

# tutorial4

May 8, 2021

## 1 Task 1

Write down the differential equation describing the system of chemical equations (assuming a volume of 1).

Using the simple technique described in the supplementary notes of the GRN lecture, we obtain the following ODEs:

$$d/dt[X] = 1 - 2.04[X] + 0.02([X]^2)[Y]$$

$$d/dt[Y] = 2[X] - 0.02([X]^2)[Y]$$

## 2 Task 2

Use a package to solve the differential equation for 500 time units starting from  $X(0) = Y(0) = 0$

The odeint package is used to solve the differential equations and then the results are plotted with matplotlib:

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

```
[27]: def solver(vals, t):
        [x,y] = vals
        dxdt = 1 - (2.04 * x) + (0.02 * (x * x) * y)
        dydt = (2 * x) - (0.02 * (x * x) * y)
        return [dxdt, dydt]

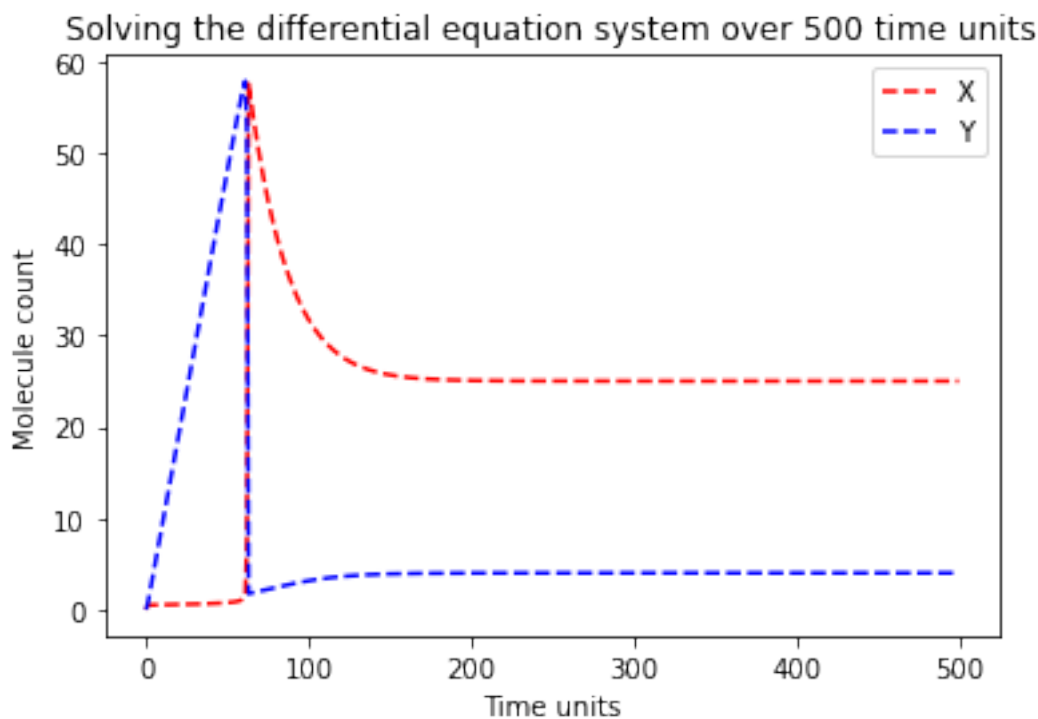
        #initially X = 0 and Y = 0
        init = [0,0]

        #solve the differential equations for 500 time units
        results = odeint(solver, init, range(500))

        #pair the results with each of their timesteps
        xvals = [(time, pair[0]) for time, pair in zip(range(500), results)]
        yvals = [(time, pair[1]) for time, pair in zip(range(500), results)]
```

```
[109]: fig, ax = plt.subplots()
ax.plot(*zip(*xvals), "r--", label = "X")
ax.plot(*zip(*yvals), "b--", label = "Y")
ax.set_xlabel("Time units")
ax.set_ylabel("Molecule count")
ax.title.set_text("Solving the differential equation system over 500 time_
↪units")
ax.legend()
```

[109]: <matplotlib.legend.Legend at 0x7f1edb7e4fd0>



As would be expected by the ODEs, the Y population initially grows at a greater rate than X, until after around 60 time periods it rapidly decreases causing both X and Y to reach an equilibrium state where the X population size stays around 27 and the Y population size stays around 3.

### 3 Task 3

Write a Gillespie algorithm to simulate the same four chemical equations and plot the results for 500 time units

```
[122]: def run_gillespie():
    #initially both 0 again
    x_amt = 0
```

```

y_amt = 0

#Model the chemical equations described
def eq1():
    nonlocal x_amt
    x_amt += 1

def eq2():
    nonlocal x_amt
    nonlocal y_amt
    x_amt -= 1
    y_amt += 1

def eq3():
    nonlocal x_amt
    nonlocal y_amt
    x_amt += 1
    y_amt -= 1

def eq4():
    nonlocal x_amt
    x_amt -= 1

equations = [eq1, eq2, eq3, eq4]

#Put the data in numpy arrays for convenience of element-wise multiplication
equation_rates = np.array([1, 2, 0.02, 0.04])

#Calculate the lhs values using the technique from the GRN lecture
#supplementary notes, make it a callable so that it uses the
#current values for x and y amounts
lhs = lambda: np.array([1, x_amt, (x_amt ** 2) * y_amt, x_amt])

def calc_weights():
    weights = equation_rates * lhs()
    total = sum(weights)
    return [weight/total for weight in weights]

sum_weights = lambda: sum(equation_rates * lhs())

#timestamps paired with population data that will be plotted
x_data = []
y_data = []

t = 0.0
T = 500.0

```

```

while t < T:

    #log of a random value between 0.0 and 1.0 as determined
    #by the algorithm
    dt = (-np.log(np.random.uniform())) / sum_weights()

    #choose the next most likely equation determined by the rates
    #timesed by current amounts, then "run" the chemical equation
    np.random.choice(equations, p = calc_weights())()

    #keep track of the x and y populations at each timestep
    x_data.append((t,x_amt))
    y_data.append((t,y_amt))

    t += dt

return (x_data, y_data)

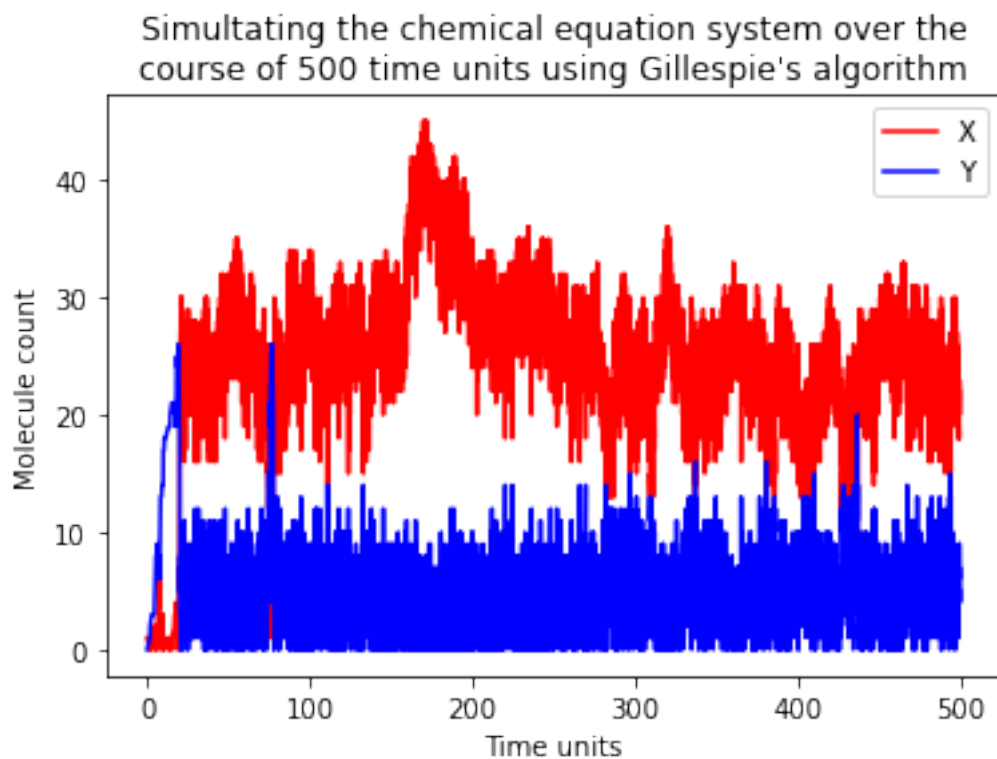
```

```

[127]: def plot_gillespie(x_data, y_data):
        fig, ax = plt.subplots()
        ax.plot(*zip(*x_data), "r", label = "X")
        ax.plot(*zip(*y_data), "b", label = "Y")
        ax.set_xlabel("Time units")
        ax.set_ylabel("Molecule count")
        ax.title.set_text("Simultating the chemical equation system over
↳the\ncourse of 500 time units using Gillespie's algorithm")
        ax.legend()

x_data, y_data = run_gillespie()
plot_gillespie(x_data, y_data)

```



Although the graph does not show the same clear, smooth lines shown by our solutions from the ODE solver, we can recognise the same trends namely the Y population initially growing much faster than X and then immediately decreasing to a smaller amount, followed by X and Y remaining in this “equilibrium” state with the Y population staying smaller than the X population. The equilibrium state is clearer for the Y molecules than the X molecules, however this is to be expected due to the fact the general X population stays larger so any slight fluctuations are much more apparent in the data.