Exercise 5 - Variance redudiction Methods

The tasks centers around similation of the integral $\int_0^1 e^x \, \mathrm{d}x$

```
In [ ]: import numpy as np
   import numpy.random as rnd
   import scipy.stats as stats
   import math as math
```

1) Simulation of the integral using a Crude Monte Carlo Method

Using from the slides that the integral can be simplified to the expectation of e^U where $U\sim Uniform(0,1)$. Simulate n=100 instances of e^x , and find the mean. To get a 95% CI, use quantiles from t distribution with n-1 dof and $\alpha=0.05$.

$$CI = \left[ar{ heta} + rac{S_{ heta}}{\sqrt{n}} t_{rac{lpha}{2}}(n-1); ar{ heta} + rac{S_{ heta}}{\sqrt{n}} t_{1-rac{lpha}{2}}(n-1)
ight]$$

```
In [ ]: def crudeMC(n):
            U = rnd.uniform(size = n)
            x = np.exp(U)
            return x
        def meanVar(x):
            mean = np.mean(x)
            var = np.var(x)
            return mean, var
        def tConf(x, alpha, string):
            mean, var = meanVar(x)
            dof = len(x) - 1
            s = np.sqrt(var / len(x))
            a = stats.t.ppf(alpha/2, dof)
            b = stats.t.ppf(1 - alpha/2, dof)
            print("For", string, ":")
            print(f"Mean is {round(mean,3)}, with 95% confidence interval [{round(mean +
        alpha = 0.05
        n = 100
        Xs_crude = crudeMC(n)
        tConf(Xs_crude, alpha, "Crude Monte Carlo")
        print(f"The true value of the integral is: {round(np.exp(1) - np.exp(0),4)}")
       For Crude Monte Carlo:
```

2) Antithetic variables

The true value of the integral is: 1.7183

We can use Anithetic variables to exploit the fact that the integral $\int_{-\infty}^{\infty} e^x \, \mathrm{d}x$ is

Mean is 1.739, with 95% confidence interval [1.64,1.841]

monotonely increasing. In practice this means we can use a single uniformly distributed value $U \sim Uniform(0,1)$, as in the Crude Monte Carlo Method, an use it to create a second uniformly distributed value U-1, almost for free.

Using these we can create 2 estimates of the integral by e^U and e^{U-1} , and take the average of these two to get a much more robust estimate of the integral

$$Y_i=rac{e^{U_i-e^{U_i-1}}}{2}.$$

The expectation of the integral is now

$$E(Y_i)$$
.

 Y_i which on the slides is proved to have variance $\frac{1}{4}Var\left(e^{U_i}\right)+\frac{1}{4}Var\left(e^{1-U_i}\right)+\frac{1}{2}Cov\left(e^{U_i},e^{1-U_i}\right)$. As e^{U_i} and e^{1-U_i} are obviosly negatively correlated, this variance is much lower than before.

By rewriting Y_i computing cost can be lowered, to only calculate a single exponential, resulting in an only marginally more expensive computation for $n\ Y$'s compared to $n\ X$'s.

$$Y_i = rac{e^{U_i + rac{e}{e^{U_i}}}}{2}.$$

Note: Had the integral not been monotonely increasing, say we had attemted to estimate some other function f, we could have had a situation where $f\left(U_i\right)$ and $f\left(U_i-1\right)$ had been positively correlated, which could lead to a higher variance on Y_i . For the exponential function one of e^{U_i} and e^{U_i-1} is always large, and one is always small.

```
In [ ]: def antithetic(n):
    U = rnd.uniform(size = n)
    t = np.exp(U)
    Ys = 0.5 * (t + np.exp(1) / t )
    return Ys
    Xs_antithetic = antithetic(100)
    tConf(Xs_antithetic, alpha, "Antithetics Variables")
```

For Antithetics Variables :
Mean is 1.717, with 95% confidence interval [1.7,1.729]

3) Control Variates

We can use the variable

$$Z = X + c \left(Y - \mu_i \right),\,$$

instead of X_i as the estimate of the integral. It can then be shown that $E\left(X\right)=E\left(Z\right)$, since

$$E\left(Z
ight) = E\left(X
ight) + E\left(c\left(Y - \mu_i
ight)
ight) \ = E\left(X
ight) + c\left(E\left(Y
ight) - E\left(\mu_i
ight)
ight) = E\left(X
ight) + c\left(\mu_i - \mu_i
ight) = E\left(X_i
ight).$$

