

Lab 2: Descriptive Statistics and ggplot2

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```
## [1] "/Users/haowang/Dropbox/2017 Spring/ 603/lab/Lab 2"
```

Working Example: Baylor's American Religious Survey

The following R chunk load data from Baylor Religious Survey. Baylor Religious Study is a comprehensive analysis on religious beliefs in the United States. For detailed explanation please refer to <http://www.thearda.com/Archive/Files/Descriptions/BAYLORW2.asp>. Total sample size is 1648 adults, with 318 variables, survey was conducted by Gallup.

```
#The first step is to load Byalor Religious Survey 2005 (I)
mydata <- read.dta13("http://www.thearda.com/download/download.aspx?file=Baylor%20Religion%20Survey,%20")
```

Subsetting Data

Because the raw file includes some missing points, I create a new dataset of potential interesting variables without missing data.

```
plotdata <- mydata[, c('religious', 'attend', 'gaymarr', 'gayborn',
                      'votefem', 'partyid')]
#religious: how religious you are
#attend: how often do you attend church
#gaymarr: gay marriage should be legal
#gayborn: people are born either homo or heterosexual
#votefem: would you vote for a female cadidate nominated by your party
#partyid: democract or republican
plotdata <- na.omit(plotdata)
```

Mean attitudes towards gay marriage

```
summary(plotdata$gaymarr)
```

```
## Strongly disagree      Disagree      Agree      Strongly agree
##           507           311           223           291
##      Undecided
##           159
```

```
mean(plotdata$gaymarr)
```

```
## Warning in mean.default(plotdata$gaymarr): argument is not numeric or
## logical: returning NA
```

```
## [1] NA
```

What happened? we got an error message since all the survey items are coded in the categorical way.

Convert categorical data into numeric

```
plotdata2=as.data.frame(sapply(plotdata, as.numeric))#sapply function returns matrix with same  
#length, and in the same time converted factors into numeric numbers by as.numeric command  
table(plotdata2$gaymarr)
```

```
##  
##    1    2    3    4    5  
## 507 311 223 291 159
```

```
mean(plotdata2$gaymarr)
```

```
## [1] 2.519785
```

- Lab Practice 1: How to interpret the mean value of plotdata2? Is it correct?

Recode variables

```
plotdata2$gaymarr1 <- plotdata2$gaymarr  
plotdata2$gaymarr1[plotdata2$gaymarr1 == 5] <- NA  
mean(plotdata2$gaymarr1, na.rm = TRUE)
```

```
## [1] 2.223724
```

```
median(plotdata2$gaymarr1, na.rm = TRUE)
```

```
## [1] 2
```

Get an overview by the summary() command

```
summary(plotdata)
```

```
##           religious           attend  
## Not at all religious:165   Never           :327  
## Not too religious   :213   Weekly           :318  
## Somewhat religious  :617   Several times a year :168  
## Very religious      :483   Once or twice a year :155  
## Don't know          : 13   Several times a week :137  
##                     Less than once a year:122  
##                     (Other)           :264  
##           gaymarr          gayborn    votefem  
## Strongly disagree:507   Strongly disagree:254   Yes:1190  
## Disagree          :311   Disagree           :259   No : 301  
## Agree             :223   Agree              :420  
## Strongly agree    :291   Strongly agree     :281
```

```
## Undecided      :159 Undecided      :277
##
##
##      partyid
## Independent    :311
## Moderate Democrat :281
## Moderate Republican:261
## Strong Democrat  :196
## Strong Republican :174
## Leaning Republican :125
## (Other)         :143
```

Frequency Table

