

R: Presenting outputs

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CRAN Task View

CRAN Task View: Reproducible Research

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URL: <https://CRAN.R-project.org/view=ReproducibleResearch>

The goal of reproducible research is to tie specific instructions to data analysis and experimental data so that scholarship can be recreated, better understood and verified. Packages in R for this purpose can be split into groups for: literate programming, package reproducibility, code/data formatting tools, format convertors, and object caching.

Literate Programming

The primary way that R facilitates reproducible research is using a document that is a combination of content and data analysis code. The Sweave function (in the base R utils package) and the [knitr](#) package can be used to blend the subject matter and R code so that a single document defines the content and the analysis. The [brew](#) and [R.rsp](#) packages contain alternative approaches to embedding R code into various markups.

The resources for literate programming are best organized by the document type/markup language:

LaTeX

Both Sweave and [knitr](#) can process LaTeX files. [lazyWeave](#) can create LaTeX documents from scratch.

Object Conversion Functions:

- *summary tables/statistics* : [Hmisc](#), [NMOF](#), [papeR](#), [quantreg](#), [rapport](#), [reporttools](#), [sparktex](#), [tables](#), [xtable](#), [ztable](#)
- *tables/cross-tabulations* : [compareGroups](#), [Hmisc](#), [lazyWeave](#), [knitLatex](#), [knitr](#), [reporttools](#), [ztable](#)
- *graphics* : [animation](#), [Hmisc](#), [grDevices:::pictex](#), [sparktex](#), [tikzDevice](#)
- *statistical models/methods* : [apsrtable](#), [memisc](#), [quantreg](#), [r2lh](#), [rms](#), [stargazer](#), [suRtex](#), [TeachingSampling](#), [texreg](#), [xtable](#), [ztable](#)
- *bibtex* : [bibtex](#) and [RefManageR](#)
- *others* : [latex2exp](#) converts LaTeX equations to plotmath expressions.

Miscellaneous Tools

- [Hmisc](#) contains a function to correctly escape special characters. [resumer](#) creates resumes. Standardized exams can be created using the [exams](#) package.

HTML

The [knitr](#) package can process HTML files directly. Sweave can also work with HTML by way of the [R2HTML](#) package. [Kmisc](#) and [lazyWeave](#) can create HTML format documents from scratch.

Object Conversion Functions:

- *summary tables/statistics* : [stargazer](#)
- *tables/cross-tabulations* : [compareGroups](#), [DT](#), [formattable](#), [htmlTable](#), [HTMLUtils](#), [hwriter](#), [Kmisc](#), [knitr](#), [lazyWeave](#), [SortableHTMLTables](#), [texreg](#), [ztable](#)
- *statistical models/methods* : [r2lh](#), [rapport](#), [stargazer](#), [xtable](#)
- *others* : [knitcitations](#), [RefManageR](#)

Figure 1:

Scatter plot: add points of different colours and symbols

class: small-code

Scatter plot: add points of different colours and symbols

```
points(x, y = NULL, type = "p", ...)  
x <- rnorm(10, sd=5, mean=20)  
y <- 2.5*x - 1.0 + rnorm(10, sd=9, mean=0)  
plot(x,y,xlab="Independent", ylab="Dependent", main="Random Stuff")  
  
x1 <- runif(8,15,25)  
y1 <- 2.5*x1 - 1.0 + runif(8,-6,6)  
??  
  
x2 <- runif(8,15,25)  
y2 <- 2.5*x2 - 1.0 + runif(8,-6,6)  
??
```

Scatter plot: legend

'legend(x, y = NULL, legend, fill = NULL, col = par("col"), border = "black", lty, lwd, pch, angle = 45, density = NULL, bty = "o", bg = par("bg"), box.lwd = par("lwd"), box.lty = par("lty"), box.col = par("fg"), pt.bg = NA, cex = 1, pt.cex = cex, pt.lwd = lwd, xjust = 0, yjust = 1, x.intersp = 1, y.intersp = 1, adj = c(0, 0.5), text.width = NULL, text.col = par("col"), text.font = NULL, merge = do.lines && has.pch, trace = FALSE, plot = TRUE, ncol = 1, horiz = FALSE, title = NULL, inset = 0, xpd, title.col = text.col, title.adj = 0.5, seg.len = 2)'

```
plot(x,y,xlab="Independent",ylab="Dependent",main="Random Stuff")  
points(x1,y1,col=2,pch=3)  
points(x2,y2,col=4,pch=5)  
???
```

Simple fitted lines

```
abline(a = NULL, b = NULL, h = NULL, v = NULL, reg = NULL, coef = NULL, untf =  
FALSE, ...)  
  
lm(formula, data, subset, weights, na.action, method = "qr", model = TRUE, x = FALSE,  
y = FALSE, qr = TRUE, singular.ok = TRUE, contrasts = NULL, offset, ...)  
  
lowess(x, y = NULL, f = 2/3, iter = 3, delta = 0.01 * diff(range(x)))  
  
x <- rnorm(10,sd=5,mean=20)  
y <- 2.5*x - 1.0 + rnorm(10,sd=9,mean=0)  
x2 <- runif(8,15,25)  
y2 <- 2.5*x2 - 1.0 + runif(8,-6,6)  
  
plot(x,y,xlab="Independent",ylab="Dependent",main="Random Stuff")  
points(x1,y1,col=2,pch=3)
```





0: 	10: 	20: 	A: A
1: 	11: 	21: 	a: a
2: 	12: 	22: 	B: B
3: 	13: 	23: 	b: b
4: 	14: 	24: 	S: S
5: 	15: 	25: 	`: `
6: 	16: 	@: 	.: .
7: 	17: 	+: 	,: ,
8: 	18: 	?: 	?: ?
9: 	19: 	#: 	*: *

Figure 2:

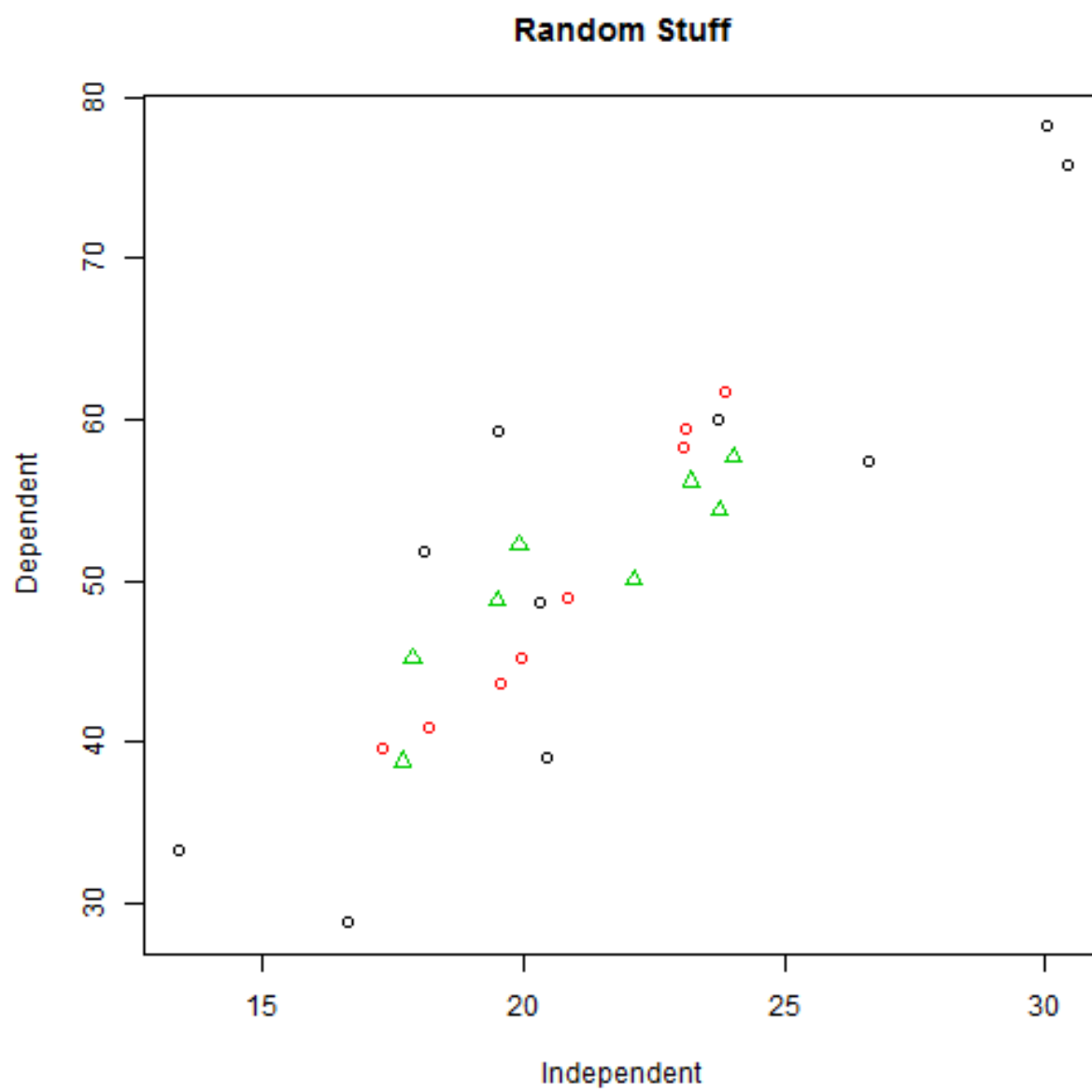


Figure 3: plot of chunk unnamed-chunk-2

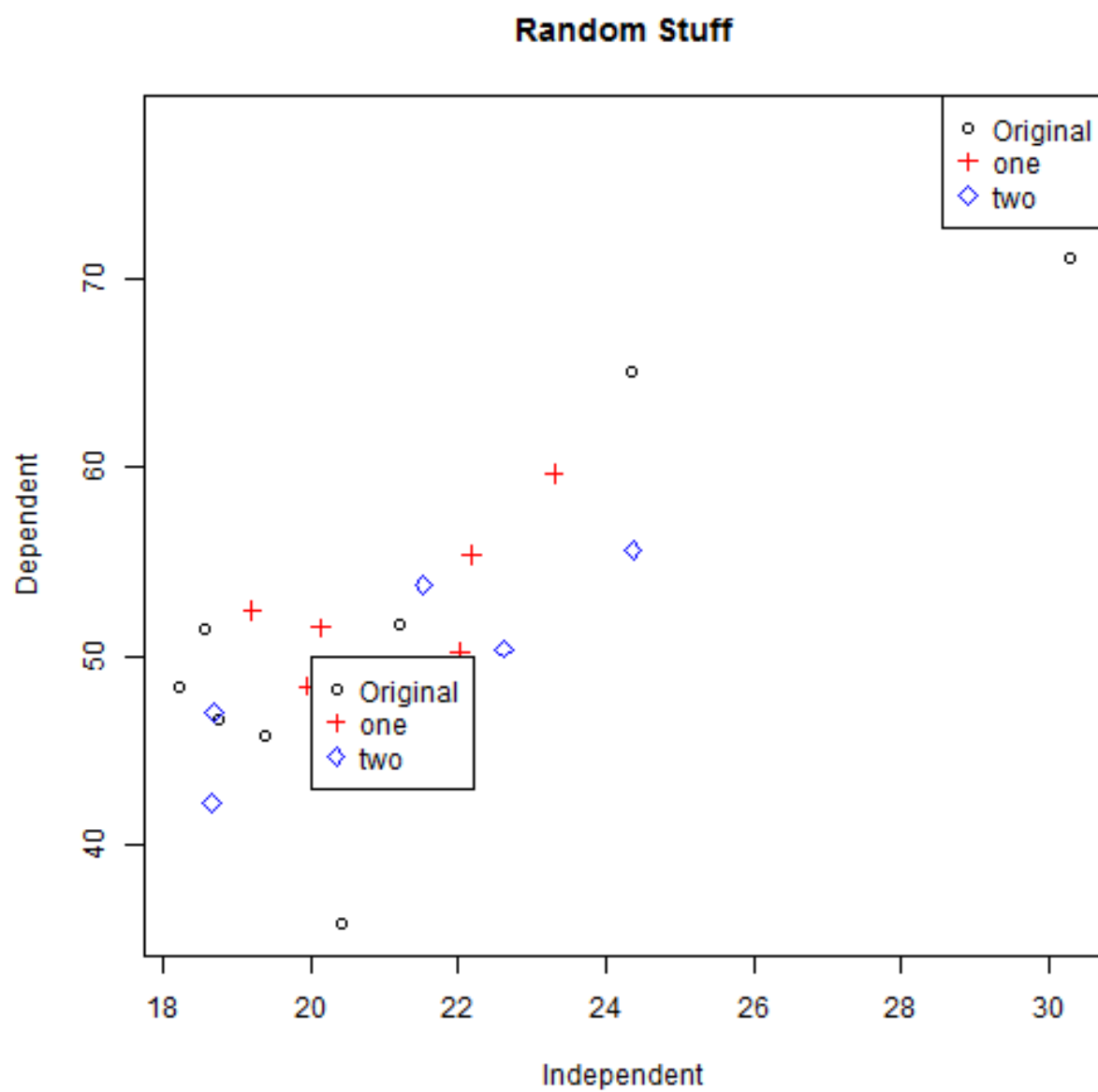


Figure 4: plot of chunk unnamed-chunk-5

```
points(x2,y2,col=4,pch=5)
legend(25,80,c("Original","one","two"),col=c(1,2,4),pch=c(1,3,5))
???
```

