

# Probability Distributions

## Part II

---

### Session 3

Module 1 Probability & Statistical Inference

**The Summer Institutes**

DEPARTMENT OF BIostatISTICS

SCHOOL OF PUBLIC HEALTH

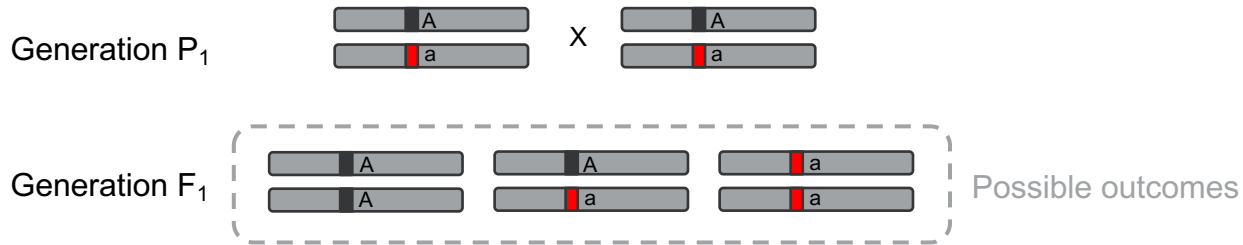
UNIVERSITY *of* WASHINGTON



# Multinomial Distribution

This distribution generalizes beyond 2 outcomes of the binomial distribution.

For example, let's consider a cross between two parents that are heterozygous carriers for a recessive trait:



We can then use the multinomial distribution to calculate the probability of observing various outcomes in the offspring generation.

Session 1




PROBABILITY AND  
INFERENCE STATISTICS

UNIVERSITY of WASHINGTON



# Multinomial Distribution

For any given offspring, the 3 possible outcomes can be represented by:

$Y_{i1} = 1$ if $i^{\text{th}}$ offspring is <b>unaffected</b> (AA), = 0 otherwise	
$Y_{i2} = 1$ if $i^{\text{th}}$ offspring is a <b>carrier</b> (Aa), = 0 otherwise	
$Y_{i3} = 1$ if $i^{\text{th}}$ offspring is <b>affected</b> (aa), = 0 otherwise	

Only one of  $Y_{i1}$ ,  $Y_{i2}$ ,  $Y_{i3}$  can be equal to 1, so  $Y_{i1} + Y_{i2} + Y_{i3} = 1$ .

For the binomial distribution with 2 outcomes, there are  $2^n$  unique outcomes in  $n$  trials. With  $n=3$  offspring, there are  $2^3 = 8$  unique outcomes.

For the multinomial distribution with 3 outcomes, the number of unique outcomes in  $n$  trials is  $3^n$ . With  $n=3$  offspring, there are  $3^3=27$  unique outcomes.

# Multinomial distribution

To calculate probabilities of interest, we can use **combinations**. For the multinomial distribution, the combinations are calculated as:

$$C_k^n = \frac{n!}{k_1! k_2! \dots k_j!}$$

where  $k_j$  ( $j=1, 2, \dots, J$ ) correspond to the totals for the different outcomes

Let's consider a scenario where:  $n = 2$  offspring  
 $J = 3$  possible outcomes (unaffected, carrier, affected)

The possible outcomes are:

Offspring 1	Offspring 2	Outcome
AA	AA	2 unaffected, 0 carrier, 0 affected
AA	Aa	1 unaffected, 1 carrier, 0 affected
Aa	AA	1 unaffected, 1 carrier, 0 affected
AA	aa	1 unaffected, 0 carrier, 1 affected
aa	AA	1 unaffected, 0 carrier, 1 affected
Aa	Aa	0 unaffected, 2 carrier, 0 affected
aa	Aa	0 unaffected, 1 carrier, 1 affected
Aa	aa	0 unaffected, 1 carrier, 1 affected
aa	aa	0 unaffected, 0 carrier, 2 affected

**Session 1**  
 PROBABILITY AND  
 INFERENCE STATISTICS

Number of unique outcomes  
 $3^n = 3^2 = 9$



# Multinomial distribution

For a defined number of offspring, what is the probability of a specific outcome? E.g., for  $n=2$ , what is the probability of observing two unaffected individuals? Or two affected? Or...?

