Stochastic predation-prey competition: biodiversity and species extinction

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historic Hudson Bay Company data

Lotka-Volterra model (A. J. Lotka, 1920; V. Volterra, 1926)

- predators: $A \longrightarrow 0$ death, rate μ
- prey: $B \longrightarrow B + B$ birth, rate σ
- predation: $A + B \longrightarrow A + A$, rate λ

mean-field rate equations for homogeneous densities:

$$egin{array}{rcl} rac{da(t)}{dt}&=&-\mu\, a(t)+\lambda\, a(t)\, b(t)\ rac{db(t)}{dt}&=&\sigma\, b(t)-\lambda\, a(t)\, b(t) \end{array}$$

conserved quantity: $K = \lambda(a + b) - \sigma \ln a - \mu \ln b$

 \longrightarrow limit cycles, population oscillations





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Generalization: multi-species Lotka-Volterra rate equations

$$\frac{dx_i(t)}{dt} = x_i(t) \left(r_i + \sum_{j=1}^n \alpha_{i,j} x_j(t) \right)$$

r_i: intrinsic growth or decay

 $\alpha_{i,j}:$ represents interaction matrix that encodes competition between species i and j

very general equations that encompass many different cases (food chains)

example: cyclic Lotka-Volterra model $A_1 + A_2 \longrightarrow 2A_1, A_2 + A_3 \longrightarrow 2A_2, \dots, A_{n-1} + A_n \longrightarrow 2A_{n-1},$ $A_n + A_1 \longrightarrow 2A_n$

original (two-species) Lotka-Volterra model

$$\begin{array}{rcl} \displaystyle \frac{da(t)}{dt} & = & -\mu \, a(t) + \lambda \, a(t) \, b(t) \\ \displaystyle \frac{db(t)}{dt} & = & \sigma \, b(t) - \lambda \, a(t) \, b(t) \end{array}$$

stationary states (fixed point)

•
$$(a^*, b^*) = (0, 0) \longrightarrow$$
 extinction

- $(a^*, b^*) = (0, \infty) \longrightarrow$ predators extinct, Malthusian prey proliferation
- $(a_c, b_c) = (\sigma/\lambda, \mu/\lambda) \longrightarrow$ species coexistence

linearization about coexistence stationary state

 \implies purely oscillatory kinetics with characteristic frequency $\omega=\sqrt{\mu\sigma}$

conservation law for K and related purely oscillatory motion are special features of the deterministic model equations

results unstable with respect to perturbations:

- model modifications
- spatial degrees of freedom
- stochasticity

Stochastic Lotka-Volterra model: no space dependence

A predators and B preys \longrightarrow discrete degrees of freedom

$$\frac{dP(A, B; t)}{dt} = \lambda(A-1)(B+1)P(A-1, B+1; t) + \mu(A+1)P(A+1, B; t) - (\mu A + \sigma B + \lambda A B)P(A, B; t)$$

only one stable steady state: $P_S(A = 0, B = 0) = 1$ and $P_S(A \neq 0, B \neq 0) = 0$ as $t \longrightarrow \infty$, empty (absorbing) state will be reached

at finite times: erratic population oscillations (*resonant amplification mechanism*) McKane/Newman '05



each lattice site either empty or occupied by a predator or a prey mean-field rate equations for the particle densities

$$\frac{da}{dt} = -\mu a(t) + \lambda a(t)b(t)$$

$$\frac{db}{dt} = \sigma[1 - a(t) - b(t)]b(t) - \lambda a(t)b(t)$$

absorbing state: $\lambda < \mu$; $a \longrightarrow 0$, $b \longrightarrow 1$

 $\lambda > \mu$: active phase: A and B coexist active/absorbing phase transition \longrightarrow nonequilibrium phase transition

Stochastic Lotka-Volterra model on a lattice

coarsening in one dimension – final state: system full of prey predator (red) and prey (blue) domains



complicated space-time pattern in two space dimensions (Mobilia, Georgiev, Täuber '07)

Cyclic dominance of competing species

real-world example: competing bacterial strains (*Escherichia coli*) (Kerr et al. '02)



three cyclically competing species: Rock-Paper-Scissors game

$$a+b \xrightarrow{k_a} a+a$$

$$b+c \xrightarrow{k_b} b+b$$

$$c+a \xrightarrow{k_c} c+c$$



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Rock-Paper-Scissors game

three cyclically competing species: Rock-Paper-Scissors game

$$\begin{array}{rrrr} a+b & \stackrel{k_a}{\longrightarrow} & a+a \\ b+c & \stackrel{k_b}{\longrightarrow} & b+b \\ c+a & \stackrel{k_c}{\longrightarrow} & c+c \end{array}$$

three ways of realizing mobility when on a lattice:

• exchange of individuals

$$a+b \stackrel{s_{ab}}{\leftarrow} b+a$$
$$b+c \stackrel{s_{bc}}{\leftarrow} c+b$$
$$c+a \stackrel{s_{ca}}{\leftarrow} a+c$$

conserved quantity: $N_a + N_b + N_c = N$

- empty sites
- multiple occupancy of sites

Formation of space-time pattern

May-Leonard model

three species on a two-dimensional lattice separation of predation and reproduction

 $\begin{array}{c} A+B \longrightarrow A+0 \\ A+0 \longrightarrow A+A \end{array}$



What about more than three species?

Simplest generalization: four species

$$a + b \xrightarrow{k_a} a + a$$
$$b + c \xrightarrow{k_b} b + b$$
$$c + d \xrightarrow{k_c} c + c$$
$$d + a \xrightarrow{k_d} d + a$$



What about more than three species?

Simplest generalization: four species

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$$c + d \xrightarrow{k_c} c + c$$
$$d + a \xrightarrow{k_d} d + a$$



formation of partner-pairs!

configuration space for four species



ac and bd pairs do not interact

 \implies final (absorbing) state displays coexistence of these pairs

every point along a - c and b - d edges represents such a state $\implies 2(N + 1)$ absorbing states

mean field approximation for the evolution of the averages of the fractions

$$A(t) \equiv \sum_{\{N_m\}} (N_a/N) P(\{N_m\}; t) \quad \text{etc.}$$

neglect all correlations and replace averages of products by the products of averages

MF equations $(k_a + k_b + k_c + k_d = 1)$:

$$\partial_t A = [k_a B - k_d D] A$$

$$\partial_t B = [k_b C - k_a A] B$$

$$\partial_t C = [k_c D - k_b B] C$$

$$\partial_t D = [k_d A - k_c C] D$$

mean field approximation for the evolution of the averages of the fractions

$$A(t) \equiv \sum_{\{N_m\}} (N_a/N) P(\{N_m\}; t) \quad \text{etc.}$$

neglect all correlations and replace averages of products by the products of averages

MF equations $(k_a + k_b + k_c + k_d = 1)$:

$$\partial_t \ln A = k_a B - k_d D$$

$$\partial_t \ln B = k_b C - k_a A$$

$$\partial_t \ln C = k_c D - k_b B$$

$$\partial_t \ln D = k_d A - k_c C$$

contributions from a single species to the growth/decay of two other species:

$$\partial_t \left[k_b \ln A + k_a \ln C \right] = \lambda D$$

$$\partial_t \left[k_c \ln A + k_d \ln C \right] = \lambda B$$

$$\partial_t [k_c \ln B + k_b \ln D] = -\lambda A$$

$$\partial_t [k_d \ln B + k_a \ln D] = -\lambda C$$

key control parameter: $\lambda \equiv k_a k_c - k_b k_d$

quantity

$$Q \equiv \frac{A^{k_b + k_c} C^{k_d + k_a}}{B^{k_c + k_d} D^{k_a + k_b}}$$

evolves in an extremely simple manner:

$$Q(t) = Q(0) e^{\lambda t}$$

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 $\lambda = 0 \longrightarrow k_a k_c = k_b k_d \longrightarrow Q$ is a constant of motion

numerator/denominator of Q are constant each defines a (generalized) hyperbolic sheet intersection is a closed loop (\sim edge of a saddle)



saddle-shaped orbits and fixed points

 $\lambda \neq 0$

$$Q(t) = Q(0) e^{\lambda t}$$
 with $Q \equiv \frac{A^{k_b + k_c} C^{k_d + k_a}}{B^{k_c + k_d} D^{k_a + k_b}}$

spirals and arrows



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starting from symmetry point with $\lambda = -0.0273$

going beyond mean field approximation: numerical simulations

 $\lambda = 0$: stochastic effects



1000 particles, $(k_a, k_b, k_c, k_d) = (0.4, 0.4, 0.1, 0.1)$ and $(A_0, B_0, C_0, D_0) = (0.02, 0.10, 0.48, 0.40)$

 $\lambda \neq 0$: extinction events



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 $(k_a,k_b,k_c,k_d)=(0.1,0.0001,0.1,0.7999)$ and $(A_0,B_0,C_0,D_0)=(0.1,0.7,0.1,0.1)$

$\lambda \neq 0$: extinction events



 $(k_a, k_b, k_c, k_d) = (0.1, 0.0001, 0.1, 0.7999)$ and $(A_0, B_0, C_0, D_0) = (0.1, 0.7, 0.1, 0.1)$

going beyond mean field approximation: numerical simulations

 $\lambda \neq 0$: extinction events



 $(k_a, k_b, k_c, k_d) = (0.1, 0.0001, 0.1, 0.7999)$ and $(A_0, B_0, C_0, D_0) = (0.1, 0.7, 0.1, 0.1)$

extinction time distributions for small systems



Symmetric interaction and swapping rates for four species space-time diagrams



 $k = 0.8, \ s = 0.2$ $k = 0.1, \ s = 0.9$ $k = 0.01, \ s = 0.99$

Symmetric interaction and swapping rates for four species average domain size (for k + s = 1)



 \rightarrow exchanges speed up the coarsening process!

four species: coexistence, but no well formed space-time patterns



k = 1 and s = 0

Coarsening in two dimensions

four species with exchanges between individuals belonging to a partner-pair

 \implies coarsening of partner-pair domains





$$k = 0.8$$
 and $s = 0.2$, $s_n = 0.2$

Coarsening in two dimensions

correlation length from the correlation function

$$C(t, ec{r}) = \sum_{i} \left[\left\langle n_i(t, ec{r}) n_i(0, ec{0})
ight
angle - \left\langle n_i(t, ec{r})
ight
angle \left\langle n_i(0, ec{0})
ight
angle
ight] \; .$$

 $n_i(t, \vec{r})$: occupation number



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dynamical exponent z = 2

Extinction times

extinction time distributions for small systems



two different paths to extinction revealed by the distribution function

N different species in two dimensions

 X_i : member of species *i*

reaction scheme:

$$egin{array}{rcl} X_i + X_j & & \stackrel{\delta_{ij}}{\longrightarrow} & \emptyset + X_i & ext{predation} \ X_i + \emptyset & \stackrel{\gamma_i}{\longrightarrow} & X_i + X_i & ext{birth} \ X_i + \emptyset & & \stackrel{\beta_i}{\longrightarrow} & \emptyset + X_i & ext{diffusion} \end{array}$$

 $X_i + X_j \stackrel{lpha_{ij}}{\longrightarrow} X_j + X_i$ swapping

model (N, r): generalized May-Leonard model with N species where each species preys on r other species in a cyclic way

(N, 1): N-species cyclic Lotka-Volterra game discussed until now

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space-time pattern and coarsening model (6,5)





space-time pattern and coarsening

model (6,4)





space-time pattern and coarsening

model (6,3)





space-time pattern and coarsening

model (6,2)





space-time pattern and coarsening model (6,1)





• domains grow as $t^{1/2}$



• interface width: (6,3)



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• square of the adjacency matrix contains all information about preferred partnership formations

Example: model (6,4)





 square of the adjacency matrix contains all information about preferred partnership formations

Example: model (6,4)



adjacency matrix



 square of the adjacency matrix contains all information about preferred partnership formations

Example: model (6,4)



square of the adjacency matrix

$$\mathbf{B} = \begin{pmatrix} 3 & 2 & 2 & 2 & 3 & 4 \\ 4 & 3 & 2 & 2 & 2 & 3 \\ 3 & 4 & 3 & 2 & 2 & 2 \\ 2 & 3 & 4 & 3 & 2 & 2 \\ 2 & 2 & 3 & 4 & 3 & 2 \\ 2 & 2 & 2 & 3 & 4 & 3 & 2 \end{pmatrix}$$

 square of the adjacency matrix contains all information about preferred partnership formations

Example: model (6,4)



square of the adjacency matrix

 b_{ij} : number of directed paths of length 2 from vertex *i* to vertex *j* $(i \longrightarrow k \longrightarrow j)$

the enemy of my enemy is my friend \implies preferred ally of species j: $\max_{i} b_{ij}$

 square of the adjacency matrix contains all information about preferred partnership formations

Example: model (6,4)



square of the adjacency matrix

$$\mathbf{B} = \begin{pmatrix} 3 & 2 & 2 & 2 & 3 & 4 \\ 4 & 3 & 2 & 2 & 2 & 3 \\ 3 & 4 & 3 & 2 & 2 & 2 \\ 2 & 3 & 4 & 3 & 2 & 2 \\ 2 & 2 & 3 & 4 & 3 & 2 \\ 2 & 2 & 2 & 3 & 4 & 3 & 2 \end{pmatrix}$$

• square of the adjacency matrix contains all information about preferred partnership formations

Example: model (6,4)





Can be generalized to very complicated food networks using additional matrices!

• analytical expressions describing space-time patterns can be obtained through a complex Ginzburg-Landau approach

starting point: mean-field rate equations

- \longrightarrow single coexistence fixed point
- $\longrightarrow \mathsf{unstable} \text{ invariant manifold}$
- $\longrightarrow {\sf Stuart-Landau \ normal \ form \ on} \\ unstable \ manifold$

$$\dot{z}_{s} = (c_{1,s} - i\omega_{s})z_{s} - c_{2,s}(1 + ic_{3,s})z_{s} |z_{s}|^{2}$$

 expressions for linear spreading velocity, wavelength and frequency of spirals



Stochastic effects very important in population dynamics

- mean-field predictions not valid for small populations
- formation of complicated space-time patterns for three or more species that compete against each other
- generalized May-Leonard systems: coarsening processes with internal dynamics inside the growing domains
- exact method to predict alliance formation and space-time patterns for very general ecological networks

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No site restriction

- d = 2: always coexistence \implies absence of active/absorbing phase transition
- *d* = 1: always coexistence diffusion-dominated



reaction-dominated



 $\lambda = 0 \longrightarrow k_a k_c = k_b k_d$

 \boldsymbol{Q} is a constant of motion



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Coarsening in two dimensions

four species with exchanges between individuals belonging to a partner-pair

 \implies coarsening of partner-pair domains





$$k = 0.2$$
 and $s = 0.8$, $s_n = 0.8$

Symmetric interaction and swapping rates for three species $$_{\rm space}$$



 $k = 0.9, \ s = 0.1$ $k = 0.1, \ s = 0.9$

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Symmetric interaction and swapping rates for three species

average domain size (for k + s = 1)



swapping rates *s* larger than $s_c \approx 0.84$: exchange mechanism very effectively mixes different species \rightarrow coexistence of species is promoted

Asymmetric interaction and swapping rates for three species asymmetry in the rates \implies dominance of a single species

Example: $k_a = 0.45$, $k_b = k_c = 0.4$, $s_{bc} = s_{ca} = 0.4$



Asymmetric interaction and swapping rates for three species dynamical phase diagram for $k_b = k_c = 0.4$, $s_{bc} = s_{ca} = 0.4$



I: A dominates, II: B dominates, III: C dominates

Cyclic dominance of competing species

real-world example: lizard populations in southern California (Sinervo/Lively '96)



