Package 'AdapSamp'

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Type Package

rARS

rARS

Adaptive Rejection Sampling Algorithm

Description

rARS generates a sequence of random numbers using the adaptive rejection sampling algorithm.

Usage

```
rARS(n, formula, min = -Inf, max = Inf, sp)
```

Arguments

n Desired sample size;

formula Kernal of the target density;

min, max Domain including positive and negative infinity of the target distribution;

sp Supporting set.

Author(s)

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```
# Example 1: Standard normal distribution
x1 \leftarrow rARS(100, "exp(-x^2/2)", -Inf, Inf, c(-2,2))
# Example 2: Truncated normal distribution
x2 \leftarrow rARS(100, "exp(-x^2/2)", -2.1, 2.1, c(-2, 2))
# Example 3: Normal distribution with mean=2 and sd=2
x3 \leftarrow rARS(100, "exp(-(x-2)^2/(2*4))", -Inf, Inf, c(-3,3))
# Example 4: Exponential distribution with rate=3
x4 <- rARS(100,"exp(-3*x)",0,Inf,c(2,3,100))
# Example 5: Beta distribution with alpha=3 and beta=4
x5 \leftarrow rARS(100, "x^2*(1-x)^3", 0, 1, c(0.4, 0.6))
# Example 6: Gamma distribution with alpha=5 and lambda=2
x6 \leftarrow rARS(100, "x^{(5-1)} + exp(-2 + x)", 0, Inf, c(1, 10))
# Example 7: Student distribution with df=10
x7 \leftarrow rARS(100,"(1+x^2/10)^(-(10+1)/2)",-Inf,Inf,c(-10,2))
# Example 8: F distribution with m=10 and n=5
x8 \leftarrow rARS(100, "x^(10/2-1)/(1+10/5*x)^(15/2)", 0, Inf, c(3,10))
```

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```
# Example 9:Cauchy distribution x9 \leftarrow rARS(100,"1/(1+(x-1)^2)",-Inf,Inf,c(-2,2,10)) # Example 10:Rayleigh distribution with lambda=1 x10 \leftarrow rARS(100,"2*x*exp(-x^2)",0,Inf,c(0.01,10))
```

rASS

Adaptive Slice Sampling Algorithm With Stepping-Out Procedures

Description

rASS generates a sequence of random numbers by the adaptive slice sampling algorithm with stepping-out procedures.

Usage

```
rASS(n, x0 = 0, formula, w = 3)
```

Arguments

n Desired sample size;

x0 Initial value;

w Length of the coverage interval.

Author(s)

Dong Zhang <dzhang0716@126.com>

References

Neal R M. Slice sampling - Rejoinder[J]. Annals of Statistics, 2003, 31(3):758-767.

```
# Example 1: Sampling from exponential distribution with bounded domain x<-rASS(100,-1,"1.114283*exp(-(4-x^2)^2)",3) plot(density(x))
```

4 rCCARS

rCCARS

Concave-Convex Adaptive Rejection Sampling Algorithm

Description

rCCARS generates a sequence of random numbers by the concave-convex adaptive rejection sampling algorithm from target distributions with bounded domain.

Usage

```
rCCARS(n, cvformula, ccformula, min, max, sp)
```

Arguments

n Desired sample size;

cvformula, ccformula

Convex and concave decompositions for -ln(p(x)) where p(x) is the kernal of

target density;

min, max Domain except positive and negative infinity;

sp Supporting set

Details

Strictly speaking, the concave-convex adaptive rejection sampling algorithm can generate samples from target distributions who have bounded domains. For distributions with unbounded domain, rCCARS can also be used for sampling approximately. For example, if we want draw a sequence from N(0,1) by the concave-convex adaptive rejection sampling algorithm. We know that $X \sim N(0,1)$ has a so small probability in two tails that we can ingore the parts at both ends. Pr(X>20)=P(X<-20)=2.753624e-89, therefore we can get random numbers approximately from N(0,1) with the bound [-20,20]. Also, you can make this bound large enough to reduce sampling error.

Author(s)

Dong Zhang <dzhang0716@126.com>

References

Teh Y W. Concave-Convex Adaptive Rejection Sampling[J]. Journal of Computational & Graphical Statistics, 2011, 20(3):670-691.

```
# Example 1: Generalized inverse bounded gaussian distribution with lambda=-1 and a=b=2 x<-rCCARS(100,"x+x^-1","2*log(x)",0.001,100,1) hist(x,breaks=20,probability =TRUE);lines(density(x,bw=0.1),col="red",lwd=2,lty=2) f <- function(x) \{x^{(-2)*exp(-x-x^{(-1))/0.2797318}}
```

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```
lines(seq(0,5,0.01),f(seq(0,5,0.01)),lwd=2,lty=3,col="blue")

#The following examples are also available;
#But it may take a few minutes to run them.

