

Rejection method for random number generation



Step by Step Explanation using R

Simulate Standard Normal Distribution (1/4)

```
fx <- function(x){  
  return((2/pi)^0.5*exp(-(x^2)/2))  
}
```

•Build the truncated normal function (divide Normal Distribution density function by 0.5)

```
b <- function(){  
  k <- rbinom(1,1,0.5)  
  if(k==0) return(-1)  
  else return(1)  
}
```

•Build function randomly generating 1 & -1
(relocate half of the accepted data to negative side)

Simulate Standard Normal Distribution (2/4)

```
set.seed(10)
```

```
n <- 10000
```

```
g <- rep(0,n)
```

```
count <- 0
```

```
for(i in 1:n){
```

```
  x <- rexp(1)
```

```
  y <- runif(1,0,(2*exp(1)/pi)^0.5*exp(-x))
```

```
  if (y<=fx(x)){
```

```
    g[i]<- x*b()
```

```
    count <- count+1
```

```
  }else{
```

```
    g[i] <- NA
```

```
  }
```

```
}
```

•Set simulation times

•Build empty vector

•Build variable record the acceptance times

•Generate random variable in positive real number field (from 0 to infinity)

•Generate the envelope

•If the value generate from the envelope smaller than the value generated from truncated normal distribution, then store into vector g and add one to count. Otherwise, store NA into vector g.

Simulate Standard Normal Distribution (3/4)

`k<-g[!is.na(g)]` → •Rule out the NA in the matrix

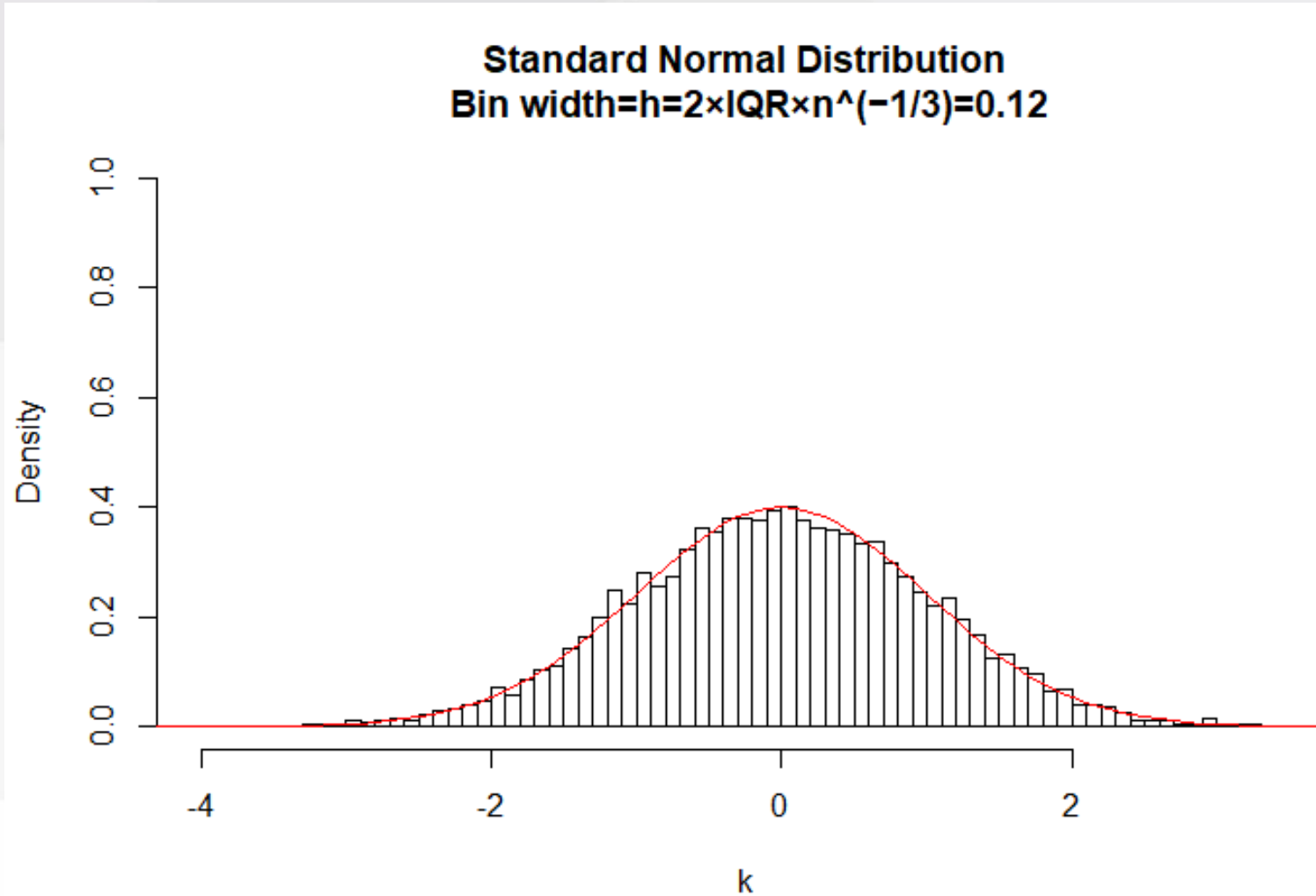
`x=seq(-5,5,by=0.1)`

`hist(k,breaks="FD",freq=F,ylim=c(0,1),main="Standard Normal Distribution")` → •Determine the bin width with Freedman-Diaconis rule

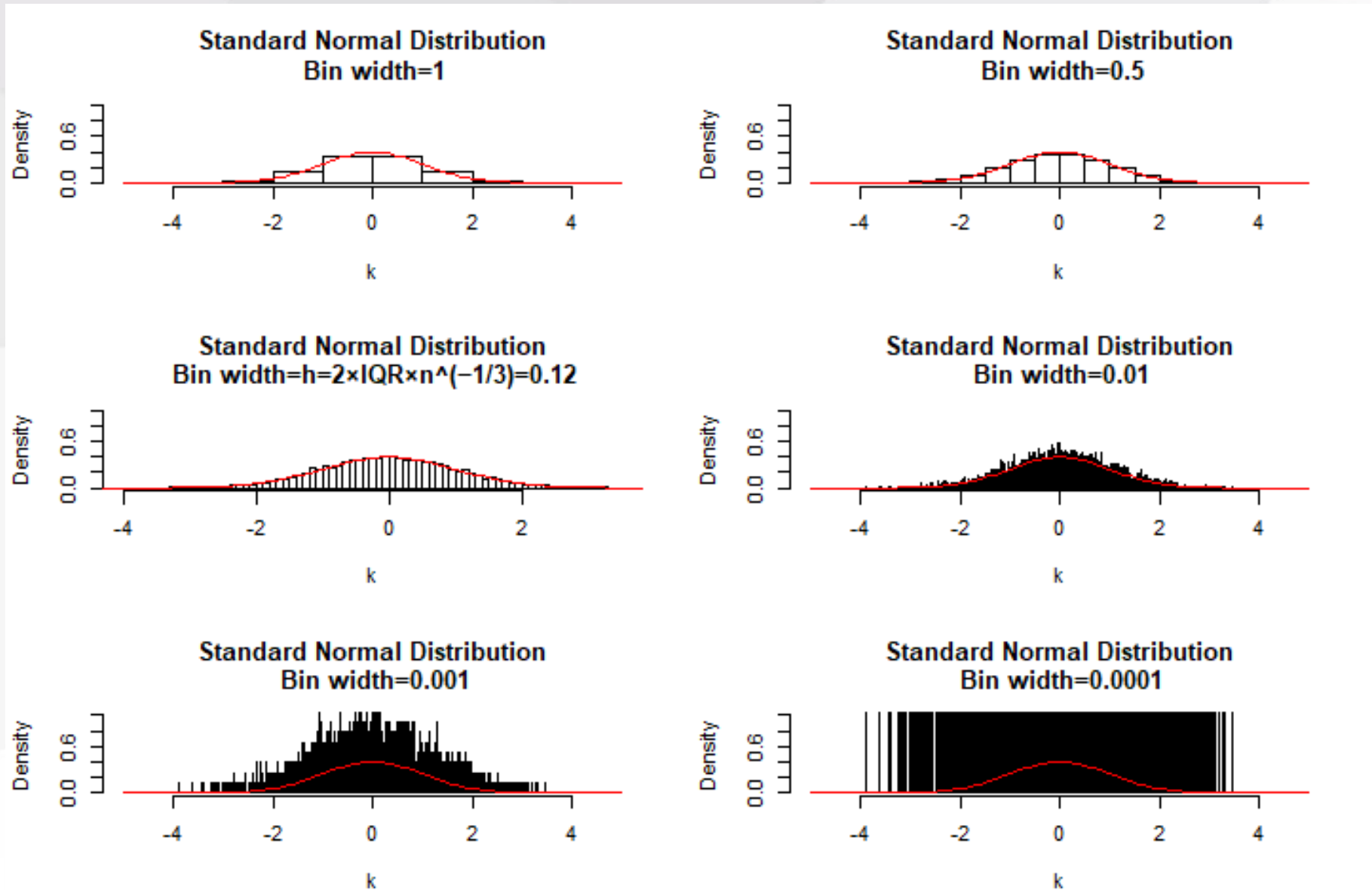
`lines(x,dnorm(x),col="yellow")` → •Plot real standard normal density

`ap_rate= count/n` → •Count acceptance rate

Simulate Standard Normal Distribution (4/4)



*Discussion: How to determine bin width?



*Discussion: How to determine bin width?

1. Sturges' formula

$$k = \lceil \log_2 n \rceil + 1$$

1. Doane's Rule

1. Scott's Rule

1. Rice's Rule

1. Freedman-Diaconis rule

$$h = \frac{3.5\hat{\sigma}}{\sqrt[3]{n}}$$

$$k = \lceil 2\sqrt[3]{n} \rceil$$

$$h = 2 \times \text{IQR} \times n^{-1/3}$$

$$\log_2(n) + 1 + \log_2\left(1 + \frac{\sqrt{b}}{\sigma\sqrt{b}}\right)$$

Where

$$\sqrt{b} = \frac{\sum_{i=1}^n (X_i - \bar{X})^3}{[\sum_{i=1}^n (X_i - \bar{X})^2]^{(3/2)}}$$

and

$$\sigma\sqrt{b} = \sqrt{\frac{6(n-2)}{(n+1)(n+3)}}$$

Application: Simulate Beta Distribution (1/3)

```
set.seed(10)
```

```
n <- 10000
```

```
g <- rep(0,n)
```

```
count <- 0
```

•Set simulation times

•Build empty vector

•Build variable record the acceptance times

```
for(i in 1:n){
```

```
  y1<-runif(1,0,1)
```

```
  y2<-runif(1,0,1)
```

```
  if (y2<=4*(y1*(1-y1))) {
```

```
    g[i]<-y1
```

```
    count <- count+1
```

```
  }else{
```

```
    g[i] <- NA
```

```
  }
```

```
}
```

•Generate random variable from uniform[0,1]

•Beta distribution with n=m

•If the value generate randomly from the niform[0,1] smaller than the value generated from Beta distribution, then store into vector g and add one to count. Otherwise, store NA into vector g.

Simulate Beta Distribution (2/3)

```
k<-g[!is.na(g)]
```

- Rule out the NA in the matrix

```
x=seq(0,1,by=0.001)
```

```
hist(k,breaks="FD",freq=F,ylim=c(0,3),main="beta")
```

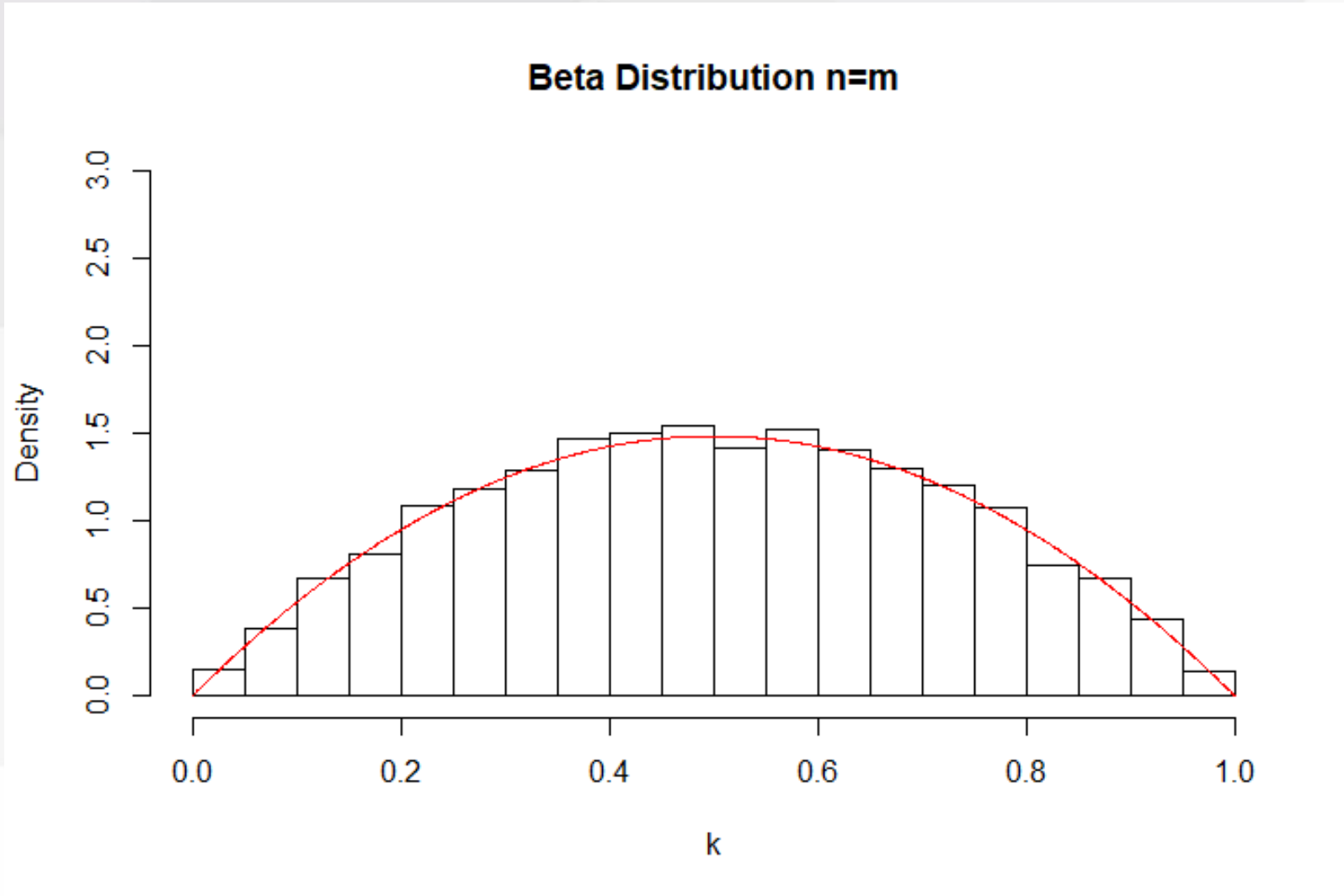
- Determine the bin width with Freedman-Diaconis rule

```
lines(x,5.93*x*(1-x),col="red")
```

```
ap_rate= count/n
```

- Plot real Beta distribution ($n=m$)
- Count acceptance rate

Simulate Beta Distribution (3/3)





THANKS!