

# Biochemical Frequency Control by Synchronisation of Coupled Repressilators

## An *In-silico* Study of Modules for Circadian Clock Systems

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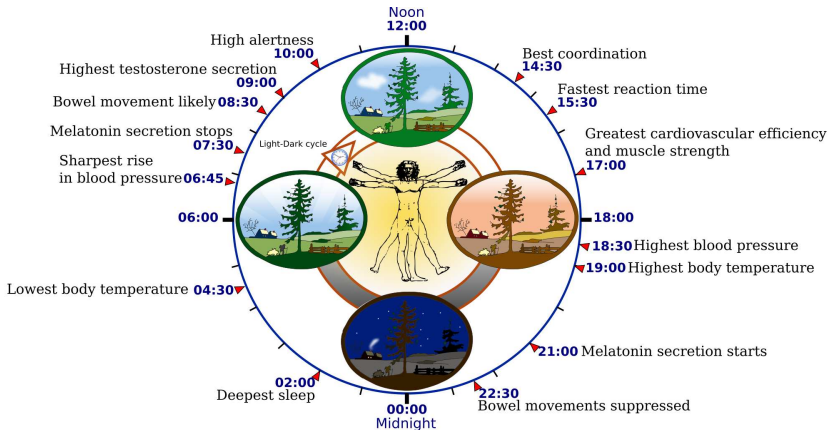
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Modelling Oscillatory  
Information Processing Group

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# Human Daily Rhythm: Trigger and Control System



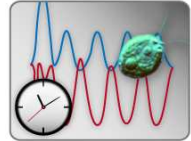
[www.wikipedia.org](http://www.wikipedia.org)

# Biological Clocks

## Significance

- **Nanoscaled oscillatory reaction systems**
- High precision and self-sustainability
- Robust and reliable control systems for manifold processes
- Adaptability to specific environmental conditions (e.g. cycles of light/darkness)
- Infradian (period  $> 1$  day), *circadian* ( $\approx 1$  day), and ultradian ( $< 1$  day) rhythms
- Several independent evolutionary origins
- Prototypes for fine-grained clock synchronisation
- Medicine, agriculture, bionics, material sciences, biology

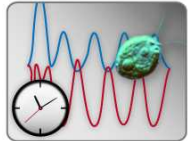
⇒ Keeping environmental time within living organisms



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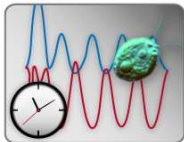


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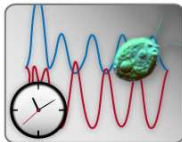


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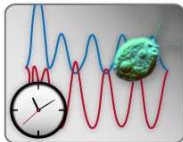


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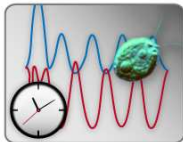


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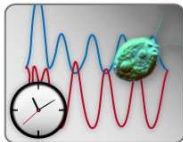
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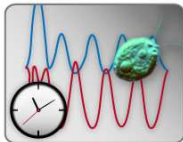


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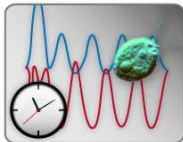


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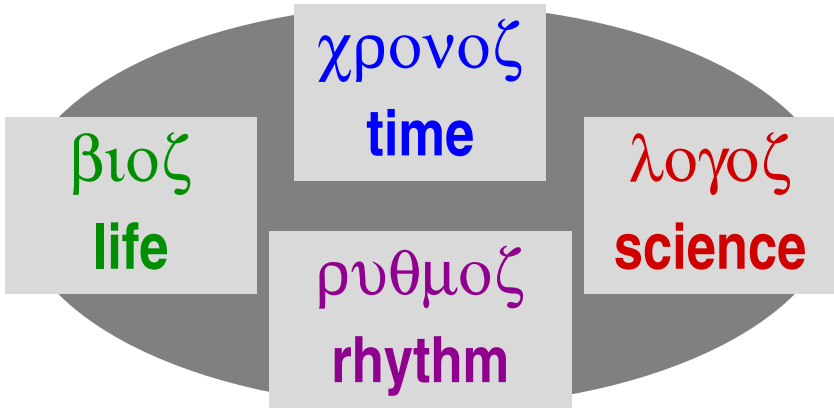
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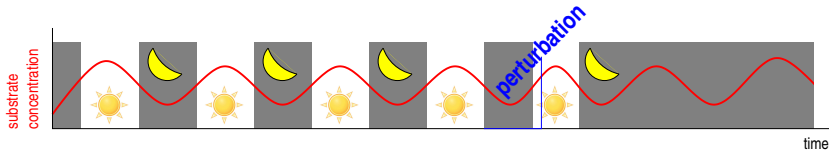
# Chronobiology



