

Ensayo 2 todo (pTet)

February 13, 2018

```
In [2]: import numpy as np
import matplotlib
import matplotlib.pyplot as plt
%matplotlib inline
from matplotlib import colors
from scipy.interpolate import UnivariateSpline
from scipy.optimize import curve_fit
from scipy import stats
import seaborn as sns
```

```
In [3]: tt=np.fromfile('t', sep=',')
```

```
#arrays replicas glucosa
cfp15211=np.fromfile('p1521gCFP1', sep=',')
rfp15211=np.fromfile('p1521gRFP1', sep=',')
yfp15211=np.fromfile('p1521gYFP1', sep=',')
od15211=np.fromfile('p1521gOD1', sep=',')
cfp15212=np.fromfile('p1521gCFP2', sep=',')
rfp15212=np.fromfile('p1521gRFP2', sep=',')
yfp15212=np.fromfile('p1521gYFP2', sep=',')
od15212=np.fromfile('p1521gOD2', sep=',')
cfp15213=np.fromfile('p1521gCFP3', sep=',')
rfp15213=np.fromfile('p1521gRFP3', sep=',')
yfp15213=np.fromfile('p1521gYFP3', sep=',')
od15213=np.fromfile('p1521gOD3', sep=',')
'''

print(cfp15211.shape)
print(rfp15211.shape)
print(yfp15211.shape)
print(od15211.shape)
print(cfp15212.shape)
print(rfp15212.shape)
print(yfp15212.shape)
print(od15212.shape)
print(cfp15213.shape)
print(rfp15213.shape)
print(yfp15213.shape)
```

```

print(od15213.shape)'''

cfp15231=np.fromfile('p1523gCFP1', sep=',')
rfp15231=np.fromfile('p1523gRFP1', sep=',')
yfp15231=np.fromfile('p1523gYFP1', sep=',')
od15231=np.fromfile('p1523gOD1', sep=',')
cfp15232=np.fromfile('p1523gCFP2', sep=',')
rfp15232=np.fromfile('p1523gRFP2', sep=',')
yfp15232=np.fromfile('p1523gYFP2', sep=',')
od15232=np.fromfile('p1523gOD2', sep=',')
cfp15233=np.fromfile('p1523gCFP3', sep=',')
rfp15233=np.fromfile('p1523gRFP3', sep=',')
yfp15233=np.fromfile('p1523gYFP3', sep=',')
od15233=np.fromfile('p1523gOD3', sep=',')

'''
print(cfp15231.shape)
print(rfp15231.shape)
print(yfp15231.shape)
print(od15231.shape)
print(cfp15232.shape)
print(rfp15232.shape)
print(yfp15232.shape)
print(od15232.shape)
print(cfp15233.shape)
print(rfp15233.shape)
print(yfp15233.shape)
print(od15233.shape)'''

cfp15261=np.fromfile('p1526gCFP1', sep=',')
rfp15261=np.fromfile('p1526gRFP1', sep=',')
yfp15261=np.fromfile('p1526gYFP1', sep=',')
od15261=np.fromfile('p1526gOD1', sep=',')
cfp15262=np.fromfile('p1526gCFP2', sep=',')
rfp15262=np.fromfile('p1526gRFP2', sep=',')
yfp15262=np.fromfile('p1526gYFP2', sep=',')
od15262=np.fromfile('p1526gOD2', sep=',')
cfp15263=np.fromfile('p1526gCFP3', sep=',')
rfp15263=np.fromfile('p1526gRFP3', sep=',')
yfp15263=np.fromfile('p1526gYFP3', sep=',')
od15263=np.fromfile('p1526gOD3', sep=',')

'''
print(cfp15261.shape)
print(rfp15261.shape)
print(yfp15261.shape)
print(od15261.shape)
print(cfp15262.shape)

```

```

print(rfp15262.shape)
print(yfp15262.shape)
print(od15262.shape)
print(cfp15263.shape)
print(rfp15263.shape)
print(yfp15263.shape)
print(od15263.shape)'''

cfp15271=np.fromfile('p1527gCFP1', sep=',')
rfp15271=np.fromfile('p1527gRFP1', sep=',')
yfp15271=np.fromfile('p1527gYFP1', sep=',')
od15271=np.fromfile('p1527gOD1', sep=',')
cfp15272=np.fromfile('p1527gCFP2', sep=',')
rfp15272=np.fromfile('p1527gRFP2', sep=',')
yfp15272=np.fromfile('p1527gYFP2', sep=',')
od15272=np.fromfile('p1527gOD2', sep=',')
cfp15273=np.fromfile('p1527gCFP3', sep=',')
rfp15273=np.fromfile('p1527gRFP3', sep=',')
yfp15273=np.fromfile('p1527gYFP3', sep=',')
od15273=np.fromfile('p1527gOD3', sep=',')

'''
print(cfp15271.shape)
print(rfp15271.shape)
print(yfp15271.shape)
print(od15271.shape)
print(cfp15272.shape)
print(rfp15272.shape)
print(yfp15272.shape)
print(od15272.shape)
print(cfp15273.shape)
print(rfp15273.shape)
print(yfp15273.shape)
print(od15273.shape)'''

#Controles
#Promedios controles glucosa
cfpcg1=np.fromfile('pcgCFP1', sep=',')
rfpcg1=np.fromfile('pcgRFP1', sep=',')
yfpcg1=np.fromfile('pcgYFP1', sep=',')
odcg1=np.fromfile('pcgOD1', sep=',')
cfpcg2=np.fromfile('pcgCFP2', sep=',')
rfpcg2=np.fromfile('pcgRFP2', sep=',')
yfpcg2=np.fromfile('pcgYFP2', sep=',')
odcg2=np.fromfile('pcgOD2', sep=',')
cfpcg3=np.fromfile('pcgCFP3', sep=',')
rfpcg3=np.fromfile('pcgRFP3', sep=',')
yfpcg3=np.fromfile('pcgYFP3', sep=',')

```

```
odcg3=np.fromfile('pcg0D3', sep=',')
'''
```

```
print(cfp1.shape)
print(rfp1.shape)
print(yfp1.shape)
print(odcg1.shape)
print(cfp1.shape)
print(rfp1.shape)
print(yfp1.shape)
print(odcg1.shape)
print(cfp1.shape)
print(rfp1.shape)
print(yfp1.shape)
print(odcg1.shape)'''
```

Out[3]: '\nprint(cfp1.shape)\nprint(rfp1.shape)\nprint(yfp1.shape)\nprint(odcg1.shape)\npr

In [4]: *#Promedios glicerol*

#arrays replicas glicerol

```
cfp1521g1=np.fromfile('p1521g1CFP1', sep=',')
rfp1521g1=np.fromfile('p1521g1RFP1', sep=',')
yfp1521g1=np.fromfile('p1521g1YFP1', sep=',')
od1521g1=np.fromfile('p1521g1OD1', sep=',')
cfp1521g2=np.fromfile('p1521g1CFP2', sep=',')
rfp1521g2=np.fromfile('p1521g1RFP2', sep=',')
yfp1521g2=np.fromfile('p1521g1YFP2', sep=',')
od1521g2=np.fromfile('p1521g1OD2', sep=',')
cfp1521g3=np.fromfile('p1521g1CFP3', sep=',')
rfp1521g3=np.fromfile('p1521g1RFP3', sep=',')
yfp1521g3=np.fromfile('p1521g1YFP3', sep=',')
od1521g3=np.fromfile('p1521g1OD3', sep=',')
'''
```

```
print(cfp1521g1.shape)
print(rfp1521g1.shape)
print(yfp1521g1.shape)
print(od1521g1.shape)
print(cfp1521g2.shape)
print(rfp1521g2.shape)
print(yfp1521g2.shape)
print(od1521g2.shape)
print(cfp1521g3.shape)
print(rfp1521g3.shape)
print(yfp1521g3.shape)
print(od1521g3.shape)'''
```

```
cfp1523g1=np.fromfile('p1523g1CFP1', sep=',')
rfp1523g1=np.fromfile('p1523g1RFP1', sep=',')
```

```

yfp1523g1=np.fromfile('p1523g1YFP1', sep=',')
od1523g1=np.fromfile('p1523g1OD1', sep=',')
cfp1523g2=np.fromfile('p1523g1CFP2', sep=',')
rfp1523g2=np.fromfile('p1523g1RFP2', sep=',')
yfp1523g2=np.fromfile('p1523g1YFP2', sep=',')
od1523g2=np.fromfile('p1523g1OD2', sep=',')
cfp1523g3=np.fromfile('p1523g1CFP3', sep=',')
rfp1523g3=np.fromfile('p1523g1RFP3', sep=',')
yfp1523g3=np.fromfile('p1523g1YFP3', sep=',')
od1523g3=np.fromfile('p1523g1OD3', sep=',')

```

```

'''
print(cfp1523g1.shape)
print(rfp1523g1.shape)
print(yfp1523g1.shape)
print(od1523g1.shape)
print(cfp1523g2.shape)
print(rfp1523g2.shape)
print(yfp1523g2.shape)
print(od1523g2.shape)
print(cfp1523g3.shape)
print(rfp1523g3.shape)
print(yfp1523g3.shape)
print(od1523g3.shape)'''

```

```

cfp1526g1=np.fromfile('p1526g1CFP1', sep=',')
rfp1526g1=np.fromfile('p1526g1RFP1', sep=',')
yfp1526g1=np.fromfile('p1526g1YFP1', sep=',')
od1526g1=np.fromfile('p1526g1OD1', sep=',')
cfp1526g2=np.fromfile('p1526g1CFP2', sep=',')
rfp1526g2=np.fromfile('p1526g1RFP2', sep=',')
yfp1526g2=np.fromfile('p1526g1YFP2', sep=',')
od1526g2=np.fromfile('p1526g1OD2', sep=',')
cfp1526g3=np.fromfile('p1526g1CFP3', sep=',')
rfp1526g3=np.fromfile('p1526g1RFP3', sep=',')
yfp1526g3=np.fromfile('p1526g1YFP3', sep=',')
od1526g3=np.fromfile('p1526g1OD3', sep=',')

```

```

'''
print(cfp1526g1.shape)
print(rfp1526g1.shape)
print(yfp1526g1.shape)
print(od1526g1.shape)
print(cfp1526g2.shape)
print(rfp1526g2.shape)
print(yfp1526g2.shape)
print(od1526g2.shape)
print(cfp1526g3.shape)

```

```

print(rfp1526g3.shape)
print(yfp1526g3.shape)
print(od1526g3.shape)'''

cfp1527g1=np.fromfile('p1527glCFP1', sep=',')
rfp1527g1=np.fromfile('p1527glRFP1', sep=',')
yfp1527g1=np.fromfile('p1527glYFP1', sep=',')
od1527g1=np.fromfile('p1527glOD1', sep=',')
cfp1527g2=np.fromfile('p1527glCFP2', sep=',')
rfp1527g2=np.fromfile('p1527glRFP2', sep=',')
yfp1527g2=np.fromfile('p1527glYFP2', sep=',')
od1527g2=np.fromfile('p1527glOD2', sep=',')
cfp1527g3=np.fromfile('p1527glCFP3', sep=',')
rfp1527g3=np.fromfile('p1527glRFP3', sep=',')
yfp1527g3=np.fromfile('p1527glYFP3', sep=',')
od1527g3=np.fromfile('p1527glOD3', sep=',')

'''

print(cfp1527g1.shape)
print(rfp1527g1.shape)
print(yfp1527g1.shape)
print(od1527g1.shape)
print(cfp1527g2.shape)
print(rfp1527g2.shape)
print(yfp1527g2.shape)
print(od1527g2.shape)
print(cfp1527g3.shape)
print(rfp1527g3.shape)
print(yfp1527g3.shape)
print(od1527g3.shape)'''

#Promedios controles glicerol
cfpcgl1=np.fromfile('pcglCFP1', sep=',')
rfpcgl1=np.fromfile('pcglRFP1', sep=',')
yfpcgl1=np.fromfile('pcglYFP1', sep=',')
odcgl1=np.fromfile('pcglOD1', sep=',')
cfpcgl2=np.fromfile('pcglCFP2', sep=',')
rfpcgl2=np.fromfile('pcglRFP2', sep=',')
yfpcgl2=np.fromfile('pcglYFP2', sep=',')
odcgl2=np.fromfile('pcglOD2', sep=',')
cfpcgl3=np.fromfile('pcglCFP3', sep=',')
rfpcgl3=np.fromfile('pcglRFP3', sep=',')
yfpcgl3=np.fromfile('pcglYFP3', sep=',')
odcgl3=np.fromfile('pcglOD3', sep=',')

'''

print(cfpcgl1.shape)
print(rfpcgl1.shape)

```

```

print(yfpcgl1.shape)
print(odcgl1.shape)
print(cfpcgl1.shape)
print(rfpcgl1.shape)
print(yfpcgl1.shape)
print(odcgl1.shape)
print(cfpcgl1.shape)
print(rfpcgl1.shape)
print(yfpcgl1.shape)
print(odcgl1.shape)'''

```

```

Out[4]: '\nprint(cfpcgl1.shape)\nprint(rfpcgl1.shape)\nprint(yfpcgl1.shape)\nprint(odcgl1.shape)

```

```

In [5]: #Funciones para ajuste Gompertz
def F_sigma(t, A, um,l):
    return ((A*np.exp(-np.exp((((um*np.exp(1))/A)*(1-t))+1))))

def Function_fit(xdata,ydata,init,end,func=F_sigma,ParamBounds=([0,0,0],[3,1,300]), titl
    Y_fit={}

    z,_=curve_fit(func,xdata[init:end], ydata[init:end], bounds=ParamBounds)

    print(z)

    evalF=func(xdata,z[0],z[1],z[2])

    plt.figure()
    plt.plot(xdata, ydata, '.',label='OD')
    plt.plot(xdata, evalF, '-',label='Ajuste')
    plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
    plt.title(title)
    plt.ylabel('Abs(nm)')
    plt.xlabel('Tiempo(min)')
    lgd=plt.legend(loc='lower right')
    plt.show()

    Y_fit=evalF,z

    return(Y_fit)

```

```

In [6]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#control glucosa rep 1
y1 = np.log(odcgl)-np.log(np.min(odcgl))
print('Min OD = %e'%((np.min(odcgl))))
evaly, params=Function_fit(tt,y1,0,-1,title = 'Ajuste control glucosa 0,4% E2R1')
A1 = params[0]

```

```

um1=params[1]
l1=params[2]
print('A=%e'%(A1))
print('um=%e'%(um1))
print('l=%e'%(l1))

#Cálculo datos para determinar extensión de la fase exponencial
tm1=((A1/(np.exp(1)*um1))+l1)
print('Tm=%e'%(tm1))
t21=((np.log(2))/um1)
print('doubpe=%e'%(t21))
extdp1=2*t21
print('ext=%e'%extdp1)
ttot1=tm1+extdp1
print('Tfinal=%e'%ttot1)

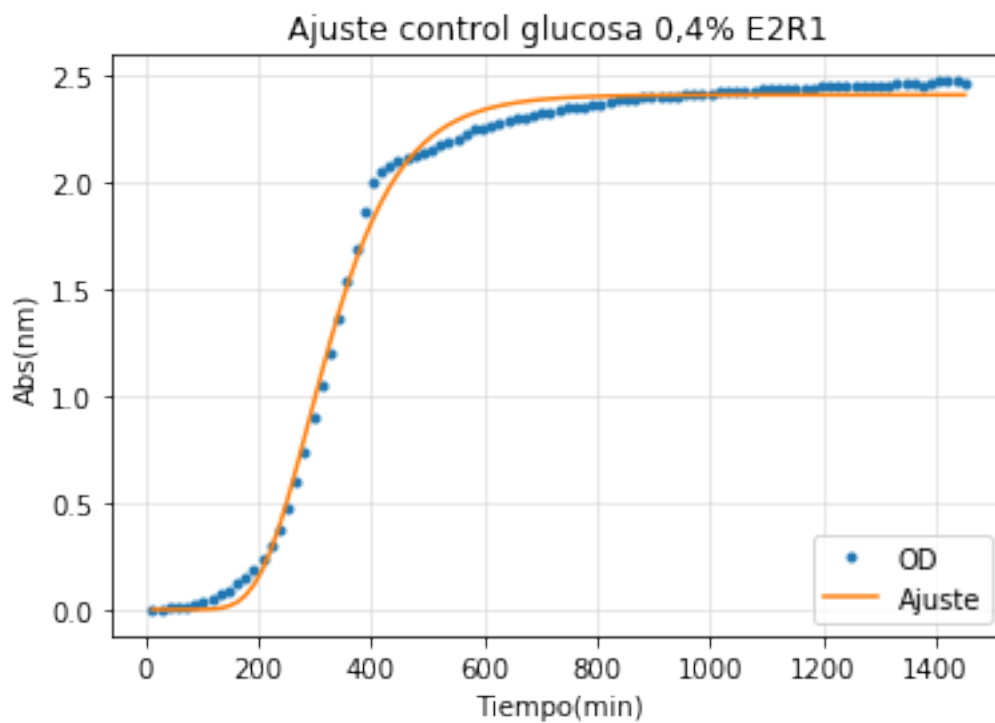
#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[19]
y2=tt[29]
plt.figure()
plt.title('Control Glucosa 0,4% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,odcg1,label='OD control E2R1 ')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[19:30],odcg1[19:30],label='OD control E2R1')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')

```

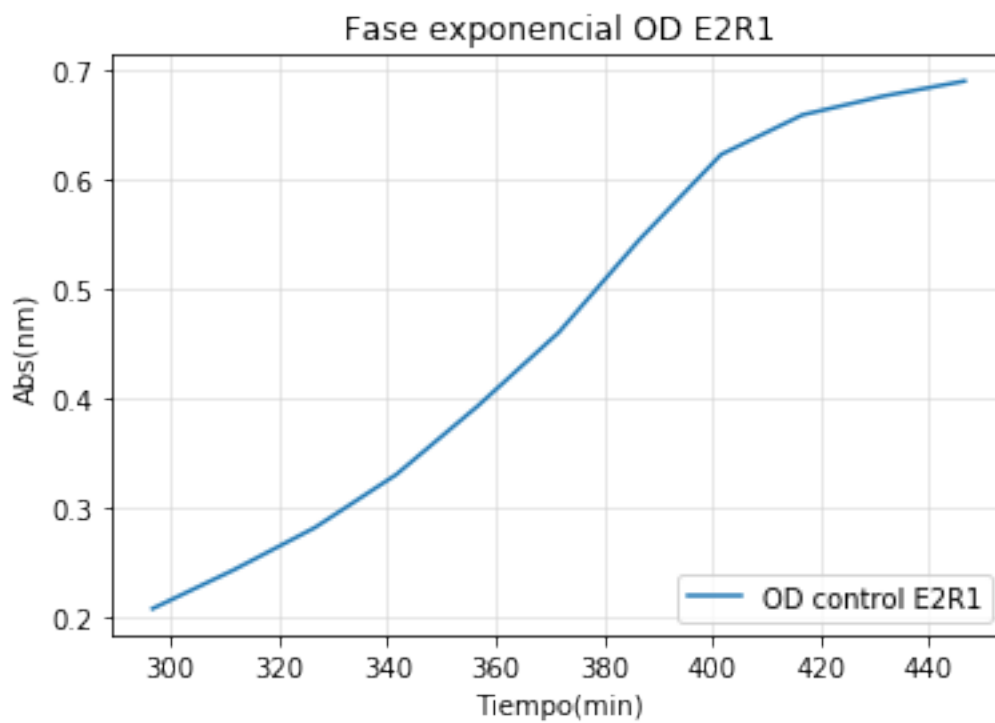
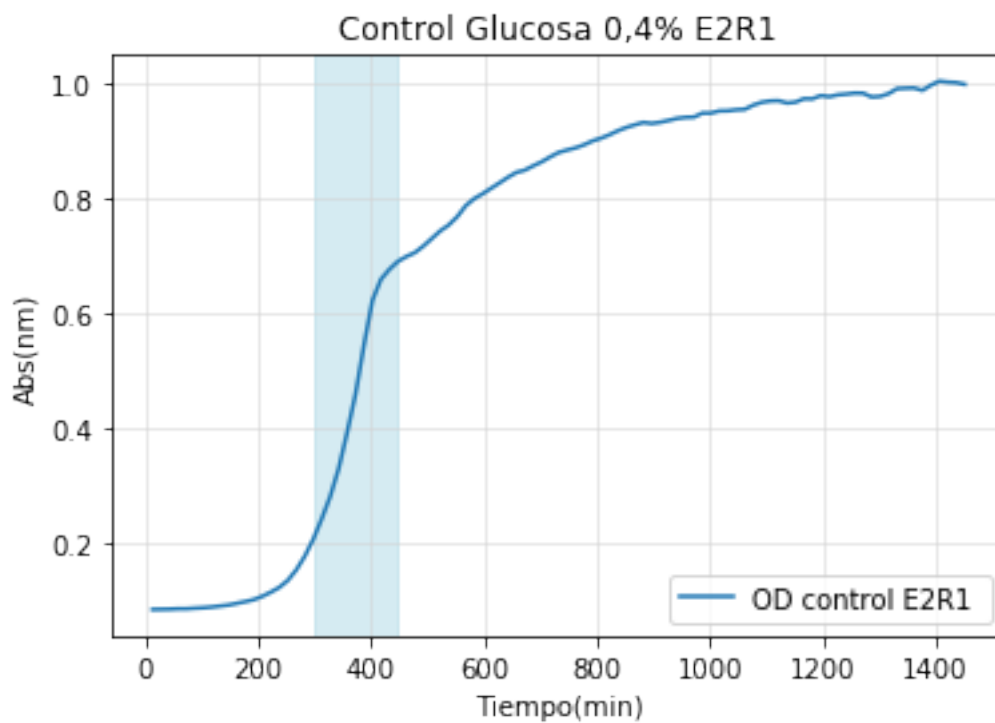
Min OD = 8.475000e-02

[2.40766598e+00 1.00663667e-02 2.01163416e+02]



```
A=2.407666e+00  
um=1.006637e-02  
l=2.011634e+02  
Tm=2.891525e+02  
doubpe=6.885773e+01  
ext=1.377155e+02  
Tfinal=4.268680e+02
```

```
Out[6]: <matplotlib.legend.Legend at 0x267e9a887f0>
```



```

In [7]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
        #control glucosa rep 2
        y2= np.log(odcg2)-np.log(np.min(odcg2))
        print('Min OD = %e'%((np.min(odcg2))))
        evaly, params=Function_fit(tt,y2,0,-1, title = 'Ajuste control glucosa 0,4% E2R2')
        A2 = params[0]
        um2=params[1]
        l2=params[2]
        print('A=%e'%(A2))
        print('um=%e'%(um2))
        print('l=%e'%(l2))

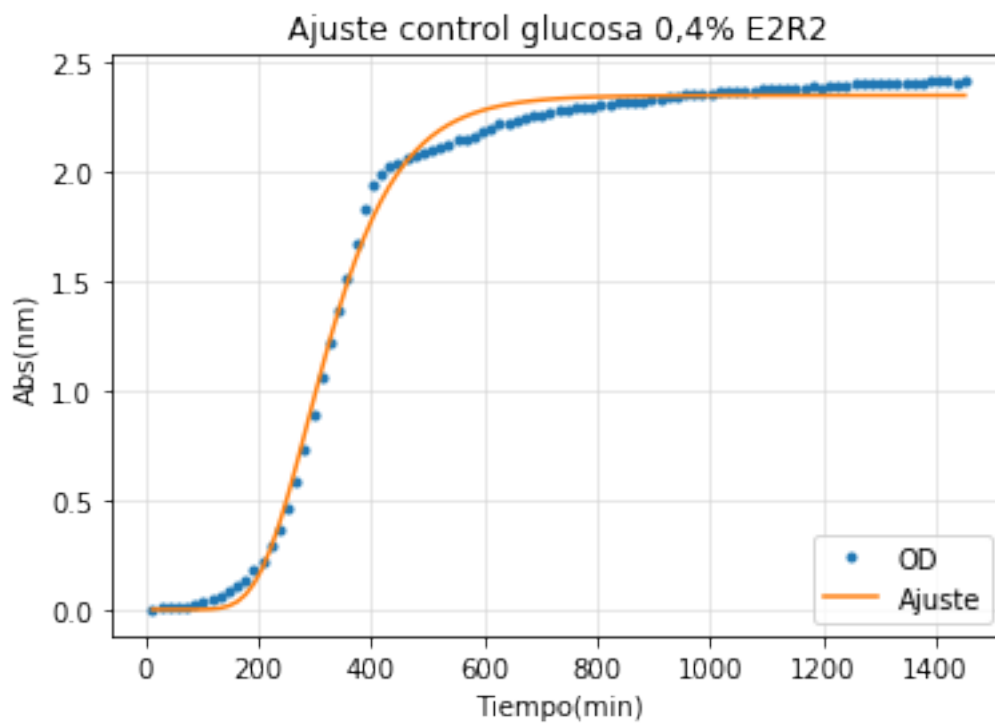
        #Cálculo datos para determinar extensión de la fase exponencial
        tm2=((A2/(np.exp(1)*um2))+l2)
        print('Tm=%e'%(tm2))
        t22=((np.log(2))/um2)
        print('doubpe=%e'%(t22))
        extdp2=2*t22
        print('ext=%e'%extdp2)
        ttot2=tm2+extdp2
        print('Tfinal=%e'%ttot2)

        #Delimitación fase exponencial en grafico con OD/tiempo
        y1=tt[19]
        y2=tt[29]
        plt.figure()
        plt.title('Control Glucosa 0,4% E2R2')
        plt.xlabel('Tiempo(min)')
        plt.ylabel('Abs(nm)')
        plt.plot(tt,odcg2,label='OD control E2R2 ')
        plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
        plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
        plt.legend(loc='lower right')

        #Fase exponencial OD/tiempo
        plt.figure()
        plt.title('Fase exponencial OD E2R2')
        plt.xlabel('Tiempo(min)')
        plt.ylabel('Abs(nm)')
        plt.plot(tt[19:30],odcg2[19:30],label='OD control E2R2')
        plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
        plt.legend(loc='lower right')

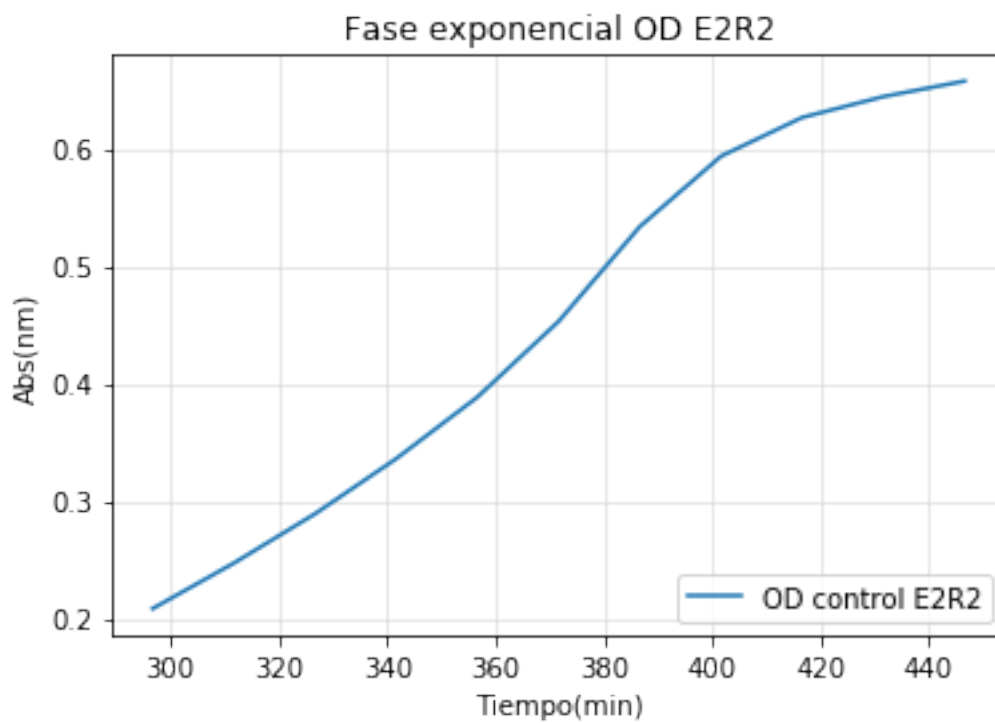
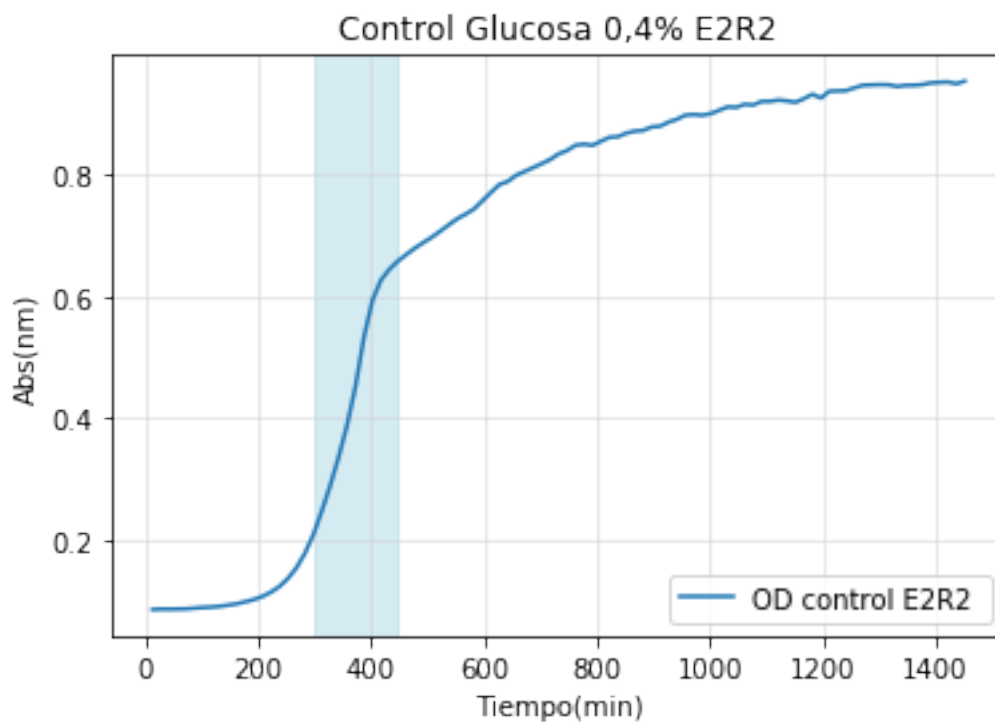
Min OD = 8.550000e-02
[ 2.34701846e+00  9.83004768e-03  1.99784486e+02]

```



```
A=2.347018e+00  
um=9.830048e-03  
l=1.997845e+02  
Tm=2.876192e+02  
doubpe=7.051310e+01  
ext=1.410262e+02  
Tfinal=4.286455e+02
```

```
Out[7]: <matplotlib.legend.Legend at 0x267eac4b748>
```



```

In [8]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
        #control glucosa rep 3
        y3= np.log(odcg3)-np.log(np.min(odcg3))
        print('Min OD = %e'%((np.min(odcg3))))
        evaly, params=Function_fit(tt,y3,0,-1, title = 'Ajuste control glucosa 0,4% E2R3')
        A3= params[0]
        um3=params[1]
        l3=params[2]
        print('A=%e'%(A3))
        print('um=%e'%(um3))
        print('l=%e'%(l3))

        #Cálculo datos para determinar extensión de la fase exponencial
        tm3=((A3/(np.exp(1)*um3))+l3)
        print('Tm=%e'%(tm3))
        t23=((np.log(2))/um3)
        print('doubpe=%e'%(t23))
        extdp3=2*t23
        print('ext=%e'%extdp3)
        ttot3=tm3+extdp3
        print('Tfinal=%e'%ttot3)

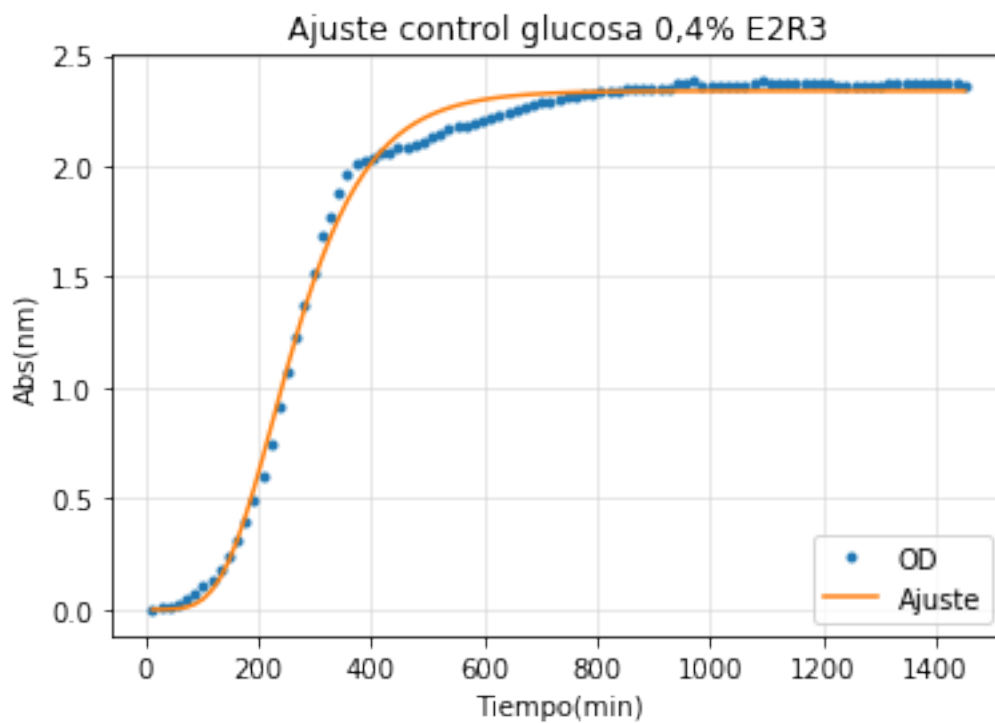
        #Delimitación fase exponencial en grafico con OD/tiempo
        y1=tt[15]
        y2=tt[26]
        plt.figure()
        plt.title('Control Glucosa 0,4% E2R3')
        plt.xlabel('Tiempo(min)')
        plt.ylabel('Abs(nm)')
        plt.plot(tt,odcg3,label='OD control E2R3')
        plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
        plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
        plt.legend(loc='lower right')

        #Fase exponencial OD/tiempo
        plt.figure()
        plt.title('Fase exponencial OD E2R3')
        plt.xlabel('Tiempo(min)')
        plt.ylabel('Abs(nm)')
        plt.plot(tt[15:27],odcg3[15:27],label='OD control E2R3')
        plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
        plt.legend(loc='lower right')

```

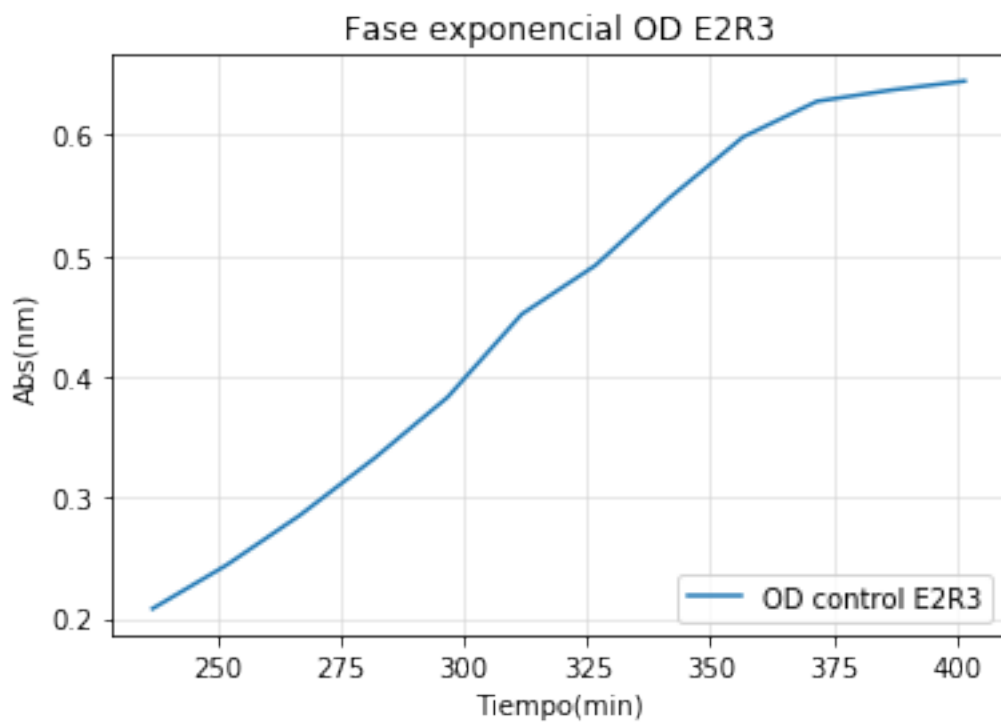
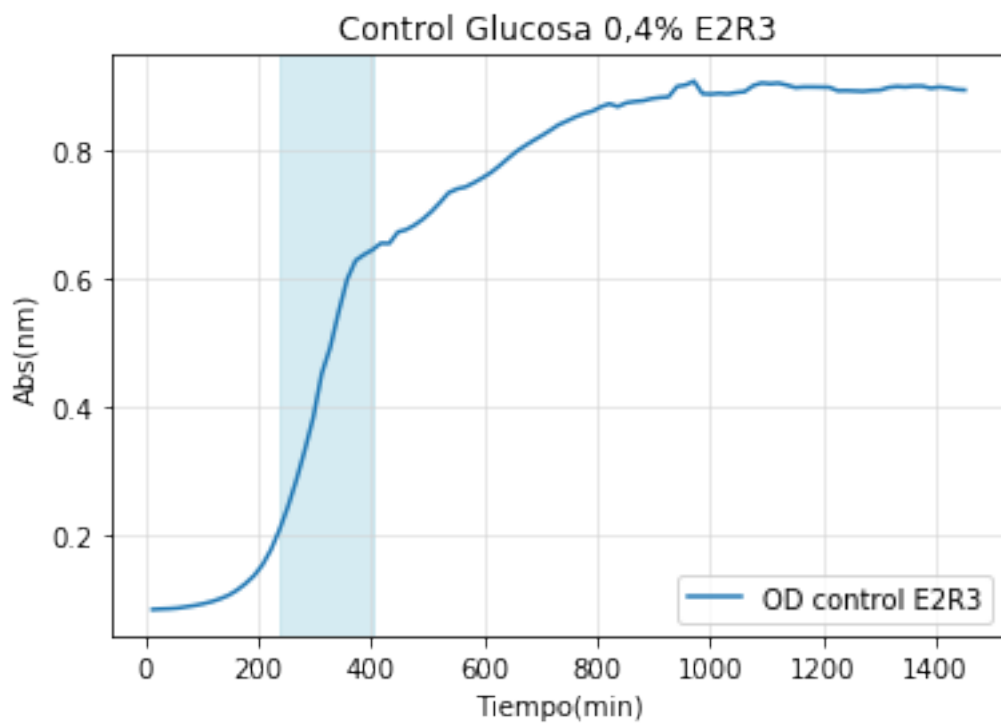
Min OD = 8.375000e-02

[2.33929203e+00 9.40512328e-03 1.34262384e+02]



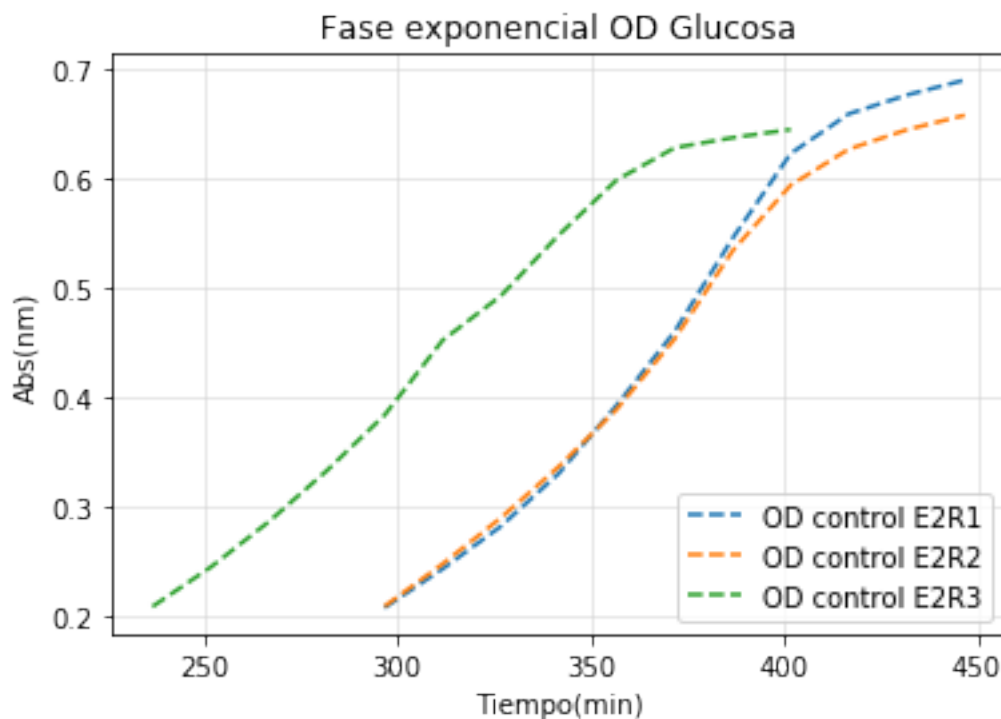
```
A=2.339292e+00  
um=9.405123e-03  
l=1.342624e+02  
Tm=2.257633e+02  
doubpe=7.369889e+01  
ext=1.473978e+02  
Tfinal=3.731611e+02
```

```
Out[8]: <matplotlib.legend.Legend at 0x267eae5cf98>
```




```
In [9]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD Glucosa')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[19:30],odcg1[19:30], '--',label='OD control E2R1')
plt.plot(tt[19:30],odcg2[19:30], '--',label='OD control E2R2')
plt.plot(tt[15:27],odcg3[15:27], '--',label='OD control E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')
```

Out[9]: <matplotlib.legend.Legend at 0x267eaf76f60>



```
In [10]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#control glicerol rep 1
y4= np.log(odcg1)-np.log(np.min(odcg1))
print('Min OD = %e'%((np.min(odcg1))))
evaly, params=Function_fit(tt,y4,0,-1, title = 'Ajuste control glicerol 0,2% E2R1')
A4= params[0]
um4=params[1]
l4=params[2]
print('A=%e'%(A4))
print('um=%e'%(um4))
print('l=%e'%(l4))
```

```

#Cálculo datos para determinar extensión de la fase exponencial
tm4=((A4/(np.exp(1)*um4))+14)
print('Tm=%e'%(tm4))
t24=((np.log(2))/um4)
print('doubpe=%e'%(t24))
extdp4=2.5*t24
print('ext=%e'%extdp4)
ttot4=tm4+extdp4
print('Tfinal=%e'%ttot4)

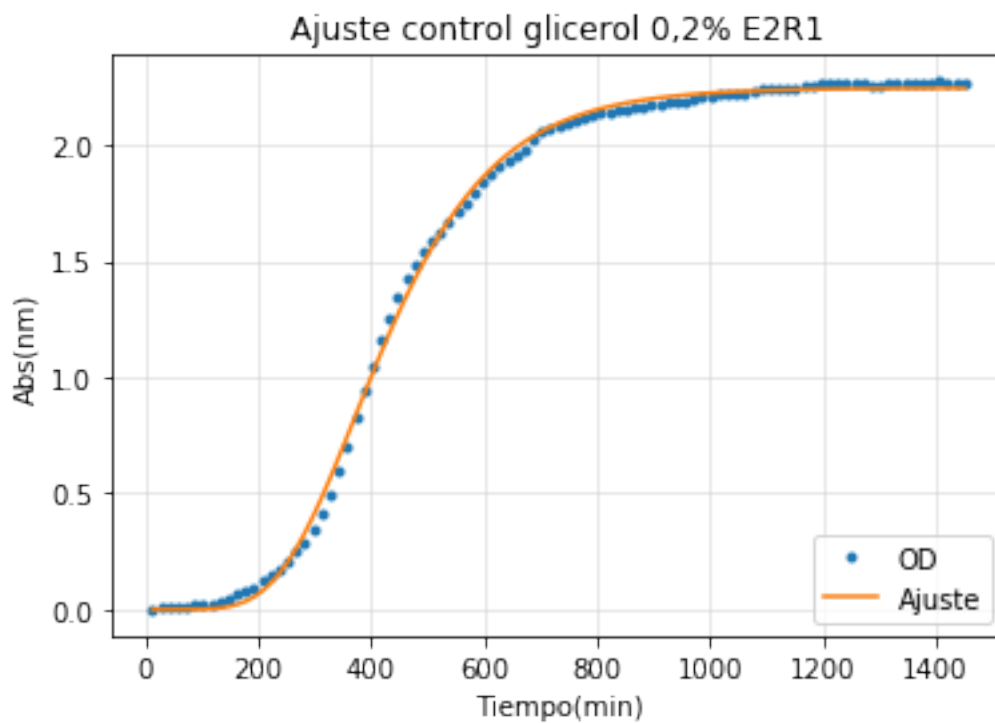
#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[24]
y2=tt[44]
plt.figure()
plt.title('Control Glicerol 0,2% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,odcgl1,label='OD control E2R1')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[24:45],odcgl1[24:45],label='OD control E2R1')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')

```

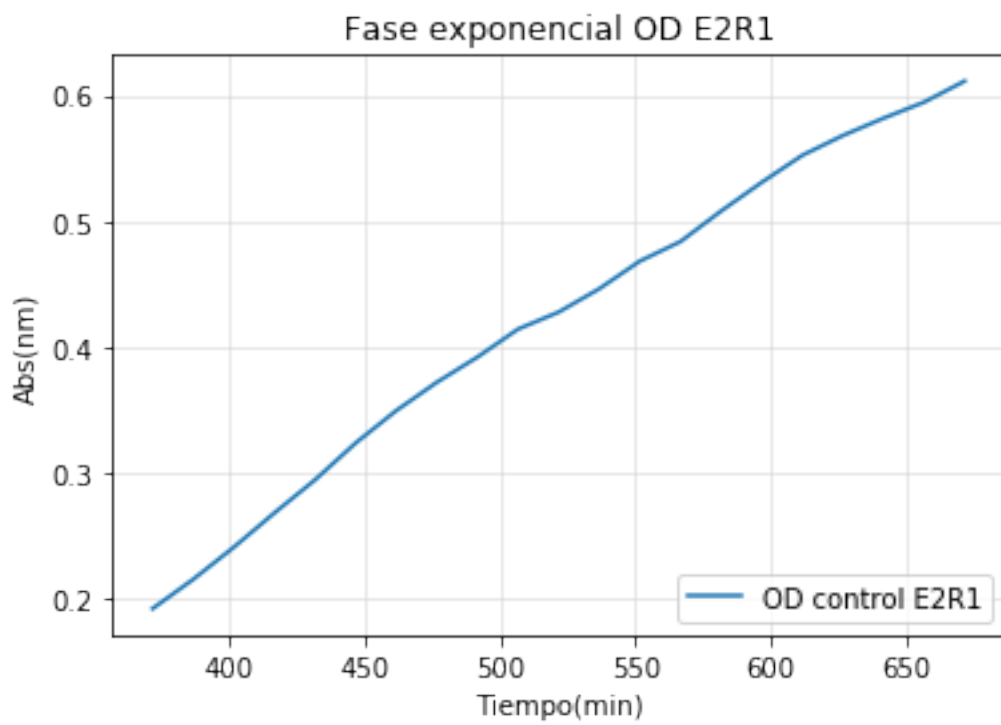
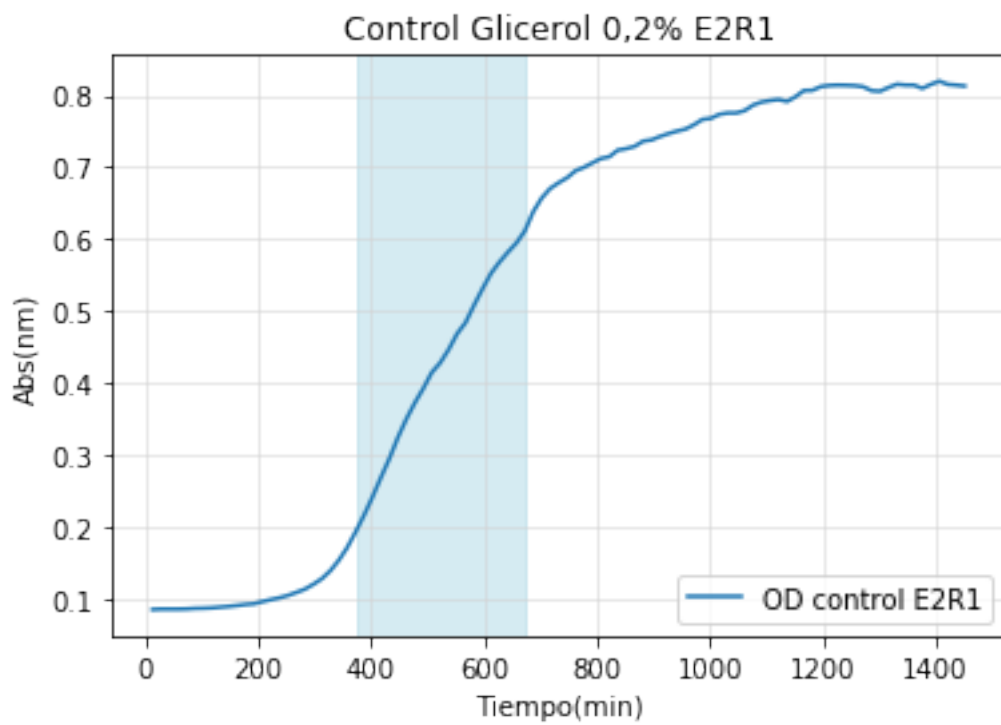
Min OD = 8.425000e-02

[2.24595733e+00 6.08336131e-03 2.34248586e+02]



```
A=2.245957e+00  
um=6.083361e-03  
l=2.342486e+02  
Tm=3.700685e+02  
doubpe=1.139415e+02  
ext=2.848537e+02  
Tfinal=6.549222e+02
```

```
Out[10]: <matplotlib.legend.Legend at 0x267eaec0f0>
```



```

In [11]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#control glicerol rep 2
y5= np.log(odcgl2)-np.log(np.min(odcgl2))
print('Min OD = %e'%((np.min(odcgl2))))
evaly, params=Function_fit(tt,y5,0,-1, title = 'Ajuste control glicerol 0,2% E2R2')
A5= params[0]
um5=params[1]
l5=params[2]
print('A=%e'%(A5))
print('um=%e'%(um5))
print('l=%e'%(l5))

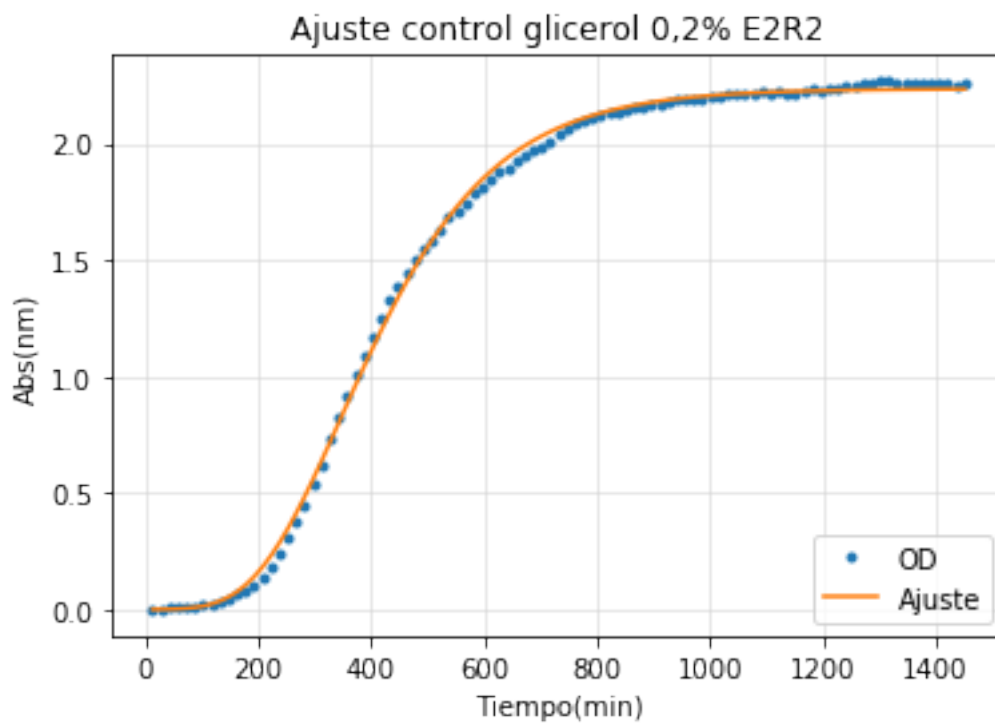
#Cálculo datos para determinar extensión de la fase exponencial
tm5=((A5/(np.exp(1)*um5))+l5)
print('Tm=%e'%(tm5))
t25=((np.log(2))/um5)
print('doubpe=%e'%(t25))
extdp5=2.5*t25
print('ext=%e'%extdp5)
ttot5=tm5+extdp5
print('Tfinal=%e'%ttot5)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[23]
y2=tt[45]
plt.figure()
plt.title('Control Glicerol 0,2% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,odcgl2,label='OD control E2R2')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[23:46],odcgl2[23:46],label='OD control E2R2')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

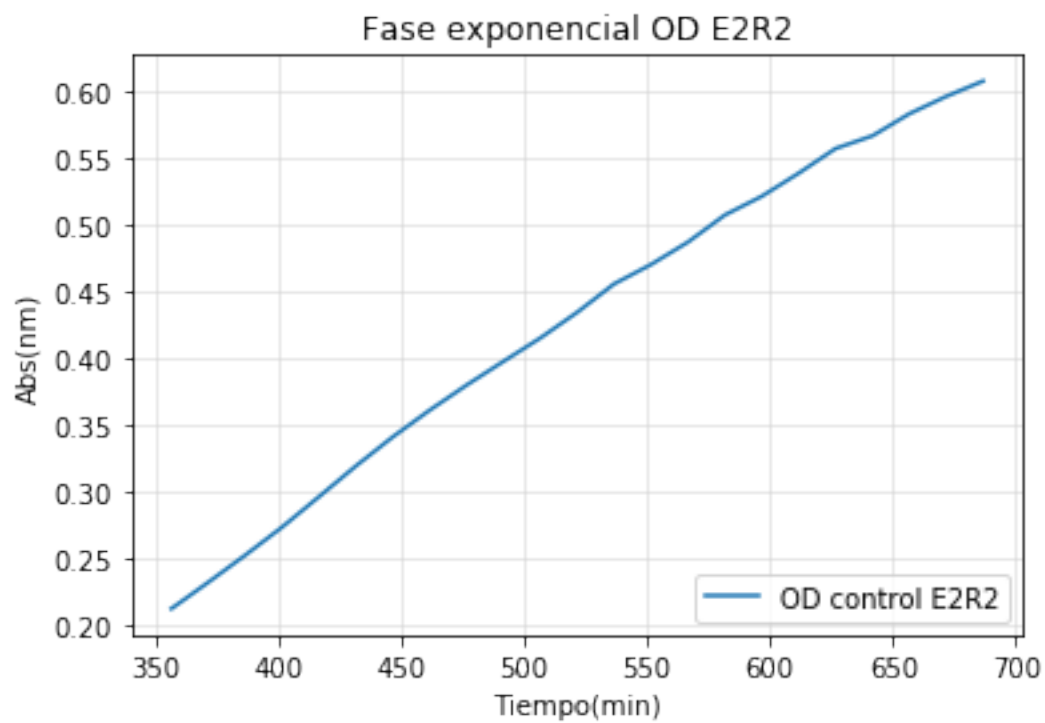
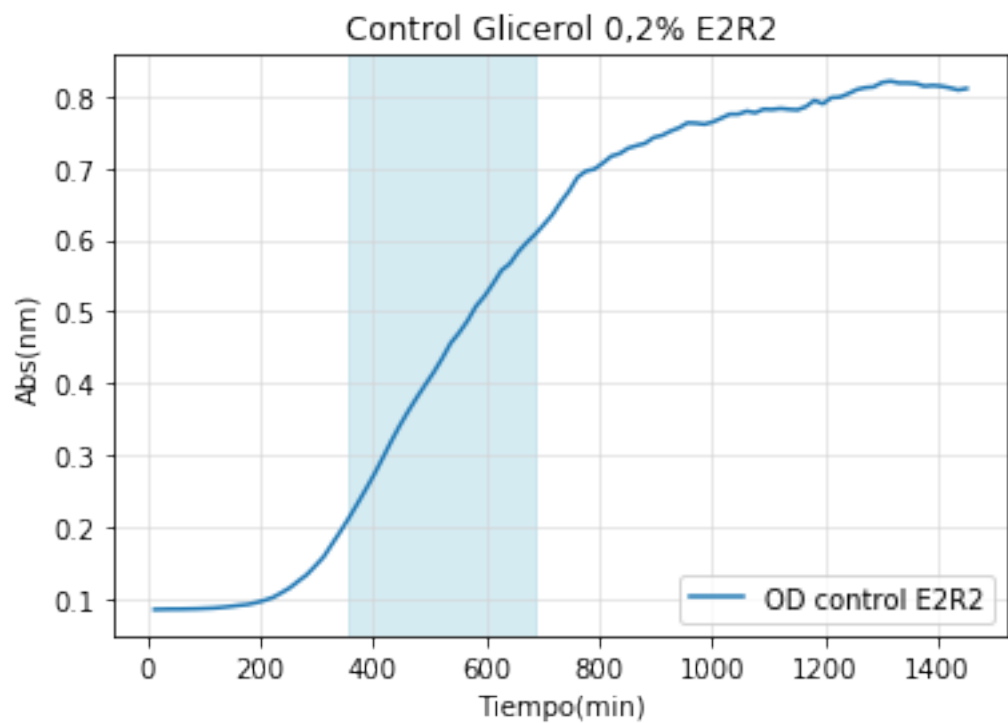
Min OD = 8.500000e-02
[ 2.23521178e+00  5.43425313e-03  1.94280385e+02]

```



```
A=2.235212e+00  
um=5.434253e-03  
l=1.942804e+02  
Tm=3.455962e+02  
doubpe=1.275515e+02  
ext=3.188788e+02  
Tfinal=6.644750e+02
```

```
Out[11]: <matplotlib.legend.Legend at 0x267e9a637f0>
```



```

In [12]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#control glicerol rep 3
y6= np.log(odcgl3)-np.log(np.min(odcgl3))
print('Min OD = %e'%((np.min(odcgl3))))
evaly, params=Function_fit(tt,y6,0,-1, title = 'Ajuste control glicerol 0,2% E2R3')
A6= params[0]
um6=params[1]
l6=params[2]
print('A=%e'%(A6))
print('um=%e'%(um6))
print('l=%e'%(l6))

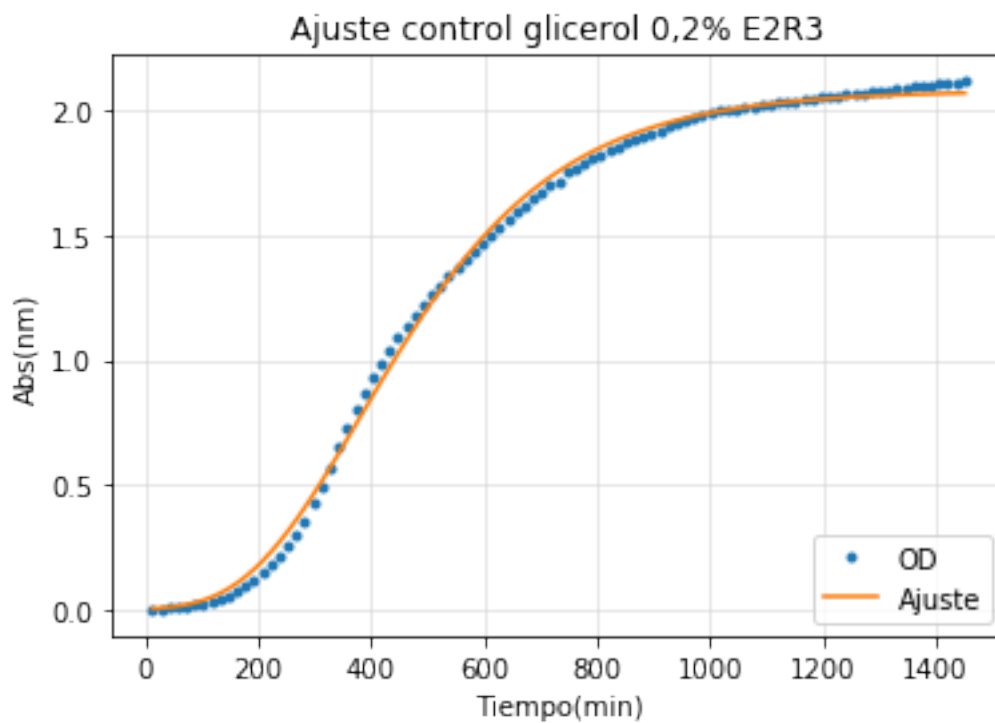
#Cálculo datos para determinar extensión de la fase exponencial
tm6=((A6/(np.exp(1)*um6))+l6)
print('Tm=%e'%(tm6))
t26=((np.log(2))/um6)
print('doubpe=%e'%(t26))
extdp6=2.5*t26
print('ext=%e'%extdp6)
ttot6=tm6+extdp6
print('Tfinal=%e'%ttot6)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[25]
y2=tt[56]
plt.figure()
plt.title('Control Glicerol 0,2% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,odcgl3,label='OD control E2R3')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[25:57],odcgl3[25:57],label='OD control E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

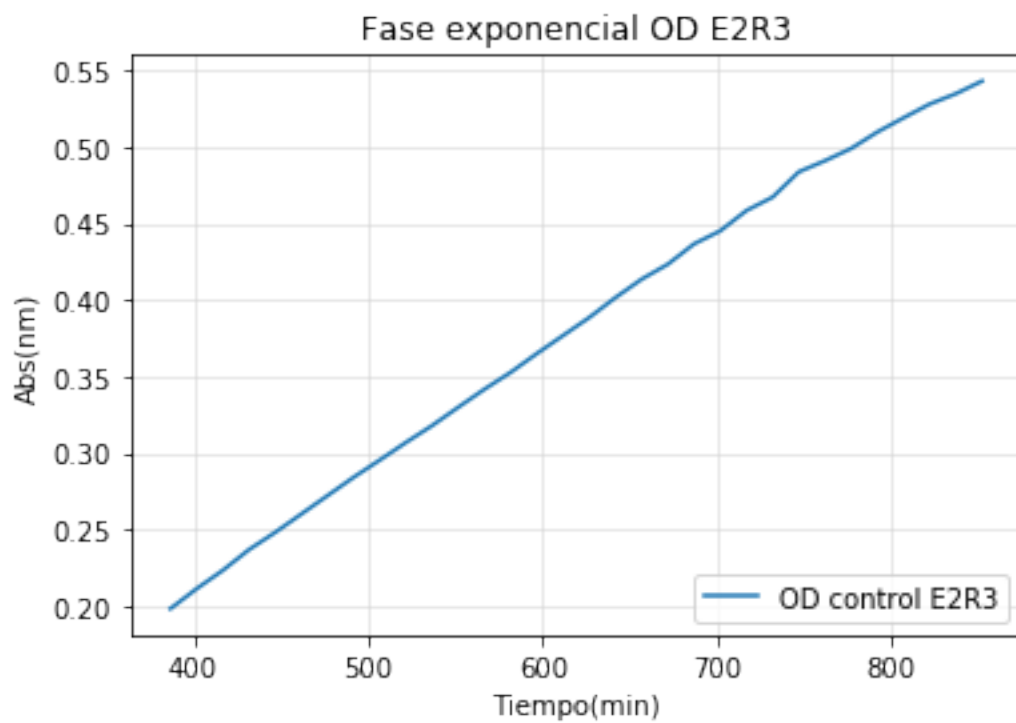
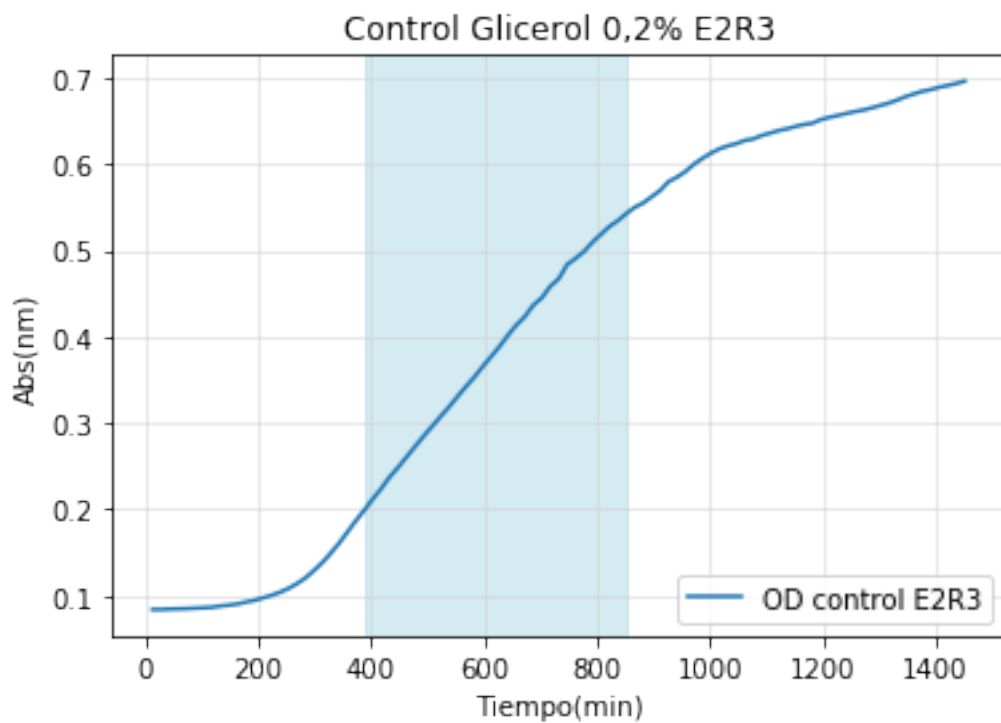
Min OD = 8.375000e-02
[ 2.07943812e+00  3.85150055e-03  1.79490460e+02]

```

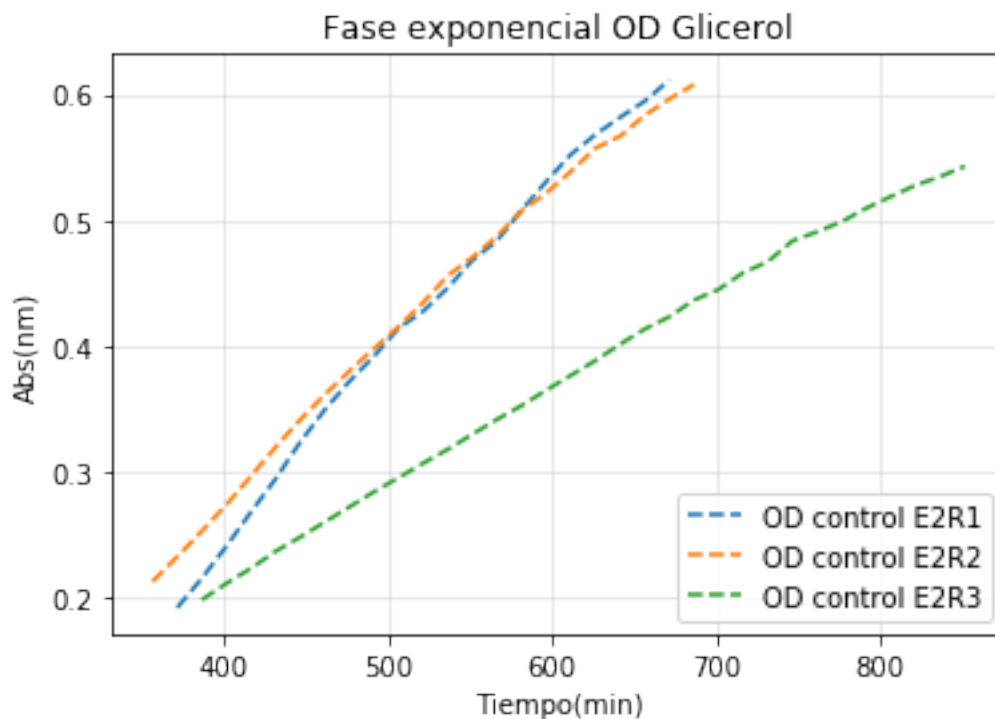
```
A=2.079438e+00  
um=3.851501e-03  
l=1.794905e+02  
Tm=3.781098e+02  
doubpe=1.799681e+02  
ext=4.499202e+02  
Tfinal=8.280300e+02
```

```
Out[12]: <matplotlib.legend.Legend at 0x267e9bbce48>
```



```
In [13]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD Glicerol')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[24:45],odcg11[24:45], '--', label='OD control E2R1')
plt.plot(tt[23:46],odcg12[23:46], '--', label='OD control E2R2')
plt.plot(tt[25:57],odcg13[25:57], '--', label='OD control E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')
```

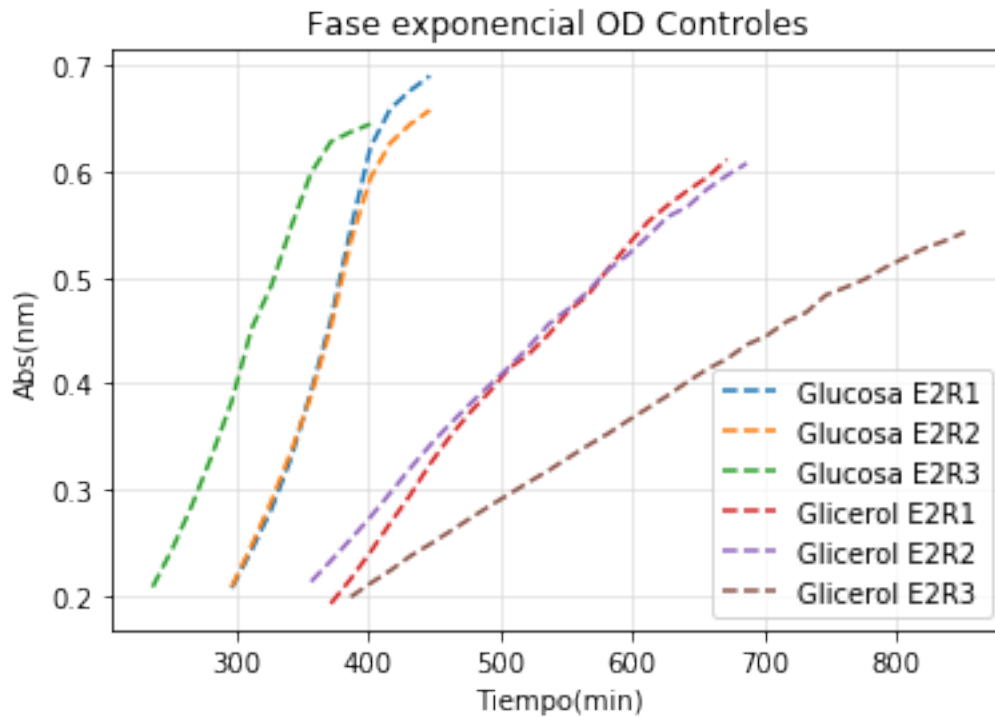
Out[13]: <matplotlib.legend.Legend at 0x267eb181e48>



```
In [14]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD Controles')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[19:30],odcg1[19:30], '--', label='Glucosa E2R1')
plt.plot(tt[19:30],odcg2[19:30], '--', label='Glucosa E2R2')
plt.plot(tt[15:27],odcg3[15:27], '--', label='Glucosa E2R3')
plt.plot(tt[24:45],odcg11[24:45], '--', label='Glicerol E2R1')
plt.plot(tt[23:46],odcg12[23:46], '--', label='Glicerol E2R2')
plt.plot(tt[25:57],odcg13[25:57], '--', label='Glicerol E2R3')
```

```
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')
```

Out[14]: <matplotlib.legend.Legend at 0x267eb23bcf8>



```
In [15]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-J23101-std glucosa rep 1
y7= np.log(od15211)-np.log(np.min(od15211))
print('Min OD = %e'%((np.min(od15211))))
evaly, params=Function_fit(tt,y7,0,-1,title = 'Ajuste pTet-J23101-J23101 glucosa 0,4% E
A7 = params[0]
um7=params[1]
l7=params[2]
print('A=%e'%(A7))
print('um=%e'%(um7))
print('l=%e'%(l7))

#Cálculo datos para determinar extensión de la fase exponencial
tm7=((A7/(np.exp(1)*um7))+l7)
print('Tm=%e'%(tm7))
t27=((np.log(2))/um7)
print('doubpe=%e'%(t27))
extdp7=2*t27
print('ext=%e'%extdp7)
```

```

ttot7=tm7+extdp7
print('Tfinal=%e'%ttot7)

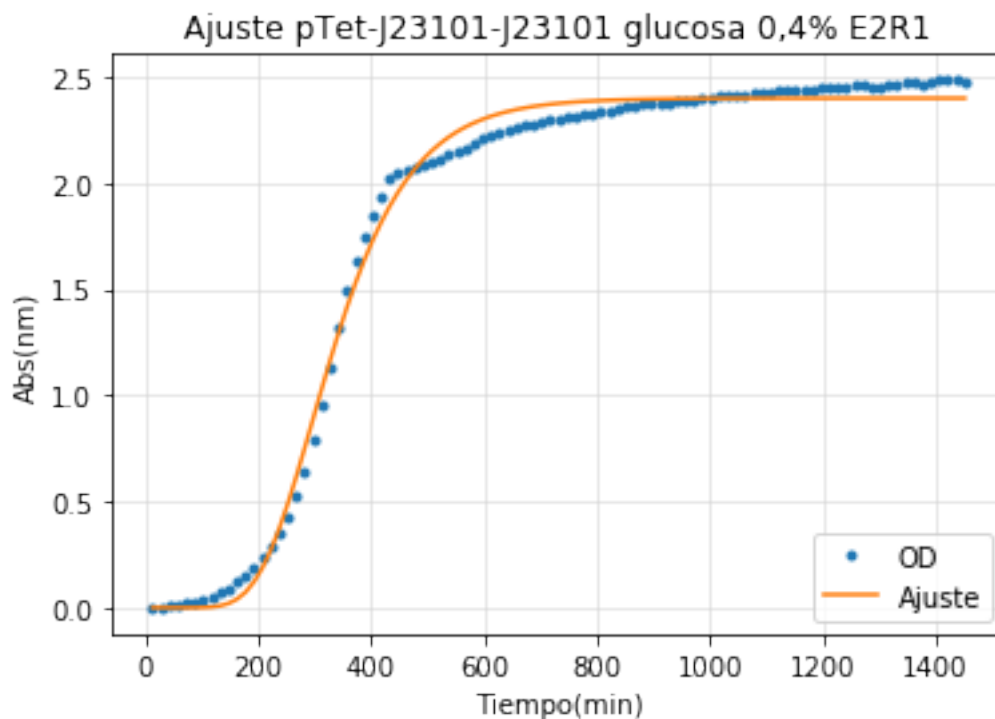
#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[19]
y2=tt[30]
plt.figure()
plt.title('pTet-J23101-J23101 Glucosa 0,4% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od15211,label='OD pTet-J23101-J23101 E2R1 ')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[19:31],od15211[19:31],label='OD pTet-J23101-J23101 E2R1')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

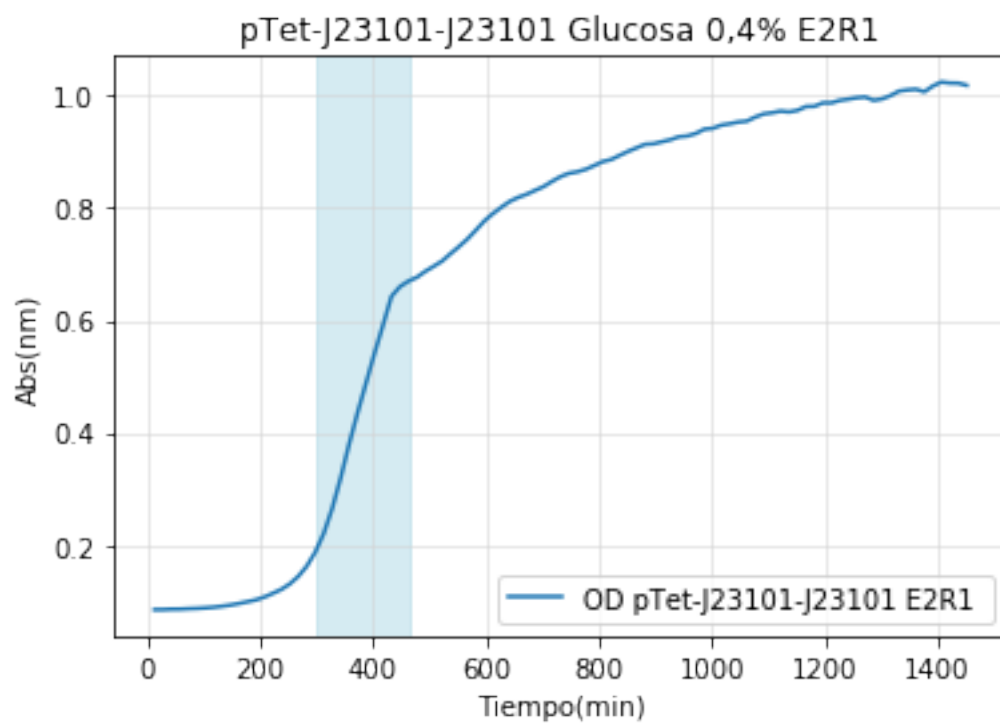
Min OD = 8.530000e-02

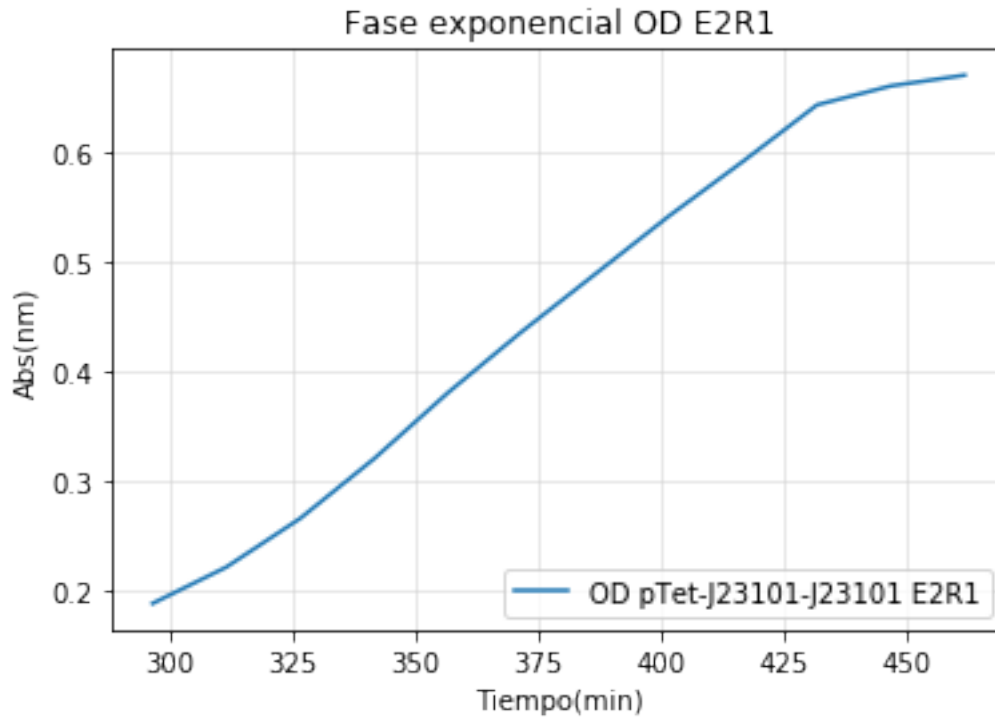
[2.39988520e+00 9.27231338e-03 2.00377623e+02]



A=2.399885e+00
um=9.272313e-03
l=2.003776e+02
Tm=2.955932e+02
doubpe=7.475450e+01
ext=1.495090e+02
Tfinal=4.451022e+02

Out[15]: <matplotlib.legend.Legend at 0x267e9657668>





```
In [16]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-std-std glucosa rep 2
y8= np.log(od15212)-np.log(np.min(od15212))
print('Min OD = %e'%((np.min(od15212))))
evaly, params=Function_fit(tt,y8,0,-1,title = 'Ajuste pTet-J23101-J23101 glucosa 0,4% E
A8= params[0]
um8=params[1]
l8=params[2]
print('A=%e'%(A8))
print('um=%e'%(um8))
print('l=%e'%(l8))

#Cálculo datos para determinar extensión de la fase exponencial
tm8=((A8/(np.exp(1)*um8))+l8)
print('Tm=%e'%(tm8))
t28=((np.log(2))/um8)
print('doubpe=%e'%(t28))
extdp8=2*t28
print('ext=%e'%extdp8)
ttot8=tm8+extdp8
print('Tfinal=%e'%ttot8)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[19]
```

```

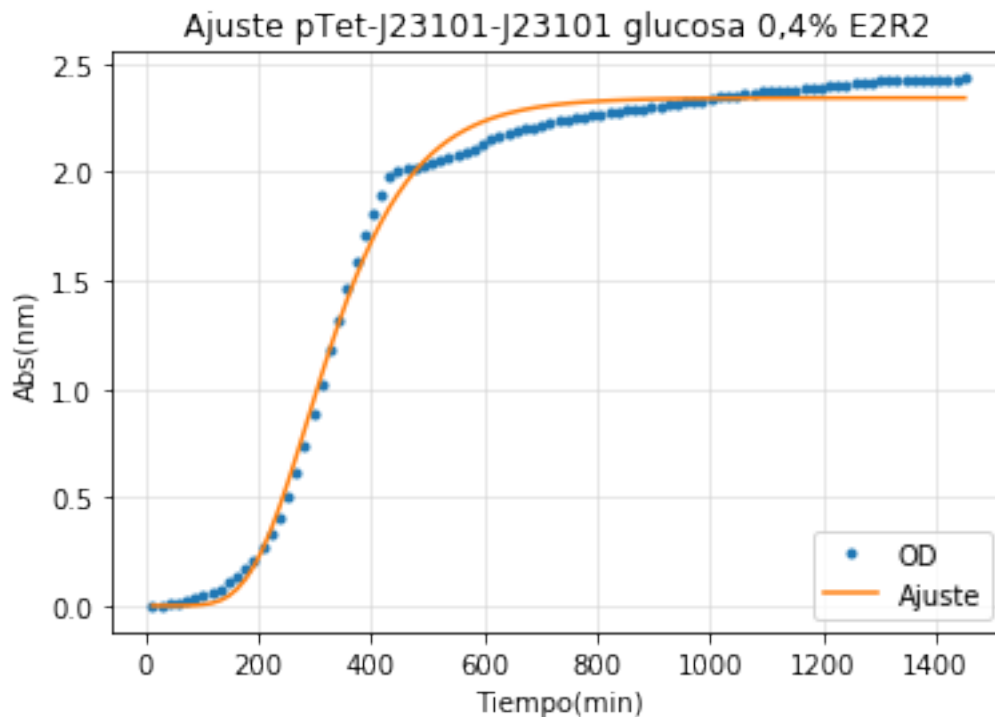
y2=tt[31]
plt.figure()
plt.title('pTet-J23101-J23101 Glucosa 0,4% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od15212,label='OD pTet-J23101-J23101 E2R2')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[19:32],od15212[19:32],label='OD pTet-J23101-J23101 E2R2')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.580000e-02

