## From data consumer to tool maker

**Winston Chang** 

**RStudio** 

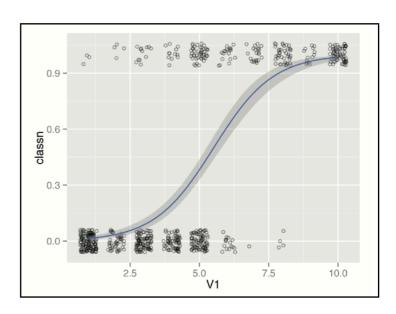
### **Tool maker**

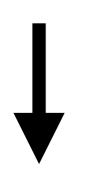
### Data analyst

### Consumer

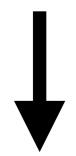


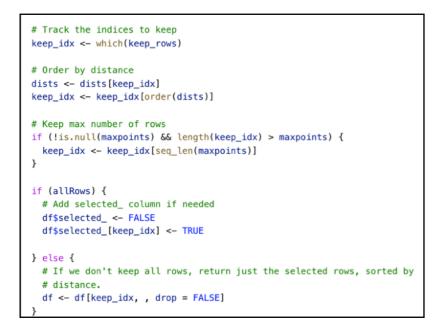
ID	V1	V2	V3	V4	class
1036172	2	1	1	1	benign
1041801	5	3	3	3	malignant
1043999	1	1	1	1	benign
1044572	8	7	5	10	malignant
1047630	7	4	6	4	malignant
1048672	4	1	1	1	benign
1049815	4	1	1	1	benign

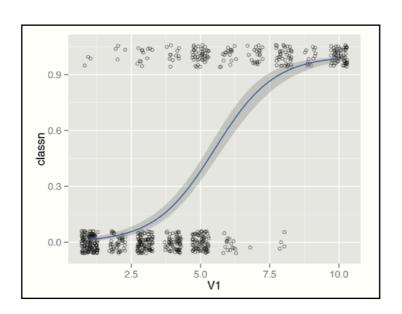












No surgery needed.

Decision/action

### Tool maker

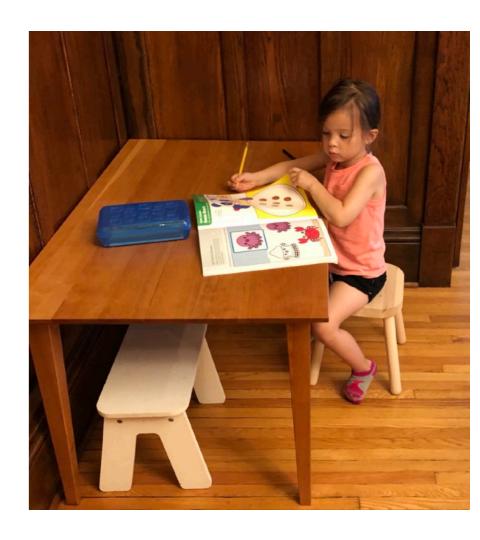
### **Furniture maker**

### Consumer







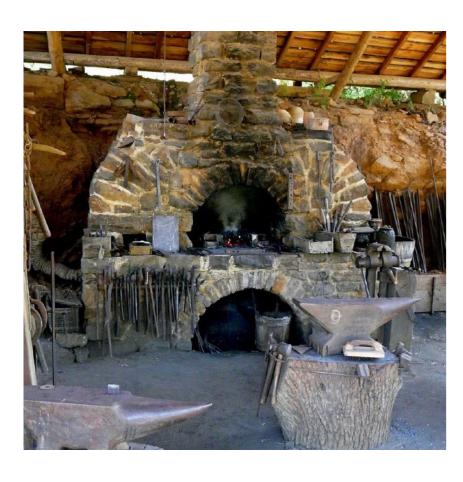


## Things you need

#### **Tool maker**

### **Furniture maker**

#### Consumer









## Things you need

**Tool maker** 



Consumer











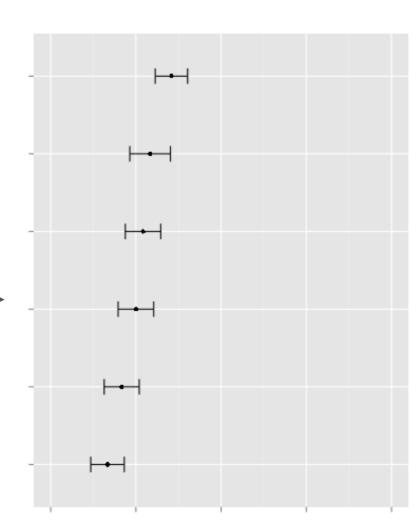


# Using R to analyze data

subject	csqForm	<b>c1</b>	<b>c2</b>	сЗ	с4	с5	с6
490	В	1	1	2	1	2	2
529	В	2	2	1	1	2	1

form	new	original
Α	1	1
Α	2	2
Α	3	3
Α	4	4
Α	5	5
Α	6	6
В	1	5
В	2	6
В	3	4

subject	csqForm	qNum.new	qNum	rating
490	В	5	1	2
490	В	6	2	2
490	В	4	3	1
529	В	5	1	2
529	В	6	2	1
529	В	4	3	1

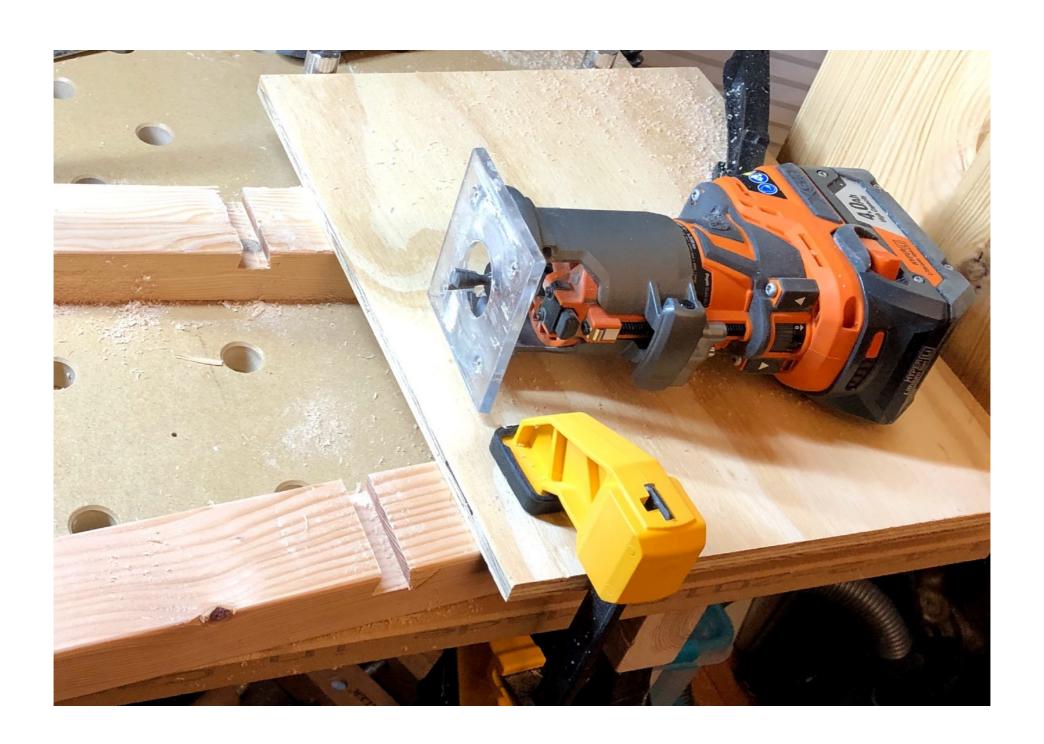


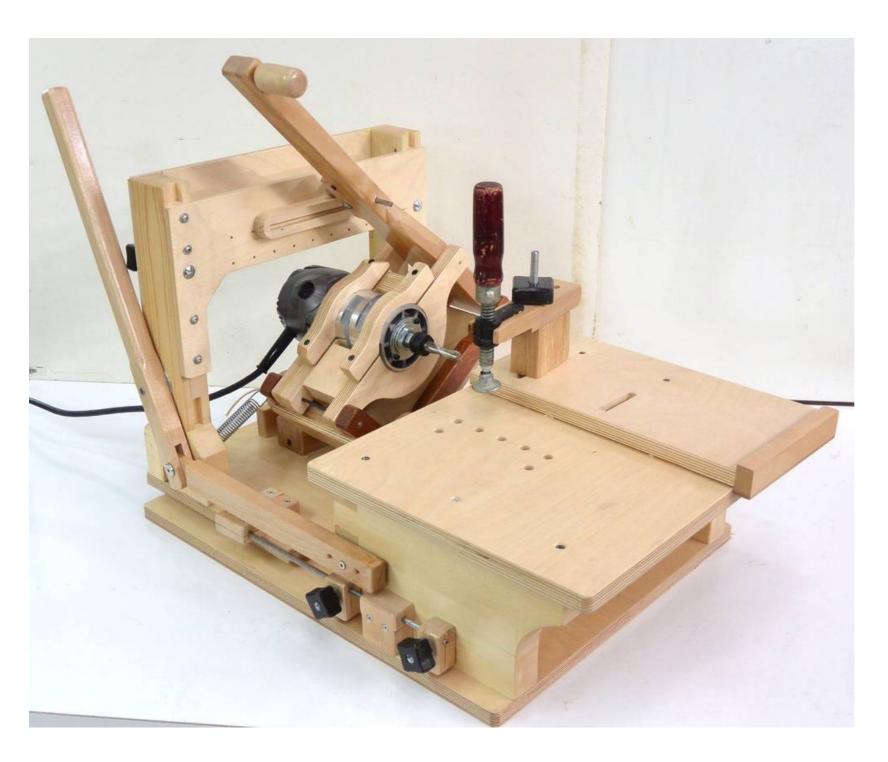
```
library(reshape)
rawdata <- read.csv('data.csv', header=T)</pre>
csq0rder <- read.csv('csq_order.csv', header=T)</pre>
csqRawData <- melt(rawdata,</pre>
    id.vars = c("subject", "csqForm"),
    measure vars = c("c1","c2","c3","c4","c5","c6"),
    variable name = "qNum.new")
names(csqRawData)[names(csqRawData)=="value"] <- "rating"</pre>
levels(csqRawData$qNum.new)[levels(csqRawData$qNum.new)=="c1"] <- "1"</pre>
levels(csqRawData$qNum.new)[levels(csqRawData$qNum.new)=="c2"] <- "2"</pre>
levels(csqRawData$qNum.new)[levels(csqRawData$qNum.new)=="c3"] <- "3"</pre>
levels(csqRawData$qNum.new)[levels(csqRawData$qNum.new)=="c4"] <- "4"</pre>
levels(csqRawData$qNum.new)[levels(csqRawData$qNum.new)=="c5"] <- "5"</pre>
levels(csqRawData$qNum.new)[levels(csqRawData$qNum.new)=="c6"] <- "6"</pre>
# Merge the two data frames to get original numbers
csqData <- merge(csqRawData, csqOrder,</pre>
    by x = c("csqForm","qNum.new"),
    by_y = c("form", "new"),
names(csqData)[names(csqData)=="original"] <- "qNum"</pre>
# Reorder the columns to something a little nicer
csqData <- csqData[, c(3,1,2,6,5,4)]
csqData <- csqData[order(csqData$subject, csqData$qNum), ]</pre>
```



```
library(tidyverse)
rawdata <- read.csv('data.csv', header=T)</pre>
csq0rder <- read.csv('csq order.csv', header=T)</pre>
csqRawData <- rawdata %>%
  pivot_longer(cols = c1:c6, names_to = "qNum.new",
               values to = "rating") %>%
  select(subject, csqForm, qNum.new, rating) %>%
  mutate(qNum.new = recode(qNum.new,
                            c1=1, c2=2, c3=3, c4=4, c5=5, c6=6)
csqData <-
  inner_join(csqRawData, csq0rder,
             by = c("csqForm" = "form", "qNum.new" = "new")) %>%
  select(subject, csqForm, qNum.new, story, qNum = "original",
         rating) %>%
  arrange(subject, qNum)
```

# Making your own tools







https://woodgears.ca/pantorouter/

```
do <- function(.data, ...) {</pre>
 UseMethod("do")
#' @export
do.default <- function(.data, ...) {</pre>
  do_(.data, .dots = compat_as_lazy_dots(...))
#' @export
#' @rdname se-deprecated
do_ <- function(.data, ..., .dots = list()) {</pre>
 UseMethod("do_")
#' @export
do.NULL <- function(.data, ...) {</pre>
 NULL
#' @export
do_.NULL <- function(.data, ..., .dots = list()) {</pre>
 NULL
# Helper functions --
label output dataframe <- function(labels, out, groups) {</pre>
  data_frame <- vapply(out[[1]], is.data.frame, logical(1))</pre>
 if (any(!data_frame)) {
    bad("Results {bad} must be data frames, not {first_bad_class}",
      bad = fmt_comma(which(!data_frame)),
      first_bad_class = fmt_classes(out[[1]][[which.min(data_frame)]])
 }
  rows <- vapply(out[[1]], nrow, numeric(1))</pre>
  out <- bind_rows(out[[1]])</pre>
  if (!is.null(labels)) {
    # Remove any common columns from labels
    labels <- labels[setdiff(names(labels), names(out))]</pre>
    # Repeat each row to match data
    labels <- labels[rep(seq_len(nrow(labels)), rows), , drop = FALSE]
    rownames(labels) <- NULL
```

#### install.packages("dplyr")

```
label_output_list <- function(labels, out, groups) {</pre>
 if (!is.null(labels)) {
    labels[names(out)] <- out</pre>
    rowwise(labels)
 } else {
   class(out) <- "data.frame"</pre>
   attr(out, "row.names") <- .set_row_names(length(out[[1]]))</pre>
   rowwise(out)
named_args <- function(args) {</pre>
 # Arguments must either be all named or all unnamed.
  named <- sum(names2(args) != "")</pre>
  if (!(named == 0 || named == length(args))) {
   abort("Arguments must either be all named or all unnamed")
  if (named == 0 && length(args) > 1) {
   bad("Can only supply one unnamed argument, not {length(args)}")
  # Check for old syntax
  if (named == 1 && names(args) == ".f") {
   abort("do syntax changed in dplyr 0.2. Please see documentation for details")
  named != 0
```

## **Functions**

A collection of functions in a .R file

R package on GitHub

R package on CRAN

# Making tools: some lessons learned

A good API is hard to design.

## Renaming factor levels

```
levels(df$col)[levels(df$col)=='c1'] <- '1'</pre>
levels(df$col)[levels(df$col)=='c2'] <- '2'</pre>
levels(df$col)[levels(df$col)=='c3'] <- '3'</pre>
library(plyr)
revalue(df$col, c(c1='1', c2='2', c3='3'))
library(dplyr)
recode(df$col, c1='1', c2='2', c3='3'))
library(forcats)
fct_recode(df$col, '1'='c1', '2'='c2', '3'='c3'))
```

## Shiny: renderImage

## Shiny: server.R and ui.R

```
# server.R
shinyServer(function(input, output) {
  # ...
})
# ui.R
shinyUI(fluidPage(
  # ...
})
# app.R
shinyApp(
  ui = fluidPage(
    #...
  server = function(input, output) {
    # ...
  }
```

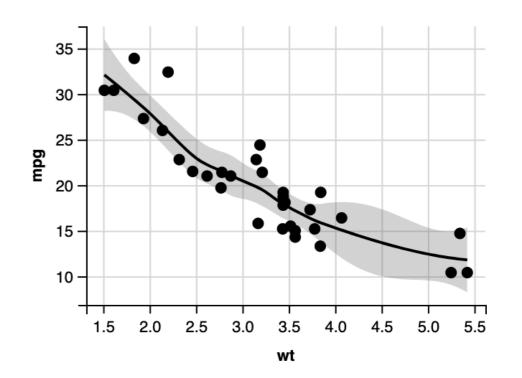
## Why are good APIs hard to design?

