



Multi-layered networks and emergence of spatio-temporal order

Evolutionary game theory approach

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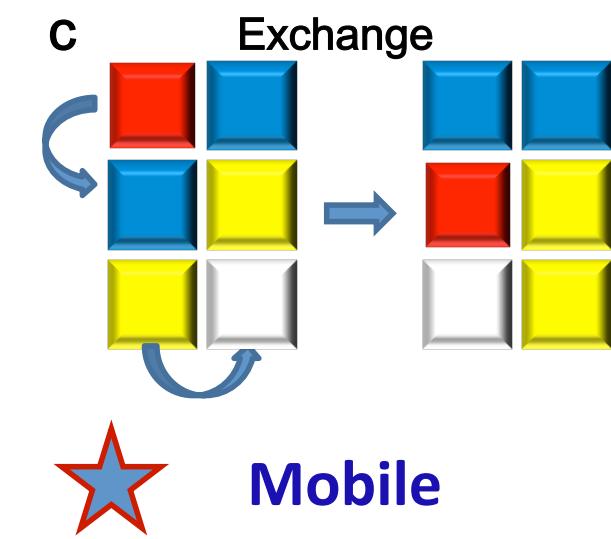
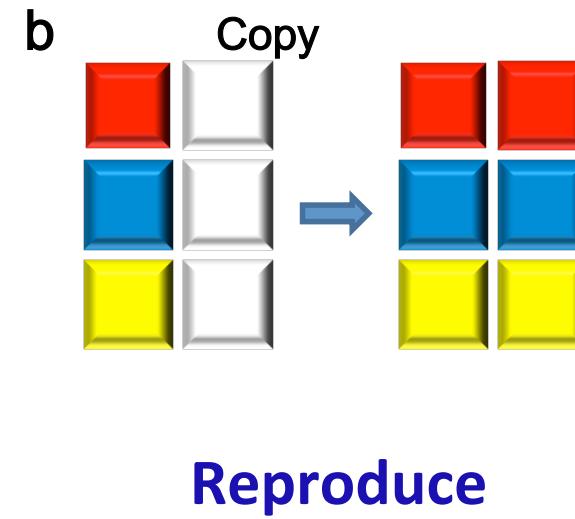
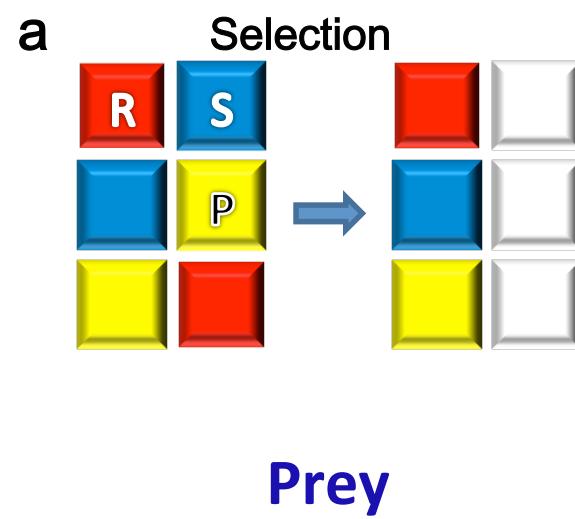
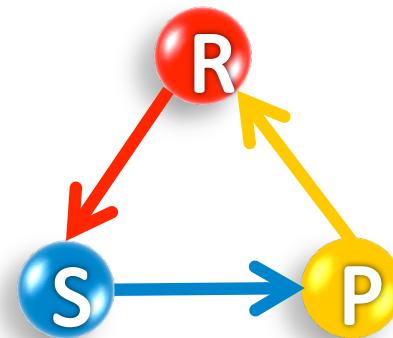
Collaborators:

Dr. W.-X. Wang, BNU; Dr. X. Ni, Arizona

Prof. Ying-Cheng Lai, Arizona and Aberdeen

RPS Game Model

Reichenbach, Mobilia & Frey, Nature (London) 448, 1046, (2007).



Role of Epidemic Spreading in Coexistence

Model

- SIR mode: portion of infected individuals will die
- Spatial neighbors can infect each other
- Offspring is always healthy
- Our findings
 - Virus spread within the same species promotes species coexistence;
 - Spread among different species jeopardizes species coexistence.

SIR Spreading Within Species

infection

I	n
I	n
I	n

rate λ

I	I
I	I
I	I

death

I	n
I	n
I	n

rate δ

	n
	n
	n

copy

I	
I	
I	

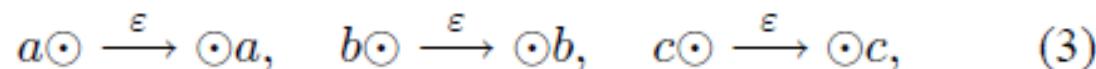
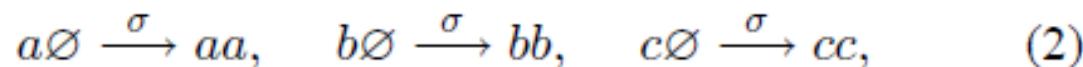
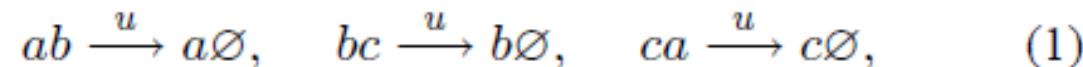
I	n
I	n
I	n

Epidemic spreading does not affect predation and mobility

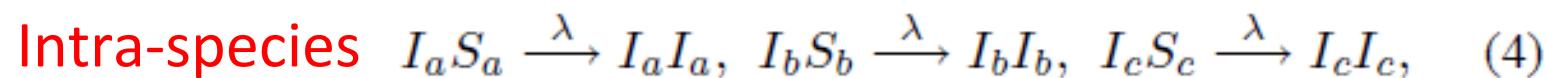
Offspring is always normal

RPS Model with Epidemic Spreading on 2D Lattice

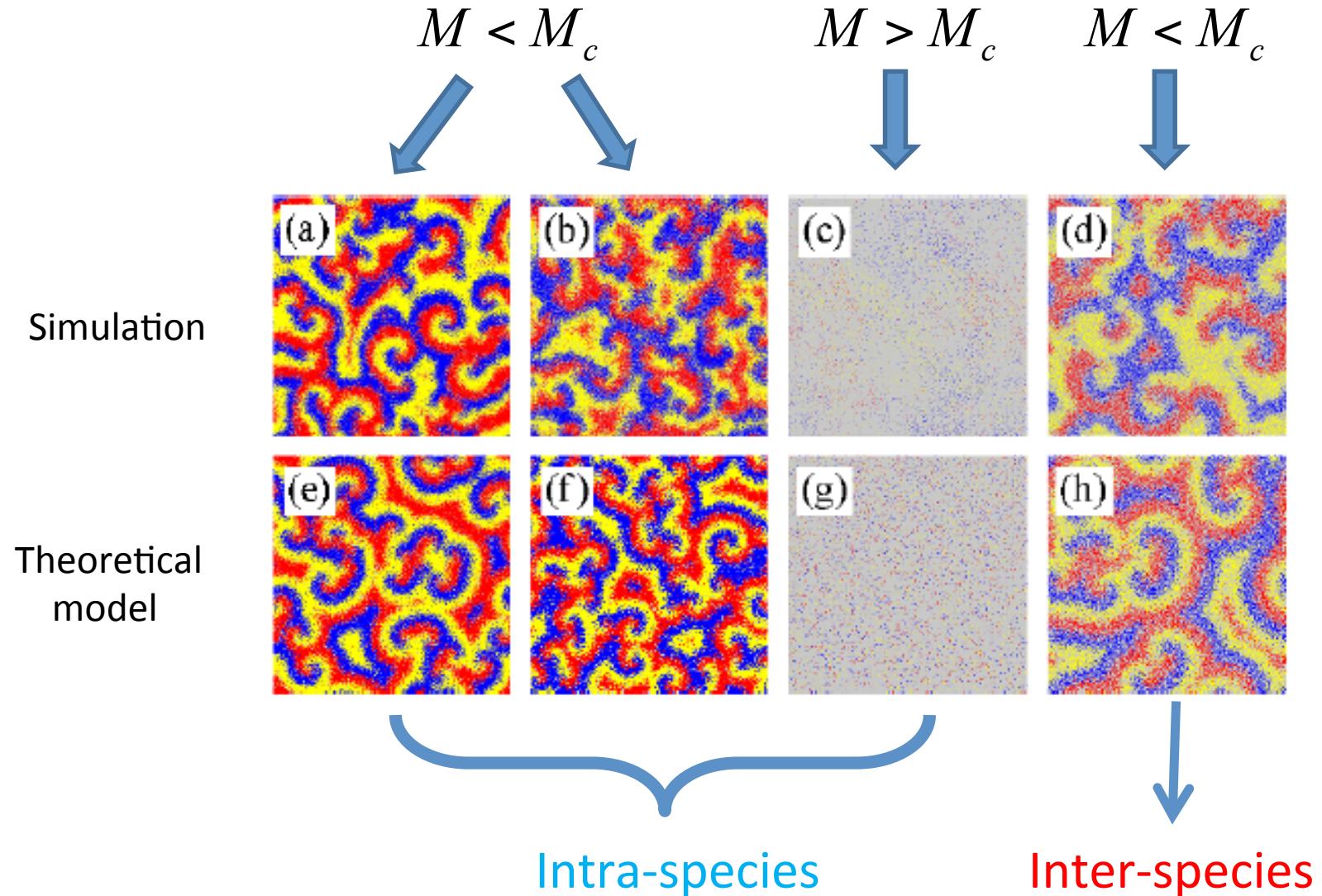
Cyclic competition, reproduction and mobility:



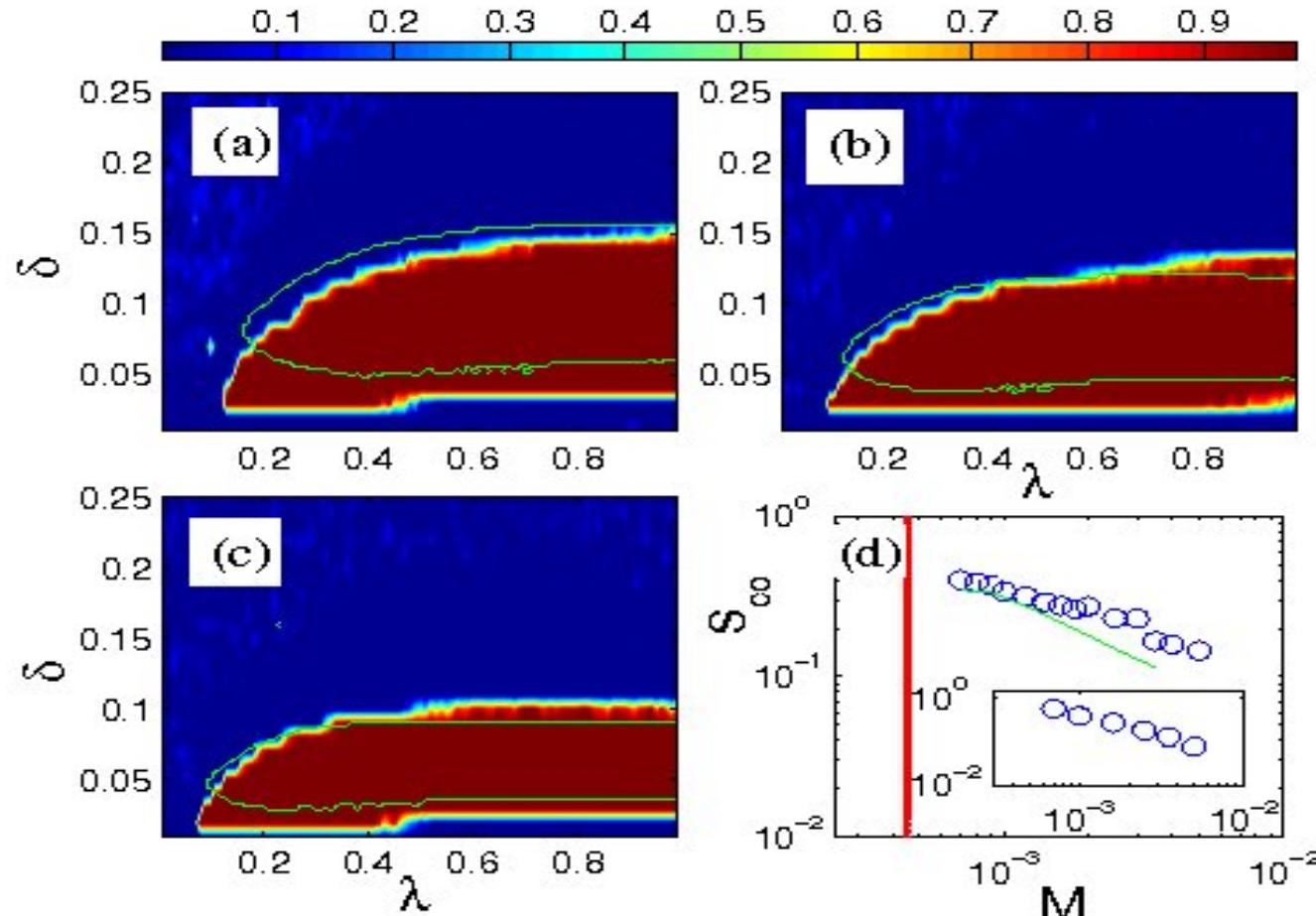
Epidemic spreading: infection and death



Spatial patterns



Parameter-Space Characterization



- (a) $M = 9 \times 10^{-4} > M_c$,
- (b) $M = 1.2 \times 10^{-3} > M_c$,
- (c) $M = 1.6 \times 10^{-3} > M_c$

Intra-species
virus spreading

Green boundaries –
Prediction from PDE
model

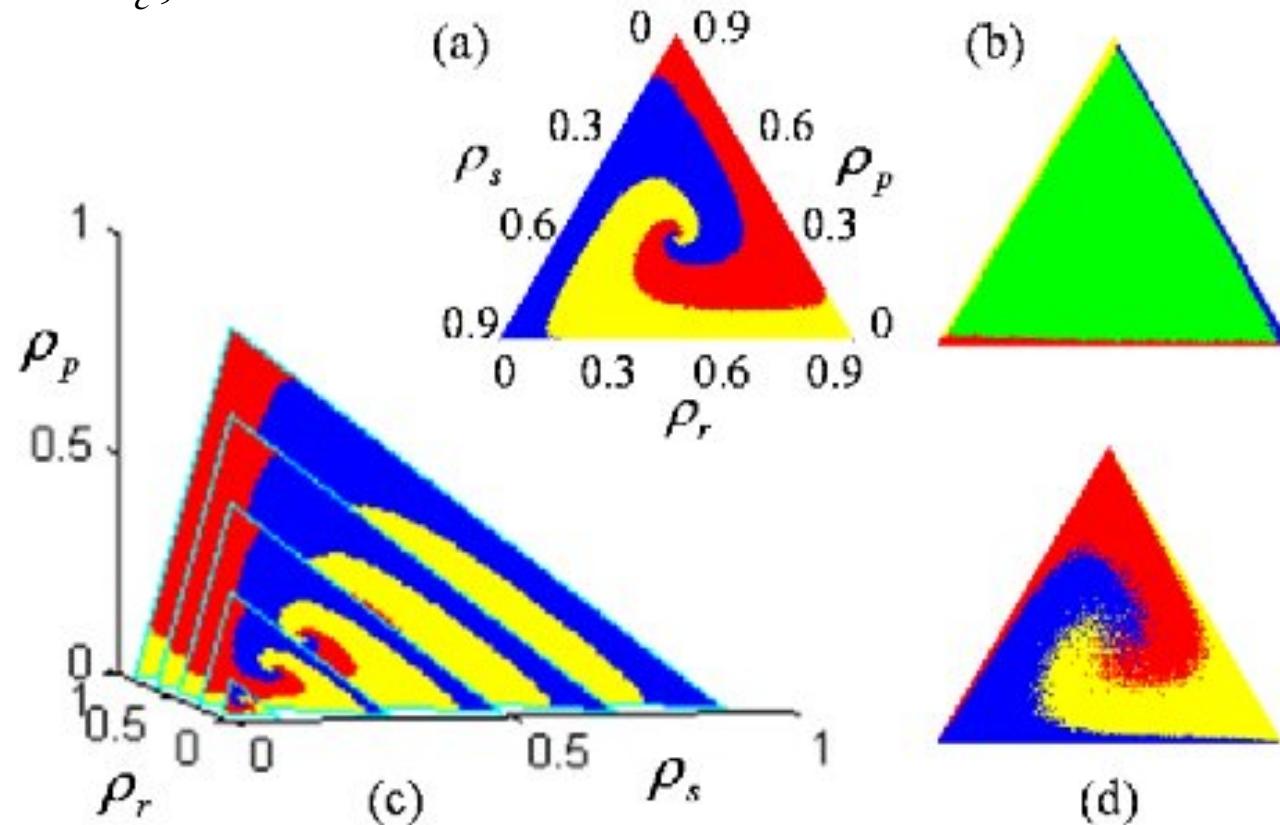
W.-X. Wang, Y.-C. Lai, and C. Grebogi, **Effect of epidemic spreading on species coexistence in rock-paper-scissors games**, Phys. Rev. E **81**, 046113(1-4) (2010)

Intra-Species Virus Spreading Induces Coexistence

$$M = 10^{-3} > M_c,$$

$$\lambda = 0.6$$

$$\delta = 0.13$$



- (a) No virus spreading; (b) With **intra-species** virus spreading;
- (c) Under **inter-species** spreading, basin structure predicted by PDE model for different values of initial density of empty sites;
- (d) Basin structure under **inter-species** virus spreading (from direct simulation)

Theoretical Model

Infection densities: $I_a(\mathbf{r}, t)$, $I_b(\mathbf{r}, t)$ and $I_c(\mathbf{r}, t)$

Susceptible densities: $S_a(\mathbf{r}, t)$, $S_b(\mathbf{r}, t)$ and $S_c(\mathbf{r}, t)$

Spatial site: $\mathbf{r} = (r_1, r_2)$ for 2d lattice

Neighboring location: $\mathbf{r} \pm \delta\mathbf{r} \cdot \mathbf{e}_i$, where \mathbf{e}_i is the basis of 2d lattice

$$\rho_a(\mathbf{r}, t) \equiv I_a(\mathbf{r}, t) + S_a(\mathbf{r}, t);$$

$$\rho_b(\mathbf{r}, t) \equiv I_b(\mathbf{r}, t) + S_b(\mathbf{r}, t); \quad \text{density of species}$$

$$\rho_c(\mathbf{r}, t) \equiv I_c(\mathbf{r}, t) + S_c(\mathbf{r}, t);$$

$$\rho(\mathbf{r}, t) \equiv \rho_a(\mathbf{r}, t) + \rho_b(\mathbf{r}, t) + \rho_c(\mathbf{r}, t); \quad \text{total density of species}$$

rescaled probabilities of

infection and death

$$\alpha = \frac{\lambda}{\varepsilon + u + \sigma}, \quad \beta = \frac{\delta}{\varepsilon + u + \sigma}$$



Evolutionary Equations

$$\partial_t I_a(\mathbf{r}, t) = \frac{1}{z} \sum_{\pm, i=1}^2 \left\{ 2\varepsilon [I_a(\mathbf{r} \pm \delta r \cdot \mathbf{e}_i, t) - I_a(\mathbf{r}, t)] + \alpha I_a(\mathbf{r} \pm \delta r \cdot \mathbf{e}_i, t) S_a(\mathbf{r}, t) - \beta I_a(\mathbf{r}, t) - u I_a(\mathbf{r}, t) \rho_c(\mathbf{r} \pm \delta r \cdot \mathbf{e}_i, t) \right\},$$

$$\begin{aligned} \partial_t S_a(\mathbf{r}, t) = & \frac{1}{z} \sum_{\pm, i=1}^2 \left\{ 2\varepsilon [S_a(\mathbf{r} \pm \delta r \cdot \mathbf{e}_i, t) - S_a(\mathbf{r}, t)] - \alpha I_a(\mathbf{r} \pm \delta r \cdot \mathbf{e}_i, t) S_a(\mathbf{r}, t) - u S_a(\mathbf{r}, t) \rho_c(\mathbf{r} \pm \delta r \cdot \mathbf{e}_i, t) \right. \\ & \left. + \sigma \rho_a(\mathbf{r} \pm \delta r \cdot \mathbf{e}_i, t) [1 - \rho(\mathbf{r}, t)] \right\} \end{aligned}$$

For $N \rightarrow \infty$ and lattice size fixed to 1, $\delta r \rightarrow 0$. Thus \mathbf{r} can be treated as a continuous variable together with the following expansion for I_a and S_a :

$$I_a(\mathbf{r} \pm \delta r \cdot \mathbf{e}_i, t) = I_a(\mathbf{r}, t) \pm \delta r \partial_i I_a(\mathbf{r}, t) + \frac{1}{2} \delta r^2 \partial_i^2 I_a(\mathbf{r}, t) + o(\delta r^2),$$

$$S_a(\mathbf{r} \pm \delta r \cdot \mathbf{e}_i, t) = S_a(\mathbf{r}, t) \pm \delta r \partial_i S_a(\mathbf{r}, t) + \frac{1}{2} \delta r^2 \partial_i^2 S_a(\mathbf{r}, t) + o(\delta r^2).$$

Up to second order: $\frac{2\varepsilon}{z} \sum_{\pm, i=1}^2 [I_a(\mathbf{r} \pm \delta r \cdot \mathbf{e}_i, t) - I_a(\mathbf{r}, t)] = \frac{\varepsilon}{2} \delta r^2 \partial_i^2 I_a(\mathbf{r}, t),$

$$\frac{2\varepsilon}{z} \sum_{\pm, i=1}^2 [S_a(\mathbf{r} \pm \delta r \cdot \mathbf{e}_i, t) - S_a(\mathbf{r}, t)] = \frac{\varepsilon}{2} \delta r^2 \partial_i^2 S_a(\mathbf{r}, t).$$

Partial Differential Equations

Rescaled exchange rate $\varepsilon = 2MN \Rightarrow \varepsilon\delta r^2 = 2M$.

