# Generating continuous random variables

Math 276 Actuarial Models

Spring 2008 semester

EA Valdez University of Connecticut - Storrs Lecture Weeks 4-5

#### Generating continuous random variables

#### FA Valdez





# The inverse transform

#### Some examples Simulating exponentials in

Gamma distribution

Simulating gamma in R Poisson distribution

### The rejection method

### Simulating half-normal in R

### Generating Normal

#### random variables Box-Muller transformations

- continued

Illustrating Box-Muller in R The polar method

Illustrating polar method in

### Generating a Poisson

Introduction

Generating the first T time units

Simulating a Poisson process in R

Alternative approach

Nonhomogeneous Poisson process

page 1



• Proposition: Let  $U \sim U(0,1)$ . For any continuous CDF F, the random variable X defined by

$$X = F^{-1}(U)$$

has distribution function F.

- Proof: in class.
- So long as we can derive the explicit form for  $F^{-1}$ , we can use this result to generate from a continuous distribution with CDF F.

Step 1: generate a random number *U*.

Step 2: set  $X = F^{-1}(U)$  and you are done.

Some examples Simulating exponentials in

Gamma distribution Simulating gamma in R Poisson distribution

The rejection method Simulating half-normal in R

#### Generating Normal random variables

Box-Muller transformations - continued Illustrating Box-Muller in R

The polar method Illustrating polar method in

## Generating a Poisson

Introduction Generating the first T time units Simulating a Poisson

process in R Alternative approach

$$F(x) = x^n, 0 < x < 1.$$

We can generate from this distribution by setting  $X = U^{1/n}$ after generating U.

**2** Exponential Suppose X comes from an  $Exp(\lambda)$  with PDF

$$f(x) = \lambda e^{-\lambda x}, x > 0.$$

We can generate from this distribution by setting  $X = -\frac{1}{3} \log U$  where *U* is the generated random number.

Weibull A random variable X is said to have a Weibull( $\alpha$ ,  $\beta$ ) distribution if its CDF has the form

$$F(x) = 1 - \exp(-\alpha x^{\beta}), \quad x > 0.$$

Describe a method to generate from this distribution.

Generating continuous random variables

FA Valdez





The inverse transform

#### Some examples

Simulating exponentials in Gamma distribution

Simulating gamma in R Poisson distribution

The rejection method Simulating half-normal in R

#### Generating Normal random variables

Box-Muller transformations - continued

Illustrating Box-Muller in R The polar method

Illustrating polar method in

## Generating a Poisson

Introduction

Generating the first T time units

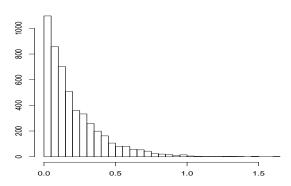
Simulating a Poisson process in R

Alternative approach

Simulating exponentials in R
The following executes commands in R to simulate from exponential with  $\lambda = 5$ .

```
> urandom <- runif(5000)
> lambda <- 5
> x.exp <- -log(urandom)/lambda
> x.exp[1:20]
 [1] 0.44388841 0.03379772 0.09910273 0.15201846 0.03336014 0.01021211
 [7] 0.22211658 0.14559882 0.49368172 0.18587106 0.01590227 0.07396419
[13] 0.19668067 0.38644089 0.05817653 0.16742193 0.10366767 0.06664743
[19] 0.21755809 0.29353129
> summary(x.exp)
     Min. 1st Qu.
                       Median
                                   Mean
                                          3rd On.
                                                       Max.
3.168e-05 5.675e-02 1.321e-01 1.975e-01 2.692e-01 1.479e+00
> hist(x.exp,br=25,xlab="",ylab="",main="5,000 simulations from Exponential")
```

#### 5,000 simulations from Exponential



#### Generating continuous random variables

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The inverse transform algorithm