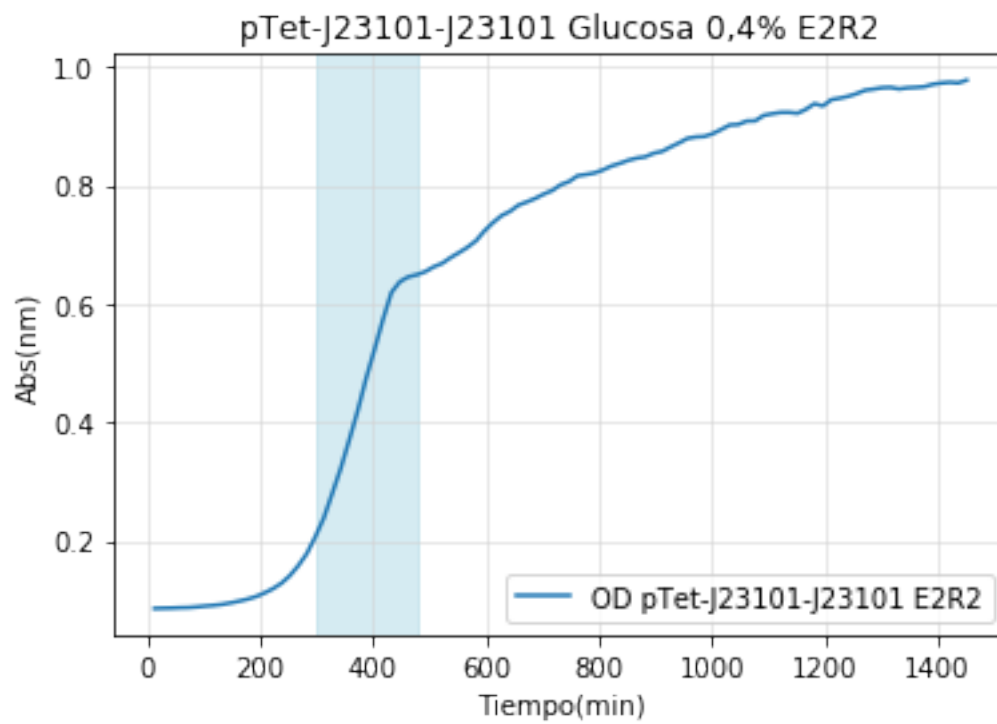
[2.34140500e+00 8.43333849e-03 1.84992926e+02]

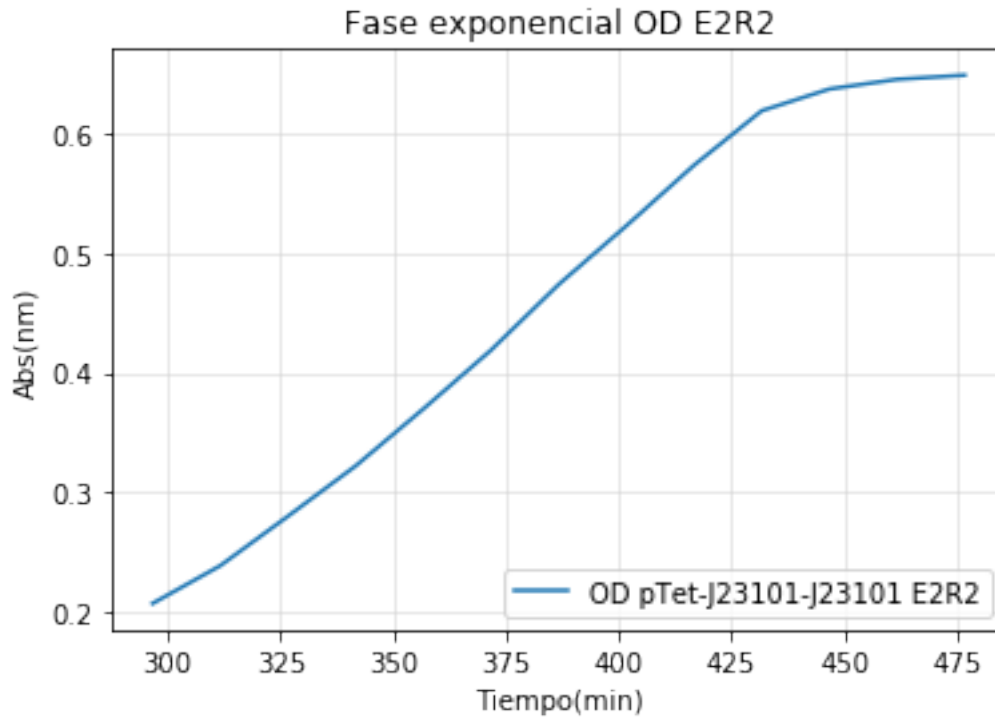


A=2.341405e+00
um=8.433338e-03

l=1.849929e+02
Tm=2.871298e+02
doubpe=8.219132e+01
ext=1.643826e+02
Tfinal=4.515124e+02

Out[16]: <matplotlib.legend.Legend at 0x267e9bc4080>





```
In [17]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-std-std glucosa rep 3
y9= np.log(od15213)-np.log(np.min(od15213))
print('Min OD = %e'%((np.min(od15213))))
evaly, params=Function_fit(tt,y9,0,-1,title = 'Ajuste pTet-J23101-J23101 glucosa 0,4% E
A9= params[0]
um9=params[1]
l9=params[2]
print('A=%e'%(A9))
print('um=%e'%(um9))
print('l=%e'%(l9))

#Cálculo datos para determinar extensión de la fase exponencial
tm9=((A9/(np.exp(1)*um9))+l9)
print('Tm=%e'%(tm9))
t29=((np.log(2))/um9)
print('doubpe=%e'%(t29))
extdp9=2*t29
print('ext=%e'%extdp9)
ttot9=tm9+extdp9
print('Tfinal=%e'%ttot9)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[16]
```

```

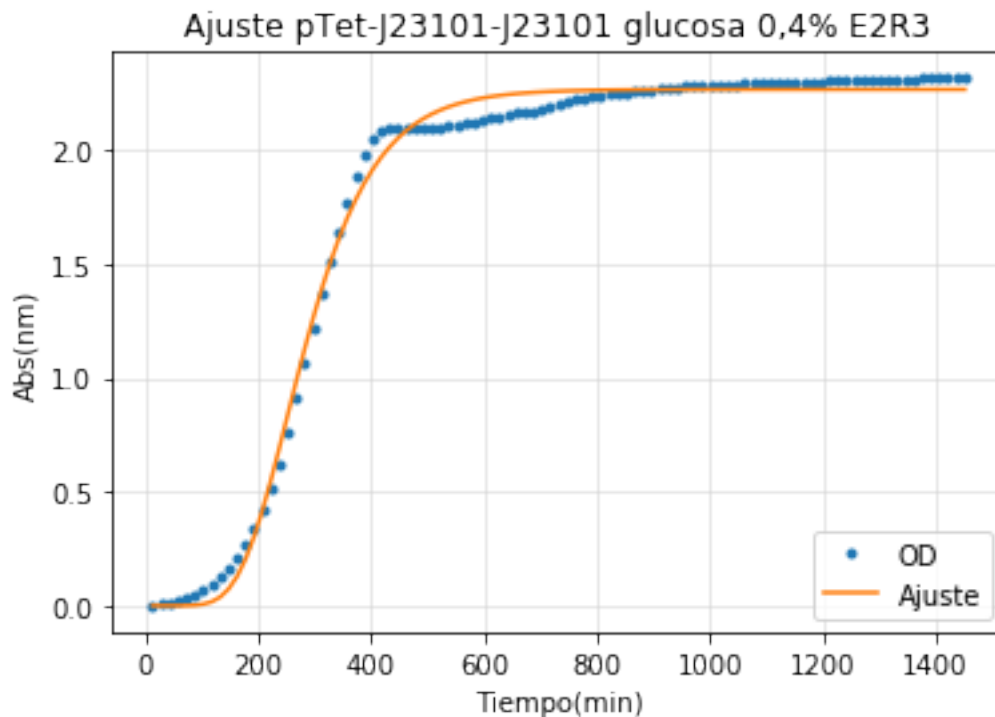
y2=tt[27]
plt.figure()
plt.title('pTet-J23101-J23101 Glucosa 0,4% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od15213,label='OD pTet-J23101-J23101 E2R3')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[16:28],od15213[16:28],label='OD pTet-J23101-J23101 E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.360000e-02

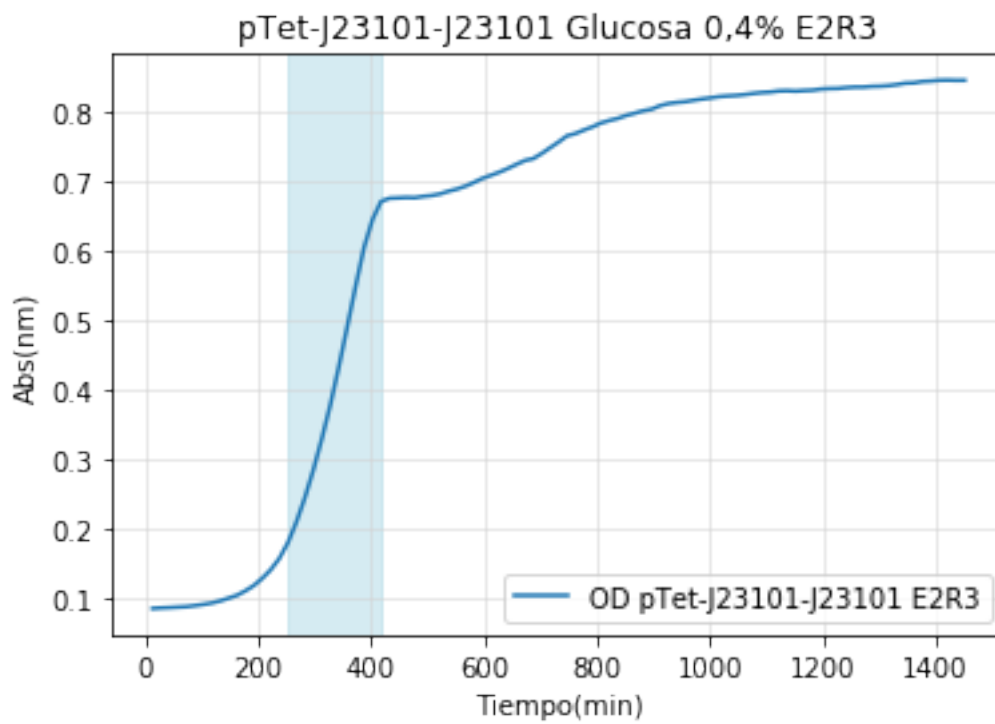
[2.26457592e+00 9.83425423e-03 1.66237345e+02]

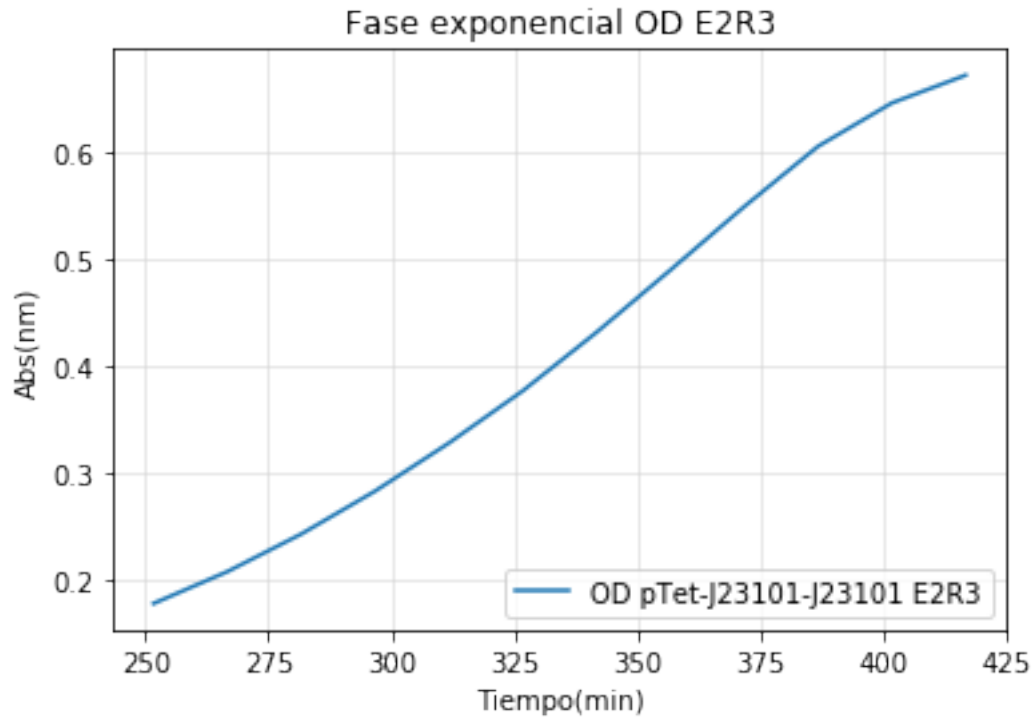


A=2.264576e+00
um=9.834254e-03

l=1.662373e+02
Tm=2.509505e+02
doubpe=7.048294e+01
ext=1.409659e+02
Tfinal=3.919164e+02

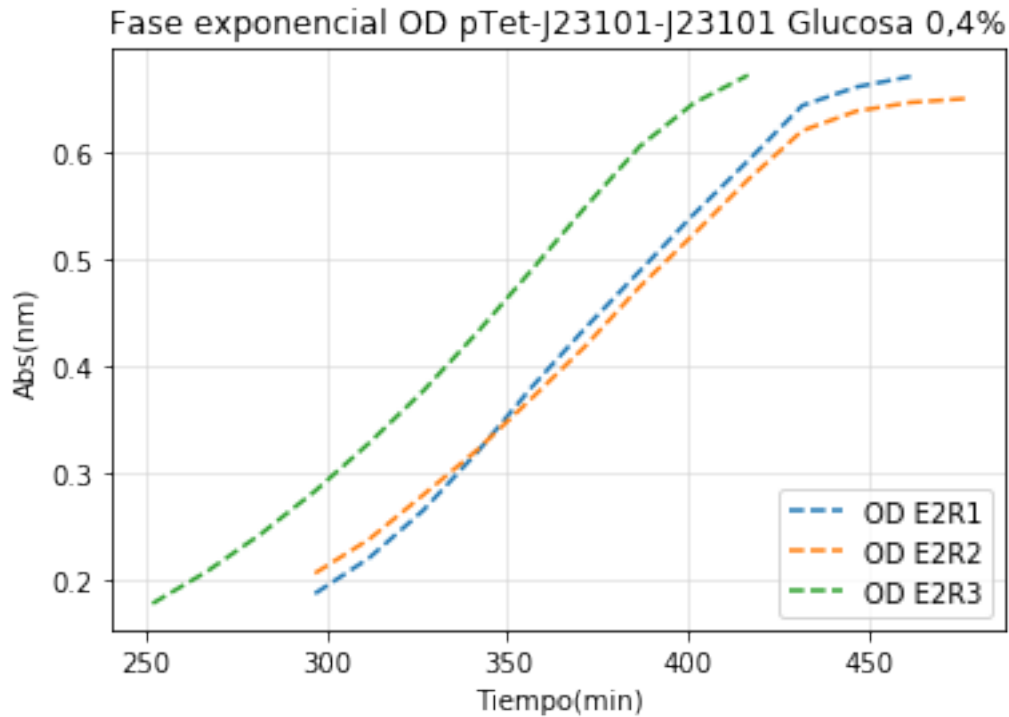
Out[17]: <matplotlib.legend.Legend at 0x267eb4690f0>





```
In [18]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD pTet-J23101-J23101 Glucosa 0,4%')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[19:31],od15211[19:31], '--',label='OD E2R1')
plt.plot(tt[19:32],od15212[19:32], '--',label='OD E2R2')
plt.plot(tt[16:28],od15213[16:28], '--',label='OD E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')
```

```
Out[18]: <matplotlib.legend.Legend at 0x267eb58a710>
```



```
In [19]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-std-std glicerol rep 1
y10= np.log(od1521g1)-np.log(np.min(od1521g1))
print('Min OD = %e'%((np.min(od1521g1))))
evaly, params=Function_fit(tt,y10,0,-1,title = 'Ajuste pTet-J23101-J23101 glicerol 0,2%')
A10= params[0]
um10=params[1]
l10=params[2]
print('A=%e'%(A10))
print('um=%e'%(um10))
print('l=%e'%(l10))

#Cálculo datos para determinar extensión de la fase exponencial
tm10=((A10/(np.exp(1)*um10))+l10)
print('Tm=%e'%(tm10))
t210=((np.log(2))/um10)
print('doubpe=%e'%(t210))
extdp10=2.5*t210
print('ext=%e'%extdp10)
ttot10=tm10+extdp10
print('Tfinal=%e'%ttot10)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[27]
```

```

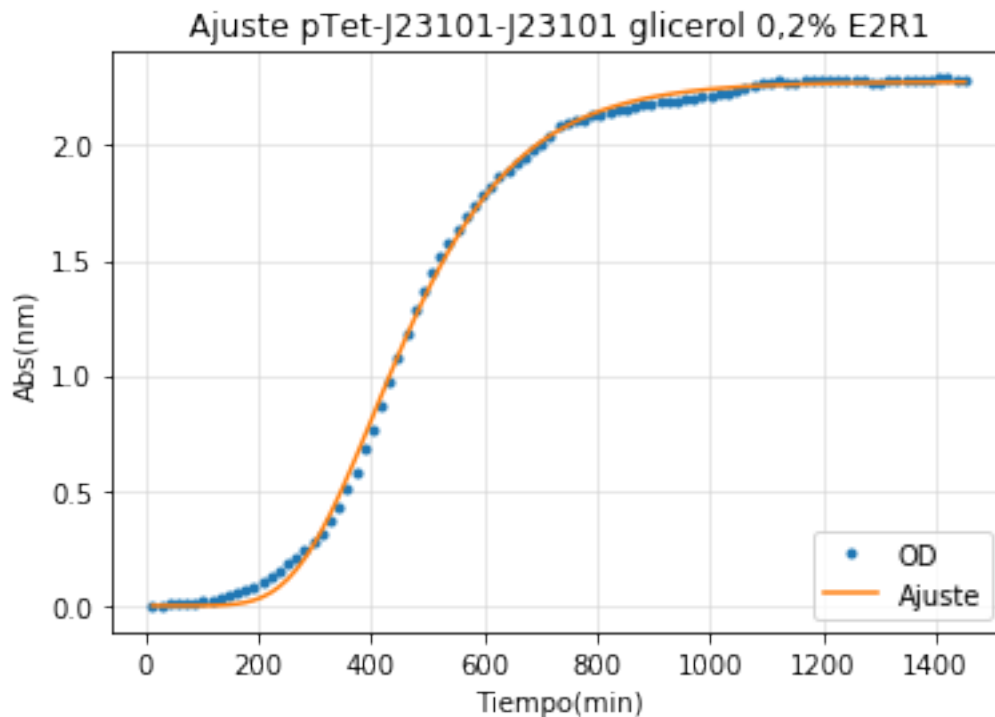
y2=tt[47]
plt.figure()
plt.title('pTet-J23101-J23101 Glicerol 0,2% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od1521g1,label='OD pTet-J23101-J23101 E2R1')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[27:48],od1521g1[27:48],label='OD pTet-J23101-J23101 E2R1')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.510000e-02

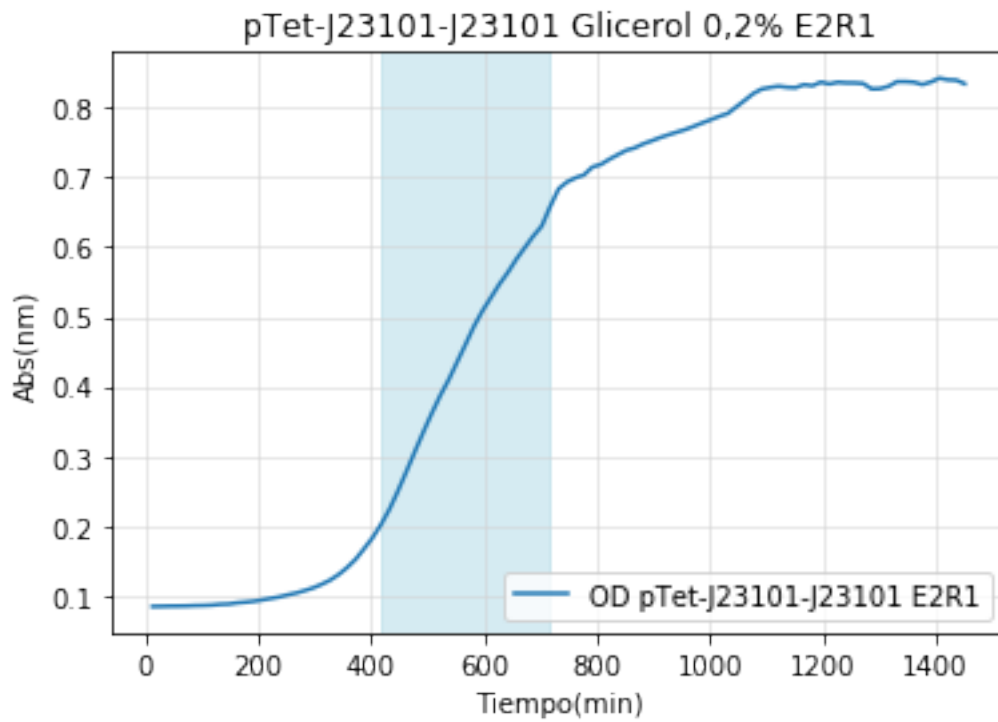
[2.27552043e+00 5.96588621e-03 2.64362418e+02]

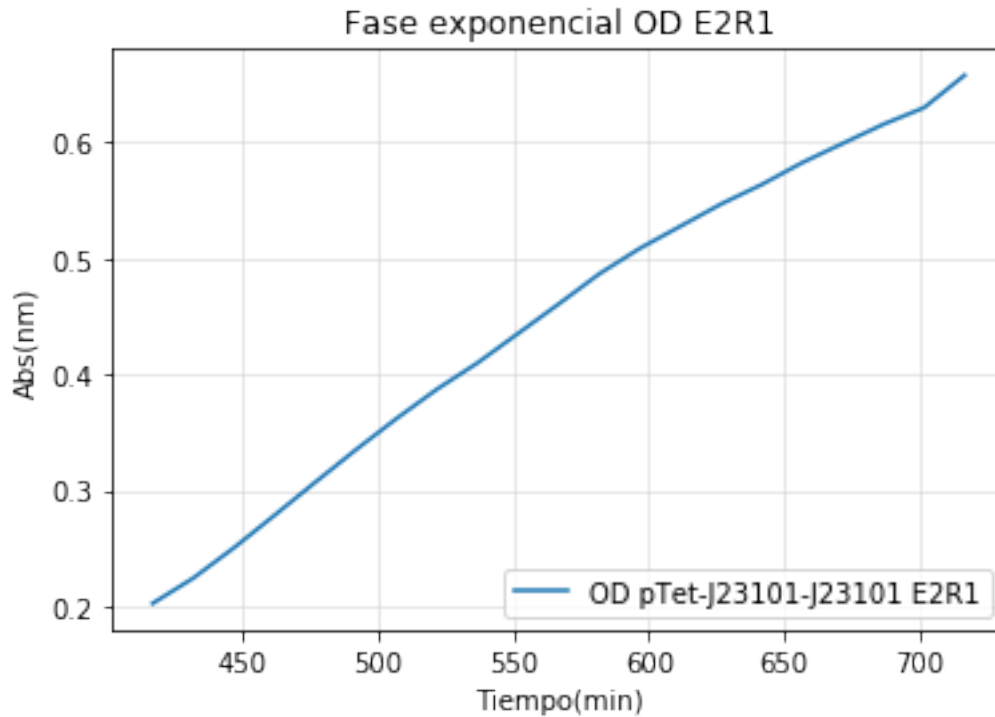


A=2.275520e+00
um=5.965886e-03

```
l=2.643624e+02  
Tm=4.046797e+02  
doubpe=1.161851e+02  
ext=2.904628e+02  
Tfinal=6.951425e+02
```

Out[19]: <matplotlib.legend.Legend at 0x267eb436358>





```
In [20]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-std-std glicerol rep 2
y11= np.log(od1521g2)-np.log(np.min(od1521g2))
print('Min OD = %e'%((np.min(od1521g2))))
evaly, params=Function_fit(tt,y11,0,-1,title = 'Ajuste pTet-J23101-J23101 glicerol 0,2%')
A11= params[0]
um11=params[1]
l11=params[2]
print('A=%e'%(A11))
print('um=%e'%(um11))
print('l=%e'%(l11))

#Cálculo datos para determinar extensión de la fase exponencial
tm11=((A11/(np.exp(1)*um11))+l11)
print('Tm=%e'%(tm11))
t211=((np.log(2))/um11)
print('doubpe=%e'%(t211))
extdp11=2.5*t211
print('ext=%e'%extdp11)
ttot11=tm11+extdp11
print('Tfinal=%e'%ttot11)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[28]
```

```

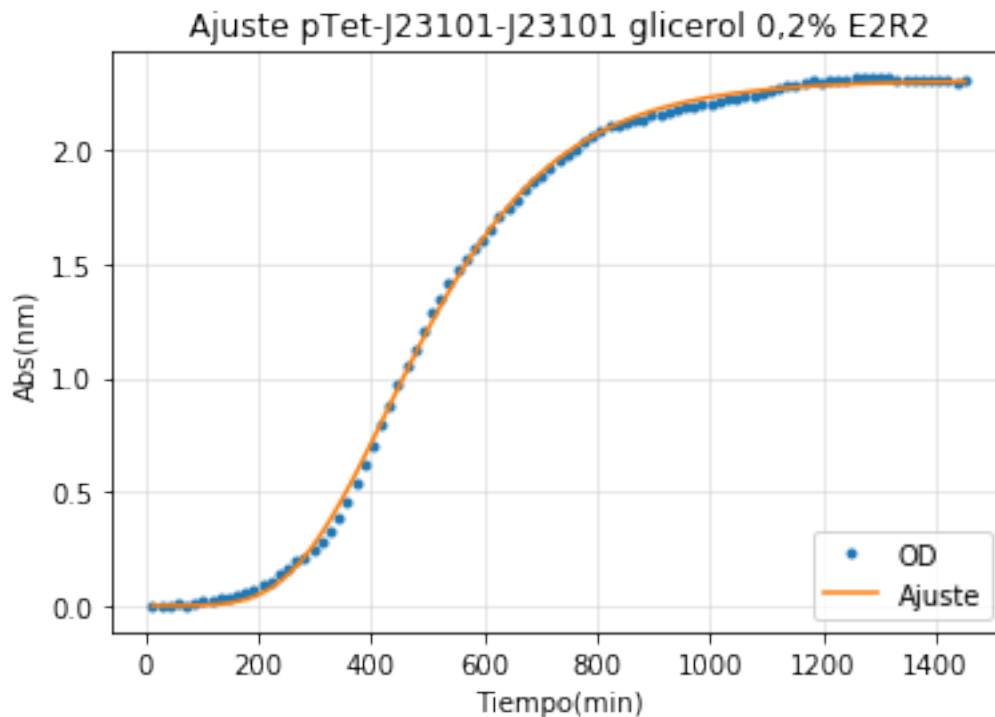
y2=tt[52]
plt.figure()
plt.title('pTet-J23101-J23101 Glicerol 0,2% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od1521g2,label='OD pTet-J23101-J23101 E2R2')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[28:53],od1521g2[28:53],label='OD pTet-J23101-J23101 E2R2')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.600000e-02

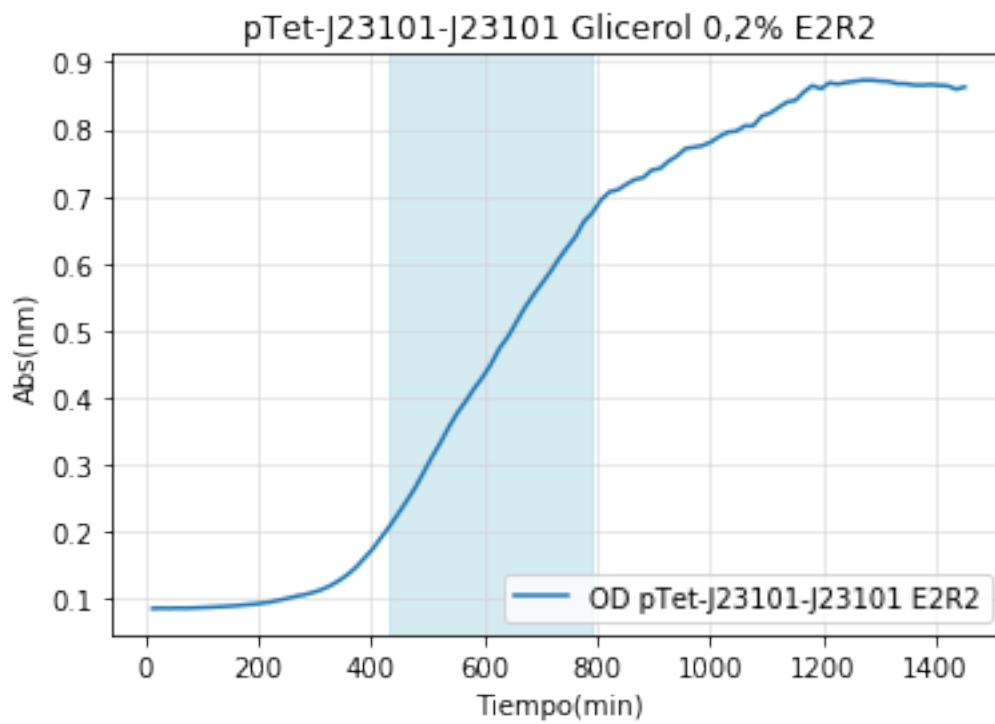
[2.30705217e+00 5.07308656e-03 2.58634310e+02]

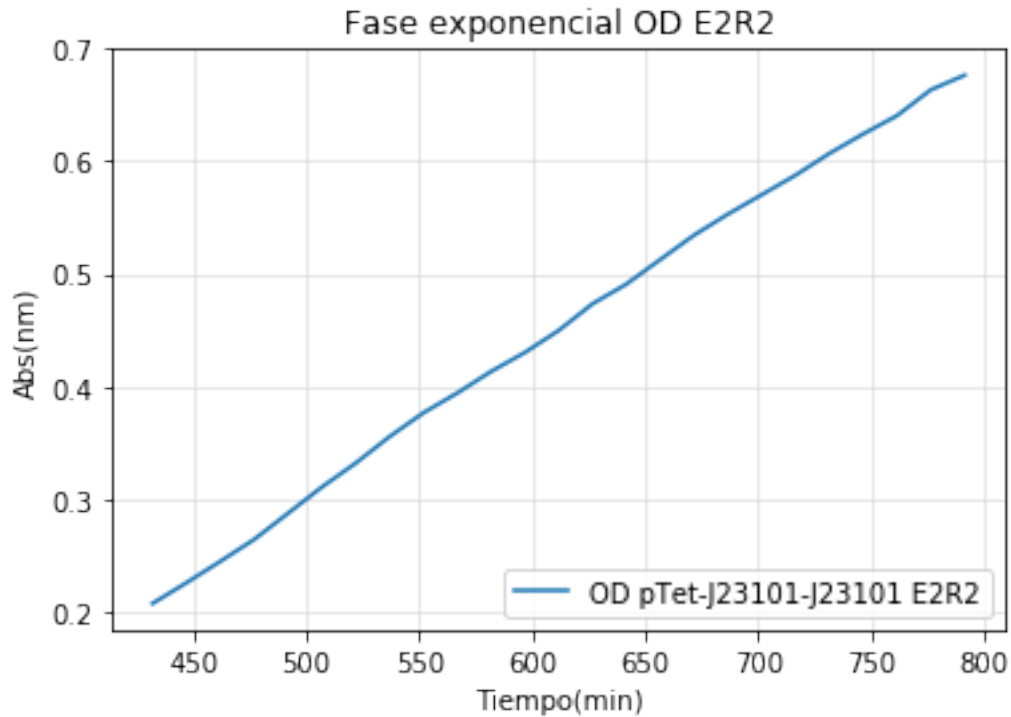


A=2.307052e+00
um=5.073087e-03

l=2.586343e+02
Tm=4.259323e+02
doubpe=1.366322e+02
ext=3.415806e+02
Tfinal=7.675129e+02

Out[20]: <matplotlib.legend.Legend at 0x267eb14b320>





```
In [21]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-std-std glicerol rep 3
y12= np.log(od1521g3)-np.log(np.min(od1521g3))
print('Min OD = %e'%((np.min(od1521g3))))
evaly, params=Function_fit(tt,y12,0,-1,title = 'Ajuste pTet-J23101-J23101 glicerol 0,2%')
A12= params[0]
um12=params[1]
l12=params[2]
print('A=%e'%(A12))
print('um=%e'%(um12))
print('l=%e'%(l12))

#Cálculo datos para determinar extensión de la fase exponencial
tm12=((A12/(np.exp(1)*um12))+l12)
print('Tm=%e'%(tm12))
t212=((np.log(2))/um12)
print('doubpe=%e'%(t212))
extdp12=2*t212
print('ext=%e'%extdp12)
ttot12=tm12+extdp12
print('Tfinal=%e'%ttot12)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[27]
```

```

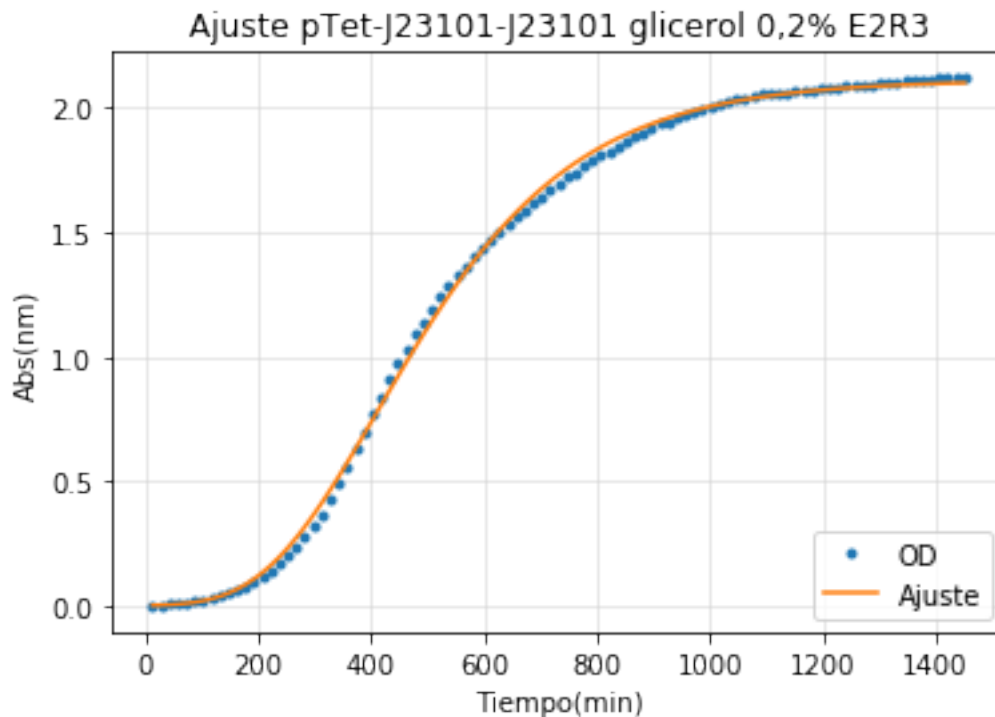
y2=tt[52]
plt.figure()
plt.title('pTet-J23101-J23101 Glicerol 0,2% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od1521g3,label='OD pTet-J23101-J23101 E2R3')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[27:53],od1521g3[27:53],label='OD pTet-J23101-J23101 E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.320000e-02

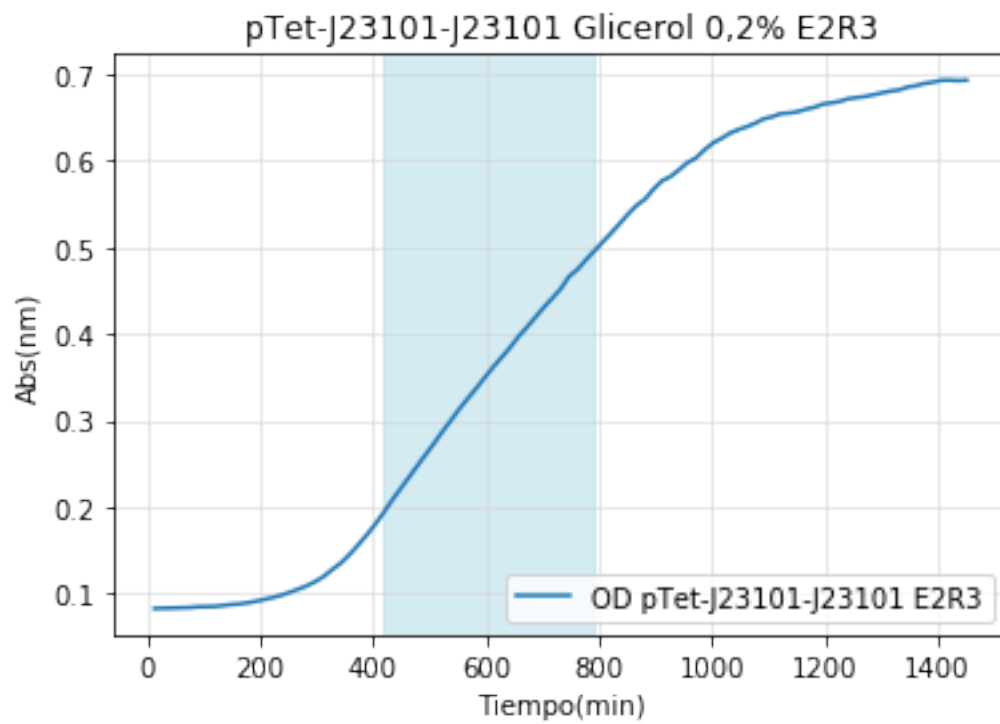
[2.11194516e+00 3.89343431e-03 2.09853636e+02]

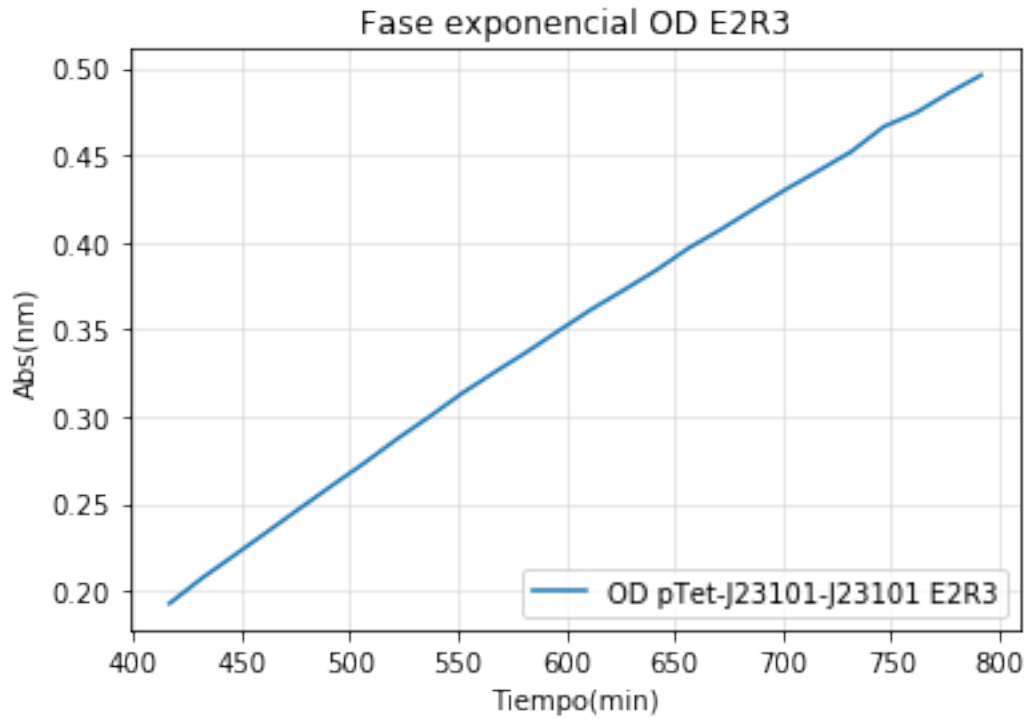


A=2.111945e+00
um=3.893434e-03

```
l=2.098536e+02  
Tm=4.094053e+02  
doubpe=1.780298e+02  
ext=3.560595e+02  
Tfinal=7.654648e+02
```

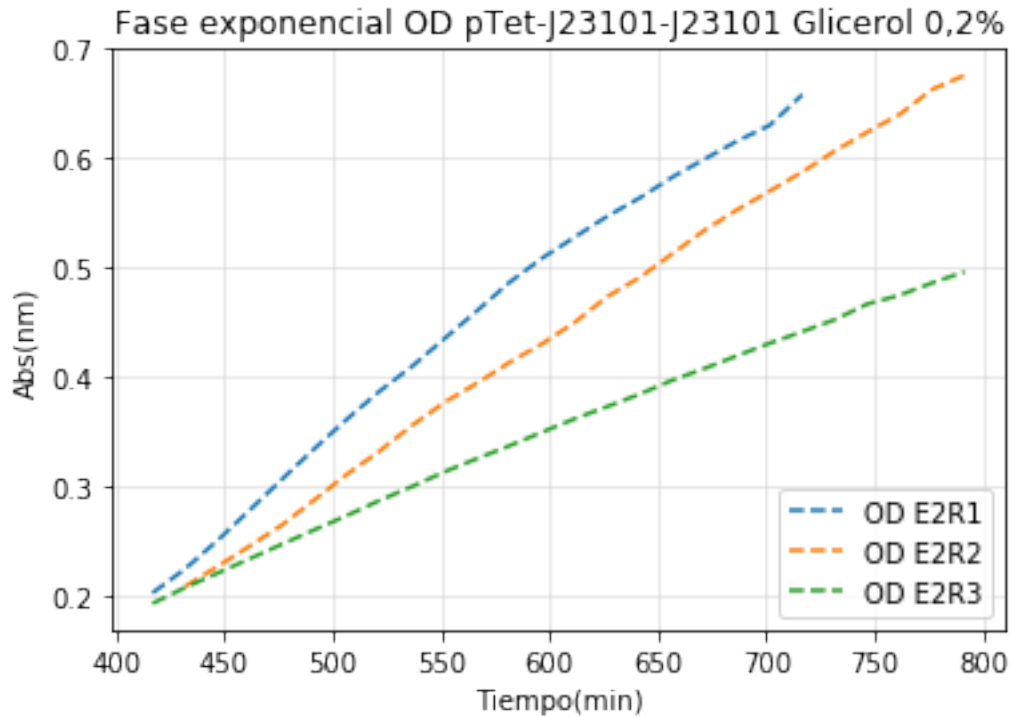
Out[21]: <matplotlib.legend.Legend at 0x267eb6395f8>





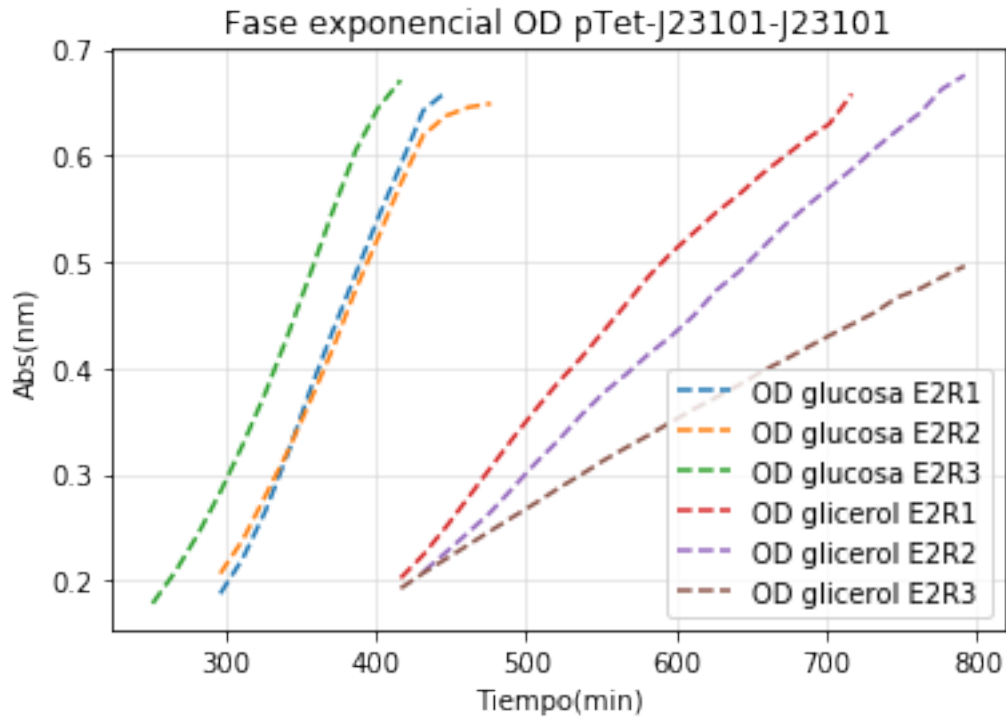
```
In [22]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD pTet-J23101-J23101 Glicerol 0,2%')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[27:48],od1521g1[27:48], '--',label='OD E2R1')
plt.plot(tt[28:53],od1521g2[28:53], '--',label='OD E2R2')
plt.plot(tt[27:53],od1521g3[27:53], '--',label='OD E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')
```

```
Out[22]: <matplotlib.legend.Legend at 0x267eb789ef0>
```



```
In [23]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD pTet-J23101-J23101')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[19:30],od15211[19:30], '--',label='OD glucosa E2R1')
plt.plot(tt[19:32],od15212[19:32], '--',label='OD glucosa E2R2')
plt.plot(tt[16:28],od15213[16:28], '--',label='OD glucosa E2R3')
plt.plot(tt[27:48],od1521g1[27:48], '--',label='OD glicerol E2R1')
plt.plot(tt[28:53],od1521g2[28:53], '--',label='OD glicerol E2R2')
plt.plot(tt[27:53],od1521g3[27:53], '--',label='OD glicerol E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')
```

```
Out[23]: <matplotlib.legend.Legend at 0x267eb85acc0>
```

```
In [24]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-107-std glucosa rep 1
y13= np.log(od15231)-np.log(np.min(od15231))
print('Min OD = %e'%((np.min(od15231))))
evaly, params=Function_fit(tt,y13,0,-1,title = 'Ajuste pTet-J23107-J23101 glucosa 0,4%')
A13 = params[0]
um13=params[1]
l13=params[2]
print('A=%e'%(A13))
print('um=%e'%(um13))
print('l=%e'%(l13))

#Cálculo datos para determinar extensión de la fase exponencial
tm13=((A13/(np.exp(1)*um13))+l13)
print('Tm=%e'%(tm13))
t213=((np.log(2))/um13)
print('doubpe=%e'%(t213))
extdp13=2*t213
print('ext=%e'%extdp13)
ttot13=tm13+extdp13
print('Tfinal=%e'%ttot13)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[21]
```

```

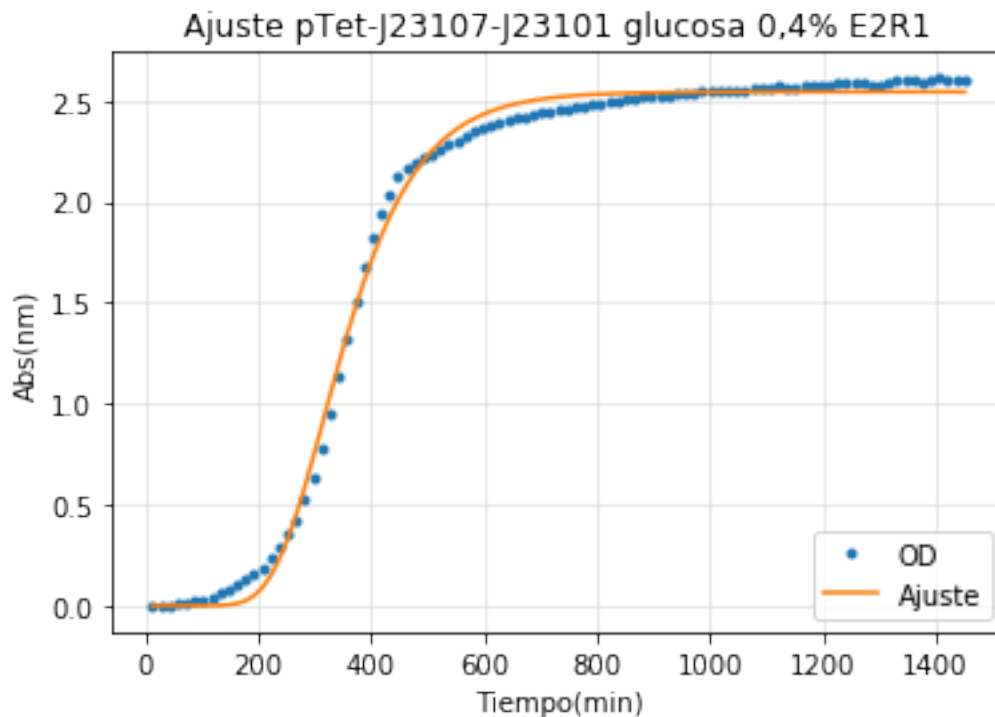
y2=tt[31]
plt.figure()
plt.title('pTet-J23107-J23101 Glucosa 0,4% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od15231,label='OD pTet-J23107-J23101 E2R1 ')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E23R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[21:32],od15231[21:32],label='OD pTet-J23107-J23101 E2R1')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.570000e-02

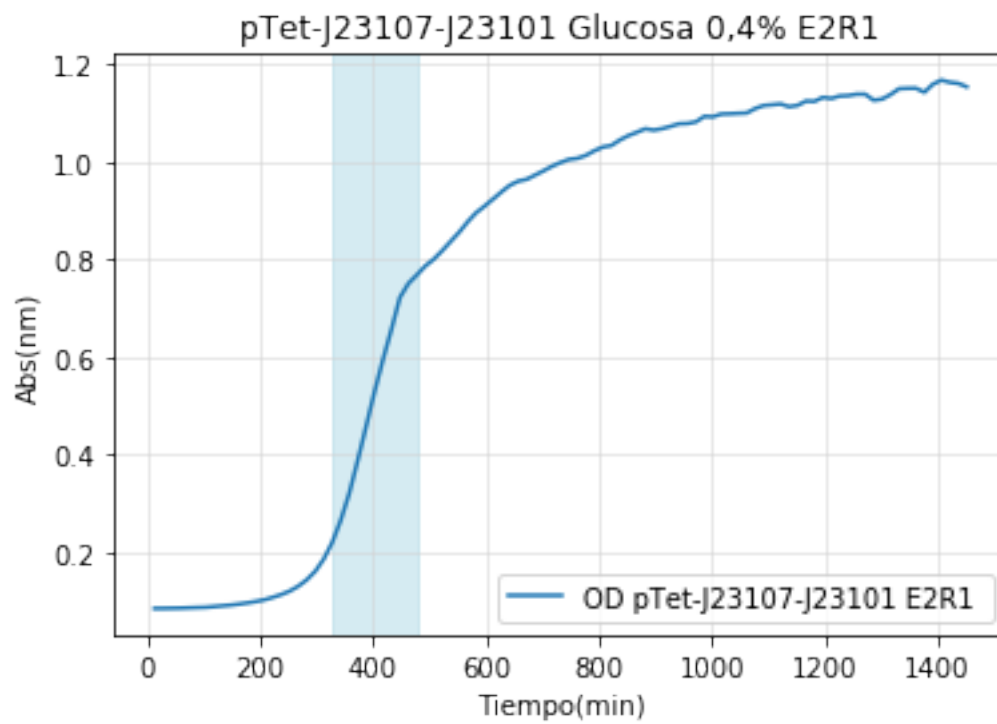
[2.54342379e+00 1.03117507e-02 2.25391606e+02]

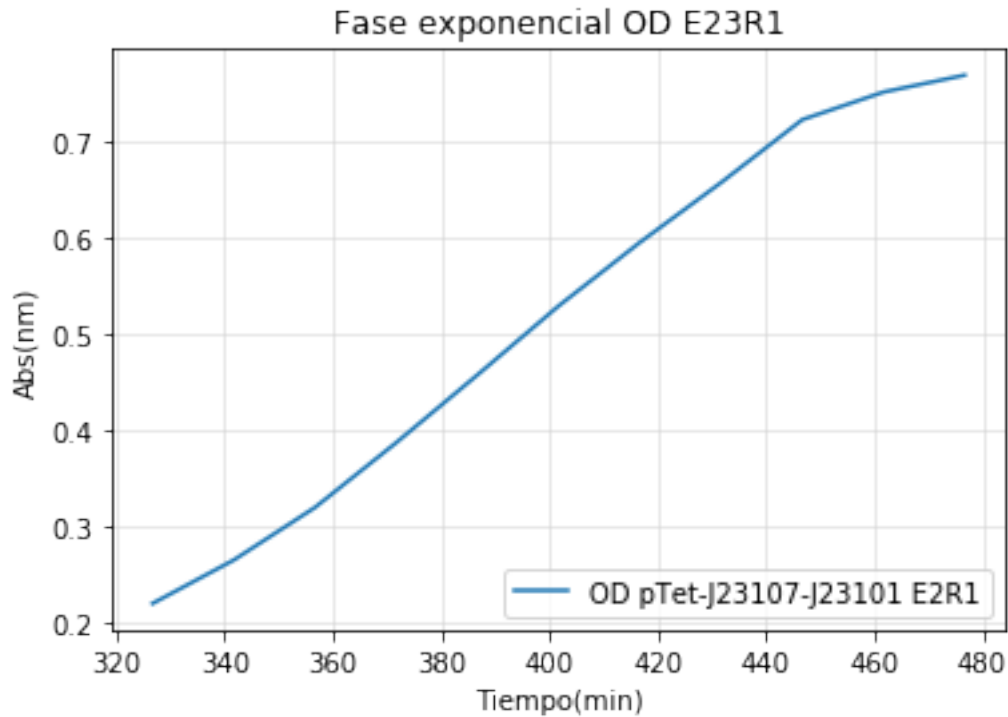


A=2.543424e+00
um=1.031175e-02

l=2.253916e+02
Tm=3.161302e+02
doubpe=6.721916e+01
ext=1.344383e+02
Tfinal=4.505685e+02

Out[24]: <matplotlib.legend.Legend at 0x267eb076048>





```
In [25]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-107-std glucosa rep 2
y14= np.log(od15232)-np.log(np.min(od15232))
print('Min OD = %e'%((np.min(od15232))))
evaly, params=Function_fit(tt,y14,0,-1,title = 'Ajuste pTet-J23107-J23101 glucosa 0,4%
A14= params[0]
um14=params[1]
l14=params[2]
print('A=%e'%(A14))
print('um=%e'%(um14))
print('l=%e'%(l14))

#Cálculo datos para determinar extensión de la fase exponencial
tm14=((A14/(np.exp(1)*um14))+l14)
print('Tm=%e'%(tm14))
t214=((np.log(2))/um14)
print('doubpe=%e'%(t214))
extdp14=2*t214
print('ext=%e'%extdp14)
ttot14=tm14+extdp14
print('Tfinal=%e'%ttot14)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[17]
```

```

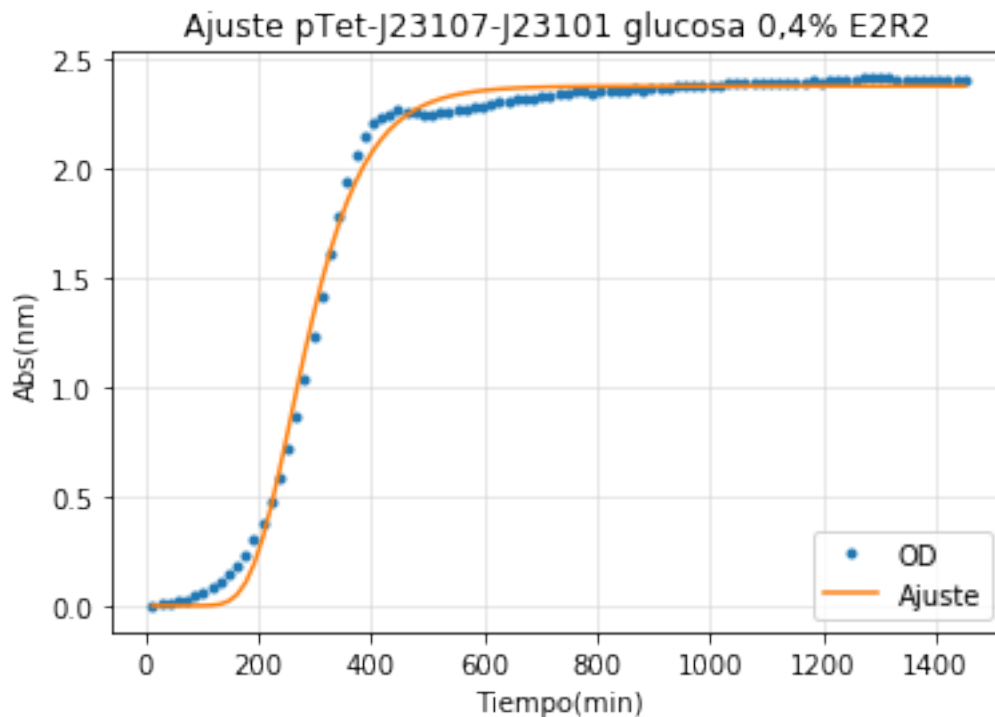
y2=tt[26]
plt.figure()
plt.title('pTet-J23107-J23101 Glucosa 0,4% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od15232,label='OD pTet-J23107-J23101 E2R2')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[17:27],od15232[17:27],label='OD pTet-J23107-J23101 E2R2')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.650000e-02

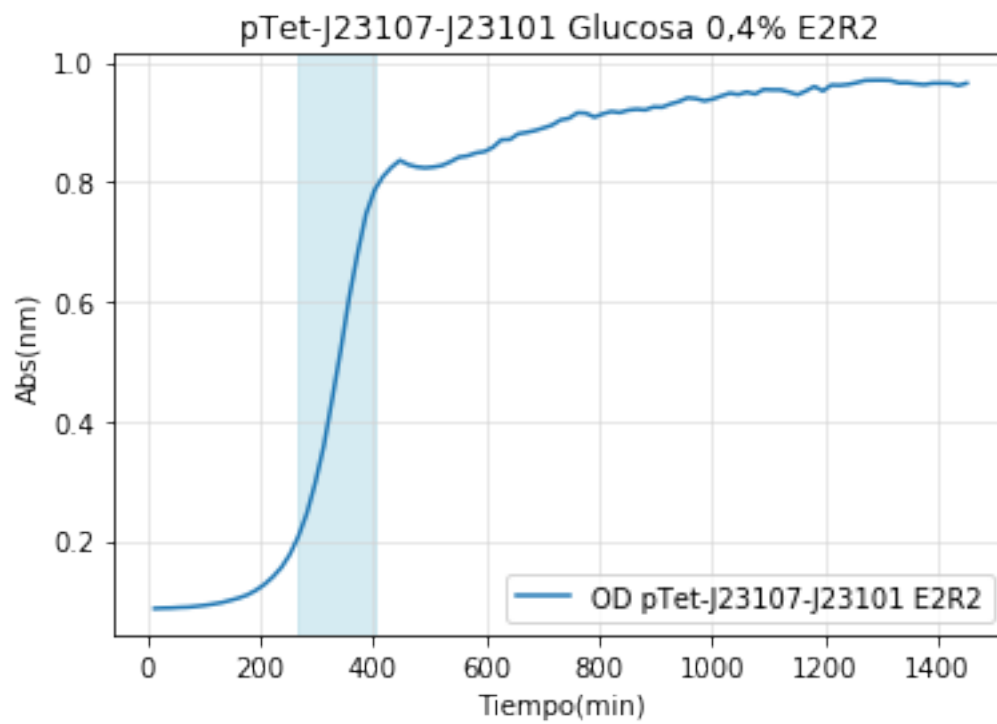
[2.38049312e+00 1.22442727e-02 1.86353375e+02]

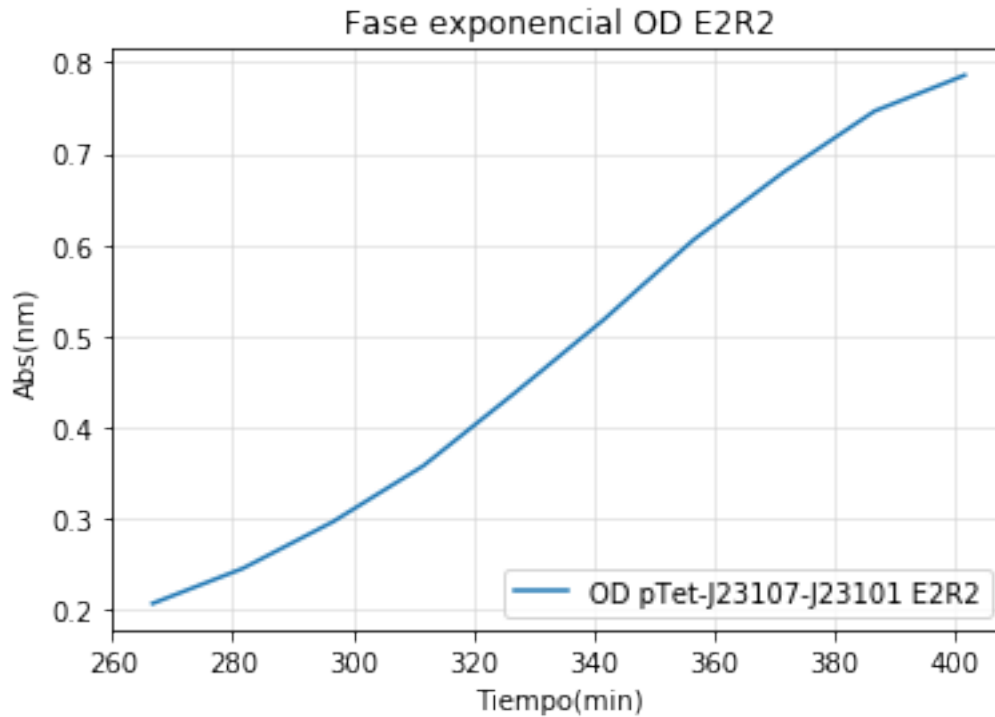


A=2.380493e+00
um=1.224427e-02

l=1.863534e+02
Tm=2.578753e+02
doubpe=5.660991e+01
ext=1.132198e+02
Tfinal=3.710952e+02

Out[25]: <matplotlib.legend.Legend at 0x267eb678a20>





```
In [26]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-107-std glucosa rep 3
y15= np.log(od15233)-np.log(np.min(od15233))
print('Min OD = %e'%((np.min(od15233))))
evaly, params=Function_fit(tt,y15,0,-1,title = 'Ajuste pTet-J23107-J23101 glucosa 0,4%')
A15= params[0]
um15=params[1]
l15=params[2]
print('A=%e'%(A15))
print('um=%e'%(um15))
print('l=%e'%(l15))

#Cálculo datos para determinar extensión de la fase exponencial
tm15=((A15/(np.exp(1)*um15))+l15)
print('Tm=%e'%(tm15))
t215=((np.log(2))/um15)
print('doubpe=%e'%(t215))
extdp15=2*t215
print('ext=%e'%extdp15)
ttot15=tm15+extdp15
print('Tfinal=%e'%ttot15)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[17]
```

```

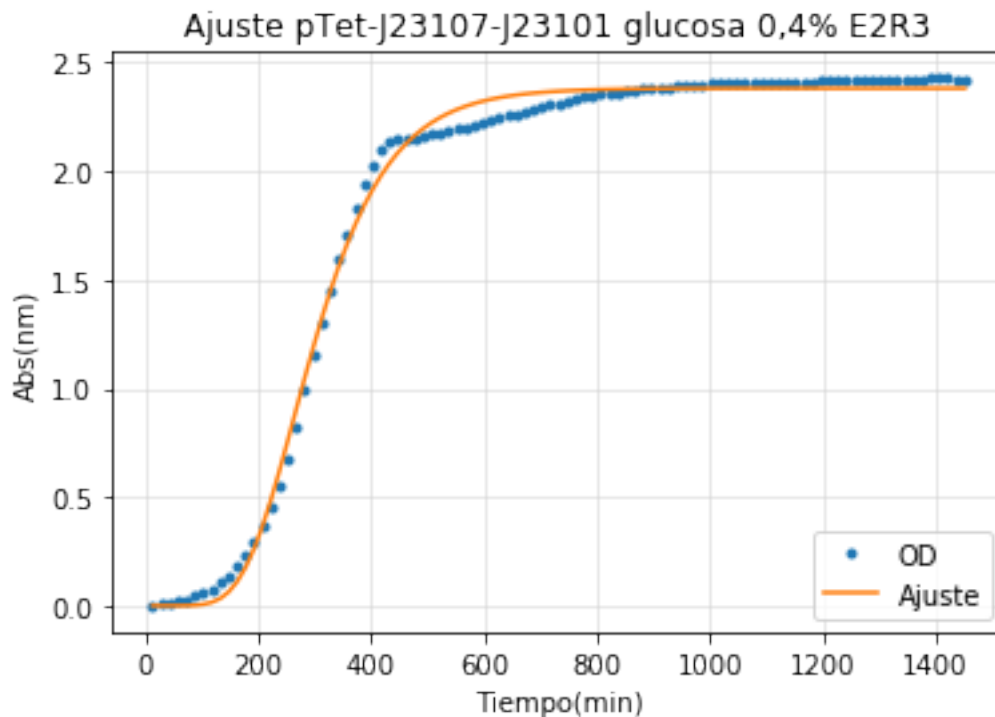
y2=tt[28]
plt.figure()
plt.title('pTet-J23107-J23101 Glucosa 0,4% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od15233,label='OD pTet-J23107-J23101 E2R3')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[17:29],od15233[17:29],label='OD pTet-J23107-J23101 E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.400000e-02

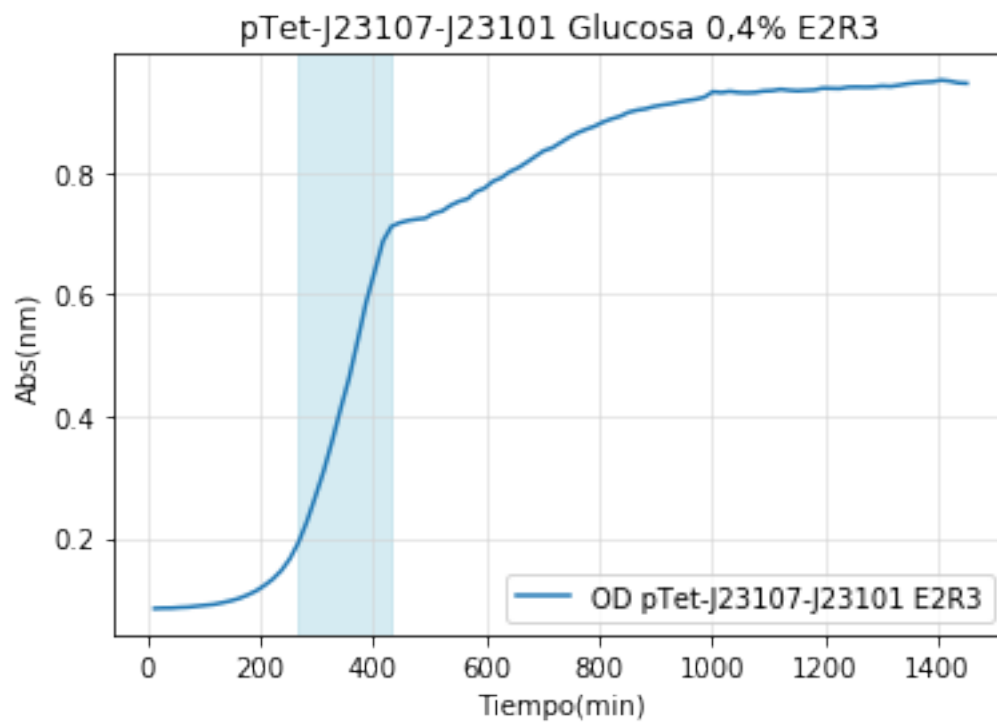
[2.38113195e+00 9.65546943e-03 1.72796102e+02]

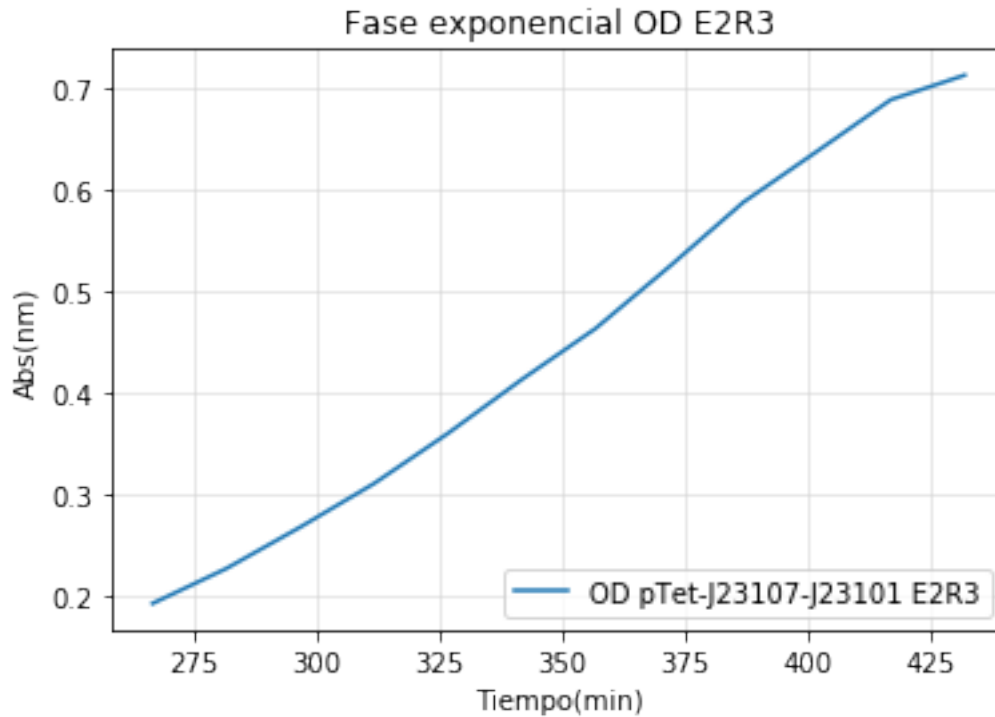


A=2.381132e+00
um=9.655469e-03

l=1.727961e+02
Tm=2.635187e+02
doubpe=7.178804e+01
ext=1.435761e+02
Tfinal=4.070948e+02

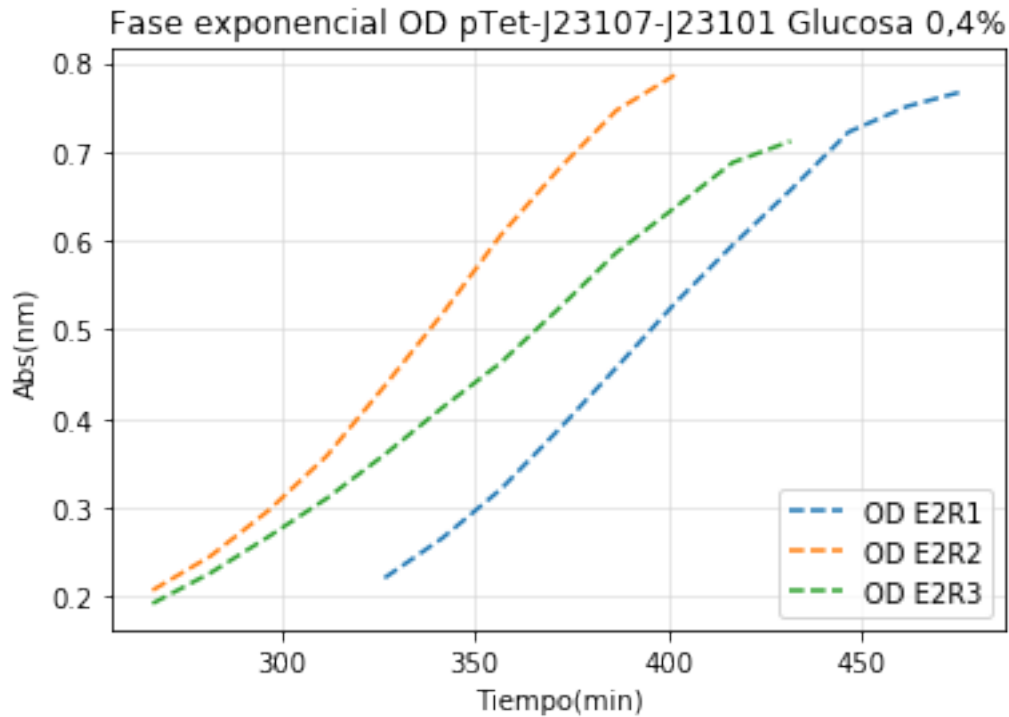
Out[26]: <matplotlib.legend.Legend at 0x267eca63e48>





```
In [27]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD pTet-J23107-J23101 Glucosa 0,4%')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[21:32],od15231[21:32], '--',label='OD E2R1')
plt.plot(tt[17:27],od15232[17:27], '--',label='OD E2R2')
plt.plot(tt[17:29],od15233[17:29], '--',label='OD E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')
```

```
Out[27]: <matplotlib.legend.Legend at 0x267ecb7d7b8>
```



```
In [28]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-107-std glicerol rep 1
y16= np.log(od1523g1)-np.log(np.min(od1523g1))
print('Min OD = %e'%((np.min(od1523g1))))
evaly, params=Function_fit(tt,y16,0,-1,title = 'Ajuste pTet-J23107-J23101 glicerol 0,2%')
A16= params[0]
um16=params[1]
l16=params[2]
print('A=%e'%(A16))
print('um=%e'%(um16))
print('l=%e'%(l16))

#Cálculo datos para determinar extensión de la fase exponencial
tm16=((A16/(np.exp(1)*um16))+l16)
print('Tm=%e'%(tm16))
t216=((np.log(2))/um16)
print('doubpe=%e'%(t216))
extdp16=2.5*t216
print('ext=%e'%extdp16)
ttot16=tm16+extdp16
print('Tfinal=%e'%ttot16)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[30]
```

```

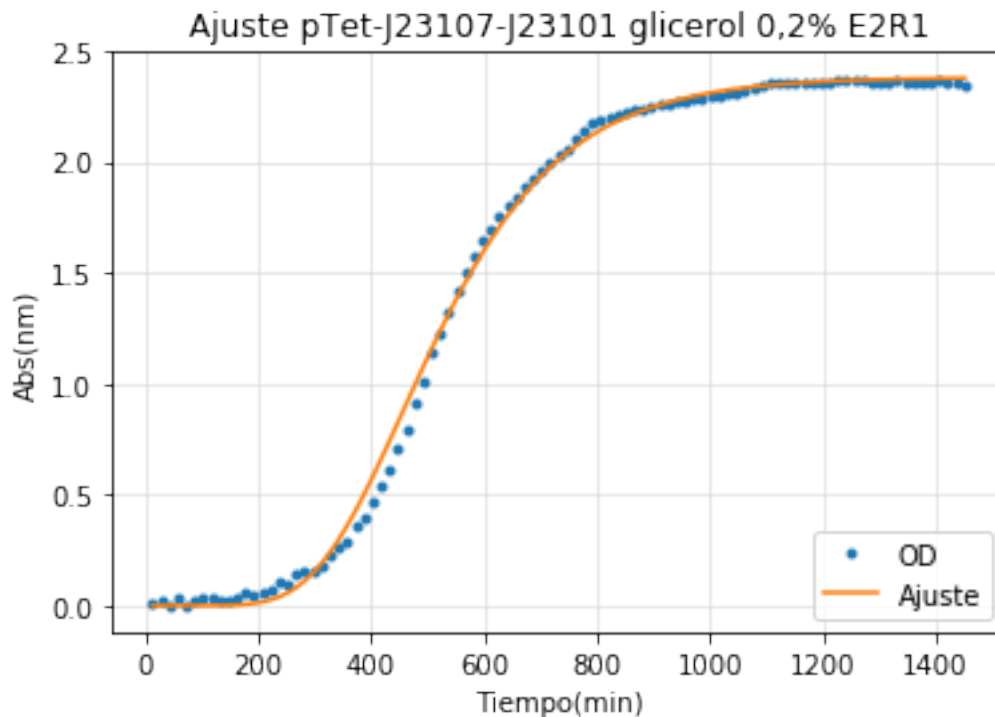
y2=tt[52]
plt.figure()
plt.title('pTet-J23107-J23101 Glicerol 0,2% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od1523g1,label='OD pTet-J23107-J23101 E2R1')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[30:53],od1523g1[30:53],label='OD pTet-J23107-J23101 E2R1')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.580000e-02

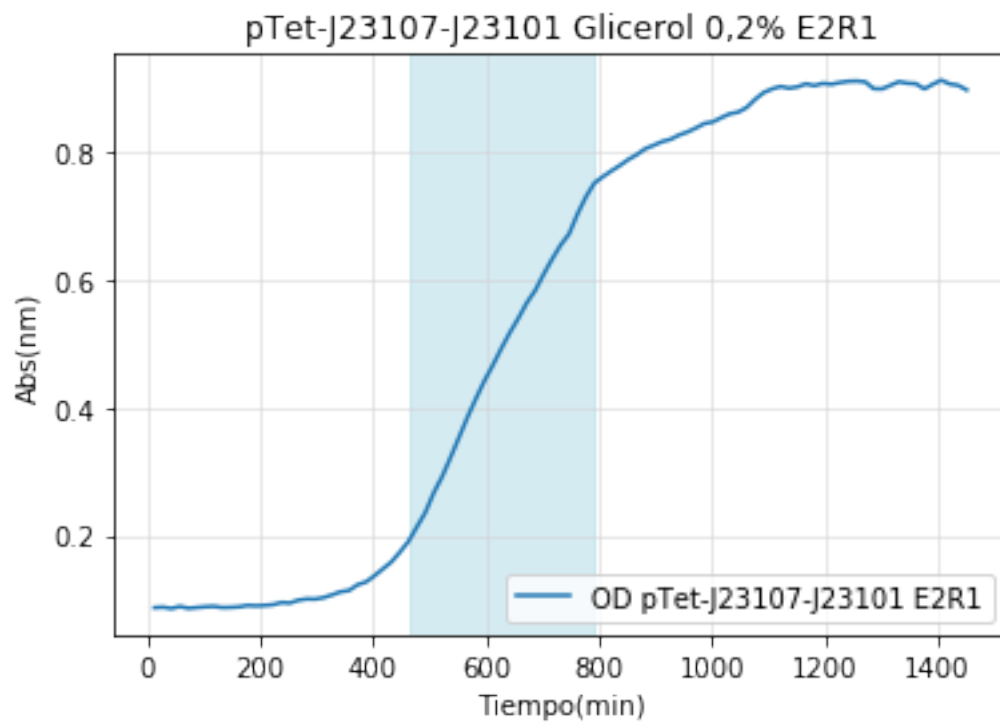
[2.37993854e+00 5.62626737e-03 3.00000000e+02]

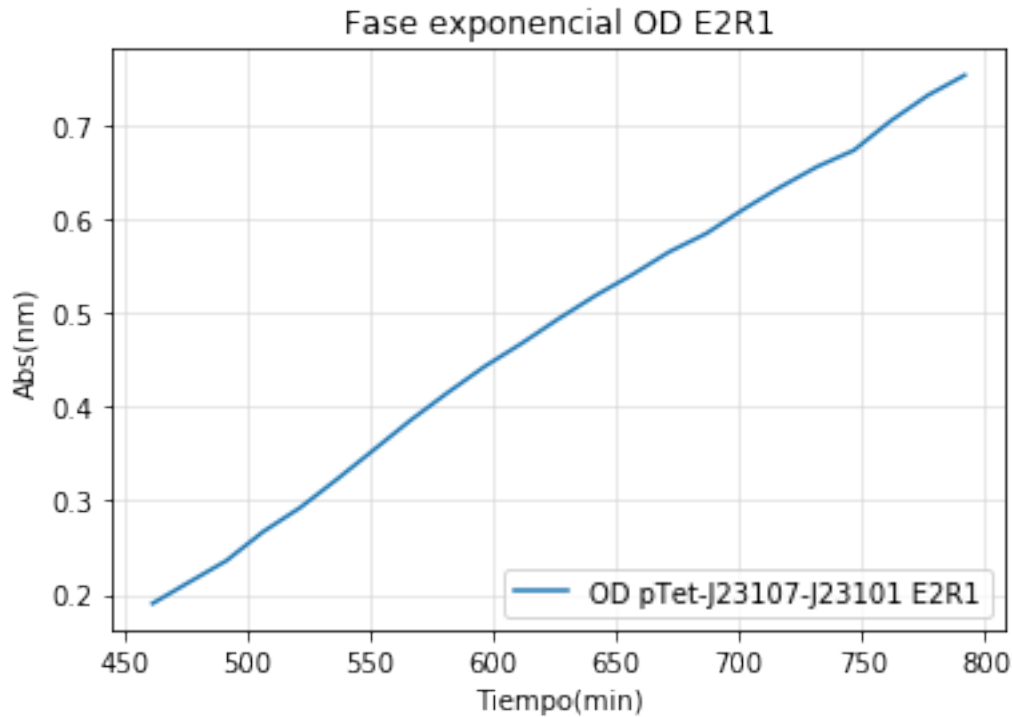


A=2.379939e+00
um=5.626267e-03

```
l=3.000000e+02  
Tm=4.556148e+02  
doubpe=1.231984e+02  
ext=3.079960e+02  
Tfinal=7.636108e+02
```

Out[28]: <matplotlib.legend.Legend at 0x267ecac6b38>





```
In [29]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-107-std glicerol rep 2
y17= np.log(od1523g2)-np.log(np.min(od1523g2))
print('Min OD = %e'%((np.min(od1523g2))))
evaly, params=Function_fit(tt,y17,0,-1,title = 'Ajuste pTet-J23107-J23101 glicerol 0,2%')
A17= params[0]
um17=params[1]
l17=params[2]
print('A=%e'%(A17))
print('um=%e'%(um17))
print('l=%e'%(l17))

#Cálculo datos para determinar extensión de la fase exponencial
tm17=((A17/(np.exp(1)*um17))+l17)
print('Tm=%e'%(tm17))
t217=((np.log(2))/um17)
print('doubpe=%e'%(t217))
extdp17=2.5*t217
print('ext=%e'%extdp17)
ttot17=tm17+extdp17
print('Tfinal=%e'%ttot17)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[27]
```

```

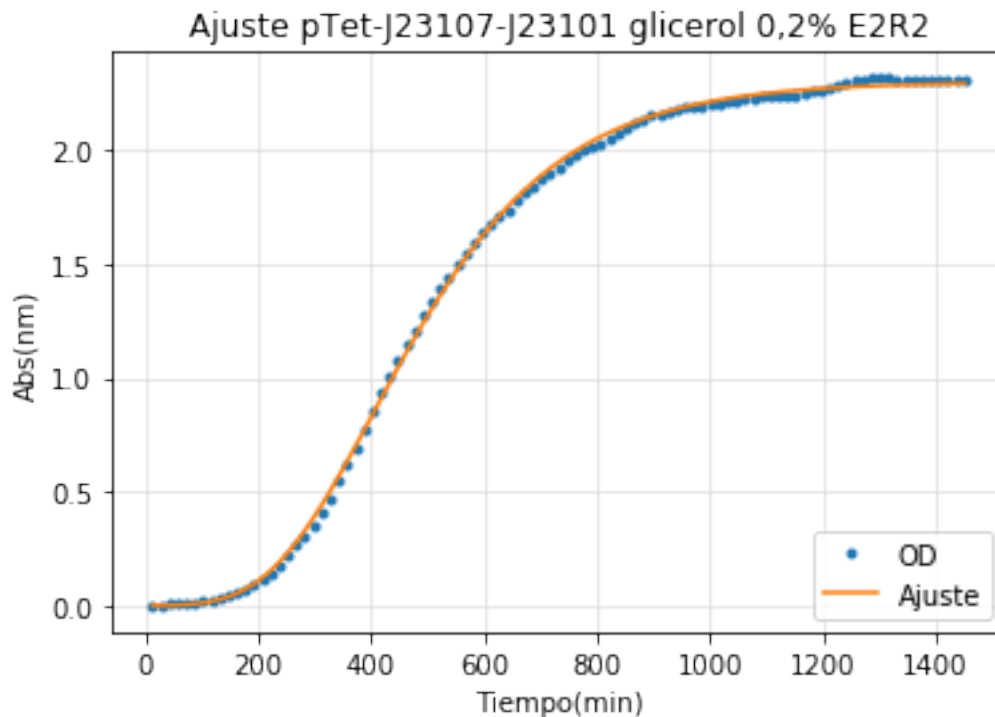
y2=tt[53]
plt.figure()
plt.title('pTet-J23107-J23101 Glicerol 0,2% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od1523g2,label='OD pTet-J23107-J23101 E2R2')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[27:54],od1523g2[27:54],label='OD pTet-J23107-J23101 E2R2')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.550000e-02

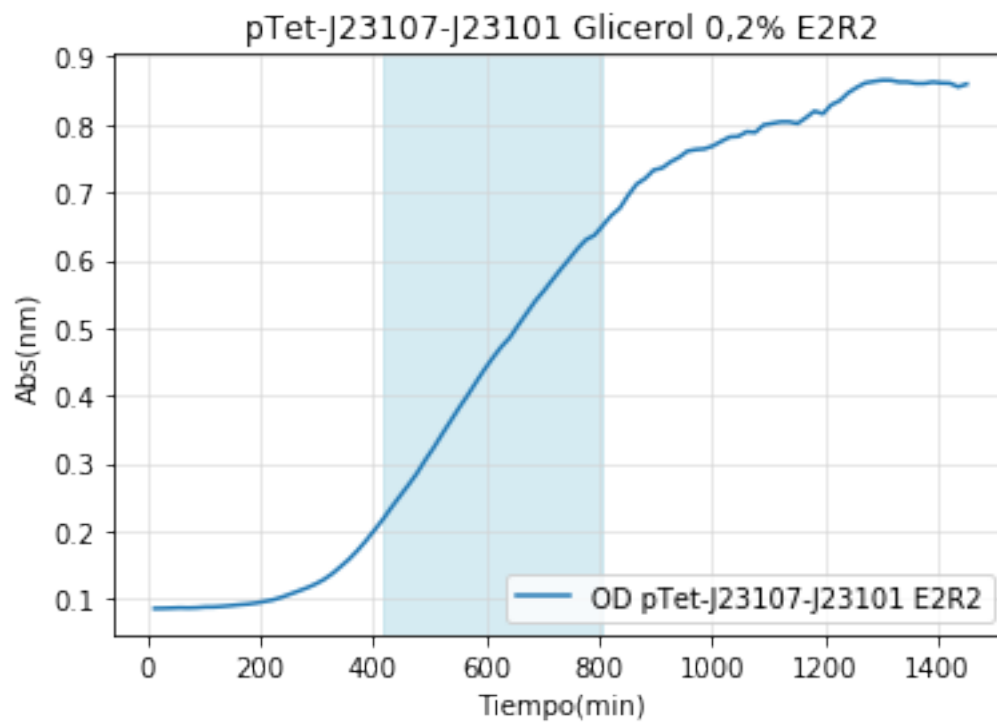
[2.29805766e+00 4.61718516e-03 2.20040295e+02]

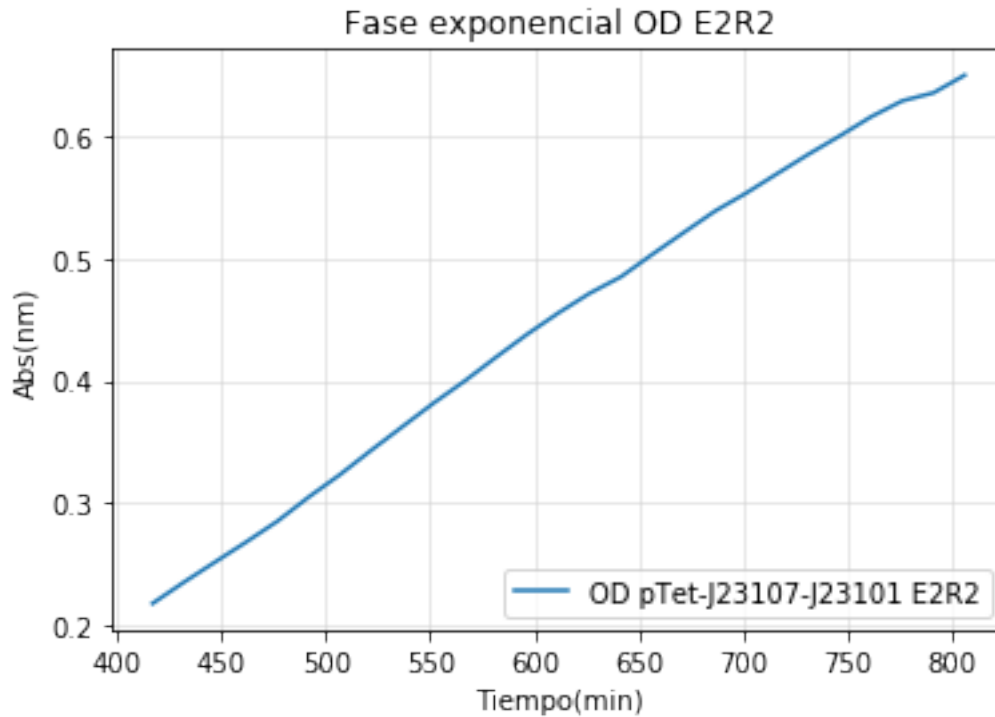


A=2.298058e+00
um=4.617185e-03

```
l=2.200403e+02  
Tm=4.031406e+02  
doubpe=1.501233e+02  
ext=3.753083e+02  
Tfinal=7.784489e+02
```

Out[29]: <matplotlib.legend.Legend at 0x267eb65dbe0>





```
In [30]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-107-std glicerol rep 3
y18= np.log(od1523g3)-np.log(np.min(od1523g3))
print('Min OD = %e'%((np.min(od1523g3))))
evaly, params=Function_fit(tt,y18,0,-1,title = 'Ajuste pTet-J23107-J23101 glicerol 0,2%')
A18= params[0]
um18=params[1]
l18=params[2]
print('A=%e'%(A18))
print('um=%e'%(um18))
print('l=%e'%(l18))

#Cálculo datos para determinar extensión de la fase exponencial
tm18=((A18/(np.exp(1)*um18))+l18)
print('Tm=%e'%(tm18))
t218=((np.log(2))/um18)
print('doubpe=%e'%(t218))
extdp18=2.5*t218
print('ext=%e'%extdp18)
ttot18=tm18+extdp18
print('Tfinal=%e'%ttot18)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[28]
```

```

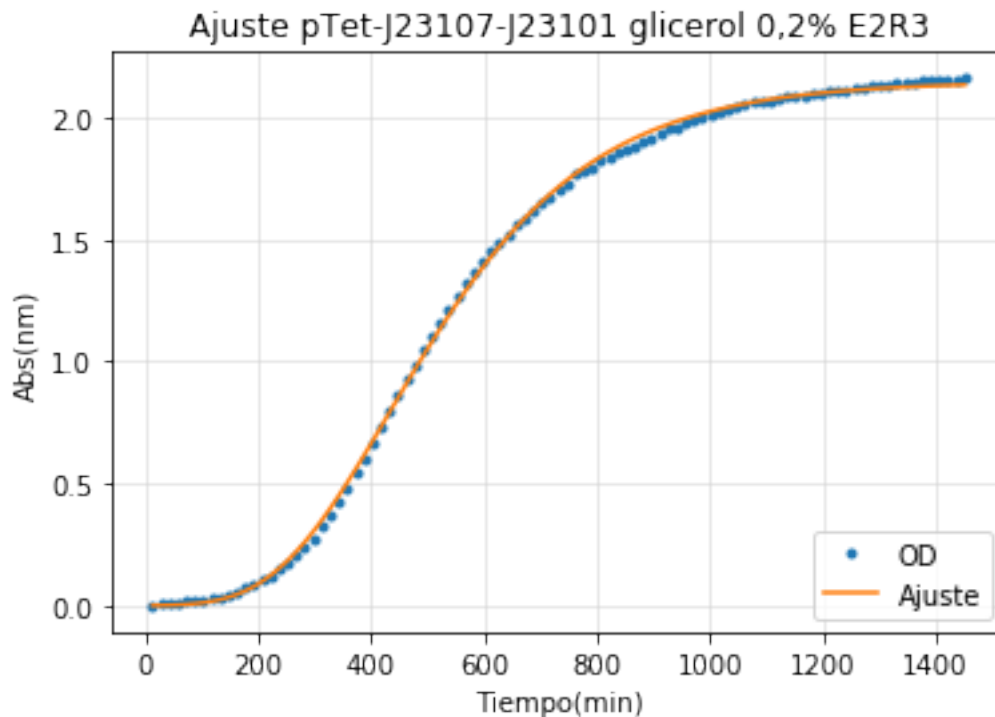
y2=tt[59]
plt.figure()
plt.title('pTet-J23107-J23101 Glicerol 0,2% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od1523g3,label='OD pTet-J23107-J23101 E3R3')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[28:60],od1523g3[28:60],label='OD pTet-J23107-J23101 E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.400000e-02

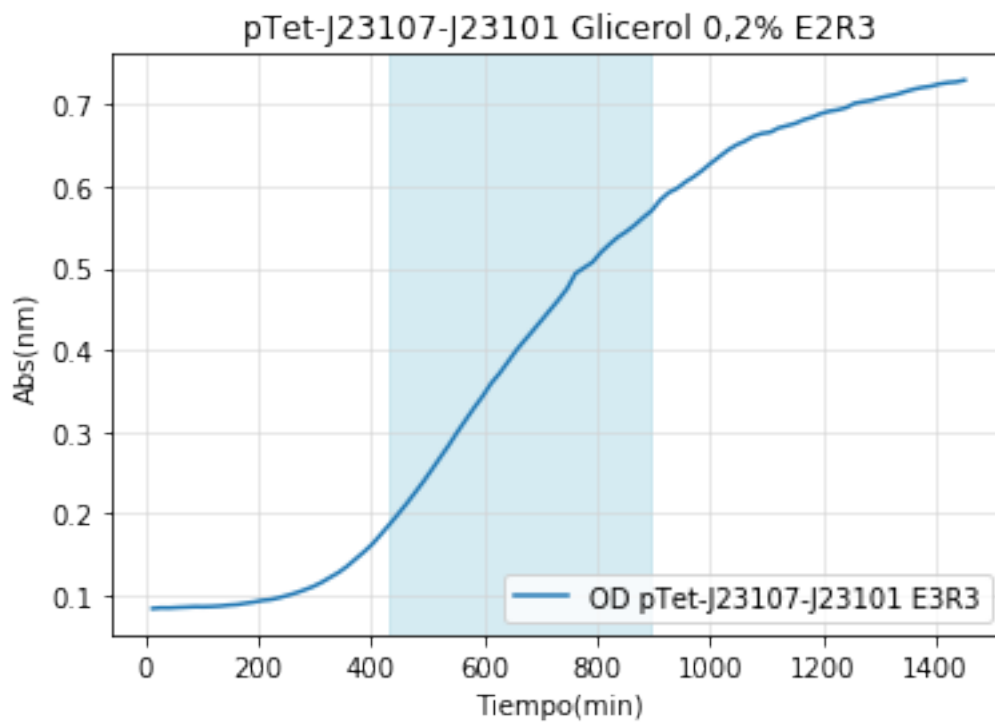
[2.14788549e+00 3.93566781e-03 2.30548233e+02]

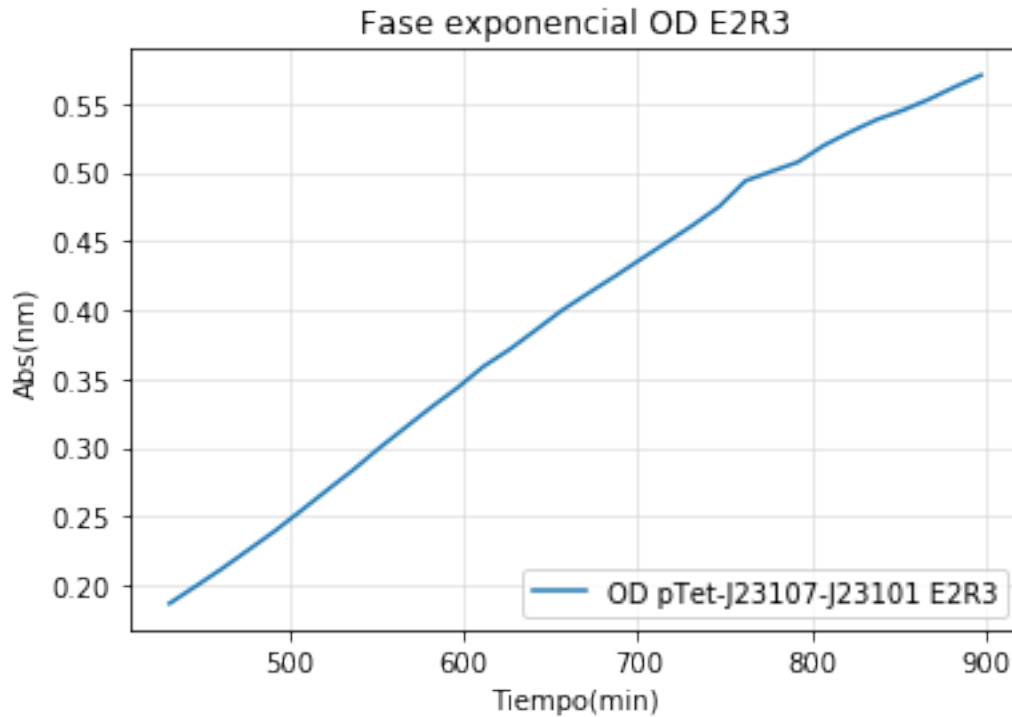


A=2.147885e+00
um=3.935668e-03

```
l=2.305482e+02  
Tm=4.313180e+02  
doubpe=1.761193e+02  
ext=4.402983e+02  
Tfinal=8.716163e+02
```

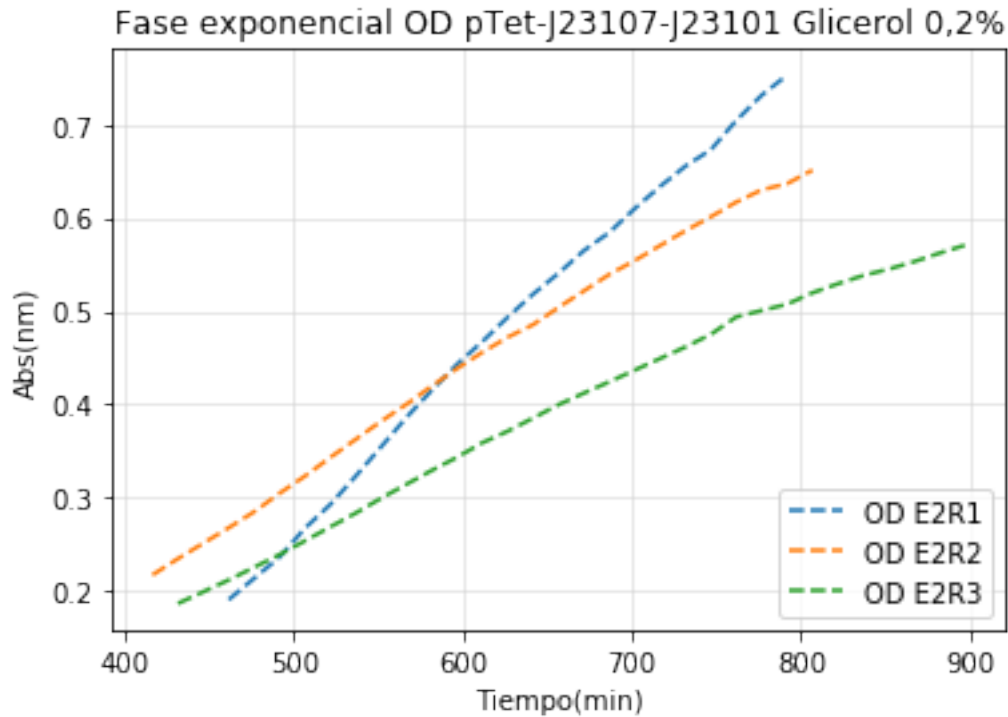
Out[30]: <matplotlib.legend.Legend at 0x267ecc6bfd0>





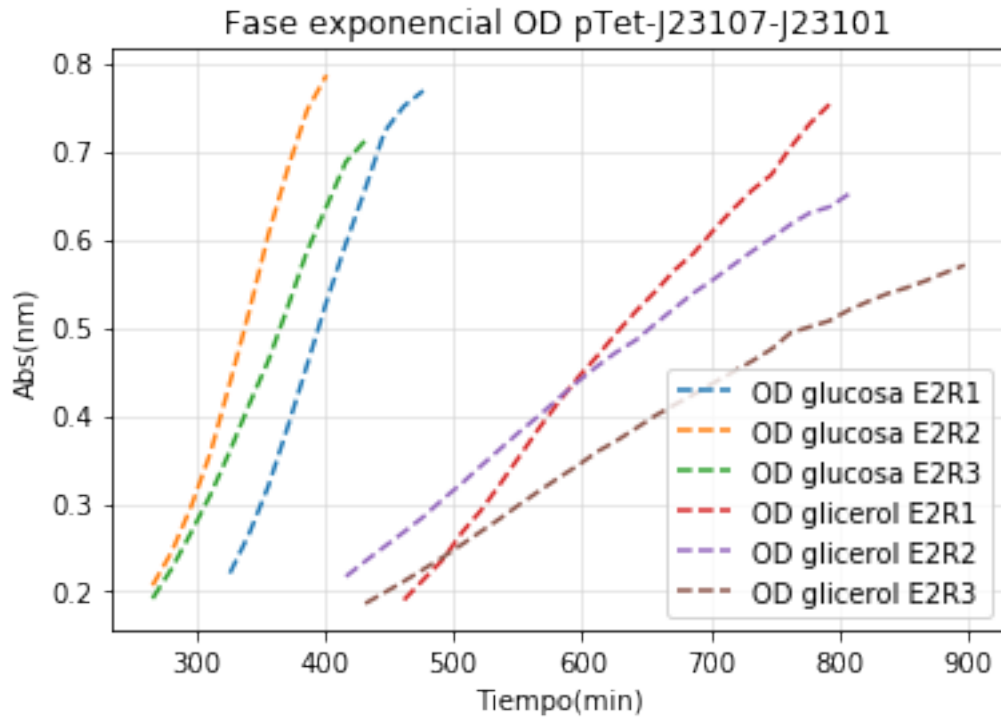
```
In [31]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD pTet-J23107-J23101 Glicerol 0,2%')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[30:53],od1523g1[30:53], '--',label='OD E2R1')
plt.plot(tt[27:54],od1523g2[27:54], '--',label='OD E2R2')
plt.plot(tt[28:60],od1523g3[28:60], '--',label='OD E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')
```

```
Out[31]: <matplotlib.legend.Legend at 0x267ecd79a58>
```



```
In [32]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD pTet-J23107-J23101')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[21:32],od15231[21:32], '--',label='OD glucosa E2R1')
plt.plot(tt[17:27],od15232[17:27], '--',label='OD glucosa E2R2')
plt.plot(tt[17:29],od15233[17:29], '--',label='OD glucosa E2R3')
plt.plot(tt[30:53],od1523g1[30:53], '--',label='OD glicerol E2R1')
plt.plot(tt[27:54],od1523g2[27:54], '--',label='OD glicerol E2R2')
plt.plot(tt[28:60],od1523g3[28:60], '--',label='OD glicerol E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')
```

```
Out[32]: <matplotlib.legend.Legend at 0x267ece3afd0>
```



```
In [33]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-plac-std glucosa rep 1
y19= np.log(od15261)-np.log(np.min(od15261))
print('Min OD = %e'%((np.min(od15261))))
evaly, params=Function_fit(tt,y19,0,-1,title = 'Ajuste pTet-pLacI-J23101 glucosa 0,4% E
A19= params[0]
um19=params[1]
l19=params[2]
print('A=%e'%(A19))
print('um=%e'%(um19))
print('l=%e'%(l19))

#Cálculo datos para determinar extensión de la fase exponencial
tm19=((A19/(np.exp(1)*um19))+l19)
print('Tm=%e'%(tm19))
t219=((np.log(2))/um19)
print('doubpe=%e'%(t219))
extdp19=2*t219
print('ext=%e'%extdp19)
ttot19=tm19+extdp19
print('Tfinal=%e'%ttot19)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[19]
```

```

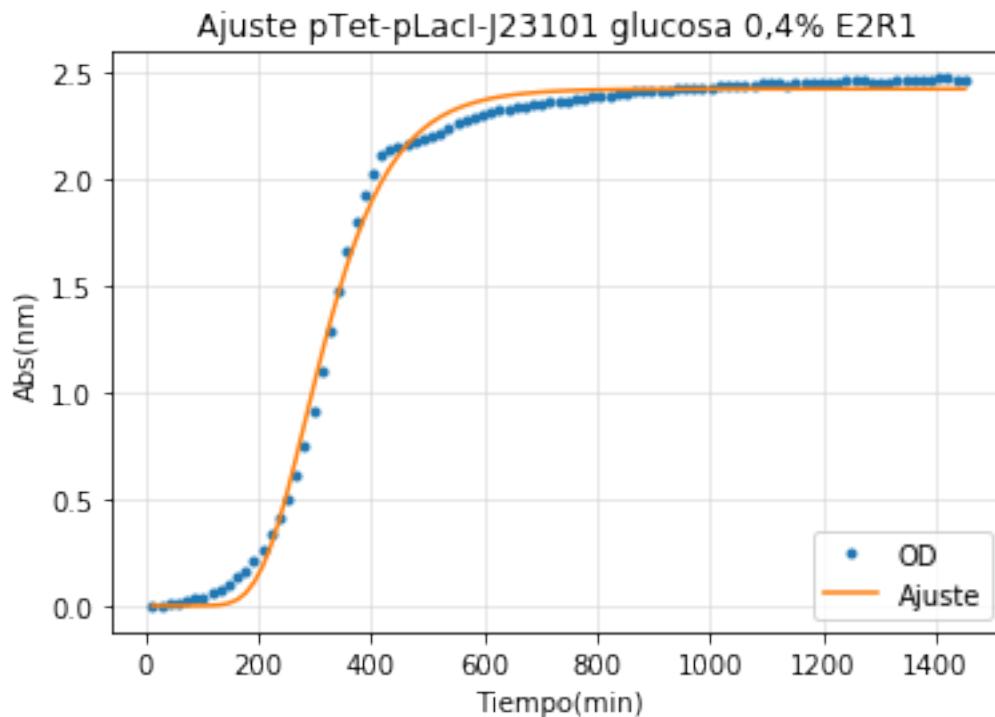
y2=tt[28]
plt.figure()
plt.title('pTet-pLacI-J23101 Glucosa 0,4% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od15261,label='OD pTet-pLacI-J23101 E2R1')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[19:29],od15261[19:29],label='OD pTet-pLacI-J23101 E2R1')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.530000e-02

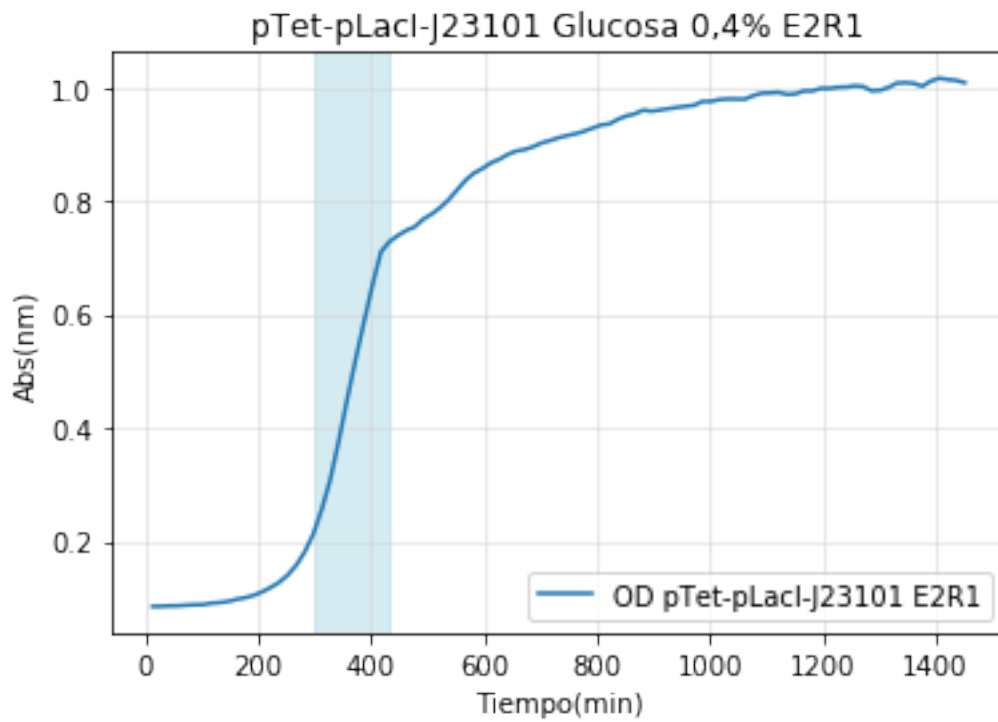
[2.42681432e+00 1.08626281e-02 2.02426630e+02]

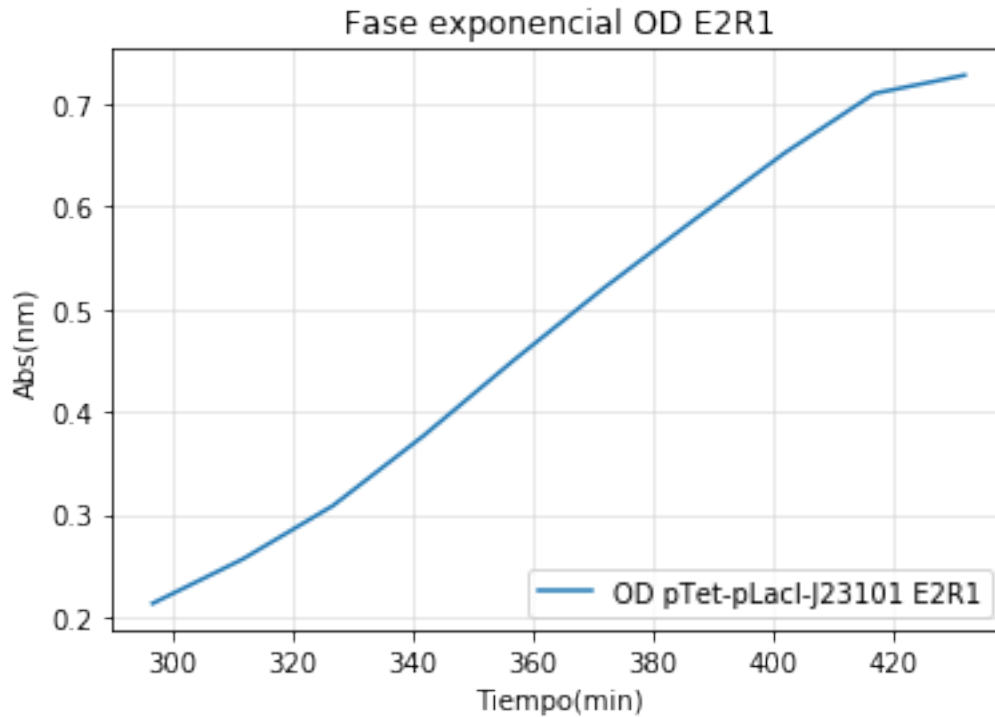


A=2.426814e+00
um=1.086263e-02

l=2.024266e+02
Tm=2.846144e+02
doubpe=6.381027e+01
ext=1.276205e+02
Tfinal=4.122349e+02

Out[33]: <matplotlib.legend.Legend at 0x267eb375ac8>





```
In [34]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-placI-std glucosa rep 2
y20= np.log(od15262)-np.log(np.min(od15262))
print('Min OD = %e'%((np.min(od15262))))
evaly, params=Function_fit(tt,y20,0,-1,title = 'Ajuste pTet-pLacI-J23101 glucosa 0,4% E
A20= params[0]
um20=params[1]
l20=params[2]
print('A=%e'%(A20))
print('um=%e'%(um20))
print('l=%e'%(l20))

#Cálculo datos para determinar extensión de la fase exponencial
tm20=((A20/(np.exp(1)*um20))+l20)
print('Tm=%e'%(tm20))
t220=((np.log(2))/um20)
print('doubpe=%e'%(t220))
extdp20=2*t220
print('ext=%e'%extdp20)
ttot20=tm20+extdp20
print('Tfinal=%e'%ttot20)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[18]
```

```

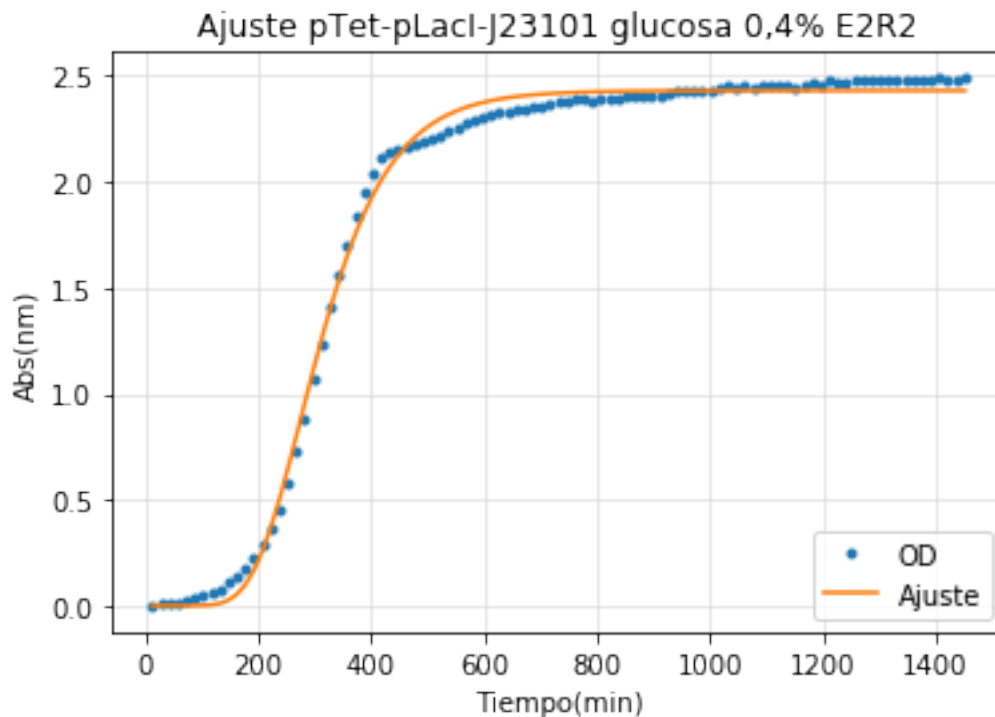
y2=tt[28]
plt.figure()
plt.title('pTet-pLacI-J23101 Glucosa 0,4% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od15262,label='OD pTet-pLacI-J23101 E2R2 ')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[18:29],od15262[18:29],label='OD pTet-pLacI-J23101 E2R2')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.650000e-02

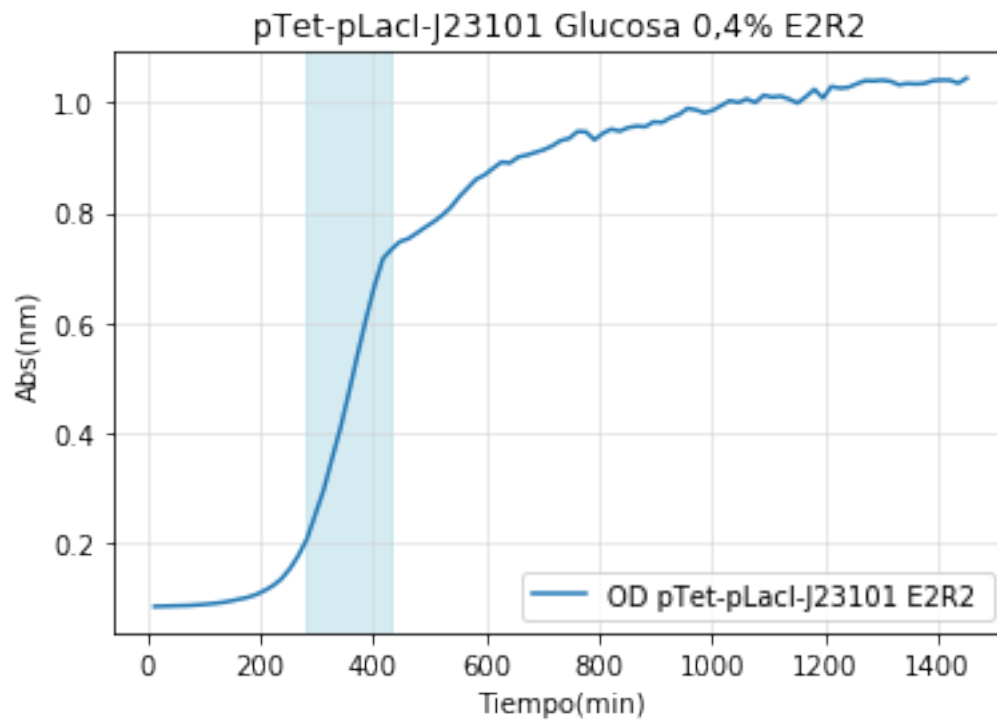
[2.43066594e+00 1.04655379e-02 1.90115277e+02]

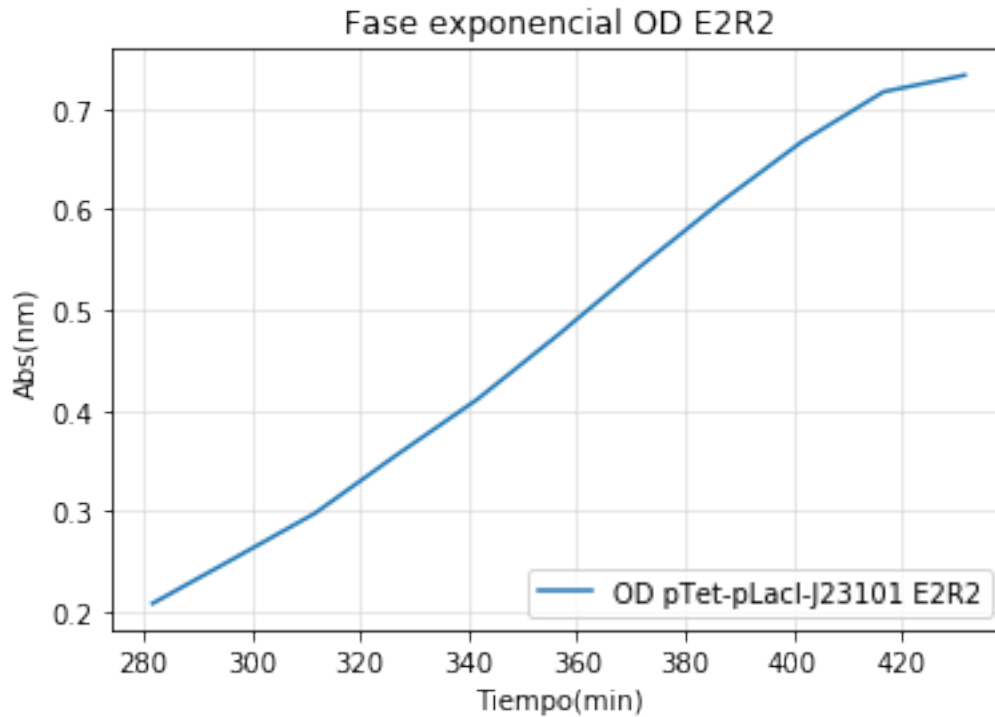


A=2.430666e+00
um=1.046554e-02

```
l=1.901153e+02  
Tm=2.755569e+02  
doubpe=6.623140e+01  
ext=1.324628e+02  
Tfinal=4.080196e+02
```

Out[34]: <matplotlib.legend.Legend at 0x267ecc57710>





```
In [35]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-plac-std glucosa rep 3
y21= np.log(od15263)-np.log(np.min(od15263))
print('Min OD = %e'%((np.min(od15263))))
evaly, params=Function_fit(tt,y21,0,-1,title = 'Ajuste pTet-pLacI-J23101 glucosa 0,4% E
A21= params[0]
um21=params[1]
l21=params[2]
print('A=%e'%(A21))
print('um=%e'%(um21))
print('l=%e'%(l21))

#Cálculo datos para determinar extensión de la fase exponencial
tm21=((A21/(np.exp(1)*um21))+l21)
print('Tm=%e'%(tm21))
t221=((np.log(2))/um21)
print('doubpe=%e'%(t221))
extdp21=2*t221
print('ext=%e'%extdp21)
ttot21=tm21+extdp21
print('Tfinal=%e'%ttot21)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[16]
```

```

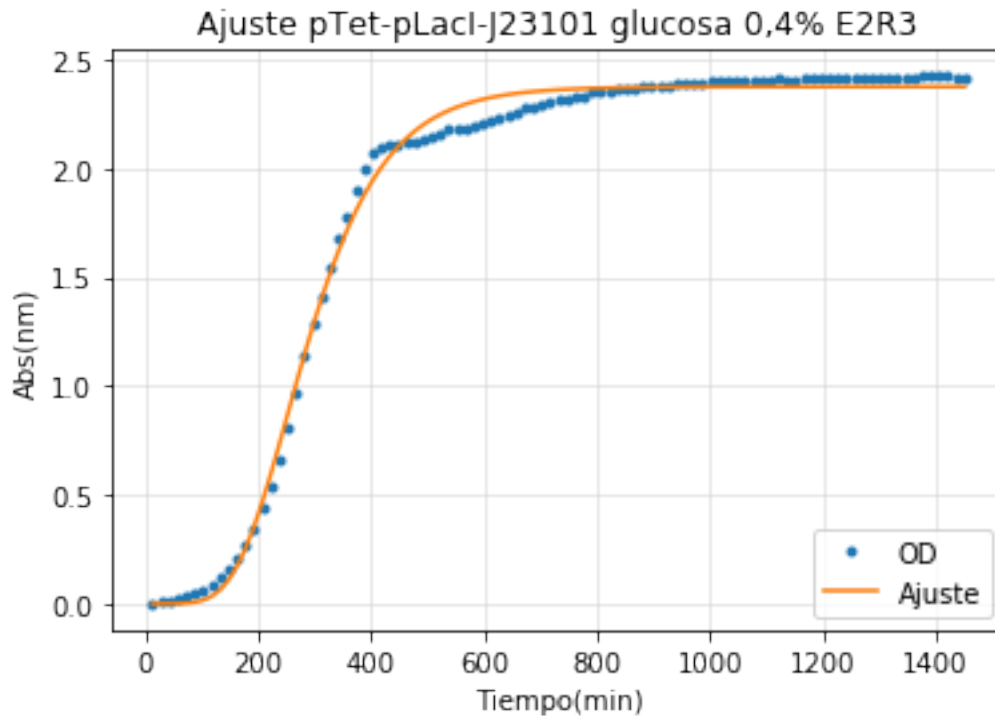
y2=tt[27]
plt.figure()
plt.title('pTet-pLacI-J23101 Glucosa 0,4% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od15263,label='OD pTet-pLacI-J23101 E2R3')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[16:28],od15263[16:28],label='OD pTet-pLacI-J23101 E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')

```

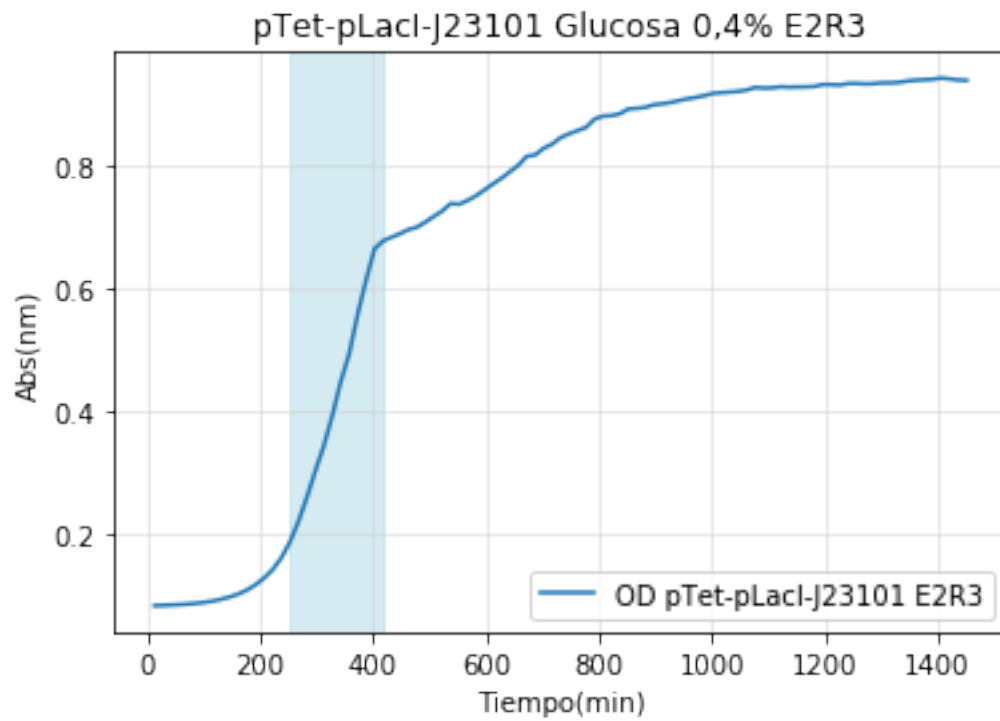
Min OD = 8.370000e-02

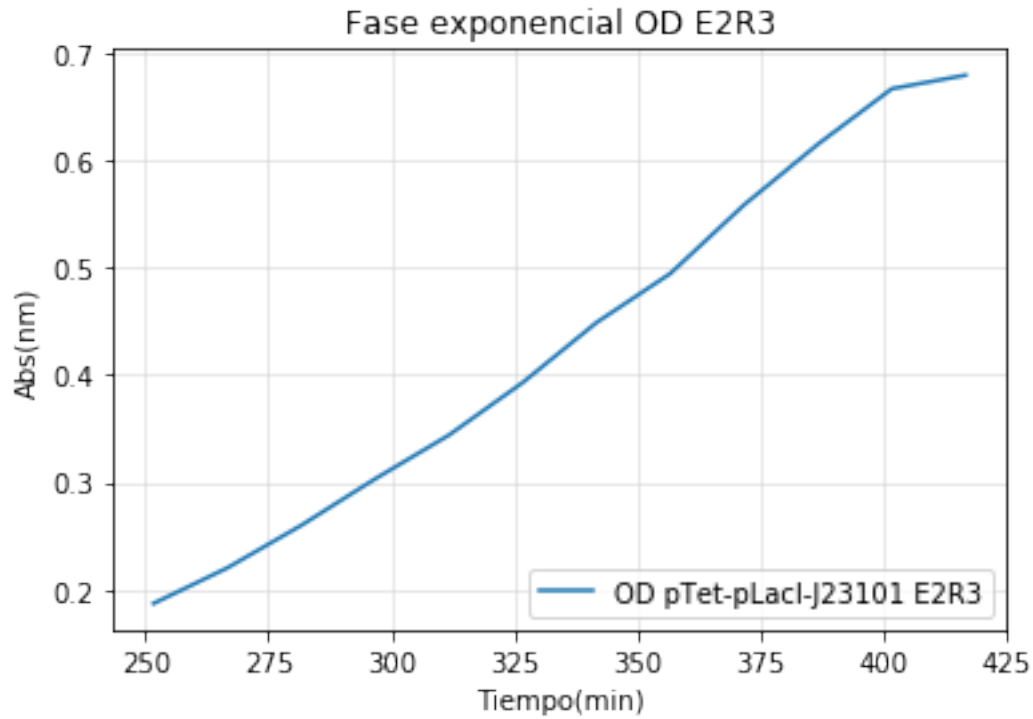
[2.37459827e+00 9.36201784e-03 1.58277013e+02]



A=2.374598e+00
um=9.362018e-03
l=1.582770e+02
Tm=2.515866e+02
doubpe=7.403822e+01
ext=1.480764e+02
Tfinal=3.996630e+02

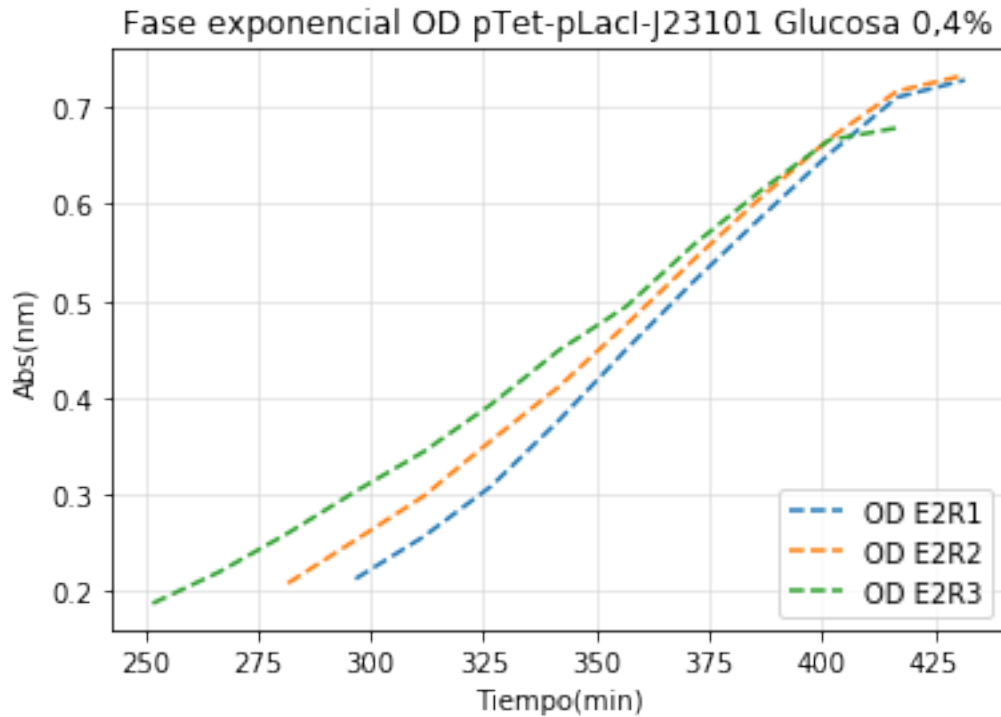
Out[35]: <matplotlib.legend.Legend at 0x267ed081518>





```
In [36]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD pTet-pLacI-J23101 Glucosa 0,4%')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[19:29],od15261[19:29], '--',label='OD E2R1')
plt.plot(tt[18:29],od15262[18:29], '--',label='OD E2R2')
plt.plot(tt[16:28],od15263[16:28], '--',label='OD E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')
```

```
Out[36]: <matplotlib.legend.Legend at 0x267ed196eb8>
```



```
In [37]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-plac-std glicerol rep 1
y22= np.log(od1526g1)-np.log(np.min(od1526g1))
print('Min OD = %e'%((np.min(od1526g1))))
evaly, params=Function_fit(tt,y22,0,-1,title = 'Ajuste pTet-pLacI-J23101 glicerol 0,2%')
A22= params[0]
um22=params[1]
l22=params[2]
print('A=%e'%(A22))
print('um=%e'%(um22))
print('l=%e'%(l22))

#Cálculo datos para determinar extensión de la fase exponencial
tm22=((A22/(np.exp(1)*um22))+l22)
print('Tm=%e'%(tm22))
t222=((np.log(2))/um22)
print('doubpe=%e'%(t222))
extdp22=2.5*t222
print('ext=%e'%extdp22)
ttot22=tm22+extdp22
print('Tfinal=%e'%ttot22)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[27]
```



```

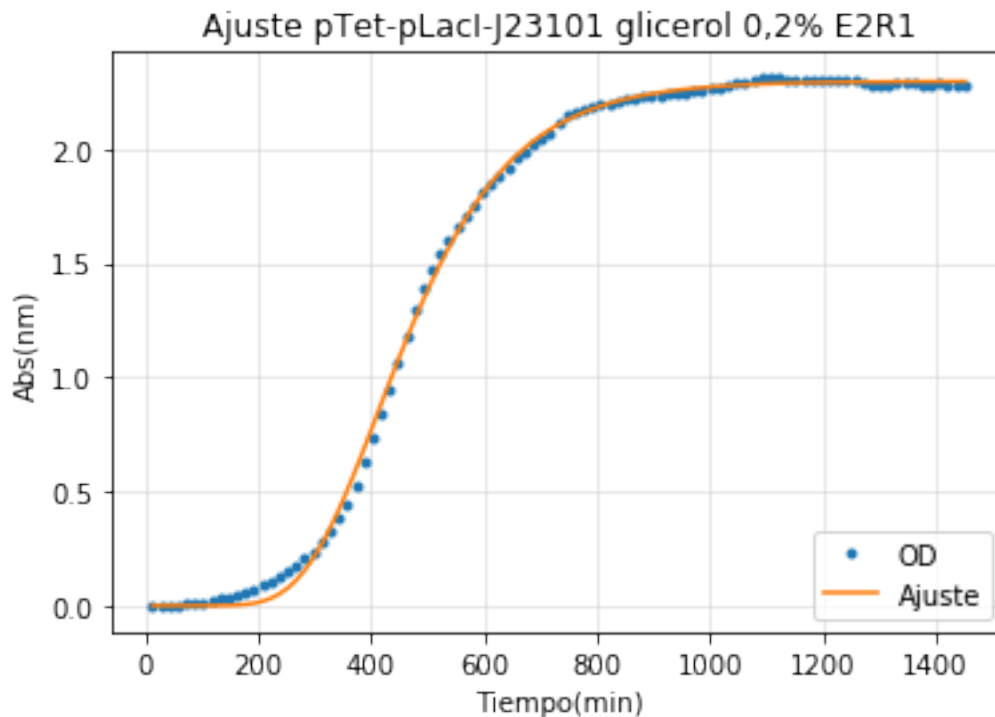
y2=tt[46]
plt.figure()
plt.title('pTet-pLacI-J23101 Glicerol 0,2% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od1526g1,label='OD pTet-pLacI-J23101 E2R1')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[27:47],od1526g1[27:47],label='OD pTet-pLacI-J23101 E2R1')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.570000e-02

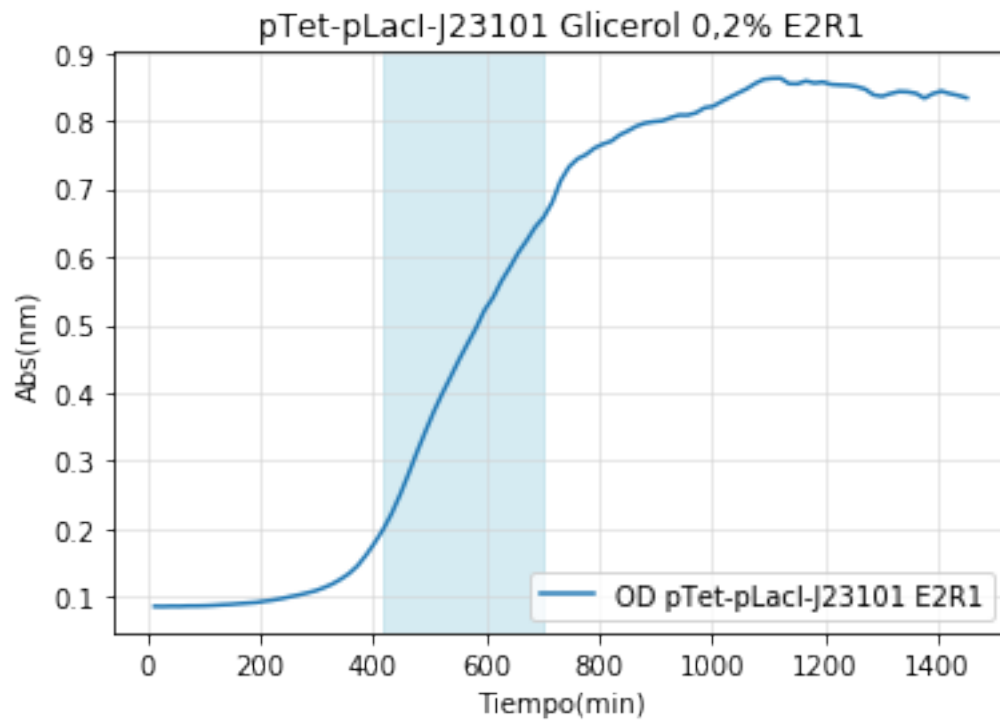
[2.29532726e+00 6.46446174e-03 2.80386578e+02]

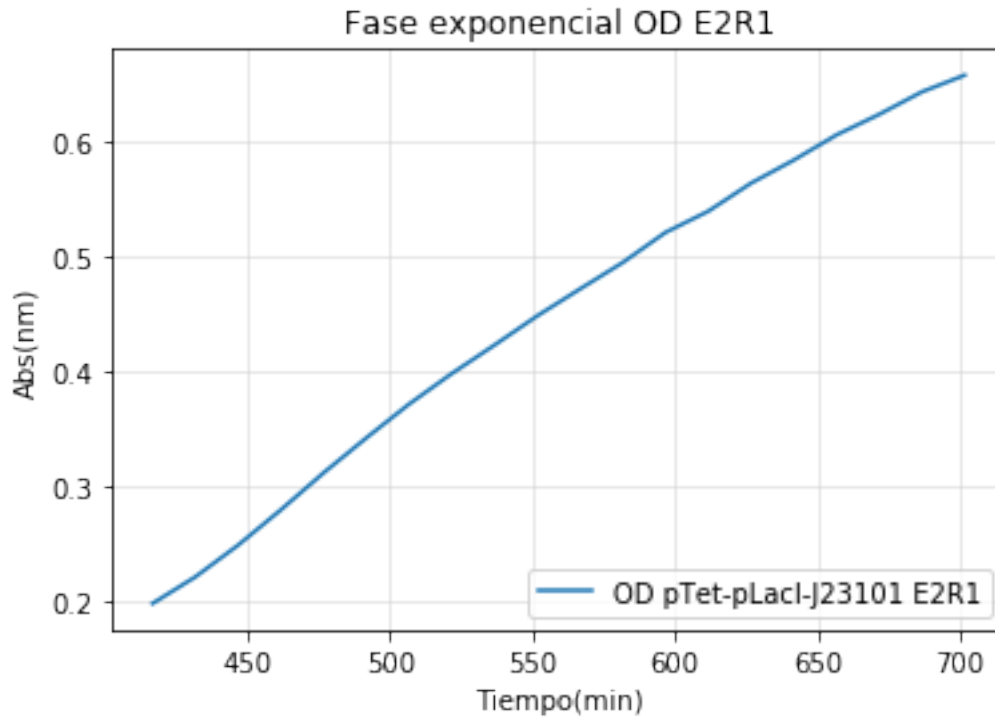


A=2.295327e+00
um=6.464462e-03

```
l=2.803866e+02  
Tm=4.110090e+02  
doubpe=1.072243e+02  
ext=2.680607e+02  
Tfinal=6.790697e+02
```

Out[37]: <matplotlib.legend.Legend at 0x267ecf91c50>





```
In [38]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-plac-std glicerol rep 2
y23= np.log(od1526g2)-np.log(np.min(od1526g2))
print('Min OD = %e'%((np.min(od1526g2))))
evaly, params=Function_fit(tt,y23,0,-1,title = 'Ajuste pTet-pLacI-J23101 glicerol 0,2%')
A23= params[0]
um23=params[1]
l23=params[2]
print('A=%e'%(A23))
print('um=%e'%(um23))
print('l=%e'%(l23))

#Cálculo datos para determinar extensión de la fase exponencial
tm23=((A23/(np.exp(1)*um23))+l23)
print('Tm=%e'%(tm23))
t223=((np.log(2))/um23)
print('doubpe=%e'%(t223))
extdp23=2.5*t223
print('ext=%e'%extdp23)
ttot23=tm23+extdp23
print('Tfinal=%e'%ttot23)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[24]
```

```

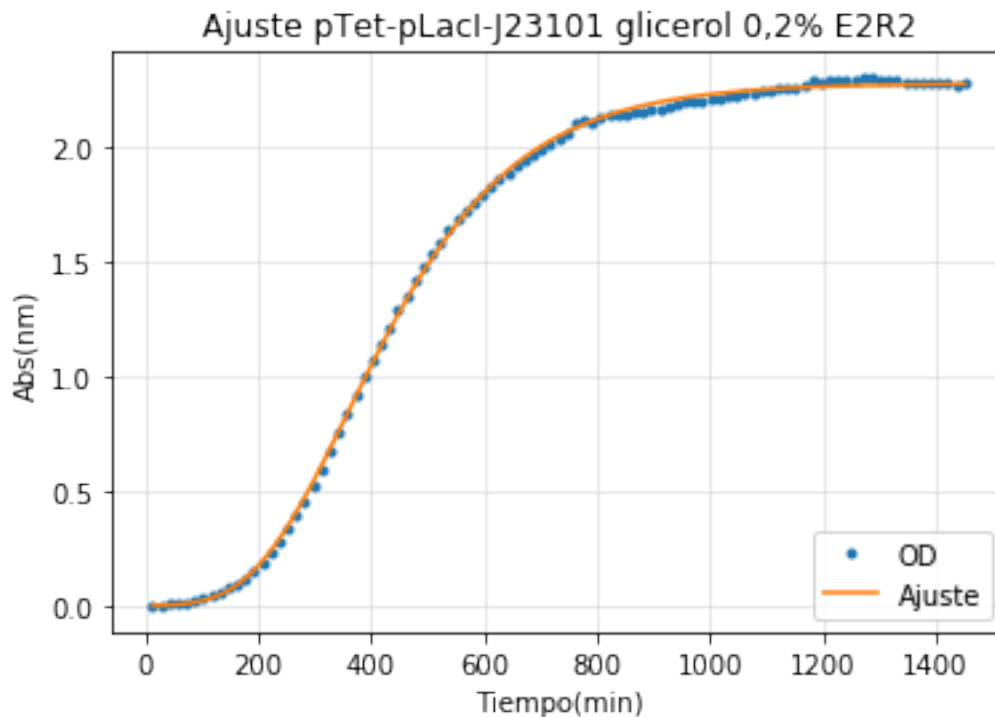
y2=tt[48]
plt.figure()
plt.title('pTet-pLacI-J23101 Glicerol 0,2% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od1526g2,label='OD pTet-pLacI-J23101 E2R2')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[24:49],od1526g2[24:49],label='OD pTet-pLacI-J23101 E2R2')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.610000e-02

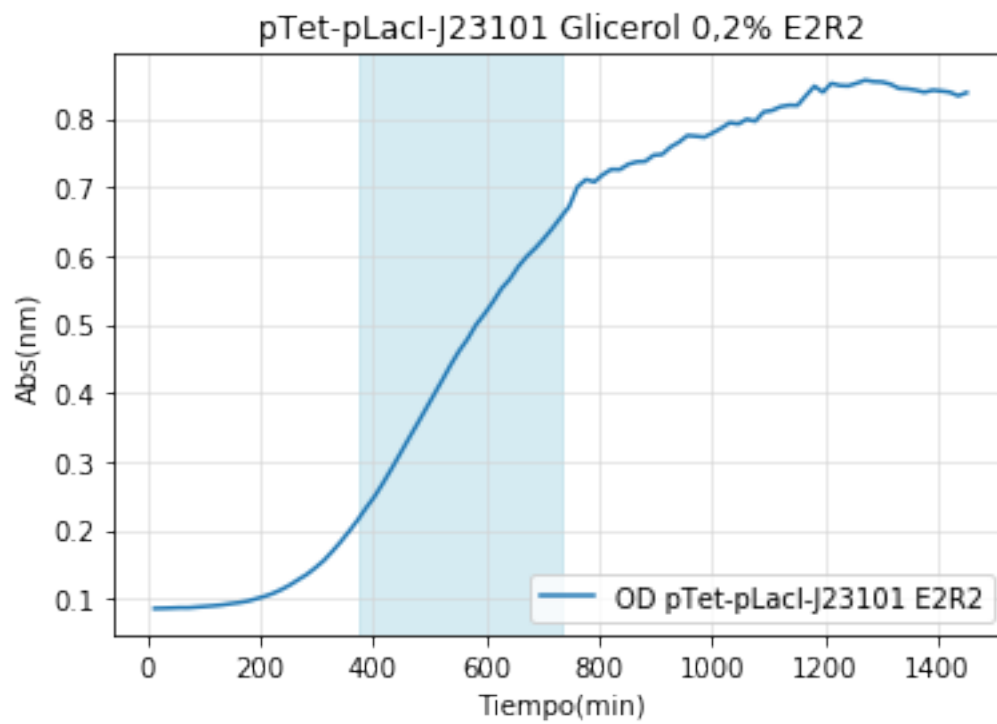
[2.27382560e+00 4.99992320e-03 1.89857406e+02]

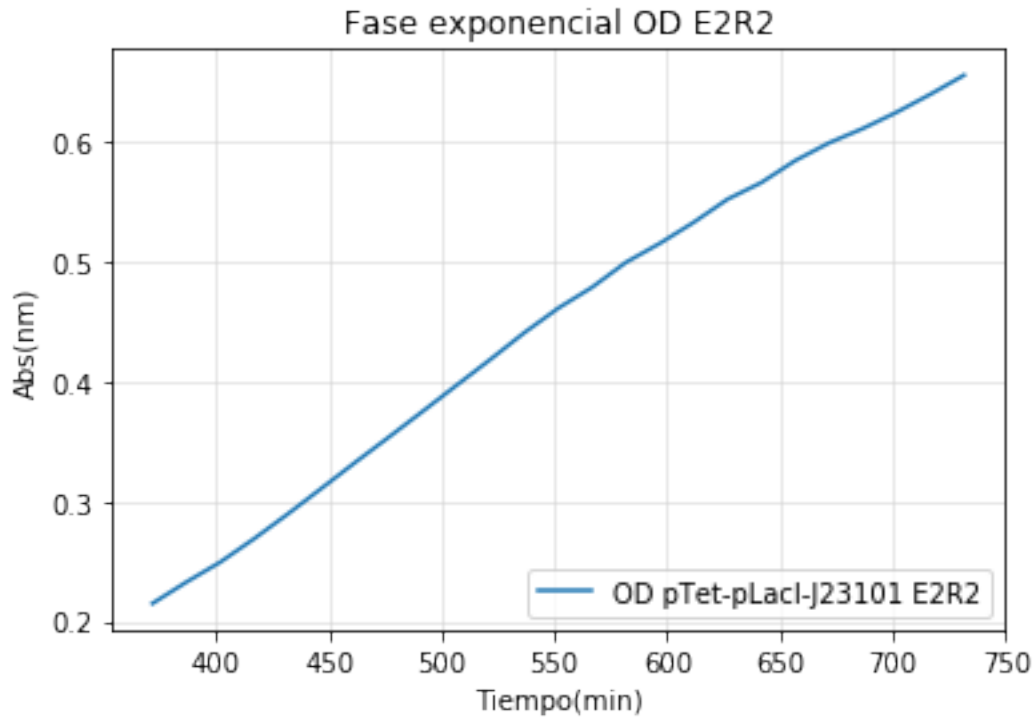


A=2.273826e+00
um=4.999923e-03

```
l=1.898574e+02  
Tm=3.571587e+02  
doubpe=1.386316e+02  
ext=3.465789e+02  
Tfinal=7.037376e+02
```

Out[38]: <matplotlib.legend.Legend at 0x267ec9e37f0>





```
In [39]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#pTet-plac-std glicerol rep 3
y24= np.log(od1526g3)-np.log(np.min(od1526g3))
print('Min OD = %e'%((np.min(od1526g3))))
evaly, params=Function_fit(tt,y24,0,-1,title = 'Ajuste pTet-pLacI-J23101 glicerol 0,2%')
A24= params[0]
um24=params[1]
l24=params[2]
print('A=%e'%(A24))
print('um=%e'%(um24))
print('l=%e'%(l24))

#Cálculo datos para determinar extensión de la fase exponencial
tm24=((A24/(np.exp(1)*um24))+l24)
print('Tm=%e'%(tm24))
t224=((np.log(2))/um24)
print('doubpe=%e'%(t224))
extdp24=2.5*t224
print('ext=%e'%extdp24)
ttot24=tm24+extdp24
print('Tfinal=%e'%ttot24)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[27]
```

```

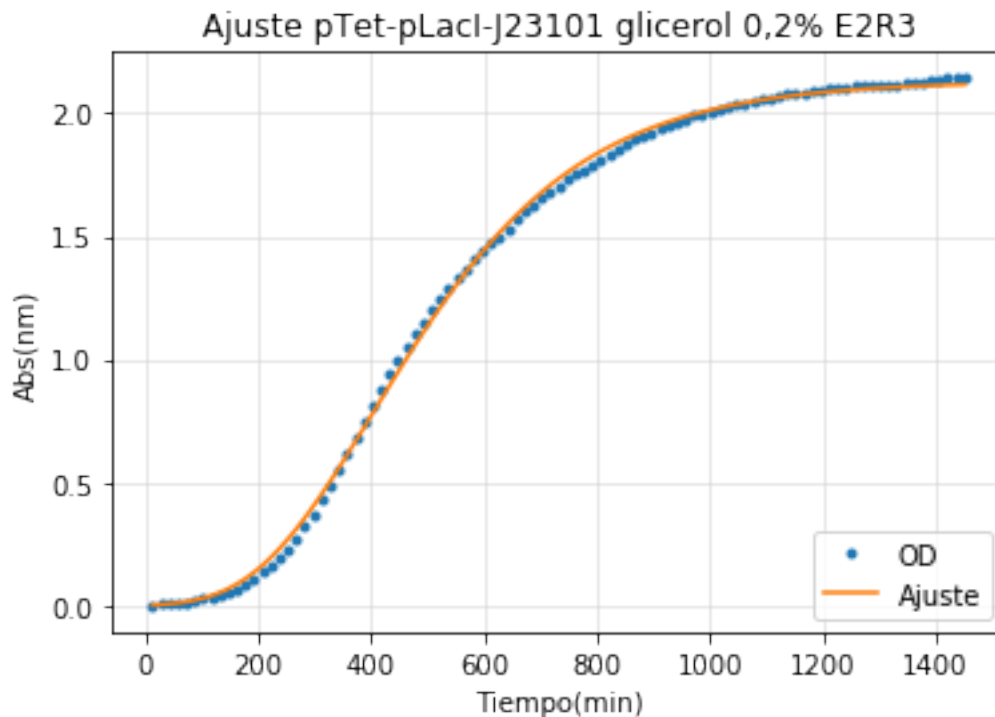
y2=tt[58]
plt.figure()
plt.title('pTet-pLacI-J23101 Glicerol 0,2% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od1526g3,label='OD pTet-pLacI-J23101 E2R3')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[27:59],od1526g3[27:59],label='OD pTet-pLacI-J23101 E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.340000e-02

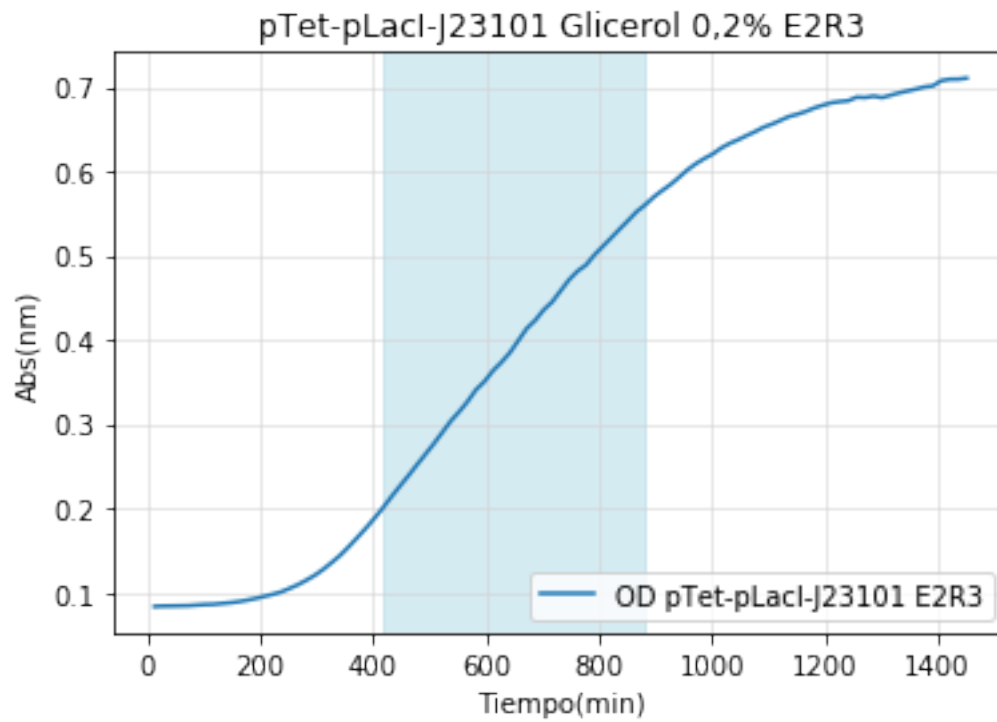
[2.13015451e+00 3.76288308e-03 1.94366098e+02]

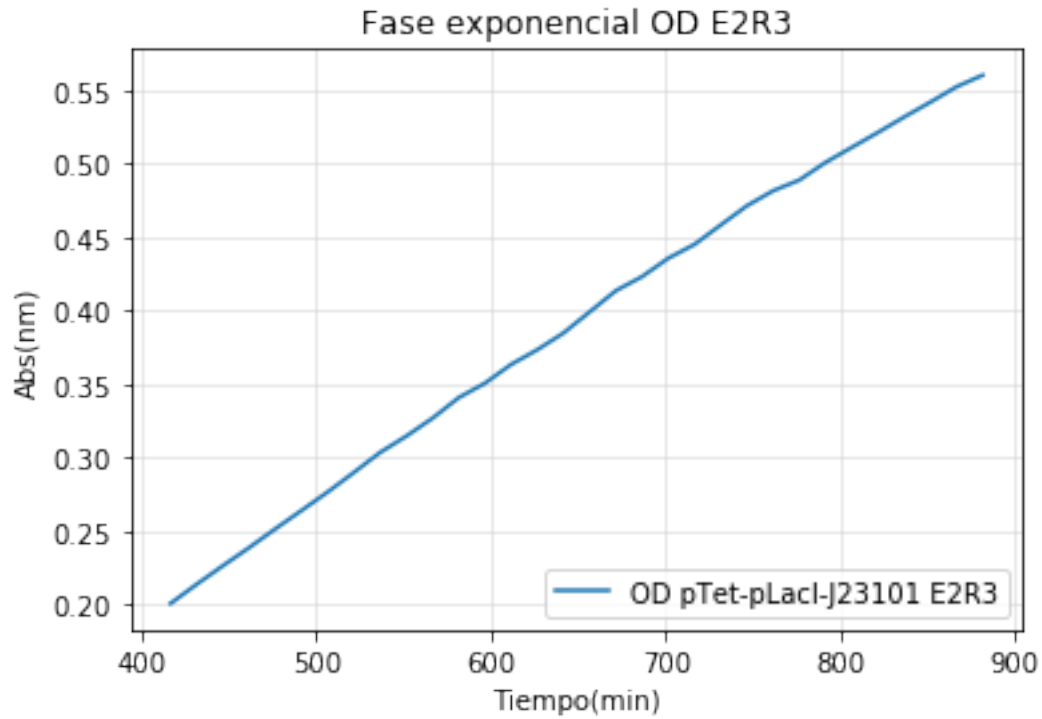


A=2.130155e+00
um=3.762883e-03

```
l=1.943661e+02  
Tm=4.026213e+02  
doubpe=1.842064e+02  
ext=4.605160e+02  
Tfinal=8.631373e+02
```

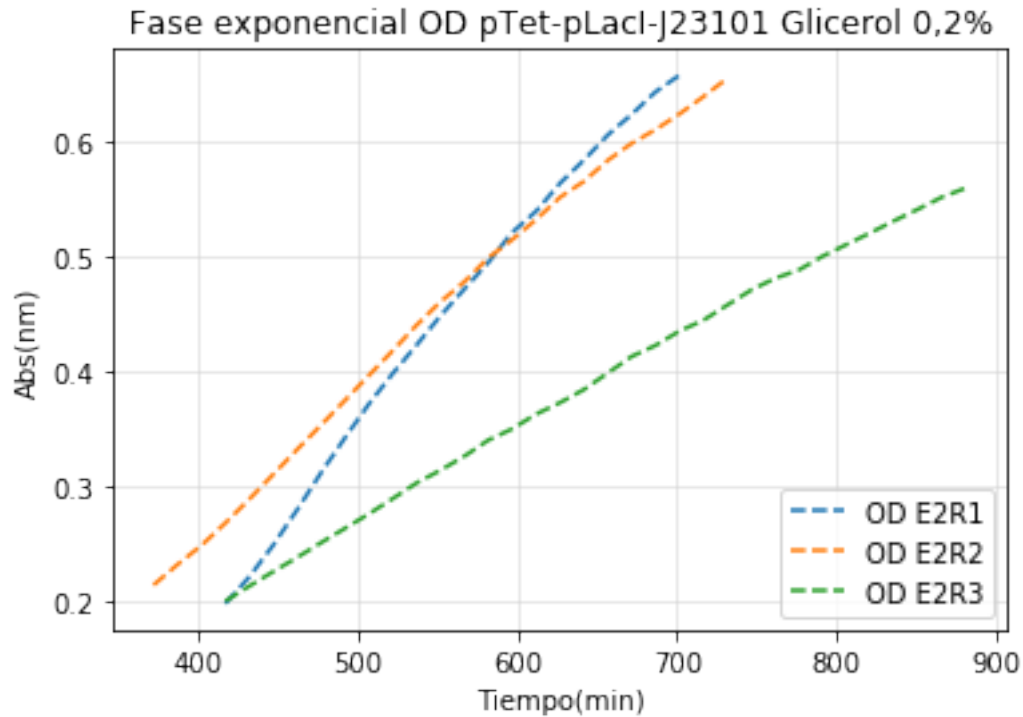
Out[39]: <matplotlib.legend.Legend at 0x267ed23d4a8>





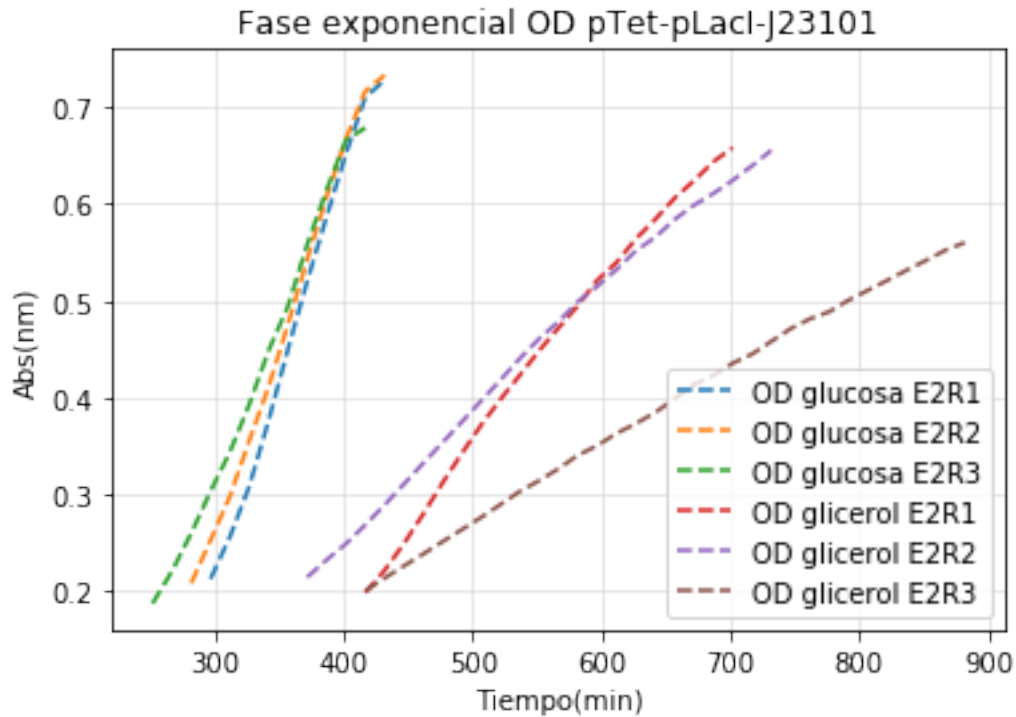
```
In [40]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD pTet-pLacI-J23101 Glicerol 0,2%')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[27:47],od1526g1[27:47], '--',label='OD E2R1')
plt.plot(tt[24:49],od1526g2[24:49], '--',label='OD E2R2')
plt.plot(tt[27:59],od1526g3[27:59], '--',label='OD E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')
```

```
Out[40]: <matplotlib.legend.Legend at 0x267ed3a9c88>
```



```
In [41]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD pTet-pLacI-J23101')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[19:29],od15261[19:29], '--',label='OD glucosa E2R1')
plt.plot(tt[18:29],od15262[18:29], '--',label='OD glucosa E2R2')
plt.plot(tt[16:28],od15263[16:28], '--',label='OD glucosa E2R3')
plt.plot(tt[27:47],od1526g1[27:47], '--',label='OD glicerol E2R1')
plt.plot(tt[24:49],od1526g2[24:49], '--',label='OD glicerol E2R2')
plt.plot(tt[27:59],od1526g3[27:59], '--',label='OD glicerol E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')
```

Out[41]: <matplotlib.legend.Legend at 0x267ed46ce80>



```
In [42]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-plas-std glucosa rep 1
y25= np.log(od15271)-np.log(np.min(od15271))
print('Min OD = %e'%((np.min(od15271))))
evaly, params=Function_fit(tt,y25,0,-1,title = 'Ajuste pTet-pLas81-J23101 glucosa 0,4%')
A25= params[0]
um25=params[1]
l25=params[2]
print('A=%e'%(A25))
print('um=%e'%(um25))
print('l=%e'%(l25))

#Cálculo datos para determinar extensión de la fase exponencial
tm25=((A25/(np.exp(1)*um25))+l25)
print('Tm=%e'%(tm25))
t225=((np.log(2))/um25)
print('doubpe=%e'%(t225))
extdp25=2*t225
print('ext=%e'%extdp25)
ttot25=tm25+extdp25
print('Tfinal=%e'%ttot25)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[19]
```

```

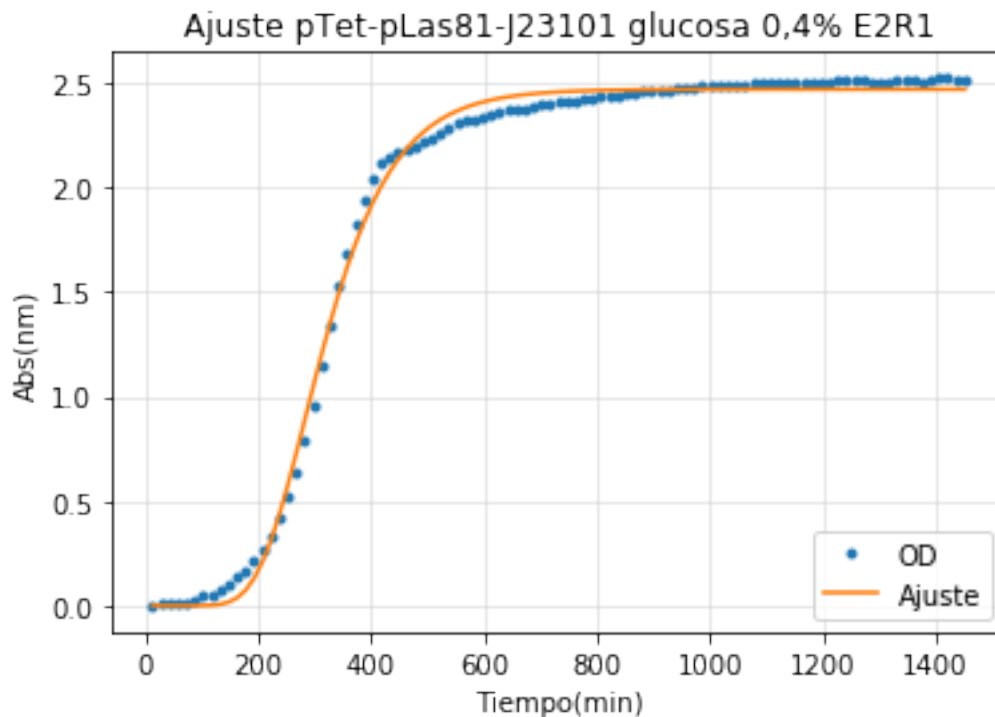
y2=tt[28]
plt.figure()
plt.title('pTet-pLas81-J23101 Glucosa 0,4% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od15271,label='OD pTet-pLas81-J23101 E2R1 ')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[18:29],od15271[18:29],label='OD pTet-pLas81-J23101 E2R1')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.510000e-02

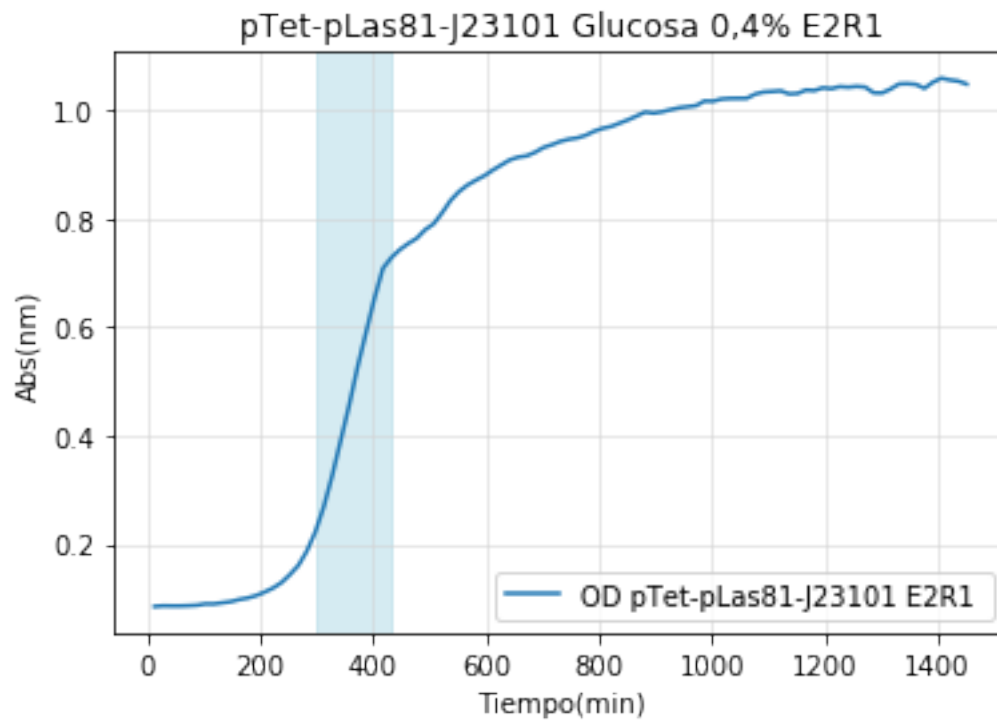
[2.46838538e+00 1.06746902e-02 1.98321599e+02]

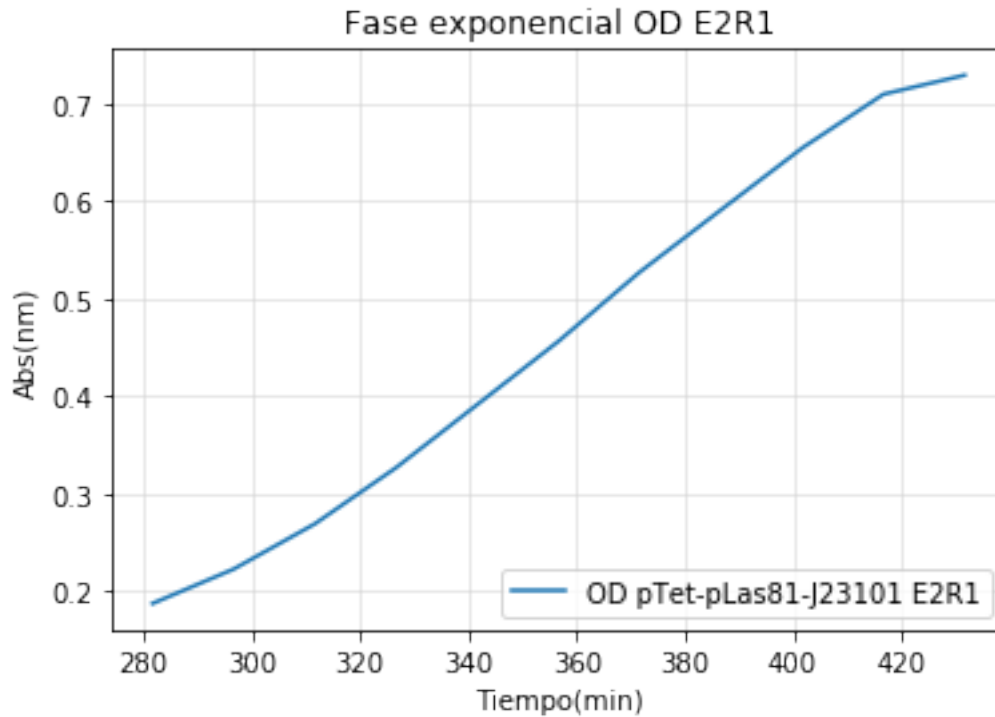


A=2.468385e+00
um=1.067469e-02

l=1.983216e+02
Tm=2.833890e+02
doubpe=6.493370e+01
ext=1.298674e+02
Tfinal=4.132564e+02

Out[42]: <matplotlib.legend.Legend at 0x267ed61a6d8>





```
In [43]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-pLas-std glucosa rep 2
y26= np.log(od15272)-np.log(np.min(od15272))
print('Min OD = %e'%((np.min(od15272))))
evaly, params=Function_fit(tt,y26,0,-1,title = 'Ajuste pTet-pLas81-J23101 glucosa 0,4%')
A26= params[0]
um26=params[1]
l26=params[2]
print('A=%e'%(A26))
print('um=%e'%(um26))
print('l=%e'%(l26))

#Cálculo datos para determinar extensión de la fase exponencial
tm26=((A26/(np.exp(1)*um26))+l26)
print('Tm=%e'%(tm26))
t226=((np.log(2))/um26)
print('doubpe=%e'%(t226))
extdp26=2*t226
print('ext=%e'%extdp26)
ttot26=tm26+extdp26
print('Tfinal=%e'%ttot26)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[16]
```

```

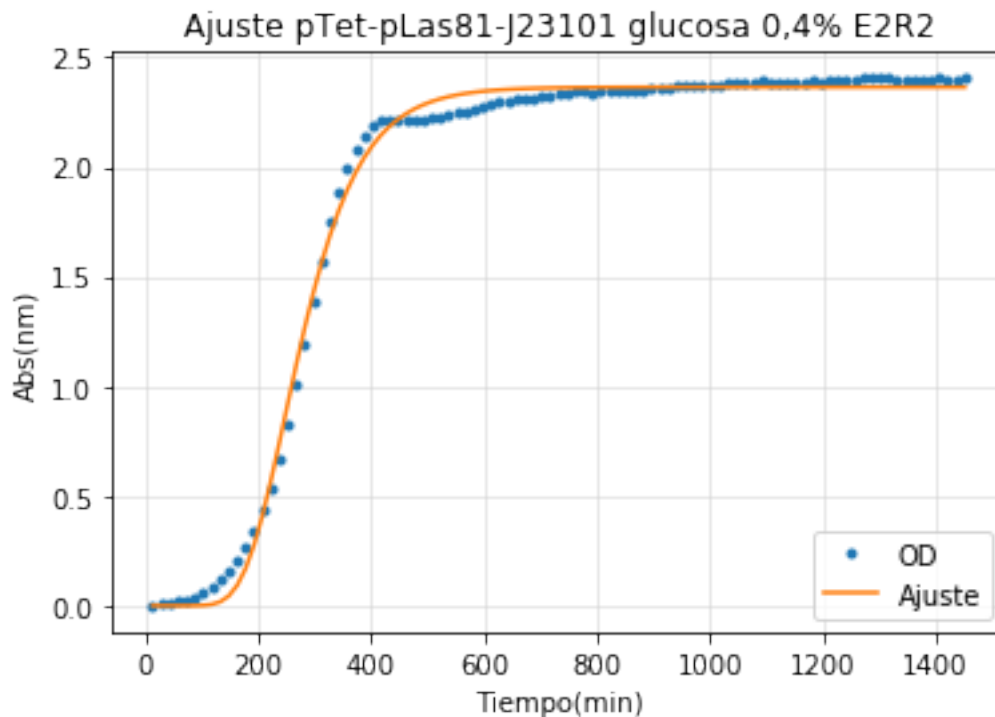
y2=tt[25]
plt.figure()
plt.title('pTet-pLas81-J23101 Glucosa 0,4% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od15272,label='OD pTet-pLas81-J23101 E2R2')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[16:26],od15272[16:26],label='OD pTet-pLas81-J23101 E2R2')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.580000e-02

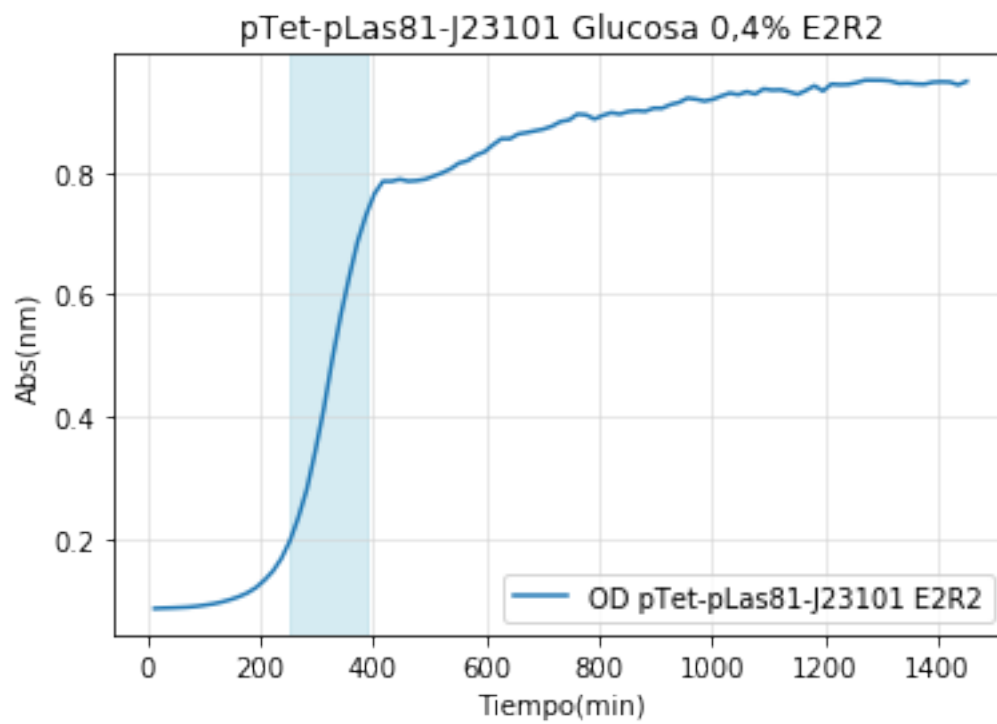
[2.36424970e+00 1.19691673e-02 1.74474154e+02]

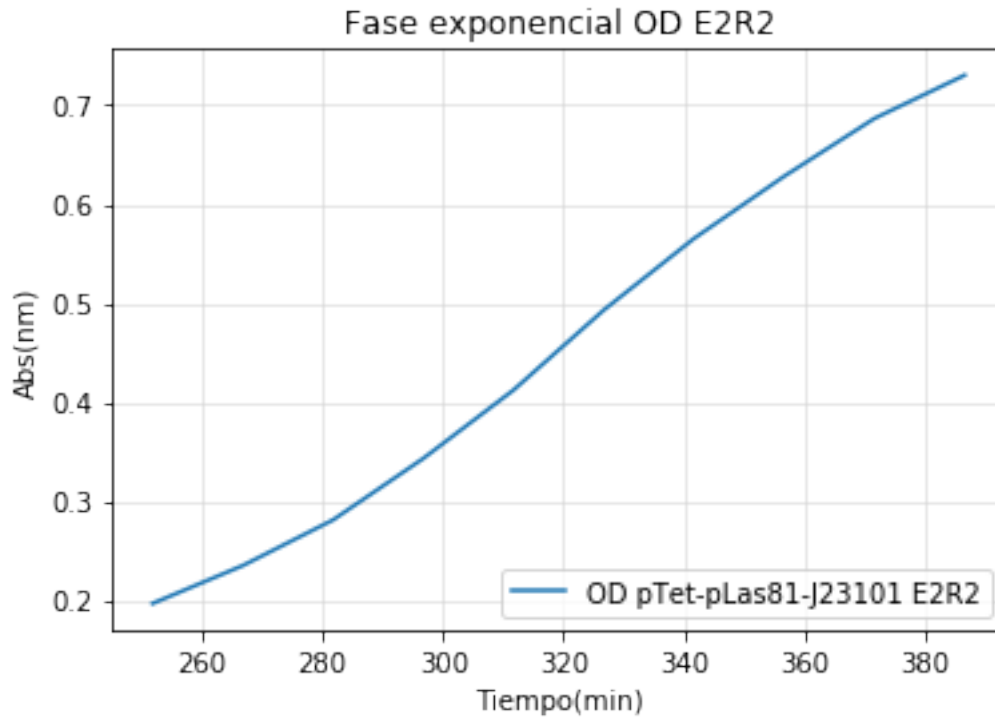


A=2.364250e+00
um=1.196917e-02

```
l=1.744742e+02  
Tm=2.471408e+02  
doubpe=5.791106e+01  
ext=1.158221e+02  
Tfinal=3.629629e+02
```

Out[43]: <matplotlib.legend.Legend at 0x267ecbb8a20>





```
In [44]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-pLas-std glucosa rep 3
y27= np.log(od15273)-np.log(np.min(od15273))
print('Min OD = %e'%((np.min(od15273))))
evaly, params=Function_fit(tt,y27,0,-1,title = 'Ajuste pTet-pLas81-J23101 glucosa 0,4%')
A27= params[0]
um27=params[1]
l27=params[2]
print('A=%e'%(A27))
print('um=%e'%(um27))
print('l=%e'%(l27))

#Cálculo datos para determinar extensión de la fase exponencial
tm27=((A27/(np.exp(1)*um27))+l27)
print('Tm=%e'%(tm27))
t227=((np.log(2))/um27)
print('doubpe=%e'%(t227))
extdp27=2*t227
print('ext=%e'%extdp27)
ttot27=tm27+extdp27
print('Tfinal=%e'%ttot27)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[16]
```

```

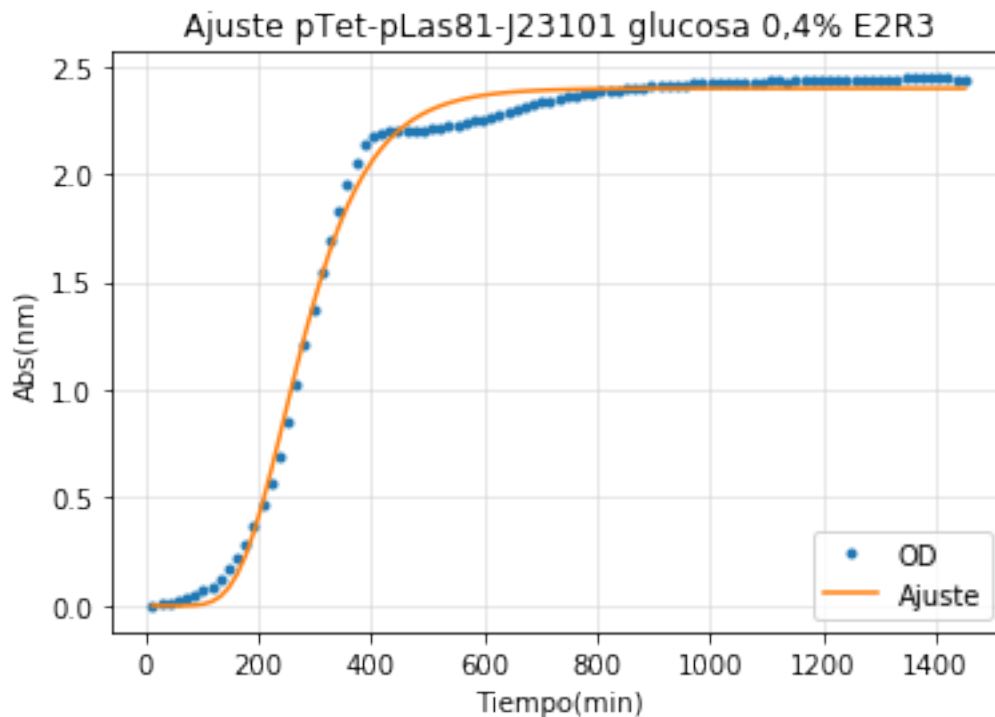
y2=tt[26]
plt.figure()
plt.title('pTet-pLas81-J23101 Glucosa 0,4% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od15273,label='OD pTet-pLas81-J23101 E2R3')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[16:27],od15273[16:27],label='OD pTet-pLas81-J23101 E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.390000e-02

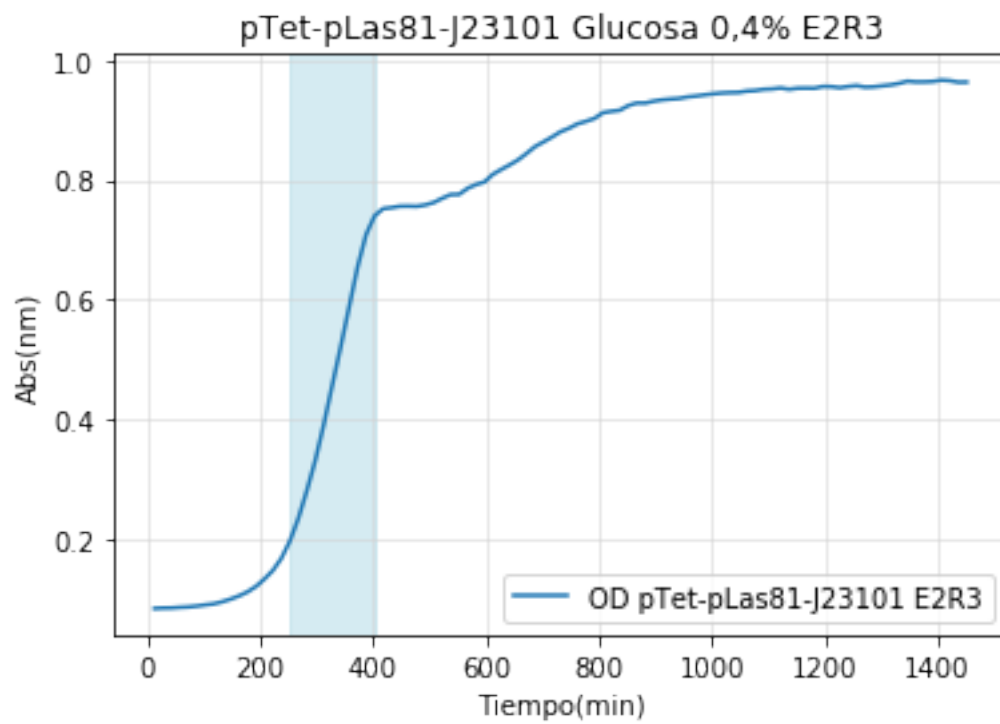
[2.39690020e+00 1.07573329e-02 1.64117750e+02]

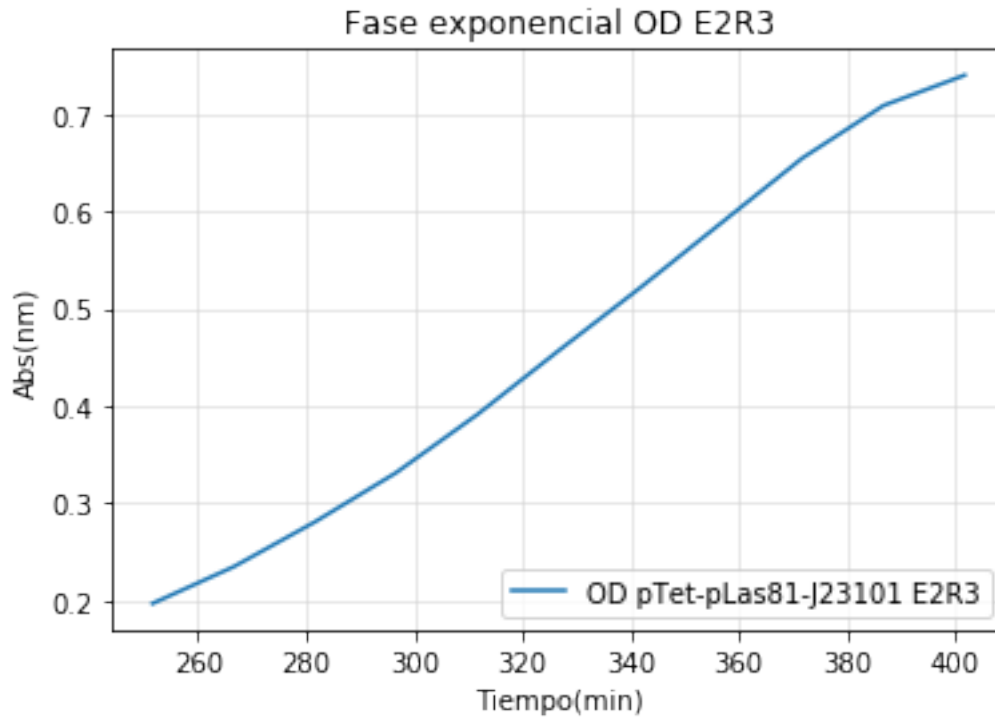


A=2.396900e+00
um=1.075733e-02

l=1.641177e+02
Tm=2.460870e+02
doubpe=6.443485e+01
ext=1.288697e+02
Tfinal=3.749567e+02

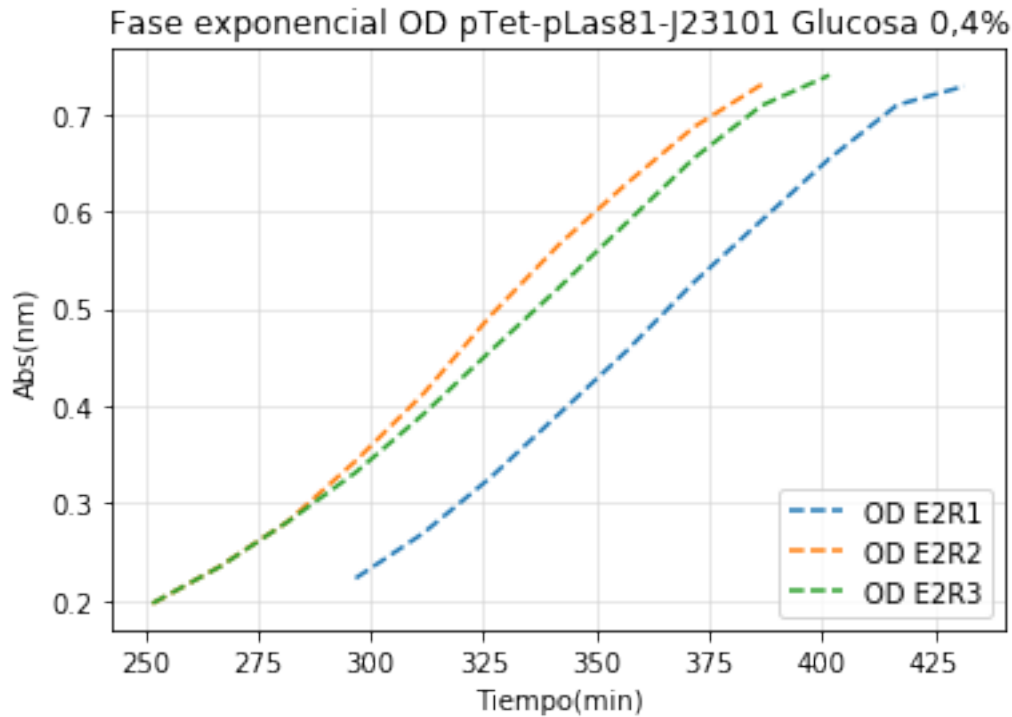
Out[44]: <matplotlib.legend.Legend at 0x267ed24cf60>





```
In [45]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD pTet-pLas81-J23101 Glucosa 0,4%')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[19:29],od15271[19:29], '--',label='OD E2R1')
plt.plot(tt[16:26],od15272[16:26], '--',label='OD E2R2')
plt.plot(tt[16:27],od15273[16:27], '--',label='OD E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5)
plt.legend(loc='lower right')
```

```
Out[45]: <matplotlib.legend.Legend at 0x267ee6db710>
```



```
In [46]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-pLas-std glicerol rep 1
y28= np.log(od1527g1)-np.log(np.min(od1527g1))
print('Min OD = %e'%((np.min(od1527g1))))
evaly, params=Function_fit(tt,y28,0,-1,title = 'Ajuste pTet-pLas81-J23101 glicerol 0,2%')
A28= params[0]
um28=params[1]
l28=params[2]
print('A=%e'%(A28))
print('um=%e'%(um28))
print('l=%e'%(l28))

#Cálculo datos para determinar extensión de la fase exponencial
tm28=((A28/(np.exp(1)*um28))+l28)
print('Tm=%e'%(tm28))
t228=((np.log(2))/um28)
print('doubpe=%e'%(t228))
extdp28=2.5*t228
print('ext=%e'%extdp28)
ttot28=tm28+extdp28
print('Tfinal=%e'%ttot28)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[29]
```

```

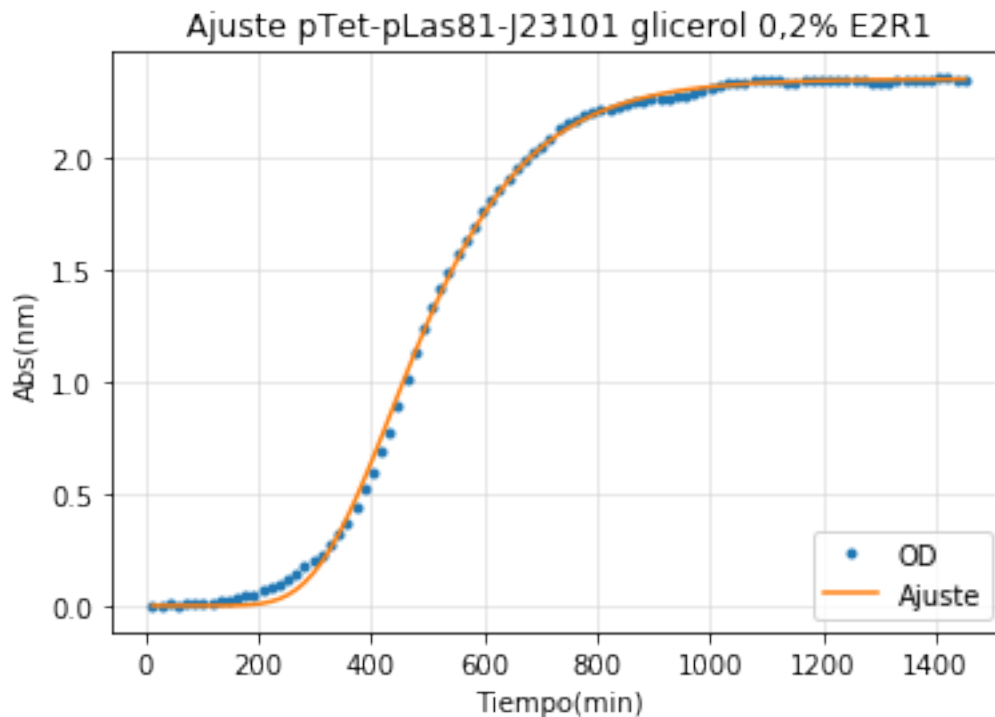
y2=tt[48]
plt.figure()
plt.title('pTet-pLas81-J23101 Glicerol 0,2% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od1527g1,label='OD pTet-pLas81-J23101 E2R1')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[29:49],od1527g1[29:49],label='OD pTet-pLas81-J23101 E2R1')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.620000e-02

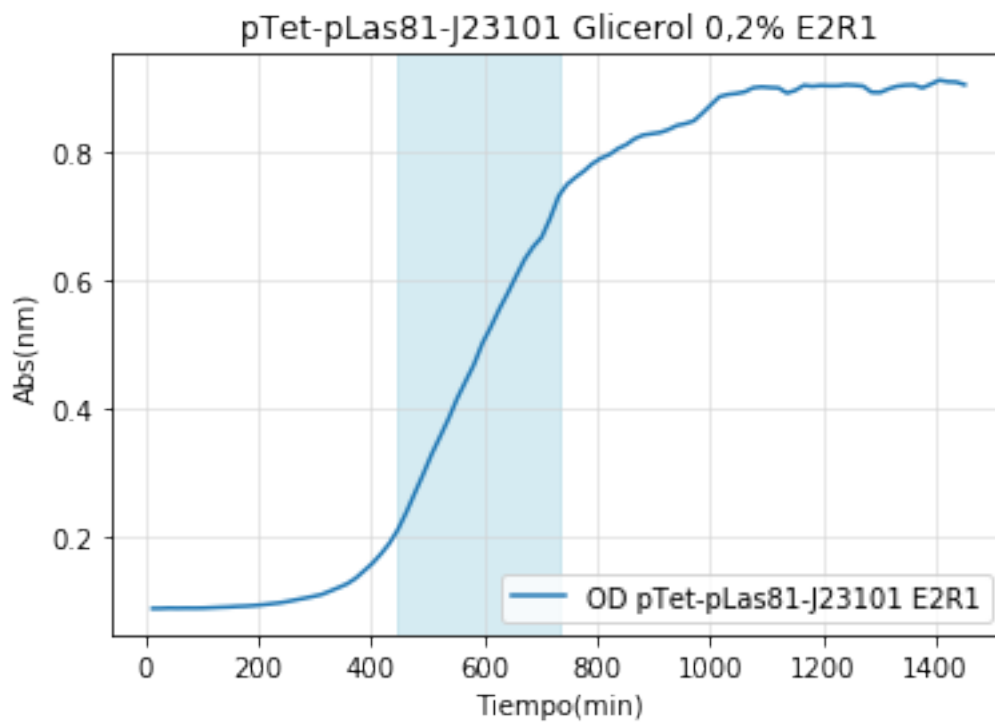
[2.35645905e+00 6.42374075e-03 3.00000000e+02]

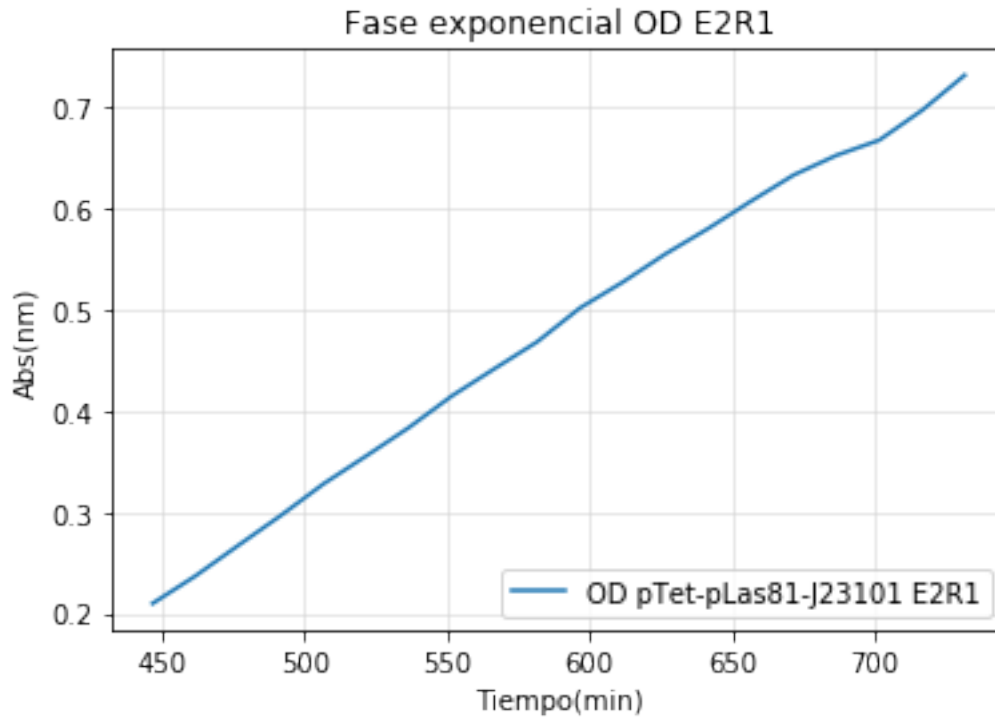


A=2.356459e+00
um=6.423741e-03

l=3.000000e+02
Tm=4.349514e+02
doubpe=1.079040e+02
ext=2.697599e+02
Tfinal=7.047114e+02

Out[46]: <matplotlib.legend.Legend at 0x267ee8719b0>





```
In [47]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-pLas-std glicerol rep 2
y29= np.log(od1527g2)-np.log(np.min(od1527g2))
print('Min OD = %e'%((np.min(od1527g2))))
evaly, params=Function_fit(tt,y29,0,-1,title = 'Ajuste pTet-pLas81-J23101 glicerol 0,2%')
A29= params[0]
um29=params[1]
l29=params[2]
print('A=%e'%(A29))
print('um=%e'%(um29))
print('l=%e'%(l29))

#Cálculo datos para determinar extensión de la fase exponencial
tm29=((A29/(np.exp(1)*um29))+l29)
print('Tm=%e'%(tm29))
t229=((np.log(2))/um29)
print('doubpe=%e'%(t229))
extdp29=2.5*t229
print('ext=%e'%extdp29)
ttot29=tm29+extdp29
print('Tfinal=%e'%ttot29)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[26]
```



```

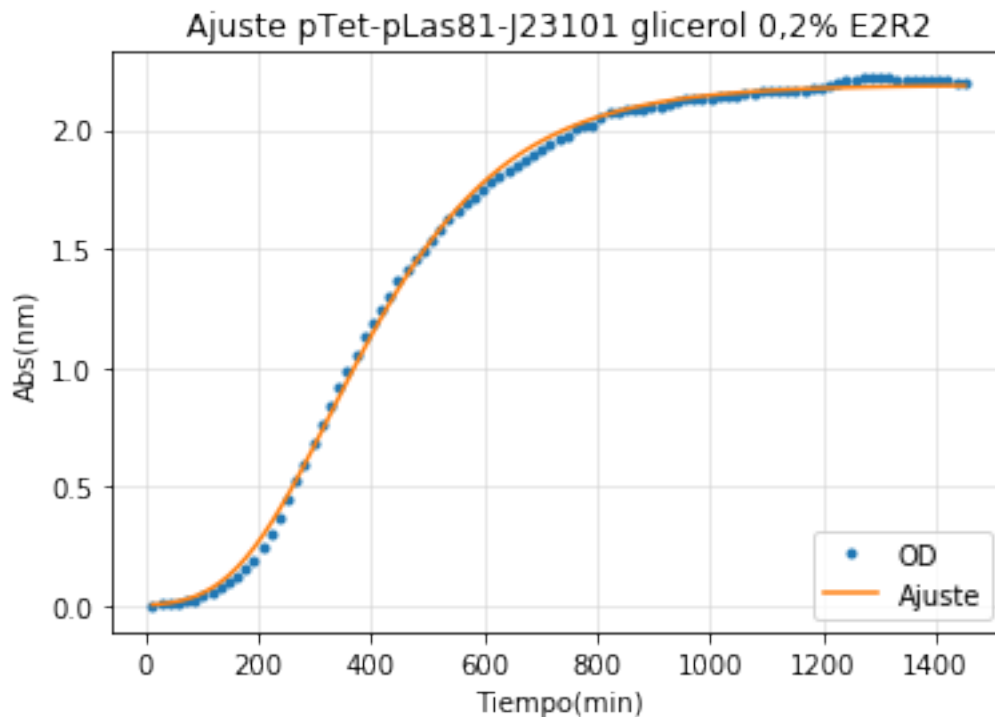
y2=tt[57]
plt.figure()
plt.title('pTet-pLas81-J23101 Glicerol 0,2% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od1527g2,label='OD pTet-pLas81-J23101 E2R2')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[26:58],od1527g2[26:58],label='OD pTet-pLas81-J23101 E2R2')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.590000e-02

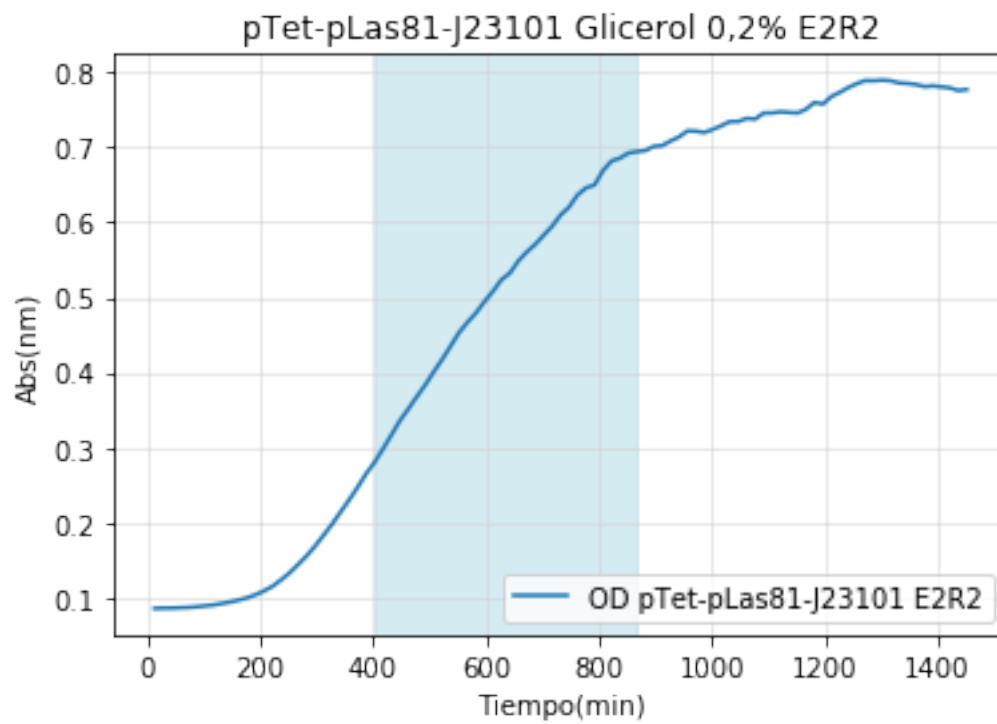
[2.18898193e+00 4.67358353e-03 1.54692670e+02]

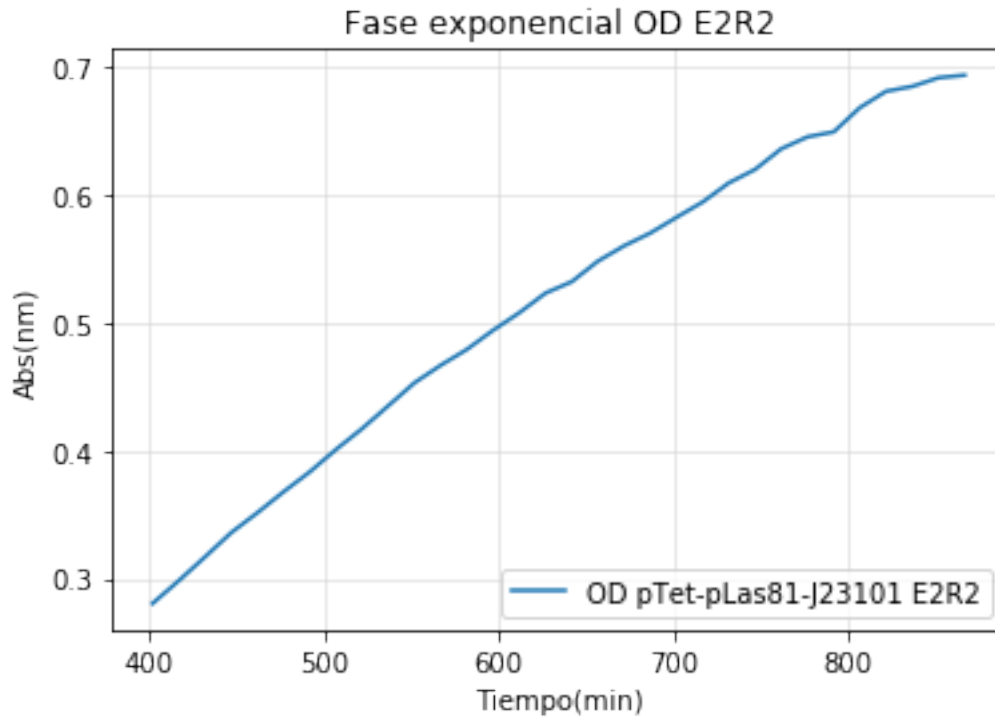


A=2.188982e+00
um=4.673584e-03

l=1.546927e+02
Tm=3.269976e+02
doubpe=1.483117e+02
ext=3.707793e+02
Tfinal=6.977769e+02

Out[47]: <matplotlib.legend.Legend at 0x267eea668d0>





```
In [48]: #Ajuste de datos con modelo Gompertz y obtención de parámetros para el ajuste
#ptet-pLas-std glicerol rep 3
y30= np.log(od1527g3)-np.log(np.min(od1527g3))
print('Min OD = %e'%((np.min(od1527g3))))
evaly, params=Function_fit(tt,y30,0,-1,title = 'Ajuste pTet-pLas81-J23101 glicerol 0,2%')
A30= params[0]
um30=params[1]
l30=params[2]
print('A=%e'%(A30))
print('um=%e'%(um30))
print('l=%e'%(l30))

#Cálculo datos para determinar extensión de la fase exponencial
tm30=((A30/(np.exp(1)*um30))+l30)
print('Tm=%e'%(tm30))
t230=((np.log(2))/um30)
print('doubpe=%e'%(t230))
extdp30=2.5*t230
print('ext=%e'%extdp30)
ttot30=tm30+extdp30
print('Tfinal=%e'%ttot30)

#Delimitación fase exponencial en grafico con OD/tiempo
y1=tt[25]
```

```

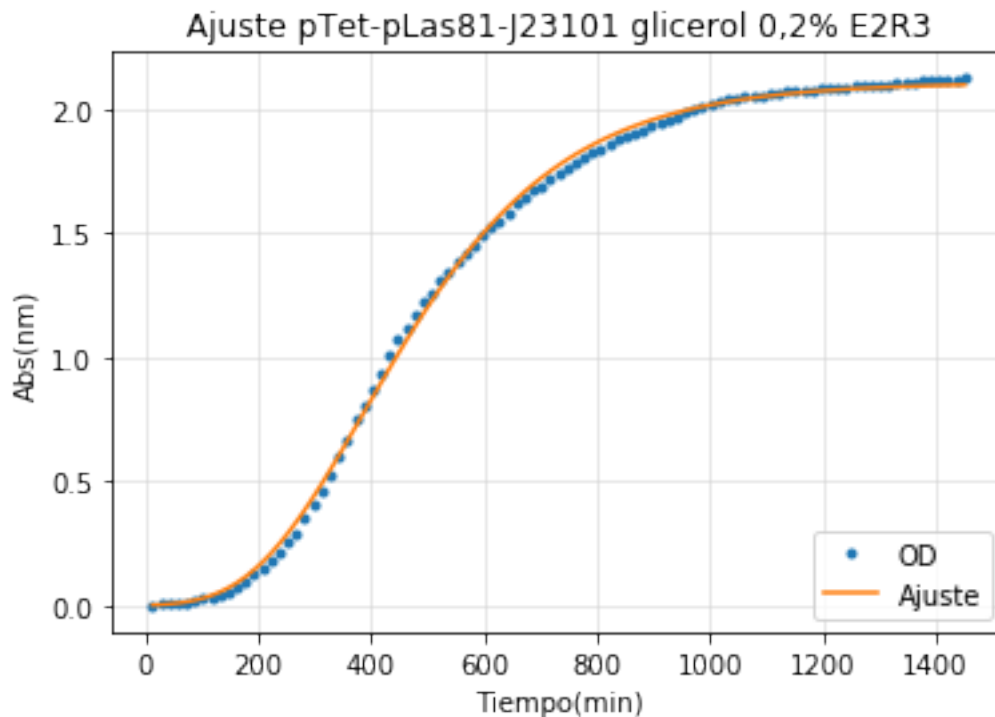
y2=tt[56]
plt.figure()
plt.title('pTet-pLas81-J23101 Glicerol 0,2% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt,od1527g3,label='OD pTet-pLas81-J23101 E2R3')
plt.axvspan(y1,y2, color='lightblue', alpha=0.5)
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

#Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[25:57],od1527g3[25:57],label='OD pTet-pLas81-J23101 E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')

```

Min OD = 8.350000e-02

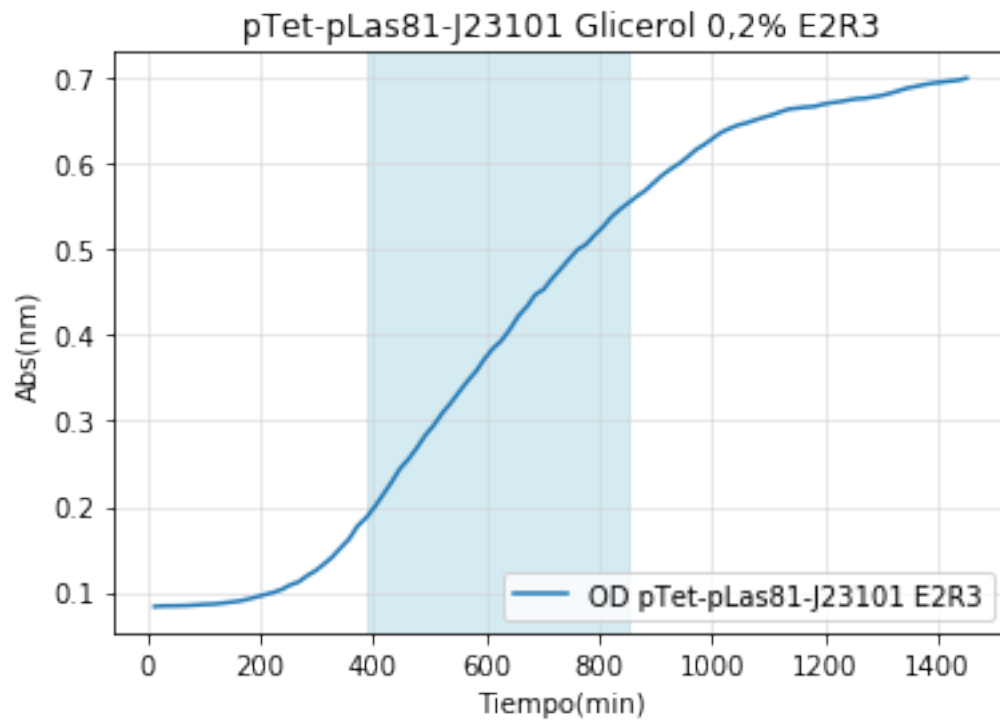
[2.10762223e+00 3.94236767e-03 1.88946108e+02]

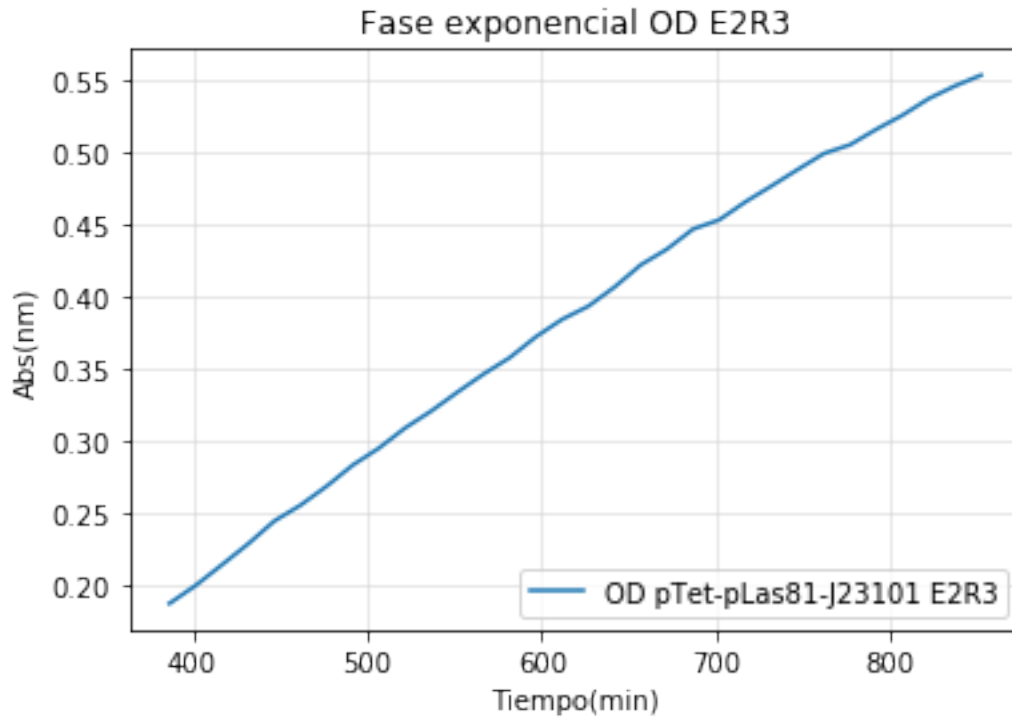


A=2.107622e+00
um=3.942368e-03

```
l=1.889461e+02  
Tm=3.856175e+02  
doubpe=1.758200e+02  
ext=4.395501e+02  
Tfinal=8.251675e+02
```

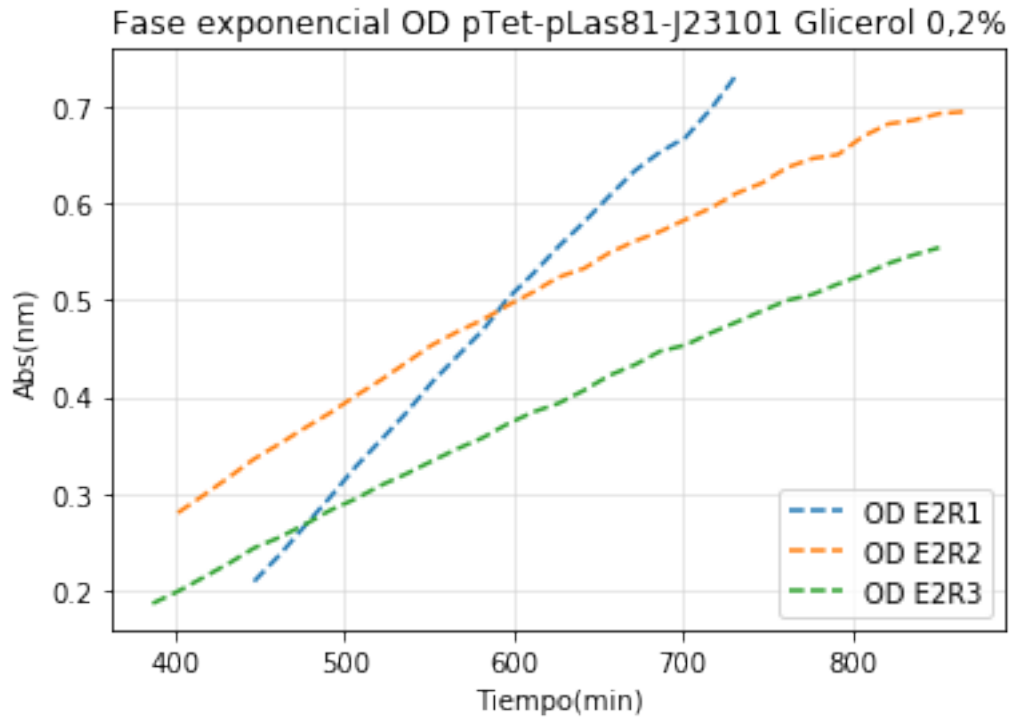
Out[48]: <matplotlib.legend.Legend at 0x267ee7adf28>





```
In [49]: #Fase exponencial OD/tiempo
plt.figure()
plt.title('Fase exponencial OD pTet-pLas81-J23101 Glicerol 0,2%')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[29:49],od1527g1[29:49], '--',label='OD E2R1')
plt.plot(tt[26:58],od1527g2[26:58], '--',label='OD E2R2')
plt.plot(tt[25:57],od1527g3[25:57], '--',label='OD E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')
```

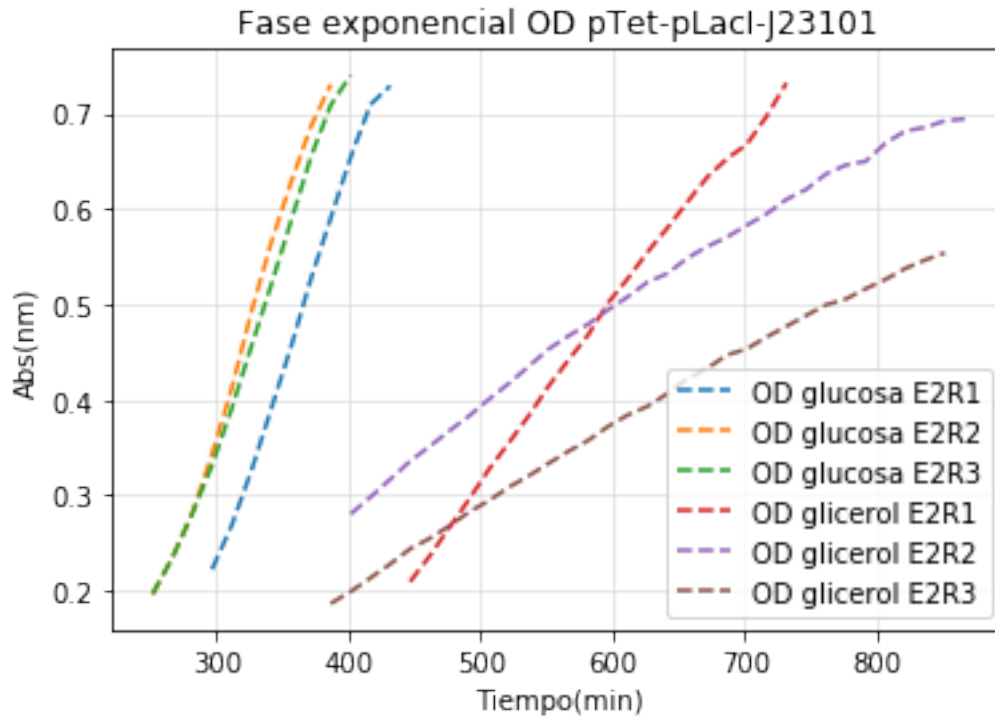
```
Out[49]: <matplotlib.legend.Legend at 0x267ecc09ba8>
```



In [50]: *#Fase exponencial OD/tiempo*

```
plt.figure()
plt.title('Fase exponencial OD pTet-pLacI-J23101')
plt.xlabel('Tiempo(min)')
plt.ylabel('Abs(nm)')
plt.plot(tt[19:29],od15271[19:29], '--',label='OD glucosa E2R1')
plt.plot(tt[16:26],od15272[16:26], '--',label='OD glucosa E2R2')
plt.plot(tt[16:27],od15273[16:27], '--',label='OD glucosa E2R3')
plt.plot(tt[29:49],od1527g1[29:49], '--',label='OD glicerol E2R1')
plt.plot(tt[26:58],od1527g2[26:58], '--',label='OD glicerol E2R2')
plt.plot(tt[25:57],od1527g3[25:57], '--',label='OD glicerol E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5)
plt.legend(loc='lower right')
```

Out[50]: <matplotlib.legend.Legend at 0x267eccdcf98>



```
In [51]: #Selección de datos en arrays, según lo determinado
#controles glucosa
o1=odcg1[19:30]
c1=cfpcg1[19:30]
r1=rfpcg1[19:30]
y1=yfpcg1[19:30]

o2=odcg2[19:30]
c2=cfpcg2[19:30]
r2=rfpcg2[19:30]
y2=yfpcg2[19:30]

o3=odcg3[15:27]
c3=cfpcg3[15:27]
r3=rfpcg3[15:27]
y3=yfpcg3[15:27]

#controles glicerol
o4=odcg11[24:45]
c4=cfpcg11[24:45]
r4=rfpcg11[24:45]
y4=yfpcg11[24:45]

o5=odcg12[23:46]
```



```
c5=cfpcg12[23:46]
r5=rfpcg12[23:46]
y5=yfpcg12[23:46]
```

```
o6=odcgl3[25:57]
c6=cfpcgl3[25:57]
r6=rfpcgl3[25:57]
y6=yfpcgl3[25:57]
```

```
#ptet-std-std glucosa
```

```
o7=od15211[19:30]
c7=cfp15211[19:30]
r7=rfp15211[19:30]
y7=yfp15211[19:30]
```

```
o8=od15212[19:32]
c8=cfp15212[19:32]
r8=rfp15212[19:32]
y8=yfp15212[19:32]
```

```
o9=od15213[16:28]
c9=cfp15213[16:28]
r9=rfp15213[16:28]
y9=yfp15213[16:28]
```

```
#ptet-std-std glycerol
```

```
o10=od1521g1[27:48]
c10=cfp1521g1[27:48]
r10=rfp1521g1[27:48]
y10=yfp1521g1[27:48]
```

```
o11=od1521g2[28:53]
c11=cfp1521g2[28:53]
r11=rfp1521g2[28:53]
y11=yfp1521g2[28:53]
```

```
o12=od1521g3[27:53]
c12=cfp1521g3[27:53]
r12=rfp1521g3[27:53]
y12=yfp1521g3[27:53]
```

```
#ptet-107-std glucosa
```

```
o13=od15231[21:32]
c13=cfp15231[21:32]
r13=rfp15231[21:32]
y13=yfp15231[21:32]
```

```
o14=od15232[17:27]
```

c14=cfp15232[17:27]
r14=rfp15232[17:27]
y14=yfp15232[17:27]

o15=od15233[17:29]
c15=cfp15233[17:29]
r15=rfp15233[17:29]
y15=yfp15233[17:29]

#ptet-107-std glycerol

o16=od1523g1[30:49]
c16=cfp1523g1[30:49]
r16=rfp1523g1[30:49]
y16=yfp1523g1[30:49]

o17=od1523g2[27:54]
c17=cfp1523g2[27:54]
r17=rfp1523g2[27:54]
y17=yfp1523g2[27:54]

o18=od1523g3[28:60]
c18=cfp1523g3[28:60]
r18=rfp1523g3[28:60]
y18=yfp1523g3[28:60]

#ptet-pLac-std glucosa

o19=od15261[19:29]
c19=cfp15261[19:29]
r19=rfp15261[19:29]
y19=yfp15261[19:29]

o20=od15262[18:29]
c20=cfp15262[18:29]
r20=rfp15262[18:29]
y20=yfp15262[18:29]

o21=od15263[16:28]
c21=cfp15263[16:28]
r21=rfp15263[16:28]
y21=yfp15263[16:28]

#ptet-pLac-std glycerol

o22=od1526g1[27:47]
c22=cfp1526g1[27:47]
r22=rfp1526g1[27:47]
y22=yfp1526g1[27:47]

o23=od1526g2[24:49]

```
c23=cfp1526g2[24:49]
r23=rfp1526g2[24:49]
y23=yfp1526g2[24:49]
```

```
o24=od1526g3[27:59]
c24=cfp1526g3[27:59]
r24=rfp1526g3[27:59]
y24=yfp1526g3[27:59]
```

```
#ptet-pLas-std glucosa
```

```
o25=od15271[19:29]
c25=cfp15271[19:29]
r25=rfp15271[19:29]
y25=yfp15271[19:29]
```

```
o26=od15272[16:26]
c26=cfp15272[16:26]
r26=rfp15272[16:26]
y26=yfp15272[16:26]
```

```
o27=od15273[16:26]
c27=cfp15273[16:26]
r27=rfp15273[16:26]
y27=yfp15273[16:26]
```

```
#ptet-pLas-std glicerol
```

```
o28=od1527g1[29:49]
c28=cfp1527g1[29:49]
r28=rfp1527g1[29:49]
y28=yfp1527g1[29:49]
```

```
o29=od1527g2[26:58]
c29=cfp1527g2[26:58]
r29=rfp1527g2[26:58]
y29=yfp1527g2[26:58]
```

```
o30=od1527g3[25:57]
c30=cfp1527g3[25:57]
r30=rfp1527g3[25:57]
y30=yfp1527g3[25:57]
```

```
In [52]: #regresion lineal de replicas
```

```
#Controles glucosa
```

```
slope, intercept, r_value, p_value, std_err = stats.linregress(o1, c1)
slopec1 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o1, r1)
sloper1 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o1, y1)
```

```

slopey1=slope

slope, intercept, r_value, p_value,std_err=stats.linregress(o2,c2)
slopec2=slope
slope, intercept, r_value, p_value,std_err=stats.linregress(o2,r2)
sloper2=slope
slope, intercept, r_value, p_value,std_err=stats.linregress(o2,y2)
slopey2=slope

slope, intercept, r_value, p_value,std_err=stats.linregress(o3,c3)
slopec3=slope
slope, intercept, r_value, p_value,std_err=stats.linregress(o3,r3)
sloper3=slope
slope, intercept, r_value, p_value,std_err=stats.linregress(o3,y3)
slopey3=slope

#Controles glicerol
slope, intercept, r_value, p_value,std_err=stats.linregress(o4,c4)
slopec4=slope
slope, intercept, r_value, p_value,std_err=stats.linregress(o4,r4)
sloper4=slope
slope, intercept, r_value, p_value,std_err=stats.linregress(o4,y4)
slopey4=slope

slope, intercept, r_value, p_value,std_err=stats.linregress(o5,c5)
slopec5=slope
slope, intercept, r_value, p_value,std_err=stats.linregress(o5,r5)
sloper5=slope
slope, intercept, r_value, p_value,std_err=stats.linregress(o5,y5)
slopey5=slope

slope, intercept, r_value, p_value,std_err=stats.linregress(o6,c6)
slopec6=slope
slope, intercept, r_value, p_value,std_err=stats.linregress(o6,r6)
sloper6=slope
slope, intercept, r_value, p_value,std_err=stats.linregress(o6,y6)
slopey6=slope

#ptet-std-std glucosa
slope, intercept, r_value, p_value,std_err=stats.linregress(o7,c7)
slopec7=slope
slope, intercept, r_value, p_value,std_err=stats.linregress(o7,r7)
sloper7=slope
slope, intercept, r_value, p_value,std_err=stats.linregress(o7,y7)
slopey7=slope

slope, intercept, r_value, p_value,std_err=stats.linregress(o8,c8)
slopec8=slope

```

```

slope, intercept, r_value, p_value, std_err=stats.linregress(o8,r8)
sloper8=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o8,y8)
slopey8=slope

slope, intercept, r_value, p_value, std_err=stats.linregress(o9,c9)
slopec9=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o9,r9)
sloper9=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o9,y9)
slopey9=slope

#ptet-std-std glicerol
slope, intercept, r_value, p_value, std_err=stats.linregress(o10,c10)
slopec10=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o10,r10)
sloper10=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o10,y10)
slopey10=slope

slope, intercept, r_value, p_value, std_err=stats.linregress(o11,c11)
slopec11=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o11,r11)
sloper11=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o11,y11)
slopey11=slope

slope, intercept, r_value, p_value, std_err=stats.linregress(o12,c12)
slopec12=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o12,r12)
sloper12=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o12,y12)
slopey12=slope

#ptet-107-std glucosa
slope, intercept, r_value, p_value, std_err=stats.linregress(o13,c13)
slopec13=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o13,r13)
sloper13=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o13,y13)
slopey13=slope

slope, intercept, r_value, p_value, std_err=stats.linregress(o14,c14)
slopec14=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o14,r14)
sloper14=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o14,y14)
slopey14=slope

```

```

slope, intercept, r_value, p_value, std_err = stats.linregress(o15, c15)
slopec15 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o15, r15)
sloper15 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o15, y15)
slopepy15 = slope

```

#ptet-107-std glycerol

```

slope, intercept, r_value, p_value, std_err = stats.linregress(o16, c16)
slopec16 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o16, r16)
sloper16 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o16, y16)
slopepy16 = slope

```

```

slope, intercept, r_value, p_value, std_err = stats.linregress(o17, c17)
slopec17 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o17, r17)
sloper17 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o17, y17)
slopepy17 = slope

```

```

slope, intercept, r_value, p_value, std_err = stats.linregress(o18, c18)
slopec18 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o18, r18)
sloper18 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o18, y18)
slopepy18 = slope

```

#ptet-pLacI-std glucosa

```

slope, intercept, r_value, p_value, std_err = stats.linregress(o19, c19)
slopec19 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o19, r19)
sloper19 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o19, y19)
slopepy19 = slope

```

```

slope, intercept, r_value, p_value, std_err = stats.linregress(o20, c20)
slopec20 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o20, r20)
sloper20 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o20, y20)
slopepy20 = slope

```

```

slope, intercept, r_value, p_value, std_err = stats.linregress(o21, c21)
slopec21 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o21, r21)

```

```

sloper21=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o21,y21)
slopey21=slope

#ptet-LacI-std glycerol
slope, intercept, r_value, p_value, std_err=stats.linregress(o22,c22)
slopec22=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o22,r22)
sloper22=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o22,y22)
slopey22=slope

slope, intercept, r_value, p_value, std_err=stats.linregress(o23,c23)
slopec23=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o23,r23)
sloper23=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o23,y23)
slopey23=slope

slope, intercept, r_value, p_value, std_err=stats.linregress(o24,c24)
slopec24=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o24,r24)
sloper24=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o24,y24)
slopey24=slope

#ptet-pLas81-std glucosa
slope, intercept, r_value, p_value, std_err=stats.linregress(o25,c25)
slopec25=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o25,r25)
sloper25=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o25,y25)
slopey25=slope

slope, intercept, r_value, p_value, std_err=stats.linregress(o26,c26)
slopec26=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o26,r26)
sloper26=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o26,y26)
slopey26=slope

slope, intercept, r_value, p_value, std_err=stats.linregress(o27,c27)
slopec27=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o27,r27)
sloper27=slope
slope, intercept, r_value, p_value, std_err=stats.linregress(o27,y27)
slopey27=slope

```

```

#ptet-pLas81-std glicerol
slope, intercept, r_value, p_value, std_err = stats.linregress(o28, c28)
slopec28 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o28, r28)
sloper28 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o28, y28)
slopepy28 = slope

slope, intercept, r_value, p_value, std_err = stats.linregress(o29, c29)
slopec29 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o29, r29)
sloper29 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o29, y29)
slopepy29 = slope

slope, intercept, r_value, p_value, std_err = stats.linregress(o30, c30)
slopec30 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o30, r30)
sloper30 = slope
slope, intercept, r_value, p_value, std_err = stats.linregress(o30, y30)
slopepy30 = slope

```

```

In [53]: pendientesc = [slopec1, slopec2, slopec3, slopec4, slopec5, slopec6, slopec7, slopec8, slopec9, slopec10]
pendientesr = [sloper1, sloper2, sloper3, sloper4, sloper5, sloper6, sloper7, sloper8, sloper9, sloper10]
pendientesy = [slopepy1, slopepy2, slopepy3, slopepy4, slopepy5, slopepy6, slopepy7, slopepy8, slopepy9, slopepy10]

```

```

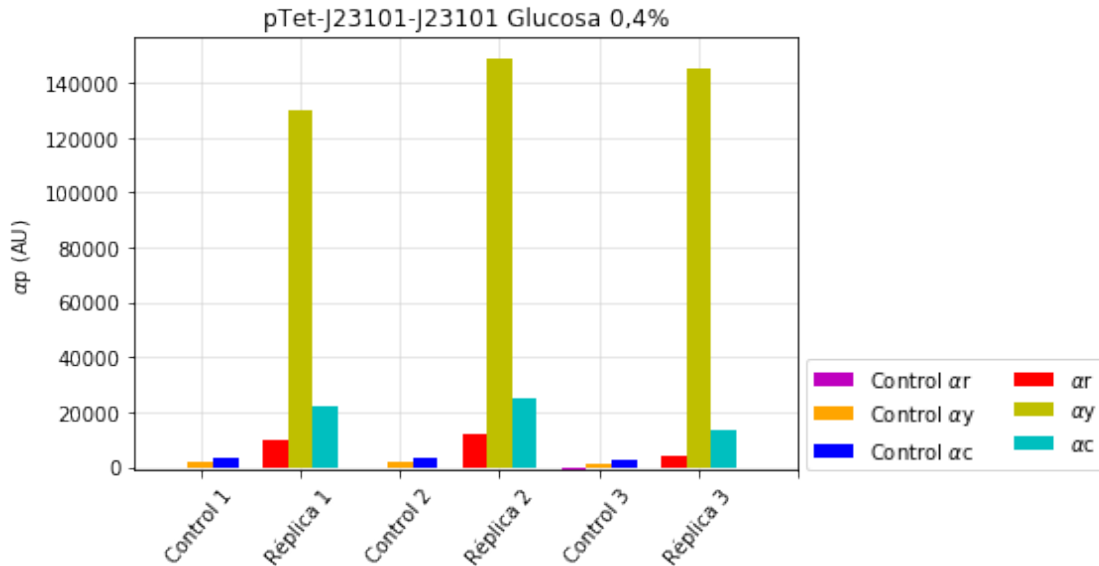
#Grafico pendientes ptet-std-std Glucosa
X = np.arange(7)
plt.figure()
plt.title('pTet-J23101-J23101 Glucosa 0,4%')
plt.ylabel(r'$\alpha$P (AU)')
plt.bar(X[0]-0.25, pendientesr[0], color='m', width=0.25, label='Control' + ' ' + r'$\alpha$r')
plt.bar(X[0]+0.00, pendientesy[0], color='orange', width=0.25, label='Control' + ' ' + r'$\alpha$y')
plt.bar(X[0]+0.25, pendientesc[0], color='b', width=0.25, label='Control' + ' ' + r'$\alpha$c')
plt.bar(X[1]-0.25, pendientesr[6], color='r', width=0.25, label=r'$\alpha$r', zorder=3)
plt.bar(X[1]+0.00, pendientesy[6], color='y', width=0.25, label=r'$\alpha$y', zorder=3)
plt.bar(X[1]+0.25, pendientesc[6], color='c', width=0.25, label=r'$\alpha$c', zorder=3)
plt.bar(X[2]-0.25, pendientesr[1], color='m', width=0.25, zorder=3)
plt.bar(X[2]+0.00, pendientesy[1], color='orange', width=0.25, zorder=3)
plt.bar(X[2]+0.25, pendientesc[1], color='b', width=0.25, zorder=3)
plt.bar(X[3]-0.25, pendientesr[7], color='r', width=0.25, zorder=3)
plt.bar(X[3]+0.00, pendientesy[7], color='y', width=0.25, zorder=3)
plt.bar(X[3]+0.25, pendientesc[7], color='c', width=0.25, zorder=3)
plt.bar(X[4]-0.25, pendientesr[2], color='m', width=0.25, zorder=3)
plt.bar(X[4]+0.00, pendientesy[2], color='orange', width=0.25, zorder=3)
plt.bar(X[4]+0.25, pendientesc[2], color='b', width=0.25, zorder=3)
plt.bar(X[5]-0.25, pendientesr[8], color='r', width=0.25, zorder=3)
plt.bar(X[5]+0.00, pendientesy[8], color='y', width=0.25, zorder=3)

```



```
plt.bar(X[5]+0.25,pendientesc[8],color='c',width=0.25,zorder=3)
plt.xticks(X, ['Control 1',"Réplica 1",'Control 2',"Réplica 2",'Control 3',"Réplica 3"])
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)
```

Out[53]: <matplotlib.legend.Legend at 0x267ed5157b8>



```
In [54]: #Grafico pendientes ptet-std-std Glicerol
X = np.arange(7)
plt.figure()
plt.title('pTet-J23101-J23101 Glicerol 0,2%')
plt.ylabel(r'$\alpha$ p (AU)')
plt.bar(X[0]-0.25,pendientesr[3],color='m',width=0.25,label='Control'+ ' '+ r'$\alpha$r',zorder=3)
plt.bar(X[0]+0.00,pendientesy[3],color='orange',width=0.25,label='Control'+ ' '+ r'$\alpha$y',zorder=3)
plt.bar(X[0]+0.25,pendientesc[3],color='b',width=0.25,label='Control'+ ' '+ r'$\alpha$c',zorder=3)
plt.bar(X[1]-0.25,pendientesr[9],color='r',width=0.25,label=r'$\alpha$r',zorder=3)
plt.bar(X[1]+0.00,pendientesy[9],color='y',width=0.25,label=r'$\alpha$y',zorder=3)
plt.bar(X[1]+0.25,pendientesc[9],color='c',width=0.25,label=r'$\alpha$c',zorder=3)
plt.bar(X[2]-0.25,pendientesr[4],color='m',width=0.25,zorder=3)
plt.bar(X[2]+0.00,pendientesy[4],color='orange',width=0.25,zorder=3)
plt.bar(X[2]+0.25,pendientesc[4],color='b',width=0.25,zorder=3)
plt.bar(X[3]-0.25,pendientesr[10],color='r',width=0.25,zorder=3)
plt.bar(X[3]+0.00,pendientesy[10],color='y',width=0.25,zorder=3)
plt.bar(X[3]+0.25,pendientesc[10],color='c',width=0.25,zorder=3)
plt.bar(X[4]-0.25,pendientesr[5],color='m',width=0.25,zorder=3)
plt.bar(X[4]+0.00,pendientesy[5],color='orange',width=0.25,zorder=3)
plt.bar(X[4]+0.25,pendientesc[5],color='b',width=0.25,zorder=3)
plt.bar(X[5]-0.25,pendientesr[11],color='r',width=0.25,zorder=3)
```

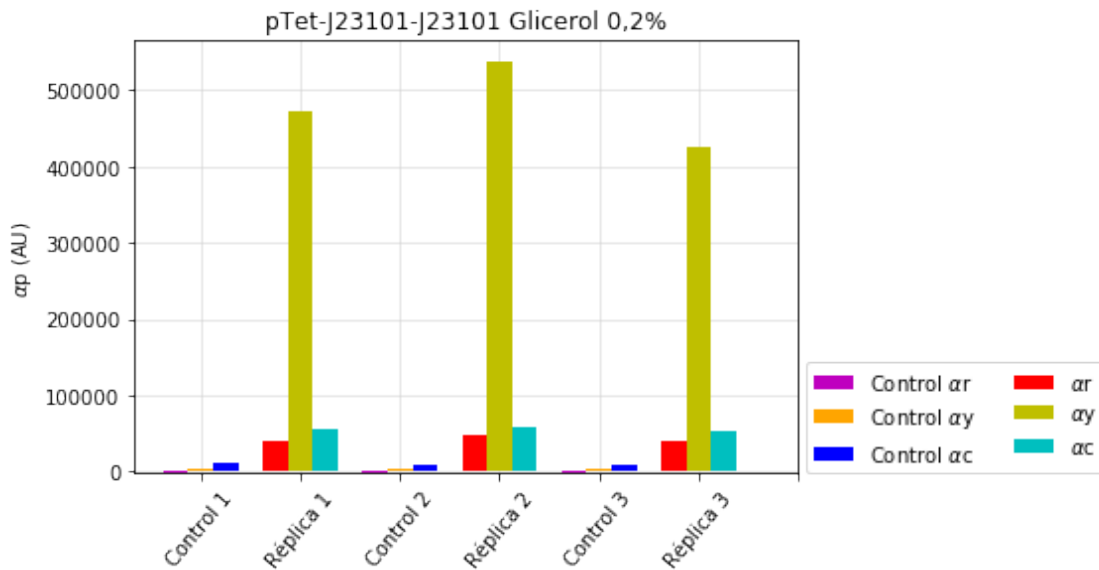
```

plt.bar(X[5]+0.00,pendientesy[11],color='y',width=0.25,zorder=3)
plt.bar(X[5]+0.25,pendientesy[11],color='c',width=0.25,zorder=3)
plt.xticks(X, ['Control 1',"Réplica 1",'Control 2',"Réplica 2",'Control 3',"Réplica 3"])

plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

```

Out[54]: <matplotlib.legend.Legend at 0x267eecf5ac8>



```

In [55]: #Grafico pendientes ptet-std-std
X = np.arange(13)
plt.figure(figsize=(10,7))
plt.title('pTet-J23101-J23101')
plt.ylabel(r'$\alpha_p$ (AU)')
plt.bar(X[0]-0.25,pendientesr[0],color='m',width=0.25,label='Control'+ ' '+ r'$\alpha$r',zorder=3)
plt.bar(X[0]+0.00,pendientesy[0],color='orange',width=0.25,label='Control'+ ' '+ r'$\alpha$y',zorder=3)
plt.bar(X[0]+0.25,pendientesy[0],color='b',width=0.25,label='Control'+ ' '+ r'$\alpha$c',zorder=3)
plt.bar(X[1]-0.25,pendientesr[6],color='r',width=0.25,label=r'$\alpha$r',zorder=3)
plt.bar(X[1]+0.00,pendientesy[6],color='y',width=0.25,label= r'$\alpha$y',zorder=3)
plt.bar(X[1]+0.25,pendientesy[6],color='c',width=0.25,label=r'$\alpha$c',zorder=3)
plt.bar(X[2]-0.25,pendientesr[3],color='m',width=0.25,zorder=3)
plt.bar(X[2]+0.00,pendientesy[3],color='orange',width=0.25,zorder=3)
plt.bar(X[2]+0.25,pendientesy[3],color='b',width=0.25,zorder=3)
plt.bar(X[3]-0.25,pendientesr[9],color='r',width=0.25,zorder=3)
plt.bar(X[3]+0.00,pendientesy[9],color='y',width=0.25,zorder=3)
plt.bar(X[3]+0.25,pendientesy[9],color='c',width=0.25,zorder=3)
plt.bar(X[4]-0.25,pendientesr[1],color='m',width=0.25,zorder=3)
plt.bar(X[4]+0.00,pendientesy[1],color='orange',width=0.25,zorder=3)

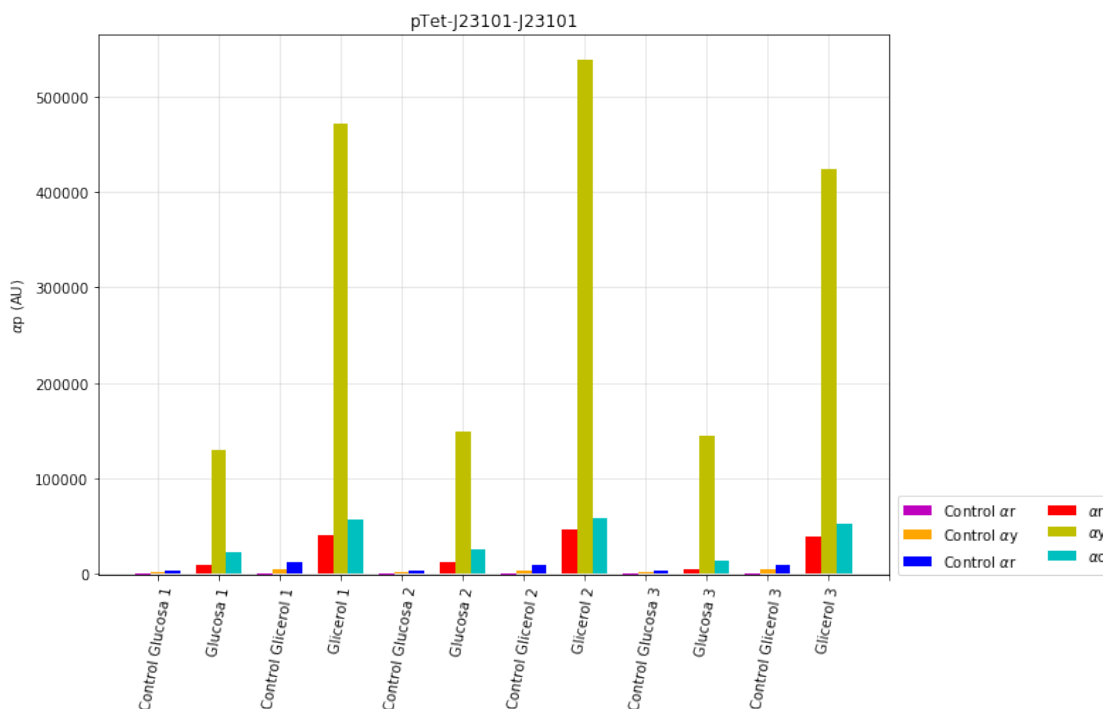
```

```

plt.bar(X[4]+0.25,pendientesc[1],color='b',width=0.25,zorder=3)
plt.bar(X[5]-0.25,pendientesr[7],color='r',width=0.25,zorder=3)
plt.bar(X[5]+0.00,pendientesy[7],color='y',width=0.25,zorder=3)
plt.bar(X[5]+0.25,pendientesc[7],color='c',width=0.25,zorder=3)
plt.bar(X[6]-0.25,pendientesr[4],color='m',width=0.25,zorder=3)
plt.bar(X[6]+0.00,pendientesy[4],color='orange',width=0.25,zorder=3)
plt.bar(X[6]+0.25,pendientesc[4],color='b',width=0.25,zorder=3)
plt.bar(X[7]-0.25,pendientesr[10],color='r',width=0.25,zorder=3)
plt.bar(X[7]+0.00,pendientesy[10],color='y',width=0.25,zorder=3)
plt.bar(X[7]+0.25,pendientesc[10],color='c',width=0.25,zorder=3)
plt.bar(X[8]-0.25,pendientesr[2],color='m',width=0.25,zorder=3)
plt.bar(X[8]+0.00,pendientesy[2],color='orange',width=0.25,zorder=3)
plt.bar(X[8]+0.25,pendientesc[2],color='b',width=0.25,zorder=3)
plt.bar(X[9]-0.25,pendientesr[8],color='r',width=0.25,zorder=3)
plt.bar(X[9]+0.00,pendientesy[8],color='y',width=0.25,zorder=3)
plt.bar(X[9]+0.25,pendientesc[8],color='c',width=0.25,zorder=3)
plt.bar(X[10]-0.25,pendientesr[5],color='m',width=0.25,zorder=3)
plt.bar(X[10]+0.00,pendientesy[5],color='orange',width=0.25,zorder=3)
plt.bar(X[10]+0.25,pendientesc[5],color='b',width=0.25,zorder=3)
plt.bar(X[11]-0.25,pendientesr[11],color='r',width=0.25,zorder=3)
plt.bar(X[11]+0.00,pendientesy[11],color='y',width=0.25,zorder=3)
plt.bar(X[11]+0.25,pendientesc[11],color='c',width=0.25,zorder=3)
plt.xticks(X, ['Control Glucosa 1',"Glucosa 1",'Control Glicerol 1',"Glicerol 1",'Contr
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

```

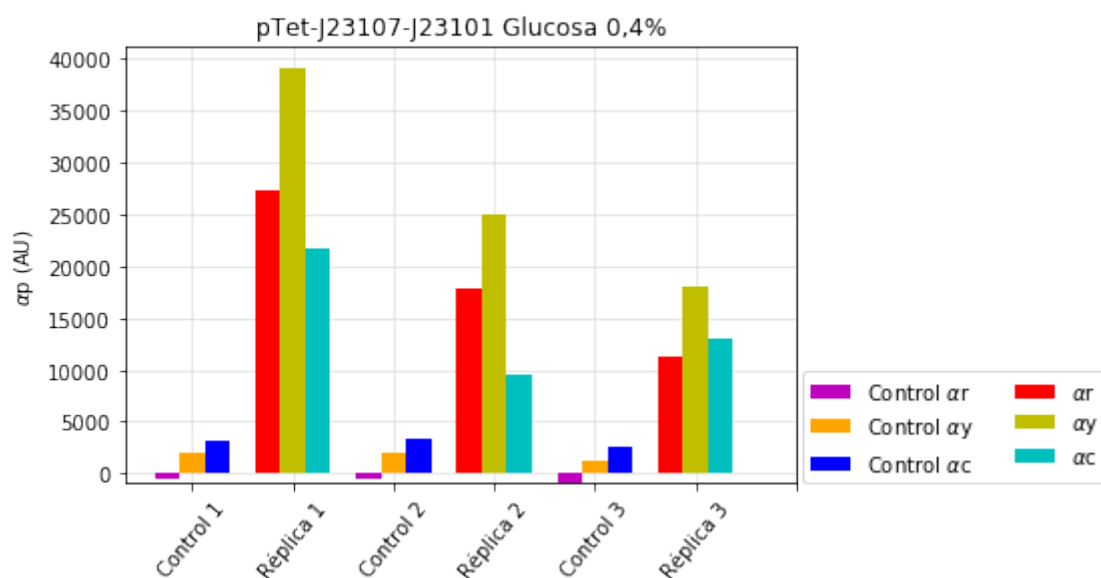
Out[55]: <matplotlib.legend.Legend at 0x267eeea2940>



