modification information for 3 nucleosomes for different values of kon and koff

April 18, 2021

```
[7]: #avg rate of modification vs kon/koff
    import numpy as np
    import math
    import random
    import matplotlib.pyplot as plt
    def mon(n,kf):
        #i = int(input("Enter the number of nucleosomes = "))
        i = 3
        x_array = np.zeros((n,i))
        t_array = np.zeros(n)
        t = 0
        mi = 0
        1 = []
        for y in range(n):
            for z in range(i):
                kon = 0.5
                koff = kf
                \#koff = 0.02
                a = 1/(kon+koff)
                pa = kon*a
                pb = koff*a
                r1 = random.uniform(0,1)
                if r1<pa:</pre>
                    mz = 1
                else:
                    mz = 0
                x_array[y,z] = mz
                r2 = random.uniform(0,1)
                dt = (-1)*a*math.log(r2)
                t = t + dt
            t_array[y] = t
        x1 = x_array[:,0].tolist()
```

```
x2 = x_array[:,1].tolist()
x3 = x_array[:,2].tolist()
\#print(x1, x2, x3)
s1 = 0; s2 = 0; s3 = 0; s4 = 0; s5 = 0; s6 = 0; s7 = 0; s0 = 0
for i in range(n):
    if x1[i]==1 and x2[i]==0 and x3[i]==0:
        s1 +=1
    else:
        s1 +=0
    if x2[i] == 1 and x1[i] == 0 and x3[i] == 0:
        s2 +=1
    else:
        s2 +=0
    if x3[i]==1 and x1[i]==0 and x2[i]==0:
        s3 +=1
    else:
        s3 +=0
    if x1[i]==0 and x2[i]==0 and x3[i]==0:
        s0 += 1
    else:
        s0 += 0
    if x1[i]==1 and x2[i]==1 and x3[i]==0:
        s4 += 1
    else:
        s4 += 0
    if x1[i]==0 and x2[i]==1 and x3[i]==1:
        s5 += 1
    else:
        s5 += 0
    if x1[i]==1 and x2[i]==0 and x3[i]==1:
        s6 += 1
    else:
        s6 += 0
    if x1[i]==1 and x2[i]==1 and x3[i]==1:
        s7 += 1
    else:
        s7 += 0
#print(s0,s1,s2,s3,s4,s5,s6,s7,s0)
```

```
p0 = s0/n; p1 = s1/n; p2 = s2/n; p3 = s3/n; p4 = s4/n; p5 = s5/n; p6 = s5/n;
\rightarrowp6 = s6/n; p7 = s7/n
   total = p0+p1+p2+p3+p4+p5+p6+p7
   \#print('probablity\ for\ 8\ microstates\ are\ :\ p0=',p0,'p1=',p1,'p2=',p2,'p3
\rightarrow=',p3,'p4=',p4,'p5=',p5,'p6=',p6,'p7=',p7)
   in0 = 0; in1 = 0; in2 = 0; in3 = 0; in4 = 0; in5 = 0; in6 = 0; in7 = 0
   if p0 == 0:
       in0 = 0
   else:
       in0 = (p0*math.log2(p0))
   if p1 == 0:
       in1 = 0
   else:
       in1 = (p1*math.log2(p1))
   if p2 ==0:
       in2 = 0
       in3 = (p2*math.log2(p2))
   if p3 ==0:
       in3 = 0
   else:
       in3 = (p3*math.log2(p3))
   if p4==0:
       in4 = 0
   else:
       in4 = (p4*math.log2(p4))
   if p5==0:
       in5 = 0
   else:
       in5 = (p5*math.log2(p5))
   if p6==0:
       in6 = 0
   else:
       in6 = (p6*math.log2(p6))
   if p7 ==0:
       in7 = 0
   else:
       in7 = (p7*math.log2(p7))
```

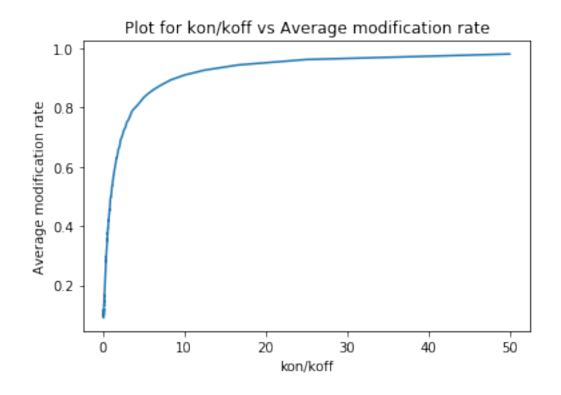
```
nc = np.count_nonzero(x_array== 1, axis=1)
    #print(nc)
    clist = nc.tolist()
    avr = sum(clist)/(n*3)
    return (kon/koff,avr,inf)
a list = []
a2_list = []
i list = []
m_list = []
r_list = []
r2_list = []
xx = []
yy = []
[] = ox
nu = int(input('enter n = '))
for i in np.arange(0.01,5.0,0.01):
    n = nu
    (ri,ai,nf) = mon(n,i)
    a_list.append(ai)
    a2_list.append(1/ai)
    r list.append(ri)
    r2_list.append(1/ri)
    i_list.append(i)
    xx.append(nf)
    if nf >2.22745703 and nf< 2.370568562:</pre>
       m_list.append(i)
print('Possible values of koff when kon = 0.5, for information of modification ⊔

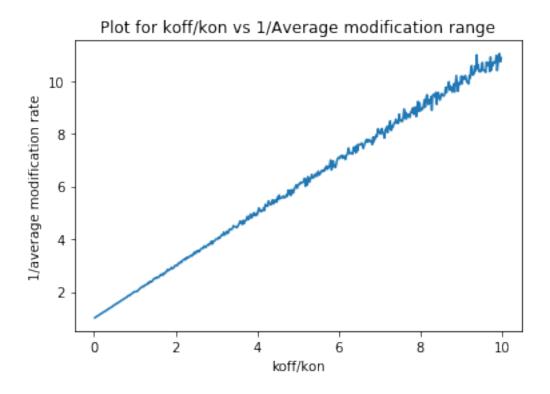
→to be in range of 2.27 to 2.32',m_list)
plt.plot(r_list,a_list)
plt.xlabel('kon/koff')
plt.ylabel('Average modification rate')
plt.title('Plot for kon/koff vs Average modification rate')
plt.show()
plt.plot(r2_list,a2_list)
plt.xlabel('koff/kon')
plt.ylabel('1/average modification rate')
plt.title('Plot for koff/kon vs 1/Average modification range')
plt.show()
plt.plot(r_list,a2_list)
plt.xlabel('kon/koff')
plt.ylabel('1/Average modification rate')
```

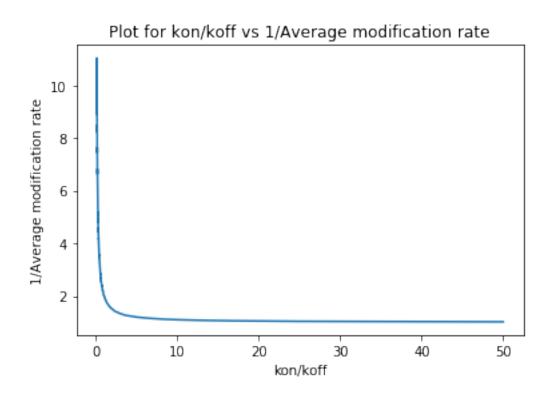
```
plt.title('Plot for kon/koff vs 1/Average modification rate')
plt.show()
plt.plot(r2_list,a_list)
plt.xlabel('koff/kon')
plt.ylabel('Average modification rate')
plt.title('Plot for koff/kon vs Average Modification rate')
plt.show()
plt.plot(r_list,xx)
plt.xlabel('kon/koff')
plt.ylabel('Information in bits')
plt.title('Plot for kon/koff vs Information value')
plt.show()
plt.plot(r2_list,xx)
plt.xlabel('koff/kon')
plt.ylabel('Information in bits')
plt.title('Plot for koff/kon vs Information value')
plt.show()
```

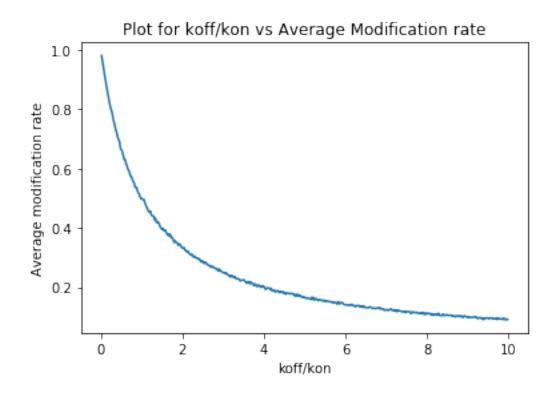
enter n = 10000

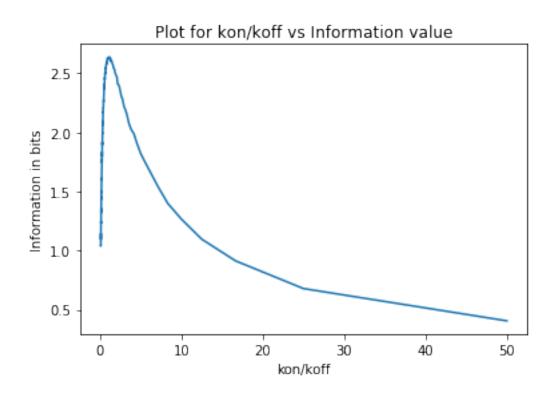
Possible values of koff when kon = 0.5, for information of modification to be in range of 2.27 to 2.32 [0.180000000000000, 0.19, 0.2, 0.99, 1.0, 1.01, 1.02, 1.03, 1.04, 1.05, 1.06, 1.07, 1.08, 1.09, 1.1, 1.11, 1.12, 1.130000000000001, 1.1400000000000001, 1.150000000000001, 1.160000000000001, 1.17, 1.19]

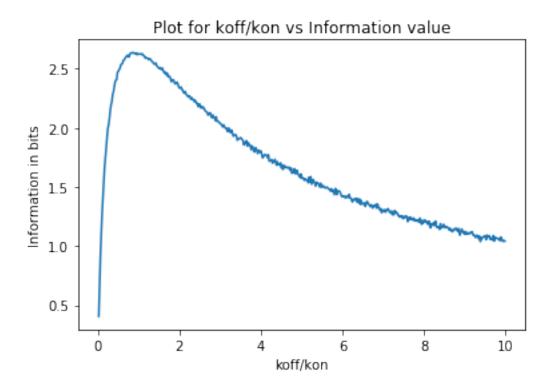












```
[5]: #avg rate of modification vs kon/koff
    import numpy as np
    import math
    import random
    import matplotlib.pyplot as plt
    %matplotlib inline
    def mon(n,kn):
        #i = int(input("Enter the number of nucleosomes = "))
        i = 3
        x_array = np.zeros((n,i))
        t_array = np.zeros(n)
        t = 0
        mi = 0
        1 = []
        for y in range(n):
            for z in range(i):
                koff = 0.5
                kon = kn
                \#koff = 0.02
                a = 1/(kon+koff)
                pa = kon*a
                pb = koff*a
```

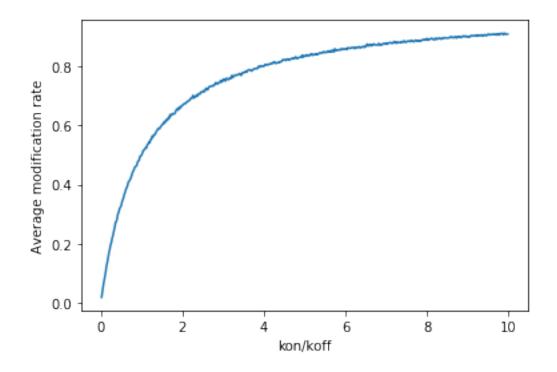
```
r1 = random.uniform(0,1)
        if r1<pa:
            mz = 1
        else:
            mz = 0
        x_{array}[y,z] = mz
        r2 = random.uniform(0,1)
        dt = (-1)*a*math.log(r2)
        t = t + dt
    t_array[y] = t
x1 = x_array[:,0].tolist()
x2 = x_array[:,1].tolist()
x3 = x_array[:,2].tolist()
\#print(x1,x2,x3)
s1 = 0; s2 = 0; s3 = 0; s4 = 0; s5 = 0; s6 = 0; s7 = 0; s0 = 0
for i in range(n):
    if x1[i]==1 and x2[i]==0 and x3[i]==0:
        s1 +=1
    else:
        s1 +=0
    if x2[i] == 1 and x1[i] == 0 and x3[i] == 0:
        s2 +=1
    else:
        s2 +=0
    if x3[i]==1 and x1[i]==0 and x2[i]==0:
        s3 +=1
    else:
        s3 +=0
    if x1[i]==0 and x2[i]==0 and x3[i]==0:
        s0 += 1
    else:
        s0 += 0
    if x1[i]==1 and x2[i]==1 and x3[i]==0:
        s4 += 1
    else:
        s4 += 0
    if x1[i]==0 and x2[i]==1 and x3[i]==1:
        s5 += 1
    else:
```

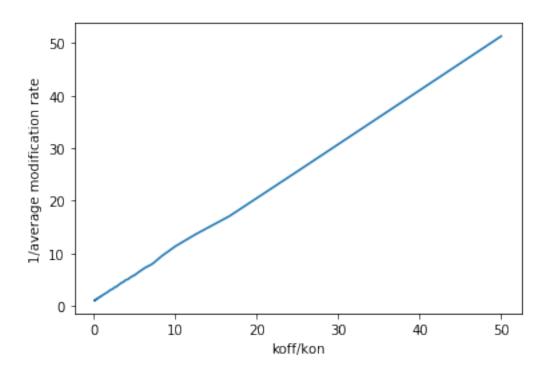
```
s5 += 0
       if x1[i]==1 and x2[i]==0 and x3[i]==1:
       else:
           s6 += 0
       if x1[i]==1 and x2[i]==1 and x3[i]==1:
           s7 += 1
       else:
           s7 += 0
   #print(s0,s1,s2,s3,s4,s5,s6,s7,s0)
   p0 = s0/n; p1 = s1/n; p2 = s2/n; p3 = s3/n; p4 = s4/n; p5 = s5/n; p6 = s5/n;
\rightarrowp6 = s6/n; p7 = s7/n
  total = p0+p1+p2+p3+p4+p5+p6+p7
   \#print('probablity\ for\ 8\ microstates\ are\ :\ p0=',p0,'p1=',p1,'p2=',p2,'p3
\rightarrow=',p3,'p4=',p4,'p5=',p5,'p6=',p6,'p7=',p7)
   in0 = 0; in1 = 0; in2 = 0; in3 = 0; in4 = 0; in5 = 0; in6 = 0; in7 = 0
   if p0 == 0:
       in0 = 0
   else:
       in0 = (p0*math.log2(p0))
   if p1 == 0:
       in1 = 0
   else:
       in1 = (p1*math.log2(p1))
   if p2 ==0:
       in2 = 0
   else:
       in3 = (p2*math.log2(p2))
   if p3 ==0:
       in3 = 0
   else:
       in3 = (p3*math.log2(p3))
   if p4==0:
       in4 = 0
       in4 = (p4*math.log2(p4))
   if p5==0:
       in5 = 0
```

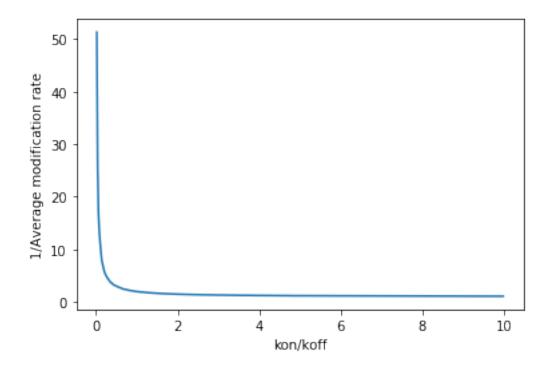
```
else:
        in5 = (p5*math.log2(p5))
    if p6==0:
        in6 = 0
    else:
        in6 = (p6*math.log2(p6))
    if p7 ==0:
        in7 = 0
    else:
        in7 = (p7*math.log2(p7))
    inf = -(in0+in1+in2+in3+in4+in5+in6+in7)
    nc = np.count_nonzero(x_array== 1, axis=1)
    #print(nc)
    clist = nc.tolist()
    avr = sum(clist)/(n*3)
    return (kon/koff,avr,inf)
a_list = []
a2_list = []
i_list = []
m_list = []
r_list = []
r2_list = []
xx = []
yy = []
[] = ox
nu = int(input('enter n = '))
for i in np.arange(0.01,5.0,0.01):
   n = nu
    (ri,ai,nf) = mon(n,i)
    a_list.append(ai)
    a2_list.append(1/ai)
    r_list.append(ri)
   r2_list.append(1/ri)
    i_list.append(i)
    xx.append(nf)
    if nf >2.22745703 and nf< 2.370568562:</pre>
        m_list.append(i)
print(m_list)
plt.plot(r_list,a_list)
plt.xlabel('kon/koff')
plt.ylabel('Average modification rate')
```

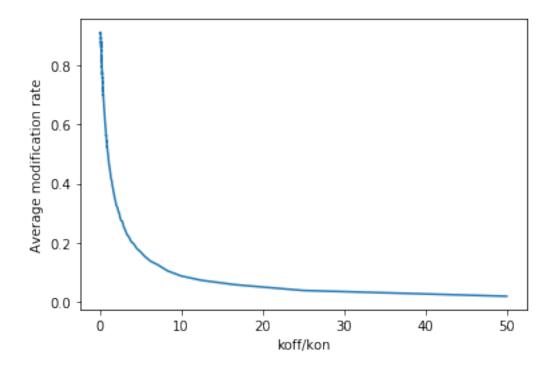
```
plt.show()
plt.plot(r2_list,a2_list)
plt.xlabel('koff/kon')
plt.ylabel('1/average modification rate')
plt.show()
plt.plot(r_list,a2_list)
plt.xlabel('kon/koff')
plt.ylabel('1/Average modification rate')
plt.show()
plt.plot(r2_list,a_list)
plt.xlabel('koff/kon')
plt.ylabel('Average modification rate')
plt.show()
plt.plot(r_list,xx)
plt.xlabel('kon/koff')
plt.ylabel('Information in bits')
plt.show()
plt.plot(r2_list,xx)
plt.xlabel('koff/kon')
plt.ylabel('Information in bits')
plt.show()
```

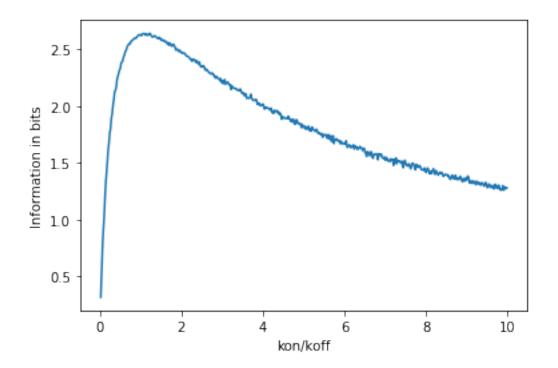
```
enter n = 10000
[0.210000000000002, 0.22, 0.23, 0.24000000000002, 0.25, 1.23, 1.24, 1.25,
1.26, 1.27, 1.28, 1.29, 1.3, 1.31, 1.32, 1.33, 1.34, 1.35, 1.36, 1.37,
1.380000000000001, 1.3900000000001, 1.4000000000001, 1.41000000000001,
1.42, 1.43, 1.44, 1.45, 1.46, 1.48, 1.53]
```

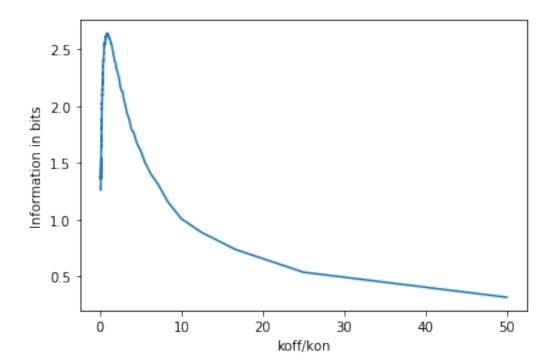












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