

# Probability!

*Probability* - The fraction of observations of an event given multiple repeated independent observations.

## A Feeding Trial Example

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45 eat.  $P(\text{eats}) = \frac{45}{50} = 0.9$



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Now you offer  
50 others a **treated** leaf.  
10 eat.  $P(\text{eats}) = \frac{10}{50} = 0.2$



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What is the probability of **not** eating if you are fed a treated leaf?



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$$P(!A) = 1 - P(A)$$



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$$P(A \text{ or } B) = P(A) + P(B)$$



## Two Events

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$$P(A \text{ and } B) = P(A)P(B)$$



## Two Conditional Events

If we are interested in the probability of eating twice - i.e. the probability of eating a second time *given* that a budworm ate once, we phrase that somewhat differently.

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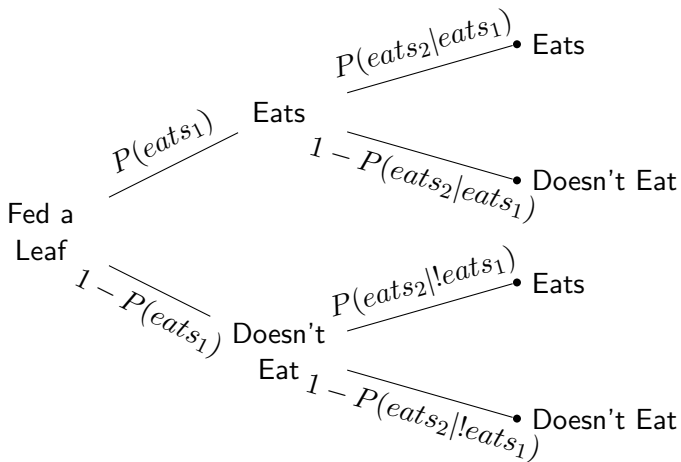
$$P(eats_2|eats_1)$$

So,  $P(A \text{ and } B) = P(A)P(B||A)$

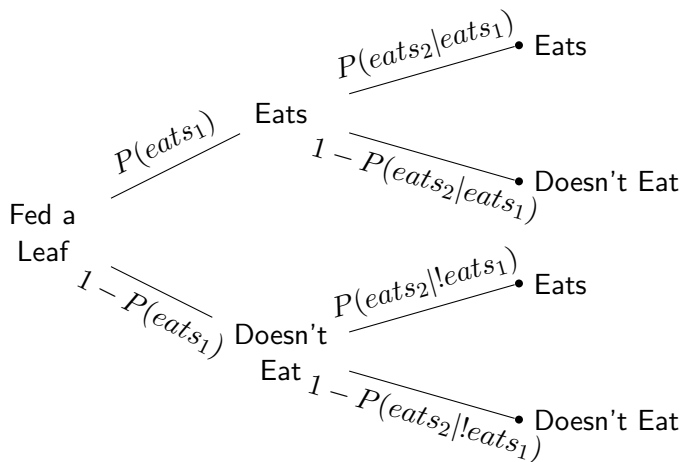




# Probability Tree



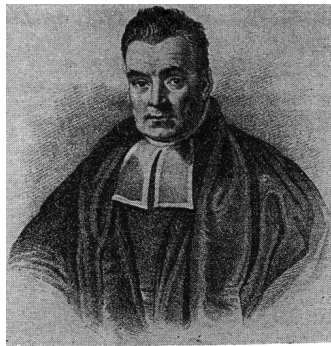
# Probability Tree



$$P(eats_2) = P(eats_2|eats_1) + P(eats_2|!eats_1)$$

# Bayes Theorem

$$P(A|B)P(B) = P(B|A)P(A)$$



REV. T. BAYES

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So...

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$



REV. T. BAYES

## GMO Collateral Damage

Let's say you had a rare but extremely harmful budworm munching ravenously through your fields. You've developed a really effective GMO tobacco leaf to help stop it. It has a 75% kill rate. And, miraculously, it only has a 15% kill rate of non-budworms. Given that the budworms make up about 10% of the insects in a field, what's porportion of dead insects WON'T be budworms?

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$1 - 0.357 = 0.643$  - the majority of the dead!

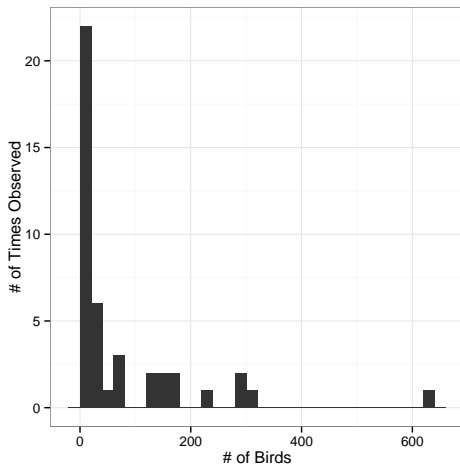
# Why are we talking about this??

As we test hypotheses in a *frequentist* framework, we'll be asking about the probability of observing data given that a hypothesis is true -  $P(\text{Data} \parallel \text{Hypothesis})$ .

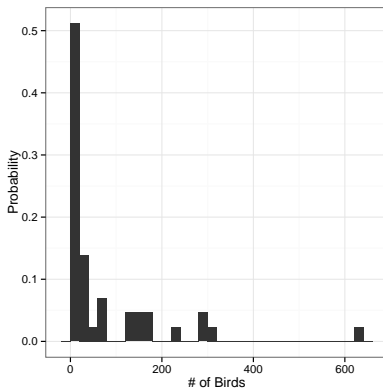
# Distributions!

(when a point probability just ain't enough)

# Frequency Distributions Make Intuitive Sense

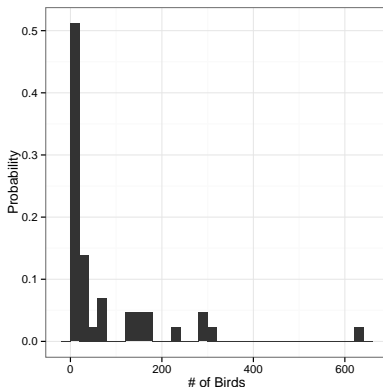


# Frequencies Can be Turned Into Probabilities



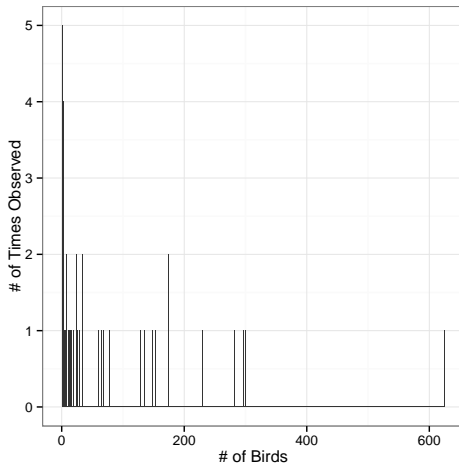
Just divide by total # of observations

# Frequencies Can be Turned Into Probabilities



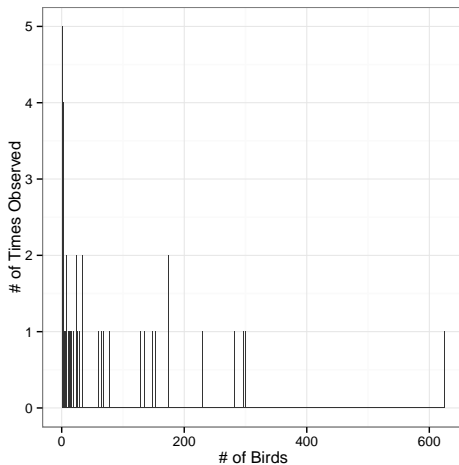
Just divide by total # of observations  
But - we have binned observations...

# Frequencies of Individual Observations

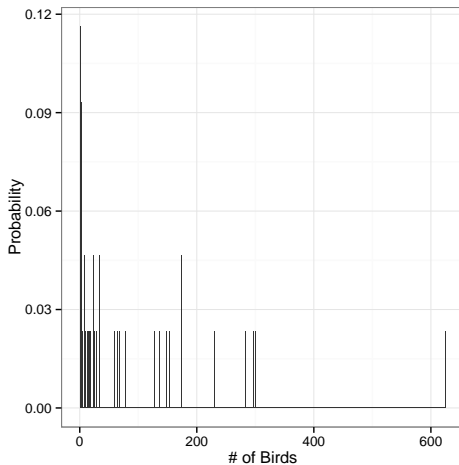




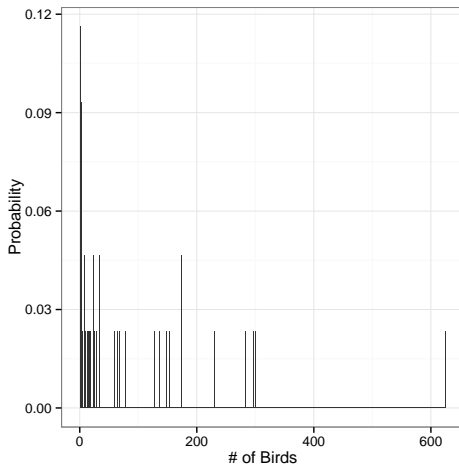
# Frequencies of Individual Observations



# Probabilities of Individual Measurements

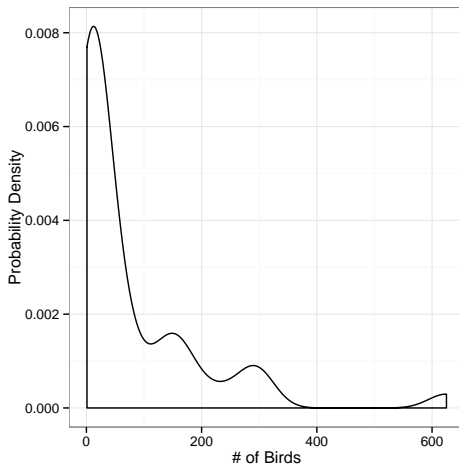


# Probabilities of Individual Measurements

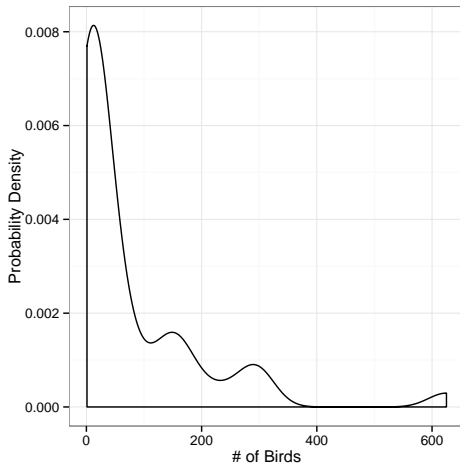


Many probabilities small, and what about the gaps?

# Continuous Probability Distributions

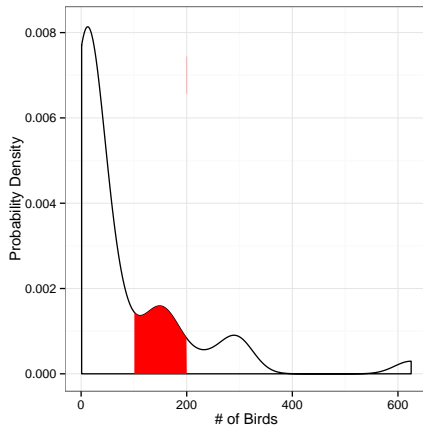


# Continuous Probability Distributions

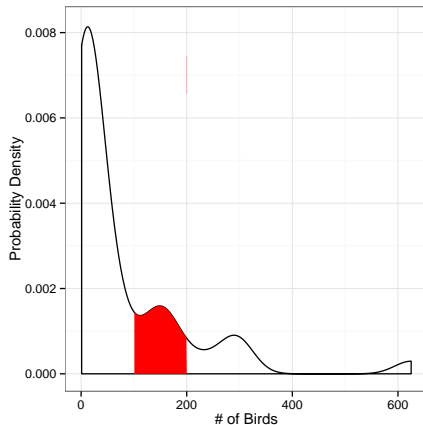


Any individual observation has a *probability density*.

# Probability as Integral Under the Curve

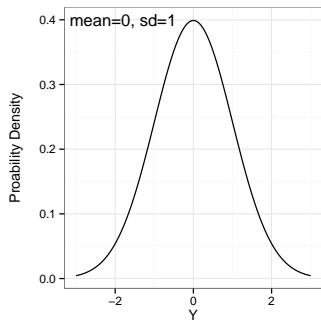


# Probability as Integral Under the Curve



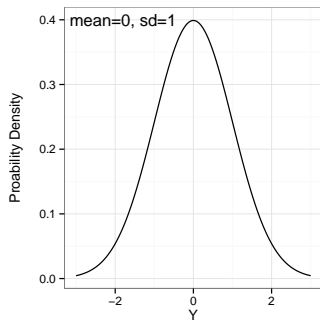
We obtain probabilities of observations between a range of values by integrating the distribution over selected values.

# The Normal Distribution



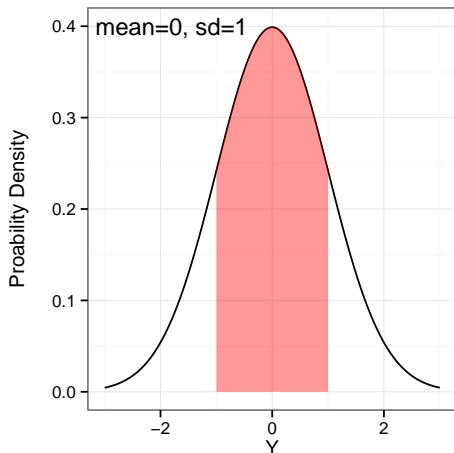


# The Normal Distribution

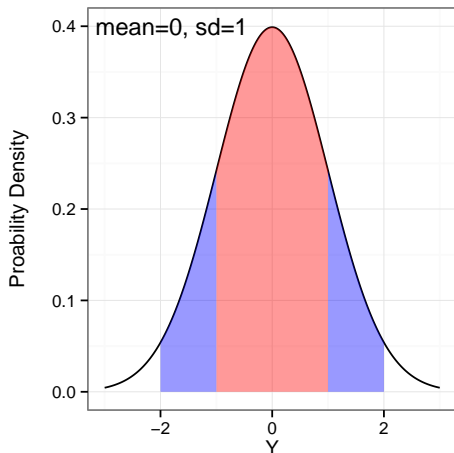


- ▶ Defined by its mean and standard deviation.
- ▶  $Y \sim N(\mu, \sigma)$
- ▶ Single mode
- ▶ Symmetric

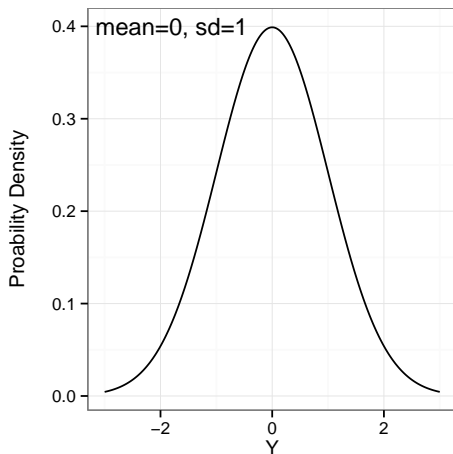
## 67% of Values within 1 SD



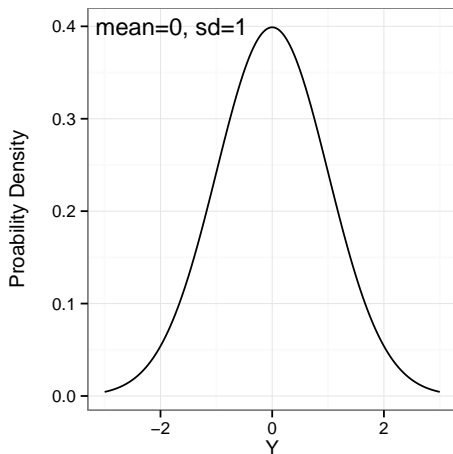
## 95% of Values within 2 (1.96) SD



# How to Get A Probability Density in R

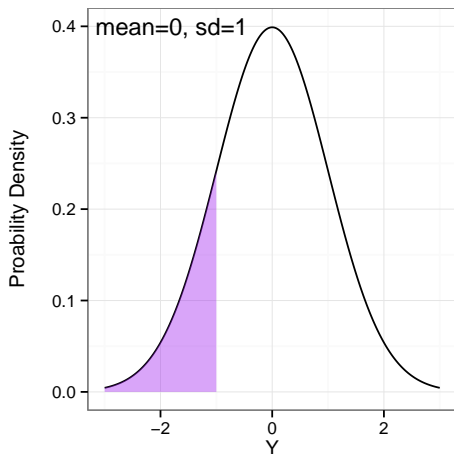


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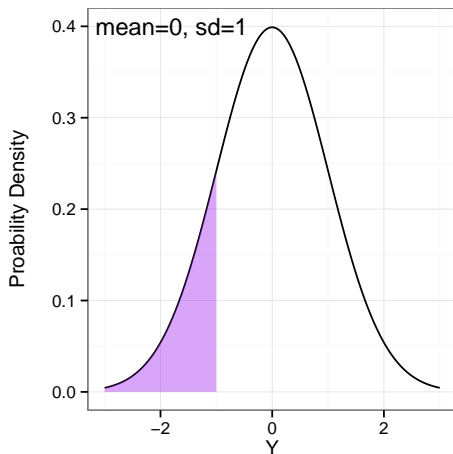


```
dnorm(Y, mean = 0, sd = 1)
```

# The Probability of a Value or More Extreme Value

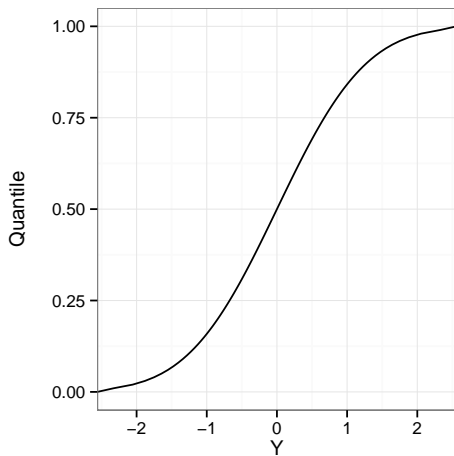


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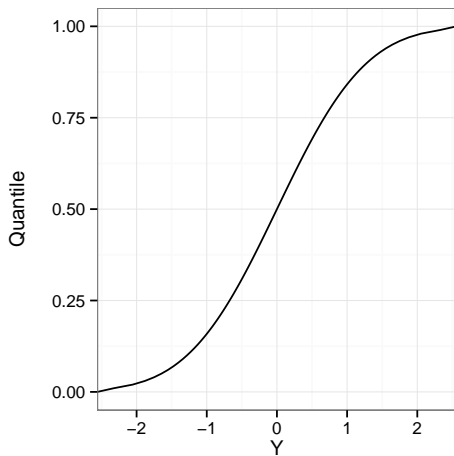
```
pnorm(Y, mean = 0, sd = 1)
```

# The Cumulative Distribution/Quantile Function





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```
qnorm(p, mean = 0, sd = 1)
```

# The Cumulative Distribution/Quantile Function

`pnorm` and `qnorm` are the inverse of one another

```
pnorm(-1)

## [1] 0.1587

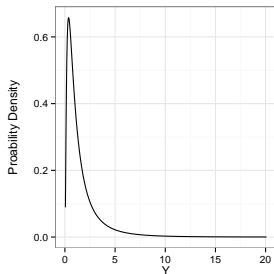
qnorm(pnorm(-1))

## [1] -1

qnorm(0.025)

## [1] -1.96
```

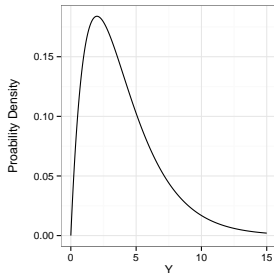
# The Lognormal Distribution



- ▶ An exponentiated normal
- ▶ Defined by the mean and standard deviation of its log.
- ▶  $Y \sim \text{LN}(\mu_{\log}, \sigma_{\log})$
- ▶ Generated by multiplicative processes

```
dlnorm(Y, meanlog = 0, sdlog = 1)
```

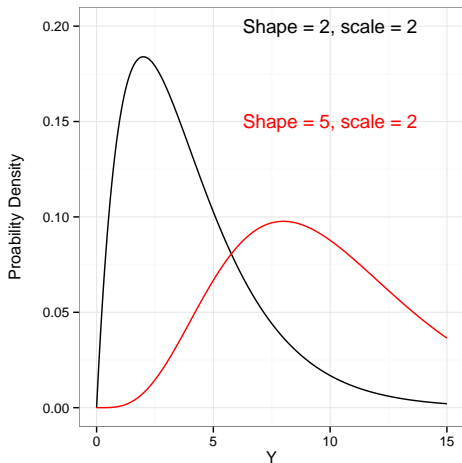
# The Gamma Distribution



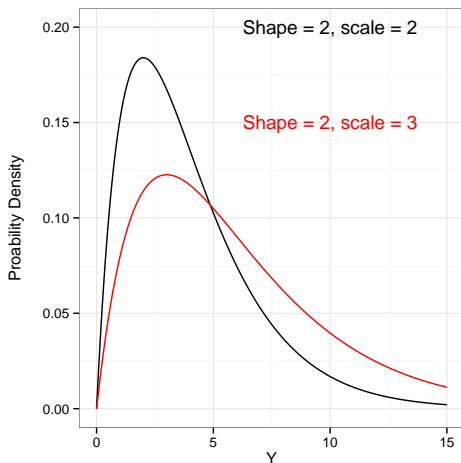
- ▶ Defined by number of events(shape) average time to an event (scale)
- ▶ Can also use rate ( $1/\text{scale}$ )
- ▶  $Y \sim G(\text{shape}, \text{scale})$
- ▶ Think of time spent waiting for a bus to arrive

```
dgamma(Y, shape = 2, scale = 2)
```

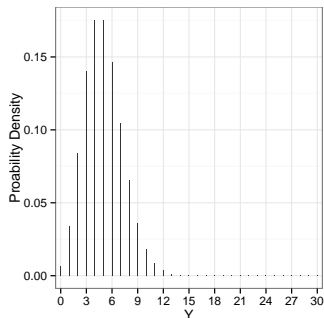
## Waiting for more events



## Longer average time per event



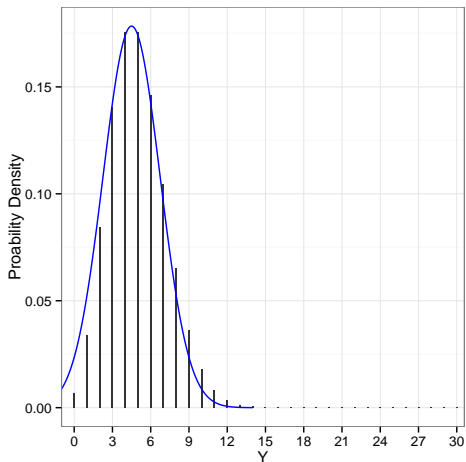
# The Poisson Distribution



- ▶ Defined by  $\lambda$  - the mean and variance
- ▶  $Y \sim P(\text{lambda})$

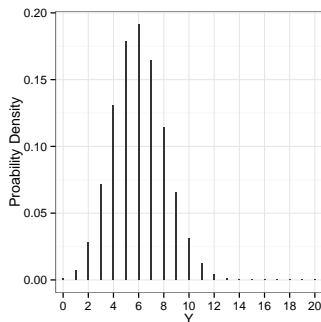
```
dpois(Y, lambda = 5)
```

## When Lambda is Large, Approximately Normal





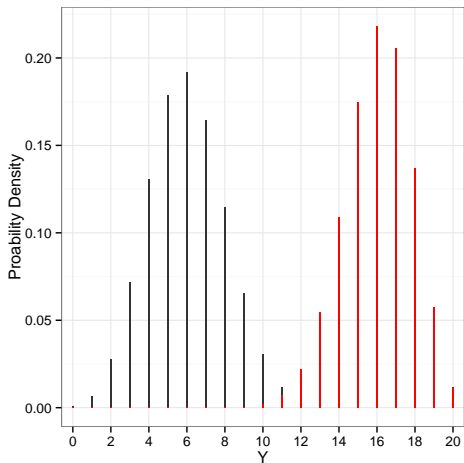
# The Binomial Distribution



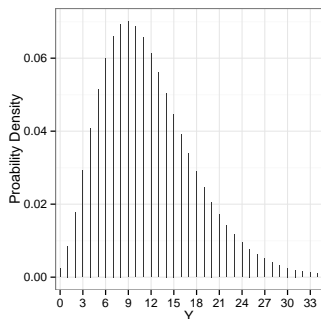
- ▶ Results from multiple coin flips
- ▶ Defined by size (# of flips) and prob (probability of heads)
- ▶  $Y \sim B(\text{size}, \text{prob})$
- ▶ bounded by 0 and size

```
dpois(Y, size, prob)
```

# Increasing Probability Shifts Distribution



# The Negative Binomial Distribution



- ▶ Distribution of number of failures before  $n$  number of successes in  $k$  trials
- ▶ Or mean  $\#$  of counts,  $\mu$ , with an overdispersion parameter, size
- ▶  $Y \sim \text{NB}(\mu, \text{size})$

```
dnbin(Y, mu, size)
```

## Exercise

- ▶ Explore the distributions we have discussed
- ▶ Examine how changing parameters shifts the output of probability function
- ▶ Compare curves generated using density functions (e.g., `dnorm`) and large number of random draws (e.g. from `rnorm`)
- ▶ Overlay these in plots if you can (hist, lines, etc.)
- ▶ Challenge: graphically show integration under the different types of distribution curves (?`polygon` or ?`geom_ribbon`)