

```
In [1]: import numpy as np
from scipy.integrate import odeint
from scipy.optimize import curve_fit
from scipy.optimize import differential_evolution
%matplotlib inline
import matplotlib.pyplot as plt
from matplotlib.backends.backend_pdf import PdfPages
import pandas as pd
```

```
In [2]: from endpoint_maker import sqres
from endpoint_maker import func
from endpoint_maker import t
from endpoint_maker import new
from endpoint_maker import lightdata
from endpoint_maker import rawdata
```

```
In [3]: if __name__ == '__main__':
def initparams():
    #bounds = ([0.00001,10000],[0,4],[0.00001,1000],[0.00001,1000],[0.00001,100
    bounds = ([0.00001,1000],[0.00001,1000],[0.00001,10000],[0.00001,10000])
    result = differential_evolution(sqres,bounds,maxiter=100,popsize=20,polish=
    return result
```

```
In [4]: initialp = initparams()
print(initialp)
```

```
message: Maximum number of iterations has been exceeded.
success: False
  fun: 0.17845065007023944
   x: [ 9.840e+02  6.203e-01  1.043e-05  9.624e+03]
  nit: 100
 nfev: 8080
```

```
In [5]: #def tester(t,Kd,n,d2,k2,k3,i):
def tester(t,d2,k2,k3,k7,i):

    inivalues = [1,0,0,0,0,0,0]
    arrayvalues = np.asarray([])

    #for i in range(len(lightdata[:,0])):
    def I(t):
        tindex = t/5
        if tindex > 12241:
            tindex = 12240
        return lightdata[i][int(tindex)]

    #def odes(z,t,Kd,n,d2,k2,k3):
    def odes(z,t,d2,k2,k3,k7):
        Pu,Pb,Pa,mRNA,mCherry1,mCherry2,mCherry3 = z
        d1 = 0.019905
        k1 = 0.08299
        Kd = 90.41
        n = 0.964487
        #d2= 486.67
```

```

#k2= 6.597
#k3= 0.0539

d3 = 0.000077
k4 = 1.25
d4 = 0.000031
k5 = 0.00283
k6 = 0.00283

Pu = z[0]
Pb = z[1]
Pa = z[2]
mRNA = z[3]
mCherry1 = z[4]
mCherry2 = z[5]
mCherry3 = z[6]

dPudt = d1*Pb - k1*I(t)*Pu + k7*I(t)**n/(Kd**n+I(t)**n)*Pa
dPbdt = k1*I(t)*Pu - k2*Pb - d1*Pb + d2*Pa
dPadt = k2*Pb - d2*Pa - k7*I(t)**n/(Kd**n+I(t)**n)*Pa
dmRNA dt = k3*Pa - d3*mRNA
dmCherry1dt = k4*mRNA-(d4 + k5)*mCherry1
dmCherry2dt = k5*mCherry1-(d4+k6)*mCherry2
dmCherry3dt = k6*mCherry2 - d4*mCherry3

return [dPudt,dPbdt,dPadt,dmRNA dt, dmCherry1dt, dmCherry2dt, dmCherry3dt]

#solver = odeint(odes,inivalues,t,args = (Kd,n,d2,k2,k3),hmax=0.1)
solver = odeint(odes,inivalues,t,args = (d2,k2,k3,k7),hmax=0.1)

mCherryout = solver[:,6]
#mCherryout = mCherryout[0:24480:240]
return mCherryout

```

In [6]: `print(initialp.x)`

```
[9.84019831e+02 6.20279494e-01 1.04268611e-05 9.62380222e+03]
```

In [7]: `popt, covt = curve_fit(func,t,new,initialp.x,maxfev=1000000)`

```

#popt, covt = curve_fit(func,t,newdata,initialp.x, maxfev=10000000, bounds=((0.0000
#popt, covt = curve_fit(func,t,newdata,initialp.x, maxfev=1000000)

#Kd,n,d2,k2,k3 = popl
#print('Kd=',Kd,'n=',n,'d2=',d2,'k2=',k2,'k3=',k3)

#d3,k4,d4,k5,k6 = popl
#print('d3=',d3,'k4=',k4,'d4=',d4,'k5=',k5,'k6=',k6)

d2,k2,k3,k7 = popl
print('d2=',d2,'k2=',k2,'k3=',k3,'k7=',k7)

```

C:\Users\Leandra\anaconda3\Lib\site-packages\scipy\integrate\\_odepack\_py.py:248: ODEintWarning: Excess work done on this call (perhaps wrong Dfun type). Run with full\_output = 1 to get quantitative information.  
 warnings.warn(warning\_msg, ODEintWarning)  
 d2= 1002.8572605746836 k2= 0.7423885493341206 k3= 9.221127849160637e-06 k7= 61715.382923276775

```
In [8]: import sys
import numpy
#params = [1,1,1,1,1,1,1]
params = [1,1,1,1]
numpy.set_printoptions(threshold=10)
#model1 = np.asarray(func(t,Kd,n,d2,k2,k3))
model1 = np.asarray(func(t,d2,k2,k3,k7))
print(len(model1))
#print(model1)

#a,b,c,d,e,f,g = params
a,b,c,d = params

ydata = np.asarray(new)
print(len(ydata))

ssr = np.sum((ydata-model1)**2)
#ssr2 = np.sum((ydata-model2)**2)
sst = np.sum((ydata - np.mean(ydata))**2)
R2 = 1 - ssr/sst
#R2_2 = 1 - ssr2/sst

print('R2 is: ', R2)
#print(R2_2)
```

102

102

R2 is: 0.8374633104557399

```
In [9]: pp = PdfPages('multipage.pdf')
ydata = np.asarray(new)

#condition = [1,2,3,4,5,6,7,8,9]
for i in range(2):
    #model = tester(t,Kd,n,d2,k2,k3,i)
    model = tester(t,d2,k2,k3,k7,i)

    #print(model)
    #a,b,c,d,e,f,g= params
    a,b,c,d= params

    plt.plot(t,model,'.', label = 'model')
    #print(model)
    #t = np.linspace(0,34800, num=6961)
    #raw = rawdata[0:13920:240]
    plt.plot(t,raw[i],'.',label = 'data')
    #print(rawdata[i])
    plt.legend()
    pp.savefig()
```

```
plt.show()
```

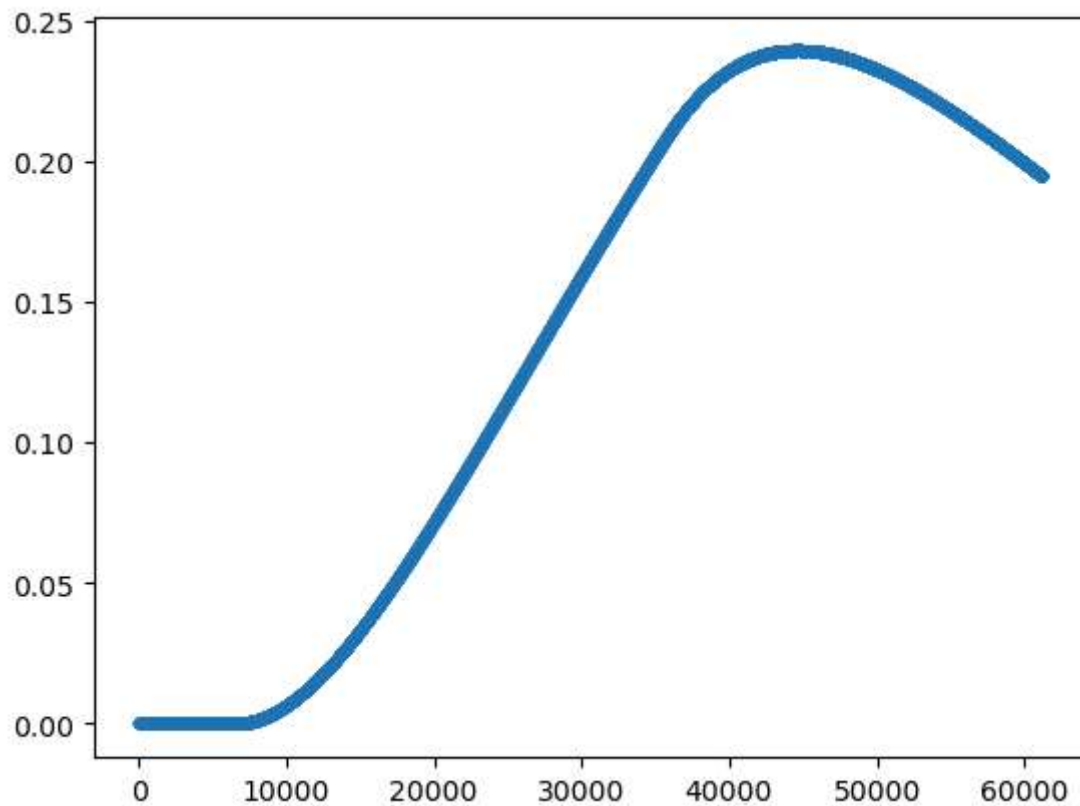
```
pp.close
```

-----  
**NameError** Traceback (most recent call last)

Cell In[9], line 17

```
13 plt.plot(t,model,'.', label = 'model')
14 #print(model)
15 #t = np.linspace(0,34800, num=6961)
16 #raw = rawdata[0:13920:240]
---> 17 plt.plot(t,raw[i],'.',label = 'data')
18 #print(rawdata[i])
19 plt.legend()
```

**NameError**: name 'raw' is not defined



In [ ]: