levels ranging from the molecular to the organismic. The precise meaning assigned to the term symbiosis may vary depending on context; a fairly general definition, given in [\[2\]](#page-21-1), is a close physical association between different species, regardless of whether that association is harmful, beneficial or neutral to any of the species involved.

In many cases of interest, the symbiosis is between a host, and one or more types of symbiont living within the host, often in large numbers. Two examples, both relating to humans, are infectious diseases [\[3,](#page-21-2) [4\]](#page-21-3) and gut microbe interactions [\[5\]](#page-21-4). A harmful symbiont is a pathogen, while a beneficial symbiont is a mutualist. Previous mathematical studies have focused on studying either the pathogen case [\[6\]](#page-21-5) or the mutualist case [\[7,](#page-21-6) [8\]](#page-21-7), however it would be useful to understand, from a unified perspective, how the nature of the host-symbiont relationship affects the range of observed interactions.

To this end, we study a simple two-variable model of host-symbiont interactions with four parameters (described in the next section) that includes both horizontal and vertical transmission of the symbiont, and naturally incorporates both the pathogen and mutualist cases, as well as the intermediate neutral (neither harmful nor beneficial) case, modelled by the symbiont's effect on the host birth rate. As well, we assume recovery is possible; that is, it is possible for the symbiont within a host to die without killing the host. For simplicity, we assume perfect vertical transmission of the symbiont, and distinguish only presence or absence of the symbiont within the host, tracking the population density of both *unassociated* hosts (those without the symbiont) and *associated* hosts. A generalization to imperfect vertical transmission is also easily implemented, though we have not considered it here.

A special case of our model appears in [\[9\]](#page-21-8), in which the authors discuss how it is possible, with a mixture of horizontal and vertical transmission, for a pathogen to successfully invade and persist within a host population, and they also give precise conditions for the pathogen to reach full occupation. In their model, only the pathogen case is considered, and recovery from the symbiont does not occur. In addition, the concern is with existence of certain equilibria and so a full description of the dynamics is not given. However, they also consider the case of imperfect transmission which is not covered here. From a different perspective, a model of mutualism [\[7\]](#page-21-6), which is similar though not identical to this model, shows a qualitatively similar nullcline structure in some cases that leads to bistability; interestingly, this similarity is only observed after a change of variables is made in our model, from associated and unassociated host density to total host density and proportion of hosts that are associated.

For our model we are able to obtain a complete description of the dynamics for all values of the four parameters. In all but a single case, every trajectory tends to one of at most three distinct equilibria, ruling out sustained oscillations and reducing the study of the model to the characterization of equilibria. In turn, the number and location of equilibria is closely related to two factors:

- the effect of the symbiont upon the host (pathogen, mutualist or neutral), and
- which of the two is larger: the difference in the birth rate of associated vs. unassociated hosts (which is positive only for mutualists), and the horizontal rate of spread of the symbiont through the host population.

The second quantity is shown to be relevant only after making the above-mentioned change of variables. One way to view it is by how the symbiont allocates resources: is it focused on increasing the host's rate of birth and thus vertical transmission, or is it focused on spreading immediately to existing hosts via horizontal transmission?

An important finding for this model is that bistability (between host-symbiont coexistence and either host extinction or host survival only) can only occur when the symbiont is a mutualist that is more focused on horizontal transmission than on improving host birth rate (i.e. when the second quantity above is positive), and that in the absence of bistability, all solutions with a positive density of both host and symbiont tend to a unique equilibrium. Moreover, in the absence of bistability the set of qualitative behaviours displayed by the system is insensitive as to whether the symbiont is a pathogen or a mutualist, except in the difference in host density between the host-without-symbiont equilibrium vs. the host-symbiont coexistence equilibrium, when both exist.

The original motivation for making a detailed study of this model was to obtain complementary results for a study of the corresponding stochastic spatial model [\[10\]](#page-21-9). When working on that paper, a preliminary analysis of the differential equations yielded partial but artifically restrictive results. The reason for this restriction was revealed only after making the change of variables mentioned above, from associated vs. unassociated host density to total host density vs. proportion of hosts that are associated, which made evident the occurrence of bistability, which previously we had thought was not possible. Thus one conclusion of this analysis is that the (biological) perspective of host-symbiont interactions as being a "stacked" system, in which hosts have their own birth-death dynamics, while symbionts within the host population have a similar birth-death dynamics taking place over the host population, is also a useful perspective when analyzing the dynamical system. It is worth noting that the stacked perspective is also used in [\[11\]](#page-21-10) and indeed in our paper [\[10\]](#page-21-9), where it assists in the analysis of the stochastic spatial model by suggesting comparisons with simpler models.

2. Model derivation and main result

For simplicity, we ignore environmental transmission of the symbiont, and distinguish only presence or absence of the symbiont within the host. Unassociated hosts reproduce at some rate, and all hosts die at some rate which is normalized to 1. Associated hosts can infect unassociated hosts, and can also recover and become unassociated. Moreover, associated hosts reproduce at some possibly different rate from associated hosts. We assume, again for simplicity, perfect vertical transmission of the symbiont, so that offspring of associated parents are always associated. The symbiont is identified as pathogen, mutualist, or neutral by its effect on the host's birth rate.

To derive the differential equations that we study, we begin with a set of *N* sites each of which can be either empty, occupied by an unassociated host, or occupied by an associated host. Each unassociated host attempts to give birth to an unassociated host onto a site chosen uniformly at random at rate λ_{10} , being successful if that site is empty. Similarly, each associated host attempts to give birth to an associated host at rate λ_{20} . Each host dies at rate 1, while each associated host becomes an unassociated host at rate δ . Each associated host attempts to transmit the symbiont to a randomly chosen site at rate host at rate δ . Each associated host attempts to transmit the symbiont to a randomly chosen site at rate
declines successful if the recipient is an unassociated host. Let $U^{(N)}(t)$, $U^{(N)}(t)$, $U^{(N)}(t)$ denote the λ_{21} , being successful if the recipient is an unassociated host. Let $U_0^{(N)}$
number of empty sites, unassociated hosts and associated hosts respect $U_0^{(N)}(t)$, $U_1^{(N)}$

(*N*) ctively $U_1^{(N)}(t)$, $U_2^{(N)}$
Since $\bar{\nabla}^2$ $t_2^{(N)}(t)$ denote the number of empty sites, unassociated hosts and associated hosts, respectively. Since $\sum_{i=0}^{2} U_i^{(N)}$ $i^{(N)}(t) = N$, we find that $(U_1^{(N)})$ $U_1^{(N)}(t)$, $U_2^{(N)}$ $\binom{N}{2}(t)$ is a Markov chain with the following transitions.

$$
(U_1, U_2) \rightarrow (U_1, U_2) + \begin{cases} (1, 0) & \text{at rate} & \lambda_{10} U_0 U_1 / N \\ (-1, 0) & \text{at rate} & U_1 \\ (0, 1) & \text{at rate} & \lambda_{20} U_0 U_2 / N \\ (0, -1) & \text{at rate} & U_2 \\ (-1, 1) & \text{at rate} & \lambda_{21} U_1 U_2 / N \\ (1, -1) & \text{at rate} & \delta U_2 \end{cases}
$$

If we rescale to proportions $u_i^{(N)}$ $U_i^{(N)}(t) = U_i^{(N)}$ If we rescale to proportions $u_i^{(N)}(t) = U_i^{(N)}(t)/N$, then a result of Kurtz [\[12\]](#page-21-11) implies that as $N \to \infty$, on any bounded time interval, sample paths of $(u_1^{(N)}(t), u_2^{(N)}(t))$ converge in probability to solutions of the $u_1^{(N)}(t)$, $u_2^{(N)}$
our object $2^{N(t)}(t)$ converge in probability to solutions of the following pair of equations, which we take as our object of study:

$$
u'_1 = \lambda_{10} u_0 u_1 - u_1 + \delta u_2 - \lambda_{21} u_1 u_2
$$

\n
$$
u'_2 = \lambda_{20} u_0 u_2 - u_2 - \delta u_2 + \lambda_{21} u_1 u_2.
$$
\n(2.1)

Mathematical Biosciences and Engineering Volume 16, Issue 4, 3047–3070.

We note of course that the asymptotic behaviour of these equations is, in general, not the same as the above stochastic process. That said, for such models the connections often run deeper than just finite-time convergence; see for example the work of [\[13\]](#page-21-12).

As it turns out, it is more productive to study the total proportion of hosts: $x_1 = u_1 + u_2$ and the proportion of hosts that are associated: $x_2 = u_2/x_1$. Letting $\lambda_a = \lambda_{20} - \lambda_{10}$ and $\lambda_b = -\lambda_a + \lambda_{21}$, after a bit of algebra we obtain the system

$$
x'_1 = G_1(x_1, x_2) = x_1((\lambda_{10} + \lambda_a x_2)(1 - x_1) - 1)
$$

\n
$$
x'_2 = G_2(x_1, x_2) = x_2((\lambda_a + \lambda_b x_1)(1 - x_2) - \delta).
$$
\n(2.2)

The coordinate change $u \to x$ is singular at $x_1 = 0$ but note the inverse $u_1 = x_1(1 - x_2)$, $u_2 = x_1x_2$ is smooth and onto. Thus, every solution of (2.1) is the image of a solution of (2.2) , so to understand (2.1) it is enough to study (2.2) . In terms of the original description, the host population survives if lim inf_{$t\rightarrow\infty$} $x_1(t) > 0$ and the symbiont survives if lim inf_{$t\rightarrow\infty$} $x_2(t) > 0$. The symbiont takes over if $\lim_{t\to\infty} x_2(t) = 1$. It should be clear that takeover is only possible if $\delta = 0$, and even then only if further conditions are satisfied.

From the form of [\(2.2\)](#page-3-0) it should be clear that (x_1, x_2) is a more natural choice of variables than (u_1, u_2) , since the right-hand sides in [\(2.2\)](#page-3-0) both factor nicely into density dependent form, with a good deal of symmetry between the two equations. This form is the previously mentioned "stacked" perspective, wherein the total host density, and the prevalence of the symbiont within the host population, are thought to be the best descriptors of the population demographics. The sign of λ_a separates the pathogen, neutral and mutualist cases by $\lambda_a < 0$, $\lambda_b = 0$ respectively. The sign of λ_b determines whether or not a higher host density is beneficial to the spread of the symbiont, measured by the proportion of hosts that are associated – since it depends on λ_{21} , we say the symbiont is *weakly infectious* if $\lambda_b < 0$, *neutrally infectious* if $\lambda_b = 0$ and *strongly infectious* if $\lambda_b > 0$. Distinguishing various cases according to the sign of λ_a and λ_b is crucial to making a complete analysis.

We first identify the regions of interest. Define the feasible region $\Lambda = [0, 1]^2$ and let

$$
\Lambda_{+} = \begin{cases} (0,1] \times (0,1) = \{(x_1, x_2) \in \Lambda : x_1 > 0, \ 0 < x_2 < 1 \} & \text{if } \delta = 0, \\ (0,1]^2 = \{(x_1, x_2) \in \Lambda : x_1, x_2 > 0 \} & \text{if } \delta > 0. \end{cases}
$$

In addition, let

$$
p_0 = (0, 0),
$$

\n
$$
p_1 = (a_1, 0) = (1 - 1/\lambda_{10}, 0),
$$

\n
$$
p_2 = (a_2, 1) = (1 - 1/\lambda_{20}, 1)
$$
 and
\n
$$
p_3 = (0, a_3) = (0, 1 - \delta/\lambda_a).
$$

Except for some corner cases, these are the only candidates for equilibria on the boundary of Λ . Define $\lambda_M = \max(\lambda_{10}, \lambda_{20})$ which is an upper bound on the host birth rate. Also define the conditions

(AinvU):
$$
\lambda_{20}(1 - a_1) + \lambda_{21}a_1 > 1 + \delta
$$
 and
(UinvA): $\lambda_{10}(1 - a_2) - \lambda_{21}a_2 > 1$.