First, write the outcomes in terms of  $k$ 's:  $k_1$ =number of unaffected  
 $k_2$ =number of carriers  
 $k_3$ =number of affected

Offspring 1	Offspring 2	Outcome	# ways
$p_1$	$p_1$	$k_1=2, k_2=0, k_3=0$	1
$p_1$	$p_2$	$k_1=1, k_2=1, k_3=0$	2
$p_2$	$p_1$	$k_1=1, k_2=1, k_3=0$	2
$p_1$	$p_3$	$k_1=1, k_2=0, k_3=1$	2
$p_3$	$p_1$	$k_1=1, k_2=0, k_3=1$	2
$p_2$	$p_2$	$k_1=0, k_2=2, k_3=0$	1
$p_3$	$p_2$	$k_1=0, k_2=1, k_3=1$	2
$p_2$	$p_3$	$k_1=0, k_2=1, k_3=1$	2
$p_3$	$p_3$	$k_1=0, k_2=0, k_3=2$	1

Formula for # ways

$$\frac{n!}{k_1!k_2!k_3!}$$

Probability for each possible outcome

$$\Pr[Y_1=k_1, Y_2=k_2, Y_3=k_3] = p_1^{k_1} p_2^{k_2} p_3^{k_3}$$

The probability of a **specified outcome** is going to be:

[prob of the possible outcome] x [number of ways for that outcome]

# Multinomial distribution

The probability that a multinomial random variable with  $n$  trials and success probabilities  $p_1, p_2, \dots, p_J$  will yield exactly  $k_1, k_2, \dots, k_J$  successes is:

Probability of a specific scenario

# ways

Probability of the outcome

$$P(Y_1 = k_1, Y_2 = k_2, \dots, Y_J = k_J) = \frac{n!}{k_1! k_2! \dots k_J!} p_1^{k_1} p_2^{k_2} \dots p_J^{k_J}$$

Defines what we are asking, e.g., for  $n=2$ , what is the probability of observing 1 unaffected ( $k_1=1$ ), 1 carrier ( $k_2=1$ ), and 0 affected ( $k_3=0$ )

The  $p$  probabilities define the "baseline" probability of success for each of the  $J$  outcomes

For heterozygous cross:

$p_1 = P(AA) = 0.25$   
 $p_2 = P(Aa) = 0.5$   
 $p_3 = P(aa) = 0.25$

Recall:  
 $k_1$ =number of unaffected  
 $k_2$ =number of carriers  
 $k_3$ =number of affected

These again are the  $k$  values, which are given based on what we are asking



# Multinomial distribution

The probability that a multinomial random variable with  $n$  trials and success probabilities  $p_1, p_2, \dots, p_j$  will yield exactly  $k_1, k_2, \dots, k_j$  successes is:

$$P(Y_1 = k_1, Y_2 = k_2, \dots, Y_j = k_j) = \frac{n!}{k_1! k_2! \dots k_j!} p_1^{k_1} p_2^{k_2} \dots p_j^{k_j}$$

## Assumptions:

- $J$  possible outcomes; only one can be a success, 1, in a given trial.
- The probability of success for each possible outcome,  $p_j$ , is the same for each trial.
- The outcome of one trial has no influence on other trials (independent trials).
- Interest is in the (sum) total number of successes over all the trials.

Session 1

PROBABILITY AND  
INFERENCE STATISTICS

UNIVERSITY of WASHINGTON



# Multinomial distribution

What is the probability that one of  $n=3$  offspring will be unaffected (AA), one will be affected (aa) and one will be a carrier (Aa)?

Our  $k$  values for this scenario are:  $k_1 = \text{number of unaffected} = 1$   
 $k_2 = \text{number of carriers} = 1$   
 $k_3 = \text{number of affected} = 1$

$$P(Y_1 = k_1, Y_2 = k_2, \dots, Y_J = k_J) = \frac{n!}{k_1! k_2! \dots k_J!} p_1^{k_1} p_2^{k_2} \dots p_J^{k_J}$$

$$\begin{aligned} P(Y_1 = 1, Y_2 = 1, Y_3 = 1) &= \frac{3!}{1!1!1!} p_1^1 p_2^1 p_3^1 \\ &= \frac{(3)(2)(1)}{(1)(1)(1)} \left(\frac{1}{4}\right)^1 \left(\frac{1}{2}\right)^1 \left(\frac{1}{4}\right)^1 \\ &= \frac{3}{16} = 0.1875. \end{aligned}$$

For heterozygous cross:



$$\begin{aligned} p_1 &= P(AA) = 0.25 \\ p_2 &= P(Aa) = 0.5 \\ p_3 &= P(aa) = 0.25 \end{aligned}$$

Session 1

PROBABILITY AND  
INFERENCE STATISTICS

UNIVERSITY of WASHINGTON





# Multinomial distribution

## Calculating the mean and variance

The marginal outcomes of the multinomial distribution are binomial.

We can obtain the means for each outcome, e.g.  $Y_j = k_j$ , the  $j^{\text{th}}$  outcome, as follows:

$$\begin{aligned}\text{Mean:} \quad E[k_j] &= E\left[\sum_{i=1}^n Y_{ij}\right] = \sum_{i=1}^n E[Y_{ij}] \\ &= \sum_{i=1}^n p_j = np_j\end{aligned}$$

$$\begin{aligned}\text{Variance:} \quad V[k_j] &= V\left[\sum_{i=1}^n Y_{ij}\right] = \sum_{i=1}^n V[Y_{ij}] \\ &= \sum_{i=1}^n p_j(1-p_j) = np_j(1-p_j)\end{aligned}$$

# Multinomial distribution

## Multinomial distribution summary

- Multinomial random variables are discrete
- Parameters are  $n, p_1, p_2, \dots, p_j$
- Each outcome  $Y_j = k_j$  is the sum of  $n$  independent Bernoulli outcomes
- Extends binomial distribution
- Seen in contingency tables, polytomous regression

# Paws



Work  
through  
questions  
1-2

# Continuous distributions

---

**Session 1**

PROBABILITY AND  
INFERENCE STATISTICS

UNIVERSITY *of* WASHINGTON



# Continuous distributions

For measurements like height or weight, it does not make sense to talk about the probability of any single value. Instead, we talk about the probability for an **interval**.

$$P[\text{weight} = 70.000\text{kg}] \approx 0$$

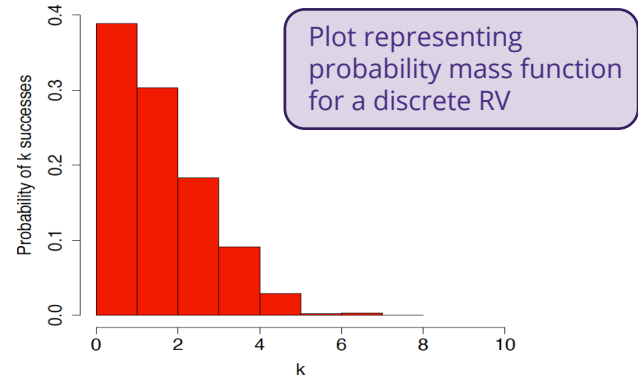
$$P[69.0\text{kg} \leq \text{weight} \leq 71.0\text{kg}] = 0.08$$

For discrete random variables, a **probability mass function** gives the probability of each possible value.

For continuous random variables, we require a **probability density function** to tell us about the probability of obtaining a value within an interval.

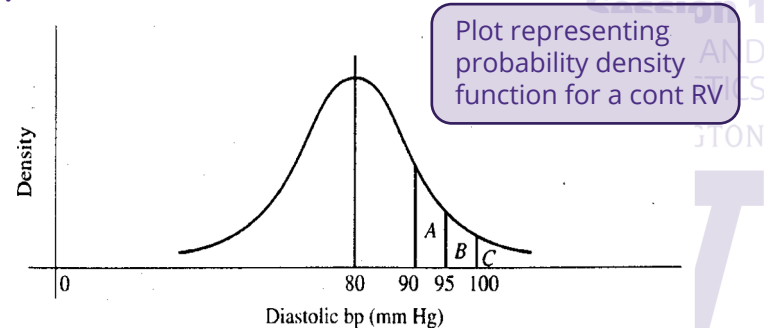
# Continuous distributions

With discrete probability distributions, we can determine the probability of a single outcome:



With continuous probability distributions, we determine the probability across a range of outcomes:

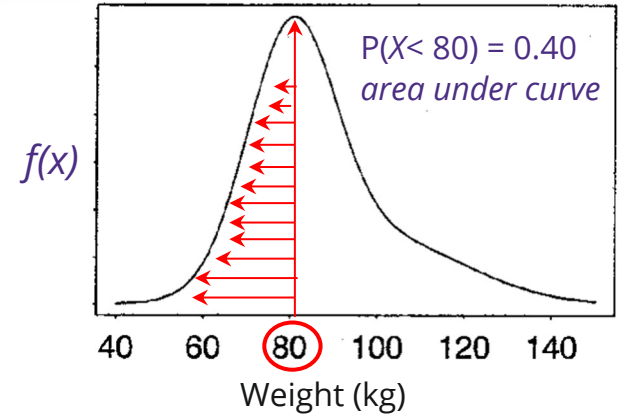
For any interval, the **area under the curve** represents the probability of obtaining a value in that interval.



# Continuous distributions

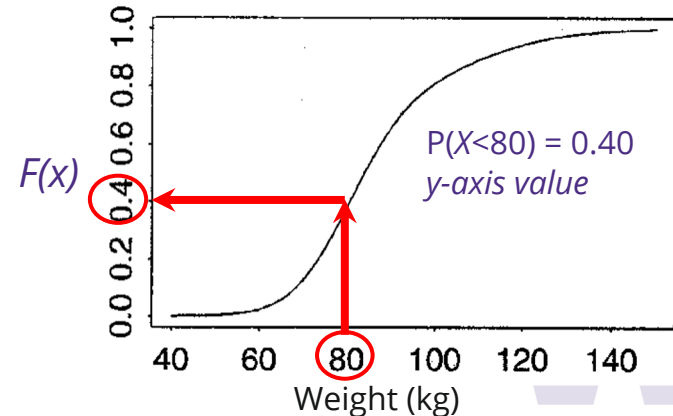
## Probability density function

- Given by  $f(x)$
- Gives probability that  $X$  falls within an interval:  
 $f(x) = P(\text{value 1} < X < \text{value 2})$
- Probability represented by area under curve
- Total area under curve is 1:  $\int f(x)dx = 1.0$



## Cumulative distribution function

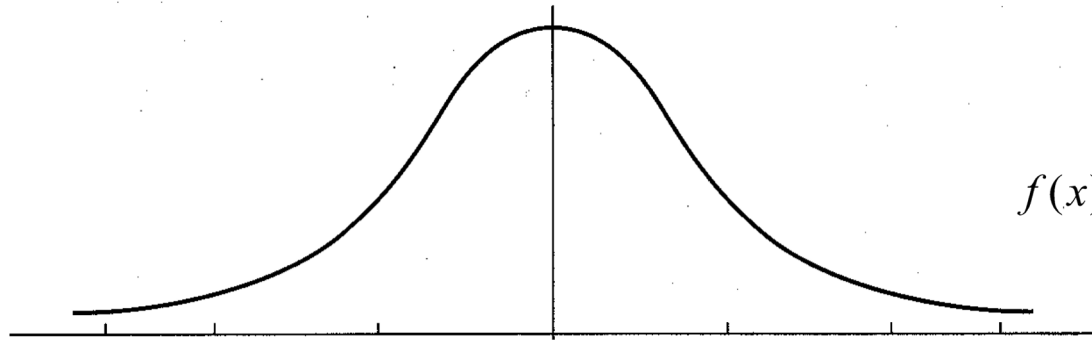
- Given by  $F(x)$
- Gives cumulative probability that  $X$  is less than some value  $x$ :
- $F(x) = P(X \leq x)$
- y-axis ceiling is 1



The **PDF** and **CDF** represent the **same information**.

# Normal distribution

The normal distribution is a well-known probability model for continuous data. It is unimodal with a “bell-shaped curve”.



PDF of the Normal distribution

$$f(x) = \frac{1}{\sigma\sqrt{2\pi}} \exp\left(-\frac{1}{2} \frac{(x-\mu)^2}{\sigma^2}\right)$$

Random variable values range from  $-\infty$  to  $+\infty$ .

Symmetric about mean: **mean = median = mode**

Common examples include human height, birth weight, blood pressure.

**Session 1**  
PROBABILITY AND  
INFERENCE STATISTICS  
UNIVERSITY of WASHINGTON





# Normal distribution

The NOR distribution is defined by its mean and variance. Note the appearance of  $\mu$  and  $\sigma$  in the probability density function for the NOR distribution:

$$f(x) = \frac{1}{\sigma\sqrt{2\pi}} \exp\left(-\frac{1}{2} \frac{(x-\mu)^2}{\sigma^2}\right)$$

Thus, the normal distribution has two parameters:

$\mu$  = the mean of X

$\sigma$  = the standard deviation of X

$$X \sim N(\mu, \sigma^2)$$

“X is normally distributed with mean  $\mu$  and variance  $\sigma^2$ ”

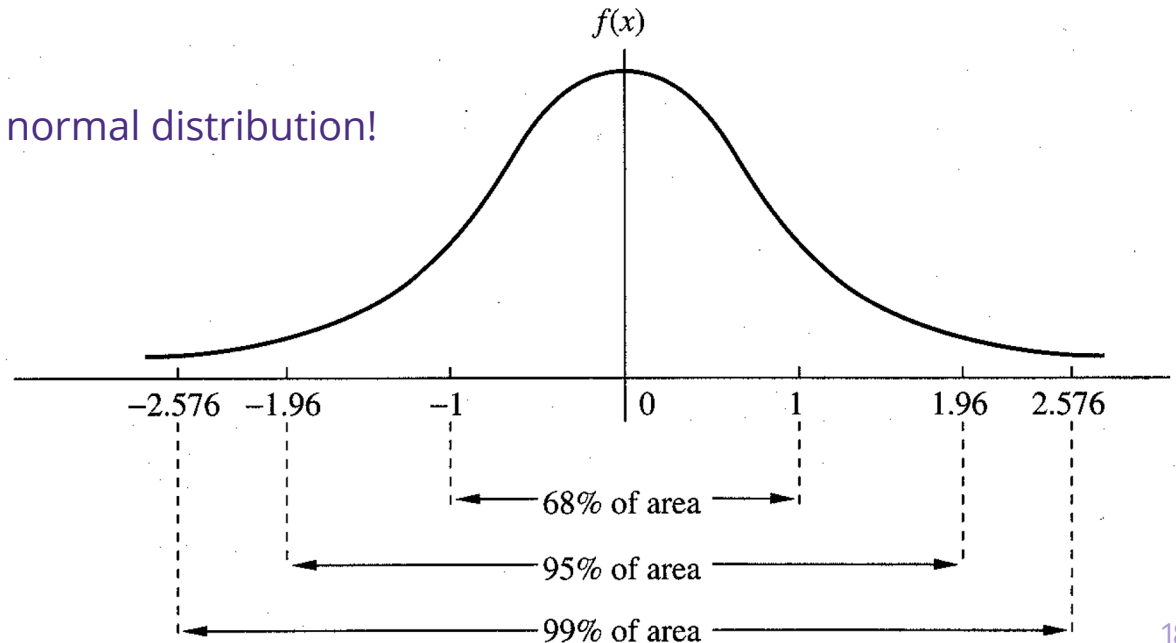
# Normal distribution

## The standard normal distribution

The standard normal is a special case of the NOR!

- AKA “z” distribution
- $Z \sim N(0, 1)$ :  $\mu = 0$  and  $\sigma^2 = 1$
- There is only ONE standard normal distribution!

Under the STD NOR, 95% of the area lies between  $\sim 2$  standard deviations of the mean. Useful!



# Standard Normal distribution

## Calculating probabilities for the standard normal distribution

Using  $z$  notation, the **probability density function** of  $Z$ , the random variable of the standard normal distribution is:

$$f(z) = \frac{1}{\sqrt{2\pi}} \exp\left(-\frac{1}{2}z^2\right)$$

The **cumulative distribution function** of  $Z$  is:

$$P(Z \leq x) = \Phi(x) = \int_{-\infty}^x \frac{1}{\sqrt{2\pi}} \exp\left(-\frac{1}{2}z^2\right) dz$$

Any computing software will give the values of  $f(z)$  and  $\Phi(x)$ .

# Standard Normal distribution

## Online calculators for standard Normal distribution probabilities

Suppose we wish to solve a probability statement for the standard Normal variable. For example, what is the probability that Z takes a value less than 0.05?

$$P(Z \leq 0.5) = ?$$



cdf normal distribution calculator

All Images Videos News Shopping More Settings Tools

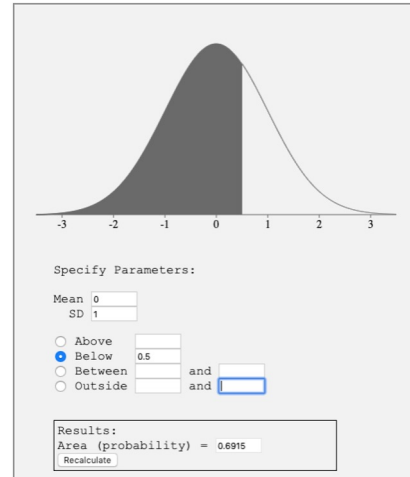
About 1,730,000 results (0.44 seconds)

[Normal Distribution Calculator - Online Stat Book](#)

[onlinestatbook.com](#) > [calculators](#) > [normal\\_dist](#)

**Normal Distribution.** Author(s). David M. Lane. Help support this free site by buying your books from Amazon following this link: Books on science and math.

[Areas under Normal Distribution](#) · [Binomial Distribution](#) · [Power Calculator](#)



$$P(Z \leq 0.5) = 0.6915$$

Session 1  
PROBABILITY AND  
INFERENCE STATISTICS  
UNIVERSITY of WASHINGTON



# Standard Normal distribution

## Properties

If the probability of ( $Z < \text{some value } z$ ) is equal to  $a$ , then the probability that ( $Z > z$ ) is equal to  $1 - a$ .

$$P(Z < z) = a$$

$$P(Z > z) = 1 - a$$

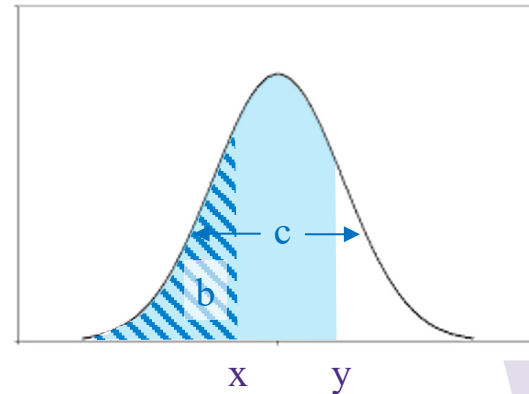
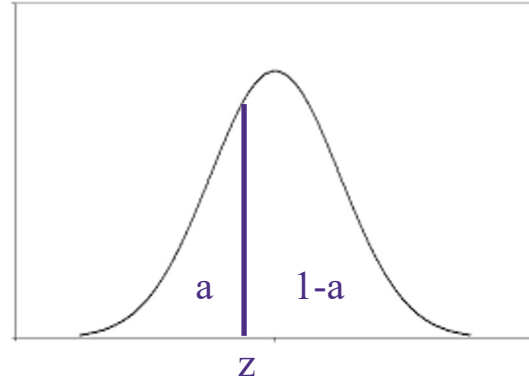
If the probability of ( $Z < \text{some value } x$ ) is equal to  $b$ , and the probability that ( $Z > y$ ) is equal to  $c$ , then the probability that ( $Z$  lies between  $x$  and  $y$ ) is equal to  $c - b$ .

$$P(Z < x) = b$$

$$P(Z < y) = c$$

$$P(x < Z < y) = c - b$$

Note that this is true of all distributions!



# Standard Normal distribution

## Properties

Because the standard Normal distribution is symmetrical around 0, the probability that  $(Z < -y)$  is equal to the probability that  $(Z > y)$ , shown here as  $d$ .

$$P(Z < -y) = P(Z > y) = d$$

Note that this is NOT true of all distributions!

