

## Entropy Production of Cyclic Population Dynamics

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Entropy serves as a central observable in equilibrium thermodynamics. However, many biological and ecological systems operate far from thermal equilibrium. Here we show that entropy production can characterize the behavior of such nonequilibrium systems. To this end we calculate the entropy production for a population model that displays nonequilibrium behavior resulting from cyclic competition. At a critical point the dynamics exhibits a transition from large, limit-cycle-like oscillations to small, erratic oscillations. We show that the entropy production peaks very close to the critical point and tends to zero upon deviating from it. We further provide analytical methods for computing the entropy production which agree excellently with numerical simulations.

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The study of complex systems with a large number of interacting particles requires global observables that characterize their behavior. Modern statistical mechanics has successfully identified, interpreted, and applied such observables for equilibrium systems. One of these observables is the entropy which allows for predictions of a system's behavior through the second law of thermodynamics—an isolated system's entropy cannot decrease. Identifying similar principles for nonequilibrium systems, however, proves elusive. Neither a characteristic global observable nor a universal principle has been identified in a general way. While also in nonequilibrium the entropy production has been proposed as a useful observable [1,2], and different principles governing its behavior have been suggested [3,4], problems arise from different employed definitions of entropy and approaches to nonequilibrium dynamics [2,5,6].

In this Letter we demonstrate that entropy production can successfully characterize ecological systems with cyclic competition. Ecological systems display a wide variety of nonlinear and nonequilibrium behavior. Random interactions between individuals and the finiteness of the population lead to intrinsic stochasticity. Nonequilibrium results when interactions between individuals of different species include cyclic dependencies where a species  $A_1$  benefits from and suppresses a species  $A_2$ , while  $A_2$  benefits from and suppresses a species  $A_3$ , and so on, with some species  $A_k$  of the resulting chain benefiting from and suppressing species  $A_1$ . Such cycles can lead to erratic or limit-cycle oscillations in the steady state of the population dynamics [7–16]. Experimental observations of cyclic dynamics and corresponding oscillations have, amongst others, been documented for mating behavior of lizards in costal California [17] and in microbial laboratory communities [18].

The dynamics of ecological systems can be conveniently described as a Markovian stochastic process through a master equation,

$$\partial_t P_i(t) = \sum_j [\omega_i^j P_j(t) - \omega_j^i P_i(t)], \quad (1)$$

in which  $P_i(t)$  denotes the probability of finding the system in a certain state  $i$  at time  $t$  and  $\omega_i^j$  is the transition probability from state  $j$  to state  $i$ . The associated mean entropy production  $\dot{S}$  of the system follows as

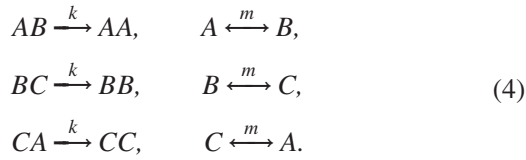
$$\dot{S} = \frac{1}{2} \sum_{i,j} [\omega_i^j P_j(t) - \omega_j^i P_i(t)] \ln \left[ \frac{\omega_i^j P_j(t)}{\omega_j^i P_i(t)} \right]. \quad (2)$$

Equation (2) can be obtained through considering the difference between forward and backward entropy per unit time of the stochastic process, Eq. (1) [19]. Equation (2) follows also as the temporal derivative of the system's Gibbs entropy together with a term describing the total increase of thermodynamic entropy in the reservoirs to which the system is coupled [20]. For steady states defined by  $\partial_t P_i(t) = 0$ , as we consider in this Letter, the entropy production simplifies to

$$\dot{S} = \frac{1}{2} \sum_{i,j} [\omega_i^j P_j - \omega_j^i P_i] \ln \left[ \frac{\omega_i^j}{\omega_j^i} \right]. \quad (3)$$

It follows from Eq. (2) that the entropy production vanishes if and only if the system obeys detailed balance,  $\omega_i^j P_j = \omega_j^i P_i$ . Indeed, detailed balance represents the notion of thermodynamic equilibrium in the framework of the master equation. Cyclic population dynamics violates detailed balance; the computation and discussion of the associated entropy production is the scope of this Letter.

