## Question 1)b)

```
1. # importing the necessary packages
import numpy as np
3. from numpy import sqrt, exp, log
4. from scipy.stats import norm
5.
7.
8. # variables given
9. S_0 = 100.
10. B0 = 1
11. mu = 0.03
12. sigma = 0.25
13. r = 0.01
14.
15. # European Call, using Black Scholes formula
16. K = 120.
17. T = 1
18.
19. d1 = (\log(S_0/K) + (r + (sigma**2)/2)*T) / (sigma * sqrt(T))
20. d2 = d1 - sigma*sqrt(T)
21.
22. Price1 = S_0*norm.cdf(d1) - K*exp(-r*T)*norm.cdf(d2)
23.
24. print("European Call option price (using Black Scholes formula) is:")
25. print(round(Price1,2))
26.
28. # question 1)
29.
30. # defining the stock generation function
31. def generateStock(S_0, r, sigma, T):
32. return S_0 * exp((r - (sigma**2)/2)*T + sigma * sqrt(T) * np.random.standard_normal
 ())
33.
34. # defining payoff function, for this case it's regular European put
35. def payoff1(S, K):
36. return max(S - K, 0)
37.
38. # defining monte carlo integration formula, to make graphing easier
39. def MonteCarlo1(runs):
40. # initialize array that will have payoffs of option
41.
      payoffs = np.zeros(0)
42.
43.
      # looping through
44.
      for i in xrange(runs):
          # generate future stock
45.
          S T = generateStock(S_0, r, sigma, T)
46.
47.
          # append to the payoffs list whatever the payoff is
48.
49.
          payoffs = np.append(payoffs,
50.
                            exp(-r*T)*payoff1(S_T, K))
51.
      return sum(payoffs)/runs, payoffs
52.
53.
54. # setting number of simulation runs
55. \text{ runs} = 10000
```

```
57. # initialize array that will have payoffs of option
58. payoffs = np.zeros(0)
60. # monte carlo calculation of option price
61. Price2, payoffs = MonteCarlo1(runs)
62.
63. var sample = 0
64. payoffs int = payoffs.astype(int)
65. # setting up iterating array
66. it = np.nditer(payoffs, flags = ['f index'])
68. while not it.finished:
       var sample = var sample + (payoffs[it.index] - Price2)**2
       it.iternext()
71. var_sample = var_sample / (runs - 1)
72.
73.
74. # from analytical calculation, we know sigma = Var(X1) = 657.973
76. # 95% confidence, z-score = 1.96
77. size estimate = (1.96**2 * 657.973) / 0.05**2
78.
79. print("European Call option price (using Monte Carlo integration) is: ")
80. print(round(Price2,2))
81. print("Approximation of how many samples needed for 95% confidence (up to dime):")
82. print(int(size estimate))
83.
84. # running Monte Carlo calculation 100 times with the above size estimate
85. # counters for counting number of times the Monte Carlo estimated price
86. # correct up to the dime or not
87. correct = 0
88. incorrect = 0
89. for j in xrange(100):
90.
       price to compare, dummypayoff = MonteCarlo1(int(size estimate))
91.
92.
       if price to compare > Price1 - 0.05 and price to compare < Price1 + 0.05:</pre>
93.
           correct += 1
94.
       else:
95.
           incorrect += 1
96.
97. print("Number of times Monte Carlo estimation was within a dime:")
98. print(correct)
99. print("Number of times Monte Carlo estimation was NOT within a dime:")
        print(incorrect)
```

European Call option price (using Black Scholes formula) is:

3.95

European Call option price (using Monte Carlo integration) is:

3.91

Approximation of how many samples needed for 95% confidence (up to dime): 1011067

#### Question 2) b)

```
1. # importing the necessary packages
2. import numpy as np
3.
6. alpha = 0.6
7. \#lamba = 1
8.
9. losses = np.zeros(0)
10.
11. # loss random variable equation
12. def loss(y):
       return 100000*y - 220000
13.
14.
15. n = 0
16. while n < 100000:
      x = loss(np.random.weibull(alpha))
18. losses = np.append(losses, x)
19.
       n += 1
20.
21. # sort losses array from largest to lowest
22. losses = np.sort(losses)
23. losses[:] = losses[::-1]
25. # select the loss (95% value at risk) that is the specified one
26. # 100000*0.05 = 5000
27.
28. print("Numerical estimation of 95% Value At Risk is:")
29. print(losses[5000])
```

#### **Output:**

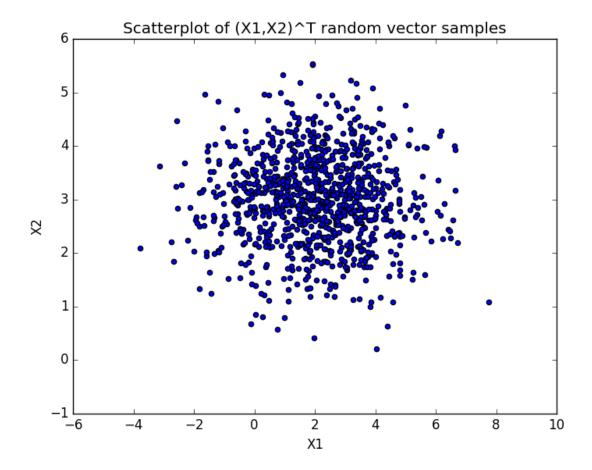
Numerical estimation of 95% Value At Risk is: 401456.414615

## Question 3)c) and d)

```
1. # importing the necessary packages
import numpy as np
3. from numpy import sqrt
4. import matplotlib.pyplot as plt
7.
8. # initial variables
9. covariance = np.zeros([2,2])
10. sigma1 = 3.2
11. sigma2 = 0.7
12. corr = 0.6
13. mu1 = 2
14. \text{ mu2} = 3
15.
16. # setting up array for actual covariance calculation
17. array1 = np.array([[sqrt(sigma1), 0],
                   [0, sqrt(sigma2)]])
19. array2 = np.array([[1, corr],
20.
      [corr, 1]])
21.
22. covariance = (array1.dot(array2)).dot(array1)
23. print("The covariance matrix is: ")
24. print(covariance)
25.
27.
28. # hand-written result of cholesky factorization of A
29. \# A = [[1.78885, 0]]
30. # [0.501996, 0.669328]]
31.
32. # cholesky factorization, from built-in default function
33. A_actual = np.linalg.cholesky(covariance)
34. print("The matrix A, based on built-in cholesky function np.linalg.cholesky(): ")
35. print(A_actual)
36.
38.
39. # cholesky factorization by python algorithm coding
40.
41. # Note, need to do A actual.tolist() to make it python list type
42.
43. def cholesky(A):
44. # establish output S matrix
45.
      n = len(A)
46.
47.
      # setup output array
48.
      S = [[0.0] * n for i in xrange(n)]
49.
50.
      # setting up calculation of individual elements for loop
51.
      for i in xrange(n):
52.
          for j in xrange(i + 1):
53.
             # sigma ij formula from class
54.
             temp = sum(S[i][k] * S[j][k] for k in xrange(j))
55.
56.
             # for the diagnoal case
57.
             if i == j:
58.
                S[i][j] = sqrt(A[i][i] - temp)
```

```
59.
              # for regular case
60.
              else:
61.
                  S[i][j] = (A[i][j] - temp) / S[j][j]
62.
63.
       return S
64.
65. A_cholesky = np.array(cholesky(covariance))
67. print("The matrix A, based on python code algorithm function: ")
68. print(A cholesky)
71.
72. # scatterplot work
73.
74. plt.clf()
75.
76. num = 0
78. # producing 1000 samples
79. while num < 1000:
80.
       X1 = mu1 + A cholesky[0,0]*np.random.standard normal()
81.
82.
       X2 = mu2 + A_cholesky[1,0]*np.random.standard_normal() + A_cholesky[1,1]*np.random.
   standard normal()
83.
84.
       plt.scatter(X1, X2)
85.
       num += 1
87. plt.title("Scatterplot of (X1,X2)^T random vector samples")
88. plt.xlabel("X1")
89. plt.ylabel("X2")
90.
91. plt.show()
```