**science of biological rhythms and clock systems**

# Circadian Clock

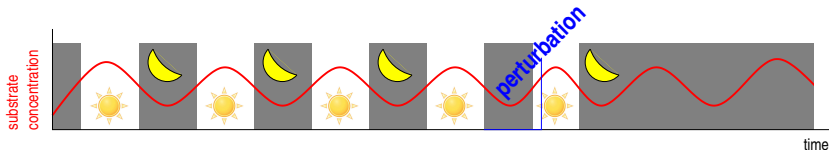
- Undamped biochemical oscillation
- Period *approx.* 24 hours persisting under constant environmental conditions (e.g. permanent darkness DD or permanent light LL)
- **Entrainment** – adaptation to external stimuli (e.g. light-dark cycles induced by sunlight)
- Temperature compensation within a physiological range
- Reaction systems with at least one feedback loop



⇒ Biological counterpart of frequency control system

# Circadian Clock

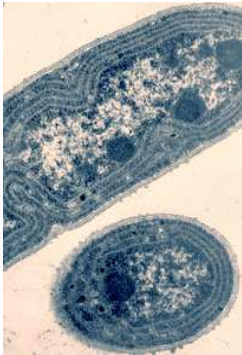
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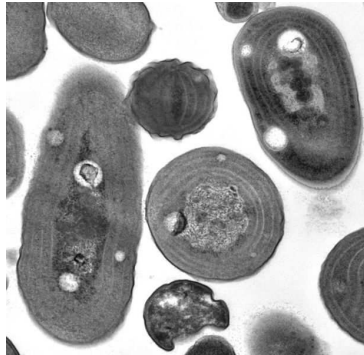
⇒ **Biological counterpart of frequency control system**

# Cyanobacterium *Synechococcus elongatus*

“Simplest and earliest cells known to exhibit circadian phenomena”



[www.genome.jgi-psf.org](http://www.genome.jgi-psf.org)



[www.wikipedia.org](http://www.wikipedia.org)

- Prokaryotic autotrophic picoplankton in tropical seas
- Assumed to be on earth for more than 3.5 billion years
- Clock: Phosphorylation cycle without gene expression

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Specifications for Synchronisation of Oscillatory Signals

# 3. Repressilator

Gene Regulatory Network with Oscillatory Behaviour

# 4. Internal Synchronisation

Simulation Studies using Coupled Repressilators

# 5. External Synchronisation

Frequency Control Systems with Phase-Locked Loop



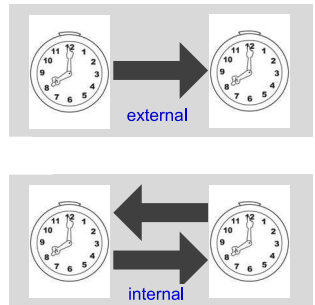
# Entrainment vs. Synchronisation

## Entrainment

- Oscillating signal (frequency, phase, and amplitude) dynamically adapts to (varying) external stimulus. External stimulus itself not influenced.

