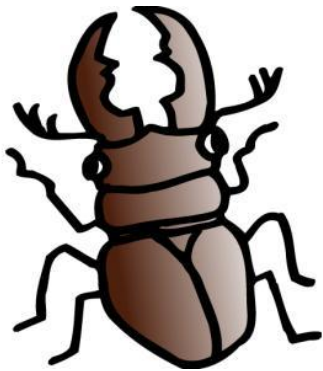
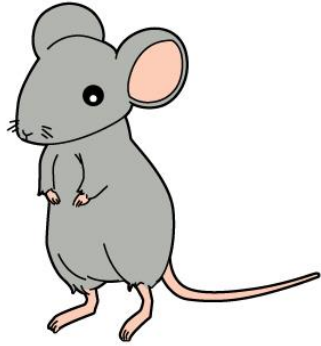
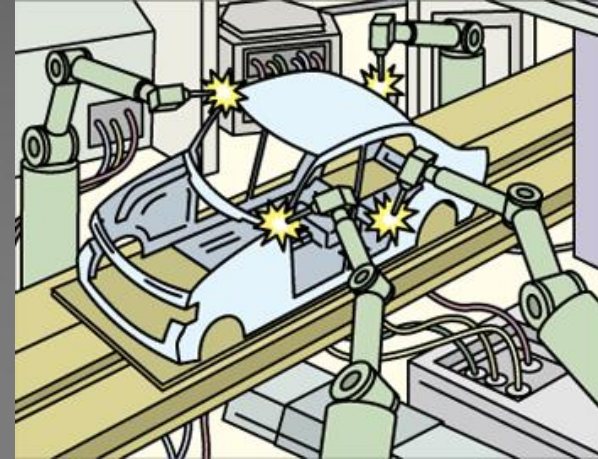
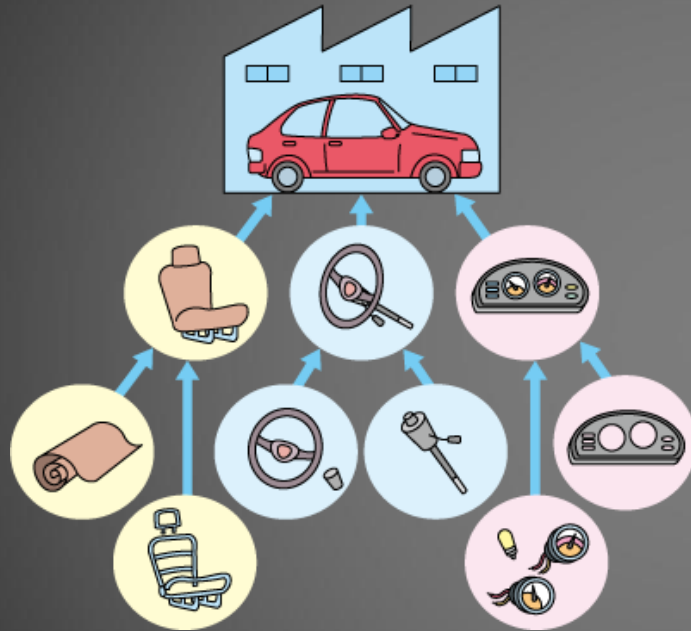


*Introduction of the study
in Pattern Formation Group
I*

*Shigeru Kondo
Osaka University, Japan*



Making a complex machine



We need to make all the parts of the machine.

We need to assemble the parts.

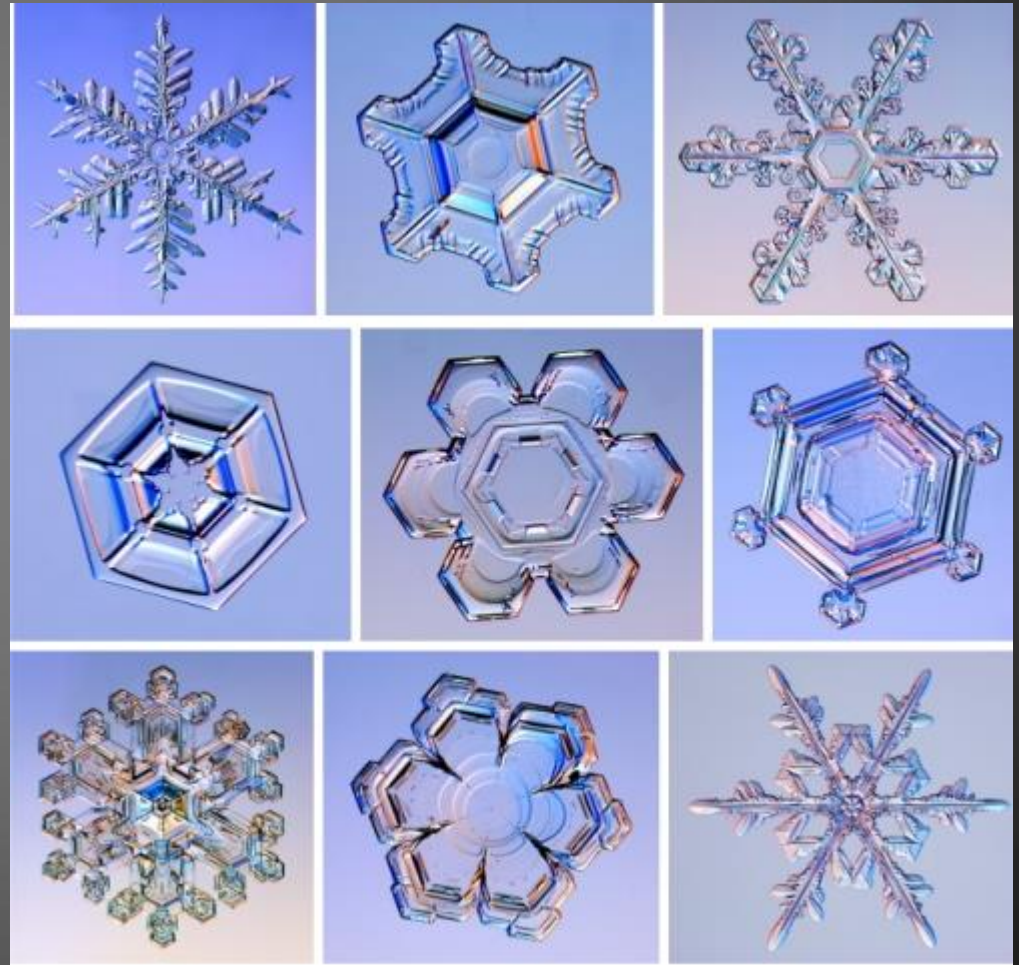
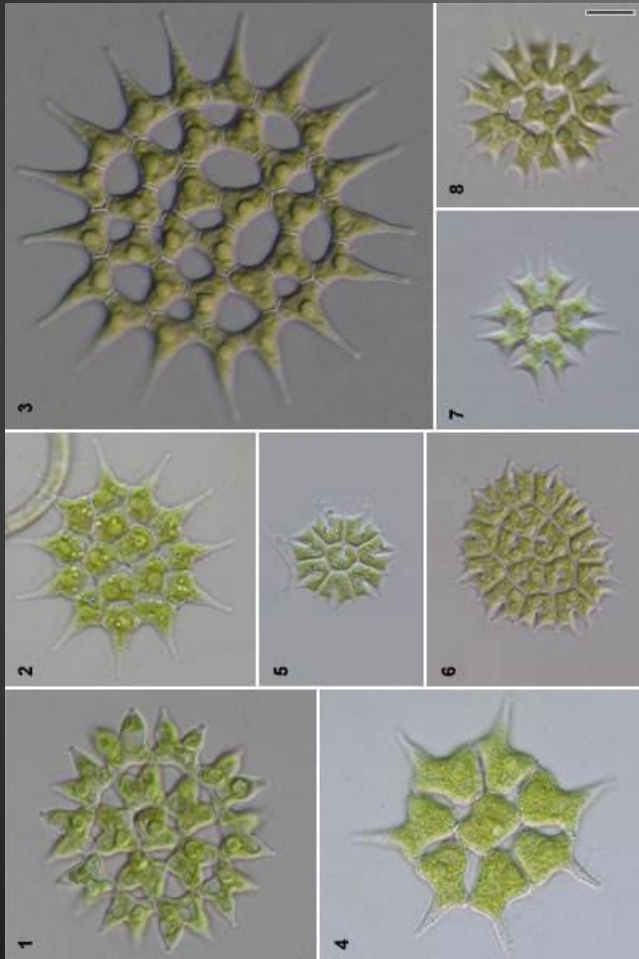
Cars never makes it self. We need to do everything.

All the process is absolutely autonomous!!



©Warren Photographic

Similar patterns develop in non-organic systems





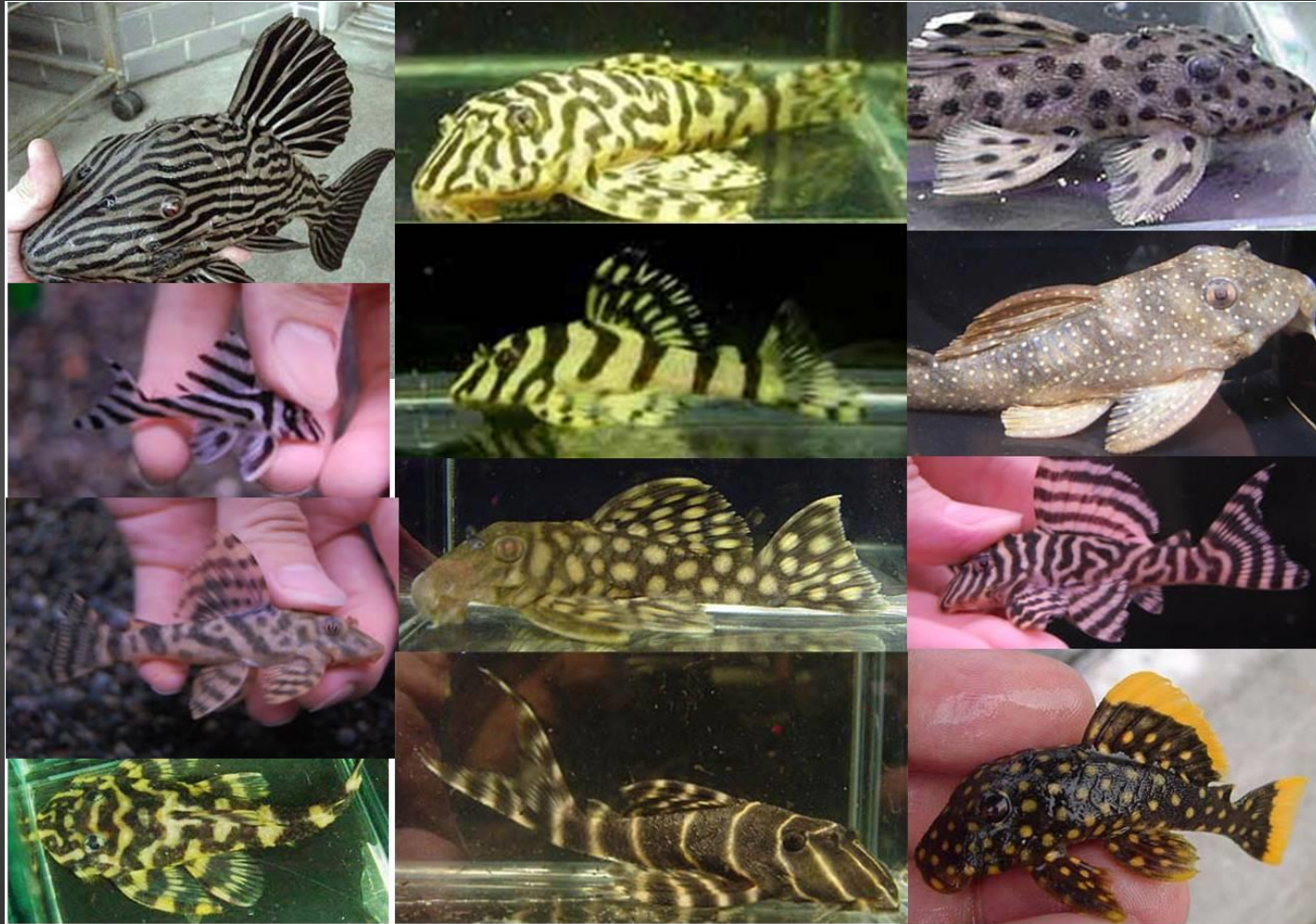
Similar mechanism???



To find out the mechanism autonomous pattern formation, we selected the animal skin markings.



How are the pigmentation patterns made ?



Genetically related species show variety of skin pattern.

>>An identical mechanism must be able to generate different pattern.

How are the pigmentation patterns made ?



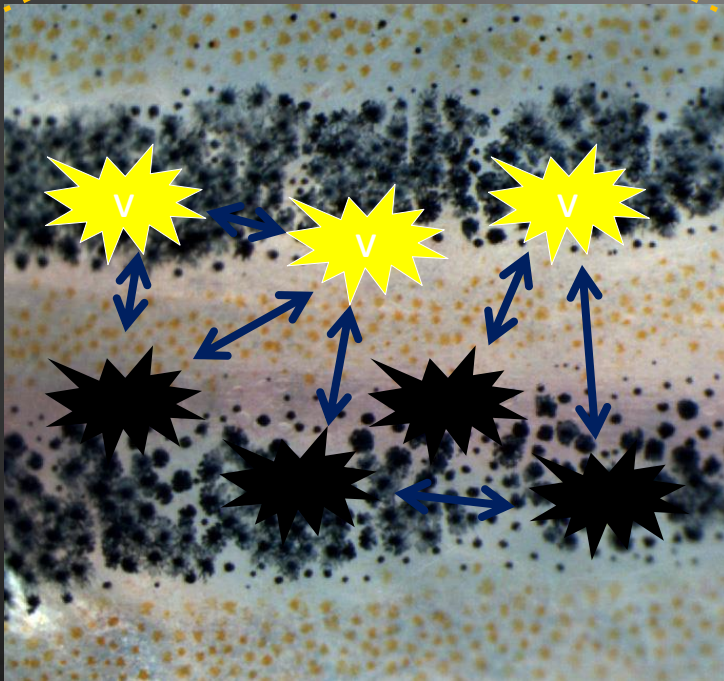
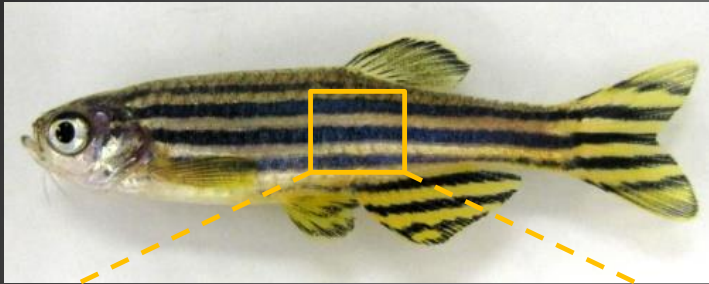
*No similar pattern exist beneath the skin.
>>Patterns are autonomously made in the skin.*

How are the pigmentation patterns made ?



*A common characteristic
>>periodicity*

What is making the skin pattern?



Patterns on the skin is the array of numerous pigment cells.

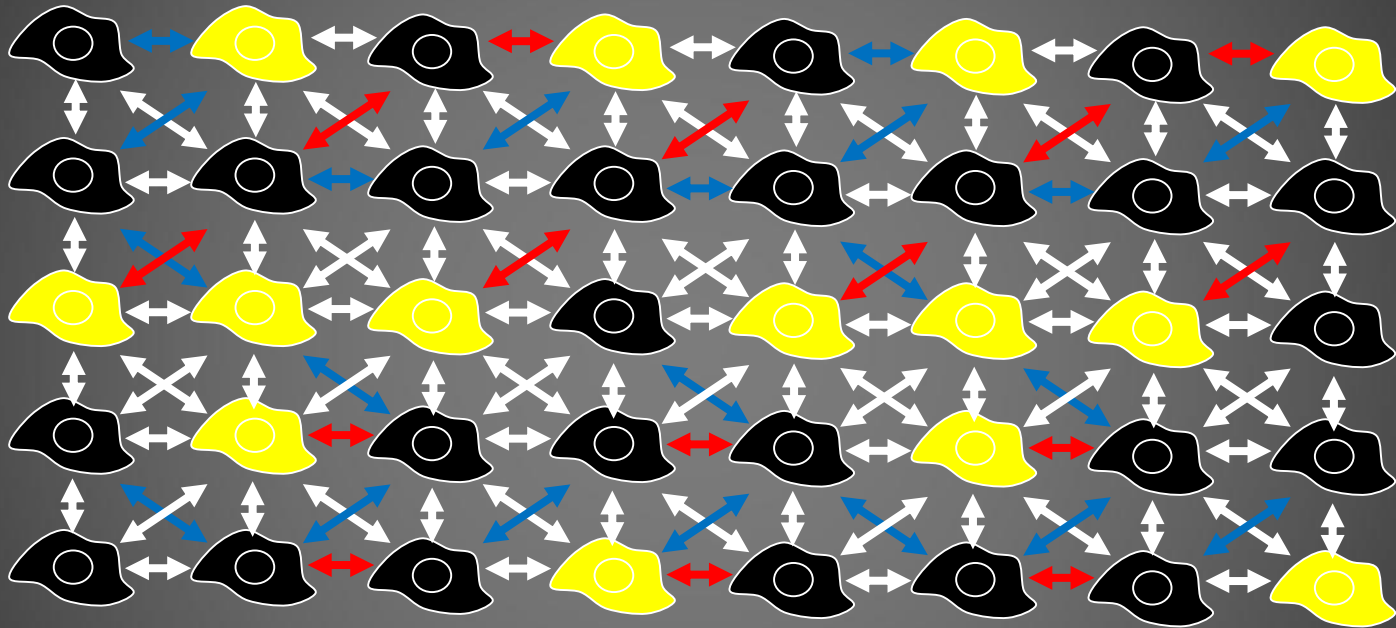
More than two types of pigment cells.

Cells interact each other.

Cells can move, develop and die.



We cannot grasp what really happen with our brain.



Huge number of interactions occur simultaneously.



Need the help of computer and mathematics.

Alan Mathison Turing



Turing presented a simple system in which interactions among the chemical (cells) and the diffusion of the chemicals signal can generate a spatial pattern.

Reaction-diffusion model

Integrating the differential equation

Stationary case. After a sufficient lapse of time $X_r - h$ and $Y_r - k$ approach asymptotically to the forms

$$\left. \begin{aligned} X_r - h &= 2\mathcal{R}A_{s_0} \exp\left[\frac{2\pi i s_0 r}{N} + It\right], \\ Y_r - k &= 2\mathcal{R}C_{s_0} \exp\left[\frac{2\pi i s_0 r}{N} + It\right]. \end{aligned} \right\} \quad (8-1)$$

Oscillatory case. After a sufficient lapse of time $X_r - h$ and $Y_r - k$ approach the forms

$$\left. \begin{aligned} X_r - h &= 2e^{It} \mathcal{R} \left\{ A_{s_0} \exp\left[\frac{2\pi i s_0 r}{N} + i\omega t\right] + A_{N-s_0} \exp\left[-\frac{2\pi i s_0 r}{N} - i\omega t\right] \right\}, \\ Y_r - k &= 2e^{It} \mathcal{R} \left\{ C_{s_0} \exp\left[\frac{2\pi i s_0 r}{N} + i\omega t\right] + C_{N-s_0} \exp\left[-\frac{2\pi i s_0 r}{N} - i\omega t\right] \right\}. \end{aligned} \right\} \quad (8-2)$$

Answer is

After long time has past,,,,

Case A: generates stable pattern

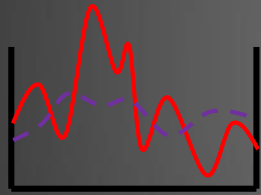
$$\left. \begin{aligned} X_r - h &= 2\Re A_{s_0} \exp\left[\frac{2\pi i s_0 r}{N} + It\right] \\ Y_r - k &= 2\Re C_{s_0} \exp\left[\frac{2\pi i s_0 r}{N} + It\right] \end{aligned} \right\}$$

Case B: generates oscillating pattern.

$$\left. \begin{aligned} X_r - h &= 2e^{\lambda t} \Re \left\{ A_{s_0} \exp\left[\frac{2\pi i s_0 r}{N} + i\alpha t\right] + A_{N-s_0} \exp\left[-\frac{2\pi i s_0 r}{N} - i\alpha t\right] \right\} \\ Y_r - k &= 2e^{\lambda t} \Re \left\{ C_{s_0} \exp\left[\frac{2\pi i s_0 r}{N} + i\alpha t\right] + C_{N-s_0} \exp\left[-\frac{2\pi i s_0 r}{N} - i\alpha t\right] \right\} \end{aligned} \right\}$$

Six different states are possible

Initial condition



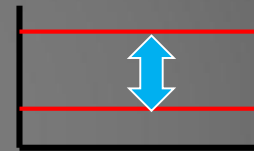
Both morphogenesis diffuse and react each other



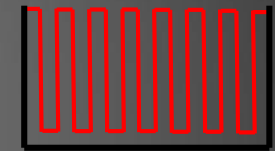
Six possible states



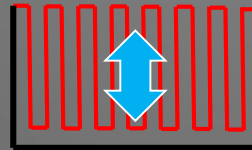
uniform, stationary



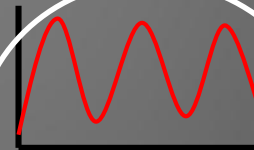
uniform, oscillating



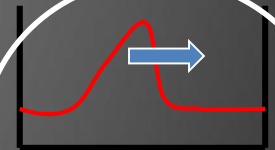
stationary waves with extreme short wave-length



Oscillatory cases with extreme short wave-length



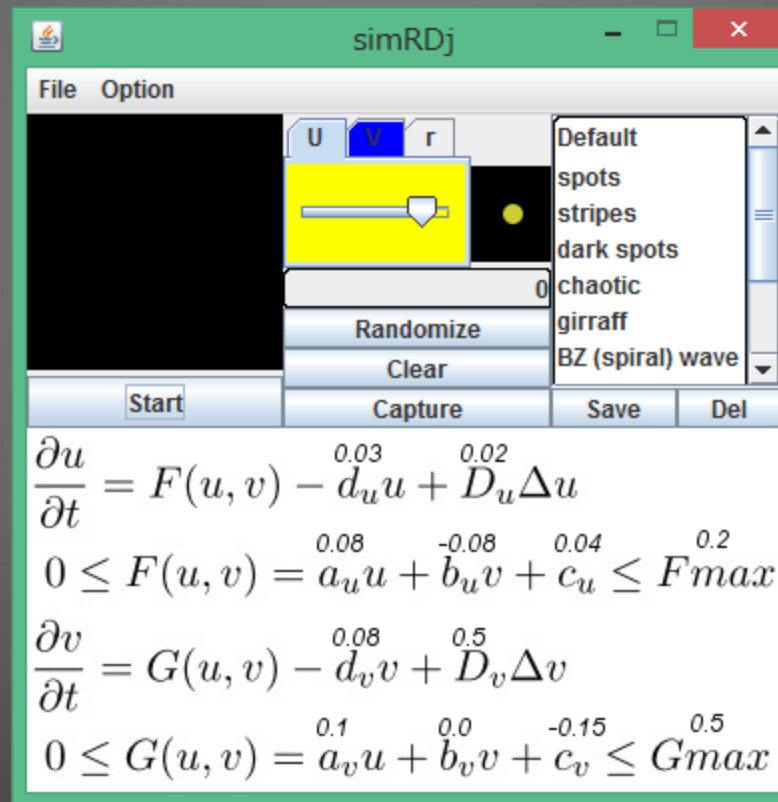
stationary waves with finite wave-length
(Turing pattern)



Oscillatory cases with finite wave-length

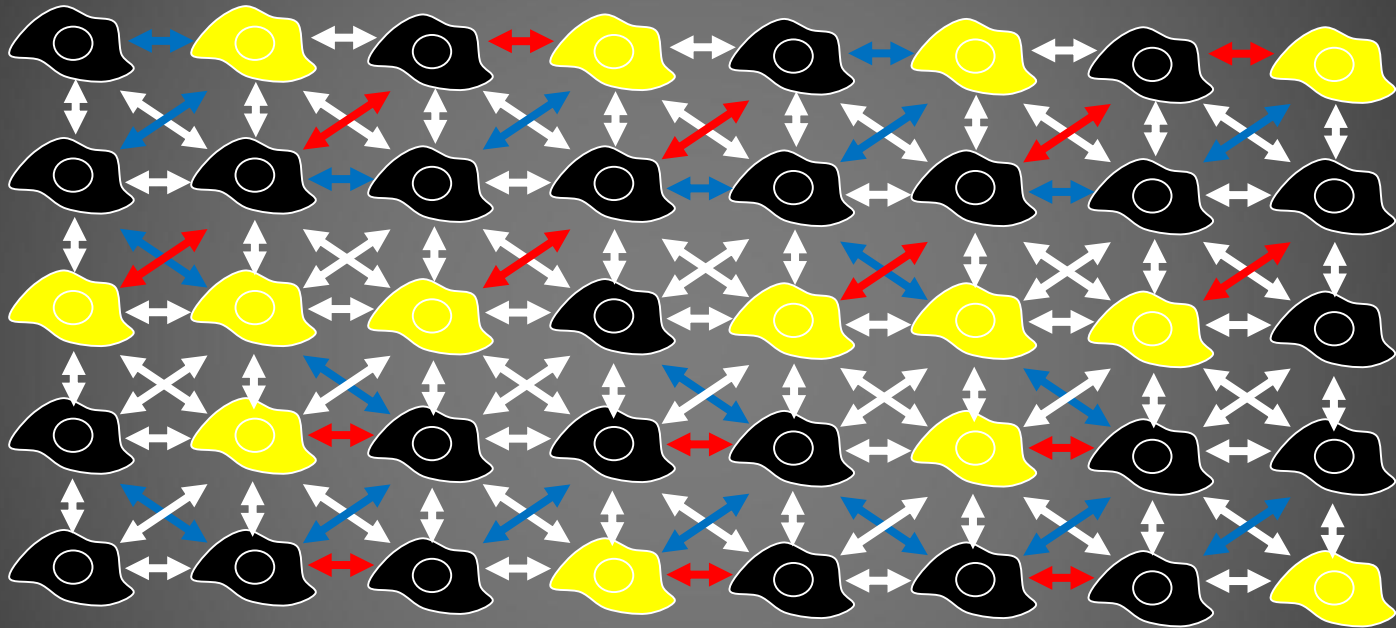
Mathematics is difficult,,,,,

- Testing the system with simulation is easy!!



The screenshot shows the 'simRDj' software window. The interface includes a menu bar with 'File' and 'Option', a main display area (currently black), and a control panel with tabs for 'U', 'V', and 'r'. The 'V' tab is active, showing a yellow slider and a small yellow circle. Below the control panel are buttons for 'Start', 'Capture', 'Save', and 'Del'. A list of simulation patterns is visible on the right: Default, spots, stripes, dark spots, chaotic, girraff, and BZ (spiral) wave. At the bottom, the following mathematical equations are displayed:

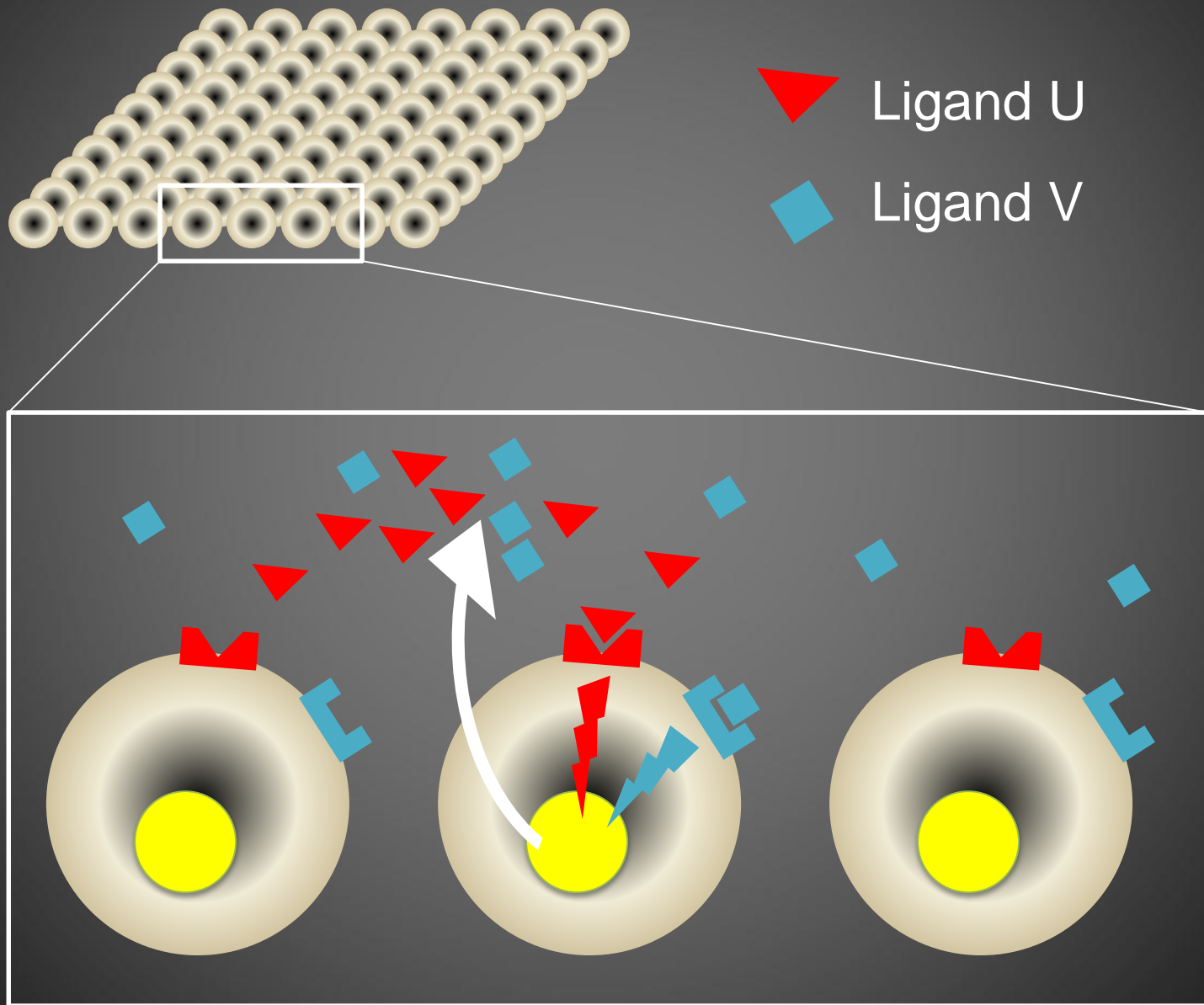
$$\frac{\partial u}{\partial t} = F(u, v) - d_u u + D_u \Delta u$$
$$0 \leq F(u, v) = a_u u + b_u v + c_u \leq Fmax$$
$$\frac{\partial v}{\partial t} = G(u, v) - d_v v + D_v \Delta v$$
$$0 \leq G(u, v) = a_v u + b_v v + c_v \leq Gmax$$



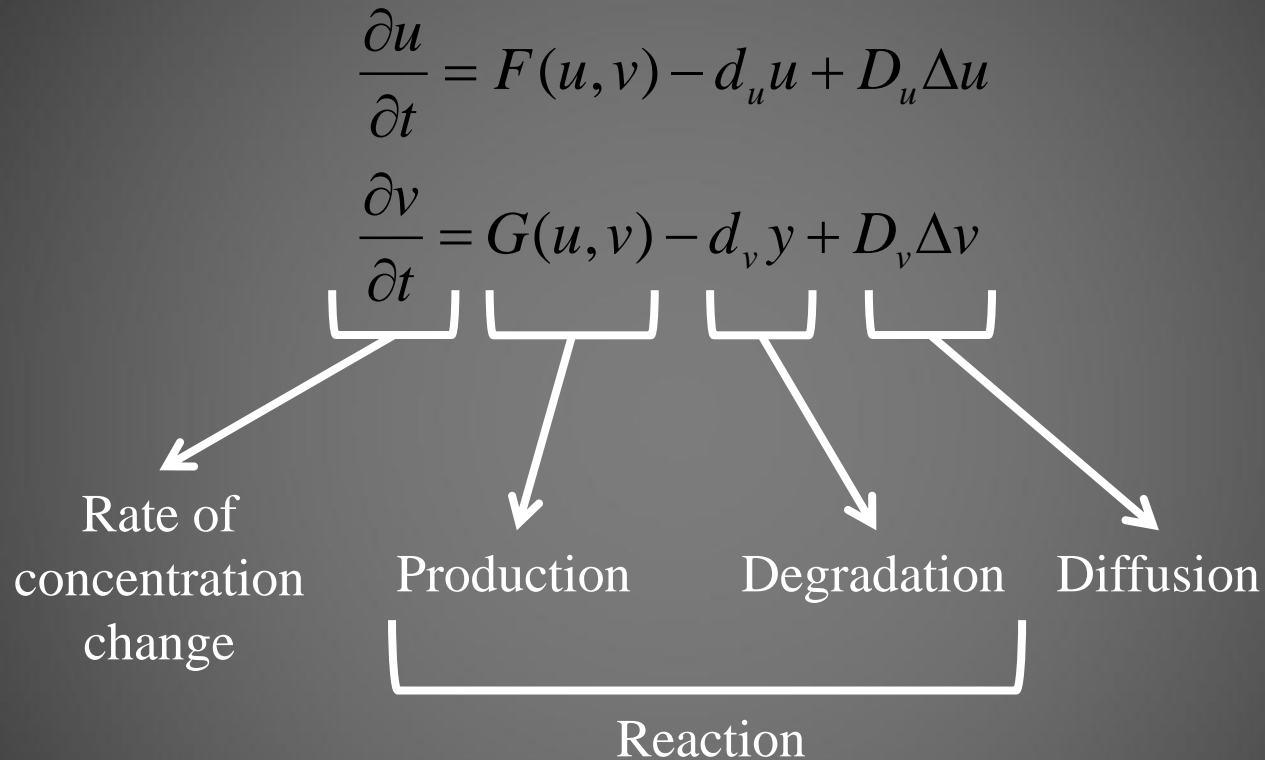
Huge number of interactions occur simultaneously.



Need the help of computer and mathematics.



数式で表すと...



$$F(u, v) = a_1 u + b_1 v + c_1$$

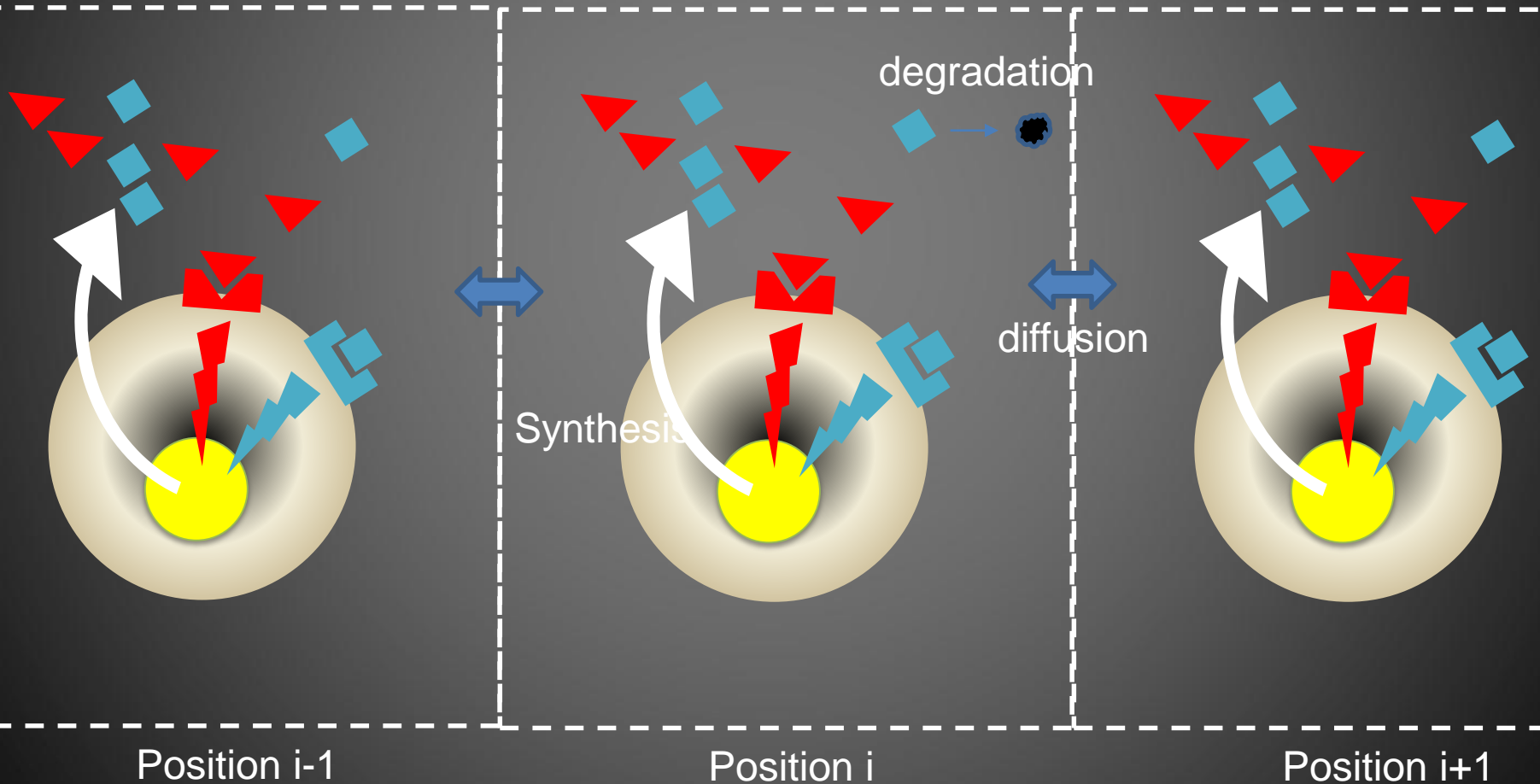
$$G(u, v) = a_2 u + b_2 v + c_2$$

$U_i(t_0)$ = Concentration of U in position i at time t

$V_i(t_0)$ = Concentration of V in position i at time t

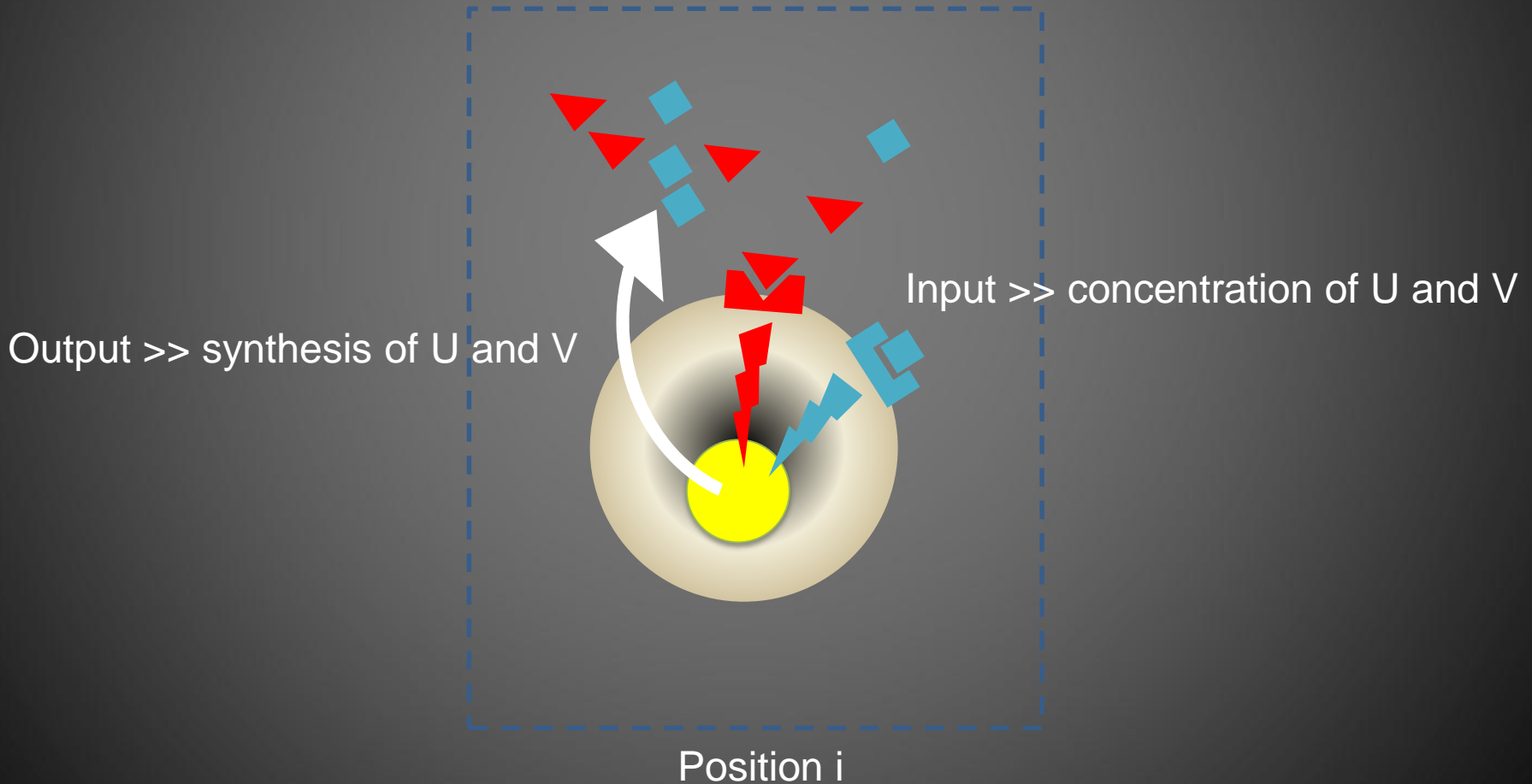
If we know $U_i(t+dt)$ and $V_i(t+dt)$, we can predict how the system behave in the future

What we need to calculate is Synthesis, degradation and diffusion.



