Data manipulation with pipes

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R Markdown

This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. For more details on using R Markdown see http://rmarkdown.rstudio.com.

When you click the **Knit** button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

summary(cars)

```
##
        speed
                         dist
##
           : 4.0
                    Min.
                            : 2.00
    1st Qu.:12.0
                    1st Qu.: 26.00
##
##
    Median:15.0
                    Median: 36.00
##
    Mean
            :15.4
                    Mean
                            : 42.98
    3rd Qu.:19.0
                    3rd Qu.: 56.00
    Max.
            :25.0
                    Max.
                            :120.00
```

Including Plots

You can also embed plots, for example:



Note that the \mbox{echo} = FALSE parameter was added to the code chunk to prevent printing of the R code that generated the plot.

Sequential Code

```
numbers <- -300:546
numbers <- 1:300
numbers_mean <- mean(numbers)
sqrt(x = numbers_mean)</pre>
```

[1] 12.26784

Nested Code

```
numbers <- 1:300
mean(numbers)

## [1] 150.5

sqrt(mean(numbers))</pre>
```

[1] 12.26784

Piping code

```
implement with 'magrittr'
library(magrittr)
pipe symbol is %>% or |
1:300 %>% mean() %>% sqrt()
## [1] 12.26784
surveys <- read.csv(file = "../data-raw/surveys.csv")</pre>
str(surveys)
## 'data.frame': 35549 obs. of 9 variables:
## $ record_id : int 1 2 3 4 5 6 7 8 9 10 ...
                    : int 7777777777...
## $ month
## $ day
                   : int 16 16 16 16 16 16 16 16 16 16 ...
: int 1977 1977 1977 1977 1978 1978 ## $ plot_id : int 2 3 2 7 3 1 2 1 1 6 ... ## $ species_id : chr "NI." "NI" "DN" "
                   : chr "M" "M" "F" "M" ...
## $ sex
## $ hindfoot_length: int 32 33 37 36 35 14 NA 37 34 20 ...
## $ weight
                    : int NA ...
surveys$year %>% mean()
## [1] 1990.475
surveys$weight %>% mean(na.rm = TRUE)
## [1] 42.67243
surveys1 <- select(surveys, year, month, day, species_id)</pre>
surveys2 <- select(surveys, year, species_id, weight) %>% filter(!is.na(weight)) %>%
mutate(weight_kg = weight/1000) %>% select(year, species_id, weight_kg) %>%
  filter(species_id == 'SH')
str(surveys2)
## 'data.frame':
                   141 obs. of 3 variables:
           : int 1978 1982 1982 1986 1987 1987 1987 1987 1988 ...
## $ year
## $ species_id: chr "SH" "SH" "SH" "SH" ...
## $ weight_kg : num 0.089 0.106 0.052 0.055 0.077 0.078 0.104 0.058 0.052 0.06 ...
surveys2_filtered <- filter(surveys2, species_id == 'SH')</pre>
str(surveys2_filtered)
## 'data.frame':
                   141 obs. of 3 variables:
## $ year
            : int 1978 1982 1982 1986 1987 1987 1987 1987 1988 ...
## $ species_id: chr "SH" "SH" "SH" "SH" ...
```

\$ weight_kg : num 0.089 0.106 0.052 0.055 0.077 0.078 0.104 0.058 0.052 0.06 ...