(AinvU) stands for "associated invades unassociated", and is relevant iff $\lambda_{10} > 1$, in which case it corresponds to parameter values for which a small introduction of associated hosts in a stable population of unassociated hosts leads to an increase in the proportion of associated hosts. Equivalently, $G_2(p_1 + \epsilon e_2) > 0$ for small $\epsilon > 0$, where $e_2 = (0, 1)$. (UinvA) is analogous, and is relevant iff $\lambda_{20} > 1$ and $\delta = 0$.

The following is the main result of this article. Since the classification is somewhat lengthy we begin with a brief summary. There are four main cases: $(E),(UH),(AH),(C)$, and two exceptional cases: (RS) and (B). In every case except for redundant symbiont (RS), every trajectory in Λ_{+} , the interior of the feasible region, tends to one of at most three equilibria, and tends to a unique equilibrium when bistability (B) does not occur. When (RS) and (B) do not occur, the system settles into one of the following four states:

- extinction of both host and symbiont (E),
- survival of host and extinction of symbiont (UH),
- survival of host and complete adoption of the symbiont (AH), which is only possible if $\delta = 0$, and
- survival of host, with stable coexistence of associated and unassociated hosts (C).

Case (RS) is defined precisely as $\lambda_{10} = \lambda_{20} > 1$, $\delta = \lambda_{21} = 0$ and would otherwise correspond to (UH). Case (B) is defined later, under section heading Subcase 5b, in terms of intersections of the nullclines. It falls strictly within the range of parameter values satisfying $\lambda_a > 0$ and $\lambda_b > 0$, that is a strongly infectious, mutualist symbiont, and would otherwise correspond to either (E) or (UH).

Theorem 2.1. *The following six cases include all parameter values. We begin with two exceptional cases.*

- *1. Redundant symbiont (RS). Suppose* $\lambda_{10} = \lambda_{20} > 1$ *and* $\delta = \lambda_a = \lambda_{21} = 0$ *, which would otherwise correspond to (UH). For all* $x \in \Lambda_+$, $\lim_{t\to\infty} \phi_1(t, x) = a_1$ *and* $t \mapsto \phi_2(t, x)$ *is constant.*
- *2. Bistability* (*B*). This occurs for a non-empty set of parameter values satisfying $min(\lambda_a, \lambda_b) > 0$, $\delta > 0$, that would otherwise correspond to (E) if $\lambda_{10} \leq 1$, or to (UH) if $\lambda_{10} > 1$. Moreover, the set *of values* (δ, λ_{21}) *that gives* (B) *is non-empty iff* $\lambda_{20} > \max(1, \lambda_{10}^2)$ *.*

There are three subcases: onset (OB), marginal (MB) and non-marginal (NB) bistability. (NB) is an open, contractible set of parameter values and (OB), (MB) are piecewise smooth and on the boundary of (NB). In each subcase there is an interior equilibrium $\bar{x} \in \Lambda_+$ *satisfying* max(0, *a*₁) < $\bar{x}_1 < a_2.$

- *i*) *Onset of bistability (OB).* $\lim_{t\to\infty} \phi(t, x) = \bar{x}$ *for* $x \in \Lambda_+$ *.*
- *ii) Marginal bistability (MB). There is a set* $U \subset \Lambda_+$ *, open in* Λ_+ *and not containing* \bar{x} *, such that*

$$
\lim_{t \to \infty} \phi(t, x) = (\max(0, a_1), 0) \text{ for } x \in U, \text{ and}
$$

$$
\lim_{t \to \infty} \phi(t, x) = \bar{x} \text{ for } x \in \Lambda_+ \setminus U.
$$

iii) *Non-marginal bistability (NB). There are* $\bar{x} \in U_2$, $\bar{y} \in \Lambda \setminus (U_1 \cup U_2)$ *, with* max $(0, a_1) < \bar{y}_1$ < $\bar{x}_1 < a_2$ *and* max $(0, a_3) < \bar{y}_2 < \bar{x}_2 < 1$ *, and disjoint sets* $U_1, U_2 \subset \Lambda_+$ *both open in* Λ_+ *, with* $\Lambda_+ \setminus (U_1 \cup U_2)$ *the stable manifold of* \bar{y} *, such that*

> lim*t*→∞ ^φ(*t*, *^x*) ⁼ (max(0, *^a*1), 0) *for x* [∈] *^U*¹, $\lim_{t\to\infty}\phi(t,x) = \bar{x}$ *for* $x \in U_2$ *, and* $\lim_{t\to\infty}\phi(t,x) = \bar{y}$ for $x \in \Lambda_+ \setminus (U_1 \cup U_2)$.

Figure 1. Diagram indicating for which values of λ_{10} , λ_{20} there exist δ , λ_{21} such that (B) (on the left) or (E) , (C) , (UH) and (AH) (on the right) can occur.

Suppose (*RS*) and (*B*) do not hold. Then, there exists $\bar{x} \in \Lambda$ such that $\lim_{t\to\infty} \phi(t, x) = \bar{x}$ for all $x \in \Lambda_+$. *Assuming (RS) and (B) do not hold, four cases are possible.*

- *1. Extinction.* (*E)* $\bar{x} = (0, \max(0, a_3))$, if $\lambda_{10} \le 1$ *and* $\lambda_{20} \le 1 + \delta$.
- *2. Survival and coexistence of associated and unassociated host. (C)*

 $max(0, min(a_1, a_2)) < \bar{x}_1$ < $max(a_1, a_2)$ *and* $max(0, a_3) < \bar{x}_2 < 1$, *if*

- *(a)* δ > ⁰ *and either*
	- *i.* $\lambda_{10} \leq 1$ *and* $\lambda_{20} > 1 + \delta$ *, or*
	- *ii.* $\lambda_{10} > 1$ *and (AinvU) holds, or*
- *(b)* δ ⁼ ⁰ *and either*
	- *i.* $\lambda_{10} \leq 1$, $\lambda_{20} > 1$ *and (UinvA) holds,*
	- *ii.* $\lambda_{10} > 1$, $\lambda_{20} \leq 1$ *and (AinvU) holds, or*
	- *iii.* min($\lambda_{10}, \lambda_{20}$) > 1, (AinvU) holds and (UinvA) holds.
- *3. Survival of unassociated host only. (UH)* $\bar{x} = p_1$ *if* $\lambda_{10} > 1$ *and (AinvU) does not hold.*
- *4. Survival of associated host only.* $(AH) \bar{x} = p_2$ *if* $\delta = 0$, $\lambda_{20} > 1$ *and (UinvA) does not hold.*

It is worth noting the behaviour under (OB) is the same as in (C). The reason why we do not include it in (C) is because it occurs for parameter values that would otherwise satisfy either (E) or (UH).

In Figure [2](#page-6-0) we show streamlines together with nullclines in a few different cases.

Theorem [2.1](#page-4-0) gives a complete picture of the dynamics, with the possible objections that

• the exact parameter ranges for (OB),(MB), and (NB) are not specified, and

Figure 2. Streamlines in the (x_1, x_2) plane for $\lambda_{10} = 0.5$, $\lambda_{20} = 4$, $\lambda_{21} = 12$ and $\delta = 2, 4$ and 6 from left to right, corresponding to coexistence, bistability subcase (NB) and extinction, respectively. x_1 nullcline in blue, x_2 nullcline in red.

• a precise formula for interior equilibria is not given.

In the case of (MB) and interior equilibria this is because they are determined by intersection conditions between curved nullclines on the interior of Λ. Despite this, in Subcase 5b we give a fairly nice formula for (OB), which leads to the iff condition $\lambda_{20} > \max(1, \lambda_{10}^2)$ for the possibility of bistability
given in Theorem 2.1. We also give a sotisfying characterization of the values of (δ, λ_{12}) such that (NB) given in Theorem [2.1.](#page-4-0) We also give a satisfying characterization of the values of (δ, λ_{21}) such that (NB) occurs.

Before moving onto the proof we unpack somewhat the result of Theorem [2.1,](#page-4-0) tying it to the original motivation. We begin with (B) since it is the most unusual case. Ignoring (OB) and (MB) since they are boundary cases, when (NB) occurs there are two stable equilibria; the first gives either extinction of the host (if $\lambda_{10} \le 1$) or survival without symbiont (if $\lambda_{10} > 1$), while the second is a coexistence equilibrium at a higher host density. The case $\lambda_{10} \le 1$ is obligate mutualism, as for example with termites and their protozoa [\[14\]](#page-21-13), since neither the host nor the symbiont can survive without the other; this has been previously observed mathematically in, for example, [\[15\]](#page-21-14). The case $\lambda_{10} > 1$ is facultative mutualism, whereby the host is able to survive without the symbiont, but still benefits from its introduction. We note that, in order for (B) to occur, not only must the symbiont be a mutualist, but because of the condition $\lambda_{20} > \max(1, \lambda_{10}^2)$, the benefit it imparts must be sufficiently high, and quite significant if λ_{10} is fairly large. On the other hand, since $\lambda_b = -\lambda_{21} + \lambda_a$, its tendency λ_{21} to spread through the host population must also exceed the benefit $\lambda_a = \lambda_{20} - \lambda_{10}$ that it imparts to the host birth rate. Thus bistability can only occur when we have a very helpful, yet selfish mutualist in the sense that the significant benefit it provides is still exceeded by its rate of spread through the host population. The latter trait can also be viewed as a strategy for survival which is more focused on horizontal transmission than on vertical transmission.

When we have either a pathogen (λ_a < 0), or a mutualist that does more to aid host survival than to spread itself through the host population $(\lambda_b < 0)$, (B) does not occur. In this case there is a single equilibrium attracting all solutions in the interior of the feasible region, which is determined by the location of boundary equilibria (survival of unassociated hosts or associated hosts, in isolation) together with the invasion conditions (AinvU) or (UinvA), and whether or not the recovery rate is zero. In particular, it is straightforward to determine whether we have host extinction (E), survival of unassociated host only (UH), coexistence (C) , or survival of associated host only (AH) , so we leave it to the reader to inspect each case. This description, of course, bears the exception of the trivial case (RS) that essentially corresponds to two equally viable species that don't interact except in their competition for space.

The remainder of the paper is organized as follows. In Section [3](#page-7-0) we recall some basic theory of plane systems of differential equations. In Section [4](#page-8-0) we take care of the dynamics on $\Lambda \setminus \Lambda_{+}$, which consists of invariant lines on the boundary of the feasible region. In Section [5](#page-8-1) we study the dynamics on the interior of Λ , in several steps. We first study special parameter values where [\(2.2\)](#page-3-0) separates into a simpler form. We then determine boundary equilibria and their stability. Next we study the nullclines, and move on to study the itinerary of trajectories on regions delineated by the nullclines. We then give a sufficient condition to rule out periodic orbits, homoclinic orbits and heteroclinic cycles, using Dulac's criterion. At this point, we have the tools to study the dynamics in the remaining cases where (2.2) does not assume a simple form, and we complete the proof.

3. Background and definitions

We first recall some basic theory that can be found in [\[16\]](#page-22-0), specialized to the present context. Let $F \in C^1(\mathbb{R}^2, \mathbb{R}^2)$. Corresponding to *F* there exists a C^1 function $\phi : S \to \mathbb{R}^2$ called the *flow*, defined on an open $S \subset \mathbb{R} \times \mathbb{R}^2$ containing $\{0\} \times \mathbb{R}^2$ and satisfying

$$
\phi(0, x) = x
$$
 and $\partial_t \phi(t, x) = F(\phi(t, x))$ for $(t, x) \in S$.