- Don't have enough experience to decide on a consistent pattern.
- Don't fully understand how people will use a function.
- Too familiar with old ways of looking at things.
- Not familiar enough with old ways of looking at things.

If it takes more time to explain how something works than it does to simplify it, then simplify it.

Sometimes a	oroject can be	e too ambitious.

- ggvis: interactive grammar of graphics
  - different syntax from ggplot2
  - rendered in the browser, using Vega (JavaScript) library





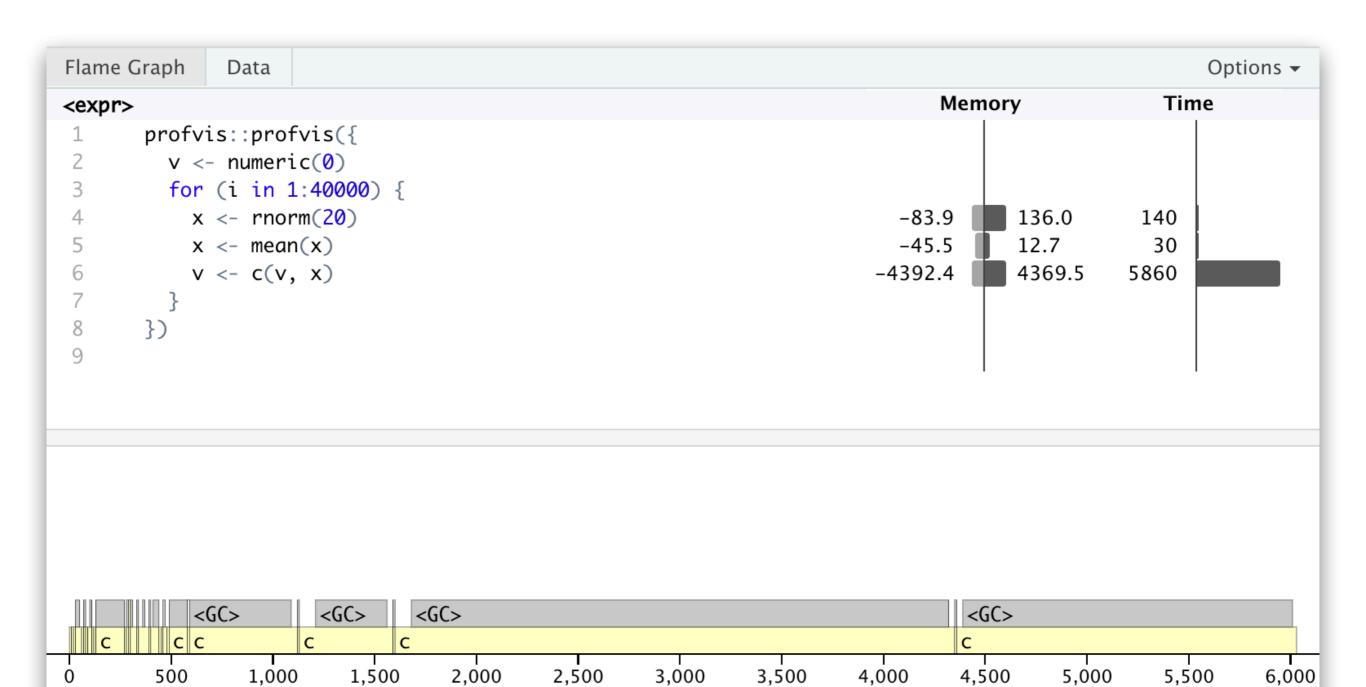
- Why didn't it work?
  - Vega made some things easy, other things impossible
  - ggvis wasn't better than ggplot2 for most users

If your code seems slower than it should be, don't guess about why; use a profiler.



```
v <- numeric(0)
for (i in 1:40000) {
    x <- rnorm(20)
    x <- mean(x)
    v <- c(v, x)
}</pre>
```

```
profvis::profvis({
   v <- numeric(0)
   for (i in 1:40000) {
      x <- rnorm(20)
      x <- mean(x)
      v <- c(v, x)
   }
})</pre>
```



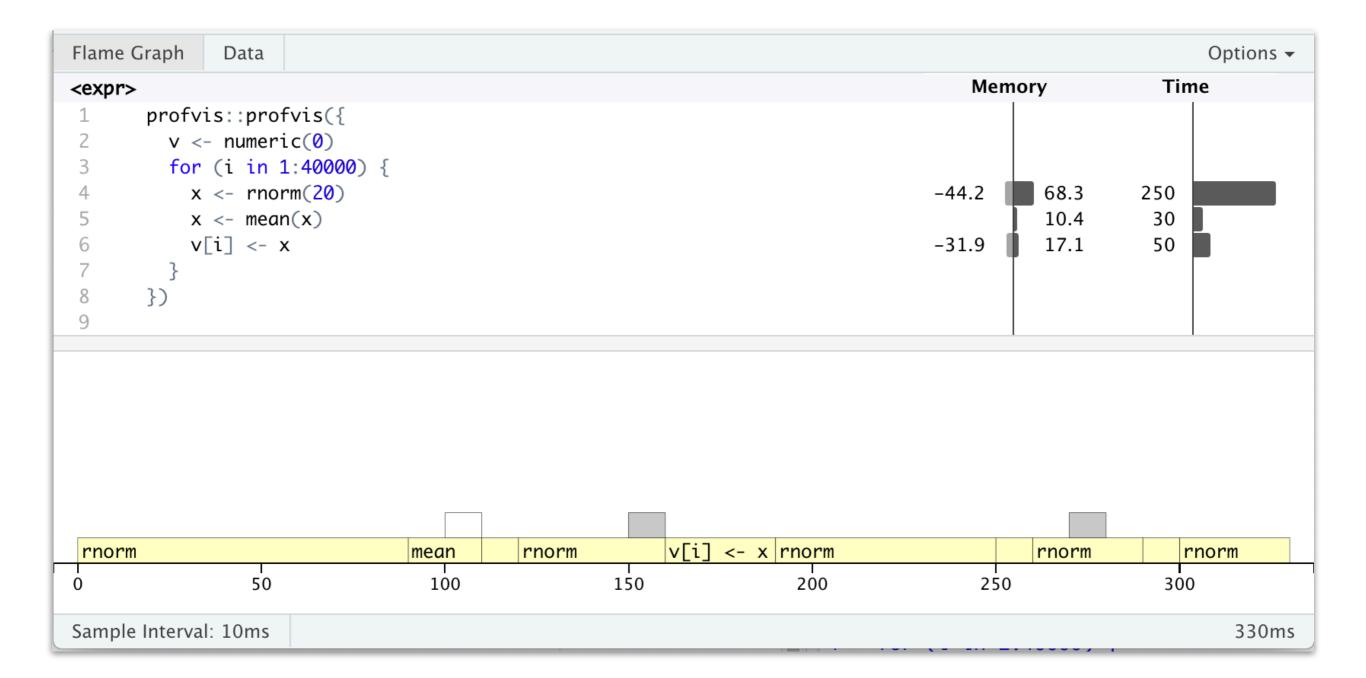
4,500

6030ms

1,500

Sample Interval: 10ms

```
v <- numeric(0)
for (i in 1:40000) {
    x <- rnorm(20)
    x <- mean(x)
    v[i] <- x
}</pre>
```



Think about s	cale when dec	iding what to w	ork on.

## Sometimes small improvements require deep digging.

## It's not a complete waste of time if you learn something from it.

## Thank you!