$$\partial_t I_a(\mathbf{r}, t) = M\nabla^2 I_a(\mathbf{r}, t) + \alpha\psi_a(\mathbf{r}, t)S_a(\mathbf{r}, t) - \beta I_a(\mathbf{r}, t) - uI_a(\mathbf{r}, t)\rho_c(\mathbf{r}, t);$$

$$\partial_t S_a(\mathbf{r}, t) = M\nabla^2 S_a(\mathbf{r}, t) - \alpha\psi_a(\mathbf{r}, t)S_a(\mathbf{r}, t) - uS_a(\mathbf{r}, t)\rho_c(\mathbf{r}, t) + \sigma\rho_a(\mathbf{r}, t)[1 - \rho(\mathbf{r}, t)];$$

$$\partial_t I_b(\mathbf{r}, t) = M\nabla^2 I_b(\mathbf{r}, t) + \alpha\psi_b(\mathbf{r}, t)S_b(\mathbf{r}, t) - \beta I_b(\mathbf{r}, t) - uI_b(\mathbf{r}, t)\rho_a(\mathbf{r}, t);$$

$$\partial_t S_b(\mathbf{r}, t) = M\nabla^2 S_b(\mathbf{r}, t) - \alpha\psi_b(\mathbf{r}, t)S_b(\mathbf{r}, t) - uS_b(\mathbf{r}, t)\rho_a(\mathbf{r}, t) + \sigma\rho_b(\mathbf{r}, t)[1 - \rho(\mathbf{r}, t)];$$

$$\partial_t I_c(\mathbf{r}, t) = M\nabla^2 I_c(\mathbf{r}, t) + \alpha\psi_c(\mathbf{r}, t)S_c(\mathbf{r}, t) - \beta I_c(\mathbf{r}, t) - uI_c(\mathbf{r}, t)\rho_b(\mathbf{r}, t);$$

$$\partial_t S_c(\mathbf{r}, t) = M\nabla^2 S_c(\mathbf{r}, t) - \alpha\psi_c(\mathbf{r}, t)S_c(\mathbf{r}, t) - uS_c(\mathbf{r}, t)\rho_b(\mathbf{r}, t) + \sigma\rho_c(\mathbf{r}, t)[1 - \rho(\mathbf{r}, t)];$$

where for intra - species transmission

$$\psi_a(\mathbf{r}, t) = I_a(\mathbf{r}, t), \quad \psi_b(\mathbf{r}, t) = I_b(\mathbf{r}, t), \quad \psi_c(\mathbf{r}, t) = I_c(\mathbf{r}, t),$$

for inter - species transmission

$$\psi_a(\mathbf{r}, t) = \psi_b(\mathbf{r}, t) = \psi_c(\mathbf{r}, t) = I_a(\mathbf{r}, t) + I_b(\mathbf{r}, t) + I_c(\mathbf{r}, t).$$

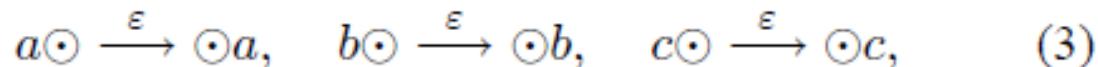
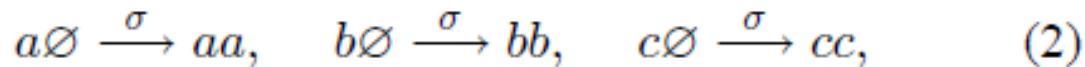
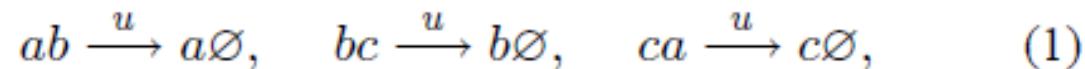
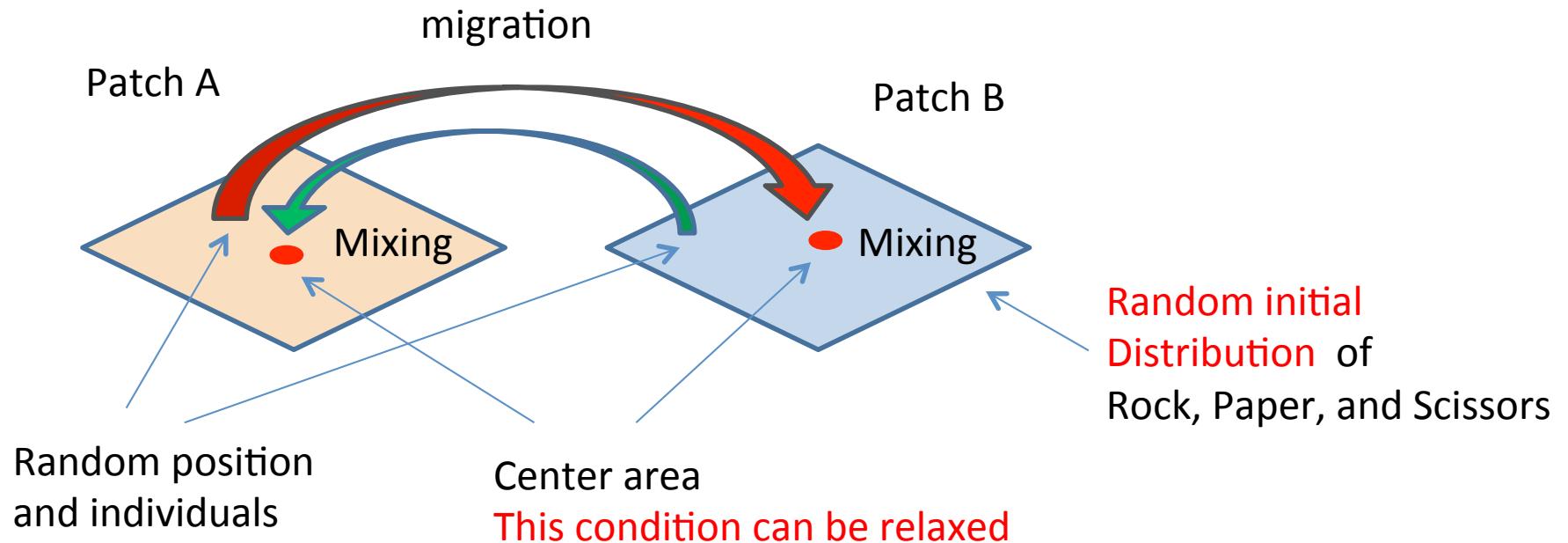


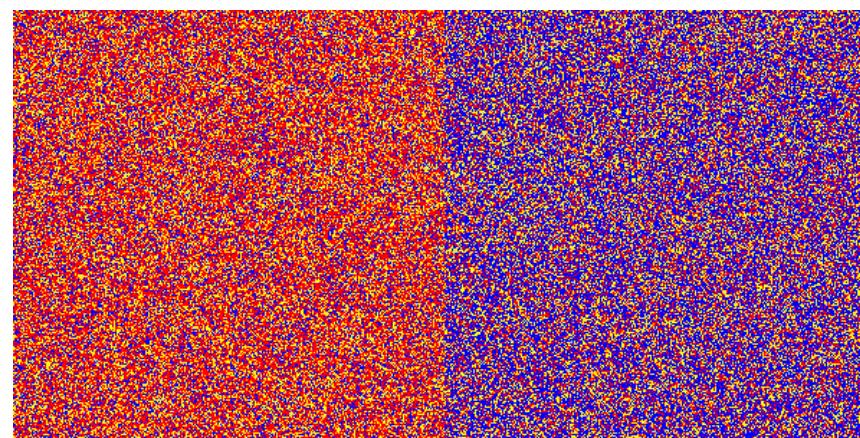
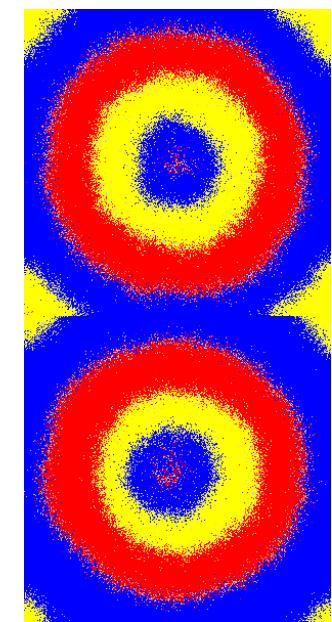
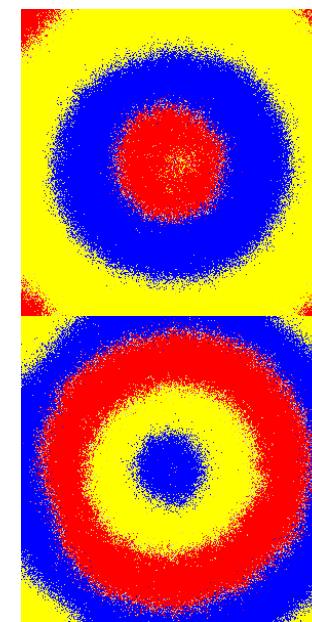
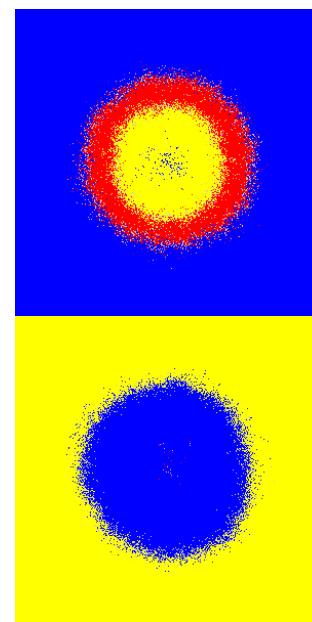
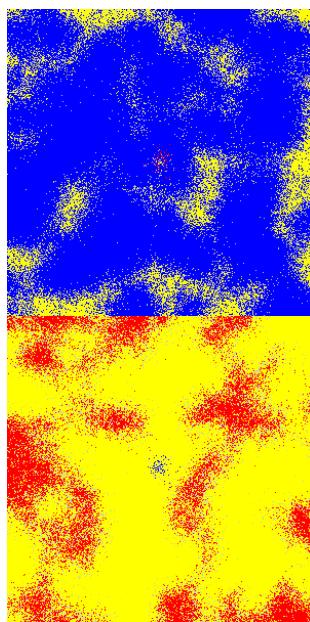
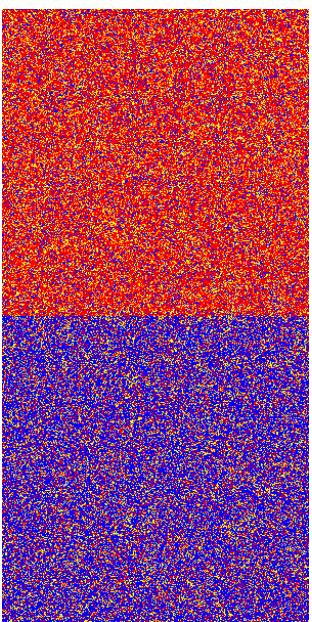
Inter-Patch Migration: Pattern Formation and Synchronization

- Target waves
- Synchronization and lag synchronization
- Coexistence of target wave and spiral waves
- Multi-target waves