## Synchronisation

- *External*: Entrainment to external stimulus (e.g. light-dark cycle induced by sunlight) + adaptation to signal shape of external stimulus
- *Internal*: oscillating signals mutually adapt, converge to a common signal

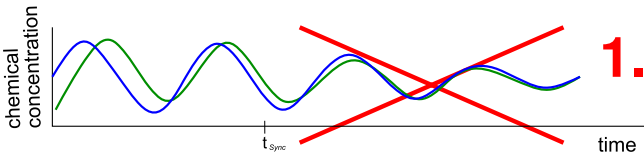


⇒ Entrainment can be seen as special case of synchronisation

# Properties of Synchronous Oscillations (I)

## Undamped oscillations

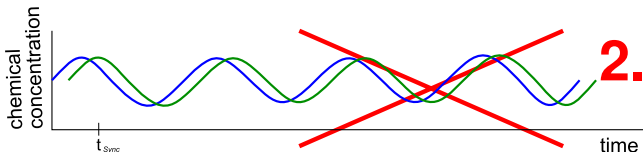
- Modelled oscillation results from solution of ordinary differential equations (ODEs) describing dynamical behaviour of the biochemical clock system
- Eigenvalues of Jacobian matrix (real parts  $< 0$ ) mostly indicate undamped oscillations
- Limit cycles (represented by orbital courses) as method of choice for numerical data



# Properties of Synchronous Oscillations (II)

## Asymptotic or total adaption

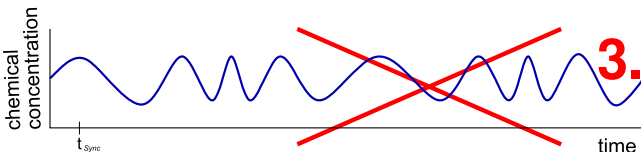
- Harmonisation of oscillating substrate concentration
  - after finite time  $t_{\text{sync}}$  within
  - arbitrarily selectable  $\varepsilon$ -neighbourhood



## Properties of Synchronous Oscillations (III)

### Monofrequential oscillation after $t_{\text{sync}}$

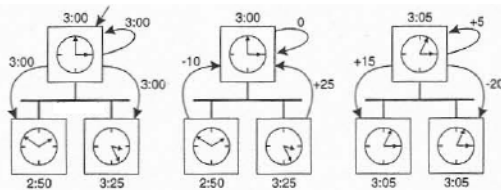
- Fast Fourier Transformation / Fourier analysis (discrete data processing and comparison of peaks)
- Laplace transform and subsequent algebraic processing (preferably for sinusoidal signals)
- Numerical exploration (e.g. sampling)



# Internal Clock Synchronisation: Technical Protocols

## Each node in a bidirectionally coupled computer network

- Comprises a specific clock (potential deviations to others)
- Can communicate with all other nodes by sending/receiving local time stamps
- Requests time stamps from others (mutually exchange)
- Successively adjusts its local clock (Lamport, Cristian, Berkeley algorithms)

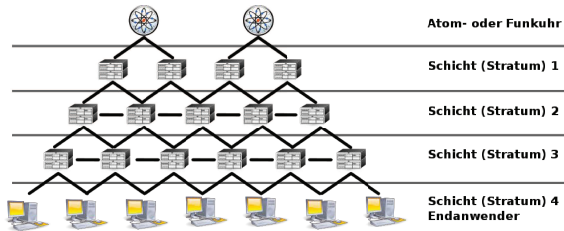


Berkeley algorithm. A. S. Tanenbaum and M. van Steen, *Distributed Systems Principles and Paradigms*, 2001

# External Clock Synchronisation: Technical Protocols

## Each node in unidirectionally coupled computer network

- Comprises a specific clock (potential deviations to others)
- Localised within hierarchial network structure
- Retrieves time stamps exclusively from upper layers (unidirectional signal transduction)
- Successively adjusts its local clock by propagating time stamps from clock(s) in root position



Network Time Protocol (NTP). [de.wikipedia.org/wiki/Network\\_Time\\_Protocol](http://de.wikipedia.org/wiki/Network_Time_Protocol)

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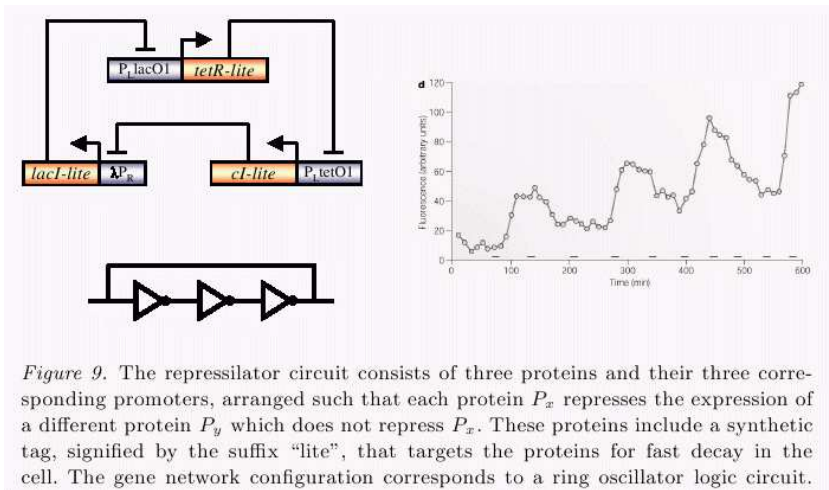
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# Repressilator Prototype

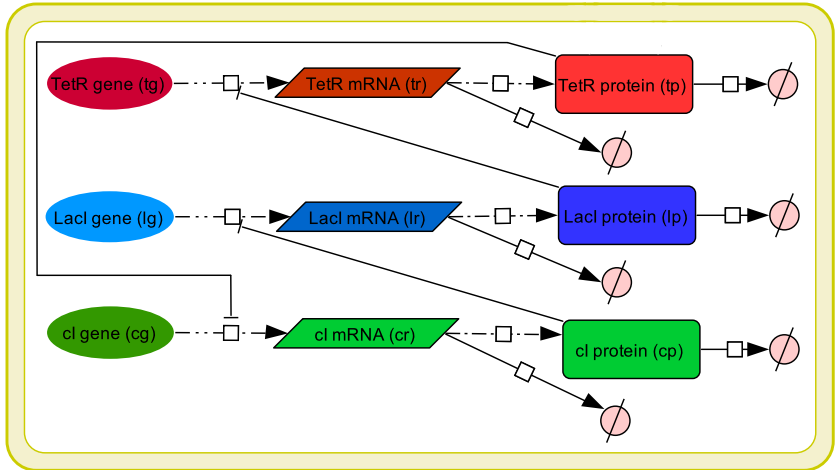
## In-vitro Oscillating Gene Regulatory Network



Eulowitz et al., Nature 403:335-338, 2000



# Repressilator Model: Network Topology



Based on M.B. Elowitz, S. Leibler. A synthetic oscillatory network of transcriptional regulators.  
Nature **403**:335-338, 2000

# ODEs Formalising Repressilator's Dynamic Behaviour

$$\frac{d \text{LacI\_Protein}}{d t} = k_{tl} \cdot \text{LacI\_mRNA} - k_p \cdot \text{LacI\_Protein}$$

$$\frac{d \text{TetR\_Protein}}{d t} = k_{tl} \cdot \text{TetR\_mRNA} - k_p \cdot \text{TetR\_Protein}$$

$$\frac{d \text{cl\_Protein}}{d t} = k_{tl} \cdot \text{cl\_mRNA} - k_p \cdot \text{cl\_Protein}$$

$$\frac{d \text{LacI\_mRNA}}{d t} = a0\_tr + \frac{a\_tr \cdot KM^n}{KM^n + \text{cl\_Protein}} - k_{tl} \cdot \text{LacI\_mRNA} - k_r \cdot \text{LacI\_mRNA}$$

$$\frac{d \text{TetR\_mRNA}}{d t} = a0\_tr + \frac{a\_tr \cdot KM^n}{KM^n + \text{LacI\_Protein}} - k_{tl} \cdot \text{TetR\_mRNA} - k_r \cdot \text{TetR\_mRNA}$$

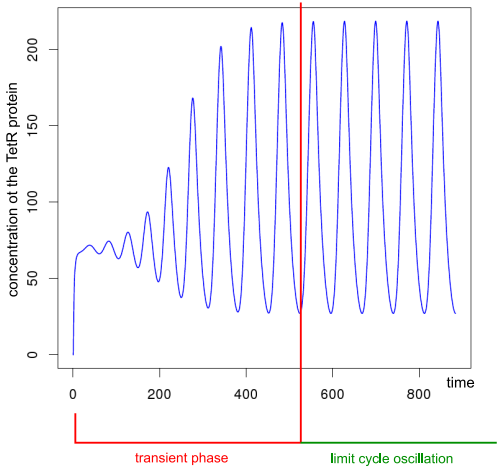
$$\frac{d \text{cl\_mRNA}}{d t} = a0\_tr + \frac{a\_tr \cdot KM^n}{KM^n + \text{TetR\_Protein}} - k_{tl} \cdot \text{cl\_mRNA} - k_r \cdot \text{cl\_mRNA}$$

Reaction rates and parameter setting:  $k_{tl} = 6.93$ ,  $k_p = 0.069$ ,  $k_r = 0.347$ ,

$a0\_tr = 0.03$ ,  $a\_tr = 29.97$ ,  $KM = 40$ ,  $n = 3$  resulted from parameter fitting based on available experimental data (Garcia-Ojalvo et al.).

System implies sustained limit-cycle oscillations after transient phase.

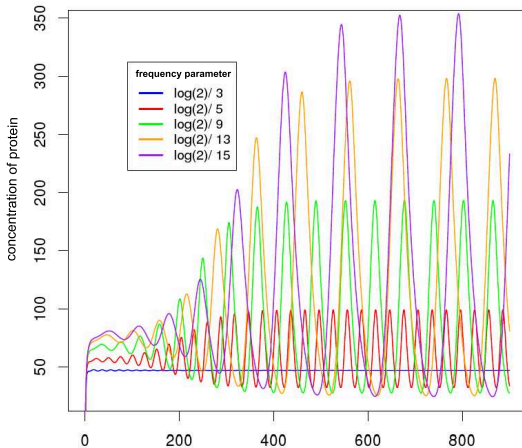
# Dynamical Behaviour of the Repressilator (TetR)



Initialisation at limit cycle avoids transient phase

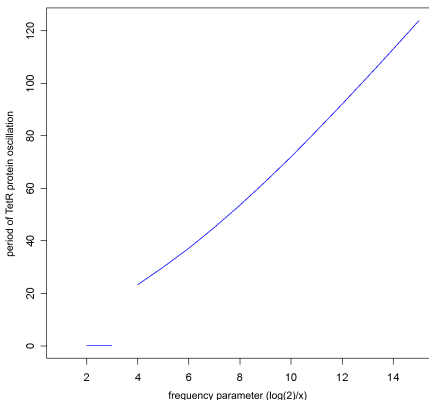
⇒ Eliminates its influence on synchronisation time

# Period Control by Velocity of Protein Degradation



Variable degradation rates  $k_p = \ln(2)/x$  (frequency parameter  $x = 3, \dots, 15$ ) of proteins sufficient for clock advance or delay.  
Frequency control: prerequisite for synchronisability.

# Repressilator's Transfer Function



Correlation between velocity of protein degradation and period.  
Identification of minimal period delimiting sustained oscillations.

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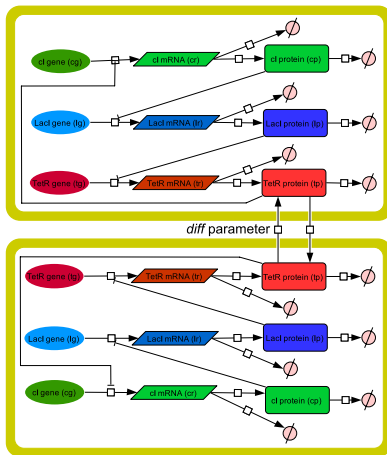
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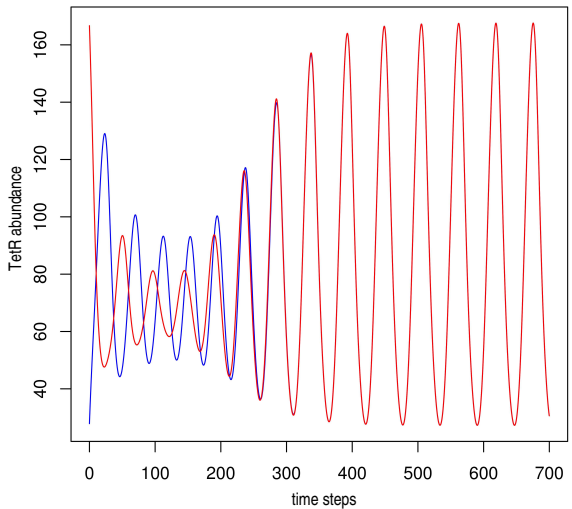
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# Coupled Repressilators for Internal Synchronisation



Bidirectional diffusion of **TetR proteins** between either repressilators enable internal synchronisation. Diffusion parameter *diff* as additional rate constant (linear kinetics)

# Typical Synchronisation Run

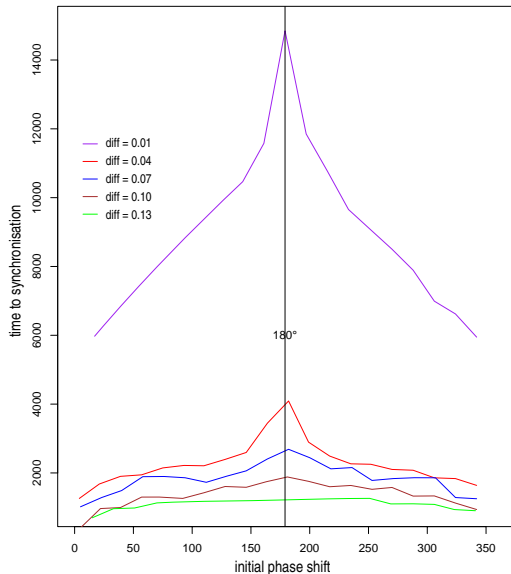


Typical synchronisation run of two TetR-coupled repressilators, coupling strength  $diff = 0.04$ , initial phase shift  $182^\circ$ .



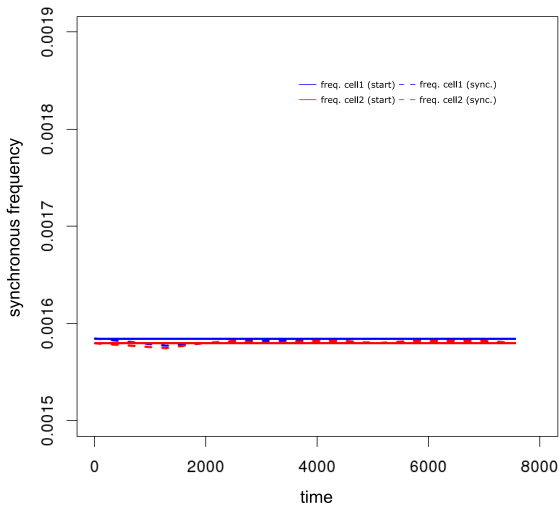


# Time to Synchronisation for Various Initial Phase Shifts



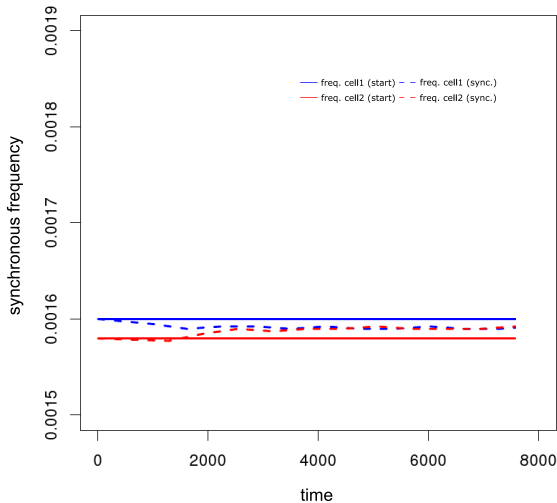
Time to synchronisation subject to various initial phase shifts. Parameter  $diff = 0.01, \dots, 0.13$  denotes coupling strength from weak to strong coupling. Initial antiphase rhythmicity (phase shift  $180^\circ$ ) between both repressilators causes the highest effort to synchronise both oscillatory signals by mutual forcing.

