# Initialize Step (Step 1/3)

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#### Algorithm:

```
Inputs: w \sim \text{Unif}(0,1)

l_k(x^*) = \log(g_l(x^*))

u_k(x^*) = \log(g_u(x^*))

h(x^*) = \log(g(x^*))

s_k(x) = \exp(u_k(x)) / \left( \int_D u_k(x') \ dx' \right) = g_u(x) / \left( \int_D g_u(x') \ dx' \right)
```

The lower bound of h(x) is  $l_k(x)$ , which connects the values of function h on abscissaes. The function of  $l_k(x)$  between two consecutive abscissaes  $x_j$  and  $x_{j+1}$  is  $l_k(x) = \frac{(x_{j+1}-x)h(x_j)+(x-x_j)h(x_{j+1})}{x_{j+1}-x_j}$ 

Let T be the domain of abscissaes, H be the domain of the realized function H at abscissaes, H\_prime be the domain of the realized first derivative of function H at abscissaes.

$$h'(x) = \frac{dlog(g(x))}{dx} = \frac{g'(x)}{g(x)}$$

#library(distr)

**Step 1**: If  $w < exp(l_k(x^*) - u_k(x^*))$ 

- Accept  $x^*$  when the condition is satisfied. Draw another  $x^*$  from  $s_k(x)$
- Reject  $x^*$  when the condition is not satisfied.

Step 2: These two procedures can be done in parallel.

- Evaluate  $h(x^*)$ ,  $h'(x^*)$ . Update  $l_k(x)$ ,  $u_k(x)$ ,  $s_k(x)$ , which are now include  $x^*$  as an element. - Accept  $x^*$  if  $w < exp(h(x^*) - u_k(x^*))$ . Otherwise, reject.

```
Example: Start with g(x) = 3*N(0,1). g(x) = \frac{3}{\sqrt{2\pi}}e^{-(x)^2/2}
```

```
#Create
#install.packages("Ryacas")
#install.packages("Deriv")
#install.packages("distr")
#install.packages("truncnorm")
library(truncnorm)
library(Ryacas)
library(Deriv)
```

```
##
## Attaching package: 'Deriv'
## The following object is masked from 'package:Ryacas':
##
## Simplify
```

```
sigma <- 1
 mu <- 0
 val <- 3/sqrt(2*pi*sigma^2)*exp(-(x-mu)^2/(2*sigma^2))</pre>
}
#Examples for function g
g(0)
g(1)
g(-1)
#Number of starting points
k_start=2
#Bounds for x
x.Bound=c(-Inf,Inf)
#Test default deriv() function + Simplify() function
sigma <- 1
mu <- 0
x \leftarrow Sym("x")
f <- 1/sqrt(2*pi*sigma^2)*exp(-(x-mu)^2/(2*sigma^2))
dx <- eval(deriv(g(x),Sym("x"))); #Derivative of f in terms of x</pre>
print(dx)
## expression(-1.1968268412 * (exp(-x^2/2) * (2 * x))/2)
dxSimp \leftarrow Ryacas::Simplify(deriv(g(x),"x")) #Simplified version of derivative (collapse terms)...
print(dxSimp)
## expression(-2.3936536824 * (exp(<math>-x^2/2) * x)/2)
s = deriv(-a^2, a', func = T) #Create function s(a) = a^2
#s(5) #5^2
#Internally created functions
g_d <- function(x){</pre>
 as.numeric(eval(Deriv(~g(X),"X")))
}
#h function
h <- function(x){
   log(g(x))
}
#Examples for function h
#h(1)
#h(-1)
#h(0)
#h' function
h_d <- function(X){</pre>
```

```
#print("X:")
    #print(X)
    #print("Original:")
    #print(h(X))
    #print("Derivative - deriv()")
    \#print(deriv(log(g(x)), "x"))
    #print("Derivative - Deriv() Unevaluated")
    \#print(Deriv(\sim log(g(X)), "X"))
    #print("Derivative - Deriv() Evaluated")
    #print(eval(fun))
    \#deriv(log(g(x),1))
    \#return(as.numeric(eval(Deriv(\sim log(g(X)),"X"))))
  return(as.numeric(1/g(X)*eval(Deriv(~g(X),"X"))))
}
\#Examples for function h'...
#h d(0)
#h_d(-0.5)
#Initial function (main)
Initial <- function(k_start,x.Bound,g,QC=FALSE){</pre>
  #print(all.equal(x.Bound,c(-Inf,Inf)))
    upBd <- x.Bound[length(x.Bound)]</pre>
    lowBd <- x.Bound[1]</pre>
    if (lowBd != -Inf){
      needPosHprime <- FALSE</pre>
    } else{
      needPosHprime <- TRUE</pre>
    if (upBd != Inf){
      needNegHprime <- FALSE</pre>
    } else{
      needNegHprime <- TRUE</pre>
    upBdInit <- upBd #Upper bound for initialization points</pre>
    lowBdInit <- lowBd #Lower bound for initialization points</pre>
    if (0==1){
      if (QC==TRUE){
        print("Lower & upper bound pairs")
        print(lowBd)
        print(upBd)
        print(upBdInit)
        print(lowBdInit)
      }
```

```
k <- 0
X \leftarrow c()
#print(pasteO("Begin: Find initialization points"))
while (k < k_start){</pre>
  samp <- rtruncnorm(1,a=lowBdInit,b=upBdInit)</pre>
  hPrime <- h d(samp)
  if (QC==TRUE){
    print(paste0("Initialization bounds: ",lowBdInit," ",upBdInit))
    print(paste0("Need positive / negative h'? ",needPosHprime," ",needNegHprime))
    print(paste0("Sampled point x: ",samp))
    print(paste0("h'(x): ",h_d(samp)))
  }
  if (hPrime >0 & needPosHprime==TRUE){
    needPosHprime <- FALSE</pre>
    X <- append(X,samp,length(X))</pre>
    k < - k+1
    \#print(pasteO("Appended X with h'(X) > O"))
    if (needNegHprime==TRUE) {
      lowBdInit <- samp</pre>
      #print(pasteO("Changed lower initialization bound to sampled point"))
    }
    else {
      lowBdInit <- lowBd</pre>
      #print(pasteO("Changed lower initialization bound to distribution lower bound"))
    }
  }
  else if (hPrime < 0 & needNegHprime==TRUE) {</pre>
    needNegHprime <- FALSE</pre>
    X <- append(X,samp,0)</pre>
    k < - k+1
    \#print(pasteO("Appended X with h'(X) < O"))
    if (needPosHprime == TRUE) {
      upBdInit <- samp
      #print(pasteO("Changed upper initialization bound to sampled point"))
    else {
      upBdInit <- upBd
      #print(paste0("Changed upper initialization bound to distribution upper bound"))
  }
  else if (k < (k_start-needPosHprime-needNegHprime)){</pre>
    X <- append(X,samp,length(X[X<samp]))</pre>
    k < - k+1
    #print(pasteO("Appended X; no h' requirements met"))
  }
  else {
```

```
#print(pasteO("Sampled point does not satisfy any initialization requirements. Sample again."))
}

#print(pasteO("Finished: Find initialization points"))

H <- h(X)
H_d <- h_d(X)
Z <- c(lowBd,upBd)
return(list(X,H,H_d,Z))
}</pre>
```

## Normal distribution (2 starting pts)

```
#User supplied g-function
g <- function(x){</pre>
  sigma <- 1
  mu <- 0
  val <- 3/sqrt(2*pi*sigma^2)*exp(-(x-mu)^2/(2*sigma^2))</pre>
Initial(k_start=2,x.Bound=c(-Inf,Inf),g
        #,QC=TRUE
        )
## [[1]]
## [1] 2.4119015 -0.1000012
##
## [[2]]
## [1] -2.7289608 0.1746736
## [[3]]
## [1] -2.4119015 0.1000012
##
## [[4]]
## [1] -Inf Inf
system.time(Initial(k_start=2,x.Bound=c(-Inf,Inf),g
        #,QC=TRUE
        ))
##
      user system elapsed
##
     0.287
           0.003 0.305
```

## Exponential distribution (2 starting pts)

```
g <- function(x){
  lam <- 1
  value <- lam*exp(-lam*x)
}</pre>
```

```
#a <- AbscontDistribution(d=g,low=0,low1=0)
#print(q(1))
Initial(k_start=2,x.Bound=c(0,Inf),g
        #,QC=TRUE
        )
## [[1]]
## [1] 0.8800005 1.1956156
##
## [[2]]
## [1] -0.8800005 -1.1956156
##
## [[3]]
## [1] -1 -1
## [[4]]
## [1]
       0 Inf
```

## Normal distribution (5 starting pts)

0.484

0.035 0.587

```
#User supplied g-function
g <- function(x){</pre>
 sigma <- 1
 mu <- 0
 val <- 3/sqrt(2*pi*sigma^2)*exp(-(x-mu)^2/(2*sigma^2))</pre>
Initial(k_start=5,x.Bound=c(-Inf,Inf),g
       #, QC=TRUE
## [1] 0.07065779 -1.46264688 -0.66467338 -0.56187684 0.33905853
## [[2]]
## [1] 0.17717749 -0.88999420 -0.04122160 0.02182096 0.12219341
##
## [[3]]
## [[4]]
## [1] -Inf Inf
print(system.time(Initial(k_start=5,x.Bound=c(-Inf,Inf),g)
       #,QC=TRUE
       )))
##
     user system elapsed
```

## Gamma distribution (5 starting pts)

```
\#User\ supplied\ g\mbox{-}function
g <- function(x){
  alpha <- 1
  beta <- 1
  val <- beta^alpha/gamma(alpha)*x^(alpha-1)*exp(-beta*x)</pre>
}
Initial(k_start=5,x.Bound=c(0,Inf),g
        #, QC=TRUE
        )
## [[1]]
## [1] 0.3903494 0.5107521 0.5438679 0.9327843 2.2745782
## [[2]]
## [1] -0.3903494 -0.5107521 -0.5438679 -0.9327843 -2.2745782
##
## [[3]]
## [1] -1 -1 -1 -1 -1
## [[4]]
## [1]
        0 Inf
```

### Beta distribution (10 starting pts)

```
## user system elapsed
## 0.838 0.015 0.879
```