Consider a simple model for cyclic population dynamics of three species  $A$ ,  $B$ , and  $C$ . Interactions are formulated as chemical reactions:



The reactions on the left describe cyclic competition:  $A$  outperforms  $B$  but is beaten by  $C$ , and  $C$  is taken over by  $B$  in turn. The reactions on the right correspond to spontaneous mutations between the three species.

The population model defined by the reactions (4) exhibits a critical mutation rate that, in the resulting nonequilibrium steady state, delineates large oscillations in the species densities from only small ones. Let us introduce this critical mutation rate first. The reactions (4) conserve the total number  $N$  of interacting individuals. The densities  $a$ ,  $b$ , and  $c$  of species  $A$ ,  $B$ ,  $C$  therefore sum up to one,  $a + b + c = 1$ , and the population's state space is the simplex  $S_3$  (see Fig. 1). Numerical simulations indicate that small values of the mutation rate  $m$  lead to large oscillations between the densities of the three species; the probability distribution is highest close to the corners of the simplex [Fig. 1(a)]. Large values of  $m$ , on the contrary, lead to an approximately Gaussian probability distribution around the simplex center [Fig. 1(b)]. Erratic oscillations occur at small amplitudes [9].

The system's behavior can be analytically described by an approximate Fokker-Planck equation. A systematic expansion in the system size  $N$  yields an equation for the temporal evolution of the probability distribution  $P(s, t)$  of the densities  $s = (a, b)$  at time  $t$ :

$$\partial_t P(s, t) = -\partial_i [\alpha_i(s) P(s, t)] + \frac{1}{2} \partial_i \partial_j [\beta_{ij}(s) P(s, t)], \quad (5)$$

in which the indices  $i, j$  run from 1 to 2; the summation

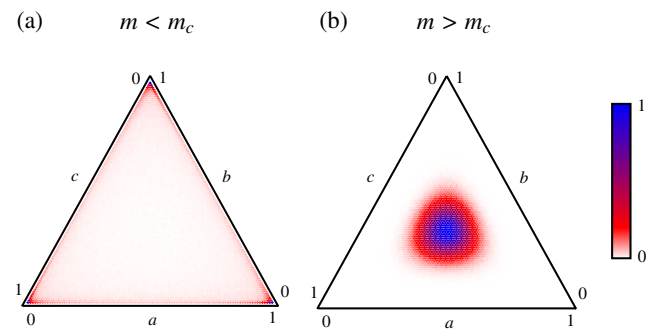


FIG. 1 (color online). Probability distributions ( $k = 1$ ,  $N = 100$ ). (a) For a mutation rate  $m = 0.003$  smaller than  $m_c = k/(2N)$  the probability distribution is concentrated near the edges and particularly near the corners of the phase space. (b) A mutation rate  $m = 0.1$  larger than  $m_c$  leads to a Gaussian distribution around the center.

convention implies summation over them. The density  $c$  follows as  $c = 1 - a - b$ . The coefficients read

$$\begin{aligned} \alpha_i(s) &= [m(1 - 3s_i) + ks_i(s_{i+1} - s_{i+2})], \\ \beta_{ii}(s) &= N^{-1}[m(1 + s_i) + ks_i(s_{i+1} + s_{i+2})], \\ \beta_{ij}(s) &= -N^{-1}[m(s_i + s_j) + ks_i s_j] \quad \text{for } i \neq j, \end{aligned} \quad (6)$$

