

IRC_Data_Analysis_Training_2

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1/11/2021

Load packages

```
# Load packages
library(tidyverse)

## Warning: package 'tidyverse' was built under R version 4.0.2
## -- Attaching packages ----- tidyverse 1.3.0 --
## v ggplot2 3.3.2      v purrr   0.3.4
## v tibble  3.0.3      v dplyr  1.0.1
## v tidyr   1.1.1      v stringr 1.4.0
## v readr   1.3.1      v forcats 0.5.0
## Warning: package 'ggplot2' was built under R version 4.0.2
## Warning: package 'tibble' was built under R version 4.0.2
## Warning: package 'tidyr' was built under R version 4.0.2
## Warning: package 'dplyr' was built under R version 4.0.2
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()    masks stats::lag()
library(here)

## here() starts at /Users/rachelkenny/Documents/IRC/R Code/IRC_Data_Analysis_Training
library(janitor)

##
## Attaching package: 'janitor'
## The following objects are masked from 'package:stats':
##
##   chisq.test, fisher.test
library(readxl)
```

Load data

```
# Load oak data
oak_data_raw <- read_csv(here("data", "Weir_Oak_Restoration_Data_winter19.csv"))
```

```
## Parsed with column specification:
## cols(
##   `Short ID` = col_character(),
##   Survival = col_logical(),
##   Quantity = col_double(),
##   `Height (cm)` = col_double(),
##   `Open Closed` = col_character(),
##   `Location UML` = col_character(),
##   `Water Yes No` = col_character(),
##   `Sampling Group` = col_character()
## )

oak_data <- clean_names(oak_data_raw)

# Load agua chinon veg data
ac_data_raw <- read_excel(here("data", "OCWR_AC_2019_Data.xlsx"))
ac_data <- clean_names(ac_data_raw)
```

Subsetting data

```
# To refer to specific column in a dataframe, use the [$] symbol
oak_data$height_cm
```

```
##      [1]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##     [15]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##     [29]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##     [43]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##     [57]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##     [71]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##     [85]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##     [99]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [113]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [127]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [141]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [155]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [169]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [183]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [197]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [211]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [225]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [239]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [253]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [267]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [281]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [295]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [309]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [323]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [337]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [351]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [365]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [379]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [393]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
##    [407]  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
```

