Customize your traces

INTERACTIVE DATA VISUALIZATION WITH PLOTLY IN R



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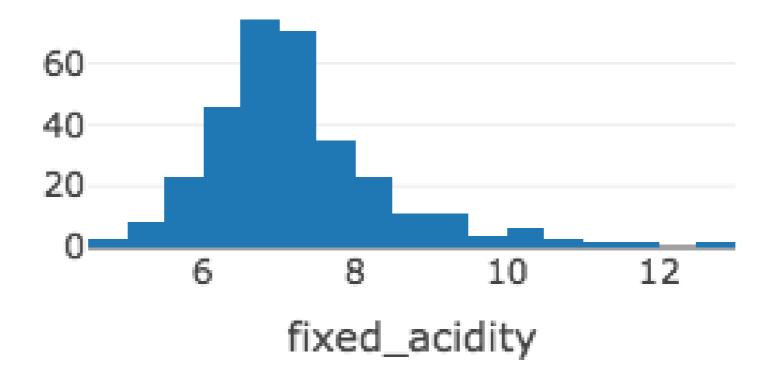
Wine quality data

glimpse(winequality)

```
Observations: 325
Variables: 14
$ type
                 <chr> "red", "red", "red", "red", "red", "...
$ fixed_acidity
                 <dbl> 8.2, 8.2, 8.0, 10.2, 8.6, 6.1, 10.7, 9.1, 7.2...
$ volatile_acidity <dbl> 0.885, 0.640, 0.715, 0.360, 0.520, 0.590, 0.6...
$ citric_acid
               <dbl> 0.20, 0.27, 0.22, 0.64, 0.38, 0.01, 0.22, 0.3...
$ residual_sugar <dbl> 1.40, 2.00, 2.30, 2.90, 1.50, 2.10, 2.70, 2.1...
$ sulphates
                  <dbl> 0.46, 0.62, 0.54, 0.66, 0.52, 0.56, 0.98, 0.8...
$ alcohol
                  <dbl> 10.0, 9.1, 9.5, 12.5, 9.4, 11.4, 9.9, 11.2, 1...
$ quality
                  <int> 5, 6, 6, 6, 5, 5, 6, 6, 6, 7, 6, 5, 4, 6, 6, ...
$ quality_label
                 <chr> "low", "medium", "medium", "medium", "low", ...
```

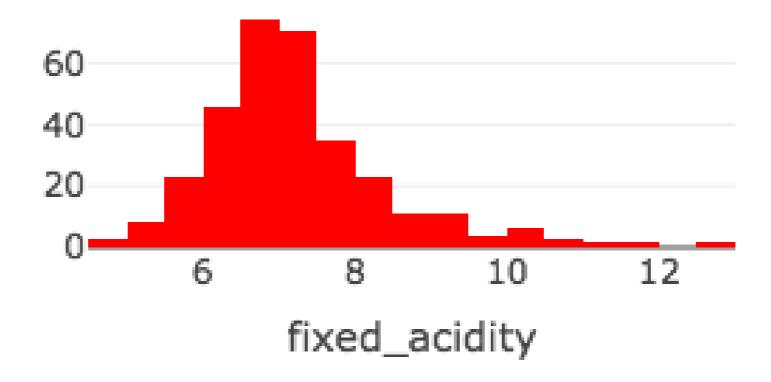


Color



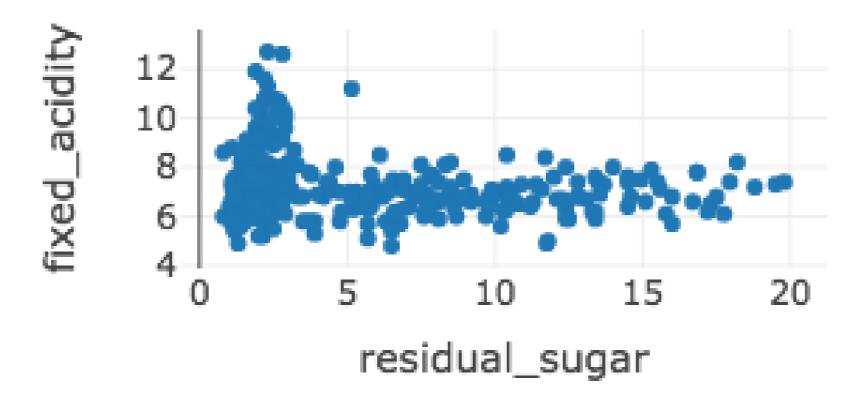
```
winequality %>%
  plot_ly(x = ~fixed_acidity) %>%
  add_histogram()
```