```
table(plotdata$gaymarr)
```

```
##
## Strongly disagree      Disagree      Agree      Strongly agree
##           507           311           223           291
## Undecided
##           159
```

Graphing Using ggplot2

The ggplot2 package, created by Hadley Wickham, offers a powerful graphics language for creating elegant and complex plots. Its popularity in the R community has exploded in recent years. Originally based on Leland Wilkinson's The Grammar of Graphics, ggplot2 allows you to create graphs that represent both univariate and multivariate numerical and categorical data in a straightforward manner. Grouping can be represented by color, symbol, size, and transparency. The creation of trellis plots (i.e., conditioning) is relatively simple.

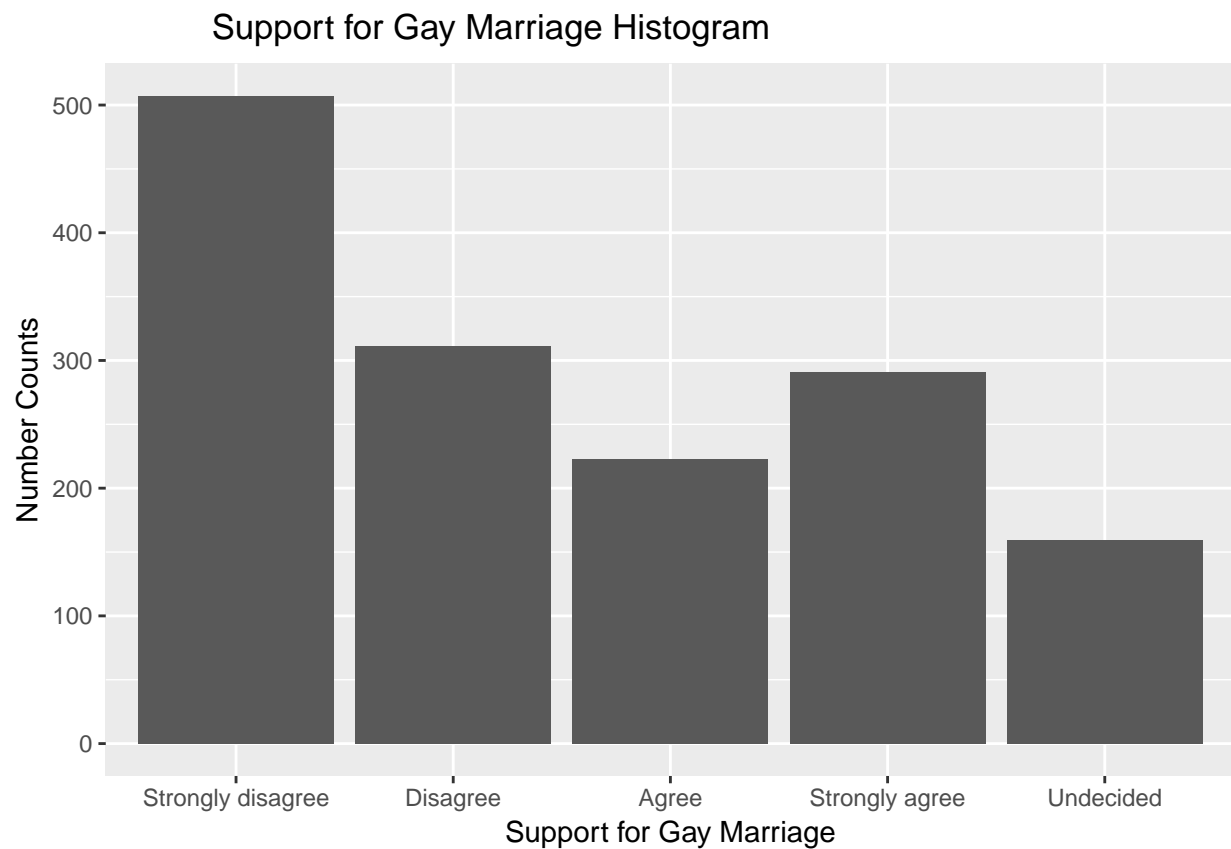
ggplot 2 reference guide: <http://docs.ggplot2.org/current/index.html#>

ggplot 2 cheatsheet: <https://www.rstudio.com/wp-content/uploads/2015/03/ggplot2-cheatsheet.pdf>

Histogram

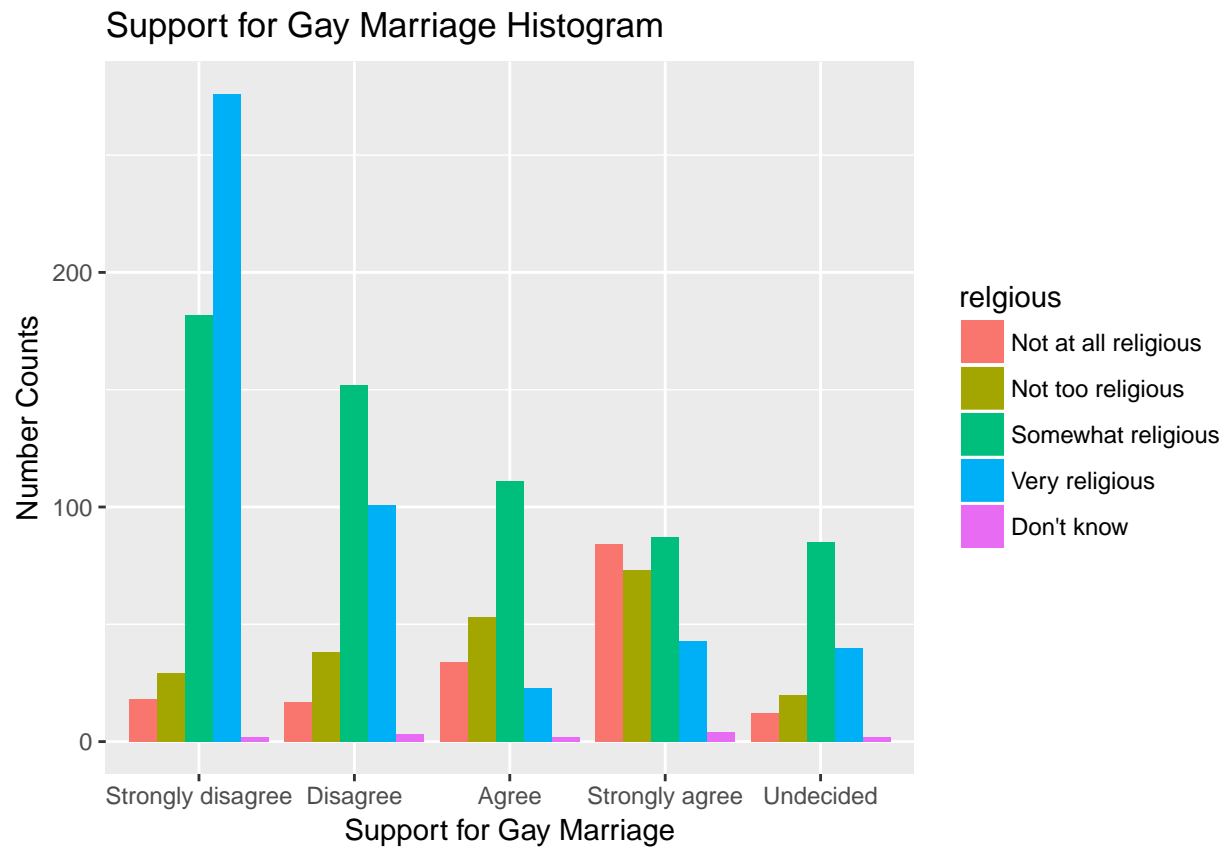
This shows the histogram of Attitudes on gay marriage

