Style And Color

Overview

ggprism and **ggsci** can respectively set the style and color of the picture generated by ggplot. This style and color are used in a large number of papers.

Package

- library(ggprism)
- library(ggsci)

ggprism

- theme_prism()

Change the style of the graph, similar to the style of the Prism software.

- scale_color_prism(palette="colors")
- scale_fill_prism(palette="colors")
- scale_colour_prism(palette="colors")

There are many optional color schemes to change the color of the picture, which is very convenient.

- scale_shape_prism()

Shape scales that approximate those used in GraphPad Prism.

- scale_x_discrete(guide = "prism_bracket")

Segment the coordinate axis.

- add_pvalue(data_frame)

Add a hypothesis test p-value to the picture

ggsci

- scale_color_aaas()
- scale_color_d3()
- scale_color_npg()
- There are too many coloring scheme to write…

This function can directly call the color scheme of various scientific journals, which is very beautiful and very convenient to use.
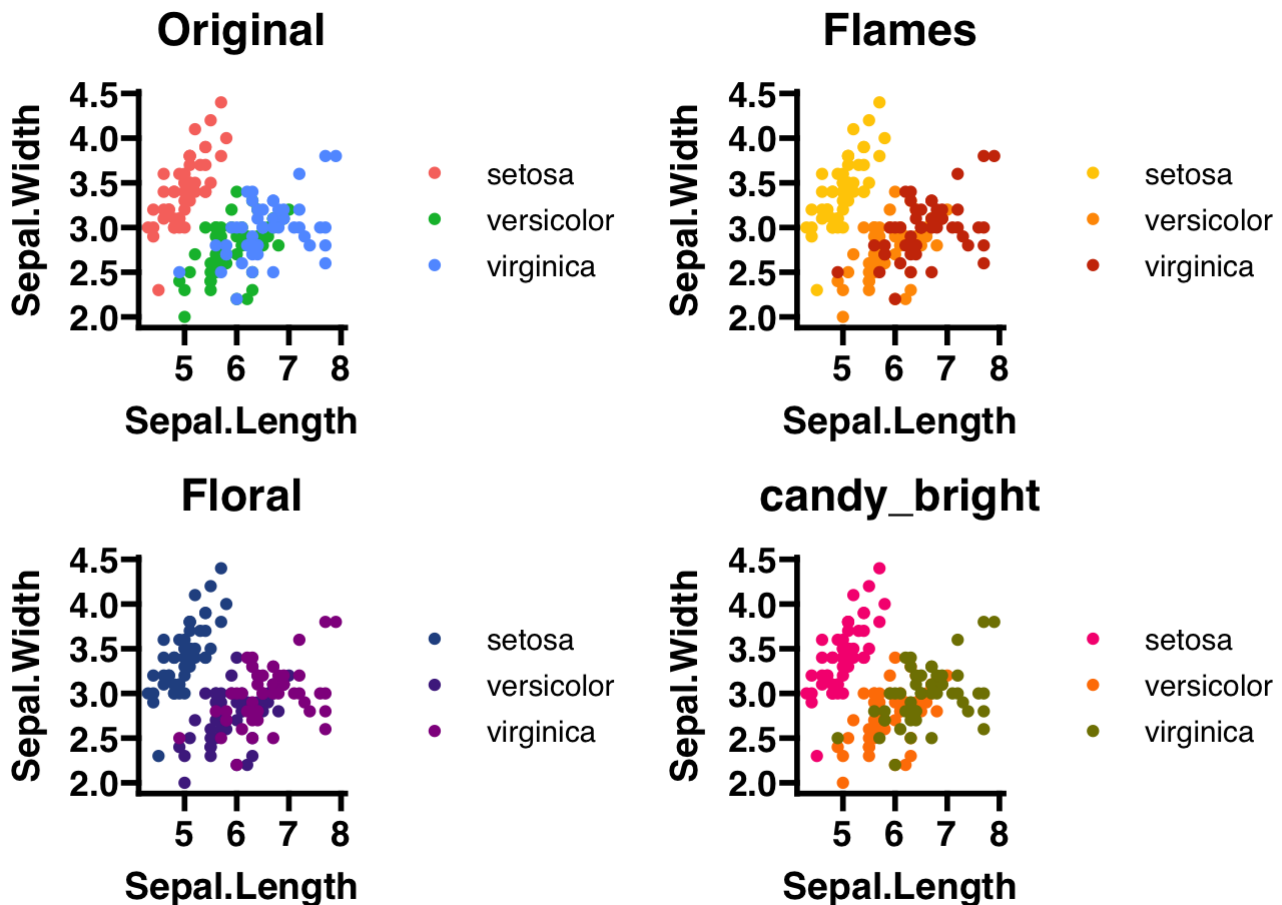
Example