Color



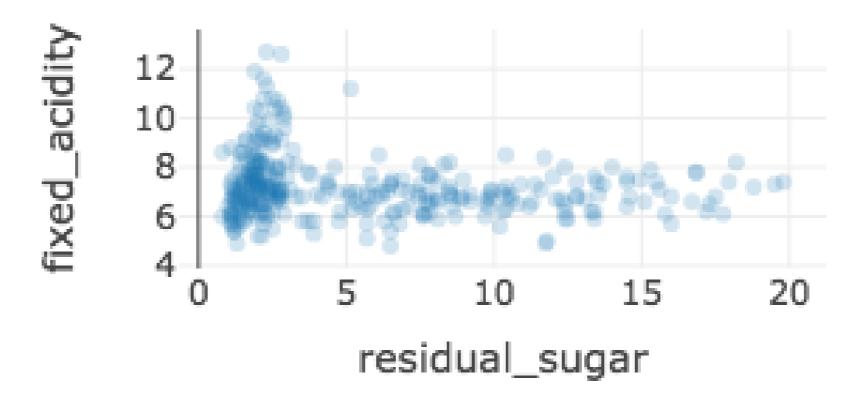
```
winequality %>%
  plot_ly(x = ~fixed_acidity) %>%
  add_histogram(color = I("red")) # Setting color
```

Opacity



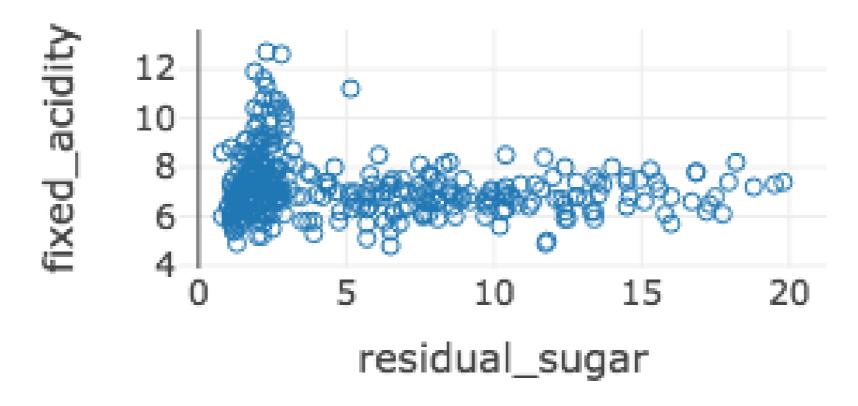
```
winequality %>%
 plot_ly(x = ~residual_sugar, y = ~fixed_acidity) %>%
 add_markers()
```

Opacity



```
winequality %>%
 plot_ly(x = ~residual_sugar, y = ~fixed_acidity) %>%
 add_markers(marker = list(opacity = 0.2)) # Adjust opacity
```

Symbols



```
winequality %>%
  plot_ly(x = ~residual_sugar, y = ~fixed_acidity) %>%
  add_markers(marker = list(symbol = "circle-open")) # Change symbol
```

Marker options

- opacity
- color
- symbol (scatter/box)
- size (scatter)
- width (bar/histogram)

Let's practice!

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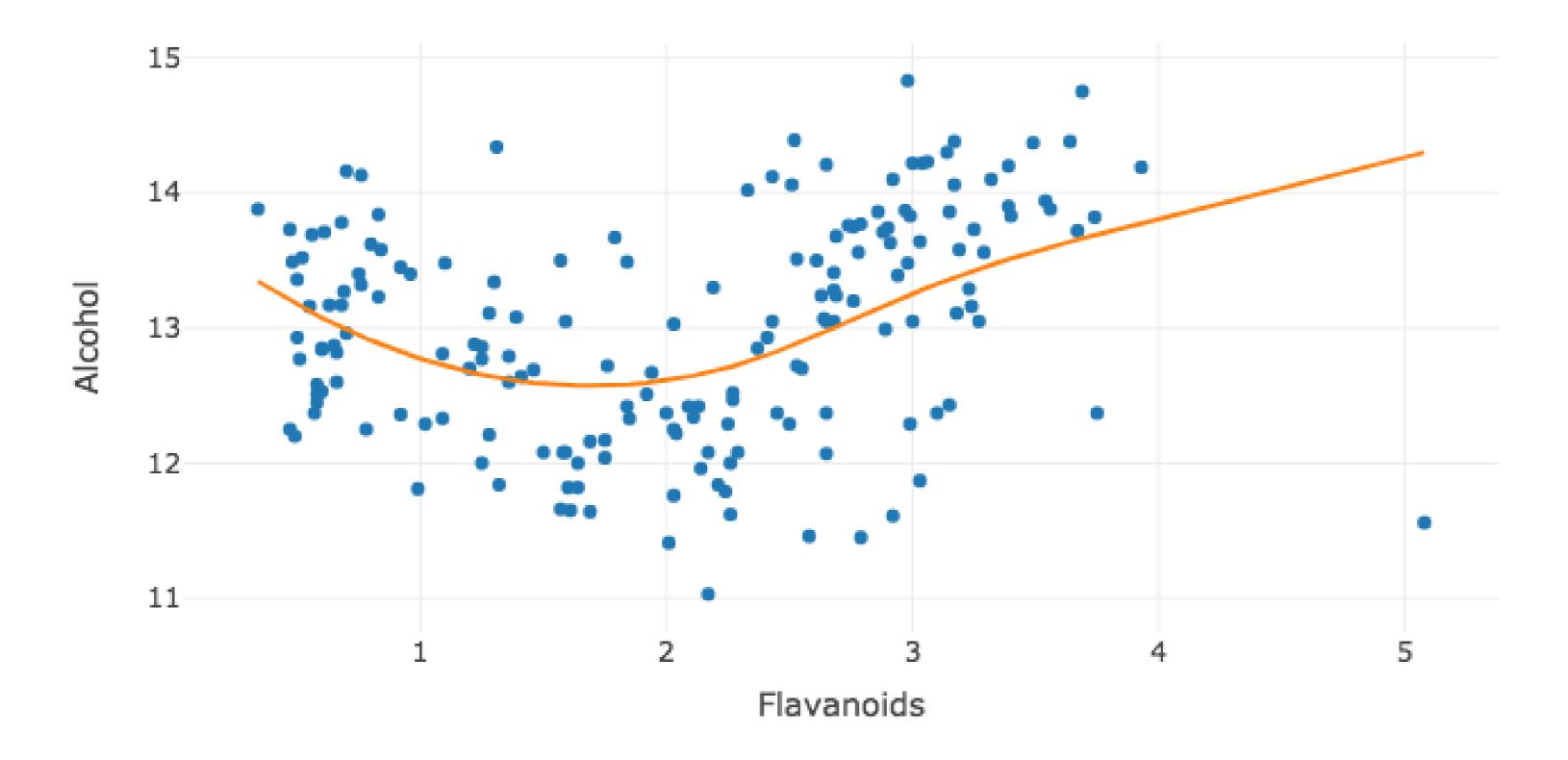
Thoughtful use of color

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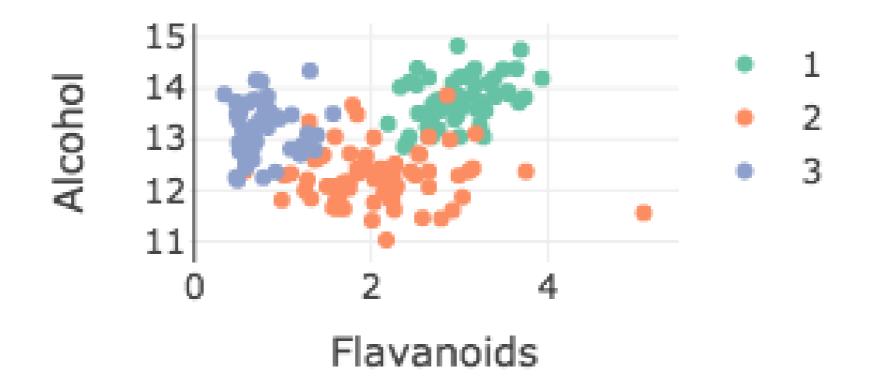


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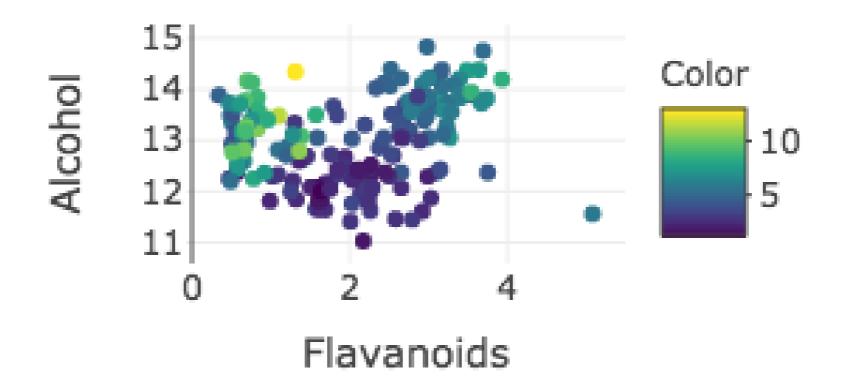


Adding a third variable



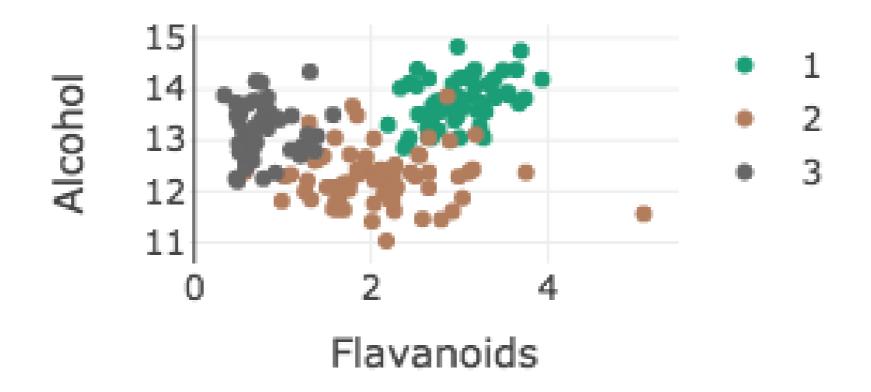
```
wine %>%
 plot_ly(x = ~Flavanoids, y = ~Alcohol, color = ~Type) %>%
 add_markers()
```

Adding a quantitative variable



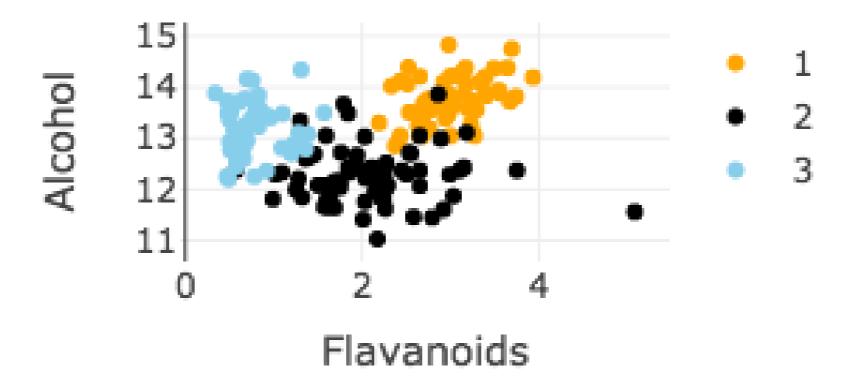
```
wine %>%
 plot_ly(x = ~Flavanoids, y = ~Alcohol, color = ~Color) %>%
 add_markers()
```

RColorBrewer palettes



```
wine %>%
 plot_ly(x = ~Flavanoids, y = ~Alcohol, color = ~Type) %>%
 add_markers(colors = "Dark2")
```

Manual palettes



```
wine %>%
  plot_ly(x = ~Flavanoids, y = ~Alcohol, color = ~Type) %>%

add_markers(colors = c("orange", "black", "skyblue"))
```



Let's practice!

INTERACTIVE DATA VISUALIZATION WITH PLOTLY IN R



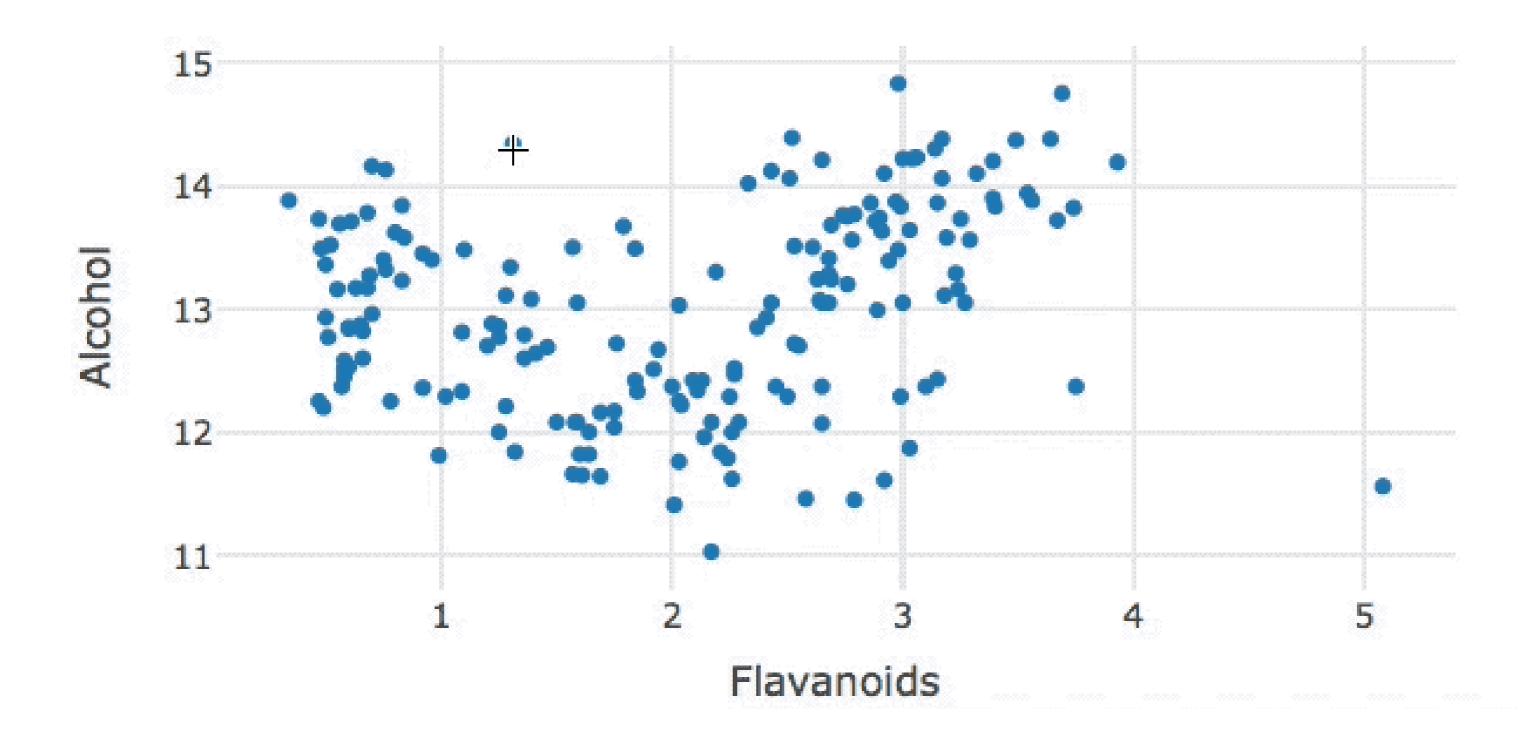
Labeling your data

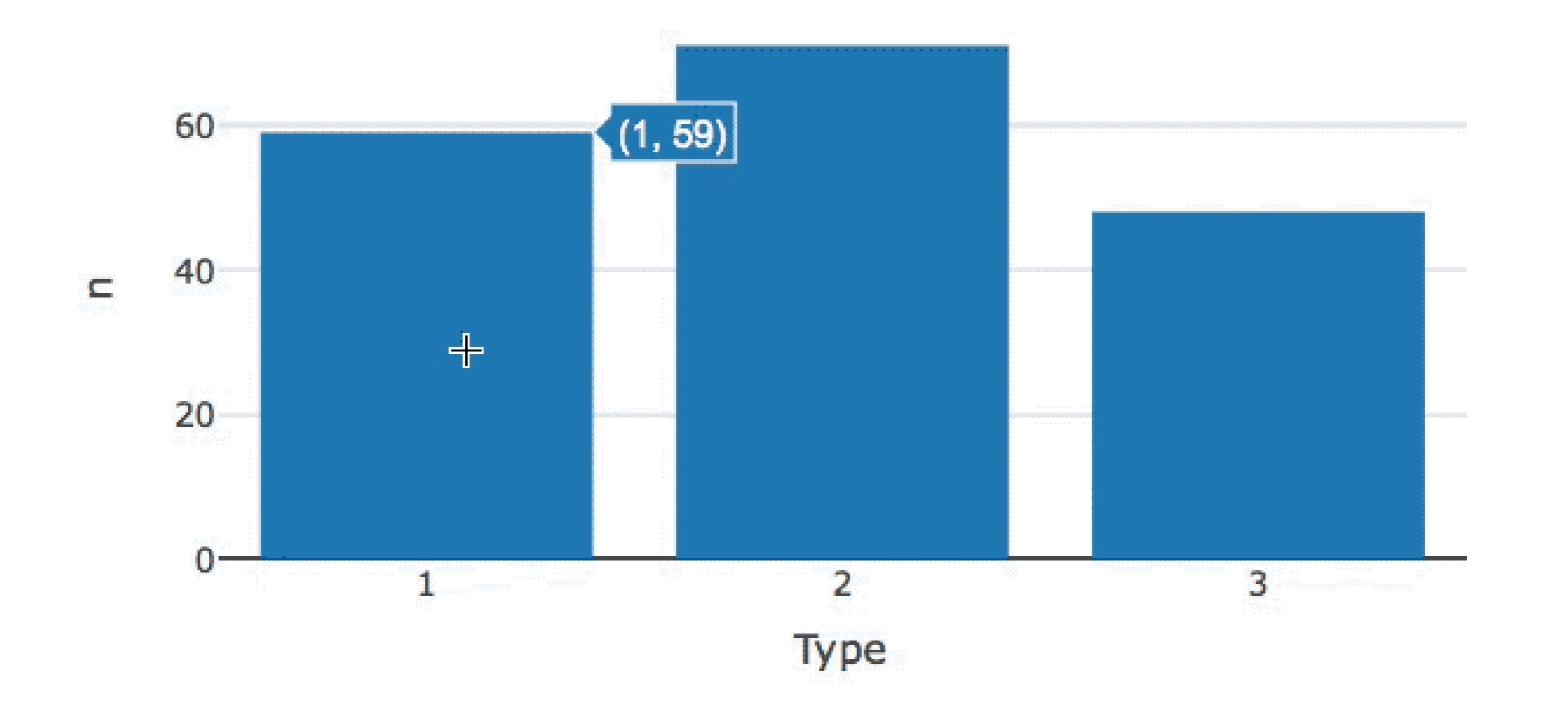
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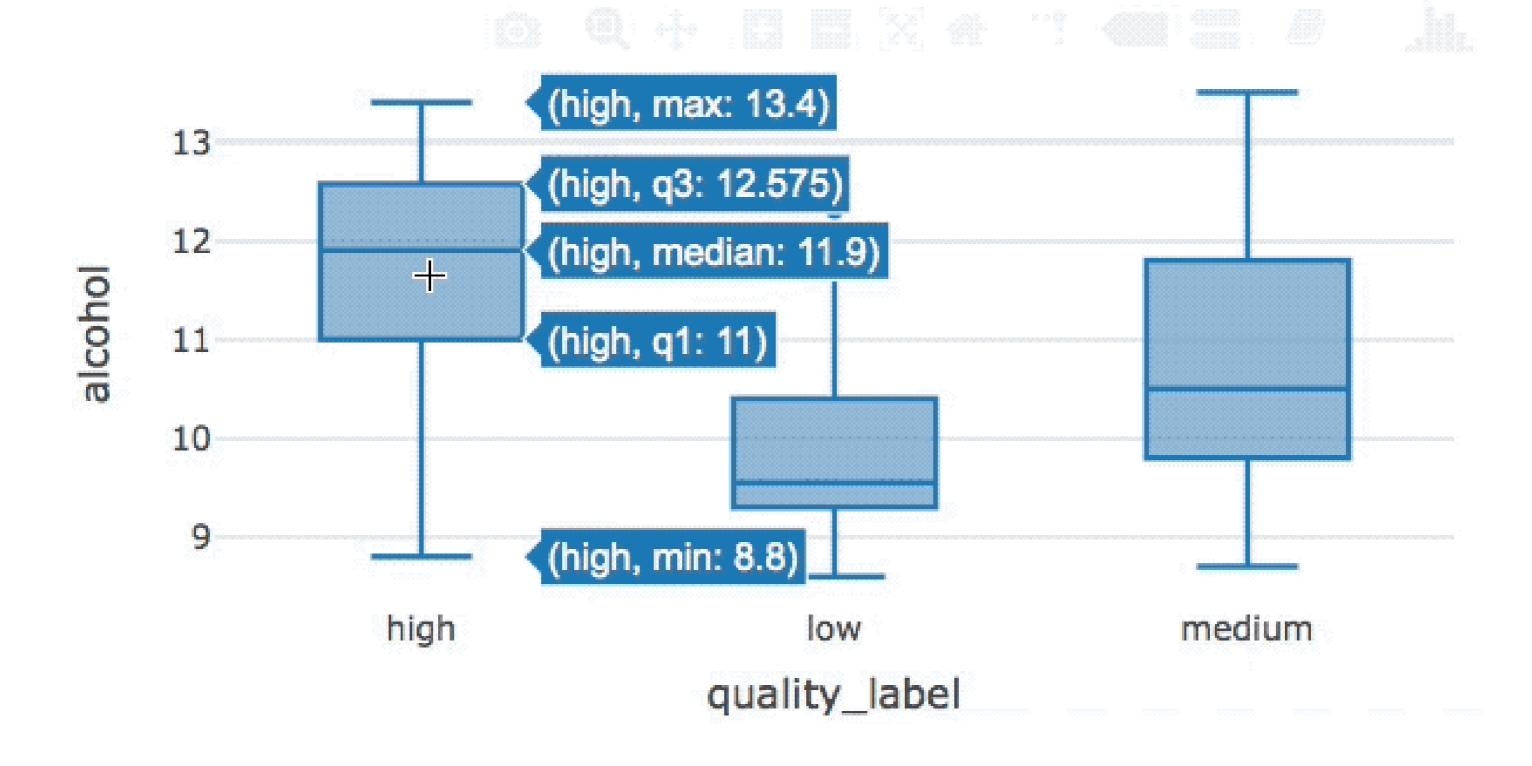


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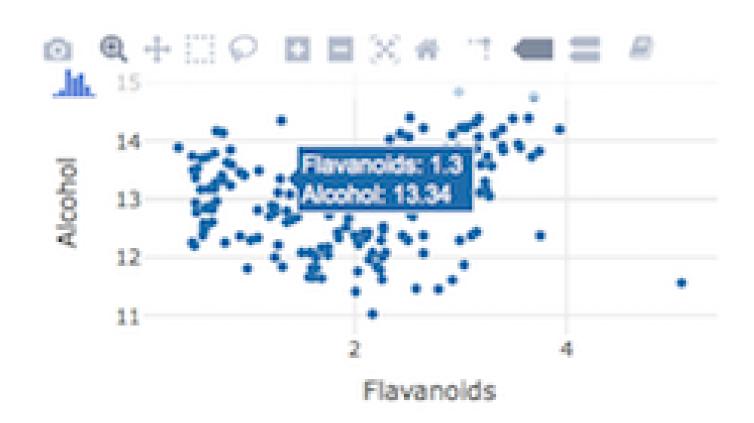
Changing the default

hoverinfo =

- "all"
- "X"
- "y"
- "x+y"
- "x+y+z"

Custom hover text

• tilde, ~ , to map columns to aesthetic parameters



Let's practice!

INTERACTIVE DATA VISUALIZATION WITH PLOTLY IN R



Customizing your layout

INTERACTIVE DATA VISUALIZATION WITH PLOTLY IN R



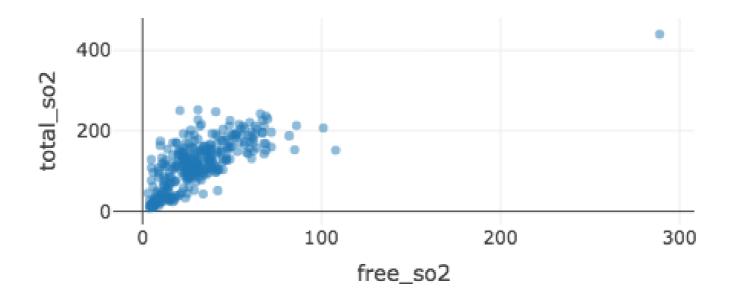
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layout()

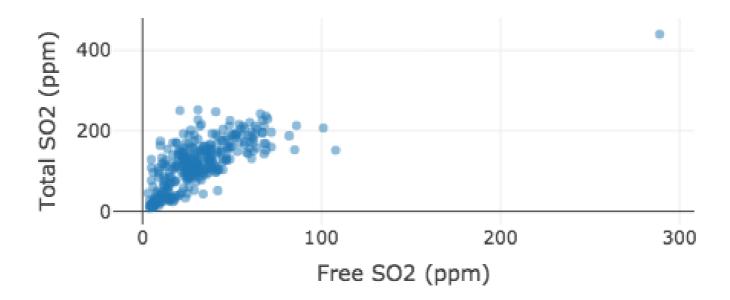
- axes: type, labels, tick marks, transformations, etc.
- **legend:** position
- canvas: grid lines, background color
- **size**: height, width, margins

Axis labels



```
winequality %>%
  plot_ly(x = ~free_so2, y = ~total_so2) %>%
  add_markers(marker = list(opacity = 0.2))
```

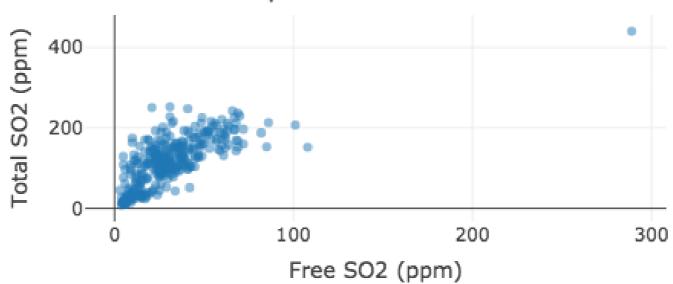
Axis labels



```
winequality %>%
  plot_ly(x = ~free_so2, y = ~total_so2) %>%
  add_markers(marker = list(opacity = 0.2)) %>%
  layout(xaxis = list(title = "Free SO2 (ppm)"),
     yaxis = list(title = "Total SO2 (ppm)"))
```

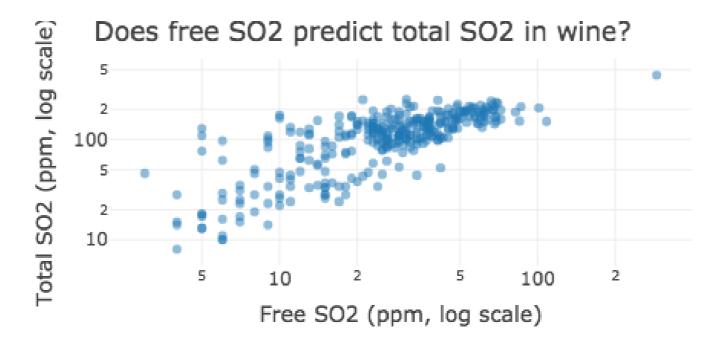
Titles

Does free SO2 predict total SO2 in wine?



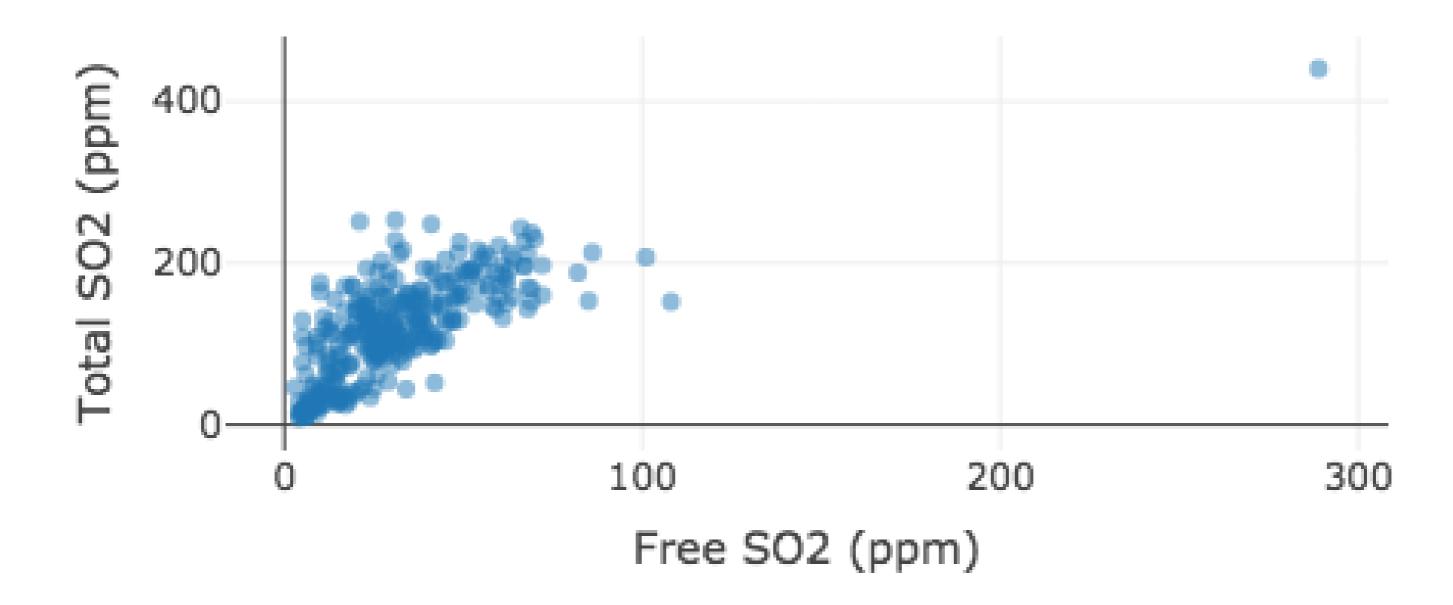
```
winequality %>%
  plot_ly(x = ~free_so2, y = ~total_so2) %>%
  add_markers(marker = list(opacity = 0.2)) %>%
  layout(xaxis = list(title = "Free SO2 (ppm)"),
      yaxis = list(title = "Total SO2 (ppm)"),
      title = "Does free SO2 predict total SO2 in wine?")
```

Transforming axes

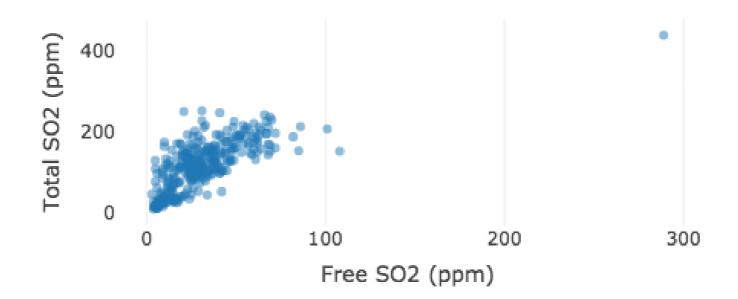


```
winequality %>%
  plot_ly(x = ~free_so2, y = ~total_so2) %>%
  add_markers(marker = list(opacity = 0.2)) %>%
  layout(xaxis = list(title = "Free SO2 (ppm, log scale)", type = "log"),
       yaxis = list(title = "Total SO2 (ppm, log scale)", type = "log"),
       title = "Does free SO2 predict total SO2 in wine?")
```

Customizing the grid

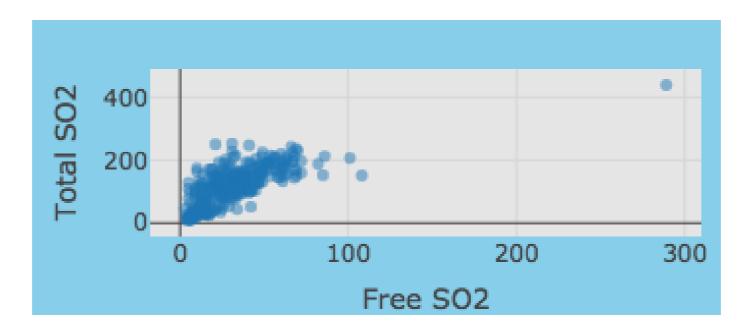


Customizing the grid



```
winequality %>%
  plot_ly(x = ~free_so2, y = ~total_so2) %>%
  add_markers(marker = list(opacity = 0.5)) %>%
  layout(xaxis = list(title = "Free SO2 (ppm)", zeroline = FALSE),
      yaxis = list(title = "Total SO2 (ppm)", zeroline = FALSE,
      showgrid = FALSE))
```

Customizing the canvas



```
plot_bgcolor = toRGB("gray90"),
paper_bgcolor = toRGB("skyblue"))
```



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