# Data Visualization with ggplot2

GITAM
(DEEMED TO BE UNIVERSITY)
(Estd. u/s 3 of the UGC Act, 1956)
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**Step1: Creating ggplot:**

- You begin a plot with the function **ggplot()**.It creates a coordinate system that you can add layers to.
- The first argument of ggplot() is the dataset to use in the graph. So **ggplot(data = iris)** creates an empty graph.
- Now complete your graph by adding one or more layers to ggplot().
- The function **geom_point()** adds a layer of points to your plot, which creates a scatterplot.
- **ggplot2** comes with many geom functions that each add a different type of layer to a plot.



- Each geom function in **ggplot2** takes a mapping argument. This defines how variables in your dataset are mapped to visual properties.
- The mapping argument is always paired with **aes(),** and the x and y arguments of aes() specify which variables to map to the x and y-axes.
- **ggplot2** looks for the mapped variable in the data argument, in this case, iris.

**Step2: Graphing Template**

- For making graphs with **ggplot2** the following code with a dataset, a geom function, or a collection of mappings and extend this template to make different types of graphs:

**ggplot(data = <DATA>) +**
**<GEOM_FUNCTION>(mapping = aes(<MAPPINGS>))**



## Aesthetic Mappings

- Graph plotting in R is of two types:

### 1. One-dimensional Plotting:

➢ In one-dimensional plotting, we plot one variable at a time.

➢ For example, we may plot a variable with the number of times each of its values occurred in the entire dataset (frequency).

➢ So, it is not compared to any other variable of the dataset. These are the 4 major types of graphs that are used for One-dimensional analysis –

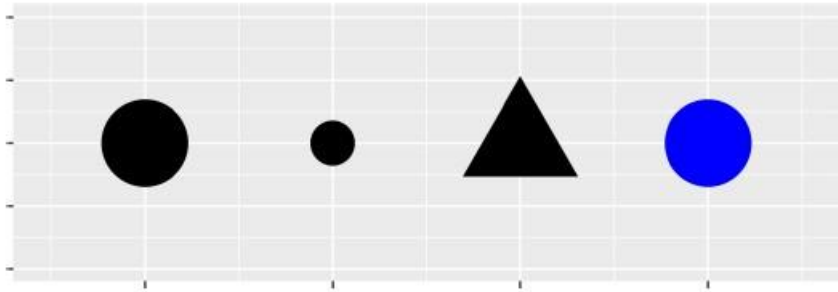
- Five Point Summary
- Box Plotting
- Histograms
- Bar Plotting



## 2. Two-dimensional Plotting:

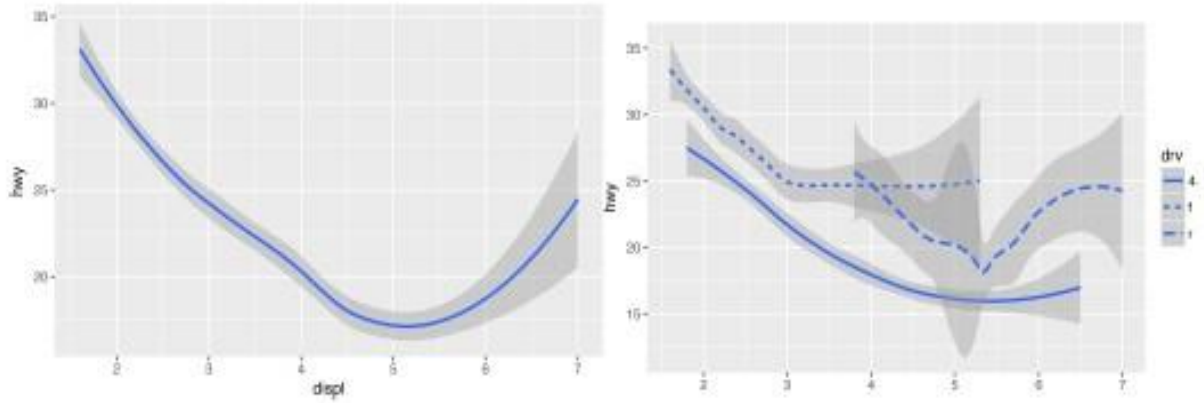
- In two-dimensional plotting, we visualize and compare one variable with respect to the other.
- For example, in Air Quality dataset, we would like to compare how the AQI varies with the temperature at a particular place.
- Temperature and AQI are two different variables and we wish to see how one changes with respect to the other.
- These are the 3 major kinds of graphs used for such kinds of analysis
  - Box Plotting
  - Histograms
  - Scatter plots

- You can add a third variable, like class, to a two-dimensional scatterplot by mapping it to an *aesthetic*.
- An aesthetic is a visual property of the objects in your plot.
- Aesthetics include things like the **size**, the **shape,** or the **color** of your points. You can display a point in different ways by changing the values of its aesthetic properties.
- By using these aesthetic properties we change the levels of a point's size, shape, and color to make the point small, triangular, or blue:



# Geometric Objects

- A *geom* is the geometrical object that a plot uses to represent data.
- Bar charts use **bar geoms**, line charts use **line geoms**, boxplots use **boxplot geoms**, and so on. Scatterplots break the trend; they use the **point geom**.
- You can use different geoms to plot the same data. To change the geom in your plot, change the geom function that you add to ggplot().
- ggplot() is providing two such geom's- **geom_point()** and **geom_smooth().**



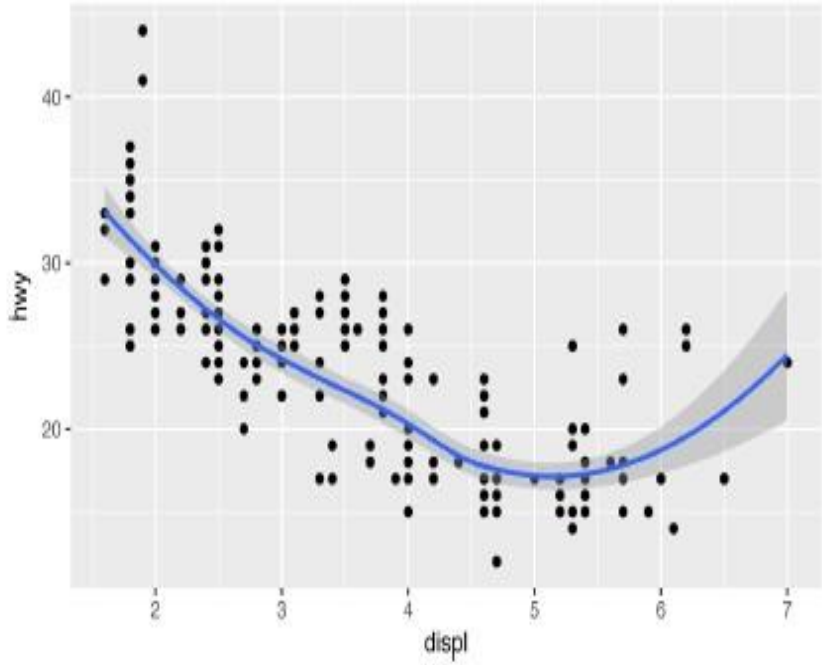
- Both plots contain the same x variable and the same y variable, and both describe the same data. But the plots are not identical.
- Each plot uses a different visual object to represent the data. **ggplot2** use different *geoms*.

```
# left
ggplot(data = mpg) +
  geom_point(mapping = aes(x = displ, y = hwy))

# right
ggplot(data = mpg) +
  geom_smooth(mapping = aes(x = displ, y = hwy))
```

- geom_smooth() will draw a different line, with a different linetype, for each unique value of the variable that you map to **linetype =.**
- To display multiple geoms in the same plot, add multiple geom functions to ggplot():

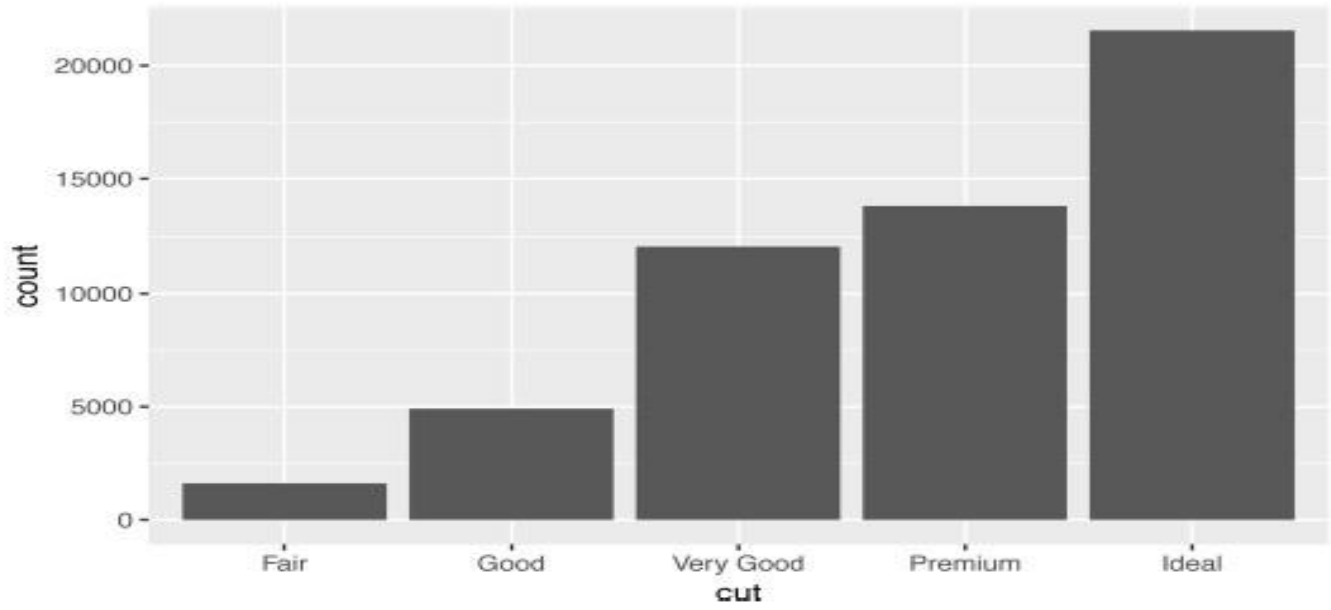
```
ggplot(data = mpg) +
geom_point(mapping = aes(x = displ, y = hwy)) +
geom_smooth(mapping = aes(x = displ, y = hwy))
```



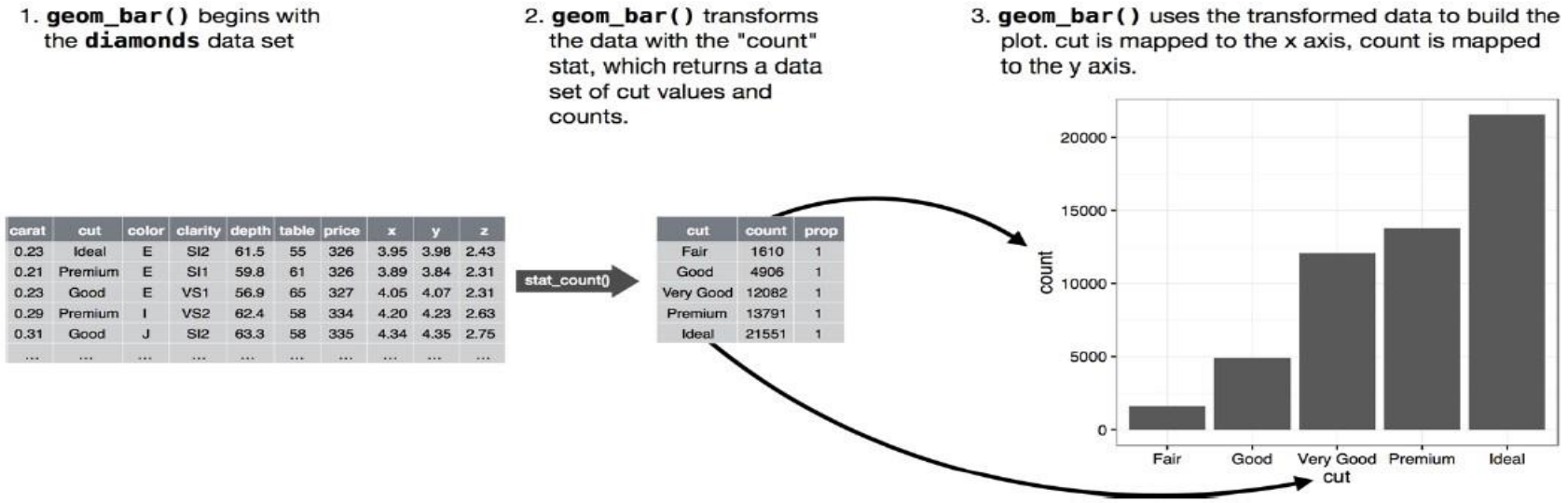
# Statistical Transformations

- Bar charts seem simple, but they are interesting because they reveal something subtle about plots.
- A basic bar chart, as drawn with **geom_bar().**

```
ggplot(data = diamonds) +
  geom_bar(mapping = aes(x = cut))
```

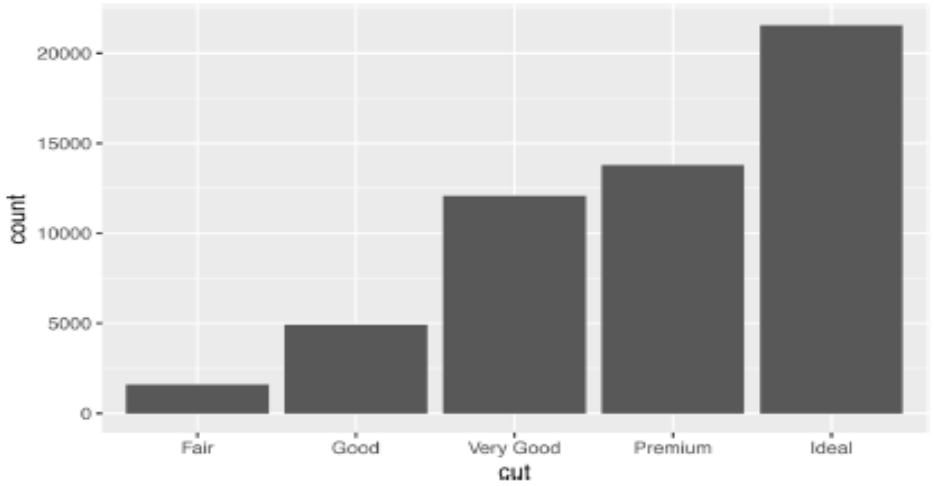


- Bar charts, histograms, and frequency polygons bin your data and then plot bin counts, the number of points that fall in each bin.
- Smoothers fit a model to your data and then plot prediction from the model.
- Boxplots compute a robust summary of the distribution and display a specially formatted box.
- The default value for stat is "count," which means that **geom_bar()** uses **stat_count().**



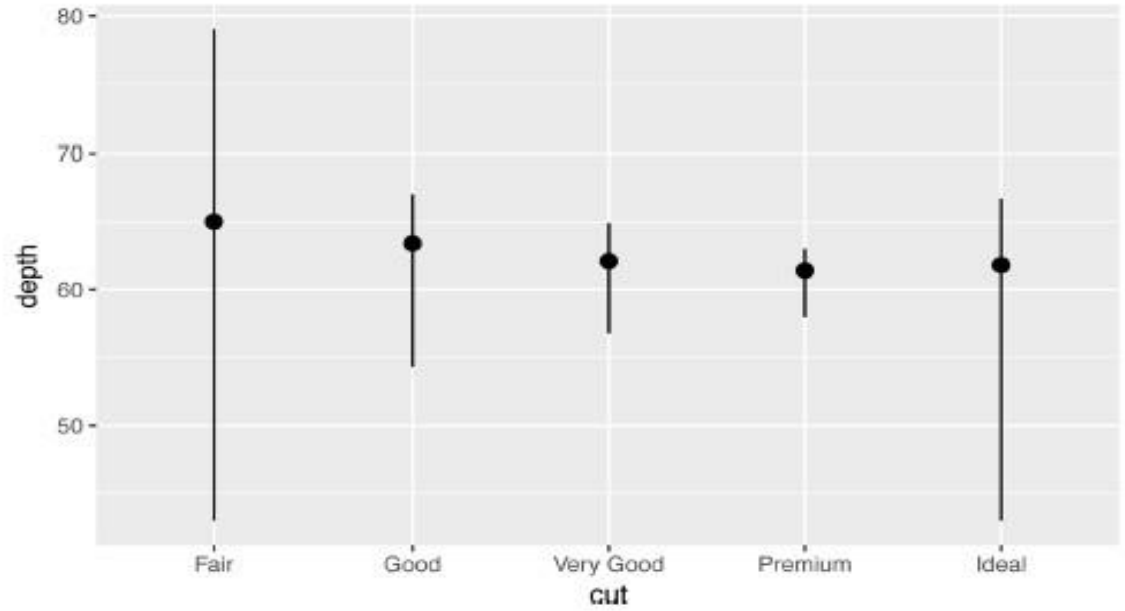
- You can re-create the previous plot using **stat_count()** instead of geom_bar():

```
ggplot(data = diamonds) +
  stat_count(mapping = aes(x = cut))
```




- You might use **stat_summary(),** which summarizes the y values for each unique x value, to draw attention.

```
ggplot(data = diamonds) +
    stat_summary(
        mapping = aes(x = cut, y = depth),
        fun.ymin = min,
        fun.ymax = max,
        fun.y = median)
```





# Position Adjustments

- You can color a bar chart using either the **color** aesthetic, or
  - ✓ by **fill = x** axes variable.
  - ✓ **clarity:** the bars are automatically stacked.
- The stacking is performed automatically by the *position adjustment* specified by the position argument. So ggplot is providing 3 such position options. They are: "identity", "dodge" or "fill".

**position = "identity“:**

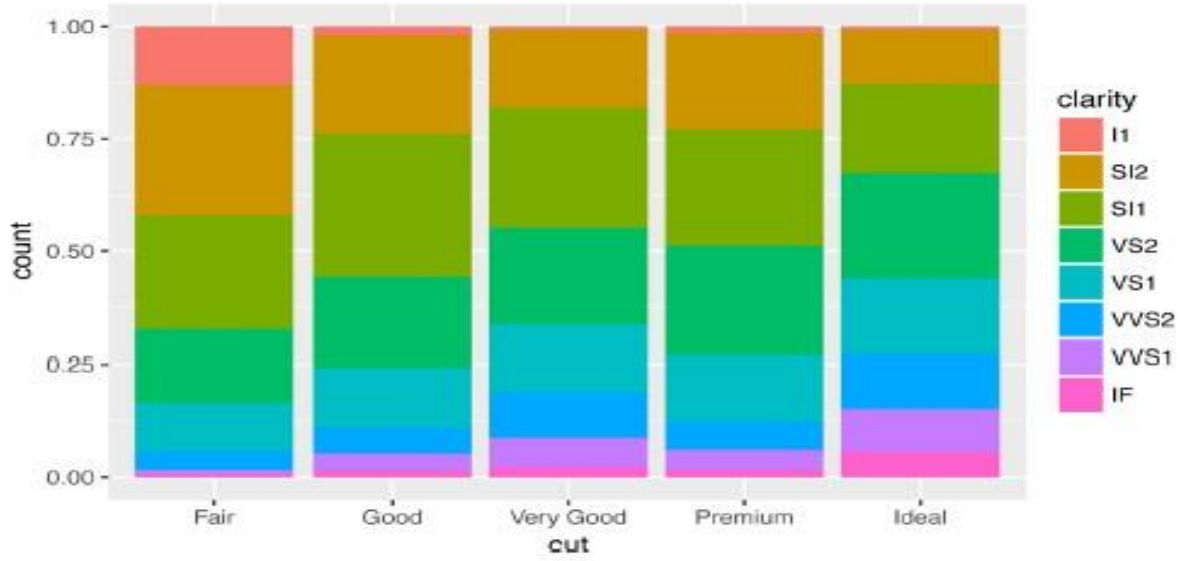
- ➢ It will place each object exactly where it falls in the context of the graph.
- ➢ This is not very useful for bars, because it overlaps them.
- ➢ To see that overlapping we either need to make the bars slightly transparent by setting **alpha** to a small value, or completely transparent by setting **fill = NA**

```
ggplot(
  data = diamonds,
  mapping = aes(x = cut, fill = clarity)
) +
  geom_bar(alpha = 1/5, position = "identity")
ggplot(
  data = diamonds,
  mapping = aes(x = cut, color = clarity)
) +
  geom_bar(fill = NA, position = "identity")
```

**position = "fill":**

- It works like stacking, but makes each set of stacked bars the same height.
- This makes it easier to compare proportions across groups:

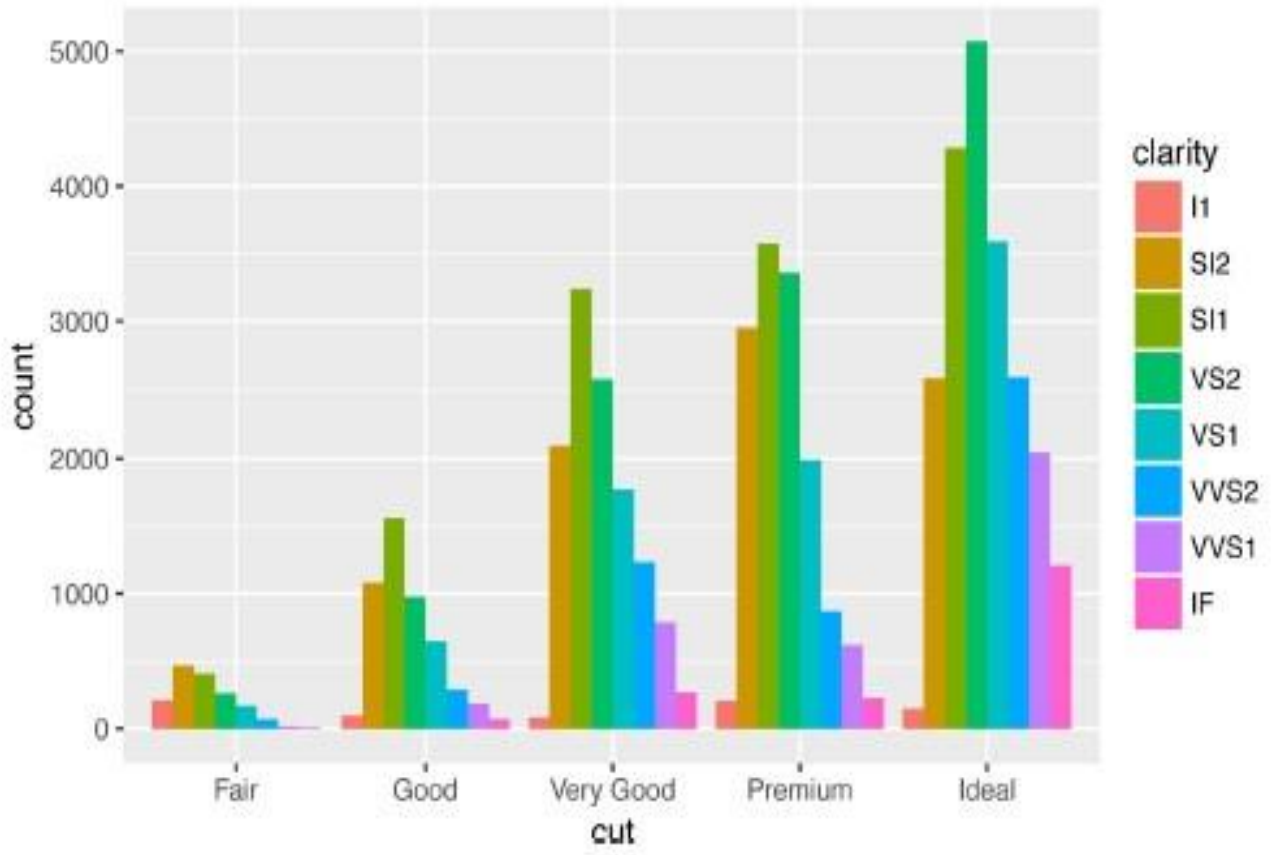
```
ggplot(data = diamonds) +
    geom_bar(mapping = aes(x = cut, fill = clarity), position = "fill")
```




**position = "dodge":**

- It places overlapping objects directly *beside* one another. This makes it easier to compare individual values:

```
ggplot(data = diamonds) +
    geom_bar(mapping = aes(x = cut, fill = clarity),
    position = "dodge")
```


5000
4000
3000
2000
1000
0
count
Fair
Good
Very Good
Premium
Ideal
cut
clarity
I1
SI2
SI1
VS2
VS1
VVS2
VVS1
IF

# Coordinate Systems

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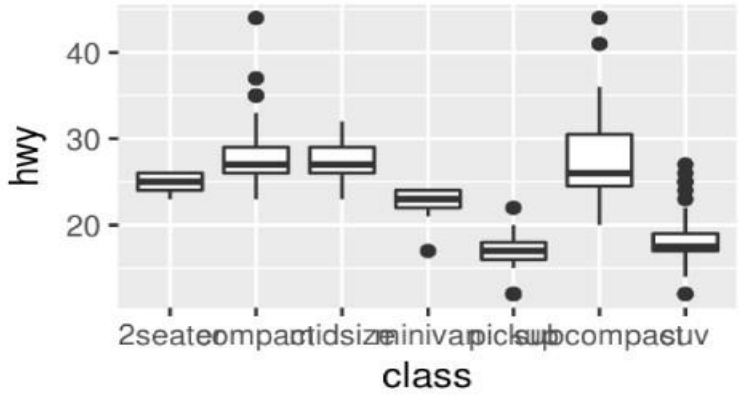
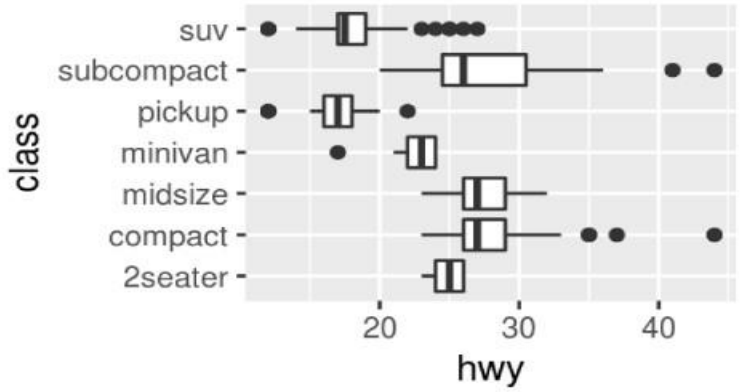
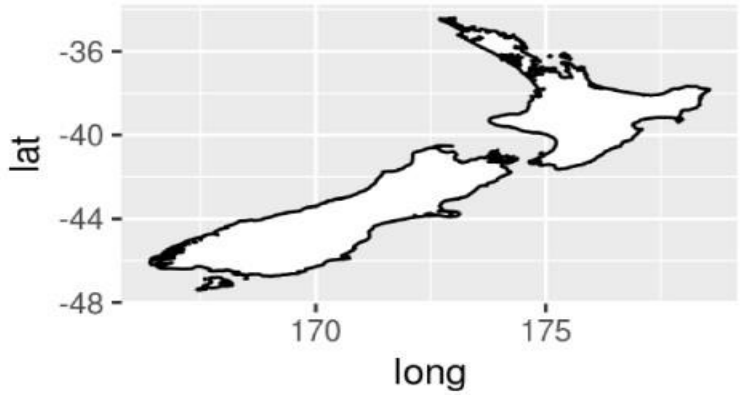
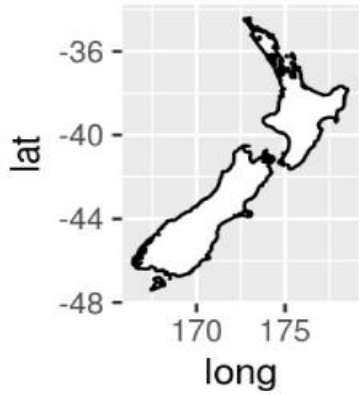
- Coordinate systems are probably the most complicated part of **ggplot2**.
- The default coordinate system is the Cartesian coordinate system where the x and y position act independently to find the location of each point.
- There are a number of other coordinate systems that are occasionally helpful:
  - ✓**coord_flip():** It switches the x- and y-axes, if you want horizontal boxplots. It's also useful for long labels.



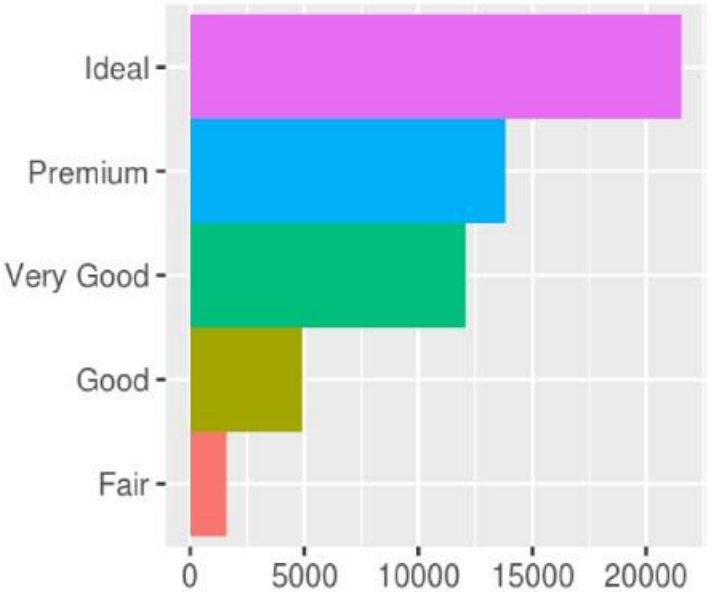
- **coord_quickmap()** sets the aspect ratio correctly for maps. This is very important if you're plotting spatial data with **ggplot2.** Example: maps




- **coord_polar()** uses polar coordinates. Polar coordinates reveal an interesting connection between a bar chart and a Coxcomb chart.




Ideal
Premium
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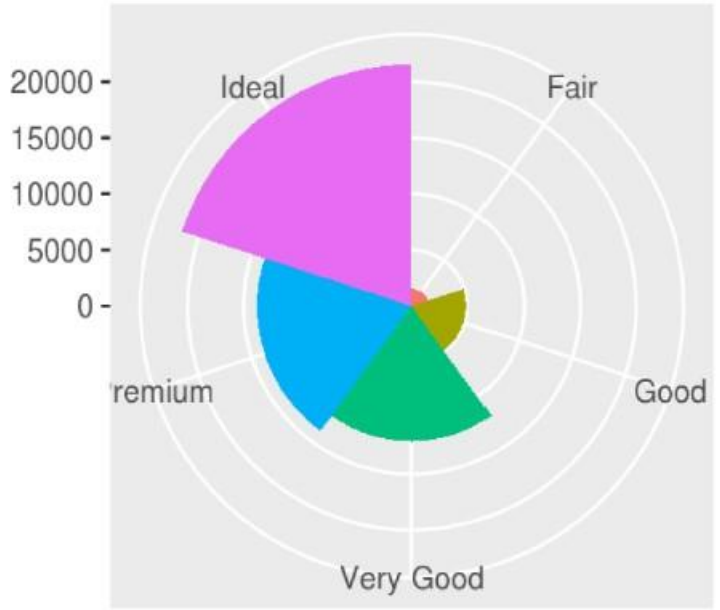
20000
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Ideal
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