practical_3_negative_feedback_solved

October 19, 2022

1 SYSTEMS AND NETWORK BIOLOGY - PRACTICAL 3

2 Negative feedback

To submit your report, answer the questions below and save the *notebook* clicking on File > Download as > iPython Notebook in the menu at the top of the page. **Rename the notebook file** to "practicalN_name1_name2.ipynb", where N is the number of the practical, and name1 and name2 are the first surnames of the two team members (only one name if the report is sent individually). Finally, **submit the resulting file through the** *Aula ESCI*.

Remember to label the axes in all the plots.

IMPORTANT REMINDER: Before the final submission, remember to **reset the kernel** and re-run the whole notebook again to check that it works.

The objectives of this practical are: - to become familiar with numerical integration of dynamical systems using scipy's odeint function. - to become familiar with the basic models of gene regulation for basal expression and transcriptional repression. - to understand the differences between unregulated gene expression and negative autoregulation. - to explore the effects of saturation in the controller of an integral feedback control system

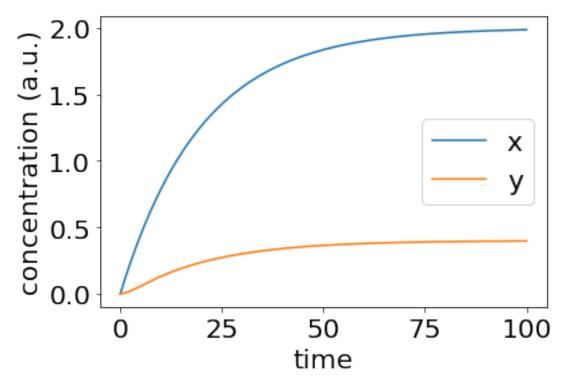
```
[1]: import numpy as np
from scipy.integrate import odeint
import matplotlib.pyplot as plt
%matplotlib inline
```

2.1 Numerical integration

We will use scipy's odeint function to numerically integrate ODEs. Here is an example:

```
[2]: def test(xy,t):
    x,y=xy
    dx=alpha1-delta1*x
    dy=alpha2*x-delta2*y
    return [dx,dy]
```

```
alpha1 = 0.1
alpha2 = 0.1
delta1 = 0.05
delta2 = 0.5
tvec = np.arange(0,100,0.01)
x0 = [0,0]
y = odeint(test,x0,tvec)
plt.figure()
plt.rc('font', size=20)
plt.plot(tvec,y[:,0],label='x')
plt.plot(tvec,y[:,1],label='y')
plt.legend()
plt.xlabel('time')
plt.ylabel('concentration (a.u.)')
plt.show()
```

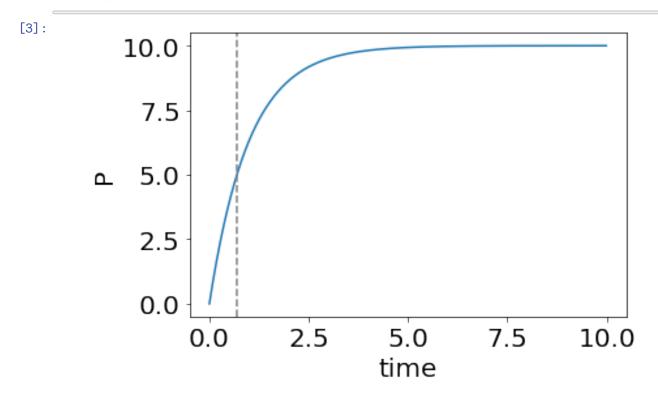


3 Constitutive expression

The following equation can be used to model the constitutive expression of a gene:

$$\frac{dP}{dt} = \alpha - \delta P$$

- 1. Consider $\alpha = 10$ and $\delta = 1$.
 - 1. What is the analytic expression for the steady state of the system?
 - 2. Plot the time evolution of the system starting from $P_0 = 0$ using scipy's odeint function.
 - 3. Does the final value of the trajectory in b) coincide with the analytical steady state?
 - 4. At what time does P reach half the steady state value? Plot a vertical line at that point.



The final value of the protein concentration is 9.999541435840129

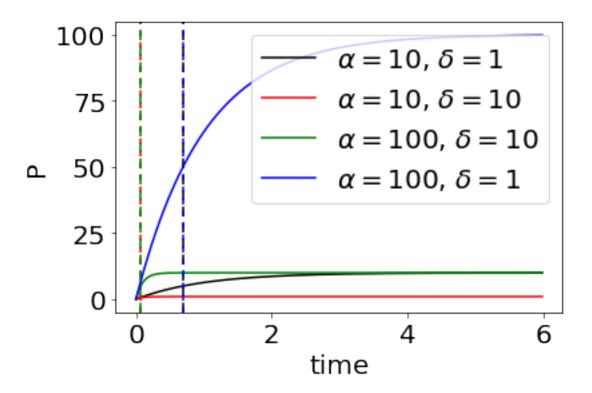
The time at which the protein concentration reaches half its steady state is 0.7

- 2. What happens for the following parameter sets, in comparison with the previous one? Plot the four of them in the same plot and compare them.
 - 1. alpha=10, delta=10
 - 2. alpha=100, delta=10
 - 3. alpha=100, delta=1
- [4]: For alpha=10, delta=1, the time to reach half of the steady state concentration is 0.7

For alpha=10, delta=10, the time to reach half of the steady state concentration is 0.07

For alpha=100, delta=10, the time to reach half of the steady state concentration is 0.07

For alpha=100, delta=1, the time to reach half of the steady state concentration is 0.7



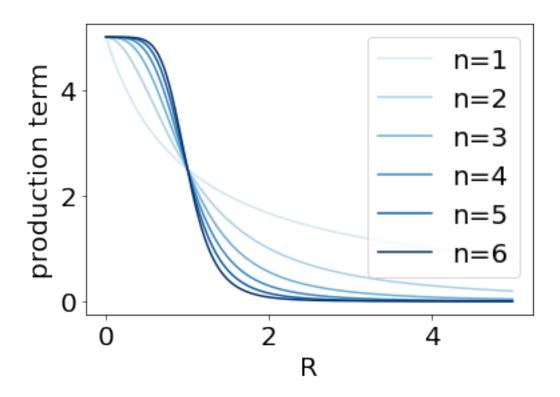
4 Negative feedback

Now consider the following negative autoregulation system:

$$\frac{dR}{dt} = \frac{\alpha}{1 + \left(\frac{R}{K}\right)^n} - \delta R$$

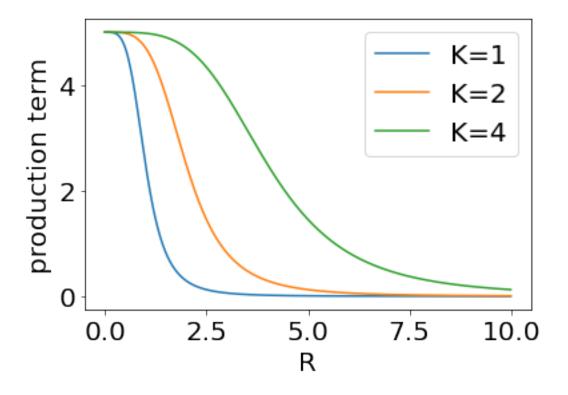
1. Plot the production term for alpha=5, K=1, and n ranging from 1 to 6, for a range of R values from 0 to 5. What is the effect of increasing n?

[5]:



2. Now fix n to 4, and plot the production term for K = 1, 2, and 4. What is the effect?

[6]:



^{**} Write your answer here **

3. If you wanted to build a repression system that is highly repressed at low values of repressor, how would you choose K and n? (in terms of low/high values)

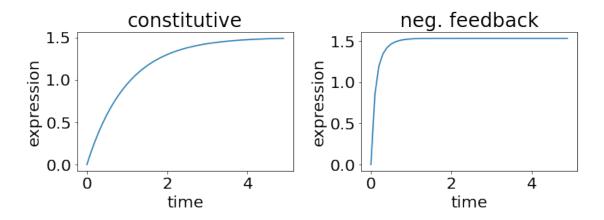
5 Study the response timescale of the system:

- 1. We will now compare how quickly the two models introduce above generate their product. To that end:
 - 1. Integrate the negative feedback model for n=4, K=1, $\alpha=10$ and $\delta=1$, starting from R=0 and compute the time to reach half the steady state concentration.
 - 2. Integrate the constitutive expression model with $\alpha = 1.5$ and $\delta = 1$ starting from R = 0 and compute the time to reach half the steady state concentration.
 - 3. Compare the results of B) and C).

[7]: Time to reach half the steady state for the constitutive model 0.7

Time to reach half the steady state for the neg. feedback model 0.1

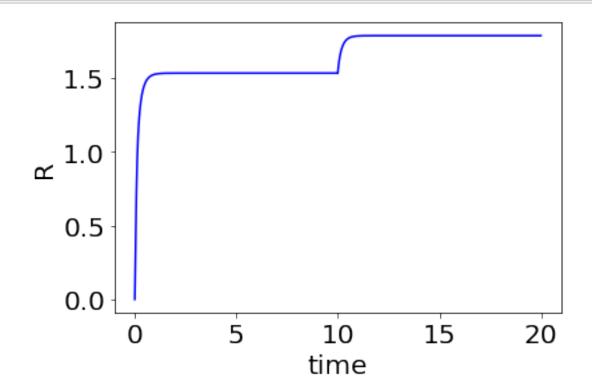
^{**} Write your answer here **



6 Adaptation to perturbations

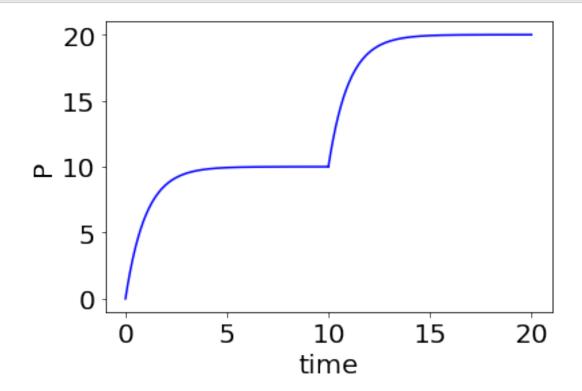
Integrate the negative feedback model for n=4, K=1, α = 10 and δ = 1, starting from R=0, for 10 time units. At that point, double the value of α and integrate for another 10 time units. Does the system return to the same steady state as before?

[8]:



Do the same for the constitutive expression model.





^{**} Write your answer here **

Interpret the results above in terms of the difference in robustness between the two models.

^{**} Write your answer here **