In [56]: *#Grafico pendientes ptet-107-std Glucosa*

```
X = np.arange(7)
plt.figure()
plt.title('pTet-J23107-J23101 Glucosa 0,4%')
plt.ylabel(r'$\alpha$ (AU)')
plt.bar(X[0]-0.25,pendientesr[0],color='m',width=0.25,label='Control'+ ' ' + r'$\alpha$r')
plt.bar(X[0]+0.00,pendientesy[0],color='orange',width=0.25,label='Control'+ ' ' + r'$\alpha$y')
plt.bar(X[0]+0.25,pendientesr[0],color='b',width=0.25,label='Control'+ ' ' + r'$\alpha$c')
plt.bar(X[1]-0.25,pendientesr[12],color='r',width=0.25,label=r'$\alpha$r',zorder=3)
plt.bar(X[1]+0.00,pendientesy[12],color='y',width=0.25,label=r'$\alpha$y',zorder=3)
plt.bar(X[1]+0.25,pendientesr[12],color='c',width=0.25,label=r'$\alpha$c',zorder=3)
plt.bar(X[2]-0.25,pendientesr[1],color='m',width=0.25,zorder=3)
plt.bar(X[2]+0.00,pendientesy[1],color='orange',width=0.25,zorder=3)
plt.bar(X[2]+0.25,pendientesr[1],color='b',width=0.25,zorder=3)
plt.bar(X[3]-0.25,pendientesr[13],color='r',width=0.25,zorder=3)
plt.bar(X[3]+0.00,pendientesy[13],color='y',width=0.25,zorder=3)
plt.bar(X[3]+0.25,pendientesr[13],color='c',width=0.25,zorder=3)
plt.bar(X[4]-0.25,pendientesr[2],color='m',width=0.25,zorder=3)
plt.bar(X[4]+0.00,pendientesy[2],color='orange',width=0.25,zorder=3)
plt.bar(X[4]+0.25,pendientesr[2],color='b',width=0.25,zorder=3)
plt.bar(X[5]-0.25,pendientesr[14],color='r',width=0.25,zorder=3)
plt.bar(X[5]+0.00,pendientesy[14],color='y',width=0.25,zorder=3)
plt.bar(X[5]+0.25,pendientesr[14],color='c',width=0.25,zorder=3)
plt.xticks(X, ['Control 1',"Réplica 1",'Control 2',"Réplica 2",'Control 3',"Réplica 3"])
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)
```

Out [56]: <matplotlib.legend.Legend at 0x267ee784e80>

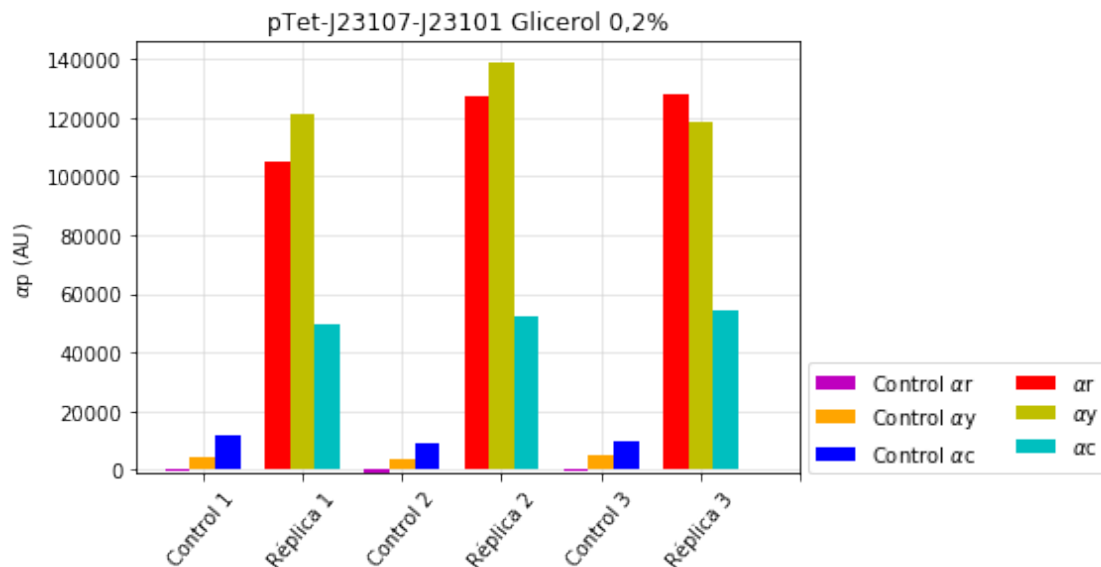


```

In [57]: #Grafico pendientes ptet-107-std Glicerol
X = np.arange(7)
plt.figure()
plt.title('pTet-J23107-J23101 Glicerol 0,2%')
plt.ylabel(r'$\alpha$ (AU)')
plt.bar(X[0]-0.25,pendientesr[3],color='m',width=0.25,label='Control'+ ' '+ r'$\alpha$r')
plt.bar(X[0]+0.00,pendientesy[3],color='orange',width=0.25,label='Control'+ ' '+ r'$\alpha$y')
plt.bar(X[0]+0.25,pendientesc[3],color='b',width=0.25,label='Control'+ ' '+ r'$\alpha$c')
plt.bar(X[1]-0.25,pendientesr[15],color='r',width=0.25,label=r'$\alpha$r',zorder=3)
plt.bar(X[1]+0.00,pendientesy[15],color='y',width=0.25,label=r'$\alpha$y',zorder=3)
plt.bar(X[1]+0.25,pendientesc[15],color='c',width=0.25,label=r'$\alpha$c',zorder=3)
plt.bar(X[2]-0.25,pendientesr[4],color='m',width=0.25,zorder=3)
plt.bar(X[2]+0.00,pendientesy[4],color='orange',width=0.25,zorder=3)
plt.bar(X[2]+0.25,pendientesc[4],color='b',width=0.25,zorder=3)
plt.bar(X[3]-0.25,pendientesr[16],color='r',width=0.25,zorder=3)
plt.bar(X[3]+0.00,pendientesy[16],color='y',width=0.25,zorder=3)
plt.bar(X[3]+0.25,pendientesc[16],color='c',width=0.25,zorder=3)
plt.bar(X[4]-0.25,pendientesr[5],color='m',width=0.25,zorder=3)
plt.bar(X[4]+0.00,pendientesy[5],color='orange',width=0.25,zorder=3)
plt.bar(X[4]+0.25,pendientesc[5],color='b',width=0.25,zorder=3)
plt.bar(X[5]-0.25,pendientesr[17],color='r',width=0.25,zorder=3)
plt.bar(X[5]+0.00,pendientesy[17],color='y',width=0.25,zorder=3)
plt.bar(X[5]+0.25,pendientesc[17],color='c',width=0.25,zorder=3)
plt.xticks(X, ['Control 1',"Réplica 1",'Control 2',"Réplica 2",'Control 3',"Réplica 3"])
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

```

Out[57]: <matplotlib.legend.Legend at 0x267ef281470>

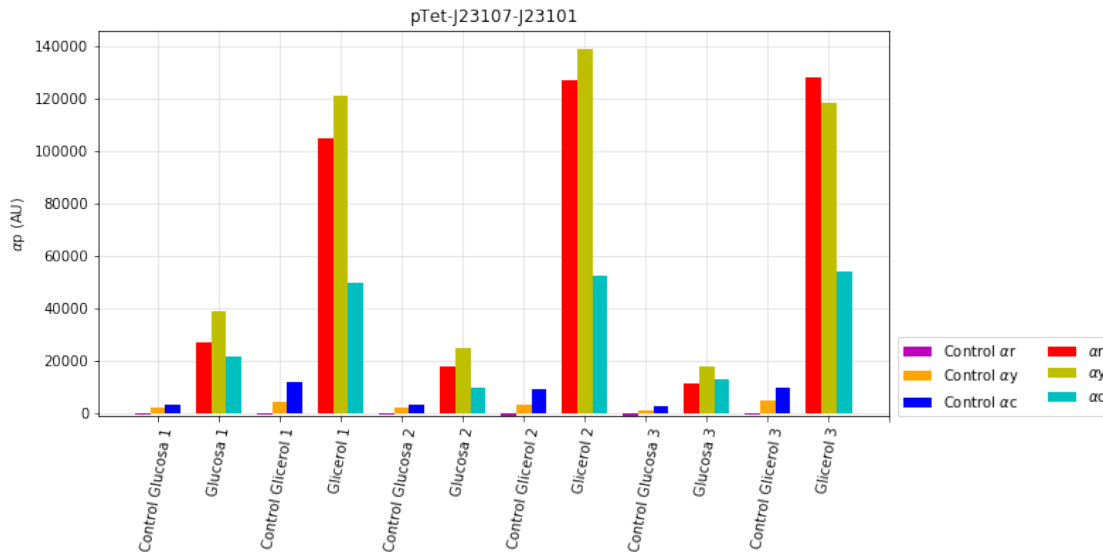


```

In [58]: #Grafico pendientes ptet-std-std
X = np.arange(13)
plt.figure(figsize=(10,5))
plt.title('pTet-J23107-J23101')
plt.ylabel(r'$\alpha$p (AU)')
plt.bar(X[0]-0.25,pendientesr[0],color='m',width=0.25,label='Control'+ ' '+ r'$\alpha$r')
plt.bar(X[0]+0.00,pendientesy[0],color='orange',width=0.25,label='Control'+ ' '+ r'$\alpha$y')
plt.bar(X[0]+0.25,pendientesr[0],color='b',width=0.25,label='Control'+ ' '+ r'$\alpha$c')
plt.bar(X[1]-0.25,pendientesr[12],color='r',width=0.25,label=r'$\alpha$r',zorder=3)
plt.bar(X[1]+0.00,pendientesy[12],color='y',width=0.25,label=r'$\alpha$y',zorder=3)
plt.bar(X[1]+0.25,pendientesr[12],color='c',width=0.25,label=r'$\alpha$c',zorder=3)
plt.bar(X[2]-0.25,pendientesr[3],color='m',width=0.25,zorder=3)
plt.bar(X[2]+0.00,pendientesy[3],color='orange',width=0.25,zorder=3)
plt.bar(X[2]+0.25,pendientesr[3],color='b',width=0.25,zorder=3)
plt.bar(X[3]-0.25,pendientesr[15],color='r',width=0.25,zorder=3)
plt.bar(X[3]+0.00,pendientesy[15],color='y',width=0.25,zorder=3)
plt.bar(X[3]+0.25,pendientesr[15],color='c',width=0.25,zorder=3)
plt.bar(X[4]-0.25,pendientesr[1],color='m',width=0.25,zorder=3)
plt.bar(X[4]+0.00,pendientesy[1],color='orange',width=0.25,zorder=3)
plt.bar(X[4]+0.25,pendientesr[1],color='b',width=0.25,zorder=3)
plt.bar(X[5]-0.25,pendientesr[13],color='r',width=0.25,zorder=3)
plt.bar(X[5]+0.00,pendientesy[13],color='y',width=0.25,zorder=3)
plt.bar(X[5]+0.25,pendientesr[13],color='c',width=0.25,zorder=3)
plt.bar(X[6]-0.25,pendientesr[4],color='m',width=0.25,zorder=3)
plt.bar(X[6]+0.00,pendientesy[4],color='orange',width=0.25,zorder=3)
plt.bar(X[6]+0.25,pendientesr[4],color='b',width=0.25,zorder=3)
plt.bar(X[7]-0.25,pendientesr[16],color='r',width=0.25,zorder=3)
plt.bar(X[7]+0.00,pendientesy[16],color='y',width=0.25,zorder=3)
plt.bar(X[7]+0.25,pendientesr[16],color='c',width=0.25,zorder=3)
plt.bar(X[8]-0.25,pendientesr[2],color='m',width=0.25,zorder=3)
plt.bar(X[8]+0.00,pendientesy[2],color='orange',width=0.25,zorder=3)
plt.bar(X[8]+0.25,pendientesr[2],color='b',width=0.25,zorder=3)
plt.bar(X[9]-0.25,pendientesr[14],color='r',width=0.25,zorder=3)
plt.bar(X[9]+0.00,pendientesy[14],color='y',width=0.25,zorder=3)
plt.bar(X[9]+0.25,pendientesr[14],color='c',width=0.25,zorder=3)
plt.bar(X[10]-0.25,pendientesr[5],color='m',width=0.25,zorder=3)
plt.bar(X[10]+0.00,pendientesy[5],color='orange',width=0.25,zorder=3)
plt.bar(X[10]+0.25,pendientesr[5],color='b',width=0.25,zorder=3)
plt.bar(X[11]-0.25,pendientesr[17],color='r',width=0.25,zorder=3)
plt.bar(X[11]+0.00,pendientesy[17],color='y',width=0.25,zorder=3)
plt.bar(X[11]+0.25,pendientesr[17],color='c',width=0.25,zorder=3)
plt.xticks(X, ['Control Glucosa 1',"Glucosa 1",'Control Glicerol 1',"Glicerol 1",'Contr
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

```

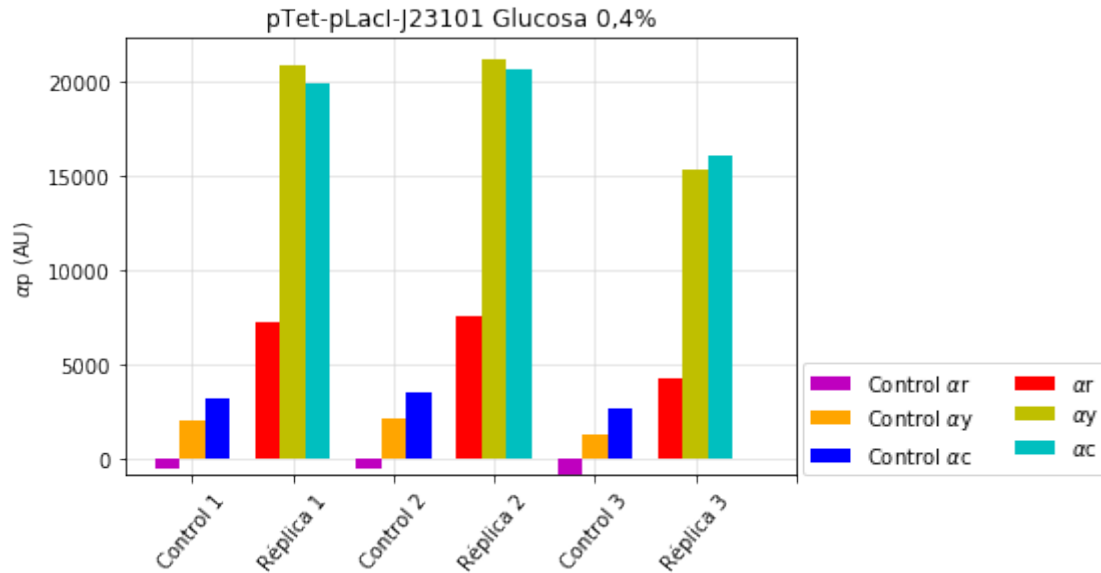
Out [58]: <matplotlib.legend.Legend at 0x267ef34a080>



In [59]: *#Grafico pendientes ptet-plac-std Glucosa*

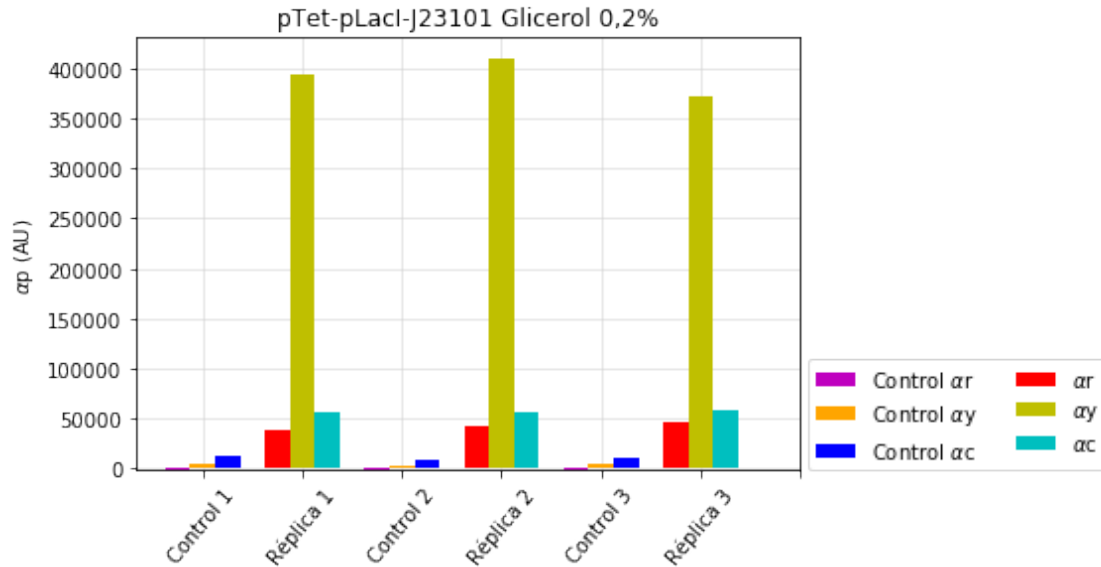
```
X = np.arange(7)
plt.figure()
plt.title('pTet-pLacI-J23101 Glucosa 0,4%')
plt.ylabel(r'$\alpha$ (AU)')
plt.bar(X[0]-0.25,pendientesr[0],color='m',width=0.25,label='Control'+ ' '+ r'$\alpha$r')
plt.bar(X[0]+0.00,pendientesy[0],color='orange',width=0.25,label='Control'+ ' '+ r'$\alpha$y')
plt.bar(X[0]+0.25,pendientesr[0],color='b',width=0.25,label='Control'+ ' '+ r'$\alpha$c')
plt.bar(X[1]-0.25,pendientesr[18],color='r',width=0.25,label=r'$\alpha$r',zorder=3)
plt.bar(X[1]+0.00,pendientesy[18],color='y',width=0.25,label=r'$\alpha$y',zorder=3)
plt.bar(X[1]+0.25,pendientesr[18],color='c',width=0.25,label=r'$\alpha$c',zorder=3)
plt.bar(X[2]-0.25,pendientesr[1],color='m',width=0.25,zorder=3)
plt.bar(X[2]+0.00,pendientesy[1],color='orange',width=0.25,zorder=3)
plt.bar(X[2]+0.25,pendientesr[1],color='b',width=0.25,zorder=3)
plt.bar(X[3]-0.25,pendientesr[19],color='r',width=0.25,zorder=3)
plt.bar(X[3]+0.00,pendientesy[19],color='y',width=0.25,zorder=3)
plt.bar(X[3]+0.25,pendientesr[19],color='c',width=0.25,zorder=3)
plt.bar(X[4]-0.25,pendientesr[2],color='m',width=0.25,zorder=3)
plt.bar(X[4]+0.00,pendientesy[2],color='orange',width=0.25,zorder=3)
plt.bar(X[4]+0.25,pendientesr[2],color='b',width=0.25,zorder=3)
plt.bar(X[5]-0.25,pendientesr[20],color='r',width=0.25,zorder=3)
plt.bar(X[5]+0.00,pendientesy[20],color='y',width=0.25,zorder=3)
plt.bar(X[5]+0.25,pendientesr[20],color='c',width=0.25,zorder=3)
plt.xticks(X, ['Control 1',"Réplica 1",'Control 2',"Réplica 2",'Control 3',"Réplica 3"])
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)
```

Out[59]: <matplotlib.legend.Legend at 0x267ef6bf780>



```
In [60]: #Grafico pendientes ptet-plac-std Glicerol
X = np.arange(7)
plt.figure()
plt.title('pTet-pLacI-J23101 Glicerol 0,2%')
plt.ylabel(r'$\alpha_p$ (AU)')
plt.bar(X[0]-0.25,pendientesr[3],color='m',width=0.25,label='Control'+ ' '+ r'$\alpha_r$')
plt.bar(X[0]+0.00,pendientesy[3],color='orange',width=0.25,label='Control'+ ' '+ r'$\alpha_y$')
plt.bar(X[0]+0.25,pendientesr[3],color='b',width=0.25,label='Control'+ ' '+ r'$\alpha_c$')
plt.bar(X[1]-0.25,pendientesr[21],color='r',width=0.25,label=r'$\alpha_r$',zorder=3)
plt.bar(X[1]+0.00,pendientesy[21],color='y',width=0.25,label=r'$\alpha_y$',zorder=3)
plt.bar(X[1]+0.25,pendientesr[21],color='c',width=0.25,label=r'$\alpha_c$',zorder=3)
plt.bar(X[2]-0.25,pendientesr[4],color='m',width=0.25,zorder=3)
plt.bar(X[2]+0.00,pendientesy[4],color='orange',width=0.25,zorder=3)
plt.bar(X[2]+0.25,pendientesr[4],color='b',width=0.25,zorder=3)
plt.bar(X[3]-0.25,pendientesr[22],color='r',width=0.25,zorder=3)
plt.bar(X[3]+0.00,pendientesy[22],color='y',width=0.25,zorder=3)
plt.bar(X[3]+0.25,pendientesr[22],color='c',width=0.25,zorder=3)
plt.bar(X[4]-0.25,pendientesr[5],color='m',width=0.25,zorder=3)
plt.bar(X[4]+0.00,pendientesy[5],color='orange',width=0.25,zorder=3)
plt.bar(X[4]+0.25,pendientesr[5],color='b',width=0.25,zorder=3)
plt.bar(X[5]-0.25,pendientesr[23],color='r',width=0.25,zorder=3)
plt.bar(X[5]+0.00,pendientesy[23],color='y',width=0.25,zorder=3)
plt.bar(X[5]+0.25,pendientesr[23],color='c',width=0.25,zorder=3)
plt.xticks(X, ['Control 1',"Réplica 1",'Control 2',"Réplica 2",'Control 3',"Réplica 3"])
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)
```

Out [60]: <matplotlib.legend.Legend at 0x267ef7e8f60>



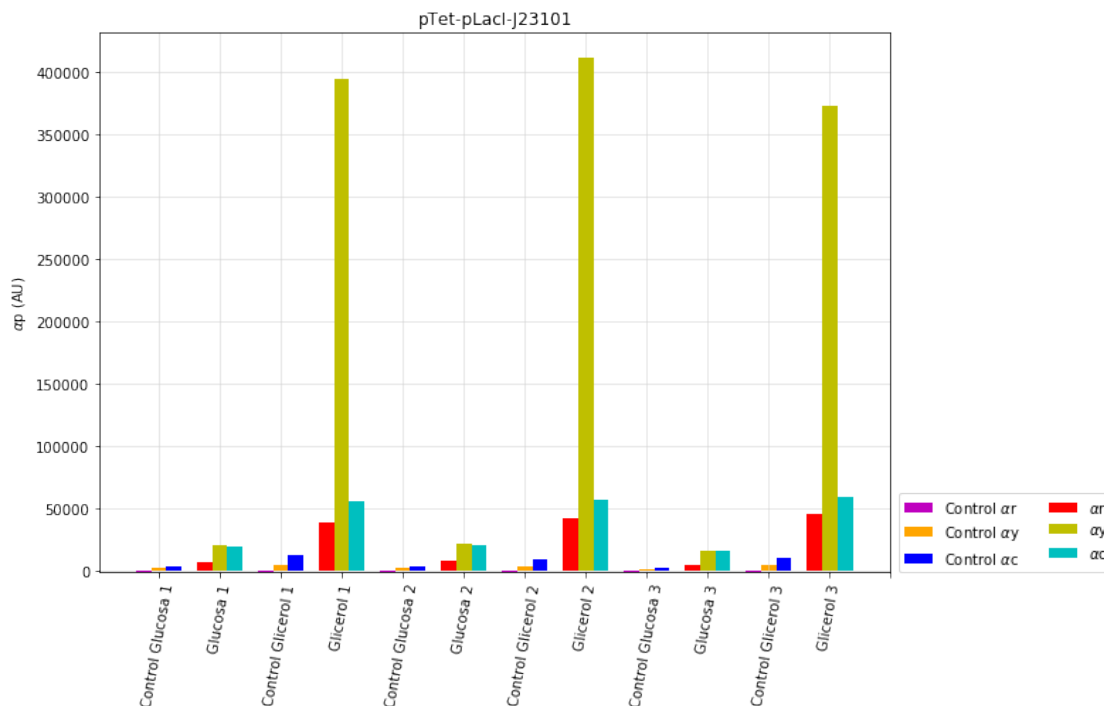
```
In [61]: #Grafico pendientes ptet-lac-std
X = np.arange(13)
plt.figure(figsize=(10,7))
plt.title('pTet-pLacI-J23101')
plt.ylabel(r'$\alpha_p$ (AU)')
plt.bar(X[0]-0.25,pendientesr[0],color='m',width=0.25,label='Control'+ ' '+ r'$\alpha$r')
plt.bar(X[0]+0.00,pendientesy[0],color='orange',width=0.25,label='Control'+ ' '+ r'$\alpha$y')
plt.bar(X[0]+0.25,pendientesr[0],color='b',width=0.25,label='Control'+ ' '+ r'$\alpha$c')
plt.bar(X[1]-0.25,pendientesr[18],color='r',width=0.25,label=r'$\alpha$r',zorder=3)
plt.bar(X[1]+0.00,pendientesy[18],color='y',width=0.25,label=r'$\alpha$y',zorder=3)
plt.bar(X[1]+0.25,pendientesr[18],color='c',width=0.25,label=r'$\alpha$c',zorder=3)
plt.bar(X[2]-0.25,pendientesr[3],color='m',width=0.25,zorder=3)
plt.bar(X[2]+0.00,pendientesy[3],color='orange',width=0.25,zorder=3)
plt.bar(X[2]+0.25,pendientesr[3],color='b',width=0.25,zorder=3)
plt.bar(X[3]-0.25,pendientesr[21],color='r',width=0.25,zorder=3)
plt.bar(X[3]+0.00,pendientesy[21],color='y',width=0.25,zorder=3)
plt.bar(X[3]+0.25,pendientesr[21],color='c',width=0.25,zorder=3)
plt.bar(X[4]-0.25,pendientesr[1],color='m',width=0.25,zorder=3)
plt.bar(X[4]+0.00,pendientesy[1],color='orange',width=0.25,zorder=3)
plt.bar(X[4]+0.25,pendientesr[1],color='b',width=0.25,zorder=3)
plt.bar(X[5]-0.25,pendientesr[19],color='r',width=0.25,zorder=3)
plt.bar(X[5]+0.00,pendientesy[19],color='y',width=0.25,zorder=3)
plt.bar(X[5]+0.25,pendientesr[19],color='c',width=0.25,zorder=3)
plt.bar(X[6]-0.25,pendientesr[4],color='m',width=0.25,zorder=3)
plt.bar(X[6]+0.00,pendientesy[4],color='orange',width=0.25,zorder=3)
plt.bar(X[6]+0.25,pendientesr[4],color='b',width=0.25,zorder=3)
plt.bar(X[7]-0.25,pendientesr[22],color='r',width=0.25,zorder=3)
plt.bar(X[7]+0.00,pendientesy[22],color='y',width=0.25,zorder=3)
```

```

plt.bar(X[7]+0.25,pendientesc[22],color='c',width=0.25,zorder=3)
plt.bar(X[8]-0.25,pendientesr[2],color='m',width=0.25,zorder=3)
plt.bar(X[8]+0.00,pendientesy[2],color='orange',width=0.25,zorder=3)
plt.bar(X[8]+0.25,pendientesc[2],color='b',width=0.25,zorder=3)
plt.bar(X[9]-0.25,pendientesr[20],color='r',width=0.25,zorder=3)
plt.bar(X[9]+0.00,pendientesy[20],color='y',width=0.25,zorder=3)
plt.bar(X[9]+0.25,pendientesc[20],color='c',width=0.25,zorder=3)
plt.bar(X[10]-0.25,pendientesr[5],color='m',width=0.25,zorder=3)
plt.bar(X[10]+0.00,pendientesy[5],color='orange',width=0.25)
plt.bar(X[10]+0.25,pendientesc[5],color='b',width=0.25,zorder=3)
plt.bar(X[11]-0.25,pendientesr[23],color='r',width=0.25,zorder=3)
plt.bar(X[11]+0.00,pendientesy[23],color='y',width=0.25,zorder=3)
plt.bar(X[11]+0.25,pendientesc[23],color='c',width=0.25,zorder=3)
plt.xticks(X, ['Control Glucosa 1',"Glucosa 1",'Control Glicerol 1',"Glicerol 1",'Contr
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

```

Out[61]: <matplotlib.legend.Legend at 0x267ef499630>



```

In [62]: #Grafico pendientes ptet-pLas-std Glucosa
X = np.arange(7)
plt.figure()
plt.title('pTet-pLas81-J23101 Glucosa 0,4%')
plt.ylabel(r'$\alpha_p$ (AU)')

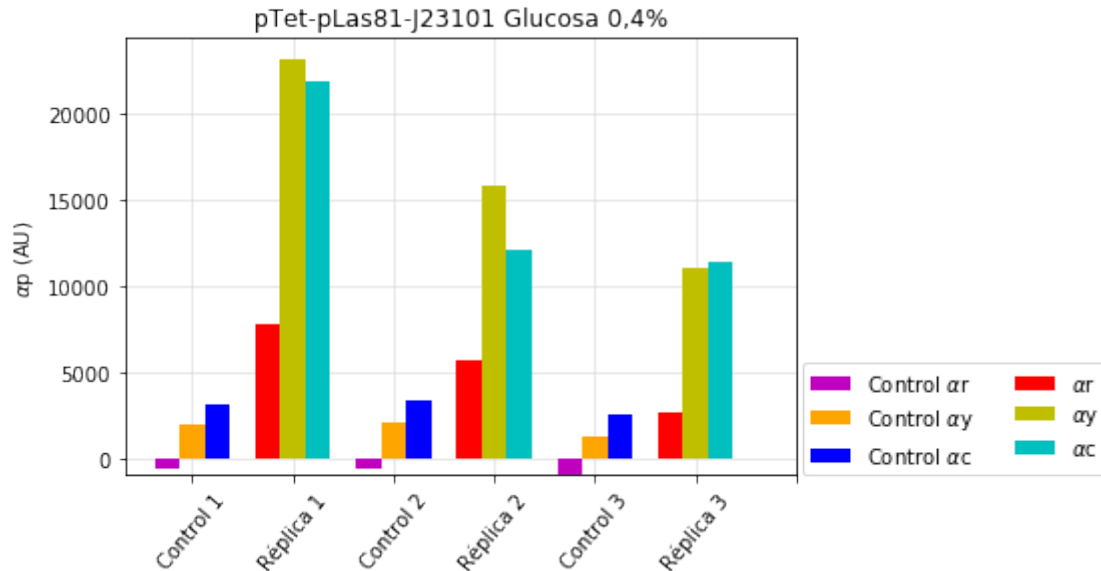
```

```

plt.bar(X[0]-0.25,pendientesr[0],color='m',width=0.25,label='Control'+ ' ' + r'$\alpha$r',zorder=3)
plt.bar(X[0]+0.00,pendientesy[0],color='orange',width=0.25,label='Control'+ ' ' + r'$\alpha$y',zorder=3)
plt.bar(X[0]+0.25,pendientesr[24],color='b',width=0.25,label='Control'+ ' ' + r'$\alpha$c',zorder=3)
plt.bar(X[1]-0.25,pendientesr[24],color='r',width=0.25,label=r'$\alpha$r',zorder=3)
plt.bar(X[1]+0.00,pendientesy[24],color='y',width=0.25,label=r'$\alpha$y',zorder=3)
plt.bar(X[1]+0.25,pendientesr[24],color='c',width=0.25,label=r'$\alpha$c',zorder=3)
plt.bar(X[2]-0.25,pendientesr[1],color='m',width=0.25,zorder=3)
plt.bar(X[2]+0.00,pendientesy[1],color='orange',width=0.25,zorder=3)
plt.bar(X[2]+0.25,pendientesr[1],color='b',width=0.25,zorder=3)
plt.bar(X[3]-0.25,pendientesr[25],color='r',width=0.25,zorder=3)
plt.bar(X[3]+0.00,pendientesy[25],color='y',width=0.25,zorder=3)
plt.bar(X[3]+0.25,pendientesr[25],color='c',width=0.25,zorder=3)
plt.bar(X[4]-0.25,pendientesr[2],color='m',width=0.25,zorder=3)
plt.bar(X[4]+0.00,pendientesy[2],color='orange',width=0.25,zorder=3)
plt.bar(X[4]+0.25,pendientesr[2],color='b',width=0.25,zorder=3)
plt.bar(X[5]-0.25,pendientesr[26],color='r',width=0.25,zorder=3)
plt.bar(X[5]+0.00,pendientesy[26],color='y',width=0.25,zorder=3)
plt.bar(X[5]+0.25,pendientesr[26],color='c',width=0.25,zorder=3)
plt.xticks(X, ['Control 1',"Réplica 1",'Control 2',"Réplica 2",'Control 3',"Réplica 3"])
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

```

Out[62]: <matplotlib.legend.Legend at 0x267eee757b8>



```

In [63]: #Grafico pendientes ptet-plas-std Glicerol
X = np.arange(7)
plt.figure()
plt.title('pTet-pLas-J23101 Glicerol 0,2%')

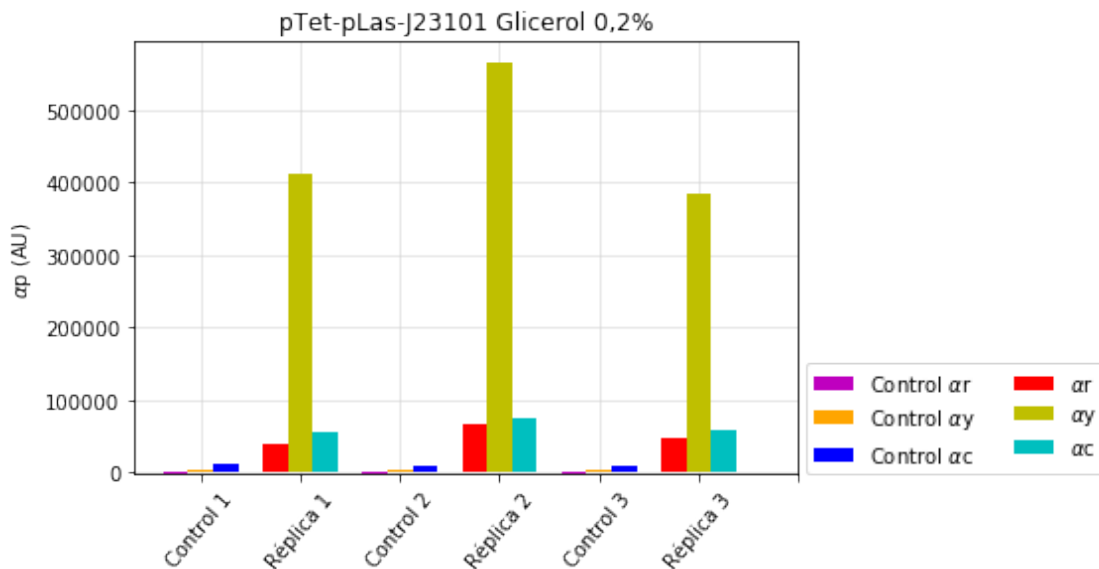
```

```

plt.ylabel(r'$\alpha$ p (AU)')
plt.bar(X[0]-0.25,pendientesr[3],color='m',width=0.25,label='Control'+ ' ' + r'$\alpha$r')
plt.bar(X[0]+0.00,pendientesy[3],color='orange',width=0.25,label='Control'+ ' ' + r'$\alpha$y')
plt.bar(X[0]+0.25,pendientesr[3],color='b',width=0.25,label='Control'+ ' ' + r'$\alpha$c')
plt.bar(X[1]-0.25,pendientesr[27],color='r',width=0.25,label=r'$\alpha$r',zorder=3)
plt.bar(X[1]+0.00,pendientesy[27],color='y',width=0.25,label=r'$\alpha$y',zorder=3)
plt.bar(X[1]+0.25,pendientesr[27],color='c',width=0.25,label=r'$\alpha$c',zorder=3)
plt.bar(X[2]-0.25,pendientesr[4],color='m',width=0.25,zorder=3)
plt.bar(X[2]+0.00,pendientesy[4],color='orange',width=0.25,zorder=3)
plt.bar(X[2]+0.25,pendientesr[4],color='b',width=0.25,zorder=3)
plt.bar(X[3]-0.25,pendientesr[28],color='r',width=0.25,zorder=3)
plt.bar(X[3]+0.00,pendientesy[28],color='y',width=0.25,zorder=3)
plt.bar(X[3]+0.25,pendientesr[28],color='c',width=0.25,zorder=3)
plt.bar(X[4]-0.25,pendientesr[5],color='m',width=0.25,zorder=3)
plt.bar(X[4]+0.00,pendientesy[5],color='orange',width=0.25,zorder=3)
plt.bar(X[4]+0.25,pendientesr[5],color='b',width=0.25,zorder=3)
plt.bar(X[5]-0.25,pendientesr[29],color='r',width=0.25,zorder=3)
plt.bar(X[5]+0.00,pendientesy[29],color='y',width=0.25,zorder=3)
plt.bar(X[5]+0.25,pendientesr[29],color='c',width=0.25,zorder=3)
plt.xticks(X, ['Control 1',"Réplica 1",'Control 2',"Réplica 2",'Control 3',"Réplica 3"])
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

```

Out [63]: <matplotlib.legend.Legend at 0x267efc7fdd8>



```

In [64]: #Grafico pendientes ptet-plas-std
X = np.arange(13)
plt.figure(figsize=(10,5))

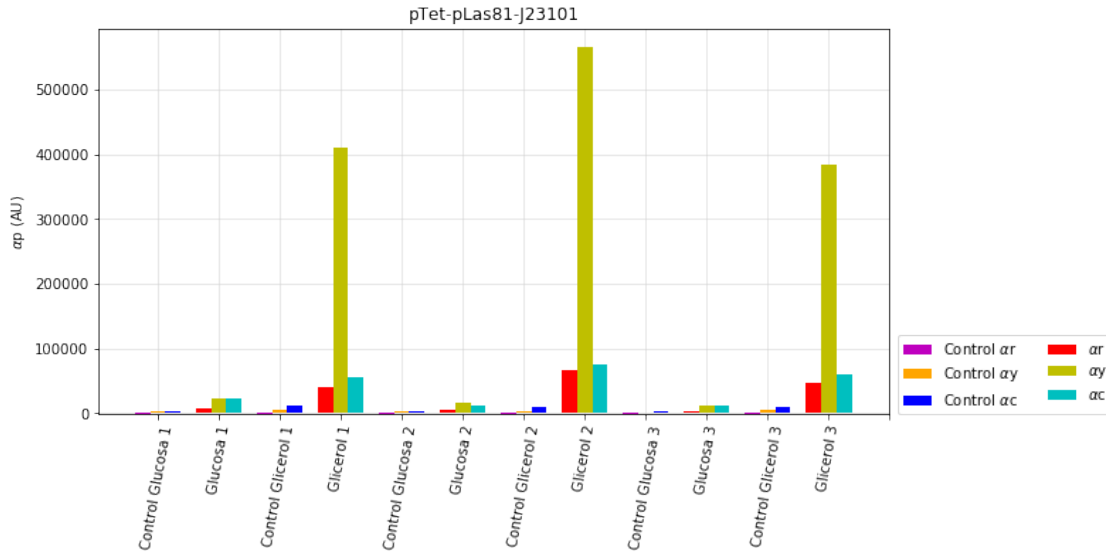
```

```

plt.title('pTet-pLas81-J23101')
plt.ylabel(r'$\alpha$ $p$  (AU)')
plt.bar(X[0]-0.25,pendientesr[0],color='m',width=0.25,label='Control'+ ' '+ r'$\alpha$ $r$ ')
plt.bar(X[0]+0.00,pendientesy[0],color='orange',width=0.25,label='Control'+ ' '+ r'$\alpha$ $y$ ')
plt.bar(X[0]+0.25,pendientesr[0],color='b',width=0.25,label='Control'+ ' '+ r'$\alpha$ $c$ ')
plt.bar(X[1]-0.25,pendientesr[24],color='r',width=0.25,label=r'$\alpha$ $r$ ',zorder=3)
plt.bar(X[1]+0.00,pendientesy[24],color='y',width=0.25,label=r'$\alpha$ $y$ ',zorder=3)
plt.bar(X[1]+0.25,pendientesr[24],color='c',width=0.25,label=r'$\alpha$ $c$ ',zorder=3)
plt.bar(X[2]-0.25,pendientesr[3],color='m',width=0.25,zorder=3)
plt.bar(X[2]+0.00,pendientesy[3],color='orange',width=0.25,zorder=3)
plt.bar(X[2]+0.25,pendientesr[3],color='b',width=0.25,zorder=3)
plt.bar(X[3]-0.25,pendientesr[27],color='r',width=0.25,zorder=3)
plt.bar(X[3]+0.00,pendientesy[27],color='y',width=0.25,zorder=3)
plt.bar(X[3]+0.25,pendientesr[27],color='c',width=0.25,zorder=3)
plt.bar(X[4]-0.25,pendientesr[1],color='m',width=0.25,zorder=3)
plt.bar(X[4]+0.00,pendientesy[1],color='orange',width=0.25,zorder=3)
plt.bar(X[4]+0.25,pendientesr[1],color='b',width=0.25,zorder=3)
plt.bar(X[5]-0.25,pendientesr[25],color='r',width=0.25,zorder=3)
plt.bar(X[5]+0.00,pendientesy[25],color='y',width=0.25,zorder=3)
plt.bar(X[5]+0.25,pendientesr[25],color='c',width=0.25,zorder=3)
plt.bar(X[6]-0.25,pendientesr[4],color='m',width=0.25,zorder=3)
plt.bar(X[6]+0.00,pendientesy[4],color='orange',width=0.25,zorder=3)
plt.bar(X[6]+0.25,pendientesr[4],color='b',width=0.25,zorder=3)
plt.bar(X[7]-0.25,pendientesr[28],color='r',width=0.25,zorder=3)
plt.bar(X[7]+0.00,pendientesy[28],color='y',width=0.25,zorder=3)
plt.bar(X[7]+0.25,pendientesr[28],color='c',width=0.25,zorder=3)
plt.bar(X[8]-0.25,pendientesr[2],color='m',width=0.25,zorder=3)
plt.bar(X[8]+0.00,pendientesy[2],color='orange',width=0.25,zorder=3)
plt.bar(X[8]+0.25,pendientesr[2],color='b',width=0.25,zorder=3)
plt.bar(X[9]-0.25,pendientesr[26],color='r',width=0.25,zorder=3)
plt.bar(X[9]+0.00,pendientesy[26],color='y',width=0.25,zorder=3)
plt.bar(X[9]+0.25,pendientesr[26],color='c',width=0.25,zorder=3)
plt.bar(X[10]-0.25,pendientesr[5],color='m',width=0.25,zorder=3)
plt.bar(X[10]+0.00,pendientesy[5],color='orange',width=0.25,zorder=3)
plt.bar(X[10]+0.25,pendientesr[5],color='b',width=0.25,zorder=3)
plt.bar(X[11]-0.25,pendientesr[29],color='r',width=0.25,zorder=3)
plt.bar(X[11]+0.00,pendientesy[29],color='y',width=0.25,zorder=3)
plt.bar(X[11]+0.25,pendientesr[29],color='c',width=0.25,zorder=3)
plt.xticks(X, ['Control Glucosa 1',"Glucosa 1",'Control Glicerol 1',"Glicerol 1",'Control Glucosa 1',"Glucosa 1",'Control Glicerol 1',"Glicerol 1"])
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

```

Out[64]: <matplotlib.legend.Legend at 0x267f0df9a20>



In [141]: *#Grafico pendientes todo*

```
X = np.arange(30)
plt.figure(figsize=(20,10))
plt.title(r'$\alpha$p Ensayo 2', fontsize=15.0)
plt.ylabel(r'$\alpha$p (AU)')
plt.bar(X[0]-0.25,pendientesr[0],color='m',width=0.25,label='Control'+ ' ' + r'$\alpha$r')
plt.bar(X[0]+0.00,pendientesy[0],color='orange',width=0.25,label='Control'+ ' ' + r'$\alpha$y')
plt.bar(X[0]+0.25,pendientesc[0],color='b',width=0.25,label='Control'+ ' ' + r'$\alpha$c')
plt.bar(X[1]-0.25,pendientesr[1],color='m',width=0.25,zorder=3)
plt.bar(X[1]+0.00,pendientesy[1],color='orange',width=0.25,zorder=3)
plt.bar(X[1]+0.25,pendientesc[1],color='b',width=0.25,zorder=3)
plt.bar(X[2]-0.25,pendientesr[2],color='m',width=0.25,zorder=3)
plt.bar(X[2]+0.00,pendientesy[2],color='orange',width=0.25,zorder=3)
plt.bar(X[2]+0.25,pendientesc[2],color='b',width=0.25,zorder=3)
plt.bar(X[3]-0.25,pendientesr[3],color='m',width=0.25,zorder=3)
plt.bar(X[3]+0.00,pendientesy[3],color='orange',width=0.25,zorder=3)
plt.bar(X[3]+0.25,pendientesc[3],color='b',width=0.25,zorder=3)
plt.bar(X[4]-0.25,pendientesr[4],color='m',width=0.25,zorder=3)
plt.bar(X[4]+0.00,pendientesy[4],color='orange',width=0.25,zorder=3)
plt.bar(X[4]+0.25,pendientesc[4],color='b',width=0.25,zorder=3)
plt.bar(X[5]-0.25,pendientesr[5],color='m',width=0.25,zorder=3)
plt.bar(X[5]+0.00,pendientesy[5],color='orange',width=0.25,zorder=3)
plt.bar(X[5]+0.25,pendientesc[5],color='b',width=0.25,zorder=3)
plt.bar(X[6]-0.25,pendientesr[6],color='r',width=0.25,label=r'$\alpha$r',zorder=3)
plt.bar(X[6]+0.00,pendientesy[6],color='y',width=0.25,label=r'$\alpha$y',zorder=3)
plt.bar(X[6]+0.25,pendientesc[6],color='c',width=0.25,label=r'$\alpha$c',zorder=3)
plt.bar(X[7]-0.25,pendientesr[9],color='r',width=0.25,zorder=3)
plt.bar(X[7]+0.00,pendientesy[9],color='y',width=0.25,zorder=3)
plt.bar(X[7]+0.25,pendientesc[9],color='c',width=0.25,zorder=3)
```

```

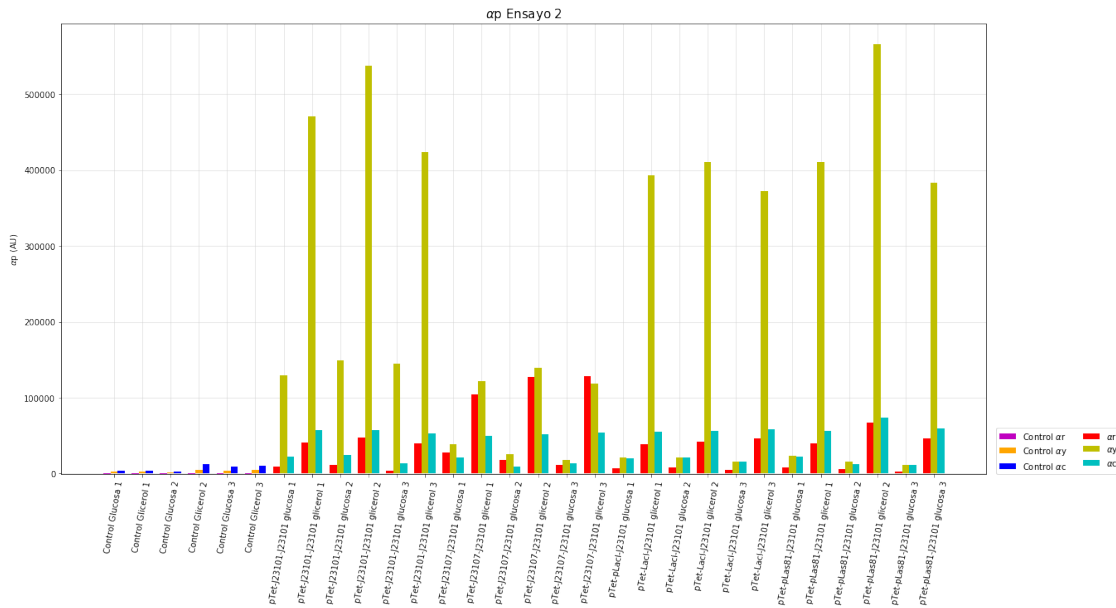
plt.bar(X[8]-0.25,pendientesr[7],color='r',width=0.25,zorder=3)
plt.bar(X[8]+0.00,pendientesy[7],color='y',width=0.25,zorder=3)
plt.bar(X[8]+0.25,pendientesr[7],color='c',width=0.25,zorder=3)
plt.bar(X[9]-0.25,pendientesr[10],color='r',width=0.25,zorder=3)
plt.bar(X[9]+0.00,pendientesy[10],color='y',width=0.25,zorder=3)
plt.bar(X[9]+0.25,pendientesr[10],color='c',width=0.25,zorder=3)
plt.bar(X[10]-0.25,pendientesr[8],color='r',width=0.25,zorder=3)
plt.bar(X[10]+0.00,pendientesy[8],color='y',width=0.25,zorder=3)
plt.bar(X[10]+0.25,pendientesr[8],color='c',width=0.25,zorder=3)
plt.bar(X[11]-0.25,pendientesr[11],color='r',width=0.25,zorder=3)
plt.bar(X[11]+0.00,pendientesy[11],color='y',width=0.25,zorder=3)
plt.bar(X[11]+0.25,pendientesr[11],color='c',width=0.25,zorder=3)
plt.bar(X[12]-0.25,pendientesr[12],color='r',width=0.25,zorder=3)
plt.bar(X[12]+0.00,pendientesy[12],color='y',width=0.25,zorder=3)
plt.bar(X[12]+0.25,pendientesr[12],color='c',width=0.25,zorder=3)
plt.bar(X[13]-0.25,pendientesr[15],color='r',width=0.25,zorder=3)
plt.bar(X[13]+0.00,pendientesy[15],color='y',width=0.25,zorder=3)
plt.bar(X[13]+0.25,pendientesr[15],color='c',width=0.25,zorder=3)
plt.bar(X[14]-0.25,pendientesr[13],color='r',width=0.25,zorder=3)
plt.bar(X[14]+0.00,pendientesy[13],color='y',width=0.25,zorder=3)
plt.bar(X[14]+0.25,pendientesr[13],color='c',width=0.25,zorder=3)
plt.bar(X[15]-0.25,pendientesr[16],color='r',width=0.25,zorder=3)
plt.bar(X[15]+0.00,pendientesy[16],color='y',width=0.25,zorder=3)
plt.bar(X[15]+0.25,pendientesr[16],color='c',width=0.25,zorder=3)
plt.bar(X[16]-0.25,pendientesr[14],color='r',width=0.25,zorder=3)
plt.bar(X[16]+0.00,pendientesy[14],color='y',width=0.25,zorder=3)
plt.bar(X[16]+0.25,pendientesr[14],color='c',width=0.25,zorder=3)
plt.bar(X[17]-0.25,pendientesr[17],color='r',width=0.25,zorder=3)
plt.bar(X[17]+0.00,pendientesy[17],color='y',width=0.25,zorder=3)
plt.bar(X[17]+0.25,pendientesr[17],color='c',width=0.25,zorder=3)
plt.bar(X[18]-0.25,pendientesr[18],color='r',width=0.25,zorder=3)
plt.bar(X[18]+0.00,pendientesy[18],color='y',width=0.25,zorder=3)
plt.bar(X[18]+0.25,pendientesr[18],color='c',width=0.25,zorder=3)
plt.bar(X[19]-0.25,pendientesr[21],color='r',width=0.25,zorder=3)
plt.bar(X[19]+0.00,pendientesy[21],color='y',width=0.25,zorder=3)
plt.bar(X[19]+0.25,pendientesr[21],color='c',width=0.25,zorder=3)
plt.bar(X[20]-0.25,pendientesr[19],color='r',width=0.25,zorder=3)
plt.bar(X[20]+0.00,pendientesy[19],color='y',width=0.25,zorder=3)
plt.bar(X[20]+0.25,pendientesr[19],color='c',width=0.25,zorder=3)
plt.bar(X[21]-0.25,pendientesr[22],color='r',width=0.25,zorder=3)
plt.bar(X[21]+0.00,pendientesy[22],color='y',width=0.25,zorder=3)
plt.bar(X[21]+0.25,pendientesr[22],color='c',width=0.25,zorder=3)
plt.bar(X[22]-0.25,pendientesr[20],color='r',width=0.25,zorder=3)
plt.bar(X[22]+0.00,pendientesy[20],color='y',width=0.25,zorder=3)
plt.bar(X[22]+0.25,pendientesr[20],color='c',width=0.25,zorder=3)
plt.bar(X[23]-0.25,pendientesr[23],color='r',width=0.25,zorder=3)
plt.bar(X[23]+0.00,pendientesy[23],color='y',width=0.25,zorder=3)
plt.bar(X[23]+0.25,pendientesr[23],color='c',width=0.25,zorder=3)

```

```

plt.bar(X[24]-0.25,pendientesr[24],color='r',width=0.25,zorder=3)
plt.bar(X[24]+0.00,pendientesy[24],color='y',width=0.25,zorder=3)
plt.bar(X[24]+0.25,pendientesc[24],color='c',width=0.25,zorder=3)
plt.bar(X[25]-0.25,pendientesr[27],color='r',width=0.25,zorder=3)
plt.bar(X[25]+0.00,pendientesy[27],color='y',width=0.25,zorder=3)
plt.bar(X[25]+0.25,pendientesc[27],color='c',width=0.25,zorder=3)
plt.bar(X[26]-0.25,pendientesr[25],color='r',width=0.25,zorder=3)
plt.bar(X[26]+0.00,pendientesy[25],color='y',width=0.25,zorder=3)
plt.bar(X[26]+0.25,pendientesc[25],color='c',width=0.25,zorder=3)
plt.bar(X[27]-0.25,pendientesr[28],color='r',width=0.25,zorder=3)
plt.bar(X[27]+0.00,pendientesy[28],color='y',width=0.25,zorder=3)
plt.bar(X[27]+0.25,pendientesc[28],color='c',width=0.25,zorder=3)
plt.bar(X[28]-0.25,pendientesr[26],color='r',width=0.25,zorder=3)
plt.bar(X[28]+0.00,pendientesy[26],color='y',width=0.25,zorder=3)
plt.bar(X[28]+0.25,pendientesc[26],color='c',width=0.25,zorder=3)
plt.bar(X[29]-0.25,pendientesr[29],color='r',width=0.25,zorder=3)
plt.bar(X[29]+0.00,pendientesy[29],color='y',width=0.25,zorder=3)
plt.bar(X[29]+0.25,pendientesc[29],color='c',width=0.25,zorder=3)
plt.xticks(X, ['Control Glucosa 1','Control Glicerol 1','Control Glucosa 2','Control G
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)
plt.savefig(filename='alfa-p ensayo 2.png', dpi=300, facecolor='w', edgecolor='w',bbox

```



```

In [66]: cglu=[[slopec1,slopec2,slopec3],[slopec7,slopec8,slopec9],[slopec13,slopec14,slopec15],
cgli=[[slopec4,slopec5,slopec6],[slopec10,slopec11,slopec12],[slopec16,slopec17,slopec18],
rglu=[[sloper1,sloper2,sloper3],[sloper7,sloper8,sloper9],[sloper13,sloper14,sloper15],
rgli=[[sloper4,sloper5,sloper6],[sloper10,sloper11,sloper12],[sloper16,sloper17,sloper18]

```



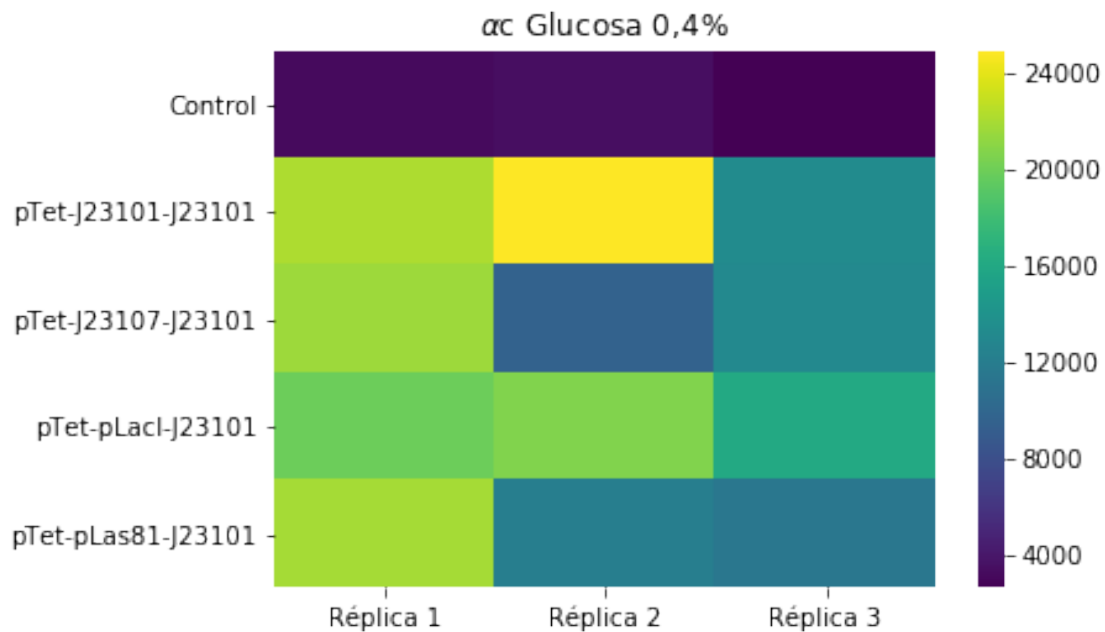
```
yglu=[[slopey1,slopey2,slopey3],[slopey7,slopey8,slopey9],[slopey13,slopey14,slopey15],
ygli=[[slopey4,slopey5,slopey6],[slopey10,slopey11,slopey12],[slopey16,slopey17,slopey18]]
```

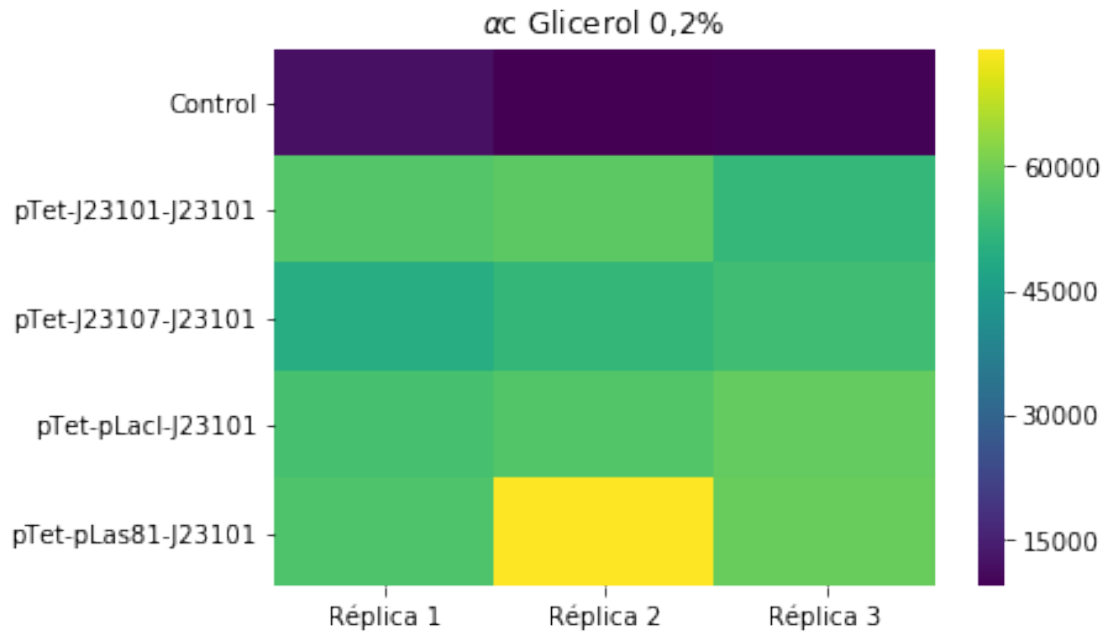
```
In [67]: xlabel=['Réplica 1','Réplica 2','Réplica 3']
ylabel=['Control','pTet-J23101-J23101','pTet-J23107-J23101','pTet-pLacI-J23101','pTet-pLas81-J23101']
```

```
plt.figure()
plt.title(r'$\alpha$ Glucosa 0,4%')
sns.heatmap(cglu,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)
```

```
plt.figure()
plt.title(r'$\alpha$ Glicerol 0,2%')
sns.heatmap(cgli,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)
```

```
Out[67]: <matplotlib.axes._subplots.AxesSubplot at 0x267faca8f28>
```



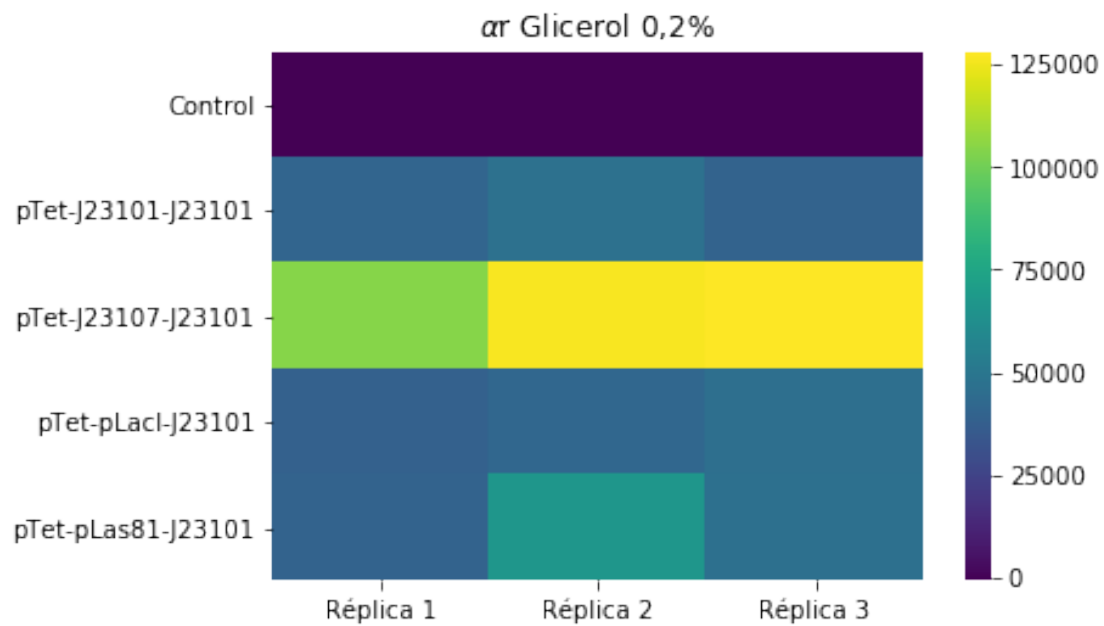
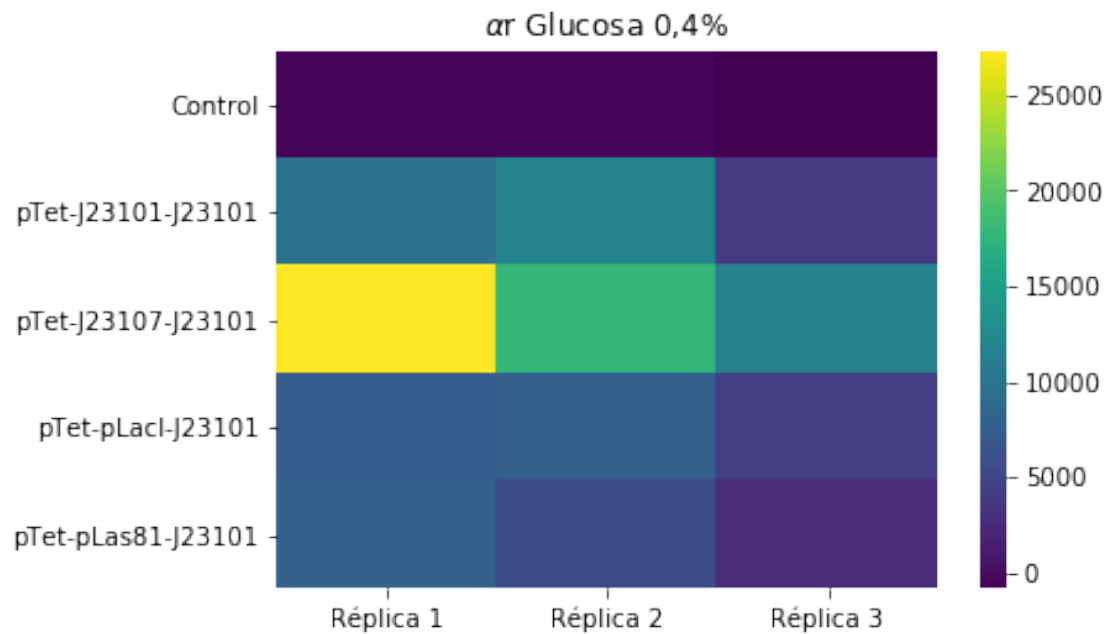


```
In [68]: xlabel=['Réplica 1', 'Réplica 2', 'Réplica 3']
        ylabel=['Control', 'pTet-J23101-J23101', 'pTet-J23107-J23101', 'pTet-pLacI-J23101', 'pTet-pLas81-J23101']

        plt.figure()
        plt.title(r'$\alpha$ Glucosa 0,4%')
        sns.heatmap(rglu, cmap='viridis', xticklabels=xlabel, yticklabels=ylabel)

        plt.figure()
        plt.title(r'$\alpha$ Glicerol 0,2%')
        sns.heatmap(rgli, cmap='viridis', xticklabels=xlabel, yticklabels=ylabel)

Out[68]: <matplotlib.axes._subplots.AxesSubplot at 0x267eeff0e10>
```

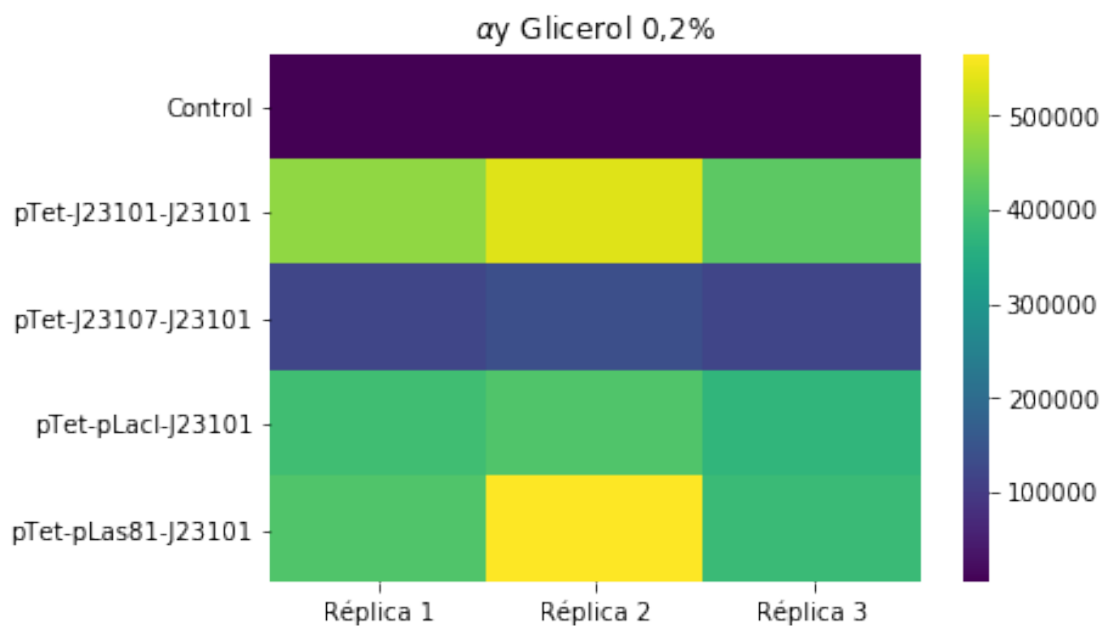
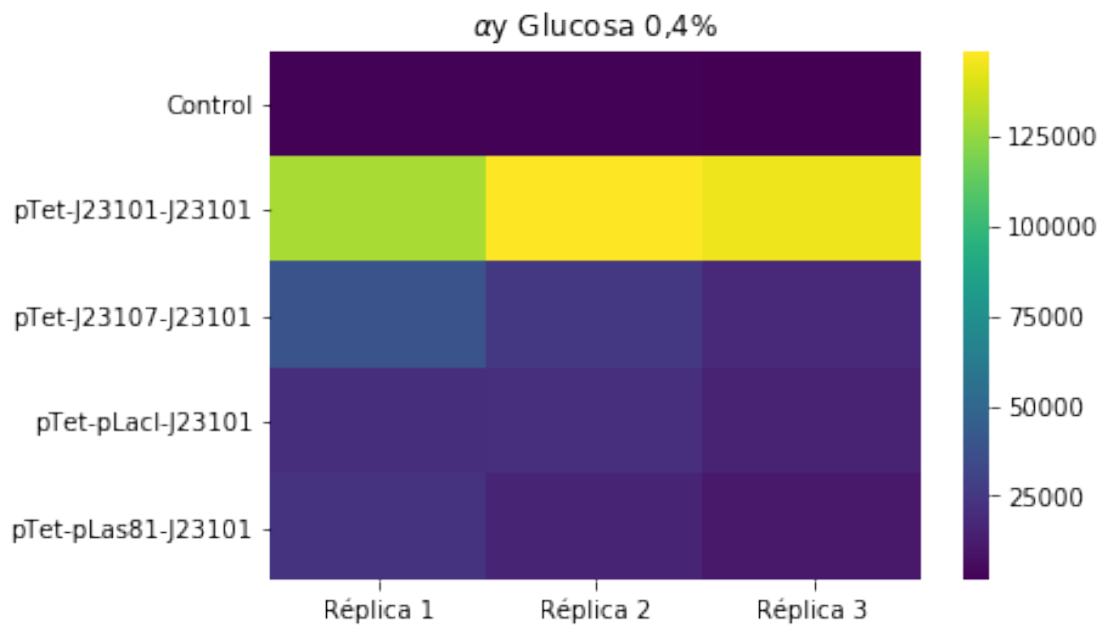


```
In [69]: xlabel=['Réplica 1','Réplica 2','Réplica 3']
        ylabel=['Control','pTet-J23101-J23101','pTet-J23107-J23101','pTet-pLacI-J23101','pTet-pLas81-J23101']
        plt.figure()
```

```
plt.title(r'\alpha$y Glucosa 0,4%')
sns.heatmap(yglu,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)

plt.figure()
plt.title(r'\alpha$y Glicerol 0,2%')
sns.heatmap(ygli,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)
```

Out[69]: <matplotlib.axes._subplots.AxesSubplot at 0x267f179f748>



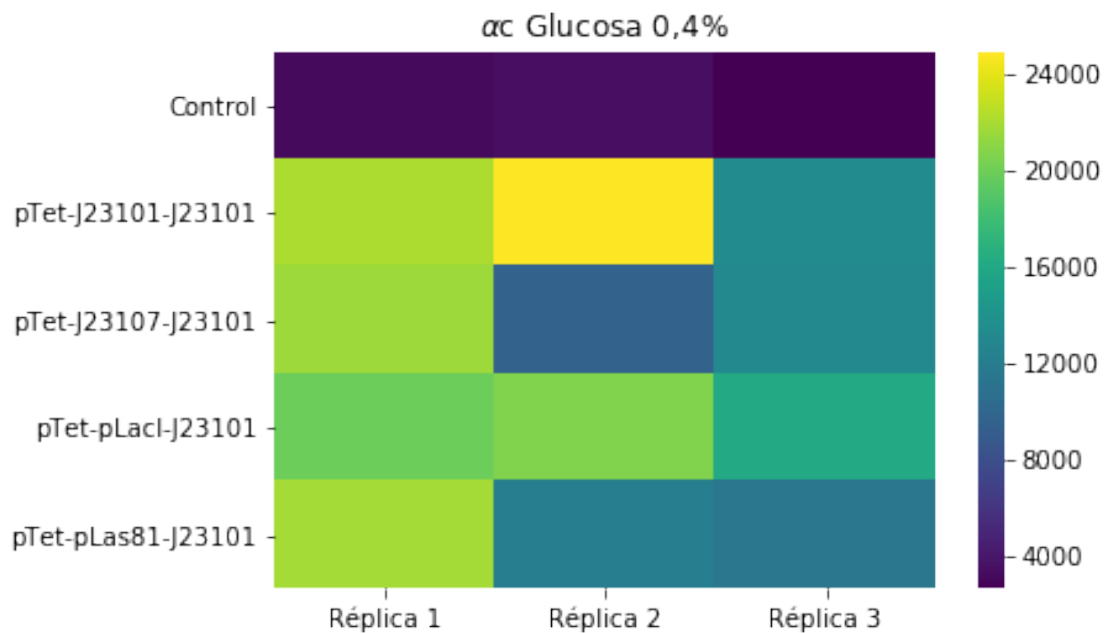
```

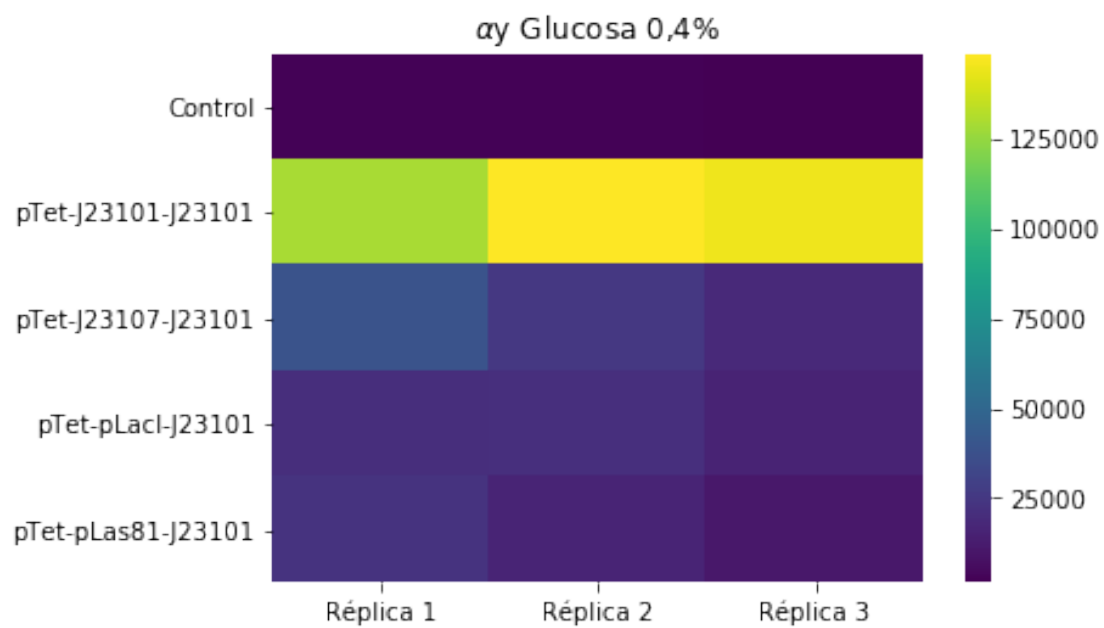
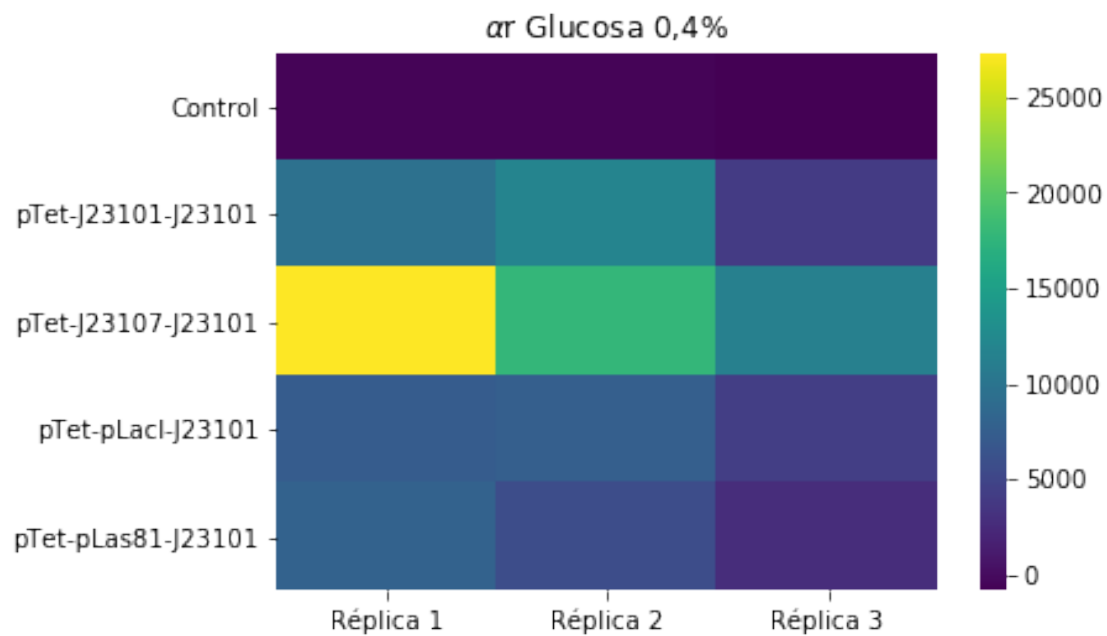
In [70]: xlabel=['Réplica 1','Réplica 2','Réplica 3']
        ylabel=['Control','pTet-J23101-J23101','pTet-J23107-J23101','pTet-pLacI-J23101','pTet-p

plt.figure()
plt.title(r'$\alpha$c Glucosa 0,4%')
sns.heatmap(cglu,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)
plt.figure()
plt.title(r'$\alpha$r Glucosa 0,4%')
sns.heatmap(rglu,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)
plt.figure()
plt.title(r'$\alpha$y Glucosa 0,4%')
sns.heatmap(yglu,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)

```

Out[70]: <matplotlib.axes._subplots.AxesSubplot at 0x267f1799518>



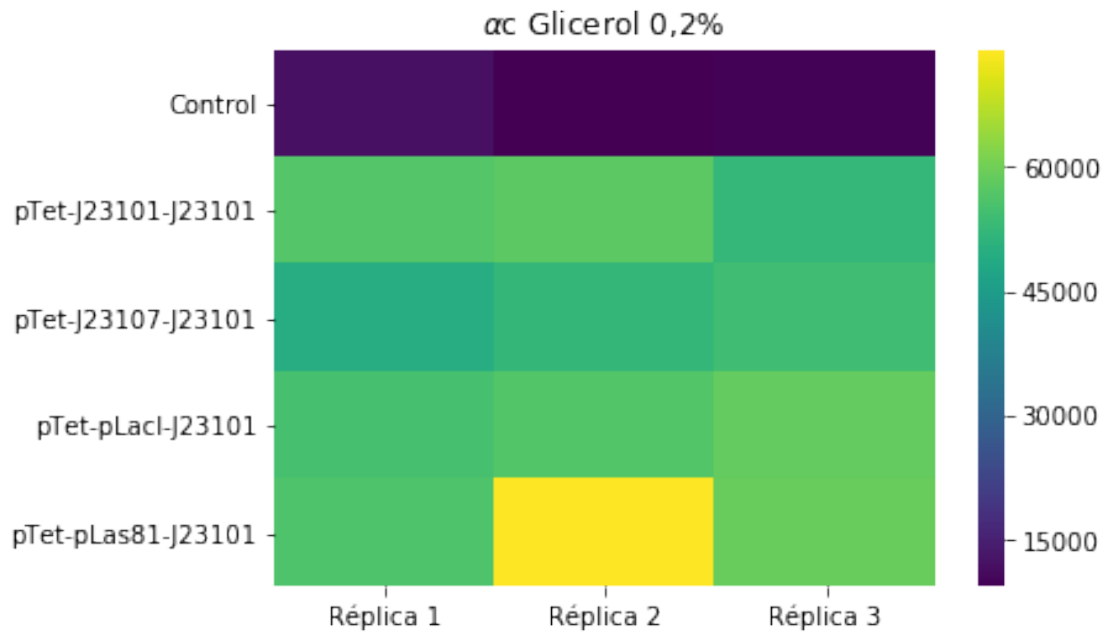


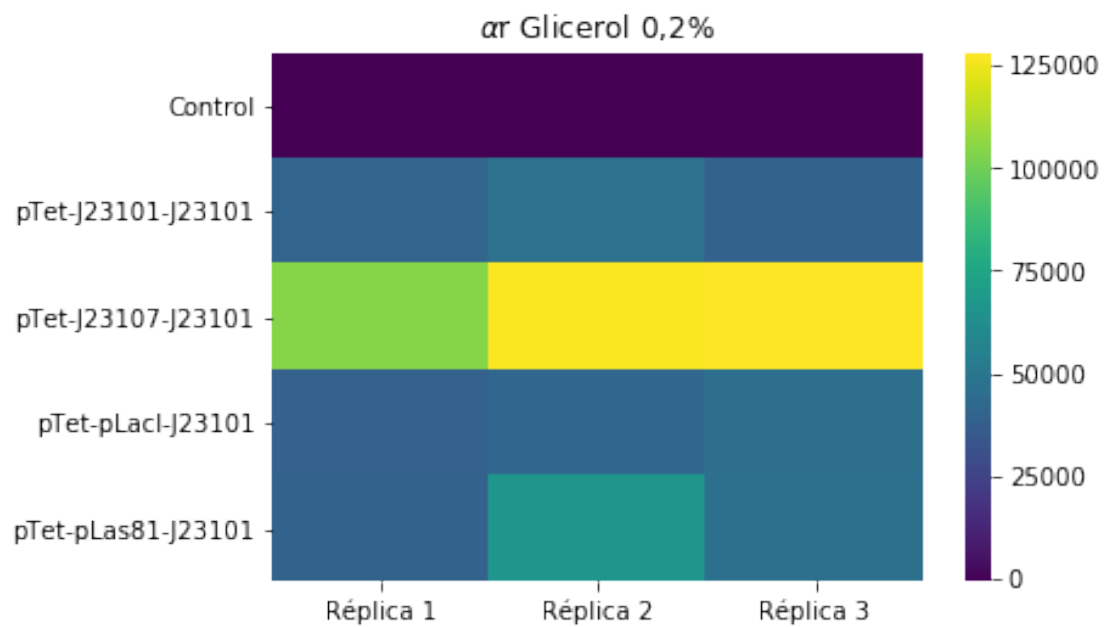
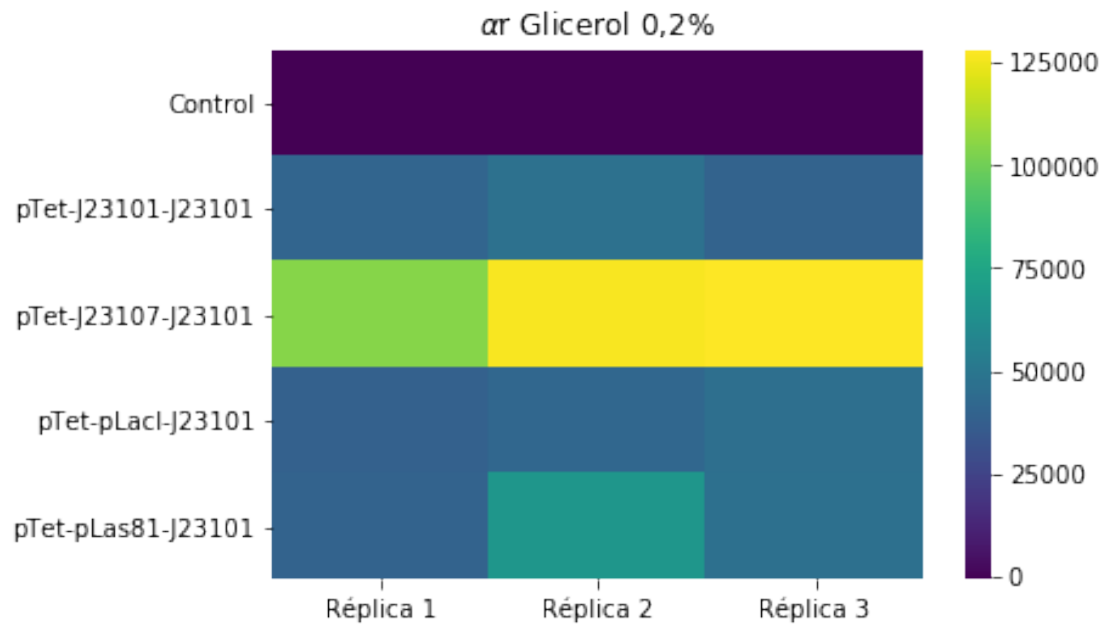
```
In [71]: xlabel=['Réplica 1','Réplica 2','Réplica 3']
        ylabel=['Control','pTet-J23101-J23101','pTet-J23107-J23101','pTet-pLacI-J23101','pTet-pLas81-J23101']

        plt.figure()
```

```
plt.title(r'\alpha$c Glicerol 0,2%')
sns.heatmap(cgli,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)
plt.figure()
plt.title(r'\alpha$r Glicerol 0,2%')
sns.heatmap(rgli,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)
plt.figure()
plt.title(r'\alpha$r Glicerol 0,2%')
sns.heatmap(rgli,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)
```

