%>% magrittr

Ceci n'est pas un pipe.

Figure 1:



http://healthyr.surgicalinformatics.org

Coding style

Our usual script preamble:

At the beginning of each script, we usually have these lines:

```
rm(list=ls())  # clears the Environment tab

library(tidyverse)  # packages: ggplot2, tibble, tidyr, readr, purrr, and dplyr
library(forcats)  # all the fct_recode() etc. functions, forcats is an anagram for factors
library(broom)  # functions: tidy(), glance(), augment()
library(magrittr)  # includes the %<>% assignment-pipe ( %>% is loaded from dplyr)
```

To see more information about any package or function, move your cursor to it and press F1.

The pipe operator - %>%:

(Image source: https://cran.r-project.org/web/packages/magrittr/vignettes/magrittr.html)

The pipe operator (%>%) sends an R object (usually a data frame) into a function. You can type it up, or you can use the keyboard short-cut of Control+Shift+M. It looks a bit scary at first, but once you get used to it you won't want to do R any other way.

Let's load some example data (the diamonds dataset that comes with ggplot2):

```
mydata = diamonds

# or equivalently
mydata <- diamonds

# or equivalently
diamonds -> mydata
```

(We tend to use the = rather than <-, but sometimes we do use the left-to-right arrow: ->.)

You can summarise a data frame like this:

```
summary(mydata)
```

Or write the same thing using a pipe:

```
mydata %>% summary()
```

This might not seem that useful at first as mydata %>% summary() is actually longer than summary(mydata). Nevertheless, piping becomes very useful when doing a few manipulations after another or filtering data before summarising it.

For example, look at this line:

```
summary(select(filter(mydata, cut %in% c('Premium', 'Ideal')), cut, price))
```

```
##
                           price
           cut
##
   Fair
             :
                  0
                      Min.
                              :
                                 326
##
   Good
                  0
                       1st Qu.: 929
   Very Good:
                  0
                      Median: 2178
##
                              : 3897
##
  Premium :13791
                      Mean
##
   Ideal
             :21551
                       3rd Qu.: 5364
##
                      Max.
                              :18823
```

```
(cut %in% c('Premium', 'Ideal') is equivalent to cut == 'Premium' | cut == 'Ideal', | means or)
```

It is quite hard to figure out what is going on there. Writing the exact same workflow out with pipes makes it (human) readable:

```
mydata %>%
  filter(cut %in% c('Premium', 'Ideal')) %>%
  select(cut, price) %>%
  summary()
```

```
##
           cut
                           price
##
    Fair
                   0
                                  326
                       Min.
##
    Good
             :
                   0
                       1st Qu.:
                                  929
                       Median: 2178
##
   Very Good:
                   0
                              : 3897
##
   Premium :13791
                       Mean
                       3rd Qu.: 5364
##
   Ideal
             :21551
##
                       Max.
                               :18823
```

Reading this: Take mydata and filter for cuts that are either "Premium" or "Ideal", select cut and price, and summarise.

->:

Once you really get into piped workflows it starts making sense to use the left-to-right assignment arrow ->:

```
mydata %>%
  filter(cut %in% c('Premium', 'Ideal')) %>%
  select(cut, price) ->
  mydata_highquality
```

is equivalent to:

```
mydata_highquality = mydata %>%
  filter(cut %in% c('Premium', 'Ideal')) %>%
  select(cut, price)
```

%<>%:

And finally, there is %<>% to send data into a function, and then save the result back into the original variable: it is a combination of %>% and <-. We don't normally use it when filtering or selecting variables from the original data frame (as you will overwrite and lose everything else), but it is very useful when doing a minor manipulation on a single column.

For example, even after filtering out the Fair, Good, and Very Good diamond cuts, they still show up with 0 counts in summary() above. This is because R knows that cut is a factor rather than just a character column:

```
mydata$cut %>% class()
```

```
## [1] "ordered" "factor"
```

If you want to get rid of the sub-par categories completely, you need to use fct_drop from library(forcats) (note the use of %<>%):

```
mydata_highquality$cut %<>% fct_drop()

mydata_highquality %>%
  summary()
```

```
##
                         price
         cut
                             : 326
##
    Premium: 13791
                     Min.
##
    Ideal :21551
                     1st Qu.: 929
##
                     Median: 2178
##
                     Mean
                            : 3897
##
                     3rd Qu.: 5364
##
                     Max.
                             :18823
```

"" or '':

In most cases, "Ideal" and 'Ideal' are equivalent. '' is one less key to press (as it doesn't require *Shift*), but "" is more widely used by the R community.