##	[421]	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
##	[435]	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
##	[449]	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
##	[463]	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
##	[477]	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
##	[491]	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
##	[505]	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
##	[519]	0.0	0.0	0.0	1.0	1.5	1.7	3.0	3.1	4.0	4.0	4.5	5.0	5.0	5.0
##	[533]	5.0	5.0	5.0	5.5	6.0	6.0	6.0	6.0	6.0	6.0	6.0	6.0	6.0	6.0
##	[547]	6.0	6.0	6.0	6.0	6.0	6.5	6.5	6.5	6.5	6.5	7.0	7.0	7.0	7.0
##	[561]	7.0	7.0	7.0	7.0	7.0	7.5	7.5	7.5	8.0	8.0	8.0	8.0	8.0	8.0
##	[575]	8.0	8.0	8.0	8.0	8.0	8.0	8.0	8.0	8.0	8.0	8.1	8.1	8.5	8.7
##	[589]	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0
##	[603]	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.5	9.5	9.5	9.5	9.5
##	[617]	9.8	9.8	9.9	9.9	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0
##	[631]	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0
##	[645]	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0	10.0
##	[659]	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.6	10.7	10.9	11.0	11.0	11.0	11.0
##	[673]	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0
##	[687]	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0	11.0
##	[701]	11.0	11.0	11.0	11.0	11.0	11.0	11.2	11.4	11.5	11.5	11.5	11.5	11.5	11.5
##	[715]	11.6	11.7	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0
##	[729]	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0
##	[743]	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.0	12.3	12.3	12.3
##	[757]	12.5	12.5	12.5	12.5	12.7	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0
##	[771]	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0
##	[785]	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0	13.0
##	[799]	13.0	13.2	13.2	13.3	13.5	13.5	13.5	13.9	14.0	14.0	14.0	14.0	14.0	14.0
##	[813]	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0
##	[827]	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0	14.0
##	[841]	14.0	14.0	14.0	14.0	14.1	14.1	14.2	14.3	14.5	14.5	14.5	14.5	14.5	14.5
##	[855]	14.5	14.5	14.5	14.9	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0
##	[869]	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0
##	[883]	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0
##	[897]	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.0	15.3
##	[911]	15.5	15.5	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0
##	[925]	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0
##	[939]	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.0	16.1	16.2	16.3	16.3	16.3
##	[953]	16.5	16.5	16.5	16.5	16.5	16.5	16.7	16.8	17.0	17.0	17.0	17.0	17.0	17.0
##	[967]	17.0	17.0	17.0	17.0	17.0	17.0	17.0	17.0	17.0	17.0	17.0	17.0	17.0	17.0
##	[981]	17.0	17.0	17.0	17.0	17.0	17.0	17.0	17.0	17.0	17.0	17.0	17.0	17.1	17.2
##	[995]	17.5	17.5	17.5	17.5	17.5	17.5	17.5	17.5	17.7	17.8	18.0	18.0	18.0	18.0
##	[1009]	18.0	18.0	18.0	18.0	18.0	18.0	18.0	18.0	18.0	18.0	18.0	18.0	18.0	18.0
##	[1023]	18.0	18.0	18.0	18.0	18.0	18.0	18.0	18.0	18.0	18.0	18.0	18.4	18.5	18.5
##	[1037]	18.5	18.5	18.5	18.5	18.6	18.9	19.0	19.0	19.0	19.0	19.0	19.0	19.0	19.0
##	[1051]	19.0	19.0	19.0	19.0	19.0	19.0	19.0	19.0	19.0	19.0	19.0	19.3	19.4	19.5
##	[1065]	19.5	19.5	19.5	19.6	20.0	20.0	20.0	20.0	20.0	20.0	20.0	20.0	20.0	20.0
##	[1079]	20.0	20.0	20.1	20.3	20.5	20.5	20.5	20.5	20.9	21.0	21.0	21.0	21.0	21.0
##	[1093]	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.5
##	[1107]	21.5	21.5	21.5	21.5	21.6	22.0	22.0	22.0	22.0	22.0	22.0	22.0	22.0	22.0
##	[1121]	22.0	22.0	22.0	22.0	22.0	22.0	22.0	22.0	22.0	22.0	22.1	22.3	22.3	22.4
##	[1135]	22.5	22.5	22.5	23.0	23.0	23.0	23.0	23.0	23.0	23.0	23.0	23.0	23.2	23.5
##	[1149]	23.8	24.0	24.0	24.0	24.0	24.0	24.0	24.0	24.0	24.0	24.0	24.0	24.2	24.5
##	[1163]	24.5	24.5	24.5	24.5	24.9	25.0	25.0	25.0	25.0	25.0	25.5	25.5	25.5	25.6

```
## [1177] 26.0 26.0 26.0 26.0 26.0 26.5 27.0 27.0 27.0 27.0 27.0 27.0 28.0 28.0
## [1191] 29.0 29.0 30.5 31.6 31.7 35.0 37.0 40.0 40.0 46.0 46.5 63.0
```

```
View(oak_data$height_cm)
```

```
View(oak_data)
```