where the indices are understood as modulus 3 and  $s_3 = c$ . The terms containing  $\alpha$  describe the deterministic part of the temporal evolution. In the absence of fluctuations, the reactions for cyclic dominance lead to neutrally stable oscillations around the internal fixed point  $s_* = (1/3, 1/3, 1/3)$ , while the spontaneous mutations render the internal fixed point stable. Demographic fluctuations are, for large system sizes  $N$ , inversely proportional to  $\sqrt{N}$  and enter the Fokker-Planck equation (5) through the terms containing  $\beta$ . They induce a stochastic drift away from the internal fixed point towards the boundaries of the phase space. The Fokker-Planck equation (5) shows that the competition between the deterministic and the stochastic effects leads, at a critical mutation rate  $m_c = k/(2N)$ , to a uniform probability distribution. Certain deviations from the uniform distribution occur near the phase space boundaries where the discreteness of the phase space becomes relevant and the continuous formulation through the Fokker-Planck equation does not hold. For a small mutation rate,  $m < m_c$ , fluctuations dominate and drive the system towards the boundary. In the absence of mutations the corner states are absorbing and the system goes extinct [21]. An arbitrary small mutation rate, however, leads to sustained species coexistence and oscillations. In the opposite case, when  $m > m_c$ , the deterministic dynamics centers the probability distribution around the internal fixed point.

The cyclic population dynamics yields a nonequilibrium steady state that is characterized by oscillations, large or small, around the internal fixed point. What is the resulting entropy production and how does it relate to the regimes of small, critical, and large mutation rates outlined above?

To tackle this question we have carried out extensive numerical simulations of the stochastic system employing the Gillespie algorithm [22]. Throughout our simulations we have considered  $k = 1$  which defines the time scale. Numerical results from computer simulations of the stochastic system show that the entropy production peaks at a certain value  $m_{\max}$  of the mutation rate (Fig. 2). The value  $m_{\max}$  approximately equals the critical mutation rate,  $m_{\max} \approx m_c$  (Fig. 2 inset). Small deviations from this behavior arise because the probability distribution at the critical mutation rate is not uniform near the boundaries as mentioned above.

Analytical understanding of the entropy production in the regimes of small, critical, and large mutation rates is feasible through the Fokker-Planck equation (5). To this end we employ a continuous version of the entropy pro-

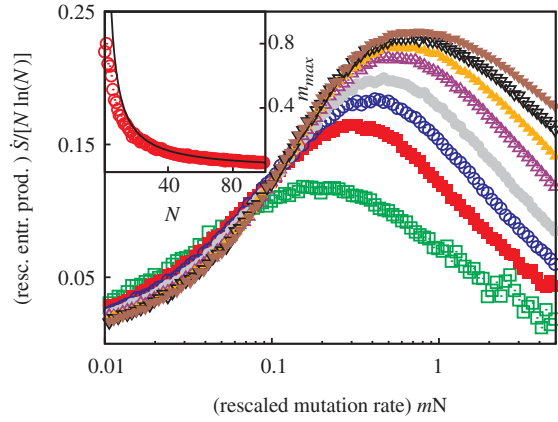


FIG. 2 (color online). Entropy production in the steady state for different system sizes ( $\square$ ,  $N = 2$ ;  $\blacksquare$ ,  $N = 5$ ;  $\circ$ ,  $N = 10$ ;  $\bullet$ ,  $N = 20$ ;  $\triangle$ ,  $N = 50$ ;  $\blacktriangle$ ,  $N = 100$ ;  $\nabla$ ,  $N = 200$ ;  $\blacktriangledown$ ,  $N = 400$ ). The entropy production vanishes for very high and very low mutation rates and exhibits a maximum at an intermediate value  $m_{\max}$ . The value  $m_{\max}$  is near the critical mutation rate as shown in the inset where the black line indicates  $m_c + 0.001$ , and red circles represent data obtained from simulations.

duction (3),

$$\dot{S} = \frac{1}{2} \int dr \int ds [\omega_s^r P(\mathbf{r}) - \omega_r^s P(\mathbf{s})] \ln \left( \frac{\omega_s^r}{\omega_r^s} \right), \quad (7)$$

where integration is over all states  $\mathbf{r}$ ,  $\mathbf{s}$  of the phase space.

The entropy production in the form of Eq. (7) can readily be evaluated at the critical mutation rate  $m_c$ . The probability distribution is uniform according to the Fokker-Planck equation (5); we obtain

$$\dot{S}_{m=m_c} = \frac{3}{144} kN [12 \ln(N) - 13 + 6 \ln(4)]. \quad (8)$$

For moderate and large  $N$  the term  $N \ln(N)$  on the right-hand side dominates the entropy production. Stochastic simulations confirm this behavior (Fig. 2).

In the regime of large mutation rates,  $m > m_c$ , we need to calculate the probability density in the steady state to compute the entropy production. We obtain the probability density by using polar coordinates  $(r, \phi)$  centered at the internal fixed point. We then simplify the Fokker-Planck equation (5) through a van Kampen approximation for the coefficients (6) in which the coefficients are approximated by their value at the internal fixed point. The resulting Fokker-Planck equation is then solved by the Gaussian distribution

$$P(r, \varphi) = \frac{1}{2\pi\sigma^2} \exp\left(\frac{-r^2}{2\sigma^2}\right), \quad (9)$$

where  $\sigma = \sqrt{(k+6m)/36mN}$ . The entropy production follows from Eq. (7) where the integral is evaluated by setting the upper boundary of the integral to  $\infty$  and an average over the angular dependence is taken:

$$\dot{S}_{m \gg m_c} = \frac{kN}{3} \ln\left(\frac{k}{3m} + 1\right). \quad (10)$$

This result agrees excellently with numerical simulations [Fig. 3(b)]. The entropy production (10) depends linearly on the system size  $N$ . This behavior arises because the typical area in phase space explored by the dynamics is proportional to  $\sigma^2 \sim 1/N$  and thus contains  $N^2 \sigma^2 \sim N$  states. The continuity approximation employed in the Fokker-Planck equation (5) holds for arbitrary large  $m$ , since the width  $\sigma$  of the probability distribution (9) remains finite as  $m \rightarrow \infty$ .

Expanding (10) for large values of  $m$  results in  $\dot{S}_{m \gg m_c} = k^2 N / (9m)$ . The entropy production vanishes as  $m/k$  increases. Indeed, only the cyclic dynamics at rate  $k$  underlies the nonequilibrium behavior and therefore entropy production, while the mutations at rate  $m$  obey detailed balance.

When the mutation rate is small,  $m \ll m_c$ , the probability distribution is concentrated near the boundaries of the phase space [Fig. 1(a)]. The dynamics occurs predominantly along the boundary and can therefore be approximately described by considering only the boundary states. Because of the threefold symmetry it suffices to regard only one edge of the simplex with periodic boundary conditions. The concentration  $x$  of one of the three species increases along this edge from 0 to 1 such that the cyclic dynamics drives the system to  $x = 1$ . The deterministic part of the dynamics is given by

$$\partial_t x = m(1-2x) - k(x-x^2) \quad (11)$$

and features a fixed point at  $x_* = [(2m+k) - \sqrt{4m^2+k^2}]/(2k)$ . In the range of  $m \ll m_c$ , this fixed point