The picture on the left does not use theme_prism, the picture on the right uses it.

```
p1 <- iris %>% ggplot(aes(Sepal.Length,Sepal.Width))+geom_point()
p2 <- iris %>% ggplot(aes(Sepal.Length,Sepal.Width))+geom_point()+theme_prism()
p1+p2
```
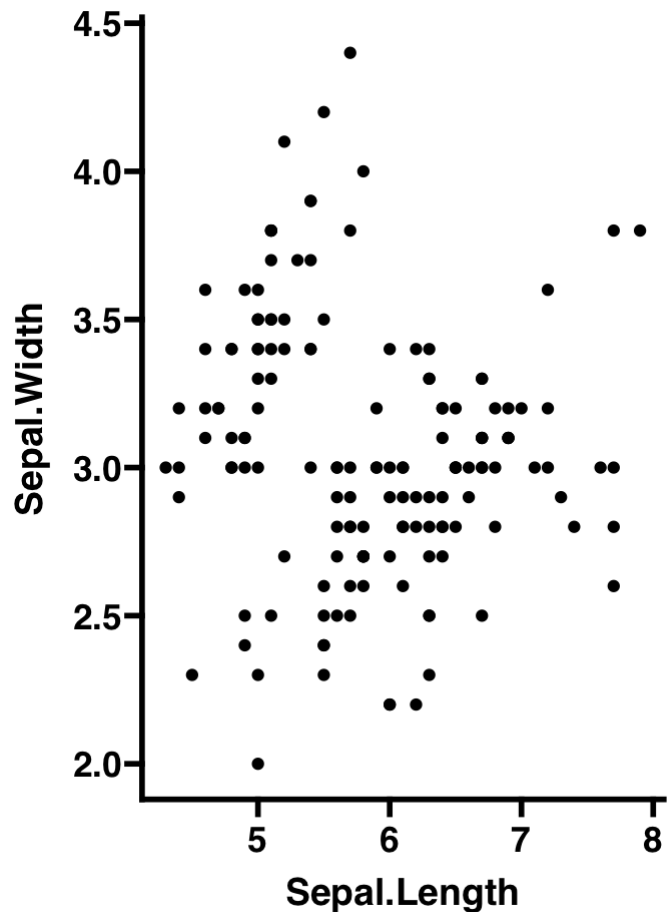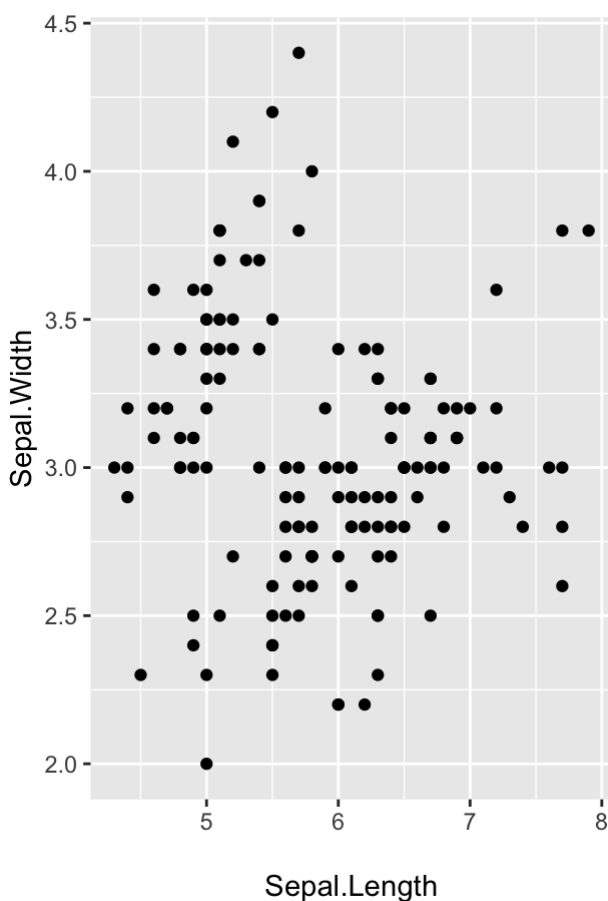
The default coordination of ggplot and the color scheme of ggprism are respectively shown. ggprism's high-contrast color matching is very suitable for scientific research articles.

```
p1 <- iris %>% ggplot(aes(Sepal.Length,Sepal.Width,color=Species))+geom_point()+theme
_prism()+ggtitle("Original")
p2 <- iris %>% ggplot(aes(Sepal.Length,Sepal.Width,color=Species))+geom_point()+theme
_prism()+scale_color_prism(palette = "flames")+ggtitle("Flames")
p3 <- iris %>% ggplot(aes(Sepal.Length,Sepal.Width,color=Species))+geom_point()+theme
_prism()+scale_color_prism(palette = "floral")+ggtitle("Floral")
p4 <- iris %>% ggplot(aes(Sepal.Length,Sepal.Width,color=Species))+geom_point()+theme
_prism()+scale_color_prism(palette = "candy_bright")+ggtitle("candy_bright")
(p1+p2)/(p3+p4)
```


The image above does not use the x-axis segmentation, and the image below uses the x-axis segmentation.

```
p1 <- iris %>% ggplot(aes(Species,Sepal.Length))+geom_jitter(aes(shape=Species))+them
e_prism()
p2 <- iris %>% ggplot(aes(Species,Sepal.Length))+geom_jitter(aes(shape=Species))+them
e_prism()+scale_x_discrete(guide = "prism_bracket")+scale_shape_prism()
p1/p2
```
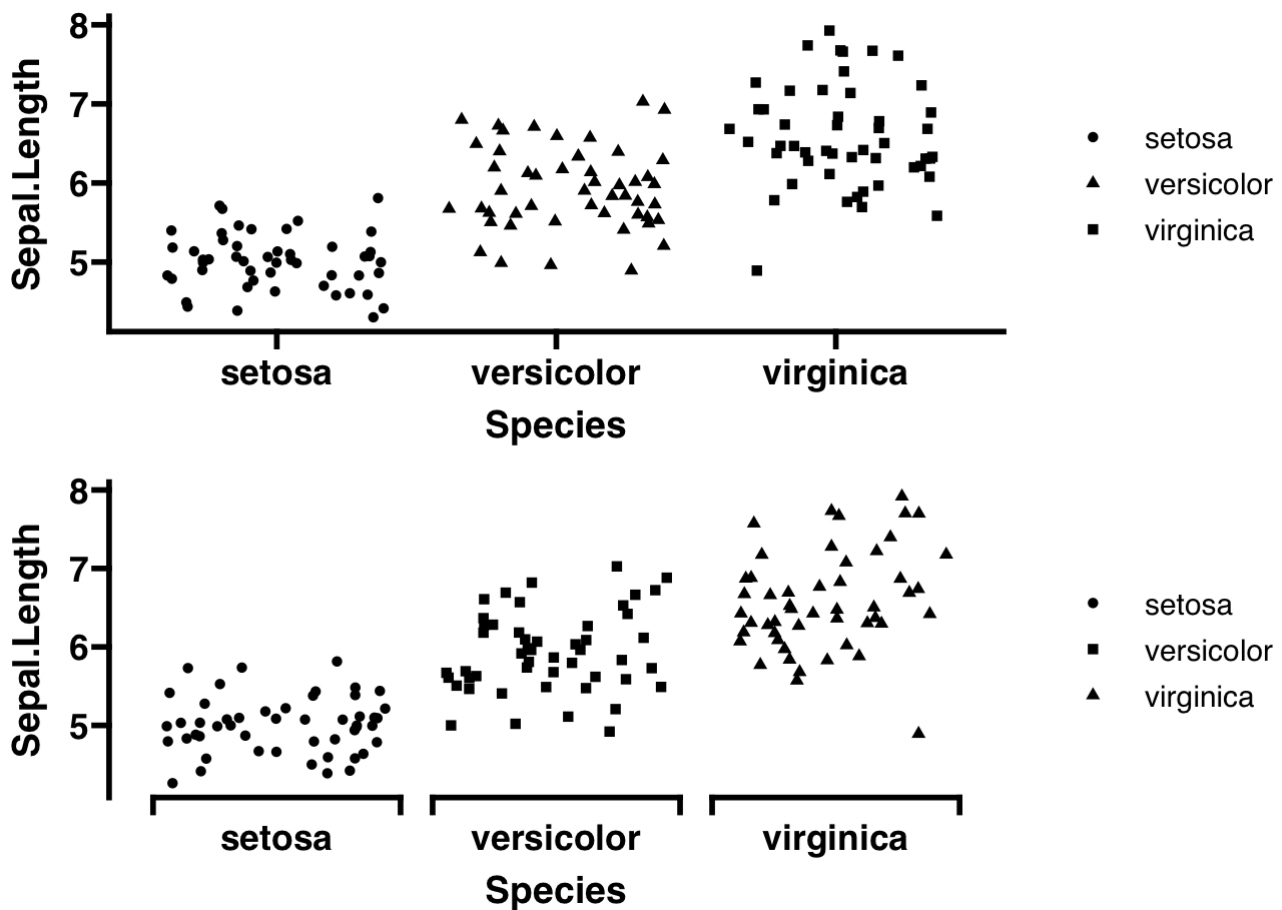