Some examples Simulating exponentials in

Gamma distribution

Simulating gamma in R Poisson distribution

The rejection method

Simulating half-normal in R Generating Normal

random variables

Box-Muller transformations

- continued Illustrating Box-Muller in R

The polar method Illustrating polar method in

# Generating a Poisson

Introduction

Generating the first T time units

Simulating a Poisson process in R

Alternative approach





The inverse transform Some examples

Simulating exponentials in

Gamma distribution Simulating gamma in R

Poisson distribution The rejection method

Simulating half-normal in R Generating Normal

random variables Box-Muller transformations

- continued Illustrating Box-Muller in R

The polar method Illustrating polar method in

## Generating a Poisson

Introduction

Generating the first T time units

Simulating a Poisson process in R

Alternative approach

Nonhomogeneous Poisson process

- For a Gamma $(n, \lambda)$  random variable, note that because we cannot write an explicit form for the expression of the  $F^{-1}$ , it is diffcult to directly apply inverse transform method.
- However, recall that the sum of independent Exponentials leads us to a Gamma distribution.
- We can generate from a Gamma distribution by generating n random numbers  $U_1, U_2, \dots, U_n$  and then setting

$$X = -\frac{1}{\lambda} \log U_1 - \dots - \frac{1}{\lambda} \log U_n$$
$$= -\frac{1}{\lambda} \log(U_1 \dots U_n).$$

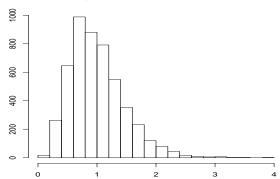
This works provided n is a positive integer.

> urandom1 <- runif(5000)

Simulating gamma in R
The following executes commands in R to simulate from gamma with n = 5 and  $\lambda = 5$ .

```
> urandom2 <- runif(5000)
> urandom3 <- runif(5000)
> urandom4 <- runif(5000)
> urandom5 <- runif(5000)
> u.prod <- urandom1*urandom2*urandom3*urandom4*urandom5
> lambda <-5
> x.gamma <- -log(u.prod)/lambda
> summary(x.gamma)
  Min. 1st Qu. Median Mean 3rd Qu.
                                          Max.
0.1140 0.6689 0.9277 0.9998 1.2490 3.9810
> hist(x.gamma,br=25,xlab="",ylab="",main="5,000 simulations from Gamma")
```

#### 5.000 simulations from Gamma



#### Generating continuous random variables

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The inverse transform

Some examples Simulating exponentials in

Gamma distribution

Simulating gamma in R

Poisson distribution

The rejection method Simulating half-normal in R

Generating Normal

random variables Box-Muller transformations

- continued Illustrating Box-Muller in R

The polar method Illustrating polar method in

# Generating a Poisson

Introduction

Generating the first T time units

Simulating a Poisson process in R

Alternative approach

### Generating a Poisson random variable

- Exploit the property of a Poisson process.
- Recall that for a Poisson process with rate  $\lambda$ , the times between successive events are independent exponentials with rate  $\lambda$ , and the number of events by time 1, N(1), is Poisson with mean  $\lambda$ .
- Thus, to generate from a Poisson with mean  $\lambda$ , we can
  - generate n successive interarrival times which are exponentials,  $X_1, X_2, \ldots, X_n$ .
  - determine the number of events by time 1 using

$$N(1) = \max\left(n: \sum_{i=1}^n X_i \le 1\right).$$

Generating continuous random variables

EA Valdez





The inverse transform algorithm

Some examples Simulating exponentials in

Gamma distribution Simulating gamma in R Poisson distribution

The rejection method

Simulating half-normal in R

Generating Normal random variables

Box-Muller transformations

- continued Illustrating Box-Muller in R

The polar method
Illustrating polar method in

### Generating a Poisson process

Introduction

Generating the first T time units

Simulating a Poisson process in R

Alternative approach

#### continued

• Equivalently, we generate  $U_1, \ldots, U_n, \ldots$  and set

$$N = \max \left( n : \sum_{i=1}^{n} -\frac{1}{\lambda} \log U_{i} \le 1 \right)$$

$$= \max \left( n : \sum_{i=1}^{n} \frac{1}{\lambda} \log U_{i} \ge -\lambda \right)$$