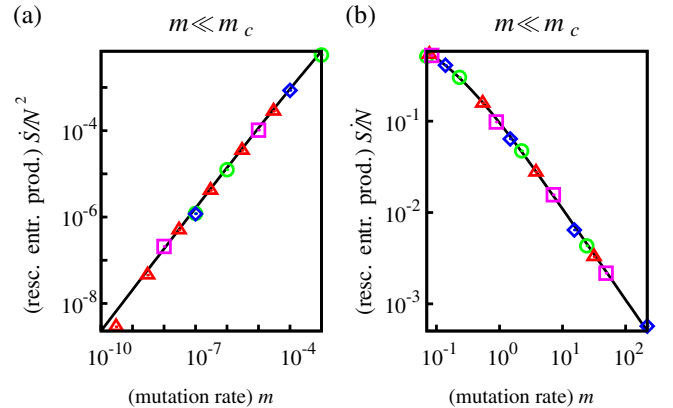


FIG. 3 (color online). Entropy production in the limiting cases  $m \ll m_c$  (a) and  $m \gg m_c$  (b). Analytical results (black lines) agree excellently with simulations ( $\circ$ ,  $N = 30$ ;  $\triangle$ ,  $N = 100$ ;  $\square$ ,  $N = 200$ ;  $\diamond$ ,  $N = 400$ ). The data confirm that the entropy production is proportional to the squared system size  $N^2$  for small mutation rates and proportional to  $N$  for large mutation rates. The simulation results further confirm that the entropy production decays as  $m$  for  $m \rightarrow 0$  and as  $1/m$  for  $m \rightarrow \infty$ .

is closer to 1 than the distance  $1/N$  between two discrete states. We conclude that fluctuations will cause the system to exhibit a constant circular current in the steady state. The probability distribution  $P(x)$  for  $x \in [\frac{1}{N}, 1 - \frac{1}{N}]$  can therefore be obtained as a solution to the Fokker-Planck equation  $0 = -\partial_x\{[m(1-2x) - k(x-x^2)]P(x)\}$  where fluctuations have been ignored:

$$P(x) = \mathcal{N} \frac{1}{m - 2mx - kx + kx^2}, \quad (12)$$

with a normalization coefficient  $\mathcal{N}$ . To determine the probability  $P_0$  of a corner state, which turns out to be finite, fluctuations have to be included.  $P_0$  can be obtained using the master equation and the values of  $P(x = 1/N)$  and  $P(x = 1 - 1/N)$ . The normalization  $\mathcal{N}$  follows from

$$3 \left[ \int_{1/N}^{1-(1/N)} P(x) dx + \frac{2}{N} P_0 \right] = 1. \quad (13)$$

The factor 3 arises because the phase space simplex possesses three edges. For moderate and large system sizes  $N$  we obtain  $P_0 = \mathcal{N}/(2m)$  which dominates the left-hand side of Eq. (13), such that  $\mathcal{N} = Nm/3$ . The resulting probability density can again be inserted into (7) to provide an analytical result for the entropy production in the regime  $m \ll \frac{k}{2N} \ll 1$ :

$$\dot{S}_{m \ll m_c} = mN^2 \ln(m/k), \quad (14)$$

in perfect agreement with simulations [Fig. 3(a)]. The entropy production for small mutation rates is proportional to the squared system size. Decreasing  $m$  lowers the entropy production in proportion because mutations are the process that restarts the cyclic dynamics once a corner state has been reached. Mutations therefore limit the dynamics to a time scale proportional to  $m$ .

In conclusion, we have examined the global entropy production in the steady state of a cyclic population model. At a critical mutation rate the system undergoes a transition from large oscillations along the phase space's boundary to small erratic oscillations around an internal fixed point. The entropy production peaks very near the critical mutation rate and decreases to zero away from it. We believe that, in a similar manner, the entropy production can yield valuable information about the nonequilibrium steady state of other stochastic systems. Indeed, in a recently studied model, because of a nonfixed system size and the extensivity of the entropy production, the *slope* of the entropy production peaks near a critical point [23]. Within our approach of a fixed system size we have investigated a stochastic system that displays a supercritical Hopf bifurcation. We found that the entropy production predicts the scale of the critical mutation rate: it peaks near the Hopf bifurcation, at a mutation rate of about 1/4 of the critical one [24]. Understanding the certain discrepancy between the maximum and the critical value may open a route to a more general understanding of the role of entropy produc-

tion. Because of the universality of the Hopf bifurcation we conclude that our approach is valid for a wide class of nonequilibrium systems, namely, those that exhibit a transition from small, erratic oscillations to limit-cycle-like ones, including systems with spatial degrees of freedom [25].

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