

# Line R-rt

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# Outline

- 1 line art
- 2 Examples
- 3 Create a Blank Sheet of Paper
- 4 Inside the Plot Region
  - points
  - arrows
  - text
  - lines, curves
  - polygon
  - rectangles
- 5 plotmath
- 6 Are you looking for skills to practice?

# Outline

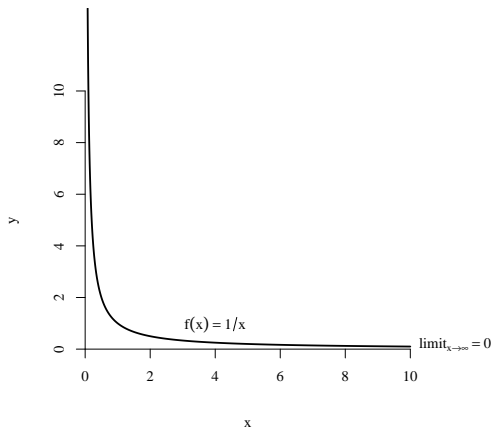
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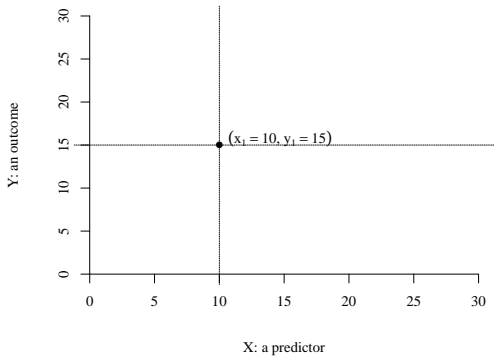
# graphics!

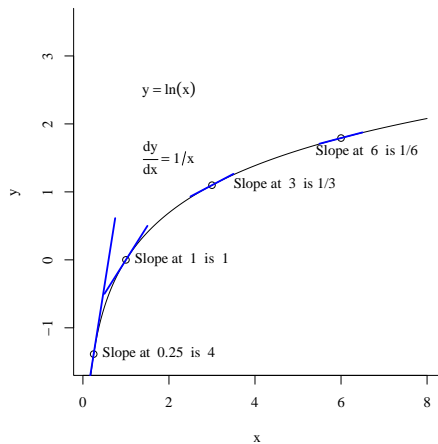
- In papers and reports, we often need technical illustrations
- Publishers refer to illustrations of this sort as “line art”, it must be supplied by authors in high-resolution graphics files (pdf, svg, tiff, etc)
- One can draw sketches by hand, of course, but almost nobody can make a publishable drawing by hand anymore
- R(R Core Team, 2017) offers a suite of functions that can be used to create artwork.

# Outline

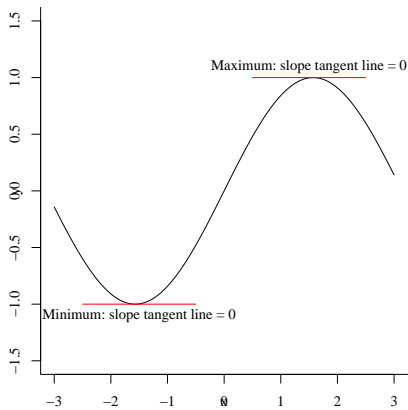
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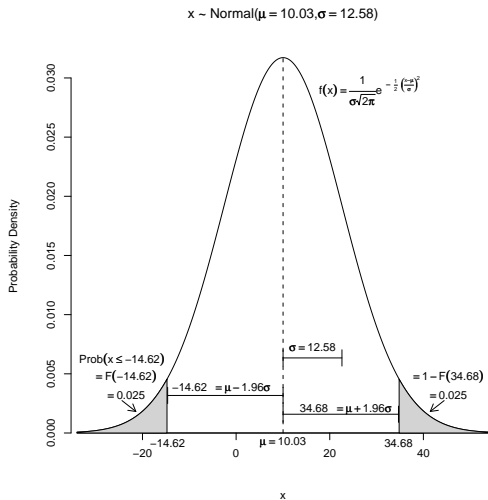