synthesis

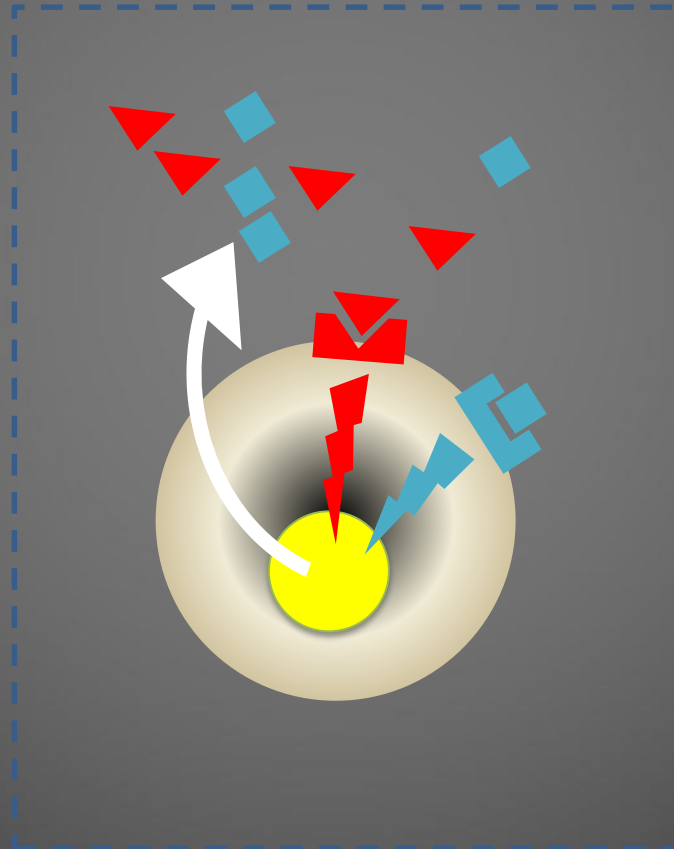
$$\begin{aligned} \text{synthesis}U &= F(u, v) = a_u u + b_u v + c_u \\ \text{synthesis}V &= G(u, v) = a_v u + b_v v + c_v \end{aligned}$$



degradation

$$\text{degradation}U = -d_u * u$$
$$\text{degradation}V = -d_v * v$$

d_u d_v are the
degradation
constant of u and v

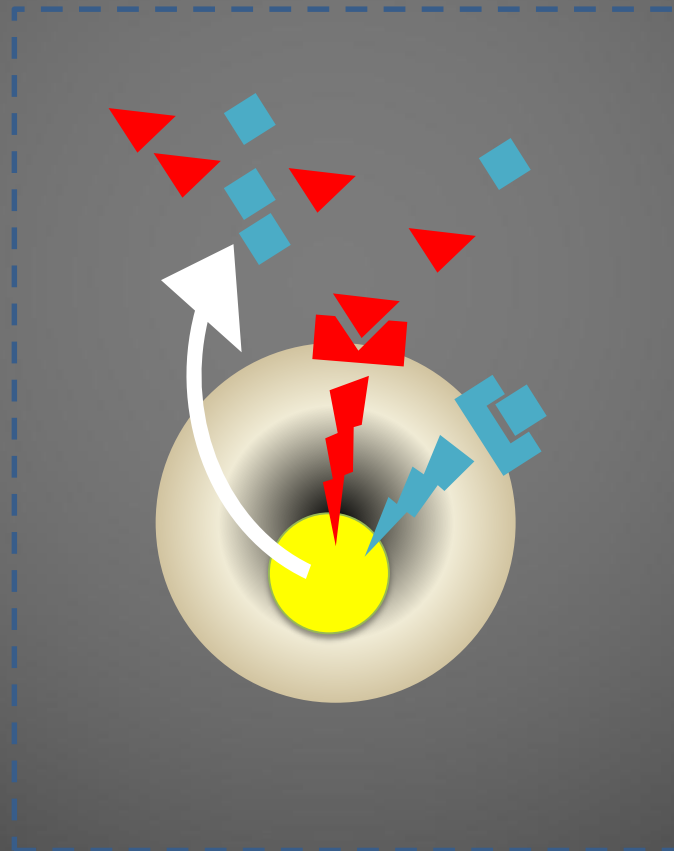


Position i

Reaction=synthesis + degradation

$$\text{reaction}U = \text{synthesis}U + \text{degradation}U = F(u, v) - d_u * u$$

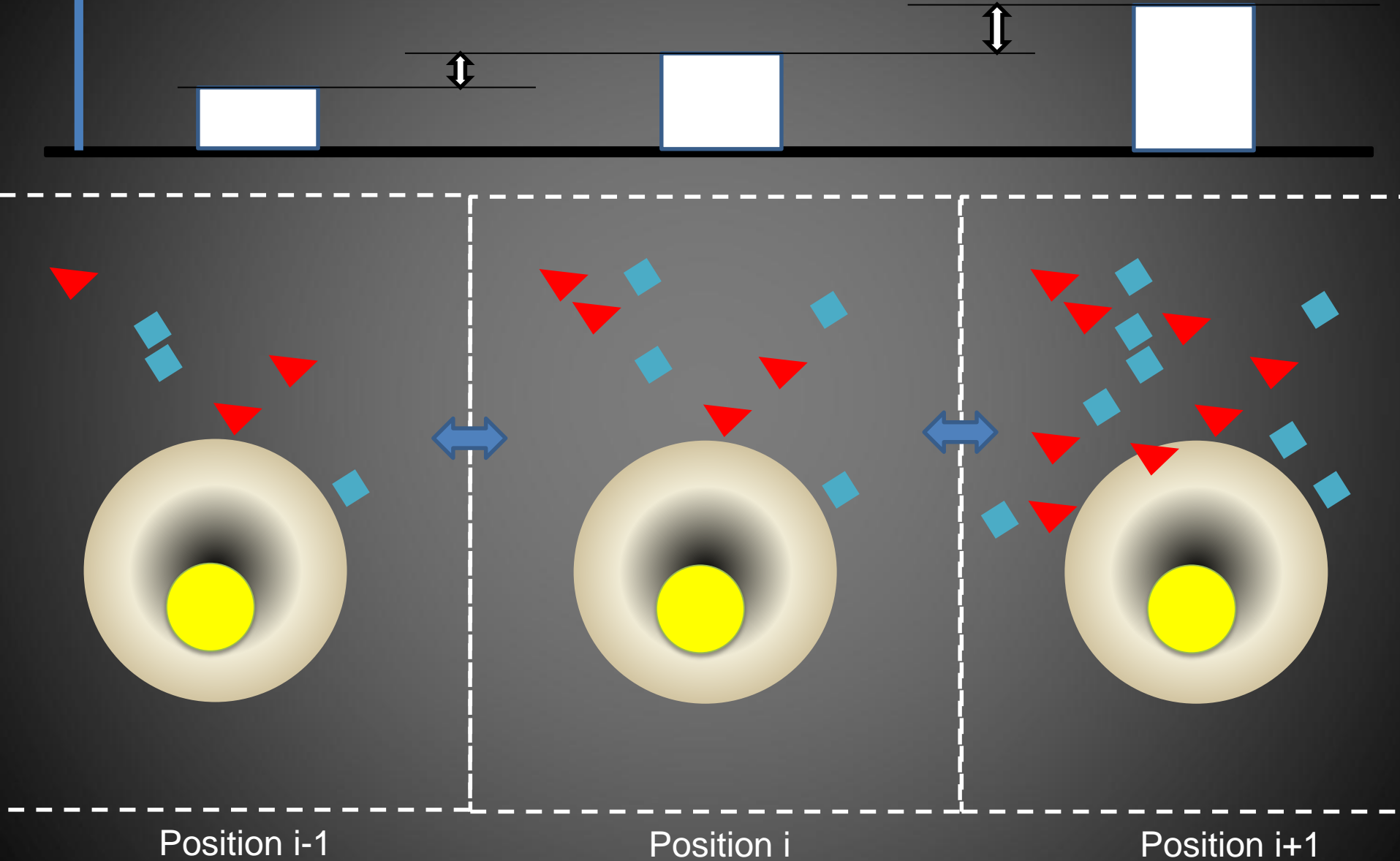
$$\text{reaction}V = \text{synthesis}V + \text{degradation}V = G(u, v) - d_v * v$$

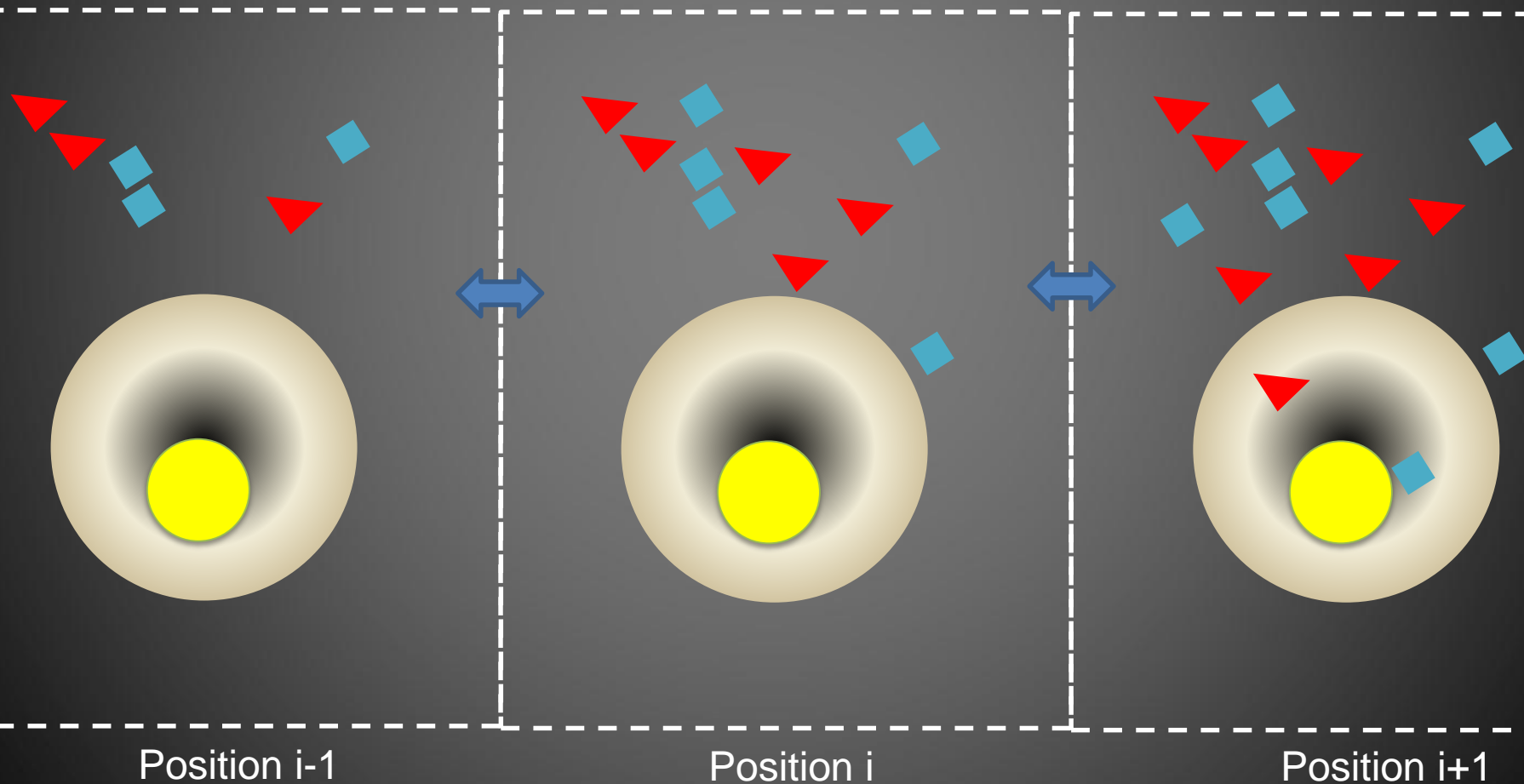
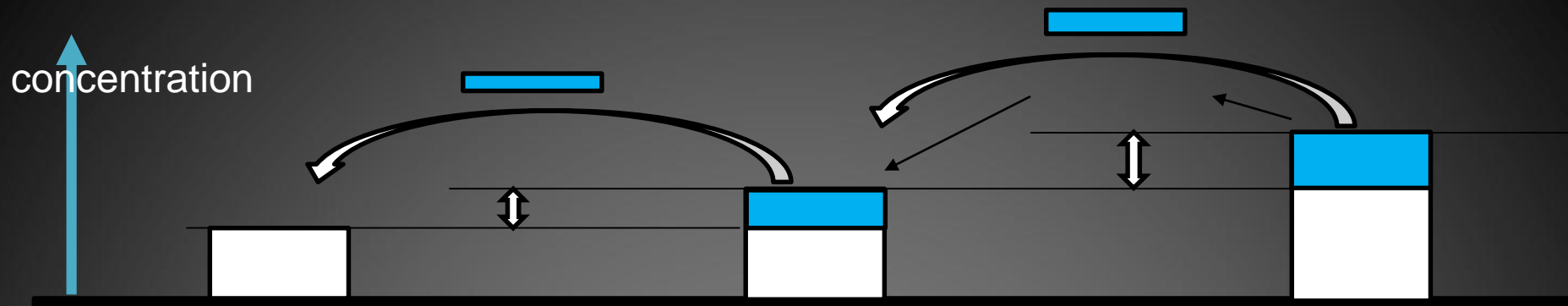


Position i

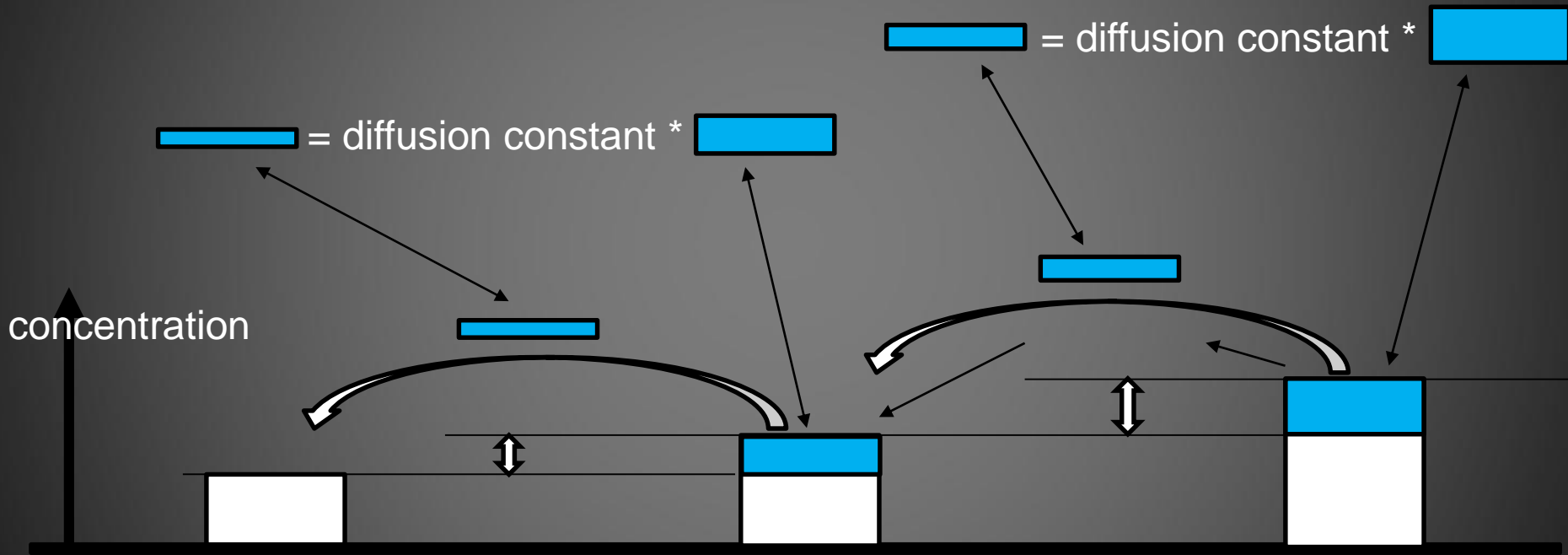
concentration

diffusion





diffusion



RD simulator

$$\frac{\partial u}{\partial t} = F(u, v) - d_u u + D_u \Delta u$$

$$0 \leq F(u, v) = a_u u + b_u v + c_u \leq Fmax$$

$$\frac{\partial v}{\partial t} = G(u, v) - d_v v + D_v \Delta v$$

$$0 \leq G(u, v) = a_v u + b_v v + c_v \leq Gmax$$

Synthesis of u

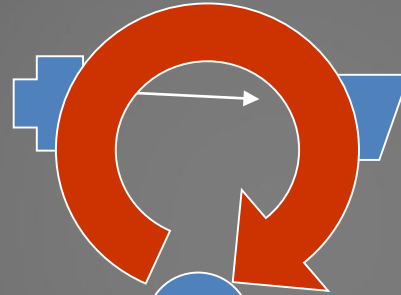
Synthesis of v

Diffusion and degradation

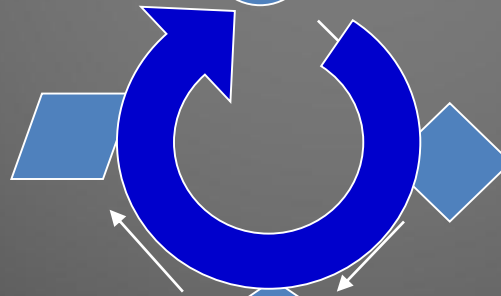
Diffusion and degradation

Conditions to make stable pattern

*Positive
feedback*



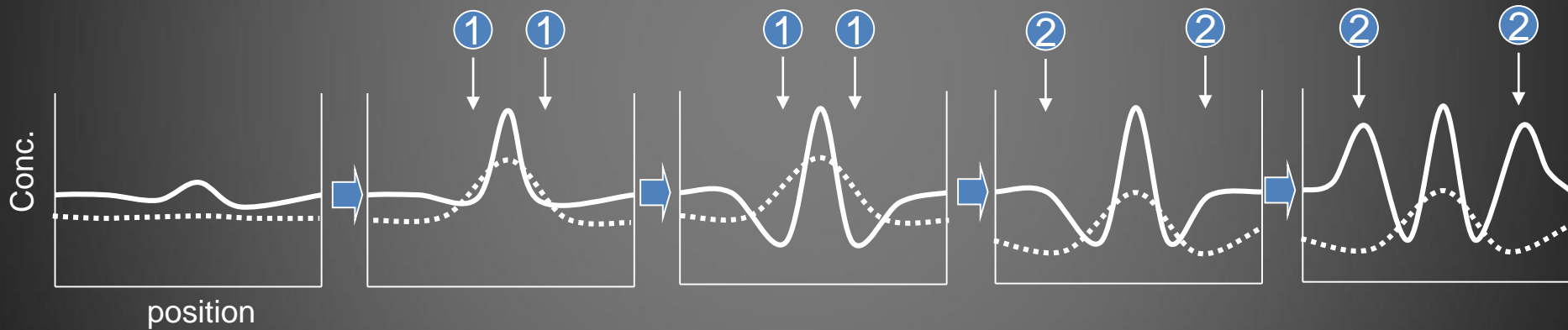
*Move slow
(local)*



*Move fast
(long range)*

*Negative
feedback*

Making a stable wave



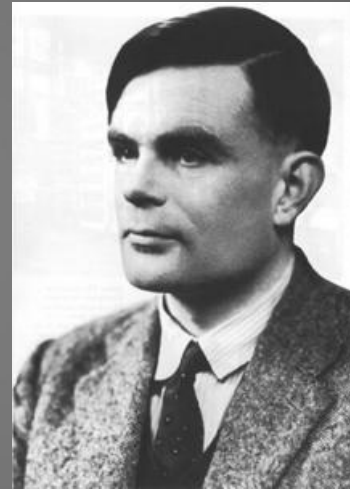
*Introduction of the study
in Pattern Formation Group
II*

*Shigeru Kondo
Osaka University, Japan*

Reaction-Diffusion Model by Alan Turing

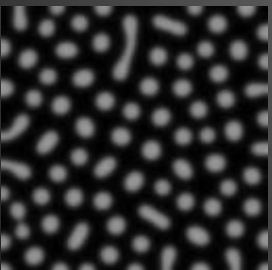
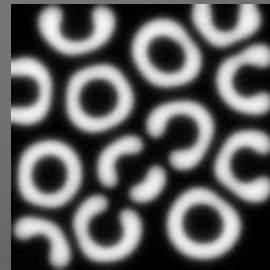
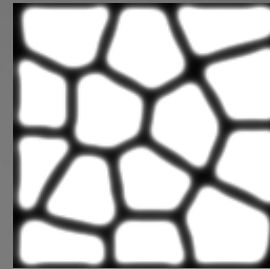
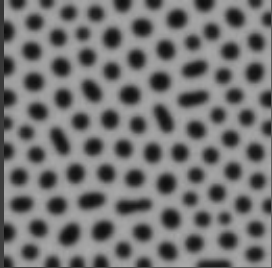
$$\frac{\partial x}{\partial t} = F(x, y) + D_x \Delta x$$

$$\frac{\partial y}{\partial t} = G(x, y) + D_y \Delta y$$

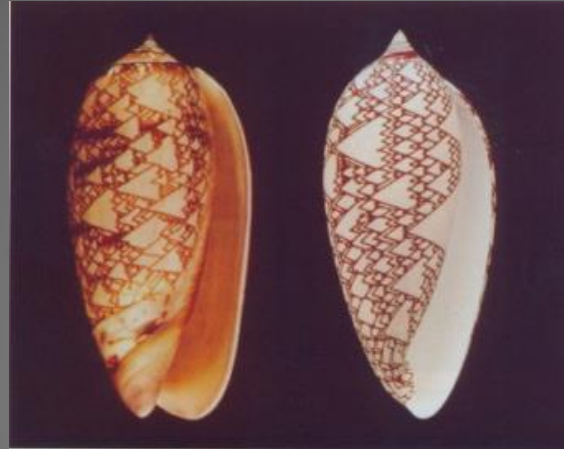


Alan Turing

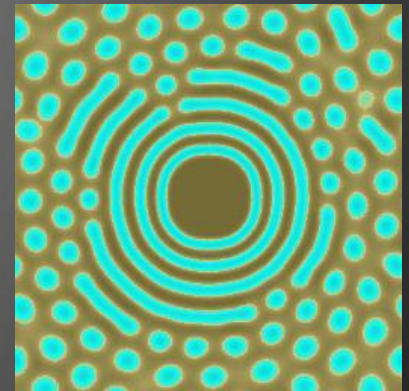
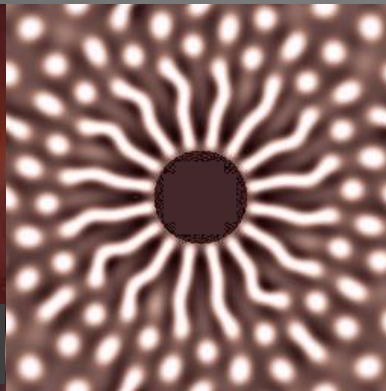
RD model reproduces animal skin patterns