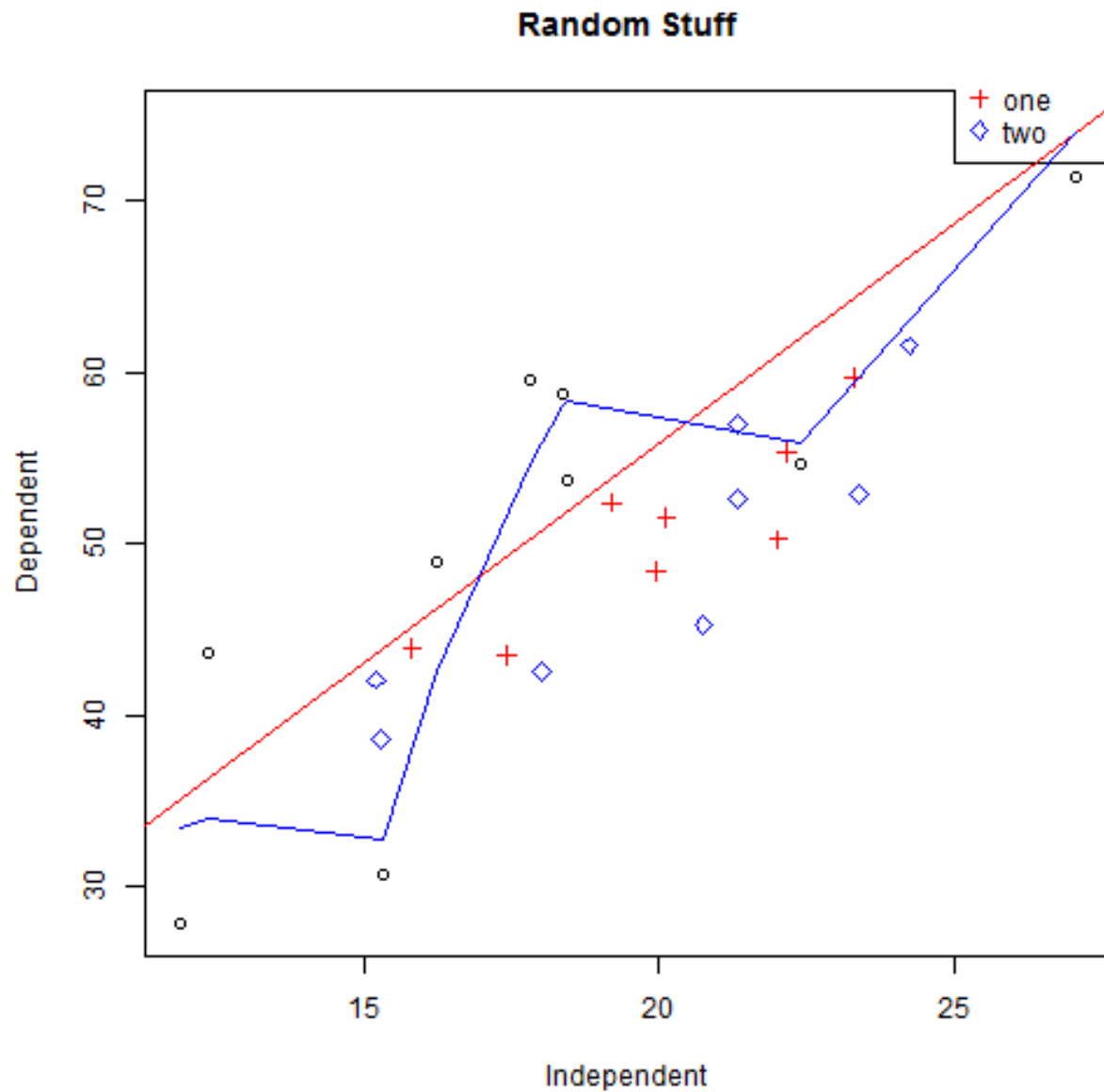


Figure 5: plot of chunk unnamed-chunk-8

With error lines

```
arrows(x0, y0, x1 = x0, y1 = y0, length = 0.25, angle = 30, code = 2, col = par("fg"),
lty = par("lty"), lwd = par("lwd"), ...)
```

```

plot(x,y,xlab="Independent",ylab="Dependent",main="Random Stuff")
xHigh <- x
yHigh <- y + abs(rnorm(10,sd=3.5))
xLow <- x
yLow <- y - abs(rnorm(10,sd=3.1))
???
```

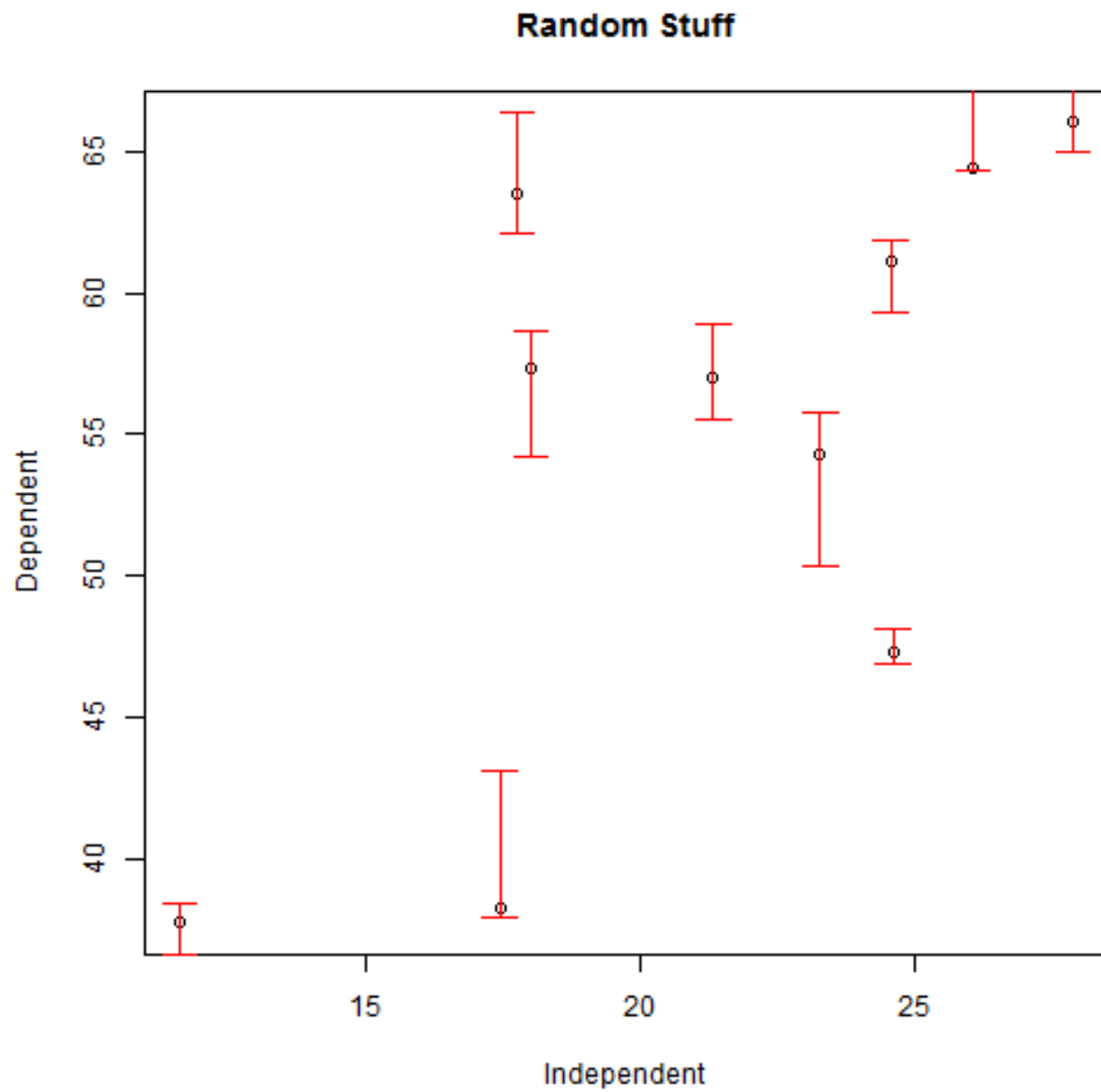


Figure 6: plot of chunk unnamed-chunk-11

Adding jitter

```
numberWhite <- ??  
numberChipped <- ??  
  
par(mfrow=c(1,2))  
plot(numberWhite,numberChipped,xlab="Number White Marbles Drawn",  
ylab="Number Chipped Marbles Drawn",main="Pulling Marbles")  
plot(jitter(numberWhite),jitter(numberChipped),xlab="Number White Marbles Drawn",  
ylab="Number Chipped Marbles Drawn",main="Pulling Marbles With Jitter")
```