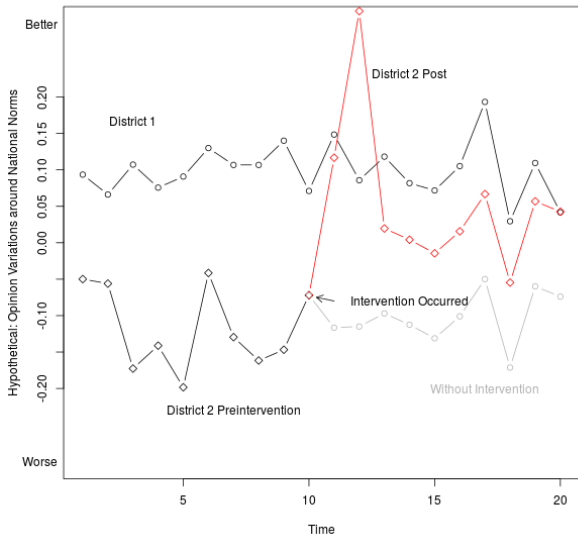


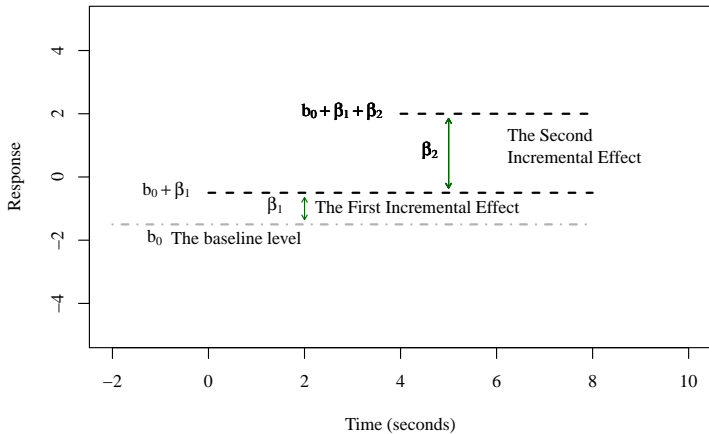




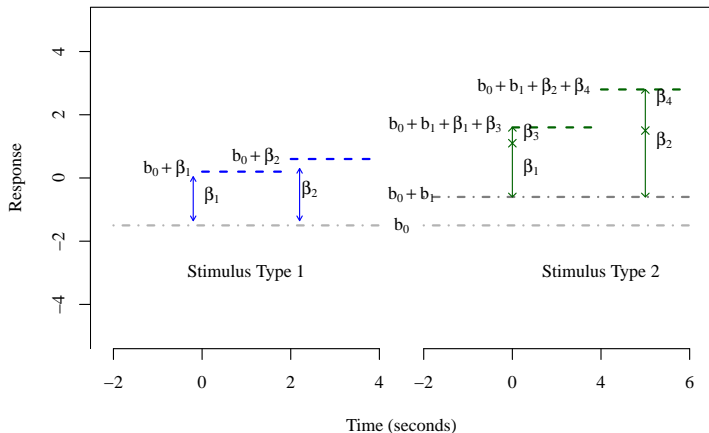


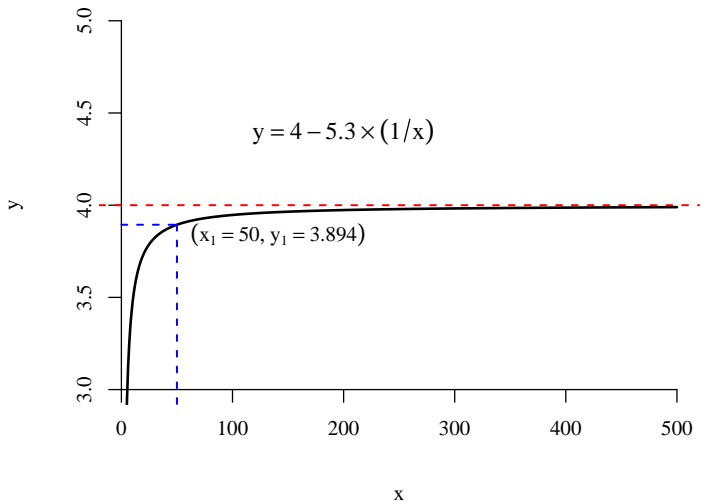




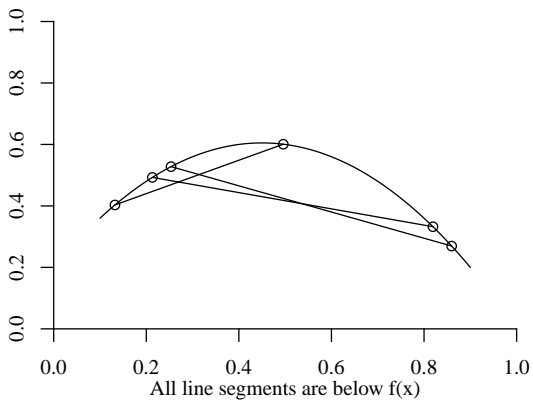


## Comparing Stimulus Types: Shared Baseline Approach

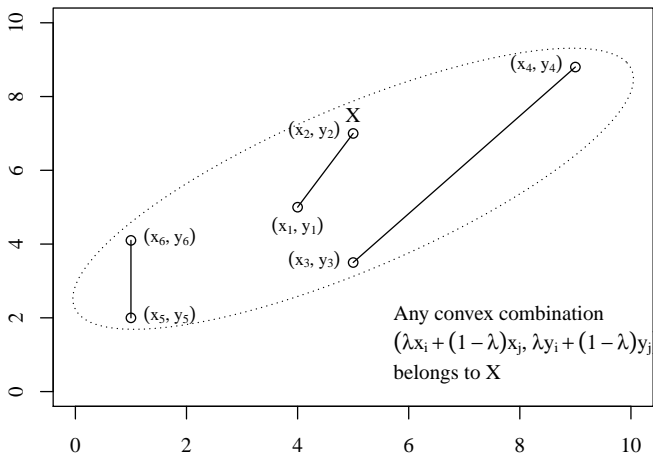




## A Concave Down Function



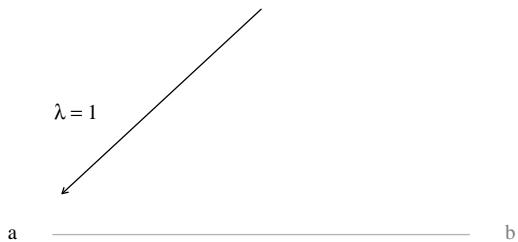
## A Convex Set





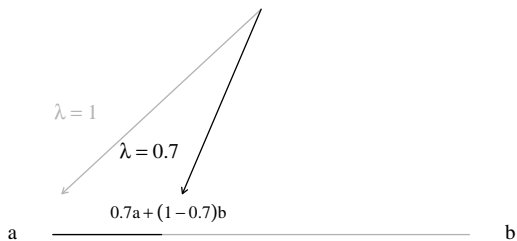
The convex combination

$$\lambda a + (1 - \lambda)b$$



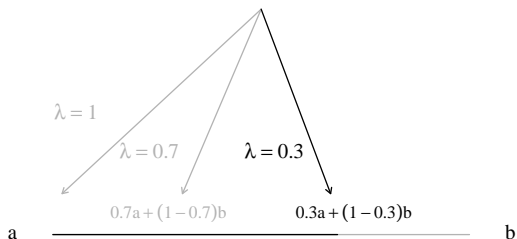
## The convex combination

$$\lambda a + (1 - \lambda)b$$



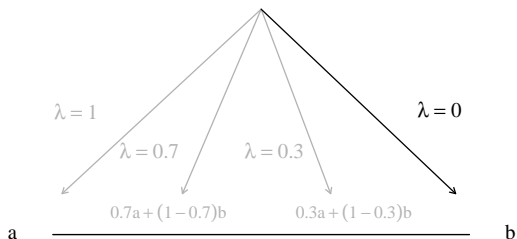
## The convex combination

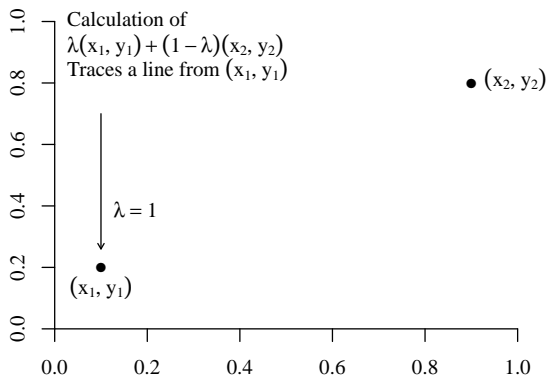
$$\lambda a + (1 - \lambda)b$$

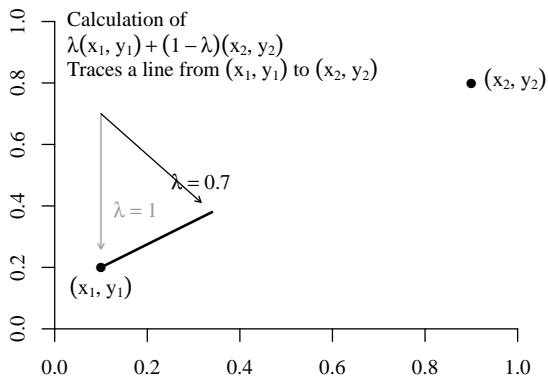


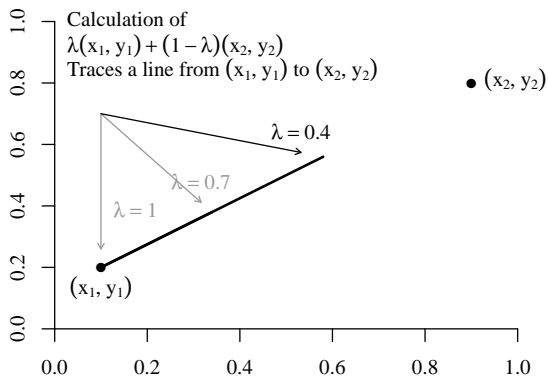
## The convex combination

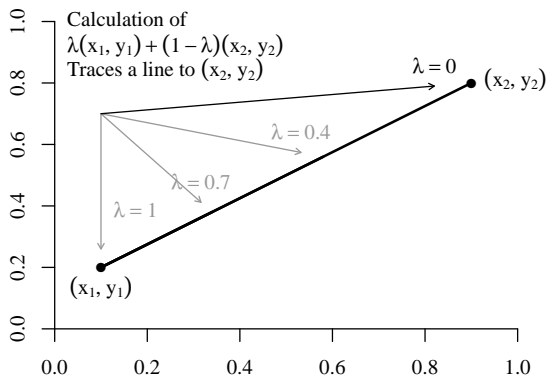
$$\lambda a + (1 - \lambda)b$$



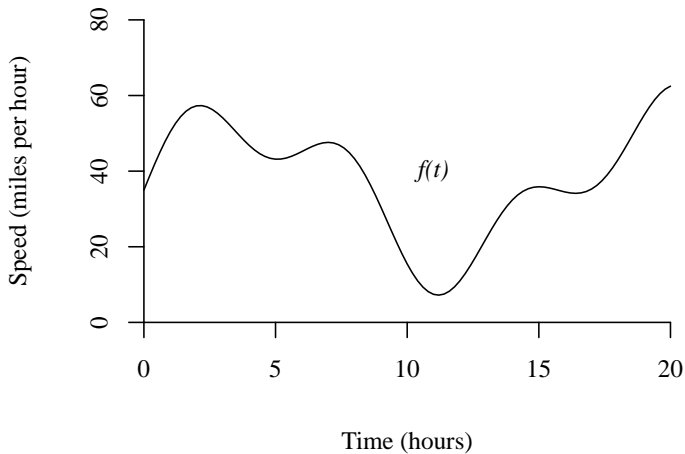


