

Modelling complexe biological systems in the context of
genomics

Ordinary Differential Equations (ODEs) and the biological switch

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Plan

- ODEs and biology: Why?
- Biological Switch
- Biological switch and ODEs
- Geometrical representation of ODEs
- Geometrical interpretation of ODEs
- Geometrical interpretation of the switch
- Conclusion

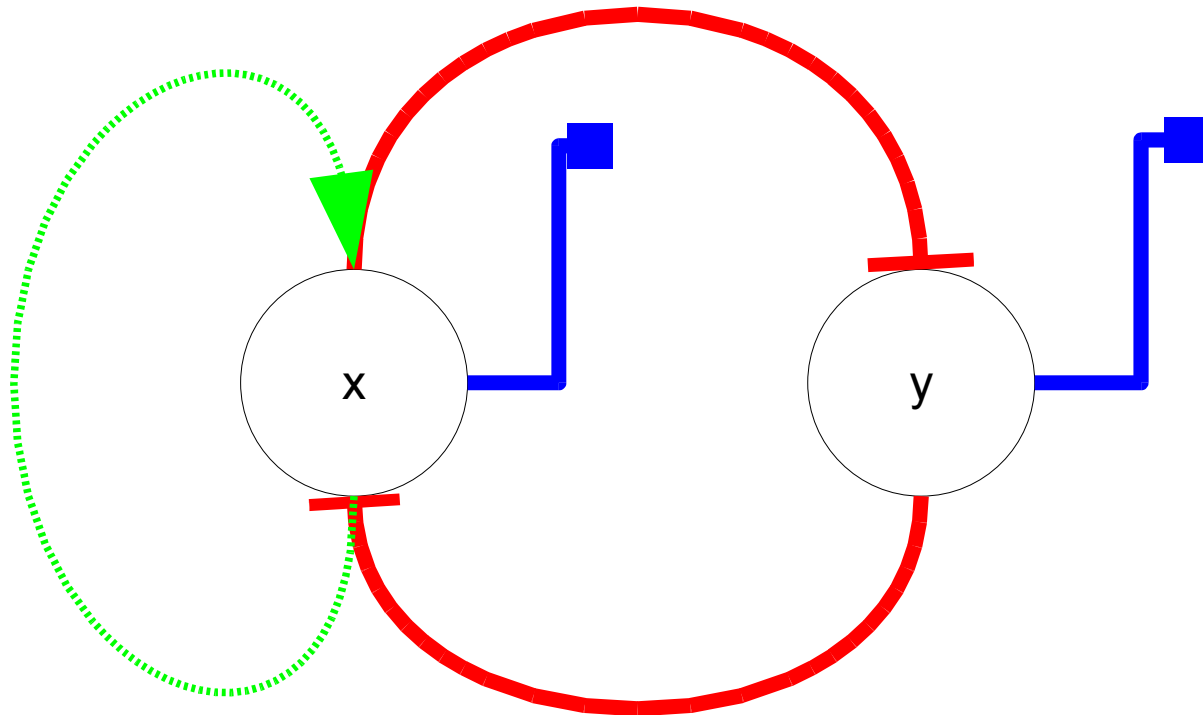
ODEs and Biology

- Assumptions for ODEs:
 - Deterministic system
 - No uncertainty
 - Populational level of description
 - ***dynamics*** (or time evolution)***of a system***
- Why ODEs:
 - Strong mathematical history and background
 - Historical relationships between ODEs and biology (bio)chemistry, enzymology, ecology, epidemiology
 - Well accepted formalism in biological communities
 - Software for *in silico* experiments for biologists

A first ODE tour for biologists

- Our aim in this tutorial is:
 - To present fundamental notions in analysing ODEs
 - Illustrate our to model a biological system (Switch)
 - To to understand the biological switch with these notions.

Biological Switch : schematic



- Two biological species: x and y
- x repress y
- y repress x
- x, y : *degradation*
- x : *auto-activation*

Modelling the Biological Switch using ODE formalism

- Fundamental idea:
 - We want the time evolution of x and y , that is $x(t)$
 - We don't know how to obtain a formula for $x(t)=???$
 - We know how to describe a small variation of the concentration of x and y during a small time interval dt
- Procedure (for each biological entity):
 - Identify each mechanism where x is involved
 - For each mechanism, give an equation describing a small variation of the concentration (dx) for a small time interval (dt)
 - Sum up to obtain $dx/dt = f(x,...)$

From dx/dt to $x(t)$

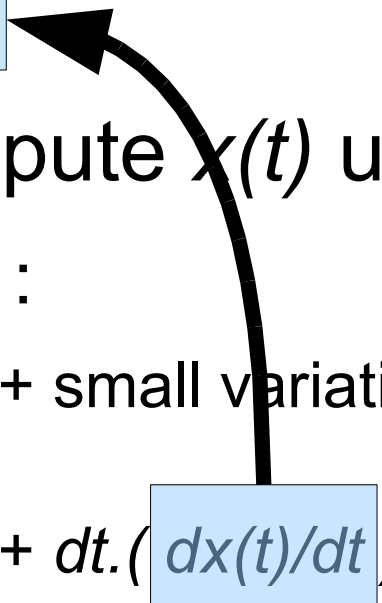
- $dx/dt = f(x, \dots)$
- One can compute $x(t)$ using dx/dt :

- By definition :

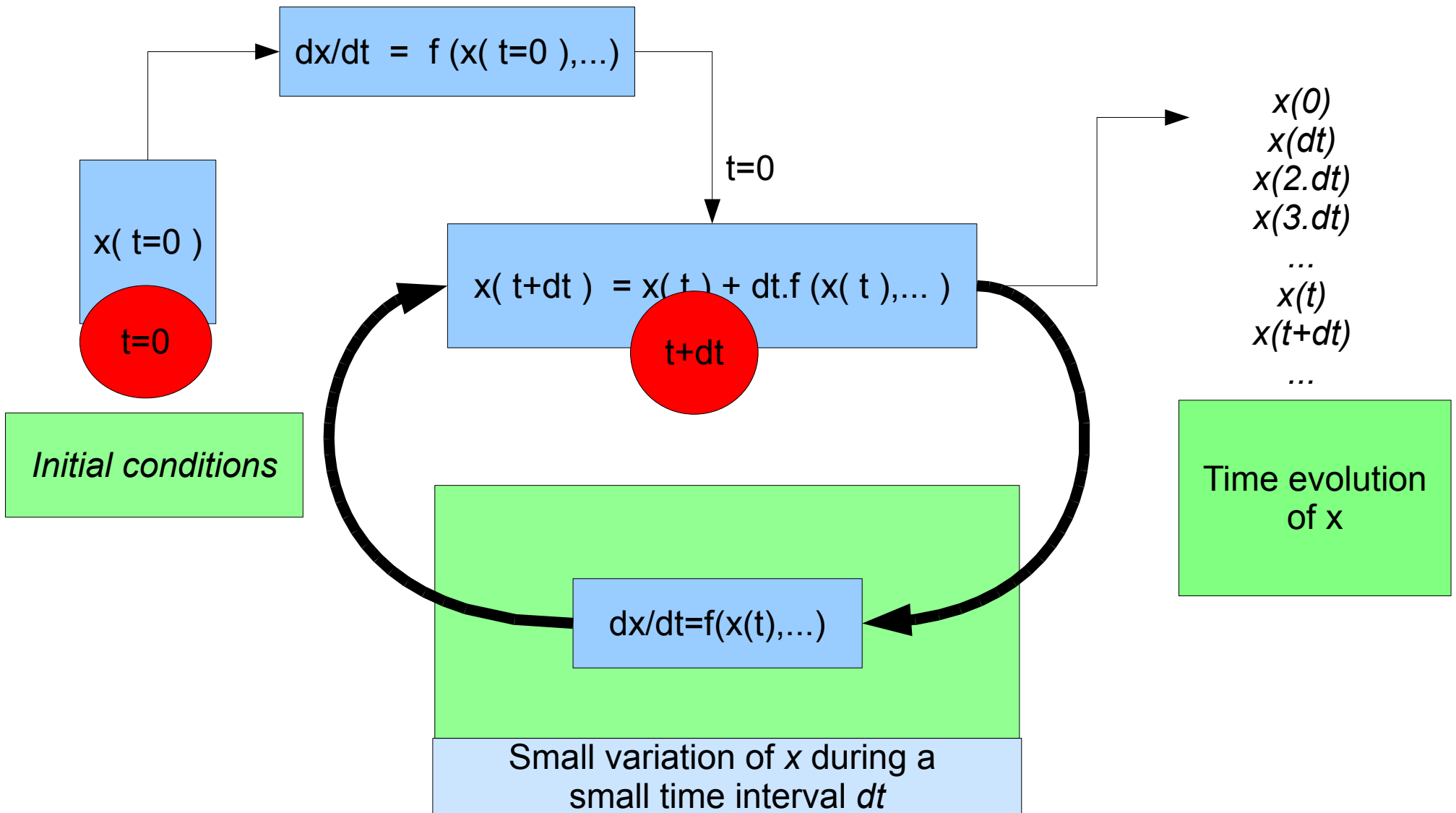
$x(t+dt) = x(t) +$ small variation of $x(t)$ during the small time interval dt .