More complex patterns

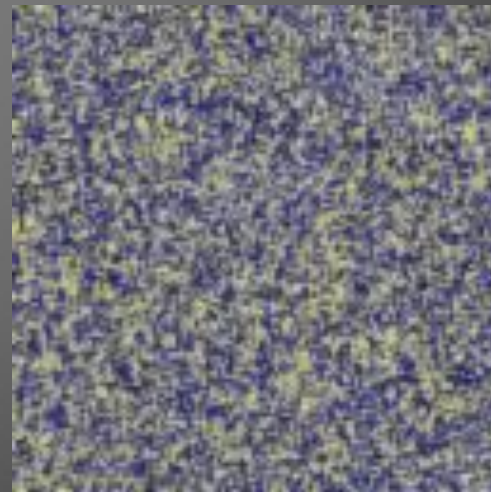
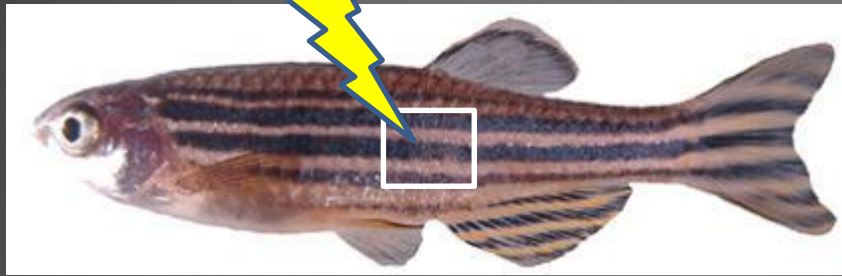


Sandersen 2006



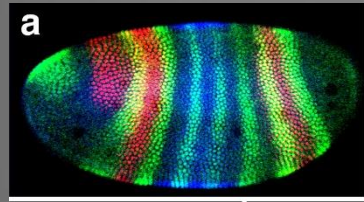
RD model reproduce the pattern regeneration

Laser ablation

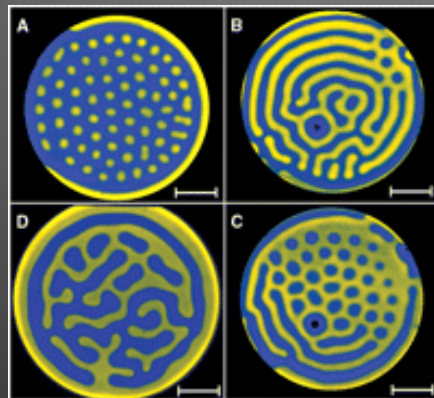


History of Turing model

- 1952 Turing presented RD model
- 1954 Turing died
- 1970 re-discovery of RD model by mathematicians
- 1972~1989 some discussions among developmental biologists



- 1990 most of biologists discarded the RD model
- 1991 generation of RD pattern in chemical system



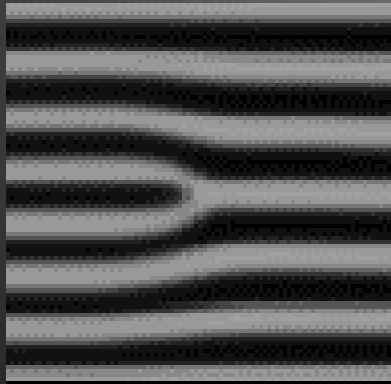
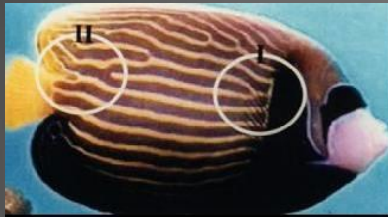
How one can prove the RD theory

- Normal strategy is too complex
 - Get mutant with disturbed pattern.
 - Identify all the molecular and cellular events related to the pattern formation
 - Measure the reaction parameters and the diffusion constants
 - Simulate the interaction network identified
- Needs more easy strategy

Easy method to prove the RD model

- Turing models says
 - Pattern is formed as a kind of “wave”.
 - Pattern regenerate when disturbed.
 - If the pattern is disturbed, it should regenerate in the way the simulation predict.

Prediction of the skin pattern change of Emperor angelfish



simulation

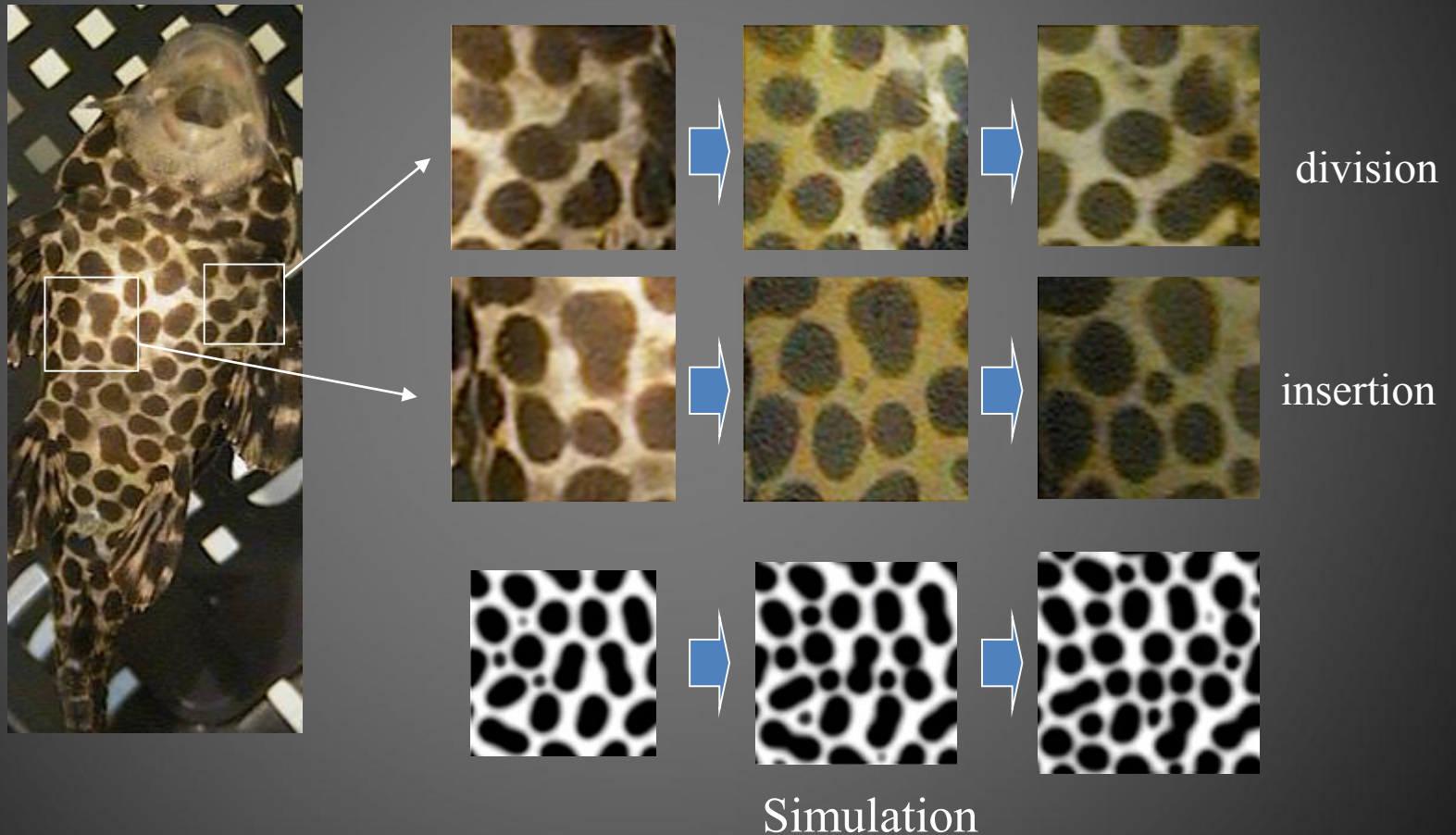


day 0

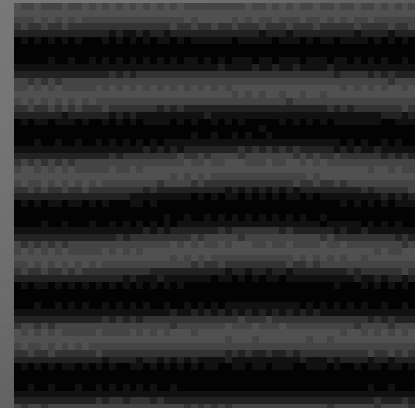
day 40

day 90

*Prediction of the change of spot pattern in a cat fish, *Plecostoms*.*



*In case of Zebrafish with artificial disturbance
of the pattern,,,,,*

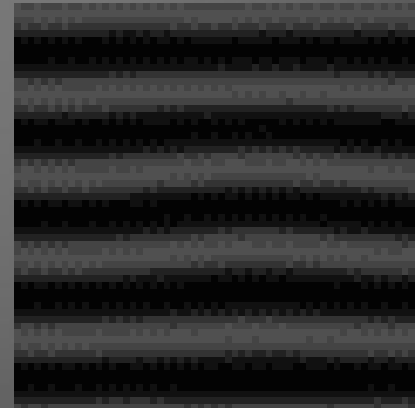


simulation

*Simulation successfully predicted the
pattern change!*



experiment

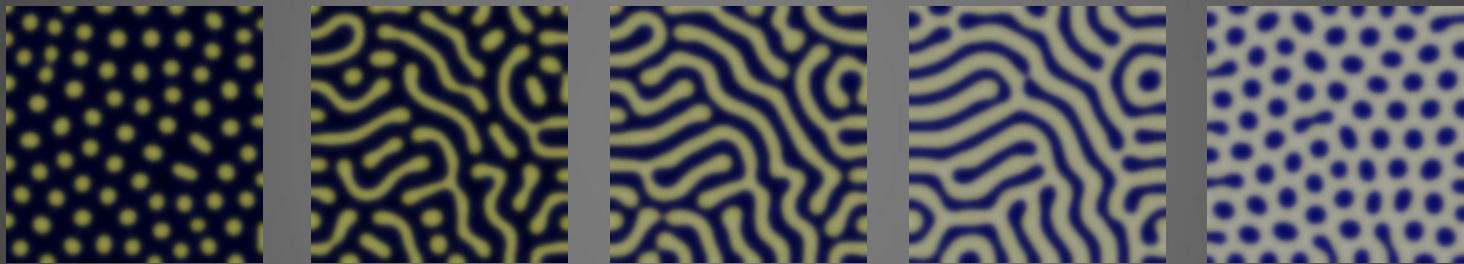


simulation

Prediction of the skin pattern of hybrid fish

$$\frac{\partial x}{\partial t} = F(x, y) + D_x \Delta x$$

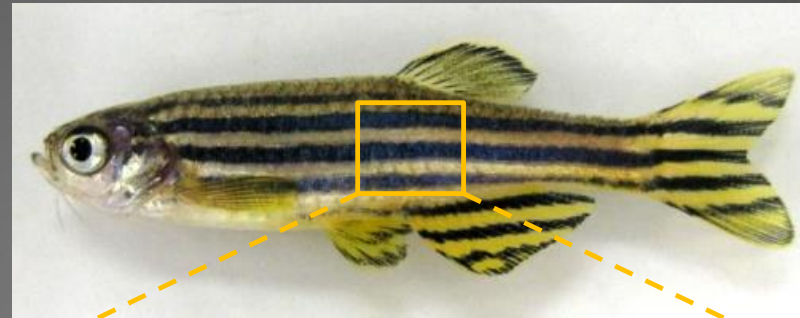
$$\frac{\partial y}{\partial t} = G(x, y) + D_y \Delta y$$



← Change of parameter value →

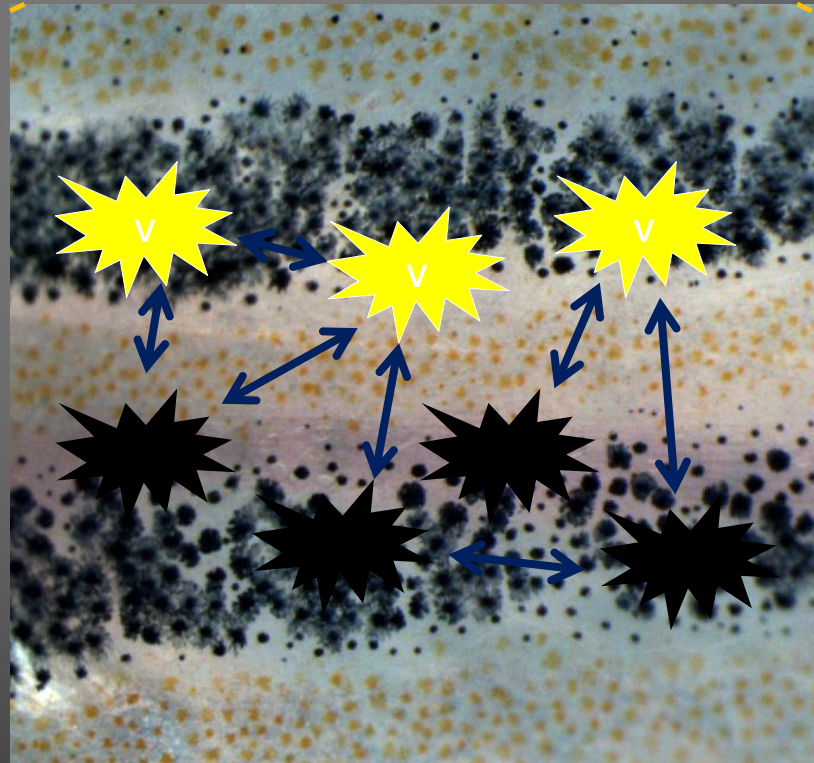


Mutual interaction between the pigment cells should be the key factor

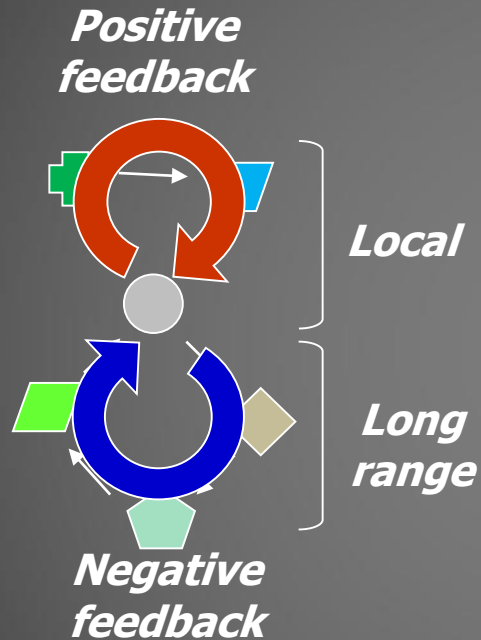


melanophore

xanthophore



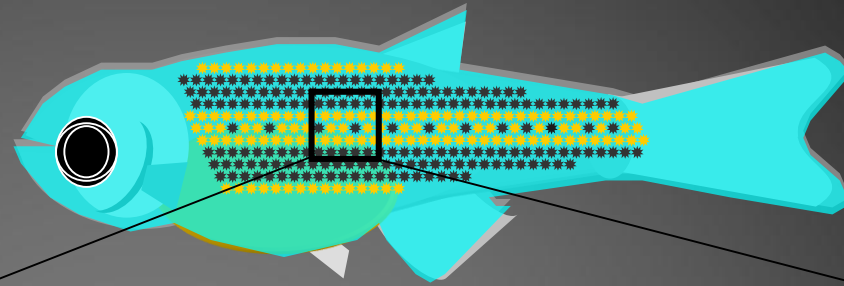
What does the mathematical theory suggest ?



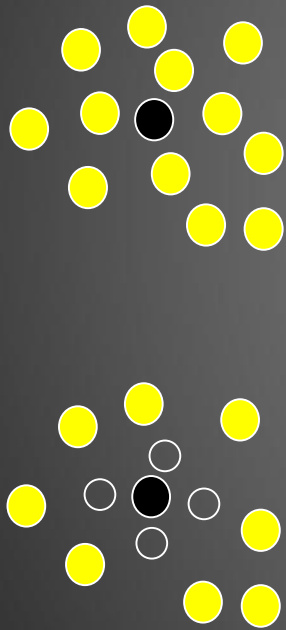
1. There should be two kinds of interactions between the pigment cells.
2. Functional distance of the interactions should be different.

local positive feedback
+
long negative feedback

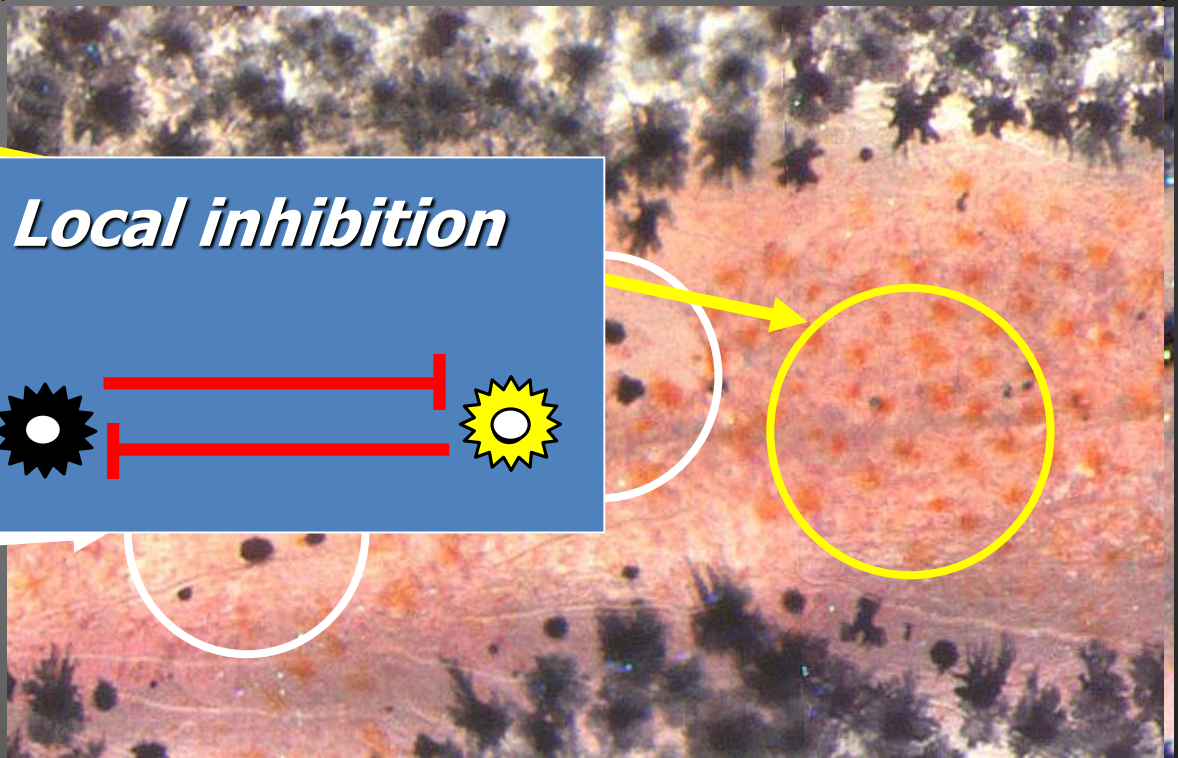
Local interaction between the pigment cells



M is surrounded by Xs

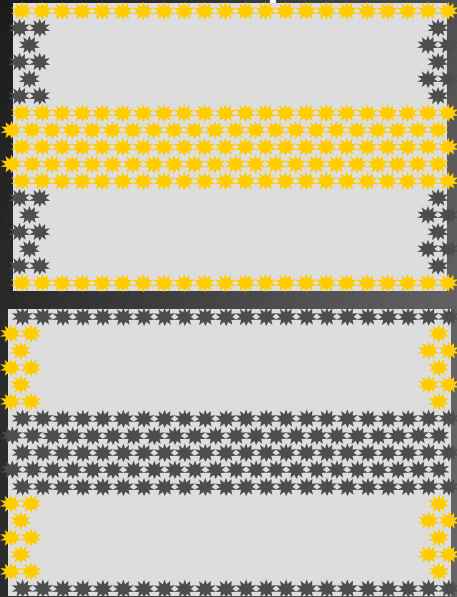


Nearest Xs are killed by laser



Long range interaction between the pigment cells

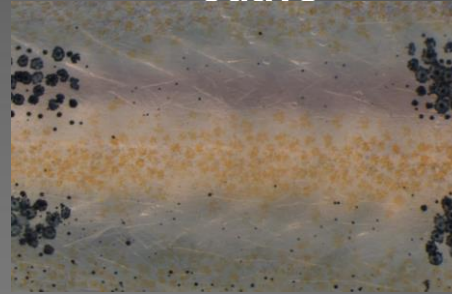
ablation pattern



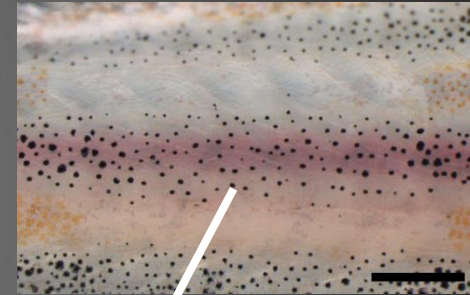
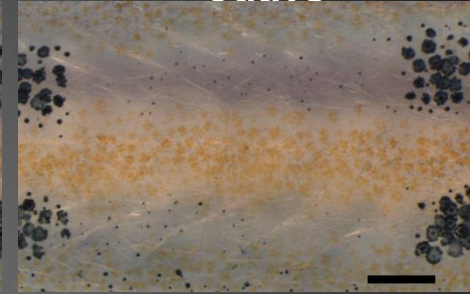
0days



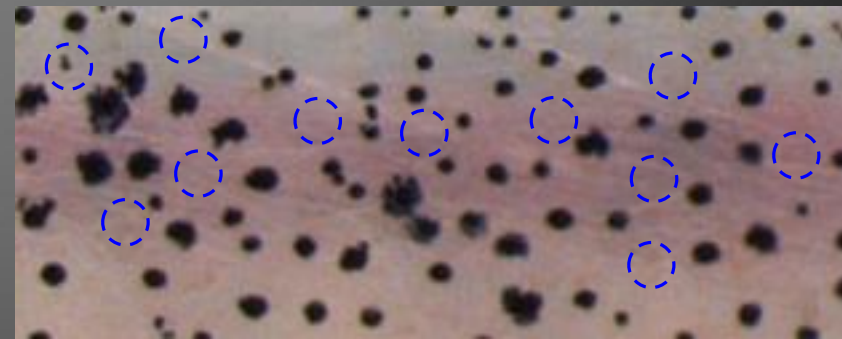
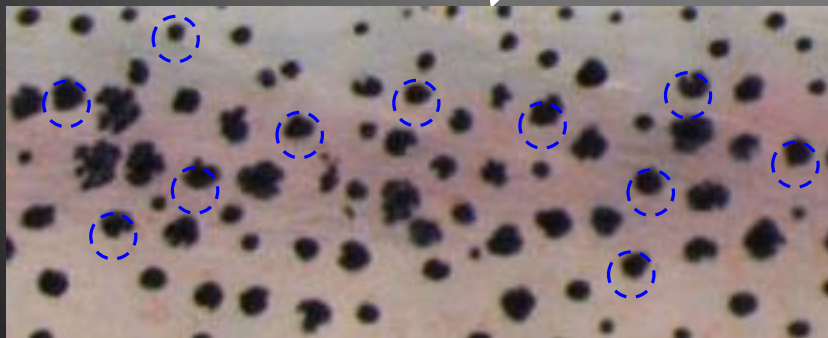
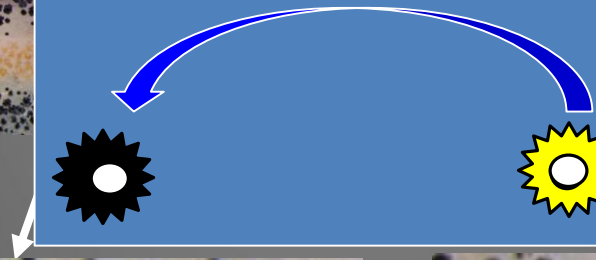
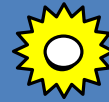
3days



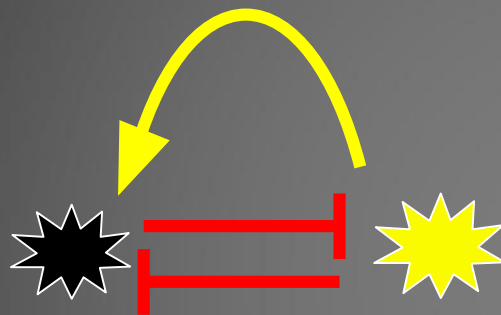
5days



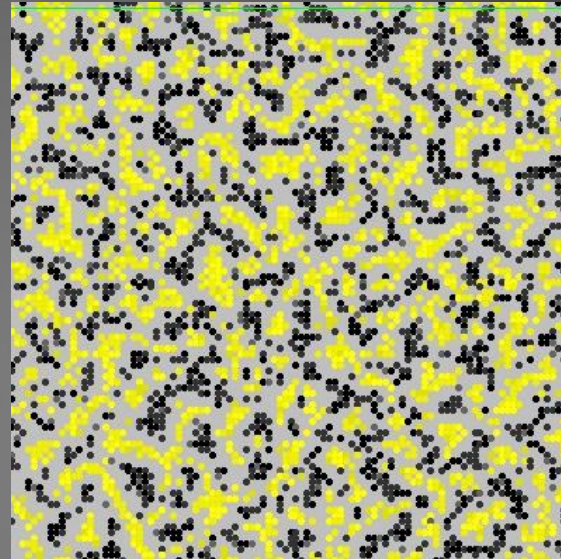
Long range help



Simulation of deduced network forms the stripes



- Activation
- ⊥ Inhibition
- Long range
- Short range



Next to do is finding the molecular and cellular basis of these three interactions

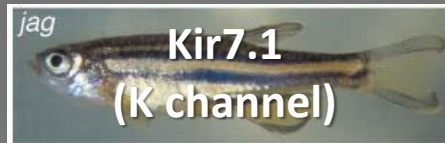
Identifying the molecular mechanism

1) Cloning of some responsible genes

+/+



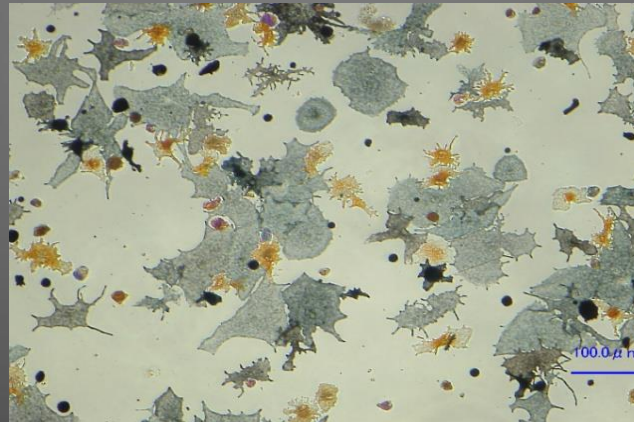
jaguar



leopard



2) Observation of in vitro interaction of pigment cells

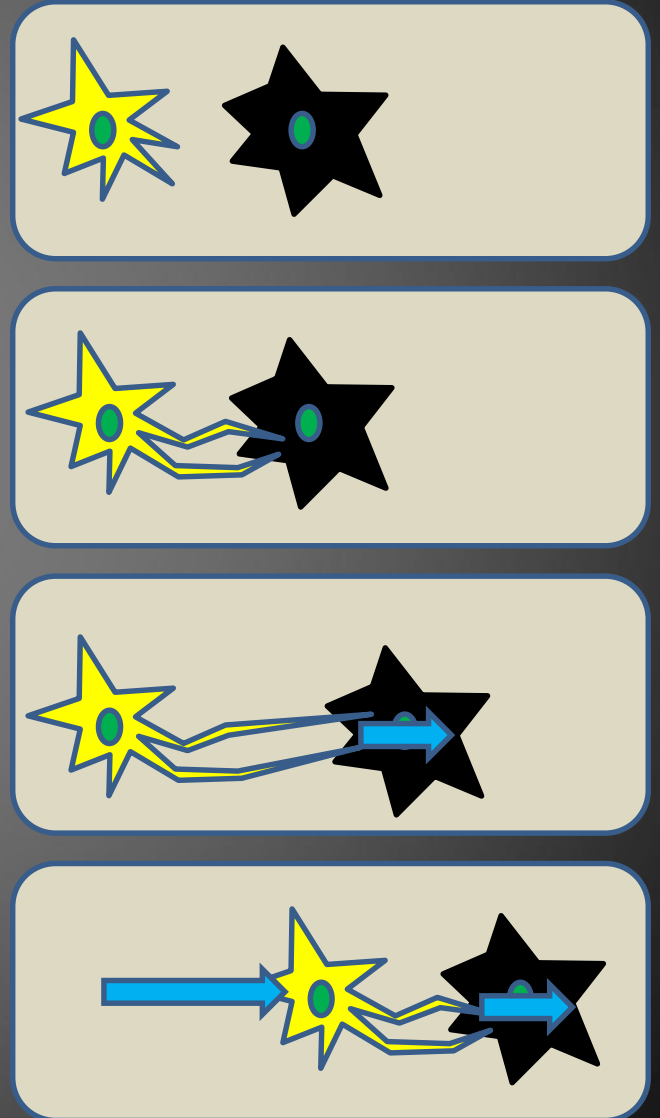


Time lapse observation

In vitro behavior of pigment cells

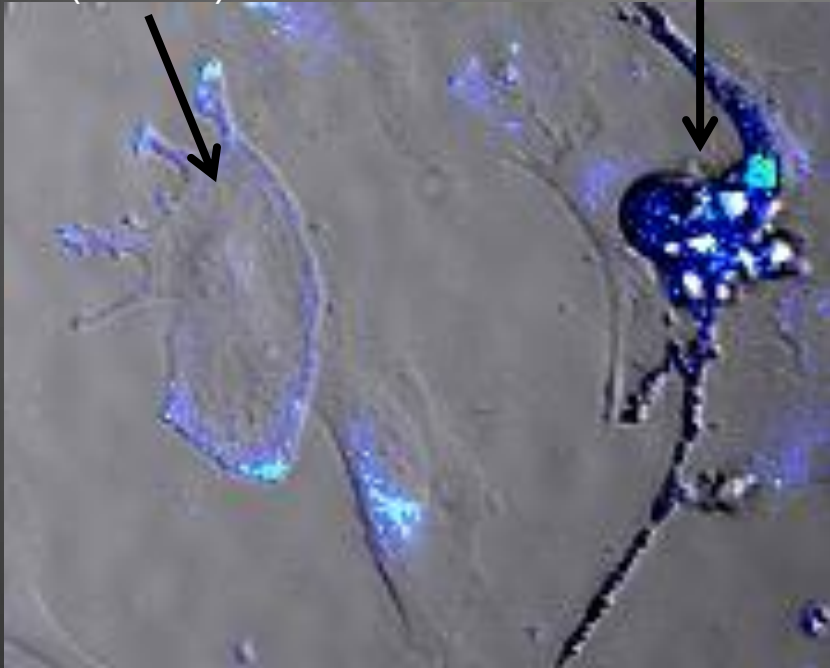


Wild type

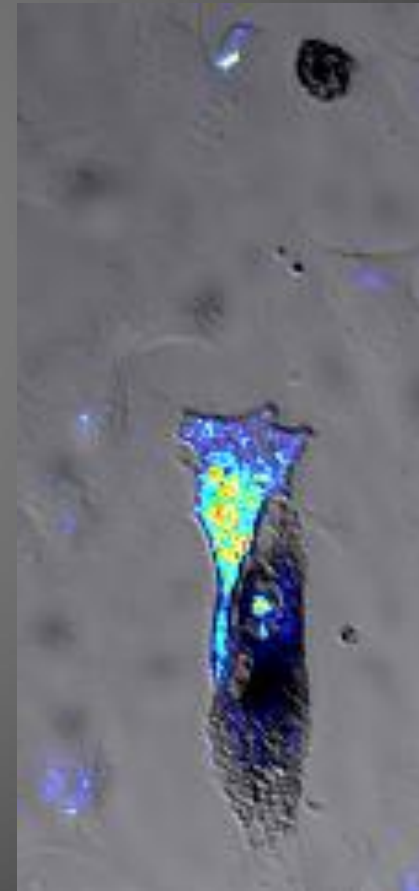


*Melanophores depolarize
when touched by xanthophores*

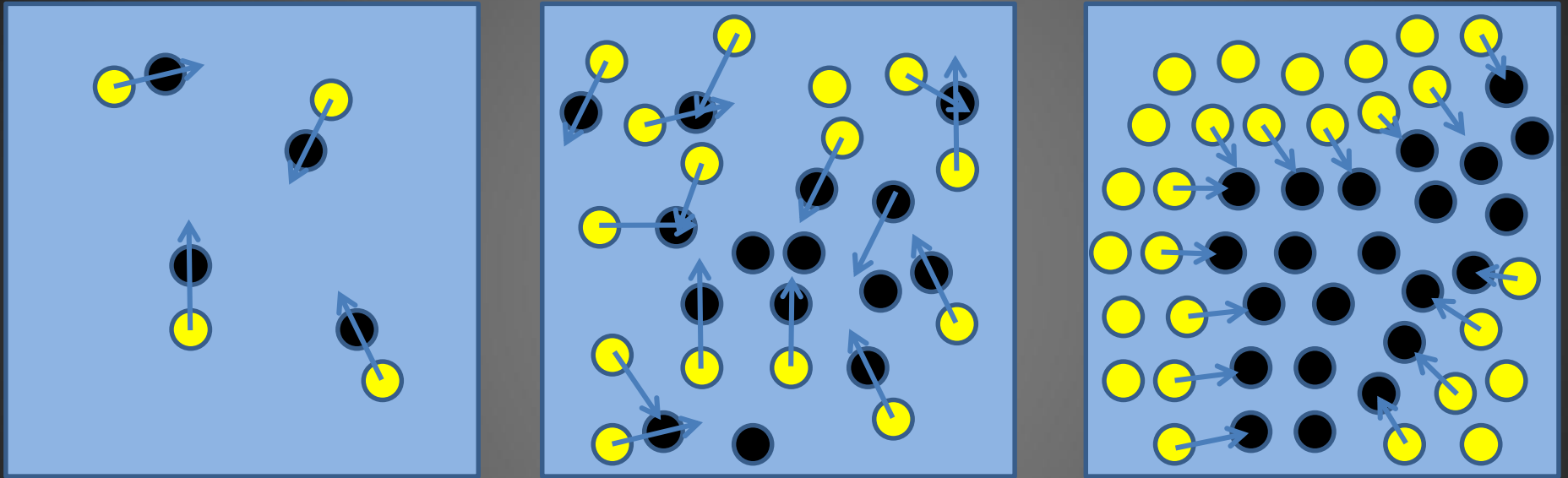
Melanophore
(Albino)



xanthophore



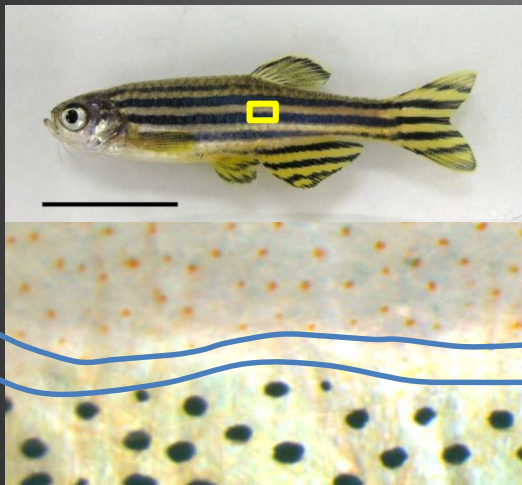
If the run-and-chase movement continues



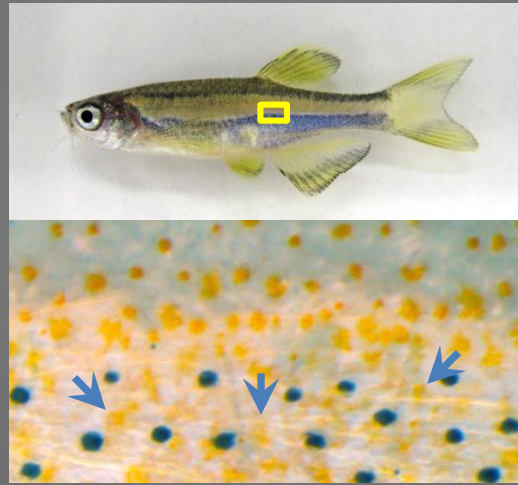
melanophores and xanthophores will segregates.

Separation is impaired in Kir7.1 mutant

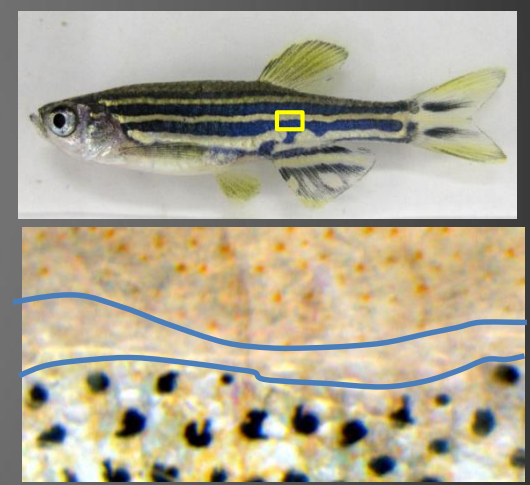
Wild type



Jaguar(Kir7.1)



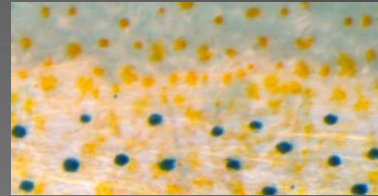
jaguar; mitfa:kir7.1



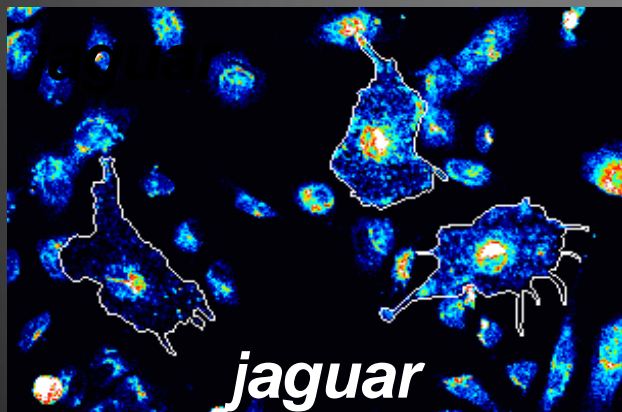
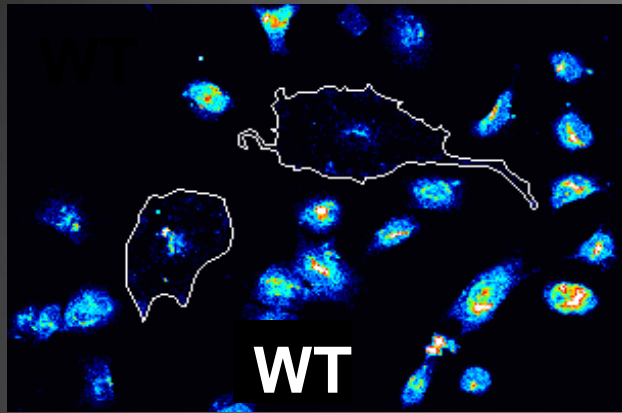
Kir7.1 is expressed in melanophore

Behavior of pigment cells in Kir7.1 mutant,

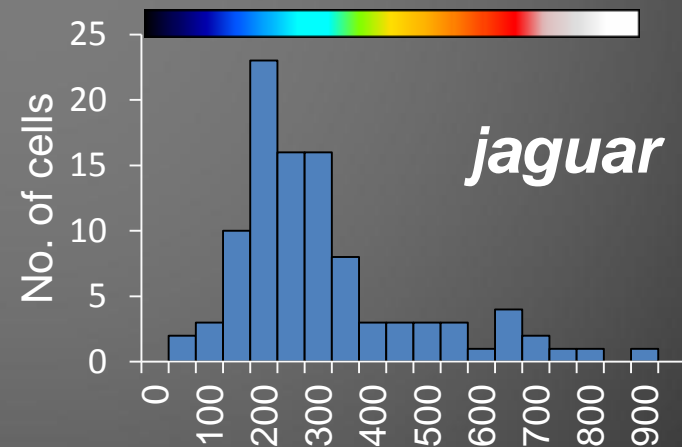
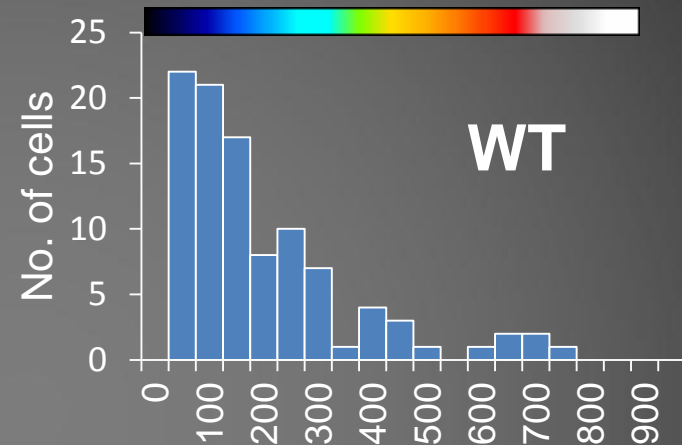
Kir7.1 is expressed only in melanophores (black cells)



Without Kir7.1, melanophores are always depolarized (=activated)

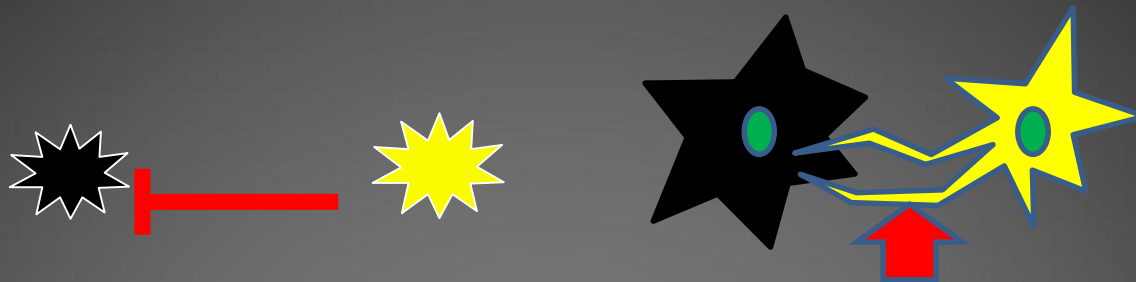


depolarization



Mean fluorescence intensity

Depolarization is detected by DiBAC reagent that accumulates in depolarized cells.

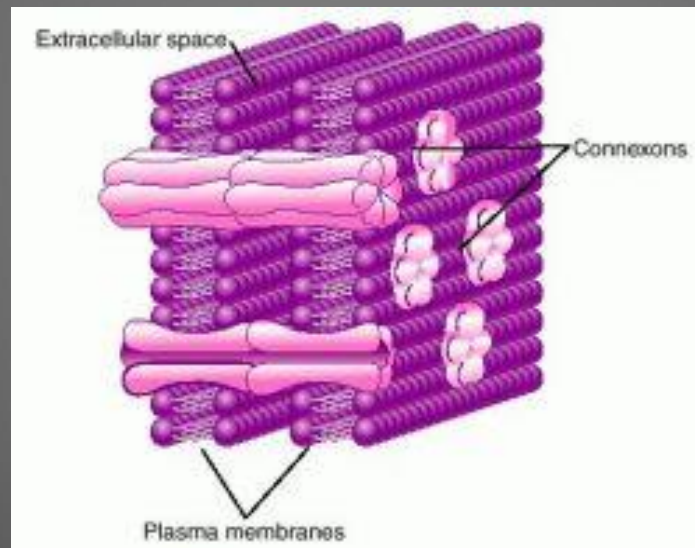


Depolarization(Kir7.1)
Migration(in vitro, in vivo)
Apoptosis(in vivo)



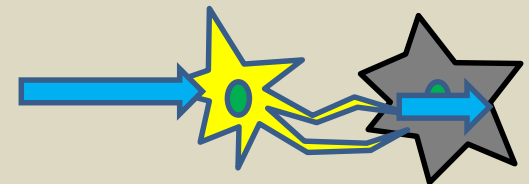
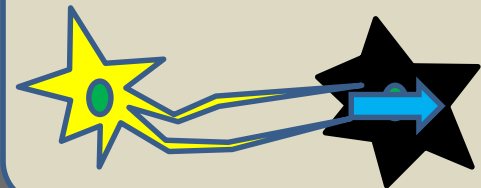
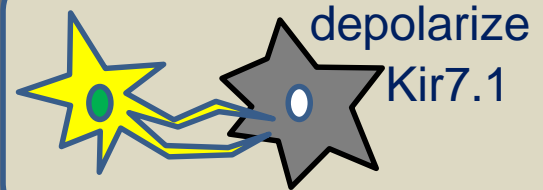
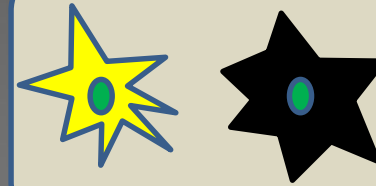


cx418 (gap junction)

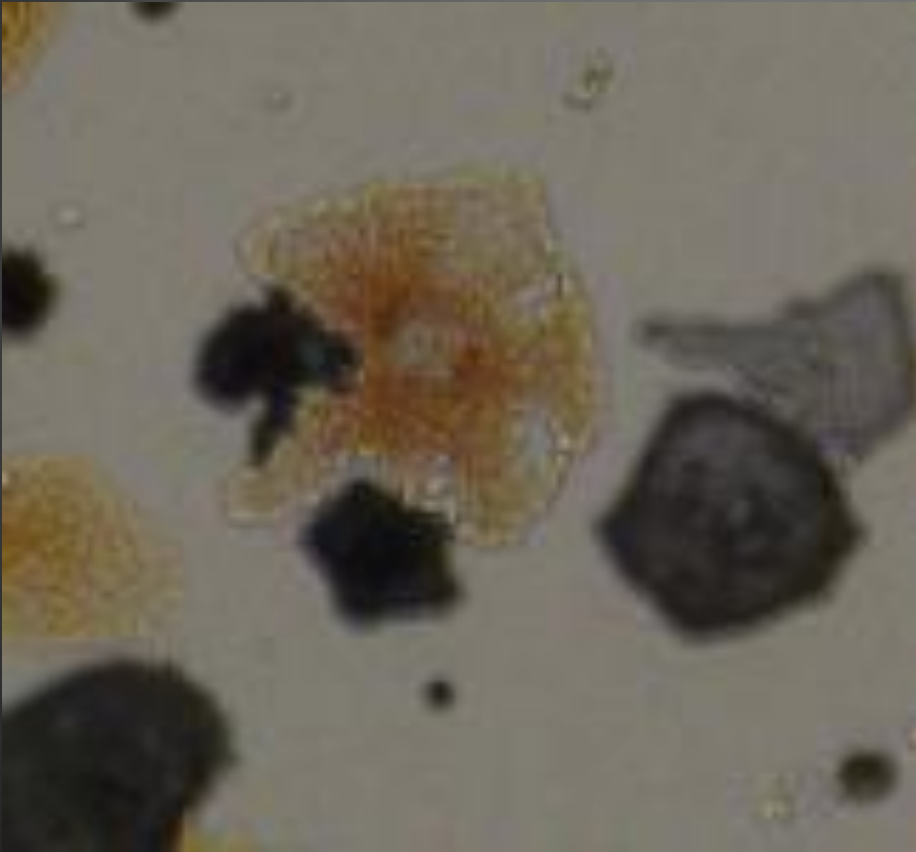


Expressed in both melanophores and xanthophores

In vitro behavior of pigment cells

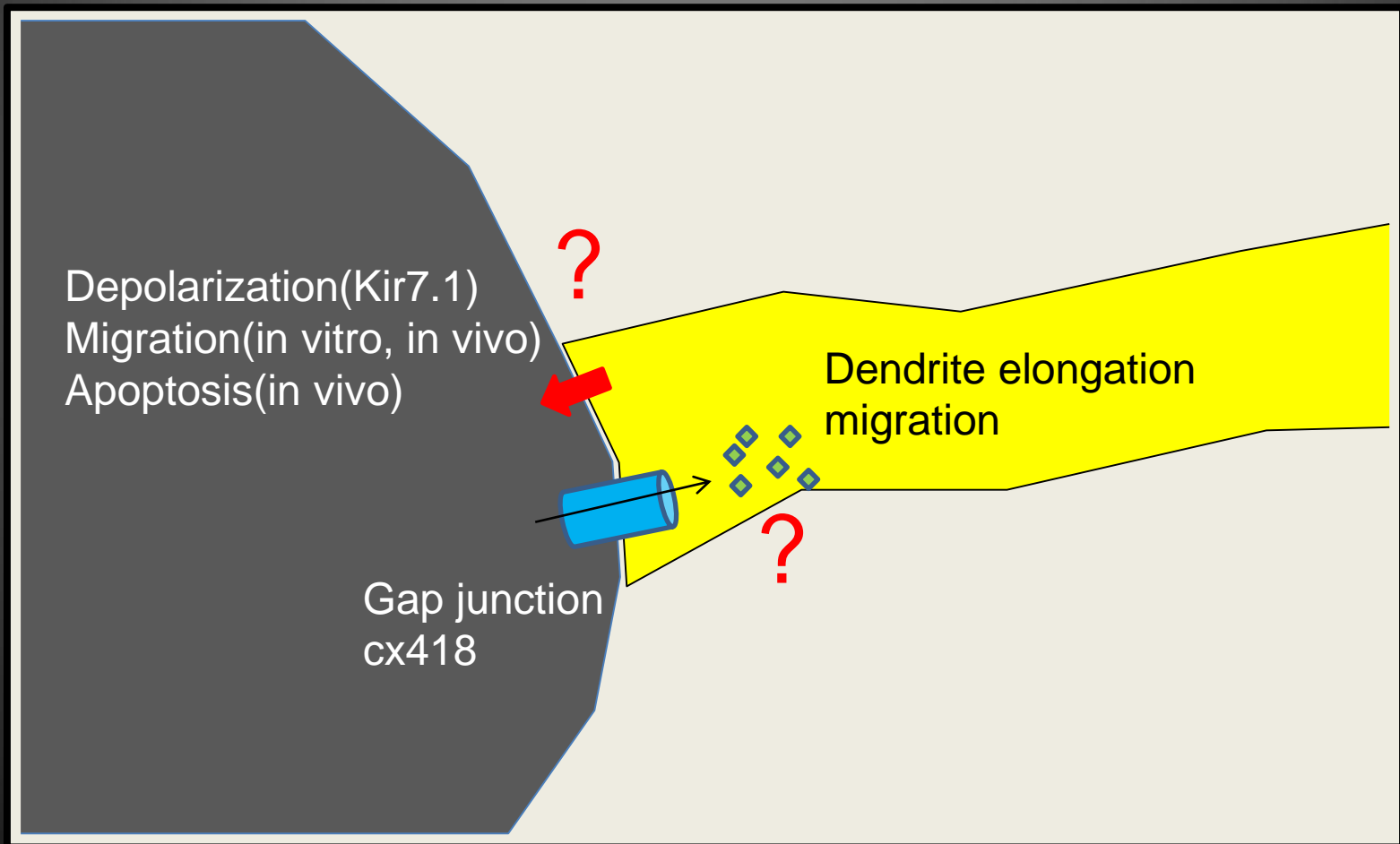
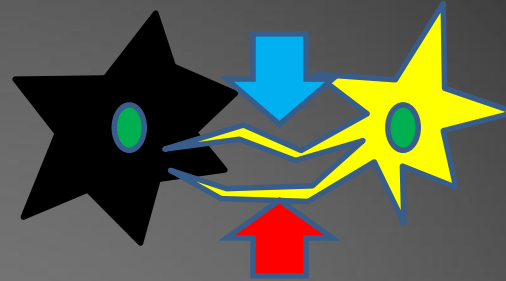


Pigment cell behavior of gap junction mutant



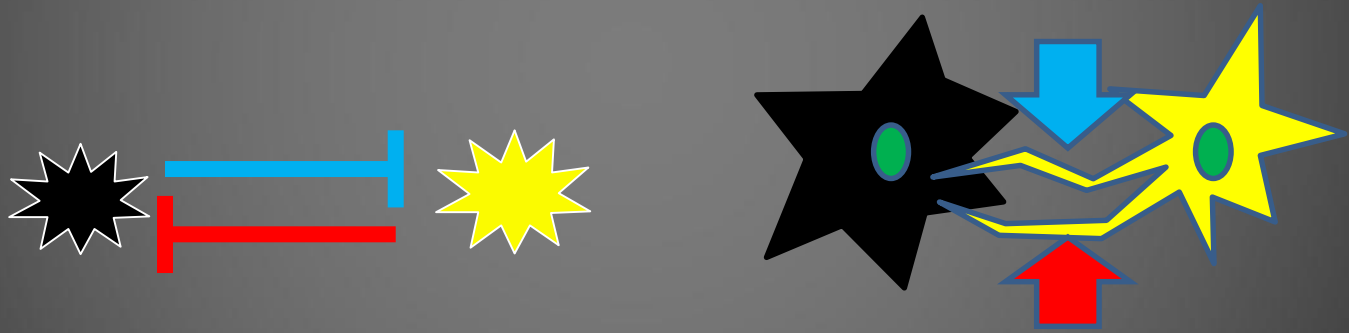
Xanthophores looks unable to sense melanophores.





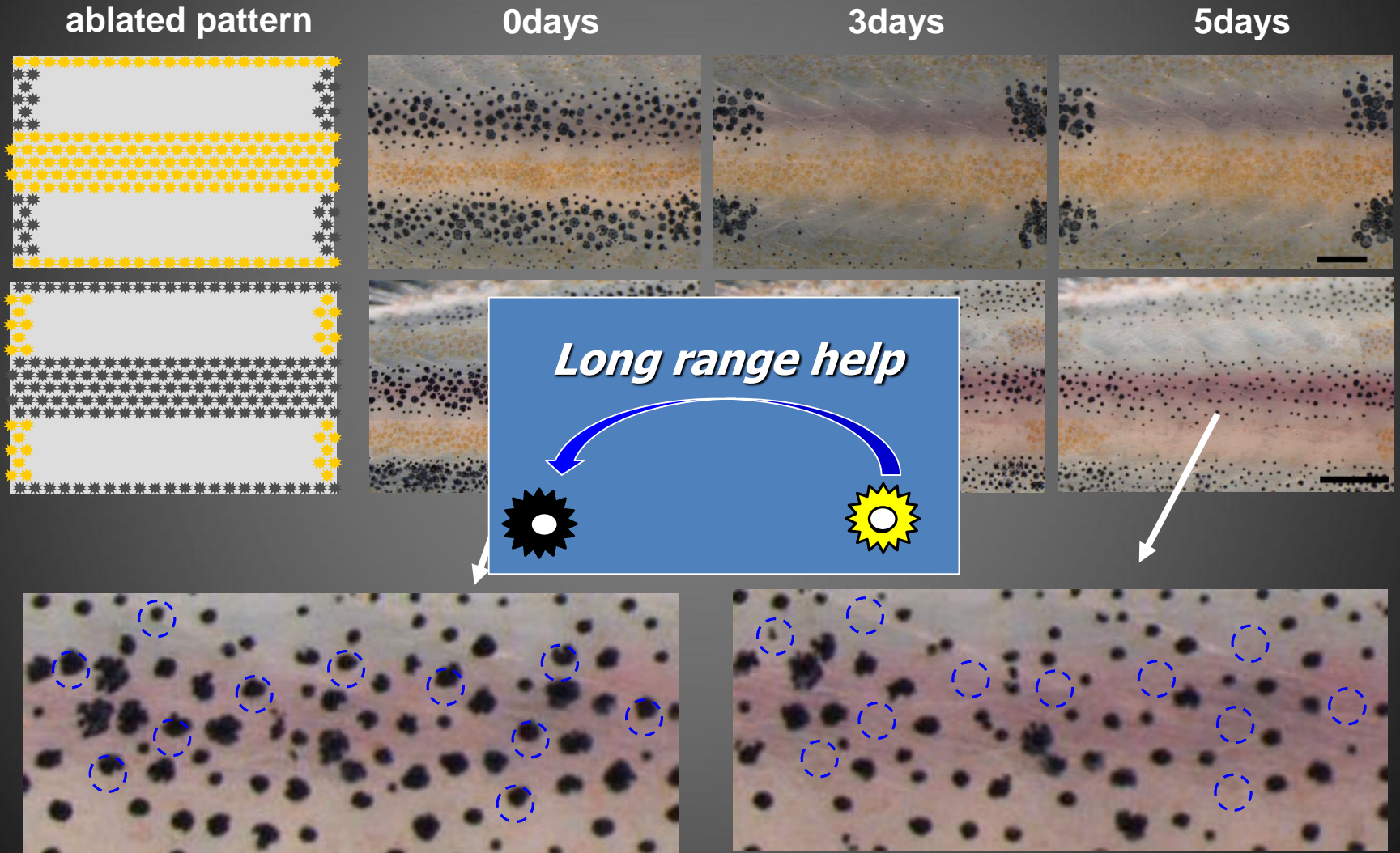
What does the mathematical theory suggest ?

1. There should be two kinds of interactions between the pigment cells.
2. Functional distance of the interactions should be different

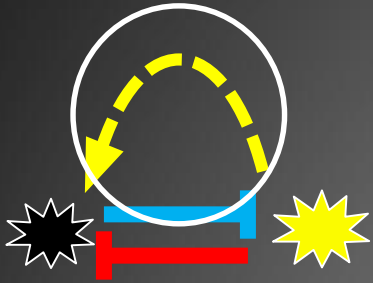


Another “ long range” signal ?

Laser ablation of xanthophores (yellow cells) induced the cell death of melanophores

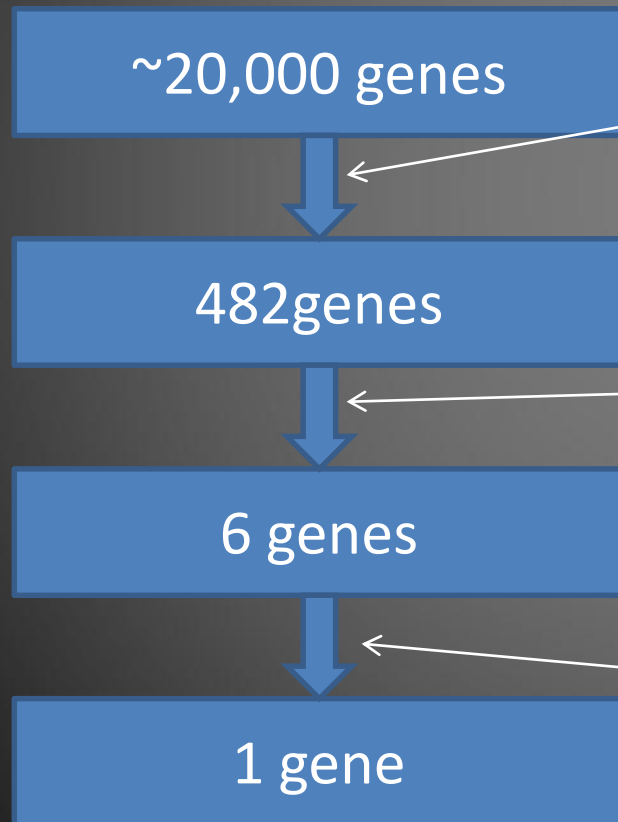


Finding the long range factor



Assumption

Ligand is expressed exclusively in xanthophores
Receptor is expressed exclusively in melanophores



Gene chip:

Xanthophores/melanophores > 3

Realtime PCR:

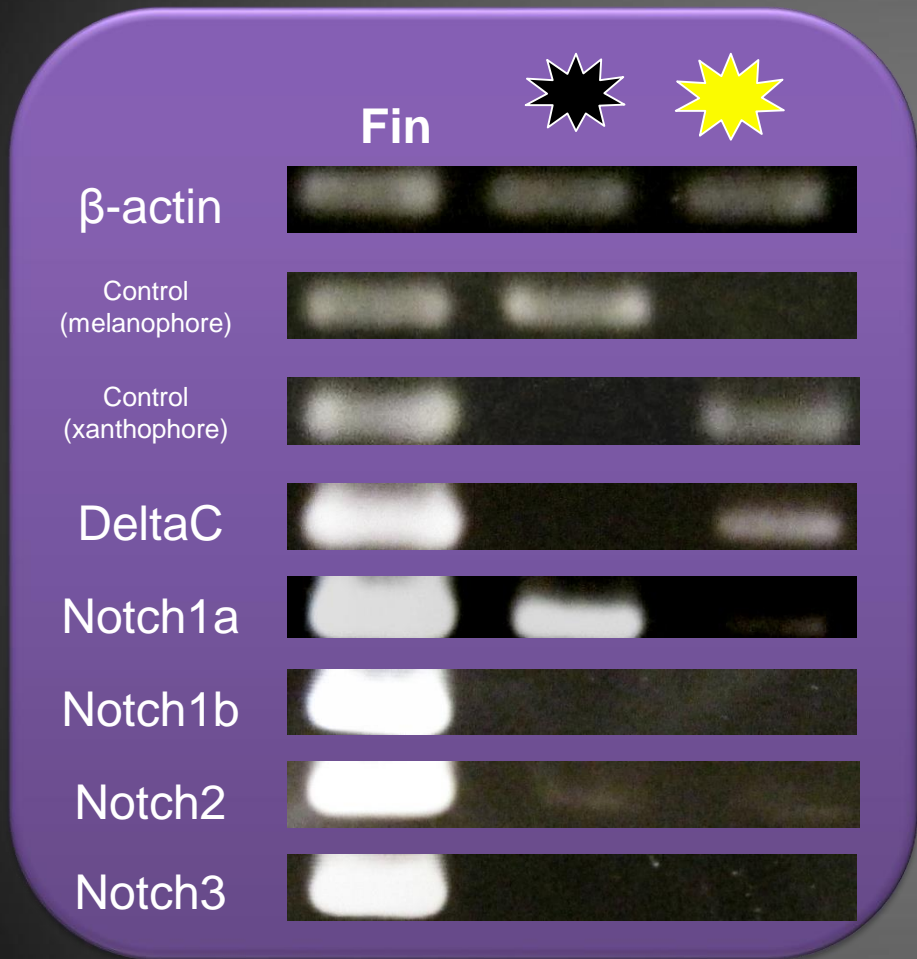
Xanthophores/melanophores > 5

+
Protein ligands or membrane proteins

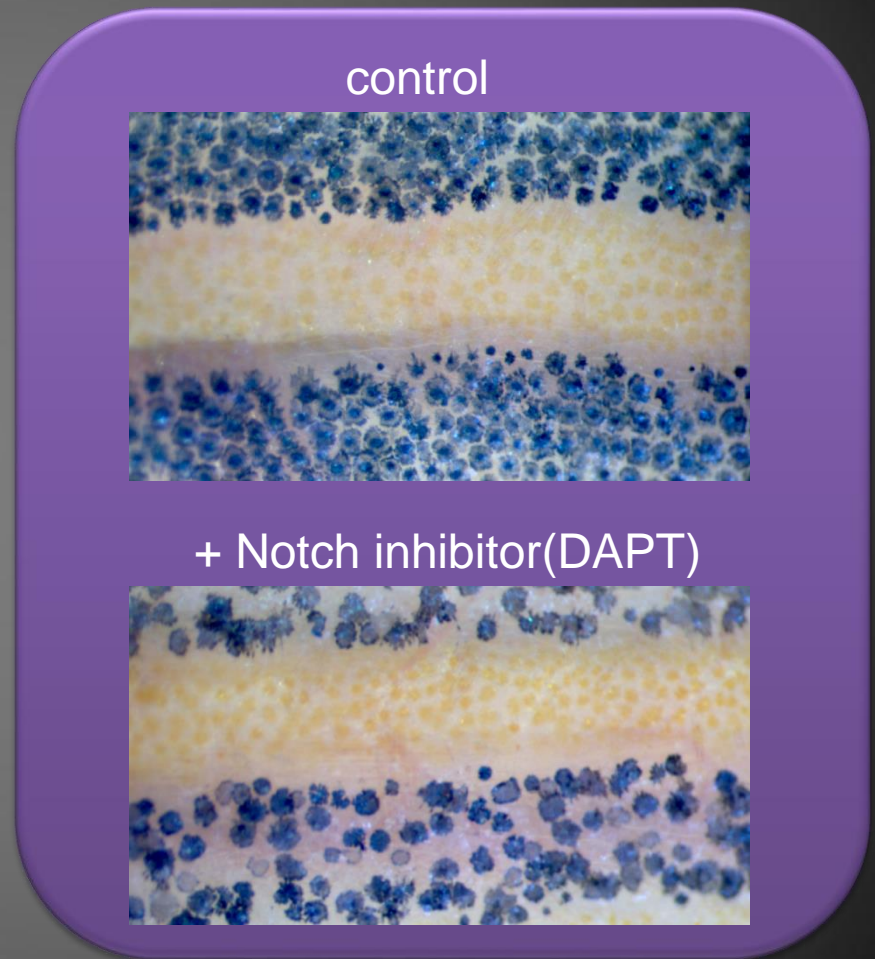
Transgenic fish:

Expression in melanophores
Alteration of the pattern?

Delta-Notch system involves in the long range effect

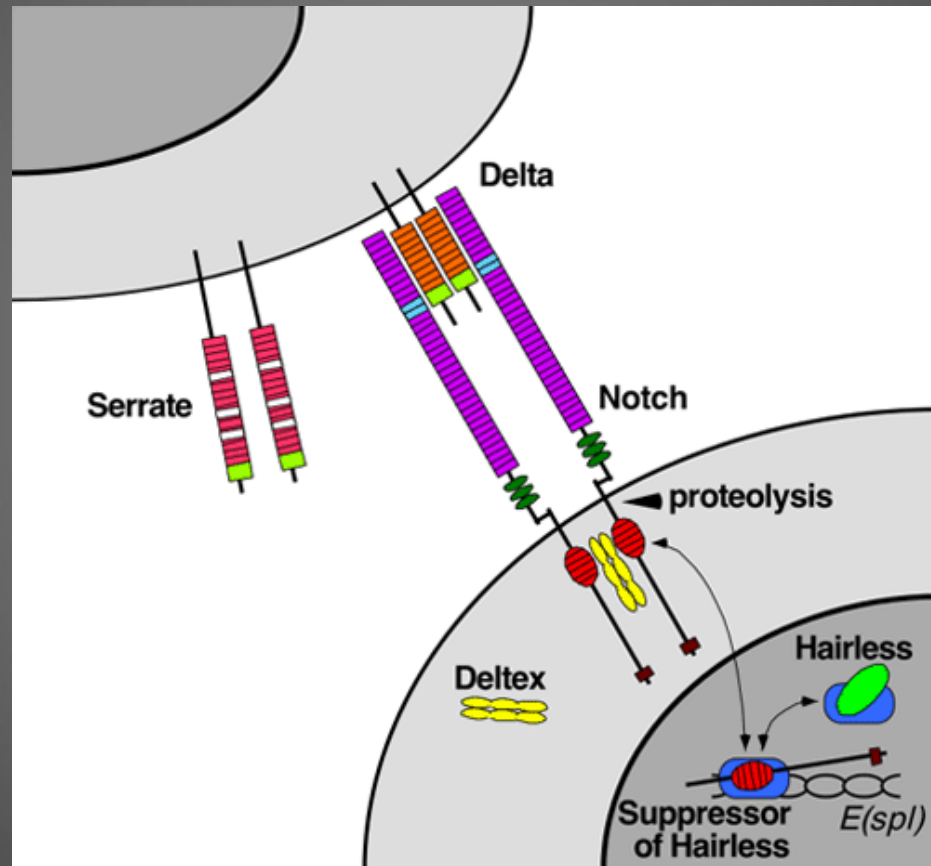


Xanthophores express DeltaC ligand
Melanophores express Notch1a receptor



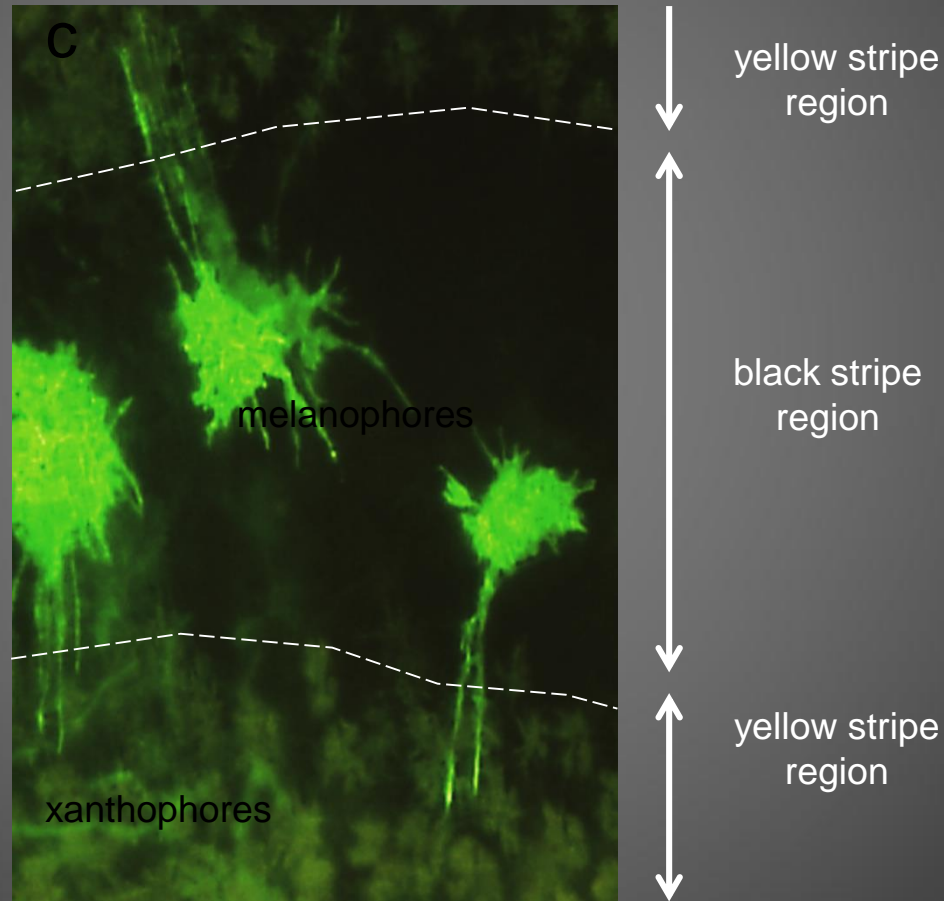
Inhibition of Notch signal
kills melanophores

Delta and Notch are the membrane bound molecules



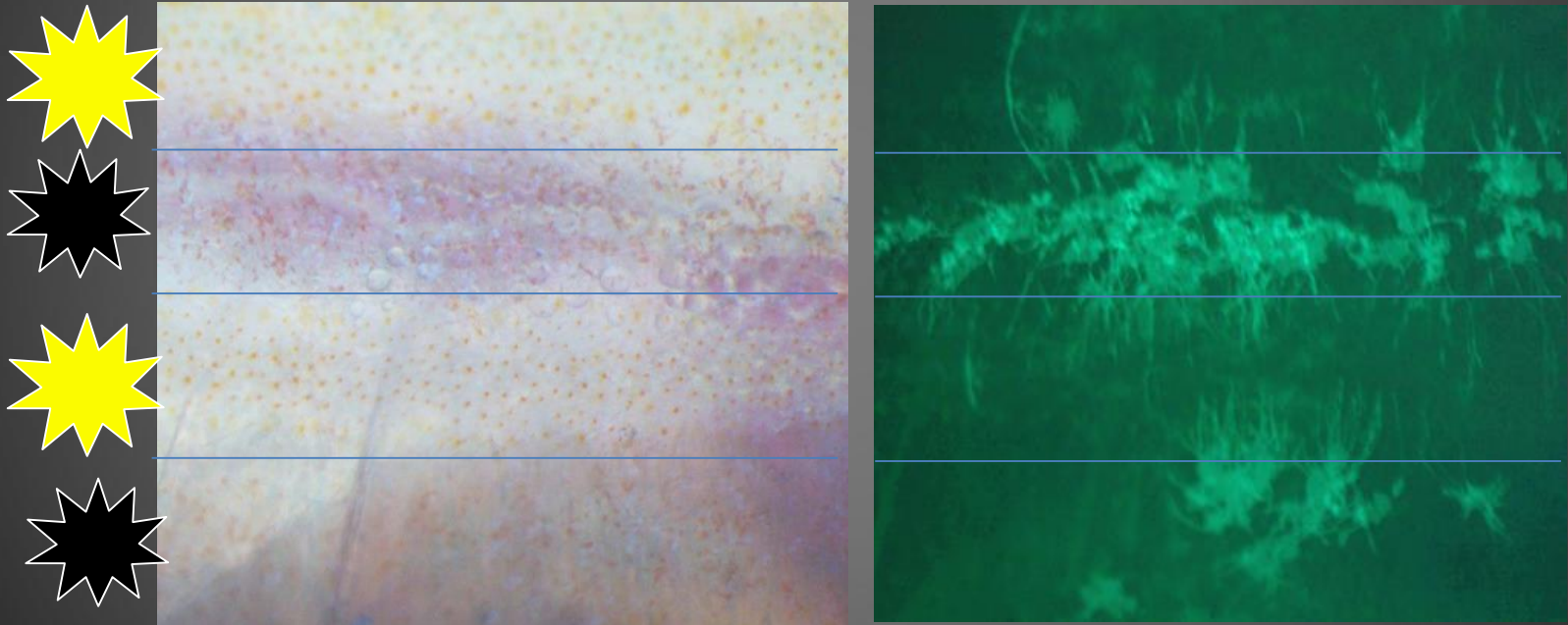
How do they work in the long range interaction ?

Melanophores elongate long projections toward xanthophores



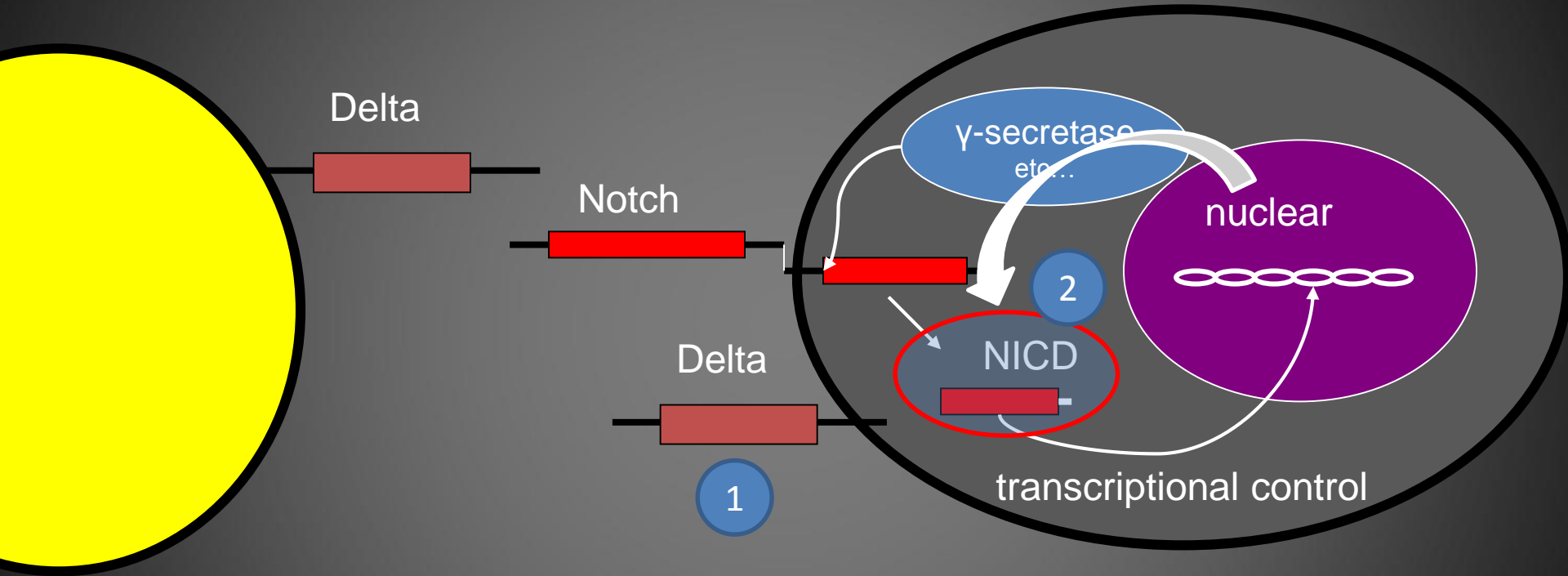
Membrane bound GFP is expressed in some of the melanophores

*Melanophores elongate long projections
toward xanthophores*



Membrane bound GFP is expressed in some of the melanophores

Transgenic fish with Delta or Notch gene



In both cases, melanophores should become less dependent to xanthophores.



Wider stripes or large spots ??

Ectopic expression of DeltaC or Notch1a-ICD caused wider stripes

WT



1



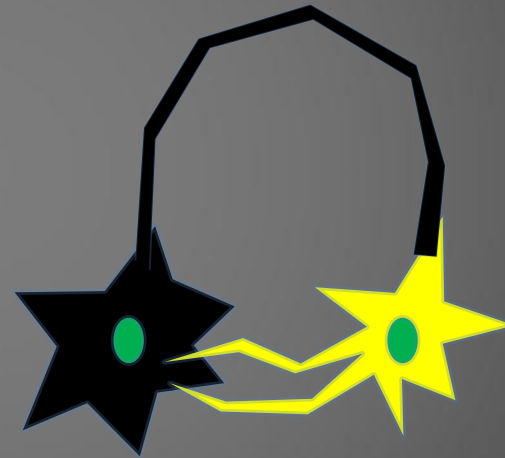
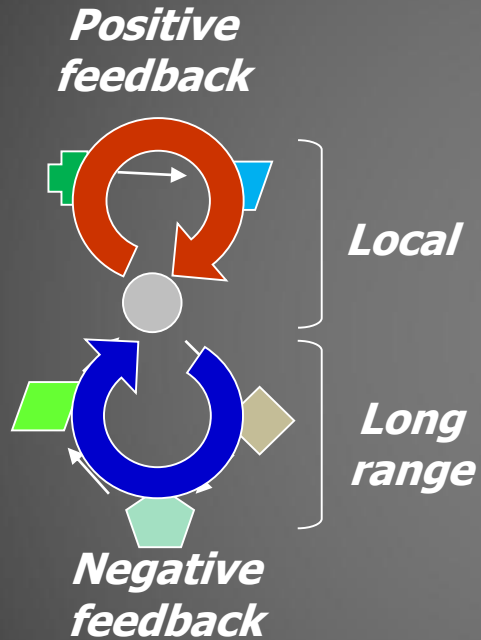
WT; DeltaC expressed in melanophores

2



Ectopic expression of Notch-ICD
in melanophore

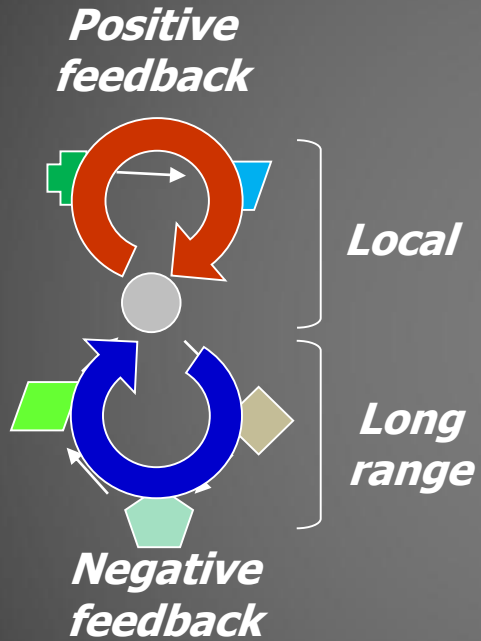
Is the identified network a “Turing mechanism” ?



local positive feedback

+

long negative feedback

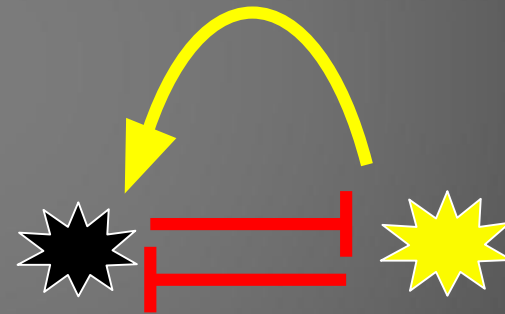


local positive feedback

+

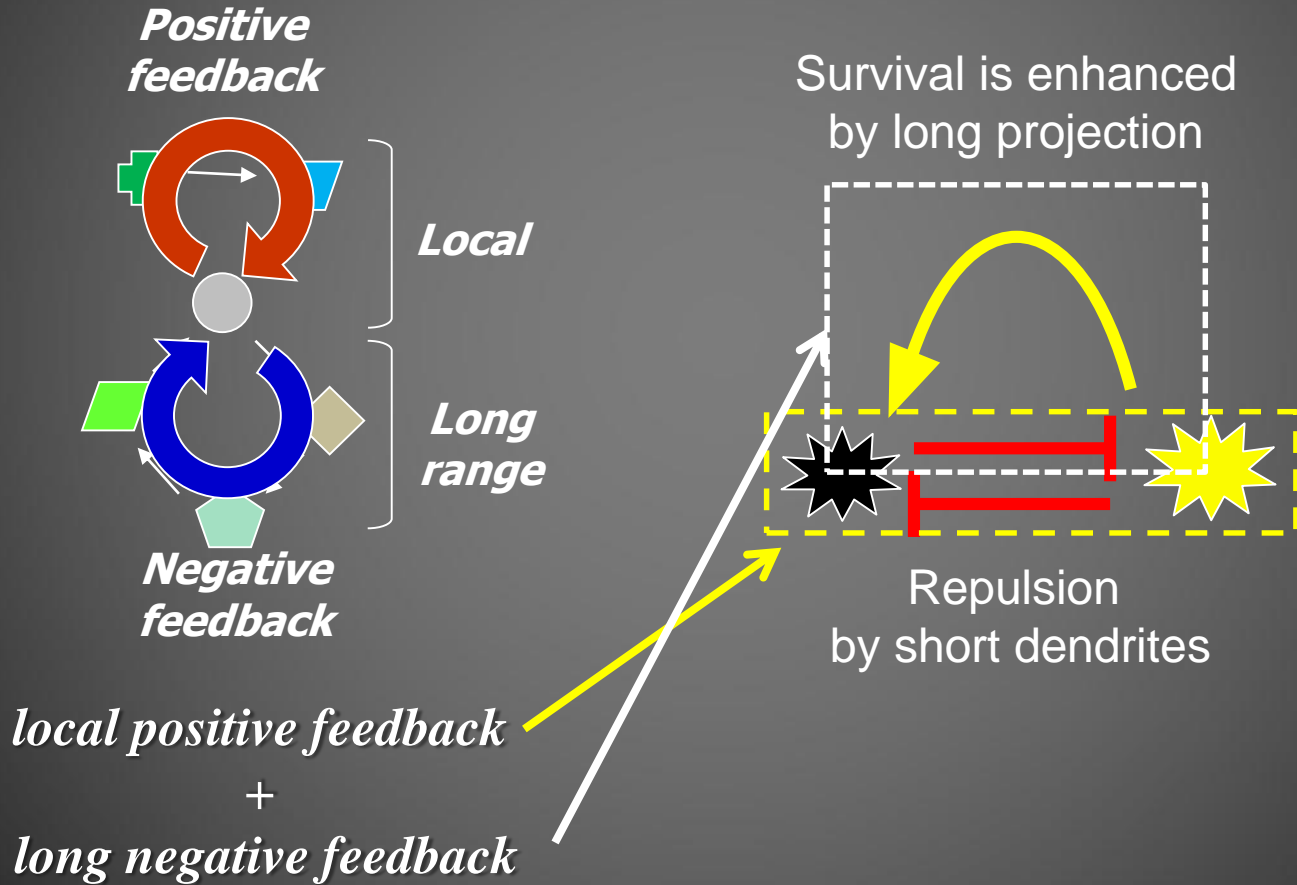
long negative feedback

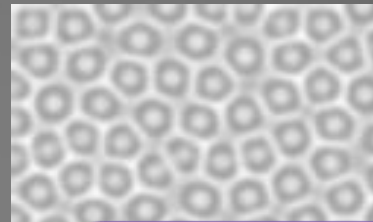
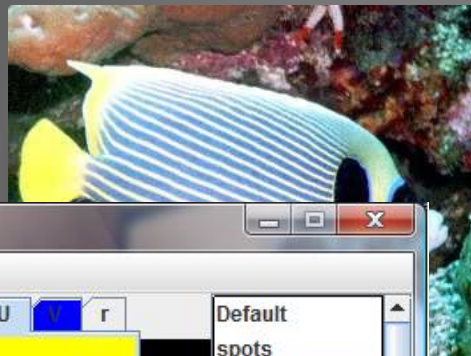
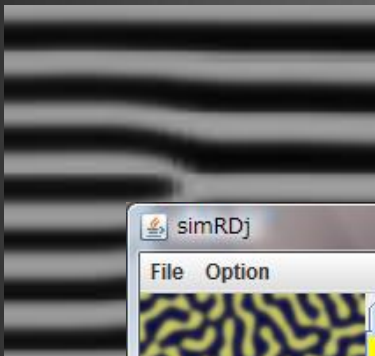
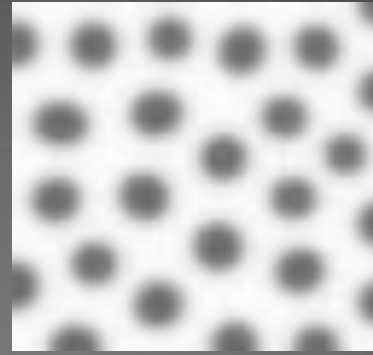
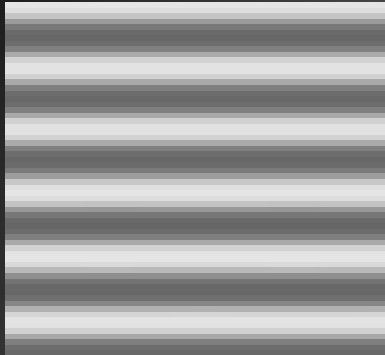
Survival is enhancement
by long projection



Repulsion
By short dendrites

Identified network is equivalent to the Turing model.





simRDj

File Option

U V r

3680

Randomize

Clear

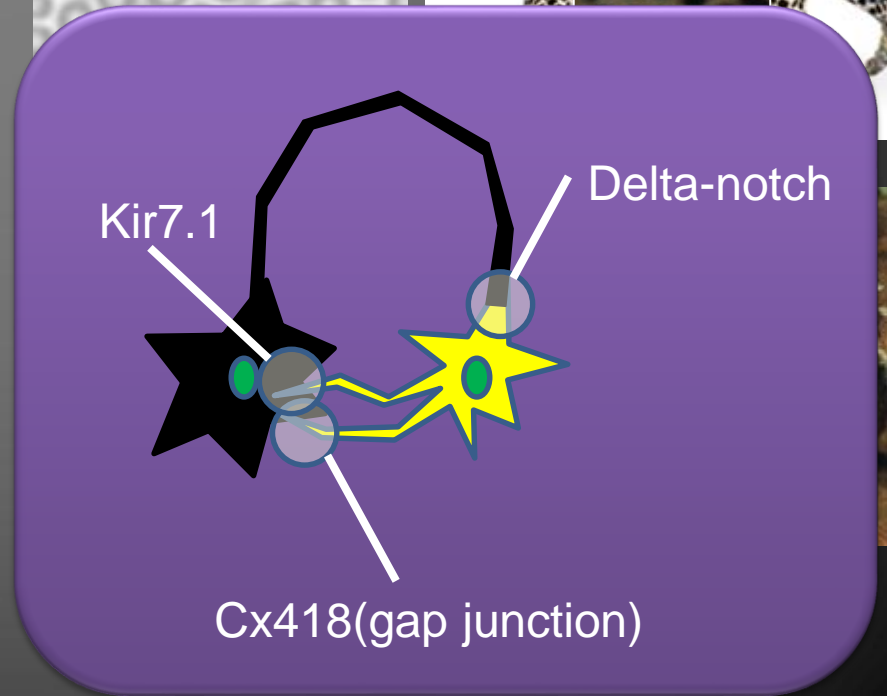
Start Capture Save Del

Default
spots
stripes
dark spots
chaotic
girraff
BZ (spiral) wave

$$\frac{\partial u}{\partial t} = F(u, v) - d_u u + D_u \Delta u$$

$$0 \leq F(u, v) = a_u u + b_u v + c_u \leq Fmax$$

$$\frac{\partial v}{\partial t} = G(u, v) - d_v v + D_v \Delta v$$

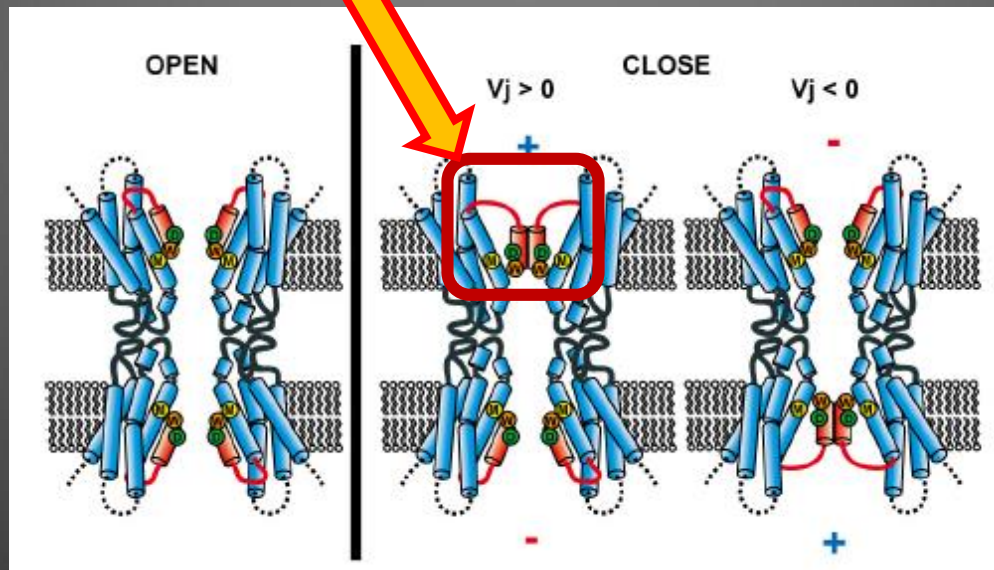
$$0 \leq G(u, v) = a_v u + b_v v + c_v \leq Gmax$$


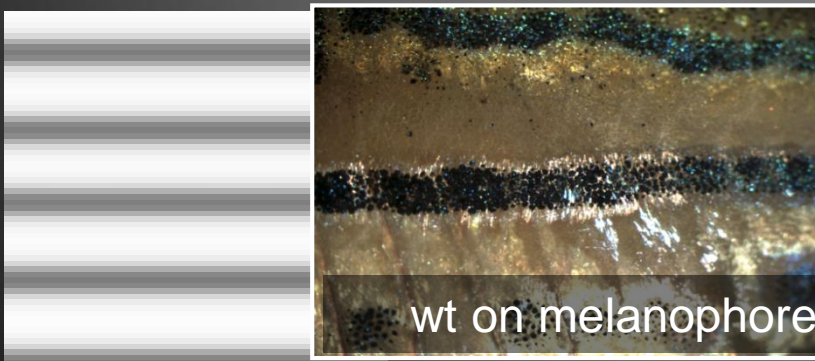
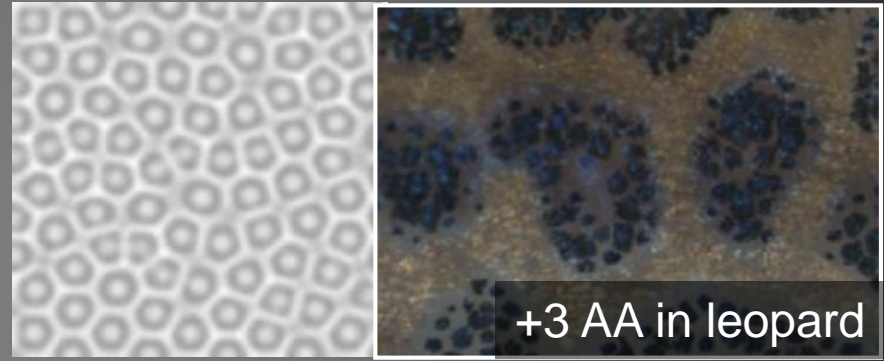
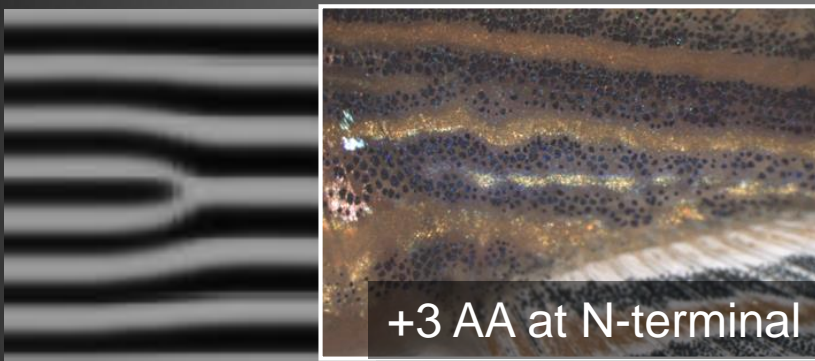
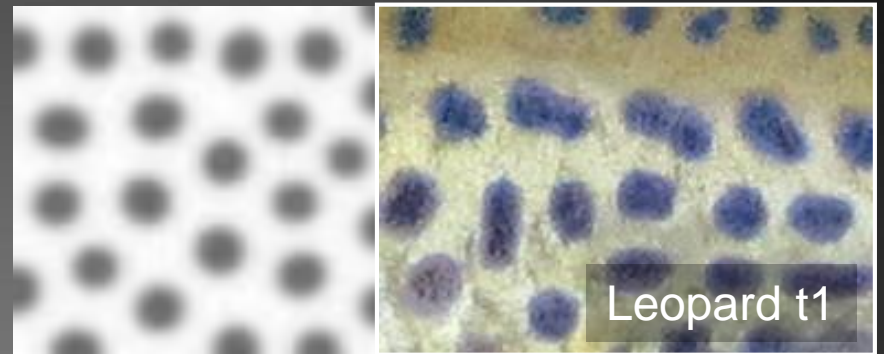
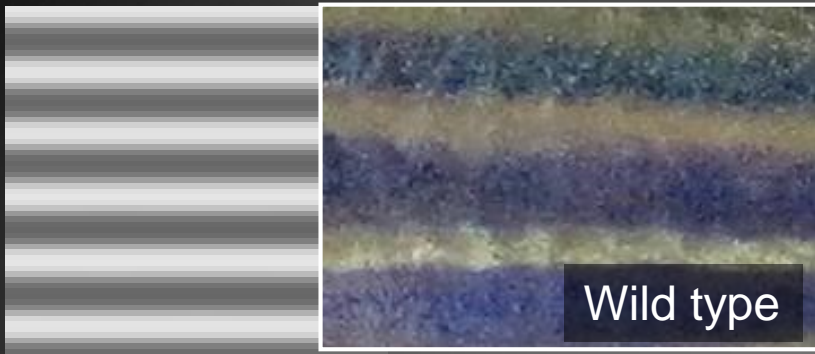
Mutation at the N-terminal of connexin changes the channel activity of gap junction

TM1

MKL418
 418 (WT)
 418del-6
 418del-10
 418del-14

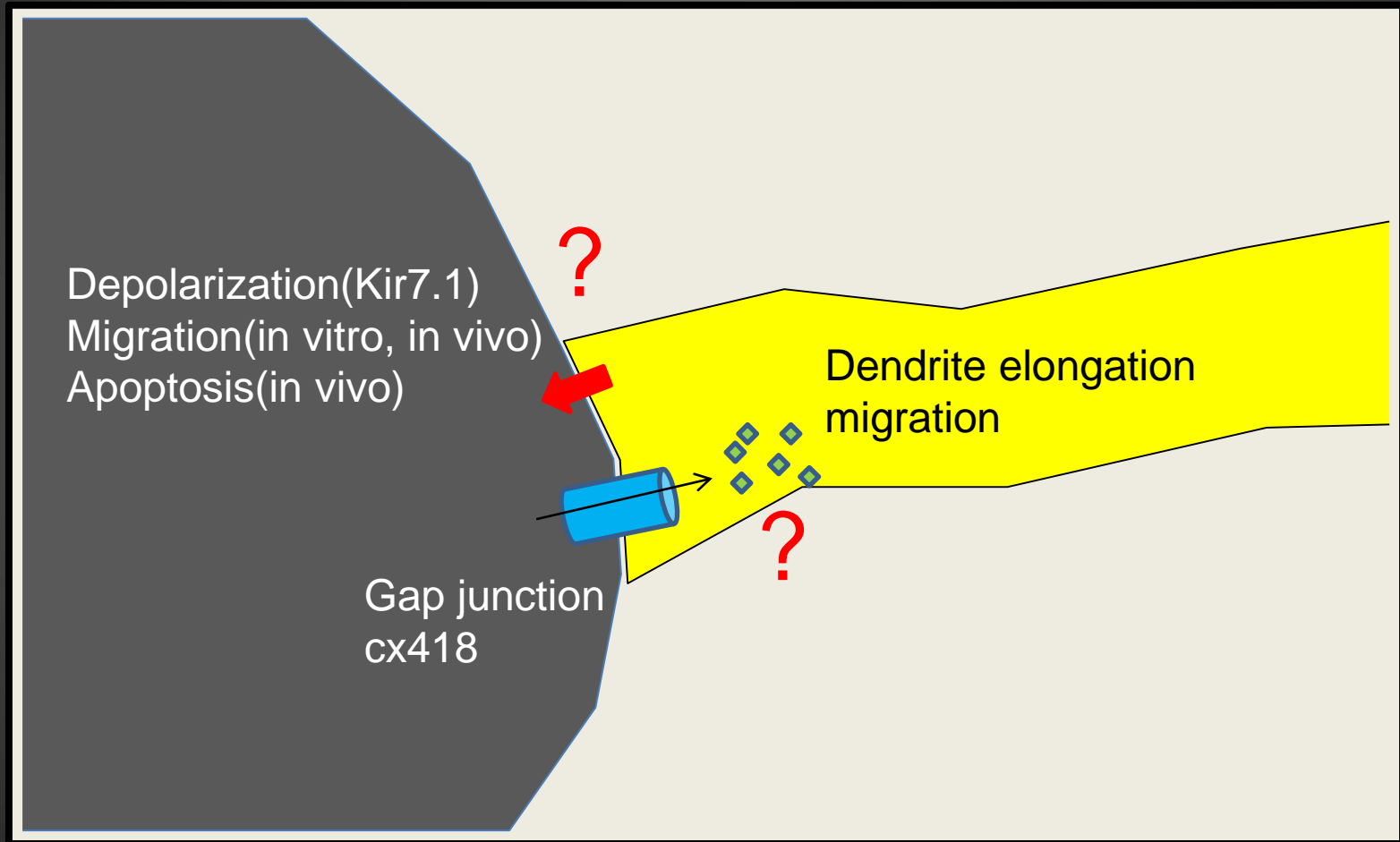
MKL	MADWSLLGSF	LEEVOEHSTS	VGKVWLTILF	IFRILVLGTA	AE
	<u>MADWSLLGSF</u>	<u>LEEVOEHSTS</u>	<u>VGKVWLTILF</u>	<u>IFRILVLGTA</u>	<u>AE</u>
	MGSF	LEEVOEHSTS	VGKVWLTILF	IFRILVLGTA	AE
		MEEVOEHSTS	VGKVWLTILF	IFRILVLGTA	AE
		MEHSTS	VGKVWLTILF	IFRILVLGTA	AE





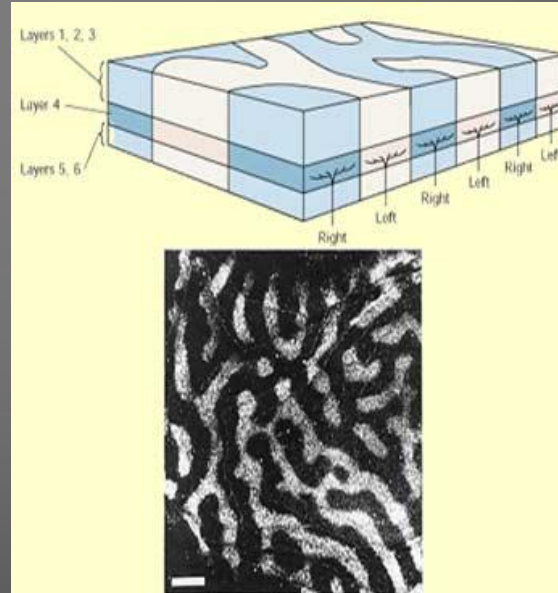
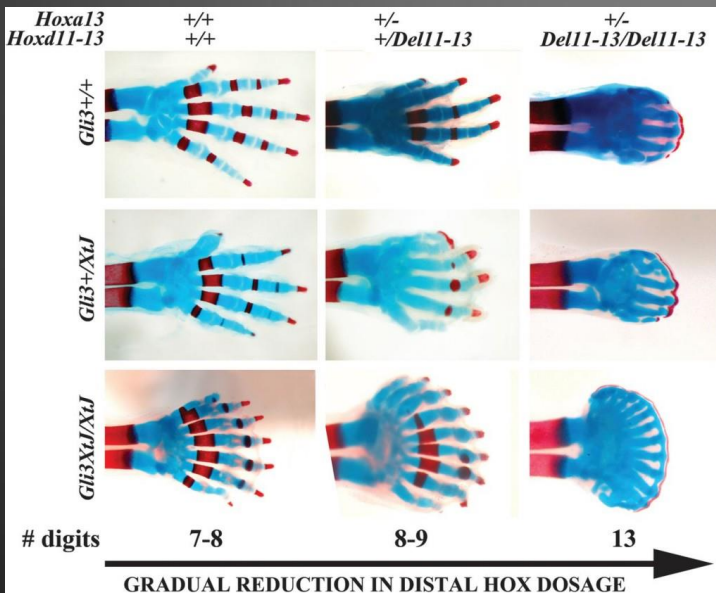
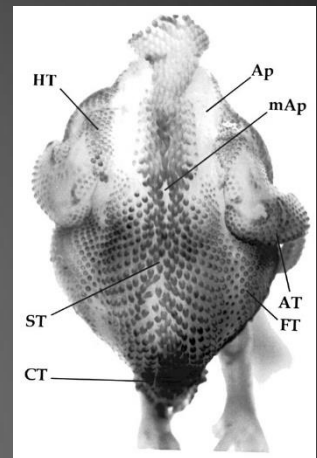
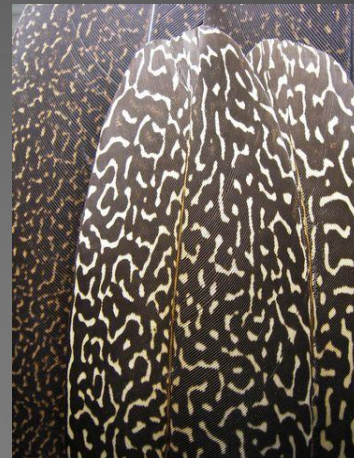
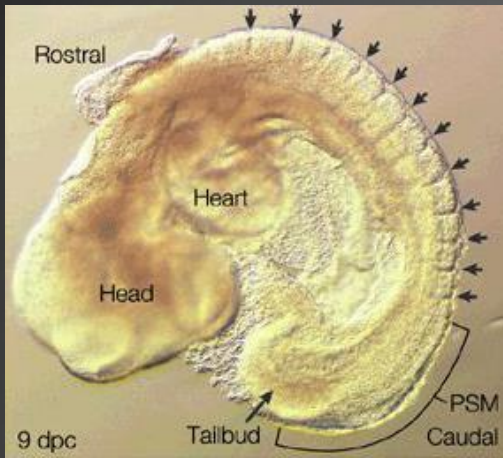
Most of the pigment patterns seen in the wild animal can be made by the modified cx418 gene.

questions to be answered



What is the signaling molecules

questions to be answered



Does Turing mechanism involve in other system?

次のフェーズへ



Bone shape ?



Wild type

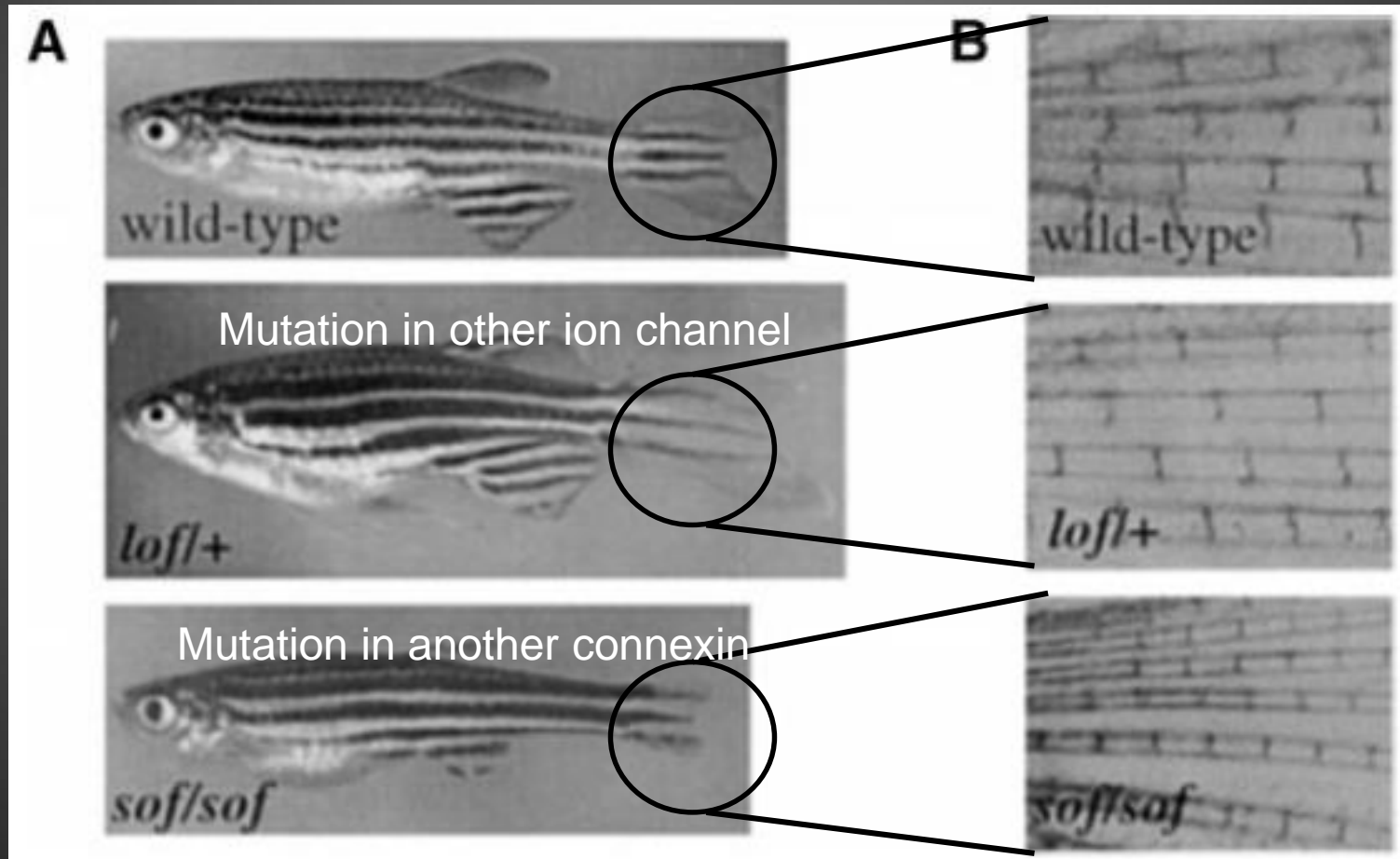


**Kir7.1
(K channel)**



**Connexin41.8
(gap junction)**

Shape of the fin-bones may be controlled by the similar mechanism.





gap junction
M. Watanabe



Delta-notch
H. Hamada



Thanks for your attention



In vitro culture
H. Yamanaka



Jaguar (Kir7.1)
M. Inaba

Institute of frontier bioscience
Pattern Formation Group

Pattern formation labo



Your visit is welcome !