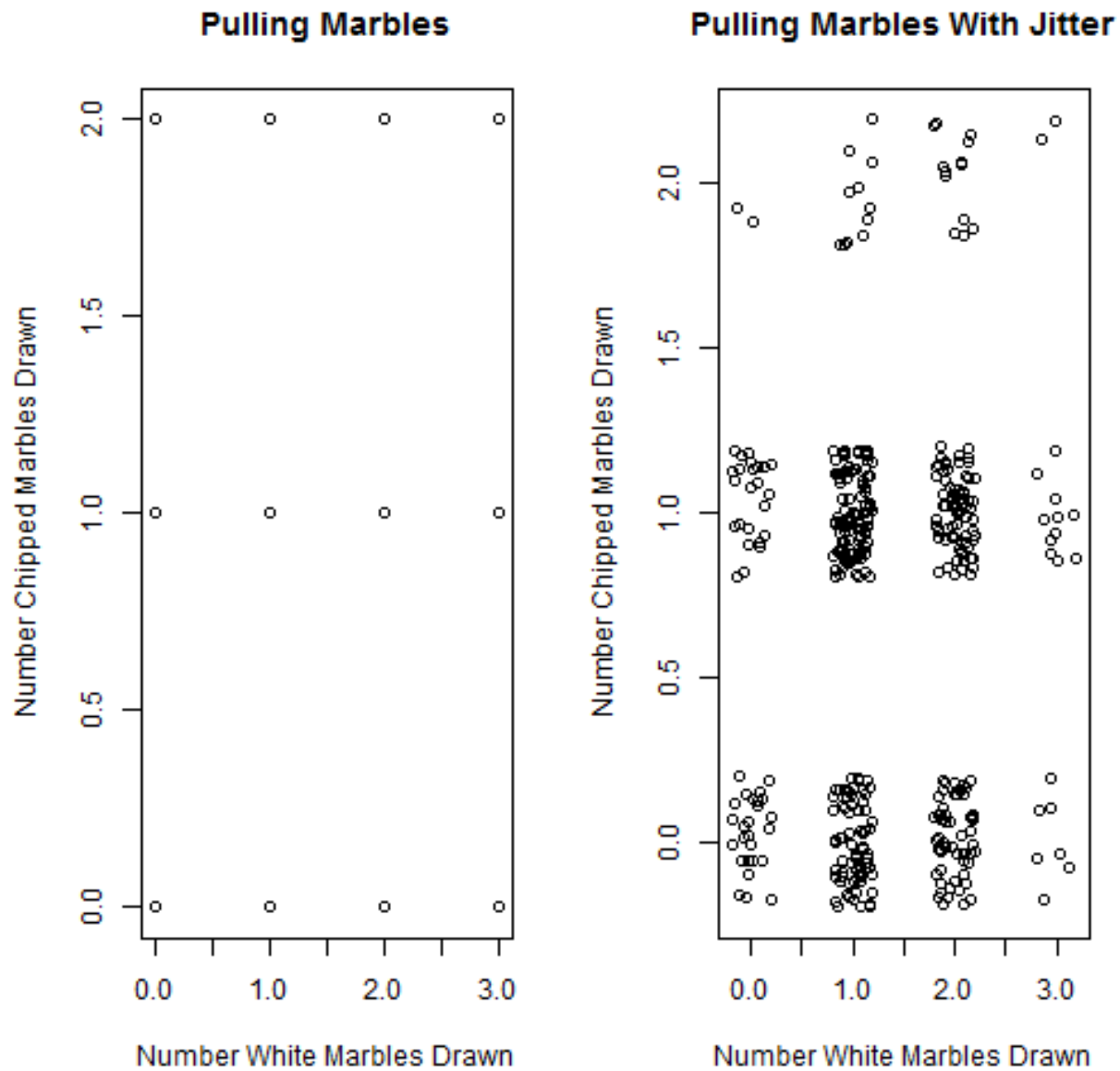


Figure 7: plot of chunk unnamed-chunk-14

Mosaic plots

```
table(...,          exclude = if (useNA == "no") c(NA, NaN),          useNA = c("no", "ifany",  
"always"),          dnn = list.names(...), deparse.level = 1)  
mosaicplot(??, main="sixth plot")
```

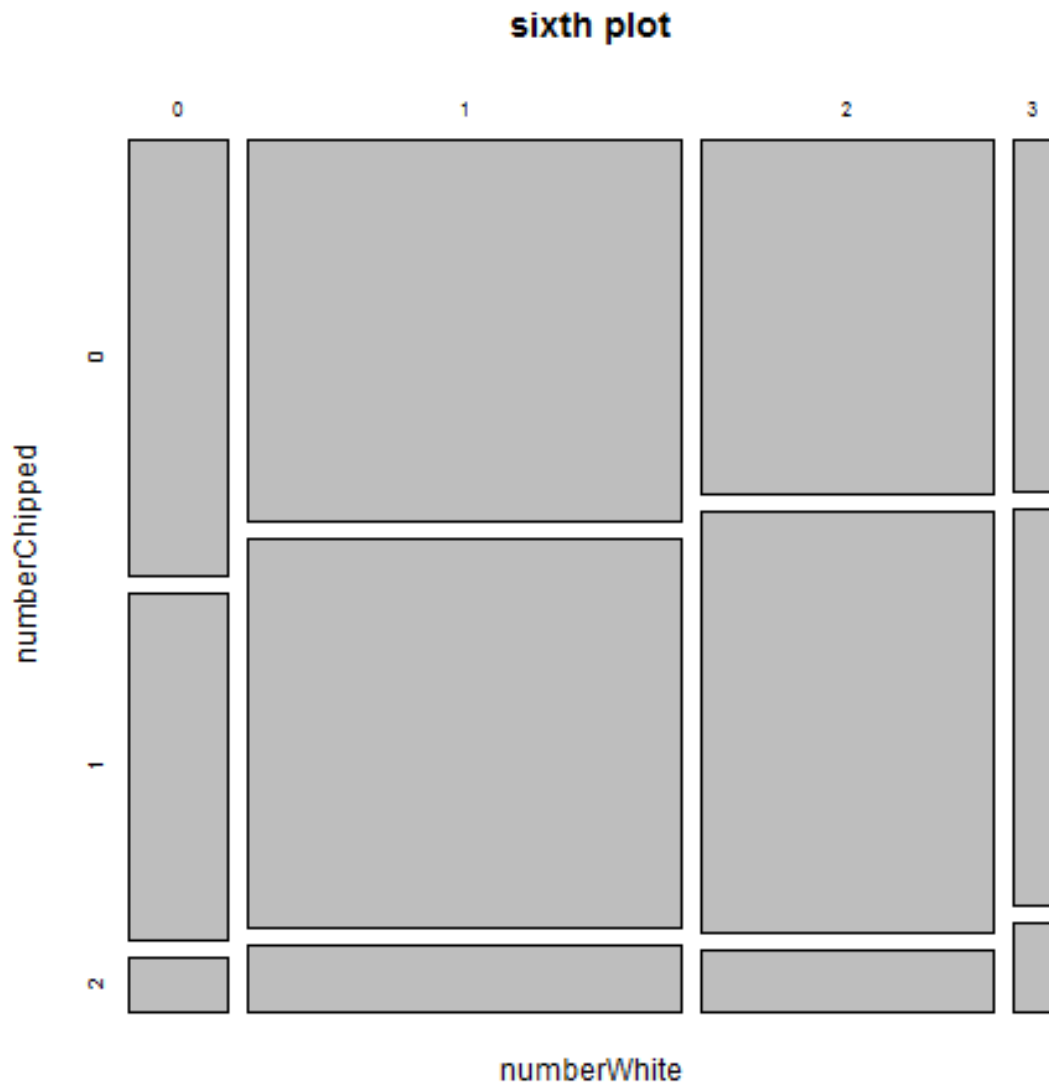


Figure 8: plot of chunk unnamed-chunk-17

Pair-wise scatter plots

```
uData <- rnorm(20)
vData <- rnorm(20,mean=5)
wData <- uData + 2*vData + rnorm(20,sd=0.5)
xData <- -2*uData+rnorm(20,sd=0.1)
yData <- 3*vData+rnorm(20,sd=2.5)

pairs(?)
```

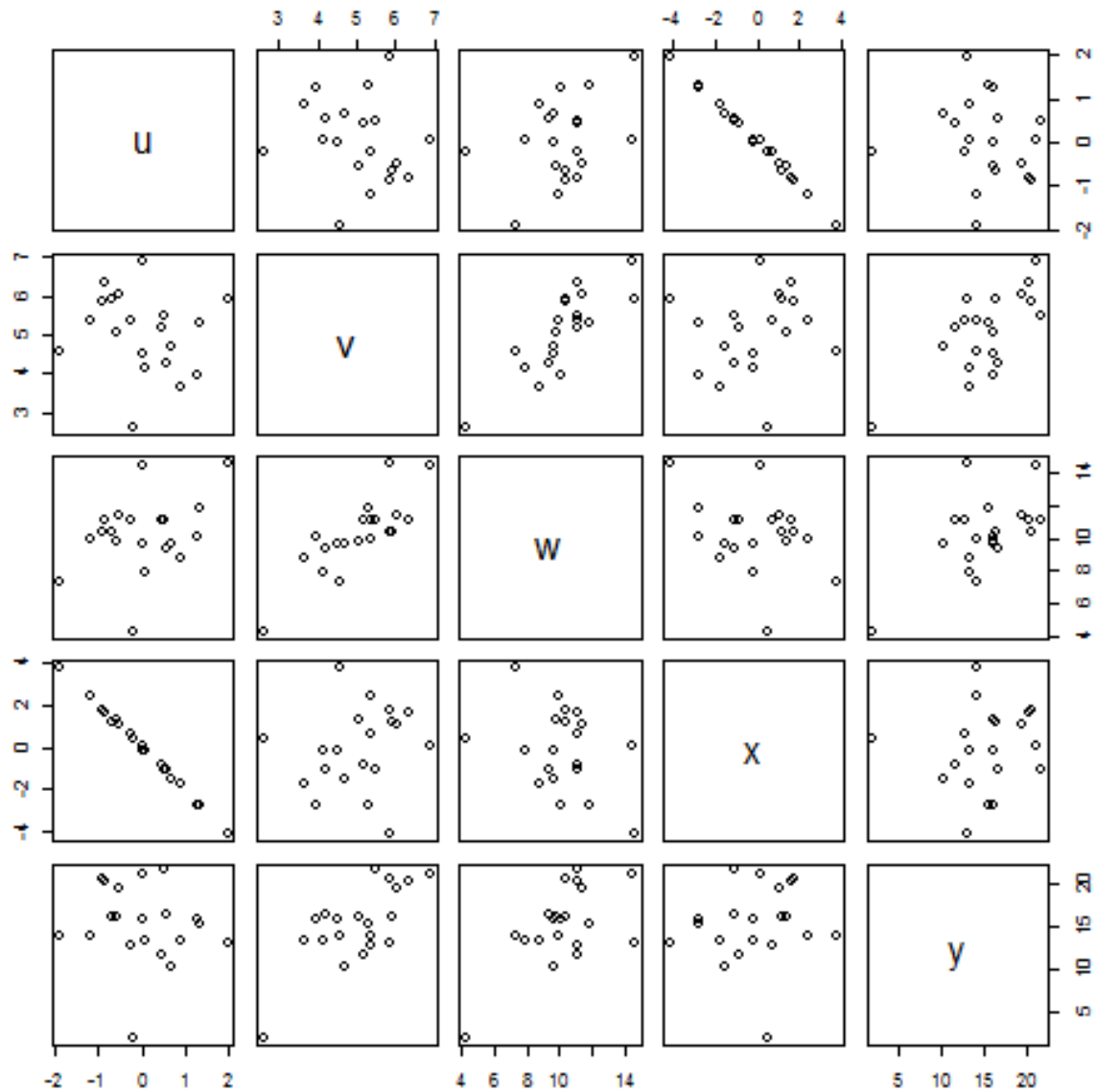


Figure 9: plot of chunk unnamed-chunk-20

Shaded areas

```

polygon(x, y = NULL, density = NULL, angle = 45,          border = NULL, col = NA, lty =
par("lty"),      ..., fillOddEven = FALSE)

stdDev <- 0.75; x <- seq(-5,5,by=0.01); y <- dnorm(x,sd=stdDev)
right <- qnorm(0.95,sd=stdDev)
plot(x,y,type="l",xaxt="n",ylab="p",xlab=expression(paste('Assumed Distribution of',bar(x))),axes=FALSE)
axis(1,at=c(-5,right,0,5), pos = c(0,0),labels=c(expression(' '),expression(bar(x)[cr]),expression(mu[0.
axis(2)
xReject <- seq(right,5,by=0.01); yReject <- dnorm(xReject,sd=stdDev)
polygon(???, col='red')