From the slides we also know that choosing the optimal $c=-\frac{Cov(X,Y)}{Var(Y)}$, results in a variance of Z

$$Var(Z) = Var(X) - rac{Cov(X,Y)^2}{Var(Y)}.$$

For this specific problem we are given $X_i=e^{U_i}$, and it is natural to choose $Y_i=U_i$. This again exploits the covariance between U_i and X_i , though this time negative correlation is not a requirement, as the covariance is squared. It should just be non-zero. As $E\left(U_i\right)=\frac{1}{2}$ we get

$$Z_i = e^{U_i} + c\left(U_i - rac{1}{2}
ight),$$

with $c \approx 0.14086$.

```
In [ ]: def controlVariates(n, alpha):
    U = rnd.uniform(size = n)
    Xs = np.exp(U)
    dof = len(Xs) - 1
    meanZ = np.mean(Xs)
    cov = np.mean(U * Xs) - np.mean(U) * np.mean(Xs)
    varZ = np.var(Xs) - cov**2 / np.var(U)
    s = np.sqrt(varZ / len(Xs))
    a = stats.t.ppf(alpha/2, dof)
    b = stats.t.ppf(1 - alpha/2, dof)
    print("For Control Variates")
    print(f"Mean is {round(meanZ,2)}, with 95% confidence interval [{round(meanZ controlVariates(n, alpha)}]
```

For Control Variates
Mean is 1.71, with 95% confidence interval [1.7,1.72]

4) Statified Sampling

Stratified sampling attemts to lower variance by (almost) gaurantying samples from the entire sample space, by cutting the sample space into m pieces, and then sampling a number of variables from each. Here we'll use m=10 even intervals, to ensure even computational cost to the other methods, 10 U_i 's are generated in each interval.

The method then takes one U_i from each interval and creates a single variable W_i

$$W_i = rac{\sum_{k=1}^m X_{i,k}}{m},$$

where k now denotes which interval the X_i comes from. Still $X_i = e^{U_i}$, or $X_{i,k} = e^{U_{i,k}}$.

```
In []: def stratSamples(n):
    num_ints = 10
    k = 1 / num_ints
    Xs = np.zeros(n)
    Us = np.zeros(num_ints)
    for j in range(n):
        for i in range(num_ints):
```

```
a = k * i
b = a + k
U = rnd.uniform(a, b, size = 1)
Us[i] = U
Xs[j] = np.mean(np.exp(Us))
return Xs
n = 10
Xs_stratified = stratSamples(n)
tConf(Xs_stratified, alpha, "Stratified Sampling")
```

```
For Stratified Sampling : Mean is 1.72, with 95% confidence interval [1.71,1.733]
```

1-4) Observations

The Crude MC method gives a very wide CI All the variance reduction methods have successfully narrowed the CI. We had the largest success with Control Variates.

5) Use control variates to reduce the variance of the estimator in exercise 4

We'll still be exploiting the smaller variance in the variable

```
In [ ]: import numpy as np
        import numpy.random as rnd
        import scipy.stats as stats
        import math as math
        from scipy.stats import expon
        import numpy as np
        from discrete_event import Customer, main_loop, confidence_intervals, erlang_b
        #arrival time differences are exponentially distributed
        lam = 1
        total customers = 10000
        m = 10
        s = 8
        repititions = 10
        #arrival time differences are exponentially distributed
        arrival_interval = lambda : np.random.exponential(1/lam, size = total_customers)
        service_time =lambda : expon.rvs(scale = s, size = total_customers)
        blocked = main_loop(arrival_interval, service_time, m, repititions = repititions
        #print("Blocking probability", blocked/total customers * 100)
        #confidence interval for the mean
        theta = np.mean(blocked)
        confint = confidence_intervals(blocked)
        print(f"Estimated blocking probability {round(theta,3)*100}%\nTrue blocking prob
        print(f"95% CI for mean of blocking probability [{round(confint[0],4)}, {round(c
        print("Interval width", round(confint[1]-confint[0],4))
       Estimated blocking probability 12.1%
       True blocking probability 12.2% (From Erlang formula)
       95% CI for mean of blocking probability [0.1185, 0.1232]
       Interval width 0.0047
```