# Example 2: Expontional bounded distribution
# x<-rCCARS(1000,"x^4","-8*x^2+16",-3,4,c(-2,1))
# hist(x,breaks=30,probability=TRUE);lines(density(x,bw=0.05),col="blue",lwd=2,lty=2)
# f <- function(x) exp(-(x^2-4)^2)/0.8974381
# lines(seq(-3,4,0.01),f(seq(-3,4,0.01)),col="red",lty=3,lwd=2)

# Example 3: Makeham bounded distribution
# x<-rCCARS(1000,"x+1/log(2)*(2^x-1)","-log(1+2^x)",0,5,c(1,2,3))
# hist(x,breaks=30,probability=TRUE);lines(density(x,bw=0.05),col="blue",lwd=2,lty=2)
# f <- function(x){(1+2^xx)*exp(-x-1/log(2)*(2^x-1))}
# lines(seq(0,5,0.01),f(seq(0,5,0.01)),col="red",lty=3,lwd=2,type="l")</pre>
```

rMARS

Modified Adaptive Rejection Sampling Algorithm

Description

rMARS generates a sequence of random numbers using the modified adaptive rejection sampling algorithm.

Usage

```
rMARS(n, formula, min = -Inf, max = Inf, sp, infp, m = 10^{-4})
```

Arguments

n	Desired sample size;
formula	Kernel of the target distribution;
min, max	Domain including positive and negative infinity of the target distribution;
sp	Supporting set;
infp	Inflexion set;
m	A parameter for judging concavity and convexity in a certain interval.

Author(s)

Dong Zhang <dzhang 0716 @126.com>

References

Martino L, Miguez J. A generalization of the adaptive rejection sampling algorithm[J]. Statistics & Computing, 2011, 21(4):633-647.

rMARS

```
# Example 1: Exponential distribution
x <- rMARS(100, "exp(-(4-x^2)^2)", -Inf, Inf, c(-2.5,0,2.5), c(-2/sqrt(3), 2/sqrt(3)))
hist(x,probability=TRUE,xlim=c(-3,3),ylim=c(0,1.2),breaks=20)
lines(density(x,bw=0.05),col="blue")
f \leftarrow function(x)(exp(-(4-x^2)^2))
lines(seq(-3,3,0.01), f(seq(-3,3,0.01))/integrate(f,-3,3)[[1]], lwd=2, lty=2, col="red")
#The following examples are also available;
#But it may take a few minutes to run them.
# Example 2: Distribution with bounded domain
# x <- rMARS(1000, "exp(-(x^2-x^3))", -3,2,c(-1,1),1/3)
# hist(x,probability=TRUE,xlim=c(-3,2.5),ylim=c(0,1.2),breaks=20)
# lines(density(x,bw=0.2),col="blue")
# f <- function(x) exp(-(x^2-x^3))
# lines(seq(-3,2,0.01),f(seq(-3,2,0.01))/integrate(f,-3,2)[[1]],lwd=2,lty=2,col="red",type="1")
# Example 3: Weibull distribution with k=3 and lambda=1
\# x < - rMARS(100, "3*x^2*exp(-x^3)", 10^-15, Inf, c(0.01, 1), (1/3)^(1/3), m=10^-4)
# hist(x,probability=TRUE,breaks=20,xlim=c(0,2))
# lines(density(x,bw=0.15),col="blue")
# f <- function(x) 3*x^2*exp(-x^3)
# lines(seq(0,2,0.01),f(seq(0,2,0.01)),lwd=2,lty=2,col="red",type="1")
# Example 4: Mixed normal distribution with p=0.3,m1=2,m2=8,sigma1=1,sigma2=2
\# x < - rMARS(100, "0.3/sqrt(2*pi)*exp(-(x-2)^2/2)+(1-0.3)/sqrt(2*pi)/2*exp(-(x-8)^2/8)", -Inf,Inf,
# c(-6,-4,0,3,6,15),c(-5.120801,-3.357761,3.357761,5.120801),m=10^-8)
# hist(x,breaks=20,probability=TRUE);lines(density(x,bw=0.45),col="blue",lwd=2)
# f <- function(x)0.3/sqrt(2*pi)*exp(-(x-2)^2/2)+(1-0.3)/sqrt(2*pi)/2*exp(-(x-8)^2/8)
# lines(seq(0,14,0.01),f(seq(0,14,0.01)),lty=3,col="red",lwd=2)
```

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