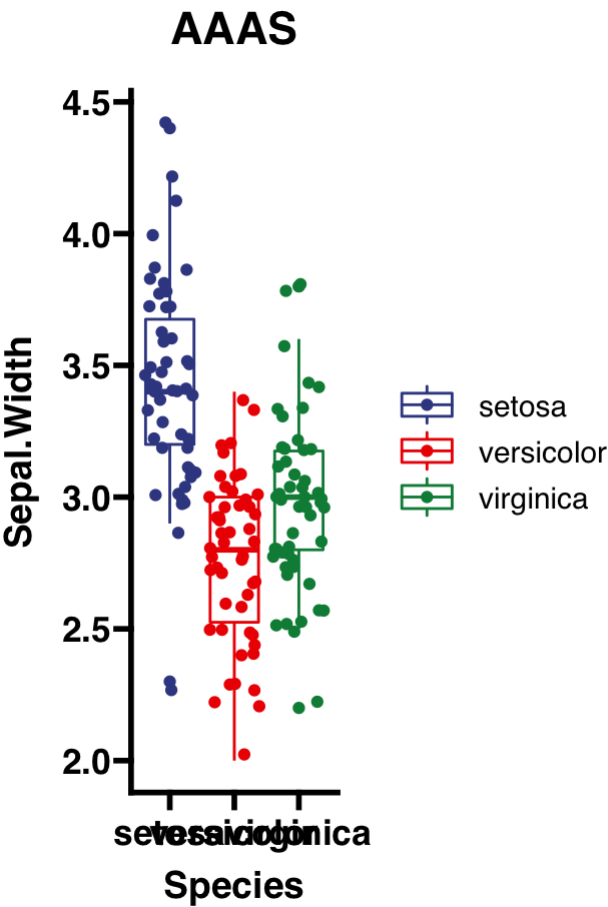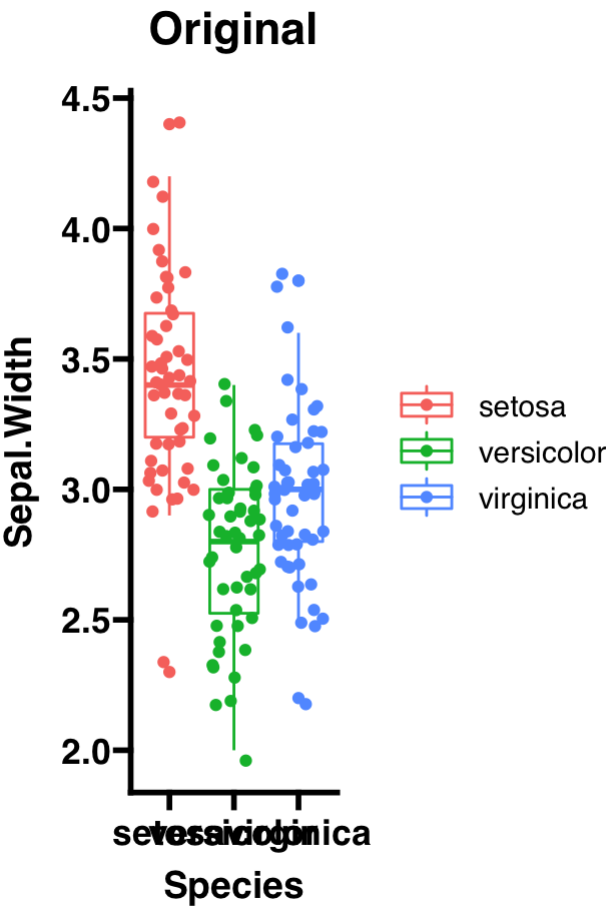The picture on the right has pvalue added, and the picture on the left does not.

```
p1 <- iris %>% filter(Species %in% c("virginica","versicolor")) %>% ggplot(aes(Specie
s,Sepal.Length,fill=Species))+geom_boxplot()+theme_prism()+scale_fill_prism("floral")
+theme(legend.position = "none")
two.means <- tibble::tribble(
  ~group1, ~group2, ~p,       ~y.position,
  "virginica",    "versicolor",    1.866e-07, 36
)
p <- data.frame(
  group1="virginica",
  group2="versicolor",
  p.adj = 1.866e-07,
  y.position = 8
)

p2 <-  iris %>% filter(Species %in% c("virginica","versicolor")) %>% ggplot(aes(facto
r(Species),Sepal.Length))+geom_boxplot(aes(fill=Species))+theme_prism()+scale_fill_pr
ism("floral")+theme(legend.position = "none")+add_pvalue(data=p)
p1+p2
```
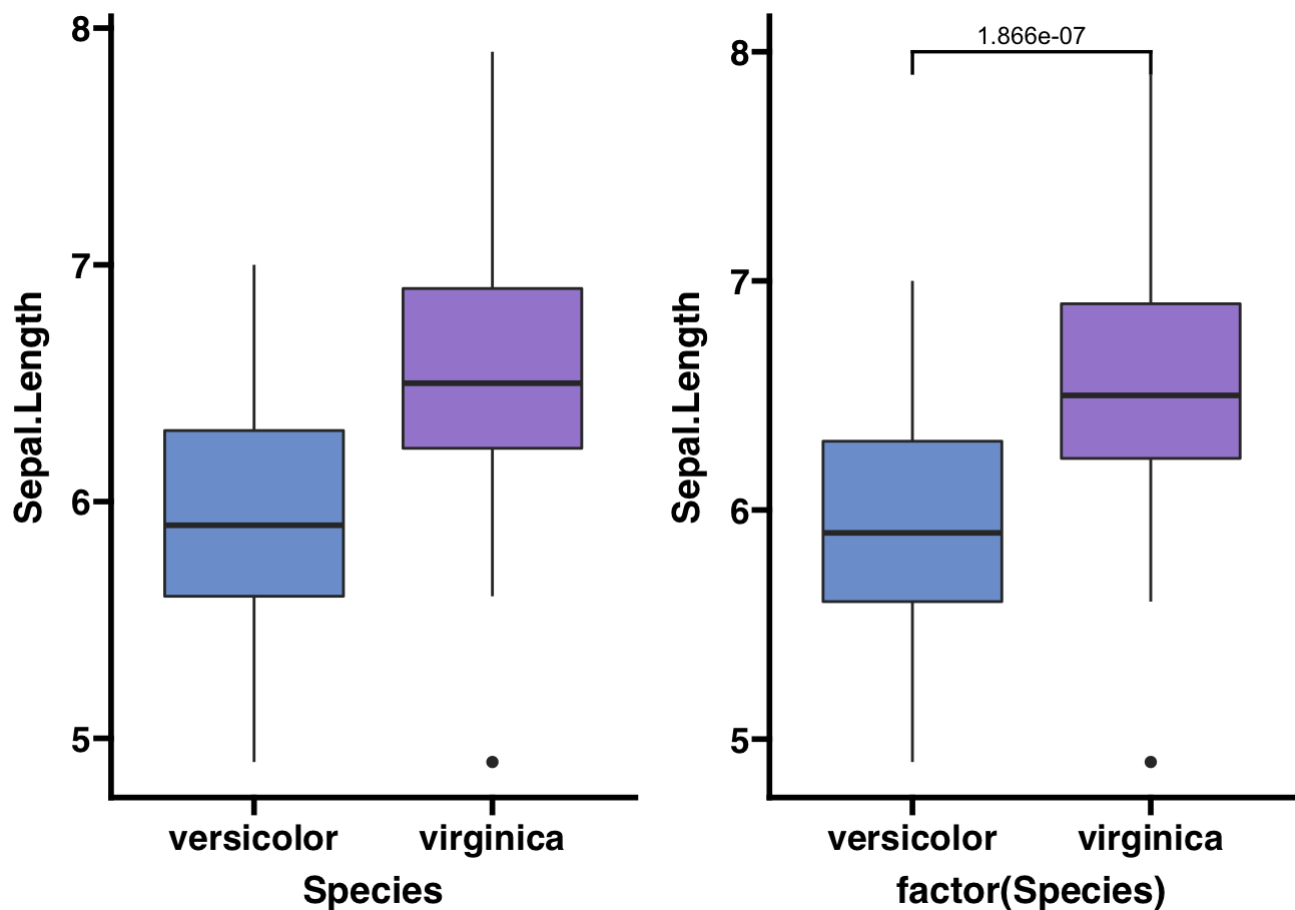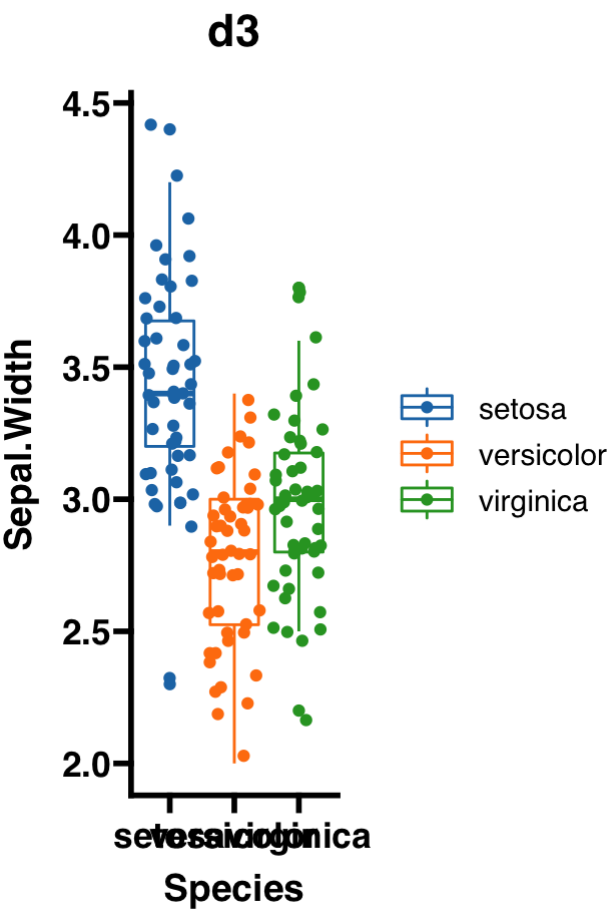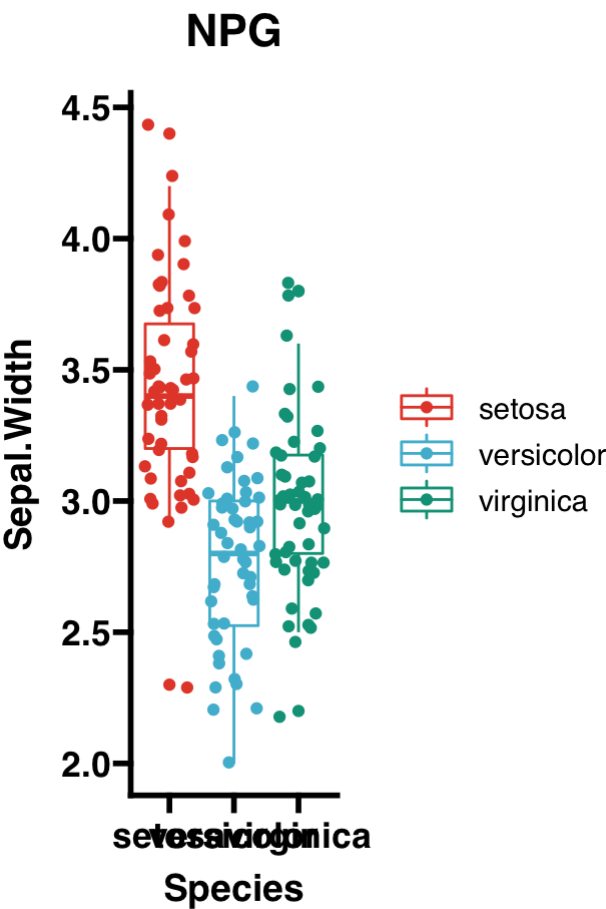
The various color schemes of ggsci are shown below.

```
p1 <- iris %>% ggplot(aes(Species,Sepal.Width,color=Species))+geom_boxplot()+geom_jit
ter()+theme_prism()+ggtitle("Original")
p2 <-  iris %>% ggplot(aes(Species,Sepal.Width,color=Species))+geom_boxplot()+geom_ji
tter()+theme_prism()+scale_color_aaas()+ggtitle("AAAS")
p3 <- iris %>% ggplot(aes(Species,Sepal.Width,color=Species))+geom_boxplot()+geom_jit
ter()+theme_prism()+scale_color_npg()+ggtitle("NPG")
p4 <- iris %>% ggplot(aes(Species,Sepal.Width,color=Species))+geom_boxplot()+geom_jit
ter()+theme_prism()+scale_color_d3()+ggtitle("d3")
p5 <- iris %>% ggplot(aes(Species,Sepal.Width,color=Species))+geom_boxplot()+geom_jit
ter()+theme_prism()+scale_color_jama()+ggtitle("JAMA")
p6 <- iris %>% ggplot(aes(Species,Sepal.Width,color=Species))+geom_boxplot()+geom_jit
ter()+theme_prism()+scale_color_lancet()+ggtitle("Lancet")
(p1+p2)
```

Original

AAAS


```
(p3+p4)
```

NPG

d3

```
(p5+p6)
```