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In []: #Empirical covariance between exponential and uniform random variables

n = 10000

U = np.random.uniform(0,1,n)

```
X = np.random.exponential(1/s,n)
#c for minimizing variance of X + c(U-1/2)
c = -np.cov(X,U)[0,1]/np.var(U)
#c

#arrival time differences are exponentially distributed
arrival_interval = lambda : np.random.exponential(1/lam, size = total_customers)
service_time =lambda : expon.rvs(scale = s, size = total_customers)
```

```
In [ ]: np.random.seed(0)
    blocked = main_loop(arrival_interval, service_time, m, repititions = repititions
    #print("Blocking probability", blocked/total_customers)
#confidence interval for the mean
    theta = np.mean(blocked)
    confint = confidence_intervals(blocked)
    print(f"Estimated blocking probability {round(theta,3)*100}%\nTrue blocking prob
    print(f"95% CI for mean of blocking probability [{round(confint[0],4)}, {round(confint[1]-confint[0],4)})
```

```
Estimated blocking probability 12.3%
True blocking probability 12.2% (From Erlang formula)
95% CI for mean of blocking probability [0.1201, 0.1257]
Interval width 0.0056
```

6) Reduce variance for the difference between solutions with poisson arrivals and hyperexponential renewal process in exercise 4

Do this using Common Random Numbers (CRN). Using CRN we should be able to reduce the width of the CI og the difference in number of blocked customers between the two arrival processes.

Using non-CRN's it should require 5 differerent processes of generating arrivals and leaves to compare the two processes. 2 generates the serice times for the processes, one generates arrivals for the poisson process and 2 generates renewals for the hyperexponential arrivals.

When using CRN's this can be reduced to 3. The services times can be generated from the same U, since they should be the same here. We can use a single U as the exponential for the poisson and hyperexponential processes. We also need one more U to flip between the two exponentials in the hyperexponential distribution.

Both test runs predict more blocked customers when assuming a hyperexponential renewal process. The width of the CI when using CRN's can be reduced to $\sim 1/3$ compared to just simulating the two processes independently.

```
In [ ]: import bisect as bisect
    class Customer:
        def __init__(self, arrival_time, service_time):
            self.service_time = service_time
            self.blocked = False

            self.event = "arrival"
            self.event_time = arrival_time
```

```
def arrive(self, servers, event_list):
        if servers < 1:</pre>
            self.blocked = True
            return servers
        else:
            servers -= 1
            servers = max(servers, 0)
            self.event = "departure"
            self.event_time += self.service_time
            bisect.insort(event_list, self, key=lambda x: x.event_time)
            return servers
    def depart(self, servers):
        servers += 1
        servers = min(servers, m)
        return servers
from distributions import getExponential, getUniform
def main_loop(event_list, m, repititions = 10):
    blocked = 0
    for i in range(repititions):
        event_list.sort(key=lambda x: x.event_time)
        open_servers = m
        while event_list:
            event = event_list.pop(0)
            if event.event == "arrival":
                open_servers = event.arrive(open_servers, event_list)
                blocked += event.blocked
            elif event.event == "departure":
                open_servers = event.depart(open_servers)
    return blocked
num_customers = 10000
lam = 1
s = 1/8
# Generate two eventlists
p1 = 0.8
lam1 = 0.8333
p2 = 0.2
lam2 = 5.0
def generateHexp(Us, Us2, p2, lam1, lam2):
    Xs = np.zeros(len(Us))
    for i in range(len(Us)):
        if Us2[i] < p2:</pre>
            Xs[i] = -np.log(Us[i]) / lam2
            Xs[i] = -np.log(Us[i]) / lam1
    return Xs
m = 10
rep = 5
blocked_hexp = np.zeros(rep)
blocked_poisson = np.zeros(rep)
blocked_poisson_common = np.zeros(rep)
blocked_hexp_common = np.zeros(rep)
for i in range(rep):
```

```
Us2 = rnd.uniform(0,1,num_customers)
            Us3 = rnd.uniform(0,1,num_customers)
            service_intervals = -np.log(Us1) / s
            poisson_arrival_times = np.cumsum(-np.log(Us2) / lam)
            hexp_arrival_times = np.cumsum(generateHexp(Us2, Us3, p2, lam1, lam2))
            event_list_poisson = [Customer(poisson_arrival_times[i], service_intervals[i
            event_list_hexp = [Customer(hexp_arrival_times[i], service_intervals[i]) for
            blocked_poisson_common[i] = main_loop(event_list_poisson, m, repititions=1)
            blocked_hexp_common[i] = main_loop(event_list_hexp, m, repititions=1)
            Us1 = rnd.uniform(0,1,num_customers)
            Us2 = rnd.uniform(0,1,num_customers)
            Us3 = rnd.uniform(0,1,num_customers)
            Us4 = rnd.uniform(0,1,num_customers)
            Us5 = rnd.uniform(0,1,num_customers)
            service_intervals1 = -np.log(Us2) / s
            service_intervals2 = -np.log(Us3) / s
            poisson_arrival_times = np.cumsum(-np.log(Us1) / lam)
            hexp_arrival_times = np.cumsum(generateHexp(Us4, Us5, p2, lam1, lam2))
            event_list_poisson = [Customer(poisson_arrival_times[i], service_intervals1[
            event_list_hexp = [Customer(hexp_arrival_times[i], service_intervals2[i]) fo
            blocked_poisson[i] = main_loop(event_list_poisson, m, repititions=1)
            blocked_hexp[i] = main_loop(event_list_hexp,m, repititions=1)
In [ ]: | theta_crn = blocked_poisson_common - blocked_hexp_common
        theta_irn = blocked_poisson - blocked_hexp
        mean_theta_crn = round(np.mean(theta_crn),0)
        var_theta_crn = np.var(theta_crn)
        s = np.sqrt(var_theta_crn / len(theta_crn))
        dof = len(theta_crn)-1
        a = stats.t.ppf(alpha/2, dof)
        b = stats.t.ppf(1 - alpha/2,dof)
        print("For common random numbers")
        print(f"Mean is {mean_theta_crn}, with confidence interval [{round(mean_theta_cr
        print(f"Width of CI = {round(np.abs(mean_theta_crn + s * a - mean_theta_crn - s*
        mean theta irn = int(np.mean(theta irn))
        var_theta_irn = np.var(theta_irn)
        s = np.sqrt(var_theta_irn / len(theta_irn))
        print("For independent random numbers")
        print(f"Mean is {mean_theta_irn}, with confidence interval [{round(mean_theta_ir
        print(f"Width of CI = {round(abs(mean_theta_irn + s * a - mean_theta_irn - s*b),
       For common random numbers
       Mean is -132.0, with confidence interval [-170.009,-93.991]
       Width of CI = 76.017
       For independent random numbers
       Mean is -152, with confidence interval [-284.092,-19.908]
       Width of CI = 264.184
```

Us1 = rnd.uniform(0,1,num_customers)

7) Using a Crude Monte Carlo estimator vs Importance

Sampling

Attempt to estimate the probality Z>a for $Z\sim N(0,1)$ for a=2,4. Do this using a crude monte carlo estimator and with importance sampling.

Using importance sampling set h(z)=z>a. We'll choose g(x) to also be a normal distribution, with the same variance as f(x), though with mean a instead. This means g samples around a more often. We thus have to sample fewer points to get an accurate estimate.

Notice n=10000 with importance sampling gives significant digits, while the Crude Estimator needs n=100000 to get a significant digit. A lot of time can be saved.

As long as a is small enough e.g. a=2 Importance Sampling does not seem nescesarry.

```
In [ ]: def crudeMonteCarloNorm(a,n):
            Us = rnd.normal(size = n)
            return sum(Us > a) / n
        n = 10000
        n2 = 100000
        p1 = crudeMonteCarloNorm(2, n2)
        p2 = crudeMonteCarloNorm(4, n2)
        print(f"With {n} samples")
        print(f"Crude probability z larger than 2 is {round(p1 * 100,5)}%")
        print(f"Crude probability z larger than 4 is {round(p2 * 100,5)}%")
        def h(a,x):
            return x > a
        sigma1 = 1
        a = 2
        ys = rnd.normal(loc = a,scale = sigma1,size = n)
        fy = stats.norm.pdf(ys)
        gy = stats.norm.pdf(ys, loc = a, scale = sigma1)
        zs = fy / gy * h(a,ys)
        print(f"Probability z larger than 2 with importance sampling {round(np.mean(zs)
        a = 4
        ys = rnd.normal(loc = a,scale = sigma1,size = n)
        fy = stats.norm.pdf(ys)
        gy = stats.norm.pdf(ys, loc = a, scale = sigma1)
        zs = fy / gy * h(a,ys)
        print(f"Probability z larger than 4 with importance sampling {round(np.mean(zs)
       With 10000 samples
       Crude probability z larger than 2 is 2.27%
       Crude probability z larger than 4 is 0.003%
       Probability z larger than 2 with importance sampling 2.3139%
```

8)

Analytically set

 $f(x) = \mathbf{1}_{0 \leq x \leq x}$

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Probability z larger than 4 with importance sampling 0.0031%

$$h(x) = e^x$$
 $g(x) = \lambda e^{-\lambda x}.$

We want to find the variance of

$$\frac{f(x)h(x)}{g(x)}.$$

Write

$$Var\left(\frac{f(x)h(x)}{g(x)}\right) = E\left(\left(\frac{f(x)h(x)}{g(x)}\right)^2\right) - E\left(\frac{f(x)h(x)}{g(x)}\right)^2$$
$$= \int_{-\infty}^{\infty} \left(\frac{f(x)h(x)}{g(x)}\right)^2 g(x) dx - \left(\int_{-\infty}^{\infty} \frac{f(x)h(x)}{g(x)} g(x) dx\right)^2.$$

The second integral is simply to rewrite

$$egin{split} \left(\int_{-\infty}^{\infty} rac{f(x)h(x)}{g(x)} g(x) \,\mathrm{d}x
ight)^2 &= \left(\int_{-\infty}^{\infty} f(x)h(x) \,\mathrm{d}x
ight)^2 \ &= \left(\int_{0}^{1} f(x)e^x \,\mathrm{d}x
ight)^2. \end{split}$$

The first integral is slightly less pretty

$$\int_{-\infty}^{\infty} \left(rac{f(x)h(x)}{g(x)}
ight)^2 g(x) \, \mathrm{d}x = \int_{-\infty}^{\infty} rac{f(x)^2 h(x)^2}{g(x)} \, \mathrm{d}x$$

$$= \int_{-\infty}^{\infty} rac{f(x)^2 e^{2x}}{\lambda e^{-\lambda x}} \, \mathrm{d}x = rac{1}{\lambda} \int_{-\infty}^{\infty} f(x)^2 e^{2x + \lambda x} \, \mathrm{d}x$$

$$= rac{1}{\lambda} \int_{0}^{1} e^{(2+\lambda)x} \, \mathrm{d}x.$$

Collecting:

$$Var\left(rac{f(x)h(x)}{g(x)}
ight) = rac{1}{\lambda} \int_0^1 e^{(2+\lambda)x} \,\mathrm{d}x - \left(\int_0^1 f(x)e^x \,\mathrm{d}x
ight)^2$$

From Maple:

$$Var\left(rac{f(x)h(x)}{g(x)}
ight) = rac{-1+e^{2+\lambda}}{2+\lambda} - (e-1)^2.$$

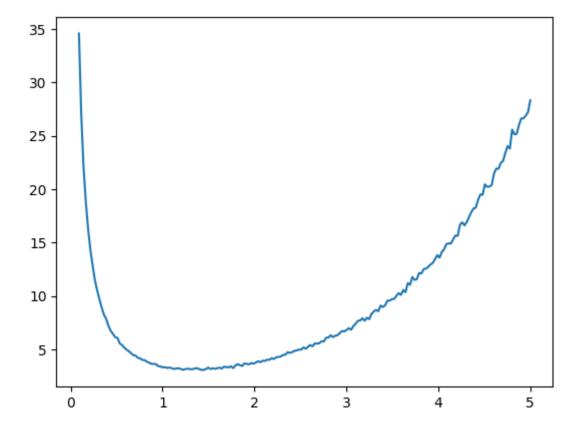
Using a numerical solver to minimize this expression gives $\lambda \approx 1.354828644$.

```
In [ ]: import matplotlib.pyplot as plt
lambda_opt = 1.354828644

def var_anal(lam):
    return (-1 + np.exp(2 + lam))/(lam*(2 + lam)) - (np.exp(1) - 1)^2
```

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```
def variance(lam, n):
     Us1 = rnd.uniform(size = n)
     Us2 = rnd.uniform(size = n)
     return 1/lam * np.mean(np.exp((2+lam)*Us1)) - np.mean(np.exp(Us2))**2
 lams = np.linspace(0.09, 5, 200)
 ys = np.zeros(200)
 for i in range(len(lams)):
     ys[i] = variance(lams[i], 10000)
 plt.plot(lams, ys)
 def g(x,lam):
     return lam * np.exp(-lam * x)
 def f(x):
     b1 = x > 0
     b2 = x < 1
     return b1 * b2
 def h(x):
     return np.exp(x)
 n = 100
 lam = lambda opt
 ys = rnd.exponential(scale = 1/lam, size = n)
 zs = f(ys) * h(ys) / g(ys, lam)
 mean = np.mean(zs)
 var = np.var(zs)
 s = np.sqrt(var / len(zs))
 a = stats.norm.ppf(alpha/2)
 b = stats.norm.ppf(1 - alpha/2)
 print("Using importance sampling")
 print(f"Mean is {round(mean,4)}, with confidence interval [{round(mean + s * a,3
 print(f"Width of CI = {round(abs(mean + s * a - mean - s*b),3)}")
 n = 100
 Xs\_crude = crudeMC(n)
 mean = np.mean(Xs_crude)
 var = np.var(Xs_crude)
 s = np.sqrt(var / len(Xs_crude))
 a = stats.norm.ppf(alpha/2)
 b = stats.norm.ppf(1 - alpha/2)
 print("Using Crude Monte Carlo")
 print(f"Mean is {round(mean,4)}, with confidence interval [{round(mean + s * a,3
 print(f"Width of CI = {round(abs(mean + s * a - mean - s*b),3)}")
Using importance sampling
Mean is 1.6713, with confidence interval [1.322,2.02]
Width of CI = 0.698
Using Crude Monte Carlo
Mean is 1.7122, with confidence interval [1.627,1.798]
Width of CI = 0.171
```



9) Deriving the IS estimator for the mean of a Pareto distribution

pdf for a arbritray Pareto distribution is $f(x)=\frac{k}{y}\left(\frac{\beta}{y}\right)^k$. From the lectures we know that the first moment distribution of a Pareto distribution is just a Pareto distribution with paramters β and k-1. Thus

$$g(x) = rac{k-1}{y} igg(rac{eta}{y}igg)^{k-1}.$$

As we're estimating the mean, set h(y)=y. Now the IS estimator for the mean is

$$\frac{f(y)h(y)}{g(y)} = \frac{\frac{\frac{k}{y}\left(\frac{\beta}{y}\right)^k \cdot y}{\frac{k-1}{y}\left(\frac{\beta}{y}\right)^{k-1}} = \frac{yk}{k-1} \frac{\left(\frac{\beta}{y}\right)^k}{\left(\frac{\beta}{y}\right)^{k-1}} = \frac{yk}{k-1} \frac{\beta}{y} = \beta \frac{k}{k-1}.$$

Which we know is just the mean of the original Pareto distribution f(x), i.e. we should be sampling exactly the mean! Which we do, with an exact CI.

```
In [ ]: k = 2
def g(x):
    return (k-1) /(x) * (1/x)**(k-1)

def f(x):
    return k /(x) * (1/x)**k

def h(x):
    return x
n = 100
```

```
lam = lambda_opt
ys = rnd.exponential(scale = 1/lam, size = n)
beta = 1
zs = f(ys) * h(ys) / g(ys)
alpha = 0.05
mean = np.mean(zs)
var = np.var(zs)
s = np.sqrt(var / len(zs))
a = stats.norm.ppf(alpha/2)
b = stats.norm.ppf(1 - alpha/2)
print("Using importance sampling to sample the mean of a pareto distribution usi
print(f"Mean is {round(mean,4)}, with confidence interval [{round(mean + s * a,3)
print(f"Width of CI = {round(abs(mean + s * a - mean - s*b),3)}")

print(f"Exact solution for the Pareto distribution mean {k/(k-1) * beta}")
```

Using importance sampling to sample the mean of a pareto distribution using its f irst moment Mean is 2.0, with confidence interval [2.0,2.0] Width of CI = 0.0 Exact solution for the Pareto distribution mean 2.0

In this case sampling values from a constant value works out, because we know the mean of the Pareto distribution. This approach could be used, but it would be nesecary to know for certain that you're sampling from the right constant. It could be possible to choose a g(x) such that $\frac{h(x)f(x)}{g(x)}$ is a constant, which is not the mean of f(x), in this case you would end ud with 100% confidence, in something which is not correct. Gauranteing that you sample from the constant mean, would require you knowing the mean, but then there would be no reason for this approach.

For the previous case $f(x)=\lambda e^{-\lambda x}$ use $g(x)=xe^{-\lambda x}$. To sample from only the mean get

$$\frac{x \cdot \lambda e^{-\lambda x}}{x e^{-\lambda x}} = \lambda.$$