

# Model-based Investigation of the Effect of the Cell Cycle on the Circadian Clock through Transcription Inhibition during Mitosis

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Sylvain Soliman

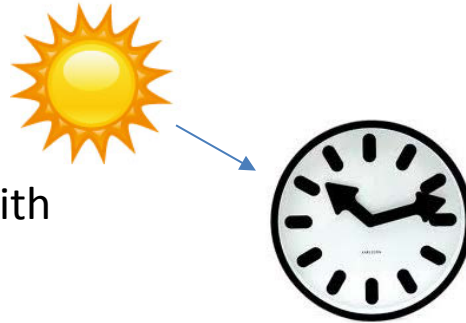
Lifeware, Inria Paris Rocquencourt



# The cell cycle and the circadian clock

## Context:

Optimizing cancer treatment with  
chronotherapy



## Two complex systems:

Circadian Clock: autonomous cellular oscillator with a  
24h period, synchronised by the central clock

Cell Cycle: divisions every 24h

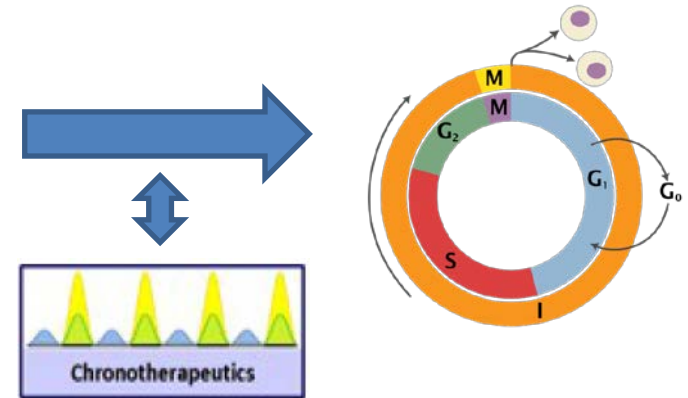
## Question:

What are the interactions between the two?

## Classical view:

Entrainment in period and phase of the cell cycle by the  
circadian clock (Matsuo et al 2003)

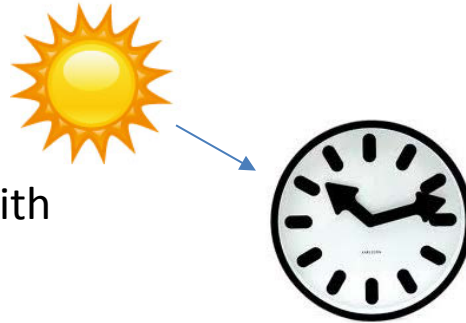
Gating by Wee1



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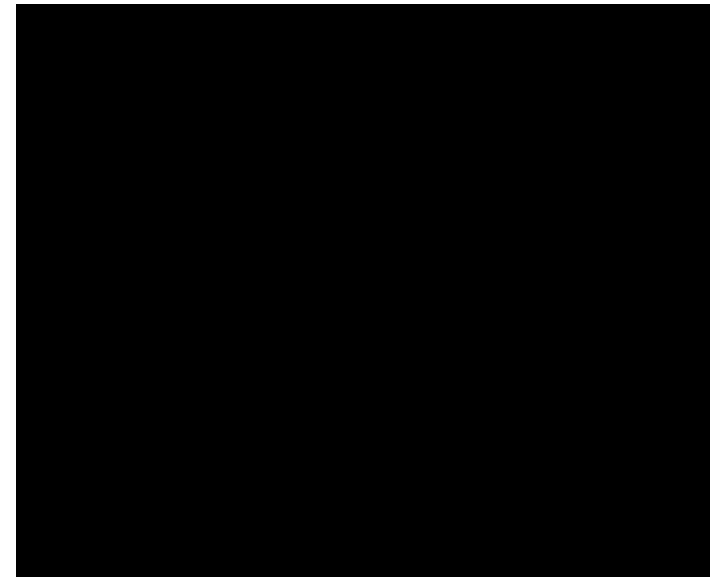
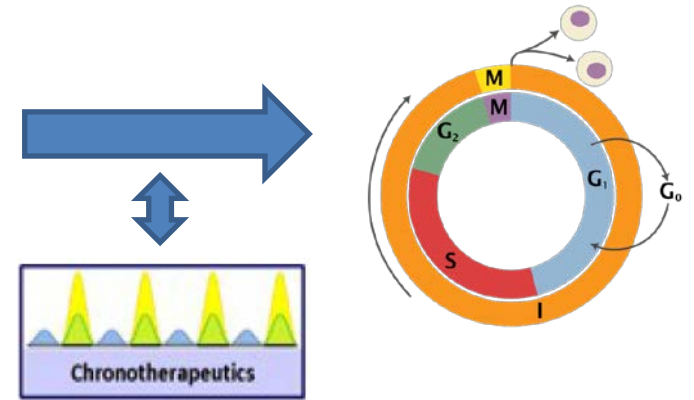
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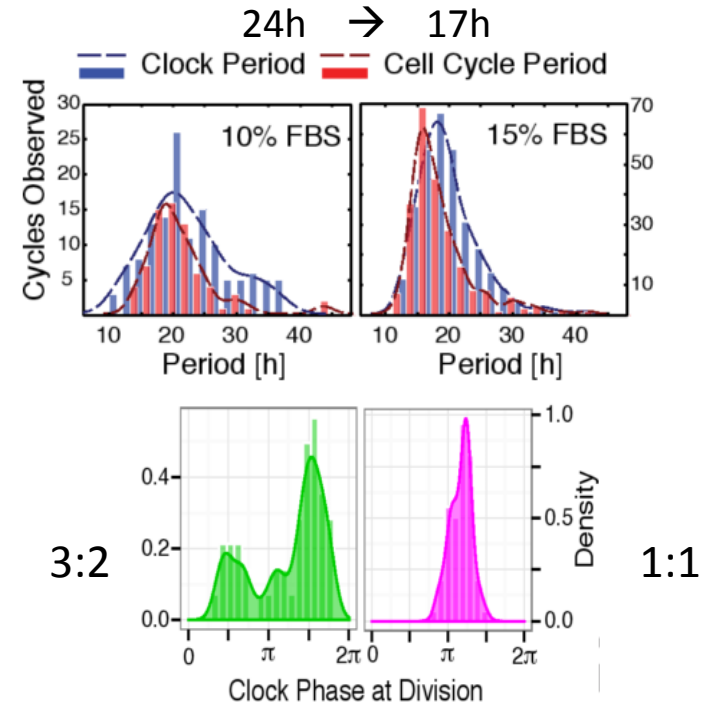
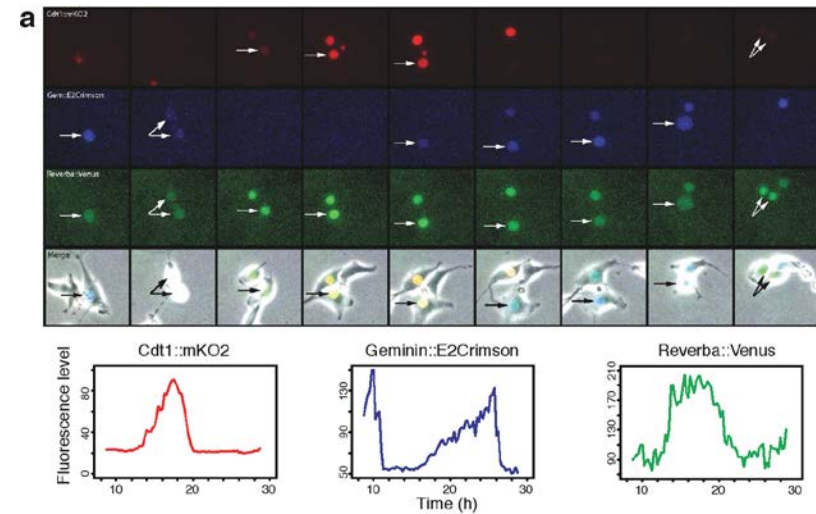
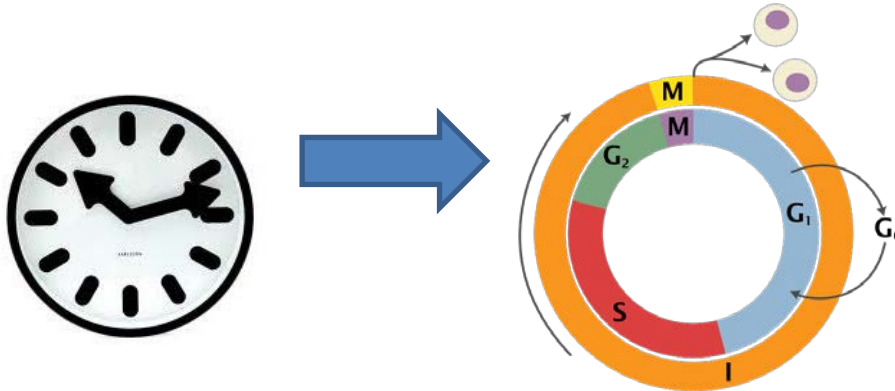


# Paradox: experimental data show an acceleration of the circadian clock at high FBS

## Experiments and observations

- Fluorescent markers of the cell cycle and the circadian clock
- Time series in individual mice fibroblasts
- Medium with various concentrations of serum (FBS) to modulate cell cycle length
  - **Acceleration** of the circadian clock in fastly dividing cells – **and not in confluent cells**
- Some experiments begin with a Dexamethasone pulse
  - **Various modes of phase-locking**

These experiments suggest that in some conditions there is **a control of the circadian clock by the cell cycle**

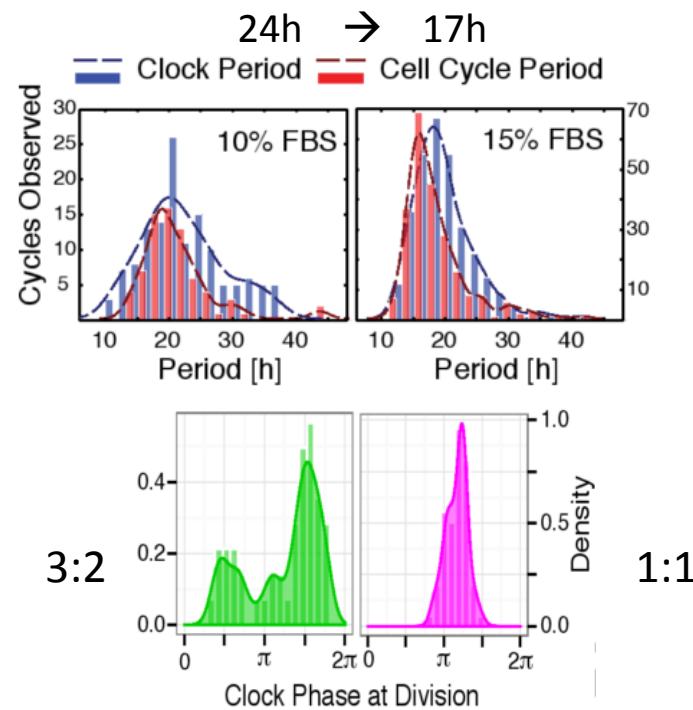
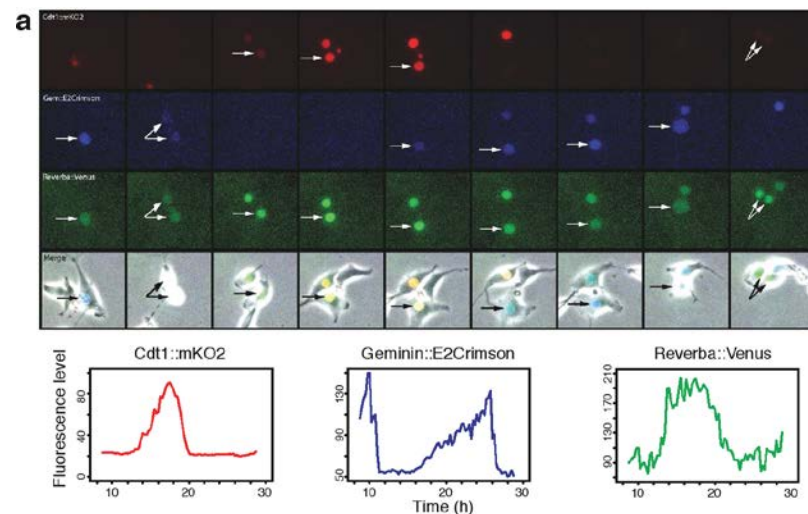


