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Using Numerical Bifurcation Analysis to Study Pattern Formation in Mussel Beds

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Abstract. Soft-bottomed mussel beds provide an important example of ecosystem-scale self-organisation. Field data from some intertidal regions shows banded patterns of mussels, running parallel to the shore. This paper demonstrates the use of numerical bifurcation methods to investigate in detail the predictions made by mathematical models concerning these patterns. The paper focusses on the “sediment accumulation model” proposed by Liu et al (Proc. R. Soc. Lond. B 14 (2012), 20120157). The author calculates the parameter region in which patterns exist, and the sub-region in which these patterns are stable as solutions of the original model. He then shows how his results can be used to explain numerical observations of history-dependent wavelength selection as parameters are varied slowly.

Key words: mussels, pattern formation, periodic travelling wave, reaction-diffusion-advection, numerical continuation, wavetrain

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1 Introduction

Pattern formation at the scale of whole ecosystems is an active sub-discipline within spatial ecology. There are now many examples of such landscape-scale patterns, and mathematical modelling

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is playing a key role in understanding them because the opportunities for empirical study are typically very limited. Examples include banded vegetation on semi-arid hillslopes [7, 37, 43], “fairy circles” in Namibia [56], patterns of ridges and hollows in peatlands [13, 14], “ribbon forests” in the Rocky mountains [1], localised patches of trees in savannah grasslands [20, 46], intertidal mudflats [54], and banded patterns in soft-bottomed mussel beds [16, 23, 49]. This paper concerns the last of these examples. It has long been known that mussels tend to self-organise into patches on rocky shores [21, 27, 55]. However, larger scale striped patterns have been reported more recently for mussel beds in both the Dutch Wadden Sea [49] and the Menai Strait (UK) [16]. These are soft-bottomed systems meaning that the mussels adhere only to one another or to debris such as shell fragments. They are also intertidal, and the mussel bands are orthogonal to the tidal flow, with a wavelength of several metres.

Two different mechanisms have been proposed for banded patterns in soft-bottomed mussel beds. One is the “reduced losses” hypothesis, which was due originally to van de Koppel et al [49]. Mussel beds are subject to disruption by predation, wave action, and ice scouring [12]. These pressures are resisted by the attachment of mussels to one another, via byssal threads. More of these attachments form when the mussels are subject to perturbations [52]. Therefore van de Koppel et al [49] hypothesised that the per capita loss rate of mussels will be reduced at higher mussel densities. This is consistent with empirical data showing that mussel density increases in response to both greater wave exposure [48] and greater predation threat [5, 26]. A mathematical model based on the reduced losses hypothesis was originally proposed by van de Koppel et al [49] and has been studied and modified in a number of subsequent papers [3, 17, 22, 35, 53].

The focus of this paper is the alternative “sediment accumulation” hypothesis proposed by Liu et al [22]. Being filter feeders mussels produce sediment (pseudo-faeces), and in the context of a mussel bed this can significantly affect the topography – thus mussels are “ecosystem engineers” [15, 51]. For a substrate with hummocks and troughs, hydrodynamics implies that the water velocity will be greater at a hummock than at a trough, and Liu et al [22] hypothesised that this will increase the supply rate of algae, which is the main food source for mussels in these ecosystems. This is consistent with empirical data, also in [22], revealing a positive correlation between elevation and mussel density. Since larger mussel densities imply more rapid sediment deposition, this hypothesis sets up a positive feedback loop that has the potential to generate patterns.

2 The sediment accumulation model

Liu et al [22] proposed a mathematical model based on their “sediment accumulation” hypothesis, formulated in terms of algal density $a(x, t)$, mussel density $m(x, t)$ and the amount of accumulated sediment $s(x, t)$:

$$\partial a / \partial t = \overbrace{1 - \alpha a}^{\text{transfer to/ from upper water layers}} - \overbrace{\beta a m (s + \eta) / (s + 1)}^{\text{depth-dependent consumption by mussels}} + \overbrace{\nu \partial a / \partial x}^{\text{advection by tide}} \quad (2.1)$$

$$\partial m / \partial t = \overbrace{\delta a m (s + \eta) / (s + 1)}^{\text{mussel growth}} - \overbrace{m}^{\text{death}} + \overbrace{\partial^2 m / \partial x^2}^{\text{random movement}} \quad (2.2)$$

$$\partial s / \partial t = \overbrace{m}^{\text{production}} - \overbrace{\theta s}^{\text{erosion}} + \overbrace{D \partial^2 s / \partial x^2}^{\text{dispersal}}. \quad (2.3)$$

This is a dimensionless version of the model: details of the dimensional version and the nondimensionalisation are given in [22]. The spatial coordinate x increases with distance from the shore; t denotes time, and $\alpha, \beta, \delta, \theta, \eta < 1, \nu$ and D are positive parameters. The only nonlinear terms in the model are the mussel growth rate $\delta a m (s + \eta) / (s + 1)$ and the corresponding rate of algal consumption: per capita growth is an increasing function of both algal density and the amount of sediment. This nonlinearity provides a short-range activation of mussel density, and the advection of algae from the open sea generates long-range inhibition, because the mussels compete spatially for food. Together these processes can lead to pattern formation, and a typical pattern solution is illustrated in Figure 1.

An important prelude to the study of patterns in (2.1–2.3) is to calculate the homogeneous steady states. For all parameter values there is a mussel-free steady state $(a, m, s) = (1/\alpha, 0, 0)$. Steady states with a non-zero mussel density require a sufficiently high level of algal supply, specifically

$$\delta \geq \delta_{\text{crit}} \equiv \left[\alpha - \beta \theta \eta + (4 \alpha \beta \theta (1 - \eta))^{1/2} \right].$$

Then there are two other steady states, $(a, m, s) = (a_{\pm}, m_{\pm}, s_{\pm})$ where

$$a_{\pm} = (s_{\pm} + 1) / [\delta (s_{\pm} + \eta)] \quad (2.4)$$

$$m_{\pm} = \theta s_{\pm} \quad (2.5)$$

$$s_{\pm} = \frac{1}{2\beta\theta} \left[\delta - \alpha - \beta\theta\eta \pm \{ (\delta - \alpha - \beta\theta\eta)^2 - 4\beta\theta(\alpha - \delta\eta) \}^{1/2} \right]. \quad (2.6)$$

General conditions on the stability of these steady states are very complicated, but for parameter values given by the estimates of Liu et al [22] $(1/\alpha, 0, 0)$ and (a_+, m_+, s_+) are stable to spatially homogeneous perturbations, while (a_-, m_-, s_-) is unstable [22].