```
library(ggplot2)
hist.gay <- ggplot(plotdata, aes(x = gaymarr)) +
  geom_histogram(stat="count", alpha =1) +
  labs(x="Support for Gay Marriage", y="Number Counts") +
  ggtitle("Support for Gay Marriage Histogram")
hist.gay
```



We can also plot the histogram according to religious degrees.

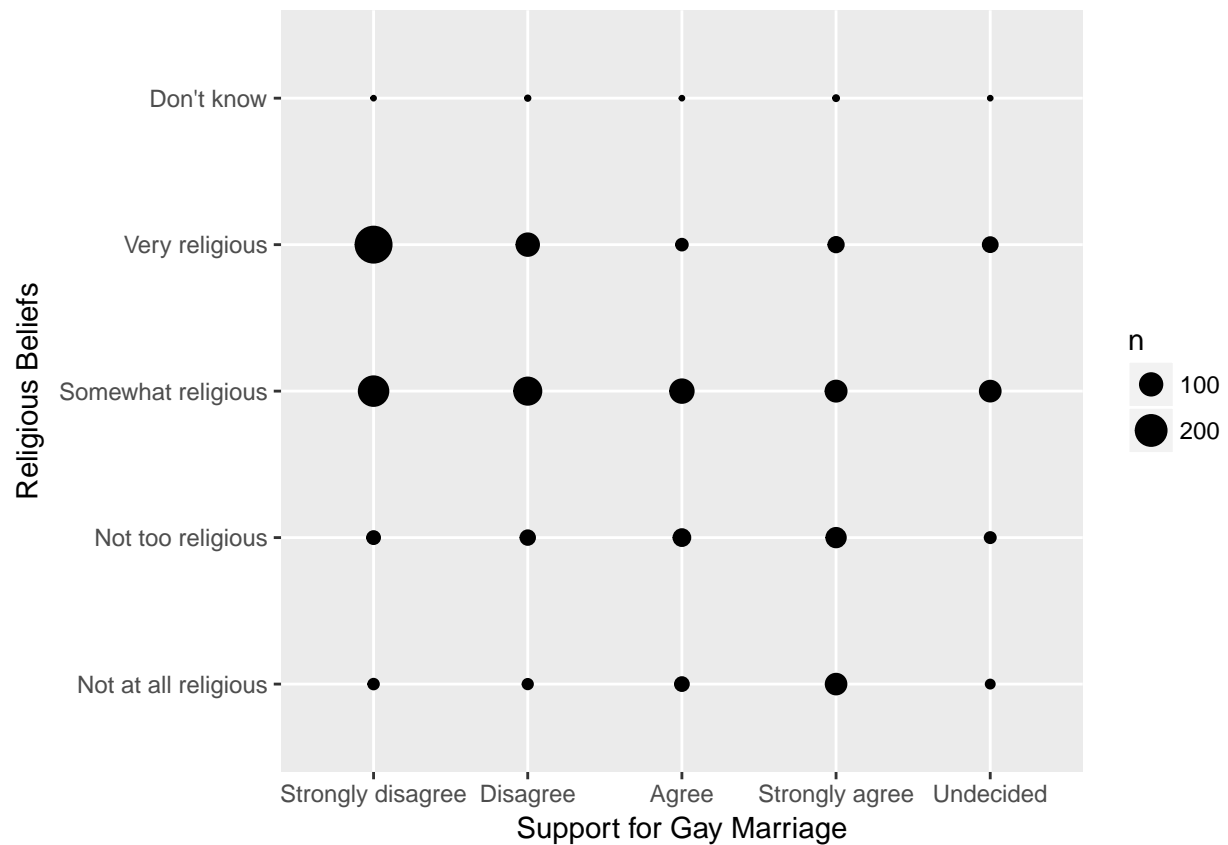
```
hist.relig.gay <- ggplot(plotdata, aes(x =gaymarr, fill=religious)) +  
  geom_histogram(stat="count",alpha =1, position = "dodge")+  
  ylab("Number Counts") +  
  xlab("Support for Gay Marriage")+  
  ggtitle("Support for Gay Marriage Histogram")  
hist.relig.gay
```



Scatter Plot

In scatter plot we write two parameters in the `aes()` option. I use additional option `geom_count()` here to illustrate the size.

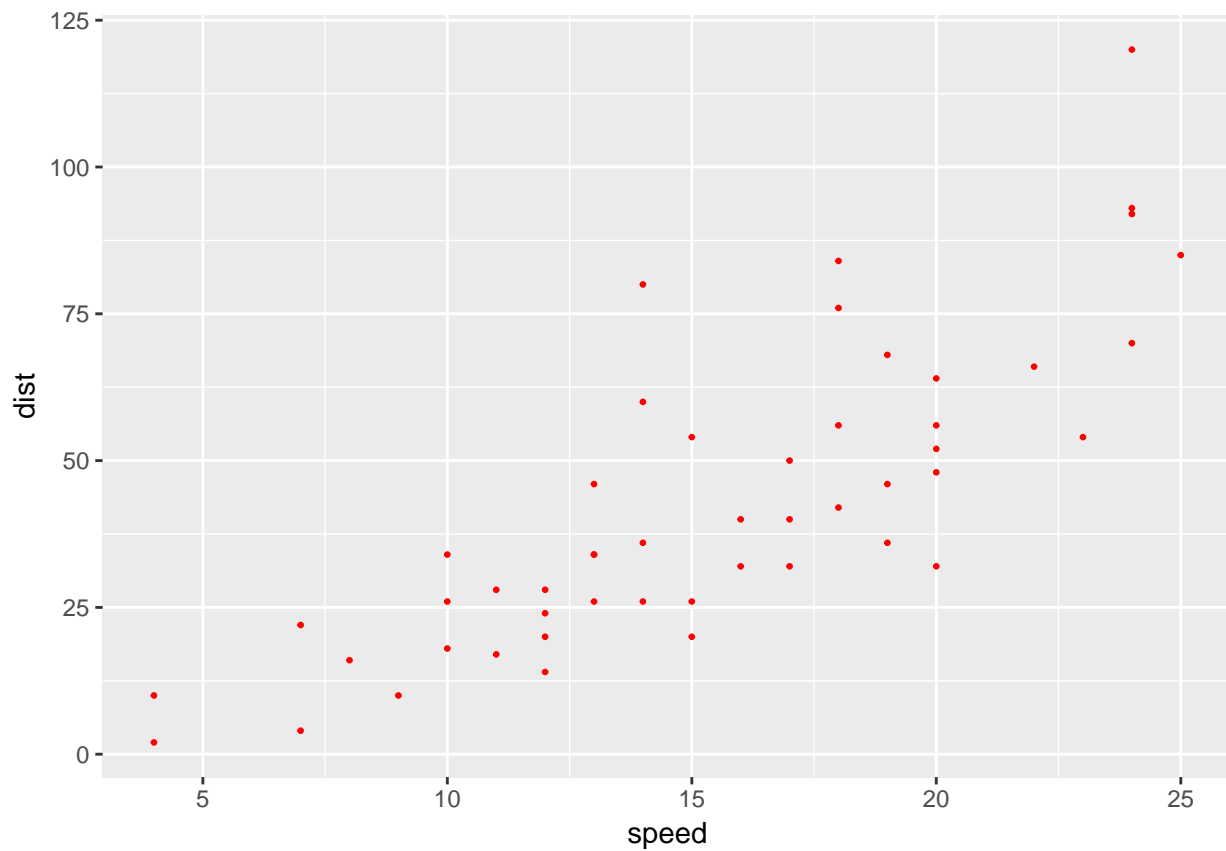
```
relig.gaymarr <- ggplot(data=plotdata, aes(y = religious, x = gaymarr)) +