$$= \max(n : \log(U_{1} \cdots U_{n}) \ge -\lambda)$$

$$= \max(n : U_{1} \cdots U_{n} \ge e^{-\lambda})$$

This is also equivalent to setting

$$N = \min(n: U_1 \cdots U_n < e^{-\lambda}) - 1.$$

Generating continuous random variables

#### FA Valdez





The inverse transform

Some examples

Simulating exponentials in

Gamma distribution Simulating gamma in R

Poisson distribution

The rejection method Simulating half-normal in R

Generating Normal random variables

Box-Muller transformations - continued

Illustrating Box-Muller in R

The polar method Illustrating polar method in

# Generating a Poisson

Introduction

Generating the first T time units

Simulating a Poisson

process in R Alternative approach





The inverse transform algorithm

Some examples

Simulating exponentials in R

Gamma distribution
Simulating gamma in R

The rejection method

### Simulating half-normal in R

Generating Normal random variables

andom variables

Box-Muller transformations

- continued

The polar method

Illustrating polar method in

Generating a Poisson process

Introduction

Generating the first T time units

Simulating a Poisson

process in R

Alternative approach

Nonhomogeneous Poisson process

- Suppose we wish to generate X from a distribution with PDF f(x).
- Assume that we are able to generate Y from a distribution with PDF g(y) and that there is a constant c such that

$$\frac{f(y)}{g(y)} \le c$$
, for all  $y$ .

- According to the rejection method, we can generate X using the following steps:
- Step 1: generate Y from distribution with density g.

Step 2: generate a random number *U*.

Step 3: if  $U \leq \frac{f(Y)}{cg(Y)}$ , set X = Y.

Step 4: else, return to step 1.





# The inverse transform algorithm

#### Some examples Simulating exponentials in

Gamma distribution

Simulating gamma in R

#### The second second second

#### Simulating half-normal in R

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### Generating Normal

random variables
Box-Muller transformations

Box-Muller transformation: - continued

Illustrating Box-Muller in R
The polar method

Illustrating polar method in

### Generating a Poisson

### process

Introduction

Generating the first T time

units Simulating a Poisson

process in R

Alternative approach

Nonhomogeneous Poisson process

#### Theorem:

- The random variable X generated by the rejection method has density f.
- The number of iterations required for the rejection algorithm has a geometric distribution with mean c.

Proof: to be discussed in class.

Generating continuous random variables

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The inverse transform algorithm

Some examples

Simulating exponentials in R

Simulating gamma in R Poisson distribution

#### The rejection method

Simulating half-normal in R
Generating Normal

## random variables Box-Muller transformations

Box-Muller transformation - continued

Illustrating Box-Muller in R The polar method Illustrating polar method in

# Generating a Poisson process

Introduction
Generating the first T time

units Simulating a Poisson

process in R

Alternative approach

Nonhomogeneous Poisson process

Example 5d: Use the rejection method to generate from

$$f(x) = 20x(1-x)^3, \ 0 < x < 1.$$

 Example 5e: Use the rejection method to generate from a Gamma(3/2, 1) density with

$$f(x) = Kx^{1/2}e^{-x}, x > 0,$$

where 
$$K = 1/\Gamma(3/2) = 2/\sqrt{\pi}$$
.

- Example 5f: Suggest a rejection method for generating from a standard Normal random variable Z ~ N(0, 1).
- Example 5g: Suggest a rejection method for generating from a truncated Gamma(2,1), conditional on its value exceeding 5.
- These examples to be discussed in details in class.

### Simulating half-normal with the rejection method in R

- The following is a routine in R to simulate from the absolute value of a standard Normal random variable, using the rejection method.
- Function is called simhalfnorm.R.

```
# function to generate from a half-Normal distribution
# i.e. absolute value of a standard Normal
simhalfnorm <- function(n.gen) {
    sim.vector <- rep(0,n.gen)
# to start, generate 2 independent exponentials with rate 1
    for(i in 1:n.gen) {
        urandom <- runif(2); y <- -log(urandom)
        while(y[2] < (y[1]-1)^2/2) {
        urandom <- runif(2); y <- -log(urandom)
    }
    sim.vector[i] <- y[1]
}
# output
sim.vector</pre>
```

# Generating continuous random variables

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The inverse transform algorithm

Some examples Simulating exponentials in

R Gamma distribution

Simulating gamma in R Poisson distribution

The rejection method

### Simulating half-normal in R

### Generating Normal

Box-Muller transformations - continued

Illustrating Box-Muller in R The polar method

Illustrating polar method in R

## Generating a Poisson process

Introduction

Generating the first T time units

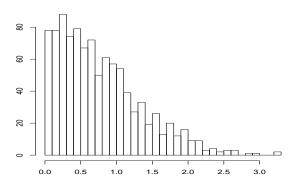
Simulating a Poisson process in R

Alternative approach

### **Executing the simulation**

```
> source("C:\\...\\Math276-Spring2008\\Rcodes-2008\\Week4\\simhalfnorm.R")
> out1 <- simhalfnorm(1000)
> out1[1:20]
 [11 0.30606099 1.03767270 0.95139082 1.89421610 0.41375272 1.09521146
 [7] 0.98360575 0.60245821 0.47552683 0.60963746 0.84500891 1.17122770
[13] 1.39765356 0.80675612 1.00840132 1.17619377 0.40388706 0.33065198
[19] 0.01615143 0.94438367
> summary(out1)
     Min.
          1st Ou.
                       Median
                                   Mean
                                          3rd Ou.
                                                       May
0.0008065 0.3058000 0.6472000 0.7694000 1.0780000 3.2830000
> sd(out1)
[11 0.5799377
> hist(out1,br=25,xlab="",ylab="",main="1,000 simulations from half-normal")
```

#### 1,000 simulations from half-normal



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### The inverse transform algorithm

Some examples Simulating exponentials in

R Gamma distribution

Simulating gamma in R

Poisson distribution

## The rejection method Simulating half-normal in R

### Generating Normal

random variables

Box-Muller transformations

- continued

Illustrating Box-Muller in R
The polar method

Illustrating polar method in R

### Generating a Poisson process

Introduction

Generating the first T time units

Simulating a Poisson

process in R

Alternative approach





The inverse transform algorithm

Some examples Simulating exponentials in

Gamma distribution
Simulating gamma in R

The rejection method
Simulating half-normal in R

Generating Normal

random variables

### Box-Muller transformations

Illustrating Box-Muller in R

Illustrating polar method in R

### Generating a Poisson process

Introduction
Generating the first T time

units Simulating a Poisson

process in R

Alternative approach

Nonhomogeneous Poisson process

 Suppose X and Y are independent standard Normal random variables with R and θ their corresponding polar coordinates:

$$R^2 = X^2 + Y^2$$
  
 $\tan \theta = \frac{Y}{X}$ 

• It can be shown that  $R^2$  and  $\Theta$  has joint density

$$f(d, \theta) = \frac{1}{2} \frac{1}{2\pi}, 0 < d < \infty, 0 < \theta < 2\pi.$$

- $R^2$  and  $\Theta$  are independent with the following marginals:
  - R<sup>2</sup> is exponential with mean 2, and
  - $\Theta$  is Uniform over  $(0, 2\pi)$ .

### - continued

• The idea then is to first generate their polar coordinates and then transform back to rectangular coordinates.

• This can be done with the following steps:

Step 1: generate two random numbers  $U_1$  and  $U_2$ .

Step 2: compute  $R^2 = -2 \log U_1$  and  $\theta = 2\pi U_2$ .

Step 3: then set

$$X = R \cos \Theta = \sqrt{-2 \log U_1} \cos(2\pi U_2)$$

$$Y = R \sin \Theta = \sqrt{-2 \log U_1} \sin(2\pi U_2)$$

These are known as Box-Muller transformations.

Generating continuous random variables

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The inverse transform algorithm

Some examples

Simulating exponentials in

Gamma distribution

Simulating gamma in R Poisson distribution

The rejection method

Simulating half-normal in R
Generating Normal

random variables

Box-Muller transformations

Illustrating Box-Muller in R The polar method Illustrating polar method in

Generating a Poisson

#### Generating a Poisson process

Introduction

Generating the first T time units

Simulating a Poisson

process in R

Alternative approach

### Simulating standard normal using Box-Muller in R

- The following is a routine in R to simulate from standard normal using the Box-Muller transformations.
- Function is called simpormbm.R.
- # function to generate standard Normal using the Box-Muller transformations

```
simnormbm <- function(n.gen) {
  urandom1 <- runif(n.gen)
  urandom2 <- runif(n.gen)
  Rsq <- -2*log(urandom1)
  theta <- 2*p1*urandom2
  x <- sqrt(Rsq)*cos(theta)
  y <- sqrt(Rsq)*sin(theta)
  y <- sqrt (xyq)*sin(theta)
  y <- (x+yy)/sqrt(2)
  # outbput
  z
  i</pre>
```

# Generating continuous random variables

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The inverse transform algorithm

Some examples Simulating exponentials in

Gamma distribution

Simulating gamma in R

The rejection method

Simulating half-normal in R

Generating Normal random variables

Box-Muller transformations

#### Illustrating Box-Muller in R

The polar method Illustrating polar method in

### Generating a Poisson process

Introduction

Generating the first T time units

> Simulating a Poisson process in R

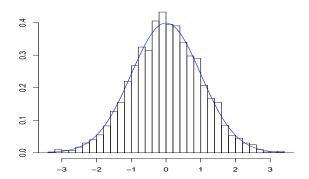
Alternative approach

### **Executing the simulation**

> curve(dnorm(x),from=-4,to=4,col="blue",add=TRUE)

```
> source("C:\\...\\Math276-Spring2008\\Rcodes-2008\\Week4\\simnormbm.R")
> out1 <- simnormbm(2000)
> out1[1:20]
[1] -0.56867837 2.20613520 -0.46835432 -0.55154500 -2.17445397 2.33841314
[71 1.12578412 0.36733281 -1.24346577 -0.41904992 -0.17311589 -0.95007900
[13] -0.04571634 2.75877598 -0.86521717 0.88587815 -0.92575967 -1.70796891
[19] -0.50454892 -1.11563401
> summary(out1)
     Min.
             1st Ou.
                         Median
                                      Mean
                                              3rd Ou.
                                                            Max.
-3.298e+00 -6.694e-01 6.473e-05 1.055e-02 6.969e-01 3.246e+00
> sd(out1)
[11 0.9935784
```

#### 2.000 simulations from Normal using Box-Muller



# Generating continuous random variables

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The inverse transform algorithm

Some examples Simulating exponentials in

[1] 0.9935784 Samma distribution from Normal using Box-Muller",freg=FALSE)

Simulating gamma in R
Poisson distribution

The rejection method
Simulating half-normal in R

Generating Normal

Box-Muller transformations

Illustrating Box-Muller in R

The polar method

Illustrating polar method in

Generating a Poisson process

Introduction

Generating the first T time

units Simulating a Poisson

process in R
Alternative approach

Nonhomogeneous Poisson

process

Some examples Simulating exponentials in

Gamma distribution

Simulating gamma in R Poisson distribution

The rejection method
Simulating half-normal in R

Generating Normal random variables

Box-Muller transformations - continued

Illustrating Box-Muller in R

The polar method Illustrating polar method in

Illustrating polar method in R

Generating a Poisson process

Introduction

Generating the first T time

units Simulating a Poisson

Simulating a Poiss process in R

Alternative approach Nonhomogeneous Poisson process

 The Box-Muller transformation is believed to be computationally inefficient because of the computation of the sine and cosine functions.

 The Polar method avoids these computations by generating random numbers U<sub>1</sub> and U<sub>2</sub> and setting

$$V_1 = 2U_1 - 1$$
 and  $V_2 = 2U_2 - 1$ .

ullet Compute  $S=R^2=V_1^2+V_2^2$  and then set

$$X = V_1 \sqrt{\frac{-2 \log S}{S}}$$

$$Y = V_2 \sqrt{\frac{-2 \log S}{S}}$$

Details in class. Algorithm is given on pages 81-82.

### Simulating standard normal using the polar method in R

- The following is a routine in R to simulate from standard normal using the polar method.
- Function is called simnormpolar.R.

```
# function to generate standard normal using the Polar method
simnormpolar <- function(n.gen) {
sim.vector <- rep(0,n.gen)
for (i in 1:n.gen) {
urandom <- runif(2)
v1 <- 2*urandom[1]-1; v2 <- 2*urandom[2]-1
s <- v1^2 + v2^2
while(s > 1){
urandom <- runif(2)
 v1 <- 2*urandom[1]-1; v2 <- 2*urandom[2]-1
s < -v1^2 + v2^2
insq < -2*log(s)/s
x <- sqrt(insq)*v1
v <- sart(insa) *v2
sim.vector[i] <- (x+y)/sqrt(2)
# output
sim.vector
```

Generating continuous random variables

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The inverse transform algorithm

Some examples

Simulating exponentials in

Gamma distribution

Simulating gamma in R

The rejection method

Simulating half-normal in R

Generating Normal random variables

Box-Muller transformations - continued

Illustrating Box-Muller in R
The polar method

Illustrating polar method in

### Generating a Poisson process

Introduction

Generating the first T time units

Simulating a Poisson process in R

Alternative approach

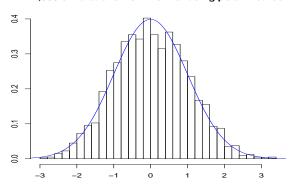
### **Executing the simulation**

> curve(dnorm(x),from=-4,to=4,col="blue",add=TRUE)

[11 1.000975

```
> source("C:\\...\\Math276-Spring2008\\Rcodes-2008\\Week4\\simnormpolar.R")
> out1 <- simnormpolar(2000)
> out1[1:20]
 [11 -0.53573046 0.58419468 -1.56107977 0.27961148 -1.37491965 -1.33070593
[71 0.09831912 -0.88206048 1.17231397 0.44666004 0.90755589 -1.38911674
    1.24109777 0.34494397 0.46953717 1.03635509 -0.53338976 1.18540414
[19] 0.29707077 -0.71940887
> summary(out1)
     Min.
          1st Ou.
                      Median
                                  Mean
                                         3rd Ou.
                                                      May
-2.926000 -0.704400 -0.008467 0.017280 0.720300 3.390000
> sd(out1)
```

#### 2.000 simulations from Normal using polar method



# Generating continuous random variables

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### The inverse transform algorithm

Some examples Simulating exponentials in

Gamma distribution
> hist(out1,br=25,xlab="",ylab="",main="2,000 simulations from Normal using polar method",freq=FALES
Smulating gamma in R

Poisson distribution

# The rejection method Simulating half-normal in R

Generating Normal

random variables
Box-Muller transformations

- continued Illustrating Box-Muller in R

The polar method

### Illustrating polar method in R

### Generating a Poisson process

Introduction

Generating the first T time units

Simulating a Poisson process in R

Alternative approach

Nonhomogeneous Poisson process

page 20





The inverse transform Some examples Simulating exponentials in

Gamma distribution

Simulating gamma in R Poisson distribution

The rejection method Simulating half-normal in R

Generating Normal random variables Box-Muller transformations

- continued Illustrating Box-Muller in R

The polar method Illustrating polar method in

Generating a Poisson

#### Introduction

Generating the first T time units

Simulating a Poisson process in R

Alternative approach

- To generate a Poisson process with rate  $\lambda$ , use the fact that the times between successive events are independent exponentials each with rate  $\lambda$ .
- Generate *n* random numbers  $U_1, \ldots, U_n$  and set  $X_i = -\frac{1}{3} \log U_i$ , the time between the (i-1)st and the *i*th event.
- The sum  $\sum_{i=1}^{j} X_i$ , for j = 1, ..., n then gives the actual time of the ith event.
- To generate then the first T time units of the process, follow the previous procedure and stopping when the sum then exceeds T.

# **B**



The inverse transform algorithm

Some examples

Simulating exponentials in R

Gamma distribution Simulating gamma in R

The rejection method
Simulating half-normal in R

Generating Normal

random variables
Box-Muller transformations

- continued Illustrating Box-Muller in R

The polar method

Illustrating polar method in

Illustrating polar r R

Generating a Poisson process

Introduction

Generating the first T time units

process

Simulating a Poisson process in R

Alternative approach
Nonhomogeneous Poisson

In the following algorithm,

t refers to time.

*I* is the number of events occurring by time t, and S(I) is the most recent event time.

 Generating the first T time units of a Poisson process with rate λ:

Step 1: start with t = 0, I = 0.

Step 2: generate a random number U.

Step 3: set  $t = t - \frac{1}{\lambda} \log U$  and STOP if t > T.

Step 4: reset I = I + 1, S(I) = t.

Step 5: return to Step 2.

• The values  $S(1), S(2), \ldots, S(I)$  are the I event times in increasing order.

### Simulating the arrival times of a Poisson process in R

Generating continuous random variables

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- The following is a routine in R to simulate the arrival times of a Poisson process.
- Function is called simpproc.R.

```
# simulating the arrival times of a Poisson process
# inputs are: number of simulations and lambda parameter
simpproc <- function(n.gen,lambda){
x \leftarrow rep(0, n.gen)
# generate exponential inter-arrival times
for(i in 1:n.gen){
urandom <- runif(1)
x[i] <- -log(urandom)/lambda
# computing the time of the n-th arrival
arrival.times <- c(0, cumsum(x))
# plotting the process
nn <- c(0:n.gen)
plot(arrival.times,nn,type="s",ylab="",xlab="arrival times", main="Simulated Poisson Process")
```

# The inverse transform

Some examples Simulating exponentials in

Gamma distribution

Simulating gamma in R

Poisson distribution

#### The rejection method

Simulating half-normal in R

#### Generating Normal random variables

Box-Muller transformations

- continued Illustrating Box-Muller in R

The polar method Illustrating polar method in

### Generating a Poisson

Introduction

Generating the first T time

units Simulating a Poisson

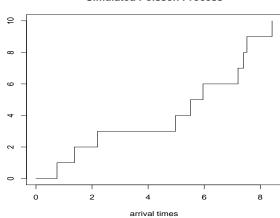
### process in R

Alternative approach

### **Executing the simulation**

- > source("C:\\...\Math276-Spring2008\\Rcodes-2008\\Week4\\simpproc.R")
- > simpproc(10,1.5)

#### Simulated Poisson Process



Generating continuous random variables

#### **EA Valdez**





#### The inverse transform algorithm

#### Some examples

Simulating exponentials in

Gamma distribution

Simulating gamma in R Poisson distribution

#### The rejection method Simulating half-normal in R

#### Generating Normal

### random variables

Box-Muller transformations - continued

Illustrating Box-Muller in R The polar method

Illustrating polar method in

### Generating a Poisson

#### Introduction

Generating the first T time units

Simulating a Poisson

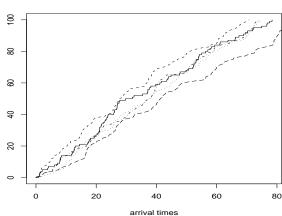
#### process in R

### Alternative approach

#### Several simulations

- > source("C:\\...\Math276-Spring2008\\Rcodes-2008\\Week4\\simpprocs.R")
- > simpprocs(100,1.2)

#### Several Simulated Poisson Processes



Generating continuous random variables

#### **EA Valdez**





#### The inverse transform algorithm

#### Some examples

Simulating exponentials in

Gamma distribution Simulating gamma in R

Poisson distribution

#### The rejection method Simulating half-normal in R

### Generating Normal

random variables Box-Muller transformations

- continued Illustrating Box-Muller in R

The polar method Illustrating polar method in

#### Generating a Poisson process

Introduction Generating the first T time

units Simulating a Poisson

### process in R

### Alternative approach



The inverse transform

Some examples Simulating exponentials in

Gamma distribution Simulating gamma in R Poisson distribution

The rejection method

Simulating half-normal in R Generating Normal random variables

Box-Muller transformations - continued Illustrating Box-Muller in R

The polar method Illustrating polar method in

# Generating a Poisson

Introduction Generating the first T time

units

Simulating a Poisson

process in R

Alternative approach

Nonhomogeneous Poisson process

• Recall that for a Poisson process with rate  $\lambda$ , the total number of events that occur by time t, N(T), is Poisson with mean  $\lambda T$ .

- The following general steps can then be followed:
  - **1** generate N(T) from a Poisson with mean  $\lambda T$ .
  - **2** generate N(T) random numbers  $U_1, \ldots, U_{N(T)}$ .
  - 0  $TU_1, TU_2, \ldots, TU_{N(T)}$  are taken as the event times by time T of the Poisson process.
- This works more efficiently than simulating from exponentials, provided all we are interested in were on the set of event times of the process.
- Often such is not the case however, we would also like the event times in increasing order.





### The inverse transform Some examples

#### Simulating exponentials in

Gamma distribution Simulating gamma in R

Poisson distribution

#### The rejection method Simulating half-normal in R

#### Generating Normal random variables

Box-Muller transformations - continued

Illustrating Box-Muller in R

The polar method Illustrating polar method in

### Generating a Poisson

Introduction

Generating the first T time

units Simulating a Poisson

process in R

Alternative approach process

Nonhomogeneous Poisson

 In the nonhomogeneous Poisson process, the rate depends on time t, where  $\lambda(t)$  is called the intensity function.

- One approach is called the thinning approach:
  - start by choosing a  $\lambda$  such that

$$\lambda(t) \leq \lambda$$
, for all  $t \leq T$ .

- randomly select the event times of a Poisson process with rate  $\lambda$ .
- If an event of a Poisson process with rate  $\lambda$  is counted with probability  $\lambda(t)/\lambda$ , then the process of counted events is the desired nonhomogeneous Poisson process.

### Algorithm with the thinning approach

Generating continuous random variables

FA Valdez





The inverse transform Some examples

Simulating exponentials in Gamma distribution

Simulating gamma in R Poisson distribution

The rejection method Simulating half-normal in R

Generating Normal random variables

Box-Muller transformations - continued

Illustrating Box-Muller in R The polar method

Illustrating polar method in

Generating a Poisson

Introduction Generating the first T time

units Simulating a Poisson

process in R Alternative approach

Nonhomogeneous Poisson process

 Applying the thinning approach, the following algorithm provides generating the first T time units of a nonhomogeneous Poisson process with intensity  $\lambda(t)$ :

Step 1: set t = 0, I = 0.

Step 2: generate a random number *U*.

Step 3: set  $t = t - \frac{1}{\lambda} \log U$  and if t > T, STOP. Step 4: generate another random number *U*.

Step 5: if  $U \le \lambda(t)/\lambda$ , set I = I + 1, S(I) = t.

Step 6: return to Step 2.

• The final value of I is the number of events by time T with the event times  $S(1), \ldots, S(I)$ .



The inverse transform algorithm

Some examples Simulating exponentials in

Gamma distribution Simulating gamma in R Poisson distribution

The rejection method Simulating half-normal in R

#### Generating Normal random variables

Box-Muller transformations - continued

Illustrating Box-Muller in R The polar method Illustrating polar method in

## Generating a Poisson

Introduction Generating the first T time

units Simulating a Poisson

process in R

Alternative approach

- An improvement to the approach, though less efficient, is to break up the whole interval into subintervals and then use the thinning procedure over each subinterval.
- In effect, determine suitable values k.  $0 = t_0 < t_1 < t_2 < \cdots < t_k < t_{k+1} = T, \lambda_1, \ldots, \lambda_{k+1}$  such that

$$\lambda(s) \leq \lambda_i$$
, if  $t_{i-1} \leq s < t_i$ ,  $i = 1, \ldots, k+1$ .

- Over each time interval  $(t_{i-1}, t_i)$ , generate exponentials with rate  $\lambda_i$  and accept the generated event with probability  $\lambda(s)/\lambda_i$ .
- This works because of the memoryless property.
- See page 85 for the algorithmic steps using subintervals.





The inverse transform Some examples

Simulating exponentials in Gamma distribution Simulating gamma in R

Poisson distribution The rejection method

Simulating half-normal in R Generating Normal

random variables Box-Muller transformations continued

Illustrating Box-Muller in R The polar method

Illustrating polar method in

Generating a Poisson

Introduction

Generating the first T time units

Simulating a Poisson

process in R

Alternative approach

- An alternative approach is to generate event times  $S_1, S_2, \ldots$  directly.
- This may be accomplished by generating first  $S_1$  from distribution function  $F_0$ ; then generate  $S_2$  by adding  $S_1$  to a generated value from  $F_{s_1}$ ; then generate  $S_3$  by adding  $S_2$ to a generated value from  $F_{s_2}$ , and so on.
- This alternative procedure works in cases where there are known efficient procedures for simulating from the F's, e.g. inverse transform.
- Suppose an event occurs at time s, additional time until the next event has distribution function:

$$F_s(x) = P(\text{time from } s \text{ until next event } \le x | \text{ event at s})$$

$$= 1 - \exp\left(-\int_0^x \lambda(s+y)dy\right).$$

### **Example 5h**

- Consider the example in Example 5h, pages 86-87.
- Suggest simulation procedure for a nonhomogeneous Poisson process with intensity function given by

$$\lambda(t) = \frac{1}{t+a}$$
, for  $t \ge 0$ .

Details in class.

Generating continuous random variables

**EA Valdez** 





The inverse transform algorithm

Some examples Simulating exponentials in

Gamma distribution

Simulating gamma in R Poisson distribution

The rejection method

Simulating half-normal in R

Generating Normal random variables

Box-Muller transformations

- continued

Illustrating Box-Muller in R
The polar method

Illustrating polar method in R

Generating a Poisson process

Introduction

Generating the first T time units

Simulating a Poisson

process in R

Alternative approach