You can also refer to the number of the column using square brackets []. Using a colon indicates subs
oak_data[4] # column 4

```
## # A tibble: 1,202 x 1
##   height_cm
##   <dbl>
## 1         0
## 2         0
## 3         0
## 4         0
## 5         0
## 6         0
## 7         0
## 8         0
## 9         0
## 10        0
## # ... with 1,192 more rows
```

```
oak_data[4:6] # column 4-6
```

```
## # A tibble: 1,202 x 3
##   height_cm open_closed location_uml
##   <dbl> <chr>         <chr>
## 1         0 Closed      Lower
## 2         0 Closed      Lower
## 3         0 Closed      Lower
## 4         0 Closed      Lower
## 5         0 Closed      Lower
## 6         0 Closed      Lower
## 7         0 Closed      Lower
## 8         0 Closed      Lower
## 9         0 Closed      Lower
## 10        0 Closed      Lower
## # ... with 1,192 more rows
```

```
oak_data[-4] # everything except for column 4
```

```
## # A tibble: 1,202 x 7
##   short_id survival quantity open_closed location_uml water_yes_no
##   <chr>    <lgl>         <dbl> <chr>         <chr>         <chr>
## 1 L_C_01_4 FALSE           0 Closed      Lower         No
## 2 L_C_02_1 FALSE           0 Closed      Lower         Yes
## 3 L_C_02_2 FALSE           0 Closed      Lower         Yes
## 4 L_C_02_3 FALSE           0 Closed      Lower         Yes
## 5 L_C_02_4 FALSE           0 Closed      Lower         No
## 6 L_C_03_3 FALSE           0 Closed      Lower         No
## 7 L_C_03_4 FALSE           0 Closed      Lower         No
## 8 L_C_04_1 FALSE           0 Closed      Lower         Yes
```

```
## 9 L_C_04_3 FALSE          0 Closed      Lower      No
## 10 L_C_05_1 FALSE         0 Closed      Lower      No
## # ... with 1,192 more rows, and 1 more variable: sampling_group <chr>

# A comma before the number indicates a column (though it is not necessary), while a comma after the number indicates a row

# FOR EXAMPLE:

# To return column 3
ac_data[3]

## # A tibble: 1,732 x 1
##   transect
##   <dbl>
## 1       1
## 2       1
## 3       1
## 4       1
## 5       1
## 6       1
## 7       1
## 8       1
## 9       1
## 10      1
## # ... with 1,722 more rows

# OR
ac_data[,3]

## # A tibble: 1,732 x 1
##   transect
##   <dbl>
## 1       1
## 2       1
## 3       1
## 4       1
## 5       1
## 6       1
## 7       1
## 8       1
## 9       1
## 10      1
## # ... with 1,722 more rows

# To return row 3
ac_data[3,]

## # A tibble: 1 x 10
##   desired_habitat polygon_id transect pin_number data_type species_code
##   <chr>          <chr>      <dbl>    <dbl> <chr>      <chr>
## 1 Weedy Control Weedy Con~      1      NA A.Belt    AMBPSI
## # ... with 4 more variables: scientific_name <chr>, native_non_native <chr>,
## #   functional_group <chr>, layer <chr>

# To return columns 1-3
ac_data[1:3]
```

```
## # A tibble: 1,732 x 3
##   desired_habitat      polygon_id  transect
##   <chr>              <chr>        <dbl>
## 1 Weedy Control      Weedy Control      1
## 2 Weedy Control      Weedy Control      1
## 3 Weedy Control      Weedy Control      1
## 4 Weedy Control      Weedy Control      1
## 5 Weedy Control      Weedy Control      1
## 6 Weedy Control      Weedy Control      1
## 7 Mulefat Scrub/sage Scrub OW-M7      1
## 8 Mulefat Scrub/sage Scrub OW-M7      1
## 9 Mulefat Scrub/sage Scrub OW-M7      1
## 10 Mulefat Scrub/sage Scrub OW-M7      1
## # ... with 1,722 more rows
```

```
# OR
```

```
ac_data[,1:3]
```

```
## # A tibble: 1,732 x 3
##   desired_habitat      polygon_id  transect
##   <chr>              <chr>        <dbl>
## 1 Weedy Control      Weedy Control      1
## 2 Weedy Control      Weedy Control      1
## 3 Weedy Control      Weedy Control      1
## 4 Weedy Control      Weedy Control      1
## 5 Weedy Control      Weedy Control      1
## 6 Weedy Control      Weedy Control      1
## 7 Mulefat Scrub/sage Scrub OW-M7      1
## 8 Mulefat Scrub/sage Scrub OW-M7      1
## 9 Mulefat Scrub/sage Scrub OW-M7      1
## 10 Mulefat Scrub/sage Scrub OW-M7      1
## # ... with 1,722 more rows
```

```
# To return rows 1-3
```

```
ac_data[1:3,]
```

```
## # A tibble: 3 x 10
##   desired_habitat polygon_id transect pin_number data_type species_code
##   <chr>          <chr>      <dbl>    <dbl> <chr>      <chr>
## 1 Weedy Control Weedy Con~      1        NA A.Belt    SOLSPP
## 2 Weedy Control Weedy Con~      1        NA A.Belt    AMBACA
## 3 Weedy Control Weedy Con~      1        NA A.Belt    AMBPSI
## # ... with 4 more variables: scientific_name <chr>, native_non_native <chr>,
## #   functional_group <chr>, layer <chr>
```

```
# Single brackets extract a list
```

```
ac_data[5]
```

```
## # A tibble: 1,732 x 1
##   data_type
##   <chr>
## 1 A.Belt
## 2 A.Belt
## 3 A.Belt
## 4 A.Belt
## 5 A.Belt
```

```
## 6 A.Belt
## 7 T.PI
## 8 T.PI
## 9 T.PI
## 10 T.PI
## # ... with 1,722 more rows
```

```
# Double brackets extract the items in a list
ac_data[[5]]
```

```
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[illegible]

[illegible]

[illegible]

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## [1689] "T.PI" "T.PI" "T.PI" "T.PI" "T.PI" "T.PI" "T.PI" "T.PI" "T.PI"
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## [1729] "A.Belt" "A.Belt" "A.Belt" "A.Belt"
```

```
ac_data[5,3]
```

```
## # A tibble: 1 x 1
##   transect
##   <dbl>
## 1       1
```

