

1D Diffusion simulation

Time step for mass diffusion.

```
In [6]: def time_step(D0, Dt, dt, D, dx):
    Dt[0] = (D0[0] + ((2*(dt)*(D)*((D0[1])-(D0[0]))))/((dx)**2)))
    Dt[-1] = (D0[-1] + ((2*(dt)*(D)*((D0[-1])-(D0[-2]))))/((dx)**2)))
    Dt[1:-1] = (D0[1:-1] + ((dt)*(D)*((D0[2:])+((D0[0:-2])-(2*(D0[1:-1])))))/((dx)**2)))
    return Dt

def stability(dt,D,dx):
    Von_neu = ((2*dt*D)/(dx**2))
    if Von_neu <= 1:
        print("Von_Neumann Stability Condition is satisfied")
        return 1
    else:
        print("Von_Neumann Stability Condition is not satisfied")
        return 0
```

Main code for system simulation for mass diffusion

```
In [ ]: import numpy as np
import time
import matplotlib.pyplot as plt
from matplotlib.colors import LogNorm

nx = 10 #Size of system [unit = A]
dx = 1 #Pos Increment [unit = A]
D = 0.260 #Diffusion coefficient [unit = (A)^2/s]
Ci = 100 #Concentration [unit = (A)^(-3)]

pos = [1,2,3,4,5,6,7,8,9,10]
D0 = np.zeros(10)
Dt = np.zeros(10)
Dt[0]= Ci #Creating high conc gradient in first position

T = 10 #Time [unit = s]
dt = 0.1 #Time increment [unit = s]
Tt = int(T/dt) #Total time steps

Von = stability(dt,D,dx)

x = 0

if Von == 1:

    for x in range(Tt+1):

        D0 = Dt

        #Plotting Position vs Concentration
        title = ("Diffusion at Time: "+str(x*dt)+"s")
        plt.title(title)
        plt.plot(Dt)
        plt.xlabel("Position [A]")
        plt.ylabel(r'Concentration [$A^{-3}$]')
        plt.show()
```

```
#Plotting Position vs Concentration for animation
plt.title(title)
plt.pcolormesh([Dt])
plt.xlabel("Position [A]")
plt.ylabel(r'Concentration [$A^{-3}$]')
ax = plt.gca()
ax.axes.yaxis.set_ticklabels([])
plt.colorbar()
plt.show()

#Diffusion Equation
Dt = time_step(D0, Dt, dt, D, dx)

#Author: Kalith M Ismail.
#Course: Mathematical Modelling of Biological Process and system.
#Organization: NRNU MEPhI__PhysBIO__Moscow__Russian Federation.
#Date: 27/10/2021.
#Mentor: Prof.Dr.Ivanov Dmitry.
```