  geom_count() +
  scale_size_area() +
  ylab("Religious Beliefs") +
  xlab("Support for Gay Marriage")
relig.gaymarr
```



Adding lines to scatter plot

Let's use the default data from the car package here. It measures the relation between car speed and stop distance.

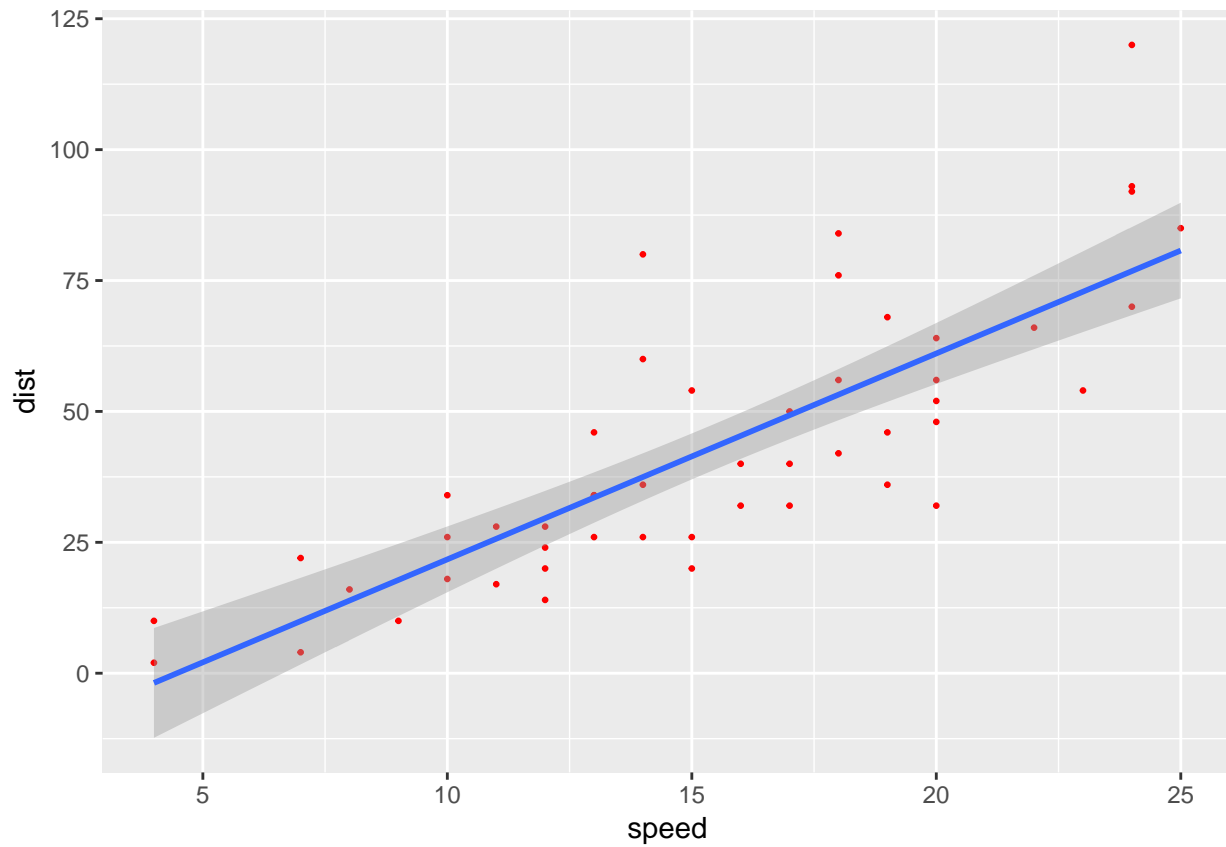
```
mydata <- cars
car <- ggplot(data = mydata, aes(x = speed, y = dist)) +
  geom_point(size=0.5, color = "red")
car
```



Let's adding a line.

Method 1, use the default `geom_smooth()`

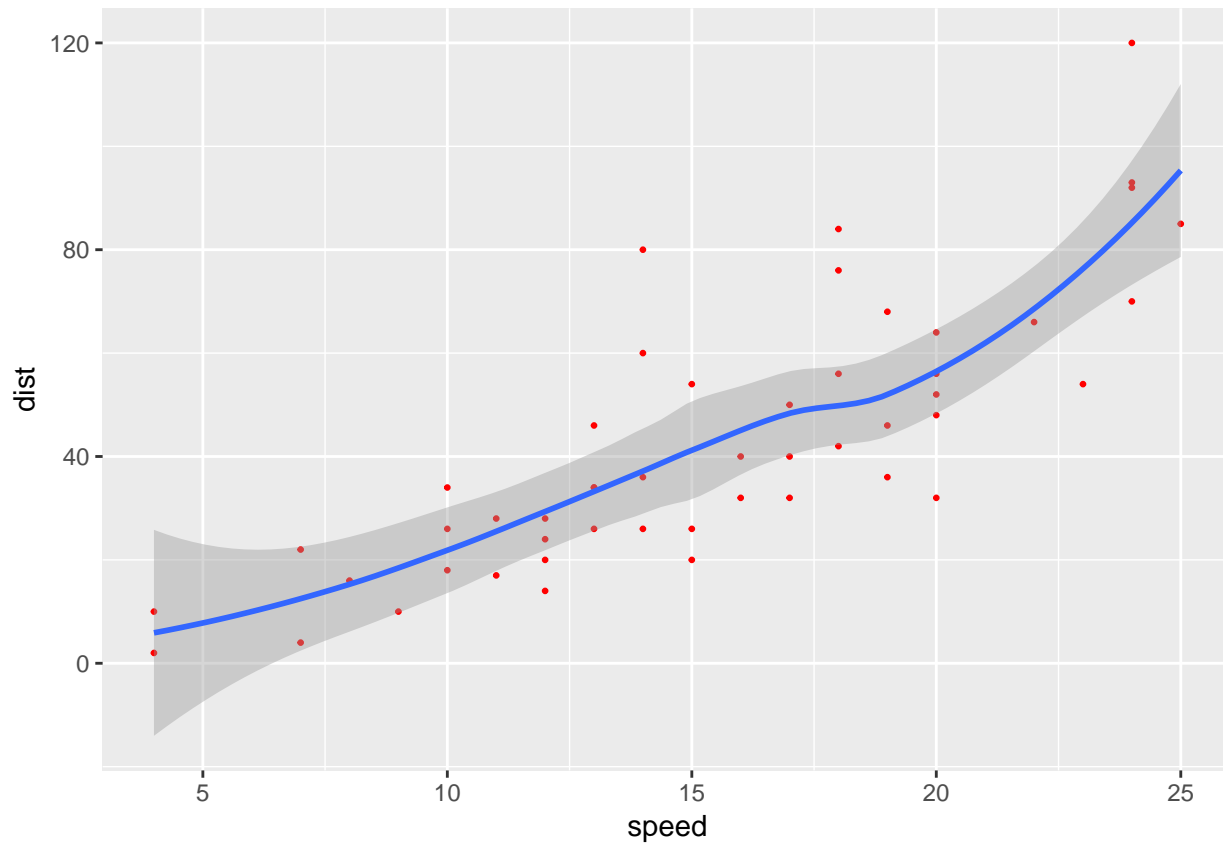
```
car <- ggplot(data = mydata, aes(x = speed, y = dist)) +  
  geom_point(size=0.5, color = "red") +  
  geom_smooth(method = 'lm', se=TRUE)  
car
```



You can try other options of `geom_smooth()`

LOESS is a nonparametric method that combine multiple regression models in a k-nearest-neighbor-based modeling.

```
car <- ggplot(data = mydata, aes(x = speed, y = dist)) +  
  geom_point(size=0.5, color = "red") +  
  geom_smooth(method = 'loess', se=TRUE)  
car
```

If you want full control over your line

In this condition you need to calculate all the parameters of the line. Let's try with linear models

```
lm <- lm(dist ~ speed, data = mydata)
summary(lm)
```

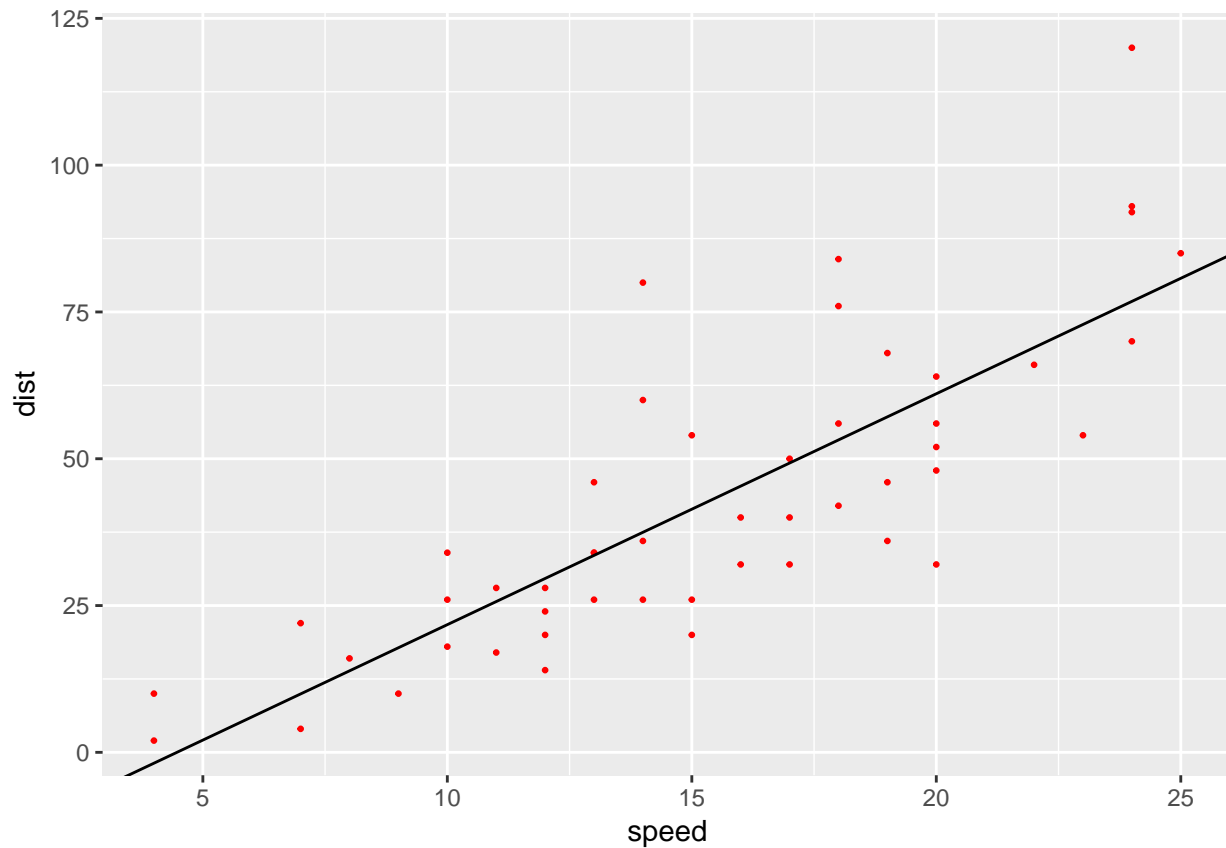
```
##
## Call:
## lm(formula = dist ~ speed, data = mydata)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -29.069  -9.525  -2.272   9.215  43.201
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -17.5791     6.7584  -2.601  0.0123 *
## speed         3.9324     0.4155   9.464 1.49e-12 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 15.38 on 48 degrees of freedom
## Multiple R-squared:  0.6511, Adjusted R-squared:  0.6438
## F-statistic: 89.57 on 1 and 48 DF,  p-value: 1.49e-12
```

```

intercept <- summary(lm)$coefficients[1,1]
slope <- summary(lm)$coefficients[2,1]

car <- ggplot(data = mydata, aes(x = speed, y = dist)) +
  geom_point(size=0.5, color = "red") +
  geom_abline(intercept = intercept, slope = slope)
car

```



The `summary(lm)` command returns coefficients of the regression. In this case we need to extract intercept in row 1, column 1; and slope in row 2, column 1.

- Lab Practice 2: In the dataset you picked,
 1. draw histograms of two variables you're interested in.
 2. Draw a scatter plot of these two variables
 3. Fit a regression line to this plot