$$x(t+dt) = x(t) + dt. (dx(t)/dt)$$

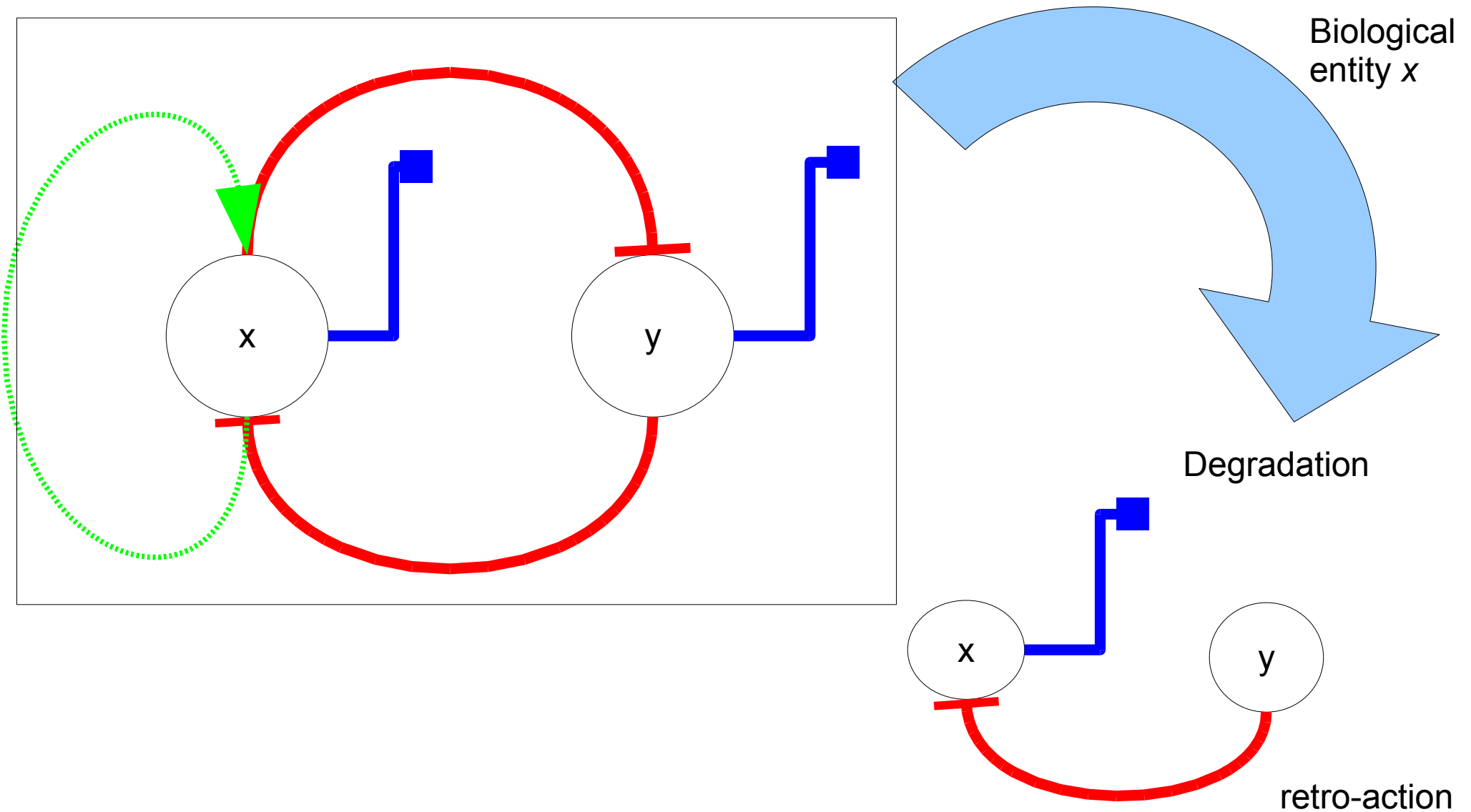
From dx/dt to $x(t)$

- $dx/dt = f(x, \dots)$
 - One can compute $x(t)$ using dx/dt :
 - By definition :
 $x(t+dt) = x(t) + \text{small variation of } x(t) \text{ during the small time interval } dt.$
 $x(t+dt) = x(t) + dt. (dx(t)/dt)$
 - We know how to compute dx/dt for a given $x(t)$
 - Then:
- 

Simple numerical integration scheme

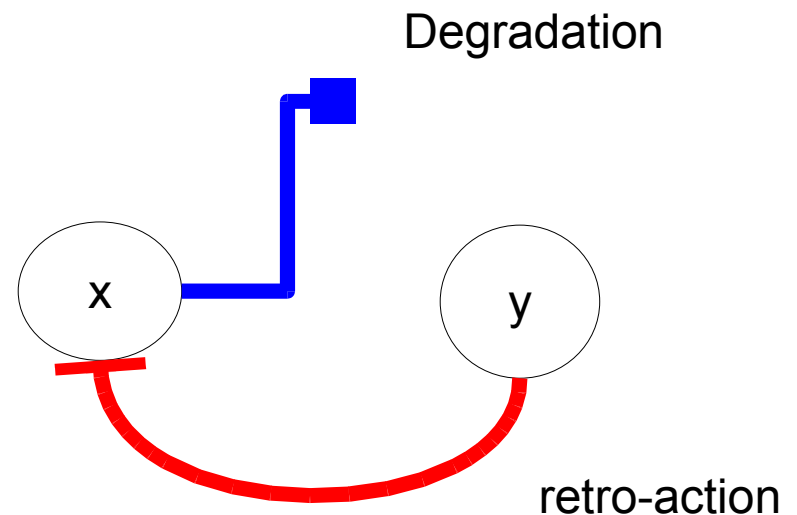


Building the biological switch ODE model



Biological switch : x proteins

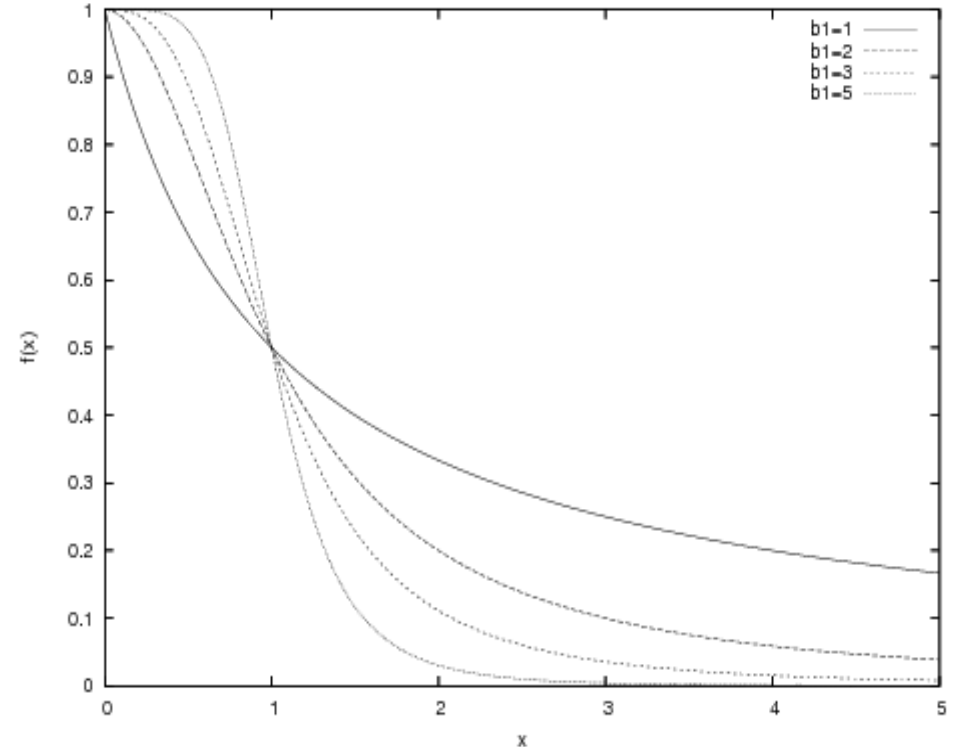
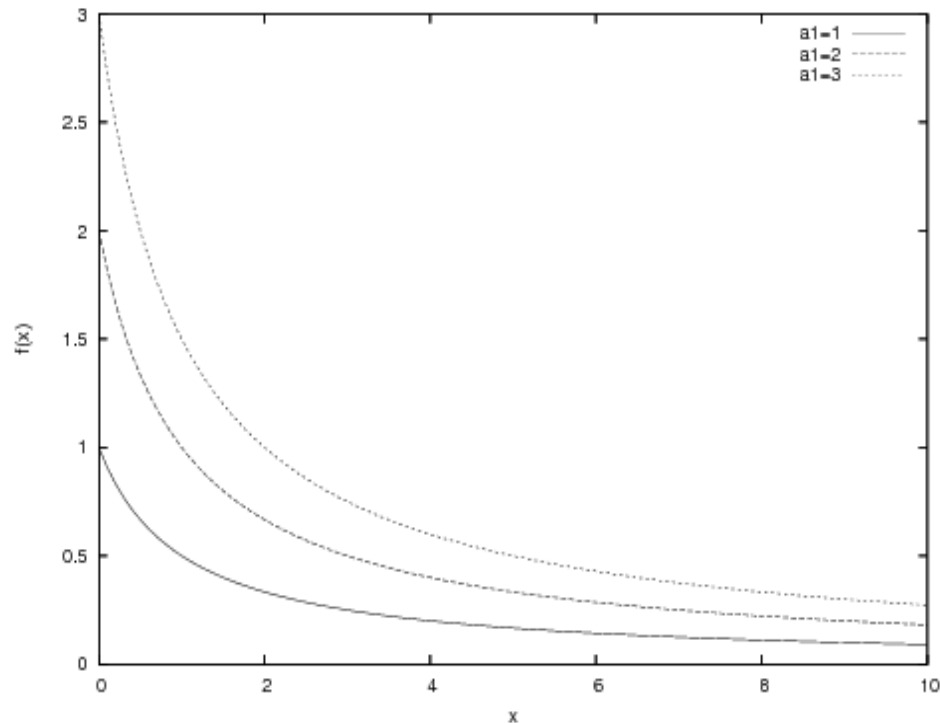
- Mechanism involving x :
 - Residual synthesis
 - Retro-inhibition by y
 - Degradation
 - Auto-activation : not included here for simplicity



Biological switch : x proteins

- Residual synthesis of x and inhibition by y:
 - Classical Hill function:

$$dx/dt = a1 / (1 + y^{b1})$$



Biological switch : x proteins

- Degradation:

$$dx/dt = - c1.x$$

- Summing all the terms:

$$dx/dt = a1 / (1 + y^{b1}) - c1.x$$

The diagram illustrates the components of the differential equation for dx/dt . A horizontal line branches into three paths: one pointing to a red box containing the term $a1 / (1 + y^{b1})$, another pointing to a green box containing the term $- c1.x$, and a third pointing to a green oval labeled 'degradation'. A fourth path, branching from the line between the red and green boxes, points to a red oval labeled 'synthesis'.