Data wrangling (part 1)

```
# Use the which() function to refer to specific observations in the code. In this example, we are replacing
ac_data$polygon_id[which(ac_data$polygon_id=="5M Buffer")] <- "5M BUFFER"
```

```
# There are a few ways to rename a column
```

```
# Method 1: use function [colnames]
```

```
df <- ac_data
```

```
colnames(df)[4:6] <- c("pin", "type_data", "code_species") # rename columns 4 through 6
```

```
# Method 2: use function [names]
```

```
df <- ac_data
```

```
names(df)[1] <- paste("habitat") # name column 1 "habitat"
```

```
View(df)
```

Data wrangling (part 2)

```
# There are a few ways to create a new column
```

```
# Method 1: Use the function [mutate]. Within the new column you can make transformations on your data.
```

```
oak_data2 <- oak_data %>%
```

```
  mutate(new_column=height_cm*2)
```

```
# Method 2: Create a new column using the [$] symbol
```

```
oak_data3 <- oak_data
```

```
oak_data3$height_rounded <- round(oak_data3$height_cm)
```

```
View(oak_data2)
```

```
View(oak_data3)
```

Creating tables

```
# To count observations by group, use the [group_by] and [count] functions. Here are two examples below

# agua chinon data example - number of observations per species code in each polygon and transect
richness1 <- ac_data %>%
  group_by(polygon_id, transect) %>%
  count(species_code)

View(richness1)

# richness2 - add column for the number of unique species per transect
richness2 <- richness1 %>%
  mutate(unique_species=n_distinct(species_code)) %>%
  select(transect, polygon_id, unique_species)
richness2 <- distinct(richness2)
View(richness2)

# oak data example - # of plantings watered or not watered by sampling group
oak_water_sampling_grp <- oak_data %>%
  group_by(sampling_group) %>%
  count(survival)

View(oak_water_sampling_grp)
View(oak_data)
```