Out[71]: <matplotlib.axes._subplots.AxesSubplot at 0x267f1ba0dd8>





```
In [72]: #grafico de ac versus Um
plt.figure(figsize=(8,5))
plt.title(r'\alpha$c Glucosa 0,4%')
plt.xlabel(r'\mu$m (min$^{-1}$)')
```



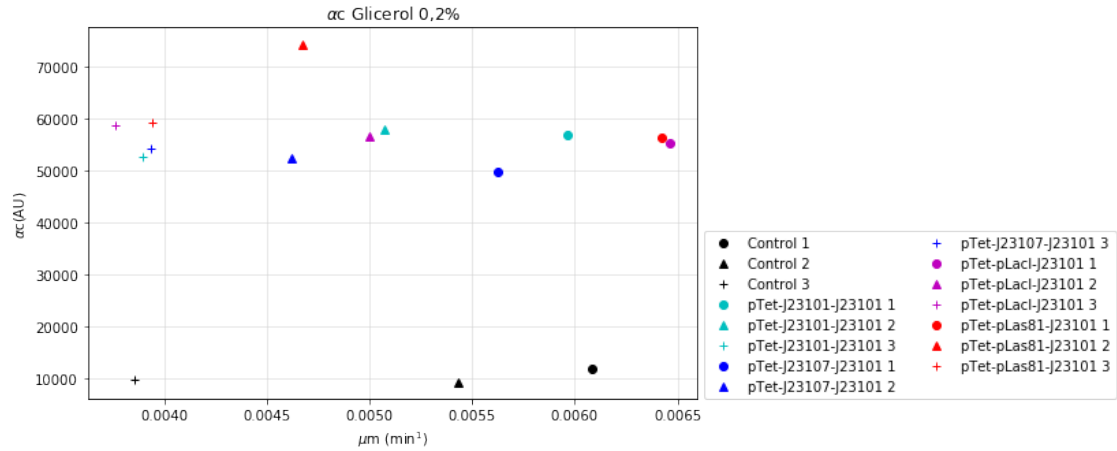
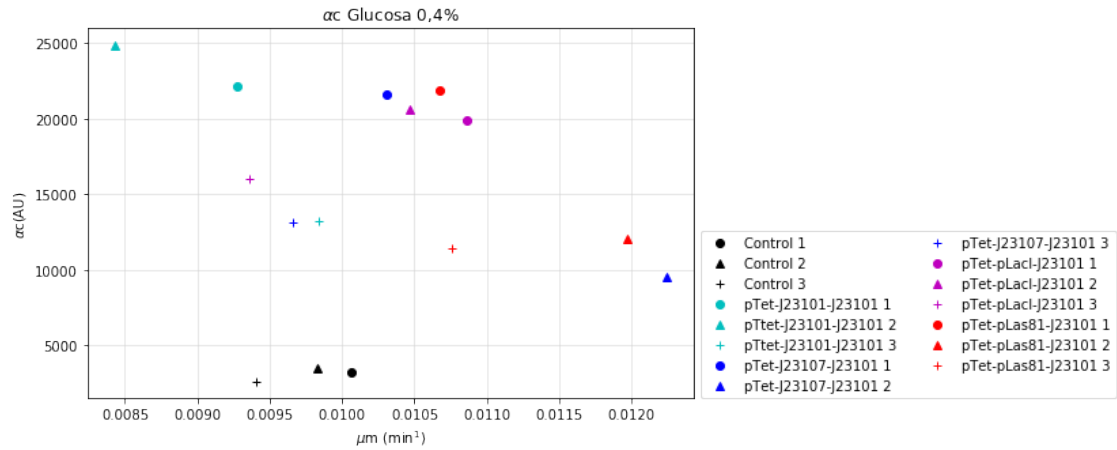
```

plt.ylabel(r'$\alpha$c(AU)')
plt.plot(um1,slopec1,'ko',label='Control 1')
plt.plot(um2,slopec2,'k^',label='Control 2')
plt.plot(um3,slopec3,'k+',label='Control 3')
plt.plot(um7,slopec7,'co',label='pTet-J23101-J23101 1')
plt.plot(um8,slopec8,'c^',label='pTtet-J23101-J23101 2')
plt.plot(um9,slopec9,'c+',label='pTtet-J23101-J23101 3')
plt.plot(um13,slopec13,'bo',label='pTet-J23107-J23101 1')
plt.plot(um14,slopec14,'b^',label='pTet-J23107-J23101 2')
plt.plot(um15,slopec15,'b+',label='pTet-J23107-J23101 3')
plt.plot(um19,slopec19,'mo',label='pTet-pLacI-J23101 1')
plt.plot(um20,slopec20,'m^',label='pTet-pLacI-J23101 2')
plt.plot(um21,slopec21,'m+',label='pTet-pLacI-J23101 3')
plt.plot(um25,slopec25,'ro',label='pTet-pLas81-J23101 1')
plt.plot(um26,slopec26,'r^',label='pTet-pLas81-J23101 2')
plt.plot(um27,slopec27,'r+',label='pTet-pLas81-J23101 3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

#grafico de ac versus Um
plt.figure(figsize=(8,5))
plt.title(r'$\alpha$c Glicerol 0,2%')
plt.xlabel(r'$\mu$m (min$^1$)')
plt.ylabel(r'$\alpha$c(AU)')
plt.plot(um4,slopec4,'ko',label='Control 1')
plt.plot(um5,slopec5,'k^',label='Control 2')
plt.plot(um6,slopec6,'k+',label='Control 3')
plt.plot(um10,slopec10,'co',label='pTet-J23101-J23101 1')
plt.plot(um11,slopec11,'c^',label='pTet-J23101-J23101 2')
plt.plot(um12,slopec12,'c+',label='pTet-J23101-J23101 3')
plt.plot(um16,slopec16,'bo',label='pTet-J23107-J23101 1')
plt.plot(um17,slopec17,'b^',label='pTet-J23107-J23101 2')
plt.plot(um18,slopec18,'b+',label='pTet-J23107-J23101 3')
plt.plot(um22,slopec22,'mo',label='pTet-pLacI-J23101 1')
plt.plot(um23,slopec23,'m^',label='pTet-pLacI-J23101 2')
plt.plot(um24,slopec24,'m+',label='pTet-pLacI-J23101 3')
plt.plot(um28,slopec28,'ro',label='pTet-pLas81-J23101 1')
plt.plot(um29,slopec29,'r^',label='pTet-pLas81-J23101 2')
plt.plot(um30,slopec30,'r+',label='pTet-pLas81-J23101 3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

```

Out[72]: <matplotlib.legend.Legend at 0x267f1cf6048>



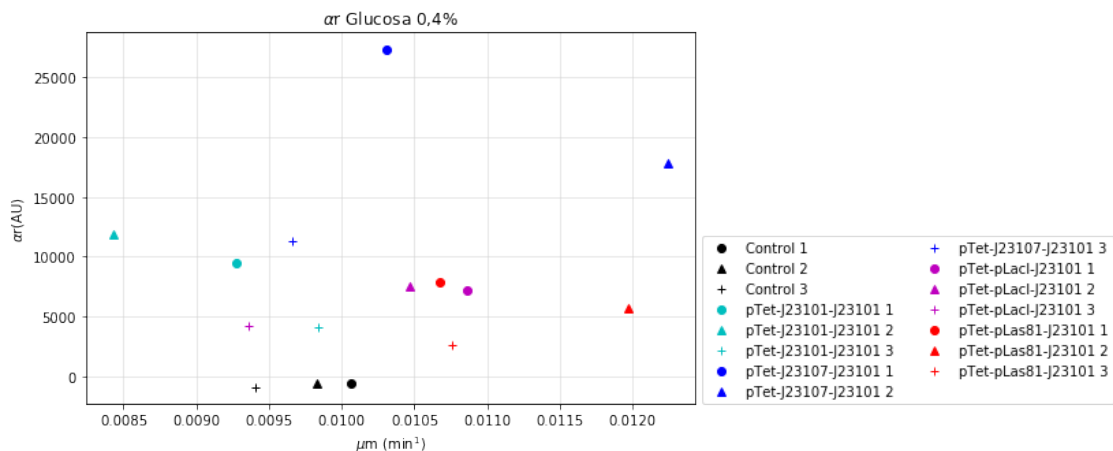
```
In [73]: #grafico de ac versus Um
plt.figure(figsize=(8,5))
plt.title(r'$\alpha$ Glucosa 0,4%')
plt.xlabel(r'$\mu$ (min$^{-1}$)')
plt.ylabel(r'$\alpha$ (AU)')
plt.plot(um1,sloper1,'ko',label='Control 1')
plt.plot(um2,sloper2,'k^',label='Control 2')
plt.plot(um3,sloper3,'k+',label='Control 3')
plt.plot(um7,sloper7,'co',label='pTet-J23101-J23101 1')
plt.plot(um8,sloper8,'c^',label='pTet-J23101-J23101 2')
plt.plot(um9,sloper9,'c+',label='pTet-J23101-J23101 3')
plt.plot(um13,sloper13,'bo',label='pTet-J23107-J23101 1')
plt.plot(um14,sloper14,'b^',label='pTet-J23107-J23101 2')
plt.plot(um15,sloper15,'b+',label='pTet-J23107-J23101 3')
plt.plot(um19,sloper19,'mo',label='pTet-pLacI-J23101 1')
```

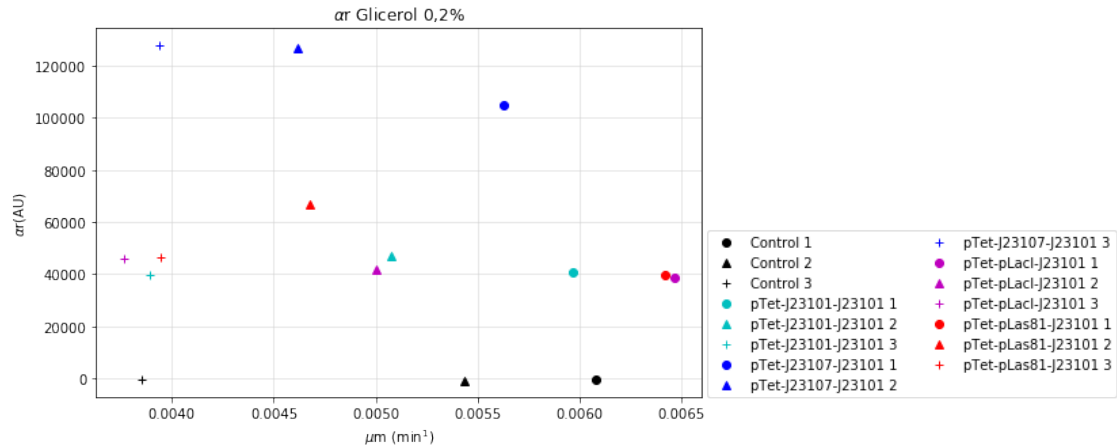
```
plt.plot(um20,sloper20,'m^',label='pTet-pLacI-J23101 2')
plt.plot(um21,sloper21,'m+',label='pTet-pLacI-J23101 3')
plt.plot(um25,sloper25,'ro',label='pTet-pLas81-J23101 1')
plt.plot(um26,sloper26,'r^',label='pTet-pLas81-J23101 2')
plt.plot(um27,sloper27,'r+',label='pTet-pLas81-J23101 3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)
```

#grafico de ac versus Um

```
plt.figure(figsize=(8,5))
plt.title(r'$\alpha$ Glicerol 0,2%')
plt.xlabel(r'$\mu$M (min$^{-1}$)')
plt.ylabel(r'$\alpha$(AU)')
plt.plot(um4,sloper4,'ko',label='Control 1')
plt.plot(um5,sloper5,'k^',label='Control 2')
plt.plot(um6,sloper6,'k+',label='Control 3')
plt.plot(um10,sloper10,'co',label='pTet-J23101-J23101 1')
plt.plot(um11,sloper11,'c^',label='pTet-J23101-J23101 2')
plt.plot(um12,sloper12,'c+',label='pTet-J23101-J23101 3')
plt.plot(um16,sloper16,'bo',label='pTet-J23107-J23101 1')
plt.plot(um17,sloper17,'b^',label='pTet-J23107-J23101 2')
plt.plot(um18,sloper18,'b+',label='pTet-J23107-J23101 3')
plt.plot(um22,sloper22,'mo',label='pTet-pLacI-J23101 1')
plt.plot(um23,sloper23,'m^',label='pTet-pLacI-J23101 2')
plt.plot(um24,sloper24,'m+',label='pTet-pLacI-J23101 3')
plt.plot(um28,sloper28,'ro',label='pTet-pLas81-J23101 1')
plt.plot(um29,sloper29,'r^',label='pTet-pLas81-J23101 2')
plt.plot(um30,sloper30,'r+',label='pTet-pLas81-J23101 3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)
```

Out[73]: <matplotlib.legend.Legend at 0x267efb64710>





```
In [74]: #grafico de ac versus Um
plt.figure(figsize=(8,5))
plt.title(r'$\alpha$y Glucosa 0,4%')
plt.xlabel(r'$\mu$m (min$^{-1}$)')
plt.ylabel(r'$\alpha$y(AU)')
plt.plot(um1,slopy1,'ko',label='Control 1')
plt.plot(um2,slopy2,'k^',label='Control 2')
plt.plot(um3,slopy3,'k+',label='Control 3')
plt.plot(um7,slopy7,'co',label='pTet-J23101-J23101 1')
plt.plot(um8,slopy8,'c^',label='pTet-J23101-J23101 2')
plt.plot(um9,slopy9,'c+',label='pTet-J23101-J23101 3')
plt.plot(um13,slopy13,'bo',label='pTet-J23107-J23101 1')
plt.plot(um14,slopy14,'b^',label='pTet-J23107-J23101 2')
plt.plot(um15,slopy15,'b+',label='pTet-J23107-J23101 3')
plt.plot(um19,slopy19,'mo',label='pTet-pLacI-J23101 1')
plt.plot(um20,slopy20,'m^',label='pTet-pLacI-J23101 2')
plt.plot(um21,slopy21,'m+',label='pTet-pLacI-J23101 3')
plt.plot(um25,slopy25,'ro',label='pTet-pLas81-J23101 1')
plt.plot(um26,slopy26,'r^',label='pTet-pLas81-J23101 2')
plt.plot(um27,slopy27,'r+',label='pTet-pLas81-J23101 3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

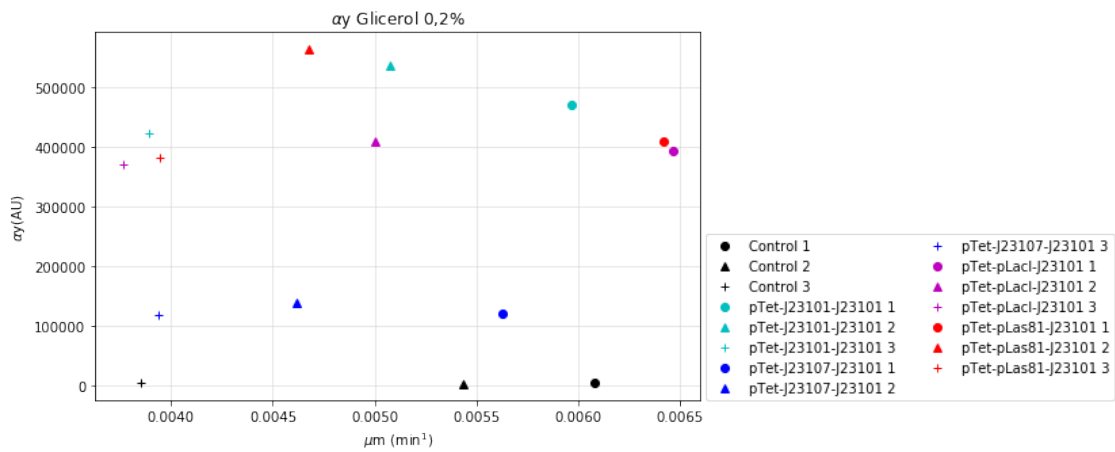
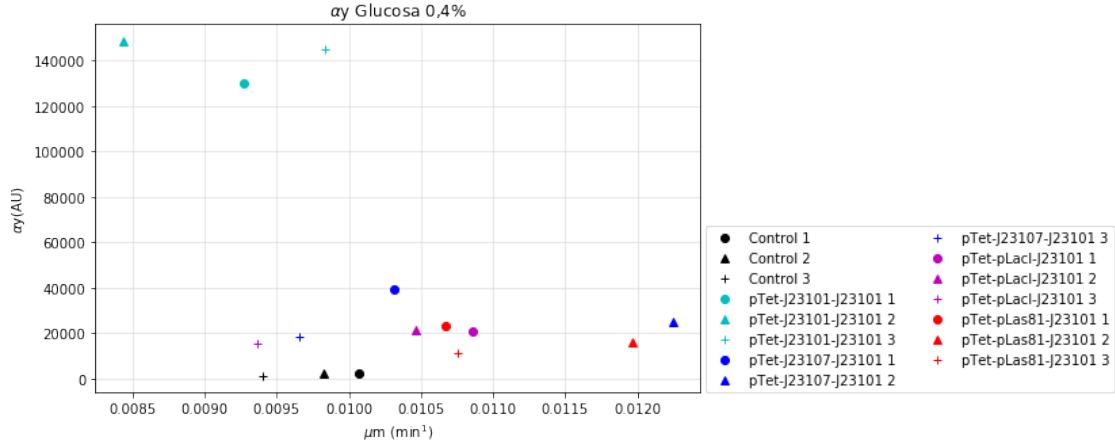
#grafico de ac versus Um
plt.figure(figsize=(8,5))
plt.title(r'$\alpha$y Glicerol 0,2%')
plt.xlabel(r'$\mu$m (min$^{-1}$)')
plt.ylabel(r'$\alpha$y(AU)')
plt.plot(um4,slopy4,'ko',label='Control 1')
plt.plot(um5,slopy5,'k^',label='Control 2')
plt.plot(um6,slopy6,'k+',label='Control 3')
```

```

plt.plot(um10,slopy10,'co',label='pTet-J23101-J23101 1')
plt.plot(um11,slopy11,'c^',label='pTet-J23101-J23101 2')
plt.plot(um12,slopy12,'c+',label='pTet-J23101-J23101 3')
plt.plot(um16,slopy16,'bo',label='pTet-J23107-J23101 1')
plt.plot(um17,slopy17,'b^',label='pTet-J23107-J23101 2')
plt.plot(um18,slopy18,'b+',label='pTet-J23107-J23101 3')
plt.plot(um22,slopy22,'mo',label='pTet-pLacI-J23101 1')
plt.plot(um23,slopy23,'m^',label='pTet-pLacI-J23101 2')
plt.plot(um24,slopy24,'m+',label='pTet-pLacI-J23101 3')
plt.plot(um28,slopy28,'ro',label='pTet-pLas81-J23101 1')
plt.plot(um29,slopy29,'r^',label='pTet-pLas81-J23101 2')
plt.plot(um30,slopy30,'r+',label='pTet-pLas81-J23101 3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

```

Out[74]: <matplotlib.legend.Legend at 0x267f1245f28>



```

In [75]: #grafico de ac versus Um
plt.figure(figsize=(8,5))
plt.title(r'$\alpha$c Ensayo 2 Réplica 1')
plt.xlabel(r'$\mu$m (min$^1$)')
plt.ylabel(r'$\alpha$c(AU)')
plt.plot(um1,slopec1,'k.',label='Control Glucosa')
plt.plot(um7,slopec7,'c.',label='pTet-J23101-J23101 Glucosa')
plt.plot(um13,slopec13,'c*',label='pTet-J23107-J23101 Glucosa')
plt.plot(um19,slopec19,'c+',label='pTet-pLacI-J23101 Glucosa')
plt.plot(um25,slopec25,'c^',label='pTet-pLas81-J23101 Glucosa')
plt.plot(um4,slopec4,'ko',label='Control Glicerol')
plt.plot(um10,slopec10,'b.',label='pTet-J23101-J23101 Glicerol')
plt.plot(um16,slopec16,'b*',label='pTet-J23107-J23101 Glicerol')
plt.plot(um22,slopec22,'b+',label='pTet-pLacI-J23101 Glicerol')
plt.plot(um28,slopec28,'b^',label='pTet-pLas81-J23101 Glicerol')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

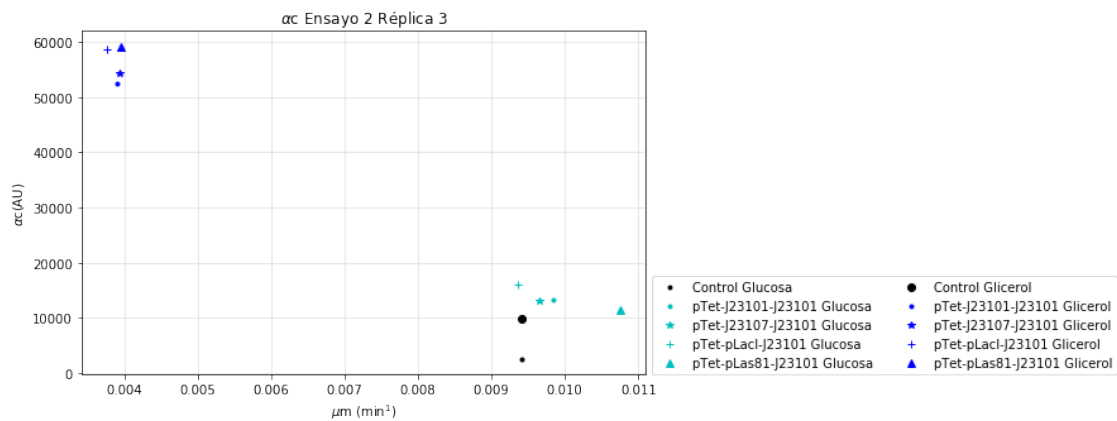
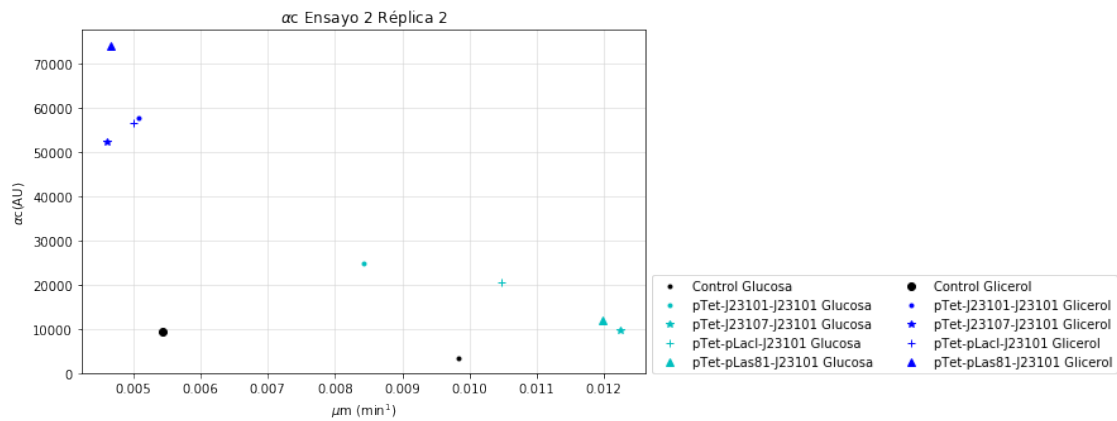
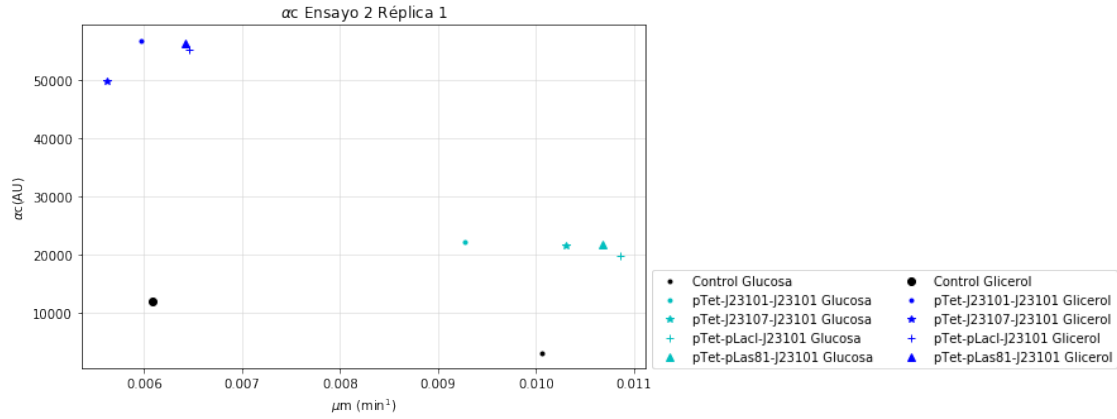
plt.figure(figsize=(8,5))
plt.title(r'$\alpha$c Ensayo 2 Réplica 2')
plt.xlabel(r'$\mu$m (min$^1$)')
plt.ylabel(r'$\alpha$c(AU)')
plt.plot(um2,slopec2,'k.',label='Control Glucosa')
plt.plot(um8,slopec8,'c.',label='pTet-J23101-J23101 Glucosa')
plt.plot(um14,slopec14,'c*',label='pTet-J23107-J23101 Glucosa')
plt.plot(um20,slopec20,'c+',label='pTet-pLacI-J23101 Glucosa')
plt.plot(um26,slopec26,'c^',label='pTet-pLas81-J23101 Glucosa')
plt.plot(um5,slopec5,'ko',label='Control Glicerol')
plt.plot(um11,slopec11,'b.',label='pTet-J23101-J23101 Glicerol')
plt.plot(um17,slopec17,'b*',label='pTet-J23107-J23101 Glicerol')
plt.plot(um23,slopec23,'b+',label='pTet-pLacI-J23101 Glicerol')
plt.plot(um29,slopec29,'b^',label='pTet-pLas81-J23101 Glicerol')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

plt.figure(figsize=(8,5))
plt.title(r'$\alpha$c Ensayo 2 Réplica 3')
plt.xlabel(r'$\mu$m (min$^1$)')
plt.ylabel(r'$\alpha$c(AU)')
plt.plot(um3,slopec3,'k.',label='Control Glucosa')
plt.plot(um9,slopec9,'c.',label='pTet-J23101-J23101 Glucosa')
plt.plot(um15,slopec15,'c*',label='pTet-J23107-J23101 Glucosa')
plt.plot(um21,slopec21,'c+',label='pTet-pLacI-J23101 Glucosa')
plt.plot(um27,slopec27,'c^',label='pTet-pLas81-J23101 Glucosa')
plt.plot(um3,slopec6,'ko',label='Control Glicerol')
plt.plot(um12,slopec12,'b.',label='pTet-J23101-J23101 Glicerol')
plt.plot(um18,slopec18,'b*',label='pTet-J23107-J23101 Glicerol')
plt.plot(um24,slopec24,'b+',label='pTet-pLacI-J23101 Glicerol')

```

```
plt.plot(um30,slopec30,'b^',label='pTet-pLas81-J23101 Glicerol')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)
```

Out[75]: <matplotlib.legend.Legend at 0x267f1139da0>



```

In [76]: #grafico de ac versus Um
plt.figure(figsize=(8,5))
plt.title(r'$\alpha$ Ensayo 2 Réplica 1')
plt.xlabel(r'$\mu$M (min$^1$)')
plt.ylabel(r'$\alpha$(AU)')
plt.plot(um1,sloper1,'k.',label='Control Glucosa ')
plt.plot(um7,sloper7,'r.',label='pTet-J23101-J23101 Glucosa ')
plt.plot(um13,sloper13,'r*',label='pTet-J23107-J23101 Glucosa ')
plt.plot(um19,sloper19,'r+',label='pTet-pLacI-J23101 Glucosa ')
plt.plot(um25,sloper25,'r^',label='pTet-pLas81-J23101 Glucosa ')
plt.plot(um4,sloper4,'ko',label='Control Glicerol ')
plt.plot(um10,sloper10,'m.',label='pTtet-J23101-J23101 Glicerol ')
plt.plot(um16,sloper16,'m*',label='pTtet-J231017-J23101 Glicerol')
plt.plot(um22,sloper22,'m+',label='pTtet-pLacI-J23101 Glicerol ')
plt.plot(um28,sloper28,'m^',label='pTtet-pLas81-J23101 Glicerol')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

plt.figure(figsize=(8,5))
plt.title(r'$\alpha$ Ensayo 2 Réplica 2')
plt.xlabel(r'$\mu$M (min$^1$)')
plt.ylabel(r'$\alpha$(AU)')
plt.plot(um2,sloper2,'k.',label='Control Glucosa')
plt.plot(um8,sloper8,'r.',label='pTet-J23101-J23101 Glucosa')
plt.plot(um14,sloper14,'r*',label='pTet-J23107-J23101 Glucosa')
plt.plot(um20,sloper20,'r+',label='pTet-pLacI-J23101 Glucosa')
plt.plot(um26,sloper26,'r^',label='pTet-pLas81-J23101 Glucosa')
plt.plot(um5,sloper5,'ko',label='Control Glicerol')
plt.plot(um11,sloper11,'m.',label='pTet-J23101-J23101 Glicerol')
plt.plot(um17,sloper17,'m*',label='pTet-J23107-J23101 Glicerol')
plt.plot(um23,sloper23,'m+',label='pTet-pLacI-J23101 Glicerol')
plt.plot(um29,sloper29,'m^',label='pTet-pLas81-J23101 Glicerol')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

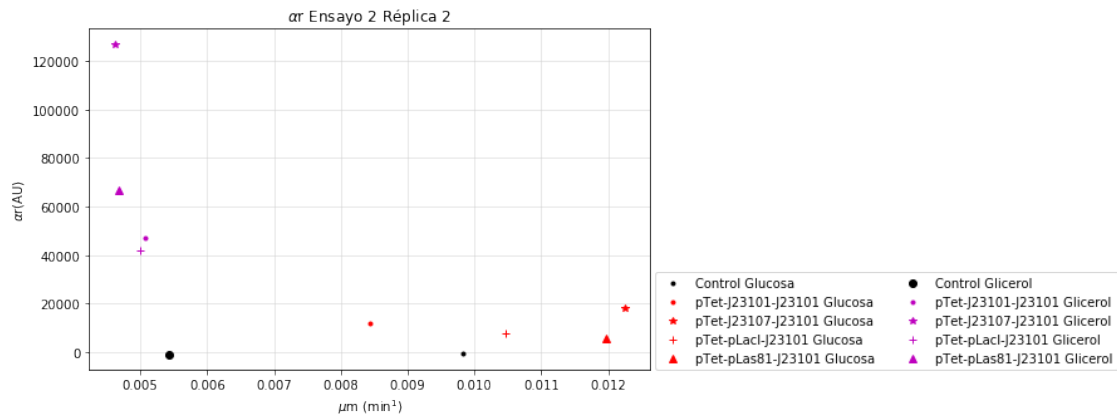
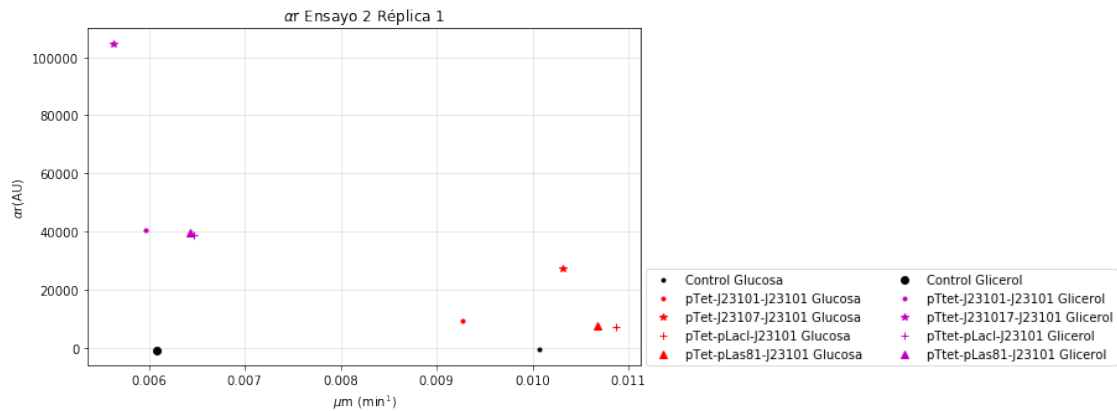
plt.figure(figsize=(8,5))
plt.title(r'$\alpha$ Ensayo 2 Réplica 3')
plt.xlabel(r'$\mu$M (min$^1$)')
plt.ylabel(r'$\alpha$(AU)')
plt.plot(um3,sloper3,'k.',label='Control Glucosa')
plt.plot(um9,sloper9,'r.',label='pTet-J23101-J23101 Glucosa')
plt.plot(um15,sloper15,'r*',label='pTet-J23107-J23101 Glucosa')
plt.plot(um21,sloper21,'r+',label='pTet-pLacI-J23101 Glucosa')
plt.plot(um27,sloper27,'r^',label='pTet-pLas81-J23101 Glucosa')
plt.plot(um6,sloper6,'ko',label='Control Glicerol')

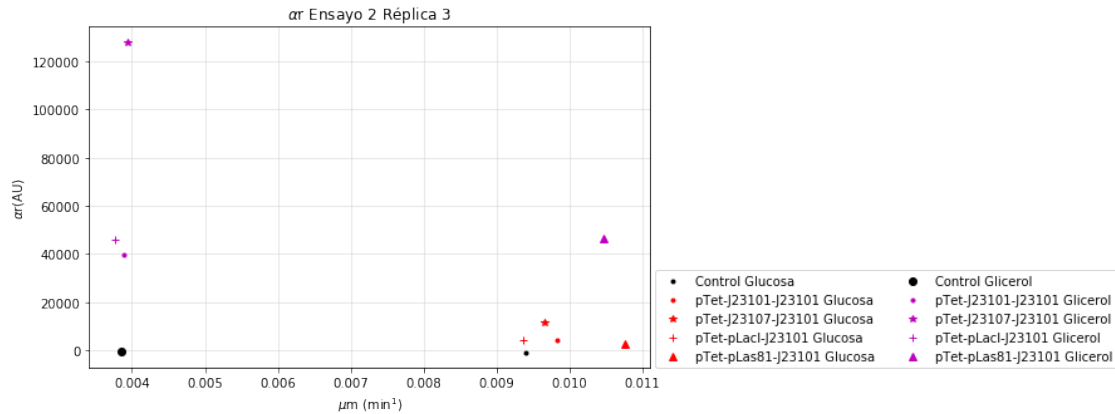
```



```
plt.plot(um12,sloper12,'m.',label='pTet-J23101-J23101 Glicerol')
plt.plot(um18,sloper18,'m*',label='pTet-J23107-J23101 Glicerol')
plt.plot(um24,sloper24,'m+',label='pTet-pLacI-J23101 Glicerol')
plt.plot(um20,sloper30,'m^',label='pTet-pLas81-J23101 Glicerol')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)
```

Out[76]: <matplotlib.legend.Legend at 0x267f391fd30>





In [77]: *#grafico de ac versus Um*

```
plt.figure(figsize=(8,5))
plt.title(r'$\alpha$ Ensayo 2 Réplica 1')
plt.xlabel(r'$\mu$ (min$^{-1}$)')
plt.ylabel(r'$\alpha$ (AU)')
plt.plot(um1,slopy1,'k.',label='Control Glucosa ')
plt.plot(um7,slopy7,'y.',label='pTet-J23101-J23101 Glucosa ')
plt.plot(um13,slopy13,'y*',label='pTet-J23107-J23101 Glucosa ')
plt.plot(um19,slopy19,'y+',label='pTet-pLacI-J23101 Glucosa ')
plt.plot(um25,slopy25,'y^',label='pTet-pLas81-J23101 Glucosa ')
plt.plot(um4,slopy4,'ko',label='Control Glicerol ')
plt.plot(um10,slopy10,'g.',label='pTet-J23101-J23101 Glicerol ')
plt.plot(um16,slopy16,'g*',label='pTet-J23107-J23101 Glicerol')
plt.plot(um22,slopy22,'g+',label='pTet-pLacI-J23101 Glicerol ')
plt.plot(um28,slopy28,'g^',label='pTet-pLas81-J23101 Glicerol')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)
```

```
plt.figure(figsize=(8,5))
plt.title(r'$\alpha$ Ensayo 2 Réplica 2')
plt.xlabel(r'$\mu$ (min$^{-1}$)')
plt.ylabel(r'$\alpha$ (AU)')
plt.plot(um2,slopy2,'k.',label='Control Glucosa')
plt.plot(um8,slopy8,'y.',label='pTet-J23101-J23101 Glucosa')
plt.plot(um14,slopy14,'y*',label='pTet-J23107-J23101 Glucosa')
plt.plot(um20,slopy20,'y+',label='pTet-pLacI-J23101 Glucosa')
plt.plot(um26,slopy26,'y^',label='pTet-pLas81-J23101 Glucosa')
plt.plot(um5,slopy5,'ko',label='Control Glicerol')
plt.plot(um11,slopy11,'g.',label='pTet-J23101-J23101 Glicerol')
plt.plot(um17,slopy17,'g*',label='pTet-J23107-J23101 Glicerol')
plt.plot(um23,slopy23,'g+',label='pTet-pLacI-J23101 Glicerol')
plt.plot(um29,slopy29,'g^',label='pTet-pLas81-J23101 Glicerol')
```

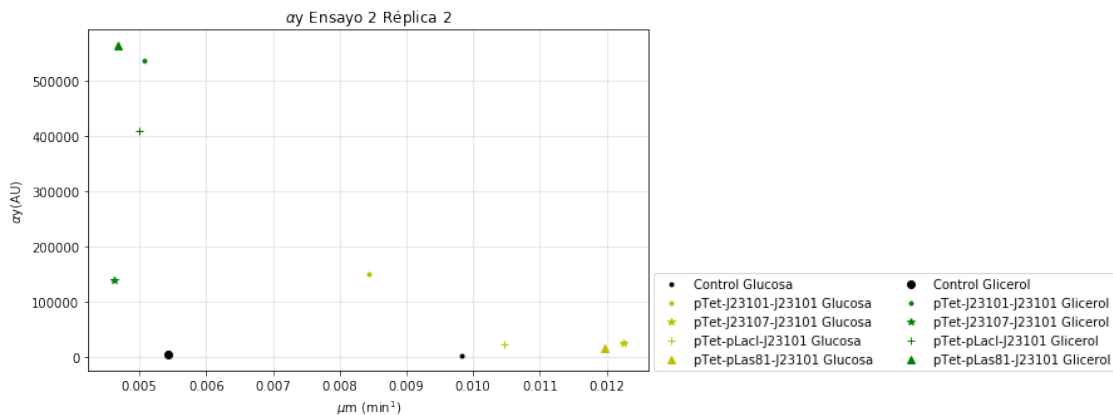
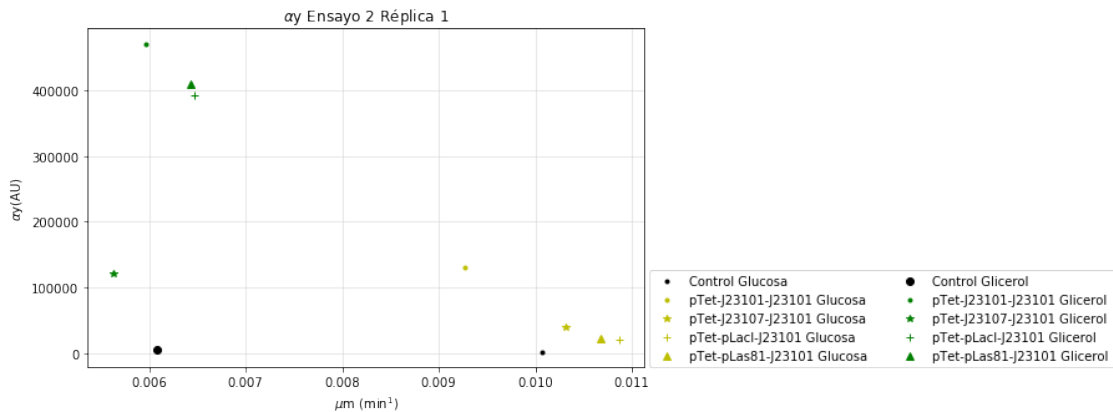
```

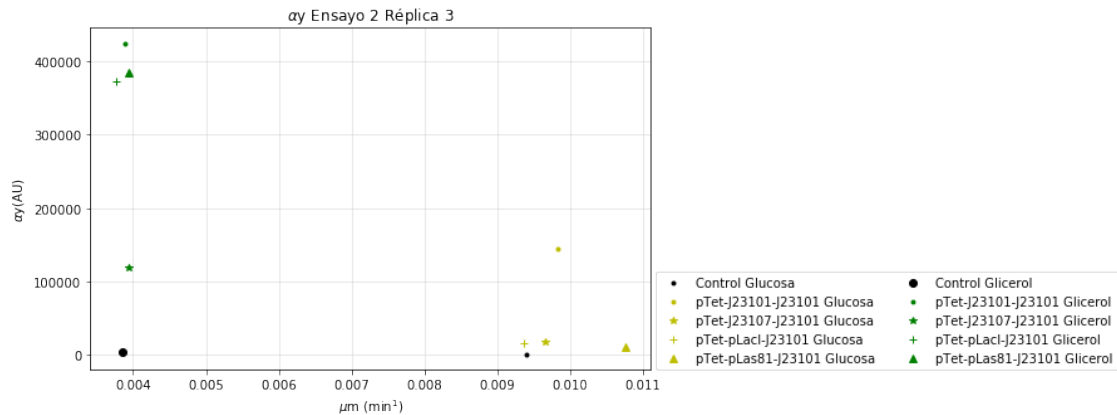
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

plt.figure(figsize=(8,5))
plt.title(r'$\alpha$ Ensayo 2 Réplica 3')
plt.xlabel(r'$\mu$ (min$^{-1}$)')
plt.ylabel(r'$\alpha$ (AU)')
plt.plot(um3,slopey3,'k.',label='Control Glucosa')
plt.plot(um9,slopey9,'y.',label='pTet-J23101-J23101 Glucosa')
plt.plot(um15,slopey15,'y*',label='pTet-J23107-J23101 Glucosa')
plt.plot(um21,slopey21,'y+',label='pTet-pLacI-J23101 Glucosa')
plt.plot(um27,slopey27,'y^',label='pTet-pLas81-J23101 Glucosa')
plt.plot(um6,slopey6,'ko',label='Control Glicerol')
plt.plot(um12,slopey12,'g.',label='pTet-J23101-J23101 Glicerol')
plt.plot(um18,slopey18,'g*',label='pTet-J23107-J23101 Glicerol')
plt.plot(um24,slopey24,'g+',label='pTet-pLacI-J23101 Glicerol')
plt.plot(um30,slopey30,'g^',label='pTet-pLas81-J23101 Glicerol')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

```

Out[77]: <matplotlib.legend.Legend at 0x267eef7dc50>





```
In [78]: #grafico de ar vs ac
plt.figure(figsize=(8,5))
plt.title(r'$\alpha$y Glucosa 0,4%')
plt.xlabel(r'$\alpha$c (AU)')
plt.ylabel(r'$\alpha$y(AU)')
plt.plot(slopec1,slopy1,'ko',label='Control 1')
plt.plot(slopec2,slopy2,'k^',label='Control 2')
plt.plot(slopec3,slopy3,'k+',label='Control 3')
plt.plot(slopec7,slopy7,'co',label='pTet-J23101-J23101 1')
plt.plot(slopec8,slopy8,'c^',label='pTet-J23101-J23101 2')
plt.plot(slopec9,slopy9,'c+',label='pTet-J23101-J23101 3')
plt.plot(slopec13,slopy13,'bo',label='pTet-J23107-J23101 1')
plt.plot(slopec14,slopy14,'b^',label='pTet-J23107-J23101 2')
plt.plot(slopec15,slopy15,'b+',label='pTet-J23107-J23101 3')
plt.plot(slopec19,slopy19,'mo',label='pTet-pLacI-J23101 1')
plt.plot(slopec20,slopy20,'m^',label='pTet-pLacI-J23101 2')
plt.plot(slopec21,slopy21,'m+',label='pTet-pLacI-J23101 3')
plt.plot(slopec25,slopy25,'ro',label='pTet-pLas81-J23101 1')
plt.plot(slopec26,slopy26,'r^',label='pTet-pLas81-J23101 2')
plt.plot(slopec27,slopy27,'r+',label='pTet-pLas81-J23101 3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

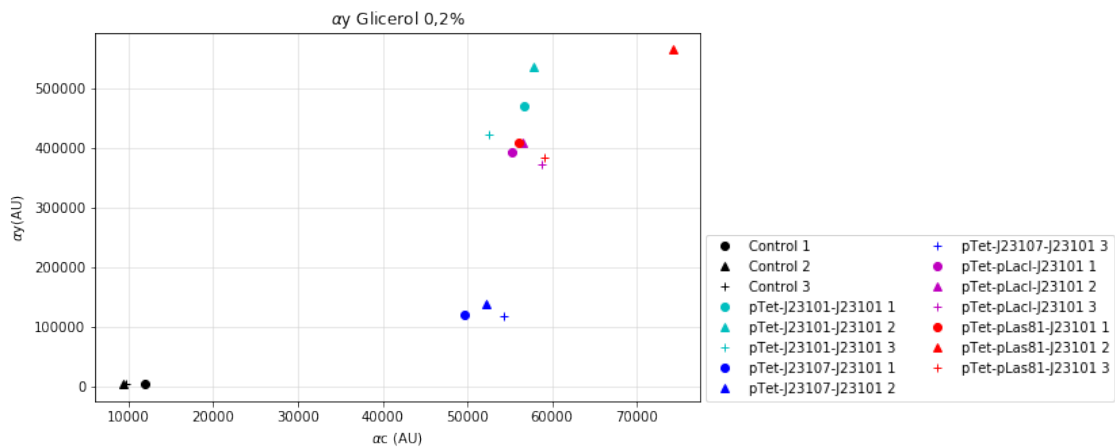
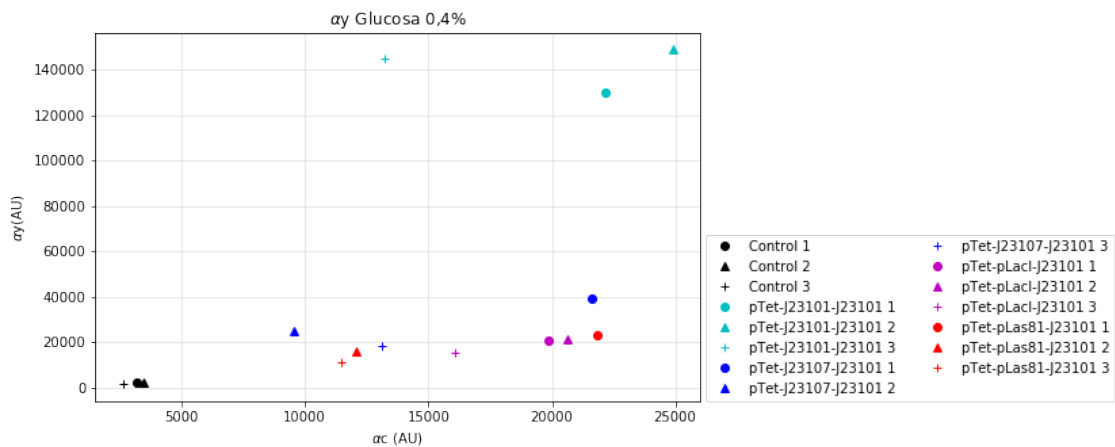
#grafico de ay vs ac
plt.figure(figsize=(8,5))
plt.title(r'$\alpha$y Glicerol 0,2%')
plt.xlabel(r'$\alpha$c (AU)')
plt.ylabel(r'$\alpha$y(AU)')
plt.plot(slopec4,slopy4,'ko',label='Control 1')
plt.plot(slopec5,slopy5,'k^',label='Control 2')
```

```

plt.plot(slopec6,slopy6,'k+',label='Control 3')
plt.plot(slopec10,slopy10,'co',label='pTet-J23101-J23101 1')
plt.plot(slopec11,slopy11,'c^',label='pTet-J23101-J23101 2')
plt.plot(slopec12,slopy12,'c+',label='pTet-J23101-J23101 3')
plt.plot(slopec16,slopy16,'bo',label='pTet-J23107-J23101 1')
plt.plot(slopec17,slopy17,'b^',label='pTet-J23107-J23101 2')
plt.plot(slopec18,slopy18,'b+',label='pTet-J23107-J23101 3')
plt.plot(slopec22,slopy22,'mo',label='pTet-pLacI-J23101 1')
plt.plot(slopec23,slopy23,'m^',label='pTet-pLacI-J23101 2')
plt.plot(slopec24,slopy24,'m+',label='pTet-pLacI-J23101 3')
plt.plot(slopec28,slopy28,'ro',label='pTet-pLas81-J23101 1')
plt.plot(slopec29,slopy29,'r^',label='pTet-pLas81-J23101 2')
plt.plot(slopec30,slopy30,'r+',label='pTet-pLas81-J23101 3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

```

Out[78]: <matplotlib.legend.Legend at 0x267f2006c50>



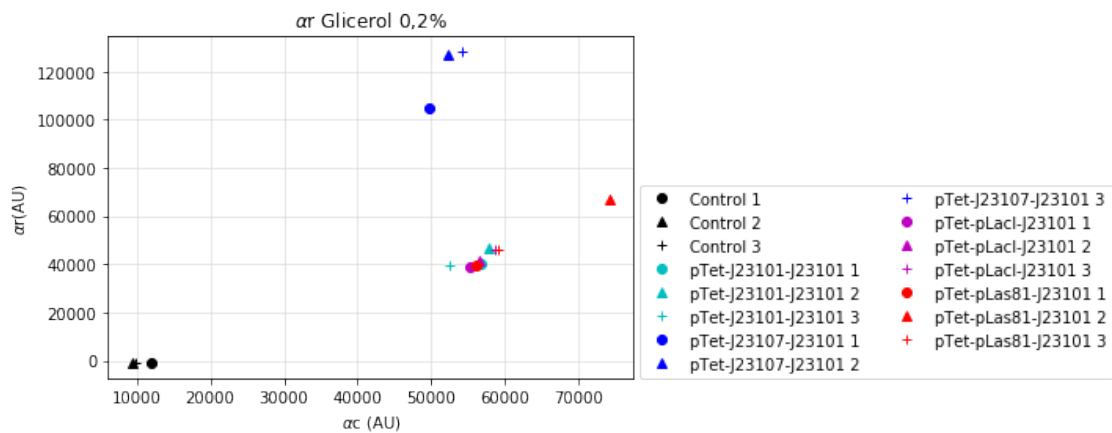
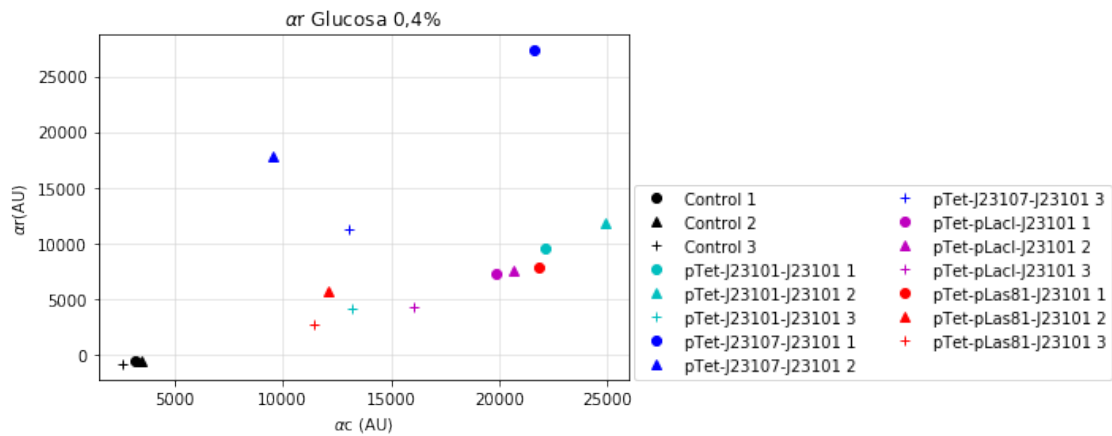
```

In [79]: #grafico de ac versus Um
plt.figure()
plt.title(r'$\alpha$r Glucosa 0,4%')
plt.xlabel(r'$\alpha$c (AU)')
plt.ylabel(r'$\alpha$r(AU)')
plt.plot(slopec1,sloper1,'ko',label='Control 1')
plt.plot(slopec2,sloper2,'k^',label='Control 2')
plt.plot(slopec3,sloper3,'k+',label='Control 3')
plt.plot(slopec7,sloper7,'co',label='pTet-J23101-J23101 1')
plt.plot(slopec8,sloper8,'c^',label='pTet-J23101-J23101 2')
plt.plot(slopec9,sloper9,'c+',label='pTet-J23101-J23101 3')
plt.plot(slopec13,sloper13,'bo',label='pTet-J23107-J23101 1')
plt.plot(slopec14,sloper14,'b^',label='pTet-J23107-J23101 2')
plt.plot(slopec15,sloper15,'b+',label='pTet-J23107-J23101 3')
plt.plot(slopec19,sloper19,'mo',label='pTet-pLacI-J23101 1')
plt.plot(slopec20,sloper20,'m^',label='pTet-pLacI-J23101 2')
plt.plot(slopec21,sloper21,'m+',label='pTet-pLacI-J23101 3')
plt.plot(slopec25,sloper25,'ro',label='pTet-pLas81-J23101 1')
plt.plot(slopec26,sloper26,'r^',label='pTet-pLas81-J23101 2')
plt.plot(slopec27,sloper27,'r+',label='pTet-pLas81-J23101 3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

#grafico de ac versus Um
plt.figure()
plt.title(r'$\alpha$r Glicerol 0,2%')
plt.xlabel(r'$\alpha$c (AU)')
plt.ylabel(r'$\alpha$r(AU)')
plt.plot(slopec4,sloper4,'ko',label='Control 1')
plt.plot(slopec5,sloper5,'k^',label='Control 2')
plt.plot(slopec6,sloper6,'k+',label='Control 3')
plt.plot(slopec10,sloper10,'co',label='pTet-J23101-J23101 1')
plt.plot(slopec11,sloper11,'c^',label='pTet-J23101-J23101 2')
plt.plot(slopec12,sloper12,'c+',label='pTet-J23101-J23101 3')
plt.plot(slopec16,sloper16,'bo',label='pTet-J23107-J23101 1')
plt.plot(slopec17,sloper17,'b^',label='pTet-J23107-J23101 2')
plt.plot(slopec18,sloper18,'b+',label='pTet-J23107-J23101 3')
plt.plot(slopec22,sloper22,'mo',label='pTet-pLacI-J23101 1')
plt.plot(slopec23,sloper23,'m^',label='pTet-pLacI-J23101 2')
plt.plot(slopec24,sloper24,'m+',label='pTet-pLacI-J23101 3')
plt.plot(slopec28,sloper28,'ro',label='pTet-pLas81-J23101 1')
plt.plot(slopec29,sloper29,'r^',label='pTet-pLas81-J23101 2')
plt.plot(slopec30,sloper30,'r+',label='pTet-pLas81-J23101 3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)

```

Out[79]: <matplotlib.legend.Legend at 0x267f34fddd8>



In [80]: #Grafico de barras um de FPs

```
uglu=[um1,um2,um3,um7,um8,um9,um13,um14,um15,um19,um20,um21,um25,um26,um27]
ugli=[um4,um5,um6,um10,um11,um12,um16,um17,um18,um22,um23,um24,um28,um29,um30]
```

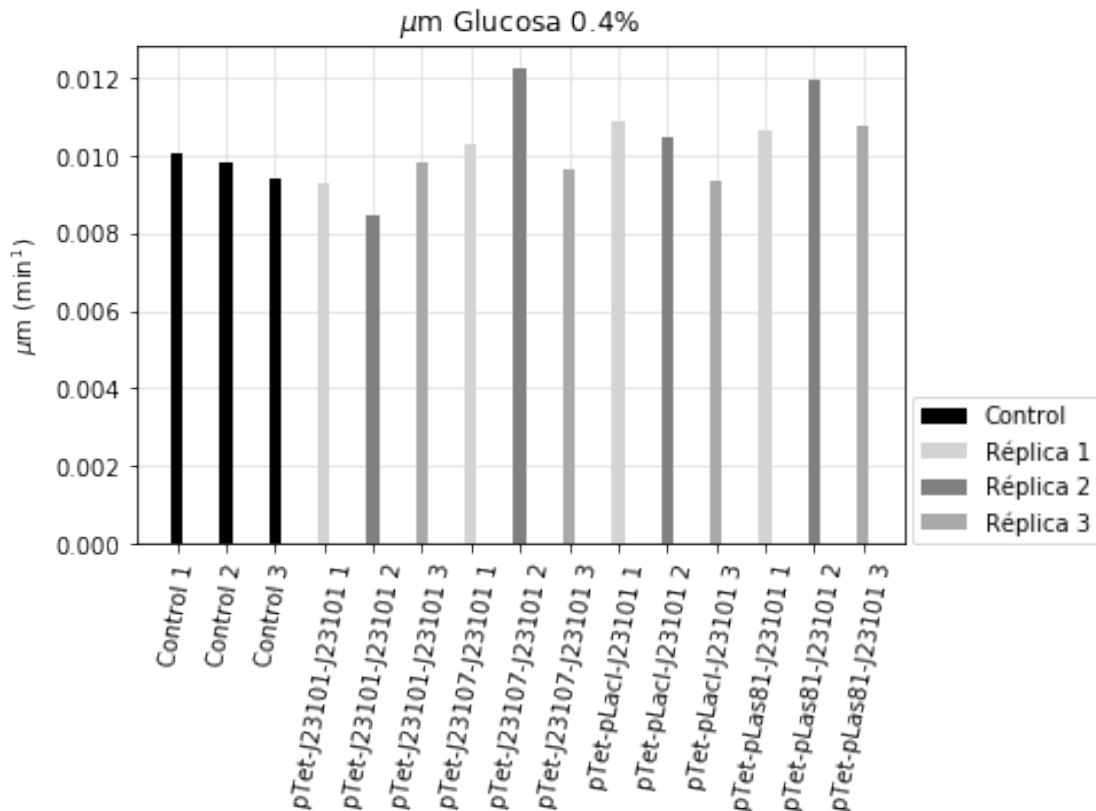
```
X = np.arange(15)
plt.figure()
plt.title(r'$\mu$ Glucosa 0.4%')
plt.ylabel(r'$\mu$ (min$^1$)')
plt.bar(X[0]+0.00,uglu[0],color='k',width=0.25,label='Control',zorder=3)
plt.bar(X[1]+0.00,uglu[1],color='k',width=0.25,zorder=3)
plt.bar(X[2]+0.00,uglu[2],color='k',width=0.25,zorder=3)
plt.bar(X[3]+0.00,uglu[3],color='lightgrey',width=0.25,label='Réplica 1',zorder=3)
plt.bar(X[4]+0.00,uglu[4],color='grey',width=0.25,label='Réplica 2',zorder=3)
plt.bar(X[5]+0.00,uglu[5],color='darkgrey',width=0.25,label='Réplica 3',zorder=3)
```

```

plt.bar(X[6]+0.00,uglu[6],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[7]+0.00,uglu[7],color='grey',width=0.25,zorder=3)
plt.bar(X[8]+0.00,uglu[8],color='darkgrey',width=0.25,zorder=3)
plt.bar(X[9]+0.00,uglu[9],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[10]+0.00,uglu[10],color='grey',width=0.25,zorder=3)
plt.bar(X[11]+0.00,uglu[11],color='darkgrey',width=0.25,zorder=3)
plt.bar(X[12]+0.00,uglu[12],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[13]+0.00,uglu[13],color='grey',width=0.25,zorder=3)
plt.bar(X[14]+0.00,uglu[14],color='darkgrey',width=0.25,zorder=3)
plt.xticks(X,['Control 1','Control 2','Control 3','pTet-J23101-J23101 1','pTet-J23101-J23101 2','pTet-J23101-J23101 3','pTet-J23107-J23107 1','pTet-J23107-J23107 2','pTet-J23107-J23107 3','pTet-pLacJ-J23101 1','pTet-pLacJ-J23101 2','pTet-pLacJ-J23101 3','pTet-pLas81-J23101 1','pTet-pLas81-J23101 2','pTet-pLas81-J23101 3'])
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0))

```

Out[80]: <matplotlib.legend.Legend at 0x267f40c35f8>



```

In [81]: X = np.arange(15)
plt.figure()
plt.title(r'$\mu$ Glicerol 0.2%')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.bar(X[0]+0.00,ugli[0],color='k',width=0.25,label='Control',zorder=3)
plt.bar(X[1]+0.00,ugli[1],color='k',width=0.25,zorder=3)

```

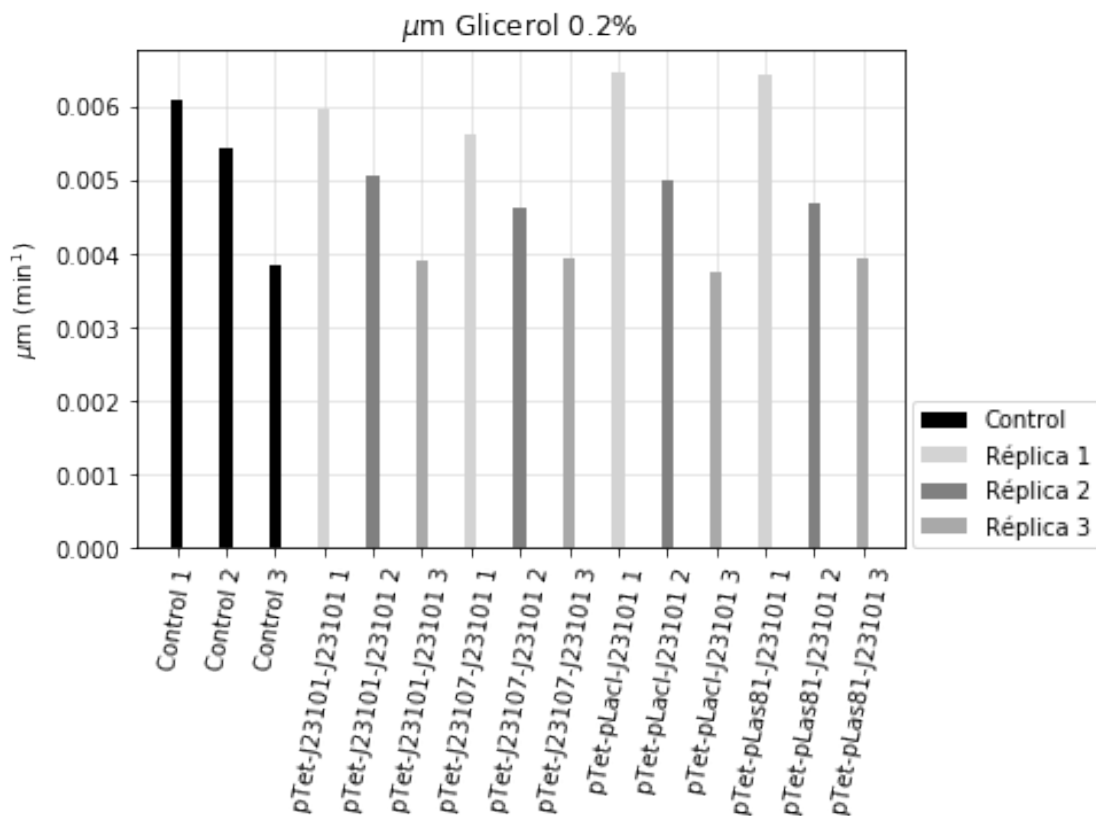


```

plt.bar(X[2]+0.00,ugli[2],color='k',width=0.25,zorder=3)
plt.bar(X[3]+0.00,ugli[3],color='lightgrey',width=0.25,label='Réplica 1',zorder=3)
plt.bar(X[4]+0.00,ugli[4],color='grey',width=0.25,label='Réplica 2',zorder=3)
plt.bar(X[5]+0.00,ugli[5],color='darkgrey',width=0.25,label='Réplica 3',zorder=3)
plt.bar(X[6]+0.00,ugli[6],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[7]+0.00,ugli[7],color='grey',width=0.25,zorder=3)
plt.bar(X[8]+0.00,ugli[8],color='darkgrey',width=0.25,zorder=3)
plt.bar(X[9]+0.00,ugli[9],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[10]+0.00,ugli[10],color='grey',width=0.25,zorder=3)
plt.bar(X[11]+0.00,ugli[11],color='darkgrey',width=0.25,zorder=3)
plt.bar(X[12]+0.00,ugli[12],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[13]+0.00,ugli[13],color='grey',width=0.25,zorder=3)
plt.bar(X[14]+0.00,ugli[14],color='darkgrey',width=0.25,zorder=3)
plt.xticks(X,['Control 1','Control 2','Control 3','pTet-J23101-J23101 1','pTet-J23101-J23101 2','pTet-J23101-J23101 3','pTet-J23107-J23107 1','pTet-J23107-J23107 2','pTet-J23107-J23107 3','pTet-pLacI-J23101 1','pTet-pLacI-J23101 2','pTet-pLacI-J23101 3','pTet-pLas81-J23101 1','pTet-pLas81-J23101 2','pTet-pLas81-J23101 3'])
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0))

```

Out [81]: <matplotlib.legend.Legend at 0x267f42b1550>



```

In [82]: X = np.arange(15)
plt.figure(figsize=(8,5))

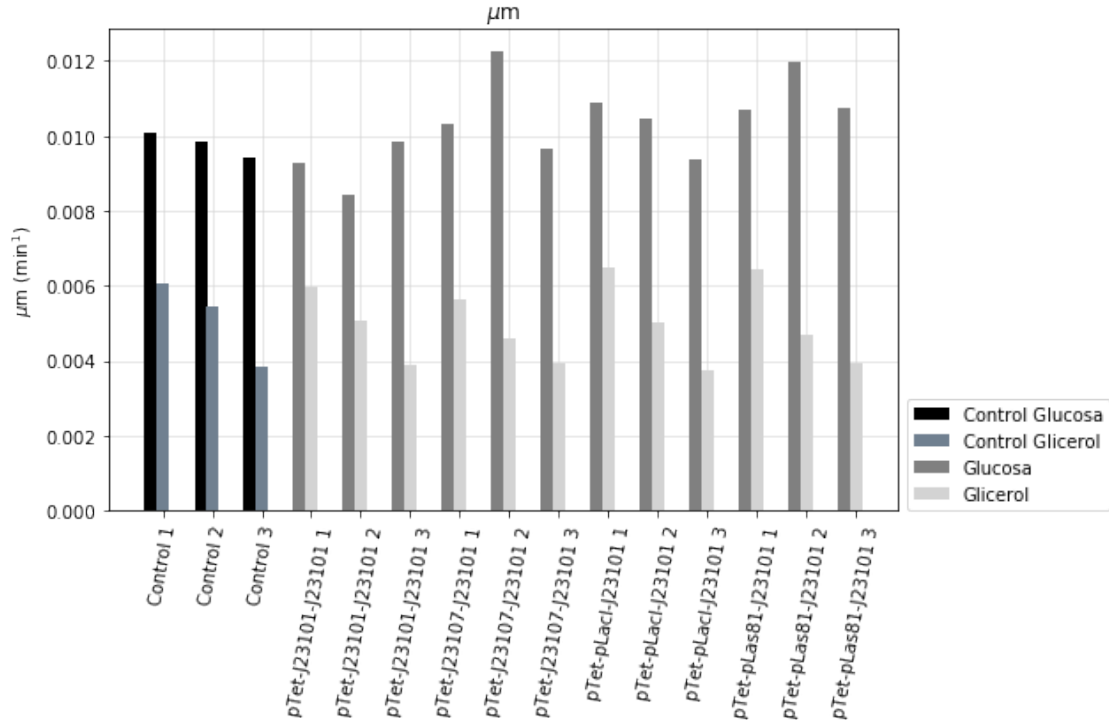
```

```

plt.title(r'$\mu$M')
plt.ylabel(r'$\mu$M (min$^{-1}$)')
plt.bar(X[0]-0.25,uglu[0],color='k',width=0.25,label='Control Glucosa',zorder=3)
plt.bar(X[0]+0.00,ugli[0],color='slategrey',width=0.25,label='Control Glicerol',zorder=3)
plt.bar(X[1]-0.20,uglu[1],color='k',width=0.25,zorder=3)
plt.bar(X[1]+0.00,ugli[1],color='slategrey',width=0.25,zorder=3)
plt.bar(X[2]-0.25,uglu[2],color='k',width=0.25,zorder=3)
plt.bar(X[2]+0.00,ugli[2],color='slategrey',width=0.25,zorder=3)
plt.bar(X[3]-0.25,uglu[3],color='grey',width=0.25,label='Glucosa',zorder=3)
plt.bar(X[3]+0.00,ugli[3],color='lightgrey',width=0.25,label='Glicerol',zorder=3)
plt.bar(X[4]-0.25,uglu[4],color='grey',width=0.25,zorder=3)
plt.bar(X[4]+0.00,ugli[4],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[5]-0.25,uglu[5],color='grey',width=0.25,zorder=3)
plt.bar(X[5]+0.00,ugli[5],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[6]-0.25,uglu[6],color='grey',width=0.25,zorder=3)
plt.bar(X[6]+0.00,ugli[6],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[7]-0.25,uglu[7],color='grey',width=0.25,zorder=3)
plt.bar(X[7]+0.00,ugli[7],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[8]-0.25,uglu[8],color='grey',width=0.25,zorder=3)
plt.bar(X[8]+0.00,ugli[8],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[9]-0.25,uglu[9],color='grey',width=0.25,zorder=3)
plt.bar(X[9]+0.00,ugli[9],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[10]-0.25,uglu[10],color='grey',width=0.25,zorder=3)
plt.bar(X[10]+0.00,ugli[10],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[11]-0.25,uglu[11],color='grey',width=0.25,zorder=3)
plt.bar(X[11]+0.00,ugli[11],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[12]-0.25,uglu[12],color='grey',width=0.25,zorder=3)
plt.bar(X[12]+0.00,ugli[12],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[13]-0.25,uglu[13],color='grey',width=0.25,zorder=3)
plt.bar(X[13]+0.00,ugli[13],color='lightgrey',width=0.25,zorder=3)
plt.bar(X[14]-0.25,uglu[14],color='grey',width=0.25,zorder=3)
plt.bar(X[14]+0.00,ugli[14],color='lightgrey',width=0.25,zorder=3)
plt.xticks(X,['Control 1','Control 2','Control 3','pTet-J23101-J23101 1','pTet-J23101-J23101 2'])
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0))

```

Out[82]: <matplotlib.legend.Legend at 0x267f445a780>



In [83]: *#Ro RFP*

```

pr1=sloper1/slopec1
pr2=sloper2/slopec2
pr3=sloper3/slopec3
pr4=sloper4/slopec4
pr5=sloper5/slopec5
pr6=sloper6/slopec6
pr7=sloper7/slopec7
pr8=sloper8/slopec8
pr9=sloper9/slopec9
pr10=sloper10/slopec10
pr11=sloper11/slopec11
pr12=sloper12/slopec12
pr13=sloper13/slopec13
pr14=sloper14/slopec14
pr15=sloper15/slopec15
pr16=sloper16/slopec16
pr17=sloper17/slopec17
pr18=sloper18/slopec18
pr19=sloper19/slopec19
pr20=sloper20/slopec20
pr21=sloper21/slopec21
pr22=sloper22/slopec22
pr23=sloper23/slopec23

```

```

pr24=sloper24/slopec24
pr25=sloper25/slopec25
pr26=sloper26/slopec26
pr27=sloper27/slopec27
pr28=sloper28/slopec28
pr29=sloper29/slopec29
pr30=sloper30/slopec30

```

```

ro_rfp=[pr1,pr2,pr3,pr4,pr5,pr6,pr7,pr8,pr9,pr10,pr11,pr12,pr13,pr14,pr15,pr16,pr17,pr18,pr19,pr20,pr21,pr22,pr23,pr24,pr25,pr26,pr27,pr28,pr29,pr30]
ro_rfpglu=[[pr1,pr2,pr3],[pr7,pr8,pr9],[pr13,pr14,pr15],[pr19,pr20,pr21],[pr25,pr26,pr27],[pr28,pr29,pr30]]
ro_rfpgli=[[pr4,pr5,pr6],[pr10,pr11,pr12],[pr16,pr17,pr18],[pr22,pr23,pr24],[pr28,pr29,pr30]]

```

In [84]: *#Ro YFP*

```

py1=slopey1/slopec1
py2=slopey2/slopec2
py3=slopey3/slopec3
py4=slopey4/slopec4
py5=slopey5/slopec5
py6=slopey6/slopec6
py7=slopey7/slopec7
py8=slopey8/slopec8
py9=slopey9/slopec9
py10=slopey10/slopec10
py11=slopey11/slopec11
py12=slopey12/slopec12
py13=slopey13/slopec13
py14=slopey14/slopec14
py15=slopey15/slopec15
py16=slopey16/slopec16
py17=slopey17/slopec17
py18=slopey18/slopec18
py19=slopey19/slopec19
py20=slopey20/slopec20
py21=slopey21/slopec21
py22=slopey22/slopec22
py23=slopey23/slopec23
py24=slopey24/slopec24
py25=slopey25/slopec25
py26=slopey26/slopec26
py27=slopey27/slopec27
py28=slopey28/slopec28
py29=slopey29/slopec29
py30=slopey30/slopec30

```

```

ro_yfp=[py1,py2,py3,py4,py5,py6,py7,py8,py9,py10,py11,py12,py13,py14,py15,py16,py17,py18,py19,py20,py21,py22,py23,py24,py25,py26,py27,py28,py29,py30]
ro_yfpglu=[[py1,py2,py3],[py7,py8,py9],[py13,py14,py15],[py19,py20,py21],[py25,py26,py27],[py28,py29,py30]]

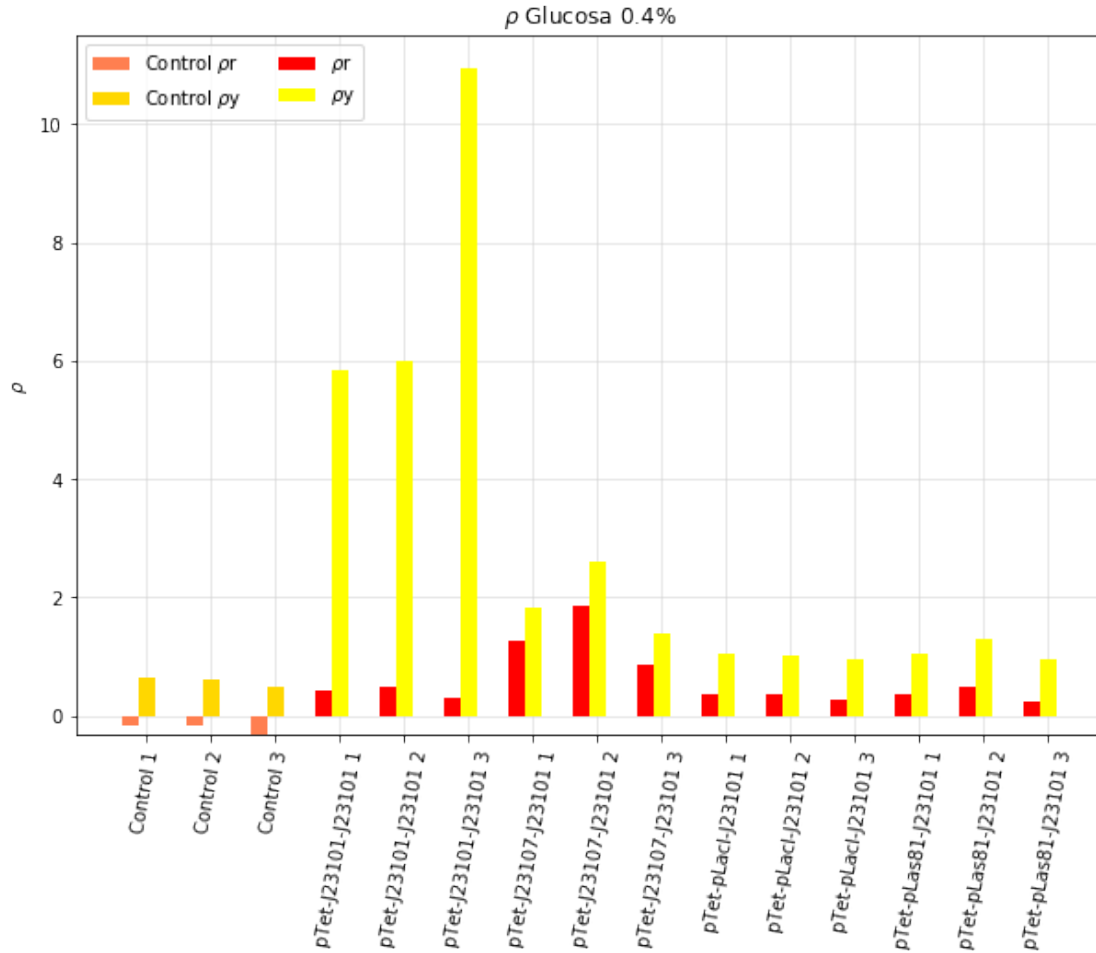
```

```
ro_yfpqli=[py4,py5,py6],[py10,py11,py12],[py16,py17,py18],[py22,py23,py24],[py28,py29,
```

```
In [85]: #grafico ro de yfp y de rfp
ro_rfp=[pr1,pr2,pr3,pr4,pr5,pr6,pr7,pr8,pr9,pr10,pr11,pr12,pr13,pr14,pr15,pr16,pr17,pr18,pr19,pr20,pr21,pr22,pr23,pr24,pr25,pr26,pr27,pr28,pr29,pr30,pr31,pr32,pr33,pr34,pr35,pr36,pr37,pr38,pr39,pr40,pr41,pr42,pr43,pr44,pr45,pr46,pr47,pr48,pr49,pr50,pr51,pr52,pr53,pr54,pr55,pr56,pr57,pr58,pr59,pr60,pr61,pr62,pr63,pr64,pr65,pr66,pr67,pr68,pr69,pr70,pr71,pr72,pr73,pr74,pr75,pr76,pr77,pr78,pr79,pr80,pr81,pr82,pr83,pr84,pr85,pr86,pr87,pr88,pr89,pr90,pr91,pr92,pr93,pr94,pr95,pr96,pr97,pr98,pr99,pr100]
ro_yfp=[py1,py2,py3,py4,py5,py6,py7,py8,py9,py10,py11,py12,py13,py14,py15,py16,py17,py18,py19,py20,py21,py22,py23,py24,py25,py26,py27,py28,py29,py30,py31,py32,py33,py34,py35,py36,py37,py38,py39,py40,py41,py42,py43,py44,py45,py46,py47,py48,py49,py50,py51,py52,py53,py54,py55,py56,py57,py58,py59,py60,py61,py62,py63,py64,py65,py66,py67,py68,py69,py70,py71,py72,py73,py74,py75,py76,py77,py78,py79,py80,py81,py82,py83,py84,py85,py86,py87,py88,py89,py90,py91,py92,py93,py94,py95,py96,py97,py98,py99,py100]

X = np.arange(15)
plt.figure(figsize=(10,7))
plt.title(r'$\rho$ Glucosa 0.4%')
plt.ylabel(r'$\rho$')
plt.bar(X[0]-0.25,ro_rfp[0],color='coral',width=0.25,label= 'Control'+ ' '+r'$\rho$r',zorder=3)
plt.bar(X[0]+0.00,ro_yfp[0],color='gold',width=0.25,label= 'Control'+ ' '+r'$\rho$y',zorder=3)
plt.bar(X[1]-0.25,ro_rfp[1],color='coral',width=0.25,zorder=3)
plt.bar(X[1]+0.00,ro_yfp[1],color='gold',width=0.25,zorder=3)
plt.bar(X[2]-0.25,ro_rfp[2],color='coral',width=0.25,zorder=3)
plt.bar(X[2]+0.00,ro_yfp[2],color='gold',width=0.25,zorder=3)
plt.bar(X[3]-0.25,ro_rfp[6],color='r',width=0.25,label=r'$\rho$r',zorder=3)
plt.bar(X[3]+0.00,ro_yfp[6],color='yellow',width=0.25,label=r'$\rho$y',zorder=3)
plt.bar(X[4]-0.25,ro_rfp[7],color='r',width=0.25,zorder=3)
plt.bar(X[4]+0.00,ro_yfp[7],color='yellow',width=0.25,zorder=3)
plt.bar(X[5]-0.25,ro_rfp[8],color='r',width=0.25,zorder=3)
plt.bar(X[5]+0.00,ro_yfp[8],color='yellow',width=0.25,zorder=3)
plt.bar(X[6]-0.25,ro_rfp[12],color='r',width=0.25,zorder=3)
plt.bar(X[6]+0.00,ro_yfp[12],color='yellow',width=0.25,zorder=3)
plt.bar(X[7]-0.25,ro_rfp[13],color='r',width=0.25,zorder=3)
plt.bar(X[7]+0.00,ro_yfp[13],color='yellow',width=0.25,zorder=3)
plt.bar(X[8]-0.25,ro_rfp[14],color='r',width=0.25,zorder=3)
plt.bar(X[8]+0.00,ro_yfp[14],color='yellow',width=0.25,zorder=3)
plt.bar(X[9]-0.25,ro_rfp[18],color='r',width=0.25,zorder=3)
plt.bar(X[9]+0.00,ro_yfp[18],color='yellow',width=0.25,zorder=3)
plt.bar(X[10]-0.25,ro_rfp[19],color='r',width=0.25,zorder=3)
plt.bar(X[10]+0.00,ro_yfp[19],color='yellow',width=0.25,zorder=3)
plt.bar(X[11]-0.25,ro_rfp[20],color='r',width=0.25,zorder=3)
plt.bar(X[11]+0.00,ro_yfp[20],color='yellow',width=0.25,zorder=3)
plt.bar(X[12]-0.25,ro_rfp[24],color='r',width=0.25,zorder=3)
plt.bar(X[12]+0.00,ro_yfp[24],color='yellow',width=0.25,zorder=3)
plt.bar(X[13]-0.25,ro_rfp[25],color='r',width=0.25,zorder=3)
plt.bar(X[13]+0.00,ro_yfp[25],color='yellow',width=0.25,zorder=3)
plt.bar(X[14]-0.25,ro_rfp[26],color='r',width=0.25,zorder=3)
plt.bar(X[14]+0.00,ro_yfp[26],color='yellow',width=0.25,zorder=3)
plt.xticks(X,['Control 1','Control 2','Control 3','pTet-J23101-J23101 1','pTet-J23101-J23101 2'])
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper left',ncol=2)
```

```
Out[85]: <matplotlib.legend.Legend at 0x267f1116c88>
```



