## Lab Course: Modelling and Simulation (SS-2016, LSI) 5th -15th September 2016

Task 1: A simple model to explain the growth of organism is the exponential model given by dN/dt = rN where N(t) is the population at time t and r > 0 is the growth rate. But exponential growth is unrealistic. Therefore, to incorporate the ideas of overcrowding and limited resources, in the exponential growth model logistic equation is used. The following one-dimensional model is such a logistic equation:

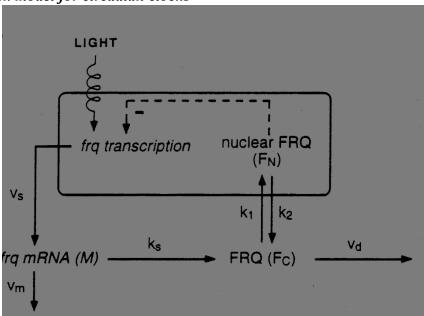
$$dN/dt = rN(1-N/K)$$

where N(t) is the population at time t, r > 0 is the growth rate and K is the carrying capacity.

- i ) Solve the above equation analytically and plot the result in a time series manner.
- ii ) Simulate it using numerical tools and compare the plots (try different solvers and step sizes)

Task 2:

Genetic feedback model for circadian clocks



We consider a **3-dimensional Goldbeter model** for expression and regulation of the *Neurospora* "frequency protein" (FRQ). For the concentrations of the mRNA (M), the "frequency protein" ( $F_C$ ) synthesized in the **cytoplasm** and the successively built up protein concentration ( $F_N$ ) in the **nucleus**, we obtain the following *system of three differential equations*:

$$dM/dt = V_s \frac{K^n}{K^n + F_N^n} - V_m \frac{M}{K_m + M}$$

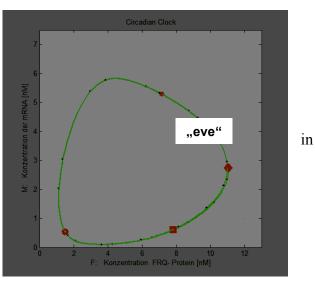
$$dF_C/dt = k_s M - V_d \frac{F_C}{K_d + F_C} - k_{in} F_C + k_{out} F_N$$

$$dF_N/dt = k_{in} F_C - k_{out} F_N$$

(1) Simulate this 3-dim Goldbeter model with suitable parameters, whereby you should start with the following parameter set and try to alter it in order to obtain the experimentally observed intrinsic period P = 21.5 h (see below)!

$$V_s = 1.5 \Leftrightarrow 2.0$$
 during dark or light, respectively  $[nM \cdot h^{-1}]$   $V_m = 1.5$   $[nM \cdot h^{-1}]$ ,  $V_d = 1.0$   $[nM \cdot h^{-1}]$ ,  $k_s = 0.5$   $[h^{-1}]$   $K = 0.2$   $[nM]$ ,  $K_m = 0.15$   $[nM]$ ,  $K_d = 0.15$   $[nM]$   $k_{in} = 0.02$   $[h^{-1}]$ ,  $k_{out} = 0.1$   $[h^{-1}]$   $n = 4$ : Hill number

- (a) Investigate the intrinsic "circadian clock" for different **transport rates**  $k_{in}$  and estimate the resulting **Period** in the time plot for M(t) and F(t). Which parameters produce a **Period**  $\approx 21.5$  h, the value for *Neurospora* in constant darkness?
- (b) During the 'time loop' determine a list of *time points* **T\_eve**, where the cytoskeletal protein level **F(t)** is **maximal**. By the mean difference **mean(diff(T\_eve))**compute the **Period** of the "clock".
- (c) Moreover, in the *state diagram* as well as the time diagram, try to mark the "quarter phases", when the intrinsic clock attains its "evening" (T\_eve), "midnight", "morning" and "noon".

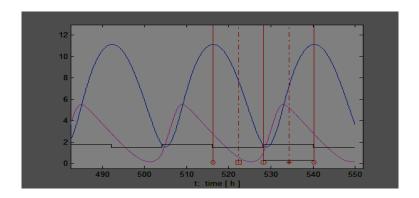


## Task 3:

(2) Perform simulations to show, under which conditions a **12-12 h light-dark stimulus** is able to entrain the intrinsic *Neurospora* clock.

Take Step = 0.5 for the increase of transcription rate altering the mRMA production speed Vs from Vs0 = 1.5 (during 12 h darkness) to Vs1 = Vs0 + Step (during the following 12 h light).

• In the time diagram the stimulus level Vs is plotted as a stair function

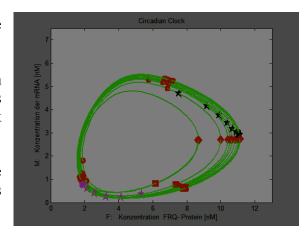


In the state diagram mark the points of light switch on and switch off (!) stimuli.

Will the intrinsic clock be entrained by the 24h cycle?

What is the *minimal step increase* in transcription rate (**Step**) that induces entrainment of the intrinsic *Neurospora* clock by the 12-12 h light-dark stimulus?

How does the outcome depend on the "phase" of first *light switch off* stimulus relative to the intrinsic clock cycle?



## *Task 4:*

Perform simulations to show, under which conditions a **18-6 h light-dark stimulus** is able to entrain the intrinsic *Neurospora* clock. Modify the simulations of task 2 using 8 h of darkness and 16 h of light.

Good luck:)