Figure:



#### Question 4)

```
1. # importing the necessary packages
import numpy as np
3.
5.
6. # initial variables
7. mu = np.array([[-1000],
8.
                         [-700],
9.
                         [300],
                         [-200]])
10.
11.
12. covariance = np.array([[144, 72, 120, 300],
                         [72, 100, 180, 230],
14.
                        [120, 180, 389, 880],
15.
                        [300, 230, 880, 4469]])
16.
17. weights = np.array([0.4, 0.2, 0.3, 0.1])
19. # setup L losses simulation
20. L = np.zeros(0)
21.
22. # get the matrix A thanks to built-in cholesky function
23. A cholesky = np.linalg.cholesky(covariance)
25. # simulation loop for 10,000 samples
26. \text{ num} = 0
27. while num < 10000:
28.
29.
       # this X matrix will hold our Assets
30. X = np.zeros like(mu)
31.
    # creating standard normal random samples matrix
32.
       # each element is a different standard normal random sample
33.
34.
       Z = np.zeros_like(A_cholesky)
35.
       for x in np.nditer(Z, op_flags = ['readwrite']):
36.
           x[...] = np.random.standard_normal()
37.
38.
       # loop through to get assets vector
39.
     for i in xrange(X.size):
40.
41.
           X[i] = mu[i] + sum(A cholesky[i]*Z[i])
42.
43.
       # loss based on calculation
44.
       L = np.append(L, weights.dot(X))
45.
46.
       num += 1
47.
48. # sort L array, from largest to smallest
49.L = np.sort(L)
50. L[:] = L[::-1]
51.
52. # we want 99% Value At Risk. So pick the 10000*0.01 = 100th loss element
53.
54. print("Numerical estimation of 99% Value At Risk is: ")
55. print(L[100])
```