# Time to Synchronisation for Various Initial Frequencies



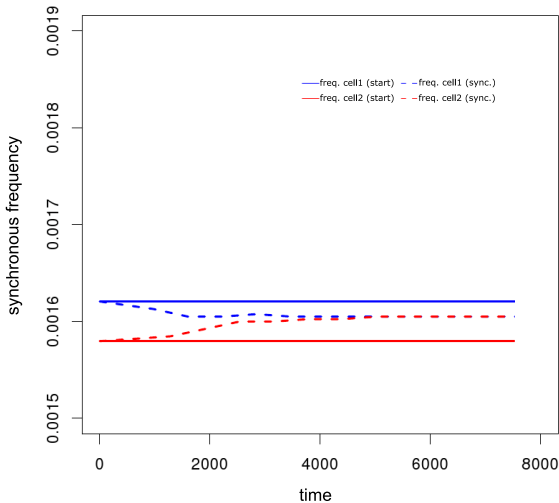
Weak diffusion,  $diff = 0.01$ , frequency parameter  $x$  ratio: 9.475 / 9.5

# Time to Synchronisation for Various Initial Frequencies



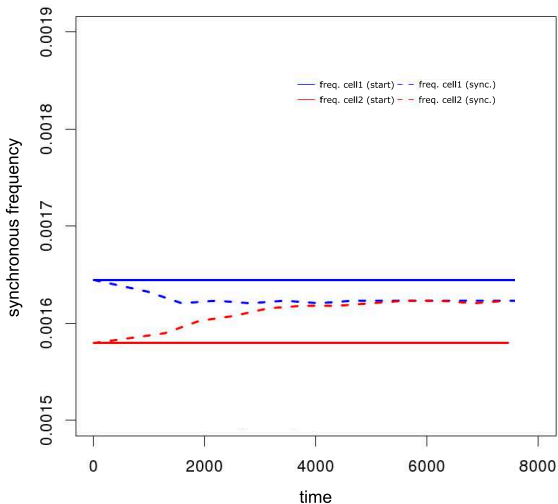
Weak diffusion,  $diff = 0.01$ , frequency parameter  $x$  ratio: 9.4 / 9.5

# Time to Synchronisation for Various Initial Frequencies



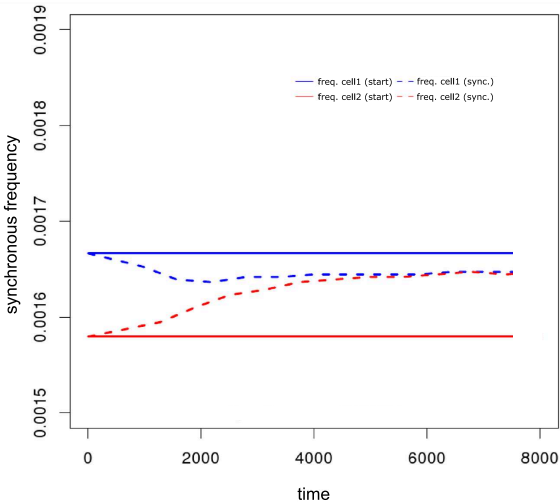
Weak diffusion,  $diff = 0.01$ , frequency parameter  $x$  ratio: 9.3 / 9.5

# Time to Synchronisation for Various Initial Frequencies



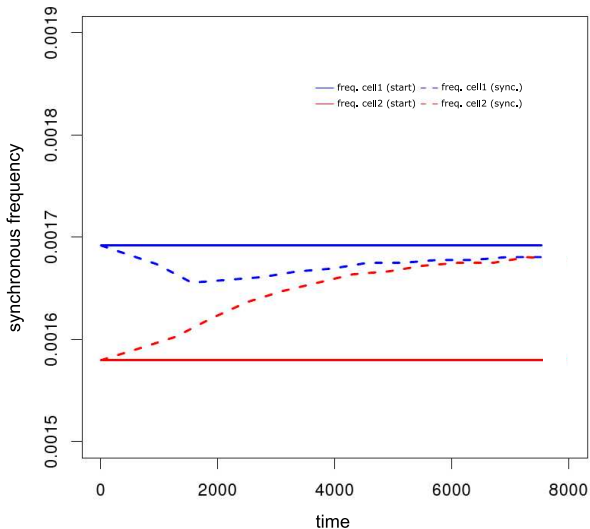
Weak diffusion,  $diff = 0.01$ , frequency parameter  $x$  ratio: 9.2 / 9.5

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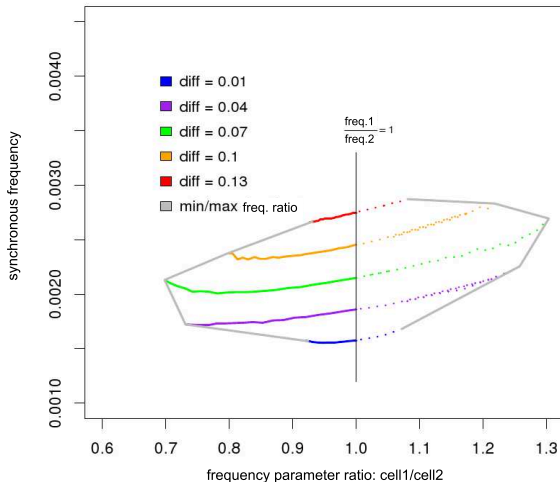
Weak diffusion,  $diff = 0.01$ , frequency parameter  $x$  ratio: 9.1 / 9.5

# Time to Synchronisation for Various Initial Frequencies



Weak diffusion,  $diff = 0.01$ , frequency parameter  $x$  ratio: 9.0 / 9.5

# Frequency Synchronisation Window



Ratios of initial frequencies subject to synchronous frequency considering variety of coupling strengths  $\text{diff} = 0.01, \dots, 0.13$ : variant of an Arnold tongue



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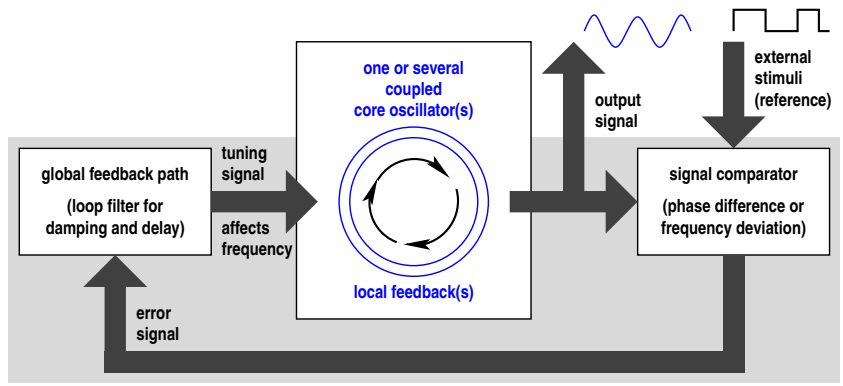
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# Frequency Control System with Phase-Locked Loop



Coupled repressilators as core oscillator of frequency control system able to manage external synchronisation to external stimuli (reference oscillation)

## Conclusions and Take Home Message

- Repressilator as promising biochemical *in-vitro* model system to explore synchronisation of circadian oscillations
- Inherent oscillation similar but not equal to sinusoidal course (hence not “symmetric”)
- Repressilator coupling by diffusion of TetR protein enables internal synchronisation.
- Arbitrary initial phase shifts (also antiphase behaviour) become harmonised while adaptation to different initial frequencies spans a synchronisation window.
- Coupled repressilators can be considered as part of a frequency control system based on phase-locked loop (PLL) utilising external synchronisation.

## Special Thanks go to . . .

### ... my coworkers

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Department Bioinformatics, FSU Jena



**Stefan Schuster**

Department Bioinformatics, FSU Jena



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Bundesministerium  
für Bildung  
und Forschung

### ... you for your attention. Questions?