# Data Types, Probability Distributions, and ggplot2

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### **Data Types**

#### **Data Types**

- In general, data can take on one of three different forms
  - 1. Metric
  - 2. Ordinal
  - 3. Nominal

 Need to consider this when developing appropriate Bayesian models

#### Data Types Metric

- When values are actual measurements that can take on a range of values (e.g., temperature, weight, counts, frequencies)
- Indicate 2 things
  - The order of values (which ones are largest and which are smallest)
  - The scale of difference between values

### Data Types Metric

#### Both metric

X	age	fat	sex
1	24	15.5	male
2	37	20.9	male
3	41	18.6	male
4	60	28	male
5	31	34.7	female
6	39	30.2	female

#### Data Types Metric

 Can tell order (which are larger and which are smaller)

 Can tell by how much values differ

#### Both metric

X	age	fat	sex
1	24	15.5	male
2	37	20.9	male
3	41	18.6	male
4	60	28	male
5	31	34.7	female
6	39	30.2	female

# Data Types Ordinal

• Provide information on order, but not scale

### Data Types Ordinal

- Provide information on order, but not scale
  - Places in a race

#### Ordinal

Person	Place	
Emily	First	
Victoria	Second	
Tasha	Third	
Ben	Fourth	

 Can tell order, but not scale (e.g, time difference between individuals)

### Data Types Nominal

- Categorical
  - Sex
  - Political party
  - Others
- Can tell what category something is in, but not order or scale

## Data Types Nominal

#### Nominal

x	age	fat	sex
1	24	15.5	male
2	37	20.9	male
3	41	18.6	male
4	60	28	male
5	31	34.7	female
6	39	30.2	female

• Can tell category, but not order or scale

# Probability Distributions In R and STAN

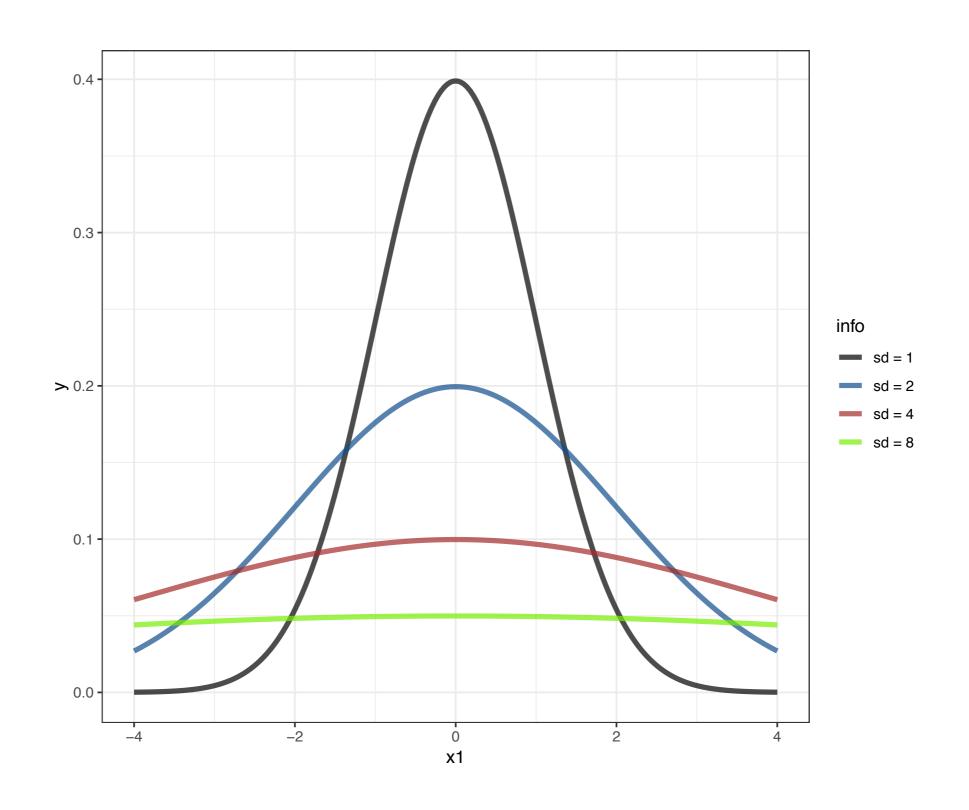
#### **Probability Distributions**

 To conduct Bayesian analyses, you have to be fairly comfortable with at least a few probability distributions

 Different ones suitable for different data, with different characteristics

# Continuous Distributions

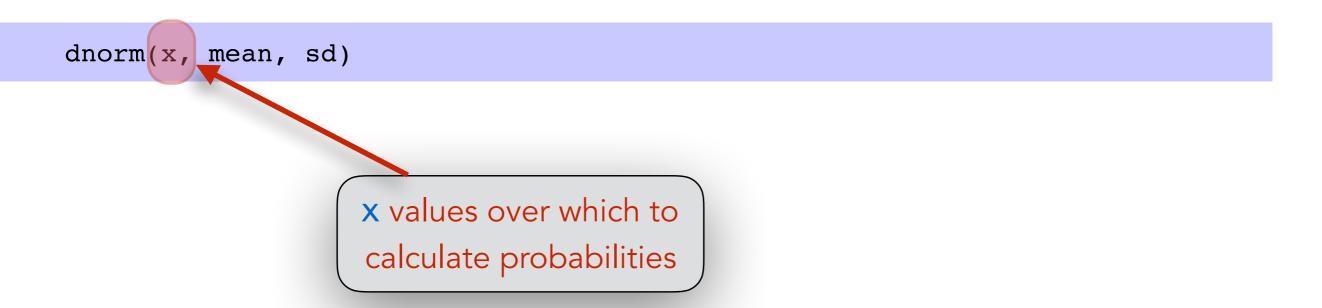
- Symmetric around the mean (mean, median, and mode are all equal)
- Range is (-∞ to ∞)
- Has two parameters:
  - Mean: determines position of peak along x-axis
  - Standard Deviation: determines how wide the peak is around the mean



• In R, use the dnorm function

dnorm(x, mean, sd)

• In R, use the dnorm function



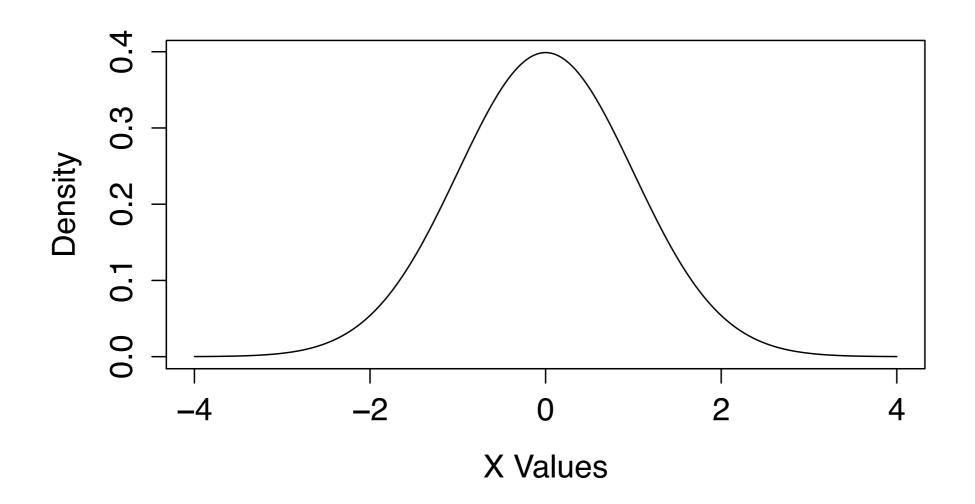
• In R, use the dnorm function

```
dnorm(x, mean, sd)
```

```
x = seq(from = -4, to = 4, length = 200)

y = dnorm(x, mean = 0, sd = 1)

plot(x, y, type = "l", lwd = 2, xlab = "X Values", ylab = "Density")
```



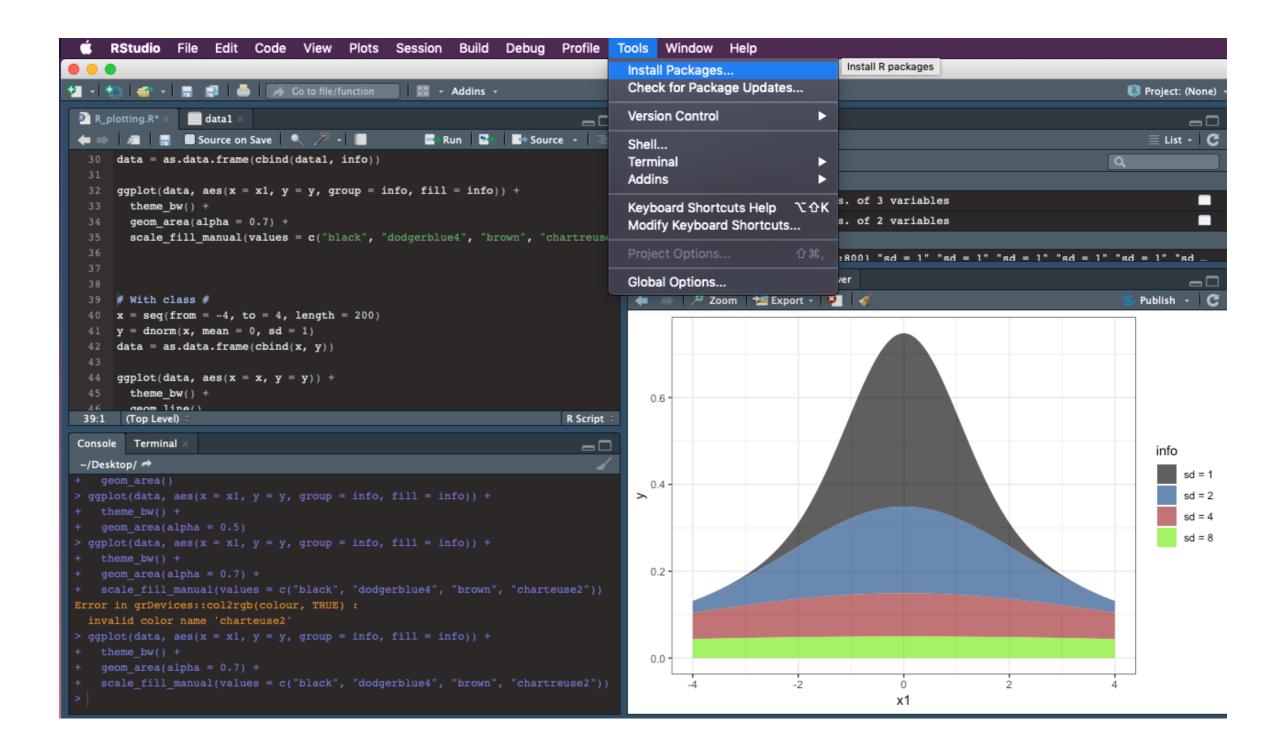
- In Stan, function is called normal, requires:
  - mu  $(\mu)$  = mean
  - sigma  $(\sigma)$  = standard deviation

normal(mu, sigma)

 Because of the central limit theorem, normal distribution is appropriate for many cases when our goal is to estimate the mean response to certain conditions

https://vimeo.com/75089338

#### Install



- "grammar of graphics"
- Build plots layer by layer, customizing as you go
  - A bit tricky to learn at first, but ultimately easier, with nicer results
- Data must be in a data frame
  - Based on the philosophy of "tidy data"

#### Tidyverse

Packages Articles Learn Help

Contribute



#### R packages for data science

The tidyverse is an opinionated **collection of R packages** designed for data science. All packages share an underlying design philosophy, grammar, and data structures.

Install the complete tidyverse with:

install.packages("tidyverse")

Load library

library(ggplot2)

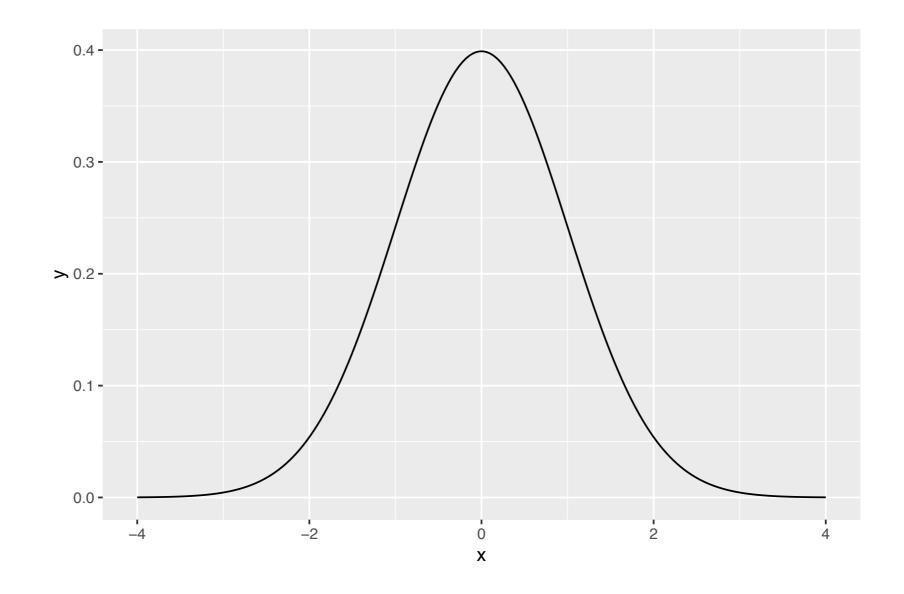
Load library

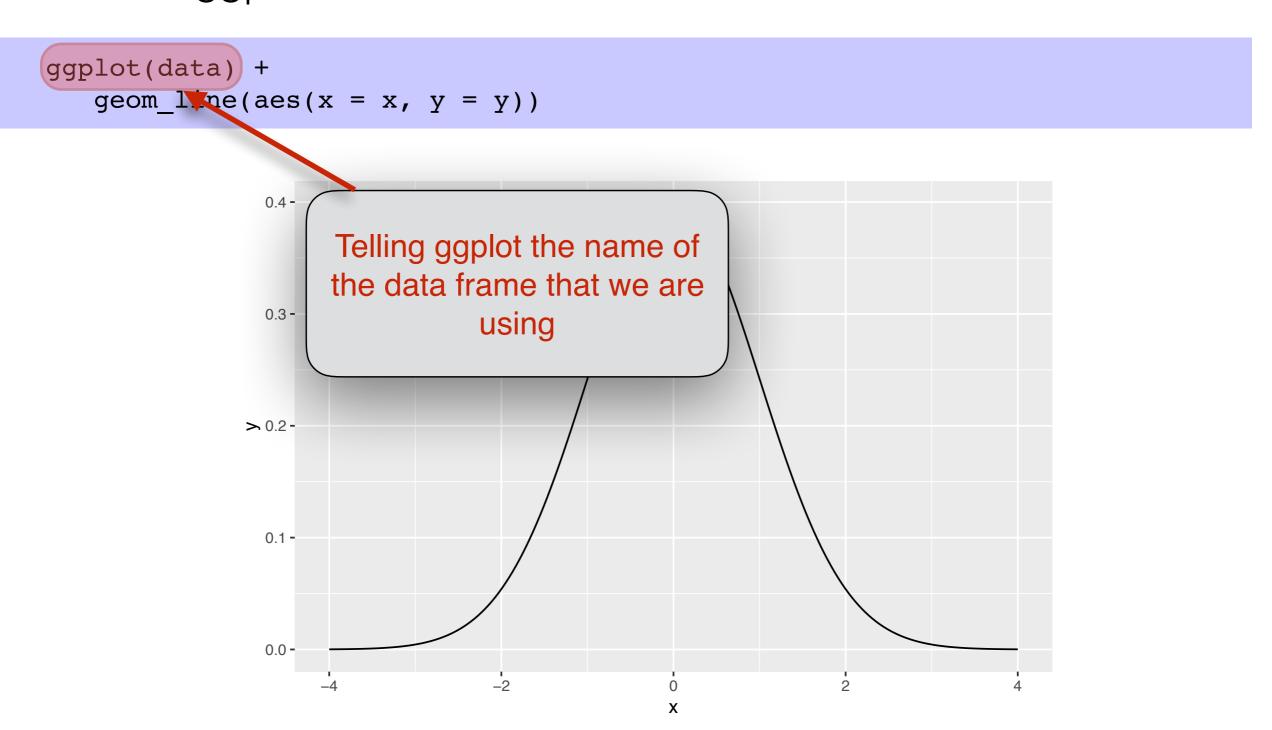
```
library(ggplot2)
```

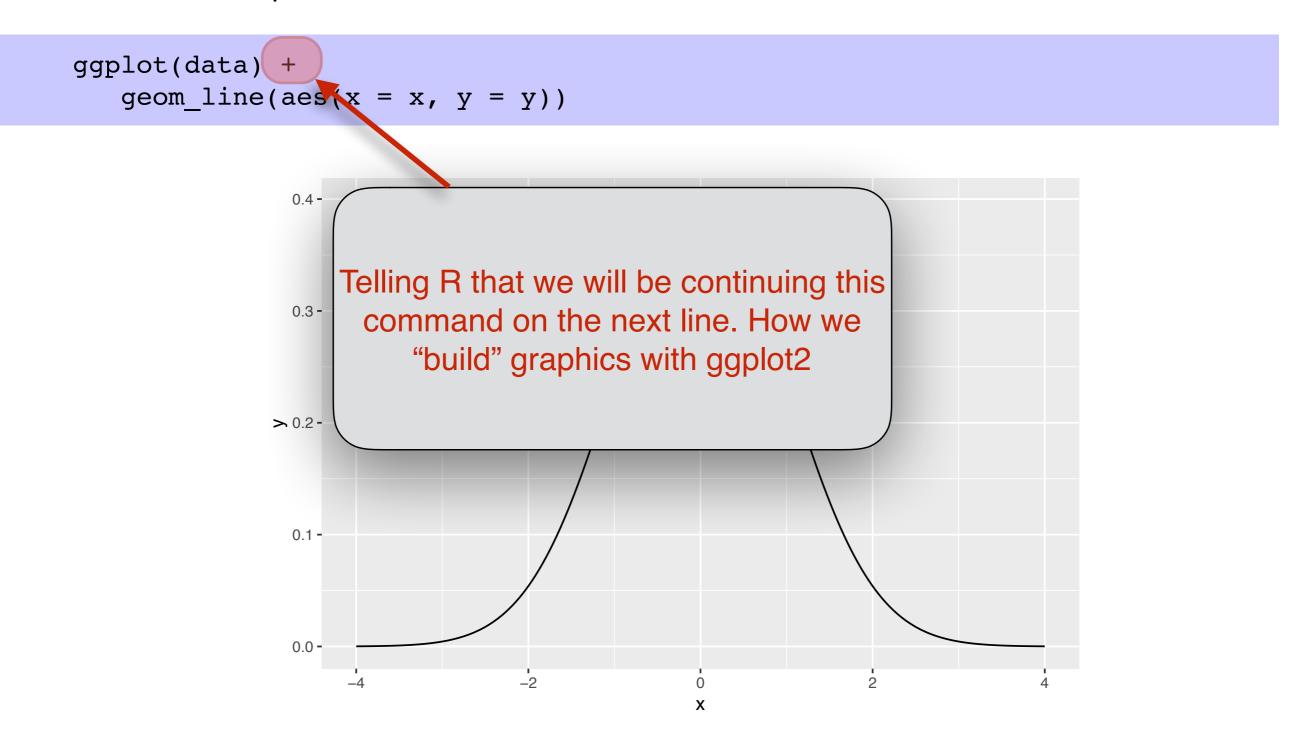
 $\bullet$  Combine x and y data into a data frame

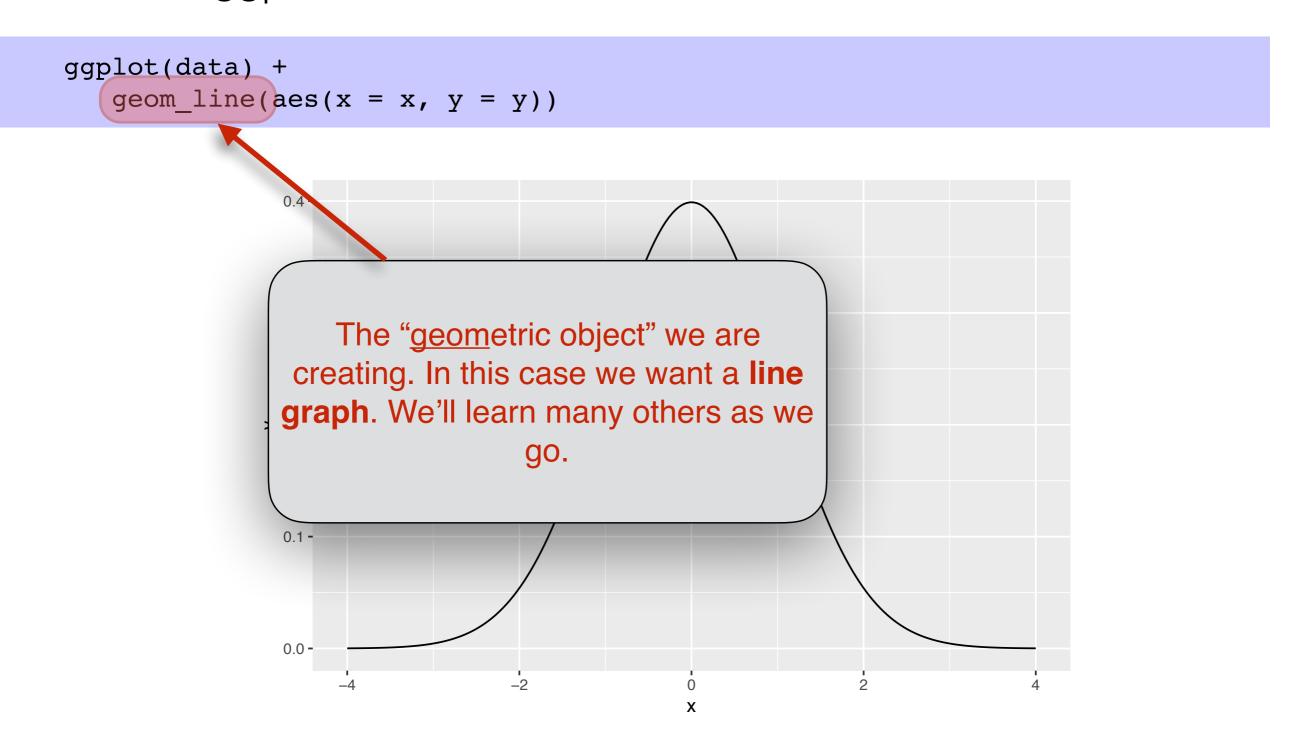
```
data = as.data.frame(cbind(x, y))
```

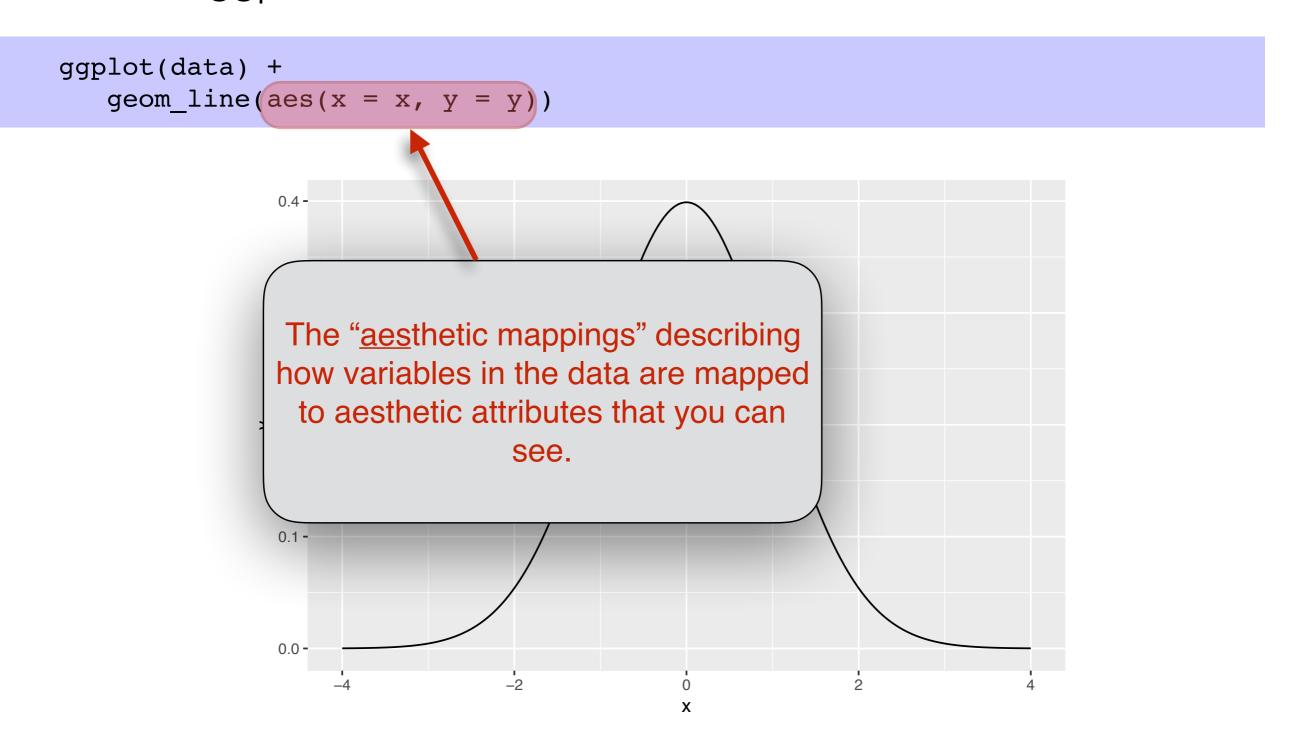
```
ggplot(data) +
  geom_line(aes(x = x, y = y))
```



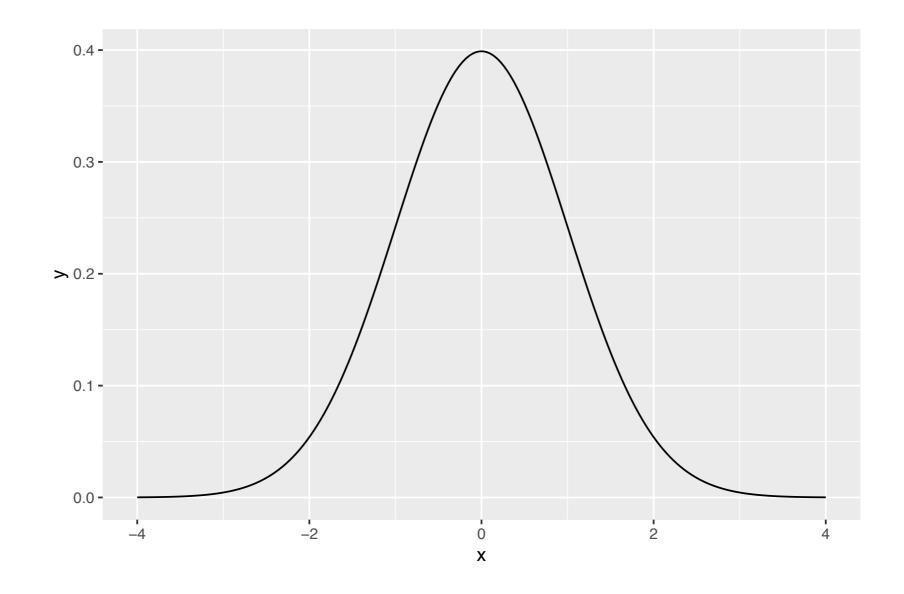






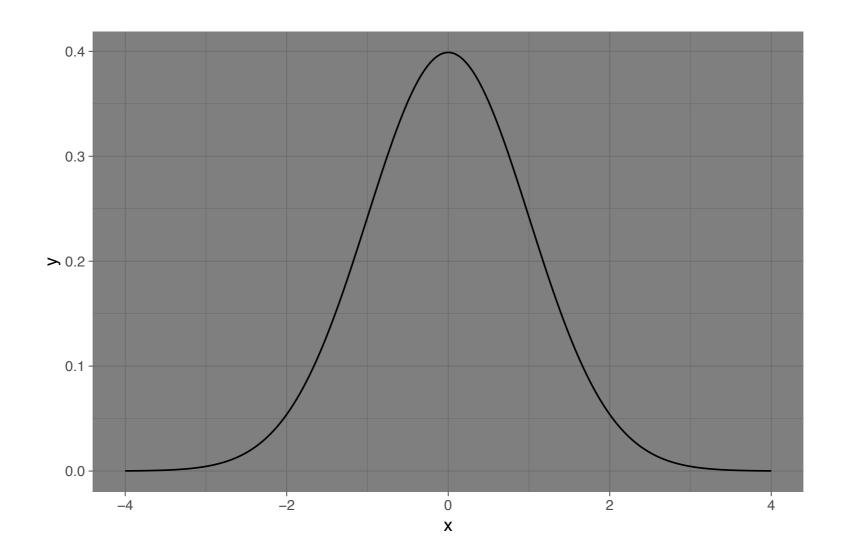


```
ggplot(data) +
  geom_line(aes(x = x, y = y))
```



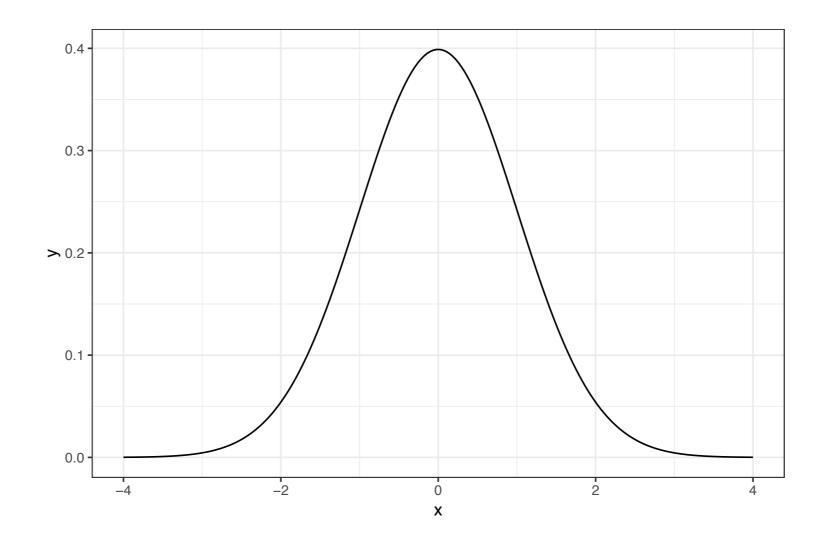
 Are many different themes in ggplot2 that influence many aspects of how a plot looks

```
ggplot(data) +
  theme_dark() +
  geom_line(aes(x = x, y = y))
```



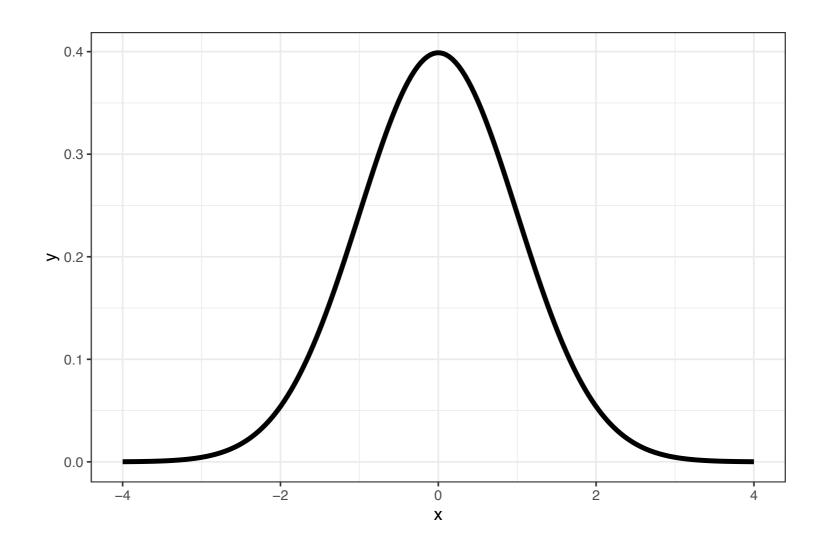
 Are many different themes in ggplot2 that influence many aspects of how a plot looks

```
ggplot(data) +
   theme_bw() +
   geom_line(aes(x = x, y = y))
```



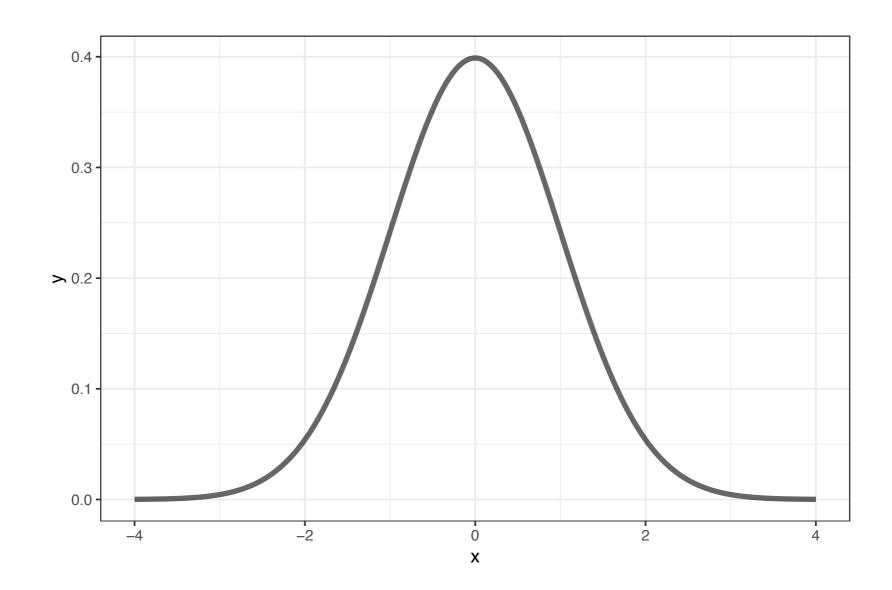
• Let's make the line a bit thicker

```
ggplot(data) +
   theme_bw() +
   geom_line(aes(x = x, y = y), size = 1.5)
```

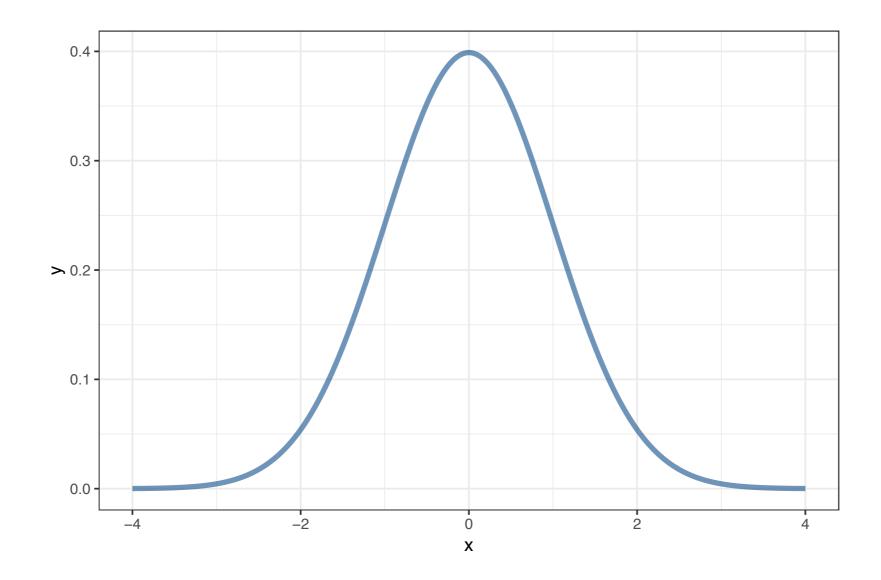


• Let's make it slightly opaque

```
ggplot(data) +
   theme_bw() +
   geom_line(aes(x = x, y = y), size = 1.5, alpha = 0.6)
```

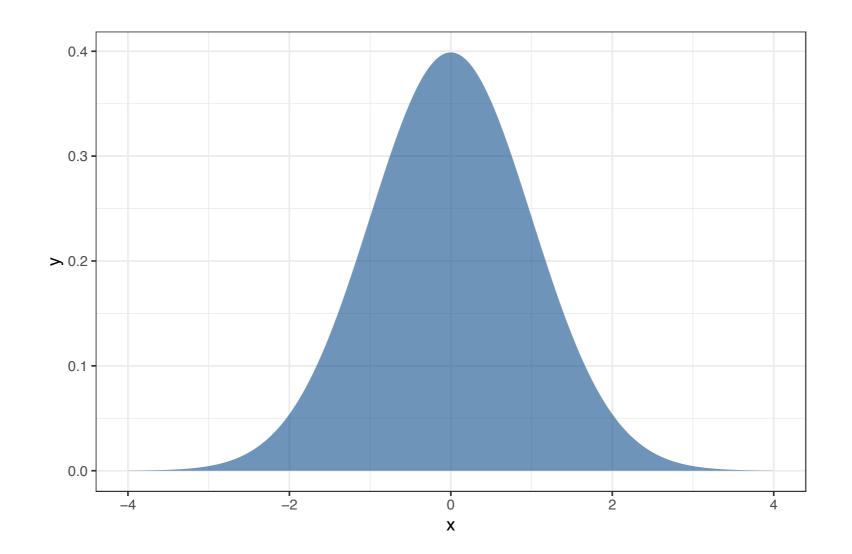


Lastly, let's customize the colour



- Let's plot it as an "area" plot
  - ymax = y values, ymin = 0

```
ggplot(data) +
   theme_bw() +
   geom_area(aes(x = x, y = y), fill = "dodgerblue4", alpha = 0.6)
```

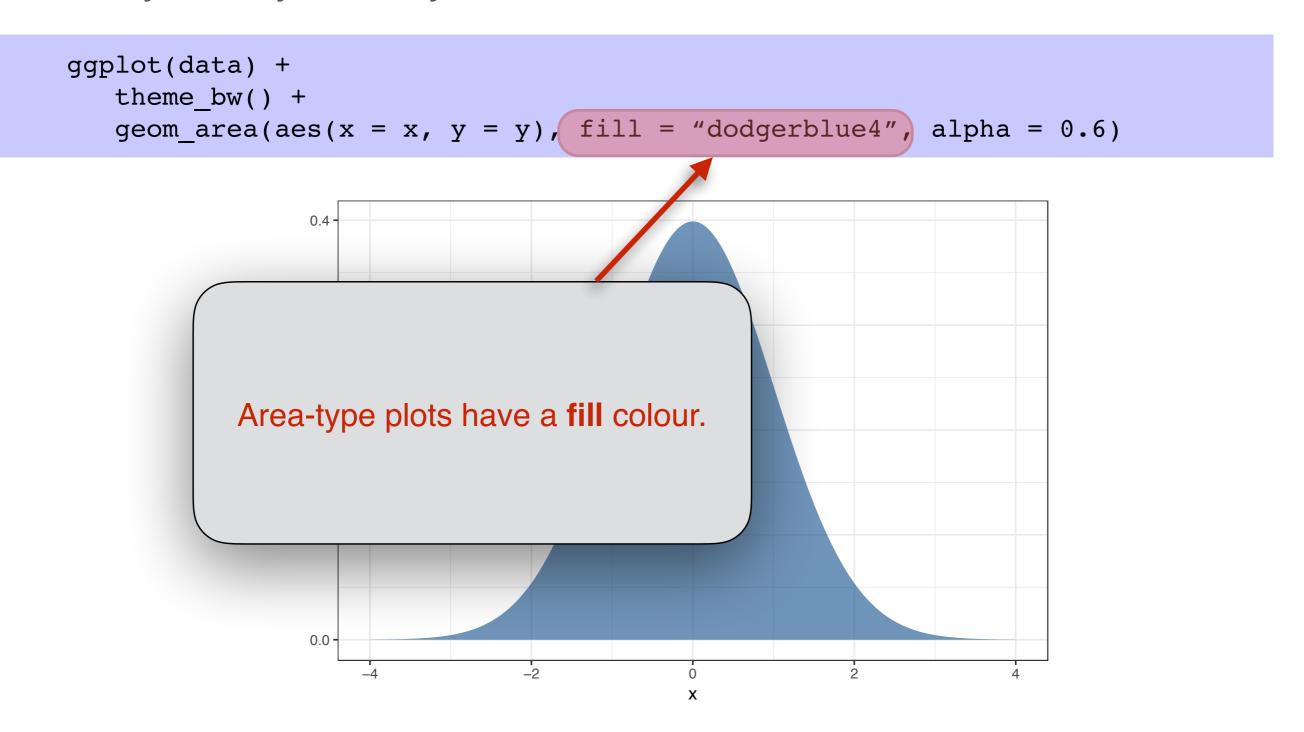


- Let's plot it as an "area" plot
  - ymax = y values, ymin = 0

```
ggplot(data) +
   theme bw() +
   geom_area(aes(x = x, y = y), fill = "dodgerblue4", alpha = 0.6)
                0.4 -
         Now making an area plot rather than
                      a line plot.
                0.0
```

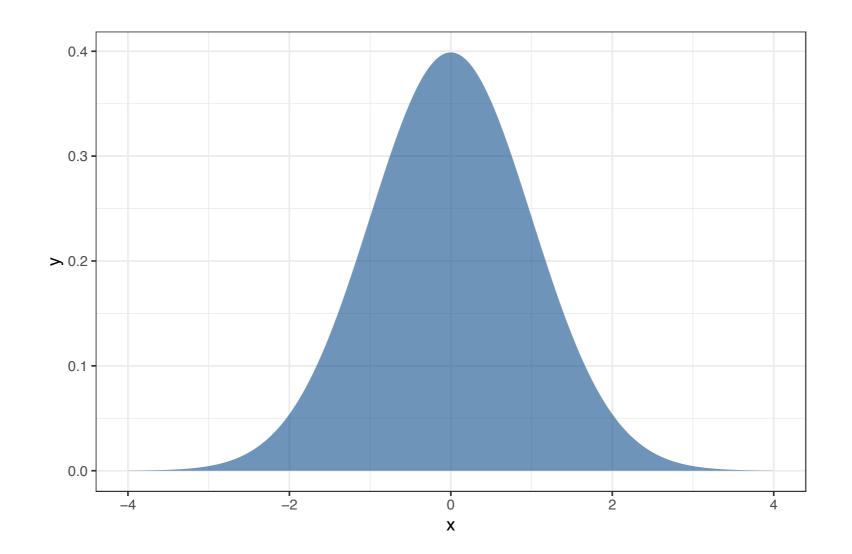
Χ

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  - ymax = y values, ymin = 0

```
ggplot(data) +
   theme_bw() +
   geom_area(aes(x = x, y = y), fill = "dodgerblue4", alpha = 0.6)
```



Google "Colors in R"

- A flat distribution (all values have the same probability)
- Range is -∞ to ∞ (you specify range)
- Has two parameters:
  - min: minimum value to consider
  - max: maximum value to consider

• In R, use the dunif function

```
dunif(x, min, max)
```

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```
dunif(x, min, max)
```

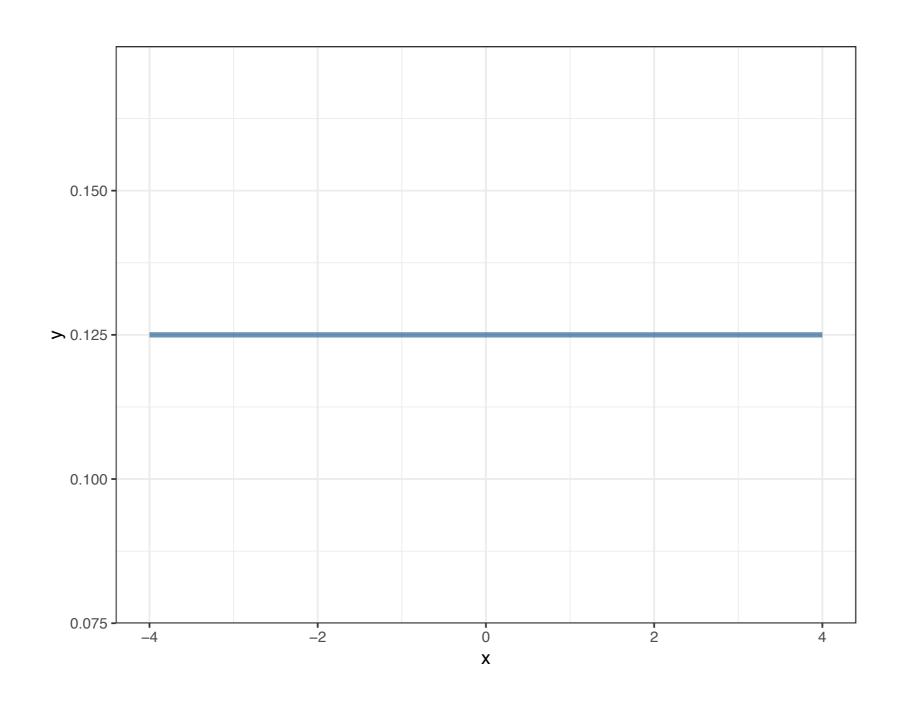
```
x = seq(from = -4, to = 4, length = 200)

y = dunif(x, min = -4, max = 4)
```

Organize and plot the data

```
data = as.data.frame(cbind(x, y))

ggplot(data) +
    theme_bw() +
    geom_line(aes(x = x, y = y), colour = "dodgerblue4", size = 1.5, alpha
    = 0.6)
```

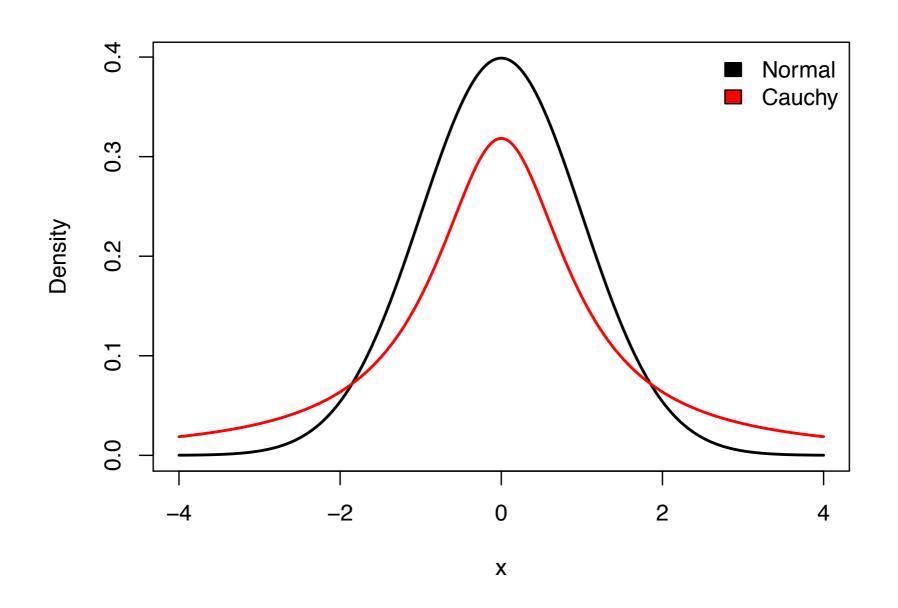


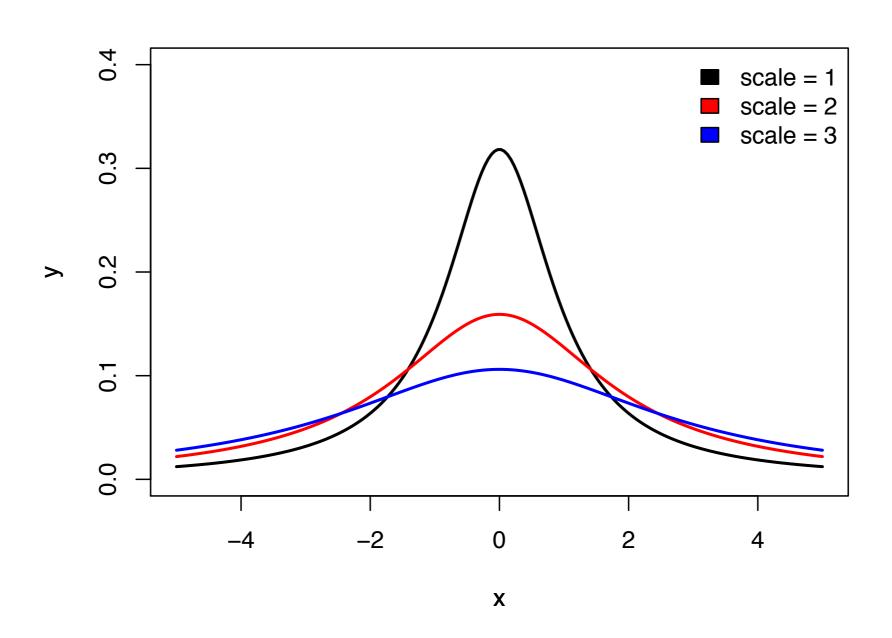
- In Stan, function is called uniform, requires:
  - alpha = minimum value
  - beta = maximum value

uniform(alpha, beta)

## Probability Distributions Cauchy distribution (pronounced *kō shee*)

- Like the normal distribution, but with heavier tails
- Symmetric around the mean (mean, median, and mode are all equal)
- Range is (-∞ to ∞)
- Has two parameters:
  - location: determines position of peak along x-axis
  - scale: determines how wide the peak is around the location





• In R, use the dcauchy function

```
dcauchy(x, location, scale)
```

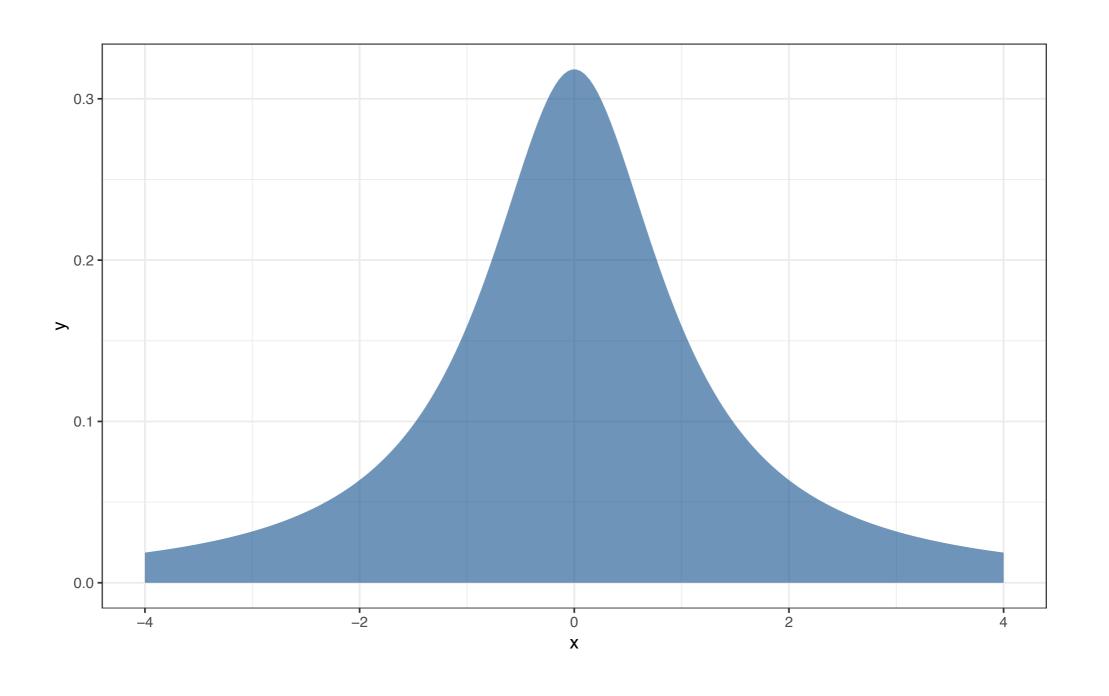
```
x = seq(from = -4, to = 4, length = 200)

y = dcauchy(x, location = 0, scale = 1)
```

Organize and plot the data (exactly the same as before)

```
data = as.data.frame(cbind(x, y))

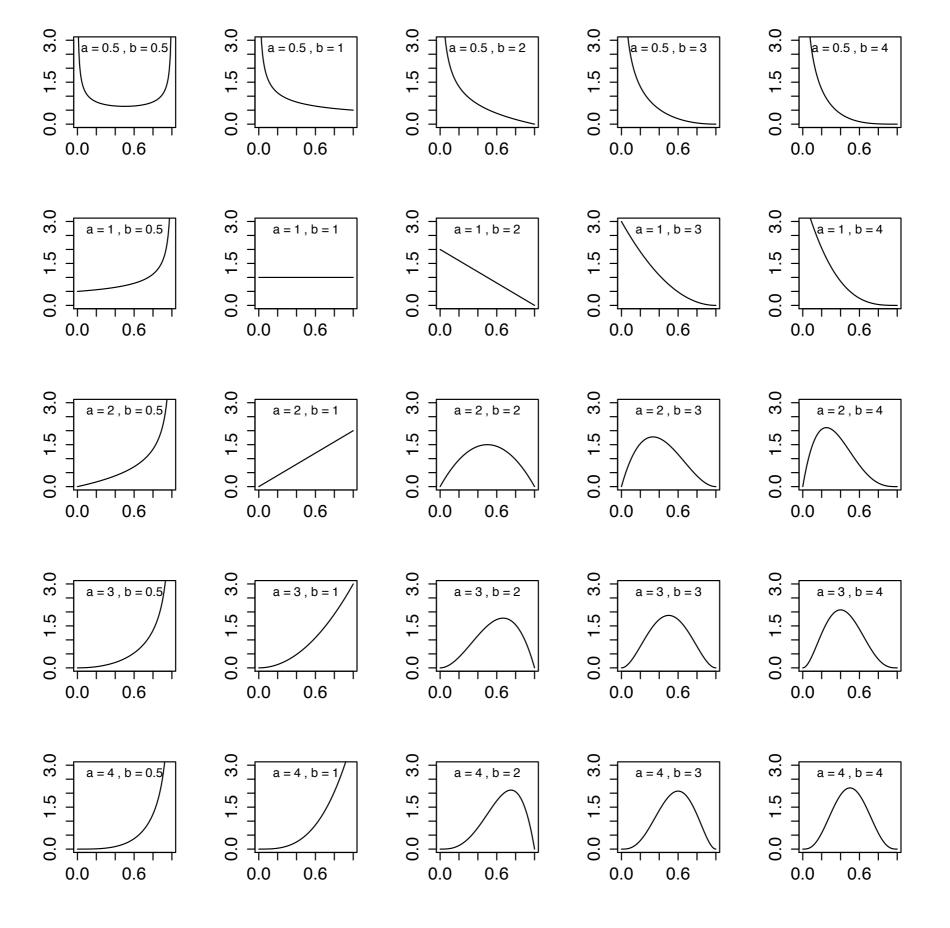
ggplot(data) +
   theme_bw() +
   geom_area(aes(x = x, y = y), fill = "dodgerblue4", alpha = 0.6)
```



- In Stan, function is called cauchy, requires:
  - location
  - scale

cauchy(location, scale)

- Can take on almost any shape, determined by two positive shape parameters
- Range is [0 to 1]
- Has two parameters:
  - First shape parameter (a): determines shape of curve
  - Second shape parameter (b): determines shape of curve



• In R, use the dbeta function

dbeta(x, shape1, shape2)

• In R, use the dbeta function

```
dbeta(x, shape1, shape2)
```

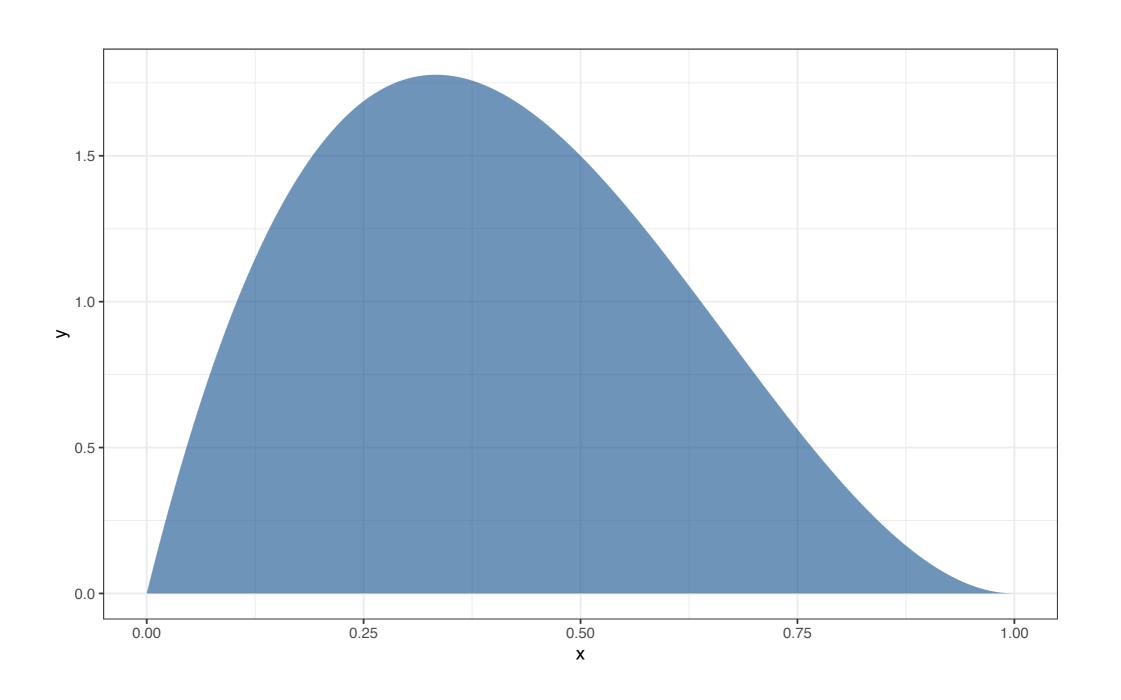
```
x = seq(from = 0, to = 1, length = 200)

y = dbeta(x, shape1 = 2, shape2 = 3)
```

Organize and plot the data

```
data = as.data.frame(cbind(x, y))

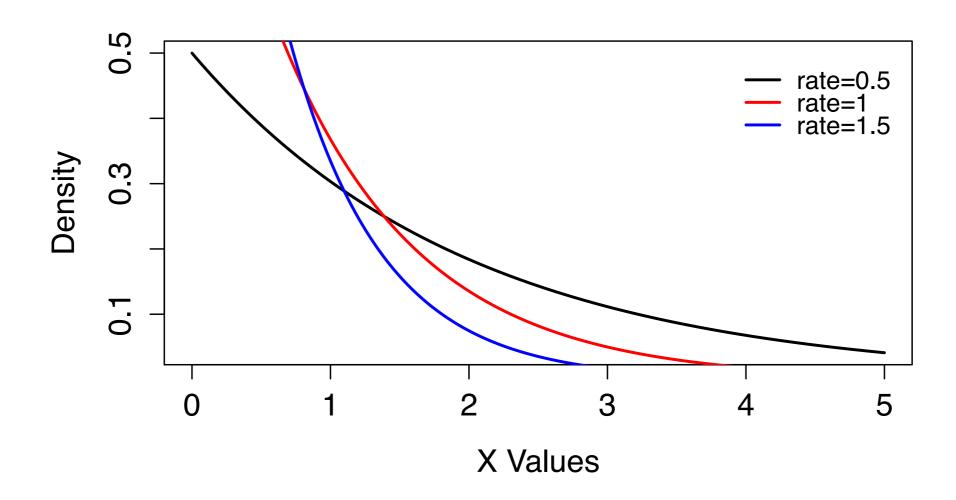
ggplot(data) +
   theme_bw() +
   geom_area(aes(x = x, y = y), fill = "dodgerblue4", alpha = 0.6)
```



- In Stan, function is called beta, requires:
  - alpha = first shape parameter
  - beta = second shape parameter

beta(alpha, beta)

- An L-shaped distribution
- Range is 0 to ∞
- Has one parameter:
  - rate: determines shape of the curve



• In R, use the dexp function

dexp(x, rate)

• In R, use the dexp function

```
dexp(x, rate)
```

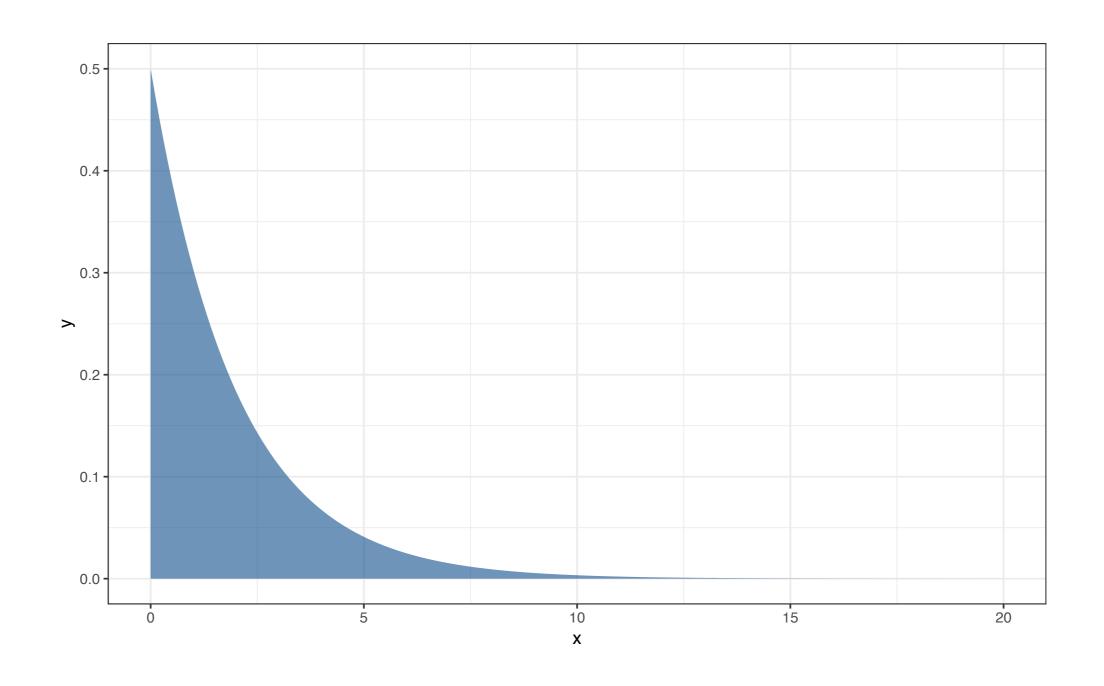
```
x = seq(from = 0, to = 20, length = 200)

y = dexp(x, rate = 0.5)
```

Organize and plot the data

```
data = as.data.frame(cbind(x, y))

ggplot(data) +
   theme_bw() +
   geom_area(aes(x = x, y = y), fill = "dodgerblue4", alpha = 0.6)
```



- In Stan, function is called exponential, requires:
  - beta = rate value

exponential(beta)

# Discrete Distributions

- Used when you have multiple trials, and only two results possible:
   0 or 1 (sometimes "failure" or "success")
- Range is integers from [0 to ∞)
  - # of "successes"
- Has two parameters:
  - Size: The number of trials
  - Prob: the probability of "success"



• In R, use the dbinom function

dbinom(x, size, prob)

• In R, use the dbinom function

```
dbinom(x, size, prob)
```

```
x = 1:50
y = dbinom(x, size = 50, prob = 0.5)
```

Organize and plot the data

```
data = as.data.frame(cbind(x, y))

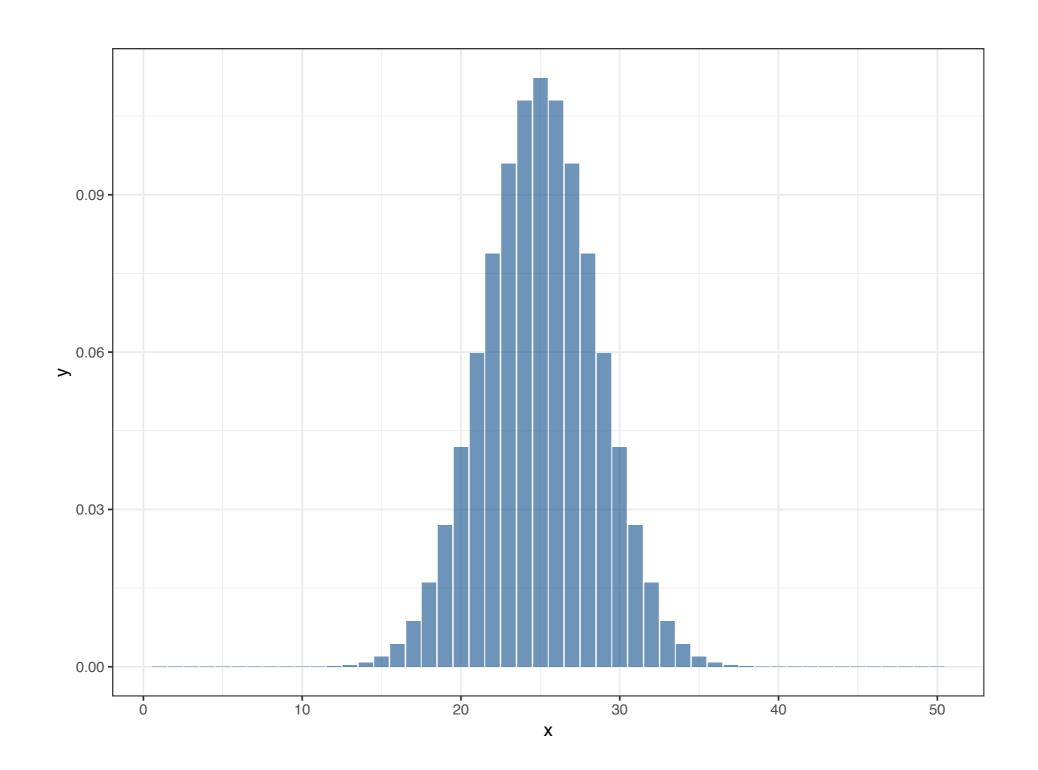
ggplot(data) +
    theme_bw() +
    geom_col(aes(x = x, y = y), fill = "dodgerblue4", alpha = 0.6)
```

Organize and plot the data

```
data = as.data.frame(cbind(x, y))

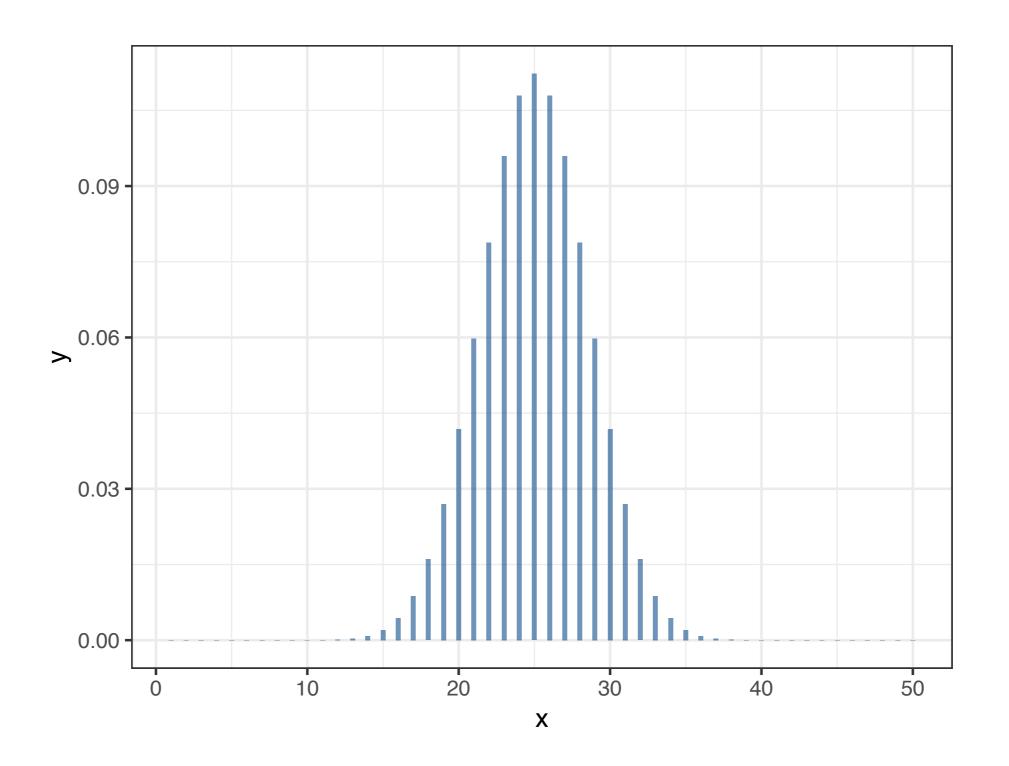
ggplot(data) +
    theme_bw() +
    geom_col(aes(x = x, y = y), fill = "dodgerblue4", alpha = 0.6)
```

Since we have discrete data (integers), area curve no longer appropriate.



Can change the width of the bars, if desired

```
ggplot(data) +
  theme_bw() +
  geom_col(aes(x = x, y = y), fill = "dodgerblue4", alpha = 0.6, width =
  0.3)
```



- In Stan, function is called binomial, requires:
  - N = number of trials
  - theta = probability of "success"

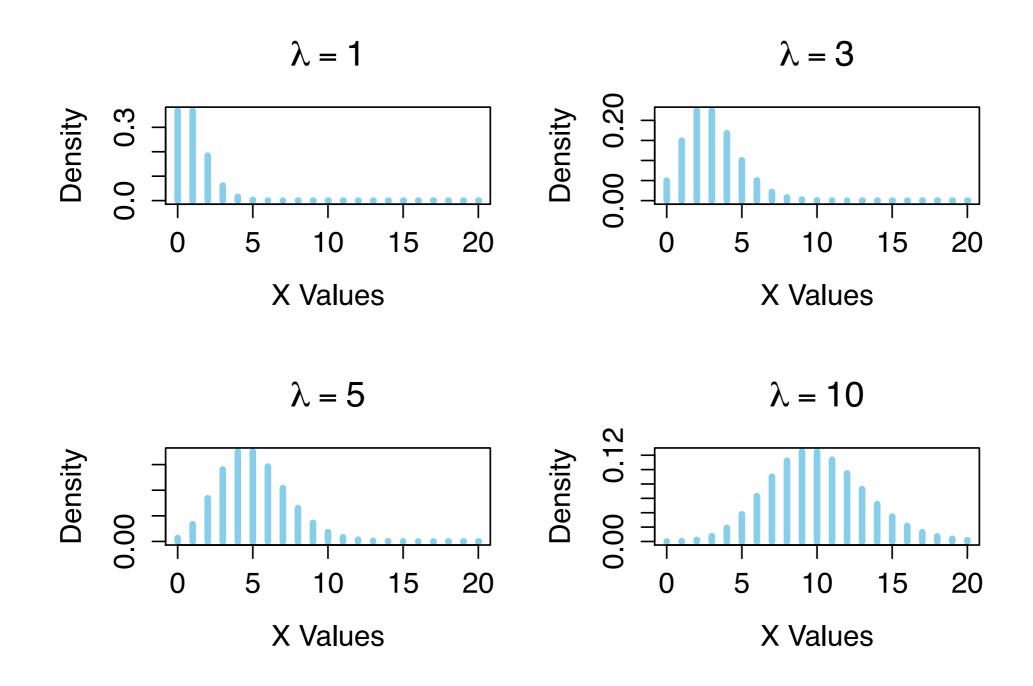
binomial(N, theta)

- One trial of a binomial situation
  - For a case-by-case analysis of a binomial process
- Many of our models could be parameterized as:
  - One group of results (size = N) from a binomial process; or
  - A series (length = N) of Bernoulli processes

- In Stan, function is called bernoulli, requires:
  - theta = probability of "success"

bernoulli(theta)

- Can range from L-shaped to a normal distribution
- Range is integers from 0 to ∞
  - Discrete values, not a continuous distribution
- Has one parameter:
  - lamdba: determines shape of the curve



• In R, use the dpois function

```
dpois(x, lambda)
```

• In R, use the dpois function

```
dpois(x, lambda)
```

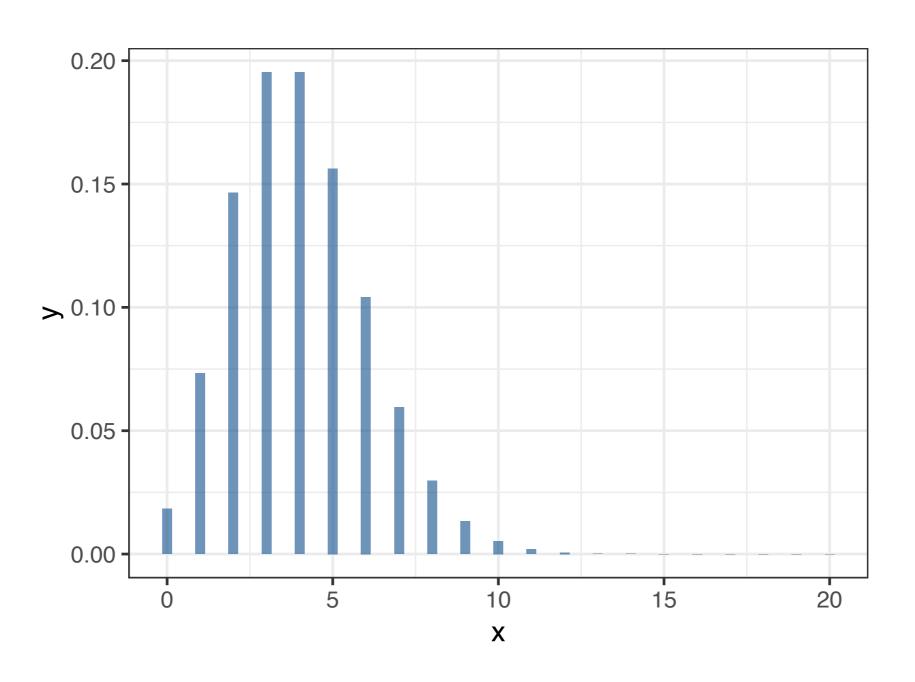
```
x = seq(from = 0, to = 20, by = 1)

y = dpois(x, lambda = 4)
```

Organize and plot the data

```
data = as.data.frame(cbind(x, y))

ggplot(data) +
   theme_bw() +
   geom_col(aes(x = x, y = y), fill = "dodgerblue4", alpha = 0.6, width = 0.3)
```



- In Stan, function is called poisson, requires:
  - lambda  $(\lambda)$  = shape value

poisson(lambda)

#### **Probability Distributions**

• See Stan manual for more, and for their proper usage

#### Questions?