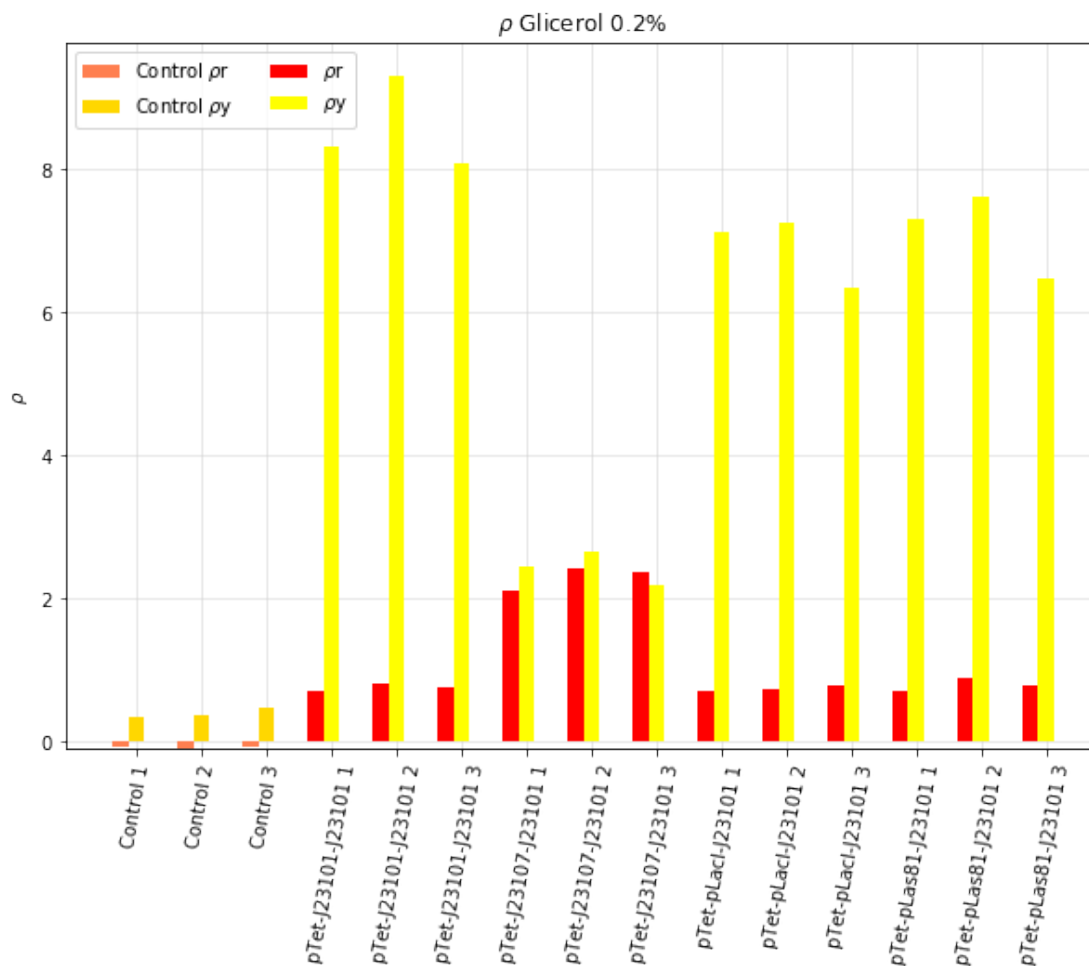
```
In [86]: X = np.arange(15)
plt.figure(figsize=(10,7))
plt.title(r'$\rho$ Glicerol 0.2%')
plt.ylabel(r'$\rho$')
plt.bar(X[0]-0.25,ro_rfp[3],color='coral',width=0.25,label= 'Control'+ ' '+r'$\rho$r',zorder=3)
plt.bar(X[0]+0.00,ro_yfp[3],color='gold',width=0.25,label= 'Control'+ ' '+r'$\rho$y',zorder=3)
plt.bar(X[1]-0.25,ro_rfp[4],color='coral',width=0.25,zorder=3)
plt.bar(X[1]+0.00,ro_yfp[4],color='gold',width=0.25,zorder=3)
plt.bar(X[2]-0.25,ro_rfp[5],color='coral',width=0.25,zorder=3)
plt.bar(X[2]+0.00,ro_yfp[5],color='gold',width=0.25,zorder=3)
plt.bar(X[3]-0.25,ro_rfp[9],color='r',width=0.25,label=r'$\rho$r',zorder=3)
plt.bar(X[3]+0.00,ro_yfp[9],color='yellow',width=0.25,label=r'$\rho$y',zorder=3)
plt.bar(X[4]-0.25,ro_rfp[10],color='r',width=0.25,zorder=3)
plt.bar(X[4]+0.00,ro_yfp[10],color='yellow',width=0.25,zorder=3)
plt.bar(X[5]-0.25,ro_rfp[11],color='r',width=0.25,zorder=3)
plt.bar(X[5]+0.00,ro_yfp[11],color='yellow',width=0.25,zorder=3)
plt.bar(X[6]-0.25,ro_rfp[15],color='r',width=0.25,zorder=3)
```

```

plt.bar(X[6]+0.00,ro_yfp[15],color='yellow',width=0.25,zorder=3)
plt.bar(X[7]-0.25,ro_rfp[16],color='r',width=0.25,zorder=3)
plt.bar(X[7]+0.00,ro_yfp[16],color='yellow',width=0.25,zorder=3)
plt.bar(X[8]-0.25,ro_rfp[17],color='r',width=0.25,zorder=3)
plt.bar(X[8]+0.00,ro_yfp[17],color='yellow',width=0.25,zorder=3)
plt.bar(X[9]-0.25,ro_rfp[21],color='r',width=0.25,zorder=3)
plt.bar(X[9]+0.00,ro_yfp[21],color='yellow',width=0.25,zorder=3)
plt.bar(X[10]-0.25,ro_rfp[22],color='r',width=0.25,zorder=3)
plt.bar(X[10]+0.00,ro_yfp[22],color='yellow',width=0.25,zorder=3)
plt.bar(X[11]-0.25,ro_rfp[23],color='r',width=0.25,zorder=3)
plt.bar(X[11]+0.00,ro_yfp[23],color='yellow',width=0.25,zorder=3)
plt.bar(X[12]-0.25,ro_rfp[27],color='r',width=0.25,zorder=3)
plt.bar(X[12]+0.00,ro_yfp[27],color='yellow',width=0.25,zorder=3)
plt.bar(X[13]-0.25,ro_rfp[28],color='r',width=0.25,zorder=3)
plt.bar(X[13]+0.00,ro_yfp[28],color='yellow',width=0.25,zorder=3)
plt.bar(X[14]-0.25,ro_rfp[29],color='r',width=0.25,zorder=3)
plt.bar(X[14]+0.00,ro_yfp[29],color='yellow',width=0.25,zorder=3)
plt.xticks(X,['Control 1','Control 2','Control 3','pTet-J23101-J23101 1','pTet-J23101-J
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper left',ncol=2)

```

Out[86]: <matplotlib.legend.Legend at 0x267f1dd0d68>



```
In [87]: X = np.arange(30)
plt.figure(figsize=(20,10))
plt.title(r'$\rho$', fontsize=15.0)
plt.ylabel(r'$\rho$')
plt.bar(X[0]-0.25,ro_rfp[0],color='coral',width=0.25,label= 'Control'+ ' '+r'$\rho$r Glu
plt.bar(X[0]+0.00,ro_yfp[0],color='gold',width=0.25,label= 'Control'+ ' '+r'$\rho$y Gluc
plt.bar(X[1]-0.25,ro_rfp[3],color='lightcoral',width=0.25,label= 'Control'+ ' '+r'$\rho$
plt.bar(X[1]+0.00,ro_yfp[3],color='palegreen',width=0.25,label= 'Control'+ ' '+r'$\rho$y
plt.bar(X[2]-0.25,ro_rfp[1],color='coral',width=0.25,zorder=3)
plt.bar(X[2]+0.00,ro_yfp[1],color='gold',width=0.25,zorder=3)
plt.bar(X[3]-0.25,ro_rfp[4],color='lightcoral',width=0.25,zorder=3)
plt.bar(X[3]+0.00,ro_yfp[4],color='palegreen',width=0.25,zorder=3)
plt.bar(X[4]-0.25,ro_rfp[2],color='coral',width=0.25,zorder=3)
plt.bar(X[4]+0.00,ro_yfp[2],color='gold',width=0.25,zorder=3)
plt.bar(X[5]-0.25,ro_rfp[5],color='lightcoral',width=0.25,zorder=3)
plt.bar(X[5]+0.00,ro_yfp[5],color='palegreen',width=0.25,zorder=3)
plt.bar(X[6]-0.25,ro_rfp[6],color='r',width=0.25,label=r'$\rho$r Glucosa',zorder=3)
```



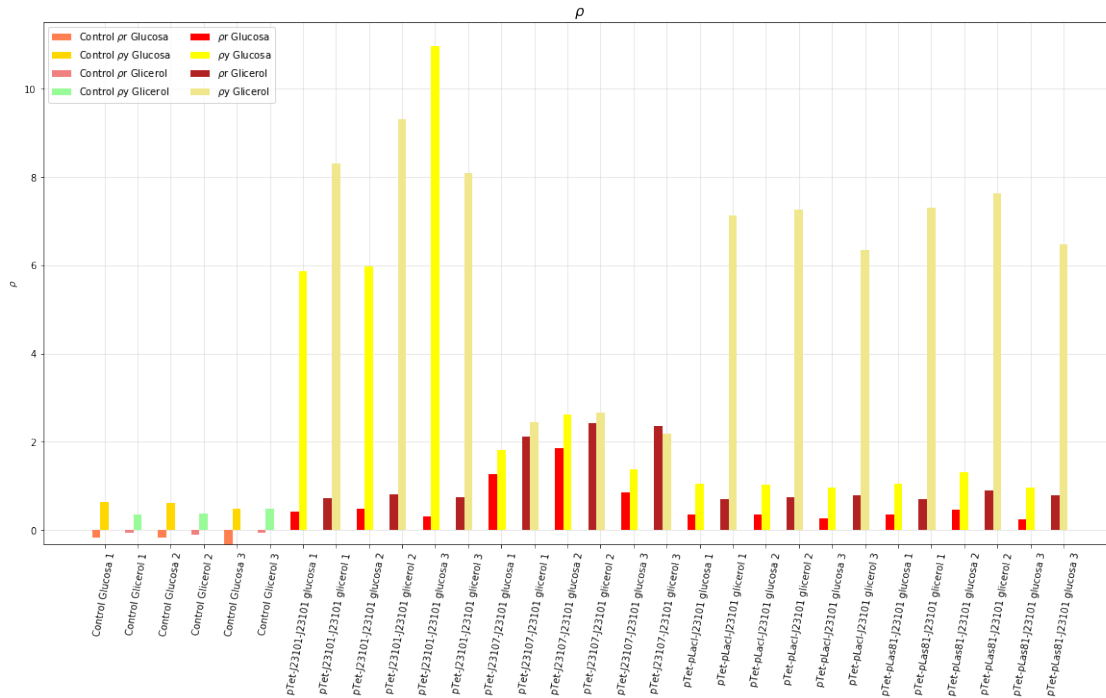
```

plt.bar(X[6]+0.00,ro_yfp[6],color='yellow',width=0.25,label=r'$\rho$y Glucosa',zorder=3)
plt.bar(X[7]-0.25,ro_rfp[9],color='firebrick',width=0.25,label=r'$\rho$r Glicerol',zord
plt.bar(X[7]+0.00,ro_yfp[9],color='khaki',width=0.25,label=r'$\rho$y Glicerol',zorder=3)
plt.bar(X[8]-0.25,ro_rfp[7],color='r',width=0.25,zorder=3)
plt.bar(X[8]+0.00,ro_yfp[7],color='yellow',width=0.25,zorder=3)
plt.bar(X[9]-0.25,ro_rfp[10],color='firebrick',width=0.25,zorder=3)
plt.bar(X[9]+0.00,ro_yfp[10],color='khaki',width=0.25,zorder=3)
plt.bar(X[10]-0.25,ro_rfp[8],color='r',width=0.25,zorder=3)
plt.bar(X[10]+0.00,ro_yfp[8],color='yellow',width=0.25,zorder=3)
plt.bar(X[11]-0.25,ro_rfp[11],color='firebrick',width=0.25,zorder=3)
plt.bar(X[11]+0.00,ro_yfp[11],color='khaki',width=0.25,zorder=3)
plt.bar(X[12]-0.25,ro_rfp[12],color='r',width=0.25,zorder=3)
plt.bar(X[12]+0.00,ro_yfp[12],color='yellow',width=0.25,zorder=3)
plt.bar(X[13]-0.25,ro_rfp[15],color='firebrick',width=0.25,zorder=3)
plt.bar(X[13]+0.00,ro_yfp[15],color='khaki',width=0.25,zorder=3)
plt.bar(X[14]-0.25,ro_rfp[13],color='r',width=0.25,zorder=3)
plt.bar(X[14]+0.00,ro_yfp[13],color='yellow',width=0.25,zorder=3)
plt.bar(X[15]-0.25,ro_rfp[16],color='firebrick',width=0.25,zorder=3)
plt.bar(X[15]+0.00,ro_yfp[16],color='khaki',width=0.25,zorder=3)
plt.bar(X[16]-0.25,ro_rfp[14],color='r',width=0.25,zorder=3)
plt.bar(X[16]+0.00,ro_yfp[14],color='yellow',width=0.25,zorder=3)
plt.bar(X[17]-0.25,ro_rfp[17],color='firebrick',width=0.25,zorder=3)
plt.bar(X[17]+0.00,ro_yfp[17],color='khaki',width=0.25,zorder=3)
plt.bar(X[18]-0.25,ro_rfp[18],color='r',width=0.25,zorder=3)
plt.bar(X[18]+0.00,ro_yfp[18],color='yellow',width=0.25,zorder=3)
plt.bar(X[19]-0.25,ro_rfp[21],color='firebrick',width=0.25,zorder=3)
plt.bar(X[19]+0.00,ro_yfp[21],color='khaki',width=0.25,zorder=3)
plt.bar(X[20]-0.25,ro_rfp[19],color='r',width=0.25,zorder=3)
plt.bar(X[20]+0.00,ro_yfp[19],color='yellow',width=0.25,zorder=3)
plt.bar(X[21]-0.25,ro_rfp[22],color='firebrick',width=0.25,zorder=3)
plt.bar(X[21]+0.00,ro_yfp[22],color='khaki',width=0.25,zorder=3)
plt.bar(X[22]-0.25,ro_rfp[20],color='r',width=0.25,zorder=3)
plt.bar(X[22]+0.00,ro_yfp[20],color='yellow',width=0.25,zorder=3)
plt.bar(X[23]-0.25,ro_rfp[23],color='firebrick',width=0.25,zorder=3)
plt.bar(X[23]+0.00,ro_yfp[23],color='khaki',width=0.25,zorder=3)
plt.bar(X[24]-0.25,ro_rfp[24],color='r',width=0.25,zorder=3)
plt.bar(X[24]+0.00,ro_yfp[24],color='yellow',width=0.25,zorder=3)
plt.bar(X[25]-0.25,ro_rfp[27],color='firebrick',width=0.25,zorder=3)
plt.bar(X[25]+0.00,ro_yfp[27],color='khaki',width=0.25,zorder=3)
plt.bar(X[26]-0.25,ro_rfp[25],color='r',width=0.25,zorder=3)
plt.bar(X[26]+0.00,ro_yfp[25],color='yellow',width=0.25,zorder=3)
plt.bar(X[27]-0.25,ro_rfp[28],color='firebrick',width=0.25,zorder=3)
plt.bar(X[27]+0.00,ro_yfp[28],color='khaki',width=0.25,zorder=3)
plt.bar(X[28]-0.25,ro_rfp[26],color='r',width=0.25,zorder=3)
plt.bar(X[28]+0.00,ro_yfp[26],color='yellow',width=0.25,zorder=3)
plt.bar(X[29]-0.25,ro_rfp[29],color='firebrick',width=0.25,zorder=3)
plt.bar(X[29]+0.00,ro_yfp[29],color='khaki',width=0.25,zorder=3)
plt.xticks(X, ['Control Glucosa 1','Control Glicerol 1','Control Glucosa 2','Control GL

```

```
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper left',ncol=2)
```

Out[87]: <matplotlib.legend.Legend at 0x267f1798c18>



In [88]: ro_rfpglu=[pr1,pr2,pr3],[pr7,pr8,pr9],[pr13,pr14,pr15],[pr19,pr20,pr21],[pr25,pr26,pr27]

ro_yfpglu=[py1,py2,py3],[py7,py8,py9],[py13,py14,py15],[py19,py20,py21],[py25,py26,py27]

xlabel=['Réplica 1','Réplica 2','Réplica 3']

ylabel=['Control','pTet-J23101-J23101','pTet-J23107-J23101','pTet-pLacI-J23101','pTet-pLacI-J23101-J23101']

```
plt.figure()
```

```
plt.title(r'$\rho$ Glucosa 0,4%')
```

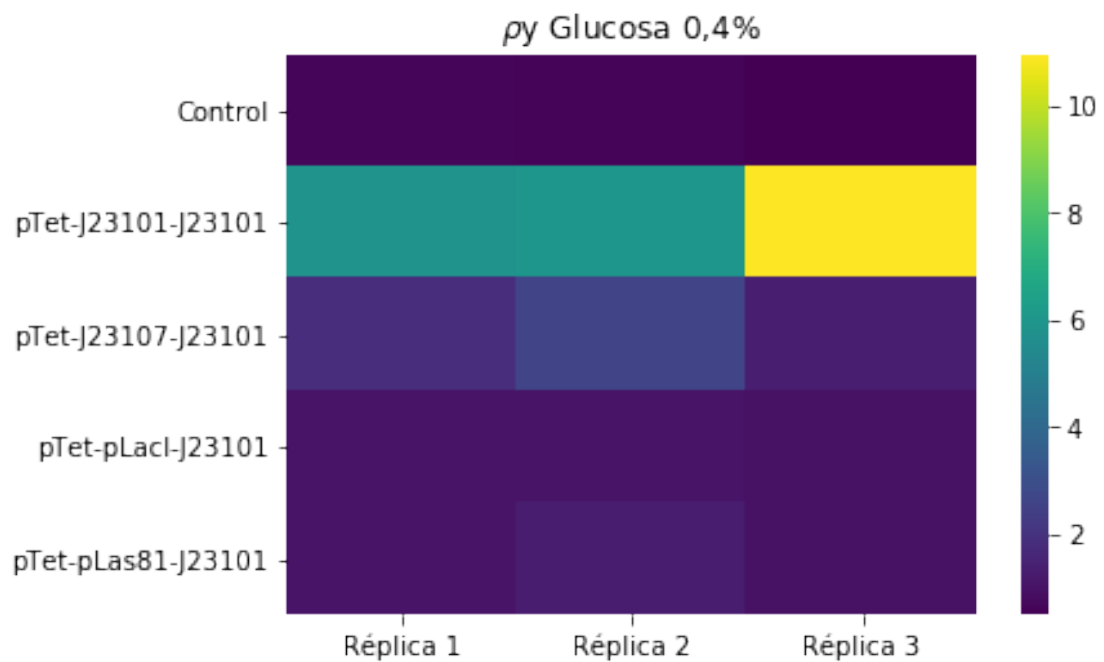
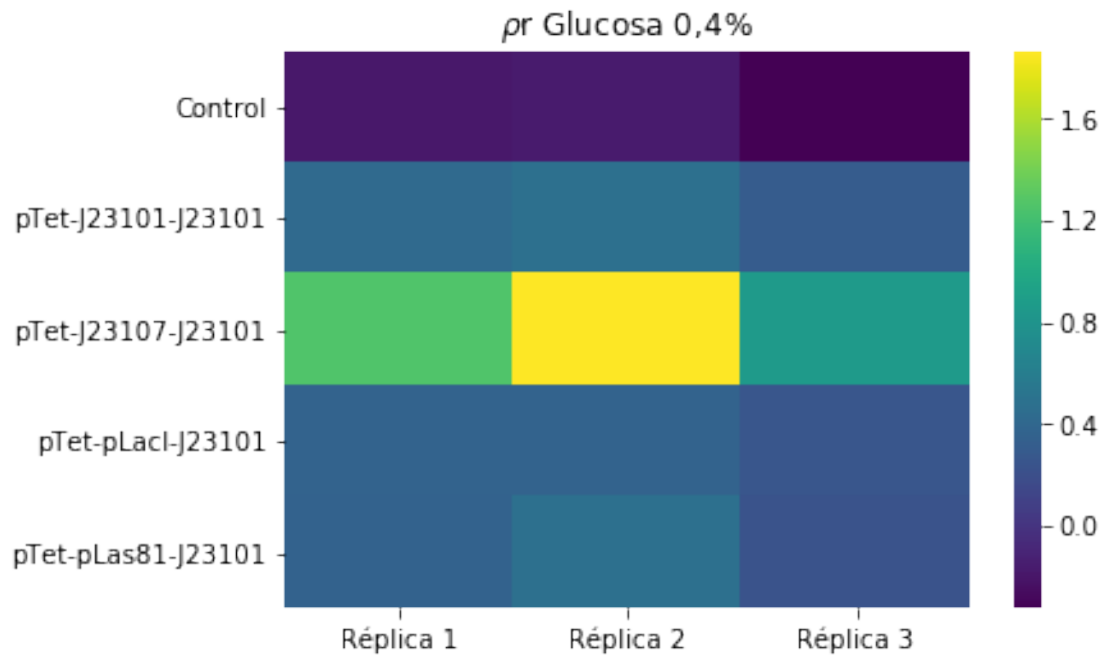
```
sns.heatmap(ro_rfpglu,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)
```

```
plt.figure()
```

```
plt.title(r'$\rho$ Glucosa 0,4%')
```

```
sns.heatmap(ro_yfpglu,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)
```

Out[88]: <matplotlib.axes._subplots.AxesSubplot at 0x267f1139b38>



```
In [89]: ro_rfpqli=[pr4,pr5,pr6],[pr10,pr11,pr12],[pr16,pr17,pr18],[pr22,pr23,pr24],[pr28,pr29,
ro_yfpqli=[py4,py5,py6],[py10,py11,py12],[py16,py17,py18],[py22,py23,py24],[py28,py29,
```

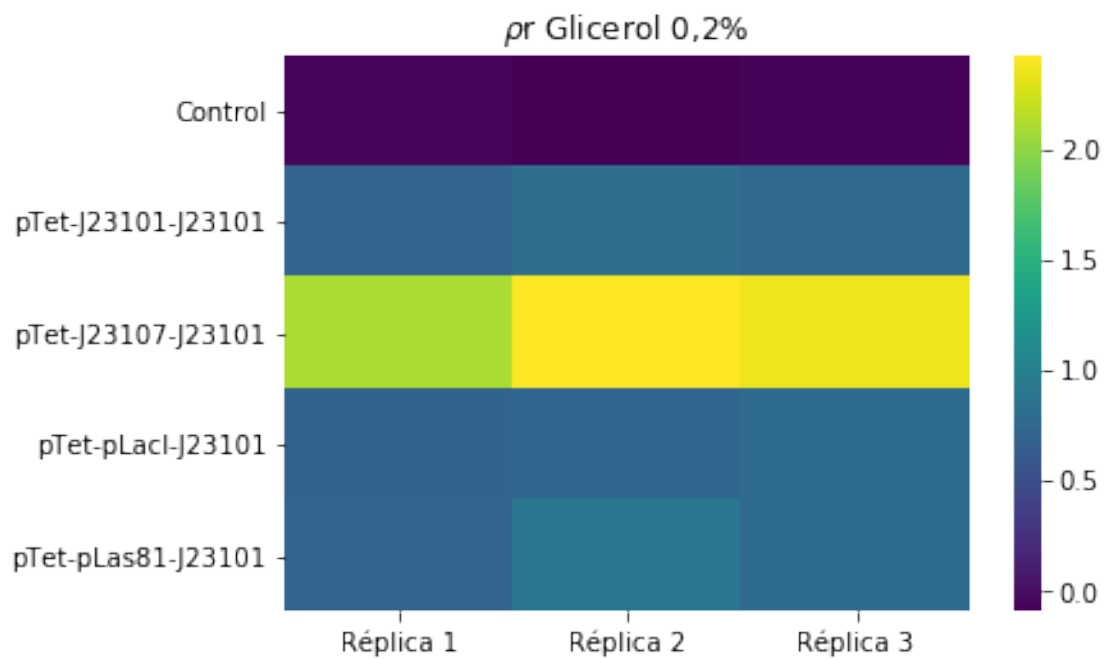
```

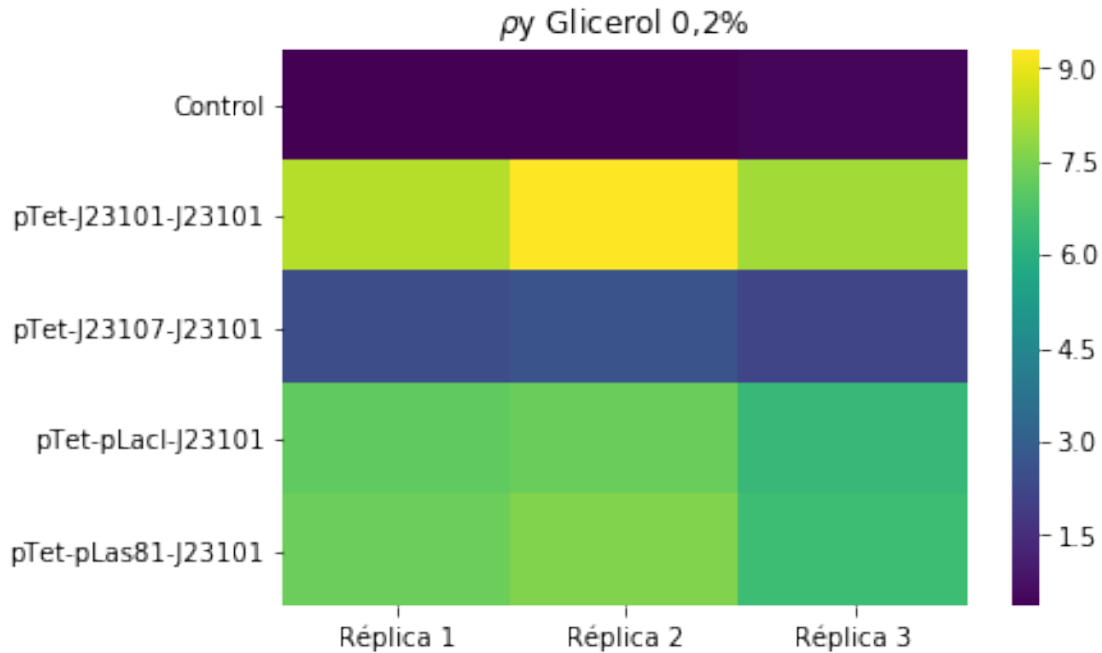
xlabel=['Réplica 1', 'Réplica 2', 'Réplica 3']
ylabel=['Control', 'pTet-J23101-J23101', 'pTet-J23107-J23101', 'pTet-pLacI-J23101', 'pTet-p
plt.figure()
plt.title(r'$\rho$ Glicerol 0,2%')
sns.heatmap(ro_rfpqli, cmap='viridis', xticklabels=xlabel, yticklabels=ylabel)

plt.figure()
plt.title(r'$\rho$ Glicerol 0,2%')
sns.heatmap(ro_yfpqli, cmap='viridis', xticklabels=xlabel, yticklabels=ylabel)

```

Out[89]: <matplotlib.axes._subplots.AxesSubplot at 0x267f155c6a0>





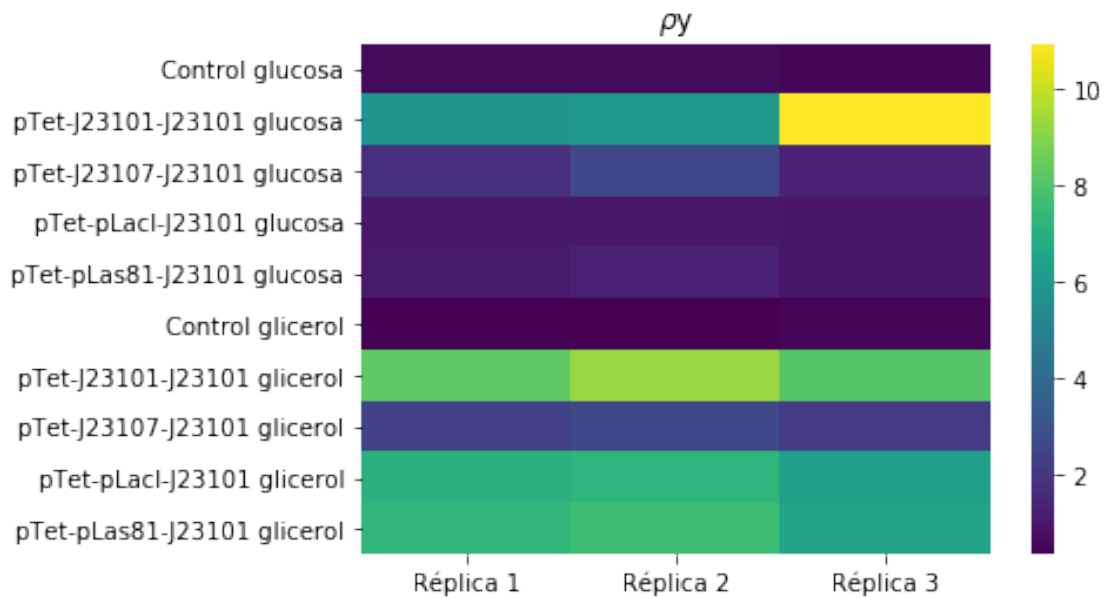
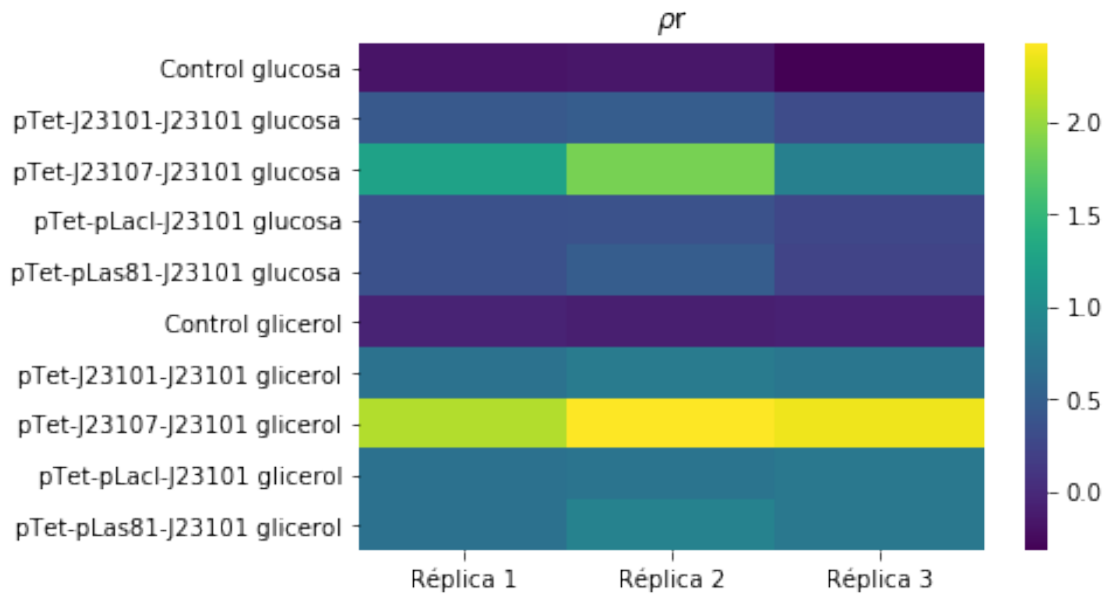
```
In [90]: ro_rfpqli=[pr4,pr5,pr6],[pr10,pr11,pr12],[pr16,pr17,pr18],[pr22,pr23,pr24],[pr28,pr29,
ro_rfpqlu=[pr1,pr2,pr3],[pr7,pr8,pr9],[pr13,pr14,pr15],[pr19,pr20,pr21],[pr25,pr26,pr27]
bothrfp=[pr1,pr2,pr3],[pr7,pr8,pr9],[pr13,pr14,pr15],[pr19,pr20,pr21],[pr25,pr26,pr27]
ro_yfpqli=[py1,py2,py3],[py7,py8,py9],[py13,py14,py15],[py19,py20,py21],[py25,py26,py27]
ro_yfpqlu=[py4,py5,py6],[py10,py11,py12],[py16,py17,py18],[py22,py23,py24],[py28,py29,py30]
bothyfp=[py1,py2,py3],[py7,py8,py9],[py13,py14,py15],[py19,py20,py21],[py25,py26,py27]

xlabel=['Réplica 1','Réplica 2','Réplica 3']
ylabel=['Control glucosa','pTet-J23101-J23101 glucosa','pTet-J23107-J23101 glucosa','pTet-pLacI-J23101 glucosa','pTet-pLas81-J23101 glucosa']

plt.figure()
plt.title(r'$\rho_r$')
sns.heatmap(bothrfp,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)

plt.figure()
plt.title(r'$\rho_y$')
sns.heatmap(bothyfp,cmap='viridis',xticklabels=xlabel,yticklabels=ylabel)
```

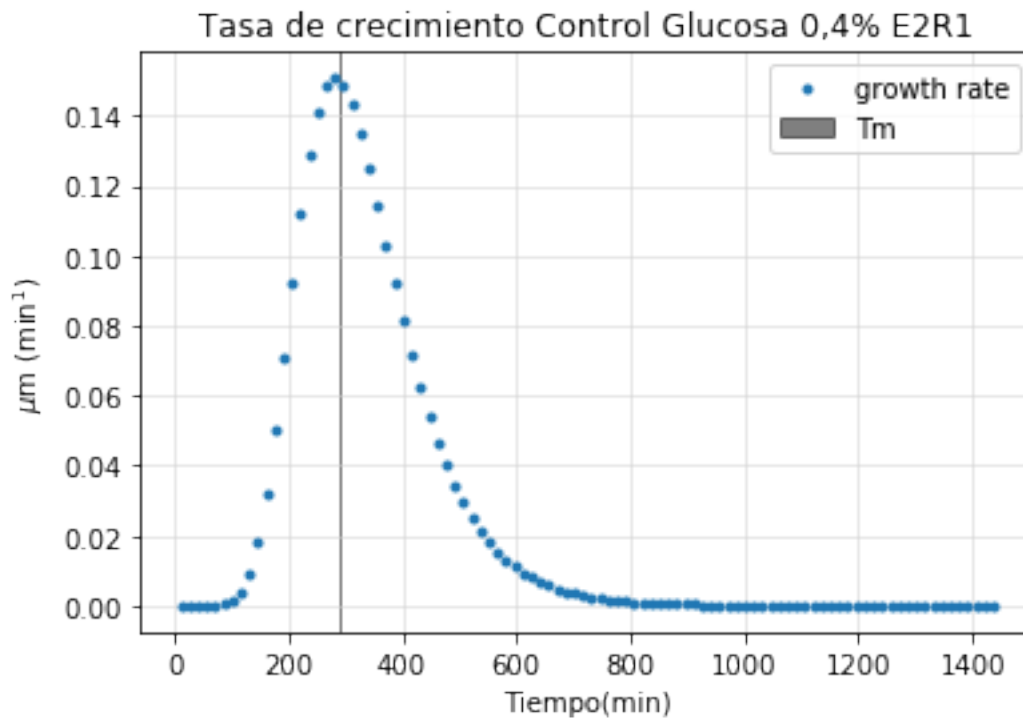
```
Out[90]: <matplotlib.axes._subplots.AxesSubplot at 0x267f34698d0>
```



```
In [91]: #tasa de crecimiento
ye1=((A1*np.exp(-np.exp((((um1*np.exp(1))/A1)*(11-tt))+1))))
#Con diff
dy1=(np.diff(ye1))
plt.figure()
plt.title('Tasa de crecimiento Control Glucosa 0,4% E2R1')
```

```
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm1,tm1, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy1,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

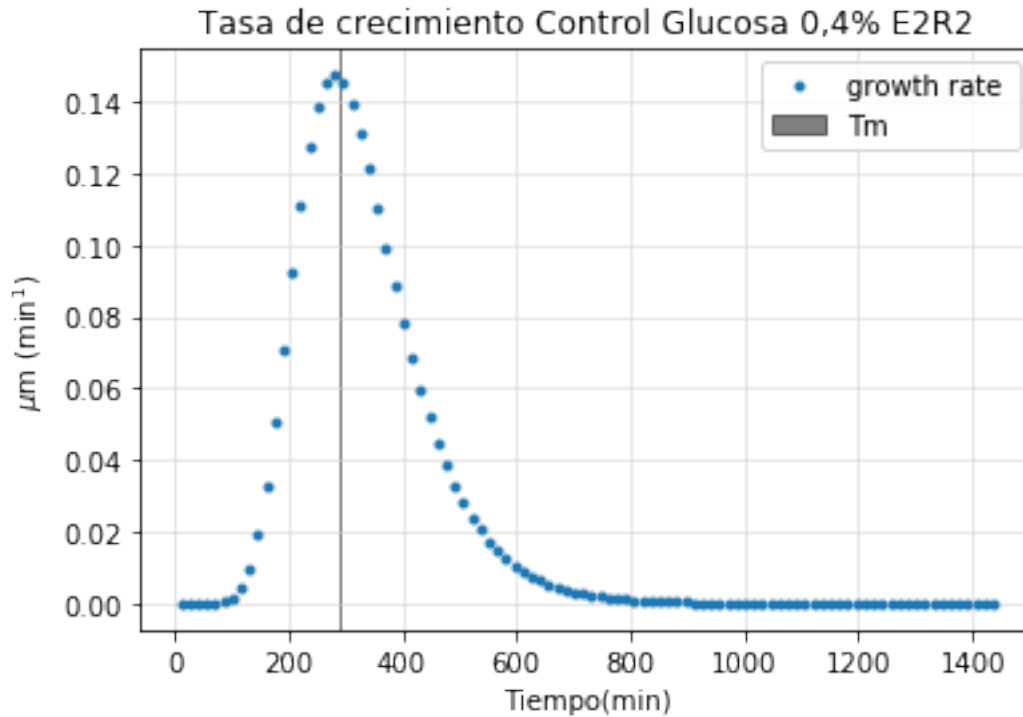
Out[91]: <matplotlib.legend.Legend at 0x267f5087c18>



In [92]: *#tasa de crecimiento*

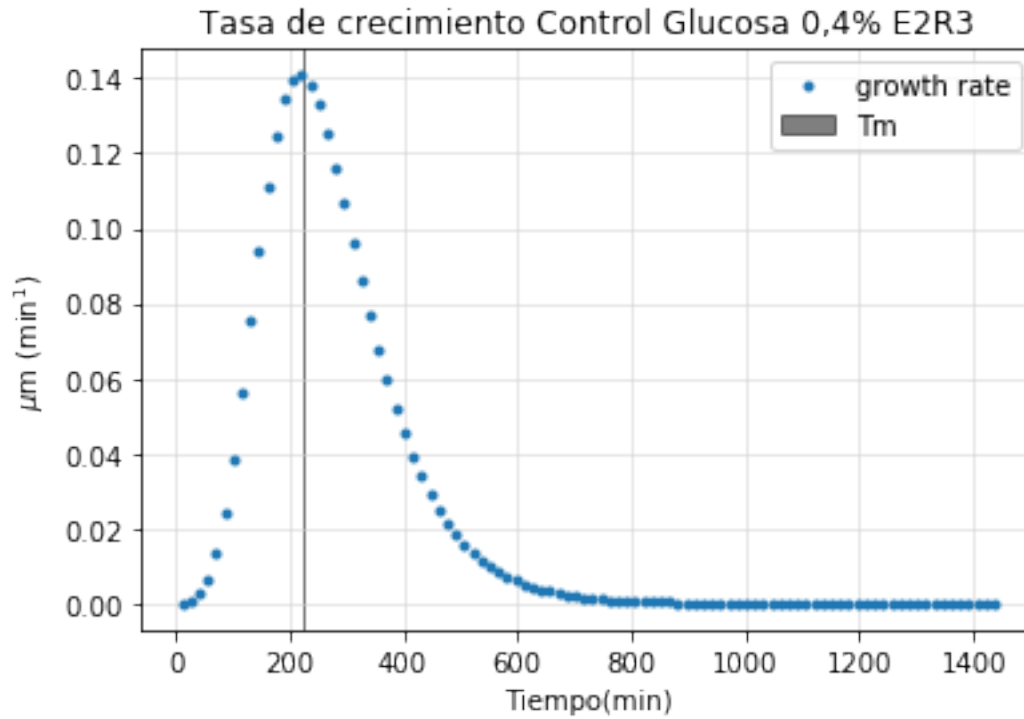
```
ye2=((A2*np.exp(-np.exp((((um2*np.exp(1))/A2)*(12-tt))+1))))
#Con diff
dy2=(np.diff(ye2))
plt.figure()
plt.title('Tasa de crecimiento Control Glucosa 0,4% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm2,tm2, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy2,'.',label='growth rate ')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

Out[92]: <matplotlib.legend.Legend at 0x267f5167d68>



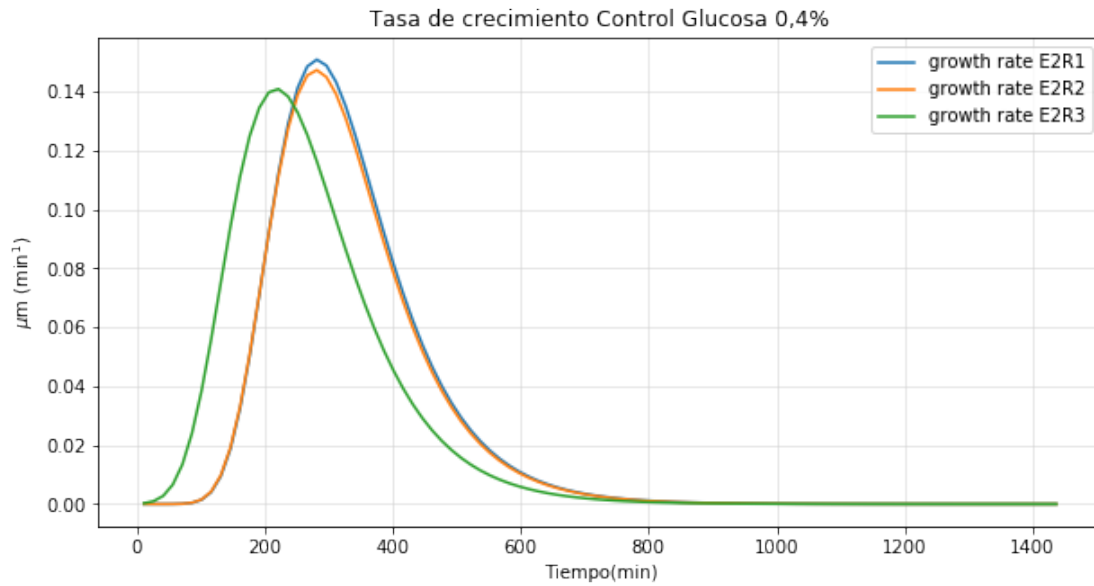
```
In [93]: #tasa de crecimiento
ye3=((A3*np.exp(-np.exp((((um3*np.exp(1))/A3)*(13-tt))+1))))
#Con diff
dy3=(np.diff(ye3))
plt.figure()
plt.title('Tasa de crecimiento Control Glucosa 0,4% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm3,tm3, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy3,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[93]: <matplotlib.legend.Legend at 0x267f5237d30>
```

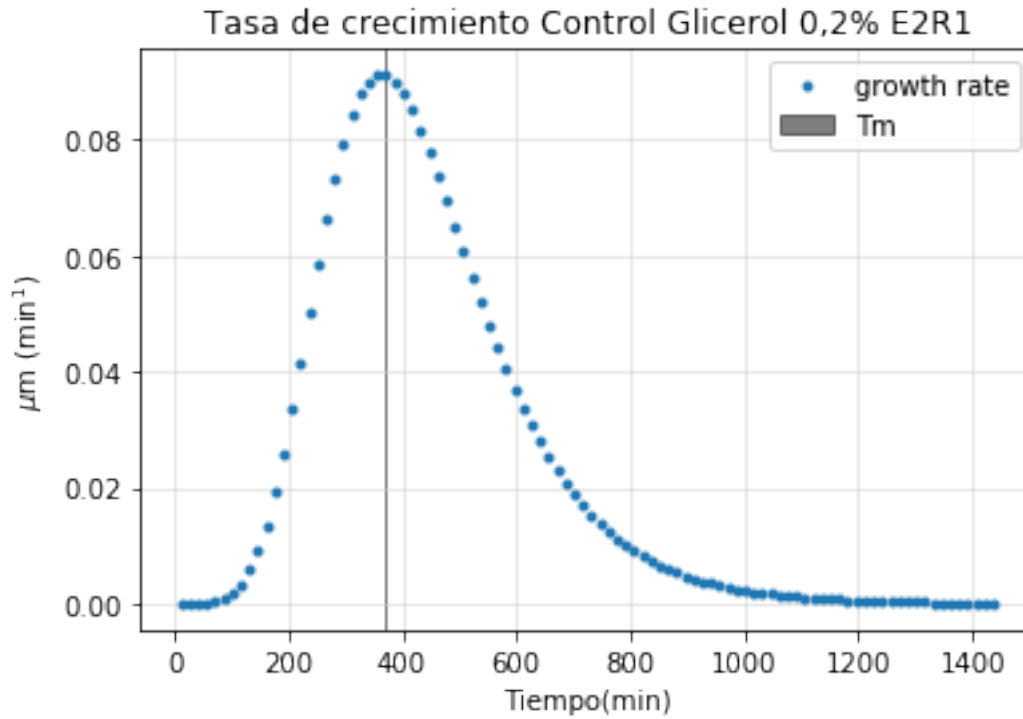
```
In [94]: #Tasas control réplicas glucosa
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento Control Glucosa 0,4%')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy1,label='growth rate E2R1')
plt.plot(tt[:-1],dy2,label='growth rate E2R2')
plt.plot(tt[:-1],dy3,label='growth rate E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[94]: <matplotlib.legend.Legend at 0x267f39fa208>
```



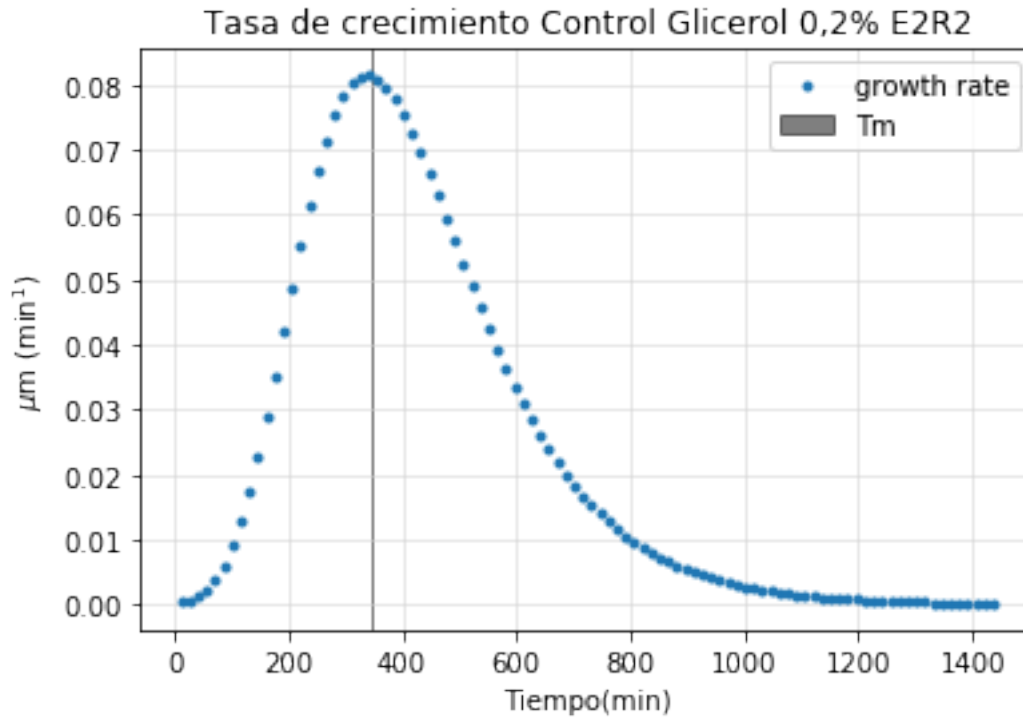
```
In [95]: #tasa de crecimiento
ye4=((A4*np.exp(-np.exp((((um4*np.exp(1))/A4)*(14-tt))+1))))
#Con diff
dy4=(np.diff(ye4))
plt.figure()
plt.title('Tasa de crecimiento Control Glicerol 0,2% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm4,tm4, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy4,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[95]: <matplotlib.legend.Legend at 0x267f1964b38>
```



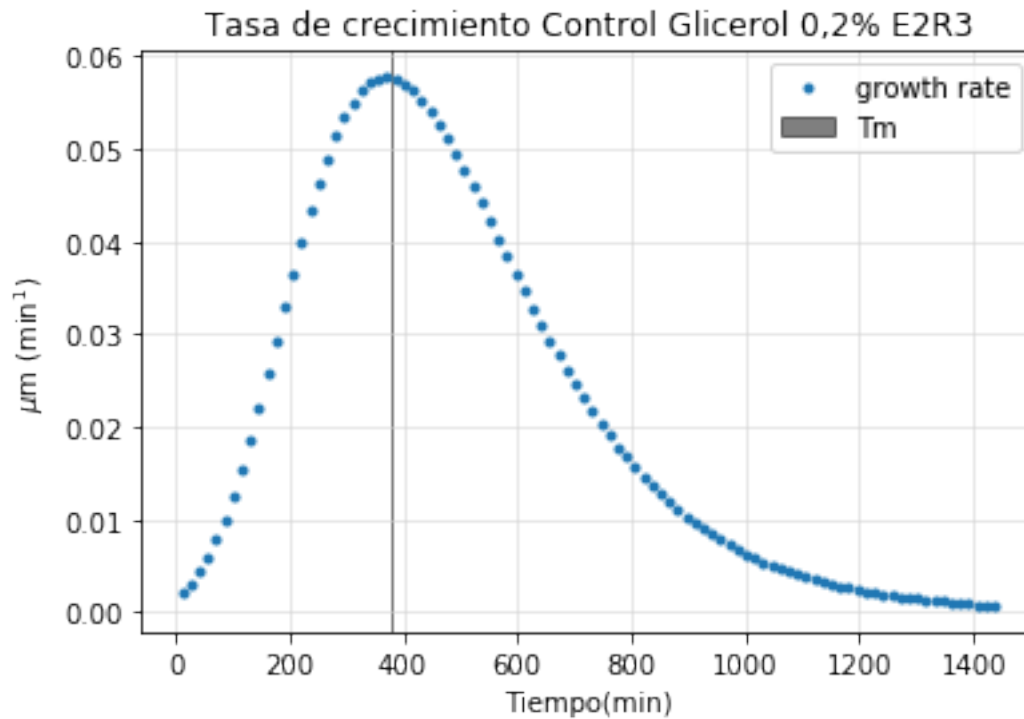
```
In [96]: #tasa de crecimiento
ye5=((A5*np.exp(-np.exp((((um5*np.exp(1))/A5)*(15-tt))+1))))
#Con diff
dy5=(np.diff(ye5))
plt.figure()
plt.title('Tasa de crecimiento Control Glicerol 0,2% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm5,tm5, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy5,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[96]: <matplotlib.legend.Legend at 0x267f3965390>
```



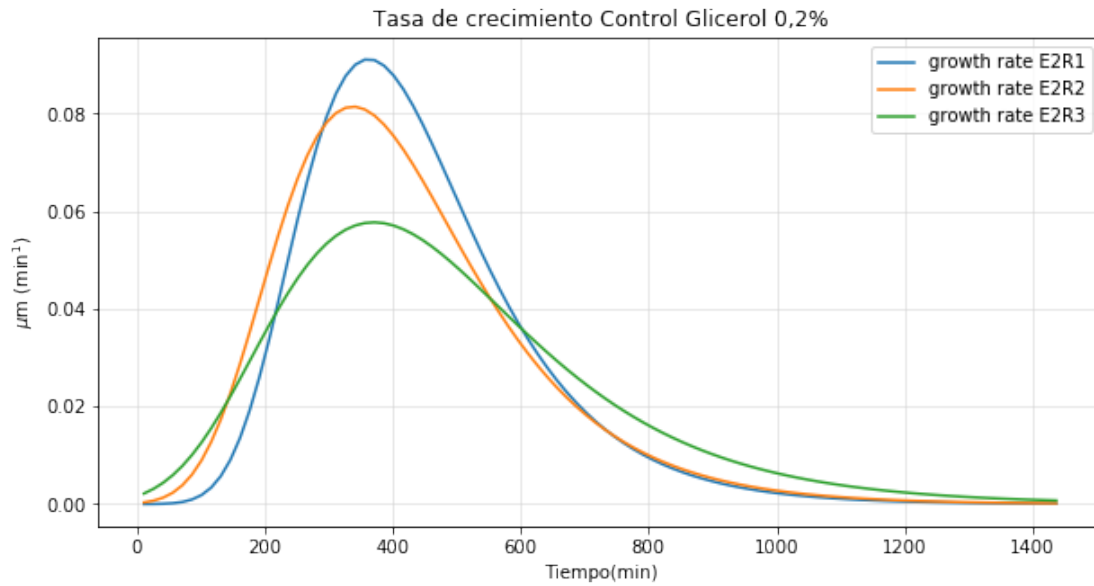
```
In [97]: #tasa de crecimiento
ye6=((A6*np.exp(-np.exp(((um6*np.exp(1))/A6)*(16-tt))+1))))
#Con diff
dy6=(np.diff(ye6))
plt.figure()
plt.title('Tasa de crecimiento Control Glicerol 0,2% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm6,tm6, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy6,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[97]: <matplotlib.legend.Legend at 0x267f1fd1b38>
```



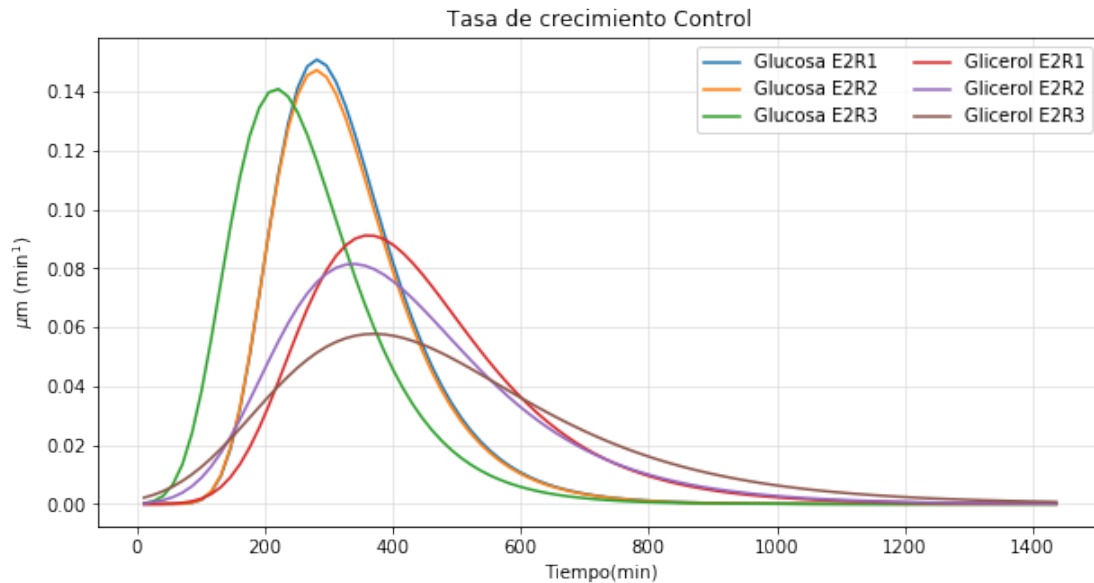
```
In [98]: #Tasas control réplicas glicerol
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento Control Glicerol 0,2%')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy4,label='growth rate E2R1')
plt.plot(tt[:-1],dy5,label='growth rate E2R2')
plt.plot(tt[:-1],dy6,label='growth rate E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[98]: <matplotlib.legend.Legend at 0x267f409d550>
```



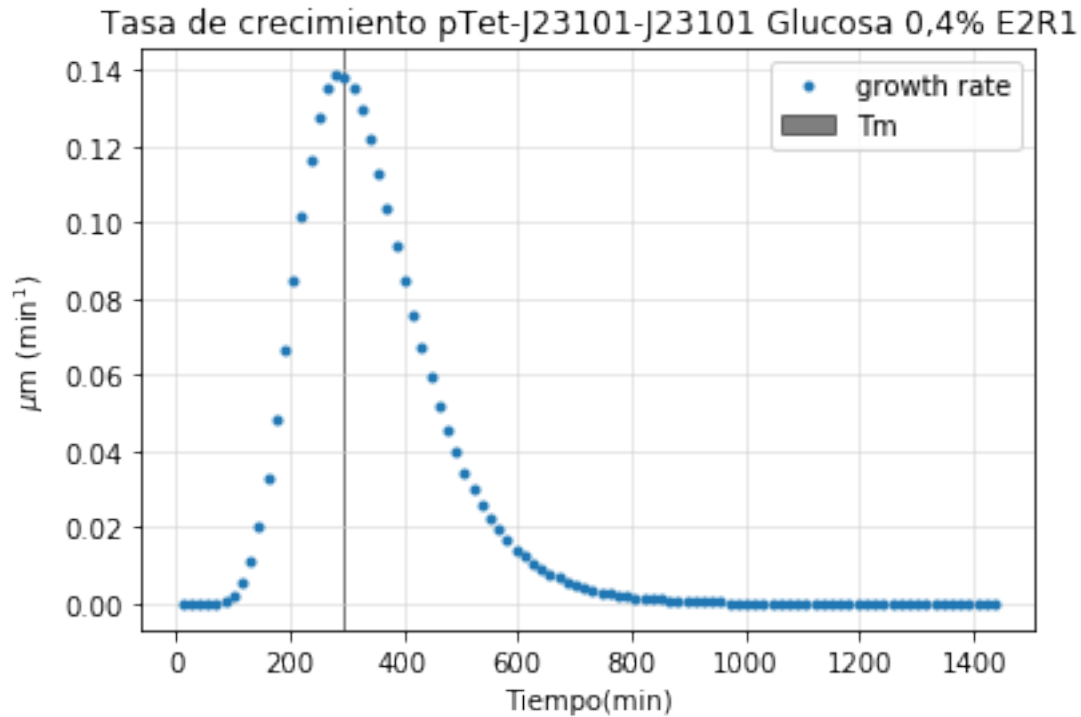
```
In [99]: #Tasas control réplicas controles
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento Control')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy1,label='Glucosa E2R1')
plt.plot(tt[:-1],dy2,label='Glucosa E2R2')
plt.plot(tt[:-1],dy3,label='Glucosa E2R3')
plt.plot(tt[:-1],dy4,label='Glicerol E2R1')
plt.plot(tt[:-1],dy5,label='Glicerol E2R2')
plt.plot(tt[:-1],dy6,label='Glicerol E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc='upper right',ncol=2)
```

```
Out[99]: <matplotlib.legend.Legend at 0x267f3342978>
```



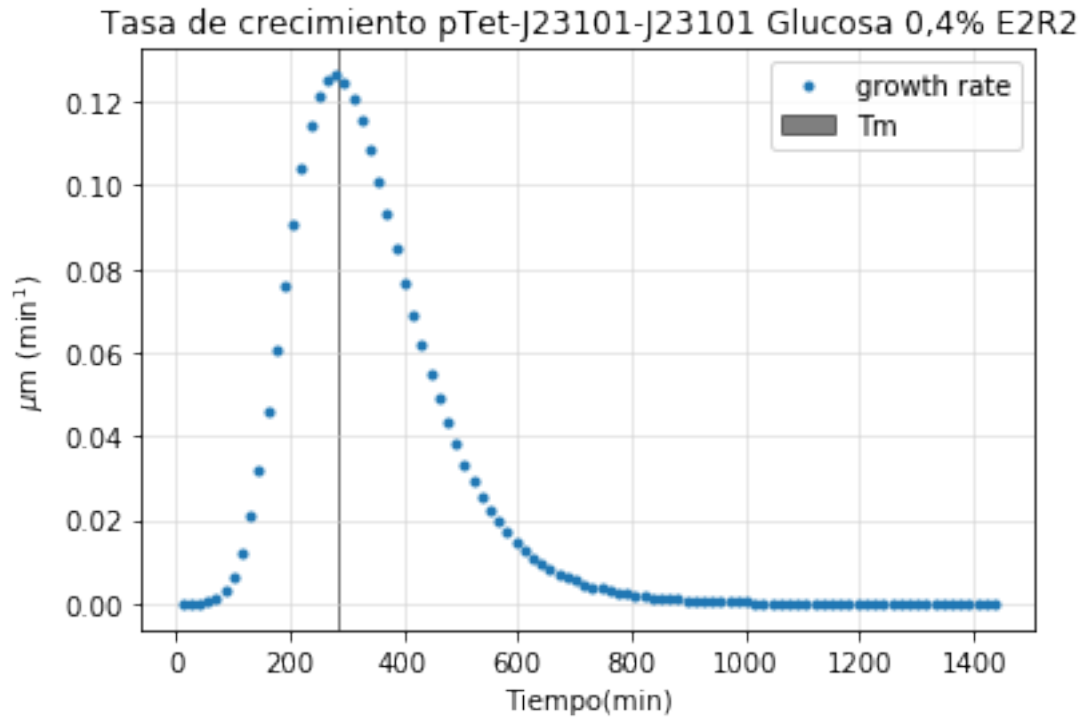
```
In [100]: #tasa de crecimiento
ye7=((A7*np.exp(-np.exp((((um7*np.exp(1))/A7)*(17-tt))+1))))
#Con diff
dy7=(np.diff(ye7))
plt.figure()
plt.title('Tasa de crecimiento pTet-J23101-J23101 Glucosa 0,4% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm7,tm7, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy7,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[100]: <matplotlib.legend.Legend at 0x267f3c8aa20>
```



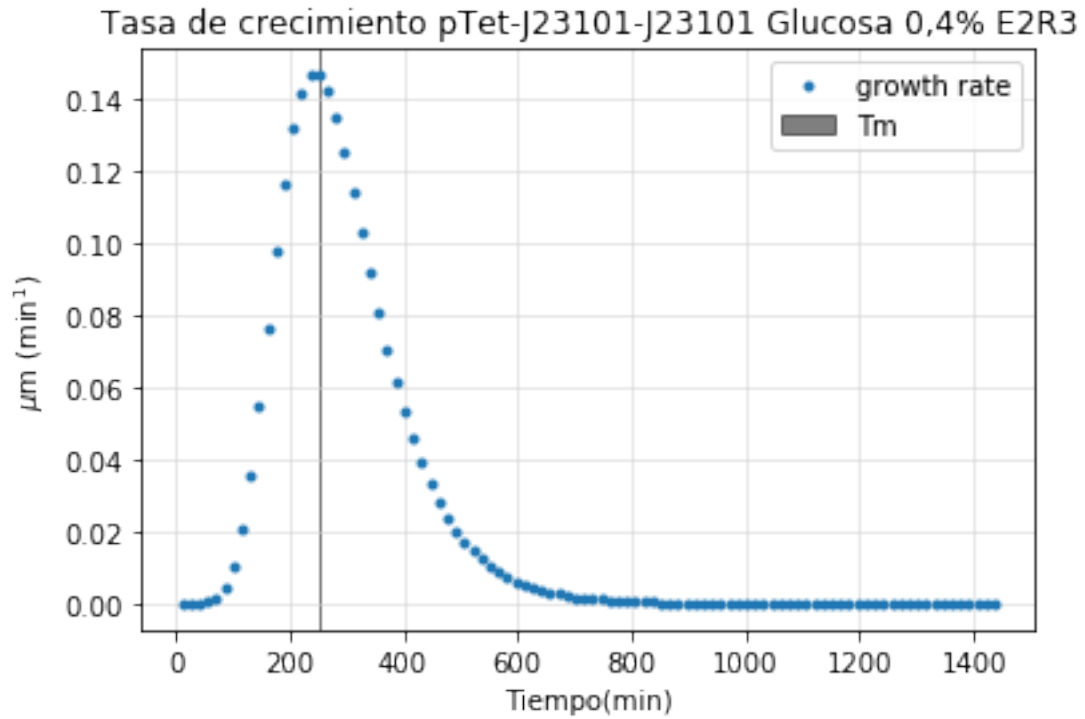
```
In [101]: #tasa de crecimiento
ye8=((A8*np.exp(-np.exp((((um8*np.exp(1))/A8)*(18-tt))+1))))
#Con diff
dy8=(np.diff(ye8))
plt.figure()
plt.title('Tasa de crecimiento pTet-J23101-J23101 Glucosa 0,4% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm8,tm8, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy8,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[101]: <matplotlib.legend.Legend at 0x267f3d57cc0>
```

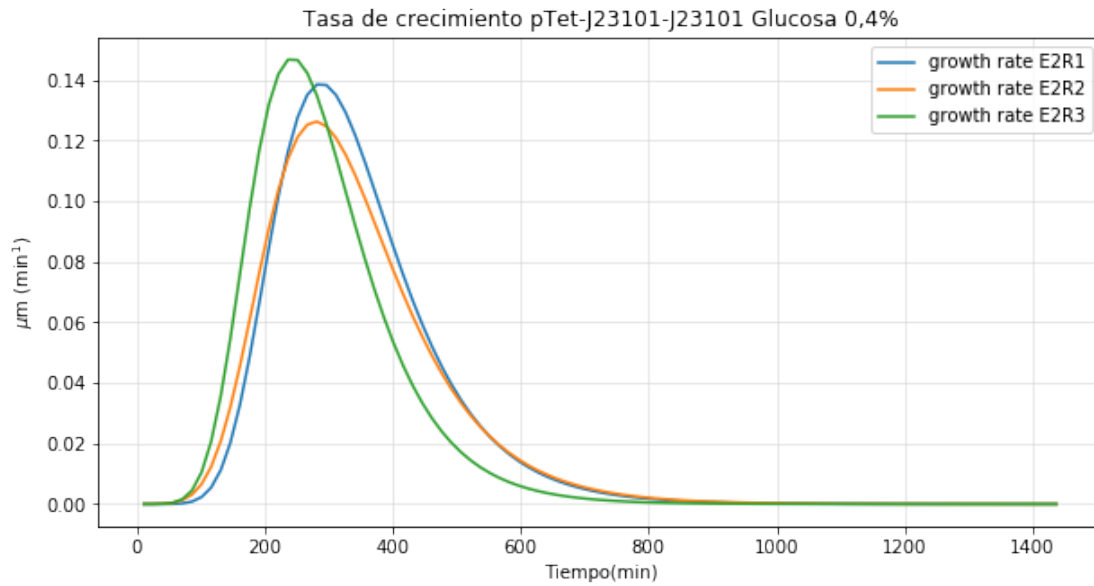
```
In [102]: #tasa de crecimiento
ye9=((A9*np.exp(-np.exp((((um9*np.exp(1))/A9)*(19-tt))+1))))
#Con diff
dy9=(np.diff(ye9))
plt.figure()
plt.title('Tasa de crecimiento pTet-J23101-J23101 Glucosa 0,4% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm9,tm9, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy9,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[102]: <matplotlib.legend.Legend at 0x267f46e3860>
```



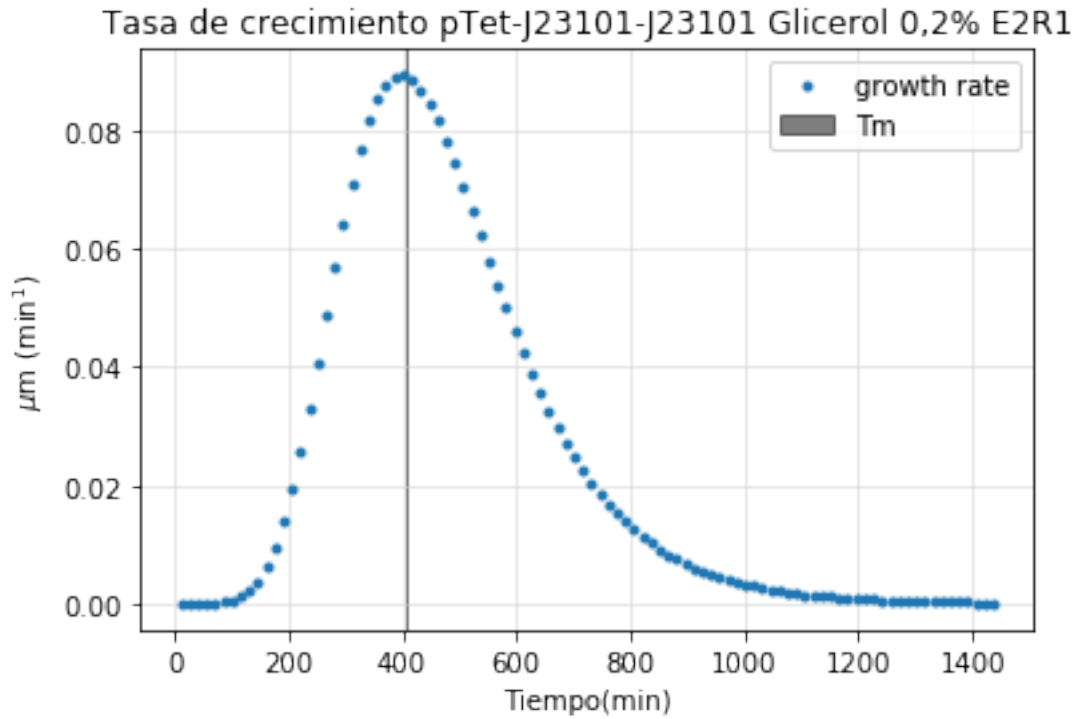
```
In [103]: #Tasas pLux-J23101-J23101 réplicas glucosa
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento pTet-J23101-J23101 Glucosa 0,4%')
plt.xlabel('Tiempo(min)')
plt.ylabel(r' $\mu\text{m (min}^{-1}\text{)}$ ')
plt.plot(tt[:-1],dy7,label='growth rate E2R1')
plt.plot(tt[:-1],dy8,label='growth rate E2R2')
plt.plot(tt[:-1],dy9,label='growth rate E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[103]: <matplotlib.legend.Legend at 0x267f49397f0>
```



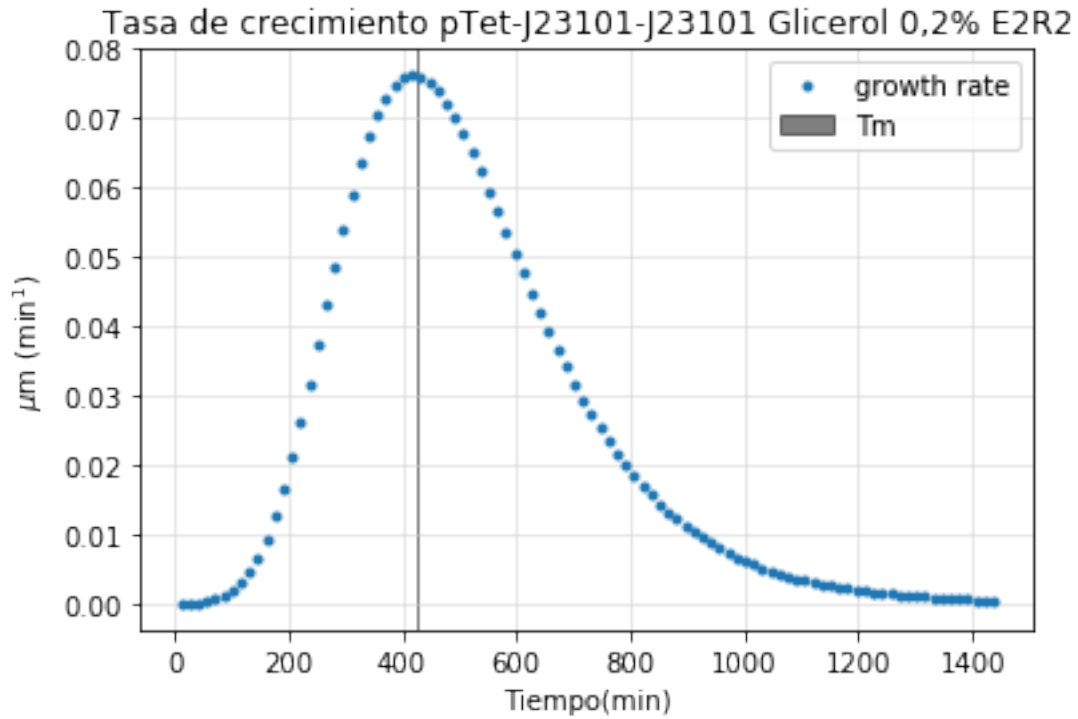
```
In [104]: #tasa de crecimiento
ye10=((A10*np.exp(-np.exp((((um10*np.exp(1))/A10)*(110-tt))+1))))
#Con diff
dy10=(np.diff(ye10))
plt.figure()
plt.title('Tasa de crecimiento pTet-J23101-J23101 Glicerol 0,2% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm10,tm10, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy10,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[104]: <matplotlib.legend.Legend at 0x267f4a1ed30>
```



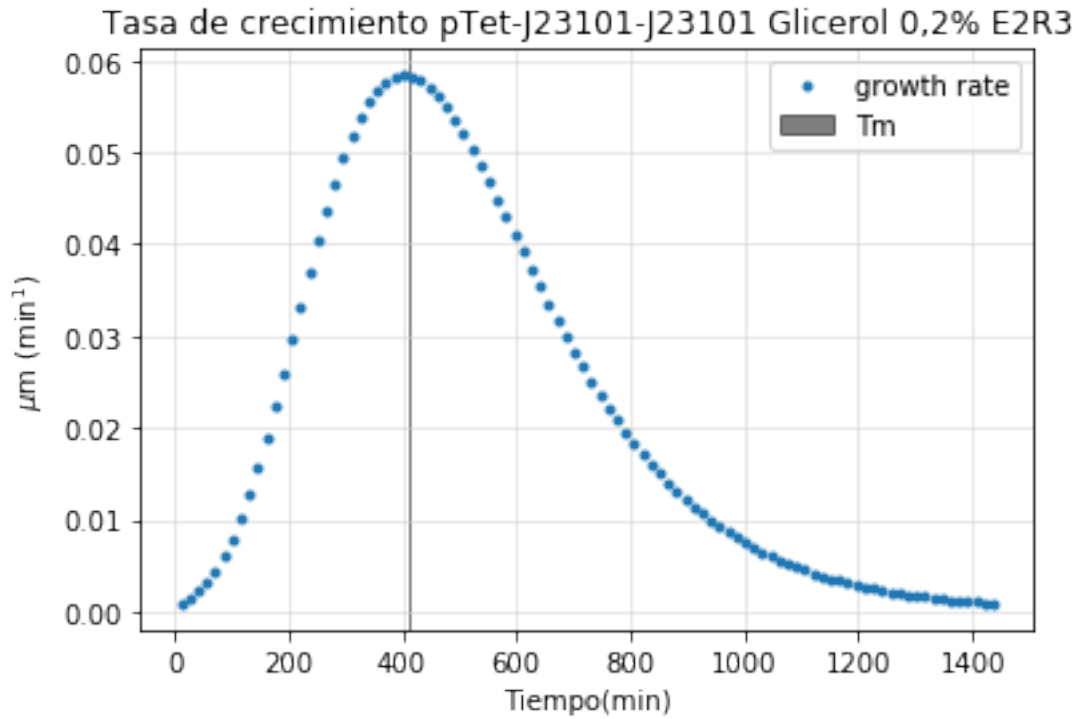
```
In [105]: #tasa de crecimiento
ye11=((A11*np.exp(-np.exp((((um11*np.exp(1))/A11)*(111-tt))+1))))
#Con diff
dy11=(np.diff(ye11))
plt.figure()
plt.title('Tasa de crecimiento pTet-J23101-J23101 Glicerol 0,2% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm11,tm11, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy11,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[105]: <matplotlib.legend.Legend at 0x267f49afa20>
```



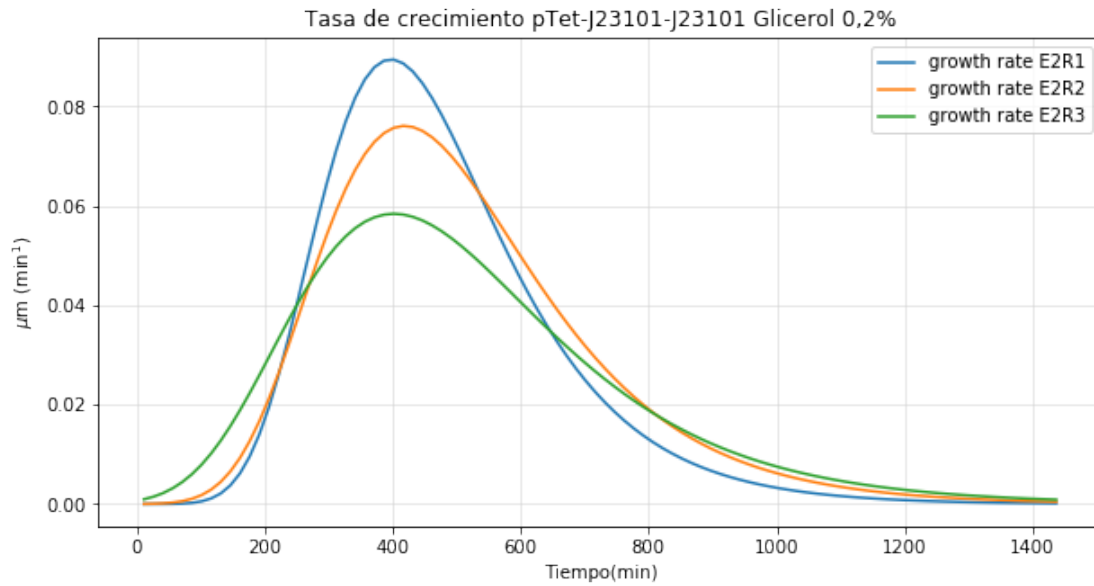
```
In [106]: #tasa de crecimiento
ye12=((A12*np.exp(-np.exp((((um12*np.exp(1))/A12)*(112-tt))+1))))
#Con diff
dy12=(np.diff(ye12))
plt.figure()
plt.title('Tasa de crecimiento pTet-J23101-J23101 Glicerol 0,2% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm12,tm12, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy12,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[106]: <matplotlib.legend.Legend at 0x267f4bc1dd8>
```



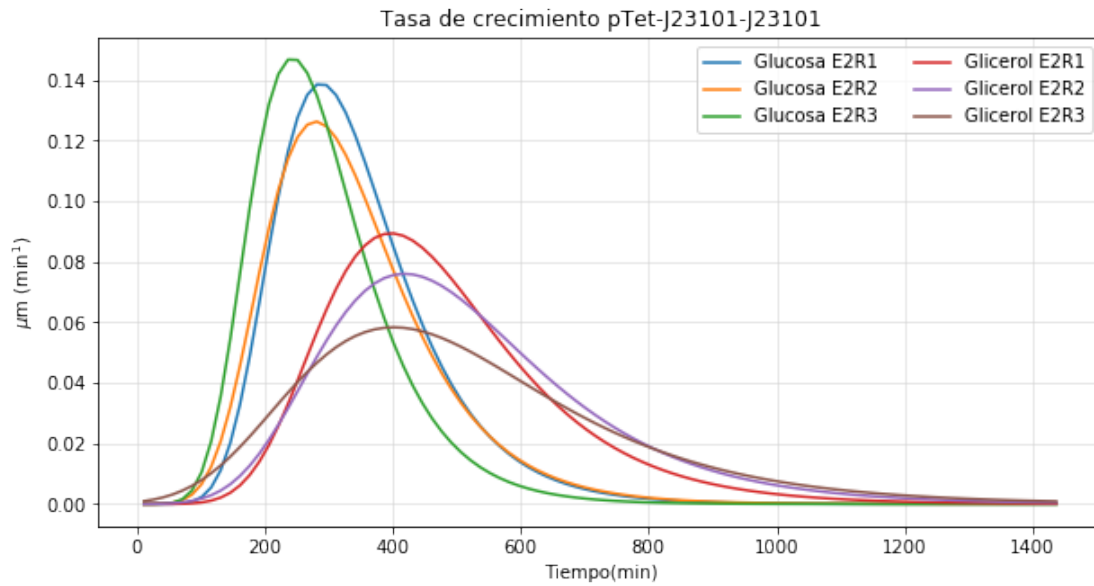
```
In [107]: #Tasas control réplicas glicerol
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento pTet-J23101-J23101 Glicerol 0,2%')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy10,label='growth rate E2R1')
plt.plot(tt[:-1],dy11,label='growth rate E2R2')
plt.plot(tt[:-1],dy12,label='growth rate E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[107]: <matplotlib.legend.Legend at 0x267f4c94b70>
```



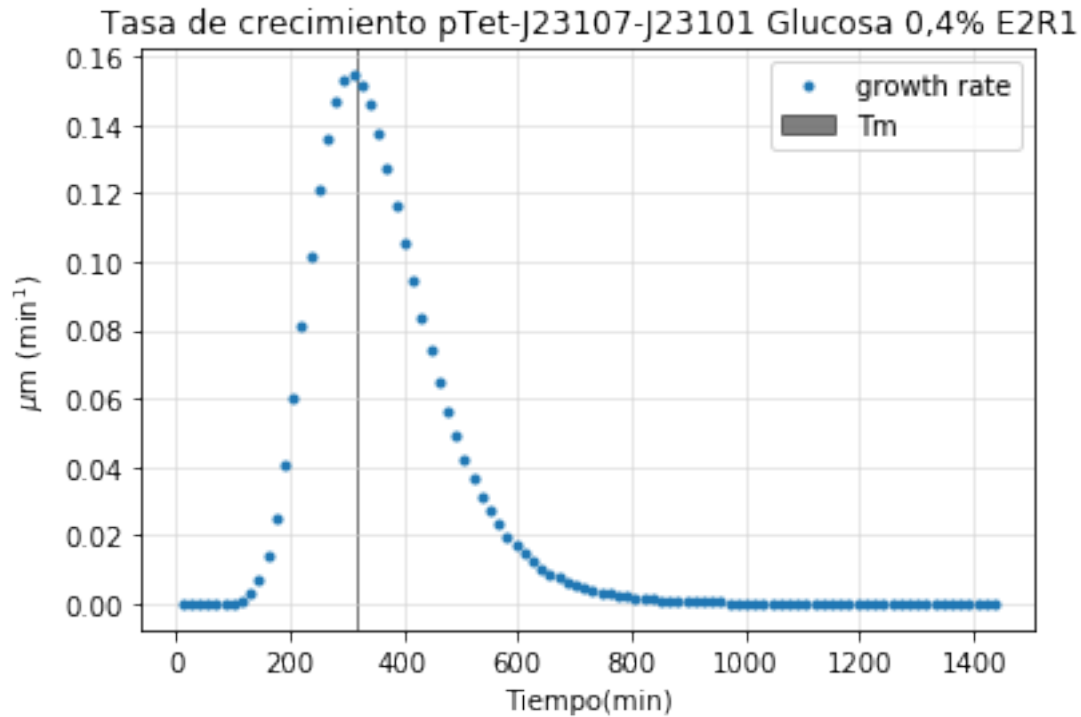
```
In [108]: #Tasas control réplicas
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento pTet-J23101-J23101')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy7,label='Glucosa E2R1')
plt.plot(tt[:-1],dy8,label='Glucosa E2R2')
plt.plot(tt[:-1],dy9,label='Glucosa E2R3')
plt.plot(tt[:-1],dy10,label='Glicerol E2R1')
plt.plot(tt[:-1],dy11,label='Glicerol E2R2')
plt.plot(tt[:-1],dy12,label='Glicerol E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc='upper right',ncol=2)
```

```
Out[108]: <matplotlib.legend.Legend at 0x267f5353780>
```



