GGPLOT

Installation and Load

```
if(!("ggplot2" %in% installed.packages())) install.packages("ggplot2")
library(ggplot2)
```

ggplot foundations

```
"...supply a dataset and aesthetic mapping (with aes()). You then add on
layers (like geom_point() or geom_histogram()), scales (like
scale_colour_brewer()), faceting specifications (like facet_wrap()) and
coordinate systems (like coord_flip())..."

(from https://ggplot2.tidyverse.org)

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

Data

```
v1 <- c(1,2,3,4,5)
v2<- v1*2
v3 <- v1*10
v4 <- v1 * 20
df1 <-data.frame(v1,v2,v3, v4); df1
rm(v1,v2,v3,v4)</pre>
```

Use of layered approach

```
# using layers to develop plot with arbitrary complexity
ggplot(df1,mapping=aes(x=v1,y=v2))
ggplot(df1,mapping=aes(x=v1,y=v2)) + geom_point()
```

Specifying Aesthetics manually

```
# specifying aesthetics
ggplot(df1,mapping=aes(x=v1,y=v2)) + geom_point(aes(v1,v3)) +
  geom point(aes(v2,v3),color='red')
ggplot(df1,mapping=aes(x=v1,y=v2)) + geom_point(aes(v1,v3)) +
  geom_point(shape=24)
# direct aesthetics mapping
ggplot(df1,mapping=aes(x=v1,y=v2)) + geom_point(aes(y=v3))+
  geom_abline(intercept = 20) + geom_point(aes(y=v4, size=v4),shape=22,
fill='red')
ggplot(df1,mapping=aes(x=v1,y=v2)) + geom_point(aes(color="red"))+
  geom_point(aes(y=v3,color="blue"))
```

Specifying Aesthetics - Scaling

```
ggplot(df1,mapping=aes(x=v1,y=v2)) + geom_point(aes(v1,v3)) +
    geom_point(aes(shape=factor(v1)))

# aesthetics mapping with variables - scaling
ggplot(data=mpg)+geom_point(mapping=aes(x=displ,y= hwy))
ggplot(data = mpg) + geom_point(mapping = aes(x = displ, y = hwy, color = class))
ggplot(data = mpg) + geom_point(mapping=aes(x=displ, y = hwy, size = class))
ggplot(data = mpg) + geom_point(mapping=aes(x=displ, y = hwy, shape = class))
ggplot(data = mpg) + geom_point(mapping=aes(x=displ, y = hwy, alpha = class))
ggplot(data = mpg) +
    geom_point(mapping=aes(x=displ, y = hwy, size = cty, color = class))
```

Plot limits

```
# use of plot limits
ggplot(df1,mapping=aes(x=v1,y=v2)) + geom_point(aes(y=v3)) +
   geom_abline(intercept = 0, slope = 1) + xlim(0,5) + ylim(0,50)
```

Mixing variable

```
# mixing variables mapping - overriding earlier specs
ggplot(df1,mapping=aes(x=v1,y=v3)) + geom_point(mapping=aes(x=v1,y=v2)) +
    geom_line(aes(y=v2))
ggplot(df1,mapping=aes(x=v1,y=v3)) + geom_point(mapping=aes(x=v1,y=v2)) +
    geom_line()
ggplot(df1,mapping=aes(x=v1,y=v3)) + geom_point(mapping=aes(x=v1,y=v2)) +
    geom_line(aes(y=v4))
```

Use of themes

```
# use of themes
ggplot(df1,mapping=aes(x=v1,y=v3)) + geom_point(mapping=aes(x=v1,y=v2)) +
  geom_line(aes(y=v4)) + theme_dark()
ggplot(df1,mapping=aes(x=v1,y=v3)) + geom_point(mapping=aes(x=v1,y=v2)) +
  geom_line(aes(y=v4)) + theme_classic()
# use of themes
mynamestheme <- theme(plot.title = element_text(family = "Helvetica", face = "bold", size = (15)),
                      legend.title = element_text(colour = "steelblue", face = "bold.italic",
family = "Helvetica"),
                      legend.text = element_text(face = "italic", colour="steelblue4",family =
"Helvetica"),
                      axis.title = element_text(family = "Helvetica", size = (10), colour =
"steelblue4"),
                      axis.text = element_text(family = "Courier", colour = "cornflowerblue", size =
(10)))
ggplot(df1,mapping=aes(x=v1,y=v2)) + geom_point(aes(v1,v3)) +
  geom_point(aes(shape=factor(v1))) +
  labs(title = "Arbitrary plot", subtitle = "multiple scatter plots") +
 xlab('v1') + ylab('v2 or v3') + mynamestheme
```

Labeling

```
# use of labs for formatting plot labeling
ggplot(df1,mapping=aes(x=v1,y=v2)) + geom_point(aes(v1,v3)) +
    geom_point(aes(shape=factor(v1))) +
    labs(title = "Arbitrary plot", subtitle = "multiple scatter plots") +
    xlab('v1') + ylab('v2 or v3')
ggplot(df1,mapping=aes(x=v1,y=v2)) + geom_point(aes(v1,v3)) +
    geom_point(aes(shape=factor(v1))) +
    labs(title = "Arbitrary plot", subtitle = "multiple scatter plots",
    x = 'v1', y = 'v2 or v3')
```

Facets and Grids

```
# facets
ggplot(data = mpg) + geom_point(mapping = aes(x = displ, y = hwy)) +
    facet_wrap(~ class)
ggplot(data = mpg) + geom_point(mapping = aes(x = displ, y = hwy)) +
    facet_wrap(~ class, nrow = 2)

ggplot(data = mpg) + geom_point(mapping = aes(x = displ, y = hwy)) +
    facet_wrap(~ class + drv)
ggplot(data = mpg) + geom_point(mapping = aes(x = displ, y = hwy)) +
    facet_grid(class ~ drv)
```

Aesthetics description

```
# check for details of aesthetics
vignette("ggplot2-specs")
```

Multiple geoms - continuous and discrete data

```
# geoms - continuous and discrete data
ggplot(data = mpg) + geom_smooth(mapping=aes(x=displ, y = hwy))
ggplot(data = mpg) + geom_point(mapping=aes(x=displ, y = hwy)) +
geom\_smooth(aes(x=displ, y = hwy))
ggplot(data = diamonds) +
  geom_bar(mapping = aes(x = cut, fill = color, y = ...count...)) #mark use of
computed variables in plot
ggplot(data = diamonds) +
  geom_bar(mapping = aes(x = cut, fill = color, y = ...count..),
           position = position_dodge())
ggplot(data = diamonds) +
  geom_bar(mapping = aes(x = cut, fill = color, y = ..prop..))
ggplot(data = diamonds, mapping = aes(x = cut, fill = clarity)) +
  geom_bar(alpha=1/2,position="dodge")
ggplot(data = diamonds) +
  geom_histogram(mapping = aes(x = carat), binwidth = 0.5)
ggplot(data = diamonds, mapping = aes(x = carat, color = cut)) +
  geom_freqpoly(binwidth = 0.1) + scale_color_hue(h=c(20,100))
ggplot(ToothGrowth, aes(x = factor(dose), y = len, fill = dose)) + geom_boxplot()
```

Multiple geoms - continuous and discrete data

```
library(ggplot2)
head(BOD)
ggplot(BOD, aes(x = Time, y = demand)) + geom_col(fill="black", color = "black")
ggplot(BOD, aes(x = factor(Time), y = demand)) +
    geom_col()
```

Breaking into parts

```
# breaking into multiple parts
base_plot <- ggplot(data=mpg,mapping=aes(x=displ,y=hwy))
base_plot + geom_point() + geom_smooth()</pre>
```

dplyr and ggplot

Use of reorder

```
my_mpg %>%
   ggplot() +
geom_bar(aes(reorder(mpg$cyl,mpg$cyl,NROW),y=after_stat(prop),group=1)) +
   xlab(levels(reorder(mpg$cyl,mpg$cyl,NROW)))
```

Scales

```
ggplot(data = diamonds, mapping = aes(x = carat, color = cut)) +
    geom_freqpoly(binwidth = 0.1) + scale_color_hue(h=c(20,100))

tg<-ToothGrowth
tg$dose <- as.factor(ToothGrowth$dose)
my_plot <- ggplot(tg, aes(x = dose, y = len, fill = dose)) + geom_boxplot()
my_plot
my_plot + scale_fill_manual(values = c("#004f71", "#465a01", "#981d97"))
my_plot + scale_fill_manual(values = c("red", "blue", "green"))

#scale_fill_manual() for box plot, bar plot, violin plot,
#scale_color_manual() for lines and points</pre>
```

Stats

Many graphs, like scatterplots, plot the raw values of your dataset. Other graphs, like bar charts, calculate new values to plot:

- 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 predictions from the model.
- boxplots compute a robust summary of the distribution and then display a specially formatted box.

The algorithm used to calculate new values for a graph is called a stat, short for statistical transformation.

You can learn which stat a geom uses by inspecting the default value for the stat argument.

geom_bar - stat_count() # prop is another computed variable

You can generally use geoms and stats interchangeably.

Stats

```
ggplot(data = diamonds) + geom_bar(mapping = aes(x = cut))
ggplot(data = diamonds) + stat_count(mapping = aes(x = cut)) #same as above

mpg_tab <- as.data.frame(table(mpg$cyl))
mpg_tab
ggplot(mpg_tab) + geom_bar(aes(Var1,Freq)) #gives error
ggplot(mpg_tab) + geom_bar(aes(Var1,Freq),stat = "identity")
ggplot(mpg) + geom_bar(aes(cyl))
ggplot(mpg) + stat_count(aes(cyl))
ggplot(mpg) + stat_count(aes(factor(cyl)))
ggplot(mpg) + stat_count(aes(cyl,y=stat(prop)))
ggplot(mpg) + geom_bar(aes(cyl,..prop..))
ggplot(mpg) + stat_count(aes(cyl,..prop..))</pre>
```

Stats

```
c <- ggplot(mpg, aes(hwy)); c2 <- ggplot(mpg)
c + stat_bin(mapping=aes(hwy,..ncount..),binwidth = 5)
c + stat_bin(mapping=aes(hwy,..density..),binwidth = 5)
c + stat_density(adjust = 1, kernel = 'gaussian')
c + stat_density(adjust = 1, kernel = 'triangular')
e <- ggplot(mpg, aes(cty, hwy))
e + stat_ellipse(level = 0.95, segments = 51, type = "t")</pre>
```

Types of plots

- One variable continuous
- one variable discrete
- two variables both continuous
- two variables both discrete
- two variables one discrete, one continuous
- three variables (contour plots, etc.)
- faceted / gridded plots

ggplot reference sheet