In particular, for fixed *x*, $t \mapsto \phi(t, x)$ solves the initial value problem

$$
y' = F(y), \quad y(0) = x
$$

in an open time interval around 0. Letting $\tau_+(x) = \sup\{t : (t, x) \in S\}$ and $\tau_-(x) = \inf\{t : (t, x) \in S\}$, if we require that for every $x \in \mathbb{R}^2$,

- 1. $\tau_-(x) = -\infty$ or $\lim_{t \to \tau_-(x)} ||\phi(t, x)|| = \infty$ and that
- 2. $\tau_+(x) = \infty$ or $\lim_{t \uparrow \tau_+(x)} ||\phi(t,x)|| = \infty$,

then ϕ exists and is the unique function with the stated properties. In particular, for every $x \in \mathbb{R}^2$ the
above initial value problem has a unique solution forward and backward in time, to either $+t = \infty$ or above initial value problem has a unique solution forward and backward in time, to either $\pm t = \infty$ or until it diverges. Given $x \in \mathbb{R}^2$, the corresponding *trajectory* Γ is defined as

$$
\Gamma(x) = \{ \phi(t, x) : \tau_-(x) < t < \tau_+(x) \},
$$

and the sets $\{\Gamma(x): x \in \mathbb{R}^2\}$ form a partition of \mathbb{R}^2 into trajectories. We are also interested in the *positive semi-trajectory*

$$
\Gamma^+(x) = \{ \phi(t, x) : 0 \le t < \tau_+(x) \}.
$$

We say that a set $E \subset \mathbb{R}^2$ is *invariant* if $x \in E$ implies $\Gamma(x) \subset E$, and *forward invariant* if $x \in E$ implies $\Gamma^+(x) \subset E$. If *E* is both bounded and forward invariant, the above implies $\tau^+(x) = \infty$ for every $x \in E$.
In this case we are interested in the *omega-limit* set of points $x \in E$ defined by In this case we are interested in the *omega-limit set* of points $x \in E$, defined by

$$
\omega(x) = \{ y : y = \lim_{n \to \infty} \phi(t_n, x) \text{ for some sequence } (t_n) \text{ with } \lim_{n \to \infty} t_n = \infty \}
$$

and contained in the closure of *E*. Omega-limit sets are known to be closed, connected, invariant, and made up of trajectories, which can be of a few types, including the following. An *equilibrium point* is a point *x* such that $F(x) = 0$. A *periodic orbit* is the positive semi-trajectory of a point *x* satisfying $\phi(T, x) = x$ for some $T > 0$, and a *separatrix cycle* is the union of finitely many equilibrium points p_1, \ldots, p_m for some $m \ge 1$ and trajectories $\Gamma(x_i)$, $i = 1, \ldots, m$ satisfying

$$
\lim_{t \to \infty} \phi(-t, x_i) = p_i \quad \text{for} \quad i = 1, \dots, m,
$$

\n
$$
\lim_{t \to \infty} \phi(t, x_i) = p_{i+1} \quad \text{for} \quad i = 1, \dots, m-1 \text{ and}
$$

\n
$$
\lim_{t \to \infty} \phi(t, x_m) = p_1.
$$
\n(3.1)

A separatrix cycle with *^m* ⁼ 1 is a *homoclinic orbit*, and with *^m* > 1 is a *heteroclinic cycle*. It should be clear from the definition that if $\omega(x)$ is a single point, then $\lim_{t\to\infty} \phi(t, x) = \omega(x)$.

4. Preliminary analysis

We return to (2.2) and begin by studying the dynamics on invariant boundary lines.

Dynamics on $\Lambda \setminus \Lambda_+$. Let $L_1 = \{x_2 = 0\}$, $L_2 = \{x_1 = 0\}$ and $L_3 = \{x_2 = 1\}$, so that

$$
\Lambda_{+} = \begin{cases} \Lambda \setminus (L_1 \cup L_2) & \text{if } \delta > 0 \\ \Lambda \setminus (L_1 \cup L_2 \cup L_3) & \text{if } \delta = 0. \end{cases}
$$

The lines L_1, L_2 are invariant, as is L_3 if $\delta = 0$. On L_1 the equilibria are p_0 and p_1 . Since $G_1 =$ $x_1(\lambda_{10}(1 - x_1) - 1)$ if $x_2 = 0$, the segment {0 < $x_1 \le 1$, $x_2 = 0$ } is forward invariant and for *x* in that segment,

$$
\lim_{t\to\infty}\phi(t,x)=(\max(0,a_1),0).
$$

On L_2 the equilibria are p_0 and p_3 . Since $G_2 = x_2(\lambda_a(1 - x_2) - \delta)$ if $x_1 = 0$, the segment { $x_1 = 0$, 0 < $x_2 \leq 1$ is forward invariant and for *x* in that segment,

$$
\lim_{t \to \infty} \phi(t, x) = (0, \max(0, a_3)).
$$

On L_3 , $G_2 = -\delta x_2$, so if $\delta = 0$ that line is invariant and the equilibria are p_3 and p_2 . Since $G_1 =$ $x_1(\lambda_{20}(1 - x_1) - 1)$ if $x_2 = 1$, the segment {0 < $x_1 \le 1$, $x_2 = 1$ } is forward invariant and for *x* in that segment,

$$
\lim_{t \to \infty} \phi(t, x) = (\max(0, a_2), 1).
$$

Forward invariance of Λ , Λ ₊. Note that on the line { $x_1 = 1$ }, $G_1 = -1 < 0$, and if $\delta > 0$ then $G_2 < 0$ on $\{x_1 > 0, x_2 = 1\}$. Therefore Λ is forward invariant. This is because every side of $[0, 1]^2$ is either
invariant or else is such that for x on that side $\phi(t, x) \in (0, 1)^2$ for small enough $t > 0$, which means invariant, or else is such that for *x* on that side, $\phi(t, x) \in (0, 1)^2$ for small enough $t > 0$, which means
that trajectories cannot exit [0, 1]² along any of its four sides. The same is true if we remove any that trajectories cannot exit $[0, 1]^2$ along any of its four sides. The same is true if we remove any number of invariant lines from Λ . In particular, Λ is also forward invariant. number of invariant lines from Λ. In particular, $Λ_+$ is also forward invariant.

5. Dynamics on Λ_+

Next we determine $\omega(x)$ for $x \in \Lambda_+$, proving Theorem [2.1.](#page-4-0) By forward invariance $\omega(x)$ \subset Cl(Λ ₊) = Λ , where Cl denotes the closure. The analysis is broken up into five broad cases. In each situation we point out which case in Theorem [2.1](#page-4-0) ((RS), (OB), (MB), (NB), (E), (C), (UH), (AH)) applies.

Case 1 – low host birth rate: $\lambda_M = \max\{\lambda_{10}, \lambda_{20}\} \le 1$ (E). If $x \in \Lambda$ then $G_1 \le x_1(\lambda_M(1 - x_1) - 1)$. So, $\lambda_M \le 1$ implies $G_1 \le -x_1^2$ \int_{1}^{2} and $\lim_{t \to \infty} \phi_1(t, x) = 0.$

Case 2 – neutral symbiont: $\lambda_M > 1$, $\lambda_a = 0$. In this case $\lambda_{10} = \lambda_{20} > 1$ and $\lambda_b = \lambda_{21}$, so

$$
G_1 = x_1(\lambda_{10}(1 - x_1) - 1)
$$
 and $G_2 = x_2(\lambda_{21}x_1(1 - x_2) - \delta)$.

Since $\lambda_{10}, \lambda_{20} > 1$, $a_1, a_2 > 0$. Moreover if $x \in \Lambda$ then $\lim_{t \to \infty} \phi_1(t, x) = a_1$. Since $\lambda_{10} = \lambda_{20}$, we have $\lambda_{20}(1 - a_1) = \lambda_{10}(1 - a_2) = 1$, so (AinvU) $\Leftrightarrow \lambda_{21}a_1 > \delta$ and (UinvA) $\Leftrightarrow \lambda_{21}a_2 > 0 \Leftrightarrow \lambda_{21} > 0$.

If $\delta = \lambda_{21} = 0$ (RS) then $G_2 = 0$, so $t \mapsto \phi_2(t, x)$ is constant. If $\delta = 0$ and $\lambda_{21} > 0$ (AH) then $G_2 > 0$ for $x_1 > 0$ and $0 < x_2 < 1$, so $\lim_{t \to \infty} \phi_2(t, x) = 1$ for $x \in \Lambda_+$. If $\delta > 0$ and $\lambda_{21} = 0$ (UH) then $G_2 < 0$ for $x_2 > 0$, so $\lim_{t \to \infty} \phi_2(t, x) = 0$ for $x \in \Lambda_+$. If δ , $\lambda_{21} > 0$, then using the limiting value of ϕ_1 we find that

$$
\lim_{t \to \infty} \phi_2(t, x) = \bar{x}_2 = \max(0, 1 - \frac{\delta}{\lambda_{21} a_1}) \text{ for } x \in \Lambda_+.
$$

If $\lambda_{21}a_1 \le \delta$ (UH) then $\bar{x}_2 = 0$ and if $\lambda_{21}a_1 > \delta$ (C) then $0 < \bar{x}_2 < 1$.

Case 3 – neutrally infectious symbiont: $\lambda_M > 1$, $\lambda_a \neq 0$, $\lambda_b = 0$. In this case $\lambda_{10} \neq \lambda_{20}$ and $\lambda_{10} = \lambda_{20} - \lambda_{21}$. This forces $\lambda_{20} > \lambda_{10}$ so that $\lambda_a > 0$ and $\lambda_{21} > 0$. Multiplying on both sides by λ_{10} ,

$$
(\text{AinvU}) \Leftrightarrow \lambda_{20} + \lambda_{21}(\lambda_{10} - 1) > (1 + \delta)\lambda_{10} \Leftrightarrow \lambda_{21}\lambda_{10} > \delta\lambda_{10}
$$

which is $\Leftrightarrow \lambda_{10} > 0$ and $\lambda_{21} > \delta$. Multiplying on both sides by λ_{20} ,

$$
(Uinv A) \Leftrightarrow \lambda_{10} - \lambda_{21}(\lambda_{20} - 1) > \lambda_{20} \Leftrightarrow -\lambda_{21}\lambda_{20} > 0,
$$

so (UinvA) does not hold. We find

$$
G_1 = x_1((\lambda_{10} + \lambda_a x_2)(1 - x_1) - 1) \text{ and } G_2 = x_2(\lambda_a(1 - x_2) - \delta),
$$

so

$$
\lim_{t \to \infty} \phi_2(t, x) = \bar{x}_2 = \max(0, 1 - \delta/\lambda_a)
$$
 for $x \in \Lambda_+$.

Then,

if
$$
\delta = 0
$$
 then $\bar{x}_2 = 1$, (AH)
if $0 < \delta < \lambda_a$ then $0 < \bar{x}_2 < 1$ (C) or (E)
and if $\delta \ge \lambda_a$ then $\bar{x}_2 = 0$ (UH) or (E).

Plugging \bar{x}_2 into the equation for G_1 we find

$$
\lim_{t \to \infty} \phi_1(t, x) = \max(0, 1 - 1/(\lambda_{10} + \lambda_a \bar{x}_2)) \text{ for } x \in \Lambda_+.
$$

Mathematical Biosciences and Engineering Volume 16, Issue 4, 3047–3070.

In particular, $\phi_1 \rightarrow 0$ exactly on the condition $\lambda_{10} + \lambda_a \bar{x}_2 \le 1$, which

if $\delta = 0$ is $\lambda_{20} \le 1$ thus empty,
if $0 < \delta < 1$ is $\lambda_{20} < 1 + \delta$ if $0 < \delta < \lambda_a$ is $\lambda_{20} \le 1 + \delta$ (E)
if $\delta > \lambda_a$ is $\lambda_a < 1$ (E) and if $\delta \ge \lambda_a$ is $\lambda_{10} \le 1$ (E).

This concludes the easy cases. To treat cases with $\lambda_a, \lambda_b \neq 0$, more effort is required. We will start by studying equilibria on the boundary of Λ_{+} , then nullclines on Λ_{+} , then itineraries on Λ_{+} . Then, we will be ready to consider the remaining cases.

Equilibria on *L*. Let $L = L_1 \cup L_2$ if $\delta > 0$ and $L = L_1 \cup L_2 \cup L_3$ if $\delta = 0$, so that $\Lambda_+ = \Lambda \setminus L$. Since $\lambda_a, \lambda_b \neq 0$, p_0, p_1, p_2, p_3 are the possible equilibria (eq) on *L*. Letting $a_1 = 1 - 1/\lambda_{10}$, $a_2 =$ $1 - 1/\lambda_{20}, a_3 = 1 - \delta/\lambda_a,$

- 1. $p_0 = (0, 0) \in L_1 \cap L_2$ is always an eq in Λ .
- 2. $p_1 = (a_1, 0) \in L_1$ is an eq in $\Lambda \setminus \{p_0\}$ if $\lambda_{10} > 1$.
- 3. $p_3 = (0, a_3) \in L_2$ is an eq in $\Lambda \setminus \{p_0\}$ if $\delta < \lambda_a$, and if $\delta = 0$ then $p_3 = (0, 1) \in L_2 \cap L_3$.
- 4. $p_2 = (a_2, 1) \in L_3$ is an eq in $\Lambda \setminus \{p_3\}$ if $\delta = 0$ and $\lambda_{20} > 1$.

To assess stability we compute the Jacobian matrix $J(x) = (\partial_i G_i(x))_{ii}$.

$$
J(x) = \begin{pmatrix} (\lambda_{10} + \lambda_a x_2)(1 - 2x_1) - 1 & \lambda_a x_1(1 - x_1) \\ \lambda_b x_2(1 - x_2) & (\lambda_a + \lambda_b x_1)(1 - 2x_2) - \delta \end{pmatrix}.
$$

If $x \in L$ then at least one of x_1, x_2 is equal to 0 or 1, and $J(x)$ is triangular, so its eigenvalues are equal to its diagonal entries. Straightforward computation shows that

- 1. *J*(p_0) has eigenvalues λ_{10} 1 and λ_a δ ,
- 2. *J*(p_1) has eigenvalues 1 λ_{10} and *J*₂₂(p_1),
- 3. *J*(p_2) has eigenvalues 1 λ_{20} and *J*₂₂(p_2) if $\delta = 0$, and
- 4. *J*(p_3) has eigenvalues λ_{20} (1 + δ) and δ λ_a .

Since $\lambda_b = -\lambda_a + \lambda_{21}$, $\lambda_a = \lambda_{20} - \lambda_{10}$, $\lambda_{10}(1 - a_1) = 1$ and $1 - a_1 = 1/\lambda_{10}$,

$$
J_{22}(p_1) = \lambda_a + \lambda_b a_1 - \delta = \lambda_a (1 - a_1) + \lambda_{21} a_1 - \delta
$$

= $\lambda_{20} (1 - a_1) + \lambda_{21} a_1 - (1 + \delta)$
= $\lambda_{20} / \lambda_{10} + \lambda_{21} (1 - 1 / \lambda_{10}) - (1 + \delta),$

and if $\delta = 0$ then since $-\lambda_a = \lambda_{10} - \lambda_{20}$ and $\lambda_{20}(1 - a_2) = 1$,

$$
J_{22}(p_2) = -\lambda_a - \lambda_b a_2 = -\lambda_a (1 - a_2) - \lambda_{21} a_2
$$

= $\lambda_{10} (1 - a_2) - \lambda_{21} a_2 - 1$
= $\lambda_{10} / \lambda_{20} - \lambda_{21} (1 - 1 / \lambda_{20}) - 1$.

Using these calculations and the fact that $G_2(x + \epsilon e_2) = \epsilon G_2(x) + o(\epsilon)$,

i) $J_{22}(p_1) > 0$ ⇔ (AinvU) ⇔ $G_2 > 0$ just above p_1 . ii) $J_{22}(p_2) > 0$ ⇔ (UinvA) ⇔ $G_2 < 0$ just below p_2 .

Mathematical Biosciences and Engineering Volume 16, Issue 4, 3047–3070.

Recall the following three types of equilibria *p*:

- i) *Attracting*: for some $\epsilon > 0$, $|x p| < \epsilon$ implies $\lim_{t \to \infty} \phi(t, x) = p$.
- ii) *Saddle point*: the *stable manifold* and *unstable manifold*

 $W_s(p) = \{x : \lim_{t \to \infty} \phi(t, x) = p\}$ and $W_u(p) = \{x : \lim_{t \to -\infty} \phi(t, x) = p\}$

are smooth curves each consisting of *p* and a pair of trajectories, and the tangent lines to W_s , W_u at *p* are eigenvectors of $I(n)$ with eigenvalues whose real part is peoplive respectively positive at *p* are eigenvectors of $J(p)$ with eigenvalues whose real part is negative, respectively positive.

iii) *Repelling*: for some $\epsilon > 0$, $|x - p| < \epsilon$ implies $\lim_{t \to -\infty} \phi(t, x) = p$.

Given equilibrium p, let μ_1, μ_2 be the eigenvalues of $J(p)$, and recall that [hartman-grobman]

- i) *p* is attracting if (μ_1) , (μ_2) < 0,
- ii) *p* is a saddle point if $(\mu_1) < 0 < (\mu_2)$, and
- iii) *p* is repelling if $0 < (\mu_1)$, (μ_2) .

If one or both of $(\mu_i) = 0$ then p could be of any type, or none.

Nullclines.

$$
\{G_1 = 0\} = \{x_1 = 0\} \cup \gamma_1 \text{ where } \gamma_1 = \{(\lambda_{10} + \lambda_a x_2)(1 - x_1) = 1\}, \text{ and } \{G_2 = 0\} = \{x_2 = 0\} \cup \gamma_2 \text{ where } \gamma_2 = \{(\lambda_a + \lambda_b x_1)(1 - x_2) = \delta\}.
$$

 γ_1 is a hyperbola with vertical asymptote $x_1 = 1$, so instead let γ_1 be the branch with $x_1 < 1$. If $\delta = 0$ then $\gamma_2 = \{x_2 = 1\} \cup \{x_1 = -\lambda_a/\lambda_b\}$ so instead let γ_2 be the vertical line $\{x_1 = -\lambda_a/\lambda_b\}$. If $\delta > 0$ then γ_2 is a hyperbola with horizontal asymptote $x_2 = 1$, so instead let γ_2 be the branch with x_2 < 1. Both y_1 and y_2 intersect each of the invariant lines at most once. The intersection points, and conditions for existence, are

- $\gamma_1 \cap \{x_2 = 0\} = p_1$ if $\lambda_{10} > 0$,
- $\gamma_1 \cap \{x_1 = 0\} = r_1 = (0, b_1)$ with $b_1 = (1 \lambda_{10})/\lambda_a$,
- $\gamma_1 \cap \{x_2 = 1\} = p_2$ if $\lambda_{20} > 0$,
- $\gamma_2 \cap \{x_2 = 0\} = r_2 = (b_2, 0)$ with $b_2 = (\delta \lambda_a)/\lambda_b$,
- $\gamma_2 \cap \{x_1 = 0\} = p_3$, and
- $\gamma_2 \cap \{x_2 = 1\} = r_3 = (b_2, 1)$ if $\delta = 0$.

By considering what happens when $x_2 = 0$ and noting that $\{x_1 = 0\}$, γ_1 are the only null sets for G_1 when $0 \le x_1 \le 1$, it follows that $G_1 > 0$ to the left of γ_1 in Λ_+ . On the other hand, since $\partial_{x_1} G_2 = \lambda_b x_2 (1 - x_2)$, if $\lambda_b < 0$ then $G_2 > 0$ to the left of γ_2 , while if $\lambda_b > 0$ then $G_2 > 0$ to the right of γ_2 , in Λ_+ .

If $\lambda_{10} > 1$ then $b_2 = a_1$ marks the boundary between (AinvU) and its complement. Similarly, if $\lambda_{20} > 1$ and $\delta = 0$ then $b_2 = a_2$ marks the boundary between (UinvA) and its complement. Also, it's easy to check that $b_1 = a_3$ is equivalent to $\lambda_{20} = 1 + \delta$.

It's useful to think of γ_1 as the graph of the function

$$
x_2 = g_1(x_1), x_1 < 1
$$
 where $g_1(x_1) = \lambda_a^{-1} \left(\frac{1}{1 - x_1} - \lambda_{10} \right)$,

and when $\delta > 0$ to think of γ_2 as the graph of the function

$$
x_2 = g_2(x_1), x_2 < 1
$$
 where $g_2(x_1) = 1 - \frac{\delta}{\lambda_a + \lambda_b x_1}$.

If $\delta = 0$ then $\gamma_1 \cap \gamma_2$ has cardinality 0 or 1 as γ_1 has at most one intersection point with any vertical line. If $\delta > 0$ then since the sign of the pairs (g_1^{\prime}) and never coincide (to see this break up according $g_1'(x_1), g_1''$
ing to (s) $\binom{n}{1}(x_1)$ and $\binom{n}{2}$ $\binom{2}{2}(x_1), g''_2$ $2''(x_1)$ do not depend on x_1 and never coincide (to see this, break up according to $(\text{sgn}(\lambda_a), \text{sgn}(\lambda_b))$, noting that $\lambda_a + \lambda_b = \lambda_{21} \ge 0$ implies $(sgn(\lambda_a), sgn(\lambda_b)) \neq (-1, -1)$), $\gamma_1 \cap \gamma_2$ has cardinality 0, 1 or 2.

Itineraries. Let *G* = (*G*₁, *G*₂) and for a pair (*a*, *b*) let sgn(*a*, *b*) = (sgn(*a*), sgn(*b*)) ∈ {−1, 0, 1}². For each sign vector $\sigma \in \{-1, 0, 1\}^2$ let

$$
R(\sigma) = \{x \in \Lambda_+ : \text{sgn}(G(x)) = \sigma\}.
$$

Since all signs are represented, $\{R(\sigma) : \sigma \in \{-1, 0, 1\}^2\}$ is a partition of Λ_+ . Note that one or more of these sets could be empty these sets could be empty.

- The set $R(0, 0) = \gamma_1 \cap \gamma_2 \cap \Lambda_+$ consists of equilibrium points and has cardinality 0, 1 or 2.
- The sets $R(0, 1) \cup R(0, -1) = \gamma_1 \cap \Lambda_+ \setminus \gamma_2$ and $R(1, 0) \cup R(-1, 0) = \gamma_2 \cap \Lambda_+ \setminus \gamma_1$, so we refer to the union of all four as *^R*γ. Since neither hyperbola has either a horizontal or vertical tangent at any point, while *G* is horizontal or vertical on each one (or if $\delta = 0$, γ_2 is vertical while *G* is horizontal on γ_2), it follows that on R_γ , *G* is transverse to the tangent space of R_γ , so for any $x \in \Lambda_+$, the set of times $\{t : \phi(t, x) \in R_{\gamma}\}\$ is a discrete set.
- The sets $R(\sigma)$ for $\sigma \in \{-1, 1\}^2$ are open subsets of $(0, 1]^2$ that, based on the number of intersection points of α , and α , each have at most 2 connected components points of γ_1 and γ_2 , each have at most 2 connected components.

If $x \in (0, 1]^2 \setminus R(0, 0)$ then since $R(0, 0)$ is invariant and $\{t : \phi(t, x) \in R_{\gamma}\}\)$ is discrete, there is possibly finite) symbolic sequence $I(x) = (x, x_0)$ called the *itingrary* of x with each $x_0 \in R_{\gamma}\}$ a (possibly finite) symbolic sequence $I(x) = (\sigma_1, \sigma_2, \dots)$ called the *itinerary* of *x*, with each $\sigma_i \in$ $\{-1, 1\}^2$, and an increasing sequence of times $t(x) = (t_1, t_2, \dots)$, with $t_0 = 0$, and with $t_i = \infty$ iff $I(x)$ has length *i* with the property that has length *i*, with the property that

$$
\phi(t,x)\in R(\sigma_i) \text{ for } t\in (t_{i-1},t_i).
$$

Define a directed graph called the *sign graph* with vertices $\{-1, 1\}^2$ by including each directed edge σ') iff there is x such that $(L(x), L(x)) = (\sigma \sigma')$ for some $i > 1$. The sign graph is *cyclic* if there is (σ, σ') iff there is *x* such that $(I_i(x), I_{i+1}(x)) = (\sigma, \sigma')$ for some $i \ge 1$. The sign graph is *cyclic* if there is a sequence of edges $(\sigma_1, \sigma_2), \ldots, (\sigma_k, \sigma_1)$, and *acyclic* if not. Appearance of (σ, σ') in the sign graph
is also represented by $P(\sigma) \rightarrow P(\sigma')$, $P(\sigma)$ is *absorbing* if pope of the edges (σ, σ) appear on the sign is also represented by $R(\sigma) \to R(\sigma')$. $R(\sigma)$ is *absorbing* if none of the edges (σ, \cdot) appear on the sign
graph. An equilibrium point $p \in C^{1}(R(\sigma))$ is a sink for $R(\sigma)$ if for some $x \in R(\sigma)$, $I(x) = \sigma$ and graph. An equilibrium point $p \in \text{Cl}(R(\sigma))$ is a *sink* for $R(\sigma)$ if for some $x \in R(\sigma)$, $I(x) = \sigma$ and lim_{$t\rightarrow\infty$} $\phi(t, x) = p$. If *p* is a sink for $R(\sigma)$ then it is a local maximum of $x \mapsto \sigma \cdot x$ on $R(\sigma)$. The following lets us seal the fate of trajectories, when the sign graph is acyclic. The proof is given in the Appendix.

Lemma 5.1. *If I(x) is finite and its last entry is* σ *then* $\omega(x)$ *is a sink for* $R(\sigma)$ *. In particular, if the sign graph is acyclic then for all* $x \in \Lambda_+$, $\omega(x)$ *is a sink.*

Periodic orbits and cycles. Lemma [5.1](#page-13-0) makes our life easier when the sign graph is acyclic. When the sign graph is cyclic, we will use the following proposition to simplify the analysis.

Proposition 5.1. *For each* $x \in \Lambda_+$, $\omega(x)$ *is either*

- *1. an equilibrium point or*
- *2. contains a separatrix cycle intersecting an equilibrium in L.*

Proof. There are three steps.

- 1. Show there are no periodic orbits in Λ and no separatrix cycles in Λ_{+} .
- 2. Show that a separatrix cycle intersecting *L* must intersect an equilibrium in *L*.
- 3. Invoke a generalized form of the Poincaré-Bendixson theorem.

Step 1. We begin with

Lemma 5.2 (Dulac's criterion). *– Suppose we have the planar system*

$$
u' = F(u)
$$
 where $u \in \mathbb{R}^2$ and $F : \mathbb{R}^2 \to \mathbb{R}^2$ is a C^1 function.

Let $R \subset \mathbb{R}^2$ be a simply connected region in the plane and $B : \mathbb{R}^2 \to \mathbb{R}^2$ be a C^1 function (called a *Dulac function) such that the divergence of BF is non-zero and has constant sign almost everywhere on R. Then, the system* $u' = F(u)$ *has no periodic orbit and no separatrix cycle on R.*

Since most proofs of Dulac's criterion only rule out periodic orbits, we include a proof in the appendix. *No periodic orbit or separatrix cycle in* Λ_+ . Since the restriction of $x \mapsto (x_1(1-x_2), x_1x_2)$ to $\{0 \le x_1 \le x_2\}$ 1, $0 < x_2 < 1$ } maps bijectively onto $\{u_1, u_2 > 0, u_1 + u_2 \le 1\}$, periodic orbits and separatrix cycles are mapped to the same. So, to rule them out on the former set, we find a Dulac function for [\(2.1\)](#page-2-0) on the latter set. The function $B(u) = 1/(u_1u_2)$ works, since

$$
\nabla \cdot (BF) = \partial_{u_1} (\lambda_{10} u_0 / u_2 - 1 / u_2 - \lambda_{21} + \delta / u_1)
$$

+ $\partial_{u_2} (\lambda_{20} u_0 / u_1 - 1 / u_1 + \lambda_{21} - \delta / u_1)$
= $-\lambda_{10} / u_2 - \delta / u_1^2 - \lambda_{20} / u_1$

is strictly negative when $u_1, u_2 > 0$ (at least one of $\lambda_{10}, \lambda_{20}$ is positive).

If $\delta = 0$ the set $\{0 < x_1 \leq 1, \ 0 < x_2 < 1\}$ is equal to Λ_+ .

If $\delta > 0$, Λ_+ includes also the segment $\{0 < x_1 \leq 1, x_2 = 1\}$. Since $G_2 < 0$ on that segment, any trajectory that intersects it also intersects Λ^c , so is not contained in Λ .

No periodic orbit in L. If a periodic orbit *C* intersects $L = \Lambda \setminus \Lambda_+$, it intersects an invariant line on *L*. Since *C* is a single trajectory, it is a subset of that invariant line. Since *C* is the continuous injective image of a circle and *L* is either a pair of orthogonal lines, or two parallel lines and an orthogonal line, this cannot occur.

Step 2. A separatrix cycle is the continuous injective image of a circle. If it intersects *L*, it intersects an invariant line on *L*, so it must contain an equilibrium point on that line. If it did not, it would consist of a single trajectory on that line.

Step 3. This step is a trivial application of the following result.

Theorem 5.1 (Generalized Poincaré-Bendixson Theorem). – Let $E \subset \mathbb{R}^2$ be an open set and $F \in$ $C^1(E, \mathbb{R}^2)$. Suppose the system $x' = F(x)$ has a positive semi-trajectory $\Gamma^+(x)$ contained in a compact set $K \subset F$ with the property that $F(x) = 0$ for at most finitely many $x \in K$. Then $\omega(x)$ is either *set* $K \subset E$ with the property that $F(x) = 0$ for at most finitely many $x \in K$. Then, $\omega(x)$ is either

- *1. an equilibrium point,*
- *2. a periodic orbit, or*
- *3. contains a separatrix cycle.*

Proof. This is given by Theorem 2 in Section 3.7 of [\[17\]](#page-22-1), and the comments that follow. □

This concludes the proof of Proposition [5.1.](#page-13-1) \Box

Trajectories. We are now ready to study dynamics in the remaining cases. We break up our analysis according to the sign of λ_a , λ_b . The interested reader will note that Case 3 above can also be covered here using the same methods, although it does not give an explicit formula for the interior equilibrium, when there is one.

Case 4 – pathogen: $\lambda_M > 1$, $\lambda_a < 0$, $\lambda_b \neq 0$. In this case, $\lambda_{10} > \max(1, \lambda_{20})$ and $\lambda_b = -\lambda_a + \lambda_{21} > 0$, so $a_1 = 1 - 1/\lambda_{10} > 0$, $b_2 = (\delta - \lambda_a)/\lambda_b > 0$ and $G_2 > 0$ to the right of γ_2 . g_1 is decreasing and concave, and if $\delta > 0$ then g_2 is increasing and concave, while if $\delta = 0$ then $\gamma_2 = \{x_1 = b_2\}$ – in either case γ_2 lies to the right of $\{x_1 < b_2\}$ so $G_2 < 0$ on this set. Since $G_1 > 0$ near p_0 it attracts no points in $Λ_+$, and since $p_3 = (0, a_3)$ with $a_3 = 1 - δ/λ_0 ≥ 1$, if $p_3 ∈ Λ$ then $δ = 0$, $p_3 = (0, 1)$ and $G_2 < 0$ in a Λ_{+} -neighbourhood of p_3 , which means p_3 attracts no points in Λ_{+} .

If *^b*² [≥] *^a*¹ then (AinvU) does not hold, which gives (UH). *^R*(−1, 1) [→] *^R*(−1, [−]1) [→] *^R*(1, [−]1) appear from right to left. If $p_2 \in \Lambda$ then since $\lambda_{20} < \lambda_{10}$, $a_2 < a_1 \le b_2$ so $G_2 < 0$ in a Λ_+ -neighbourhood of p_2 . Since the sign graph is acyclic, every solution tends to a sink, and the only candidate is p_1 .

If $\delta > 0$ and $b_2 < a_1$ (AinvU) which gives (a.ii) of (C), or $\delta = 0$ and $a_2 < b_2 < a_1$ which gives (AinvU) and (UinvA) and thus (b.iii) of (C) if $a_2 > 0$ or (b.ii) of (C) if $a_2 \le 0$, then (C) holds. $\gamma_1 \cap \gamma_2$ is a unique point $p_4 \in \Lambda_+$ and p_1 is a saddle, as is p_2 if $\delta = 0$. The sign graph is cyclic, so we use Proposition [5.1](#page-13-1) to conclude that solutions tend to p_4 provided no separatrix cycles on Λ contain an equilibrium on *L* – let *C* denote a possible candidate. Since p_1 is a saddle, $W_s(p_1) = \{0 < x_1, x_2 = 0\}$ so $p_1 \in C$ implies p_0 ∈ *C*. Since p_0 is a saddle with $W_s(p_0) = \{x_1 = 0, x_2 < a_3 = 1 - \delta/\lambda_a\}$, p_0 ∈ *C* implies $p_3 \in C$. If $\delta > 0$, $p_3 \notin \Lambda$ so *C* cannot exist, otherwise $\delta = 0$ and $p_0 \in C$ implies $p_3 = (0, 1) \in C$. If $\lambda_{20} < 1$ then $W_s(p_3) = \{a_2 < x_1, x_2 = 1\}$ and *C* cannot exist. If $\lambda_{20} = 1$ then since γ_2 does not intersect p_3 , still $G_2 < 0$ in a Λ_+ -neighbourhood of p_3 , so $W_s(p_3)$ is disjoint from Λ_+ which means $W_s(p_3) = \{0 \le x_1, x_2 = 1\}$ and *C* cannot exist. If $\lambda_{20} > 1$ then p_3 is repelling, and *C* cannot exist. If $δ > 0$ then *p*₂ ∉ Λ, and if $δ = 0$ and *p*₂ ∈ Λ then *p*₂ is a saddle with *W_s*(*p*₂) = {0 < *x*₁, *x*₂ = 1}, so

 $p_2 \in C$ implies $p_3 \in C$. Since p_3 is repelling, *C* cannot exist.

If $\delta = 0$ and $b_2 \le a_2 < a_1$ then (AinvU) holds but (UinvA) does not, which gives (AH). $R(1, -1) \rightarrow R(1, 1) \rightarrow R(-1, 1)$ appear from left to right, and p_1 is a saddle. Since the sign graph is acyclic, solutions tend to a sink, and the only candidate is *p*2.

Case 5 – mutualist: $\lambda_M > 1$, $\lambda_a > 0$, $\lambda_b \neq 0$. In this case $\lambda_{20} > \max(1, \lambda_{10})$ so $a_2 > 0$, and g_1 is increasing and convex. Notice that g_1 intersects the set $\{x_1 = 0, 0 \le x_2 < 1\} \cup \{0 \le x_1 < a_2, x_2 = 0\}$ at exactly one point, which is p_1 if $\lambda_{10} > 1$, p_0 if $\lambda_{10} = 1$ and $r_1 = (0, b_1) = (0, (1 - \lambda_{10})/\lambda_a)$ if $\lambda_{10} < 1$.

Subcase 5a – weakly infectious mutualist: $\lambda_b < 0$. If $\delta > 0$ then g_2 is decreasing and concave, and recall $\gamma_2 \cap \{x_2 = 0\}$ contains only the single point $(b_2, 0)$ with $b_2 = (\delta - \lambda_a)/\lambda_b$, and $\gamma_2 \cap \{x_1 = 0\}$ contains only the single point $p_3 = (0, a_3)$ with $a_3 = 1 - \delta/\lambda_a$. $\gamma_1 \cap \gamma_2 \cap \Lambda_+$ contains exactly one point if either $0 < a_1 < b_2$ or $a_1 \le 0$ and $a_3 > b_1$, both of which correspond to (C), and otherwise is empty. Notice that

$$
a_3 > b_1 \Leftrightarrow (\lambda_a - \delta)/\lambda_a > (1 - \lambda_{10})/\lambda_a \Leftrightarrow \lambda_a - \delta > 1 - \lambda_{10} \Leftrightarrow \lambda_{20} > 1 + \delta.
$$

If $\delta = 0$ then $b_2 = -\lambda_a/\lambda_b > 0$ and $p_3 = (0, 1)$ is a saddle, so attracts no points in Λ_+ . If $a_1 < b_2 < a_2$, which corresponds to (C), then $\gamma_1 \cap \gamma_2 \cap \Lambda_+$ contains exactly one point, otherwise $\gamma_1 \cap \gamma_2 \cap \Lambda_+$ is empty.

If $\delta > 0$ and $\gamma_1 \cap \gamma_2 \cap \Lambda_+$ is empty then either $a_1 > 0$ and $a_1 \ge b_2$ (UH), or $a_1 \le 0$ and $a_3 \le b_1$ which implies $\lambda_{10} \leq 1$ and $\lambda_{20} \leq 1 + \delta$ (E). There are at most three sign regions in Λ (two if $b_2 \leq 0$): $R(1, 1) \rightarrow R(1, -1) \rightarrow R(-1, -1)$, appearing in that order from left to right. In case of (UH), p_0 is a saddle and $p_1 \in \Lambda$, and in case of (E), p_0 is the only equilibrium in Λ .

If $\gamma_1 \cap \gamma_2 \cap \Lambda_+$ is not empty then it is a unique point $p_4 \in \Lambda_+$. If $a_1 \leq 0$ then p_0 is a saddle, otherwise *p*₀ is repelling and *p*₁ is a saddle, and if $\delta = 0$ then *p*₂, *p*₃ are saddles. A similar argument as before shows there is no periodic orbit or separatrix cycle in Λ. Thus, all solutions tend to *p*4.

If $\delta = 0$ and $0 < b_2 \le a_1 < a_2$ (UH), $R(1, 1) \rightarrow R(1, -1) \rightarrow R(-1, -1)$ appear from left to right. Since $a_1 > 0$, p_0 is a saddle and $p_1 \in \Lambda$, while p_2, p_3 are saddles, so solutions tend to p_1 .

If $\delta = 0$ and $a_1 < a_2 \le b_2$ (AH), then $R(-1, -1) \rightarrow R(-1, 1) \rightarrow R(1, 1)$ appear from right to left, p_0 is repelling, and p_1 , p_3 are saddles, so solutions tend to p_2 .

Subcase 5b – strongly infectious mutualist: $\lambda_b > 0$. In this case (B) can occur, so we take a moment to define precisely the three conditions in terms of the nullclines:

- 1. (OB) occurs when g_1, g_2 intersect at two points in Λ , one in Λ_+ and the other on the subset ${x_1 = 0, 0 \le x_2 \le 1} \cup {0 \le x_1 \le 1, x_2 = 0}$ of the boundary.
- 2. (MB) occurs when g_1, g_2 intersect tangentially at one point in Λ_+ .
- 3. (NB) occurs when g_1, g_2 intersect at two points, both in Λ_{+} .

As we go through the various cases we will point out when one of the above might occur; in particular, we'll see that they all occur for values that would otherwise correspond to (E) or (UH). We'll also describe the dynamics in all possible cases. We'll then show that each of (OB),(NB),(MB) does occur, make some effort to characterize them, and show (NB) is simply connected.

If $\delta = 0$ then $G_2 = x_2((\lambda_a + \lambda_b x_1)(1 - x_2) > 0$ on Λ_+ , so (AH) holds. $b_1 = -\lambda_a/\lambda_b < 0$ and $R(-1, 1) \rightarrow R(1, 1)$ appear from right to left. Because of the sign of G_2 , p_0 (and p_1 if $a_1 > 0$) attract no points, and $p_3 = (0, 1)$ is a saddle. Since the sign graph is acyclic, solutions tend to a sink, and the only candidate is p_2 .

For the rest we assume $\delta > 0$. g_2 is increasing and concave, so γ_2 intersects the set ${x_1 = 0, 0 ≤ x_2 < 1} ∪ {x_1 ≥ 0, x_2 = 0}$ at exactly one point, which is $p_3 = (0, a_3) = (0, 1 - \delta/\lambda_a)$ if $\delta < \lambda_a$, $p_0 = (0, 0)$ if $\delta = \lambda_a$ and $r_2 = (b_2, 0) = ((\delta - \lambda_a)/\lambda_b, 0)$ if $\delta > \lambda_a$.

If either $a_1 > \max(b_2, 0)$, or $a_1 \le 0$ and $a_3 > \max(b_1, 0)$, then $\gamma_1 \cap \gamma_2 \cap \Lambda_+$ is a unique point $p_4 \in \Lambda_+$. A similar argument as before shows that all solutions tend to p_4 . The first option gives $\lambda_{10} > 1$ and (AinvU) so corresponds to (C). For the second option, note that $a_1 \le 0 \Leftrightarrow \lambda_{10} \le 1$ and

$$
a_3 > b_1 \Leftrightarrow \lambda_a - \delta > 1 - \lambda_{10} \Leftrightarrow \lambda_{20} > 1 + \delta,
$$

so it also corresponds to (C).

If $b_2 = a_1 > 0$ (UH), then γ_1, γ_2 intersect at p_1 . If g'_1
regions $R(1, -1)$, $R(-1, -1)$, $R(-1, 1)$ appear from $y'_1(a_1) \geq g'_2$ $\gamma_2'(a_1)$, then $\gamma_1 \cap \gamma_2 \cap \Lambda_+$ is empty, and the sign
the with $R(1, -1)$, $R(-1, 1) \rightarrow R(-1, -1)$, so regions $R(1, -1)$, $R(-1, -1)$, $R(-1, 1)$ appear from left to right with $R(1, -1)$, $R(-1, 1)$ → $R(-1, -1)$, so the sign graph is acyclic. Since $b_2 > 0$ and γ_2 is increasing, $a_3 < 0$ so $p_3 \notin \Lambda_+$ and $G_1 > 0$ near p_0 so solutions tend to p_1 . If g'_1 $y'_1(a_1) < g'_2$
ur sign reg $\gamma_2'(a_1)$, which as we'll see corresponds to (OB), then $\gamma_1 \cap \gamma_2 \cap \Lambda_+$ is gions now appear, with $R(1, -1)$, $R(-1, 1)$, $R(-1, 1)$, $R(-1, -1)$, To a single point p_4 . All four sign regions now appear, with $R(1, -1)$, $R(-1, 1) \rightarrow R(1, 1)$, $R(-1, -1)$. To show all solutions tend to p_4 it remains to show that p_1 is not a sink for any sign region. p_1 is in the closure of $R(1, -1)$, $R(-1, 1)$ and $R(1, 1)$. Since $G_2 > 0$ on $R(-1, 1)$ p_1 cannot be a sink, and p_1 is a local minimum of $x \mapsto x(1, 1)$ on $R(1, 1)$, whereas a sink is a local maximum. The only remaining candidate is $R(1, -1)$. G_2 has a double root at p_1 , which can be seen from either the transverse intersection of two *x*₂-nullclines ({*x*₂ = 0} and γ ₂) or by writing *G*₂ as a function of $\tilde{x} = x - p_1$. Writing *G* in these coordinates,

$$
G_1 = (\tilde{x}_1 + a_1)((\lambda_{10} + \lambda_a \tilde{x}_2)(1 - (\tilde{x}_1 + a_1)) - 1)
$$

= (\tilde{x}_1 + a_1)(\lambda_{10} + \lambda_a \tilde{x}_2)(\lambda_{10}^{-1} - \tilde{x}_1) - 1)
= (\tilde{x}_1 + a_1)(-\lambda_{10}\tilde{x}_1 + \lambda_a \tilde{x}_2(\lambda_{10}^{-1} - \tilde{x}_1))

and

$$
G_2 = \tilde{x}_2(\lambda_a + \lambda_b(\tilde{x}_1 + (\delta - \lambda_a)/\lambda_b)(1 - \tilde{x}_2) - \delta)
$$

= $\lambda_b \tilde{x}_1 \tilde{x}_2 (1 - \tilde{x}_2) - \delta \tilde{x}_2^2$

If a trajectory in *R*(1, −1) tends to *p*₁ then $\tilde{x}_1 \to 0^-$ and $\tilde{x}_2 \to 0^+$. If $\tilde{x}_1 < 0 < \tilde{x}_2$ are small in magnitude then by 1st-order approximation $G_1 > c_1(-\tilde{x}_1 + \tilde{x}_2) > 0$ for some $c_1 > 0$ and by 2^{nd magnitude then by 1st-order approximation $G_1 \ge c_1(-\tilde{x}_1 + \tilde{x}_2) > 0$ for some $c_1 > 0$, and by 2^{nd} -order approximation, $G_2 \le c_2 \tilde{x}_1 \tilde{x}_2 - c_3 \tilde{x}_2^2 < 0$ with $c_3, c_4 > 0$. Writing a solution to [\(2.2\)](#page-3-0) as $\tilde{x}_2 = h(\tilde{x}_1)$,

$$
h' = \frac{\frac{d}{dt}\tilde{x}_2}{\frac{d}{dt}\tilde{x}_1} = \frac{G_2}{G_1} \ge \frac{c_2\tilde{x}_1\tilde{x}_2 - c_3\tilde{x}_2^2}{c_1(-\tilde{x}_1 + x_2)} = -\frac{c_2}{c_1}\tilde{x}_2\frac{-\tilde{x}_1}{-\tilde{x}_1 + \tilde{x}_2} - \frac{c_3}{c_1}\tilde{x}_2\frac{\tilde{x}_2}{-\tilde{x}_1 + \tilde{x}_2} \ge -\frac{c_2 + c_3}{c_1}\tilde{x}_2,
$$

Mathematical Biosciences and Engineering Volume 16, Issue 4, 3047–3070.

noting that $0 < -\tilde{x}_1, x_2 \le -\tilde{x}_1 + x_2$ on the last step. So, given initial value $\tilde{x}_1(0), \tilde{x}_2(0)$, the corresponding solution has $h(\tilde{x}_1) \ge \tilde{x}_1(0)e^{-c\tilde{x}_1}$ with $c = (c_2 + c_3)/c_1$, so intersects { $\tilde{x}_1 = 0$ } at a positive value of \tilde{x}_2 .
Thus no trajectories tend to n. Thus no trajectories tend to p_1 .

If $a_3 = b_1 \ge 0$ (E), then γ_1, γ_2 intersect at p_3 . If g'_1
and $B(1, -1)$, $B(-1, 1) \rightarrow B(-1, -1)$. Since $g_1 \le$ $y'_1(a_3) \geq g'_2$ $\gamma_2'(a_3)$, then as above, $\gamma_1 \cap \gamma_2 \cap \Lambda_+$ is empty
- 0 then p_2 is the only equilibrium, and if and $R(1, -1)$, $R(-1, 1) \rightarrow R(-1, -1)$. Since $a_1 \le 0$, if $a_3 = 0$ then p_0 is the only equilibrium, and if $a_3 > 0$ then $G_2 > 0$ near p_0 and $p_1 \notin \Lambda_+$ so solutions tend to p_3 . If g'_1
is a single point p_1 and $B(1, -1)$, $B(-1, 1)$, $B(-1, 1)$, $B(-1, -1)$, Λ_3 $g'_1(a_3) < g'_2$
similar ar $\gamma_2'(a_3)$, then $\gamma_1 \cap \gamma_2 \cap \Lambda_+$ is a single point p_4 , and $R(1, -1)$, $R(-1, 1)$ → $R(1, 1)$, $R(-1, -1)$. A similar argument as above shows that p_3 attracts no points in Λ_+ – this time, G_1 has a double root, so write a solution as $\tilde{x}_1 = h(\tilde{x}_2)$ and perform a similar estimate. This situation corresponds to (OB).

If $0 < a_1 < b_2$ (UH) then $\gamma_1 \cap \gamma_2 \cap \Lambda_+$ has cardinality 0, 1 or 2, with a saddle node bifurcation when the cardinality is 1.

- 1. If the cardinality is 0, then $R(1, -1)$, $R(-1, 1) \rightarrow R(-1, -1)$ and solutions tend to p_1 , similar to the case g'_1 $y'_1(a_1) \geq g'_2$ $'_{2}(a_1)$ above.
- 2. If the cardinality is 1, then we have (MB). To see this, note that p_1 is attracting, and there is a unique interior equilibrium p_4 . Three sign regions $R(1, -1)$, $R(-1, -1)$, $R(-1, 1)$ appear, with $R(1, -1), R(-1, 1) \rightarrow R(-1, -1)$ and $R(-1, -1)$ disconnected by p_4 into two components, the lower of which with sink p_1 and the upper with sink p_4 . Since p_1 is locally stable it attracts an open neighbourhood $S \ni p_1$ of points in Λ_+ . Since the basin of attraction of p_1 is $\bigcup_{t \leq 0} \phi(t, S)$ it is open in Λ is open in Λ_{+} .
- 3. If the cardinality is 2, we have (NB). To see this, note that p_1 is again attracting, and there are two interior equilibria p_4 , p_5 , letting p_5 denote the upper one. All sign regions appear, with $R(1, -1), R(-1, 1) \rightarrow R(-1, -1), R(1, 1)$ and $R(-1, -1)$ split in two as before. p_1 is the unique sink for the lower part of $R(-1, -1)$, and p_5 the unique sink for $R(1, 1)$ and the upper part of $R(-1, -1)$. Since γ_1, γ_2 intersect transversally at p_4 and p_5 and are the only nullclines through those points, the Jacobian is non-singular so both equilibria are hyperbolic. Since $p₅$ attracts more than a single curve of points it is attracting, and since p_4 fails to attract nearby points in $R(-1, -1)$ and $R(1, 1)$ it is not attracting. As argued above, both p_1 and p_5 have open basins of attraction in Λ_+ . Since $\Lambda_+ \setminus \{p_4\}$ is open in Λ_+ and connected, it cannot be the combined basins of attraction of p_1 and p_5 since that would imply a disconnection. The only option is that some points other than p_4 tend to *p*4, so *p*⁴ is a saddle and its basin of attraction is its stable manifold, which is a smooth curve.

If $a_1 \leq 0$ and $0 < a_3 < b_1$ (E) then $\gamma_1 \cap \gamma_2 \cap \Lambda_+$ has cardinality 0, 1 or 2 as before. Similar observations show that (E), (MB), (NB) occur, respectively.

Properties of (B). We now show that each of (OB), (MB) and (NB) occurs for some parameter values, that (MB) and (OB) lie on the boundary of (NB), and that (NB) is simply connected. For this it is helpful to view the parameter space in terms of the four independent variables $(\lambda_{10}, \lambda_a, \delta, \lambda_b)$, subject to the general constraints λ_{10} , $\delta \ge 0$, $\lambda_a \ge -\lambda_{10}$ and $\lambda_b \ge -\lambda_a$. Here, of course, we may assume further that δ , λ_a , $\lambda_b > 0$, since (B) has been ruled out in other cases. We'll let \bar{x} and \bar{y} , \bar{x} denote $\gamma_1 \cap \gamma_2 \cap \Lambda$

Figure 3. Transition from (OB) to (NB) to (MB) with $\lambda_{10} = 0.5$, $\lambda_a = 1.5$, $\lambda_b = 5$ and $\delta = 1, 1.1, 1.25$ from left to right. g_1 in blue, g_2 in red.

when its cardinality is, respectively, 1 or 2, with $\bar{y} < \bar{x}$ in the elementwise order.

We recall g_1, g_2 and their derivatives:

$$
g_1(x_1) = \frac{1}{\lambda_a} \left(\frac{1}{1 - x_1} - \lambda_{10} \right), \quad g_2(x_1) = 1 - \frac{\delta}{\lambda_a + \lambda_b x_1},
$$

$$
g'_1(x_1) = \frac{1}{\lambda_a} \frac{1}{(1 - x_1)^2} \quad \text{and} \quad g'_2(x_1) = \frac{\delta \lambda_b}{(\lambda_a + \lambda_b x_1)^2}.
$$

Existence. Our approach is to first characterize (OB), then show a small change in δ leads to (NB), then show a large enough change in δ gives (MB). We treat separately the cases $a_1 \leq 0$ and $a_1 > 0$, beginning with $a_1 \le 0$. To have (OB) we need $g_1(0) = g_2(0)$, i.e., $b_1 = a_3$, and g'_1 $g'_1(0) < g'_2$ $y_2'(0)$. Earlier we showed that $a_3 = b_1 \Leftrightarrow \lambda_{20} = 1 + \delta$, that we write as $\delta = \lambda_a + \lambda_{10} - 1$, which requires $\lambda_a + \lambda_{10} > 1$, i.e., $\lambda_{20} > 1$, since $\delta > 0$ by assumption. If g'_1
25 $\lambda_{\rm s} > \lambda_{\rm s}$ (δ Thus if $\lambda_{\rm s} < 1$ and $\lambda_{\rm s}$ are fixed $g'_1(0) < g'_2$
d (OB) $\lambda_2^2(0)$ then $\lambda_a^{-1} < \delta \lambda_b / \lambda_a^2$, so $\lambda_a < \delta \lambda_b$, that we write as $\lambda_b > \lambda_a/\delta$. Thus if $\lambda_{10} \le 1$ and λ_a are fixed, (OB) occurs iff $\lambda_a + \lambda_{10} > 1$, $\lambda_b \in (\lambda_a/(\lambda_a + \lambda_{10} - 1), \infty)$ and $\delta = \lambda_a + \lambda_{10} - 1$, which is a non-empty set of parameters. Take some such choice of parameters, and note that $g_1(0) - g_2(0) = 0$ and g'_1 $y'_1(0) - g'_2$ $Z_2(0) < 0$. Since g_2 decreases with δ while g_1 is unchanged,
increased enough then $g_2(1) < g_1(0)$ and $\chi_1 \cap \chi_2 \cap \Lambda$ is a small increase in δ then gives (NB). If δ is increased enough, then $g_2(1) < g_1(0)$ and $\gamma_1 \cap \gamma_2 \cap \Lambda_+$ is empty, so there is a unique intermediate value of δ that gives (MB). This is depicted in Figure [3.](#page-18-0)

If $a_1 > 0$, to have (OB) we need $a_1 = b_2 > 0$ and g'_1
t we write as $\delta = \lambda_1 a_1 + \lambda_2$ poting *a*, depends only $y'_1(a_1) < g'_2$
 $y'_2(a_1) \leq g'_3$ $\chi_2'(b_2)$. The former gives $a_1 = (\delta - \lambda_a)/\lambda_b$, that we write as $\delta = \lambda_b a_1 + \lambda_a$ noting a_1 depends only on λ_{10} , while the latter gives $\lambda_{10}^2/\lambda_a < \delta \lambda_b / \delta^2$, or $\delta < \lambda_a$, λ_a^2 . Plugging in δ and putting λ_a terms to one side gives the condition λ_a $\delta < \lambda_a \lambda_b / \lambda_{10}^2$. Plugging in δ and putting λ_b terms to one side gives the condition $\lambda_b (\lambda_a / \lambda_{10}^2 - a_1) > \lambda_a$.
Given λ_b and λ_c and λ_b can be chosen large enough to satisfy this inequality iff Given λ_{10} and λ_a , λ_b can be chosen large enough to satisfy this inequality iff

$$
\lambda_a/\lambda_{10}^2 > a_1 \Leftrightarrow \lambda_a > a_1 \lambda_{10}^2 = \lambda_{10}^2 - \lambda_{10} \Leftrightarrow \lambda_{20} > \lambda_{10}^2.
$$

Thus for $a_1 > 0$, (OB) occurs iff $\lambda_a > \lambda_{10}^2 - \lambda_{10}$, $\lambda_b \in (\lambda_a/(\lambda_a/\lambda_{10}^2 - a_1), \infty)$ and $\delta = \lambda_b a_1 + \lambda_a$. Again, this is a non-empty set of parameters. As before, we can vary δ to obtain (NB) and (MB).

A consequence of the above existence arguments is that they specify for which values of λ_{10} and λ_a bistability can be achieved by appropriate choice of δ , λ_b , namely, for $\lambda_{20} > \max(1, \lambda_{10}^2)$. They also show that for any choice of λ_{10} , λ_a , the set of values of δ that gives (B) is a bounded set show that for any choice of λ_{10} , λ_a , λ_b , the set of values of δ that gives (B) is a bounded set, with (OB)

Figure 4. Typical (δ , λ_b) slice of the phase portrait when $\lambda_{10} \le 1$ (left) and $\lambda_{10} > 1$ (right) and bistability is possible. Left: $\lambda_{10} = 0.5$ and $\lambda_a = 1.5$, and right: $\lambda_{10} = 1.25$ and $\lambda_a = 10$.

at the lower endpoint, (MB) at the upper endpoint and (NB) in the interior. A slice of the phase portrait is shown in Figure [4.](#page-19-0)

Geometry. It's clear that (NB) is an open set in parameter space. Define the functions $\tilde{\lambda}_a = \lambda_{10} + \text{max}(1, \lambda^2)$ and $max(1, \lambda_{10}^2)$ and

$$
\tilde{\lambda}_b = \begin{cases}\n\lambda_a / (\lambda_a + \lambda_{10} - 1) & \text{if } \lambda_{10} \le 1, \\
\lambda_a / (\lambda_a / \lambda_{10}^2 - (1 - 1/\lambda_{10})) & \text{if } \lambda_{10} > 1.\n\end{cases}
$$

Note that both functions are continuous. From the above discussion we obtain functions $\delta^{-}(\lambda_{10}, \lambda_a, \lambda_b)$
and $\delta^{+}(\lambda_{10}, \lambda_a, \lambda_b)$ with domain $\lambda_{10} > 0$, $\lambda_{10} > 0$, and $\lambda_{10} > 0$, with the (OR) (NR) and (MR) o and $\delta^+(\lambda_{10}, \lambda_a, \lambda_b)$ with domain $\lambda_{10} \ge 0$, $\lambda_a > \tilde{\lambda}_a$ and $\lambda_b > \tilde{\lambda}_b$ such that (OB),(NB) and (MB) occur iff $(\lambda_{10}, \lambda_a, \lambda_b)$ are in the domain and $\delta = \delta^-$, $\delta^- < \delta < \delta^+$ and $\delta = \delta^+$, respectively. The function δ^- is given above explicitly piecewise: given above explicitly, piecewise:

$$
\delta^{-}(\lambda_{10}, \lambda_a, \lambda_b) = \begin{cases} \lambda_a + \lambda_{10} - 1 & \text{if } \lambda_{10} \leq 1 \\ \lambda_a + \lambda_b (1 - 1/\lambda_{10}) & \text{if } \lambda_{10} > 1, \end{cases}
$$

and one easily verifies its continuity. The function δ^+ is not given explicitly, but we can show it is
smooth, using the implicit function theorem. (MB) corresponds to a unique $\bar{x}_i \in (0, 1)$ such that smooth, using the implicit function theorem. (MB) corresponds to a unique $\bar{x}_1 \in (0, 1)$ such that $g_1(\bar{x}_1) - g_2(\bar{x}_1) = g'_1$ $y'_{1}(\bar{x}_{1}) - g'_{2}$ $y_2'(\bar{x}_1) = 0$. We have

$$
\partial_{(x_1,\delta)}(g_1-g_2,g'_1-g'_2) = \begin{pmatrix} g'_1 - g'_2 & \partial_{\delta}(g_1 - g_2) \\ g''_1 - g''_2 & \partial_{\delta}(g'_1 - g'_2) \end{pmatrix}
$$

where the ' indicates ∂_{x_1} . At (MB), $g_1(\bar{x}_1)' - g_2(\bar{x}_1)' = 0$, moreover $g_1'' - g_2'' > 0$ since $g_1 - g_2$ is
convex. Also, $\partial_x(g_1 - g_2) = -1/(3 + \lambda x_1)^2 > 0$ since $\lambda > 0$ by assumption. Thus the above matrix convex. Also, $\partial_{\delta}(g_1 - g_2) = -1/(\lambda_a + \lambda_b x_1)^2 > 0$ since $\lambda_a > 0$ by assumption. Thus the above matrix is invertible. Since $(g_1 - g_2, g'_1)$ is a smooth function of parameters, and of x, when $\lambda > 0$ is invertible. Since $(g_1 - g_2, g'_1 - g'_2)$
and $g_1 \in (0, 1)$ by the implicit funct 2) is a smooth function of parameters, and of x_1 , when $\lambda_a > 0$
tion theorem (\bar{x}_1 , δ^{+}) is a smooth function of $(\lambda_1, \lambda_2, \lambda_3)$ on the and $x_1 \in (0, 1)$, by the implicit function theorem, (\bar{x}_1, δ^+) is a smooth function of $(\lambda_{10}, \lambda_a, \lambda_b)$, on the domain of δ^+ domain of δ^+ .

We now show (NB) is a contractible set, via a sequence of homotopies that gives a deformation retraction. Define $\tilde{\delta} = (\delta^- + \delta^+)/2$, which is continuous, and note that (NB) occurs when $\delta = \tilde{\delta}$. Notice

that $\tilde{\lambda}_a$, $\tilde{\lambda}_b$ and $\tilde{\delta}$ are functions of λ_{10} , λ_{10} and λ_a , and λ_{10} , λ_a and λ_b respectively. Define the sequence of homotopies as follows, at each step letting $\tilde{\lambda}$. and $\tilde{\delta}$ t of homotopies as follows, at each step letting $\tilde{\lambda}_a$, $\tilde{\lambda}_b$ and $\tilde{\delta}$ take the previous values in the list as their input:

- (i) $H_1 = (\lambda_{10}, \lambda_a, \lambda_b, (1-t)\delta + t\delta),$
- (ii) $H_2 = (\lambda_{10}, \lambda_a, (1-t)\lambda_b + t(\tilde{\lambda}_b + 1), \tilde{\delta}),$

(iii) $H_1 = (\lambda_{10}, (1-t)\lambda_b + t(\tilde{\lambda}_b + 1), \tilde{\lambda}_b + 1)$
- (iii) $H_3 = (\lambda_{10}, (1 t)\lambda_a + t(\tilde{\lambda}_a + 1), \tilde{\lambda}_b + 1, \tilde{\delta}),$
(iv) $H_4 = ((1 t)\lambda_{10} + t\tilde{\lambda}_a + 1, \tilde{\lambda}_b + 1, \tilde{\delta})$
- (iv) $H_4 = ((1 t)\lambda_{10} + t, \tilde{\lambda}_a + 1, \tilde{\lambda}_b + 1, \tilde{\delta}).$

Note that $t \in [0, 1]$ is the deformation parameter. In words,

- (i) H_1 acts only on δ , centering its value to δ ,
- (ii) H_2 acts on λ_b and δ , centering λ_b to $\tilde{\lambda}_b + 1$ while keeping $\delta = \tilde{\delta}$,
iii) H_1 centers λ_a to $\tilde{\lambda}_b + 1$ while keeping $\lambda_b = \tilde{\lambda}_b + 1$ and $\delta = \tilde{\delta}$ and
- (iii) H_3 centers λ_a to $\tilde{\lambda}_a + 1$ while keeping $\lambda_b = \tilde{\lambda}_b + 1$ and $\delta = \tilde{\delta}$ and (iv) H_4 centers λ_a to 1 while keeping other parameters centered
- (iv) H_4 centers λ_{10} to 1 while keeping other parameters centered.

We have already checked the continuity of component functions, and it's clear they keep points in (NB). Applying the *H_i* in sequence deforms (NB) to the single point with $\lambda_{10} = 1$ and the corresponding values of \tilde{J} , \tilde{J} , and $\tilde{\delta}$ values of $\tilde{\lambda}_a, \tilde{\lambda}_b$ and $\tilde{\delta}$.

6. Discussion and conclusion

In this article we consider a simple differential equation model of host-symbiont dynamics, that includes both vertical and horizontal transmission of the host, as well as recovery (i.e. transition from associated [with symbiont] to unassociated [without symbiont] host). The model naturally incorporates the pathogen, neutral, and mutualist cases through the choice of parameter values, providing a unified setting for the study of all three cases.

Our analysis indicates that the most natural way to view the process is through the total host density (both associated and unassociated) and the proportion of hosts that are associated. Moreover, the two main factors governing the dynamics are (i) the effect of the symbiont on the host (i.e. whether it is pathogen, neutral or mutualist) and (ii) the difference between the additional birth rate of associated vs. unassociated hosts (which is positive only for mutualists) and the rate of horizontal transmission of the symbiont.

We obtain a complete picture of the model's dynamic phases, as follows. Except for the very special case when the symbiont is completely redundant (neutral, no recovery and no horizontal transmission), populations that initially contain both host and symbiont, and not located at an unstable equilibrium, always settle down to one of at most three equilibria. The limiting equilibrium is unique when the symbiont is either a pathogen, is neutral, or is a mutualist but focuses more on increasing the host birth rate than on horizontal transmission. Bistability (exactly two locally attracting equilibria) can occur when the symbiont is a sufficiently beneficial mutualist that focuses more on horizontal transmission than on increasing the host birth rate. There are three limiting equilibria only in a boundary case at the onset of bistability. When the limiting equilibrium is unique, survival and coexistence are determined by straightforward invasibility conditions, i.e., whether one type can invade the extinction equilibrium (for survival) or the stable equilibrium containing only the other type (for coexistence).

Interestingly, the dynamical behaviour, and in particular the number of stable equilibria, hinges less on whether the symbiont is pathogen or mutualist, and more on whether the symbiont is highly beneficial (increases host birth rate), versus highly infectious (has a high rate of horizontal transmission). Moreover, we have shown that, at least when including only the simple mechanisms considered in this model, stable periodic behaviour is not possible.

Conflict of interest

All authors declare that there is no conflicts of interest in this paper.

References

- 1. N. Moran, The ubiquitous and varied role of infection in the lives of animals and plants, *Amer. Nat.* 160 (2002), S1–S8.
- 2. T. Fenchel, *Ecology of Protozoa.* Springer-Verlag, 1987.
- 3. C. C. Khor and M. L. Hibberd, Host-pathogen interactions revealed by human genome-wide surveys, *Trends Genet.* 28 (2012), 233–243.
- 4. D. Harvell, Ecology and evolution of host-pathogen interactions in nature, *Amer. Nat.* 164 (2004), S1–S5.
- 5. M. Li, B. Wang, M. Zhang, et al., Symbiotic gut microbes modulate human metabolic phenotypes, *PNAS* 105 (2008) 2117–2122.
- 6. R. M. Anderson and R. M. May, Infectious Diseases of Humans: Dynamics and Control. Oxford University Press, 1991.
- 7. C. L. Wolin and L. R. Lawlor, Models of facultative mutualism: density effects, *Am. Nat.*, 124 (1984), 843–862.
- 8. D. H. Wright, A simple, stable model of mutualism incoporating handling time, *Am. Nat.*, 134 (1989), 664–667.
- 9. M. Lipsitch, M. A. Nowak, D. Ebert, et al., The population dynamics of vertically and horizontally transmitted parasites, *P. Roy. Soc. B-Biol. Sci.*, 260 (1995).
- 10. E. Foxall and N. Lanchier, Generalized stacked contact process with variable host fitness, *arXiv:1511.01184* (2016).
- 11. S. J. Court, R. A. Blythe and R. J. Allen, Parasites on parasites: coupled fluctuations in stacked contact processes, *EPL* 101 (2013).
- 12. T. G. Kurtz, Strong approximation theorems for density dependent Markov chains, *Stoch. Proc. Appl.* 6 (1978), 223–240.
- 13. M. I. Freidlin and A. D. Wentzell, Random Perturbations of Dynamical Systems Third Edition, Springer-Verlag, 2012.
- 14. G. O. Poinar, Description of an early Cretaceous termite (Isoptera: Kalotermitidae) and its associated intestinal protozoa, with comments on their co-evolution, *Parasite. Vector.*, 2 (2009).
- 15. C. L. Wolin, The population dynamics of mutualistic systems. In: D. H. Boucher (ed.) *The Biology of Mutualism*, Oxford University Press, New York, (1985), 248–269.
- 16. H. Amann, *Ordinary Di*ff*erential Equations: an Introduction to Nonlinear Analysis.* de Gruyter Studies in Mathematics, 13, 1990.
- 17. L. Perko, *Di*ff*erential Equations and Dynamical Systems.* Third edition. Texts in Applied Mathematics, 7. Springer-Verlag, New York, 2001. xiv+553 pp.

Appendix

Proof of Lemma 5.1. For a definite sign vector $\sigma \in \{-1, 1\}^2$ and $\epsilon > 0$ let

$$
R(\sigma, \epsilon) = \{x \in \Lambda_+ : \text{sgn}(G(x)) = \sigma \text{ and } ||G(x)|| > \epsilon\},\
$$

noting that $R(\sigma) = \bigcup_{\epsilon > 0} R(\sigma, \epsilon)$ and $\sigma \cdot G(x) \ge \epsilon$ for $x \in R(\sigma, \epsilon)$. If $x \in R(\sigma, \epsilon)$ for definite σ and letting

$$
\tau(x,\epsilon) = \inf\{t : \phi(t,x) \notin R(\sigma,\epsilon)\},\
$$

 $\phi(t, x) \in R(\sigma, \epsilon)$ for $t < \tau(x, \epsilon)$, so using the second fact above,

$$
\partial_t \sigma \cdot \phi(t, x) = \sigma \cdot G(\phi(t, x)) \ge \epsilon.
$$

Since for $y, z \in \Lambda_+$, $|\sigma \cdot y - \sigma \cdot z| \leq 2$, it follows that $\tau(x) \leq 2/\epsilon$. Thus, letting $E(\sigma, \epsilon) = \text{Cl}(R(\sigma) \setminus R(\sigma, \epsilon))$ and

$$
E(\sigma) = \bigcap_{\epsilon > 0} E(\sigma, \epsilon)
$$

where Cl denotes the closure, $E(\sigma)$ consists of equilibrium points in Cl($R(\sigma)$). Letting $\tau(x) = \inf\{t :$ $\phi(t, x) \notin R(\sigma)$, the only way that $\tau(x) = \infty$ is if for any $\epsilon > 0$, eventually $\phi(t, x) \in E(\sigma, \epsilon)$, which means that $\omega(x) \in E(\sigma)$. Since $\omega(x)$ is connected and (for the cases that remain) the set of equilibrium points in Λ is finite, the first claim follows. The second claim follows since acyclic implies itineraries are finite. \Box

Proof of Lemma [5.2.](#page-13-2) Suppose Γ is a periodic orbit or a separatrix cycle, both of which are the image of a circle in \mathbb{R}^2 , under a piecewise C^1 function. Since trajectories do not cross, the image is injective, so let *D* be the interior of Γ, whose existence is given by the Jordan curve theorem. By Green's theorem,

$$
\iint_D \nabla \cdot (BF)dxdy = \oint -BF_2dx + BF_1dy.
$$

where $F = (F_1, F_2)$. First suppose Γ is a periodic orbit with solution curve $u(t)$ satisfying $u(T) = u(0)$. Ignoring the orientation, since it only changes the sign, the latter integral can be parametrized with $dx = \dot{u}_1(s)ds$, $dy = \dot{u}_2(s)ds$ as

$$
\int_0^T -B(u(s))(F_2(u(s))\dot{u}_1(s) + F_1(u(s))\dot{u}_2(s))ds.
$$

However, $F_2(u(s)) = \dot{u}_2(s)$ and $F_1(u(s)) = \dot{u}_1(s)$ so the integrand is identically zero, thus so is the integral, and this contradicts the assumption on $\nabla \cdot (BF)$. Suppose instead that Γ is a separatrix cycle and let $u^{(1)}(t), \ldots, u^{(m)}(t)$ denote the corresponding heteroclinic solutions. Since solutions are arranged

tip-to-tail as described by [\(3.1\)](#page-8-2), it follows that $\{u^{(i)}(t) : t \in \mathbb{R}, i = 1, ..., m\}$ orients the tangent space for Γ so up to its sign, we can parametrize the above contour integral as Γ, so up to its sign, we can parametrize the above contour integral as

$$
\sum_{i=1}^{m} \int_{-\infty}^{\infty} -B(u(s))(F_2(u(s))\dot{u}_1(s) + F_1(u(s))\dot{u}_2(s))ds
$$

and the same observation shows the integral is zero, which is a contradiction.

 \Box

 c 2019 the Author(s), licensee AIMS Press. This is an open access article distributed under the terms of the Creative Commons Attribution License (http://[creativecommons.org](http://creativecommons.org/licenses/by/4.0)/licenses/by/4.0)