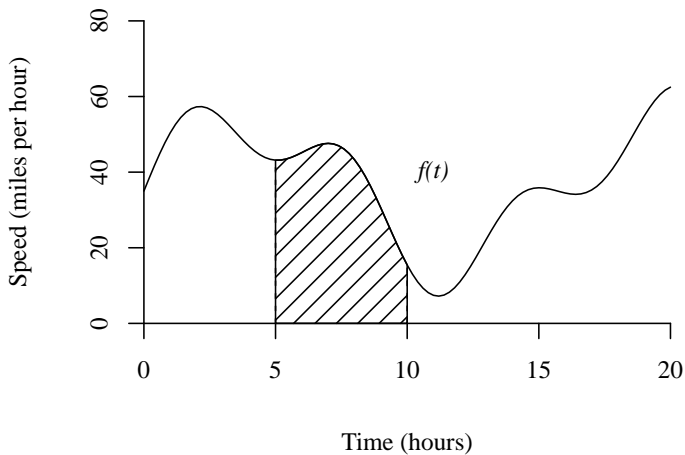


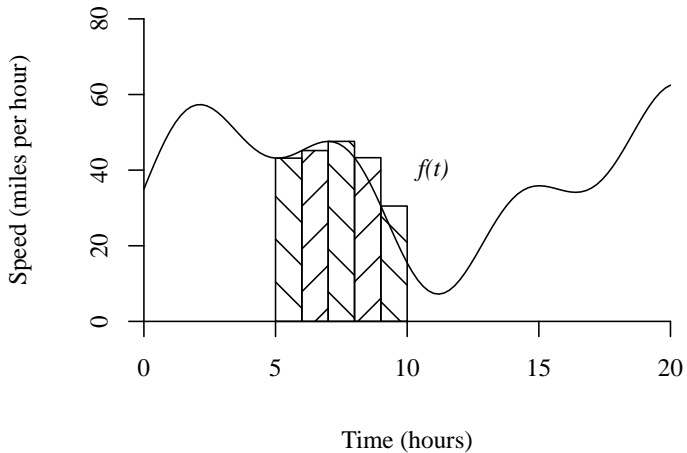


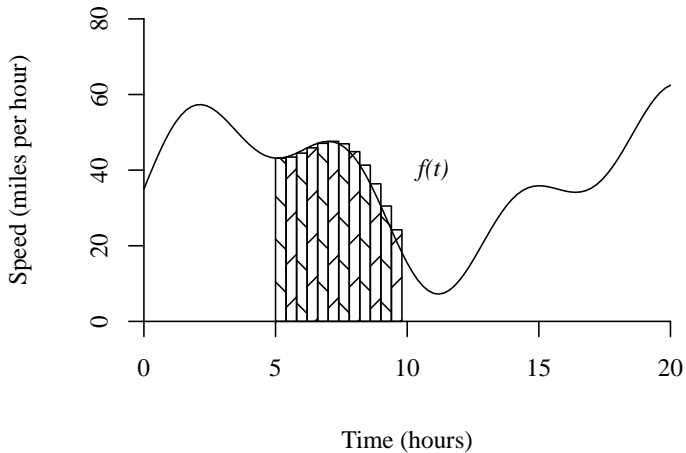


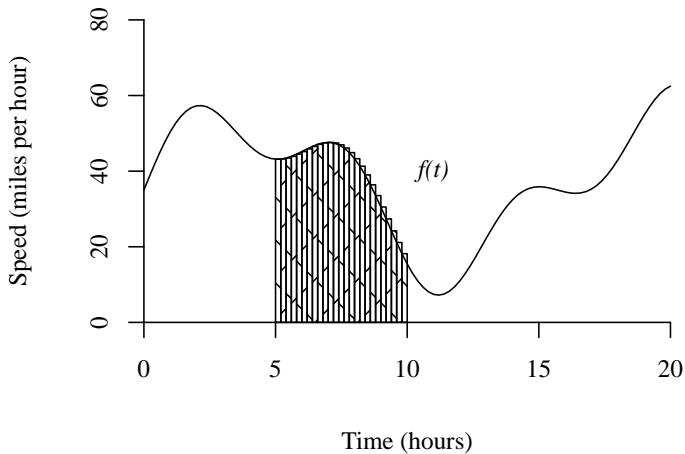




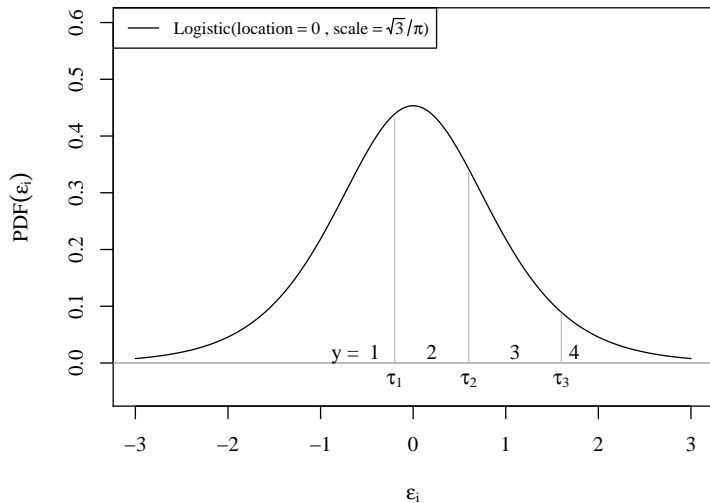




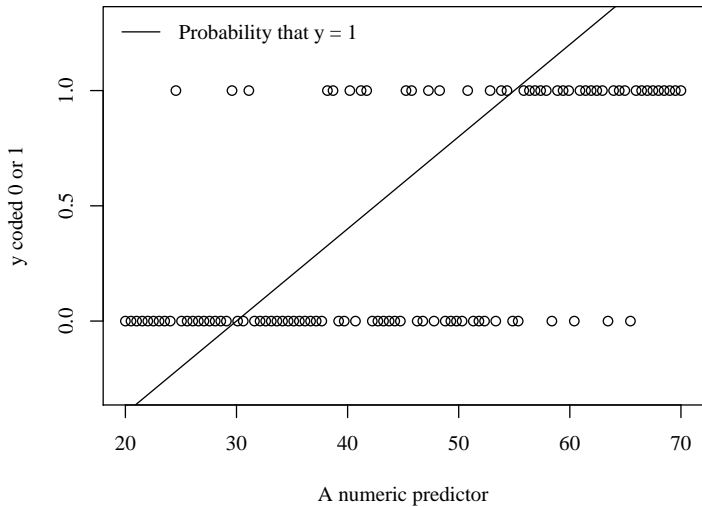




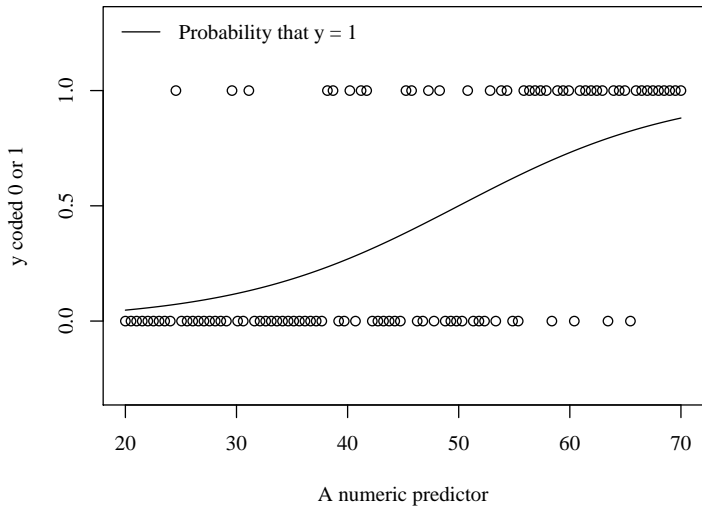
# From a Logistic Regression Lecture



## Straight Line is Not Right. Right?

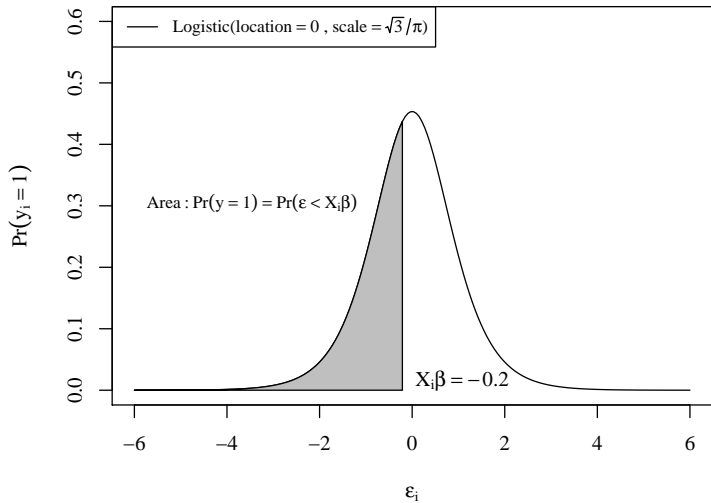


## I'd Rather Have An S-shaped Curve

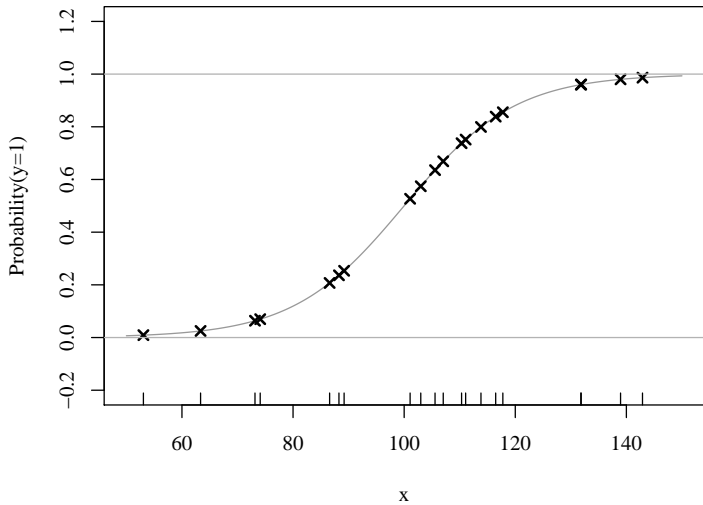


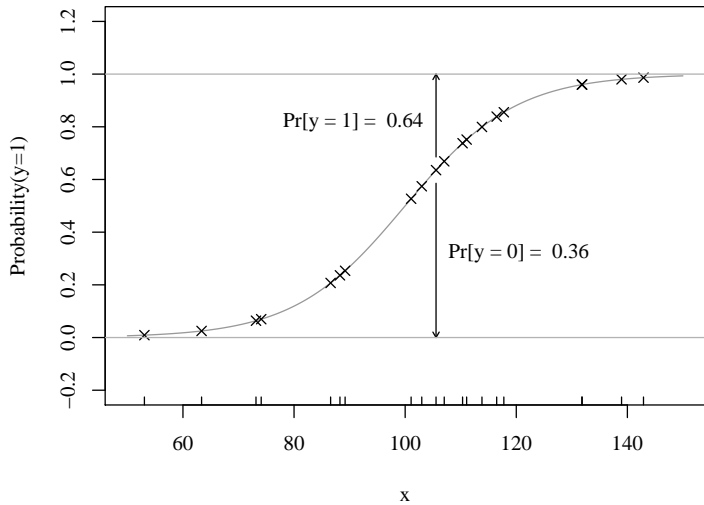


## Logistic Probability that $Y = 1$

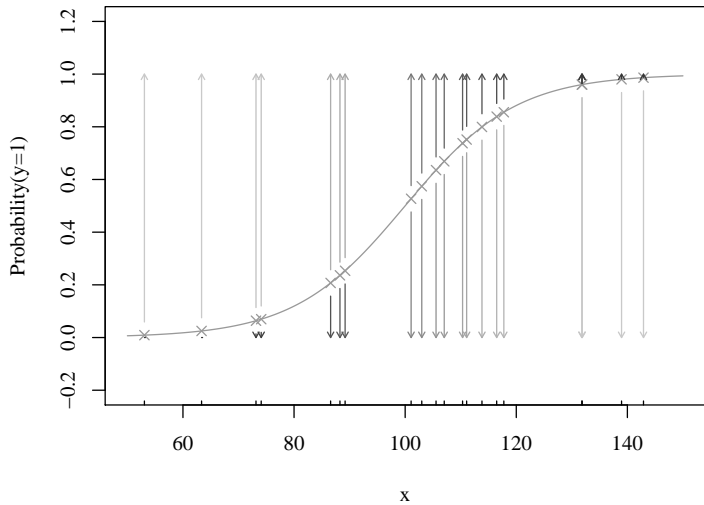


## The "true" probability that $y = 1$





## Darker Arrow Points to More Likely Outcome



# Source Code Available

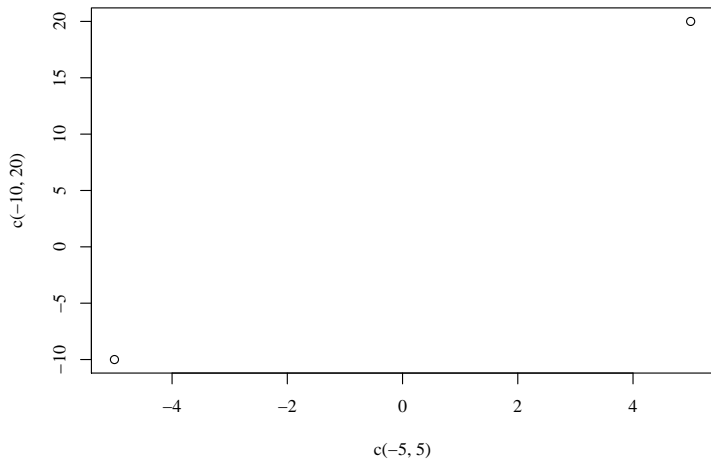
- The R files I used to produce these graphs are in the R folder distributed with this project
- The output files (displayed above) are in the output folder

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- Get a separate “device” display window
  - `dev.new()`
  - If in RStudio, `dev.new()` is blocked, must run
    - MS windows: `windows()`
    - Mac: `quartz()`
    - Linux: `X11()`
- Create a drawing region inside there.
  - I choose to have x scale go from -5 to +5 and y from -10 to 20

```
plot(x = c(-5, 5), y = c(-10, 20))
```



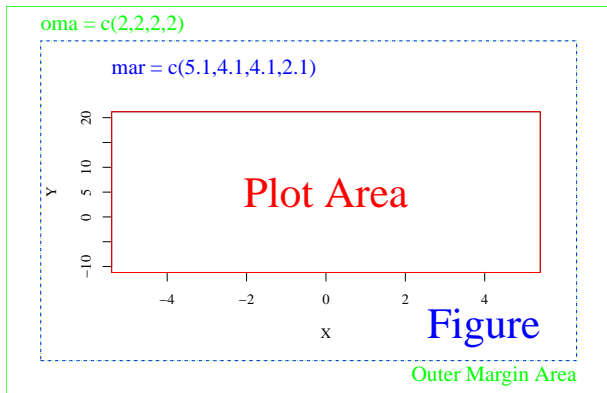


# Oops, I forgot to hide border, axes, labels, and points

```
plot(x = c(-5, 5), y = c(-10, 20), type = "n",  
     axes = FALSE, xlab = "", ylab = "")
```

# Result: blank sheet of paper

# Draw inside the Plot Area



## Defaults

- margins asymmetric (measure: lines of text)
- most commands we use write only in the Plot Area

# Here is the plan of attack

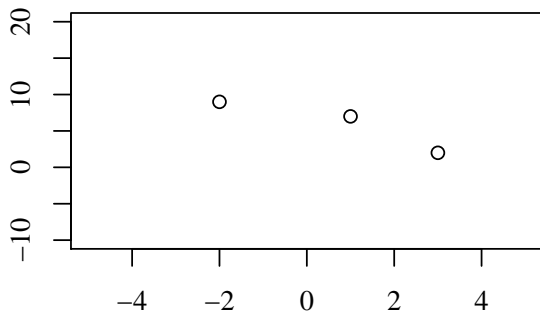
Demonstrate various drawing functions in R. For each we need to

- 1 Run blank sheet creator
- 2 Draw on the sheet
- 3 Save or Throw away that sheet.
- 4 Start over.  
(There is no eraser!)