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Model-based investigation of that **reverse coupling**  
from the cell cycle to the circadian clock

- Reusing already existing **differential models from the literature**:  
Cell cycle: Qu et al 2003  
Circadian clock: Relogio et al 2011

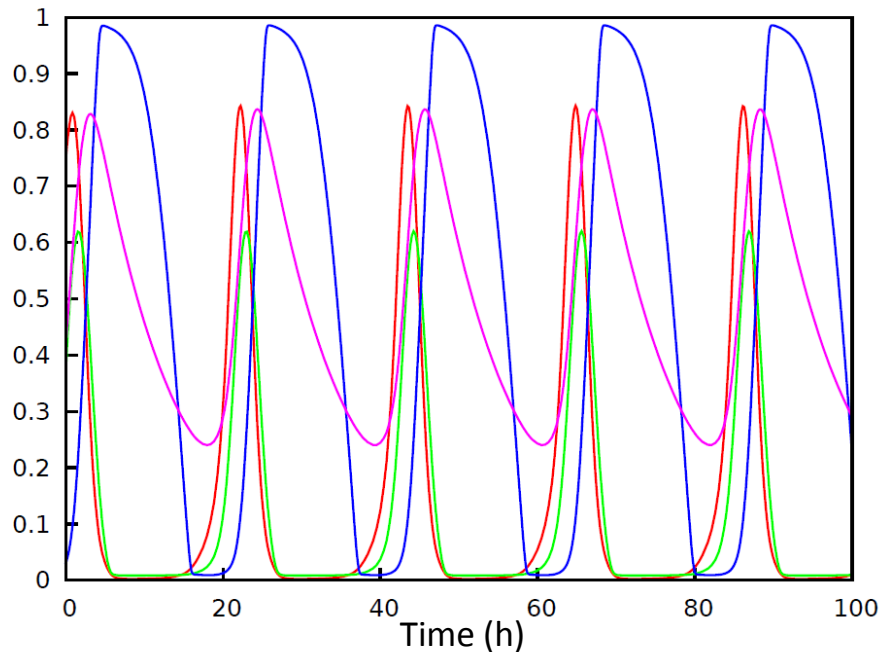
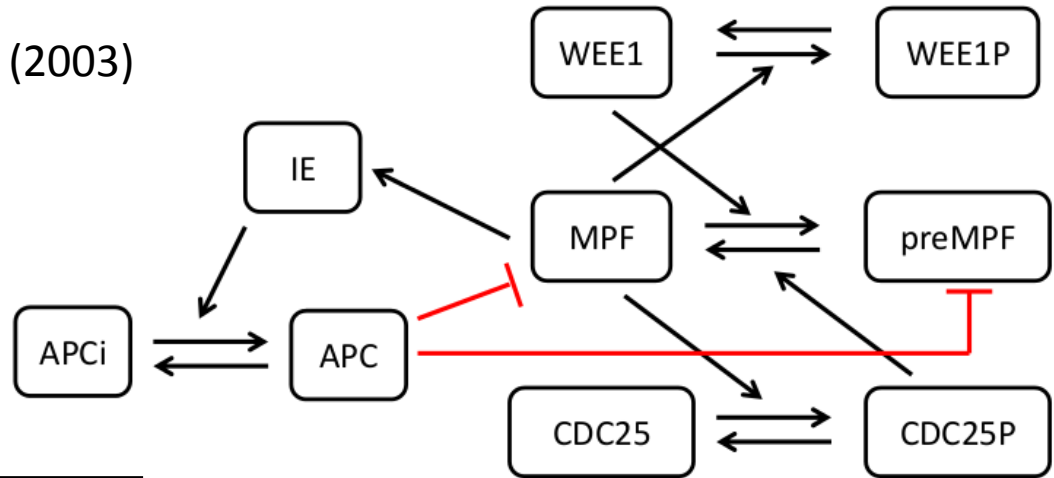
- Coupling by inhibition of transcription during mitosis

$$S * \frac{J^n}{J^n + ([MPF]/[preMPF])^n}$$

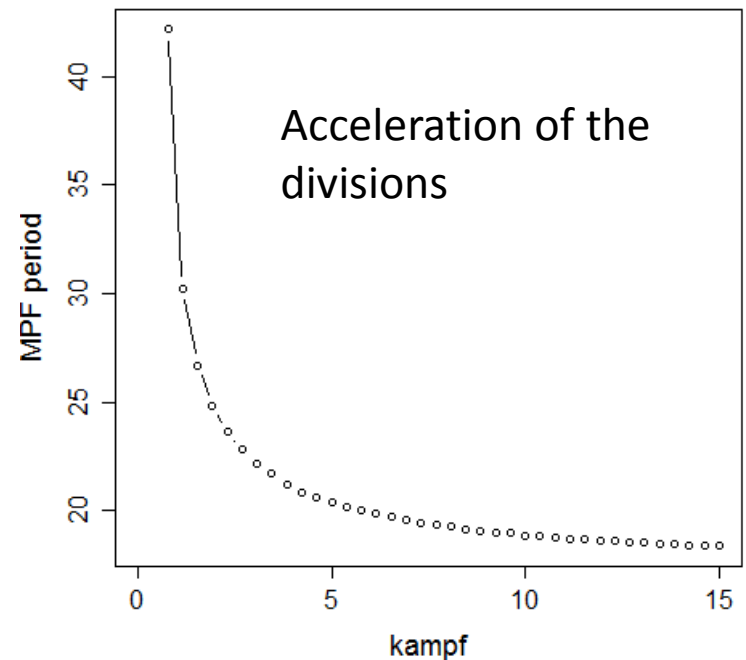
- **Model analysis** assisted with formal methods [CMSB2014]  
Parameter inference
- **Predictions**: mechanisms and perturbations, treatment optimization

# Modeling the control of the circadian clock by the cell cycle

- Cell cycle : Qu, McLellan, Weiss (2003)
  - Mitosis phase
  - 10 equations, 31 parameters

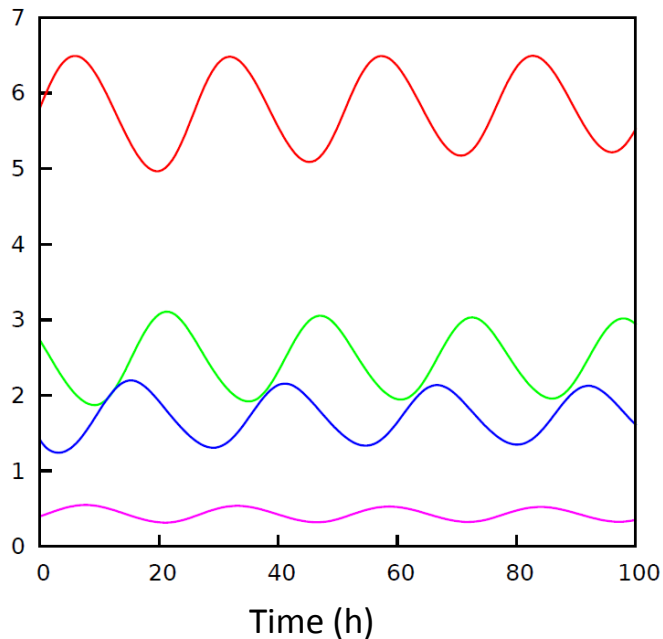


— MPF  
— C25P  
— APC  
— IE



# Modeling the control of the circadian clock by the cell cycle

- Circadian clock : Relogio Herzel (2011)
  - 20 species, 71 parameters
  - **The values of 60 parameters were determined to fit experimental data (amplitude and phase) in liver cells**
  - 5 genes : Per, Cry, RevErb, Ror, Bmal.
  - 2 negative feedback loops
  - 2 positive feedback loops



— Ror::nucl  
— RevErb::nucl  
— Bmal-Clock::nucl  
— Cry-Per::nucl

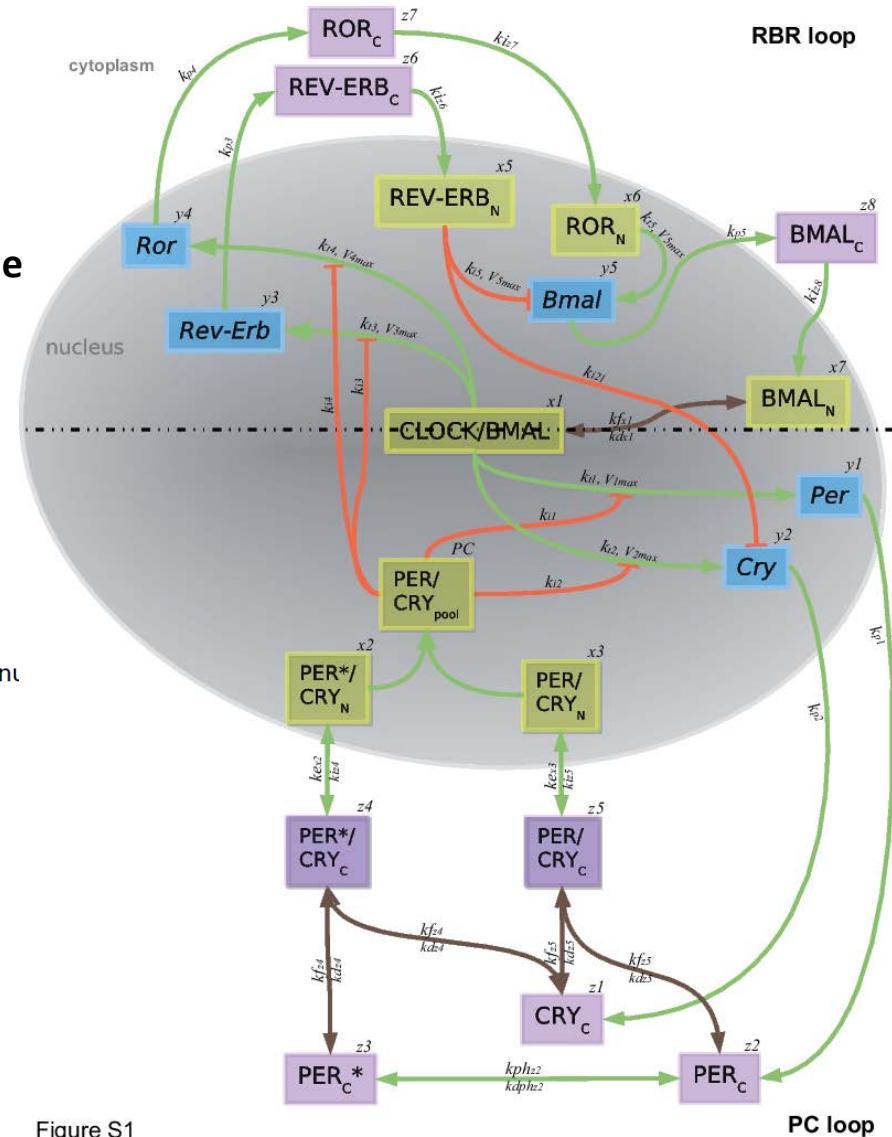
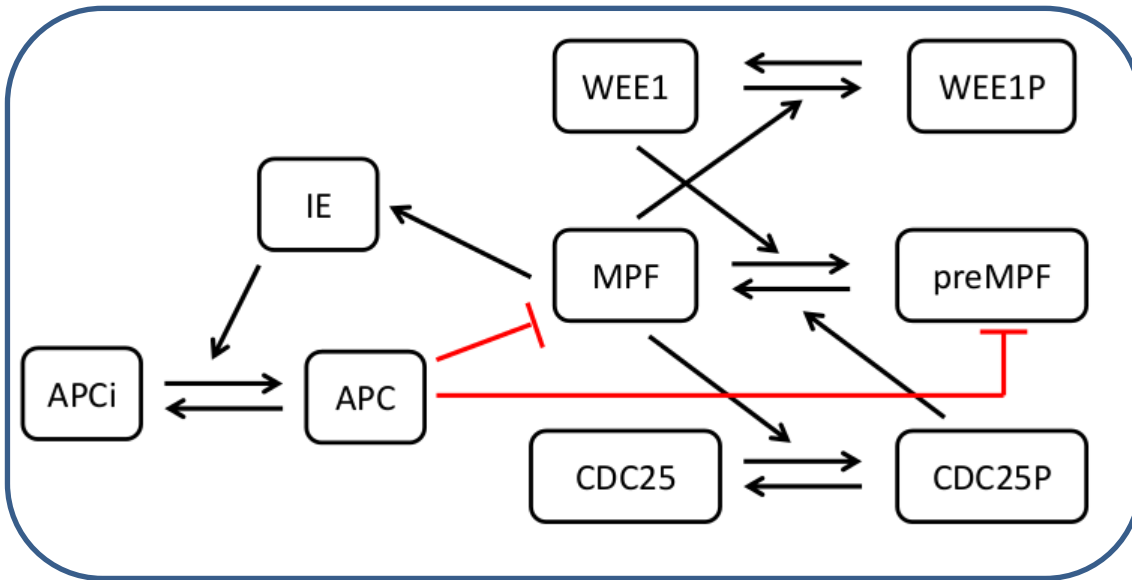


Figure S1

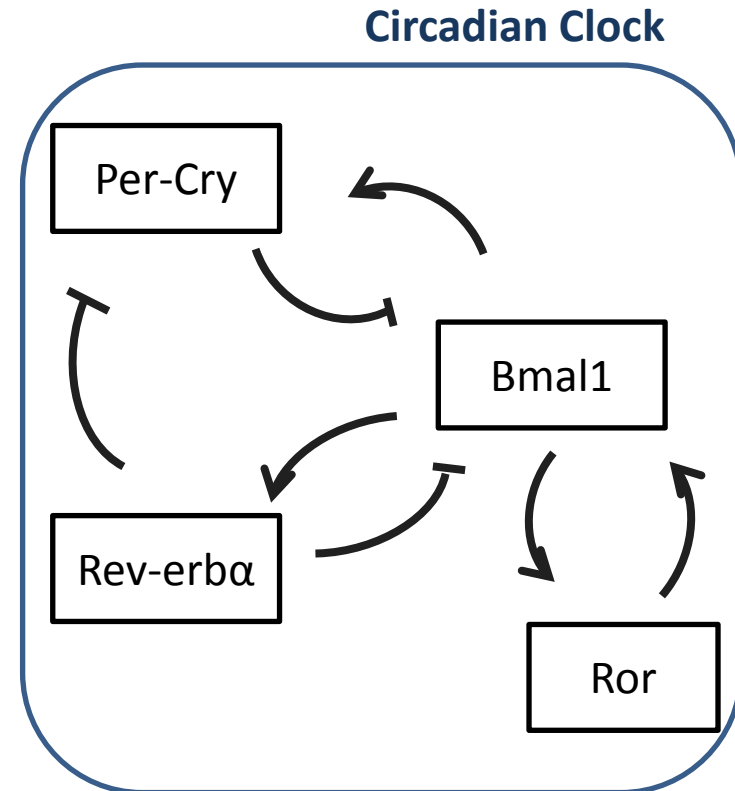
Relógio, A., Westermark, P. O., Wallach, T., Schellenberg, K., Kramer, A., & Herzel, H. (2011). Tuning the mammalian circadian clock: robust synergy of two loops. PLoS Computational Biology.



# Modeling the control of the circadian clock by the cell cycle



Cell Cycle



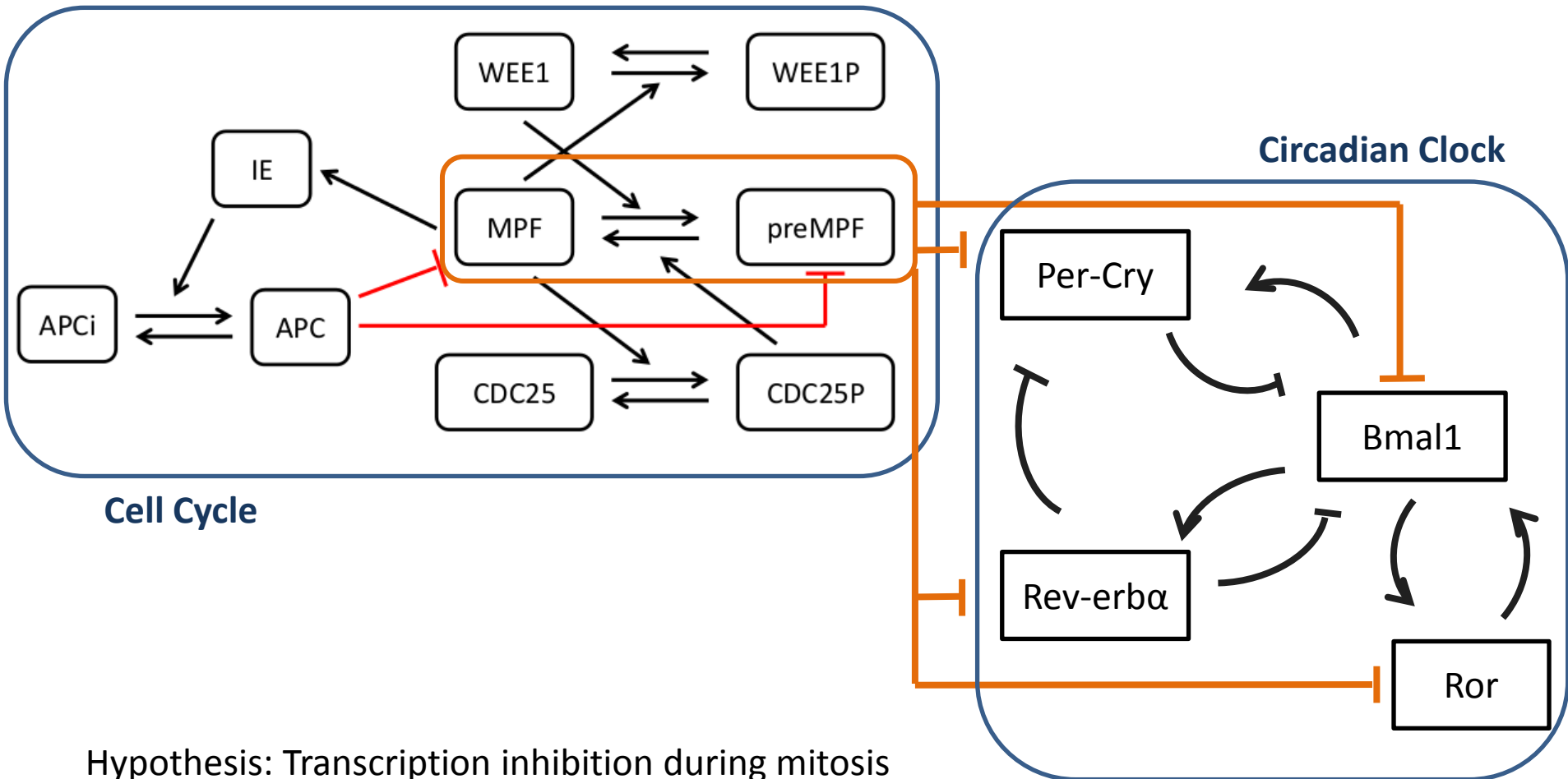
Circadian Clock

Hypothesis: Transcription inhibition during mitosis

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for each synthesis expression S in the clock

# Modeling the control of the circadian clock by the cell cycle



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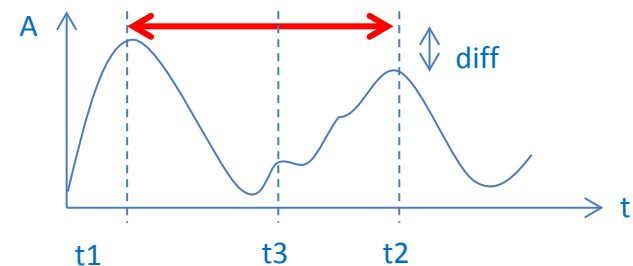
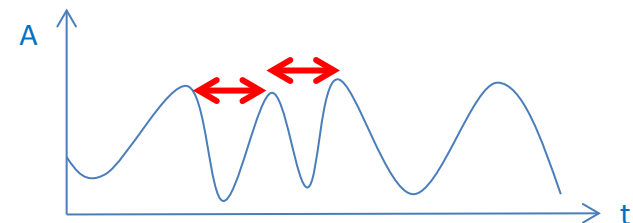
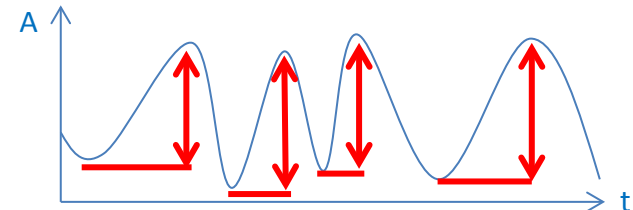
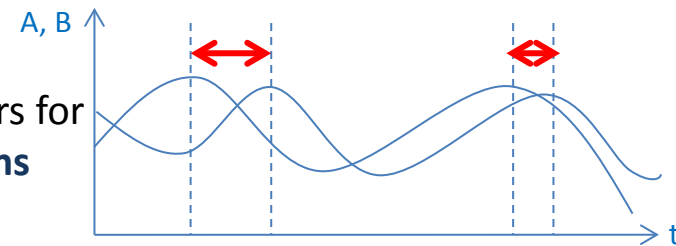
for each synthesis expression S in the clock

# Formalizing the dynamical behaviors with temporal logic QFLTL(R)

- **QFLTL**: expressing qualitative or quantitative dynamical behaviors for oscillatory systems such as **period, amplitude, phase, oscillations regularity**
- **Applications:**
  - Data analysis, Model checking, Model analysis, Parameter inference (calibration)
  - Tools implemented in **BIOCHAM**
- **Built-in functions:**  
Ex: **amplitude**(Mol,amp), **period**(Mol,per),  
**phase**(Mol1,Mol2,phase)

**Temporal specification: regular oscillations with a fixed period**

```
period([RevErb::nucl],[period])  
& Exists([maxdiff1,maxdiff2,maxpeak],  
  maxDiffDistancePeaks([RevErb::nucl],[maxdiff1])  
  & maxDiffAmplPeaks([RevErb::nucl],[maxdiff2])  
  & maxAmplPeaks([RevErb::nucl],[maxpeak]))  
& 4*maxdiff1 < period + errordiff1  
& 10*maxdiff2 < maxpeak + errordiff2  
& maxpeak > 0.1 + errorampl)
```

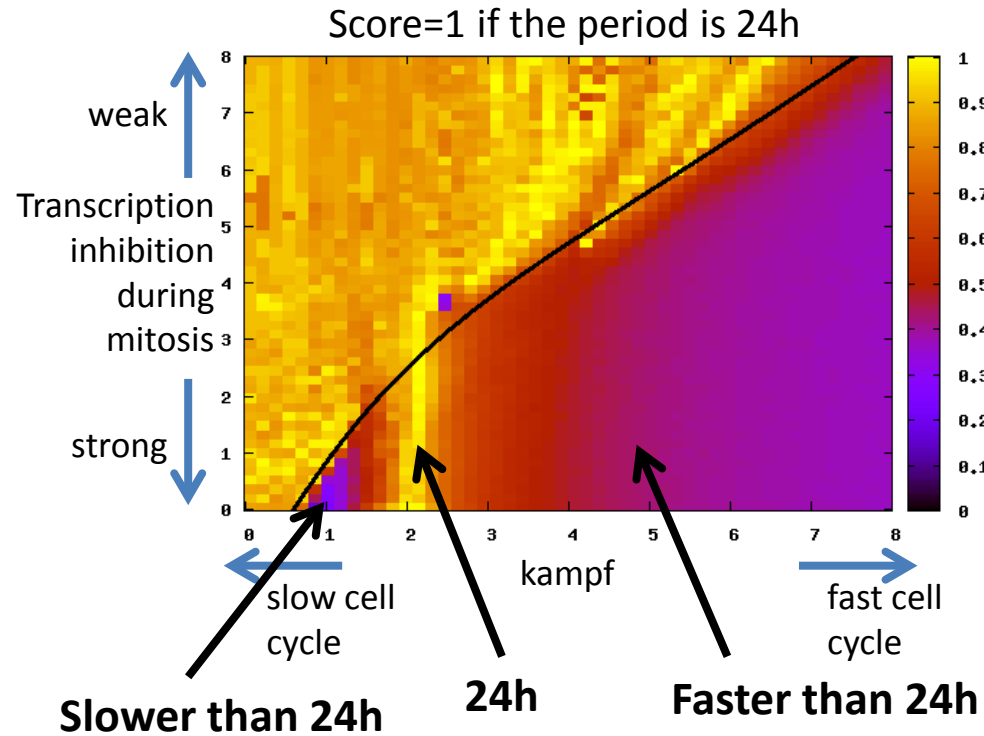


# Results without Dexamethasone

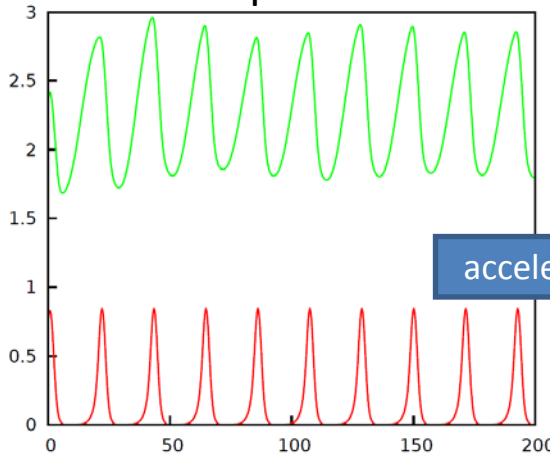
**Landscape of the score for an objective period of 24h for the circadian clock:**

**Domain of entrainment of the clock period by the cell cycle**

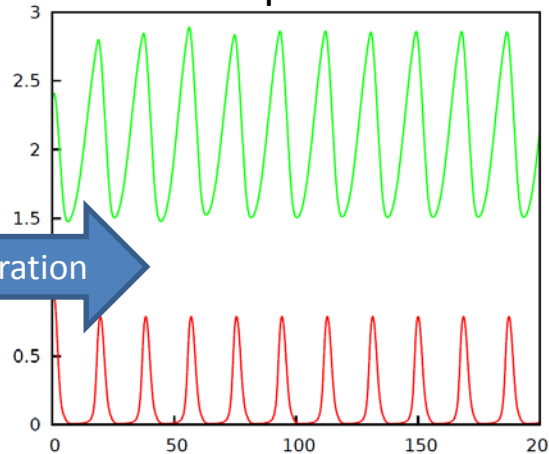
Kampf	FBS (%)	Circadian clock period (h)	Cell division period (h)
3.75	10	21.43	21.30
12.1	15	18.60	18.60
1.6	5 ?	26.16	26.32



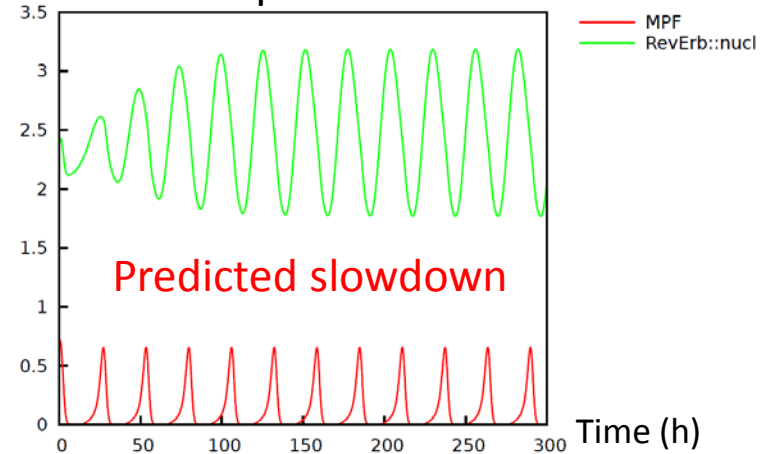
kampf=3.75



kampf=12.1



kampf=1.6



# Results with Dexamethasone

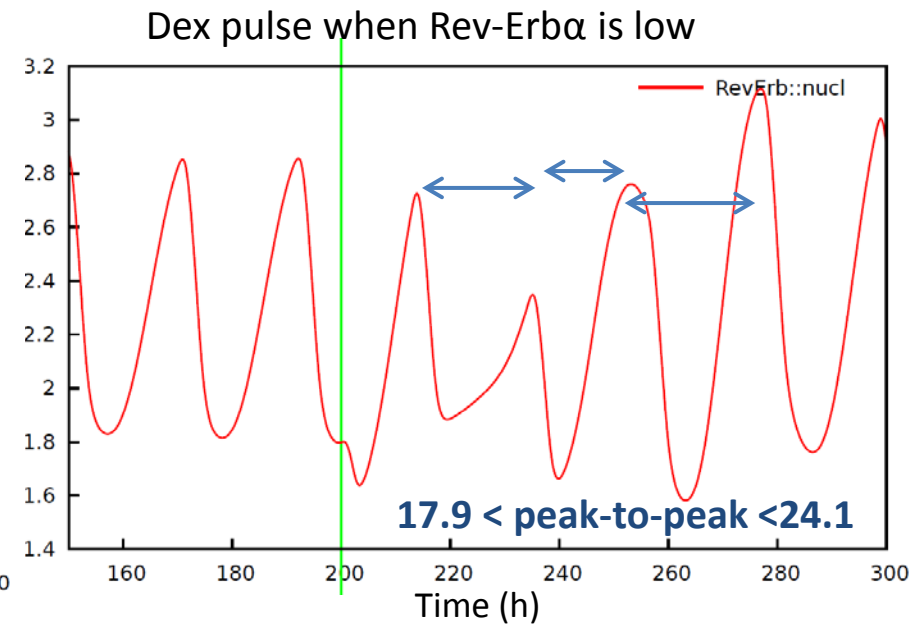
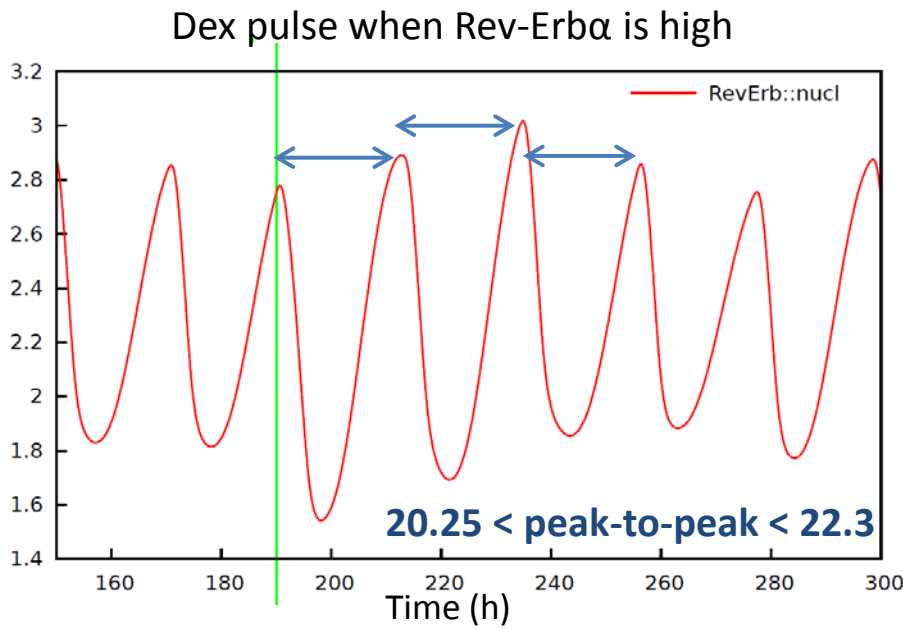
Dexamethasone pulses induce the expression of *per1*

→ extension of the model with a **2h event** increasing the synthesis of mPER

→ **The clock is disrupted** and then returns to the observed entrainment, regardless of the medium but **depending on the time of the pulse**

→ **Transitory variations** could explain the noisy data

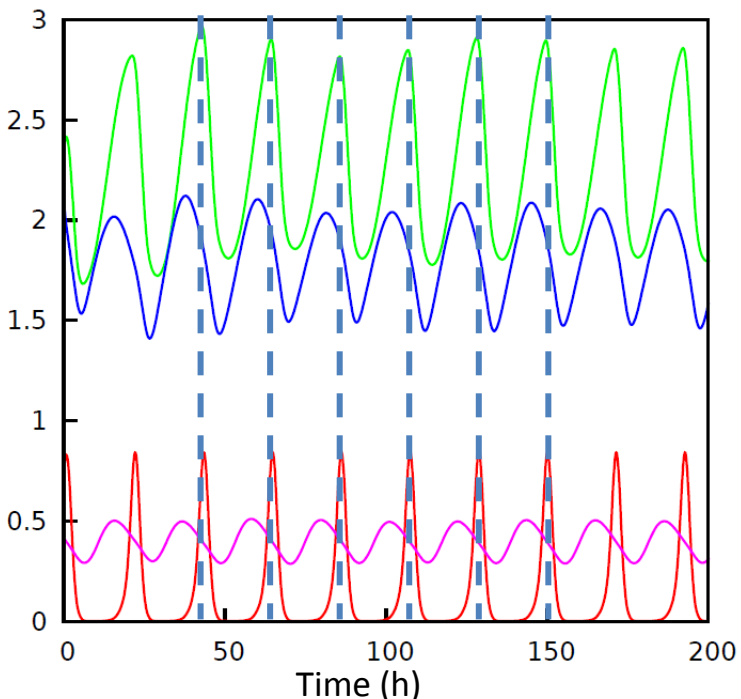
Med	No dexamethasone		Dexamethasone	
	Clock period	Division period	Clock period	Division period
FBS 10	21.9h ± 1.1h	21.3h ± 1.3h	24.2h ± 0.5h	20.1h ± 0.94h
FBS 15	19.4h ± 0.5h	18.6h ± 0.6h	NA	NA
FBS 20	NA	NA	21.25h	19.5h
FBS 20	NA	NA	29h	16h



# Conclusion 1

- Explanation hypothesis for the control of the circadian clock: **inhibition of the transcription during mitosis**
- The hypothesis reproduces the **acceleration of the circadian clock** by a fast cell cycle
- **Prediction** of the model: **entrainment in period of the circadian clock** even in slowly dividing cells

# Discrepancies on the phases



— MPF  
— RevErb::nucl  
— Bmal-Clock::nucl  
— Cry-Per::nucl

## Phase between MPF and Rev-Erb $\alpha$

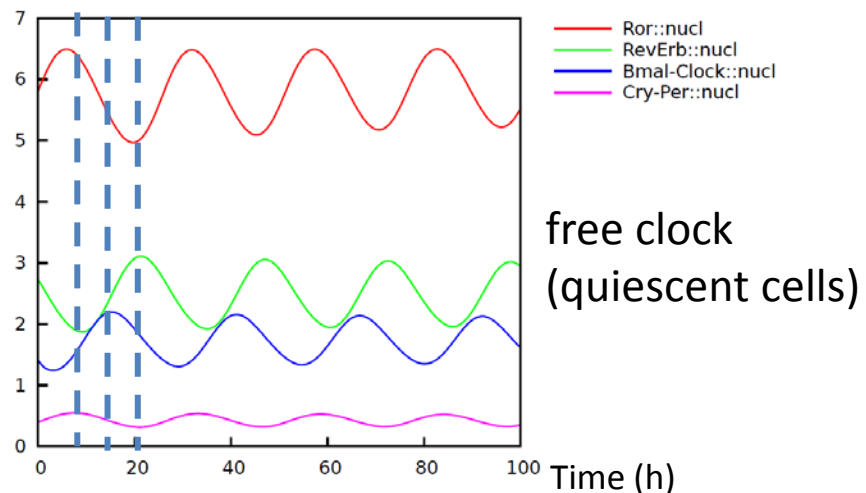
Medium	Experimental data	Model simulation
FBS 5	NA	18.6h
FBS 10	3.82h	20.7h
FBS 15	3.98	17.8h

- The phase between the mitosis and the circadian clock does not match the data
- Same problem in published models for the control of the cell cycle by the circadian clock

→ Need to revise the model of the circadian clock?  
 → Consider other coupling hypothesis?

# Prediction of the model

In quickly dividing cells, the phase shifts between the different components of the clock are shorter than in quiescent cells



free clock  
(quiescent cells)

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# Conclusion 2

- Remaining discrepancies on the **circadian phase at division**
- **Prediction** of the model: shorter phase shifts between the components of the clock



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# Acknowledgements

- **Lifeware Team**

François Fages  
Sylvain Soliman  
Grégory Batt  
Thierry Martinez  
François-Marie Floch  
Denis Thieffry

- **Collaborations**

C5Sys (ERASysBio)  
ANR HyClock  
Franck Delaunay  
Céline Feillet

- **GENCI cluster**

Public high performance computing resources

