# Systems and models in chronobiology A delay model for the circadian rhythm

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24th Benelux Meeting on Systems and Control Houffalize, 22-24 March 2005

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## Outline

#### Introduction

State of the art in mathematical modeling of the circadian rhythm Biological modeling for the circadian oscillator Results

Systems viewpoint: Why is it useful?

Application to the analysis of the circadian rhythm

Conclusion



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# Circadian rhythms (1) Definition and roles

## Definition

Biological rhythms with a period  $au \sim$  24h

### Roles

Circadian rhythms control

- Sleep
- Muscular activity and metabolism
- Food ingestion

► ...

Moreover, there are possible links with various pathologies



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# Circadian rhythms (2) Properties

Property	Comment
Ubiquitous	All eukaryotes and some prokaryotes
Entrainment	Zeitgeber=light/temperature cycles
Genetic	Single-gene clock
mechanisms	mutants isolated
Precision	$\Delta au < 0.1\%$
Robustness	T°, IC
	A cell-autonomous circadian oscillator
Cellular nature	mechanism exists and appears to be a fundamental
	unit even among multicellular organisms



Systems and models in chronobiology —State of the art in mathematical modeling of the circadian rhythm

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└─ Biological modeling for the circadian oscillator

Simple model Biochemical principle



from Goldbeter, *Biochemical oscillations and cellular rhythms*, Cambridge University Press, 1996



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Biological modeling for the circadian oscillator

# Simple model Equations

$$\dot{M} = v_s \frac{K_l^n}{K_l^n + P_N^n} - v_m \frac{M}{k_m + M}$$

$$\dot{P}_0 = k_s M - V_1 \frac{P_0}{K_1 + P_0} + V_2 \frac{P_1}{K_2 + P_1}$$

$$\dot{P}_1 = V_1 \frac{P_0}{K_1 + P_0} - V_2 \frac{P_1}{K_2 + P_1} - V_3 \frac{P_1}{K_3 + P_1} + V_4 \frac{P_2}{K_4 + P_2}$$

$$\dot{P}_2 = V_3 \frac{P_1}{K_3 + P_1} - V_4 \frac{P_2}{K_4 + P_2} - k_1 P_2 + k_2 P_N - v_d \frac{P_2}{k_d + P_2}$$

$$\dot{P}_N = k_1 P_2 - k_2 P_N$$

- 5 nonlinear ODEs
- 17 parameters



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## Evolutions of the model

### Drosophilia



- ▶ 10 nonlinear ODEs
- ► 34 parameters



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# Evolutions of the model

## Drosophilia



- ▶ 10 nonlinear ODEs
- ► 34 parameters

### Mammals



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- ▶ 19 nonlinear ODEs
- 59 parameters



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└─State of the art in mathematical modeling of the circadian rhythm

Results

Results (1) Oscillations and limit cycle

### Oscillations



### Limit Cycle





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Results (2)Entrainment





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Results

Results (3) Bifurcation diagram





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Systems and models in chronobiology └─Systems viewpoint: Why is it useful?

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Systems and models in chronobiology └─Systems viewpoint: Why is it useful?

# Limitations of current mathematical modeling

- Very complicated systems (strongly nonlinear, several variables and parameters)
  - $\rightarrow$  restricted to numerical simulations
- Impossible to test all parameter combinations
- Mathematically (and computationally) difficult to study interconnections
- Exhibit some properties, but NOT explain them



Systems and models in chronobiology — Systems viewpoint: Why is it useful?



Developing or using system models and tools for the analysis of biological system (and particularly oscillatory systems)



Systems and models in chronobiology └─Systems viewpoint: Why is it useful?

# System approach Links with system question

By nature, cells are open systems, i.e. with *inputs* and *outputs* Many unexplained properties are related to fundamental systems questions:

- Why is there an entrainment?
- Why is the system robust to certain parameters variation (T°, initial conditions...)? Why is it robust to molecular noise? And to external disturbance?
- How can we guess from observed properties, the values of unknown parameters?
- Why is there a synchronization in networks of oscillators? What kind of synchronized behaviour may we expect given a particular network configuration?



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# Recent approaches

## Classical methods for the analysis of limit cycles <u>Limitations</u>: either useless in high dimension (and so for interconnections) or unable to handle global treatment

#### Abstract models

<u>Limitations</u>: Acts more as a "black box" i.e. it is possible to obtain qualitative information, but no quantitative one.

### Development of new methods

- Monotone Systems (P. de Leenheer, D. Angeli, E. Sontag)
- Piecewise Linear Systems PLS (J. Goncalves)
- Dissipative Systems (R. Sepulchre, G.B. Stan)
- = dedicated I/O approaches



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dash Application to the analysis of the circadian rhythm

### A delay model for the circadian rhythm Variation on Goldbeter's model: 2 variables



from T. olde Scheper and al., A Mathematical Model for the Intracellular Circadian Rhythm Generator, J. of Neuroscience, 1999



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# A delay model for the circadian rhythm Further simplification : 1 variable



$$\dot{M} = v_s \frac{K_I^n}{K_I^n + M(t-\tau)^n} - v_m M$$

This model is equivalent to the famous Mackey and Glass' model for Red Blood Cell regulation



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## Conclusion and perspectives

There remains many **unsolved questions** in the exploding field of mathematical biology.

Most of these are **natural** in the framework of **systems modeling**: synchronization, robustness, entrainment...

Our research group currently develops methods based on an input/output approach to answer (some of) those questions and **classify** basic oscillatory mechanisms.

We are tempting to analyze a new mechanism based on a **delay** and a specific **regulatory nonlinearity** in the feedback loop.



Systems and models in chronobiology └─ Conclusion

# Thank you

Thank you for your attention... :-)

