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A short report on PYTHON implementation of subset simulation

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Since Python is freely accessible to almost all the researchers worldwide, I thought of sharing my Python implementation of the subset simulation method [1] in this report. The code is an exact replica of my Matlab code [2]. The Markov Chains (MC) in each conditional region can be propagated in two way - (a) one chain at a time (b) all the chains take a step at once. To clarify, consider 5 MCs each required to transition through 10 states in the first intermediate conditional region. Strategy (a): MC 1 first propagates through 10 states before MC 2 takes its first step. Strategy (b): MC 1 to 5 each take a single step at once. The performance of both the strategies is more or less the same. A highly nonlinear two-dimensional problem [3] is chosen for the purpose of illustration.

Numerical example

$$G(\mathbf{U}) = 4 - \frac{U_1}{4} + \sin(5U_1) - U_2; U_1, U_2 \sim N(0, 1) \quad (1)$$

Brute force Monte Carlo simulation

$$\hat{P}_F = \frac{1}{N} \sum_{k=1}^N I[G(\mathbf{u}^k) \leq 0] \quad (2)$$

where $I[\bullet]$ is the indicator function. Python implementation of MCS is given below:

```
import random
import numpy as np
import scipy
import math

# Performance function
def pfn(x):
    G=4-x[0]/4+math.sin(5*x[0])-x[1]
    return G

n=2                                     # Dimension of the problem
N=10000                                # Number of samples
G=np.zeros((N))
for i in range(0,N):
    ux=np.random.normal(0,1,size=(n,1)) # Generating random numbers
    G[i]=pfn(ux)                         # Performance function evaluations

pf_mcs=(G < 0).sum()/N                  # Estimate of the probability of failure
pf_mcs
```

Subset simulation

$$\bar{P}_F = P(F_1) \prod_{k=1}^{M-1} P(F_{k+1}|F_k) \quad (3)$$

Here M is the number of intermediate subsets, and the conditional probabilities, $P(F_{k+1}|F_k)$, are obtained using the Markov Chain Monte Carlo simulation with the Modified Metropolis-Hastings (MMH) algorithm [1].

Importing the required packages

```
import random
import numpy as np
import time
import scipy
import math
from operator import itemgetter
from scipy.stats import multivariate.normal
```

Defining the performance function

```
def pfn(x):
    G=4-x[0]/4+math.sin(5*x[0])-x[1]
    return G
```

Defining the parameters

```
n=2                                # Dimension of the problem
N_ss=500                           # Number of samples per subset
pp=0.1                             # Intermediate failure probability
```

Determining subset properties

```
nf_ini=int(N_ss*pp)                # Initial failure points
nf_ite=int(1/pp)                    # Number of samples generated from each Markov Chain
```

Estimating $P(F_1)$

Brute force Monte Carlo simulation is used in obtaining this quantity.

$$P(F_1) = \frac{1}{N_1} \sum_{k=1}^{N_1} I[G(\mathbf{u}^k) \leq 0] \quad (4)$$

```
ufailure=np.random.normal(0,1,size=(n,N_ss))
gfailure=np.zeros((N_ss))
for i in range(0,N_ss):
    gfailure[i]=pfn(ufailure[:,i])
pfl=nf_ini/N_ss                    # Initial pf, P(F1)
```

Choosing the initial state for the Markov Chain

```
index, g = zip(*sorted(enumerate(gfailure), key=itemgetter(1))) # Sorting the g-values
gl=g[nf_ini-1]                                                  # First intermediate threshold level, gl
if gl<0:                                                         # Check if failure region has been reached
    gl=0
    pfl=(gfailure<0).sum()/N_ss
x1=ufailure[:,index[0:nf_ini]] # Sort the samples according to ascending g-values
gx1=g[0:nf_ini]           # Choose the samples whose g-value is less than gl
sno=1                     # Counter for the number of subsets
gcollect[0]=gfinal        # This vector will store intermediate threshold values
gfinal=gl
```

Propagating the Markov Chains till failure region is reached – strategy (a)

```
while (gfinal>=0):
    ua=x1
    ga=list(gx1)
    for i in range(0,nf_ini): # For each Markov Chain
        x_seed=x1[:,i]       # Initial state for the Markov Chain
        g_seed=gx1[i]        # Corresponding g-value
        for j in range(1,nf_ite): # Till nf_ite number of samples are obtained from each Chain
            ux= (x_seed - 0.5) + np.random.uniform(0,1,size=(1,n)) # Uniform proposal density function
            m1=multivariate.normal.pdf(ux[0])
            m2=multivariate.normal.pdf(x_seed)
            alpha=np.minimum(np.ones((1,n)),m1/m2)
            r=np.random.uniform(0,1,size=(1,n))
            for ii in range(0,n): # Modified Metropolis-Hastings algorithm acceptance criterion
                if alpha[0][ii]<r[0][ii]:
                    ux[0][ii]=x_seed[ii]
            check=np.sum(alpha[0]>r[0])
```

```

        if check>0: # Performance function is evaluated only when an unique sample is obtained
            gx=pfn(ux[0])
            if gx<g1:
                x.seed=ux[0]
                g.seed=gx
            ua=np.append(ua,np.transpose([x.seed]),axis=1)
            ga.append(g.seed)
    ufailure=ua
    gfailure=ga
    index, g = zip(*sorted(enumerate(gfailure), key=itemgetter(1))) # Sorting the g-values
    x1=ufailure[:,index[0:nf_ini]] # Sort the samples according to ascending g-values
    gx1=g[0:nf_ini]
    g1=g[nf_ini-1] # Threshold level for the sno-th subset
    gfinal=g1
    gcollect[sno]=gfinal
    sno+=1

```

Propagating the Markov Chains till failure region is reached – strategy (b)

```

while (gfinal>=0):
    ua=x1
    ga=list(gx1)
    for j in range(1,nf_ite): # Till nf_ite number of samples are obtained from each Chain
        for i in range(0,nf_ini): # For each Markov Chain
            x.seed=ua[(j-1)*nf_ini+i] # Initial state for the Markov Chain
            g.seed=ga[(j-1)*nf_ini+i] # Corresponding g-value
            ux= (x.seed - 0.5) + np.random.uniform(0,1,size=(1,n))
            m1=multivariate.normal.pdf(ux[0])
            m2=multivariate.normal.pdf(x.seed)
            alpha=np.minimum(np.ones((1,n)),m1/m2)
            r=np.random.uniform(0,1,size=(1,n))
            for ii in range(0,n): # Modified Metropolis-Hastings algorithm acceptance criterion
                if alpha[0][ii]<r[0][ii]:
                    ux[0][ii]=x.seed[ii]
            check=np.sum(alpha[0]>r[0])
            if check>0: # Performance function is evaluated only when an unique sample is obtained
                gx=pfn(ux[0])
                if gx<g1:
                    x.seed=ux[0]
                    g.seed=gx
                ua=np.append(ua,np.transpose([x.seed]),axis=1)
                ga.append(g.seed)
    ufailure=ua
    gfailure=ga
    index, g = zip(*sorted(enumerate(gfailure), key=itemgetter(1)))
    x1=ufailure[:,index[0:nf_ini]]
    gx1=g[0:nf_ini]
    g1=g[nf_ini-1]
    gfinal=g1
    gcollect[sno,kkk]=gfinal
    sno+=1

```

Determining the \bar{P}_F

```

if sno>1:
    pf_final_subset=sum(gi < 0 for gi in gfailure)/(N-ss)
    pf_subset=pf1*pf1**((sno-2)*pf_final_subset)
    pf_ss=pf_subset # Estimate of the probability of failure
else :
    pf_ss=pf1

```

References

- [1] S.-K. Au, J. L. Beck, Estimation of small failure probabilities in high dimensions by subset simulation, Probabilistic Engineering Mechanics 16 (4) (2001) 263–277.
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