Pattern solutions of (2.1–2.3) move away from the shore, as shown in Figure 1. Intuitively this is because the model predicts higher algal densities on the off-shore side of a mussel band compared to the on-shore side, due to consumption in the band, and this causes a net growth of mussels on the off-shore side and a net loss on the on-shore side. This results in a gradual net off-shore migration. Mathematically the movement is a consequence of the advection term, which means that pattern onset occurs via a Turing-Hopf rather than a Turing bifurcation [24, 28, 31]. In fact, there is no evidence of large scale migration of banded patterns in real mussel

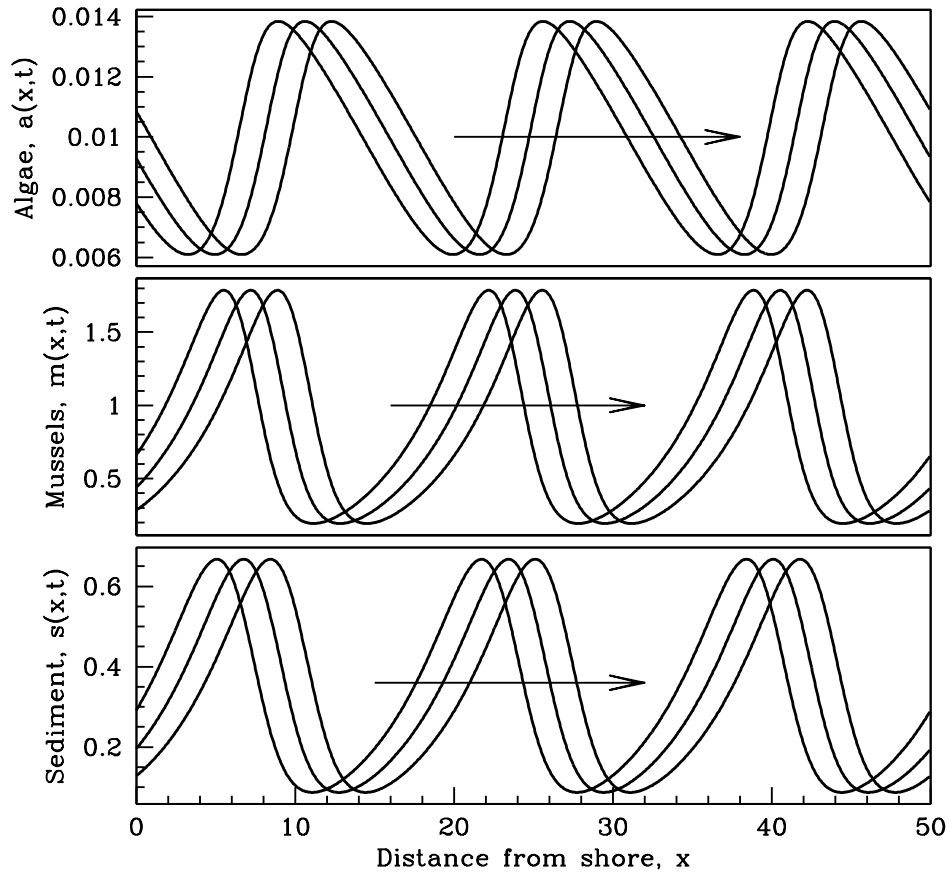


Figure 1: An illustration of a patterned solution of the model (1). The solution is plotted at $t = 500, 502$ and 504 ; the arrows show the direction of pattern movement, which is in the positive x direction, i.e. away from the shore. The equations were solved using a semi-implicit finite difference scheme on a domain of length 50 with periodic boundary conditions. The initial condition was a small perturbation to the uniform steady state (a_+, m_+, s_+) , with the perturbation consisting of a sum of small-amplitude sine functions of period $50/N$, $N = 1, \dots, 10$. The parameters were $\alpha = 50$, $\beta = 200$, $\delta = 320$, $\eta = 0.1$, $\nu = 360$, $\theta = 2.5$, $D = 1$.

beds. A natural explanation for this is the oscillatory nature of tidal flow, which would imply bidirectional oscillating advection of algae rather than the unidirectional term in (2.1). This has been investigated recently for an alternative model of banded pattern formation [39], showing that bidirectional advection can generate patterns that do not show large scale migration. However for mathematical simplicity we restrict attention in this paper to the unidirectional advection case.

Since pattern solutions of (2.1–2.3) move with constant shape and speed, they are periodic travelling waves: $a(x, t) = \tilde{a}(z)$, $m(x, t) = \tilde{m}(z)$ and $s(x, t) = \tilde{s}(z)$ where $z = x - ct$. Here $c > 0$ is the migration speed. Solutions of this form satisfy the ordinary differential equations

$$(\nu + c) \partial \tilde{a} / \partial z + 1 - \alpha \tilde{a} - \beta \tilde{a} \tilde{m} (\tilde{s} + \eta) / (\tilde{s} + 1) = 0 \quad (2.7)$$

$$\partial^2 \tilde{m} / \partial z^2 + c \partial \tilde{m} / \partial z + \delta \tilde{a} \tilde{m} (\tilde{s} + \eta) / (\tilde{s} + 1) - \tilde{m} = 0 \quad (2.8)$$

$$D \partial^2 \tilde{s} / \partial z^2 + c \partial \tilde{s} / \partial z + \tilde{m} - \theta \tilde{s} = 0 \quad (2.9)$$

with patterns corresponding to limit cycle solutions. The key ecological control parameter is δ , which reflects the algal concentration in the upper water layers, and I consider changes in δ with the other ecological parameters fixed; I take the fixed values from Liu et al (2012). However while varying δ one must also consider changes in the wave speed c , which is an additional parameter introduced by the travelling wave ansatz. Therefore my initial aim is to determine the part of the δ - c plane in which there are patterns (i.e. periodic travelling waves, or equivalently limit cycle solutions of (2.7–2.9)).

3 The parameter regions giving patterns

A useful first step in calculating the parameter region giving patterns is to construct bifurcation diagrams for the equations (2.7–2.9) as δ is varied with c fixed. The starting point for the bifurcation diagrams is the steady state $(\tilde{a}, \tilde{m}, \tilde{s}) = (a_+, m_+, s_+)$ since this corresponds to a steady state of (2.1–2.3) that is stable to spatially homogeneous perturbations: one expects patterns to develop when it becomes unstable to spatially inhomogeneous perturbations. An example bifurcation diagram is shown in Figure 2. At large values of δ there are no patterns, but as δ is decreased there is a Hopf bifurcation in (2.7–2.9) from which a branch of limit cycle (pattern) solutions emanates. This solution branch terminates at a homoclinic point, meaning that the period of the limit cycle tends to infinity; thus as δ is decreased towards this point the mussel bands become more and more widely separated. For values of the speed c close to that used in Figure 2 it is the loci of the Hopf bifurcation and homoclinic point that enclose the region of the δ - c parameter plane giving patterns.

The simple form of the bifurcation diagram in Figure 2 does not apply for all values of c : two complications can arise. First, the branch of pattern solutions may have a fold, rather than proceeding monotonically from the Hopf bifurcation to the homoclinic point (e.g. Figure 3). In some parts of the δ - c parameter plane the locus of this fold forms one boundary of the parameter region giving patterns. Note that in general, for any system of ODEs, the part of a parameter plane in which limit cycles exist is bounded by segments of three types of loci: Hopf bifurcations, homoclinic points, and folds in a limit cycle solution branch.

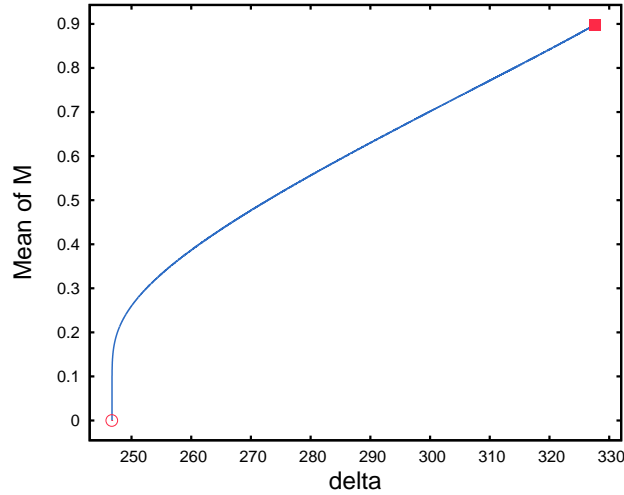


Figure 2: Bifurcation diagram for (2.7–2.9) showing the mean mussel density as a function of δ with the migration speed $c = 0.6$. The other parameters are $\alpha = 50$, $\beta = 200$, $\eta = 0.1$, $\nu = 360$, $\theta = 2.5$, $D = 1$. The square indicates a Hopf bifurcation point, and the open circle indicates a homoclinic point. All calculations and plotting were done using the software package WAVETRAIN (www.ma.hw.ac.uk/wavetrain) [34, 36]. Full details of the WAVETRAIN input files, run commands and plot commands are given at www.ma.hw.ac.uk/~jas/supplements/mmp-bif/.

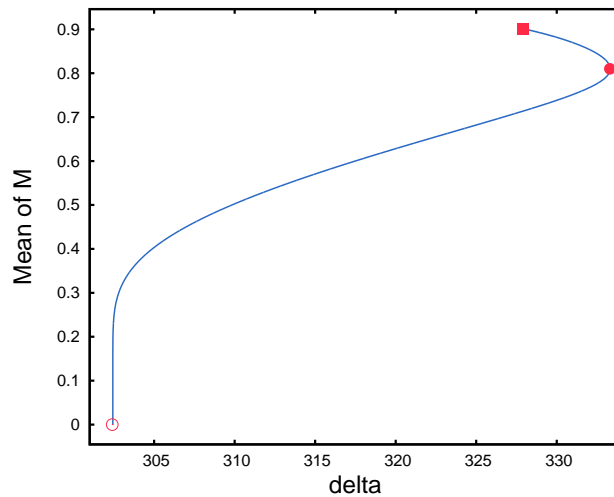


Figure 3: Bifurcation diagram for (2.7–2.9) showing the mean mussel density as a function of δ with the migration speed $c = 1.2$. The other parameters are as in Figure 2. The square indicates a Hopf bifurcation point, the open circle indicates a homoclinic point, and the closed circle indicates a fold in the solution branch. All calculations and plotting were done using the software package WAVETRAIN (www.ma.hw.ac.uk/wavetrain) [34, 36]. Full details of the WAVETRAIN input files, run commands and plot commands are given at www.ma.hw.ac.uk/~jas/supplements/mmp-bif/.

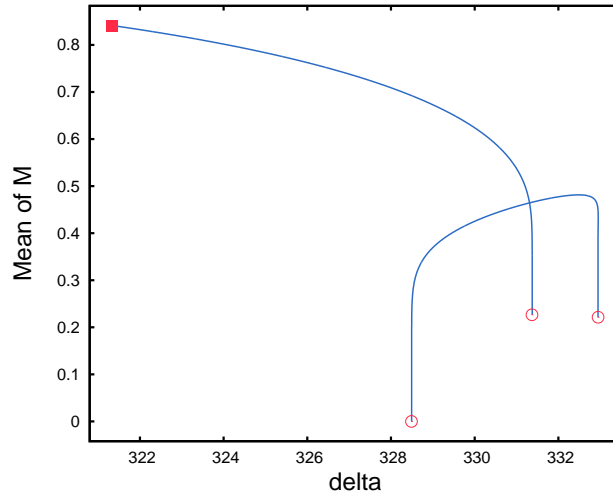


Figure 4: Bifurcation diagram for (2.7–2.9) showing the mean mussel density as a function of δ with the migration speed $c = 1.375$. The other parameters are as in Figure 2. The square indicates a Hopf bifurcation point and the open circles indicate homoclinic points. All calculations and plotting were done using the software package WAVETRAIN (www.ma.hw.ac.uk/wavetrain) [34, 36]. Full details of the WAVETRAIN input files, run commands and plot commands are given at www.ma.hw.ac.uk/~jas/supplements/mmp-bif/.

The other potential complication is that there may be a second, disconnected branch of patterns (e.g. Figure 4). This solution branch does not arise from a Hopf bifurcation point in (2.7–2.9); rather, it connects two different homoclinic points. In some parts of the parameter plane it is loci of these homoclinic points that form the boundary of the parameter region giving patterns.

I calculated these various loci using WAVETRAIN [34] (www.ma.hw.ac.uk/wavetrain); the result is shown in Figure 5. WAVETRAIN is a software package specifically designed for the study of periodic travelling wave solutions of partial differential equations. It determines the loci by numerical continuation, for which it uses the software package AUTO [8–10]. WAVETRAIN is also able to determine whether the patterns are stable as solutions of the original model (2.1–2.3), via numerical continuation of the essential spectrum: this methodology was originally developed by Rademacher and coworkers [30]. Boundaries in the parameter plane between stable and unstable waves can also be traced, by numerical continuation of special points in the spectrum [30, 36]. The subdivision between stable and unstable patterns is also shown in Figure 5; note that the region of parameter space giving stable patterns is sometimes known as the Busse balloon [2]. There are stable patterns for $244.0 < \delta < 334.7$, and for any δ within this range there is a stable pattern for a range of values of the migration speed c . There are also unstable patterns for a range of migration speed values, and in some cases these ranges overlap, implying the coexistence of stable and unstable patterns with the same migration speed. Note that the stability of periodic travelling wave solutions of PDEs is determined entirely by the continuous part of the spectrum (the essential spectrum): there is no point spectrum [32, §3.4.2]. Changes in stability can be either of Eckhaus (sideband) or Hopf type [29, 36]. In the former case there is a change in the sign of the curvature

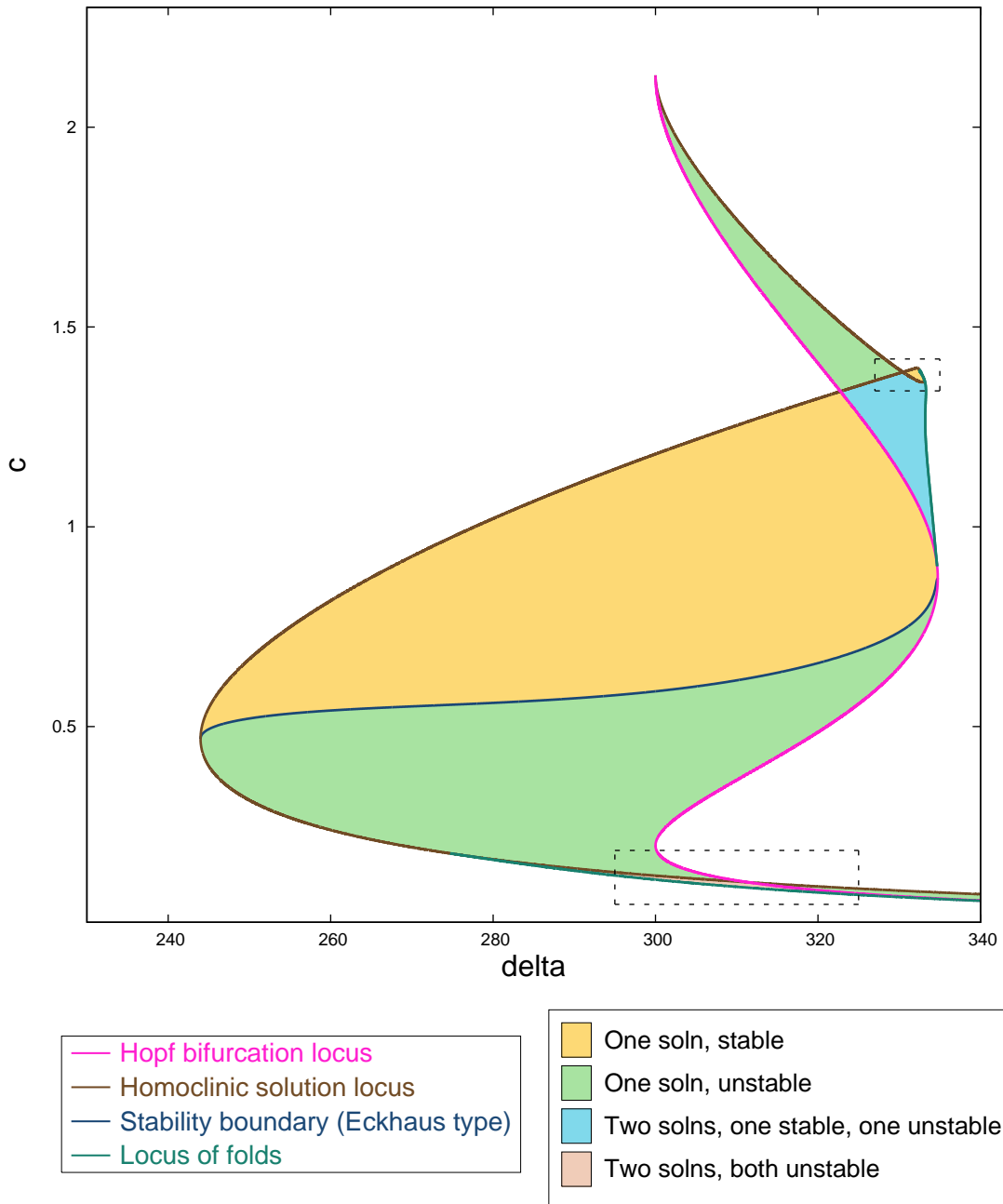


Figure 5: Existence and stability of pattern solutions of the model (2.1–2.3), as a function of the algal supply parameter δ and the migration speed c . The other parameters are $\alpha = 50$, $\beta = 200$, $\eta = 0.1$, $\nu = 360$, $\theta = 2.5$, $D = 1$ (as in Figures 2–4). The key shows the interpretation of the various line and shading types; there are no pattern solutions in the unshaded regions of the parameter plane. The thin dashed lines indicate regions that are shown at higher magnification in Figure 7. The region giving two unstable solutions is hard to see in this figure: it is clearer in Figure 7b. All calculations and plotting were done using the software package WAVETRAIN (www.ma.hw.ac.uk/wavetrain) [34, 36]. Full details of the WAVETRAIN input files, run commands and plot commands are given at www.ma.hw.ac.uk/~jas/supplements/mnnp-bif/.

of the spectrum at the origin, while in the latter a fold in the spectrum crosses the imaginary axis at a point away from the origin; note that the spectrum always passes through the origin because the patterns are neutrally stable to translations. Investigation of the spectrum, again using WAVETRAIN, shows that in this case the stability change is of Eckhaus type (Figure 6).

The plot in Figure 5 is particularly complicated in the vicinity of the small “loop” in the homoclinic locus, which is shown at greater magnification in Figure 7a. To provide a clearer account of the behaviour in this part of the parameter plane, Figure 8 shows a series of bifurcation diagrams, which plot the period of patterns as a function of δ for various fixed values of the migration speed c . This shows that the solution branch inside the “loop” connects two different homoclinic points, as in Figure 4, and is separate from that emanating from the Hopf bifurcation point. All patterns on this separate solution branch are stable as solutions of the model equations (2.1–2.3). Figure 9 shows bifurcation diagrams in which the period of patterns is plotted as a function of c for $\delta = 328$ and 332. For these values of δ there are Hopf bifurcations at three different values of c . The speed c increases monotonically along the solution branch emanating from the largest of these values, until a homoclinic point is reached. Patterns on this branch are unstable as PDE solutions. For the middle Hopf bifurcation value, the solution branch also proceeds in the direction of increasing c , and again terminates at a homoclinic solution. Patterns on this branch are initially unstable, with a switch to stability part way along the branch. On the solution branch emanating from the smallest of the three Hopf bifurcation points (not shown in Figure 9), c initially decreases before a fold is reached, after which c increases up to a homoclinic end point. This is illustrated in Figure 7b by the magnified plot of this part of the δ – c parameter plane. Note that the fold in the Hopf bifurcation locus at $c \approx 0.2$ and $\delta = 300$ corresponds to a transition from Hopf bifurcations of (a_+, m_+, s_+) to those of (a_-, m_-, s_-) ; for the parameter values used in the figure, $\delta_{\text{crit}} = 300$. Since (a_-, m_-, s_-) is unstable in (2.1–2.3) to even spatially uniform perturbations, one expects intuitively that this instability will be inherited by the bifurcating patterns, and indeed all patterns in the part of the parameter plane with $c < 0.2$ are unstable as PDE solutions.

4 History-dependent changes in pattern wavelength

A good illustration of the value of detailed results on the existence and stability of patterns, such as is provided by Figure 5, is given by studying the changes in pattern form as a parameter is varied slowly. Since we used the algal supply level δ as the main control parameter in §3, I consider slow changes in this parameter. A typical result is illustrated in Figure 10a. I selected a starting value of δ and used WAVETRAIN to calculate a pattern with a particular wavelength; for Figure 10 I started at $\delta = 310$ with a pattern of wavelength 15. I then decreased δ in small steps, solving for a fixed (and relatively long) time at each δ value. Note that I did not alter the solution in any way when changing δ : the numerical results show in Figure 10a are of a single very long numerical simulation. As δ is decreased the patterns initially retain the same wavelength – though of course the form of the pattern changes to reflect the new parameter value. One can see this in Figure 10a because the dots denoting the numerical model solutions lie on the contour of constant pattern period. However at $\delta \approx 290$ the period contour leaves the parameter region in which patterns

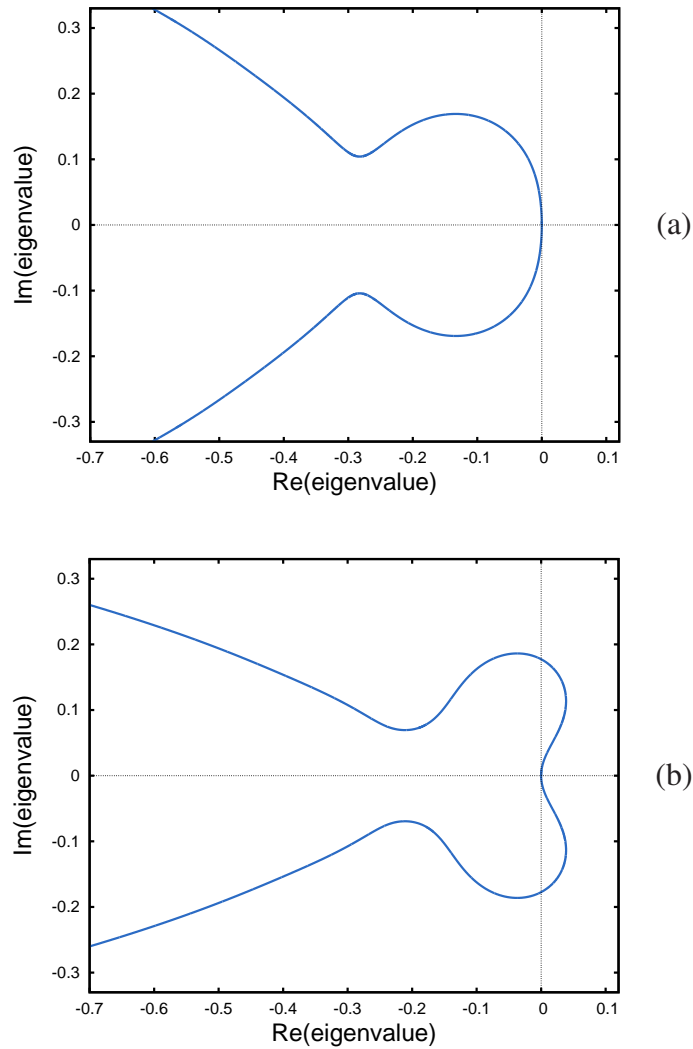


Figure 6: The essential spectra of two pattern solutions of (2.1–2.3), for $\delta = 280$ and (a) $c = 0.6$, (b) $c = 0.4$ with other parameter values as in Figure 5. These parameter values lie either side of the stability boundary (see Figure 5) and comparison of (a) and (b) shows that the stability change is of Eckhaus type, meaning that there is a change in the sign of the curvature of the spectrum at the origin. All calculations and plotting were done using the software package WAVETRAIN (www.ma.hw.ac.uk/wavetrain) [34, 36]. Full details of the WAVETRAIN input files, run commands and plot commands are given at www.ma.hw.ac.uk/~jas/supplements/mnnp-bif/.

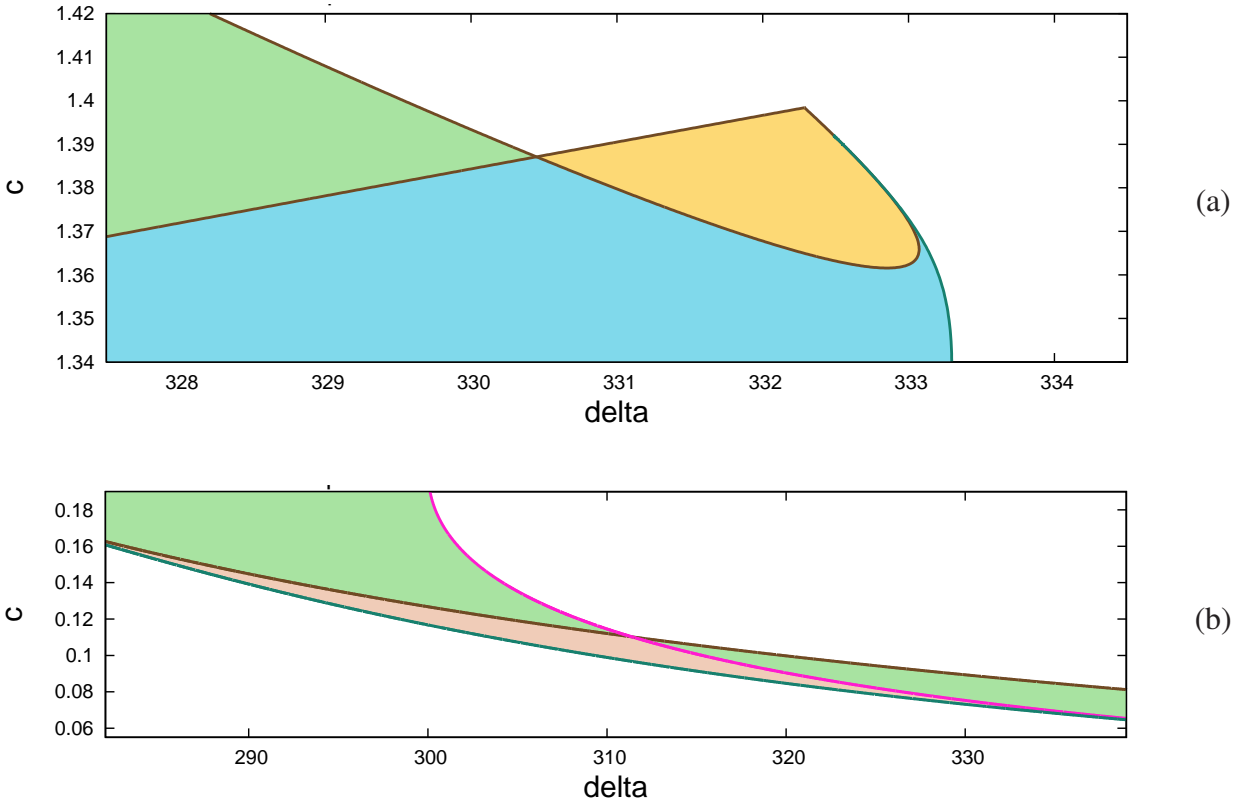


Figure 7: Details of two parts of the δ - c parameter plane, showing the existence and stability of pattern solutions of the model (2.1–2.3). The two regions are indicated by thin dashed lines in Figure 5, and the values of the other parameters and the interpretation of the various line and shading types are as in that figure. All calculations and plotting were done using the software package WAVETRAIN (www.ma.hw.ac.uk/wavetrain) [34, 36]. Full details of the WAVETRAIN input files, run commands and plot commands are given at www.ma.hw.ac.uk/~jas/supplements/mnnp-bif/.

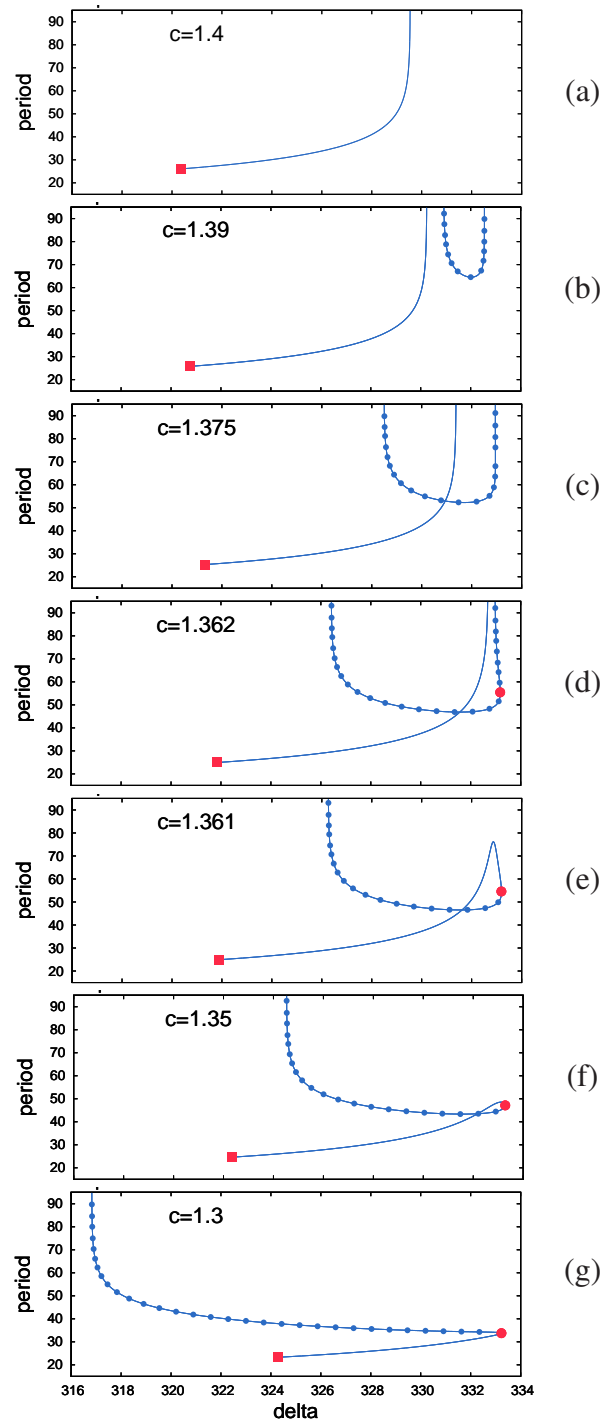


Figure 8: Caption on next page.

Figure 8: (Figure on previous page). Bifurcation diagrams for (2.7–2.9) showing the period of pattern solutions as a function of the algal supply parameter δ . The different panels correspond to different fixed values of the migration speed c ; the other parameters are as in Figure 5. The squares indicate Hopf bifurcation points and the circles indicate folds in the solution branch. Lines with dots indicate patterns that are stable as solutions of (2.1–2.3); lines without dots indicate unstable patterns. These plots should be viewed in parallel with Figure 7a. All calculations and plots were done using the software package WAVETRAIN (www.ma.hw.ac.uk/wavetrain) [34, 36]. Full details of the WAVETRAIN input files, run commands and plot commands are given at www.ma.hw.ac.uk/~jas/supplements/mmp-bif/.

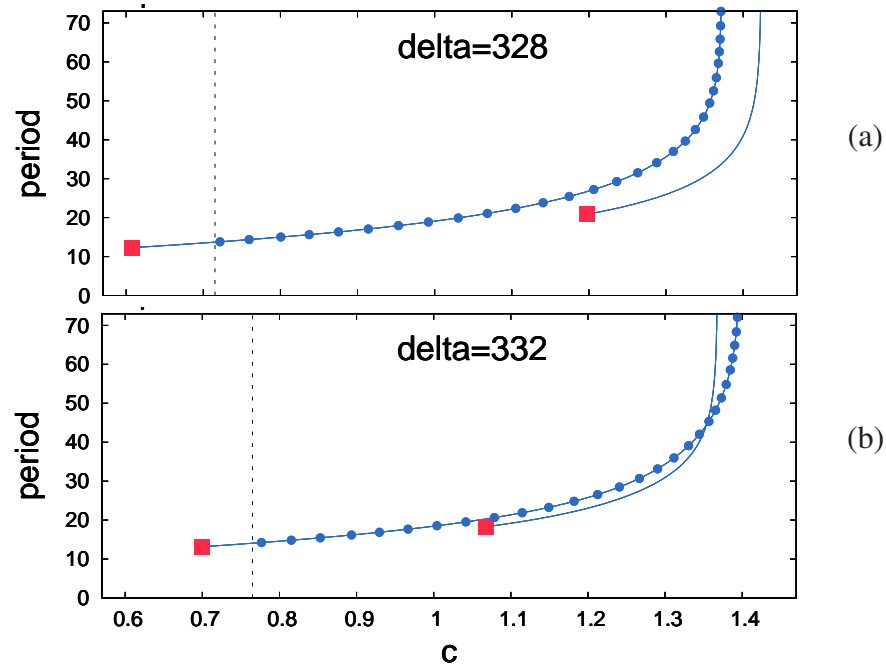


Figure 9: Bifurcation diagrams for (2.7–2.9) showing the period of pattern solutions as a function of the migration speed c . The two panels correspond to different fixed values of the algal supply parameter δ ; the other parameters are as in Figure 5. The squares indicate Hopf bifurcation points. Lines with dots indicate patterns that are stable as solutions of (2.1–2.3); lines without dots indicate unstable patterns. The dashed lines indicate the values of c at which the stability boundary crosses the two values of δ . These plots should be viewed in parallel with Figures 5 and 7a. All calculations and plotting were done using the software package WAVETRAIN (www.ma.hw.ac.uk/wavetrain) [34, 36]. Full details of the WAVETRAIN input files, run commands and plot commands are given at www.ma.hw.ac.uk/~jas/supplements/mmp-bif/.

are stable as solutions of (2.1–2.3), and the observed period changes to 20. When δ reaches 274 (chosen arbitrarily) I reversed the direction of change and increased δ , in the same small steps. The pattern remains of period 20: thus there is hysteresis (history dependence) in pattern wavelength.

In Figure 10a the time spent at each δ value is very long (2000 dimensionless units). Figure 10b shows the corresponding result when this time is halved. In this case the transition to the new wavelength does not occur until there have been two subsequent decreases in δ . Halving the time again (to 500 dimensionless units) results in δ reaching the minimum value of 274, starting to increase again, and then re-entering the region in which the pattern of wavelength 15 is stable, all before the transition to a longer wavelength would have occurred: hence the pattern remains of wavelength 15 and there is no hysteresis. Note that the results shown in Figure 10 rely heavily on the accuracy of my numerical solutions. Therefore in the Appendix I present the results of convergence tests of my numerical method, which is a simple finite difference scheme. On the basis of these tests, I fixed the spatial grid spacing at 0.025, with a time step of 5.56×10^{-5} ; these imply a CFL number of 0.8, and give an error of about 0.03% in the solution (see Table A.1).

History-dependence of pattern wavelength, such as that illustrated in Figure 10a,b, is not a new result for patterned ecosystems. Similar behaviour has been demonstrated both for models of semi-arid vegetation [35, 43] and for models of soft-bottomed mussel beds based on the “reduced losses” hypothesis [35]. Its occurrence in this additional case of Liu et al’s model of the “sediment accumulation” hypothesis for soft-bottomed mussel beds gives an increased sense that such hysteresis may be a widespread feature within patterned ecosystems. The practical applications of this are significant: parameter estimation alone is not enough to predict pattern wavelength, however carefully it is done. In addition, one requires detailed data on past values of the wavelength.

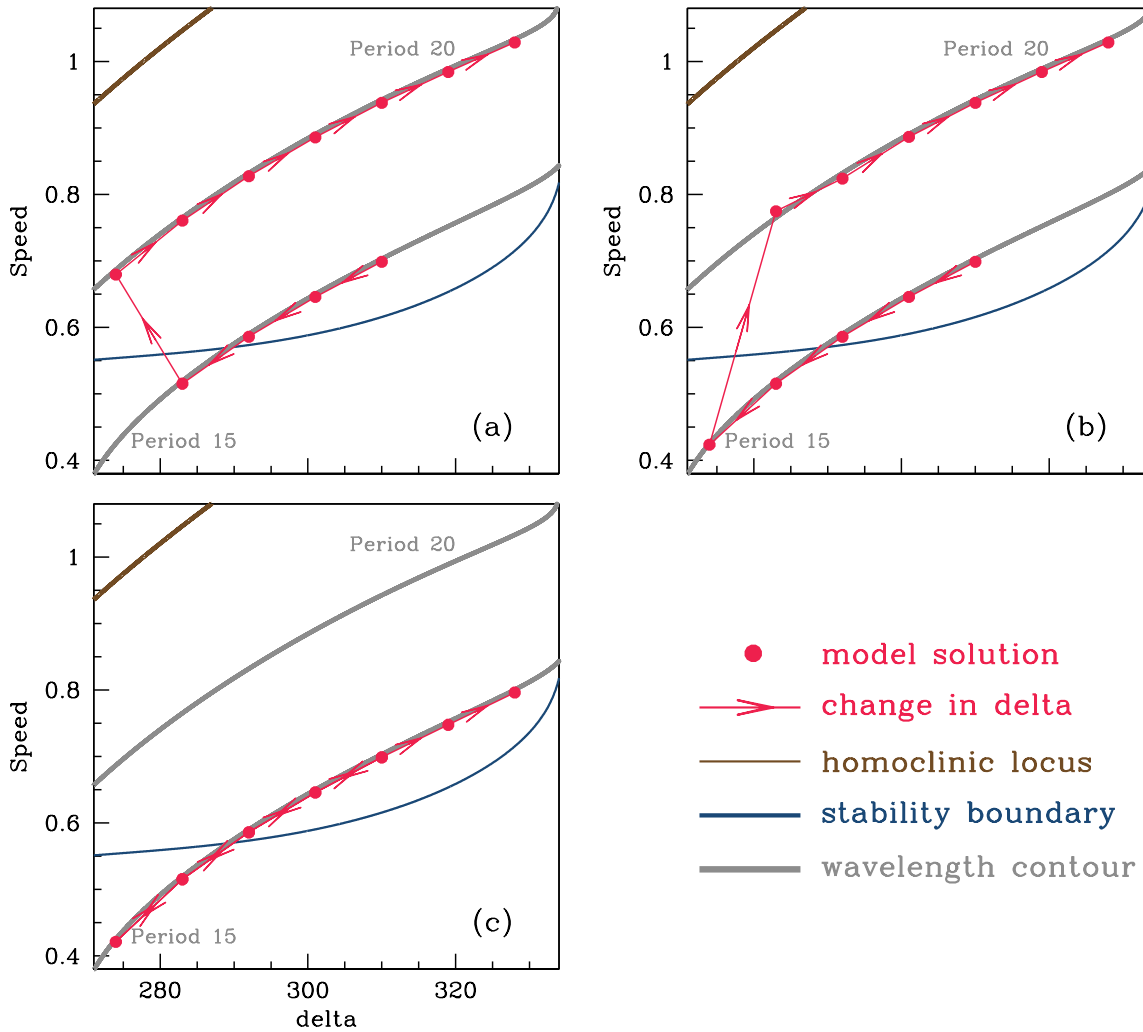


Figure 10: Changes in patterns in the model (2.1–2.3) for sediment-driven mussel bed patterns as the algal supply parameter δ is varied slowly, with parameter values as in Figure 5, on a domain of length 300. I decreased δ from 310 to 274 in steps of 9, and then increased it again. The length of (dimensionless) time spent at each δ value was (a) 2000; (b) 1000; (c) 500. There is no resetting of initial conditions when δ is changed, and the results in the each part of the figure are for a single long simulation. The boundary conditions are periodic, and the initial condition is the pattern of wavelength 15 for $\delta = 310$, which I calculated using WAVETRAIN [34]. The dots show the wavelength and speed immediately before each change in δ ; the arrows indicate the direction in which δ is changing. In (a) the solution follows the wavelength 15 contour until this leaves the parameter region giving stable patterns, when there is a transition to the new wavelength of 20. In (b) the behaviour is similar although the change in δ is sufficiently rapid that the transition to the new wavelength does not occur until there have been two subsequent decreases in δ . In (c), δ is changed sufficiently fast that it has re-entered the region in which the pattern of wavelength 15 is stable before the transition to a longer wavelength would have occurred: hence the pattern remains of wavelength 15 and there is no hysteresis. Details of the numerical method are given in the Appendix. Readers considering reproducing this figure should note that the simulation in parts (a), (b) and (c) took about 18, 9 and 4.5 days respectively on a Linux PC with a 2.83GHz Intel Core 2 Quad Q9500 processor.

5 Discussion

Numerical bifurcation analysis is a powerful tool for understanding patterned solutions of partial differential equations. In this paper I have used it to study Liu et al's [22] model (2.1–2.3) of the “sediment accumulation” hypothesis for pattern formation in soft-bottomed mussel beds. Adopting the implementation of numerical bifurcation analysis in the software package WAVETRAIN [34,36], I have been able to unravel the complicated dependence on parameter values of pattern existence and stability; this includes multiple coexisting patterns and disconnected branches of pattern solutions. The most important aspect of the results is the calculation of the region of parameter space in which there are stable patterns, sometimes known as the Busse balloon [2]. As an illustration of the value of this calculation, I showed that it enables one to understand the jumps in pattern wavelength and its history-dependence as a parameter value is varied slowly.

Although my results on pattern existence and stability are detailed, they are not exhaustive, and (at least) two important features remain to be explored. The first is stability in two space dimensions. My designation of patterns as stable or unstable refers to their stability in the one-dimensional PDE system (2.1–2.3). In reality mussel beds occupy a two-dimensional domain, and although the patterns themselves are one-dimensional (striped patterns) they could be unstable to perturbations along the stripes (transverse perturbations) even when they are stable as solutions of (2.1–2.3), i.e. stable to perturbations orthogonal to the stripes. Calculation of stability to transverse perturbations is more difficult than for one-dimensional perturbations: it is outside the scope of the software package WAVETRAIN, and to my knowledge there is only one published paper that calculates this type of stability, namely [42]. In that paper, Siero and coworkers used numerical bifurcation analysis to calculate the spectrum of striped patterns to transverse perturbations for the Klausmeier-Gray-Scott model, which has been well studied in the context of applications to chemistry [4, 11, 18] and semi-arid vegetation [19, 38, 50]. They showed that the parameter region giving patterns that are stable in one space dimension subdivides, with some of these patterns being stable to transverse perturbations while others are unstable. The break-up of unstable patterns tends to generate two-dimensional patterns with a rhombic geometry. Application of Siero et al's [42] approach is a natural direction for future work on the model (2.1–2.3) since it would add a new level of detail that would be necessary for the proper interpretation of two-dimensional numerical simulation results.

A second aspect of pattern existence and stability that I have not explored is absolute stability. In any spatiotemporal system, unstable solutions may be either absolutely unstable, or convectively unstable but absolutely stable. In the latter case, perturbations applied to the solution will grow but at the same time they will move away from the original site of the perturbation: typically the growing perturbation eventually reaches a boundary and is then lost. By contrast, “absolute instability” means that perturbations grow at their original location. (See [33, 41] for full discussions). Any solution that is not absolutely unstable can be an important long-term component of a solution of a PDE model, even if it is (convectively) unstable [6, 25, 40, 44]. Therefore it would be extremely valuable to subdivide the parameter region giving unstable patterns according to whether or not the patterns are absolutely unstable. However current numerical methods for calculating absolute stability are either too laborious to be practical in the context of a parameter plane (e.g. [47]) or

only apply to spatially uniform solutions [30,45]. The development of new methods for calculating absolute stability that would enable subdivision of the parameter region giving unstable patterns for a model such as (2.1–2.3) is a natural but very challenging area for future research.

Appendix: Error estimation for numerical simulations

The calculations in §4 depend heavily on numerical simulations of (2.1–2.3), and therefore a detailed estimation of numerical errors is essential. My numerical scheme is of a simple finite difference type, with diffusion terms evaluated semi-implicitly, an explicit upwinding representation of advection, and explicit evaluation of the reaction terms. I used a uniform grid spacing Δx and time step Δt . Experiments showed that convergence requires the Courant number \mathcal{C} to satisfy the CFL condition $1 > \mathcal{C} = \nu \Delta t / \Delta x$, as is common in reaction-diffusion-advection systems. To investigate numerical accuracy when this condition is satisfied, I used a test problem. This was formulated on a domain of length 100 with periodic boundary conditions, and parameter values $\alpha = 50$, $\beta = 200$, $\delta = 315$, $\eta = 0.1$, $\nu = 360$, $\theta = 2.5$, $D = 1$. The initial conditions were as in Figure 1: a small perturbation (the same for each run) to the homogeneous steady state (2.4–2.6). I solved for $0 \leq t \leq 300$. As a measure of error, I used the spatial average of the L_2 norm of the relative error of the final solution, compared to a reference solution calculated with high accuracy. Typically the difference in numerical discretisation between the solution being considered and the reference solution results in a spatial translation of the pattern. It is not appropriate to regard this “phase shift” as numerical error: as mentioned in the main text, the patterns are neutrally stable to translations. Therefore I performed an appropriate spatial translation before comparing the solutions.

Table A.1 shows the way in which the error varies with the space and time steps. As is typical for reaction-diffusion-advection problems, the error decreases linearly with Δx and Δt when these are varied in a constant ratio, so that the Courant number is constant (< 1). Moreover, for given Δx the error is essentially independent of the Courant number provided this is < 1 (compare the three columns of Table A.1). For fixed Courant number (< 1) the error is approximately proportional to Δx .

For the long-term simulations used for Figure 10, the choice of numerical parameters is of course a trade-off between accuracy and CPU time, and I used a CFL number of 0.8, with $\Delta x = 0.025$. In my test problem, this gives errors of about 0.03%.

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	CFL= 0.90	CFL= 0.60	CFL= 0.30
$\delta x = 0.40000$	0.00633592	0.00635213	0.00636858
$\delta x = 0.20000$	0.00294851	0.00295517	0.00296197
$\delta x = 0.10000$	0.00152884	0.00153179	0.00153494
$\delta x = 0.05000$	0.00073414	0.00073551	0.00073683
$\delta x = 0.02500$	0.00031846	0.00031916	0.00031988
$\delta x = 0.01250$	0.00010679	0.00010713	0.00010748
$\delta x = 0.00625$	0.00000064	0.00000032	0.00000000

Table A.1: Convergence of my numerical scheme for the model (2.1–2.3). I tabulate the spatial average of the L_2 norm of the relative error of the solution, compared to the most accurate solution, as a function of the grid spacing Δx and the CFL number. Details of the test problem are given in the text.

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