# Outline

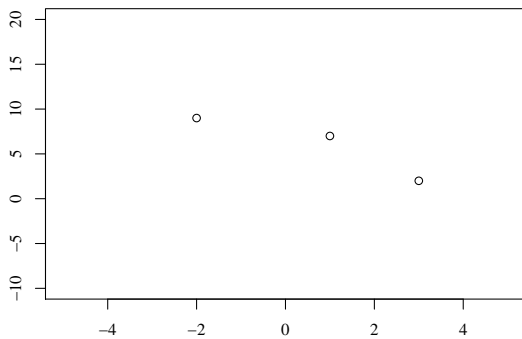
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```
plot(x = c(-5, 5), y = c(-10, 20),  
     type = "n", xlab = "", ylab = "")  
points(x = c(-2, 1, 3), y = c(9, 7, 2))
```



Create x and y vectors separately

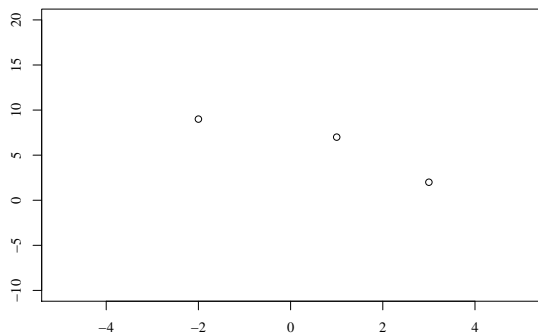
```
plot(x = c(-5, 5), y = c(-10, 20), type = "n",  
      xlab = "", ylab = "")  
x <- c(-2, 1, 3)  
y <- c(9, 7, 2)  
points(x = x, y = y)  
## Same result as  
## points(x, y)  
## because of R positional matches
```



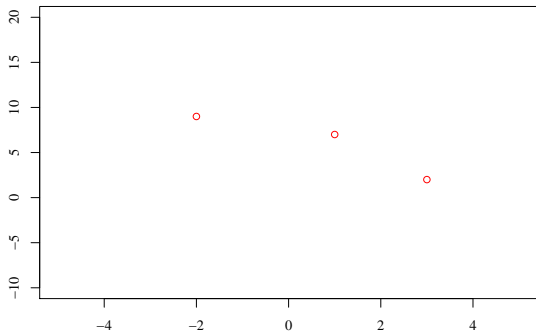


`bdraw()` is a little function, it re-draws the graph area for me (same as typing plot command)

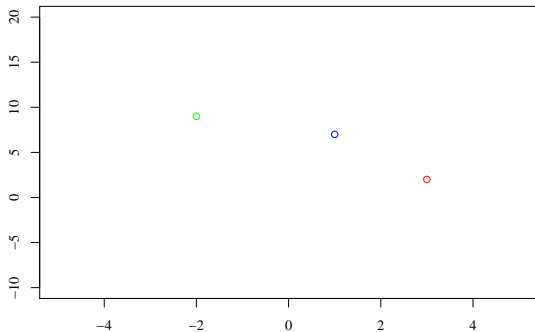
```
bdraw <- function() {  
  plot(x = c(-5, 5), y = c(-10, 20),  
       type = "n", xlab = "", ylab = "")  
}  
5 bdraw()  
points(x = x, y = y)
```



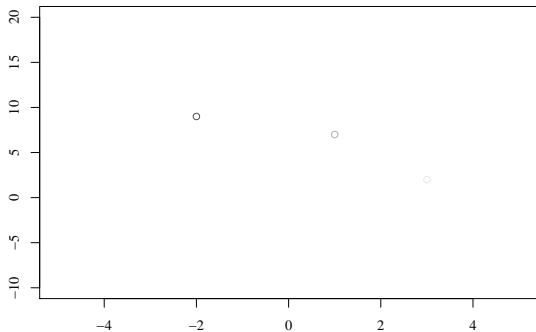
```
bdraw()  
points(x = x, y = y, col = "red")
```



```
bdraw()  
points(x = x, y = y, col = c("green", "blue",  
  "red"))
```



```
bdraw()  
points(x = x, y = y, col = gray.colors(3))
```



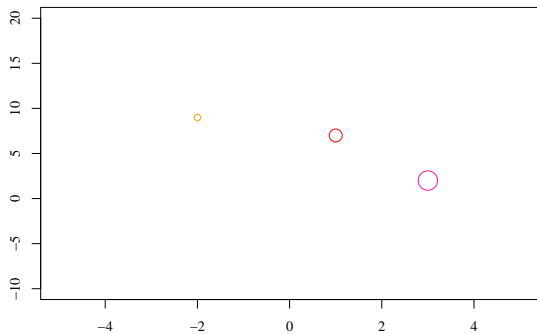
# RTFM

- `?points`. See arguments
  - `pch`: plot symbol
  - `lwd`: thickness of line in drawing
  - `cex`: character expansion: 1 is default
  - `bg`: background color for outline symbols
- Run `example(points)`
- `?points.formula`
  - allows syntax like `points(y ~ x, data = dat)`

# Try some practices

```
plot(x = c(-5, 5), y = c(-10, 20), type = "n",  
     xlab = "", ylab = "")  
x <- c(-2, 1, 3)  
y <- c(9, 7, 2)  
points(x = x, y = y, cex = c(1, 2, 3), col =  
       c("orange", "red", "deeppink"))
```

# Try some practices ...





# The plot function shortcut . . .

- I want you to understand you can draw points on top of any plot.
- But if you only want points, there is a shortcut

```
plot(y ~ x, axes = FALSE, xlab = "", ylab = "")
```

# The plot function shortcut . . . .

○

○

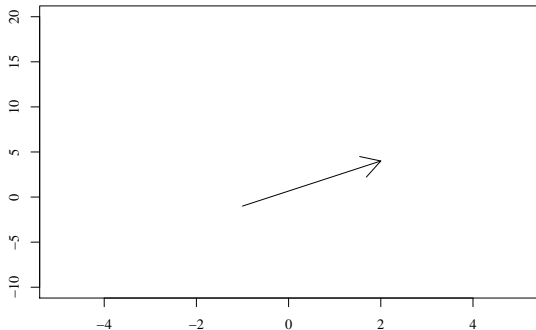
○

# Points worth mentioning

- `points()` are drawn centered on the coordinates in `x` and `y`
- for larger symbols, adjust `cex`
- for darker lines in outlines of symbols, adjust `lwd`

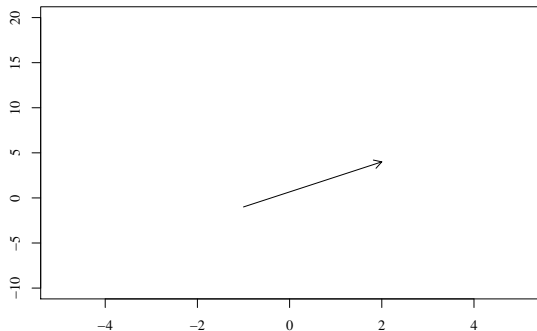
# Arrows. Learn by doing!

```
bdraw()  
arrows(x0 = -1, y0 = -1, x1 = 2, y1 = 4)
```



# Arrows. Smaller fins

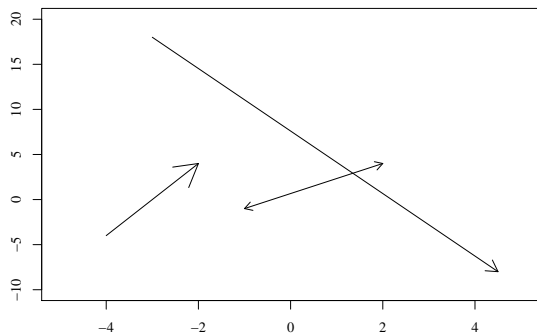
```
bdraw()  
arrows(x0 = -1, y0 = -1, x1 = 2, y1 = 4, length =  
       0.1)
```



# Code 1 2 3

```
bdraw()  
# code 3 is both ways  
arrows(x0 = -1, y0 = -1, x1 = 2, y1 = 4, length =  
       0.1, code = 3)  
# code 2 points forwards  
arrows(x0 = -4, y0 = -4, x1 = -2, y1 = 4, length  
       = 0.3, code = 2)  
# code 1 is backwards  
arrows(x0 = 4.5, y0 = -8, x1 = -3, y1 = 18,  
       length = 0.15, code = 1)
```

## Code 1 2 3 ...



# Use one `arrows()` command

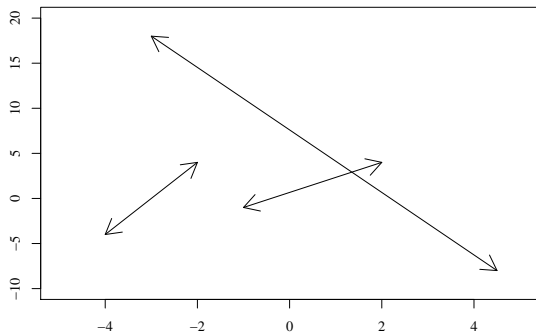
- I tried to show off, but discovered something that looks rather like a weakness in `arrows()`, possibly even a bug.
- My idea was to stack together the input data

```
x0 <- c(-1, -4, 4.5)
y0 <- c(-1, -4, -8)
x1 <- c(2, -2, -3)
y1 <- c(4, 4, 18)
mylengths <- c(0.2, 0.3, 0.15)
mycodes <- c(3, 2, 1)
bdraw()
arrows(x0 = x0, y0 = y0, x1 = x1, y1 = y1, length
       = mylengths, code = mycodes)
```



# Use one `arrows()` command ...

only the first elements in `mylengths` and `mycodes` obeyed.

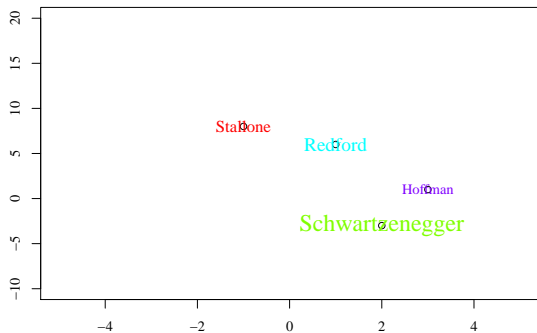


# Text is like points, except ...

- text needs labels, one for each location
- positioning text can be tricky because sometimes we want text above, below, or on side of a point.

```
x <- c(-1, 2, 1, 3); y <- c(8, -3, 6, 1)
labels <- c("Stallone", "Schwarzenegger",
            "Redford", "Hoffman")
mycolors <- rainbow(4)
bdraw()
points(x, y)
text(x = x, y = y, labels = labels, col =
     mycolors, cex = c(1.2, 1.7, 1.3, 1))
```

# Text is like points, except . . . .



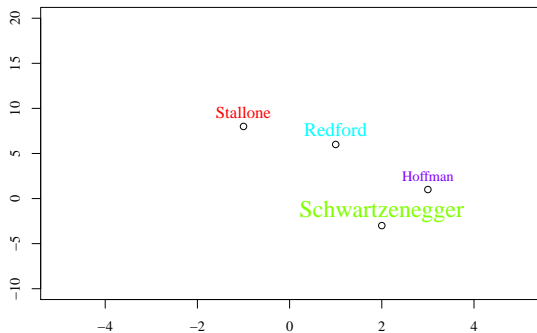
# Text labels overlap points, if you are not careful

Default `text()` has `pos = 1`.

`pos = 3` moves text above the point

```
bdraw()  
points(x, y)  
text(x = x, y = y, labels = labels, col =  
      mycolors, cex = c(1.2, 1.7, 1.3, 1), pos = 3)
```

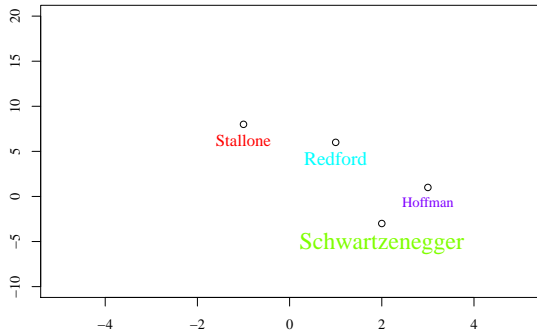
# Text labels overlap points, if you are not careful ...



# offset needed to write "under" the points

```
bdraw()  
points(x, y)  
text(x = x, y = y, labels = labels, col =  
      mycolors, cex = c(1.2, 1.7, 1.3, 1), pos = 1,  
      offset = 0.7)
```

# offset needed to write "under" the points ...



offset units are "character widths"

# offset needed to write "under" the points

- `lines()` : will "connect the dots" and do so with some smoothing for pleasant curve
- `segments()` : straight line connect the dots, no smoothing
- `abline()` : a "shortcut" function to draw some commonly used straight lines
- `curve()` : a "shortcut" function for drawing curves for functions of  $x$ .

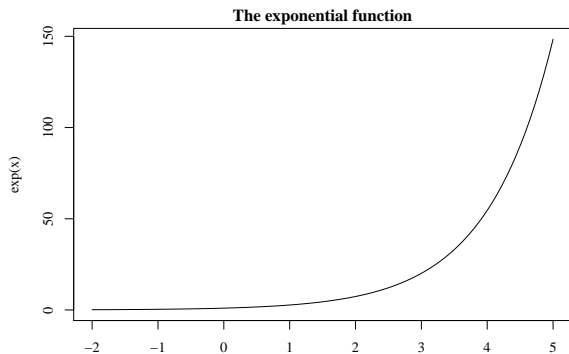


# Plotting Functions

- In statistics, we often find transformations like  $\exp(x)$  or  $\log(x)$
- A good way to learn about them is to plot them with R's curve function
- `curve()` creates its own graphic device, so we don't need to run `plot` first.

```
curve(exp(x), from = -2, to = 5, xlab = "Don't  
set x max too large", main = "The exponential  
function")
```

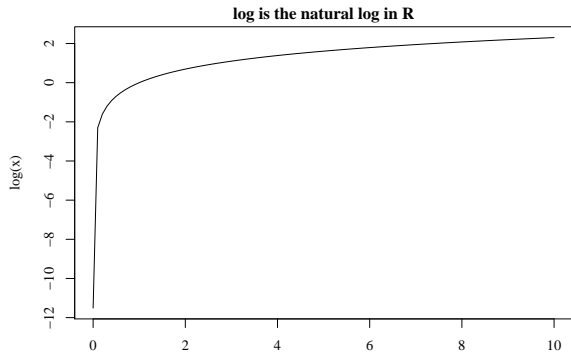
# Plotting Functions ...



Don't set x max too large

# The Natural Logarithm

```
curve(log(x), from = 0.00001, to = 10, xlab =  
      "Note minimum x is 0.00001. Guess why?", main  
      = "log is the natural log in R")
```



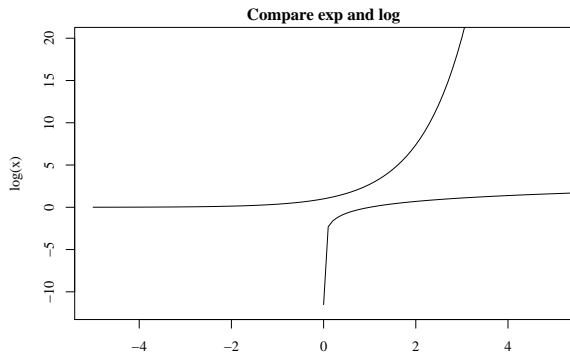
Note minimum x is 0.00001. Guess why?

# Can "Overlay" curves

- The first curve we draw sets the scale.
- `xlim`, `ylim`: arguments so that the scale is big enough to show the interesting parts of both curves.

```
curve(log(x), from = 0.00001, to = 10, xlab =  
      "The domain is now -5, 5", main = "Compare  
      exp and log", xlim = c(-5, 5), ylim = c(-12,  
      20))  
curve(exp(x), from = -5, to = 5, add = TRUE)
```

# Can "Overlay" curves ...

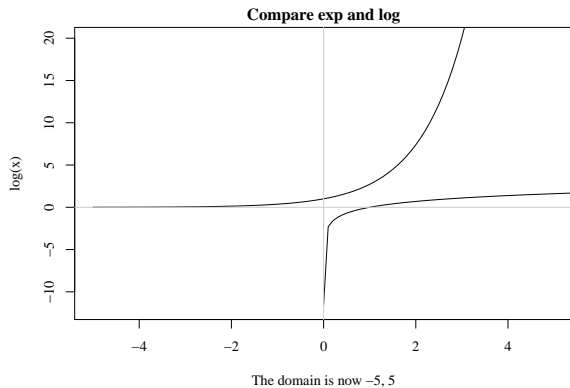


The domain is now  $-5, 5$

# Insert light reference lines with abline

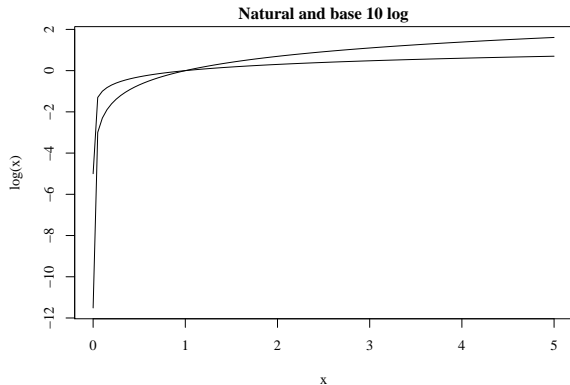
```
curve(log(x), from = 0.00001, to = 10, xlab =  
      "The domain is now -5, 5", main = "Compare  
      exp and log", xlim = c(-5, 5), ylim = c(-12,  
      20))  
curve(exp(x), from = -5, to = 5, add = TRUE)  
abline(v = 0, col = "gray80")  
abline(h = 0, col = "gray80")
```

# Insert light reference lines with abline ...



# What is the Natural Logarithm?

```
curve(log(x), from = 0.00001, to = 5, xlab = "x",  
      main = "Natural and base 10 log")  
curve(log(x, 10), from = 0.00001, to = 5, add =  
      TRUE)
```

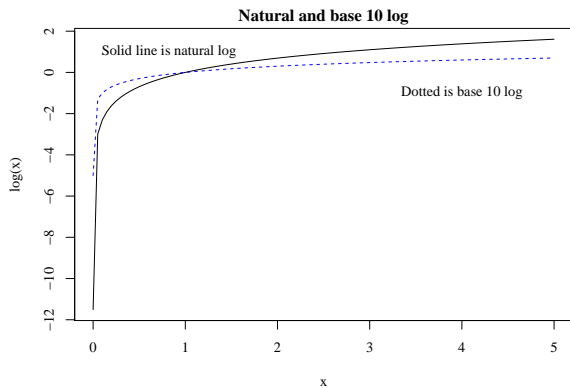




# I cannot tell those apart!

```
curve(log(x), from = 0.00001, to = 5, xlab = "x",  
      main = "Natural and base 10 log")  
curve(log(x, 10), from = 0.00001, to = 5, add =  
      TRUE, lty = 2, col = "blue")  
text(4, -1, "Dotted is base 10 log")  
text(0, 1, "Solid line is natural log", pos = 4)
```

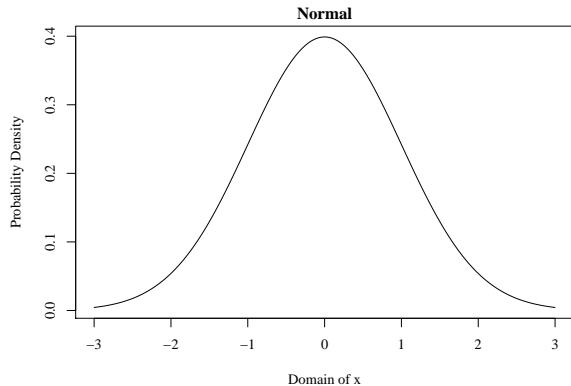
# I cannot tell those apart! ...



# My favorite use of lines(): draw probability density functions

```
x <- seq(-3, 3, length.out = 200)
xprob <- dnorm(x, m = 0, s = 1)
plot(xprob ~ x, type = "n", xlab = "Domain of x",
      ylab = "Probability Density", main = "Normal")
lines(xprob ~ x)
```

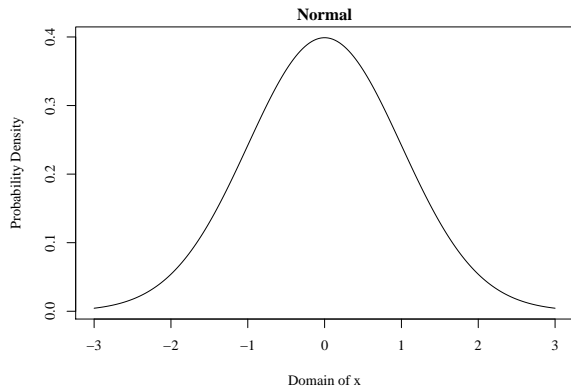
# My favorite use of `lines()`: draw probability density functions ...



`dnorm` is R's function to calculate probability density of the normal

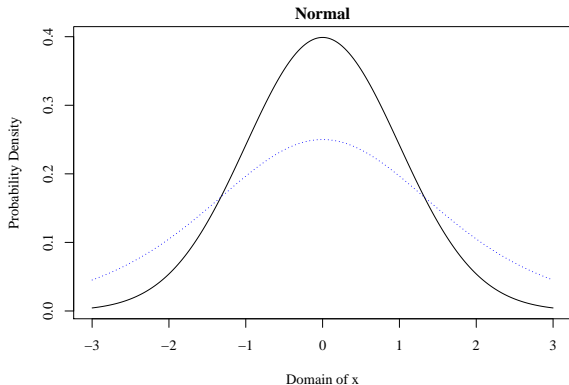
plot type = "l" is a shortcut for that

```
plot(xprob ~ x, type = "l", xlab = "Domain of x",  
     ylab = "Probability Density", main = "Normal")
```



# Compare densities of 2 different distributions

```
plot(xprob ~ x, type = "l", xlab = "Domain of x",  
     ylab = "Probability Density", main = "Normal")  
x2prob <- dlogis(x, location = 0, scale = 1)  
lines(x2prob ~ x, lty = 3, col = "blue")
```

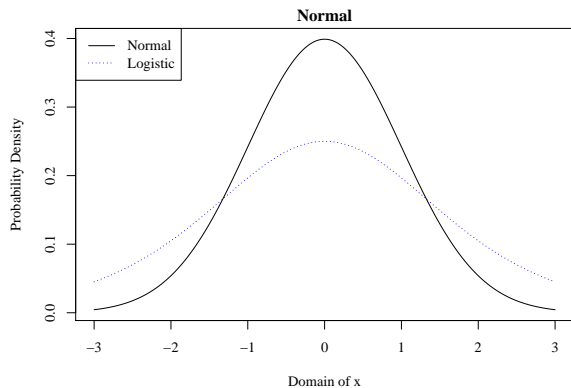


# Insert a legend

I've found that getting a legend "just right" can be very frustrating.

```
plot(xprob ~ x, type = "l", xlab = "Domain of x",  
     ylab = "Probability Density", main = "Normal")  
x2prob <- dlogis(x, location = 0, scale = 1)  
lines(x2prob ~ x, lty = 3, col = "blue")  
legend("topleft", legend = c("Normal",  
                              "Logistic"), lty = c(1, 3), col = c("black",  
                              "blue"))
```

# Insert a legend ...



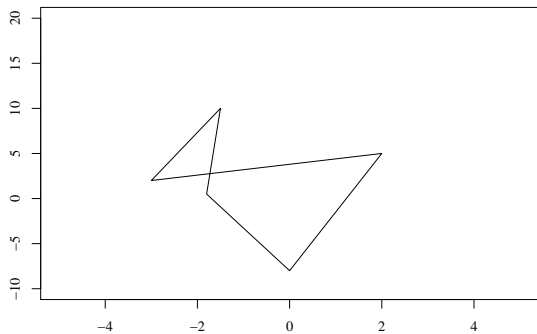


## color in shapes

- If you can supply the points, R can draw a smooth, “connect-the-dots” curve, and decorate the insides.

```
bdraw()  
x <- c(-3, -1.5, -1.8, 0, 2, -3)  
y <- c(2, 10, 0.5, -8, 5, 2)  
polygon(x, y)
```

## color in shapes ...

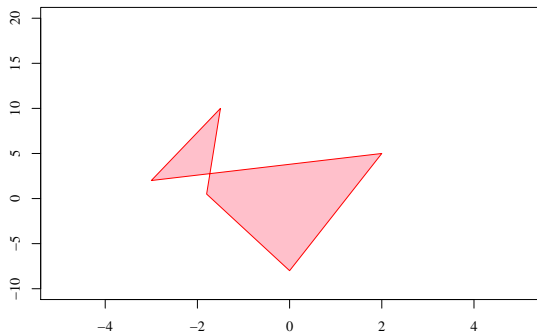


# Whoops! I forgot that Splash of Color!

- If you can supply the points, R can draw a smooth, “connect-the-dots” curve, and decorate the insides.

```
bdraw()  
polygon(x, y, col = "pink", border = "red")
```

# Whoops! I forgot that Splash of Color! ...

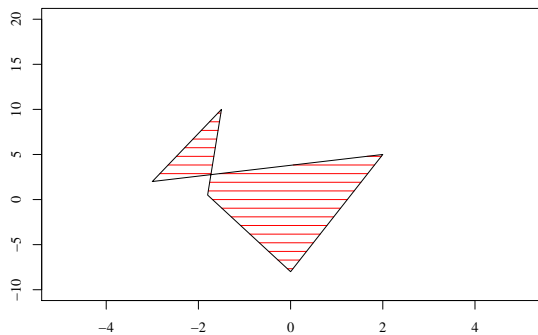


# Play with polygons

- density: Instead of coloring background, can draw lines on it.
- angle: direction of lines inside polygon
- If you can supply the points, R can draw a smooth, “connect-the-dots” curve, and decorate the insides.

```
bdraw()  
polygon(x, y, col = "red", border = "black",  
        density = 10, angle = 0)
```

# Play with polygons ...

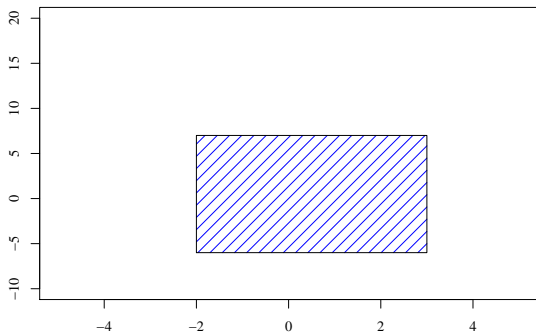


# The `rect()` function is almost identical to `polygon`

- `rect` wants 4 arguments, the corner coordinates

```
bdraw()  
rect(xleft = -2, ybottom = -6, xright = 3, ytop =  
    7, col = "blue", border = "black", density =  
    10, angle = 45)
```

The `rect()` function is almost identical to `polygon` ...





# Outline

- 1 line art
- 2 Examples
- 3 Create a Blank Sheet of Paper
- 4 Inside the Plot Region
  - points
  - arrows
  - text
  - lines, curves
  - polygon
  - rectangles
- 5 plotmath
- 6 Are you looking for skills to practice?

# Sometimes a well placed $\sigma$ or $\psi$ pushes your plot over the top

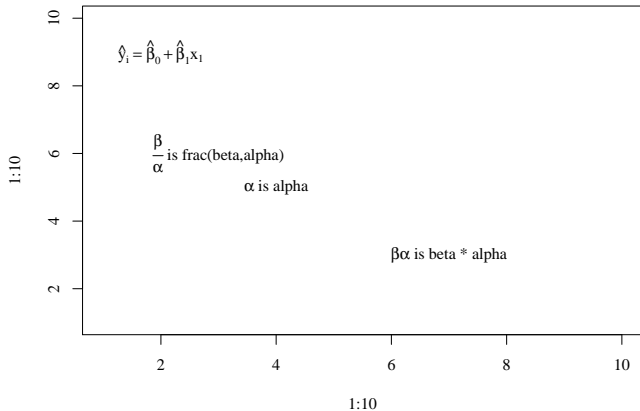
- I don't want to spend a lot of time on this, because it is almost mind-numbingly complicated in some ways, but let's just run an example.

```

plot(1:10, 1:10, type="n")
text(4, 5, expression(paste(alpha, " is alpha")))
text(7, 3, expression(paste(beta * alpha, " is
    beta * alpha")))
text(3, 6, expression(paste(frac(beta, alpha), "
    is frac(beta, alpha)")))
text(2,9, expression(paste(hat(y)[i] ==
    hat(beta)[0]+hat(beta)[1]*x[1])))

```

# I also like $\beta$ , $\alpha$ and $\Sigma$



# A Few plotmath Tips

- Two Equal Signs (`==` gives back `=`)
- Use hard brackets `[]` for subscripts, `^` for superscripts
- Want `*` to show? Type `%*%`
- Want centered `·` for multiplication? Type `cdot`
- Want  $(x - 1, y_1)$ ? Type `group("(" , list(x[1], y[1]), ")")`

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# What To Practice Today?

- Maybe this will get you started

```

plot(1:10, 1:10, type = "n")
abline(h = 2:9, v = c(3, 5, 7), col =
      "gray80")
arrows(x0 = 2, y0 = 3, x1 = 9, y1 = 2, length
      = 0.1)
text(3, 7, "Kansas in Summer is like Paris",
     pos = 4)
5 text(3.2, 6.6, "if Paris were hot and humid",
     pos = 4)

```

- Sketch a technical illustration on paper
  - Figure out how to draw it by starting with a blank device and adding lines, rectangles, etc.
- Step through the code that generates the graphs in section 1 of this presentation.
  - Leave `SAVEME <- FALSE` if you want on-screen graphics.
- If you have R for Windows or Macintosh, lets find the keystrokes to “step next” through one of these examples

# References

R Core Team (2017). *R: A Language and Environment for Statistical Computing*. R Foundation for Statistical Computing, Vienna, Austria.

# Session

```
sessionInfo()
```

```
R version 3.4.4 (2018-03-15)
Platform: x86_64-pc-linux-gnu (64-bit)
Running under: Ubuntu 18.04 LTS

5 Matrix products: default
BLAS: /usr/lib/x86_64-linux-gnu/blas/libblas.so.3.7.1
LAPACK: /usr/lib/x86_64-linux-gnu/lapack/liblapack.so.3.7.1

10 locale:
   [1] LC_CTYPE=en_US.UTF-8      LC_NUMERIC=C
       LC_TIME=en_US.UTF-8
   [4] LC_COLLATE=en_US.UTF-8   LC_MONETARY=en_US.UTF-8
       LC_MESSAGES=en_US.UTF-8
   [7] LC_PAPER=en_US.UTF-8     LC_NAME=C           LC_ADDRESS=C
  [10] LC_TELEPHONE=C          LC_MEASUREMENT=en_US.UTF-8
       LC_IDENTIFICATION=C

15 attached base packages:
   [1] stats      graphics  grDevices  utils      datasets   base

loaded via a namespace (and not attached):
   [1] compiler_3.4.4 tools_3.4.4
```