

# Differential Equations - Assignment

MME - Bioinformatics

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## 0. Naming the files

Please name your .py files in the following way and don't use special characters (like ö,ß,ñ,á):

assignment\_assignment#\_exercise#\_lastname.py

e.g.: assignment\_2\_1\_fuesschen.py for assignment 2, exercise 1 of student Füßchen

**Submit one file for each exercise!**

## 1. IVP Solver

Write a single python module that:

**A)** can solve the general initial value problem (IVP):

$$\mathbf{y}' = \mathbf{f}(t, \mathbf{y}); \mathbf{y}(t_0) = \mathbf{y}_0$$

using the explicit 4-th order Runge-Kutta method (RK4). (Note:  $\mathbf{y}$ ,  $\mathbf{y}_0$  and  $\mathbf{f}$  are vectors!)

**B)** solves the following concrete IVP in the interval [0,10] with your RK4 implementation:

$$y'' = -4y - y'$$

$$y(0) = 1; y'(0) = -1$$

and compares the result with the RK45 solution obtained from `scipy.integrate.solve_ivp`.<sup>1</sup>

**The module should contain the following:**

- A function `RK4_step(t, y, dt, f)` that performs a single RK4 integration step and returns the next time and (approximate) function values (`t_next, y_next`).

$$RK4\_step(t_n, y_n, dt, f) \rightarrow return(t_{n+1}, y_{n+1})$$

The input arguments are:

- `t`: current time (scalar)
- `y`: current function values (vector)
- `dt`: time step (scalar)
- `f`: function to calculate the derivative (function).

Note: it should work for an arbitrary function

$$f: \mathbb{R} \times \mathbb{R}^n \rightarrow \mathbb{R}^n; (t, \mathbf{y}) \rightarrow \mathbf{f}(t, \mathbf{y})$$

- A function `RK4integrator(y0, f, N, t0, dt)` that performs:
  - `N` integration steps
  - of length `dt`
  - starting at initial conditions `(t0, y0)`

by calling the `RK4_step` function `N`-times. It should return `(t, Y)`:

- `t`: an array containing the time grid
- `Y`: an array containing the numerical solution to the general IVP.

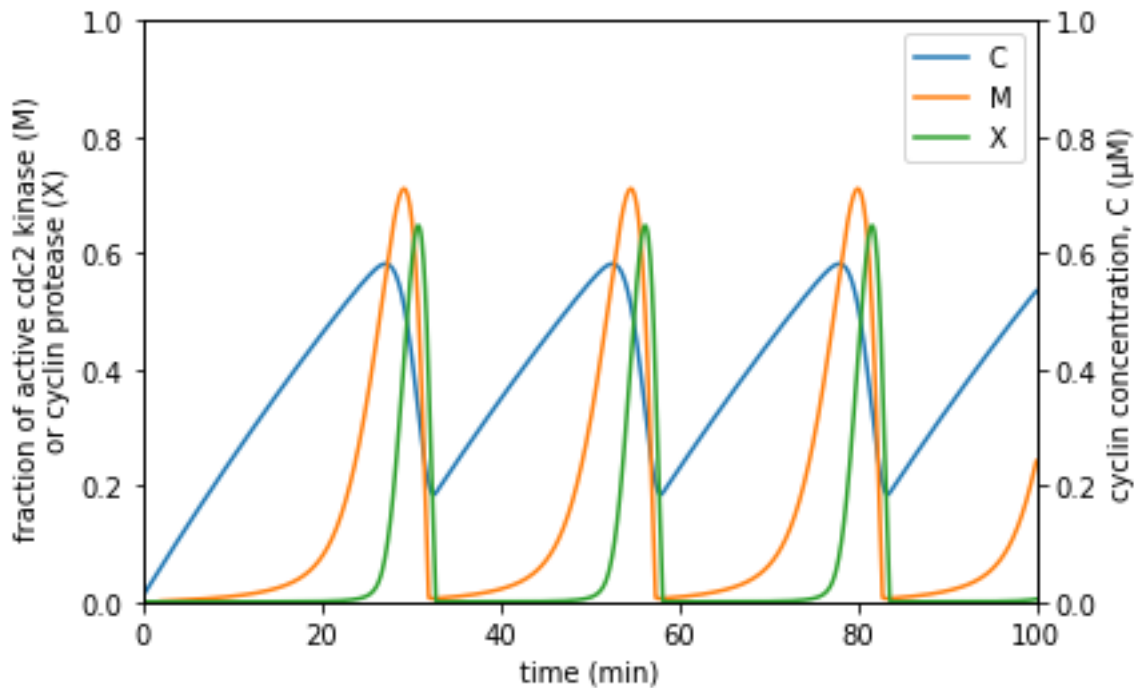
$$Y[n, :] = \mathbf{y}_n$$

- A function `test_integration()` for task B. It should plot both solutions in a single figure.

<sup>1</sup> RK45 (Dormand-Prince [2], [3]) is the default method of `scipy.integrate.solve_ivp`.  
[https://docs.scipy.org/doc/scipy/reference/generated/scipy.integrate.solve\\_ivp.html](https://docs.scipy.org/doc/scipy/reference/generated/scipy.integrate.solve_ivp.html)

## 2. Minimal Cascade Model

Write a python module that reproduces figure 3 of reference [1]. The output could look like this:



- [1] A. Goldbeter, "A minimal cascade model for the mitotic oscillator involving cyclin and cdc2 kinase.," *Proc. Natl. Acad. Sci.*, vol. 88, no. 20, pp. 9107–9111, Oct. 1991, doi: 10.1073/pnas.88.20.9107.
- [2] J. R. Dormand and P. J. Prince, "A family of embedded Runge-Kutta formulae," *J. Comput. Appl. Math.*, vol. 6, no. 1, pp. 19–26, Mar. 1980, doi: 10.1016/0771-050X(80)90013-3.
- [3] L. F. Shampine, "Some Practical Runge-Kutta Formulas," *Math. Comput.*, vol. 46, no. 173, p. 135, Jan. 1986, doi: 10.2307/2008219.