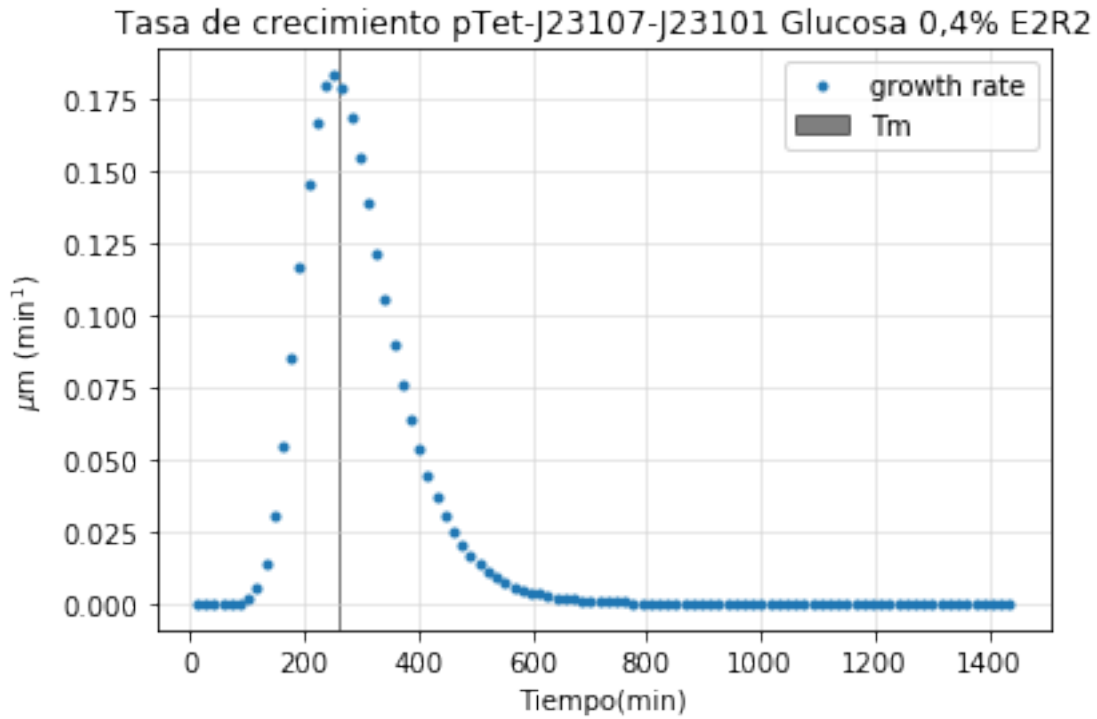
```
In [109]: #tasa de crecimiento
ye13=((A13*np.exp(-np.exp((((um13*np.exp(1))/A13)*(113-tt))+1))))
#Con diff
dy13=(np.diff(ye13))
plt.figure()
plt.title('Tasa de crecimiento pTet-J23107-J23101 Glucosa 0,4% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm13,tm13, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy13,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[109]: <matplotlib.legend.Legend at 0x267f542d940>
```

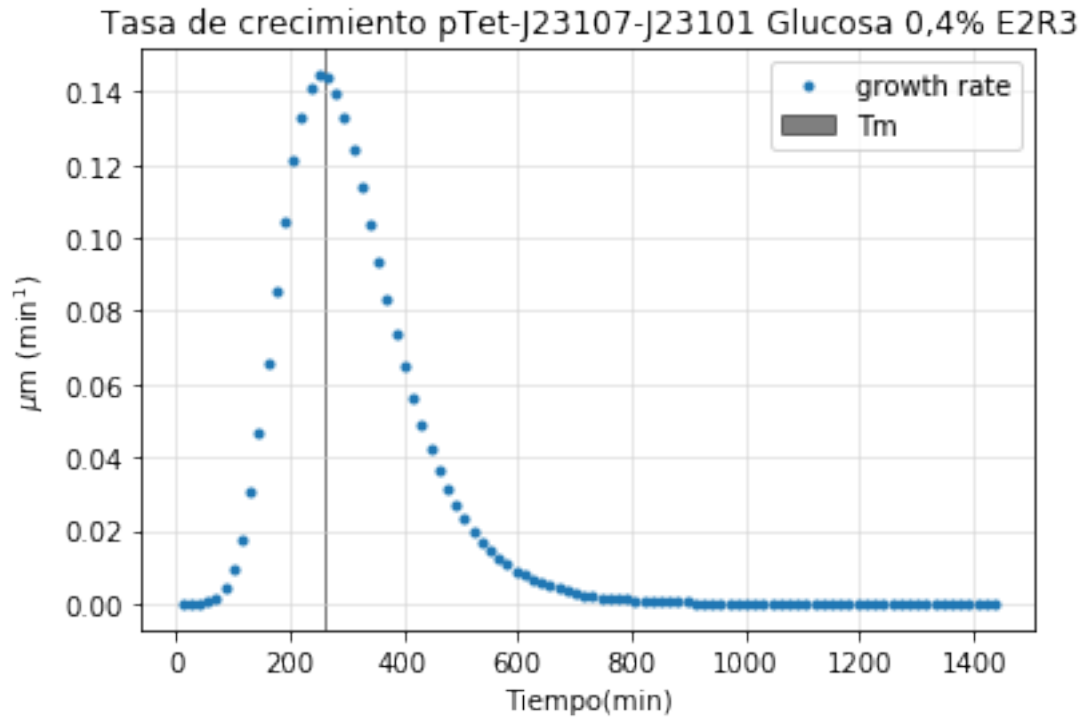
```
In [110]: #tasa de crecimiento
ye14=((A14*np.exp(-np.exp((((um14*np.exp(1))/A14)*(114-tt))+1))))
#Con diff
dy14=(np.diff(ye14))
plt.figure()
plt.title('Tasa de crecimiento pTet-J23107-J23101 Glucosa 0,4% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm14,tm14, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy14,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[110]: <matplotlib.legend.Legend at 0x267f5505e80>
```



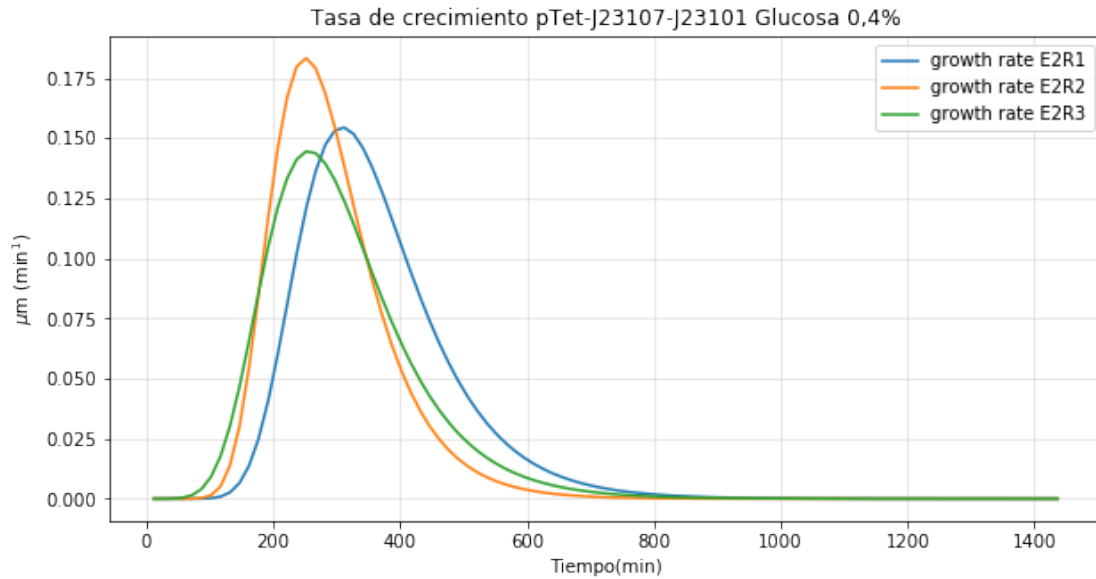
```
In [111]: #tasa de crecimiento
ye15=((A15*np.exp(-np.exp((((um15*np.exp(1))/A15)*(115-tt))+1))))
#Con diff
dy15=(np.diff(ye15))
plt.figure()
plt.title('Tasa de crecimiento pTet-J23107-J23101 Glucosa 0,4% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm15,tm15, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy15,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[111]: <matplotlib.legend.Legend at 0x267f55e6d68>
```



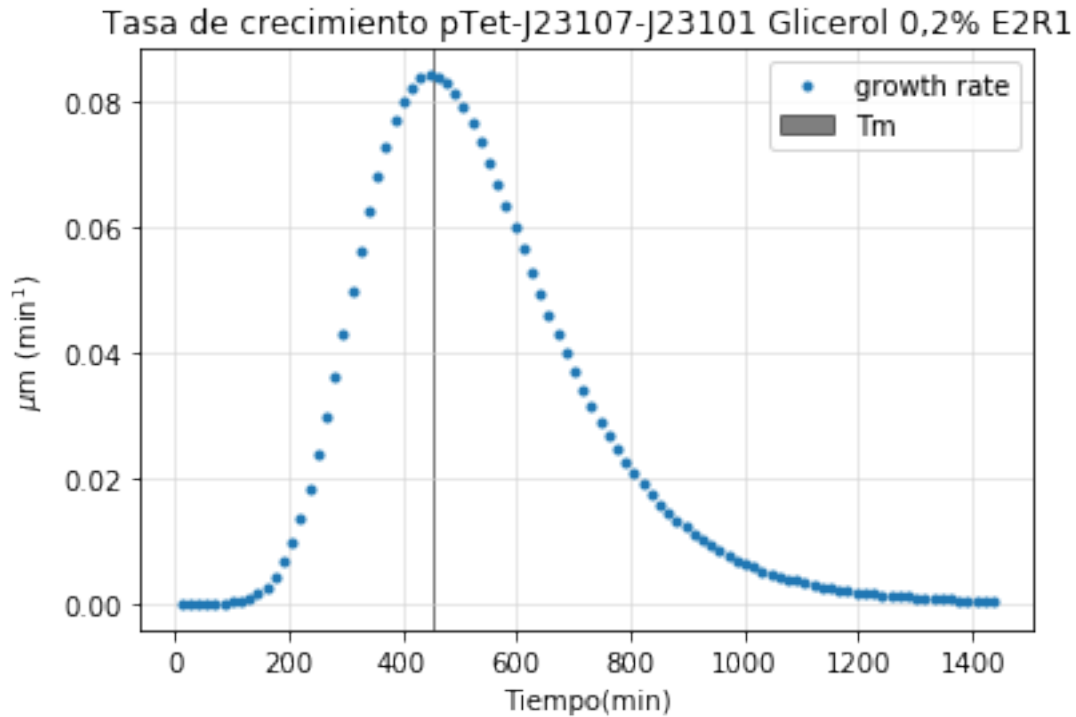
```
In [112]: #Tasas plux76-J23101-J23101 réplicas glucosa
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento pTet-J23107-J23101 Glucosa 0,4%')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy13,label='growth rate E2R1')
plt.plot(tt[:-1],dy14,label='growth rate E2R2')
plt.plot(tt[:-1],dy15,label='growth rate E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[112]: <matplotlib.legend.Legend at 0x267f56b6cc0>
```



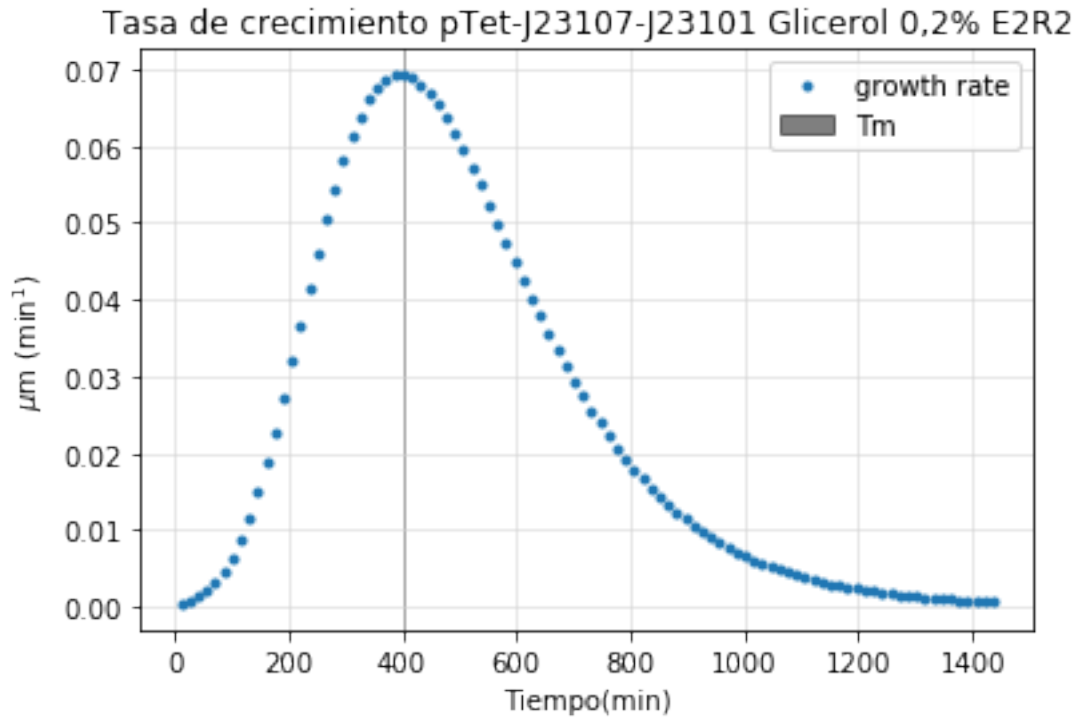
```
In [113]: #tasa de crecimiento
ye16=((A16*np.exp(-np.exp((((um16*np.exp(1))/A16)*(116-tt))+1))))
#Con diff
dy16=(np.diff(ye16))
plt.figure()
plt.title('Tasa de crecimiento pTet-J23107-J23101 Glicerol 0,2% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm16,tm16, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy16,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[113]: <matplotlib.legend.Legend at 0x267f77b9ac8>
```



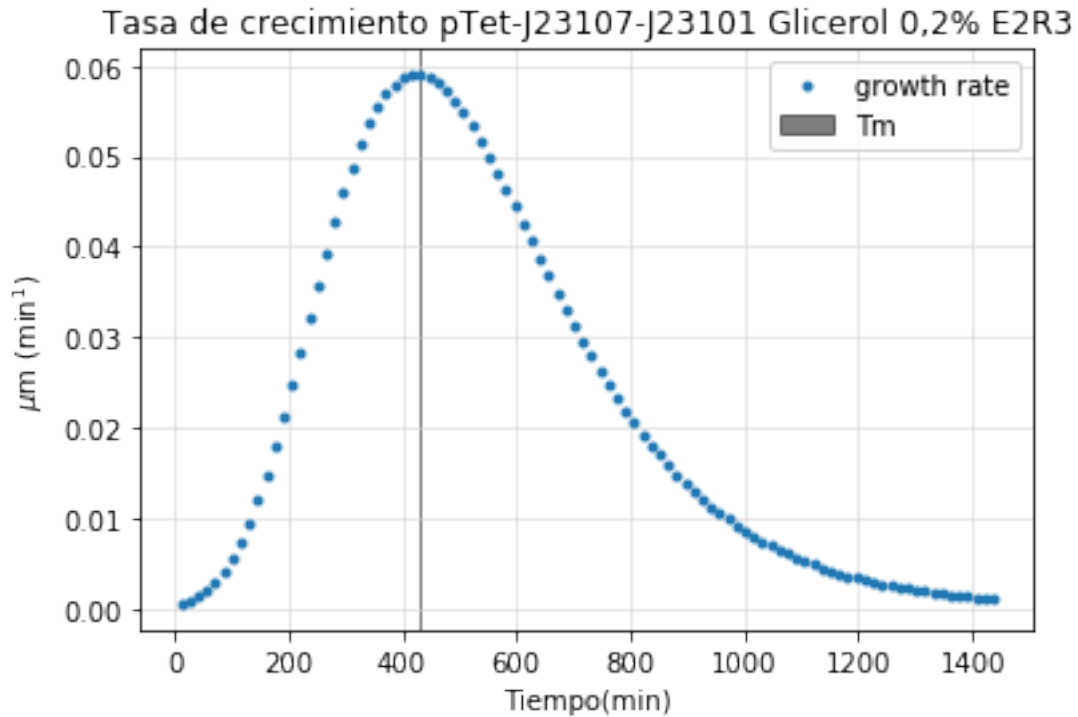
```
In [114]: #tasa de crecimiento
ye17=((A17*np.exp(-np.exp((((um17*np.exp(1))/A17)*(117-tt))+1))))
#Con diff
dy17=(np.diff(ye17))
plt.figure()
plt.title('Tasa de crecimiento pTet-J23107-J23101 Glicerol 0,2% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm17,tm17, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy17,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[114]: <matplotlib.legend.Legend at 0x267f788aeb8>
```



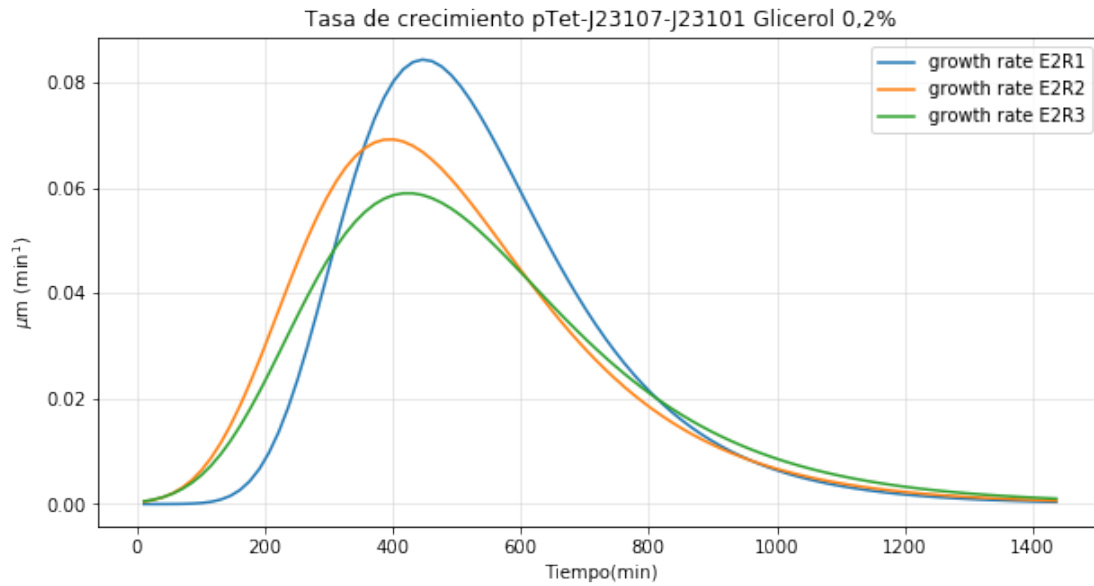
```
In [115]: #tasa de crecimiento
ye18=((A18*np.exp(-np.exp((((um18*np.exp(1))/A18)*(118-tt))+1))))
#Con diff
dy18=(np.diff(ye18))
plt.figure()
plt.title('Tasa de crecimiento pTet-J23107-J23101 Glicerol 0,2% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm18,tm18, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy18,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[115]: <matplotlib.legend.Legend at 0x267f4cee048>
```



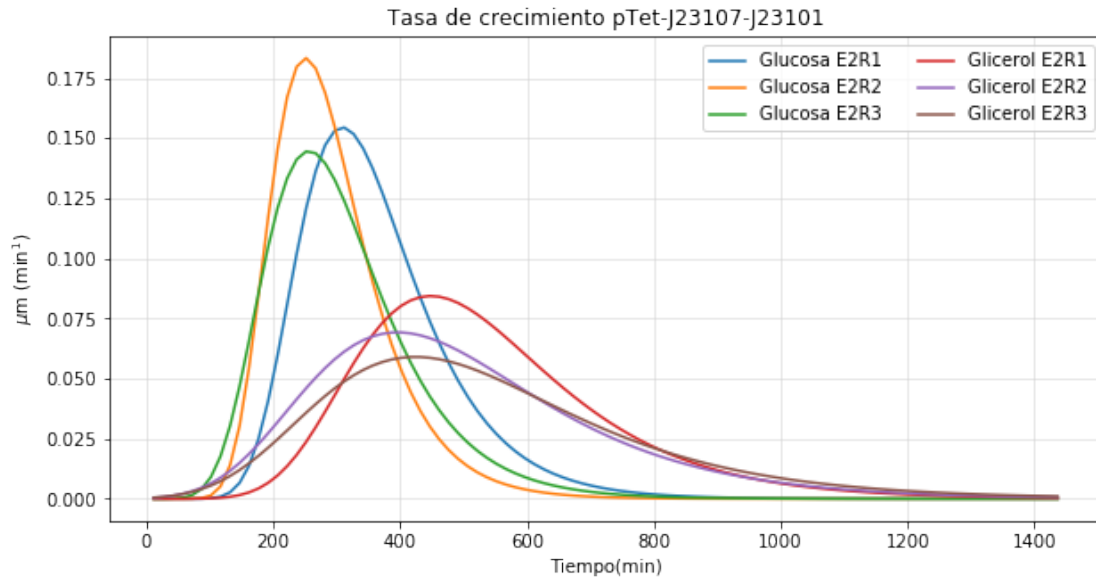
```
In [116]: #Tasas control réplicas glicerol
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento pTet-J23107-J23101 Glicerol 0,2%')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy16,label='growth rate E2R1')
plt.plot(tt[:-1],dy17,label='growth rate E2R2')
plt.plot(tt[:-1],dy18,label='growth rate E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[116]: <matplotlib.legend.Legend at 0x267f44743c8>
```



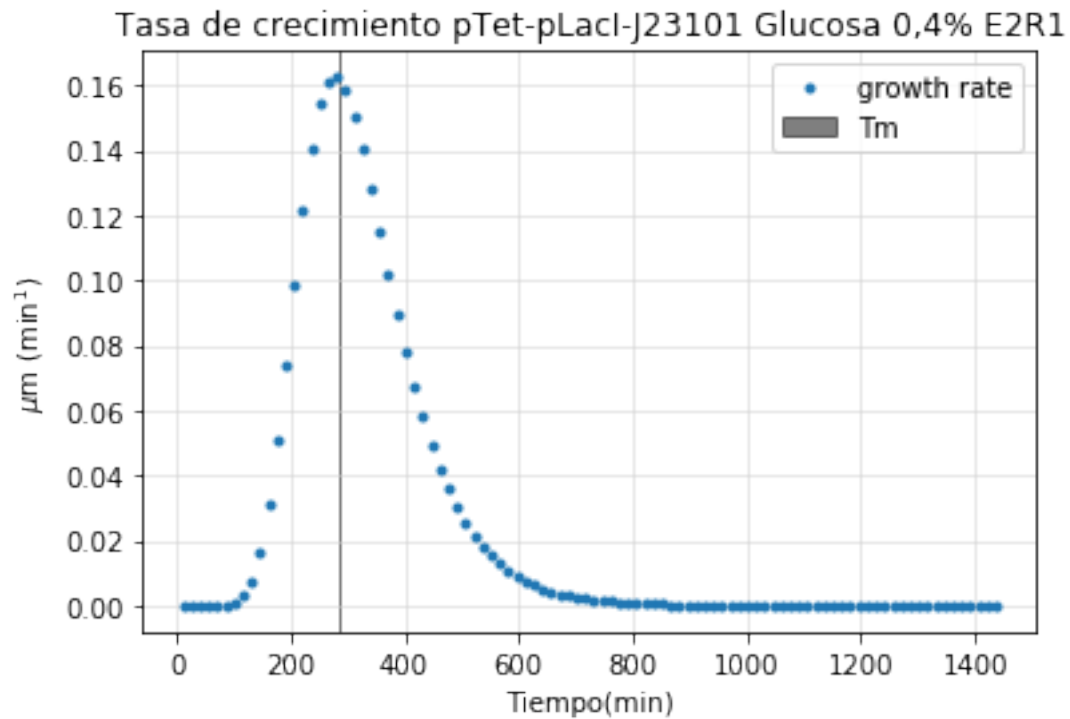
```
In [117]: plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento pTet-J23107-J23101')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$m (min$^{-1}$)')
plt.plot(tt[:-1],dy13,label='Glucosa E2R1')
plt.plot(tt[:-1],dy14,label='Glucosa E2R2')
plt.plot(tt[:-1],dy15,label='Glucosa E2R3')
plt.plot(tt[:-1],dy16,label='Glicerol E2R1')
plt.plot(tt[:-1],dy17,label='Glicerol E2R2')
plt.plot(tt[:-1],dy18,label='Glicerol E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right',ncol=2)
```

```
Out[117]: <matplotlib.legend.Legend at 0x267f792ee10>
```

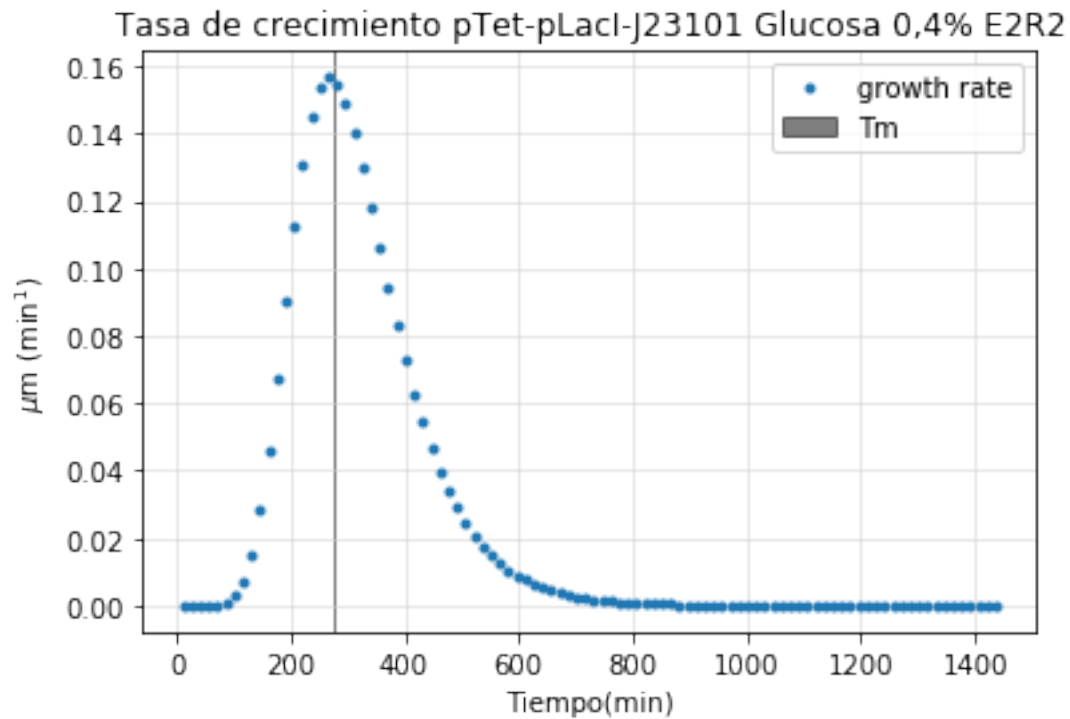
```
In [118]: #tasa de crecimiento
ye19=((A19*np.exp(-np.exp((((um19*np.exp(1))/A19)*(119-tt))+1))))
#Con diff
dy19=(np.diff(ye19))
plt.figure()
plt.title('Tasa de crecimiento pTet-pLacI-J23101 Glucosa 0,4% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm19,tm19, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy19,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[118]: <matplotlib.legend.Legend at 0x267f79d4f60>
```



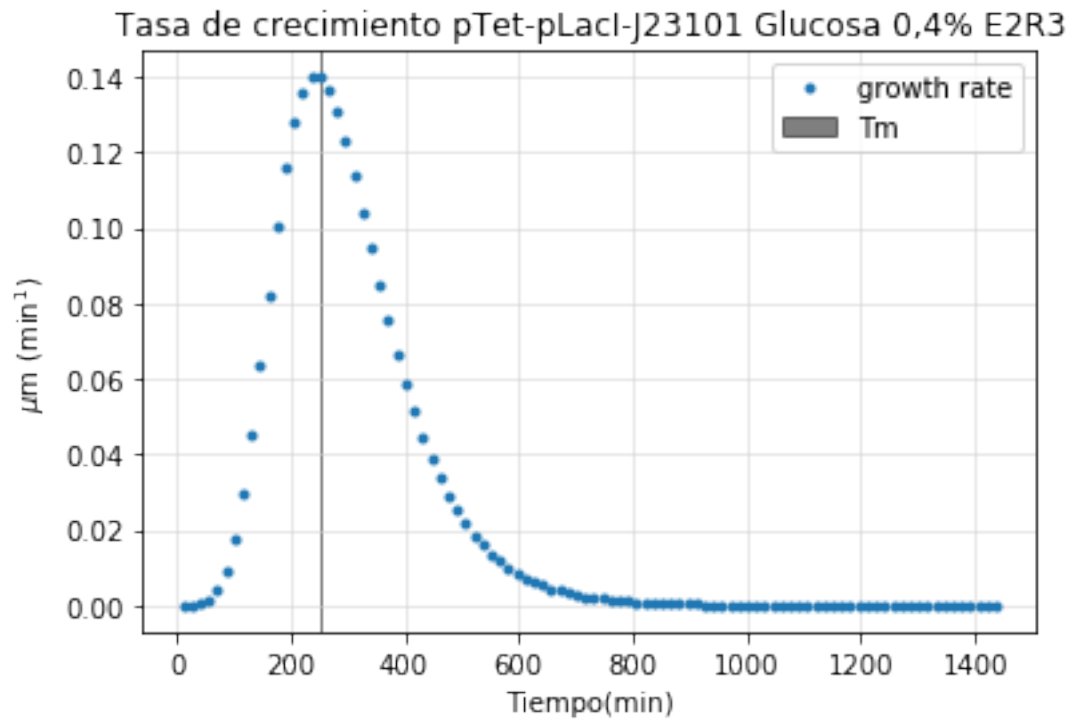
```
In [119]: #tasa de crecimiento
ye20=((A20*np.exp(-np.exp(((um20*np.exp(1))/A20)*(120-tt))+1))))
#Con diff
dy20=(np.diff(ye20))
plt.figure()
plt.title('Tasa de crecimiento pTet-pLacI-J23101 Glucosa 0,4% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm20,tm20, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy20,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[119]: <matplotlib.legend.Legend at 0x267f7aa5b00>
```



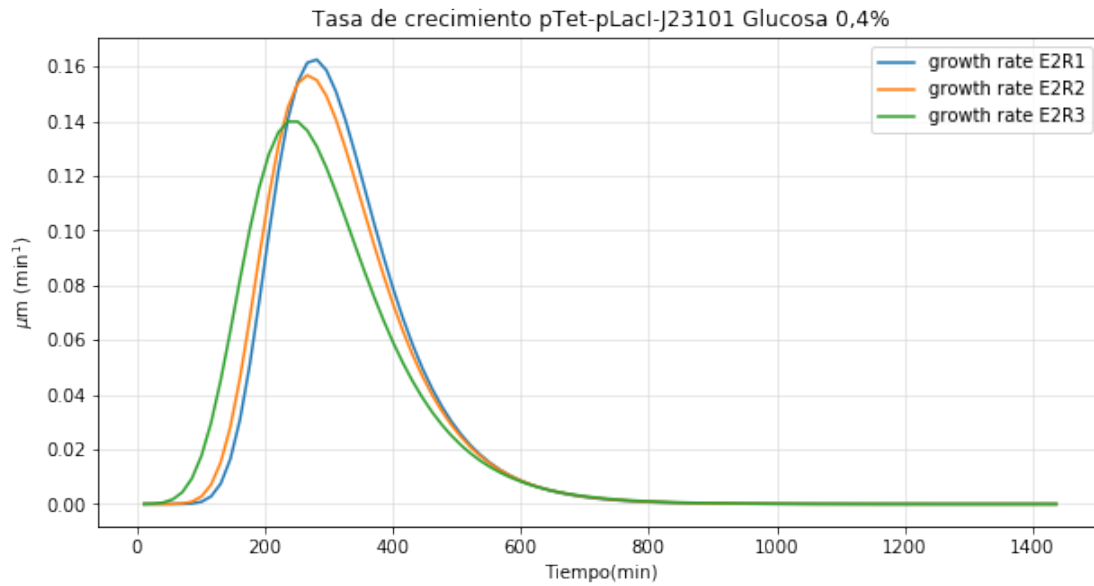
```
In [120]: #tasa de crecimiento
ye21=((A21*np.exp(-np.exp((((um21*np.exp(1))/A21)*(121-tt))+1))))
#Con diff
dy21=(np.diff(ye21))
plt.figure()
plt.title('Tasa de crecimiento pTet-pLacI-J23101 Glucosa 0,4% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm21,tm21, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy21,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[120]: <matplotlib.legend.Legend at 0x267f7b3beb8>
```



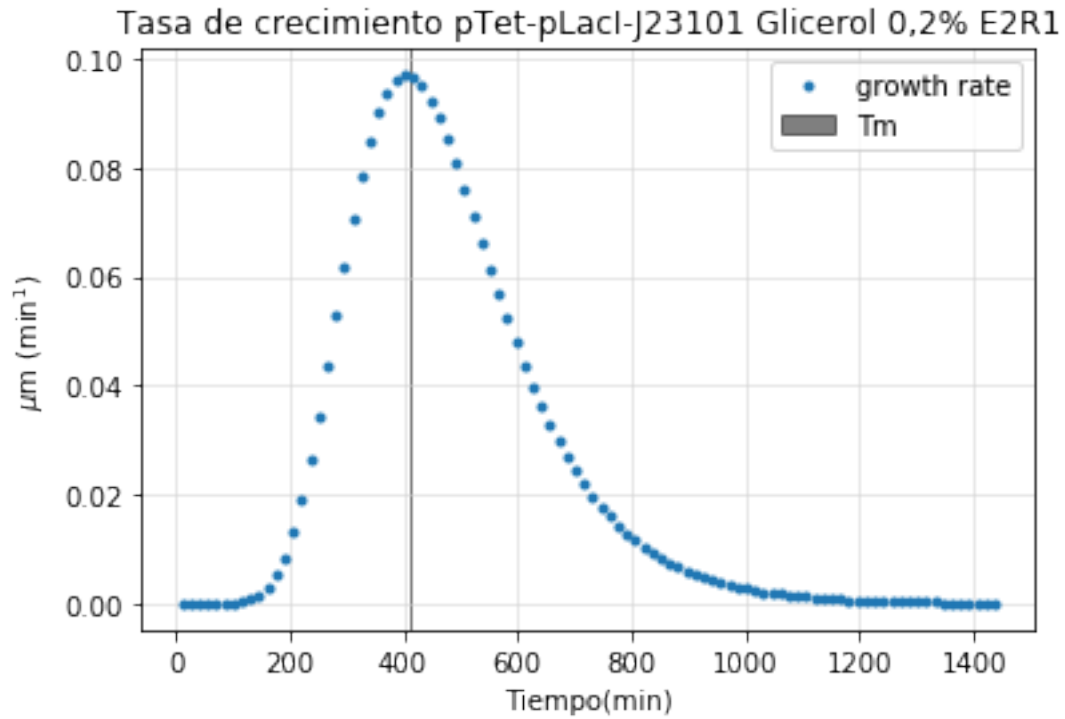
```
In [121]: #Tasas plux76-ptet-J23101 réplicas glucosa
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento pTet-pLacI-J23101 Glucosa 0,4%')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy19,label='growth rate E2R1')
plt.plot(tt[:-1],dy20,label='growth rate E2R2')
plt.plot(tt[:-1],dy21,label='growth rate E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[121]: <matplotlib.legend.Legend at 0x267f7c389e8>
```



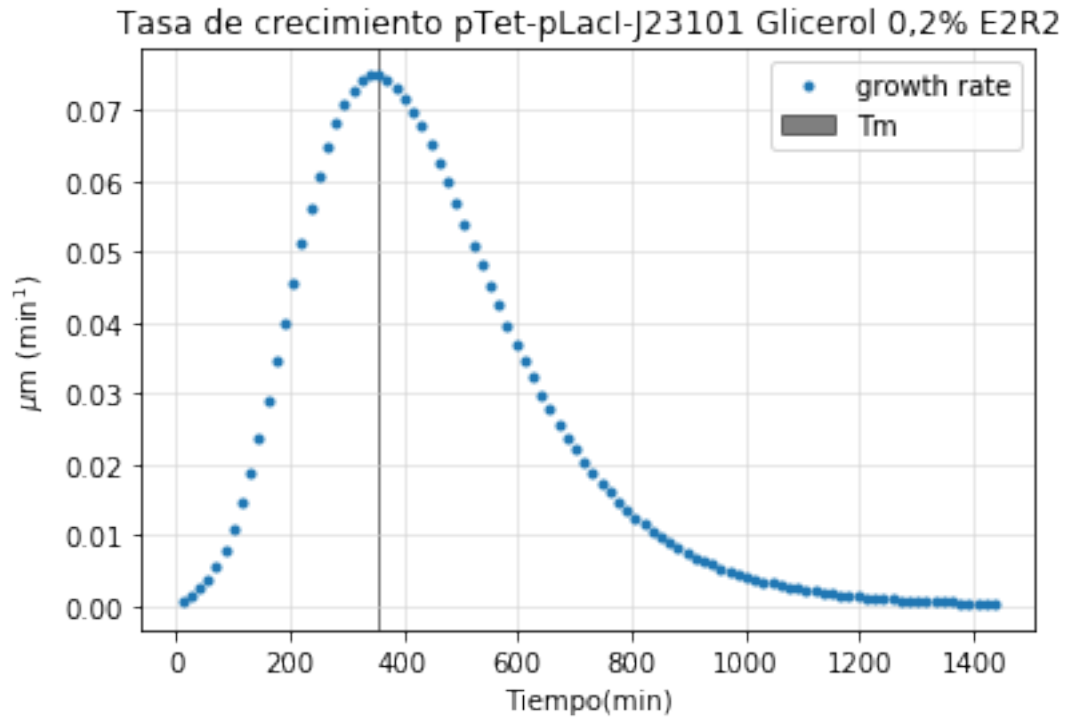
```
In [122]: #tasa de crecimiento
ye22=((A22*np.exp(-np.exp((((um22*np.exp(1))/A22)*(122-tt))+1))))
#Con diff
dy22=(np.diff(ye22))
plt.figure()
plt.title('Tasa de crecimiento pTet-pLacI-J23101 Glicerol 0,2% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm22,tm22, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy22,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[122]: <matplotlib.legend.Legend at 0x267f7d027b8>
```



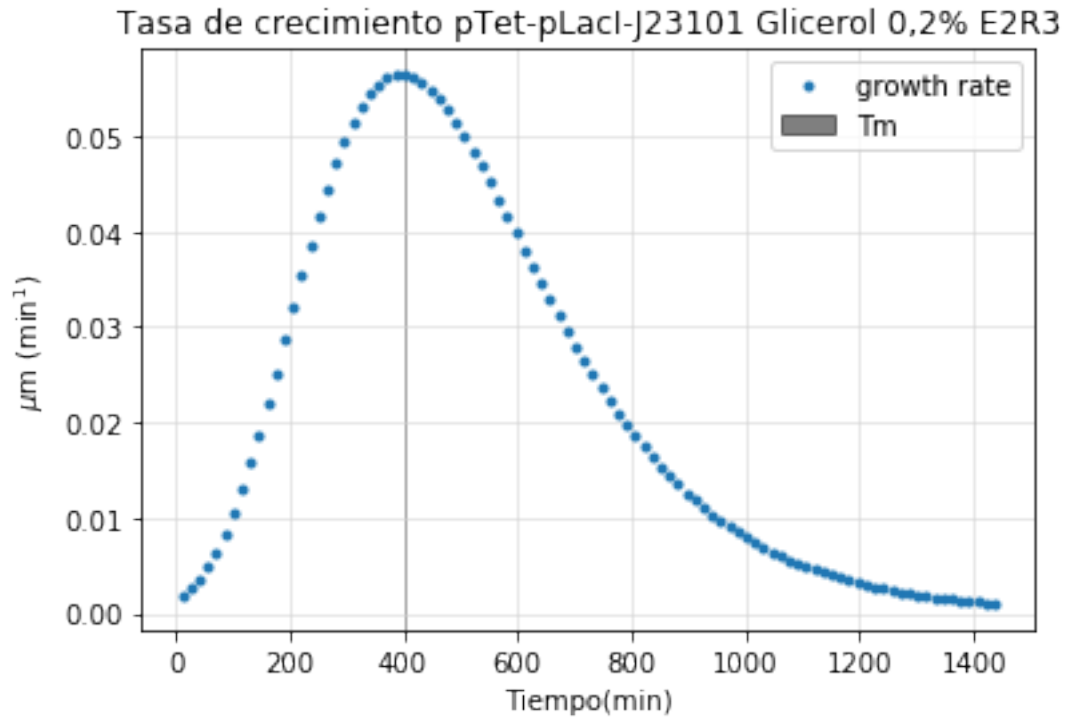
```
In [123]: #tasa de crecimiento
ye23=((A23*np.exp(-np.exp((((um23*np.exp(1))/A23)*(123-tt))+1))))
#Con diff
dy23=(np.diff(ye23))
plt.figure()
plt.title('Tasa de crecimiento pTet-pLacI-J23101 Glicerol 0,2% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm23,tm23, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy23,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[123]: <matplotlib.legend.Legend at 0x267f7dd57f0>
```



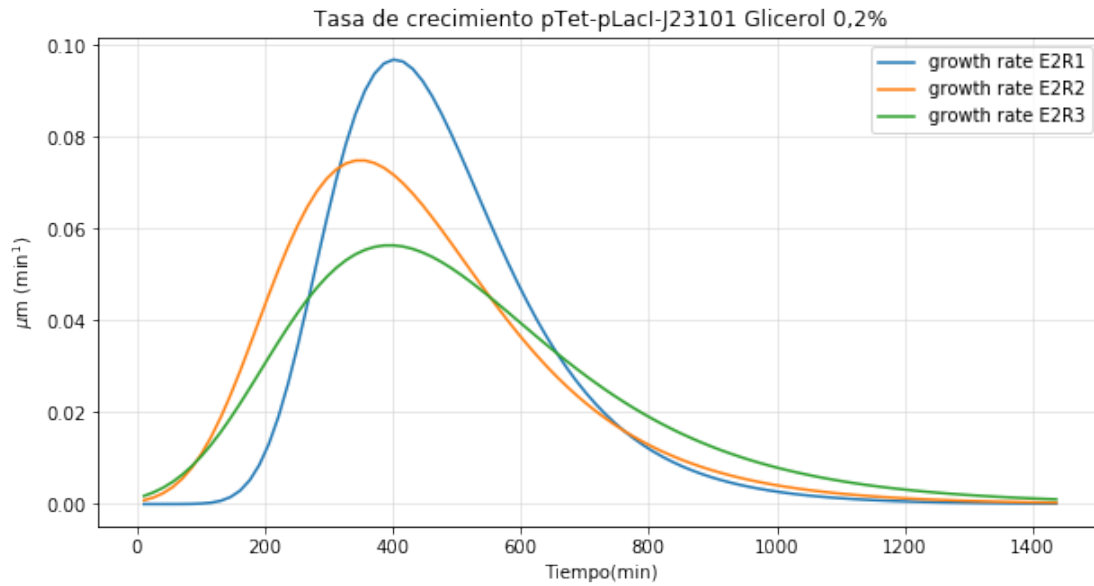
```
In [124]: #tasa de crecimiento
ye24=((A24*np.exp(-np.exp((((um24*np.exp(1))/A24)*(124-tt))+1))))
#Con diff
dy24=(np.diff(ye24))
plt.figure()
plt.title('Tasa de crecimiento pTet-pLacI-J23101 Glicerol 0,2% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm24,tm24, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy24,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[124]: <matplotlib.legend.Legend at 0x267f7ea5fd0>
```



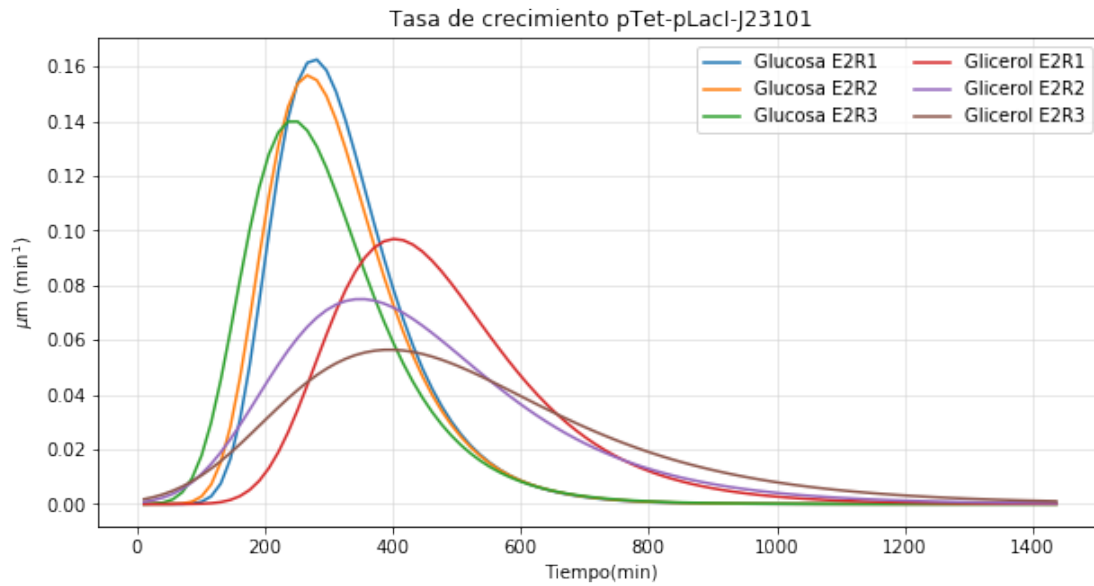
```
In [125]: #Tasas control réplicas glicerol
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento pTet-pLacI-J23101 Glicerol 0,2%')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy22,label='growth rate E2R1')
plt.plot(tt[:-1],dy23,label='growth rate E2R2')
plt.plot(tt[:-1],dy24,label='growth rate E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[125]: <matplotlib.legend.Legend at 0x267f7f76d68>
```

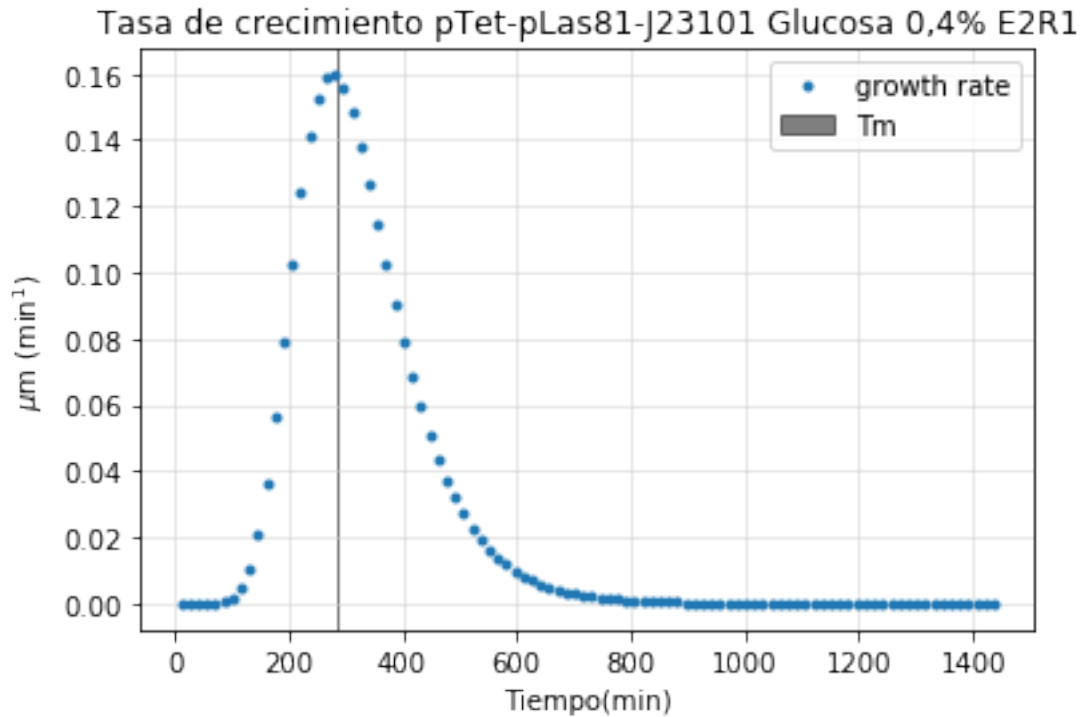
```
In [126]: #Tasas control réplicas
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento pTet-pLacI-J23101')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy19,label='Glucosa E2R1')
plt.plot(tt[:-1],dy20,label='Glucosa E2R2')
plt.plot(tt[:-1],dy21,label='Glucosa E2R3')
plt.plot(tt[:-1],dy22,label='Glicerol E2R1')
plt.plot(tt[:-1],dy23,label='Glicerol E2R2')
plt.plot(tt[:-1],dy24,label='Glicerol E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc='upper right',ncol=2)
```

```
Out[126]: <matplotlib.legend.Legend at 0x267f805ed68>
```



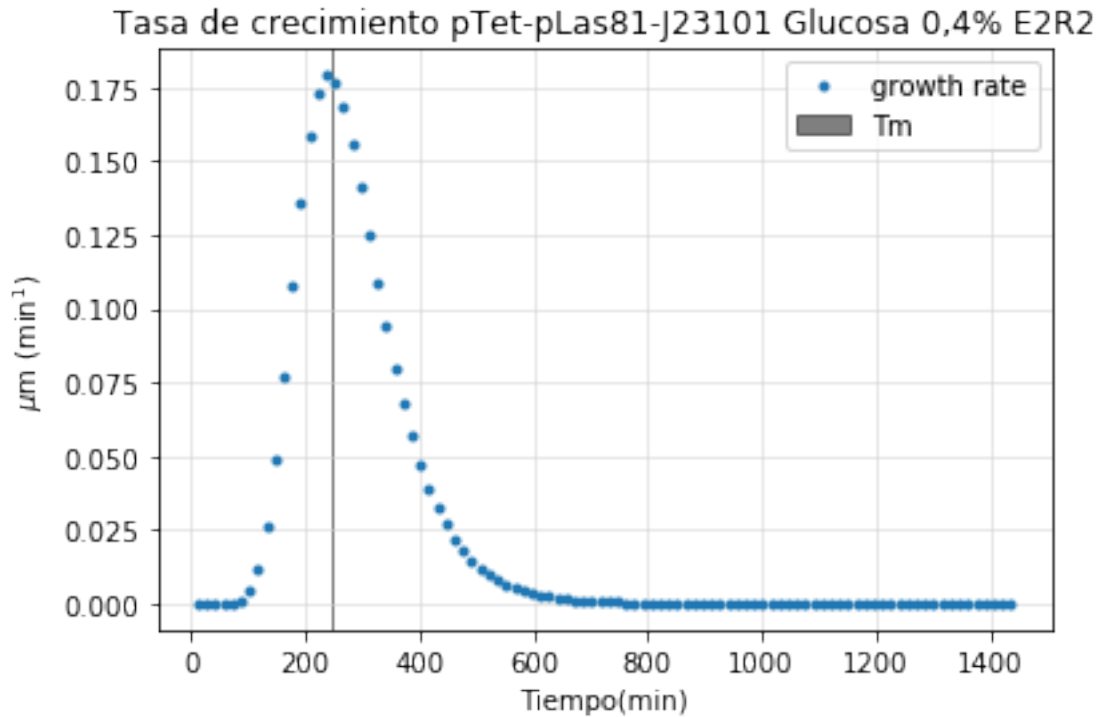
```
In [127]: #tasa de crecimiento
ye25=((A25*np.exp(-np.exp((((um25*np.exp(1))/A25)*(125-tt))+1))))
#Con diff
dy25=(np.diff(ye25))
plt.figure()
plt.title('Tasa de crecimiento pTet-pLas81-J23101 Glucosa 0,4% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm25,tm25, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy25,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

Out[127]: <matplotlib.legend.Legend at 0x267f8140dd8>



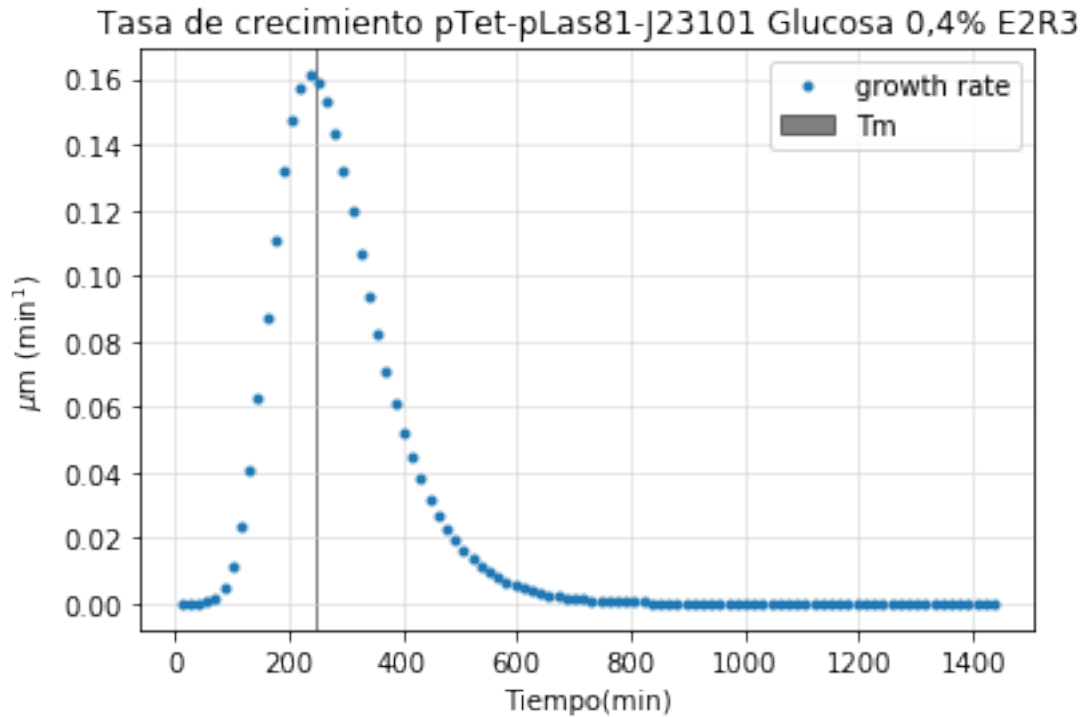
```
In [128]: #tasa de crecimiento
ye26=((A26*np.exp(-np.exp((((um26*np.exp(1))/A26)*(126-tt))+1))))
#Con diff
dy26=(np.diff(ye26))
plt.figure()
plt.title('Tasa de crecimiento pTet-pLas81-J23101 Glucosa 0,4% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm26,tm26, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy26,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[128]: <matplotlib.legend.Legend at 0x267f8218978>
```



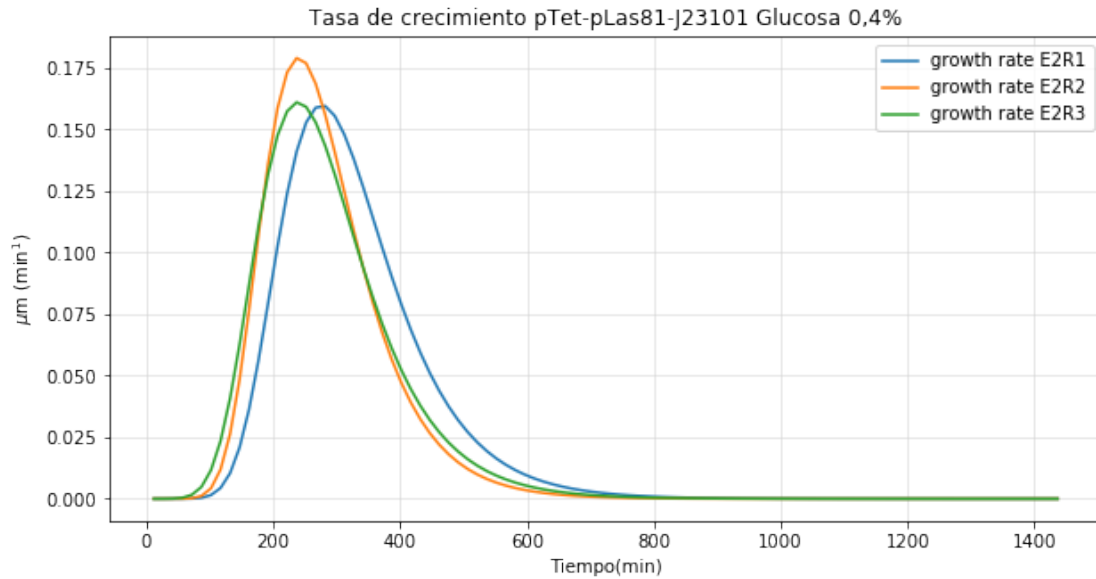
```
In [129]: #tasa de crecimiento
ye27=((A27*np.exp(-np.exp((((um27*np.exp(1))/A27)*(127-tt))+1))))
#Con diff
dy27=(np.diff(ye27))
plt.figure()
plt.title('Tasa de crecimiento pTet-pLas81-J23101 Glucosa 0,4% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm27,tm27, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy27,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[129]: <matplotlib.legend.Legend at 0x267f92c0940>
```



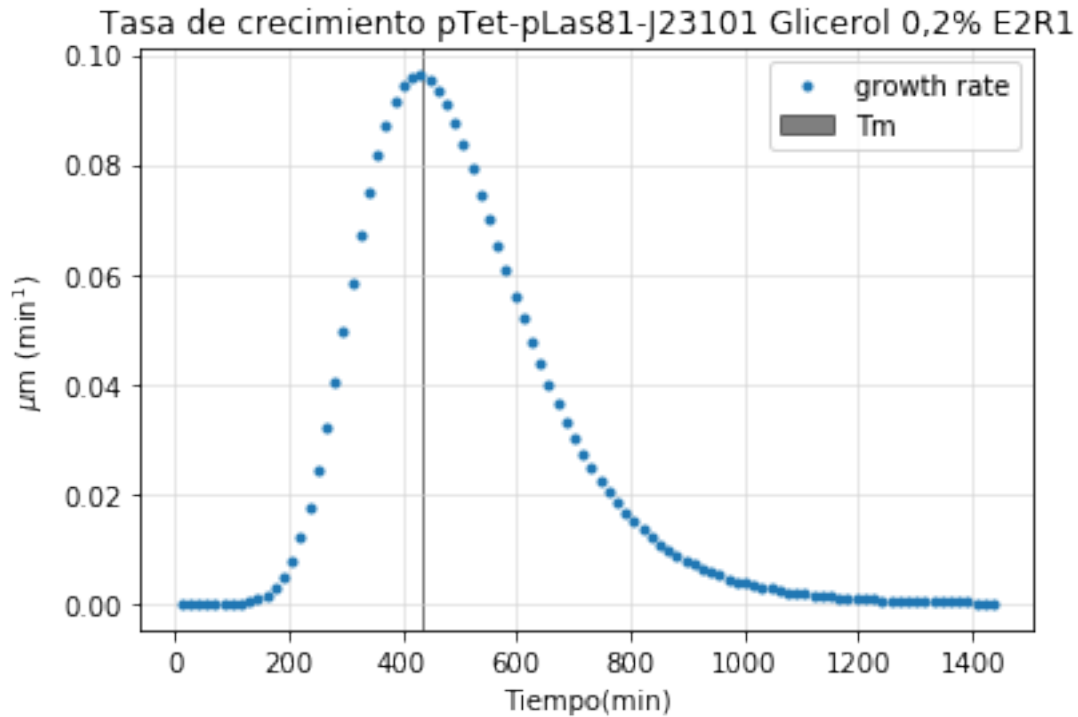
```
In [130]: #Tasas plux76-plac-J23101 réplicas glucosa
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento pTet-pLas81-J23101 Glucosa 0,4%')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy25,label='growth rate E2R1')
plt.plot(tt[:-1],dy26,label='growth rate E2R2')
plt.plot(tt[:-1],dy27,label='growth rate E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[130]: <matplotlib.legend.Legend at 0x267f937eb38>
```



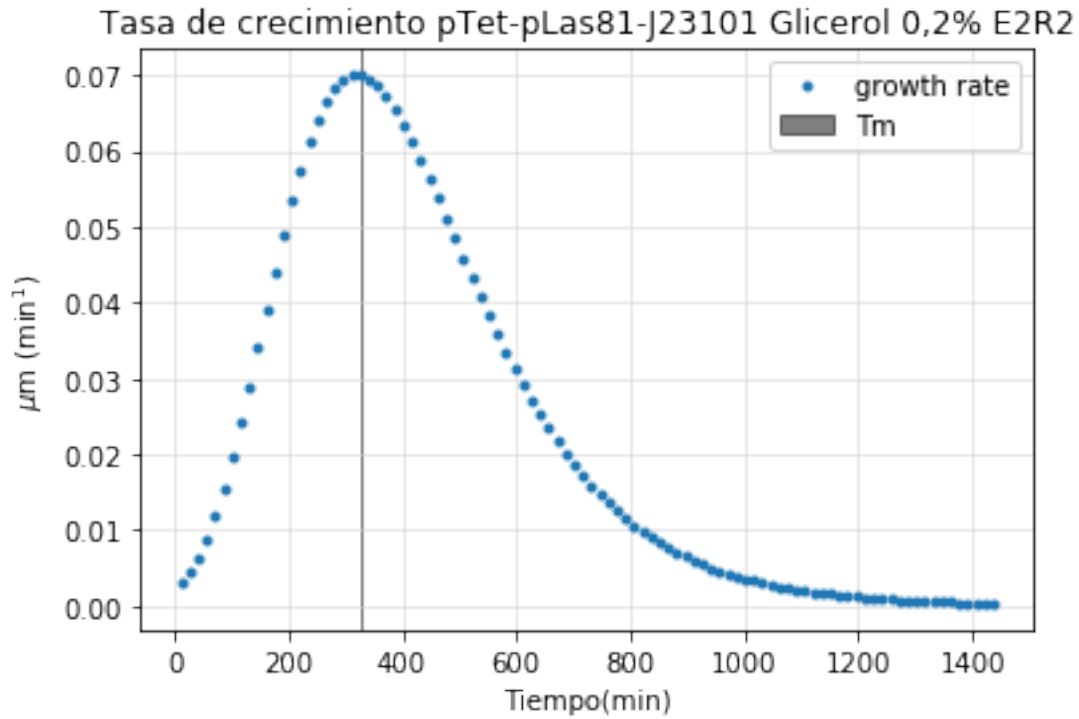
```
In [131]: #tasa de crecimiento
ye28=((A28*np.exp(-np.exp((((um28*np.exp(1))/A28)*(128-tt))+1))))
#Con diff
dy28=(np.diff(ye28))
plt.figure()
plt.title('Tasa de crecimiento pTet-pLas81-J23101 Glicerol 0,2% E2R1')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm28,tm28, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy28,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[131]: <matplotlib.legend.Legend at 0x267f945bef0>
```



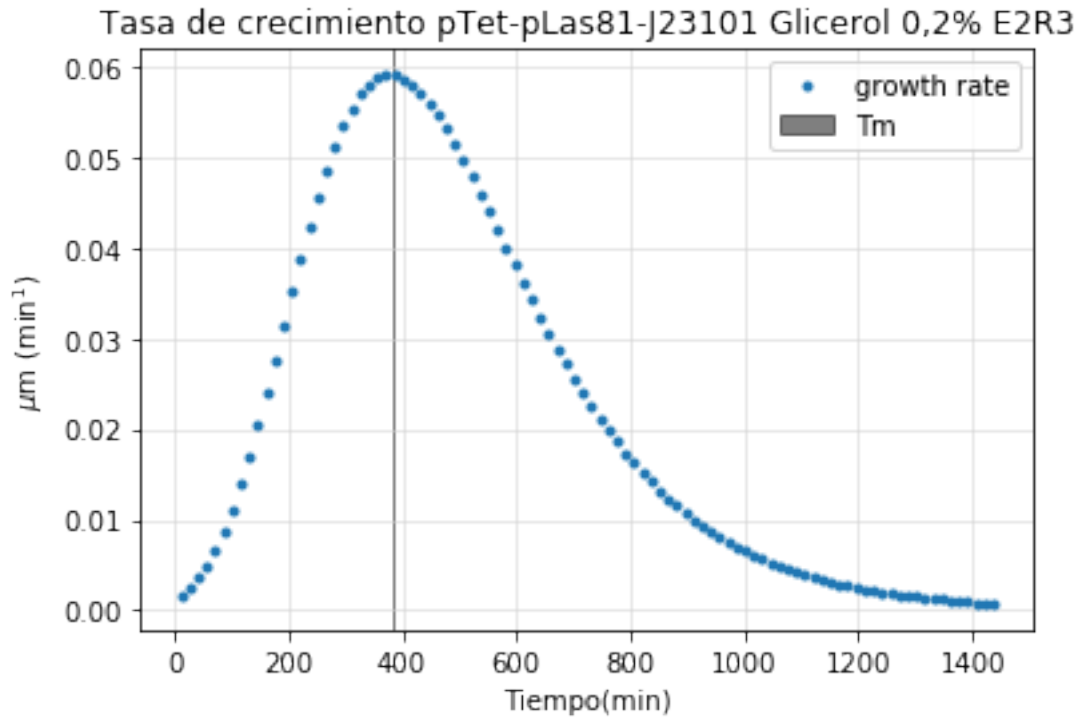
```
In [132]: #tasa de crecimiento
ye29=((A29*np.exp(-np.exp((((um29*np.exp(1))/A29)*(129-tt))+1))))
#Con diff
dy29=(np.diff(ye29))
plt.figure()
plt.title('Tasa de crecimiento pTet-pLas81-J23101 Glicerol 0,2% E2R2')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm29,tm29, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy29,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[132]: <matplotlib.legend.Legend at 0x267f9533f60>
```



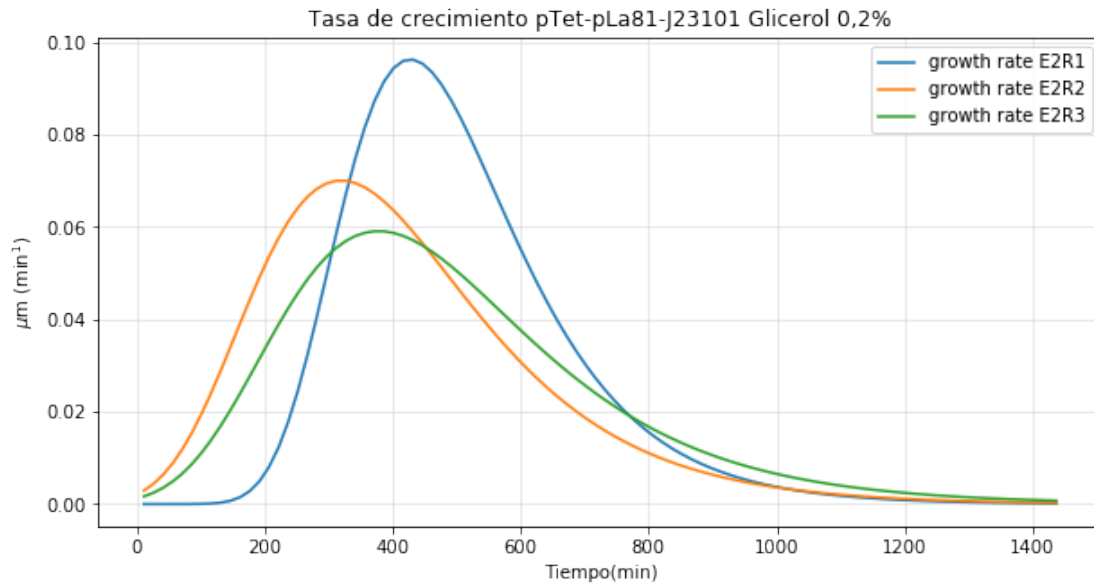
```
In [133]: #tasa de crecimiento
ye30=((A30*np.exp(-np.exp((((um30*np.exp(1))/A30)*(130-tt))+1))))
#Con diff
dy30=(np.diff(ye30))
plt.figure()
plt.title('Tasa de crecimiento pTet-pLas81-J23101 Glicerol 0,2% E2R3')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.axvspan(tm30,tm30, color='k', alpha=0.5, label="Tm")
plt.plot(tt[:-1],dy30,'.',label='growth rate')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[133]: <matplotlib.legend.Legend at 0x267f9612d30>
```

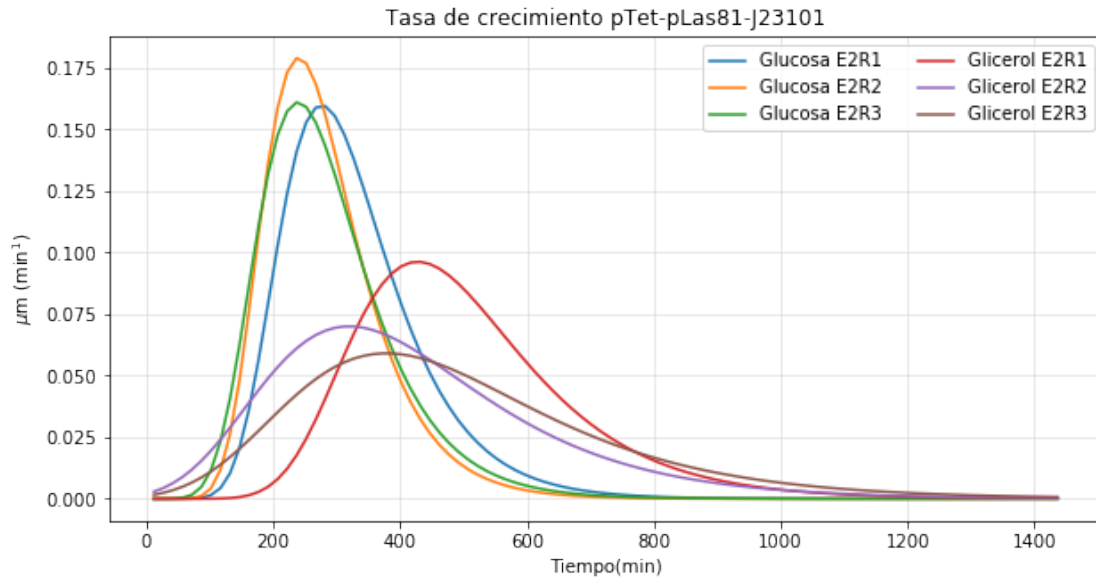
```
In [134]: #Tasas plux76-plac-J23101 réplicas glucosa
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento pTet-pLa81-J23101 Glicerol 0,2%')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy28,label='growth rate E2R1')
plt.plot(tt[:-1],dy29,label='growth rate E2R2')
plt.plot(tt[:-1],dy30,label='growth rate E2R3')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[134]: <matplotlib.legend.Legend at 0x267f96e4ac8>
```



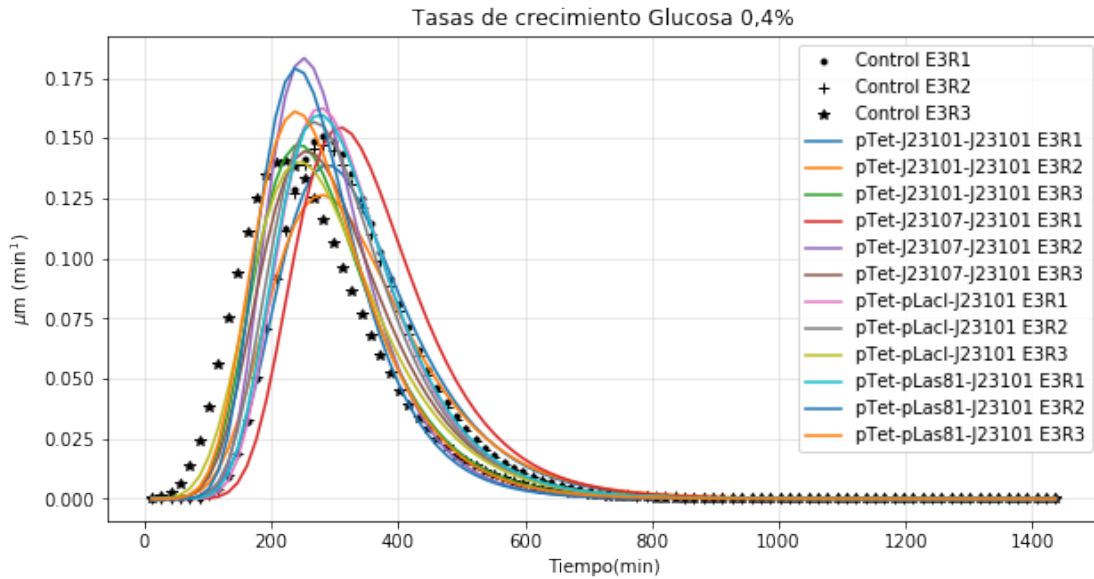
```
In [135]: #Tasas control réplicas
plt.figure(figsize=(10,5))
plt.title('Tasa de crecimiento pTet-pLas81-J23101')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy25,label='Glucosa E2R1')
plt.plot(tt[:-1],dy26,label='Glucosa E2R2')
plt.plot(tt[:-1],dy27,label='Glucosa E2R3')
plt.plot(tt[:-1],dy28,label='Glicerol E2R1')
plt.plot(tt[:-1],dy29,label='Glicerol E2R2')
plt.plot(tt[:-1],dy30,label='Glicerol E2R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc='upper right',ncol=2)
```

```
Out[135]: <matplotlib.legend.Legend at 0x267f97d3ac8>
```



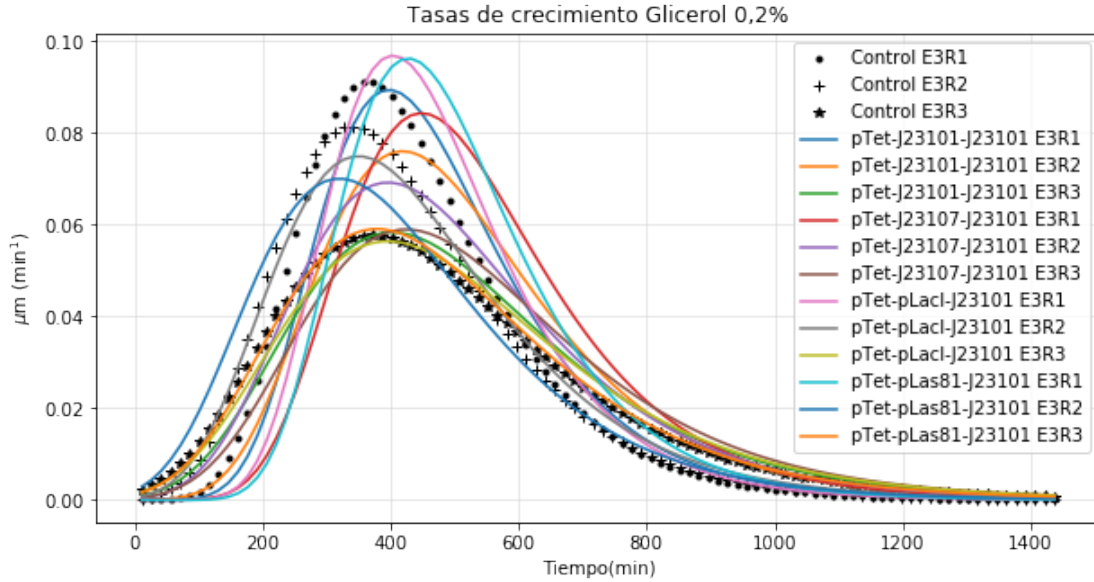
```
In [136]: #Tasas réplicas glucosa
plt.figure(figsize=(10,5))
plt.title('Tasas de crecimiento Glucosa 0,4%')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy1,'k.',label='Control E3R1')
plt.plot(tt[:-1],dy2,'k+',label='Control E3R2')
plt.plot(tt[:-1],dy3,'k*',label='Control E3R3')
plt.plot(tt[:-1],dy7,label='pTet-J23101-J23101 E3R1')
plt.plot(tt[:-1],dy8,label='pTet-J23101-J23101 E3R2')
plt.plot(tt[:-1],dy9,label='pTet-J23101-J23101 E3R3')
plt.plot(tt[:-1],dy13,label='pTet-J23107-J23101 E3R1')
plt.plot(tt[:-1],dy14,label='pTet-J23107-J23101 E3R2')
plt.plot(tt[:-1],dy15,label='pTet-J23107-J23101 E3R3')
plt.plot(tt[:-1],dy19,label='pTet-pLacI-J23101 E3R1')
plt.plot(tt[:-1],dy20,label='pTet-pLacI-J23101 E3R2')
plt.plot(tt[:-1],dy21,label='pTet-pLacI-J23101 E3R3')
plt.plot(tt[:-1],dy25,label='pTet-pLas81-J23101 E3R1')
plt.plot(tt[:-1],dy26,label='pTet-pLas81-J23101 E3R2')
plt.plot(tt[:-1],dy27,label='pTet-pLas81-J23101 E3R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

```
Out[136]: <matplotlib.legend.Legend at 0x267f98e57f0>
```



```
In [137]: #Tasas réplicas glicerol
plt.figure(figsize=(10,5))
plt.title('Tasas de crecimiento Glicerol 0,2%')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy4,'k.',label='Control E3R1')
plt.plot(tt[:-1],dy5,'k+',label='Control E3R2')
plt.plot(tt[:-1],dy6,'k*',label='Control E3R3')
plt.plot(tt[:-1],dy10,label='pTet-J23101-J23101 E3R1')
plt.plot(tt[:-1],dy11,label='pTet-J23101-J23101 E3R2')
plt.plot(tt[:-1],dy12,label='pTet-J23101-J23101 E3R3')
plt.plot(tt[:-1],dy16,label='pTet-J23107-J23101 E3R1')
plt.plot(tt[:-1],dy17,label='pTet-J23107-J23101 E3R2')
plt.plot(tt[:-1],dy18,label='pTet-J23107-J23101 E3R3')
plt.plot(tt[:-1],dy22,label='pTet-pLacI-J23101 E3R1')
plt.plot(tt[:-1],dy23,label='pTet-pLacI-J23101 E3R2')
plt.plot(tt[:-1],dy24,label='pTet-pLacI-J23101 E3R3')
plt.plot(tt[:-1],dy28,label='pTet-pLas81-J23101 E3R1')
plt.plot(tt[:-1],dy29,label='pTet-pLas81-J23101 E3R2')
plt.plot(tt[:-1],dy30,label='pTet-pLas81-J23101 E3R3')
plt.grid(color='lightgray', linestyle='--', linewidth=0.5,zorder=0)
plt.legend(loc='upper right')
```

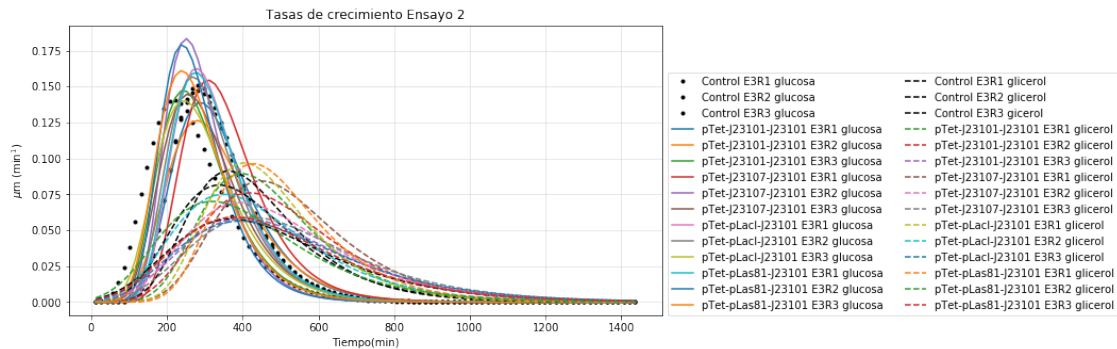
Out[137]: <matplotlib.legend.Legend at 0x267f9a54cf8>



```
In [138]: #Tasas rélicas glucosa
plt.figure(figsize=(10,5))
plt.title('Tasas de crecimiento Ensayo 2')
plt.xlabel('Tiempo(min)')
plt.ylabel(r'$\mu$ (min$^{-1}$)')
plt.plot(tt[:-1],dy1,'k.',label='Control E3R1 glucosa')
plt.plot(tt[:-1],dy2,'k.',label='Control E3R2 glucosa')
plt.plot(tt[:-1],dy3,'k.',label='Control E3R3 glucosa')
plt.plot(tt[:-1],dy7,label='pTet-J23101-J23101 E3R1 glucosa')
plt.plot(tt[:-1],dy8,label='pTet-J23101-J23101 E3R2 glucosa')
plt.plot(tt[:-1],dy9,label='pTet-J23101-J23101 E3R3 glucosa')
plt.plot(tt[:-1],dy13,label='pTet-J23107-J23101 E3R1 glucosa')
plt.plot(tt[:-1],dy14,label='pTet-J23107-J23101 E3R2 glucosa')
plt.plot(tt[:-1],dy15,label='pTet-J23107-J23101 E3R3 glucosa')
plt.plot(tt[:-1],dy19,label='pTet-pLacI-J23101 E3R1 glucosa')
plt.plot(tt[:-1],dy20,label='pTet-pLacI-J23101 E3R2 glucosa')
plt.plot(tt[:-1],dy21,label='pTet-pLacI-J23101 E3R3 glucosa')
plt.plot(tt[:-1],dy25,label='pTet-pLas81-J23101 E3R1 glucosa')
plt.plot(tt[:-1],dy26,label='pTet-pLas81-J23101 E3R2 glucosa')
plt.plot(tt[:-1],dy27,label='pTet-pLas81-J23101 E3R3 glucosa')
plt.plot(tt[:-1],dy4,'k--',label='Control E3R1 glicerol')
plt.plot(tt[:-1],dy5,'k--',label='Control E3R2 glicerol')
plt.plot(tt[:-1],dy6,'k--',label='Control E3R3 glicerol')
plt.plot(tt[:-1],dy10,'--',label='pTet-J23101-J23101 E3R1 glicerol')
plt.plot(tt[:-1],dy11,'--',label='pTet-J23101-J23101 E3R2 glicerol')
plt.plot(tt[:-1],dy12,'--',label='pTet-J23101-J23101 E3R3 glicerol')
plt.plot(tt[:-1],dy16,'--',label='pTet-J23107-J23101 E3R1 glicerol')
plt.plot(tt[:-1],dy17,'--',label='pTet-J23107-J23101 E3R2 glicerol')
```

```
plt.plot(tt[:-1],dy18,'--',label='pTet-J23107-J23101 E3R3 glicerol')
plt.plot(tt[:-1],dy22,'--',label='pTet-pLacI-J23101 E3R1 glicerol')
plt.plot(tt[:-1],dy23,'--',label='pTet-pLacI-J23101 E3R2 glicerol')
plt.plot(tt[:-1],dy24,'--',label='pTet-pLacI-J23101 E3R3 glicerol')
plt.plot(tt[:-1],dy28,'--',label='pTet-pLas81-J23101 E3R1 glicerol')
plt.plot(tt[:-1],dy29,'--',label='pTet-pLas81-J23101 E3R2 glicerol')
plt.plot(tt[:-1],dy30,'--',label='pTet-pLas81-J23101 E3R3 glicerol')
plt.grid(color='lightgray', linestyle='-', linewidth=0.5,zorder=0)
plt.legend(loc=(1.01,0.0),ncol=2)
```

Out[138]: <matplotlib.legend.Legend at 0x267f82180f0>



In []: