

Probability: Random Variables in R

In this document we show how to find probabilities of Binomial and Normal random variables in R.

Binomial Random Variable

We start with this example: Suppose in one hospital 10 babies are born each day. The probability that a new born baby is a Boy is 50%. We are interested in the distribution of the number of boys born each day in this hospital. Let

Y = Number of Boys born on a day

We know that distribution of Y is Binomial with n = 10 and p = 0.50

 $Y \sim Bin(10, 0.5)$

Let's find 'Probability Mass Function' for this random variable. First we construct a variable 'x' that stores values 0, 1, 2...10 (since these values random variable Y can take). We can do it conveniently using 'seq' function ('by' argument specifies distance between adjacent values):

x=seq(from=0,to=10,by=1)
x

[1] 0 1 2 3 4 5 6 7 8 9 10

To calculate probabilities of 'Binomial' random variable we use 'dbinom' function ('size' argument determines the number of trials and 'prob' probability of success)

prob=dbinom (x,size=10,prob=0.5)

We can round these probabilities to 3 decimal places using 'round' function, and put 'x' and 'prob' variables together using 'rbind' to get a nice table

rbind(x,round(prob,3))

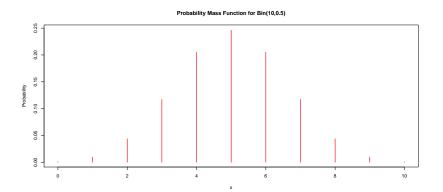
[,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10] [,11] x 0.000 1.00 2.000 3.000 4.000 5.000 6.000 7.000 8.000 9.00 1e+01 0.001 0.01 0.044 0.117 0.205 0.246 0.205 0.117 0.044 0.01 1e-03

From this table we can say for example that probability that 7 boys are born on a given day is 0.117:

$$P(Y=7) = 0.117$$

Next we want to plot 'Probability Mass Function', which is just 'prob' variable versus 'x':

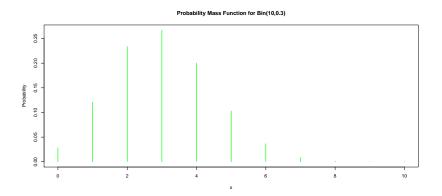
plot(x,prob,ylab='Probability',type='h',col='red',lwd=2, main='Probability Mass Function for Bin(10,0.5)')



Here 'type='h" produces vertical bars and 'lwd' specifies the width of these bars.

If the random variable Y had Bin(10, 0.3) then 'Probability Mass Function' would look like that:

```
x=seq(from=0,to=10,by=1)
prob=dbinom (x,size=10,prob=0.3)
plot(x,prob,ylab='Probability',type='h',col='green',lwd=2,
    main='Probability Mass Function for Bin(10,0.3)')
```



Observe that distribution is no longer symmetric but shifted to the left.

Suppose for the $Y \sim Bin(10, 0.3)$ we want to find probability of being less than or equal to 2. Then we can just add probabilities:

$$P(Y \le 2) = P(Y = 0) + P(Y = 1) + P(Y = 2)$$

dbinom(0,size=10,prob=0.3) + dbinom(1,size=10,prob=0.3) +
dbinom(2,size=10,prob=0.3)

[1] 0.3827828

So $P(Y \le 2) \approx 0.3828$. Instead of finding this probability by adding, it is much more convenient to use 'pbinom' function that calculates probability of being less than or equal of any value:

pbinom(2,size=10,prob=0.3)

[1] 0.3827828

Now we immediately get that $P(Y \le 2) \approx 0.3828$ without adding probabilities.

Normal Random Variable

Consider the next problem:

Suppose the birth weights of the infants born at a hospital follow a Normal distribution with mean of 3700 g and a standard deviation of 350 g.

We will work out the answers to these questions:

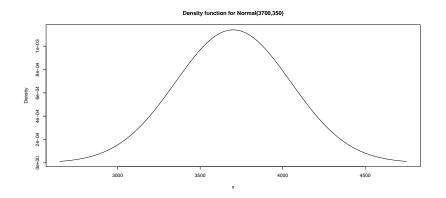
- 1. What percent of infants born at the hospital weigh less than 3000 grams?
- 2. What percent of infants weigh between 4000 and 4500 grams?
- 3. What birth weight is the first quartile?

Let X be a birth weight of an infant then we know:

 $X \sim N(3700, 350)$

Which means X has a normal distribution with mean 3700g and standard deviation 350g. Let's first plot the density of this normal using 'dnorm' function. First we create a variable 'x' that stores values from 2650 (mean -3 standard deviations) to 4750 (mean +3 standard deviations), then find density of N(3700, 350) evaluated at 'x' and finally make a plot:

```
x=seq(from=2650,to=4750,by=1)
y=dnorm(x,mean=3700,sd=350)
plot(x,y,ylab='Density',type='l',
    main='Density function for Normal(3700,350)')
```



This has a familiar bell shaped curve centered at 3700. Now let's look at the questions:

1. What percent of infants born at the hospital weigh less than 3000 grams?

In this question we need to find P(X < 3000). We can easily find this quantity using 'pnorm' function.

pnorm(q=3000, mean=3700, sd=350)

[1] 0.02275013

So we have the answer:

$$P(X < 3000) \approx 0.0228$$

Equivalently we can first standardize 3000 by subtracting the mean and divide by standard deviation, and then find probability for **Standard** Normal random variable to be less than this standardized number:

```
z=(x-3700)/350
pnorm(z, mean=0, sd=1)
```

[1] 0.02275013

Exactly the same probability!

2. What percent of infants weigh between 4000 and 4500 grams?

Here we need $P(4000 \le X < 4500)$. Observe that we can find this probability by calculating P(X < 4000) and P(X < 4500) since

$$P(4000 \le X < 4500) = P(X < 4500) - P(X < 4000)$$

We can find these two probabilities as in the first question.

pnorm(4500, mean=3700, sd=350) - pnorm(4000, mean=3700, sd=350)

[1] 0.1845475

Hence:

 $P(4000 \le X < 4500) \approx 0.1846$

3. What birth weight is the first quartile?

Here we have an opposite problem, we need x such that P(X < x) = 0.25. This value is found from 'qnorm' function.

qnorm(p=0.25, mean=3700, sd=350)

[1] 3463.929

So we get

P(X < 3463.929) = 0.25

And hence the first quartile is about 3464g.

Equivalently we can first use 'qnorm' for standard normal and then multiply the result by standard deviation and add the mean:

```
qnorm(p=0.25, mean=0, sd=1) * 350 +3700
```

[1] 3463.929

Summary of R Functions

We give a short summary of all new and/or important R functions [and arguments] that we used in this Module:

Probabilities

```
dbinom() [x,size,prob]
pbinom() [q,size,prob]
dnorm() [x,mean,sd]
pnorm() [q,mean,sd]
qnorm() [p,mean,sd]
```

Miscellaneous

rbind()
round()
seq() [from,to,by]