Pattern formation, synchronisation and outbreak of biodiversity in cyclically competing games, W.-X. Wang, X. Ni, Y.-C. Lai, and C. Grebogi, Phys. Rev. E 83, 011917(1-9) (2011)

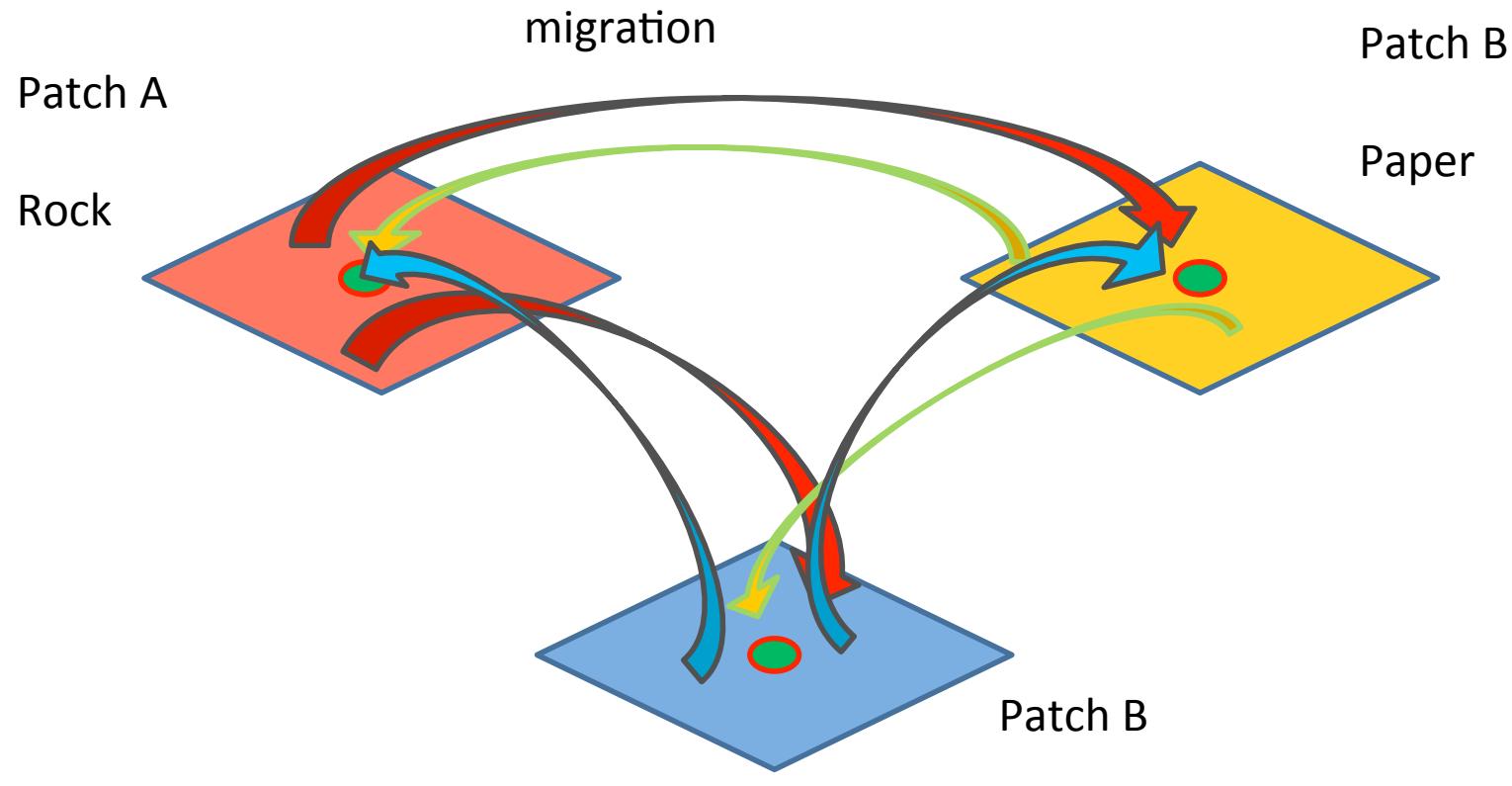
Inter-Patch Migration Model for Two Patches





\rightarrow t

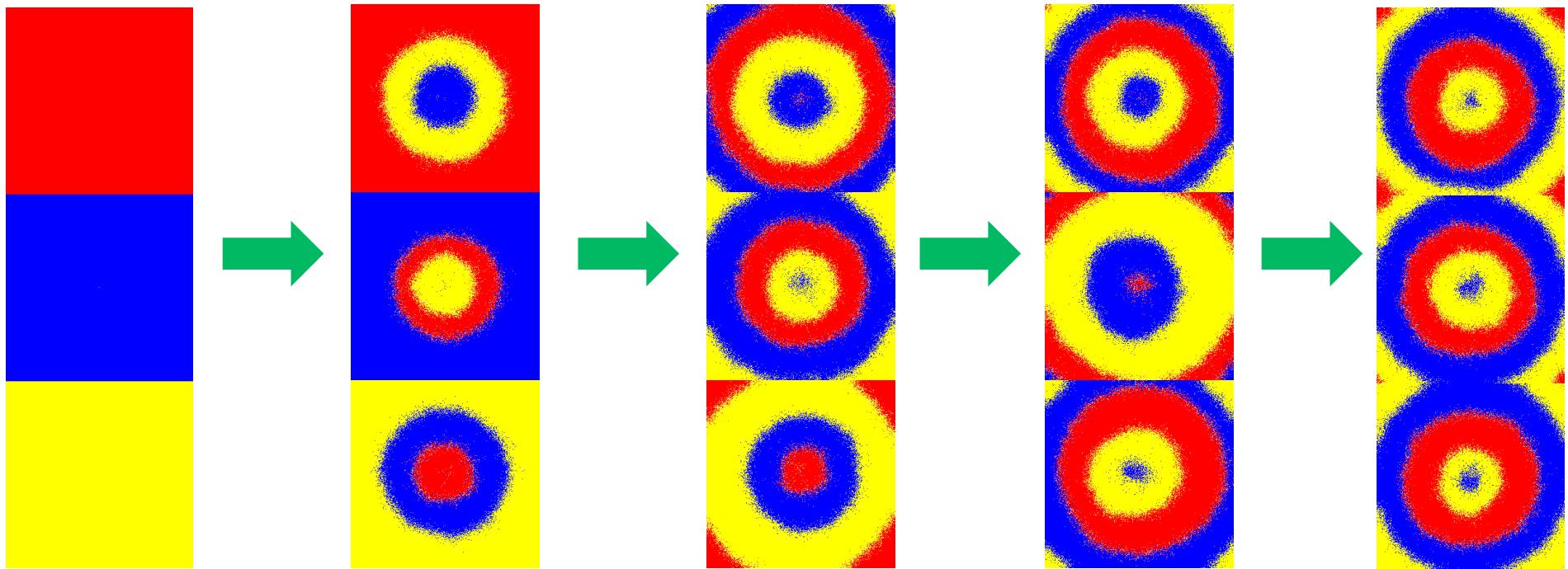
Migration Among Three Patches



Random origin of migration

Center destination point

Condition of destination can be relaxed

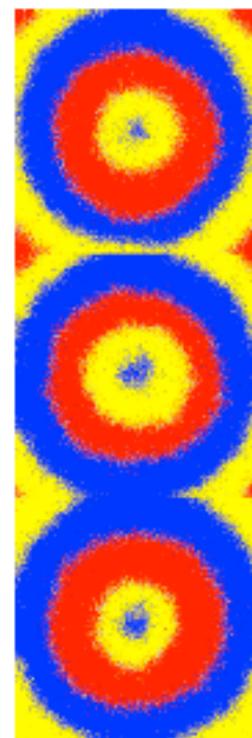


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Target Waves and Synchronisation

Three centred

A	rock
B	paper
C	scissors

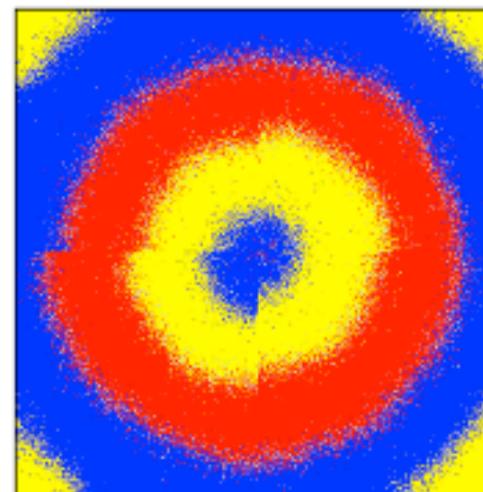


(a)

Four de-centred

A	B
rock	paper
C	D
scissors	empty

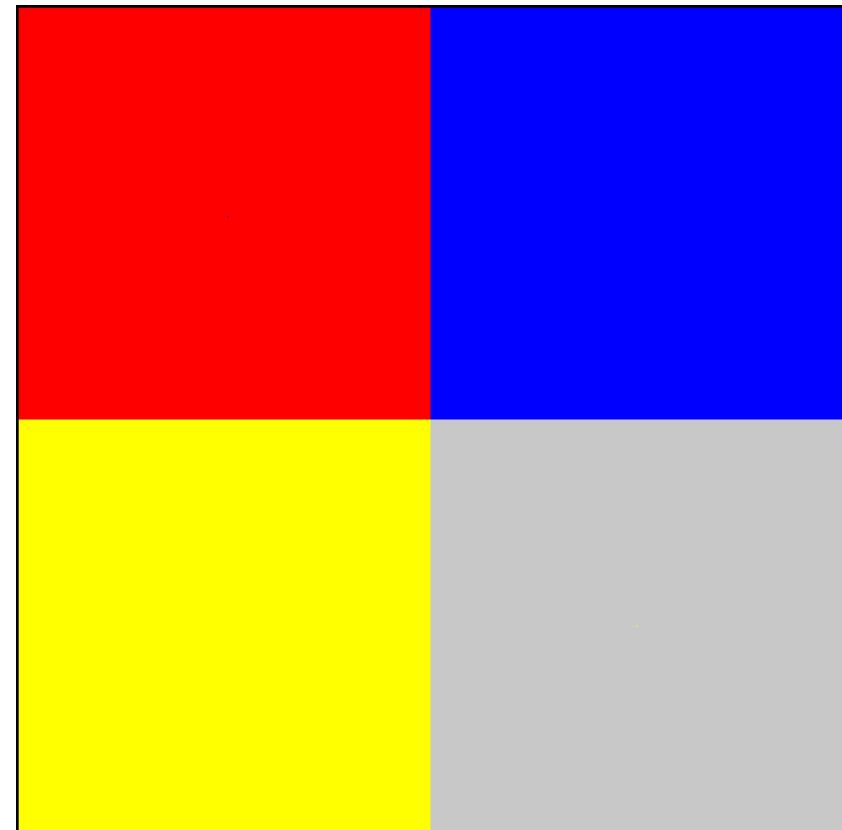
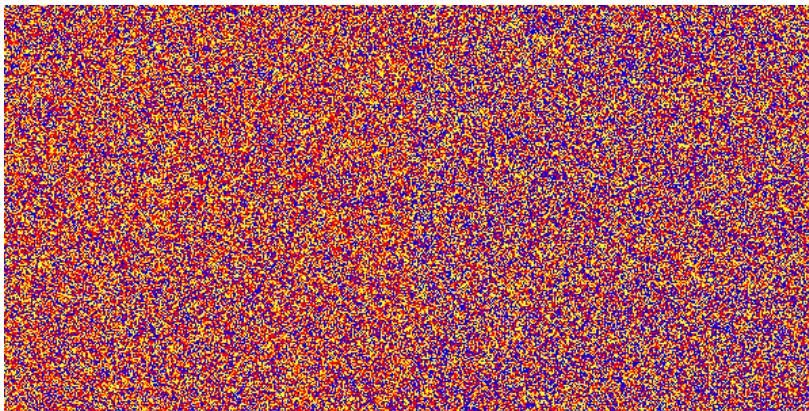
targets



(b)

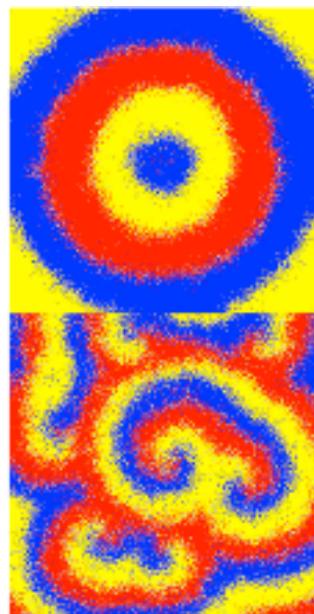


Coexistence of Target and Spiral Waves



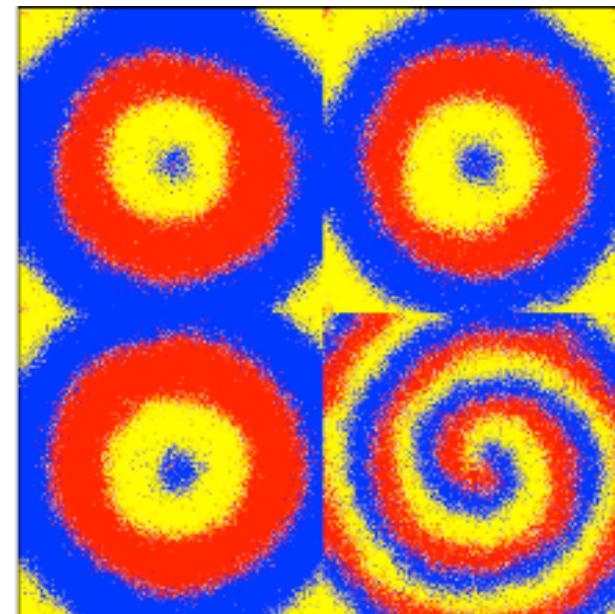
Coexistence of Target and Spiral Waves

Two-patch system



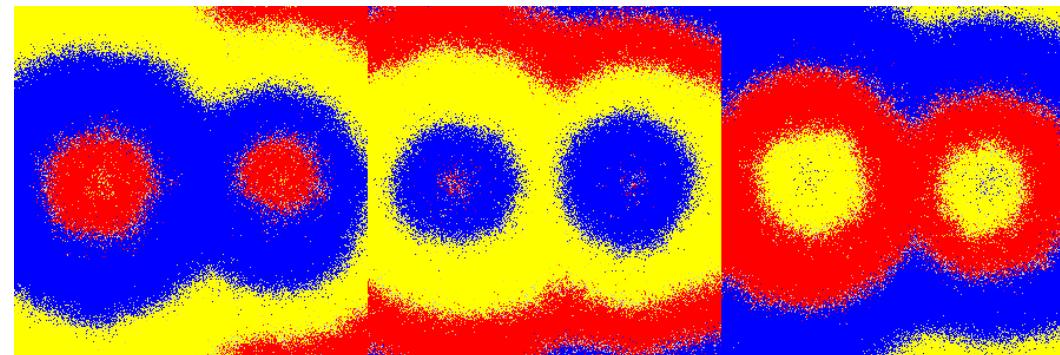
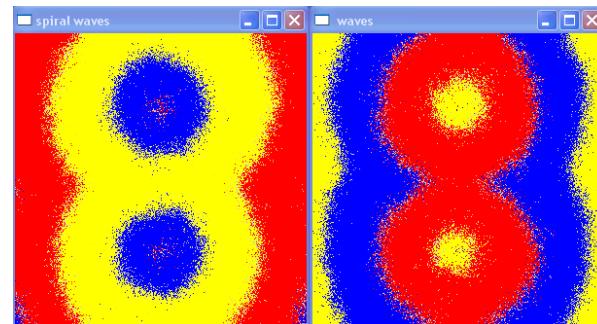
(a)

Four-patch system



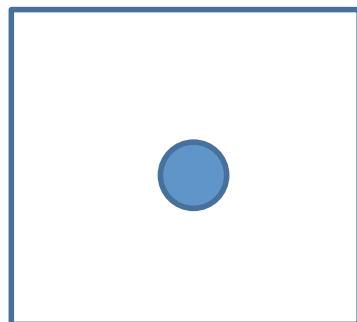
(b)

Multi-Target Waves

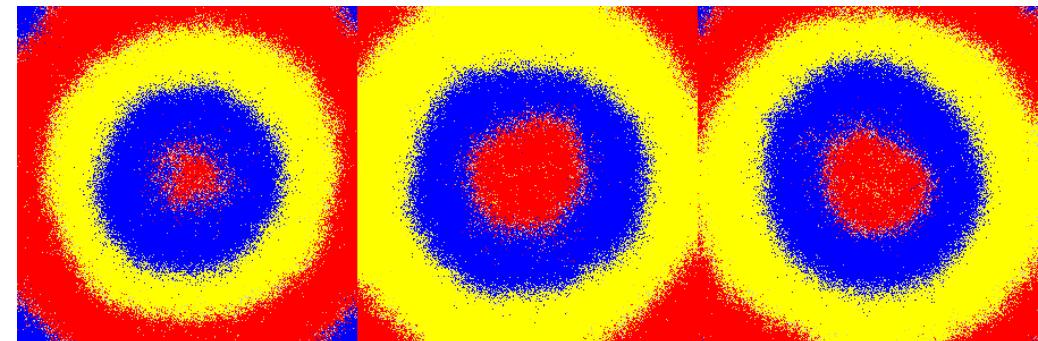


Migration to a target range

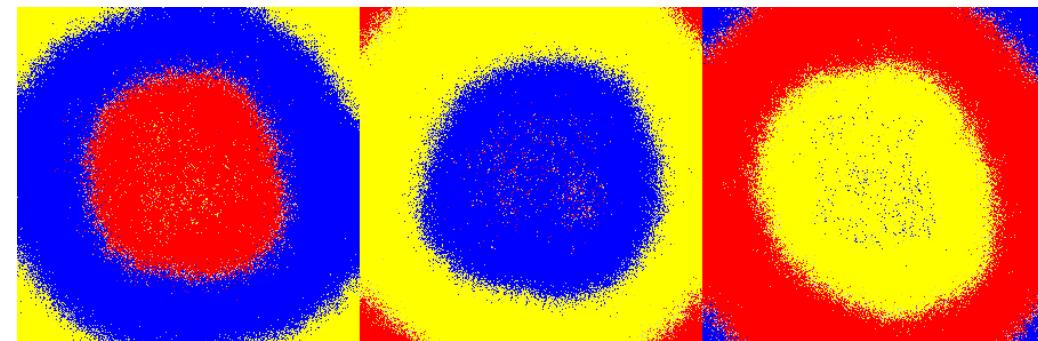
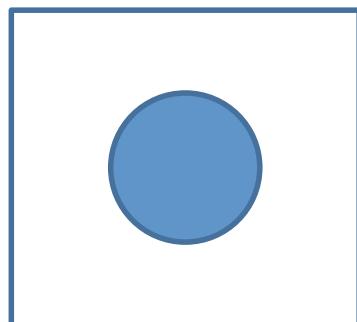
Small target range



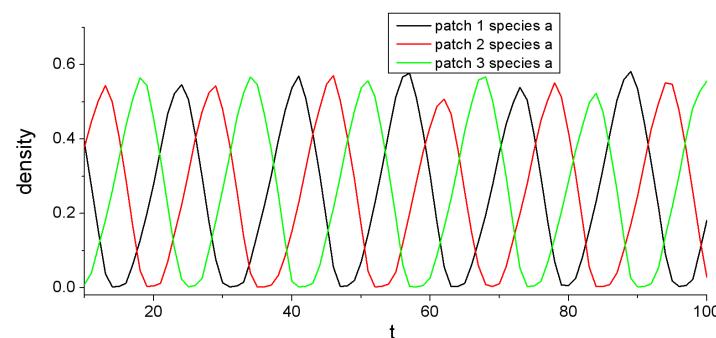
synchronization



Large target range



Lag synchronization

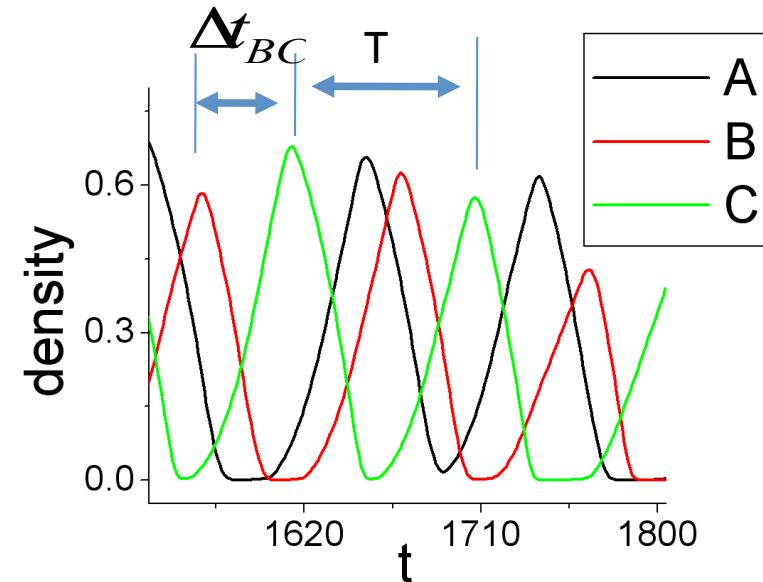
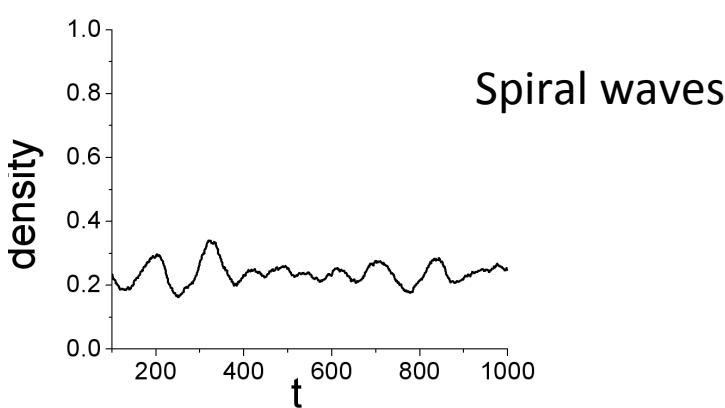
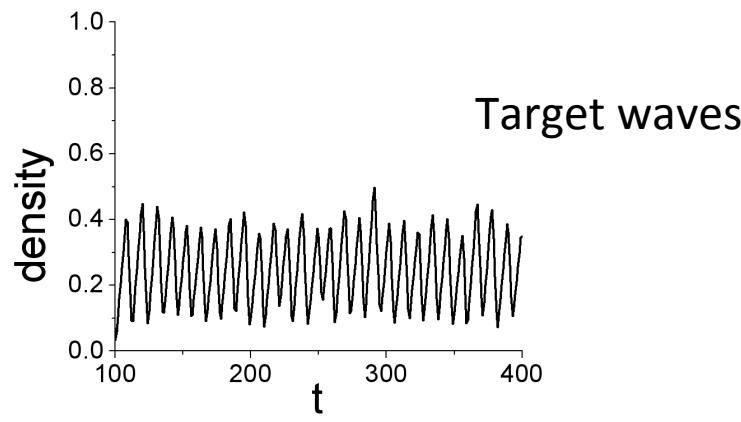