- Same reasoning for y :

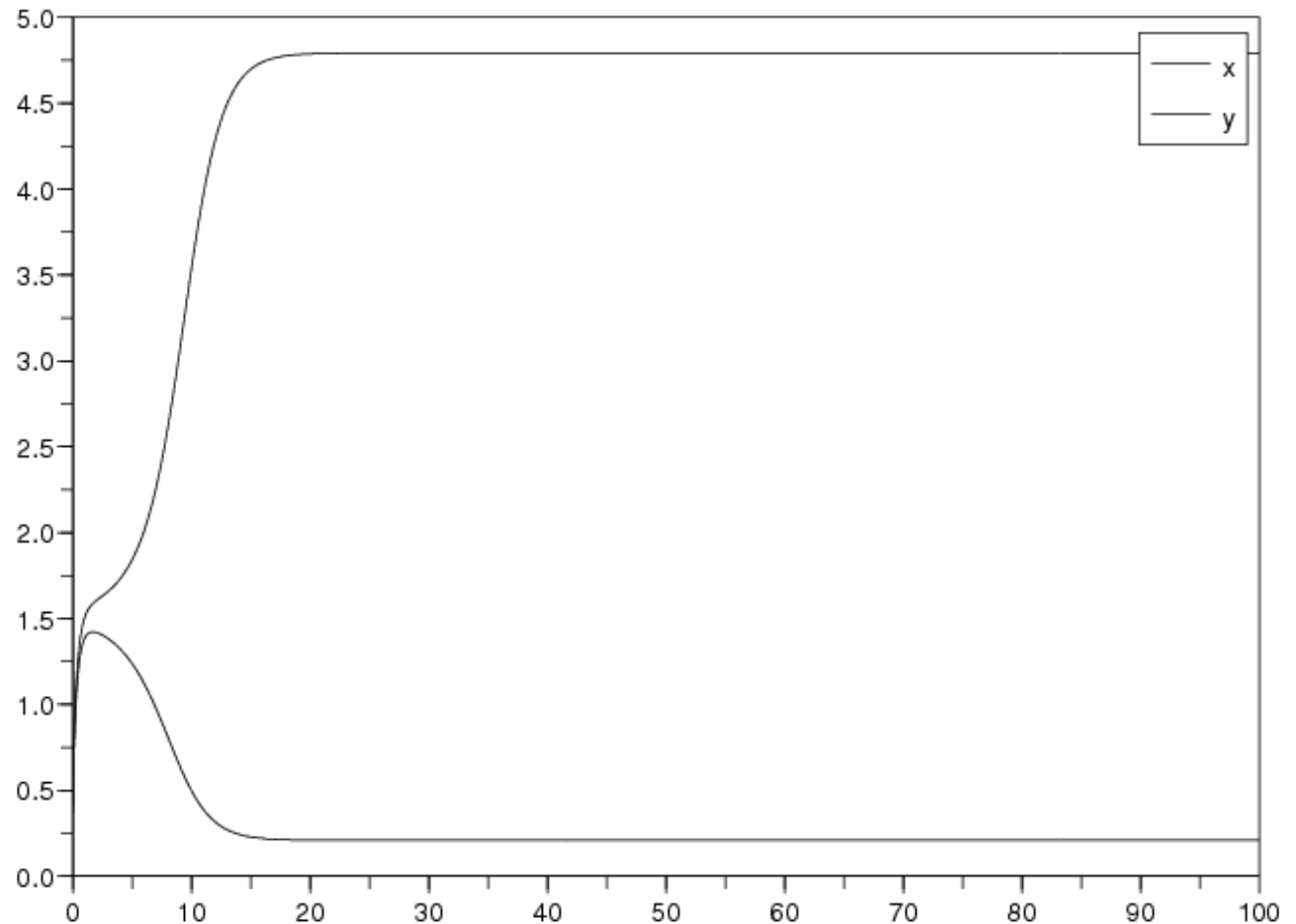
$$dy/dt = a2 / (1 + x^{b2}) - c2.x$$

Time evolution of the model: example

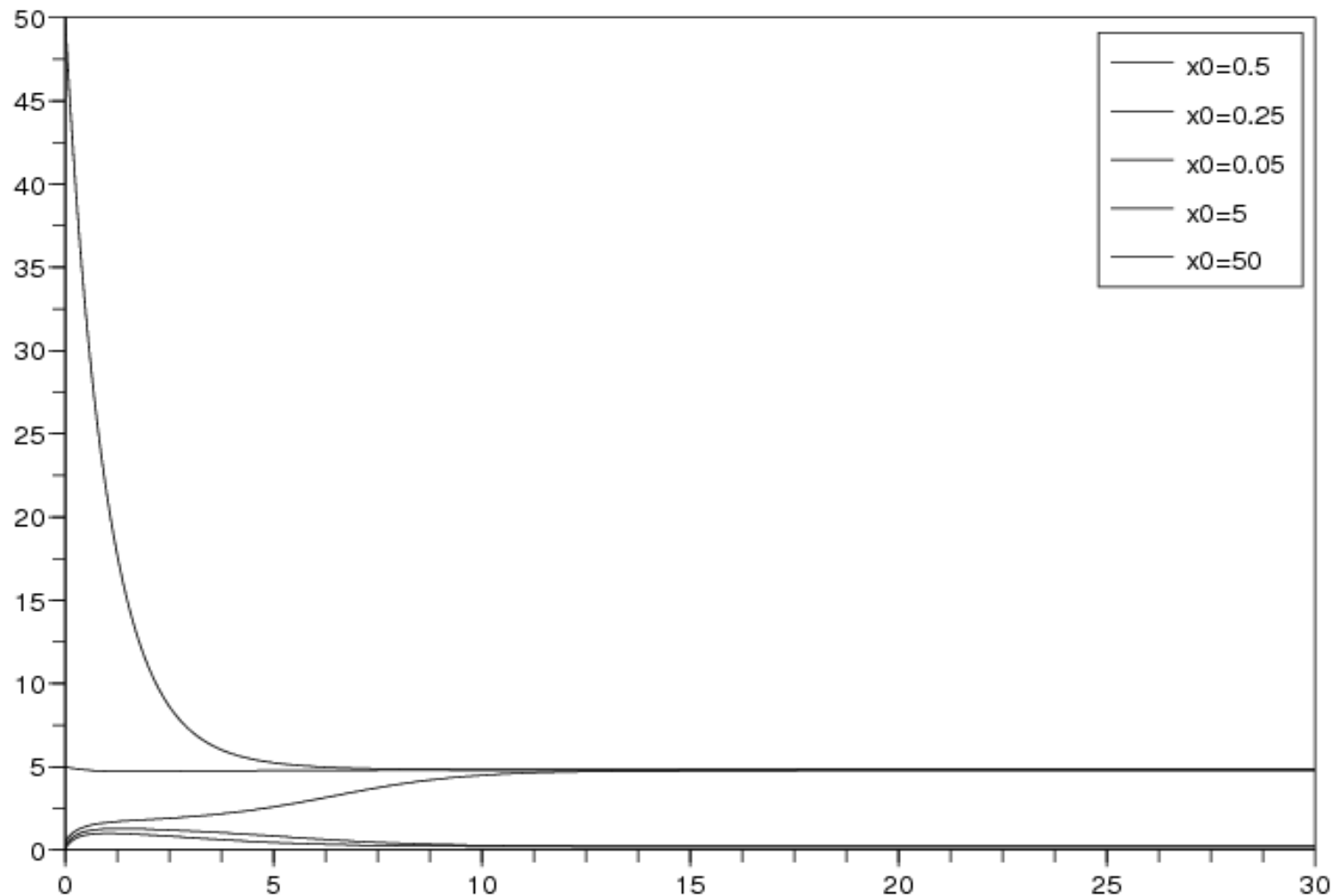
$$\begin{aligned} dx/dt &= f(x,y) \\ dy/dt &= g(x,y) \end{aligned}$$

$x(t=0), y(t=0)$: initial conditions
dt: integration step
Tmax: time of simulation

$x(0), x(dt), x(2.dt), \dots x(t), x(t+dt), \dots, x(t_{\max})$



Time evolution of x : different initial conditions



Equilibrium
state

x do not change
in time

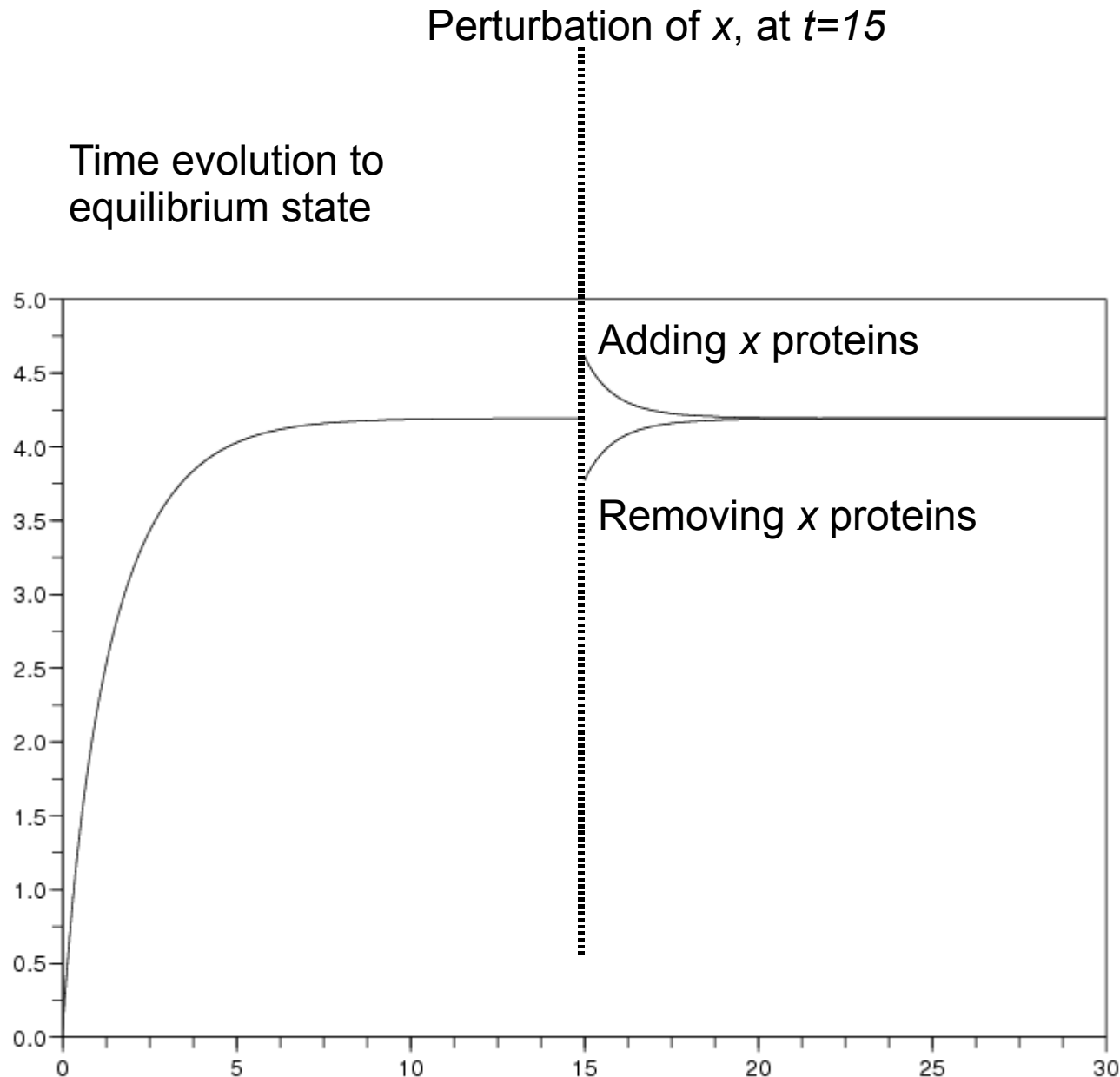
From biology to ODEs: equilibrium state

- Biological fact:
 - the concentration of the x (and y) proteins do not change in time
- ODEs « translation »:
 - A small variation of the concentration, dx (and dy), over a small time interval, dt , is null.

Equilibrium state $\Leftrightarrow dx/dt=0$ and $dy/dt=0$

- Analytically: $f(x,y)=0$ / $g(x,y)=0$ (*Fixe point*)
- What about a small perturbation of this point?

Stability of the equilibrium state?



The concentration of the x protein return to its equilibrium state

The equilibrium state is ***stable***.

Geometrical representation of ODEs

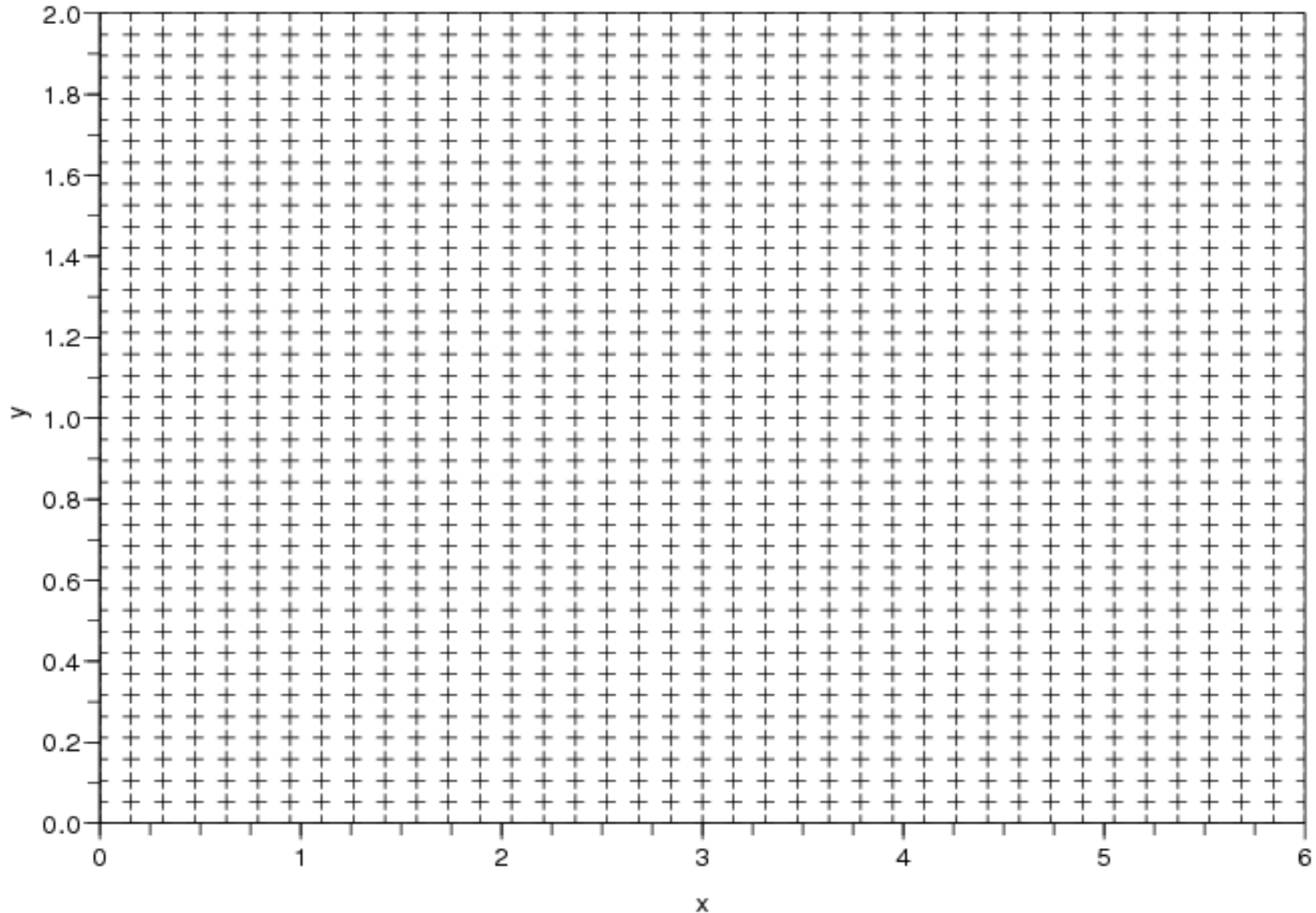
- All analytical calculus can be formulated into geometry
- Geometrical representation is « visual » and more intuitive
- We restrict this tutorial to the geometrical aspects in order to present notions of ODEs.

Geometrical representation of ODEs: Phase Plane

- We adopt a numerical point of view here
- Select a domain of values for x and y
- Select a space of n coordinates:
 - one coordinates for each variables ($n=2$: x and y)
- Each couple $(x(t), y(t))$ represent the state of the system at time t
- A point this space = the state of the system

PHASE SPACE

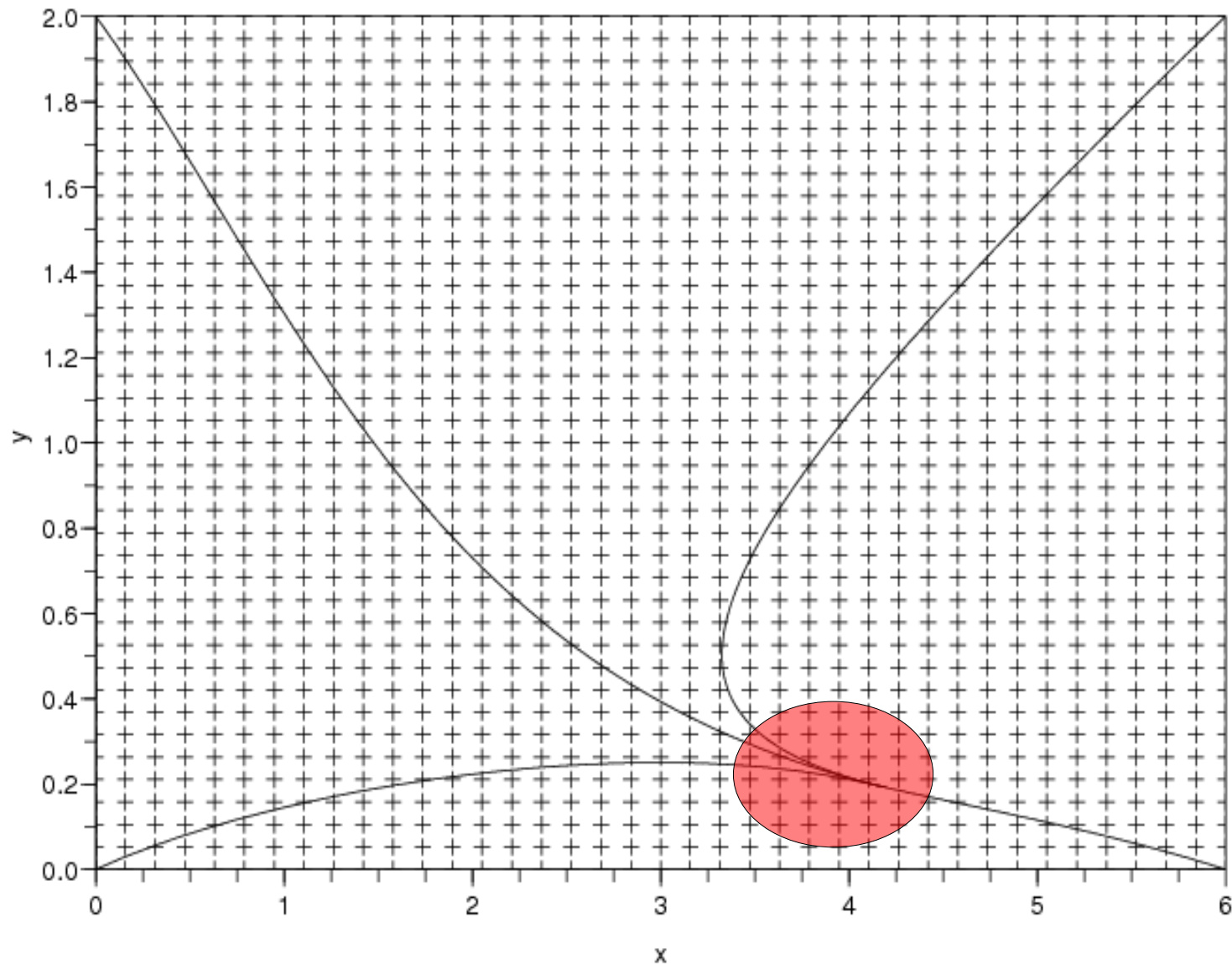
Geometrical representation of ODEs: Phase Plane



Phase trajectory

- Starting for one initial condition, one can compute the sequence of $x(t)$ and $y(t)$
- This is a sequence of states = a sequence of point in the phase plane
- Phase trajectory

