

# Visible aesthetics

INTRODUCTION TO DATA VISUALIZATION WITH GGPLOT2

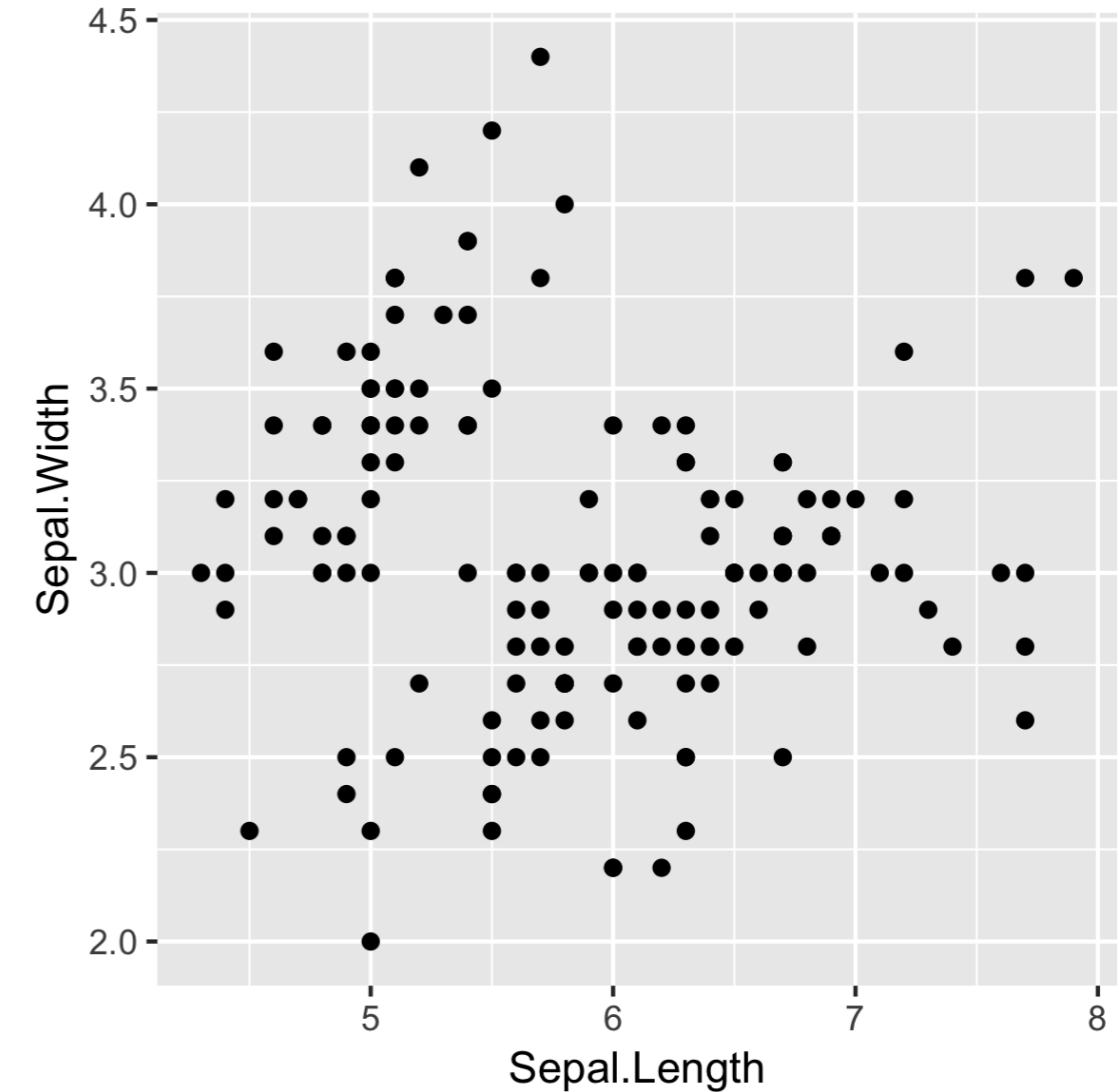


**Rick Scavetta**

Founder, Scavetta Academy

# Mapping onto the X and Y axes

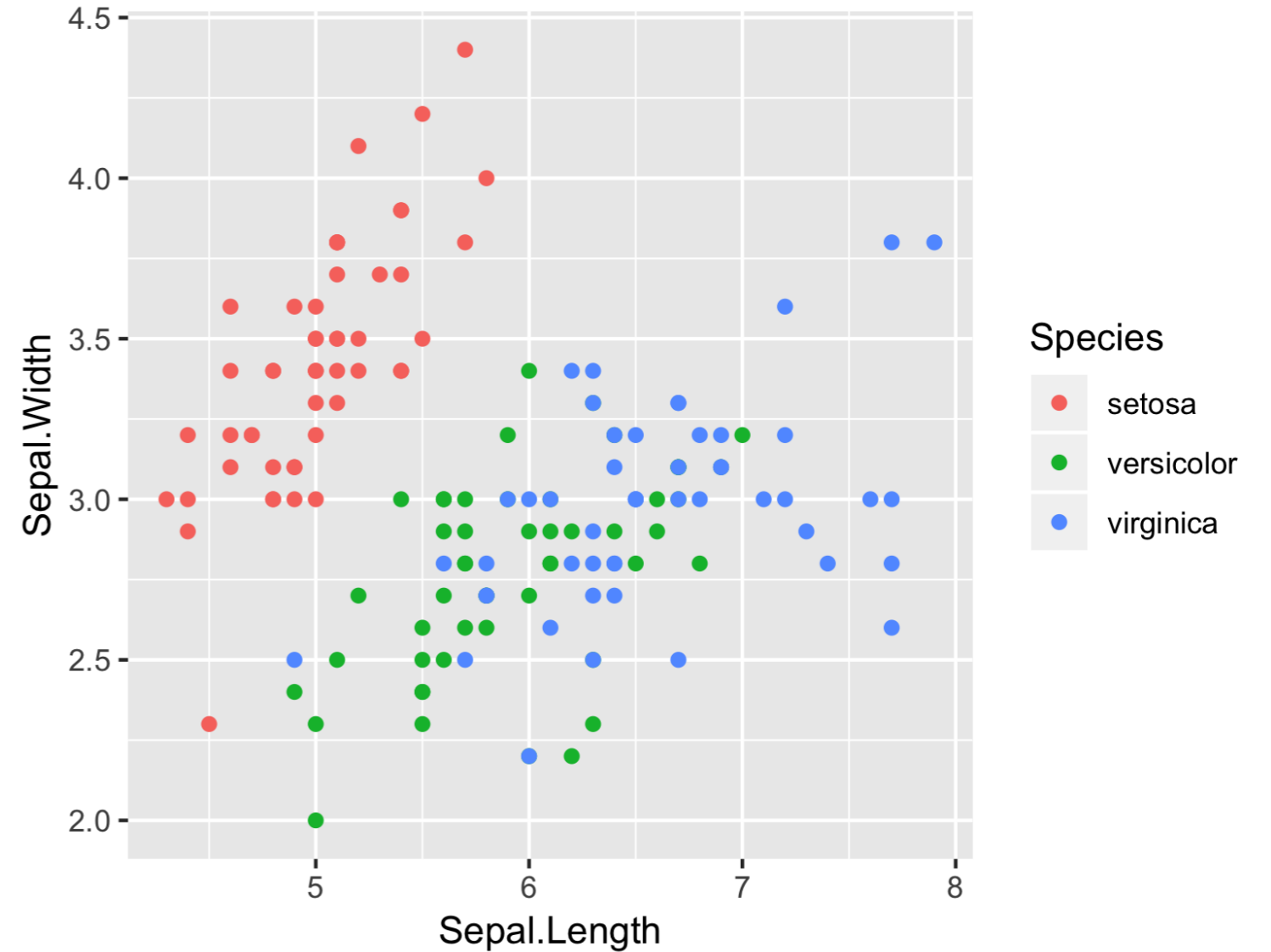
```
ggplot(iris, aes(x = Sepal.Length,  
                 y = Sepal.Width)) +  
  geom_point()
```



# Mapping onto color

```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point()
```

Type	Variable
Color	Species

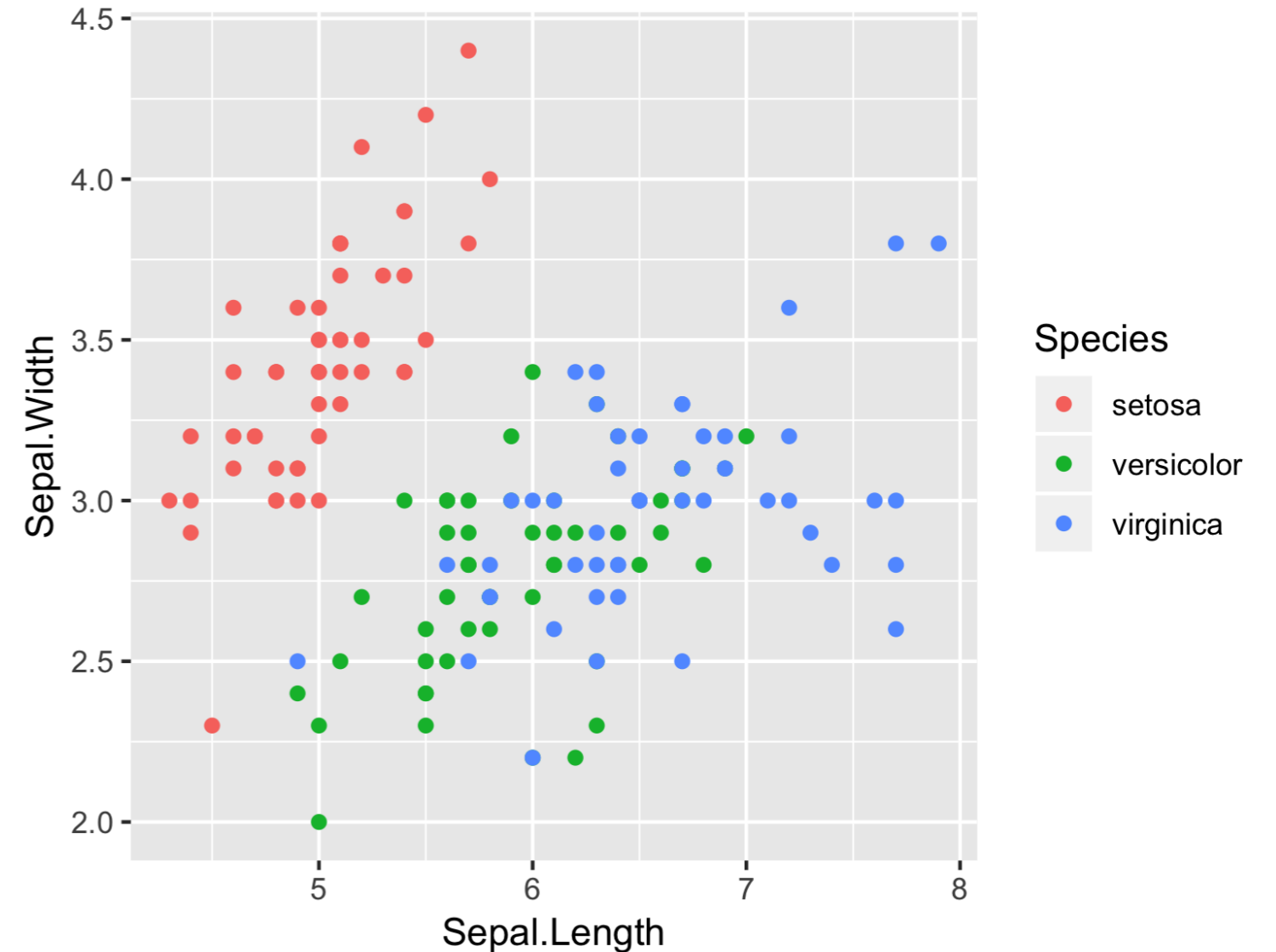


# Mapping onto the color aesthetic

```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point()
```

Type	Variable
Color	Species

**Species**, a dataframe column, is *mapped onto color*, a visible aesthetic.



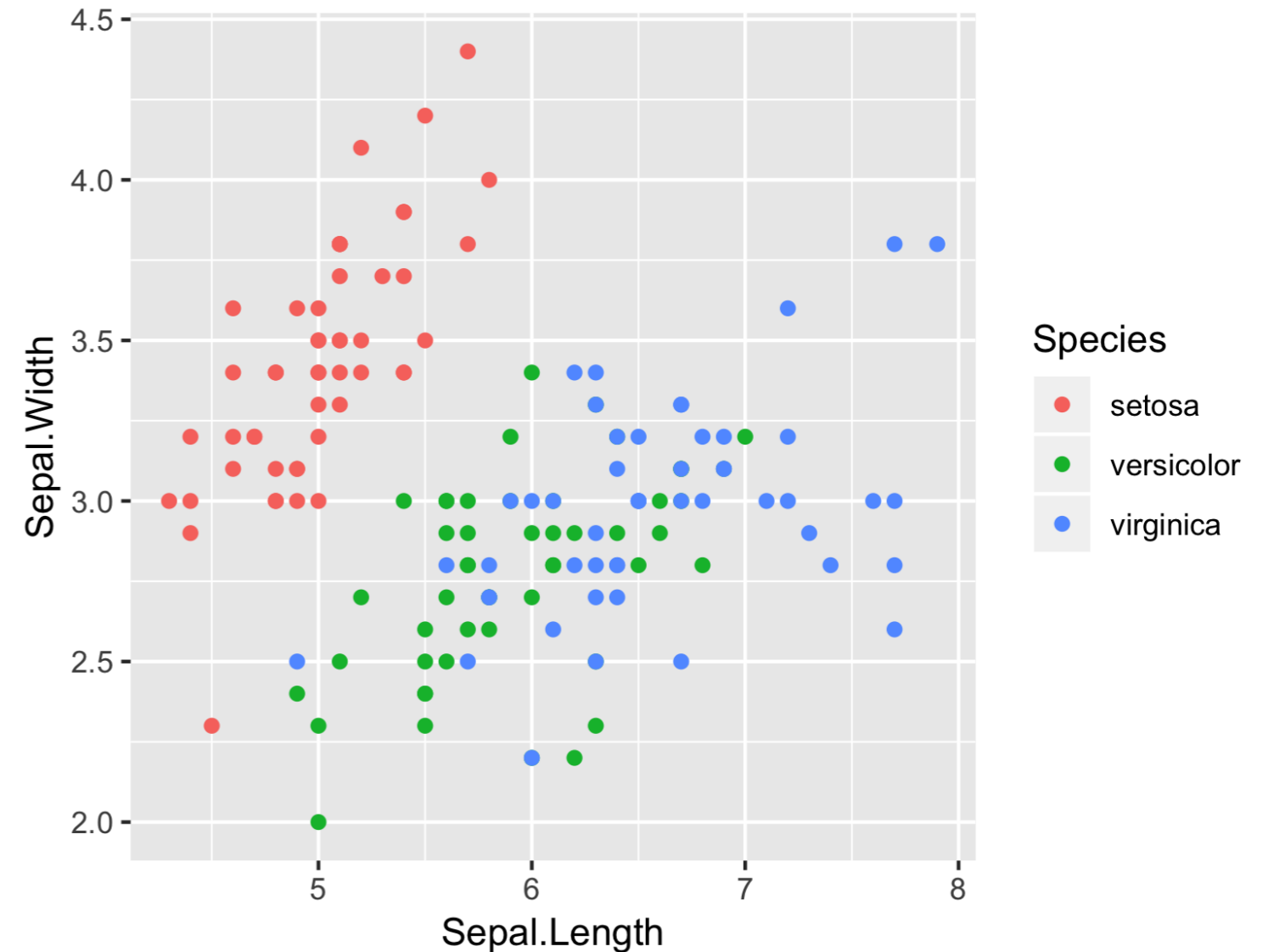
# Mapping onto the color aesthetic

```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point()
```

Type	Variable
Color	Species

**Species**, a dataframe column, is *mapped onto color*, a visible aesthetic.

Map aesthetics in `aes()`.

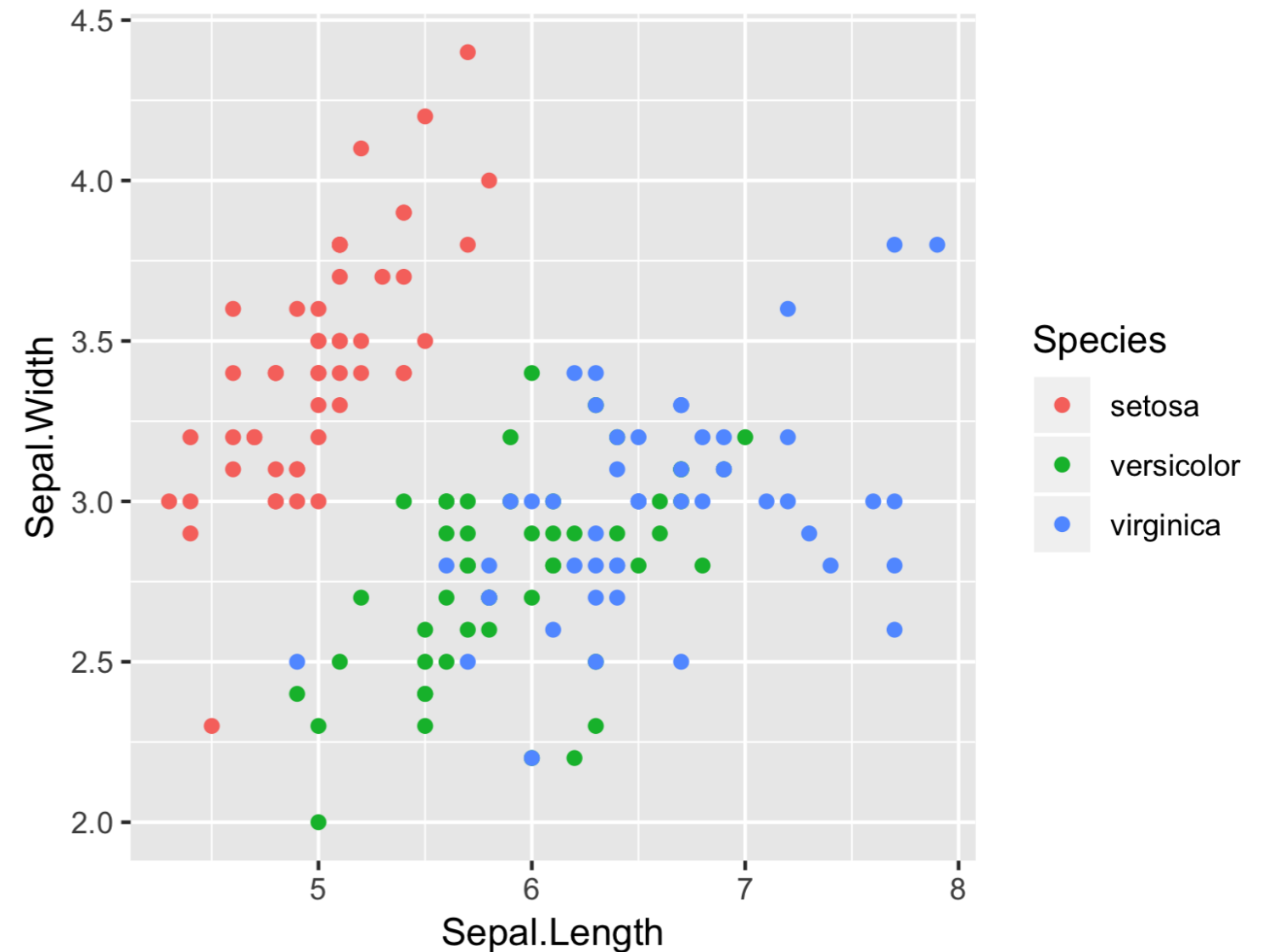


# Mapping onto the color aesthetic in geom

```
ggplot(iris) +  
  geom_point(aes(x = Sepal.Length,  
                y = Sepal.Width,  
                col = Species))
```

Only necessary if:

- All layers should *not* inherit the same aesthetics
- Mixing different data sources



# Typical visible aesthetics

Aesthetic	Description
x	X axis position
y	Y axis position

# Typical visible aesthetics

Aesthetic	Description
x	X axis position
y	Y axis position
fill	Fill color



# Typical visible aesthetics

Aesthetic	Description
x	X axis position
y	Y axis position
fill	Fill color
color	Color of points, outlines of other geoms

# Typical visible aesthetics

Aesthetic	Description
x	X axis position
y	Y axis position
fill	Fill color
color	Color of points, outlines of other geoms
size	Area or radius of points, thickness of lines

# Typical visible aesthetics

Aesthetic	Description
x	X axis position
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fill	Fill color
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size	Area or radius of points, thickness of lines

Aesthetic	Description
alpha	Transparency

# Typical visible aesthetics

Aesthetic	Description
x	X axis position
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fill	Fill color
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size	Area or radius of points, thickness of lines

Aesthetic	Description
alpha	Transparency
linetype	Line dash pattern

# Typical visible aesthetics

Aesthetic	Description
x	X axis position
y	Y axis position
fill	Fill color
color	Color of points, outlines of other geoms
size	Area or radius of points, thickness of lines

Aesthetic	Description
alpha	Transparency
linetype	Line dash pattern
labels	Text on a plot or axes

# Typical visible aesthetics

Aesthetic	Description
x	X axis position
y	Y axis position
fill	Fill color
color	Color of points, outlines of other geoms
size	Area or radius of points, thickness of lines

Aesthetic	Description
alpha	Transparency
linetype	line dash pattern
labels	Text on a plot or axes
shape	Shape

# Let's Practice

INTRODUCTION TO DATA VISUALIZATION WITH GGPLOT2

# Using attributes

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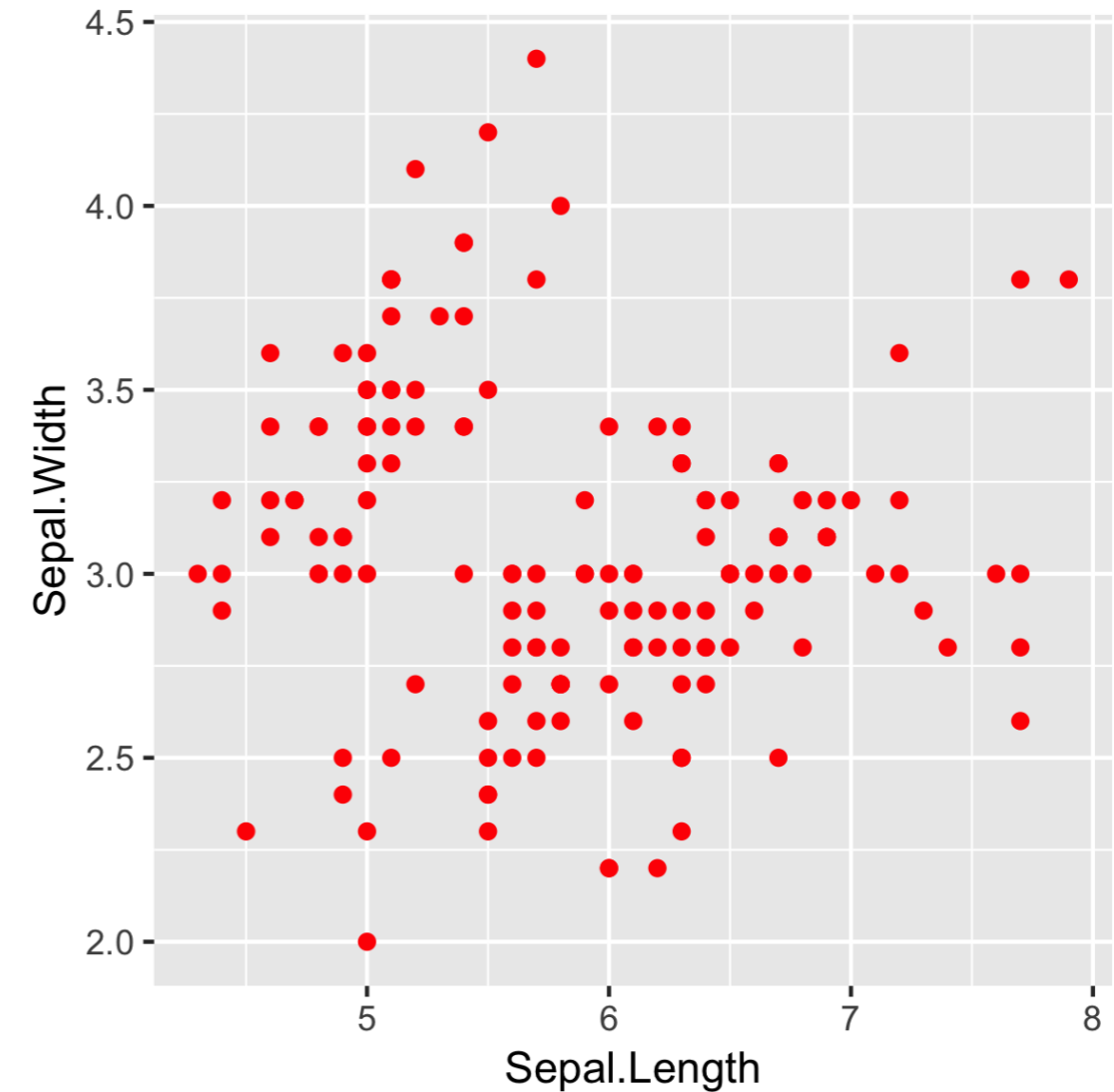
# Aesthetics? Attributes!

```
ggplot(iris, aes(x = Sepal.Length,  
                 y = Sepal.Width)) +  
  geom_point(color = "red")
```

Type	Property
Color	"red"

Set attributes in `geom_*()`.

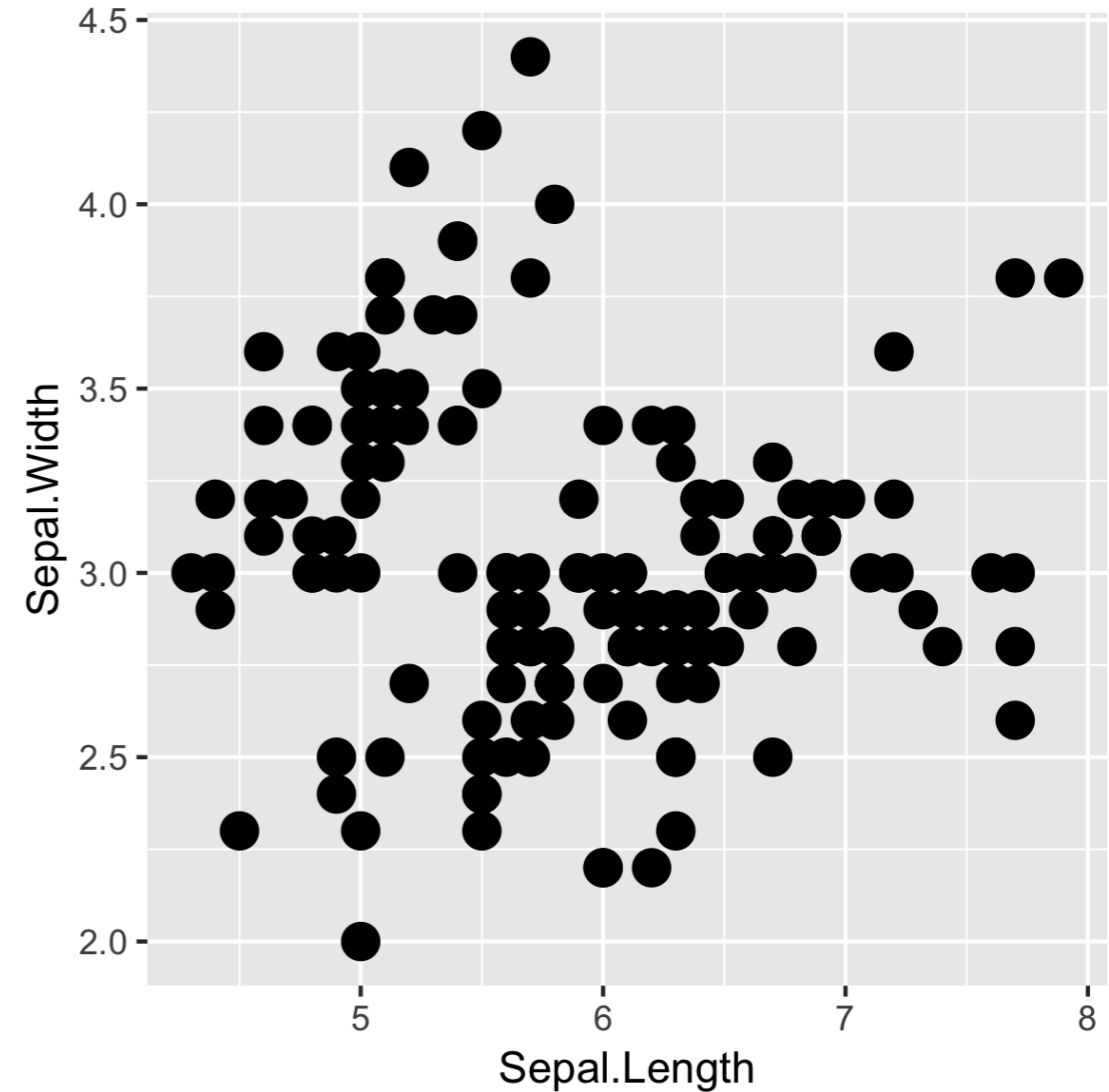
The **color** attribute is *set* to "red".



# Aesthetics? Attributes!

```
ggplot(iris, aes(x = Sepal.Length,  
                 y = Sepal.Width)) +  
  geom_point(size = 10)
```

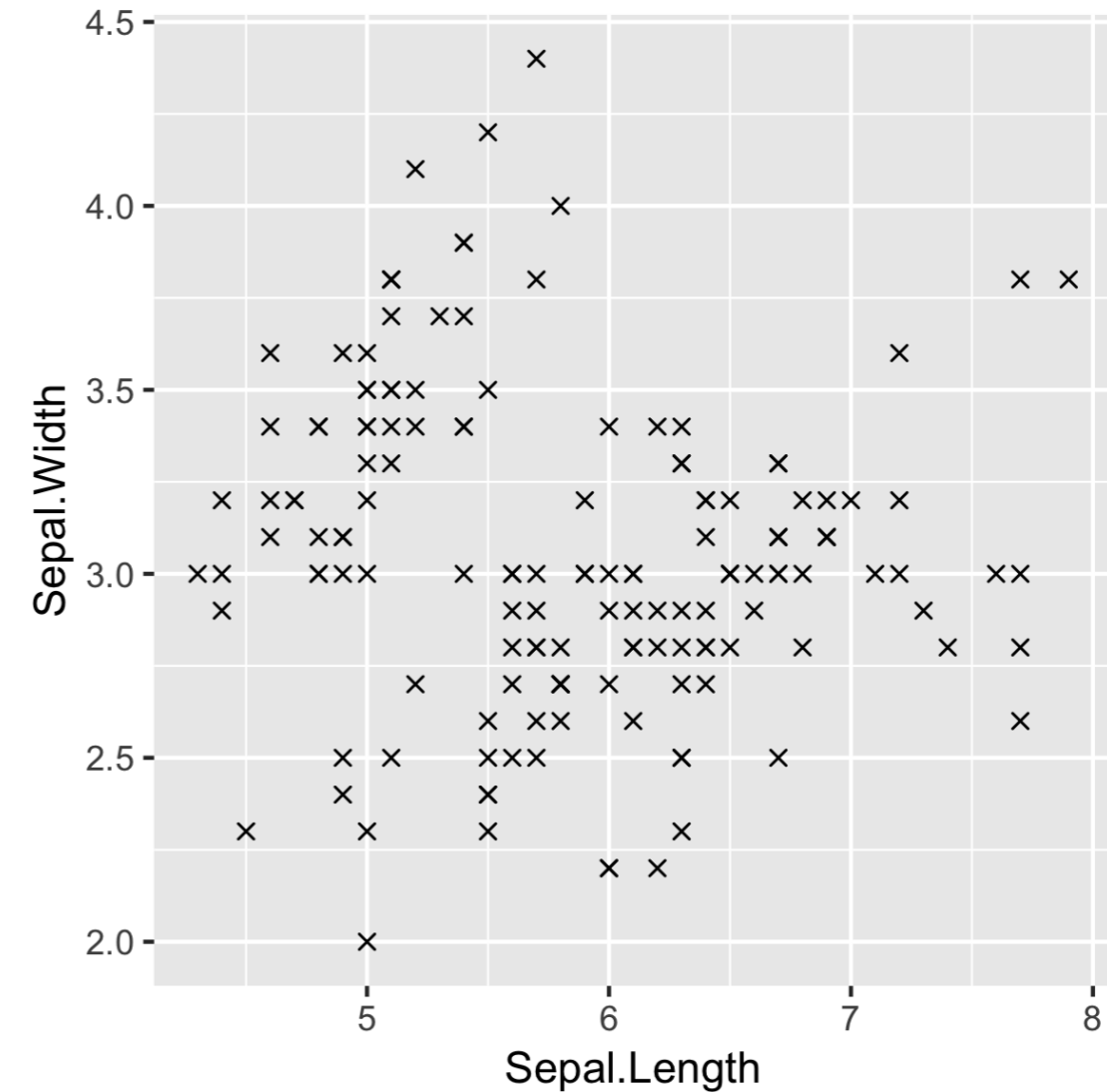
Type	Property
Size	4



# Aesthetics? Attributes!

```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width)) +  
  geom_point(shape = 4)
```

Type	Property
Shape	4



# Let's practice!

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# Modifying Aesthetics

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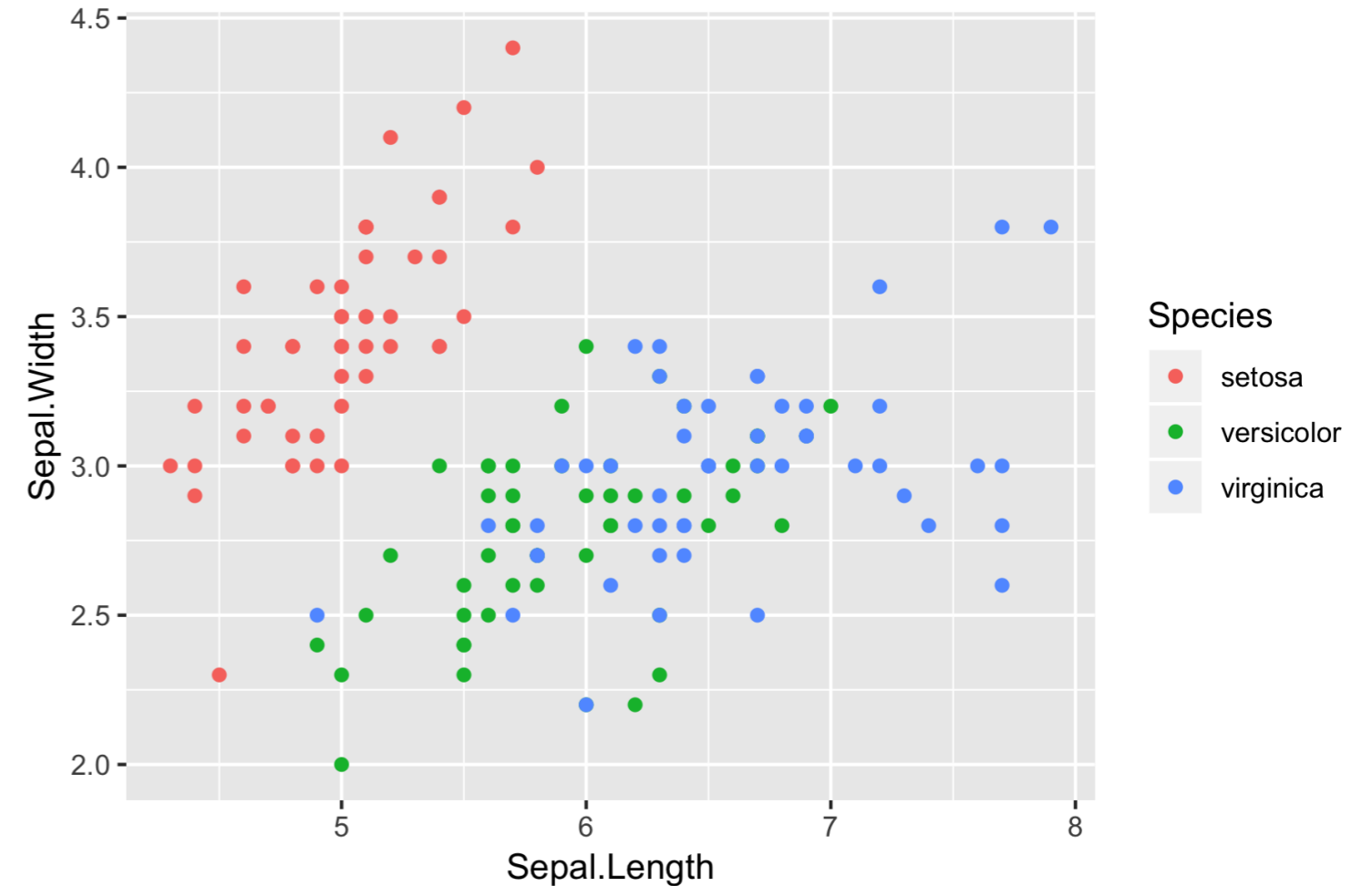
# Positions

Adjustment for overlapping

- identity
- dodge
- stack
- fill
- jitter
- jitterdodge
- nudge

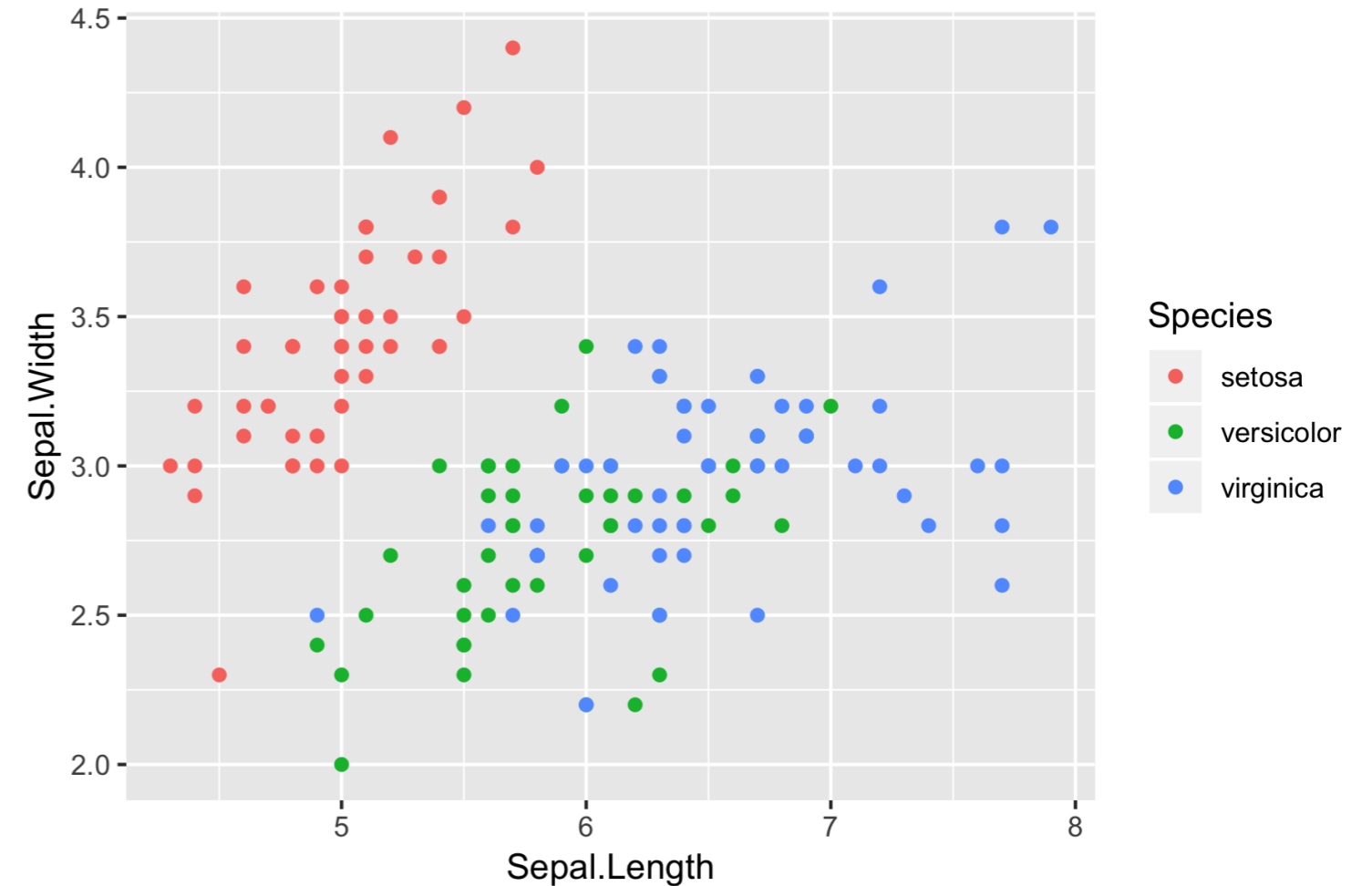
# position = "identity" (default)

```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point()
```



# position = "identity" (default)

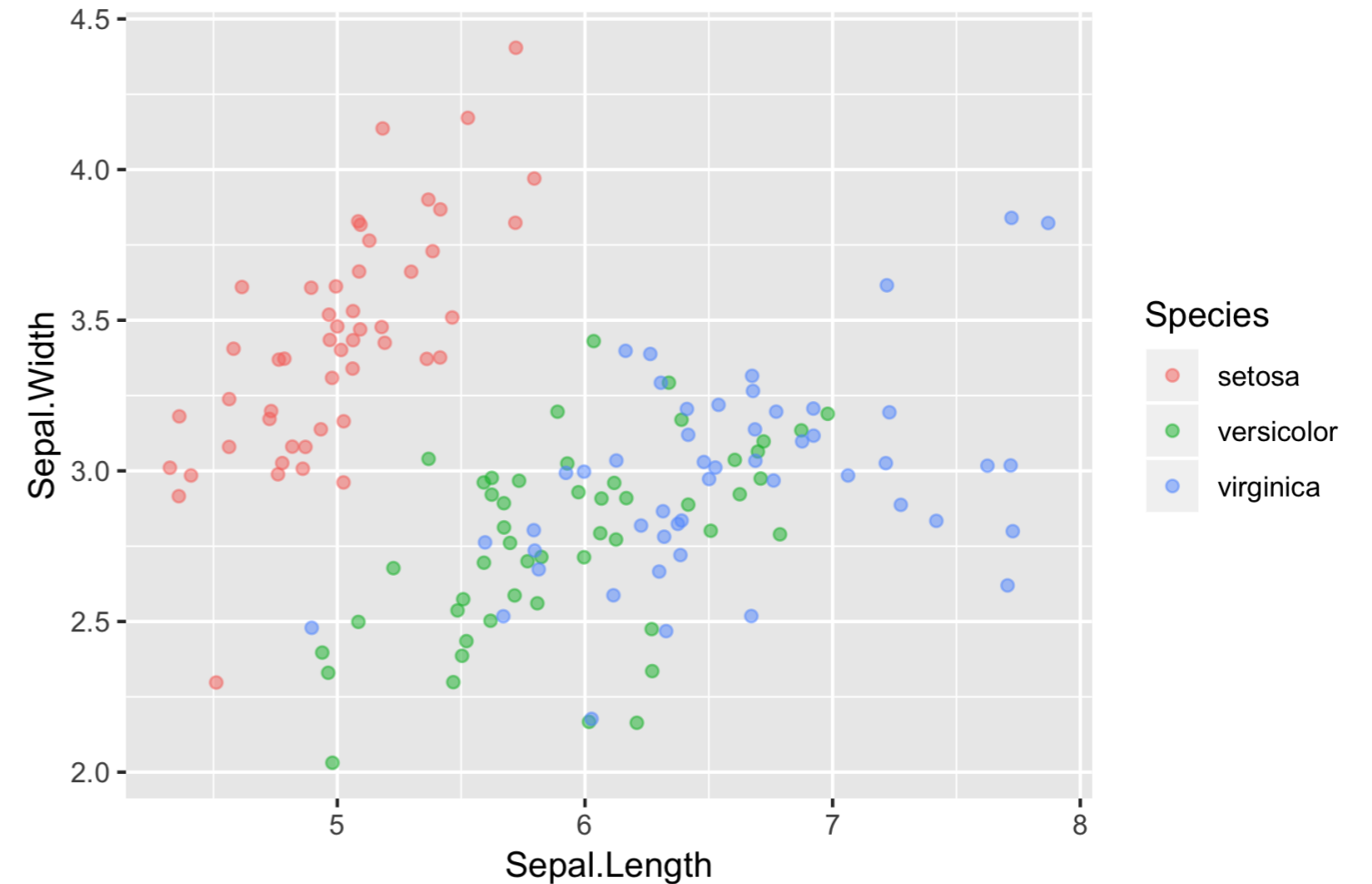
```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point(position = "identity")
```





# position = "jitter"

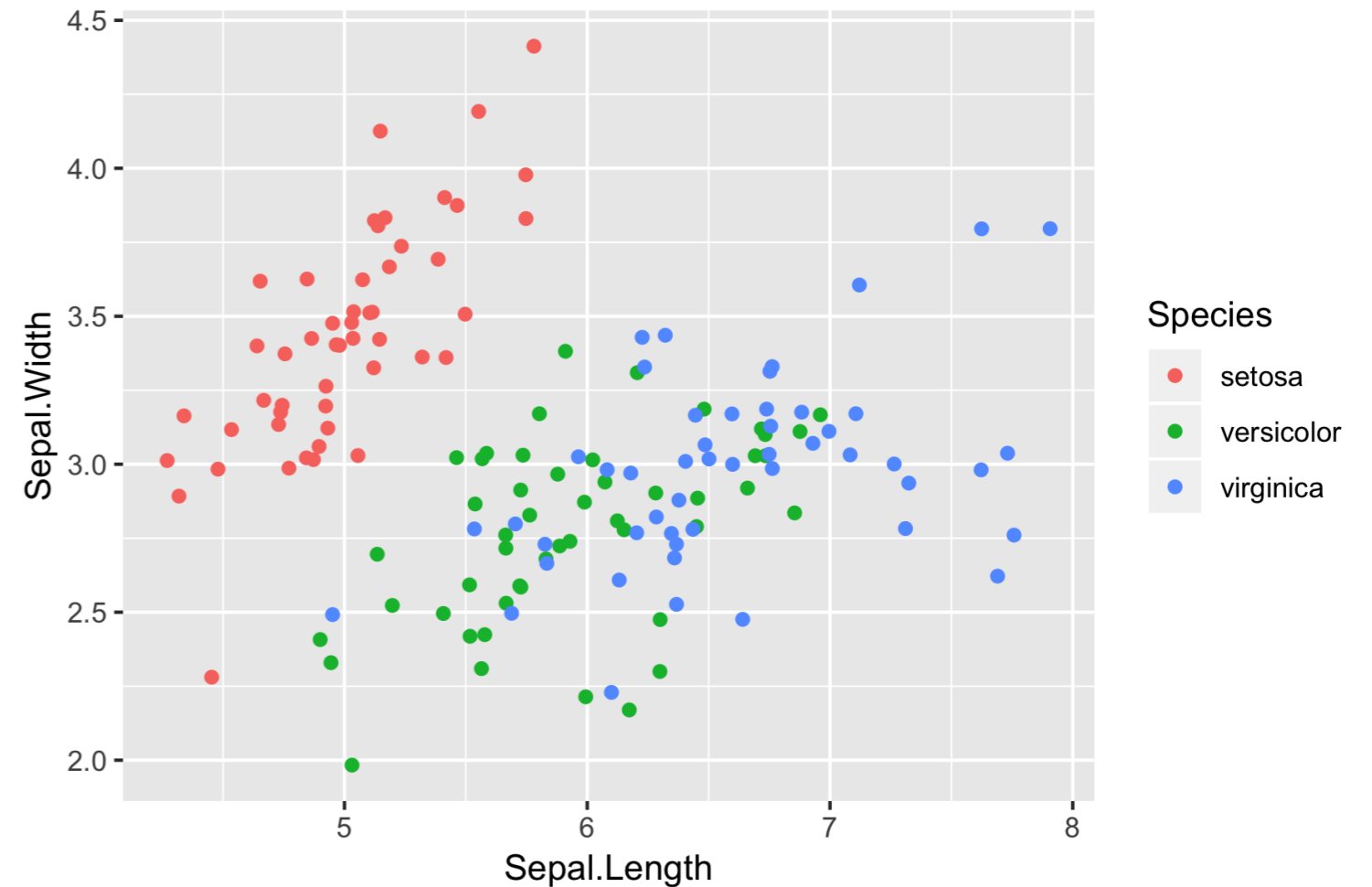
```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point(position = "jitter")
```



# position\_jitter()

```
posn_j <- position_jitter(0.1)

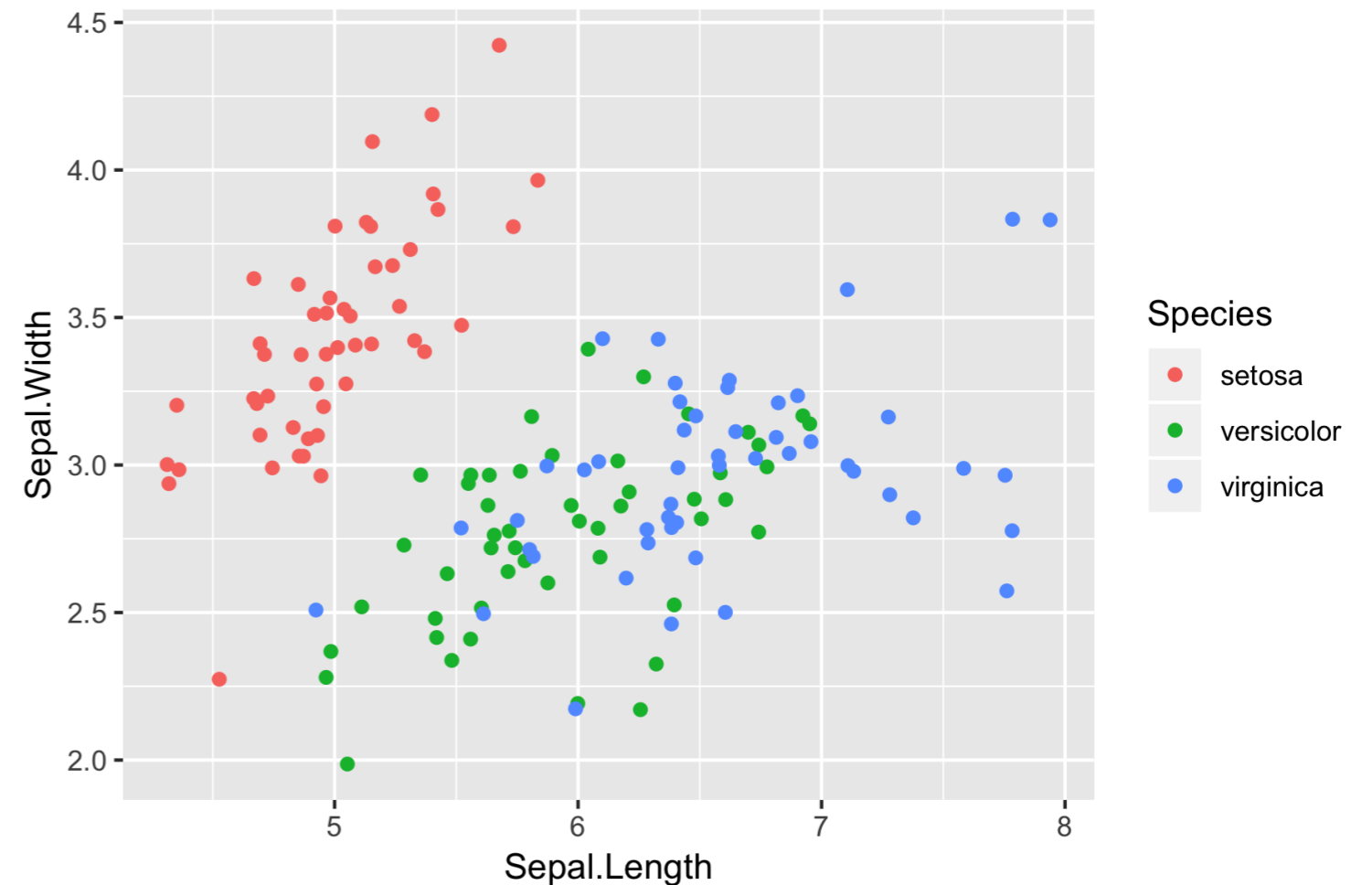
ggplot(iris, aes(x = Sepal.Length,
                 y = Sepal.Width,
                 col = Species)) +
  geom_point(position = posn_j)
```



# position\_jitter()

```
posn_j <- position_jitter(0.1,  
                          seed = 136)  
  
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point(position = posn_j)
```

- Set arguments for the position
- Consistency across plots & layers



# Scale functions

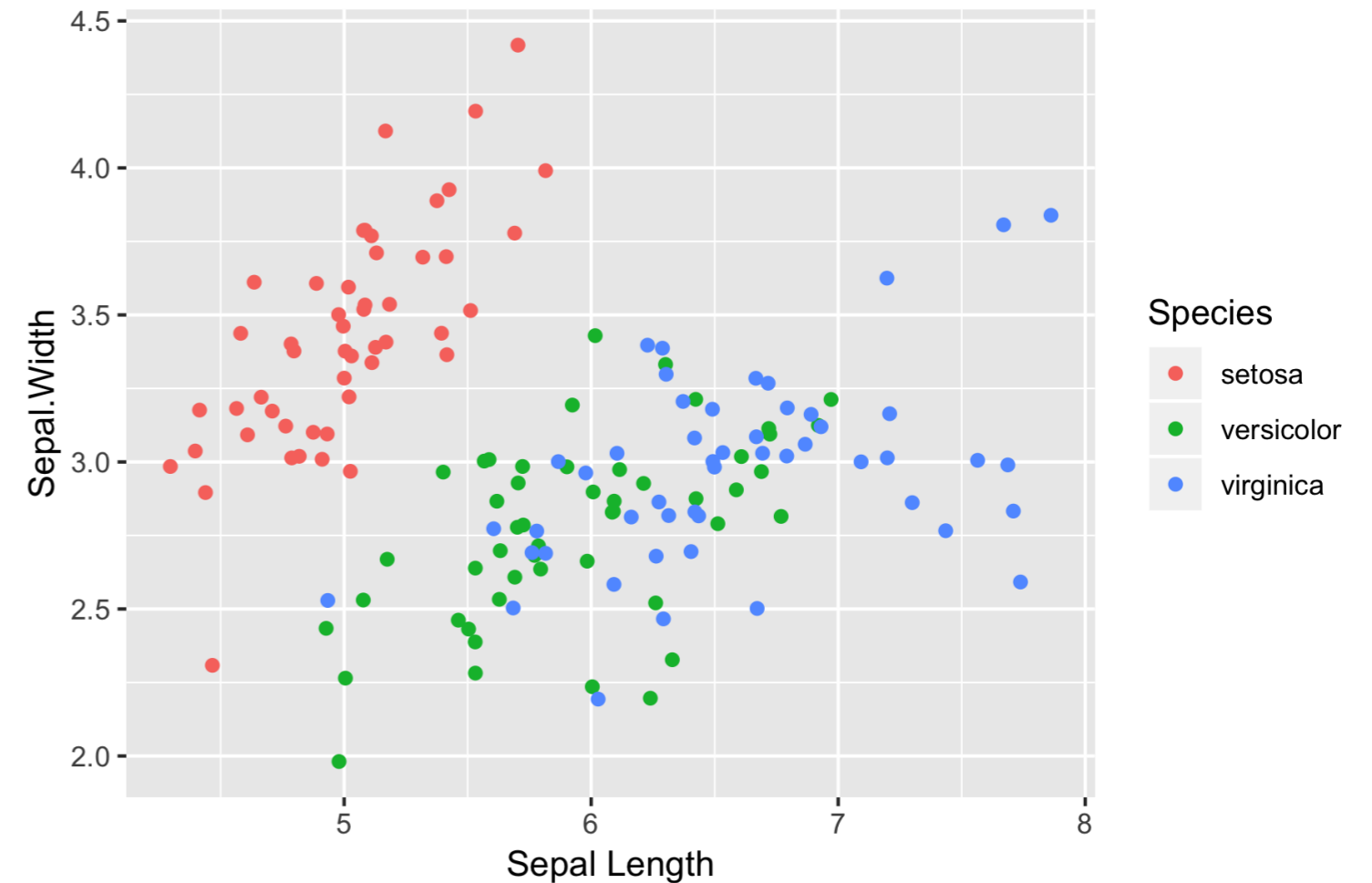
- `scale_x_*`()
- `scale_y_*`()
- `scale_color_*`()
  - Also `scale_colour_*`()
- `scale_fill_*`()
- `scale_shape_*`()
- `scale_linetype_*`()
- `scale_size_*`()

# Scale functions

- `scale_x_continuous()`
- `scale_y_*()`
- `scale_color_discrete()`
  - Alternatively, `scale_colour_*()`
- `scale_fill_*()`
- `scale_shape_*()`
- `scale_linetype_*()`
- `scale_size_*()`

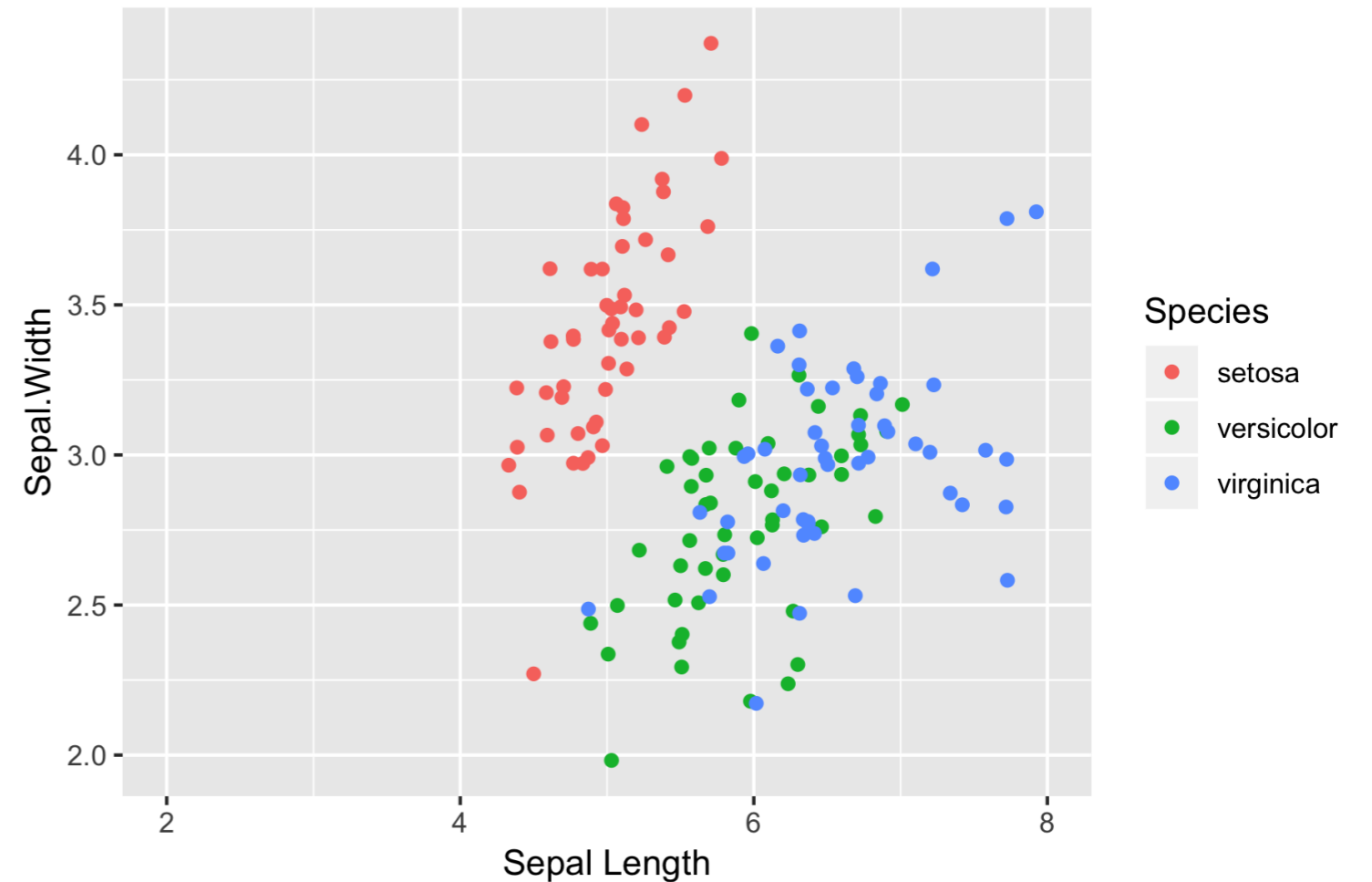
# scale\_\*\_\*()

```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point(position = "jitter") +  
  scale_x_continuous("Sepal Length") +  
  scale_color_discrete("Species")
```



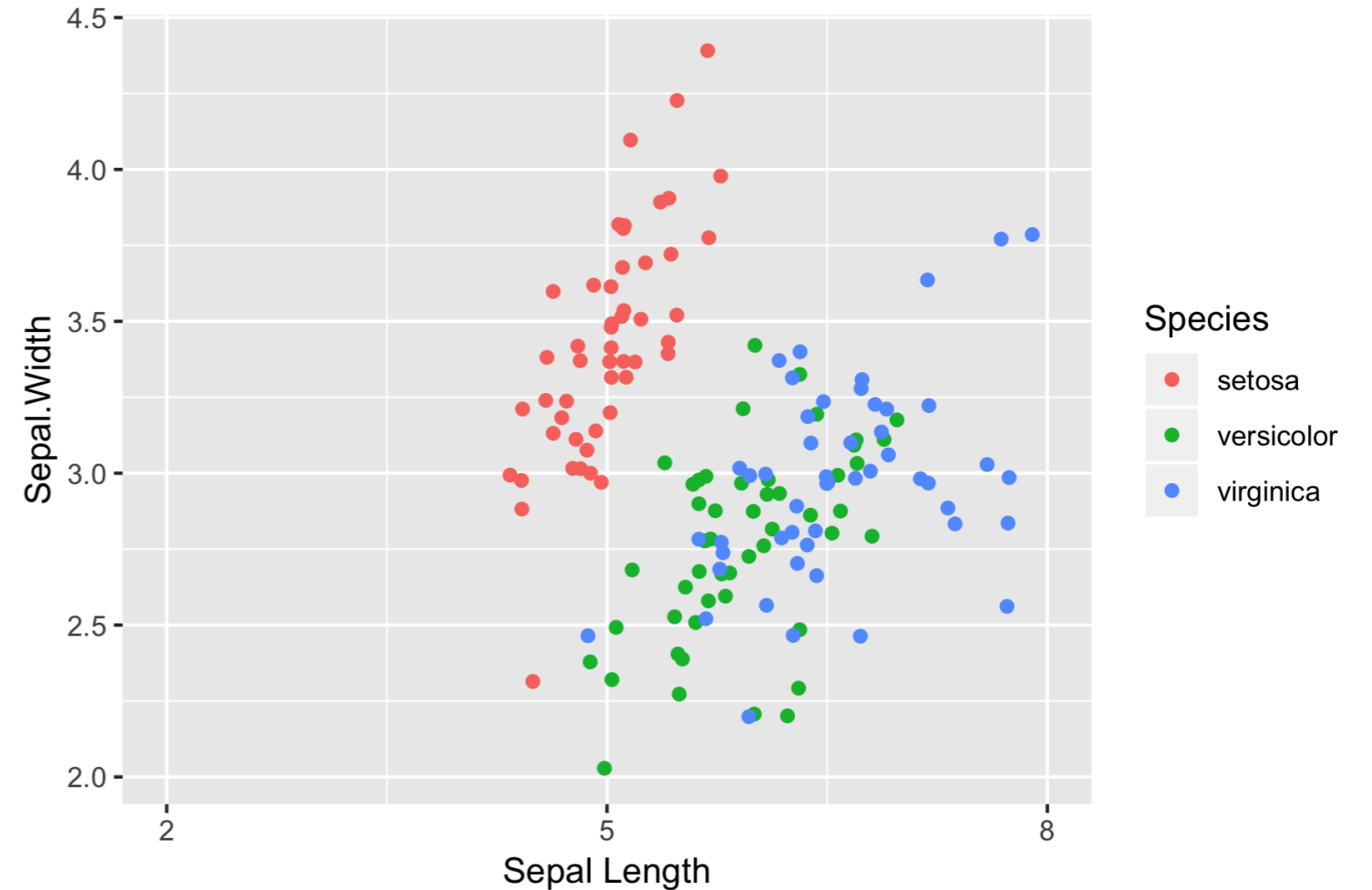
# The limits argument

```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point(position = "jitter") +  
  scale_x_continuous("Sepal Length",  
                    limits = c(2,8)) +  
  scale_color_discrete("Species")
```



# The breaks argument

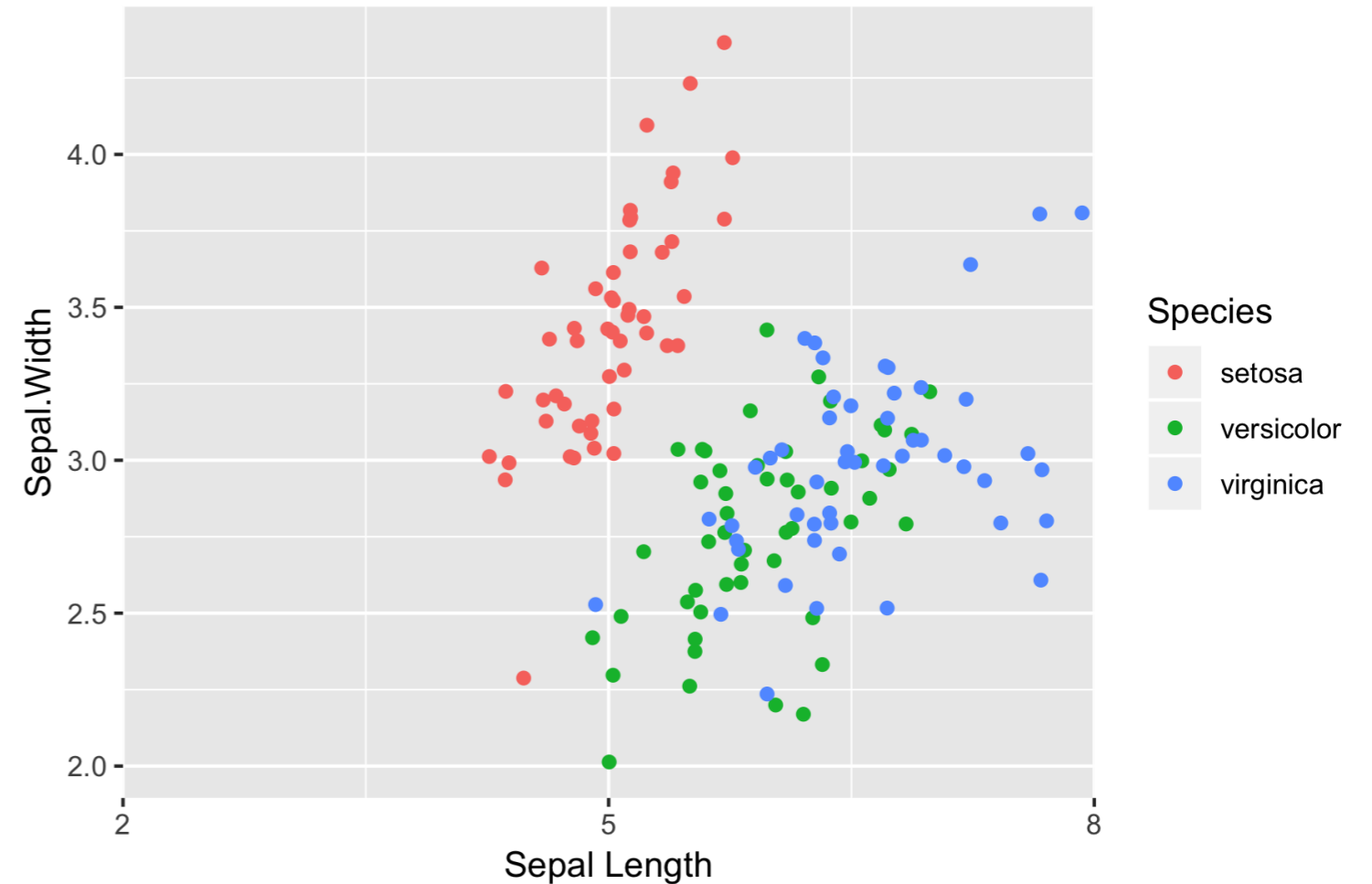
```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point(position = "jitter") +  
  scale_x_continuous("Sepal Length",  
                    limits = c(2, 8),  
                    breaks = seq(2, 8, 3)) +  
  scale_color_discrete("Species")
```





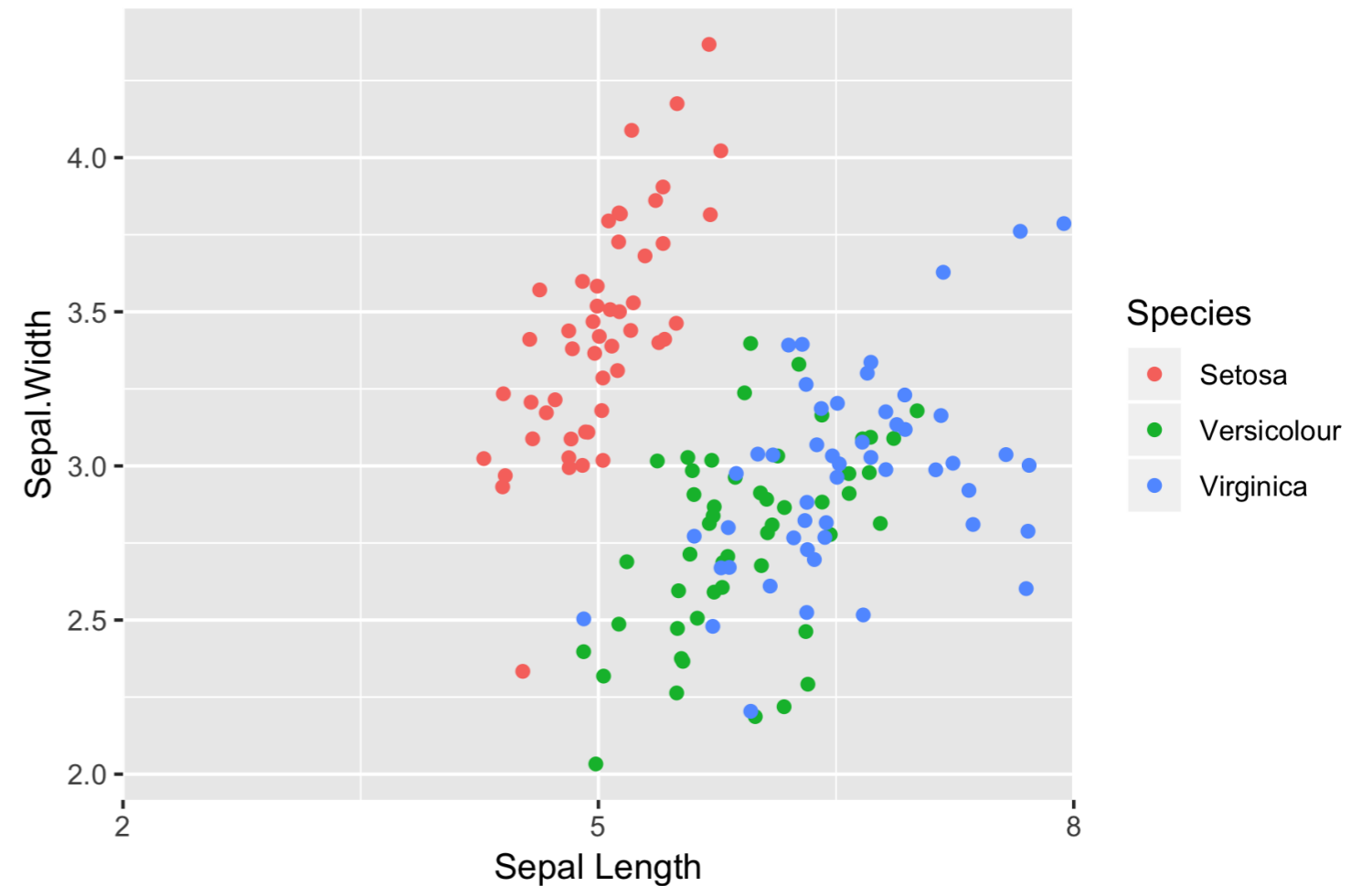
# The expand argument

```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point(position = "jitter") +  
  scale_x_continuous("Sepal Length",  
                    limits = c(2, 8),  
                    breaks = seq(2, 8, 3),  
                    expand = c(0, 0)) +  
  scale_color_discrete("Species")
```



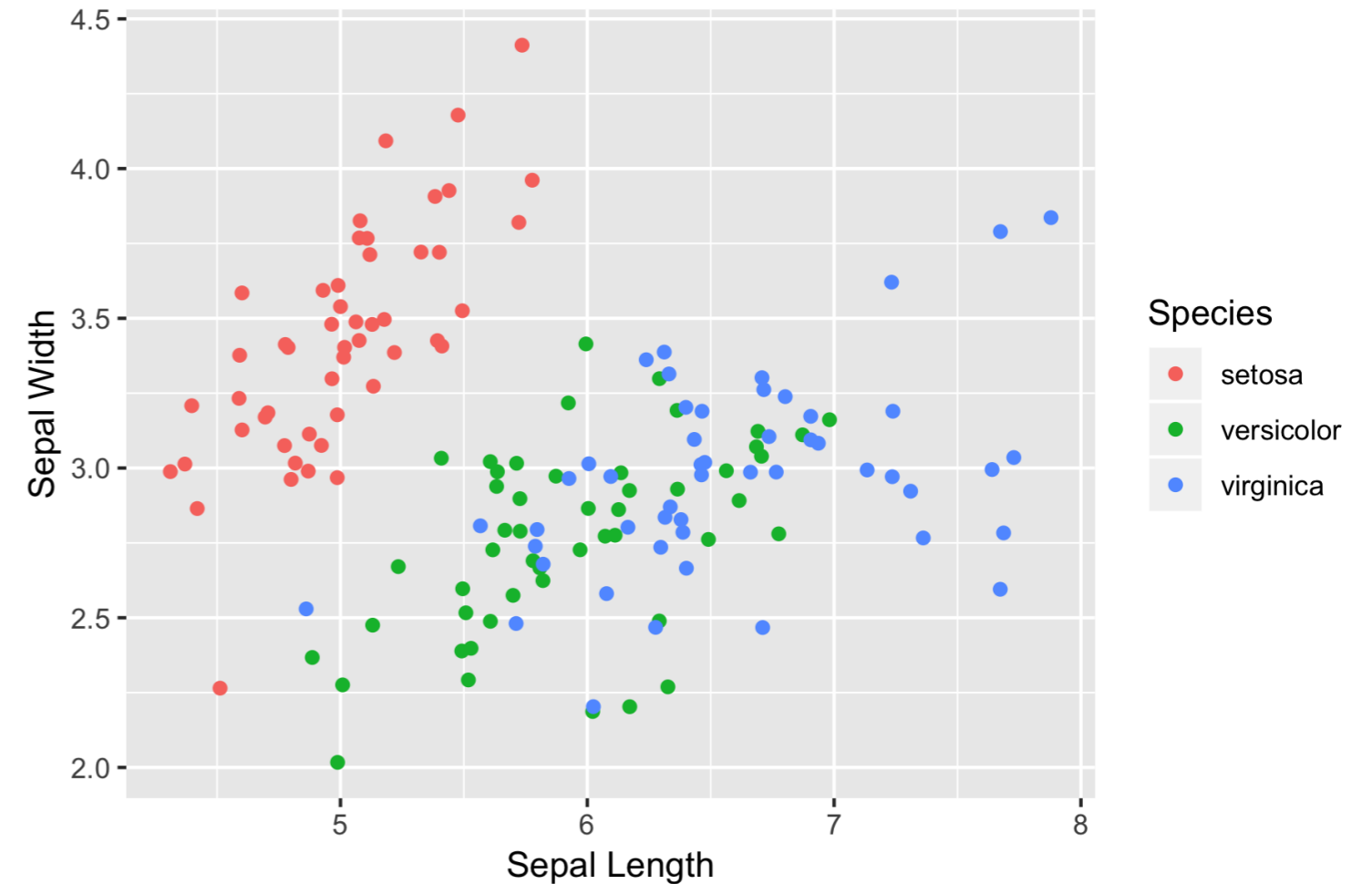
# The labels argument

```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point(position = "jitter") +  
  scale_x_continuous("Sepal Length",  
                    limits = c(2, 8),  
                    breaks = seq(2, 8, 3),  
                    expand = c(0, 0),  
                    labels = c("Setosa",  
                               "Versicolor",  
                               "Virginica")) +  
  scale_color_discrete("Species")
```



# labs()

```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point(position = "jitter") +  
  labs(x = "Sepal Length",  
       y = "Sepal Width",  
       color = "Species")
```



# Let's try it out!

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# Aesthetics best practices

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# Which aesthetics?

- Use your creative know-how, and
- Follow some clear guidelines
- Jacques Bertin
  - The Semiology of Graphics, 1967
- William Cleveland
  - The Elements of Graphing Data, 1985
  - Visualizing Data, 1993

# Form follows function



# Form follows function

## Function

### Primary:

- Accurate and efficient representations

### Secondary:

- Visually appealing, beautiful plots

## Guiding principles

### Never:

- Misrepresent or obscure data
- Confuse viewers with complexity

### Always:

- Consider the audience and purpose of every plot



x	y = f(x)	Group
51	3.5	A
4.9	3.0	A
4.7	3.2	A
4.6	3.1	A
⋮	⋮	⋮
70	3.2	B
6.4	3.2	B
6.9	3.1	B
5.5	2.3	B
⋮	⋮	⋮

x	y = f(x)	Group
5.1	3.5	A
4.9	3.0	A
4.7	3.2	A
4.6	3.1	A
⋮	⋮	⋮
7.0	3.2	B
6.4	3.2	B
6.9	3.1	B
5.5	2.3	B
⋮	⋮	⋮

Difficult directly  
from data

For each group:

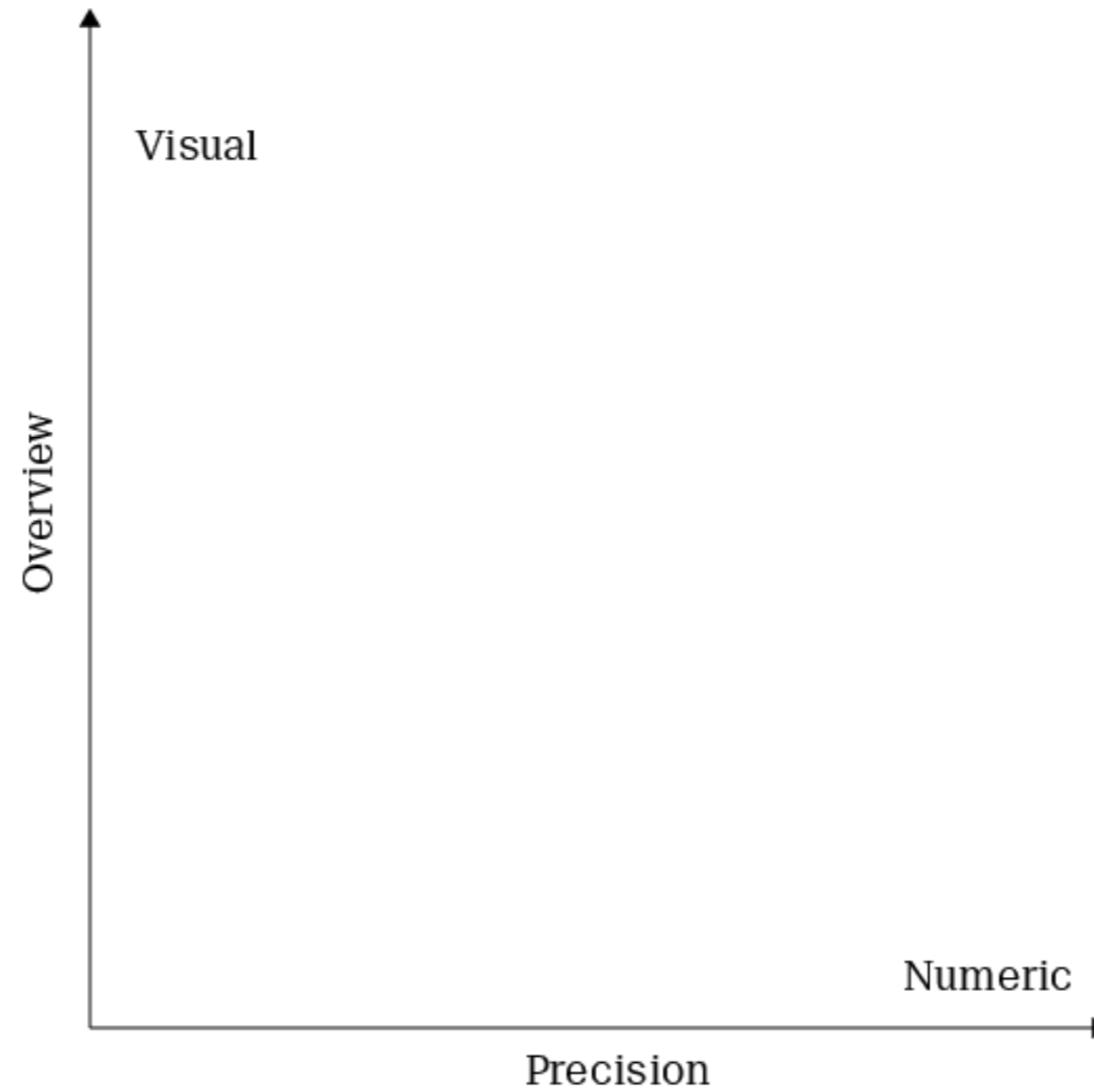
$n$  range outliers

$\bar{x}$   $s_x$   $\bar{y}$   $s_y$

$y = f(x) = \beta_0 + \beta_1 x$

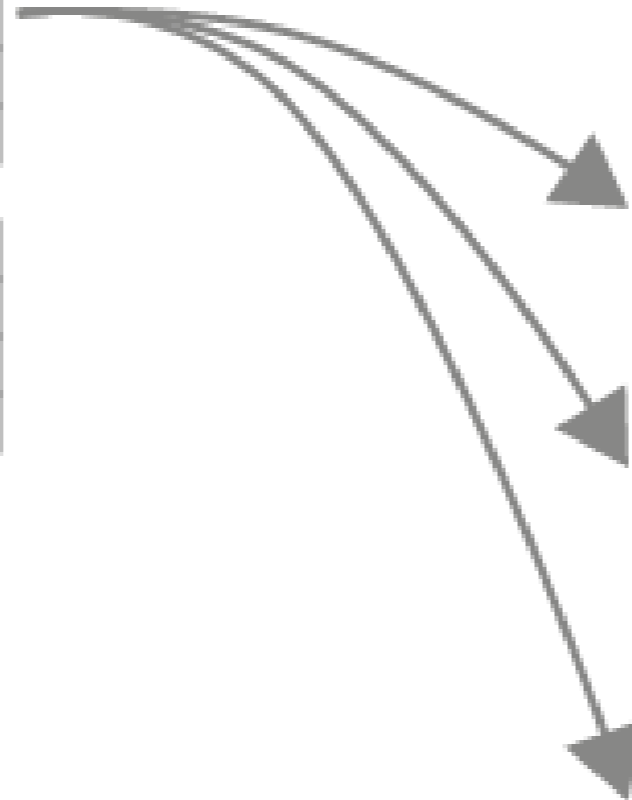
$r$   $R^2$

# Extracting information from Data



x	y = f(x)	Group
5.1	3.5	A
4.9	3.0	A
4.7	3.2	A
4.6	3.1	A
⋮	⋮	⋮
7.0	3.2	B
6.4	3.2	B
6.9	3.1	B
5.5	2.3	B
⋮	⋮	⋮

ENCODE



For each group:

$n$  range outliers

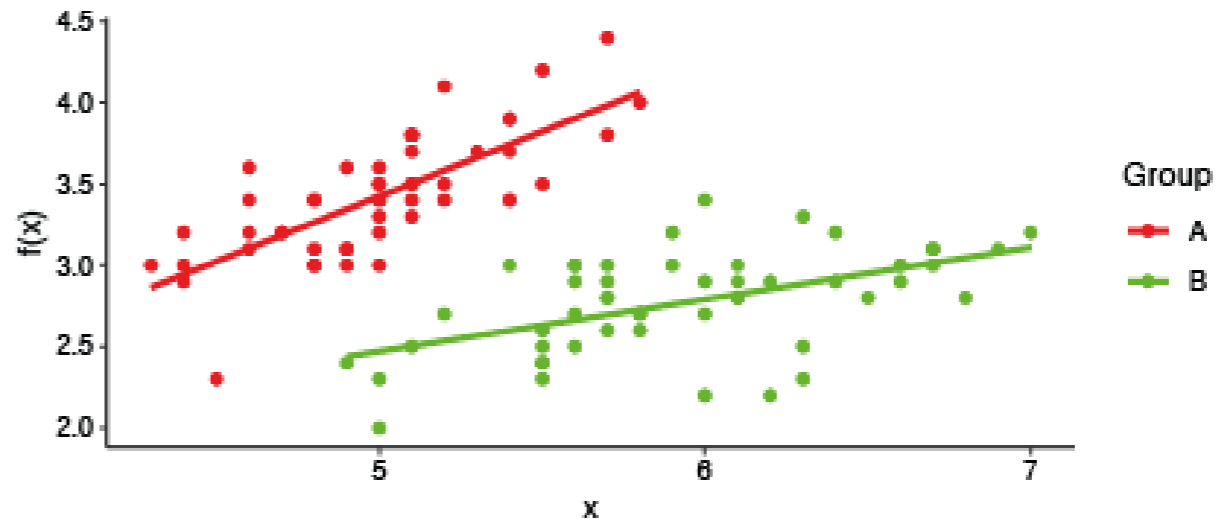
$\bar{x}$   $s_x$   $\bar{y}$   $s_y$

$y = f(x) = \beta_0 + \beta_1 x$

$r$   $R^2$

x	y = f(x)	Group
5.1	3.5	A
4.9	3.0	A
4.7	3.2	A
4.6	3.1	A
...	...	...
7.0	3.2	B
6.4	3.2	B
6.9	3.1	B
5.5	2.3	B
...	...	...

ENCODE



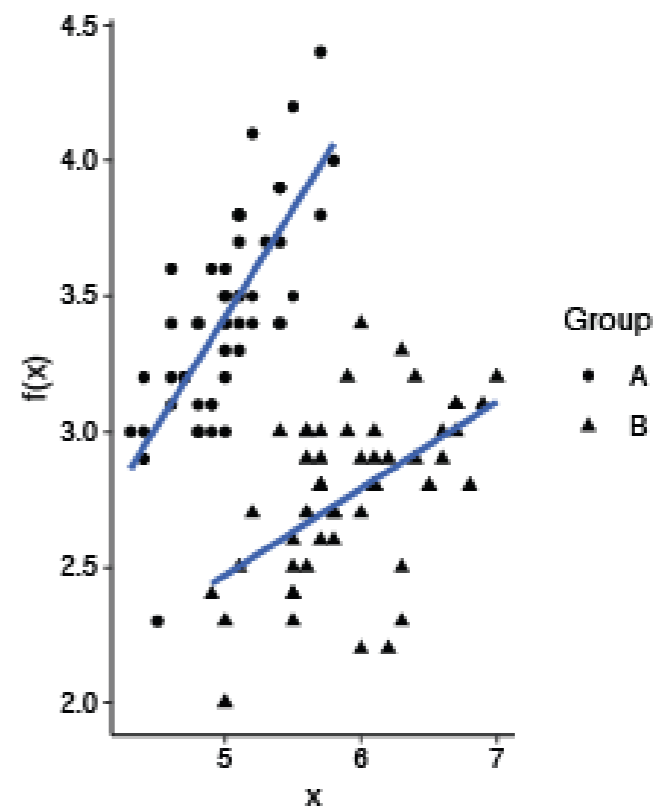
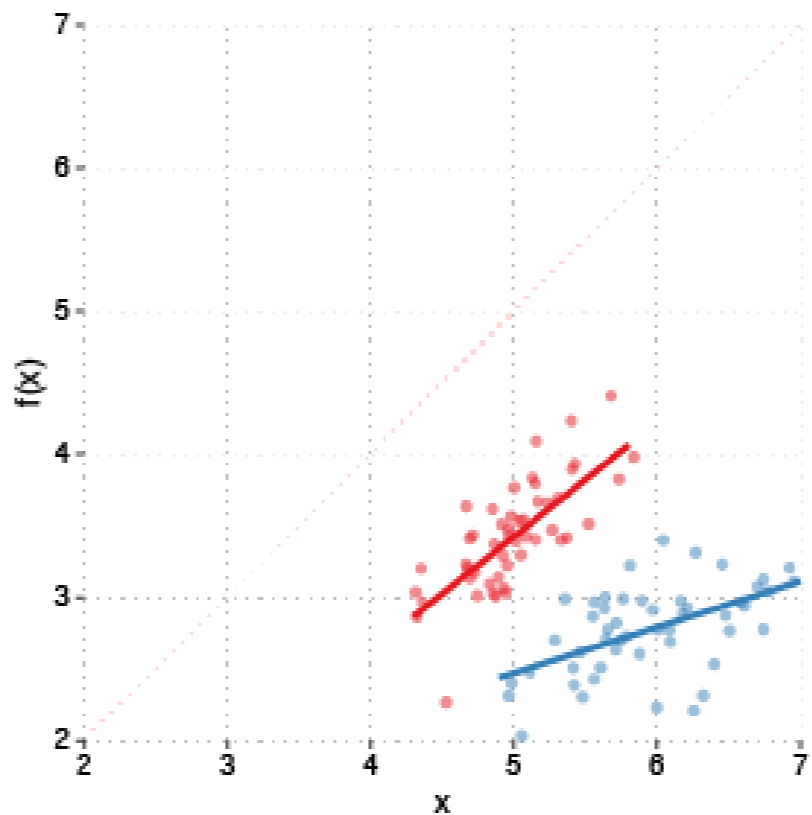
For each group:

$n$  range outliers

$\bar{x}$   $s_x$   $\bar{y}$   $s_y$

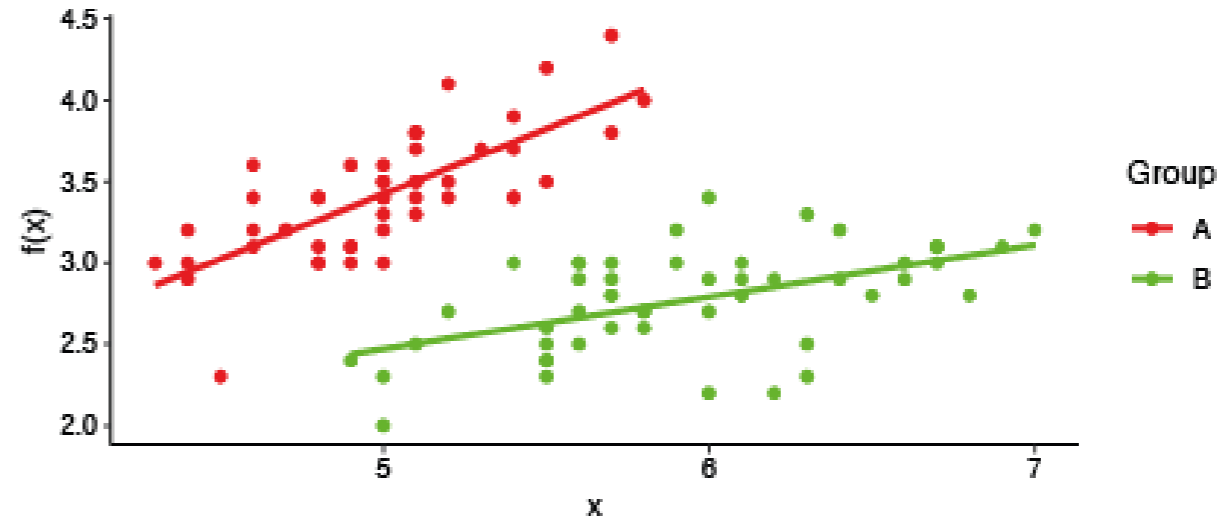
$y = f(x) = \beta_0 + \beta_1 x$

$r$   $R^2$



x	y = f(x)	Group
5.1	3.5	A
4.9	3.0	A
4.7	3.2	A
4.6	3.1	A
...	...	...
7.0	3.2	B
6.4	3.2	B
6.9	3.1	B
5.5	2.3	B
...	...	...

ENCODE



For each group:

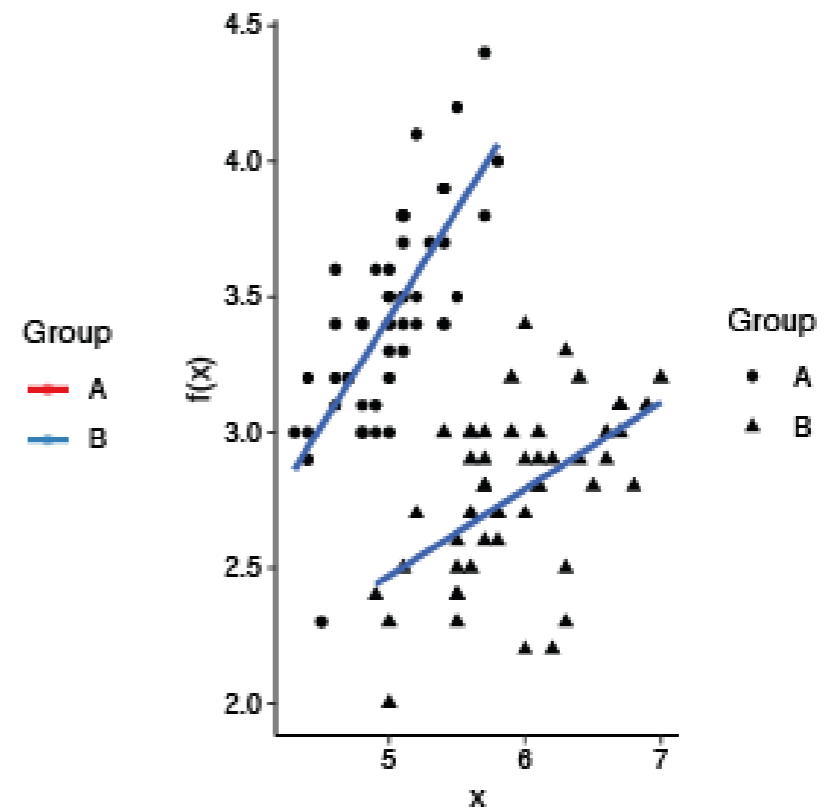
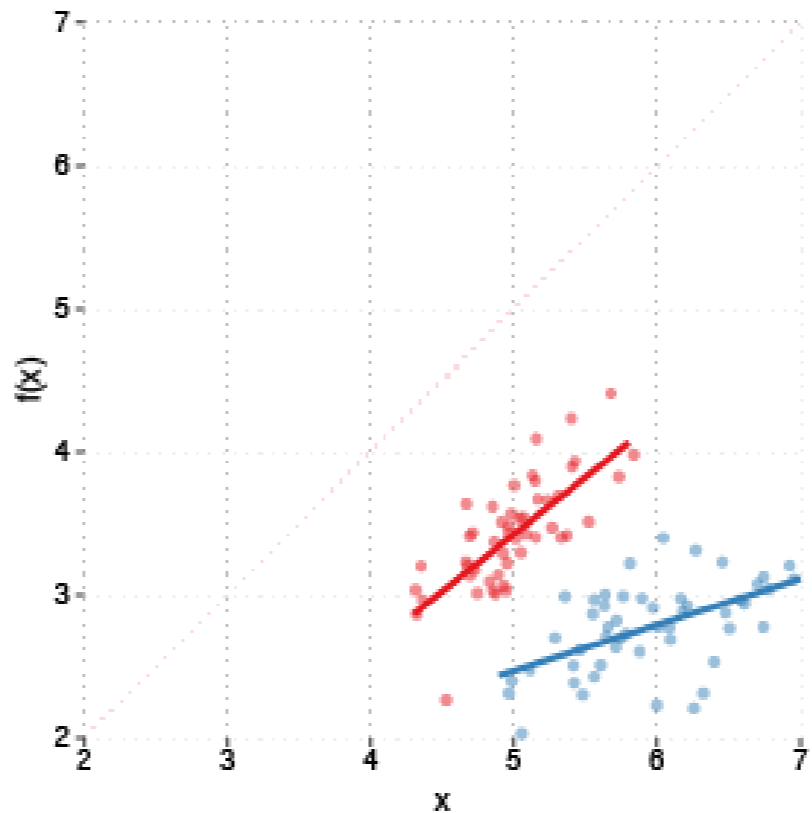
$n$  range outliers

$\bar{x}$   $s_x$   $\bar{y}$   $s_y$

$y = f(x) = \beta_0 + \beta_1 x$

$r$   $R^2$

DECODE

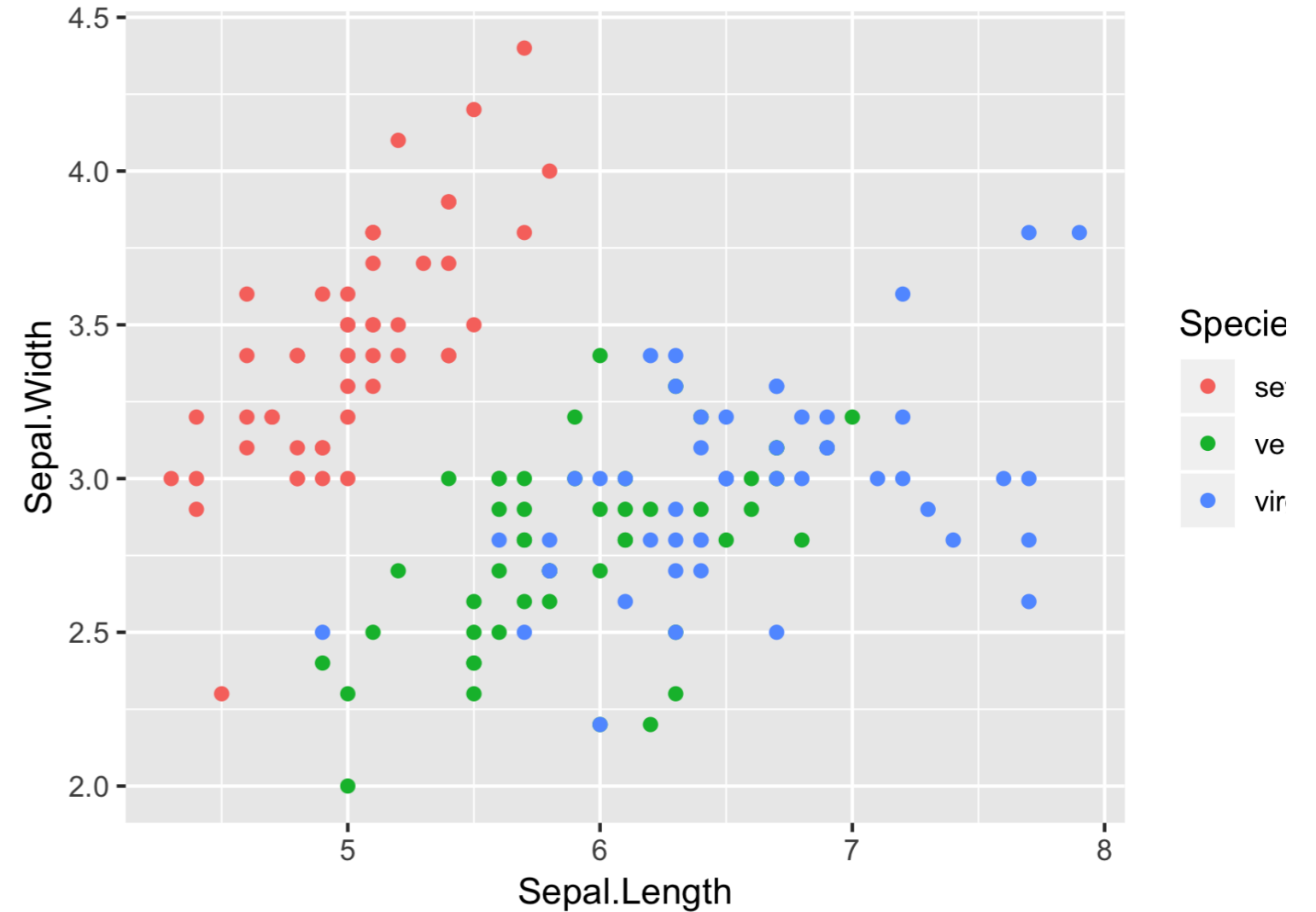


# The best choices for aesthetics

- Efficient
  - Provides a faster overview than numeric summaries
- Accurate
  - Minimizes information loss

# Aesthetics - continuous variables

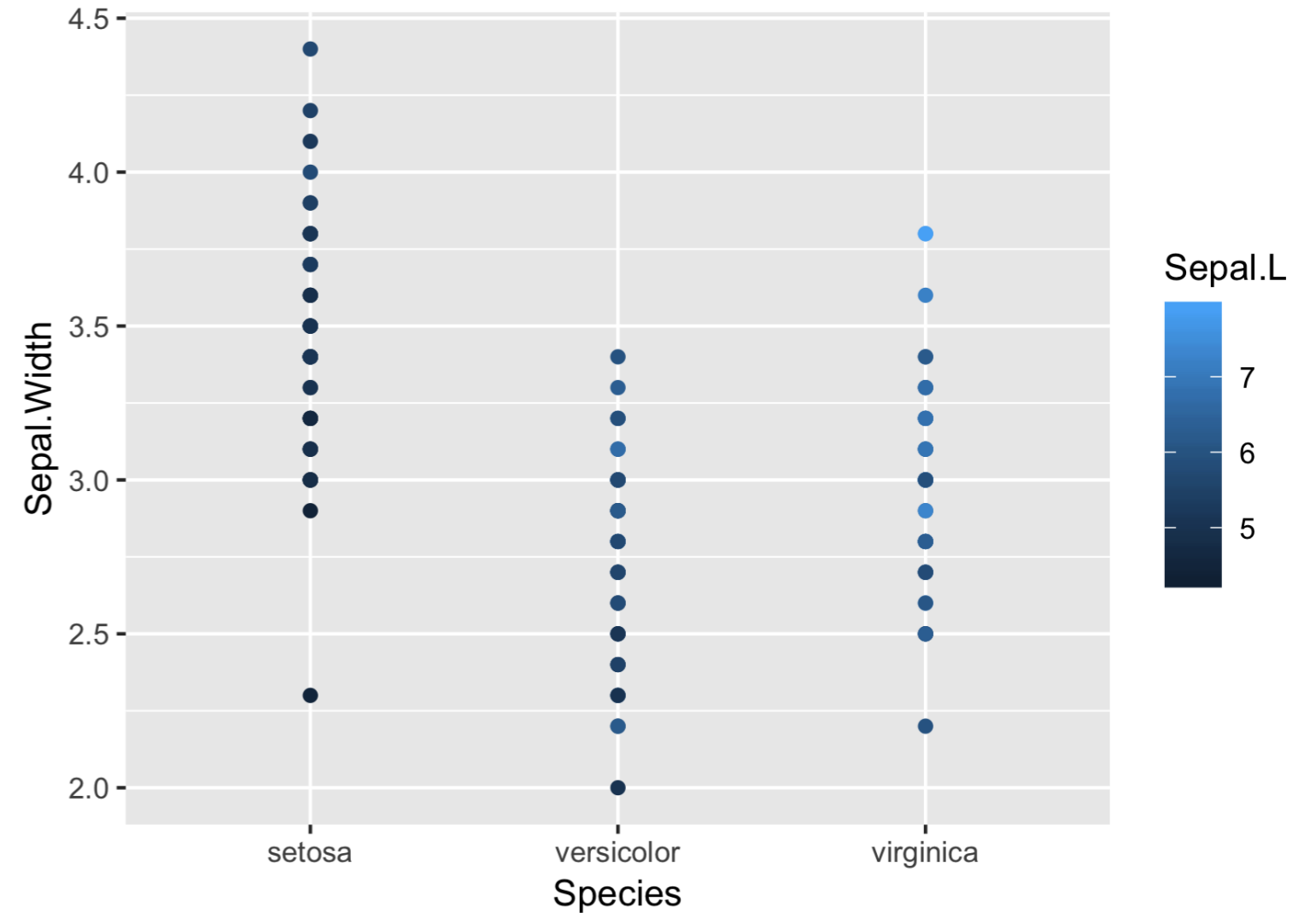
```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                color = Species)) +  
  geom_point()
```





# Aesthetics - continuous variables

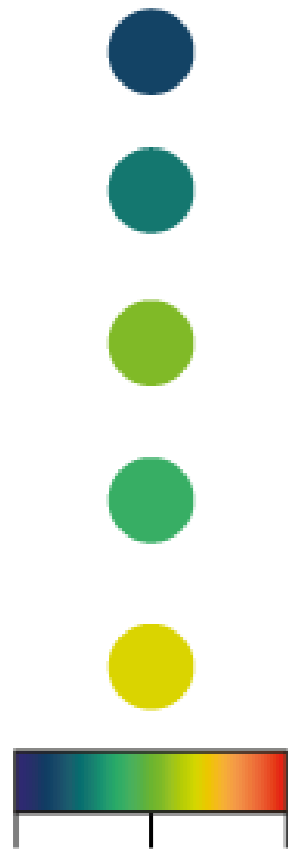
```
ggplot(iris, aes(color = Sepal.Length,  
                y = Sepal.Width,  
                x = Species)) +  
  geom_point()
```



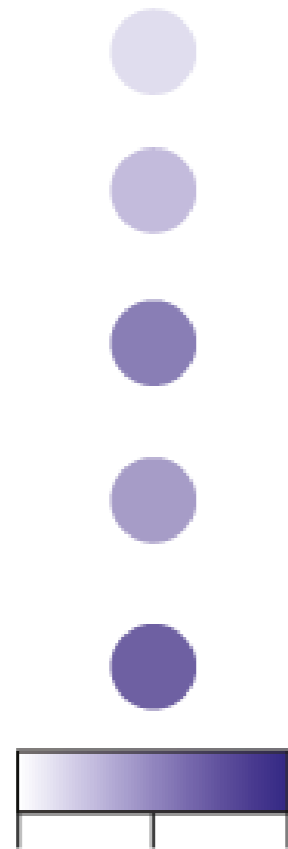
# Efficiency and Accuracy of Decoding

Low  High

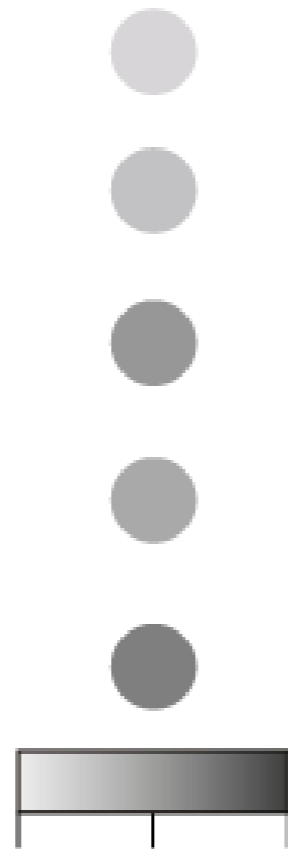
Colour  
Spectrum



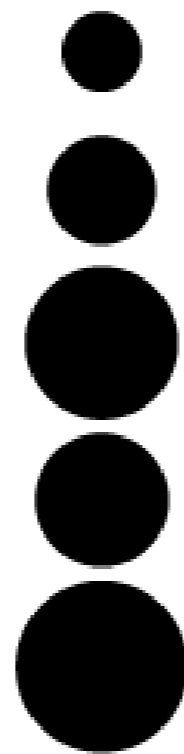
Monochromatic  
Spectrum



Grey-Scale  
Spectrum



Area



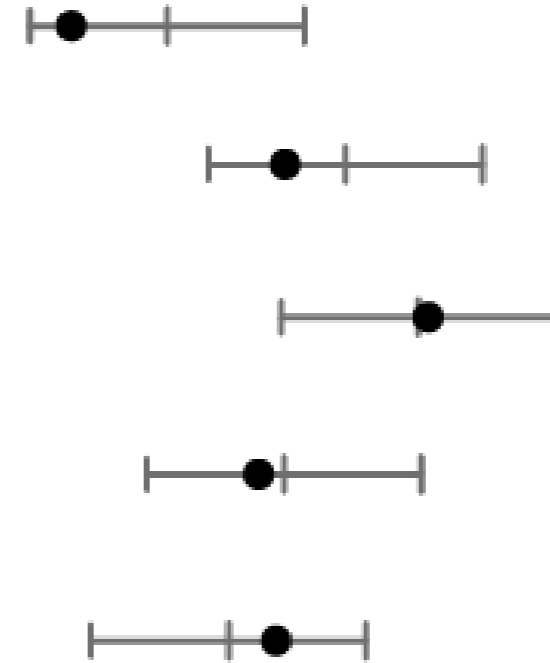
Angle



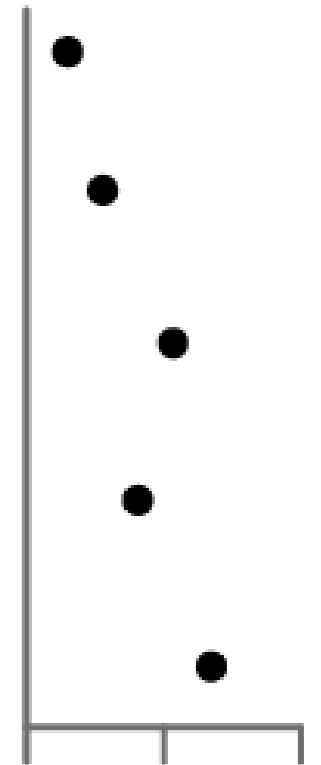
Length



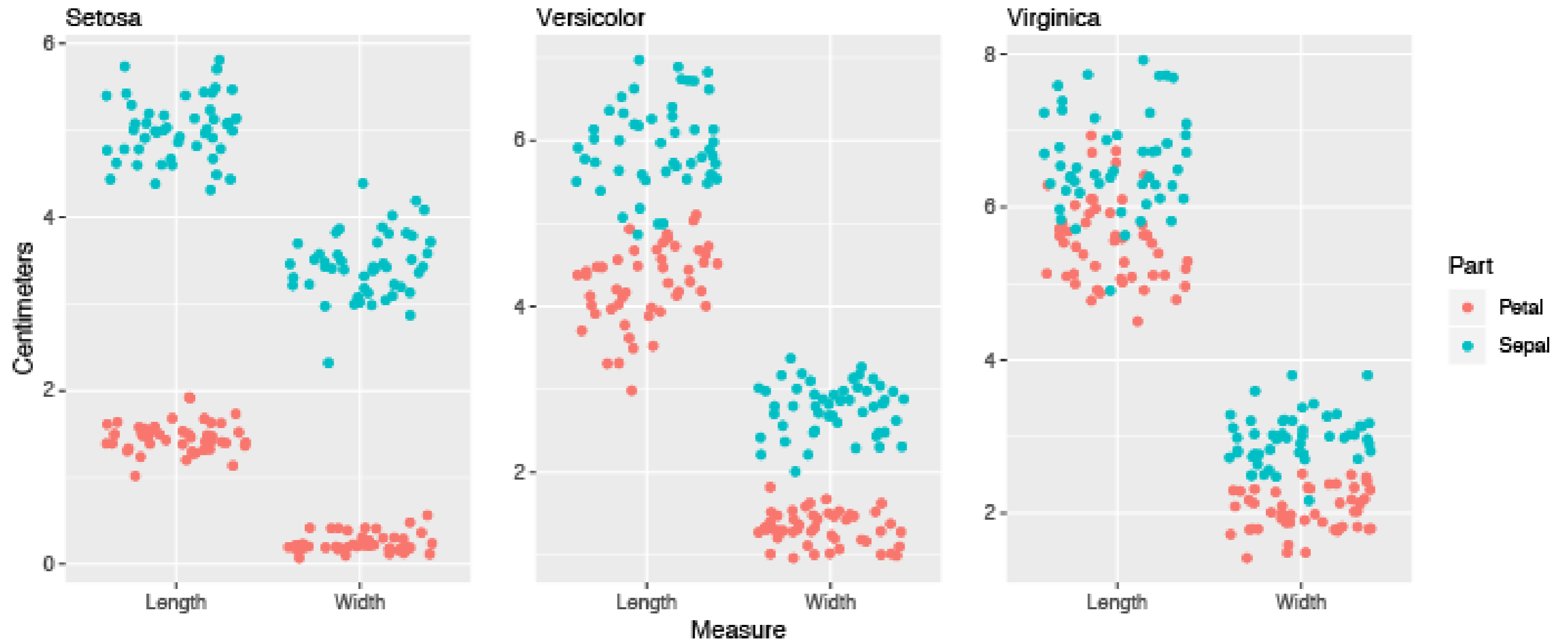
Position -  
on same, but  
unaligned, scale



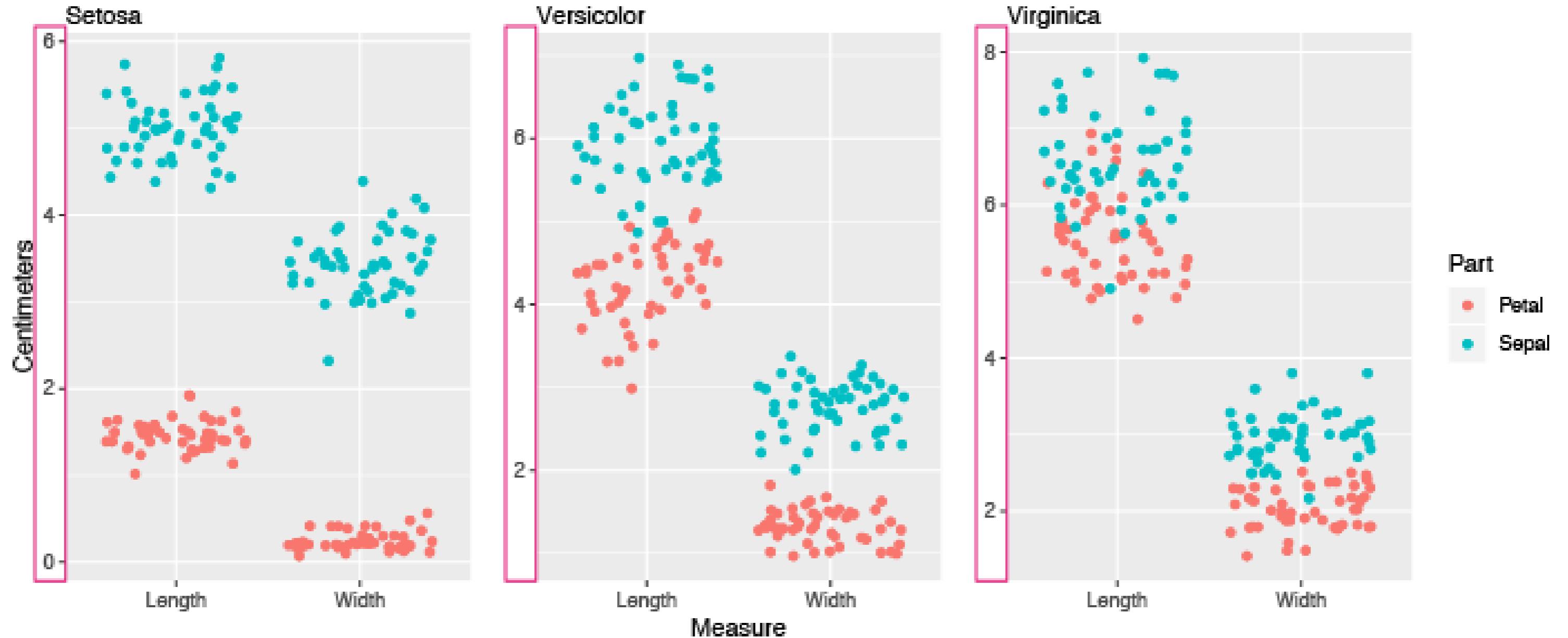
Position -  
on a  
common scale



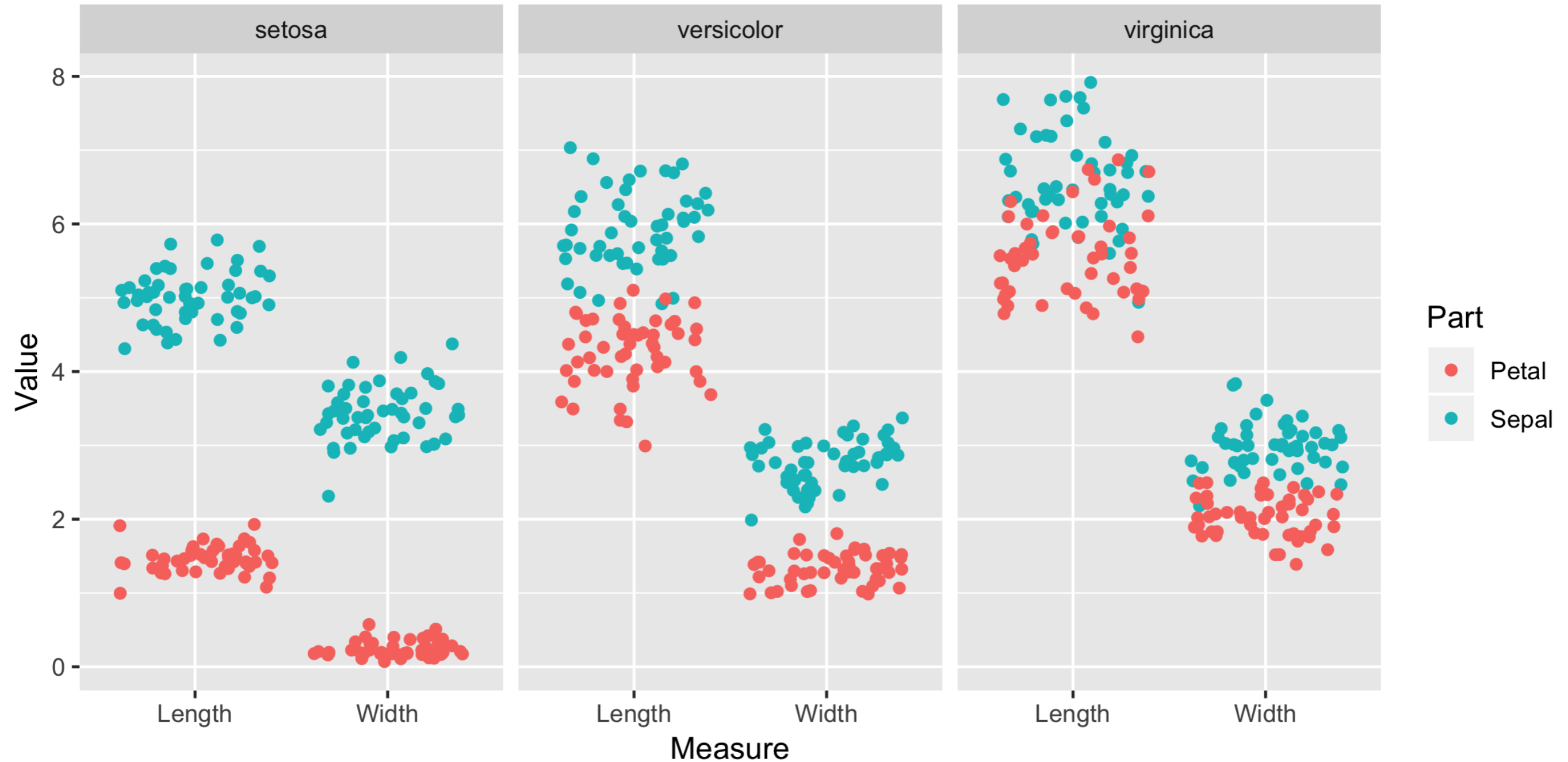
# Three iris scatter plots



# Three iris scatter plots, unaligned y-axes



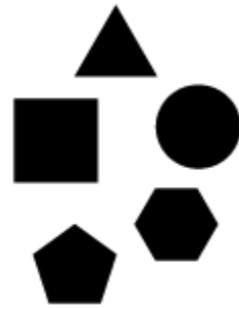
# Single faceted plot, common y-axis



# Efficiency in Decoding Separate Groups

Low  High

Filled Shapes



Sequential Colours



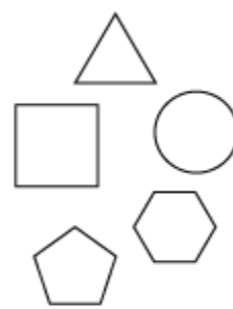
Qualitative Colours



Hatching



Shape Outlines



Labels

ANT1  
FRG2  
FRG1  
Gapdh  
DUX4

Line Width



Line Type

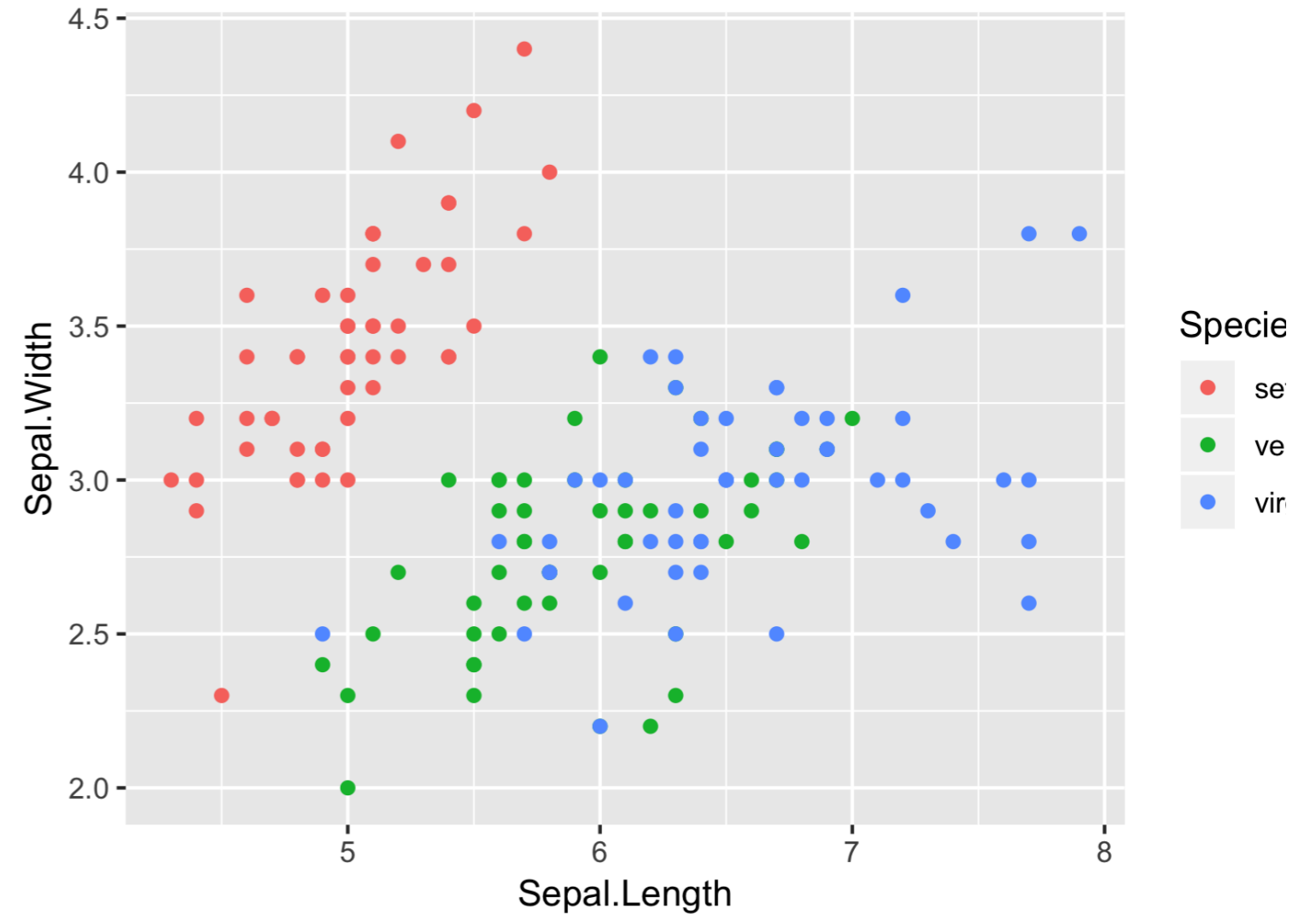


Line Colours



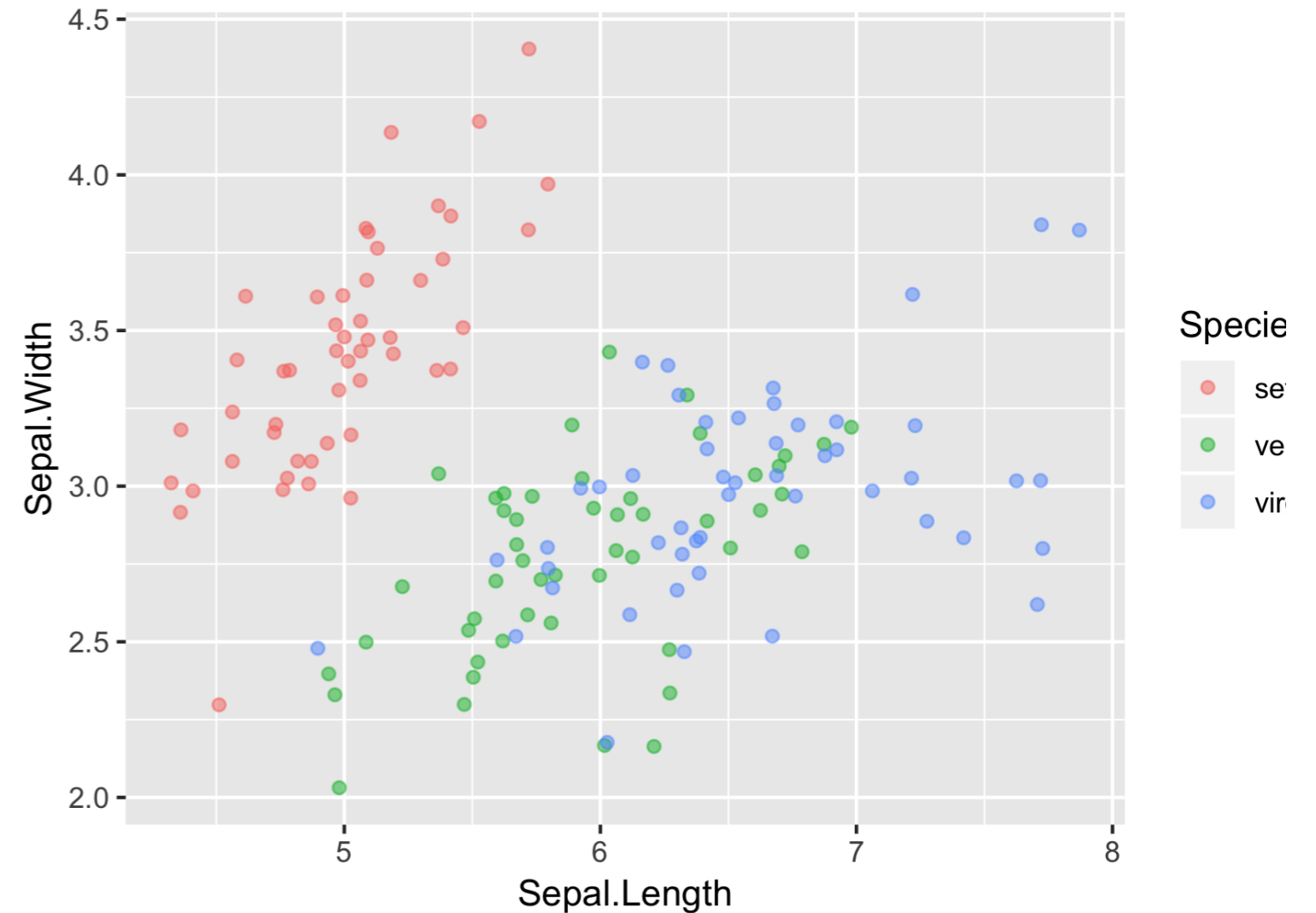
# Aesthetics - categorical variables

```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                col = Species)) +  
  geom_point()
```



# Aesthetics - categorical variables

```
ggplot(iris, aes(x = Sepal.Length,  
                y = Sepal.Width,  
                col = Species)) +  
  geom_point(position = "jitter",  
            alpha = 0.5)
```





# Now it's your turn

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