Output: Numerical estimation of 99% Value At Risk is: -445.8

## Question 5) Q5\_scaling.py

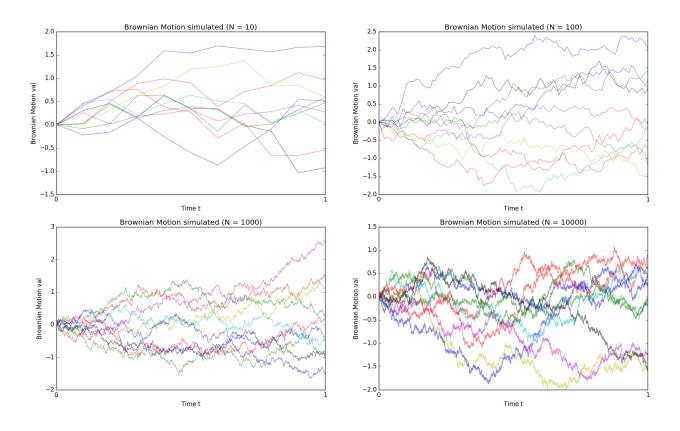
```
1. # importing the necessary packages
import numpy as np
3. from numpy import sqrt
import matplotlib.pyplot as plt
5. import timeit
6.
8.
9. # starting timer
10. start = timeit.default timer()
11.
12. # increment step method
13. def SimBMStep(T, N):
14. # initialize W brownian motion array
      W = np.zeros(N)
15.
16. W = np.append(W, 0) # initial 0 value
17.
18. # loop
19.
      num = 1
    while num <= N:
20.
21.
          W[num] = W[num - 1] + sqrt(T/N)*np.random.standard_normal()
22.
23.
24.
25.
      return W
26.
28.
29. plt.clf()
30.
31. plt.subplot(2,2,1)
33. # setting up t x-axis variable
34. t = np.arange(0.0, 11.0, 1.0)
36. # loop for 10 different paths
37. n = 0
38. while n < 10:
    W = SimBMStep(1., 10)
40. plt.plot(t, W, alpha = 0.5)
41.
      n += 1
42.
43. plt.title("Brownian Motion simulated (N = 10)")
45. ticks = np.arange(t.min(), t.max() + 1, 10)
46. labels = range(ticks.size)
47. plt.xticks(ticks, labels)
48. plt.xlabel("Time t")
50. plt.ylabel("Brownian Motion val")
52. ###################################
53. plt.subplot(2,2,2)
55. # setting up t x-axis variable
56. t = np.arange(0.0, 101.0, 1.0)
```

```
58. # loop for 10 different paths
59. n = 0
60. while n < 10:
61.
       W = SimBMStep(1., 100)
       plt.plot(t, W, alpha = 0.5)
62.
63.
       n += 1
64.
65. plt.title("Brownian Motion simulated (N = 100)")
67. ticks = np.arange(t.min(), t.max() + 1, 100)
68. labels = range(ticks.size)
69. plt.xticks(ticks, labels)
70. plt.xlabel("Time t")
71.
72. plt.ylabel("Brownian Motion val")
74. ##################################
75. plt.subplot(2,2,3)
76.
77. # setting up t x-axis variable
78. t = np.arange(0.0, 1001.0, 1.0)
80. # loop for 10 different paths
81. n = 0
82. while n < 10:
       W = SimBMStep(1., 1000)
       plt.plot(t, W, alpha = 0.5)
85.
       n += 1
86.
87. plt.title("Brownian Motion simulated (N = 1000)")
89. ticks = np.arange(t.min(), t.max() + 1, 1000)
90. labels = range(ticks.size)
91. plt.xticks(ticks, labels)
92. plt.xlabel("Time t")
93.
94. plt.ylabel("Brownian Motion val")
97. plt.subplot(2,2,4)
98.
99. # setting up t x-axis variable
100. t = np.arange(0.0, 10001.0, 1.0)
101.
102.
          # loop for 10 different paths
103.
           n = 0
104.
          while n < 10:
              W = SimBMStep(1., 10000)
105.
              plt.plot(t, W, alpha = 0.5)
106.
107.
              n += 1
108.
           plt.title("Brownian Motion simulated (N = 10000)")
109.
110.
           ticks = np.arange(t.min(), t.max() + 1, 10000)
111.
          labels = range(ticks.size)
112.
           plt.xticks(ticks, labels)
113.
           plt.xlabel("Time t")
114.
115.
           plt.ylabel("Brownian Motion val")
116.
```

```
117.
118. plt.show()
119.
120. # stopping timer
121. stop = timeit.default_timer()
122.
123. print "Step-size increment method runtime = " , stop - start, " seconds"
```

Step-size increment method runtime = 1.27352404594 seconds

## Figure:



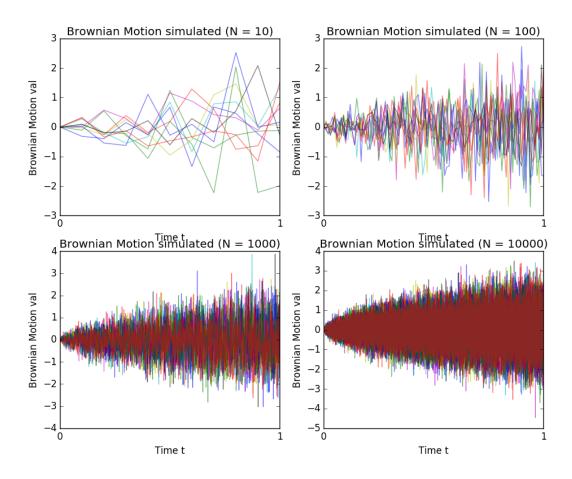
# Q5\_cholesky.py

```
12. # cholesky method
13.
14. def SimBMChol(T, N):
15.
       # initialize W brownian motion array
       W = np.zeros(N)
16.
       W = np.append(W, 0)
                              # initial 0 value
17.
18.
19.
       # looping and iterating through
20.
       for i in xrange(N):
21.
           # create matrix of Z random standard normal samples
22.
23.
           Z = np.zeros(i+1)
24.
           for x in np.nditer(Z, op flags = ['readwrite']):
25.
               x[...] = np.random.standard normal()
26.
27.
           val = sum(sqrt(T/N)*Z)
28.
29.
           W[i+1] = val
30.
31.
       return W
32.
35. plt.clf()
36.
37. plt.subplot(2,2,1)
39. # setting up t x-axis variable
40. t = np.arange(0.0, 11.0, 1.0)
41.
42. # loop for 10 different paths
43. n = 0
44. while n < 10:
       W = SimBMChol(1., 10)
       plt.plot(t, W, alpha = 0.5)
47.
       n += 1
48.
49. plt.title("Brownian Motion simulated (N = 10)")
51. ticks = np.arange(t.min(), t.max() + 1, 10)
52. labels = range(ticks.size)
53. plt.xticks(ticks, labels)
54. plt.xlabel("Time t")
56. plt.ylabel("Brownian Motion val")
58. ###################################
59. plt.subplot(2,2,2)
61. # setting up t x-axis variable
62. t = np.arange(0.0, 101.0, 1.0)
63.
64. # loop for 10 different paths
65. n = 0
66. while n < 10:
       W = SimBMChol(1., 100)
68.
       plt.plot(t, W, alpha = 0.5)
69.
       n += 1
70.
```

```
71. plt.title("Brownian Motion simulated (N = 100)")
72.
73. ticks = np.arange(t.min(), t.max() + 1, 100)
74. labels = range(ticks.size)
75. plt.xticks(ticks, labels)
76. plt.xlabel("Time t")
78. plt.ylabel("Brownian Motion val")
81. plt.subplot(2,2,3)
83. # setting up t x-axis variable
84. t = np.arange(0.0, 1001.0, 1.0)
85.
86. # loop for 10 different paths
87. n = 0
88. while n < 10:
       W = SimBMChol(1., 1000)
       plt.plot(t, W, alpha = 0.5)
91.
       n += 1
92.
93. plt.title("Brownian Motion simulated (N = 1000)")
95. ticks = np.arange(t.min(), t.max() + 1, 1000)
96. labels = range(ticks.size)
97. plt.xticks(ticks, labels)
98. plt.xlabel("Time t")
99.
          plt.ylabel("Brownian Motion val")
100.
101.
          102.
103.
          plt.subplot(2,2,4)
104.
105.
          # setting up t x-axis variable
106.
          t = np.arange(0.0, 10001.0, 1.0)
107.
          # loop for 10 different paths
108.
109.
          n = 0
110.
          while n < 10:
              W = SimBMChol(1., 10000)
111.
112.
              plt.plot(t, W, alpha = 0.5)
113.
              n += 1
114.
115.
          plt.title("Brownian Motion simulated (N = 10000)")
116.
          ticks = np.arange(t.min(), t.max() + 1, 10000)
117.
          labels = range(ticks.size)
118.
          plt.xticks(ticks, labels)
119.
          plt.xlabel("Time t")
120.
121.
          plt.ylabel("Brownian Motion val")
122.
123.
          plt.show()
124.
125.
126.
          # stopping timer
          stop = timeit.default_timer()
127.
128.
129.
          print "Cholesky method runtime = " , stop - start, " seconds"
```

Cholesky method runtime = 548.683270931 seconds

## Figure:



#### **Comments:**

The initial scaled simulation method gave a runtime of  $^{\sim}1$  second, whereas the cholesky simulation method gave a runtime of  $^{\sim}9$  minutes. There can be factors such as my algorithm implementation, but the cholesky methodology is noticeably slower than the first scaling & stepsize method.

# Question 6) 6) a) and b)

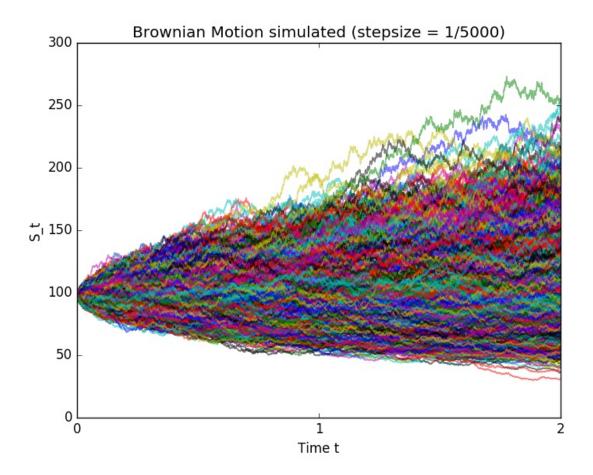
```
1. # importing the necessary packages
import numpy as np
from numpy import sqrt, exp
4. import matplotlib.pyplot as plt
5. import timeit
6.
8.
9. # starting timer
10. start = timeit.default timer()
11.
12. # initial variables
13. S 0 = 100.
14. B 0 = 1
15. mu = 0.17
16. sigma = 0.25
17. r = 0.02
18.
19. # We want to price an up-and-out barrier put option
20. T = 2.
21. K = 105.
22. B = 120.
23.
24. # increment step method defined function from Q5 scaling.py file
25. def SimBMStep(T, N):
26. # initialize W brownian motion array
27.
      W = np.zeros(N)
28. W = np.append(W, 0) # initial 0 value
29.
30. # loop
31.
      num = 1
32.
    while num <= N:</pre>
33.
          W[num] = W[num - 1] + sqrt(T/N)*np.random.standard normal()
34.
          num += 1
35.
36.
37.
      return W
38.
39. # stock generation defined function from Q1.py file
40. # note that we're multiplying by W value, instead of sqrt(T)*random normal
41. def generateStock(S_0, r, sigma, T, W):
42. return S_0 * exp((r - (sigma**2)/2)*T + sigma * W)
43.
44. # defining payoff function, for this case it's regular European put
45. def payoff_func(S, K):
46. return max(K - S, 0)
48.
49. plt.clf()
50.
51. # note, stepsize is 1/5000, but since T = 2, step size N = 10,000
52.
53. # setting up t x-axis variable
54. t = np.arange(0.0, 10001.0, 1.0)
55.
56. # keeping track of payoffs of up-and-out barrier put
```

```
57. payoffs = np.zeros(0)
58.
59. # tally for number of times barrier was hit
60. countBarrier = 0
61.
62. # loop for 10,000 different paths, for stepsize = 1/5000
63. n = 0
64. while n < 10000:
       # simulate brownian motion
67.
       W = SimBMStep(1., 10000)
68.
69.
       # generate future stock, using the simulated W value
       S_T = generateStock(S_0, r, sigma, T, W)
70.
71.
       plt.plot(t, S_T, alpha = 0.5)
72.
73.
       # check if barrier condition was hit
74.
       if np.any(S T > B):
75.
            # if past barrier at ANY point in time previously, payoff = 0
76.
            payoffs = np.append(payoffs, 0)
77.
            # keep track of how many times we hit barrier condition
78.
            countBarrier += 1
79.
       else:
80.
           payoffs = np.append(payoffs, exp(-r*T)*payoff func(S T[S T.size - 1], K))
81.
82.
83.
       n += 1
84.
85. # displaying the simulation brownian motion W
87. plt.title("Brownian Motion simulated (stepsize = 1/5000)")
89. ticks = np.arange(t.min(), t.max() + 1, 5000)
90. labels = range(ticks.size)
91. plt.xticks(ticks, labels)
92. plt.xlabel("Time t")
94. plt.ylabel("Brownian Motion val")
95.
96. plt.show()
97.
98. # calculating price
99. MonteCarloPrice = sum(payoffs) / 10000
           print("Monte Carlo integration estimation of price is: ")
101.
           print(MonteCarloPrice)
102.
           print("And number of times barrier was hit were: ")
103.
           print(countBarrier)
104.
105.
           # stopping timer
106.
           stop = timeit.default timer()
107.
           print "Barrier 120 method runtime = " , stop - start, " seconds"
108.
```

Monte Carlo integration estimation of price is: 10.8047400331

And number of times barrier was hit were: 4117 Barrier 120 method runtime = 192.690494776 seconds

## Figure:



6)c)
Number of times barrier was hit for when Barrier = 120: 4117
Number of times barrier was hit for when Barrier = 105: 7769
Number of times barrier was hit for when Barrier = 180: 155

6) d)

To improve the speed of the algorithm, I would check the barrier condition upon each S\_t simulation calculation. This is so that the inner simulation loop would end as soon the barrier condition gets hit.