

Problem1(e)

November 5, 2015

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In [13]: from __future__ import division
import numpy as np

def simulate_genome(length, alpha):
    read_arr = np.random.choice([0,1], length, p=[1-alpha, alpha])
    reads = ''
    for x in read_arr:
        reads +=str(x)
    return reads

def countoccurences(word, read):
    return read.count(word)

NN = 100000
L = 100
alpha = 0.5
max_iterations = 1000
word = '101'
nn = 100000
c = NN*L/nn

counts = {}
#for n in range(1, nn):
counts[nn] = []
for i in range(1, max_iterations):
    genome = simulate_genome(nn, alpha)
    read_start_positions = np.random.choice(nn-L-1, NN)
    occurences = 0
    for p in read_start_positions:
        read = genome[p:p+L]
        occurences += countoccurences(word, read)
    counts[nn].append(occurences)
```

1 Parameters used:

$\alpha = 0.5$
 $\beta = 0.5$
 $N = 10000$
 $n = 100000$
 $L = 100$
 $c = 100$

$$\lim_{n \rightarrow \infty} \frac{\text{Var}(Y_n)}{n} = c^2 \frac{\alpha^2 \beta (\alpha + \beta - 3\alpha^2 \beta + 2\alpha \beta^2 + 2\alpha \beta + 2\beta^2)}{(\alpha + \beta)^2}$$

$$\lim_{n \rightarrow \infty} \frac{Y_n}{n} = c \frac{\alpha^2 \beta}{\alpha + \beta}$$

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In [20]: a = alpha
         b = 1-alpha
         var = c**2 * ((a**2)*b *(a+b-3*(a**2)*b+2*a*(b**2)+2*a*b+2*b**2))/(nn*(a+b)**2)
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In [25]: print 'Yn/n\t Simulated:{} \t Analytical:{}'.format(np.mean(counts[nn])/nn,c*alpha**2*(1-alpha))
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Yn/n          Simulated:9.84286574575          Analytical:12.5
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In [26]: print 'Var(Yn)/n\t Simulated={} \t Analytical:{}'.format(np.var(counts[nn])/nn**2, var)
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Var(Yn)/n          Simulated=0.0061746138811          Analytical:0.0234375
```

Thus, as seen from above, our analytical and simulated estimate of $\frac{Y_n}{n}$ are 12.83 and 9.83 respectively. I used 1000 iterations, so the bound can improve by increasing the iterations. Similarly the estimate for $\frac{\text{Var}(Y_n)}{n}$ is pretty close (simulated: 0.006, analytical: 0.02)