Phase trajectory: 4 examples



Nullclines

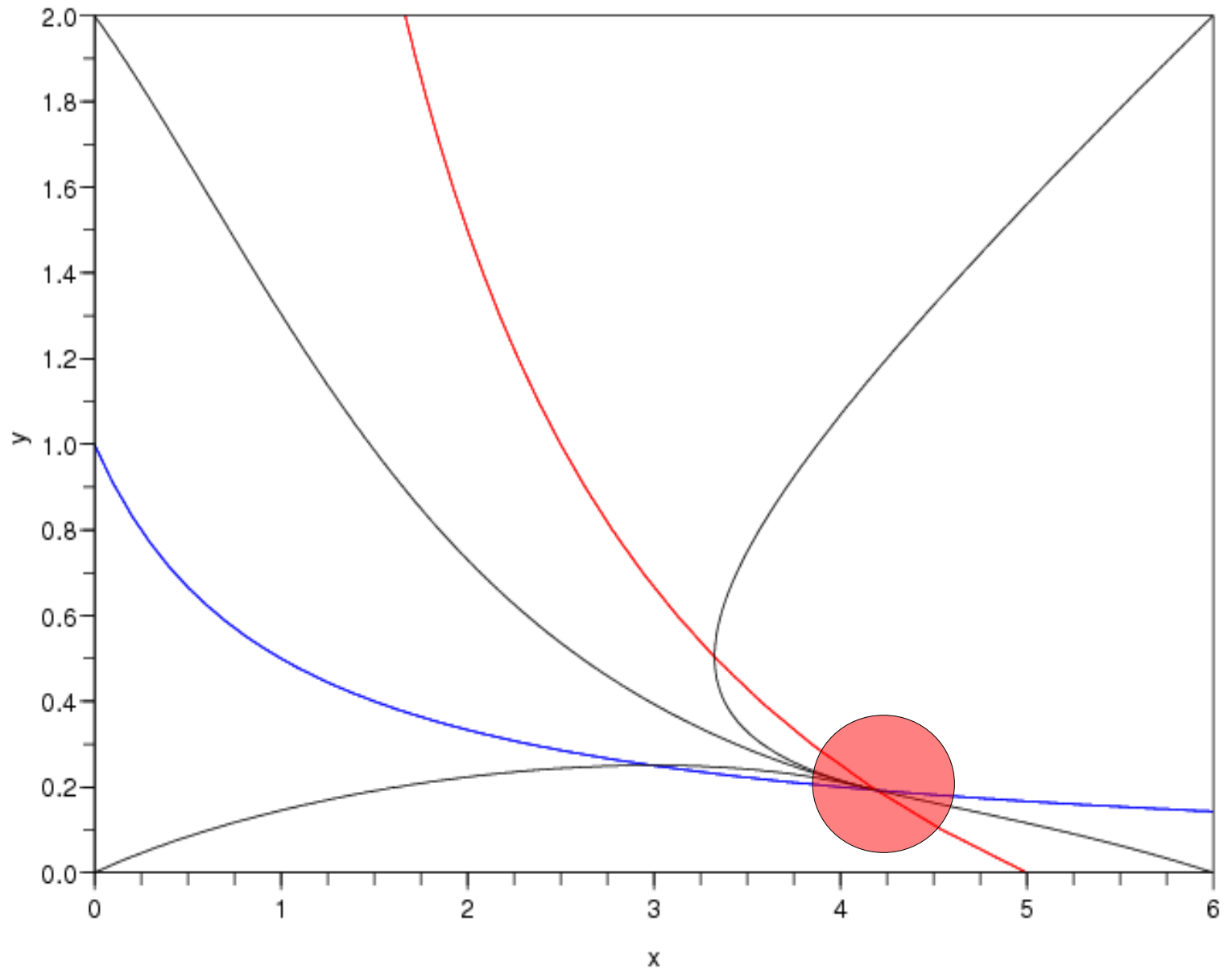
- We show an equivalence:
 - equilibrium state $\Leftrightarrow dx/dt=0$ and $dy/dt=0$
- We can draw each couples (x,y) where $dx/dt=0$
 - x-nullcline
- Same for y : y-nullcline
- The intersection points of x- and y-nullclines defines a equilibrium point.
- Geometrical criteria = biological fact

Nullclines

Red: x-nullcline

Blue: y-nullcline

Black: phase traj.

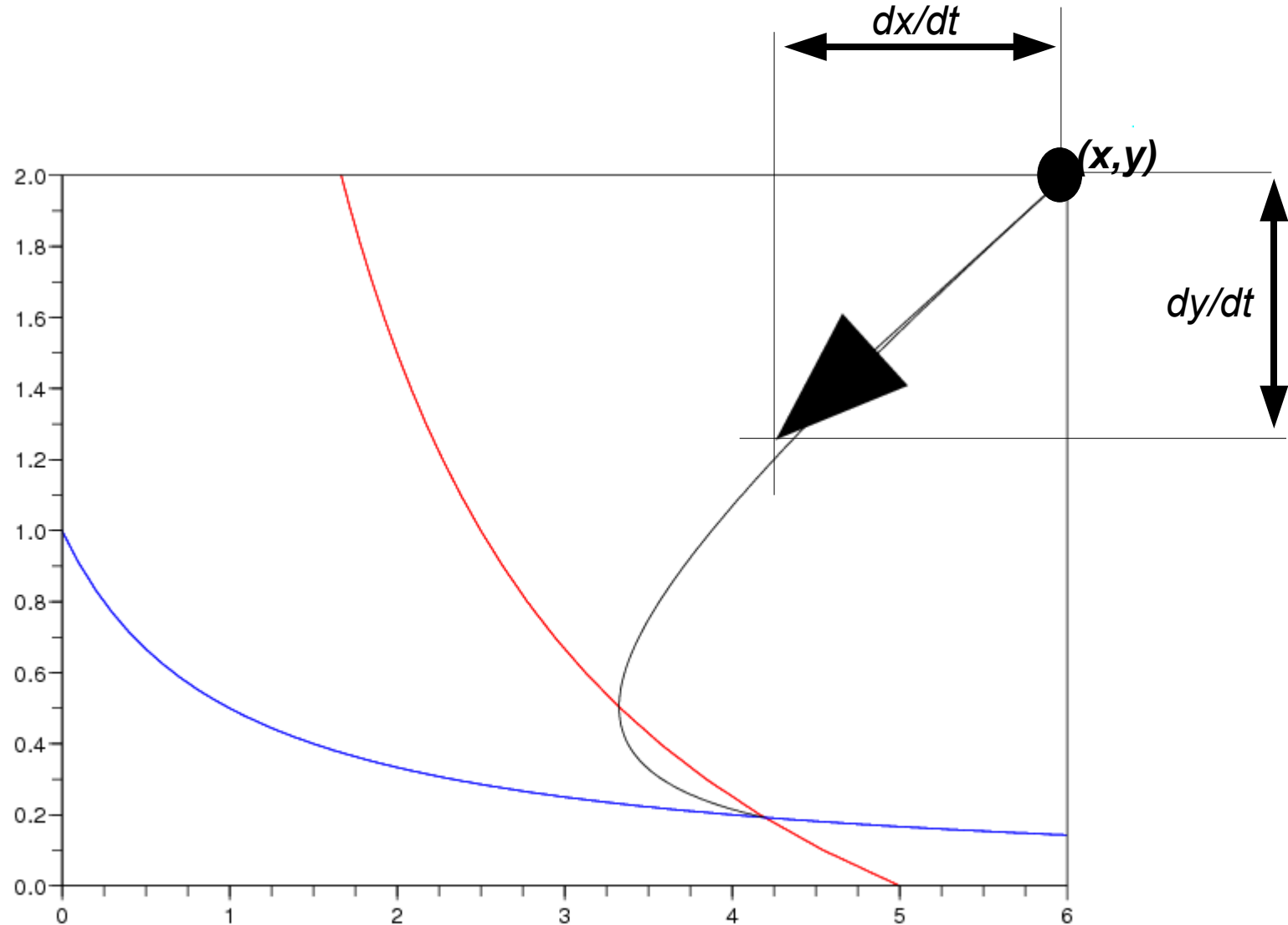


Vector field

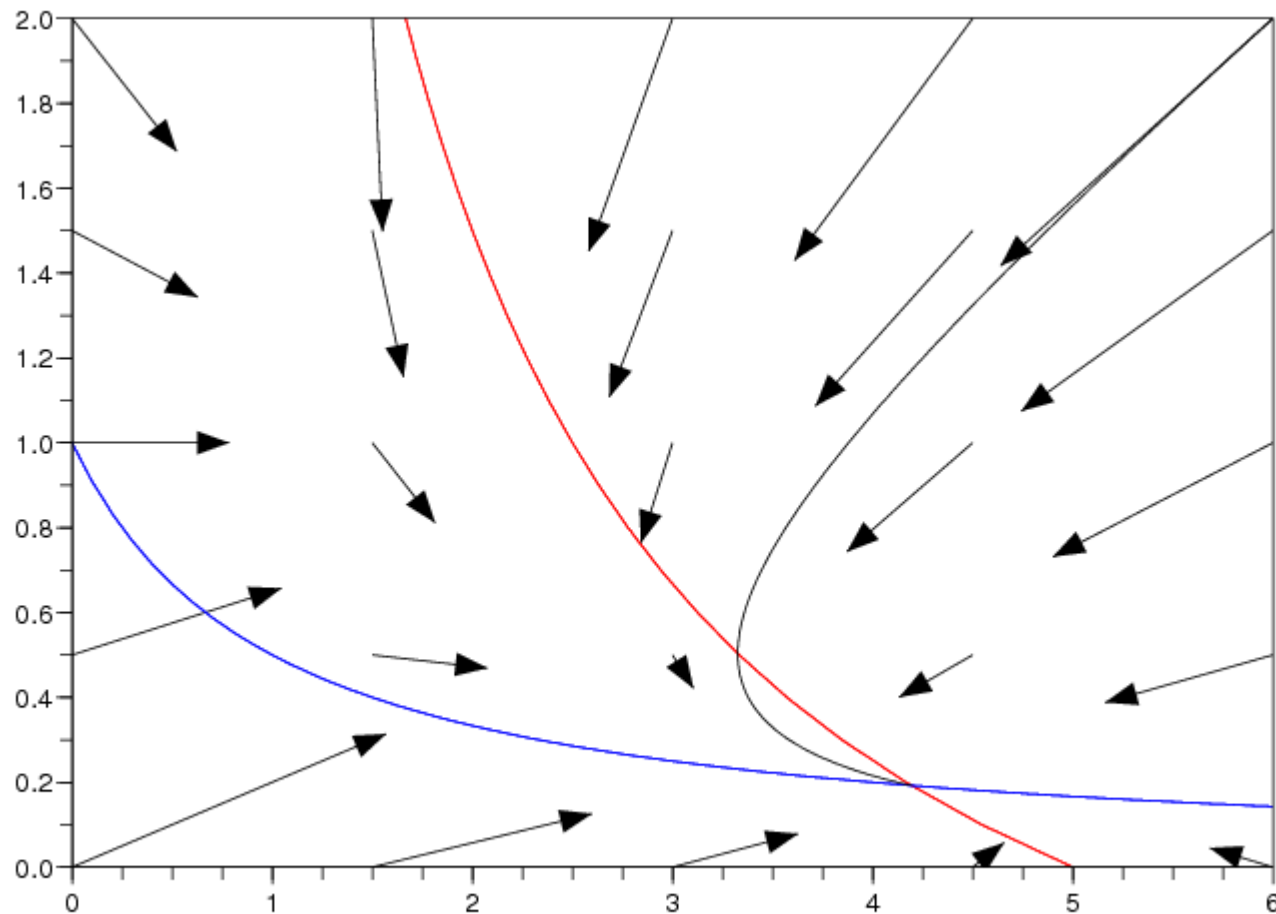
- Last informations: *derivatives*
- We can compute dx/dt and dy/dt for each couple (x,y)
 - Used (for example) in the numerical integration scheme
- We can represent these two values using an « arrow », *i.e.* a vector (here in 2D)
-
- For each couples (x,y) of the phase space

VECTOR FIELD

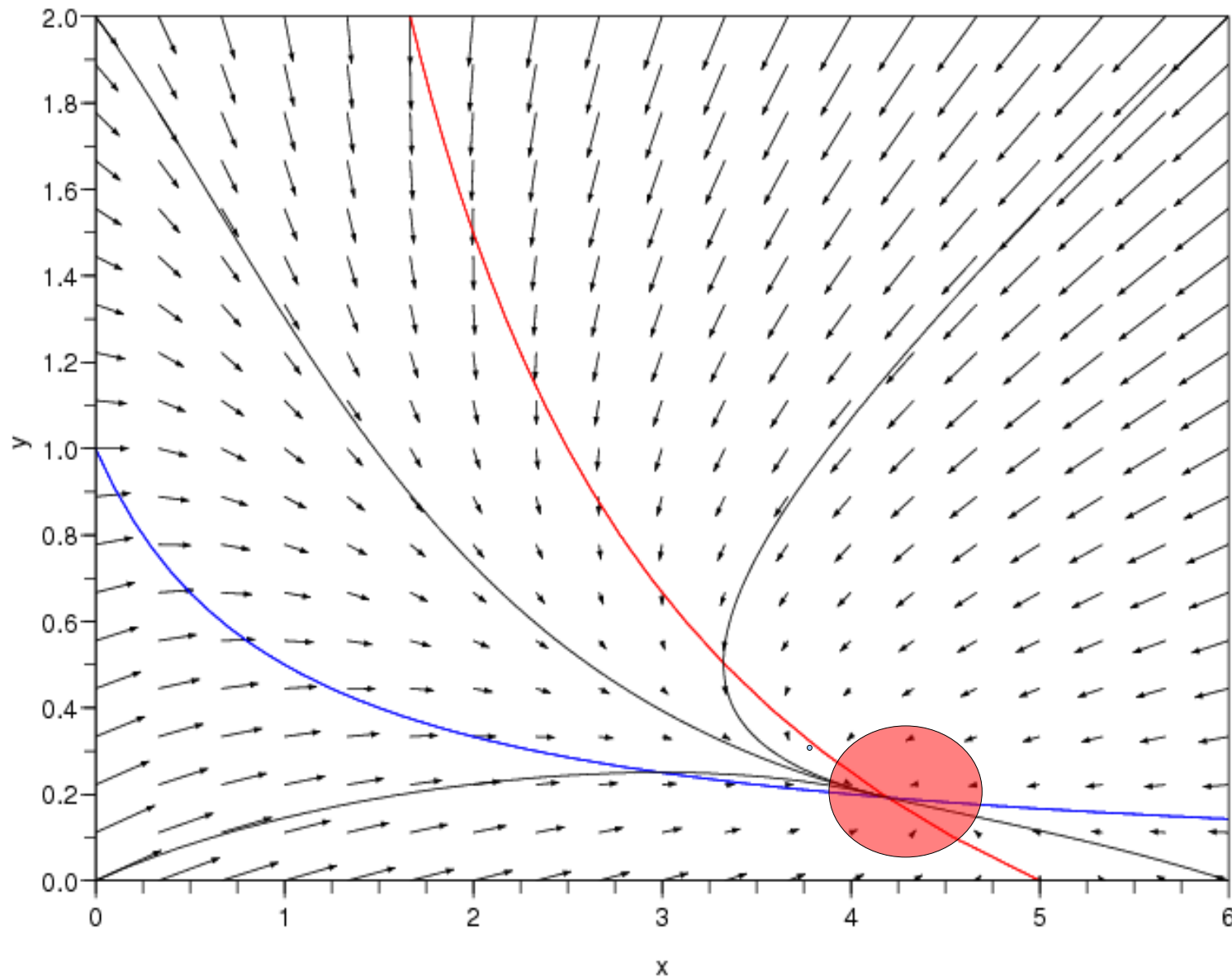
Vector field: example of computation



Vector field: example



Phase plane and vector field

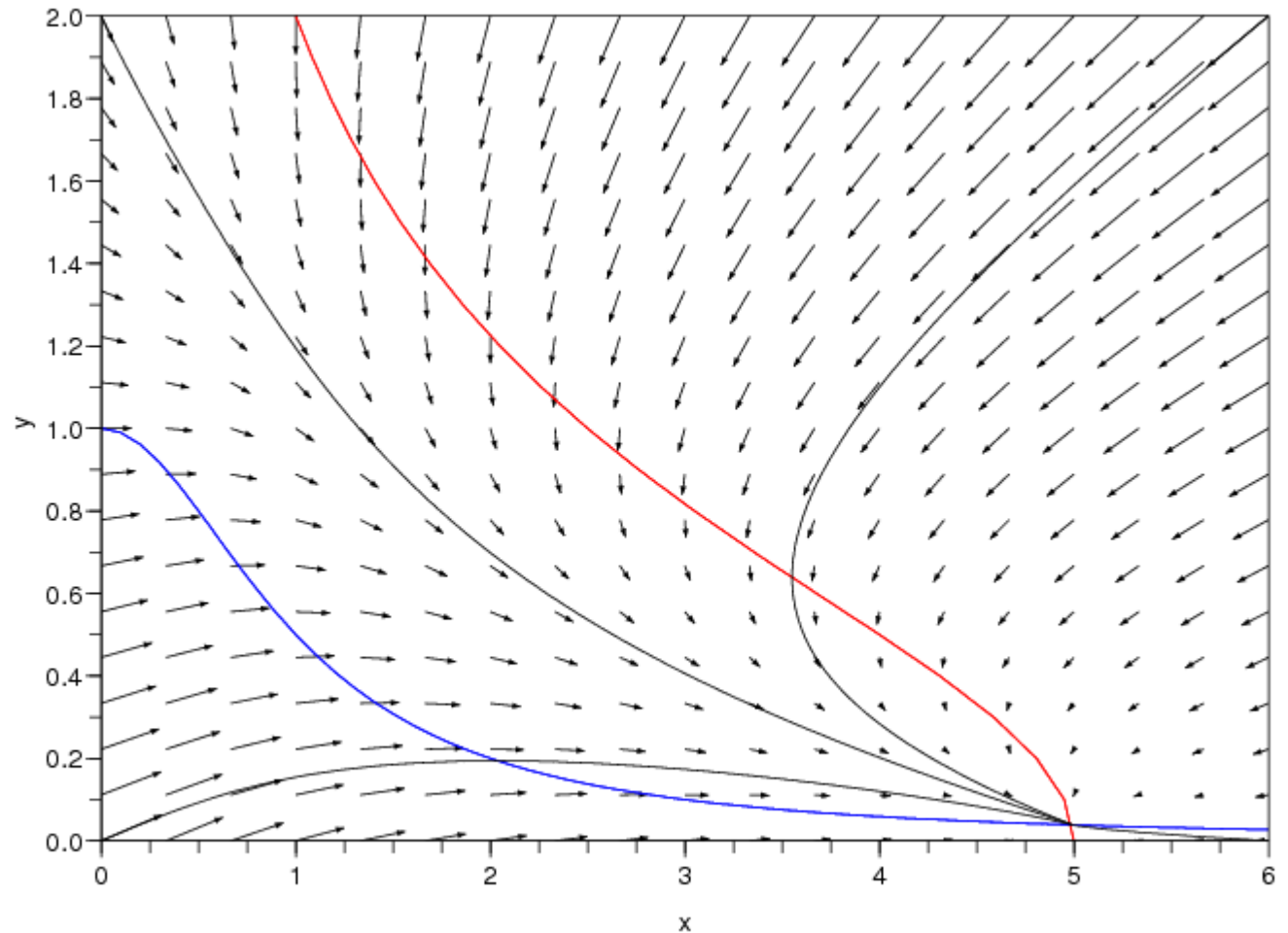
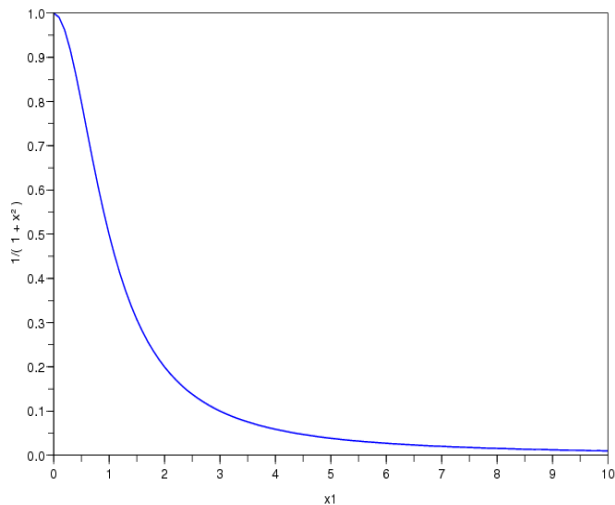


Geometry, ODEs and biology

- Phase plane: plane of each possible states of the system
- Phase trajectory: a time evolution starting from one initial condition
- Nullcline: location where derivatives are null
- Vector field: « *direction/intensity* » of small variations of the proteins concentration over a small time interval

The biological switch and ODEs: cooperativity, $a_2=1$

$$dy/dt = a_2/(1+x^{b_1}) - c_2 y$$



The biological switch and ODEs: cooperativity, $a_2=5$

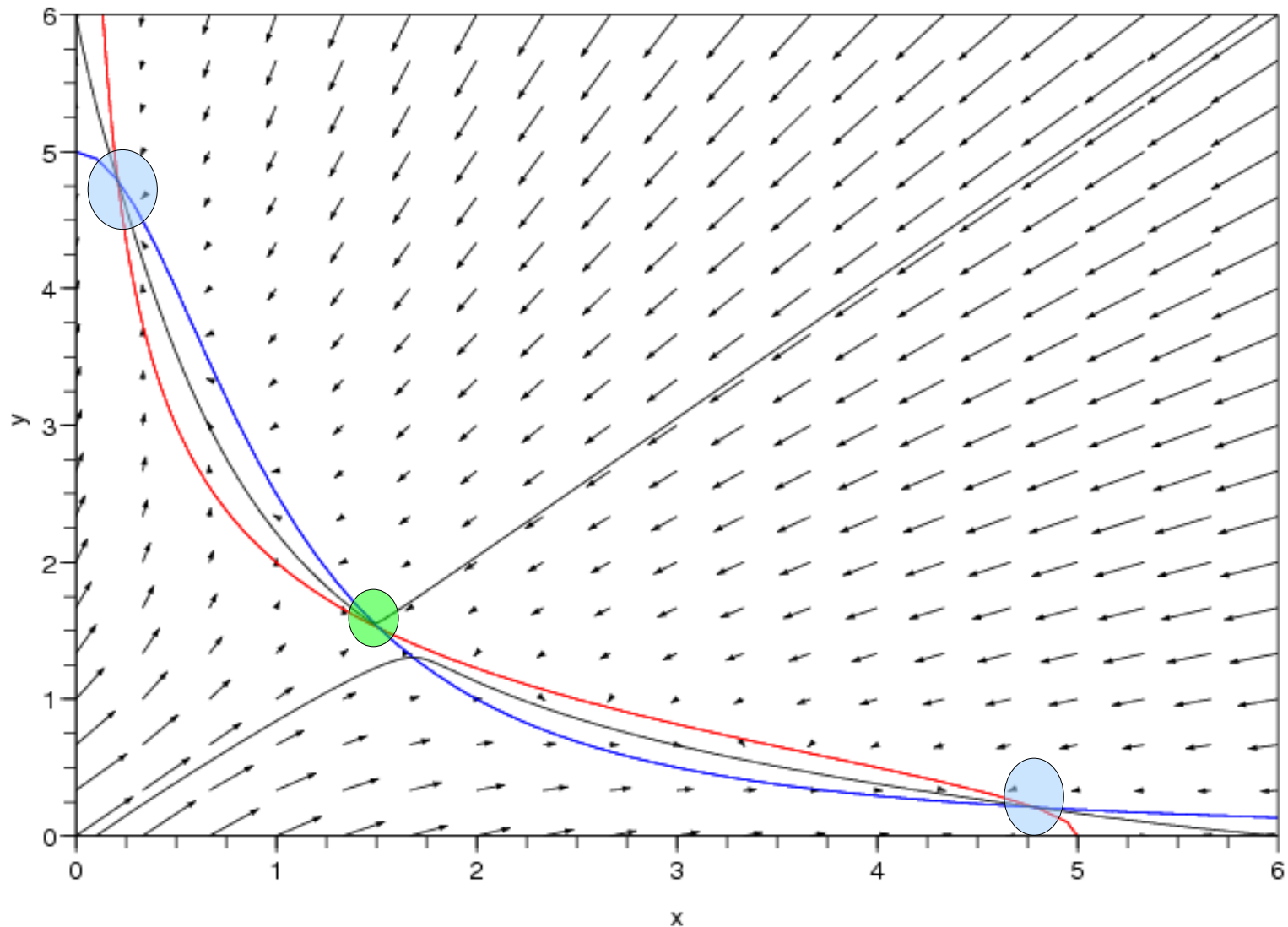
x- and y-nullclines
intersect at 3 points

Three equilibrium states

Two stables

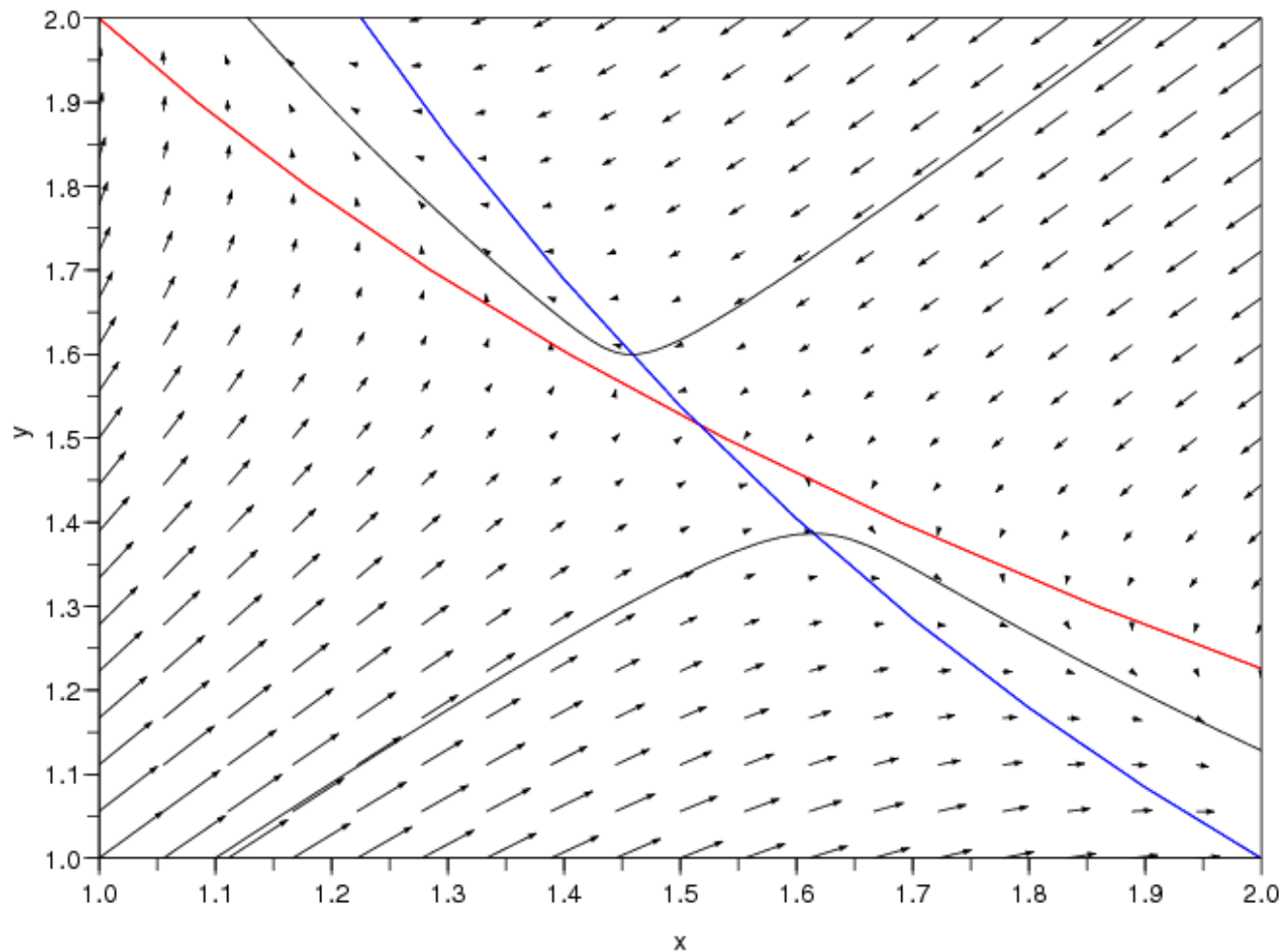


One instable

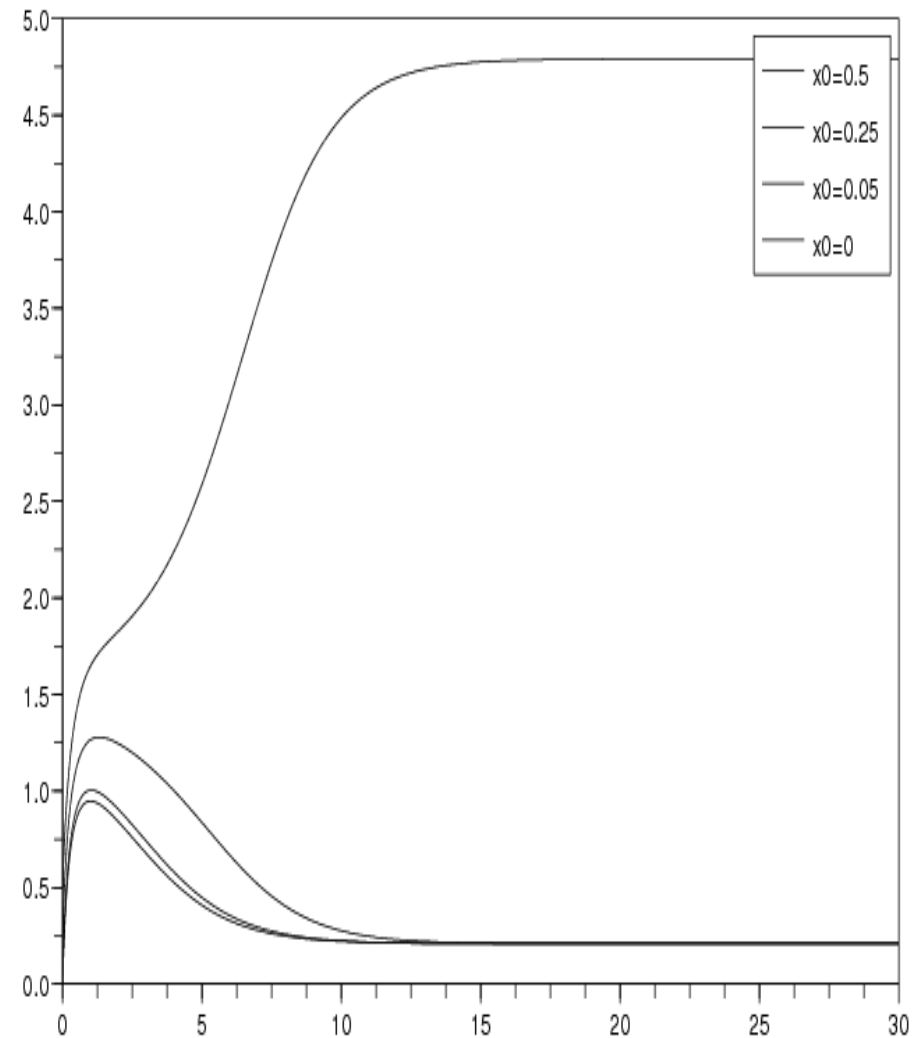
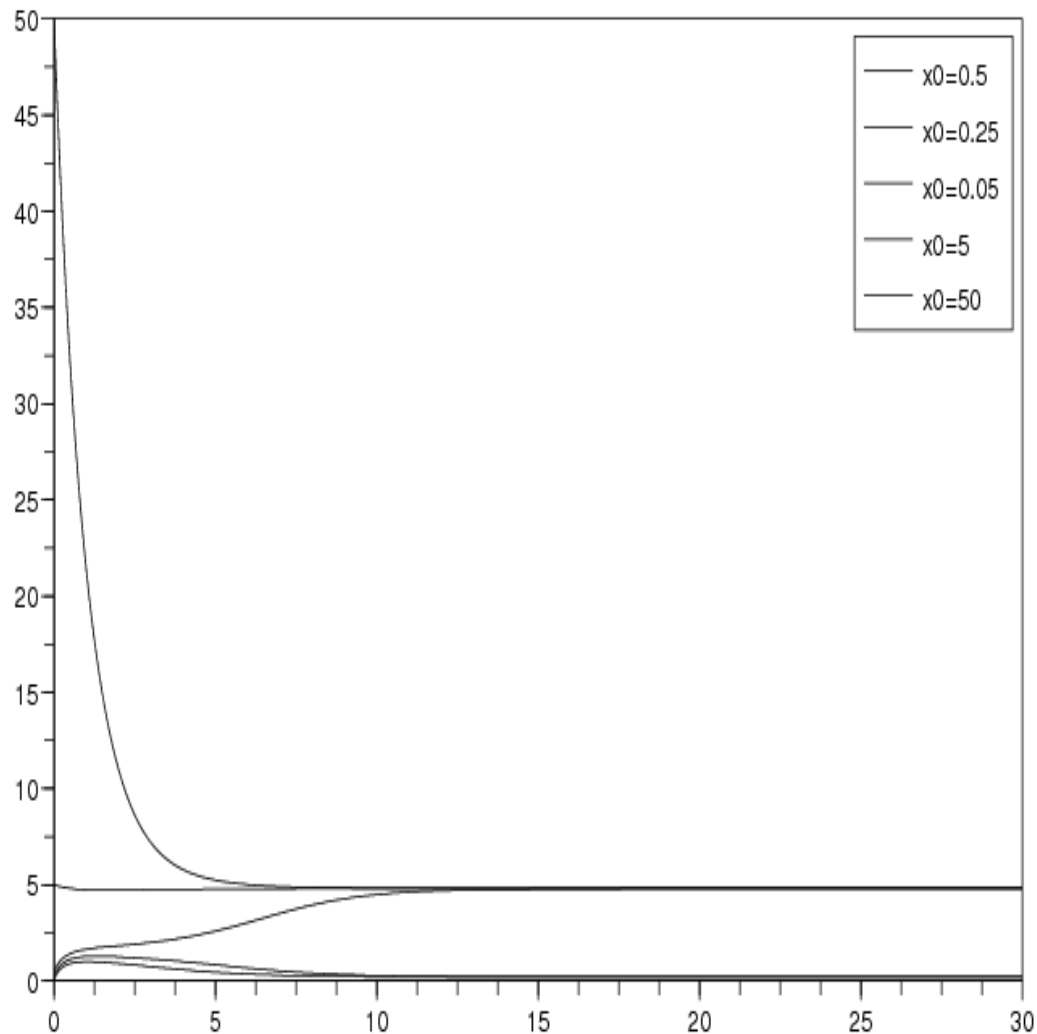


The biological switch and ODEs: cooperativity, $a_2=5$ -ZOOM

Unstable equilibrium state



Dynamical time evolution



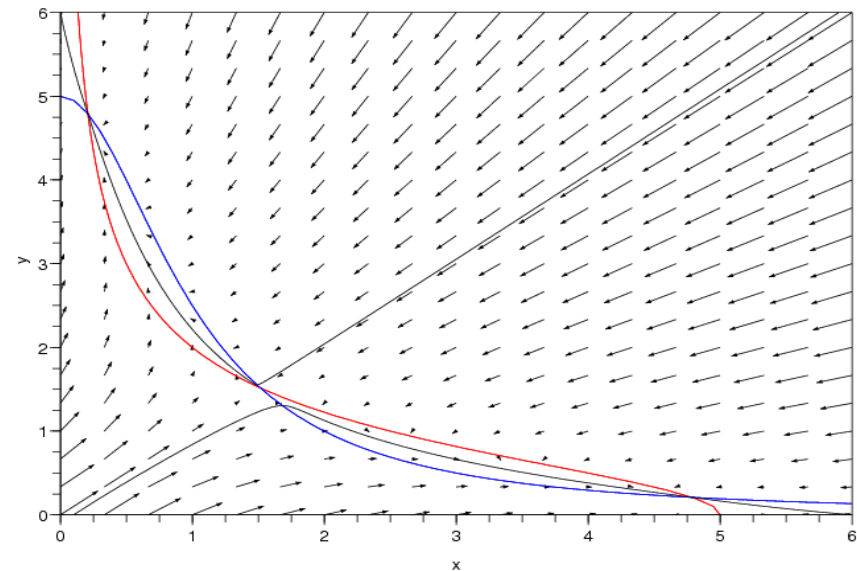
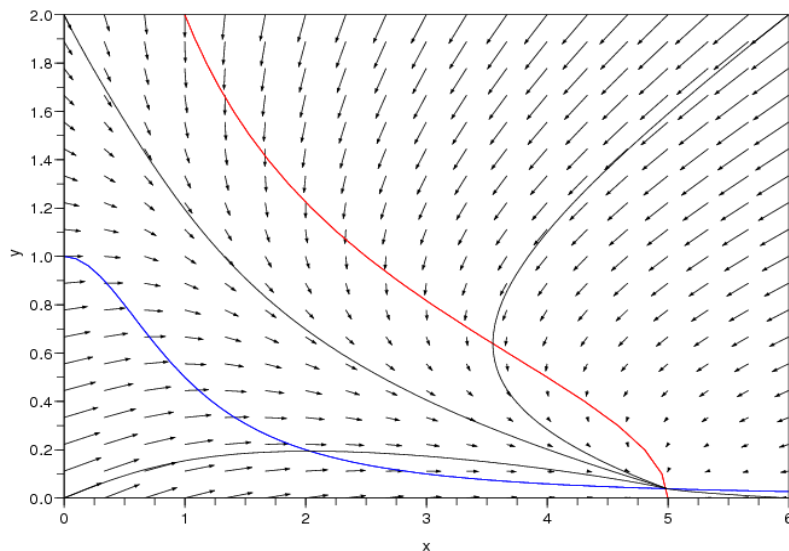
Bistable system: depending on initial conditions = one different equilibrium state

Understanding the biological switch using ODEs notions

- For specific values of parameters, the system has three equilibrium states:
 - Two *stable* equilibrium states : bi-stability
 - One *unstable* equilibrium state
- The unstable equilibrium state is very important:
 - it prevent one stable equilibrium state to switch to the other
 - It divide the phase plane into two domains: one domain (i.e. Initial conditions) is attracted by one point and the other domain is attracted to the other equilibrium point.

Understanding the biological switch using ODEs notions

- Thus, the unstable point has a structural rôle in the dynamic of the system.
- When we change parameters ($a_2: 5 \rightarrow 1$), the nullclines intersect only once: the unstable point disappear, and the system had only on equilibrium state.



Understanding the biological switch using ODEs notions

- The structural change in the geometrical properties reflect a structural change in the dynamical properties of the system.
- The biological switch is then sustained when we switch back and forth, from one geometry (3 equilibrium states) to the other (1 equilibrium state).
- We can hypothesis that if we introduced a « correct » variation of the a_2 parameter, the system will switch between these two situations, and oscillations will occur

Designing a biological switch

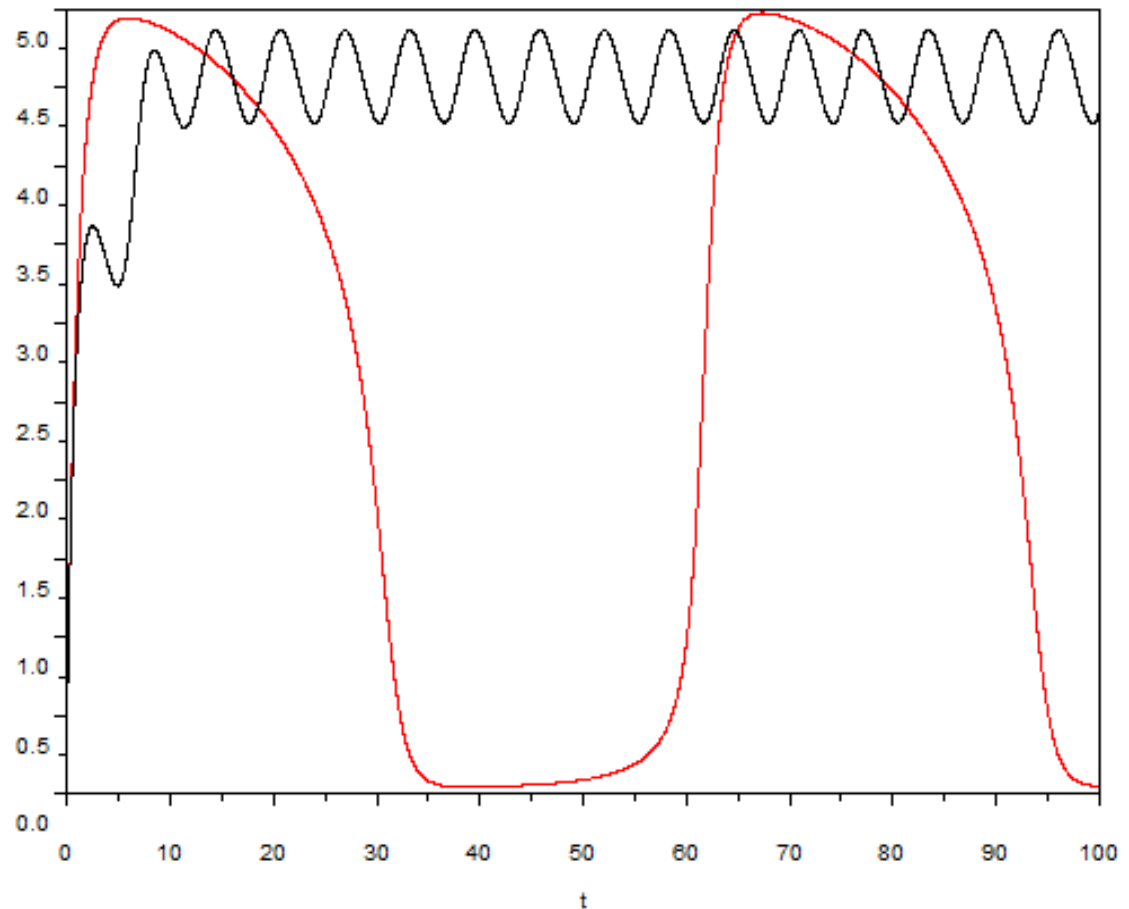
$$dx/dt = a1/(1 + y^2) - c1.x$$

$$dy/dt = a2*\sin(w.t)/(1 + x^2) - c2.y$$

$$a2=15$$

Black: $w=0.5$, fast variations
of parameter $a2$.

Red: $w=0.05$, slow variations
of parameter $a2$



Conclusions

- ODEs are a well developed mathematical formalism
- « Transcribing » a biological system into ODEs open the door to vast mathematical literature, and an active community
- Biologists can develop their model, and do some « *in silico* » experiments thanks to software
- ... provided that they understand fundamental notions
- A geometrical view of an ODEs system is a first step

Conclusions

- ODEs can be extended:
 - Take into account stochastic variations in the derivatives: Stochastic Differential Equations
 - Take into account space: Partial Differential Equations
- For biological applications:
 - Constructing biological switch: Gardner et al. (2000)
 - Understanding transition during Cell Cycle: Tyson and Novak.