Quantitative Characterization

Distinguish target waves from spiral waves

Order parameter for
phase synchronization of target waves

Fourier transform

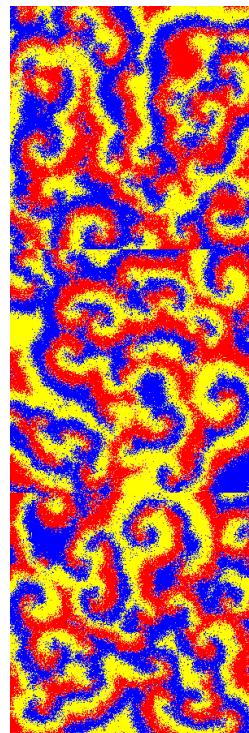


$$\eta_{BC} = 1 - \frac{\langle \min_{BC} \langle T \rangle - \Delta t_{BC} \rangle}{\langle \frac{T}{2} \rangle},$$

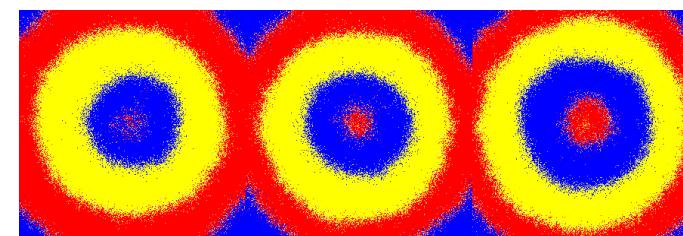
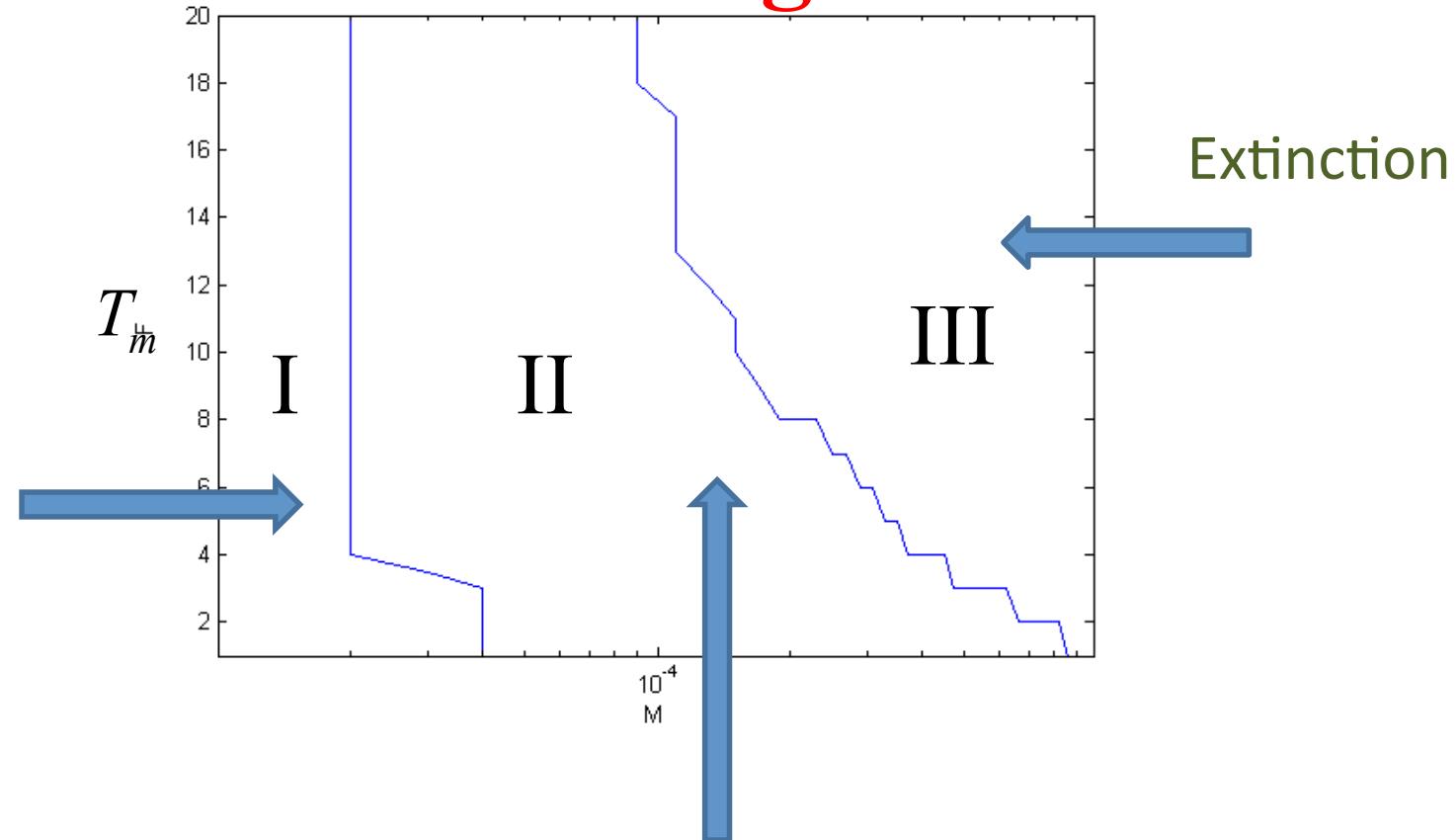
$$\eta = \frac{\eta_A + \eta_B + \eta_C}{3} \Leftarrow \text{order parameter}$$

$0 < \eta < 1$

$\eta = 1$: complete synchronization

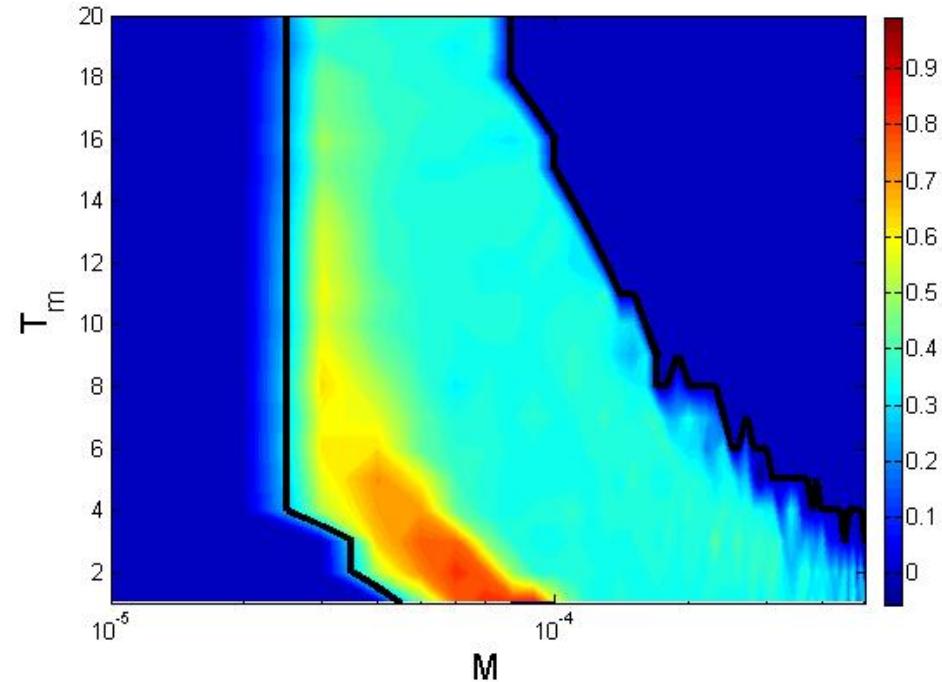


Phase diagram

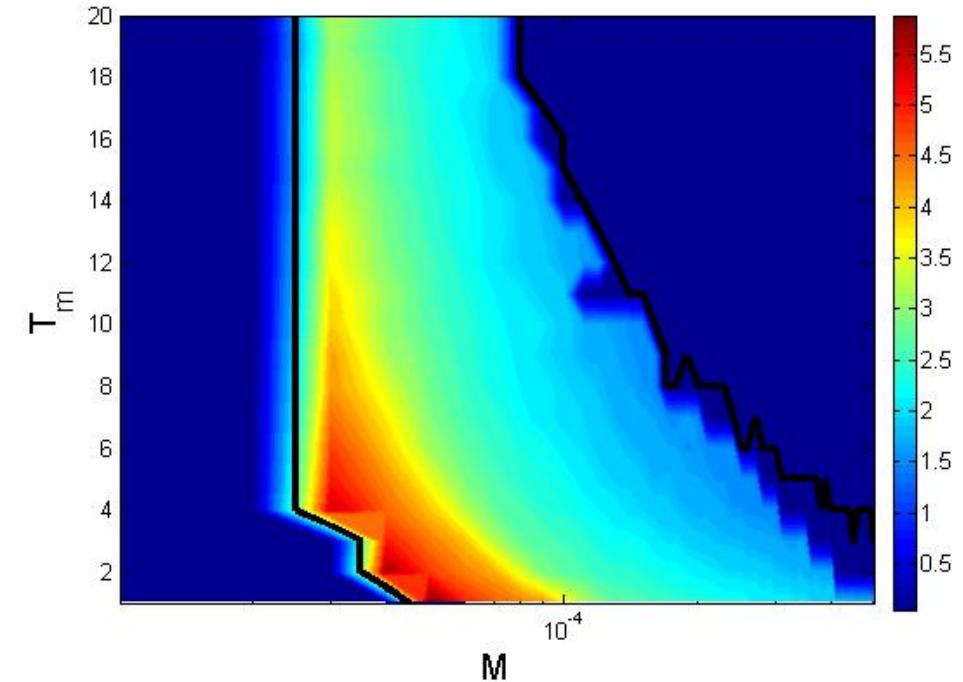


Synchronisability and Number of Rings

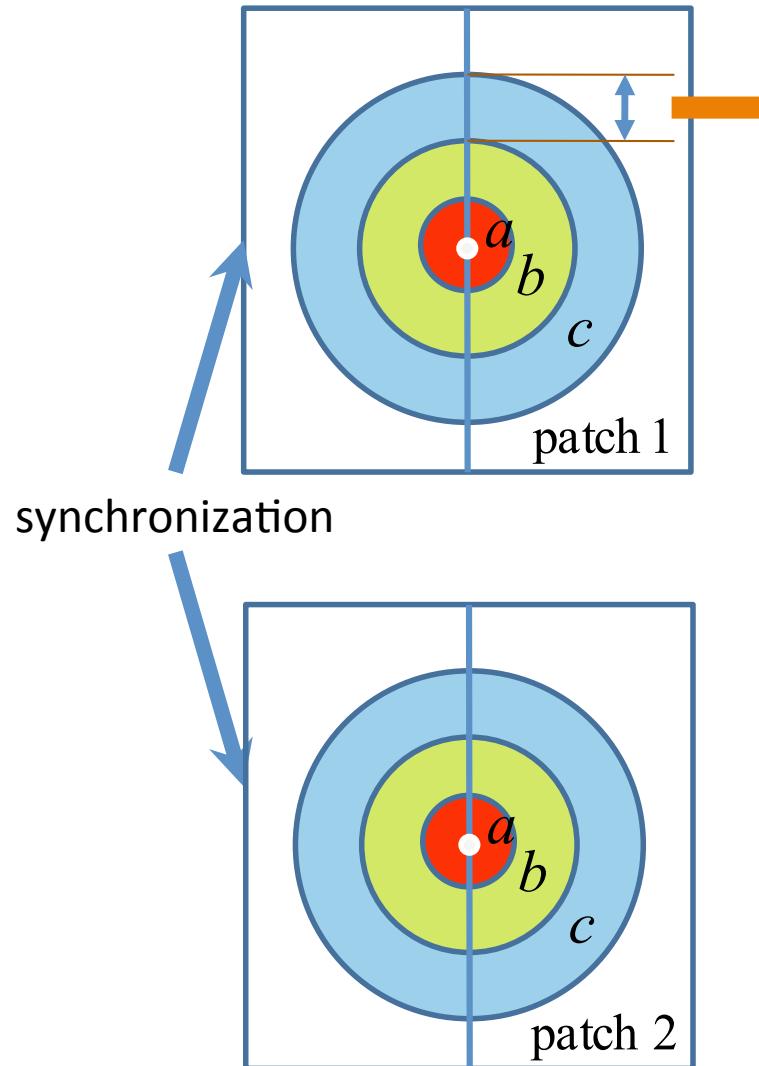
Synchronisability



Number of rings



Predicting the Number of Rings



L_r : length of rings

$$L_r = \frac{T_m}{\langle \rho_c \rangle_{patch2} - \langle \rho_b \rangle_{patch2}} \cdot V,$$

$$n_r = \frac{L}{\sqrt{2} L_r},$$

where V is the velocity of front propagation, and T_m is the period of inter-patch migration.

$V \Leftarrow \text{CGLE}$

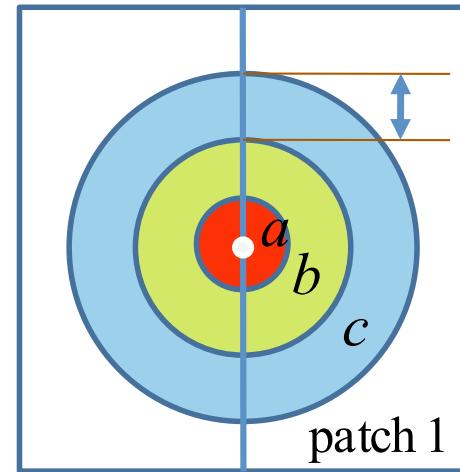
$$\partial_t z(r, t) = M \Delta z(r, t) + (c_1 - i\omega) z(r, t) - c_2 (1 - i c_3) |z(r, t)|^2 z(r, t),$$

linearize the CGLE around the state $z = 0$,

$$\partial_t z(r, t) = M \Delta z(r, t) + (c_1 - i\omega) z(r, t) + o(z^2),$$

$$\Rightarrow V = 2\sqrt{c_1 M},$$

$$\text{where } c_1 = \frac{1}{2} \frac{\mu\sigma}{3\mu + \sigma}.$$



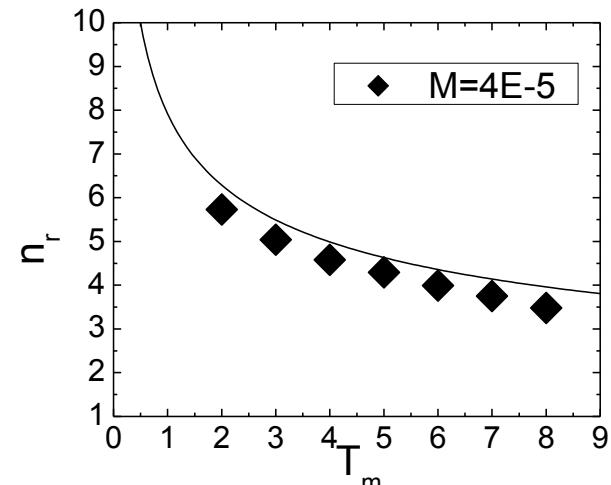
Due to synchronization, $\langle \rho_c \rangle_{patch1} = \langle \rho_c \rangle_{patch2}$,

and $\langle \rho_b \rangle_{patch1} = \langle \rho_b \rangle_{patch2}$.

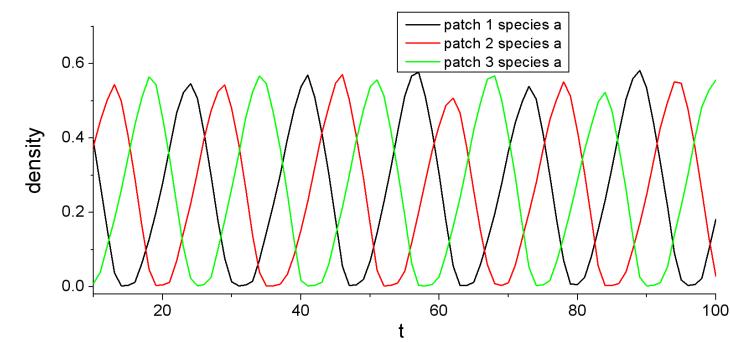
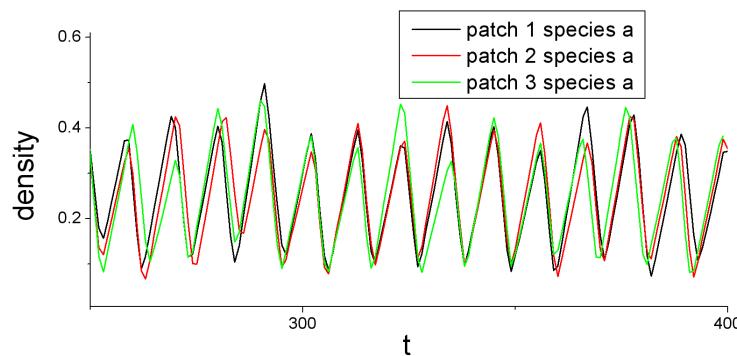
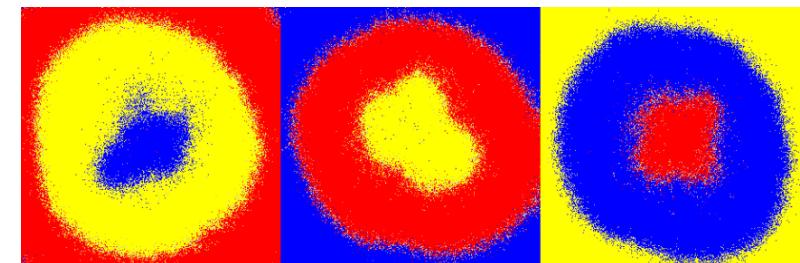
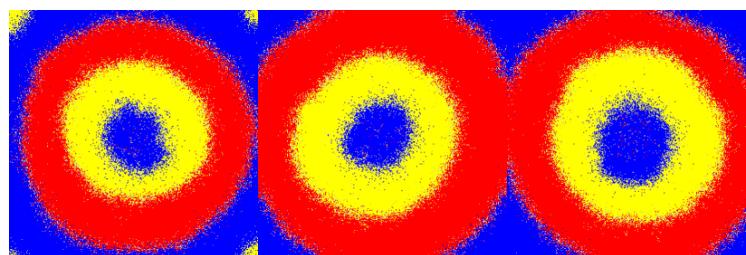
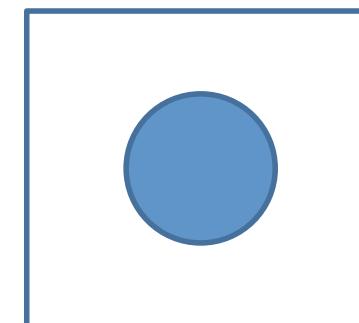
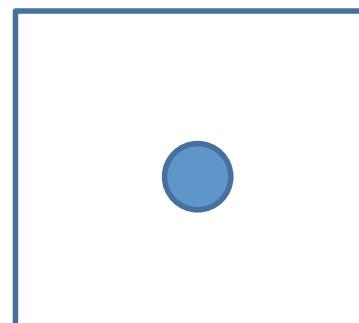
$$\langle \rho_c \rangle_{patch1} = \int_{L_r}^{2L_r} \frac{\pi(x + L_r)^2 - \pi x^2}{L_r \cdot L^2} dx = \frac{4\pi L_r^2}{L^2}.$$

$$\langle \rho_b \rangle_{patch1} = \int_0^{L_r} \frac{\pi(x + L_r)^2 - \pi x^2}{L_r \cdot L^2} dx = \frac{2\pi L_r^2}{L^2}.$$

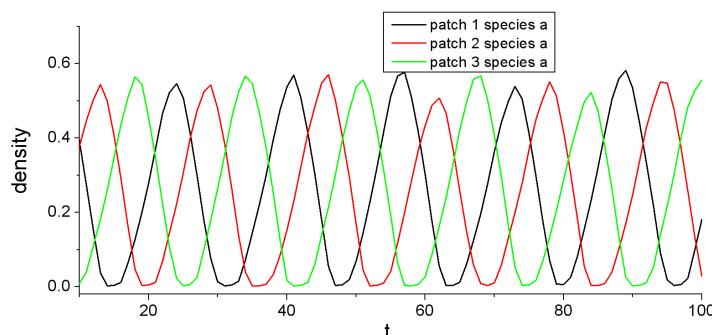
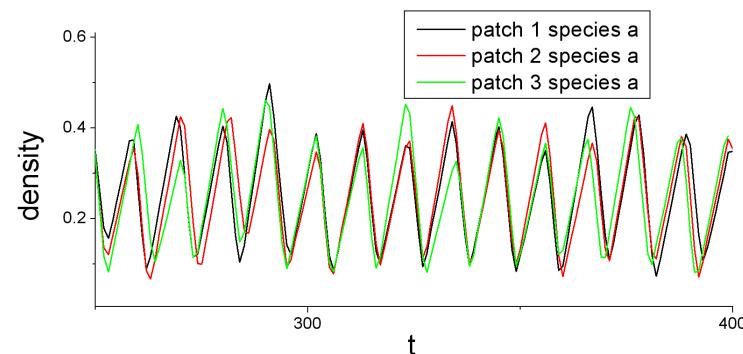
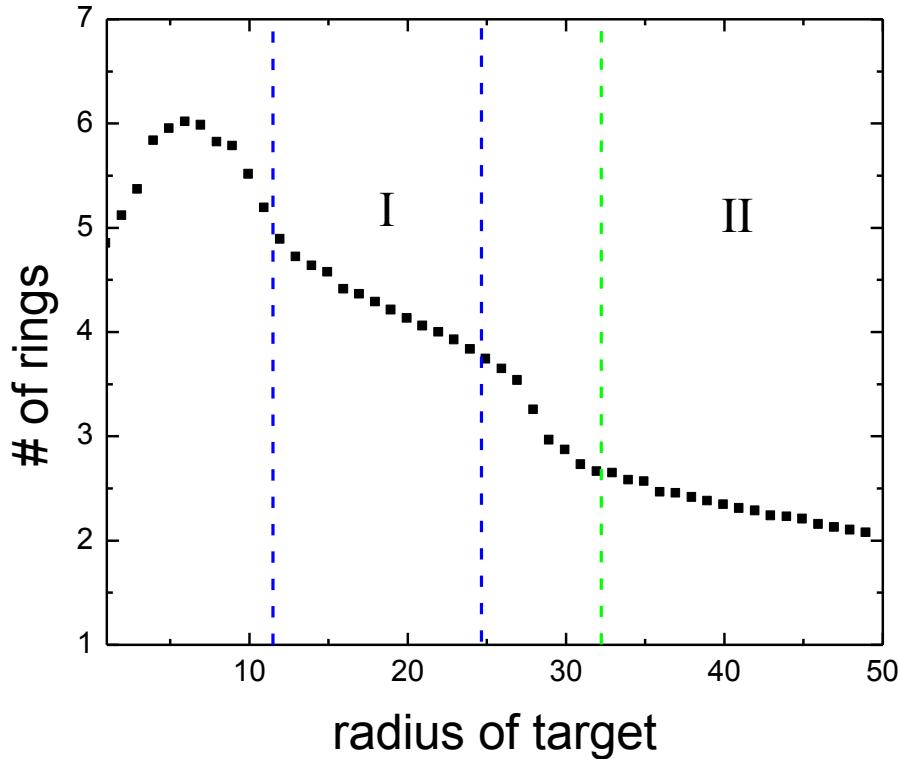
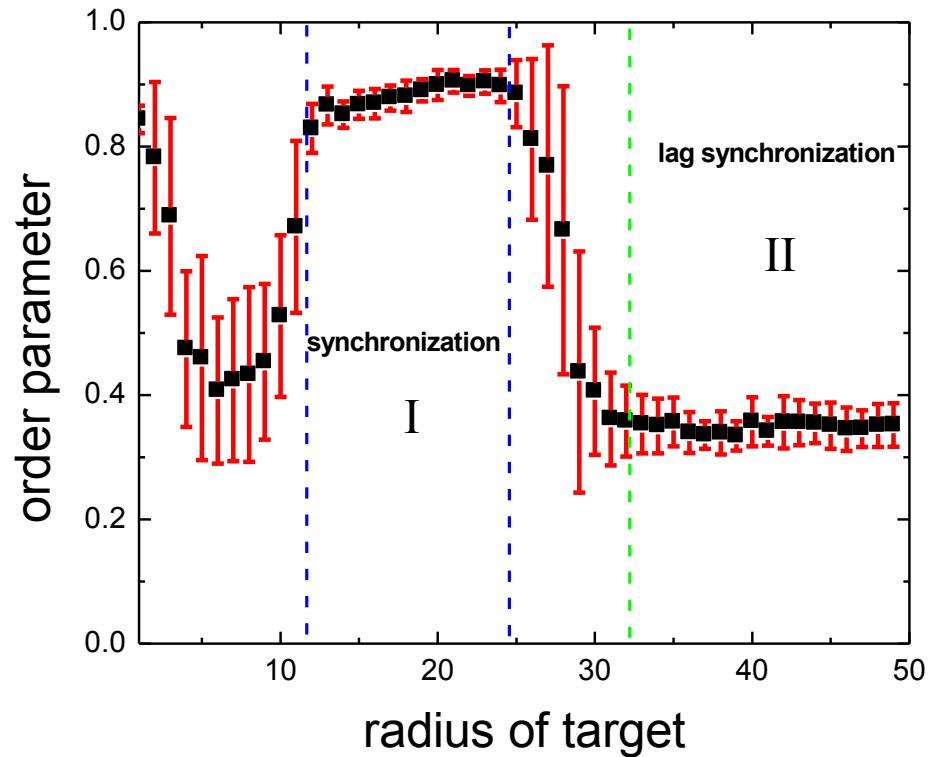
$$\Rightarrow L_r = \left(\frac{T_m L^2}{2\pi} \sqrt{\frac{2\mu\sigma}{3\mu + \sigma} M} \right)^{\frac{1}{3}} \quad \& \quad n_r = \frac{L}{\sqrt{2} L_r}.$$



Transition from Complete to Lag Synchronisation

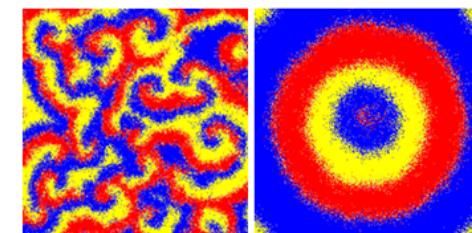
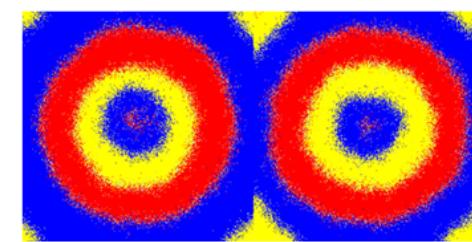
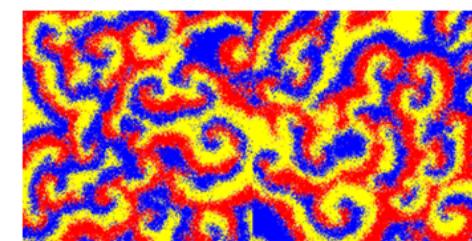
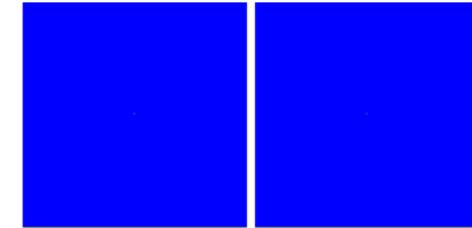
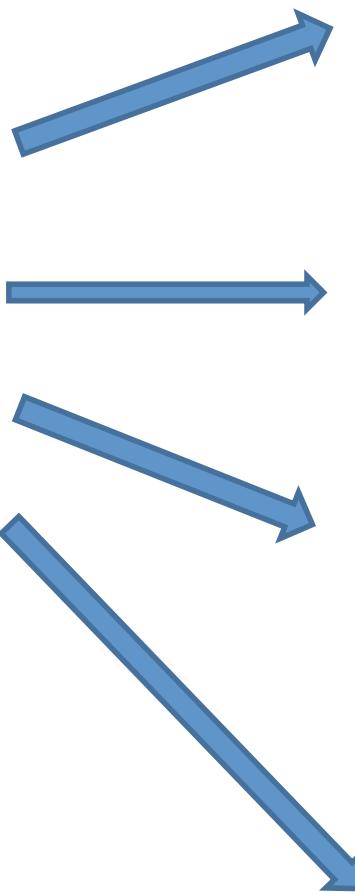
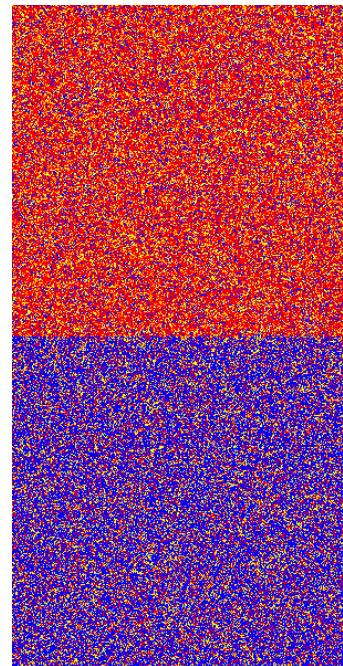


Transition to Lag Synchronisation



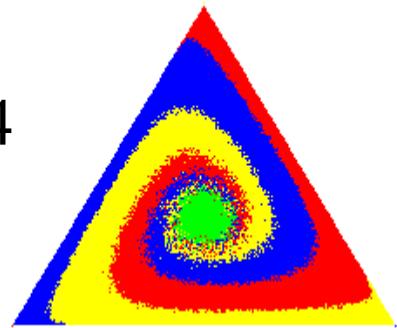
Basins of Extinction and Coexistence

$$\begin{aligned}\rho_1(a) &= \rho_2(b) \\ \rho_1(b) &= \rho_2(c) \\ \rho_1(c) &= \rho_2(a)\end{aligned}$$

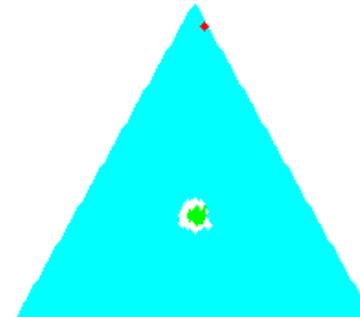
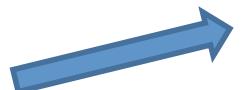


Basins

$M = 1E - 4$



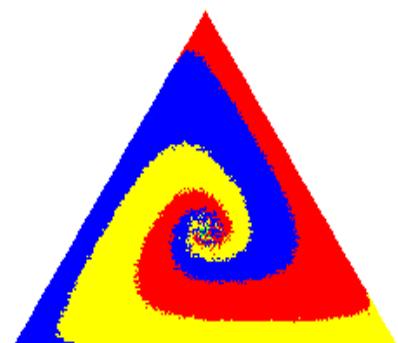
$T_m = 1$



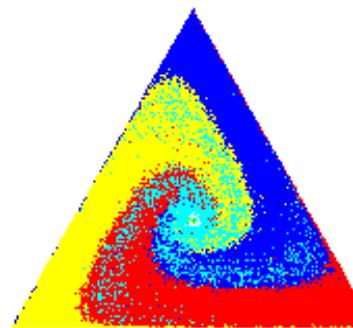
extinction



$M = 3E - 4$



$T_m = 1$



Two spiral waves

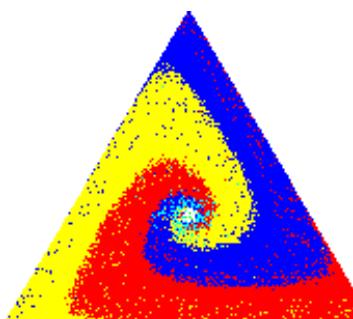


Two target waves



One spiral
One target
wave

$T_m = 15$



Conclusions

- Virus spreading within species can promote coexistence
- Virus spreading across different species tend to hamper coexistence
- Inter-patch migration can induce coexistence in the form of novel target waves
- All these are obtained from **microscopic** model of evolutionary dynamics