

Assignment

read and View data in R

```
library(readr)
fish_data <- read_csv("fish_data.csv")

## Rows: 2000 Columns: 8
## -- Column specification -----
## Delimiter: ","
## chr (2): habitat, color
## dbl (5): id, average_length, average_weight, ph_of_water, life_span
## lgl (1): Gender
##
## i Use 'spec()' to retrieve the full column specification for this data.
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
```

```
View(fish_data)
print(fish_data)
```

```
## # A tibble: 2,000 x 8
##       id average_length average_weight habitat ph_of_water color      Gender
##   <dbl>         <dbl>         <dbl> <chr>         <dbl> <chr>      <lgl>
## 1     1          14.7           5.87 ponds         6.2 Reddish_Ora~ FALSE
## 2     2           1.32           3.86 idlewater     6.8 Calico      TRUE
## 3     3          14.2          12.1 lakes         7.9 Reddish_Ora~ TRUE
## 4     4           2.54           3.2  rivers         6.7 White       FALSE
## 5     5          13.1           9.81 lakes         7.8 Orange      TRUE
## 6     6          15.2           8.99 lakes         7.8 White       FALSE
## 7     7          16.2           5.08 ponds         6.3 Red_and_Sil~ FALSE
## 8     8          13.7          13.0 rivers         6.7 White       FALSE
## 9     9          13.2           5.22 lakes         7.6 Black_and_0~ FALSE
## 10    10          19.0          15.5 rivers         6.7 Calico      TRUE
## # i 1,990 more rows
## # i 1 more variable: life_span <dbl>
```

The first, second and third, fifth, eighth columns are numerical, fourth and sixth columns are character and seventh column is logical

The dimension of the dataset is 2000x8. This means the dataset has 2000 rows and 8 columns

```
fish_data[ , 'ph_of_water' ]
```

Selecting a column using square brackets

```
## # A tibble: 2,000 x 1
##   ph_of_water
##   <dbl>
## 1         6.2
## 2         6.8
## 3         7.9
## 4         6.7
## 5         7.8
## 6         7.8
## 7         6.3
## 8         6.7
## 9         7.6
## 10        6.7
## # i 1,990 more rows
```

```
fish_data[ fish_data$ph_of_water > 7 , 'ph_of_water']
```

Selecting a column using logical statements

```
## # A tibble: 969 x 1
##   ph_of_water
##   <dbl>
## 1         7.9
## 2         7.8
## 3         7.8
## 4         7.6
## 5         7.2
## 6         7.6
## 7         7.9
## 8         7.3
## 9         7.1
## 10        7.8
## # i 959 more rows
```

```
summary(fish_data)
```

summary of data

```
##      id      average_length average_weight      habitat
## Min.   : 1.0    Min.   : 1.000    Min.   : 2.000 Length:2000
## 1st Qu.: 500.8  1st Qu.: 5.857    1st Qu.: 6.138 Class :character
```

```
## Median :1000.5   Median :10.660   Median :10.455   Mode  :character
## Mean   :1000.5   Mean   :10.557   Mean   :10.449
## 3rd Qu.:1500.2   3rd Qu.:15.172   3rd Qu.:14.665
## Max.   :2000.0   Max.   :20.000   Max.   :18.960
## ph_of_water      color      Gender      life_span
## Min.   :6.000   Length:2000   Mode :logical   Min.   : 1.00
## 1st Qu.:6.500   Class :character FALSE:1007   1st Qu.: 7.80
## Median :7.000   Mode  :character TRUE :969   Median :14.40
## Mean   :7.015           NA's :24   Mean   :14.37
## 3rd Qu.:7.500           3rd Qu.:20.90
## Max.   :8.000           Max.   :28.00
```

Summary of this data gives a simple statistics of each column. The statistics includes Max, Median, Mean, Min, 1st Quartile and 3rd quartile of 1st, 2nd, 3rd, 5th, 8th columns; length, class, mode of 4th and 6th columns; mode, false true and NA's of 7th column.

```
fish_data$double_ph_of_water = fish_data$ph_of_water * 2
head(fish_data)
```

Calculation of data with adding a column

```
## # A tibble: 6 x 9
##       id average_length average_weight habitat ph_of_water color Gender life_span
##   <dbl>      <dbl>         <dbl> <chr>      <dbl> <chr> <lgl>      <dbl>
## 1     1        14.7           5.87 ponds        6.2 Redd~ FALSE      10.9
## 2     2         1.32           3.86 idlewa~    6.8 Cali~ TRUE       5.2
## 3     3        14.2          12.1 lakes        7.9 Redd~ TRUE      25.3
## 4     4         2.54           3.2  rivers        6.7 White FALSE     16.4
## 5     5        13.1           9.81 lakes        7.8 Oran~ TRUE       3.2
## 6     6        15.2           8.99 lakes        7.8 White FALSE     21.6
## # i 1 more variable: double_ph_of_water <dbl>
```

```
aggregate(ph_of_water ~ habitat, data = fish_data, FUN = mean)
```

Use base R to aggregate data

```
##       habitat ph_of_water
## 1    idewater  6.983117
## 2      lakes  7.014115
## 3     ponds  7.039163
## 4     rivers  7.032405
## 5 slowmovingwaters 7.004545
```

```
aggregate( average_length ~ habitat, data = fish_data, FUN = mean)
```

```
##          habitat average_length
## 1      idlewater      10.40330
## 2         lakes      10.64957
## 3         ponds      10.44638
## 4         rivers      11.21332
## 5 slowmovingwaters      10.06803
```

```
aggregate( average_length ~ habitat, data = fish_data, FUN = sum)
```

```
##          habitat average_length
## 1      idlewater      4005.27
## 2         lakes      4451.52
## 3         ponds      4241.23
## 4         rivers      4429.26
## 5 slowmovingwaters      3986.94
```

```
aggregate( average_length ~ habitat, data = fish_data, FUN = max)
```

```
##          habitat average_length
## 1      idlewater      19.96
## 2         lakes      20.00
## 3         ponds      19.97
## 4         rivers      20.00
## 5 slowmovingwaters      19.95
```

```
aggregate( average_length ~ habitat, data = fish_data, FUN = min)
```

```
##          habitat average_length
## 1      idlewater       1.00
## 2         lakes       1.00
## 3         ponds       1.07
## 4         rivers       1.03
## 5 slowmovingwaters       1.01
```