```

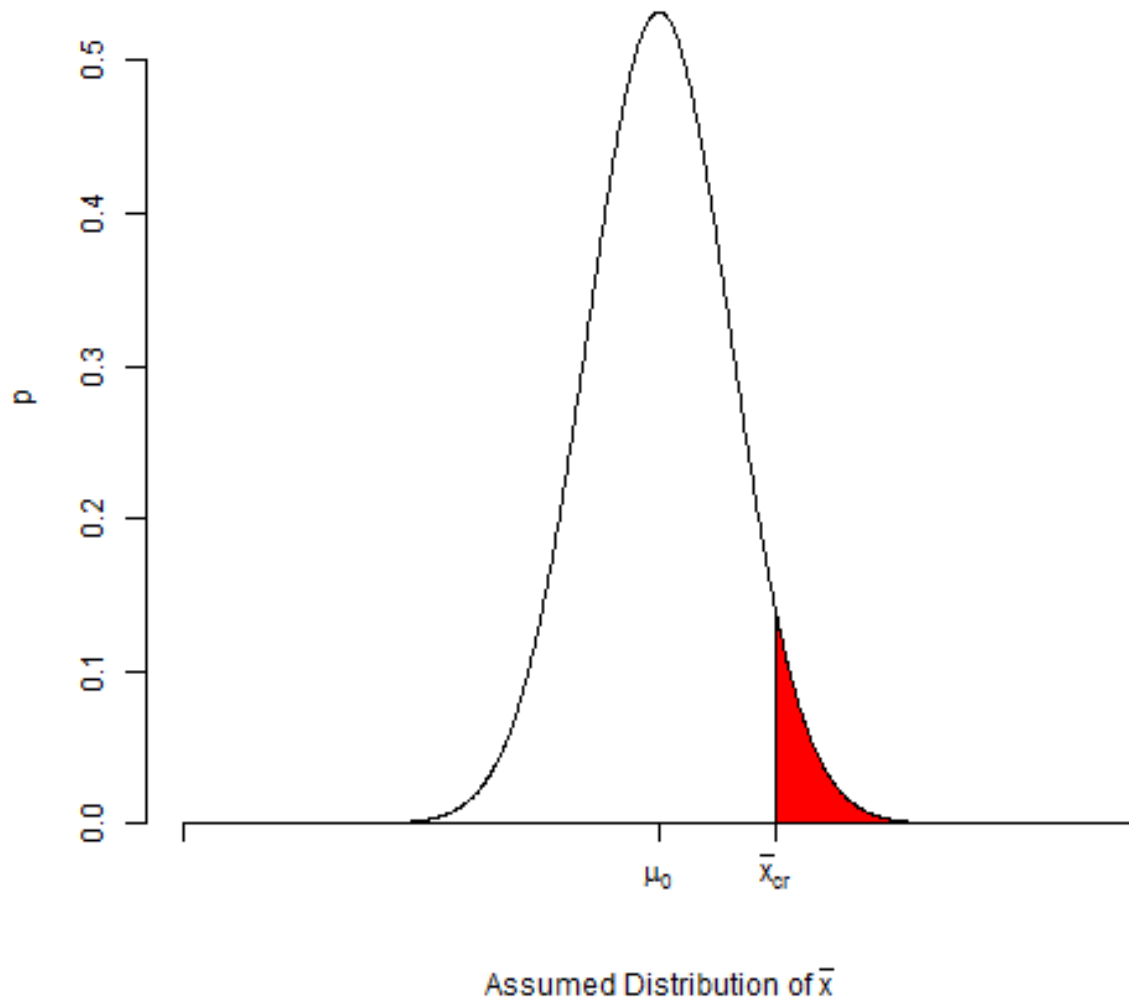


Figure 10: plot of chunk unnamed-chunk-23

Different types of lines

```
par(..., no.readonly = FALSE)
x <- c(1:5); y <- x
par(pch=22, col="red")
par(mfrow=c(2,4))
opts = ???
for(i in 1:length(opts)){
  heading = paste("type=",opts[i])
  plot(x, y, type="n", main=heading)
  lines(x, y, type=opts[i])
}
```

Barplot

```
barplot(height, width = 1, space = NULL, names.arg = NULL, legend.text = NULL,
beside = FALSE, horiz = FALSE, density = NULL, angle = 45, col = NULL,
border = par("fg"), main = NULL, sub = NULL, xlab = NULL, ylab = NULL, xlim
= NULL, ylim = NULL, xpd = TRUE, log = "", axes = TRUE, axisnames = TRUE, cex.axis
= par("cex.axis"), cex.names = par("cex.axis"), inside = TRUE, plot = TRUE,
axis.lty = 0, offset = 0, add = FALSE, args.legend = NULL, ...)
```

```
numberWhite <- rhyper(30,4,5,3)
numberWhite <- as.factor(numberWhite)
```

???

```
numberWhite
0  1  2  3
5 12 12  1
```

plotly

plotly website here

ggplot2: structure

```
ggplot(data = <default data set>, aes(x = <default x axis variable>, y
= <default y axis variable>, ... <other default aesthetic mappings>), ...
<other plot defaults>) + geom_<geom type>(aes(size = <size variable for this
geom>, ... <other aesthetic mappings>), data =
<data for this point geom>, stat = <statistic string or function>,
position = <position string or function>, color = <"fixed color specification">,
<other arguments, possibly passed to the _stat_ function> + scale_<aesthetic>_<type>(name
= <"scale label">, breaks = <where to put tick marks>, labels
= <labels for tick marks>, ... <other options for the scale>) +
theme(plot.background = element_rect(fill = "gray"), ... <other theme elements>)
```

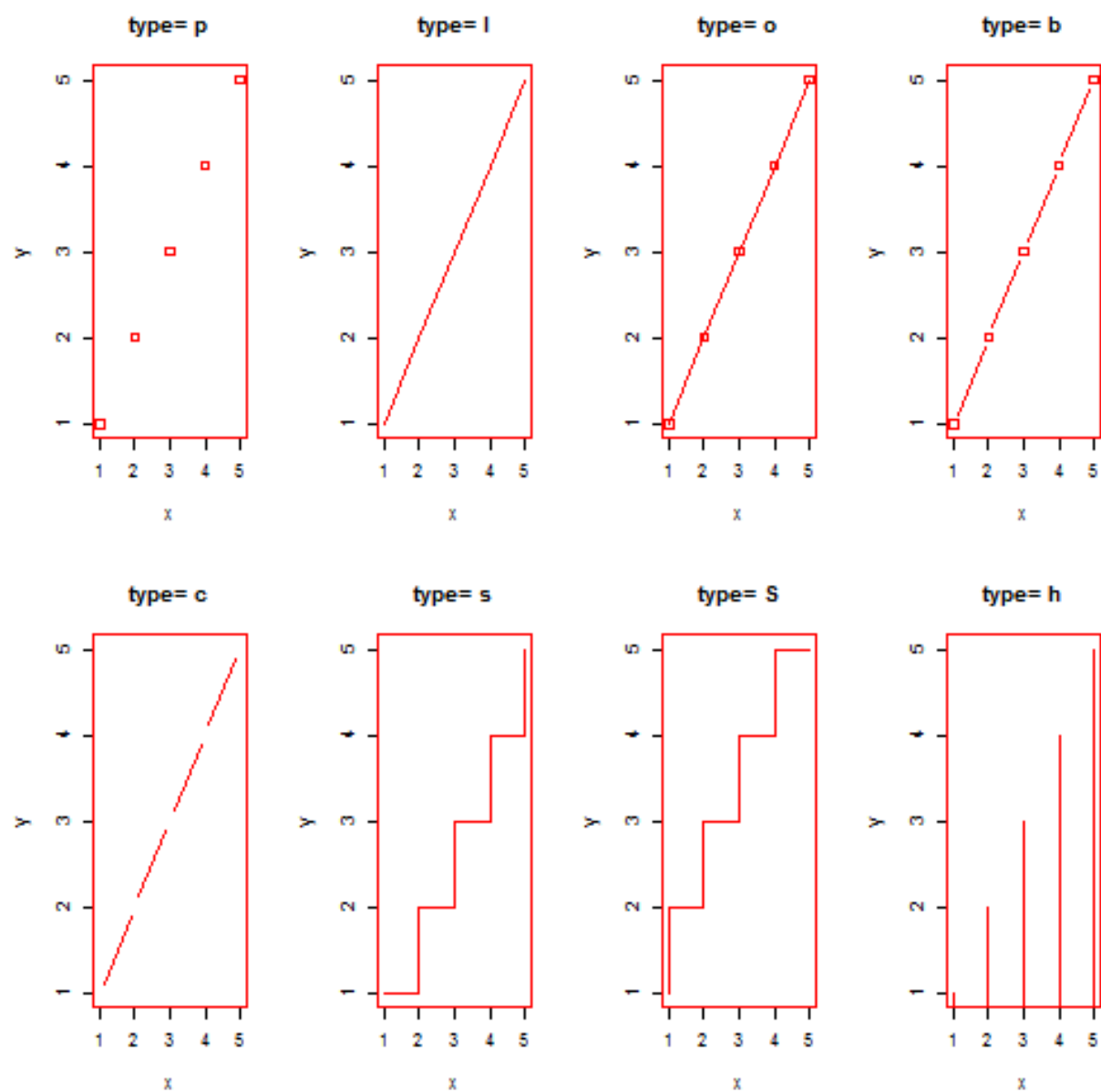


Figure 11: plot of chunk unnamed-chunk-26

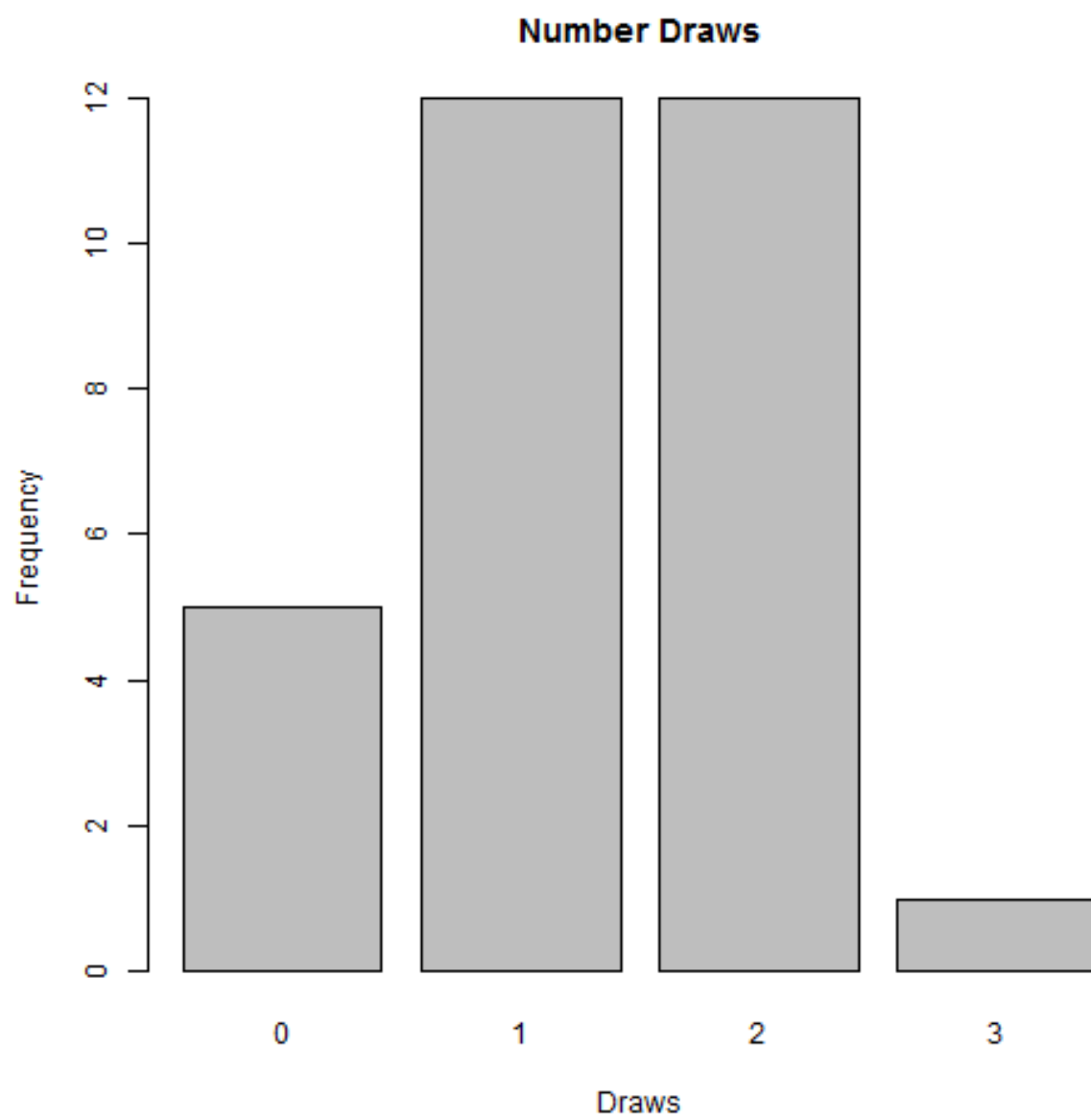


Figure 12: plot of chunk unnamed-chunk-29

qplot: overlapping densities

```
qplot(x, y = NULL, ..., data, facets = NULL, margins = FALSE, geom = "auto", xlim =  
c(NA, NA), ylim = c(NA, NA), log = "", main = NULL, xlab = deparse(substitute(x)),  
ylab = deparse(substitute(y)), asp = NA, stat = NULL, position = NULL)
```

```
library(ggplot2)  
data("mtcars")  
mtcars$gear <- factor(mtcars$gear, levels=c(3,4,5), labels=c("3gears", "4gears", "5gears"))  
mtcars$am <- factor(mtcars$am, levels=c(0,1), labels=c("Automatic", "Manual"))  
mtcars$cyl <- factor(mtcars$cyl, levels=c(4,6,8), labels=c("4cyl", "6cyl", "8cyl"))  
  
qplot(mpg,  
      data = ??,  
      geom = ??,  
      fill = ??,  
      alpha = ??,  
      main = "Distribution of Gas Milage", xlab="Miles Per Gallon", ylab="Density")
```

qplot: facets and points

```
library(ggplot2)  
data("mtcars")  
mtcars$gear <- factor(mtcars$gear, levels=c(3,4,5), labels=c("3gears", "4gears", "5gears"))  
mtcars$am <- factor(mtcars$am, levels=c(0,1), labels=c("Automatic", "Manual"))  
mtcars$cyl <- factor(mtcars$cyl, levels=c(4,6,8), labels=c("4cyl", "6cyl", "8cyl"))  
  
qplot(??,  
      ??,  
      data=mtcars, shape=, color=, facets=gear~cyl, size=I(3), xlab="Horsepower", ylab="Miles per Gallon")
```

hist vs geom_histogram

Rgraphics tutorial [here](#)

```
par(mfrow=c(2,1))  
hist(diamonds$carat)  
hist(diamonds$carat, breaks = 500)
```

ggplot2: Box plot and points with legend

```
plot(carat ~ clarity,  
     data=subset(diamonds, cut == "Good"))  
points(carat ~ clarity, col="red",  
       data=subset(diamonds, cut == "Ideal"))  
legend(0,3,  
      c("Good", "Ideal"), title="cut",
```

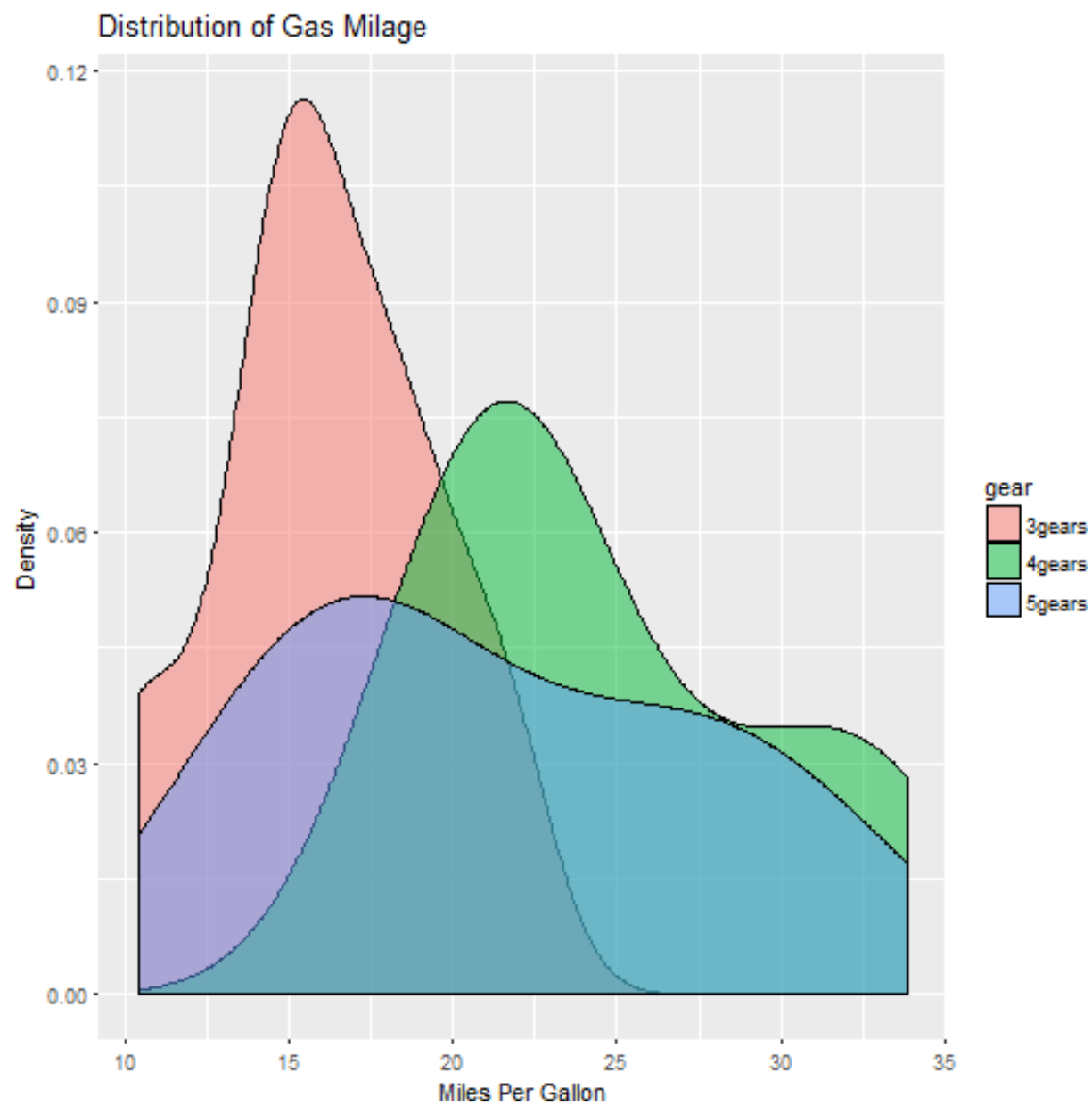


Figure 13: plot of chunk unnamed-chunk-32

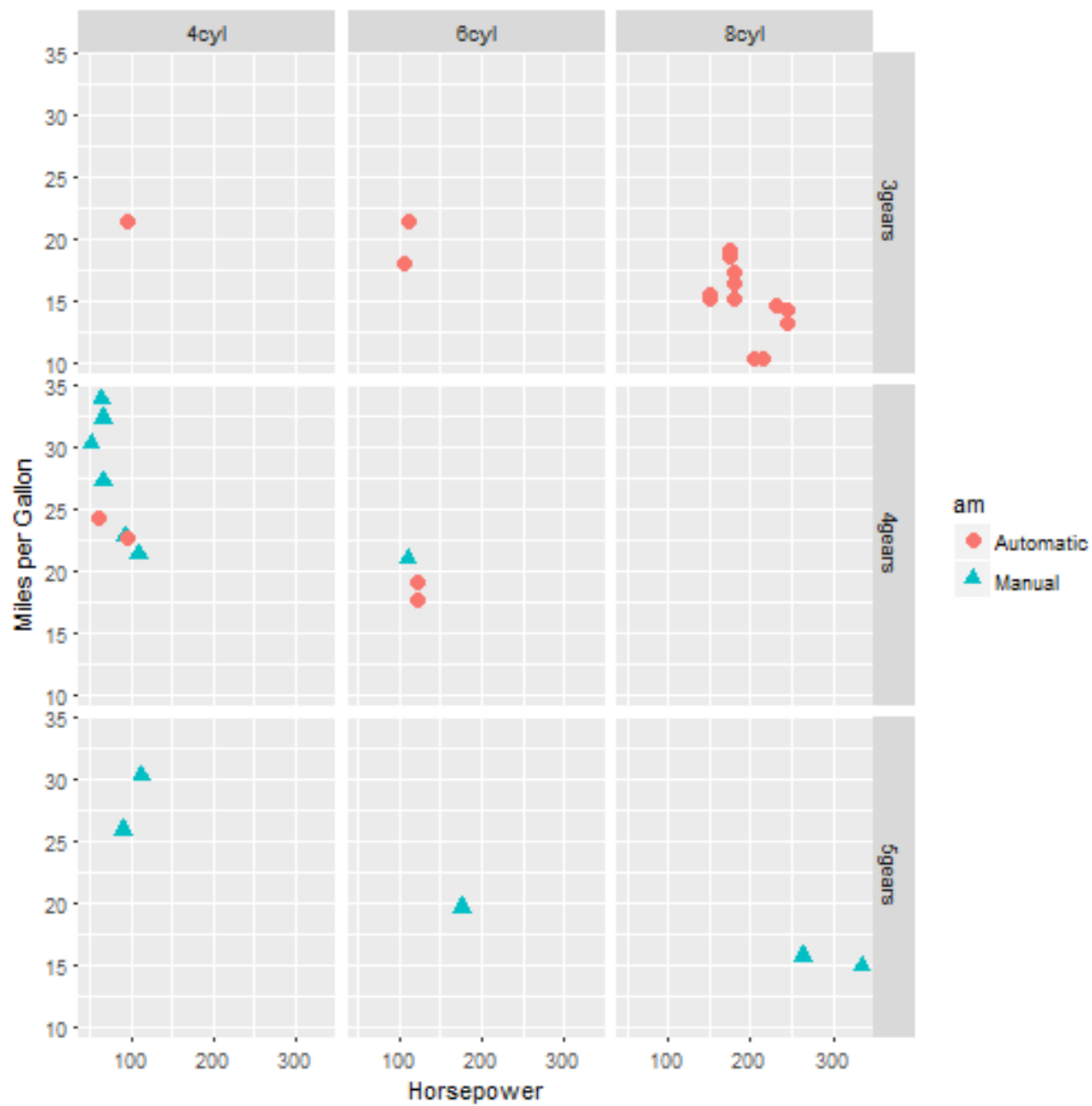


Figure 14: plot of chunk unnamed-chunk-35

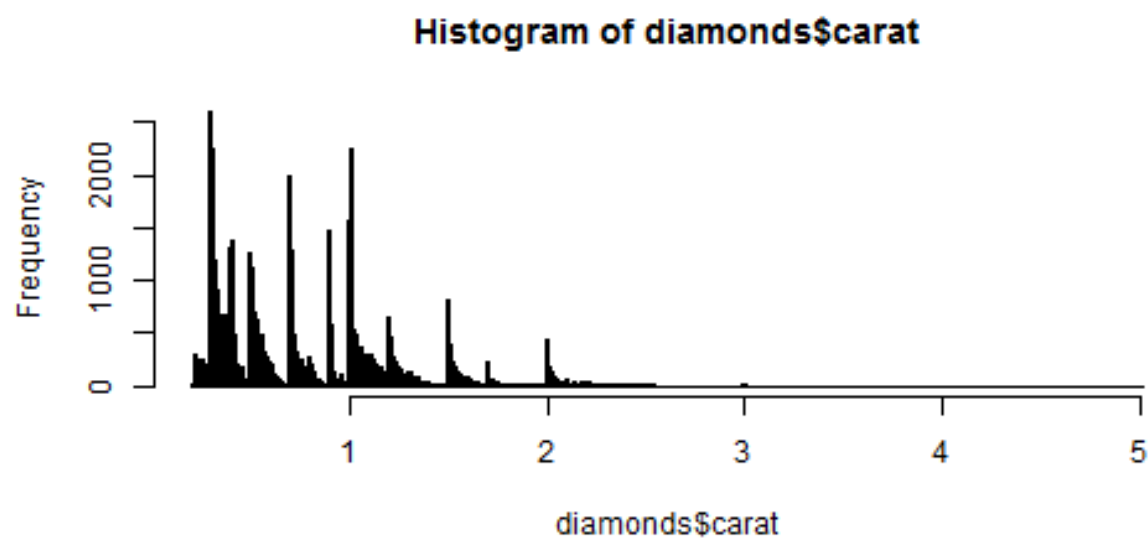
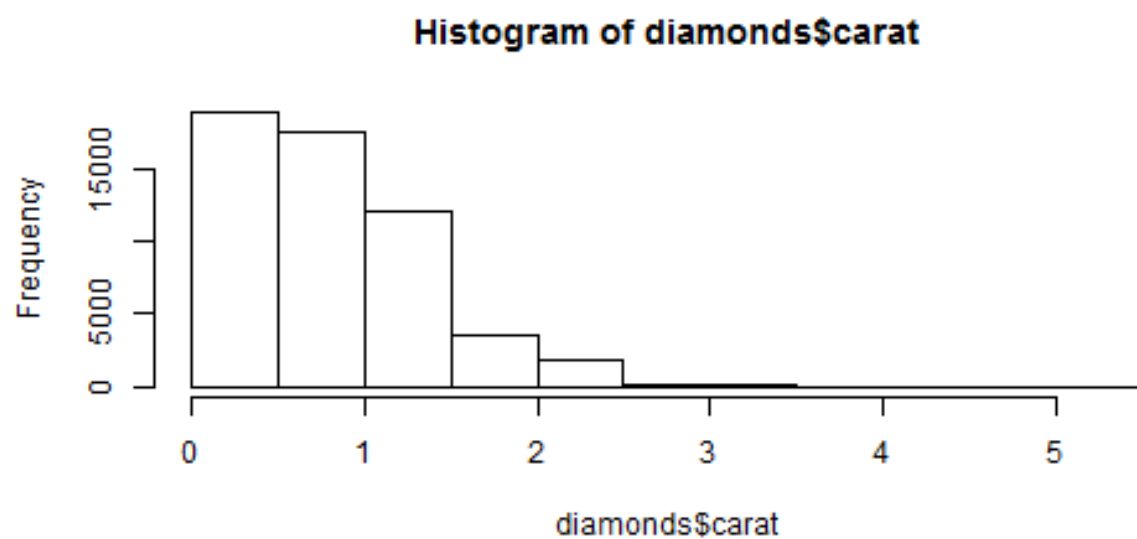


Figure 15: plot of chunk unnamed-chunk-37

```
col=c("black", "red"),
pch=c(1, 1))
```

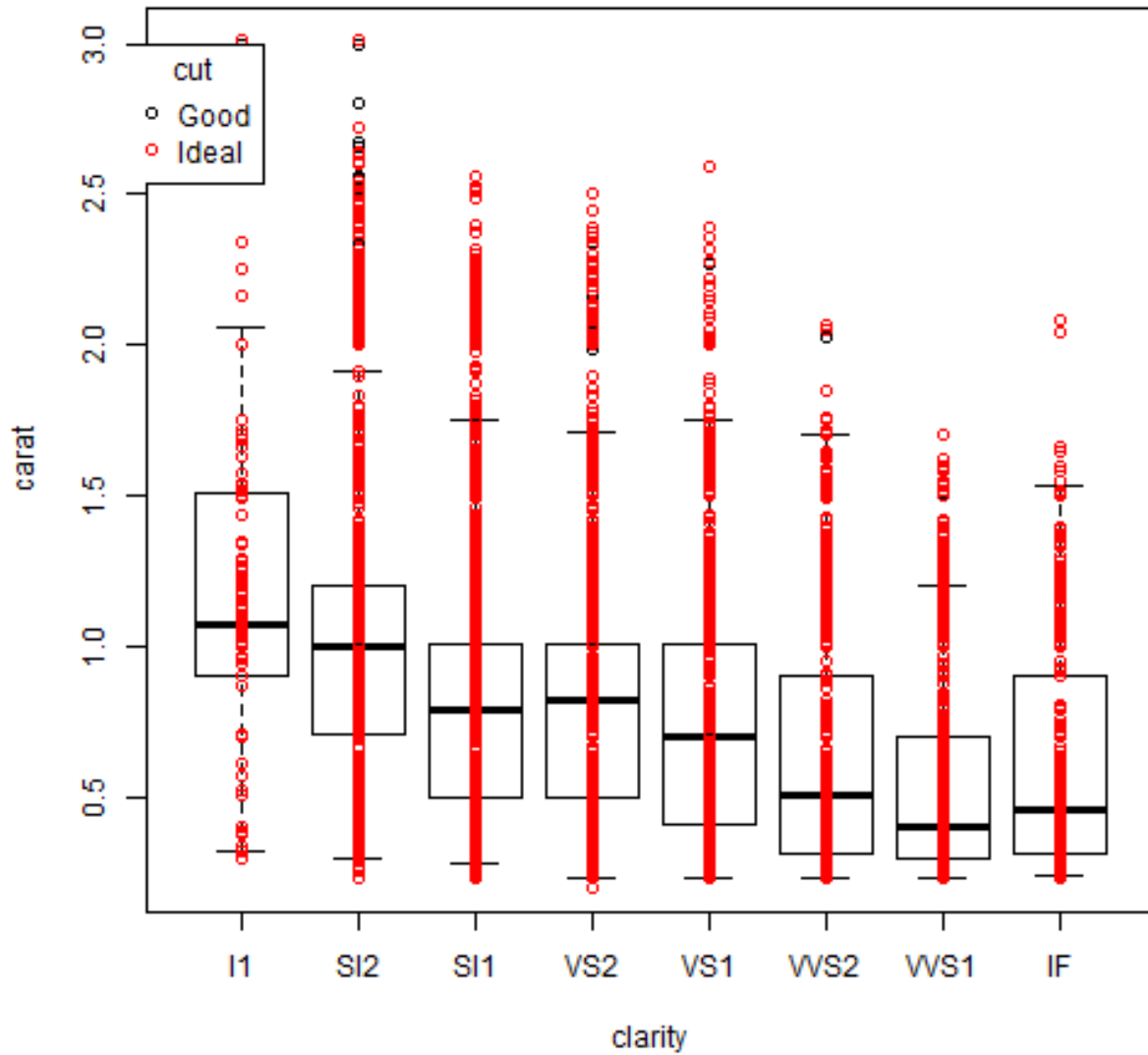


Figure 16: plot of chunk unnamed-chunk-39

ggplot2

```
ggplot(subset(diamonds, cut %in% c("Good", "Ideal")),
  aes(x=clarity,
    y=carat,
```

```
color=cut))+  
geom_point()
```

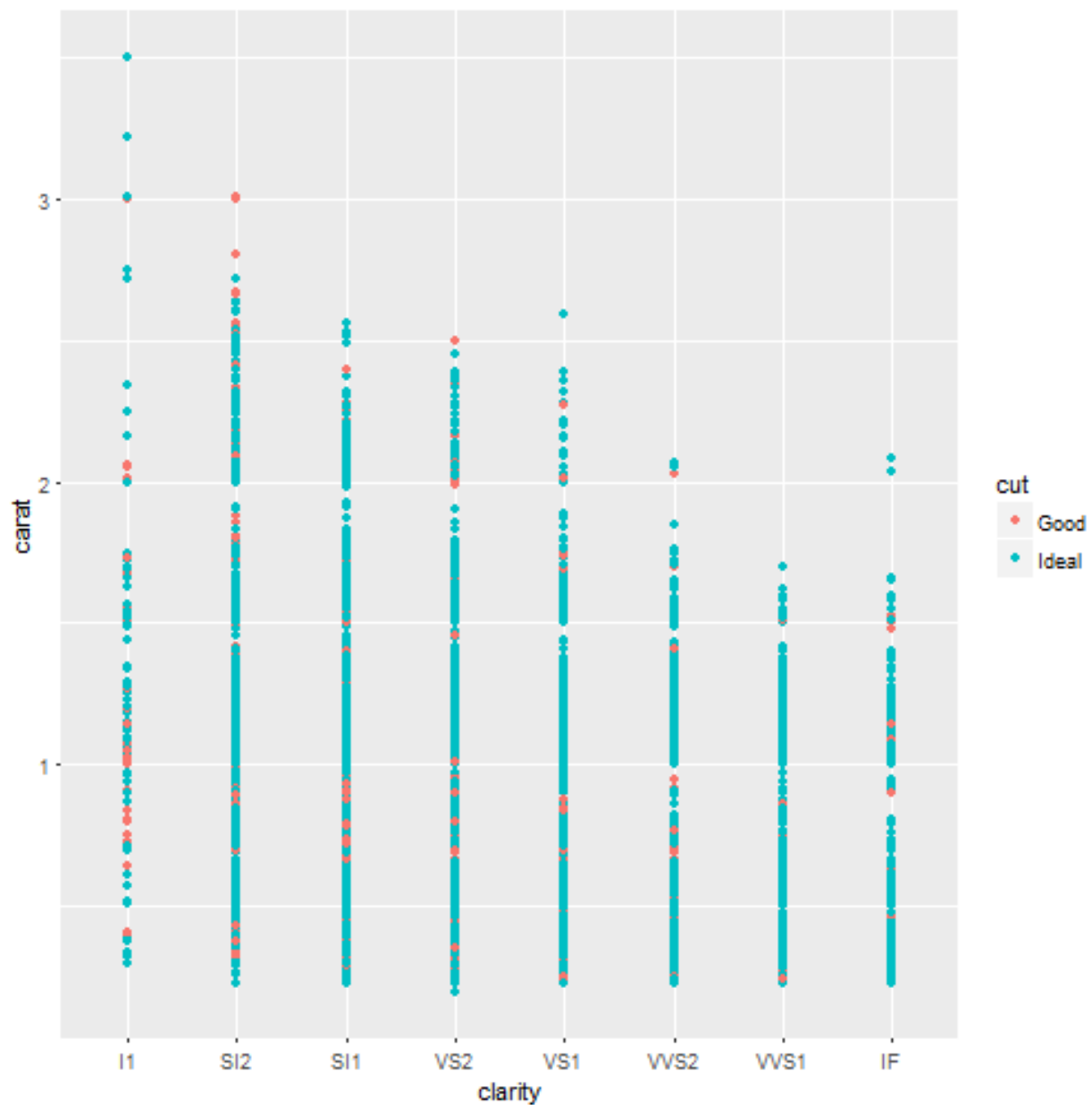


Figure 17: plot of chunk unnamed-chunk-40

Symbols

```
## A look at all 25 symbols  
df2 <- data.frame(x = 1:5 , y = 1:25, z = 1:25)  
s <- ggplot(df2, aes(x = x, y = y))  
s + geom_point(aes(shape = z), size = 4) + scale_shape_identity()
```

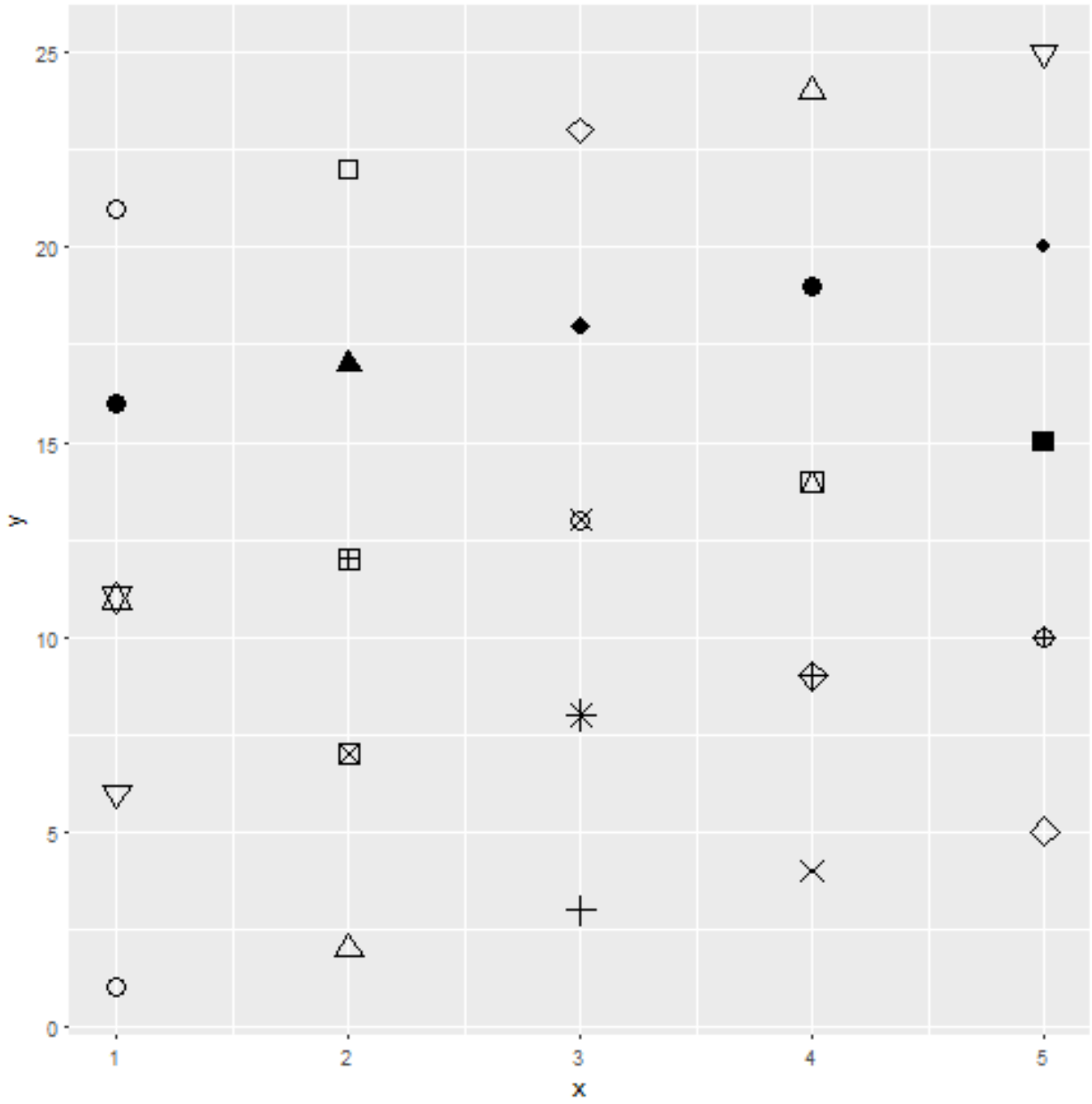


Figure 18: plot of chunk unnamed-chunk-41

```
## While all symbols have a foreground colour, symbols 19-25 also take a  
## background colour (fill)
```

Symbols (2)

```
s + geom_point(aes(shape = z), size = 4, colour = "Red") +  
  scale_shape_identity()
```

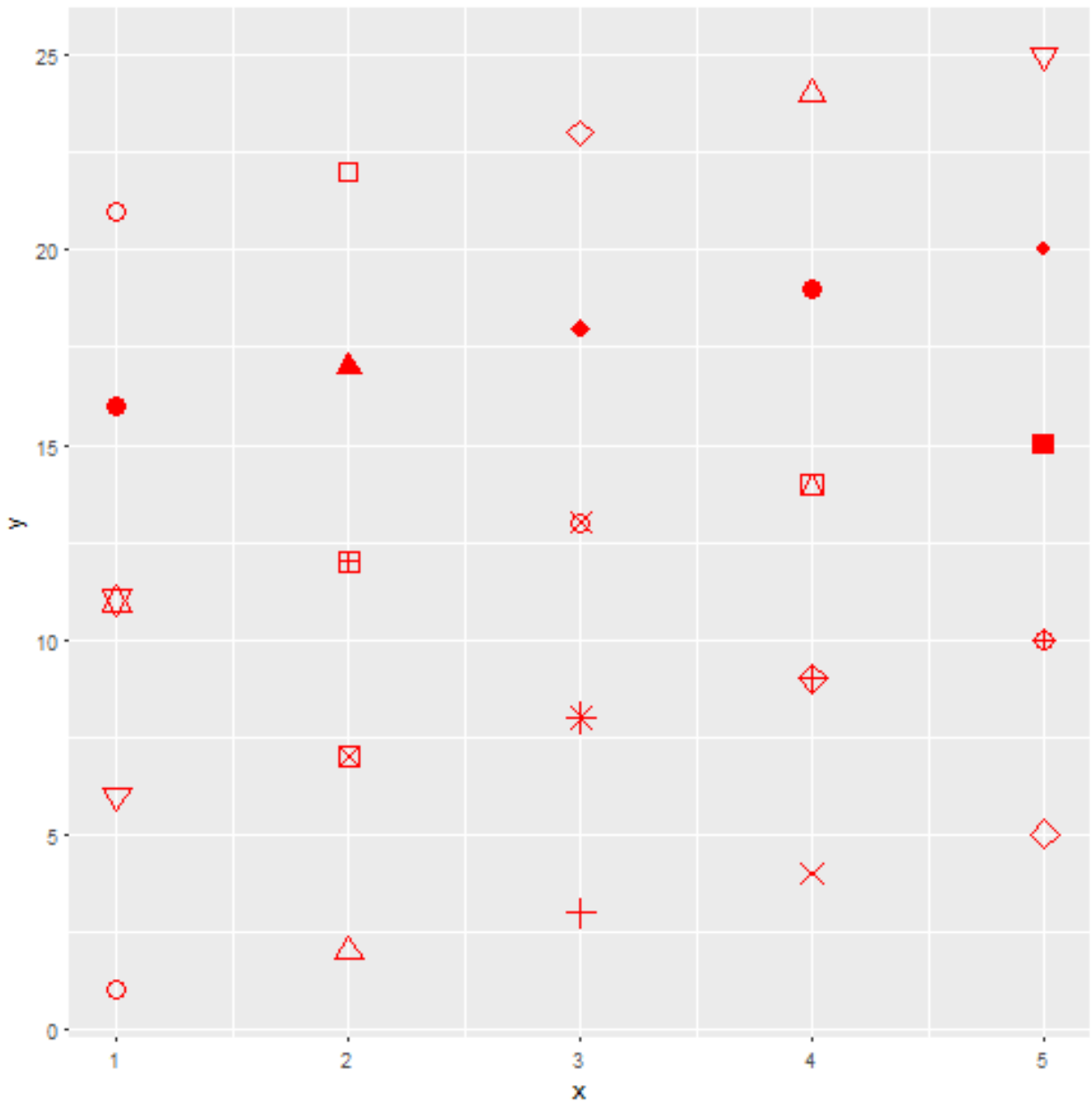


Figure 19: plot of chunk unnamed-chunk-42

Symbols (3)

```
s + geom_point(aes(shape = z), size = 4, colour = "Red", fill = "Black") +  
  scale_shape_identity()
```

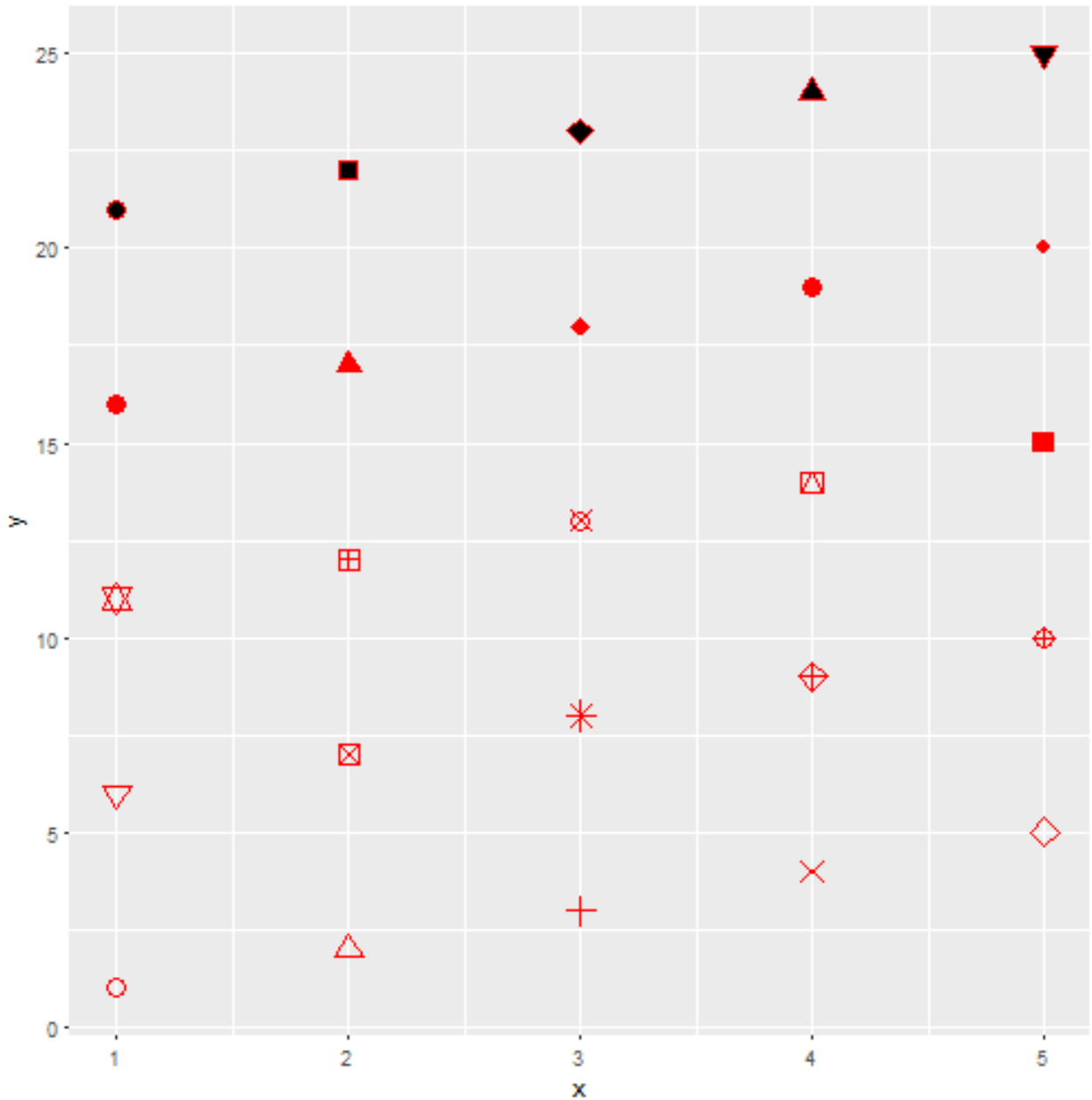


Figure 20: plot of chunk unnamed-chunk-43

ColorBrewer2

ColorBrewer2

Manipulate

manipulate examples

Options()

- Sets and reports options. Lots of them.
-
- `digits` from 3.1415927 using `options(digits = 2)` to 3.14
- Report value using e.g. `getOption("digits")`
-
- `scipen()`

```
R> ran2 <- c(1.810032e+09, 4)
R> options("scipen"=-100, "digits"=4)
R> ran2
[1] 1.81e+09 4.00e+00
R> options("scipen"=100, "digits"=4)
R> ran2
[1] 1810032000
```

Kable::

```
library(knitr)
kable(head(iris), format = "latex")
```

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
5.1	3.5	1.4	0.2	setosa
4.9	3.0	1.4	0.2	setosa
4.7	3.2	1.3	0.2	setosa
4.6	3.1	1.5	0.2	setosa
5.0	3.6	1.4	0.2	setosa
5.4	3.9	1.7	0.4	setosa

```
kable(head(iris), format = "html")
```

Sepal.Length

Sepal.Width

Petal.Length

Petal.Width

Species

5.1

3.5

1.4

0.2

setosa

4.9

3.0

1.4

0.2

setosa

4.7

3.2

1.3

0.2

setosa

4.6

3.1

1.5

0.2

setosa

5.0

3.6

1.4

0.2

setosa

5.4

3.9

1.7

0.4

setosa

printr::

```
x = matrix(rnorm(40), 5)

dimnames(x) = list(NULL, head(LETTERS, ncol(x)))

knitr::kable(x, digits = 2,
              caption = "A table produced by printr.")
```

A	B	C	D	E	F	G	H
0.65	-2.14	0.43	0.09	-0.69	-1.74	-1.05	0.12
-0.60	1.00	-1.08	-0.16	1.53	1.77	0.36	-1.18

A	B	C	D	E	F	G	H
-0.59	0.39	0.36	1.67	0.43	0.33	1.69	-0.08
0.07	-0.31	-0.25	0.28	1.41	-1.87	-1.87	-0.74
-1.51	-0.74	-2.01	-1.20	-0.03	0.23	0.28	0.90

Tables

```
library(xtable)

options(xtable.floating = FALSE)
options(xtable.timestamp = "")

data(tli)

xtable(tli[1:10, ])

% latex table generated in R 3.3.3 by xtable 1.8-2 package
%
\begin{tabular}{rrlllr}
\hline
& grade & sex & disadvg & ethnicity & tlimth \\
\hline
1 & 6 & M & YES & HISPANIC & 43 \\
2 & 7 & M & NO & BLACK & 88 \\
3 & 5 & F & YES & HISPANIC & 34 \\
4 & 3 & M & YES & HISPANIC & 65 \\
5 & 8 & M & YES & WHITE & 75 \\
6 & 5 & M & NO & BLACK & 74 \\
7 & 8 & F & YES & HISPANIC & 72 \\
8 & 4 & M & YES & BLACK & 79 \\
9 & 6 & M & NO & WHITE & 88 \\
10 & 7 & M & YES & HISPANIC & 87 \\
\hline
\end{tabular}
```

GLMs

```
fm3 <- glm(disadvg ~ ethnicity*grade, data = tli, family = binomial)
xtable(fm3)

% latex table generated in R 3.3.3 by xtable 1.8-2 package
%
\begin{tabular}{rrrrrr}
\hline
& Estimate & Std. Error & z value & Pr(>|z|) \\
\hline
(Intercept) & 3.1888 & 1.5966 & 2.00 & 0.0458 \\
ethnicityHISPANIC & -0.2848 & 2.4808 & -0.11 & 0.9086 \\
ethnicityOTHER & 212.1701 & 22122.7093 & 0.01 & 0.9923 \\
\hline
\end{tabular}
```

```

ethnictyWHITE & -8.8150 & 3.3355 & -2.64 & 0.0082 \\
grade & -0.5308 & 0.2892 & -1.84 & 0.0665 \\
ethnictyHISPANIC:grade & 0.2448 & 0.4357 & 0.56 & 0.5742 \\
ethnictyOTHER:grade & -32.6014 & 3393.4687 & -0.01 & 0.9923 \\
ethnictyWHITE:grade & 1.0171 & 0.5185 & 1.96 & 0.0498 \\
\hline
\end{tabular}

```

Flat tables

```

data(mtcars)
mtcars$cyl <- factor(mtcars$cyl, levels = c("4","6","8"),
labels = c("four","six","eight"))
tbl <- ftable(?, row.vars = c(2, 4), dnn = c("Cylinders", "V/S", "Transmission", "Gears"))
tbl
xftbl <- xtableFtable(tbl)
print.xtableFtable(xftbl)

```

		Cylinders	four	six	eight			
		Transmission	0	1	0	1	0	1
V/S	Gears							
0	3		0	0	0	0	12	0
	4		0	0	0	2	0	0
	5		0	1	0	1	0	2
1	3		1	0	2	0	0	0
	4		2	6	2	0	0	0
	5		0	1	0	0	0	0

% latex table generated in R 3.3.3 by xtable 1.8-2 package

```

%
\begin{tabular}{llll |rrrrrr}
\hline
& & & & Cylinders & & \multicolumn{1}{l}{four} & & \multicolumn{1}{l}{ } & & \multicolumn{1}{l}{ } & & \multicolumn{1}{l}{ } \\
& & & & Transmission & & \multicolumn{1}{l}{0} & & \multicolumn{1}{l}{1} & & \multicolumn{1}{l}{ } & & \multicolumn{1}{l}{ } \\
V/S & Gears & & & & & \multicolumn{1}{l}{ } & & \multicolumn{1}{l}{ } & & \multicolumn{1}{l}{ } & & \multicolumn{1}{l}{ } \\
\hline
0 & 3 & & & & & 0 & 0 & 0 & 0 & 12 & 0 & \\
& 4 & & & & & 0 & 0 & 0 & 2 & 0 & 0 & \\
& 5 & & & & & 0 & 1 & 0 & 1 & 0 & 2 & \\
1 & 3 & & & & & 1 & 0 & 2 & 0 & 0 & 0 & \\
& 4 & & & & & 2 & 6 & 2 & 0 & 0 & 0 & \\
& 5 & & & & & 0 & 1 & 0 & 0 & 0 & 0 & \\
\hline
\end{tabular}

```

Flat tables

```

data(mtcars)
mtcars$cyl <- factor(mtcars$cyl, levels = c("4","6","8"),
labels = c("four","six","eight"))

```

```
tbl <- ftable(mtcars$cyl, mtcars$vs, mtcars$am, mtcars$gear, row.vars = c(2, 4), dnn = c("Cylinders", "Transmission", "V/S", "Gears"))
tbl
```

		Cylinders		four		six		eight	
		Transmission		0	1	0	1	0	1
V/S	Gears								
0	3			0	0	0	0	12	0
	4			0	0	0	2	0	0
	5			0	1	0	1	0	2
1	3			1	0	2	0	0	0
	4			2	6	2	0	0	0
	5			0	1	0	0	0	0

```
xftbl <- xtableFtable(tbl)
print.xtableFtable(xftbl)
```

```
% latex table generated in R 3.3.3 by xtable 1.8-2 package
```

```
%
\begin{tabular}{llll |rrrrrr}
\hline
& & & & \multicolumn{2}{four} & \multicolumn{2}{ } & \multicolumn{2}{ } \\
& & & & \multicolumn{2}{0} & \multicolumn{2}{1} & \multicolumn{2}{ } \\
V/S & Gears & & & \multicolumn{2}{ } & \multicolumn{2}{ } & \multicolumn{2}{ } \\
\hline
0 & 3 & & & 0 & 0 & 0 & 0 & 12 & 0 \\
& 4 & & & 0 & 0 & 0 & 2 & 0 & 0 \\
& 5 & & & 0 & 1 & 0 & 1 & 0 & 2 \\
1 & 3 & & & 1 & 0 & 2 & 0 & 0 & 0 \\
& 4 & & & 2 & 6 & 2 & 0 & 0 & 0 \\
& 5 & & & 0 & 1 & 0 & 0 & 0 & 0 \\
\hline
\end{tabular}
```

Markdown tables

```
library(pander)
pandoc.table(tli)
```

grade	sex	disadv	ethnicity	timth
6	M	YES	HISPANIC	43
7	M	NO	BLACK	88
5	F	YES	HISPANIC	34
3	M	YES	HISPANIC	65
8	M	YES	WHITE	75
5	M	NO	BLACK	74

6	F	NO	BLACK	82
4	M	NO	WHITE	69
3	F	YES	HISPANIC	17
3	M	NO	HISPANIC	37
7	M	NO	WHITE	83
6	M	YES	HISPANIC	78
6	F	NO	WHITE	84

```
pandoc.table(tli, style="rmarkdown")
```

grade	sex	disadv	ethnicity	timth
6	M	YES	HISPANIC	43
7	M	NO	BLACK	88
5	F	YES	HISPANIC	34
6	F	NO	WHITE	91
5	F	NO	WHITE	50
7	M	NO	WHITE	83
4	F	YES	BLACK	58
4	M	YES	HISPANIC	85
7	F	NO	WHITE	52
5	M	NO	WHITE	86
4	F	YES	BLACK	79
8	M	NO	WHITE	48
5	M	NO	WHITE	91
3	M	YES	HISPANIC	89
7	F	NO	WHITE	91
5	F	YES	WHITE	79
7	M	NO	WHITE	83
6	M	YES	HISPANIC	78
6	F	NO	WHITE	84

Other HTML table

```
library("htmlTable")
htmlTable(tli)
```

```
grade
sex
disadv
ethnicity
```



```
addHeader(rtf,title="Section Header", subtitle="This is the subheading or section text.")
```

add paragraph

```
addParagraph(rtf,"This is a new self-contained paragraph.\n")
```

```
addNewLine(rtf)
```

```
addParagraph(rtf,"Normal, \\b this is bold\\b0, normal.\n")
```

```
addParagraph(rtf,"Normal, {\\b\\i bold-italic}, normal.\n")
```

add a table

```
tab <- as.data.frame(head(iris)) # create a data.frame  
colnames(tab)<-gsub("\\.", " ",colnames(tab)) # format column names  
addTable(rtf,tab,font.size=9,row.names=FALSE,NA.string="-")
```

R and Word (2)

add plot

```
addPlot(rtf,plot.fun=plot,width=6,height=6,res=300, iris[,1],iris[,2])
```

```
addPageBreak(rtf, width=8.5, height=11, omi=c(1,1,1,1))
```

```
addSessionInfo(rtf)
```

```
done(rtf)
```

rtf output

ReporteRs::

- <http://davidgohel.github.io/ReporteRs/index.html>

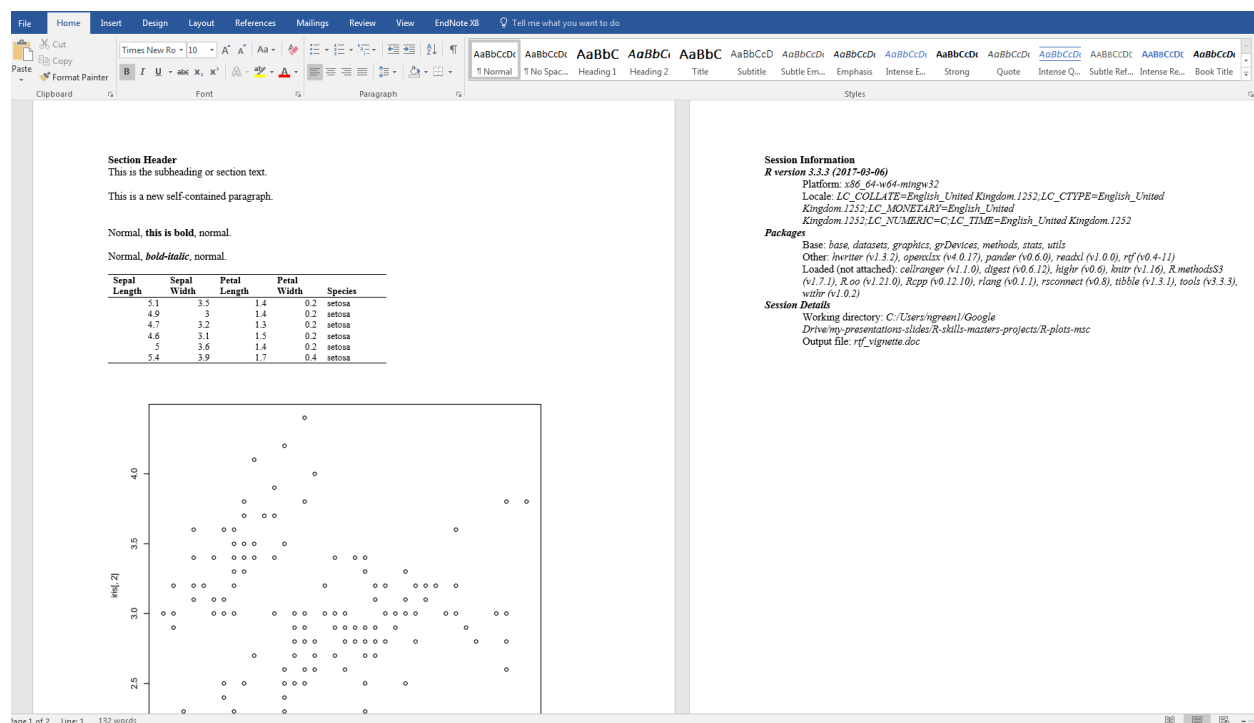


Figure 21: