RWorksheet_Cabico#5

Krestal Joy Cabico

2023-12-21

```
#..
install.packages("htmltools")
## Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.3'
## (as 'lib' is unspecified)
install.packages("pastecs")
## Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.3'
## (as 'lib' is unspecified)
install.packages("AppliedPredictiveModeling")
## Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.3'
## (as 'lib' is unspecified)
#1
library(Hmisc)
##
## Attaching package: 'Hmisc'
## The following objects are masked from 'package:base':
##
##
       format.pval, units
library(pastecs)
student score <- data.frame (</pre>
  Student = c(1,2,3,4,5,6,7,8,9,10),
   'Pre-test' = c(55, 54, 47, 57, 51, 61, 57, 54, 63, 58),
   'Post-test' = c(61, 60, 56, 63, 56, 63, 59, 56, 62, 61)
)
student_score
##
     Student Pre.test Post.test
## 1
        1
                 55
                             61
## 2
           2
                  54
                              60
           3
                   47
                              56
## 3
## 4
           4
                   57
                              63
           5
## 5
                  51
                              56
           6
                              63
## 6
                   61
## 7
           7
                   57
                              59
## 8
           8
                   54
                              56
           9
## 9
                   63
                              62
```

```
## 10
          10
                   58
                            61
summary(student score)
                                   Post.test
##
      Student
                     Pre.test
## Min. : 1.00 Min. :47.00 Min. :56.00
## 1st Qu.: 3.25 1st Qu.:54.00
                                  1st Qu.:56.75
## Median : 5.50 Median :56.00
                                  Median :60.50
## Mean : 5.50 Mean :55.70
                                  Mean :59.70
## 3rd Qu.: 7.75
                  3rd Qu.:57.75
                                  3rd Qu.:61.75
## Max. :10.00 Max. :63.00
                                  Max. :63.00
stat.desc(student score)
##
                  Student
                             Pre.test
                                         Post.test
## nbr.val
             10.0000000 10.00000000 10.00000000
## nbr.null
               0.0000000 0.00000000 0.00000000
## nbr.na
               0.0000000
                           0.00000000
                                       0.00000000
               1.0000000 47.00000000 56.00000000
## min
## max
              10.0000000 63.00000000 63.00000000
              9.0000000 16.00000000
                                       7.00000000
## range
## sum
               55.0000000 557.00000000 597.00000000
## median
              5.5000000 56.00000000 60.50000000
## mean
                5.5000000 55.70000000 59.70000000
## SE.mean
                0.9574271
                          1.46855938 0.89504811
## CI.mean.0.95 2.1658506
                          3.32211213
                                       2.02473948
## var
                9.1666667 21.56666667 8.01111111
                3.0276504 4.64399254
## std.dev
                                        2.83039063
## coef.var
                0.5504819
                           0.08337509 0.04741023
#2
agri_data <- c(10, 10, 10, 20, 20, 50, 10, 20, 10, 50, 20, 50, 20, 10)
fertilizer_levels <- factor(agri_data, levels = c(10, 20, 50), ordered = TRUE)
fertilizer_levels
## [1] 10 10 10 20 20 50 10 20 10 50 20 50 20 10
## Levels: 10 < 20 < 50
#3
#The following represents the 10 exercise levels in which "n" stands for none, "l" stands for light, an
exercise_levels <- c("l", "n", "n", "i", "l", "l", "n", "n", "i", "l")
exercise_levels
## [1] "1" "n" "n" "i" "1" "1" "n" "n" "i" "1"
#4
state <- c("tas", "sa", "qld", "nsw", "nsw", "nt", "wa", "qld",</pre>
"vic", "nsw", "vic", "qld", "qld", "sa", "tas", "sa", "nt",
"wa", "vic", "qld", "nsw", "nsw", "wa", "sa", "act", "nsw",
"vic", "vic", "act")
state_factor <- factor(state, levels = unique (state))</pre>
state factor
```

```
## [1] tas sa qld nsw nsw nt wa wa qld vic nsw vic qld qld sa tas sa nt wa
## [20] vic qld nsw nsw wa sa act nsw vic vic act
## Levels: tas sa qld nsw nt wa vic act
#5
incomes \leftarrow c(60, 49, 40, 61, 64, 60, 59, 54,
62, 69, 70, 42, 56, 61, 61, 61, 58, 51, 48,
65, 49, 49, 41, 48, 52, 46, 59, 46, 58, 43)
incmeans <- tapply(incomes, state_factor, mean)</pre>
incmeans
##
                          qld
                                   nsw
                                                               vic
        tas
                  sa
                                              nt
                                                       พล
                                                                        act
## 60.50000 55.00000 53.60000 57.33333 55.50000 52.25000 56.00000 44.50000
                       qld
              sa
                                nsw
                                           nt
#60.50000 55.00000 53.60000 57.33333 55.50000 52.25000
    vic
#56.00000 44.50000
#The average income for tax accountants varies across different states in Australia. In Tasmania, tax a
#6a
stdError <- function(x) sqrt(var(x)/length(x))</pre>
incster <- tapply(incomes, state_factor, stdError)</pre>
incster
##
        tas
                          qld
                  sa
## 0.500000 2.738613 4.106093 4.310195 4.500000 2.657536 5.244044 1.500000
#The accuracy of the average income estimate is very good for Tasmania, with a small standard error of
#7
data (Titanic)
Titanic
## , , Age = Child, Survived = No
##
         Sex
## Class Male Female
             0
##
     1st
##
     2nd
             0
                    0
##
     3rd
            35
                   17
##
    Crew
                    0
             0
##
   , , Age = Adult, Survived = No
##
##
##
         Sex
## Class Male Female
           118
                   4
##
     1st
##
     2nd
           154
                   13
##
           387
                   89
     3rd
##
     Crew 670
                   3
##
```

```
## , , Age = Child, Survived = Yes
##
##
         Sex
## Class Male Female
##
     1st
             5
##
     2nd
            11
                    13
##
     3rd
            13
                    14
##
             0
                    0
     Crew
##
##
   , , Age = Adult, Survived = Yes
##
##
         Sex
## Class Male Female
##
            57
     1st
                  140
##
     2nd
            14
                   80
##
     3rd
            75
                   76
##
     Crew 192
                   20
#7b
library(dplyr)
##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:pastecs':
##
##
       first, last
## The following objects are masked from 'package:Hmisc':
##
##
       src, summarize
## The following objects are masked from 'package:stats':
##
##
       filter, lag
## The following objects are masked from 'package:base':
##
       intersect, setdiff, setequal, union
##
Titanic_df <- as.data.frame(Titanic)</pre>
survivors <- Titanic_df %>%filter(Survived == "Yes")
head(survivors)
     Class
                    Age Survived Freq
              Sex
## 1
       1st
             Male Child
                              Yes
                                     5
## 2
       2nd
             Male Child
                              Yes
                                    11
## 3
       3rd
             Male Child
                              Yes
                                    13
             Male Child
      Crew
                              Yes
                                     0
## 5
       1st Female Child
                              Yes
                                     1
       2nd Female Child
## 6
                              Yes
                                    13
tail(survivors)
##
      Class
               Sex
                      Age Survived Freq
## 11
        3rd
              Male Adult
                               Yes
## 12 Crew
              Male Adult
                               Yes 192
## 13
        1st Female Adult
                               Yes 140
```

```
## 14
       2nd Female Adult
                             Yes
                                   80
                             Yes
## 15
       3rd Female Adult
                                   76
## 16 Crew Female Adult
                             Yes
                                   20
non_survivors <- Titanic_df %>%filter(Survived == "No")
head(non_survivors)
     Class
             Sex
                   Age Survived Freq
## 1
      1st
           Male Child
                             No
      2nd Male Child
                                   0
                             No
     3rd Male Child
## 3
                             No 35
## 4 Crew Male Child
                             No
## 5
     1st Female Child
                             No
                                   0
## 6 2nd Female Child
                             No
                                   0
tail(survivors)
##
     Class
             Sex Age Survived Freq
## 11
       3rd Male Adult
                             Yes
                             Yes 192
## 12 Crew Male Adult
## 13
      1st Female Adult
                             Yes 140
## 14
       2nd Female Adult
                             Yes 80
      3rd Female Adult
                                   76
## 15
                             Yes
## 16 Crew Female Adult
                             Yes
                                   20
#8
library(readr)
breastcancer_wisconsin <- read_csv("/cloud/project/RWorksheet_Cabico#6.R/breastcancer_wisconsin.csv", s
breastcancer wisconsin
## # A tibble: 699 x 11
##
          id clump_thickness size_uniformity shape_uniformity marginal_adhesion
##
        <dbl>
                       <dbl>
                                       <dbl>
                                                        <dbl>
## 1 1000025
                           5
                                                            1
                                                                              1
## 2 1002945
                           5
                                           4
                                                            4
## 3 1015425
                           3
                                           1
                                                            1
                                                                              1
## 4 1016277
                           6
                                           8
                                                            8
                                                                              1
## 5 1017023
                           4
                                                                              3
                                           1
                                                            1
## 6 1017122
                           8
                                          10
                                                           10
                                                                              8
## 7 1018099
                           1
                                           1
                                                            1
                                                                              1
## 8 1018561
                           2
                                                            2
                                                                              1
                                           1
## 9 1033078
                                           1
                                                            1
                                                                              1
## 10 1033078
                                           2
                                                            1
## # i 689 more rows
## # i 6 more variables: epithelial_size <dbl>, bare_nucleoli <chr>,
      bland_chromatin <dbl>, normal_nucleoli <dbl>, mitoses <dbl>, class <dbl>
#id: A unique identifier for each biopsy. clump_thickness: Describes the thickness of the clumps in the
#8d.1
sd_clump_thickness <- sd(breastcancer_wisconsin$clump_thickness)</pre>
sample_size <- length(breastcancer_wisconsin$clump_thickness)</pre>
sem_clump_thickness <- sd_clump_thickness / sqrt(sample_size)</pre>
```

```
cat("Standard Error of the Mean for Clump Thickness:", sem_clump_thickness, "\n")
## Standard Error of the Mean for Clump Thickness: 0.1065011
mean_marginal_adhesion <- mean(breastcancer_wisconsin$marginal_adhesion)</pre>
sd_marginal_adhesion <- sd(breastcancer_wisconsin$marginal_adhesion)</pre>
cv_marginal_adhesion <- (sd_marginal_adhesion / mean_marginal_adhesion) * 100</pre>
cat("Coefficient of Variation for Marginal Adhesion:", cv marginal adhesion, "%\n")
## Coefficient of Variation for Marginal Adhesion: 101.7283 %
\#8d.3
num null bare nuclei <- sum(is.na(breastcancer wisconsin$bare nucleoli))</pre>
cat("Number of null values for Bare Nuclei:", num_null_bare_nuclei, "\n")
## Number of null values for Bare Nuclei: 15
\#8d.4
mean_bland_chromatin <- mean(breastcancer_wisconsin$bland_chromatin, na.rm = TRUE)</pre>
sd bland chromatin <- sd(breastcancer wisconsin$bland chromatin, na.rm = TRUE)
cat("Mean for Bland Chromatin:", mean_bland_chromatin, "\n")
## Mean for Bland Chromatin: 3.437768
cat("Standard Deviation for Bland Chromatin:", sd_bland_chromatin, "\n")
## Standard Deviation for Bland Chromatin: 2.438364
\#8d.5
mean_value <- mean(breastcancer_wisconsin$shape_uniformity, na.rm = TRUE)</pre>
se <- sd(breastcancer_wisconsin$shape_uniformity, na.rm = TRUE) / sqrt(length(breastcancer_wisconsin$sh
confidence_level <- 0.95</pre>
margin_of_error <- qt((1 + confidence_level) / 2, df = length(breastcancer_wisconsin$shape_uniformity)</pre>
confidence interval <- c(mean value - margin of error, mean value + margin of error)
cat("Confidence Interval (", confidence_level * 100, "%) :", confidence_interval, "\n")
## Confidence Interval ( 95 %) : 2.986741 3.428138
#8d
column_names <- names(breastcancer_wisconsin)</pre>
column_names
## [1] "id"
                             "clump_thickness"
                                                 "size_uniformity"
## [4] "shape uniformity"
                             "marginal adhesion" "epithelial size"
## [7] "bare nucleoli"
                             "bland chromatin"
                                                 "normal_nucleoli"
## [10] "mitoses"
                             "class"
```

```
#8e
malignant_count <- sum(breastcancer_wisconsin$class == 4)</pre>
malignant_percentage <- (malignant_count / nrow(breastcancer_wisconsin)) * 100</pre>
cat("Percentage of respondents who are malignant:", malignant_percentage, "%\n")
## Percentage of respondents who are malignant: 34.47783 %
#9
library(readr)
abalone <- read csv("/cloud/project/RWorksheet Cabico#6.R/abalone.csv")
## Rows: 4177 Columns: 9
## -- Column specification -----
## Delimiter: ","
## chr (1): Sex
## dbl (8): Length, Diameter, Height, Whole weight, Shucked weight, Viscera wei...
## 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.
abalone
## # A tibble: 4,177 x 9
##
           Length Diameter Height `Whole weight` `Shucked weight` `Viscera weight`
      <chr> <dbl>
                     <dbl> <dbl>
                                           <dbl>
                                                            <dbl>
                                                                             <dbl>
## 1 M
            0.455
                     0.365 0.095
                                           0.514
                                                           0.224
                                                                            0.101
                     0.265 0.09
## 2 M
            0.35
                                           0.226
                                                           0.0995
                                                                           0.0485
## 3 F
            0.53
                     0.42 0.135
                                           0.677
                                                           0.256
                                                                           0.142
## 4 M
            0.44
                     0.365 0.125
                                           0.516
                                                           0.216
                                                                           0.114
## 5 I
           0.33
                     0.255 0.08
                                           0.205
                                                           0.0895
                                                                           0.0395
## 6 I
           0.425
                     0.3
                            0.095
                                           0.352
                                                           0.141
                                                                           0.0775
## 7 F
            0.53
                     0.415 0.15
                                           0.778
                                                           0.237
                                                                           0.142
## 8 F
            0.545
                     0.425 0.125
                                           0.768
                                                           0.294
                                                                           0.150
## 9 M
            0.475
                     0.37
                            0.125
                                           0.509
                                                           0.216
                                                                           0.112
## 10 F
            0.55
                     0.44
                            0.15
                                           0.894
                                                           0.314
                                                                           0.151
## # i 4,167 more rows
## # i 2 more variables: `Shell weight` <dbl>, Rings <dbl>
head(abalone)
## # A tibble: 6 x 9
##
          Length Diameter Height `Whole weight` `Shucked weight` `Viscera weight`
##
    <chr> <dbl>
                    <dbl> <dbl>
                                          <dbl>
                                                           <dbl>
                                                                            <dbl>
## 1 M
           0.455
                    0.365 0.095
                                          0.514
                                                          0.224
                                                                           0.101
## 2 M
           0.35
                    0.265 0.09
                                          0.226
                                                          0.0995
                                                                           0.0485
## 3 F
           0.53
                    0.42
                                          0.677
                                                          0.256
                          0.135
                                                                           0.142
## 4 M
           0.44
                    0.365 0.125
                                          0.516
                                                          0.216
                                                                           0.114
## 5 I
           0.33
                    0.255 0.08
                                          0.205
                                                          0.0895
                                                                           0.0395
           0.425
                    0.3
                           0.095
                                          0.352
                                                          0.141
                                                                           0.0775
## # i 2 more variables: `Shell weight` <dbl>, Rings <dbl>
summary(abalone)
```

Length

Diameter

Height

##

Sex

```
## Length:4177
                     Min. :0.075
                                    Min. :0.0550
                                                    Min. :0.0000
## Class :character
                     1st Qu.:0.450
                                    1st Qu.:0.3500
                                                    1st Qu.:0.1150
## Mode :character
                     Median :0.545
                                    Median :0.4250
                                                    Median :0.1400
##
                     Mean :0.524
                                    Mean :0.4079
                                                    Mean :0.1395
                     3rd Qu.:0.615
                                    3rd Qu.:0.4800
                                                     3rd Qu.:0.1650
##
##
                     Max.
                            :0.815
                                    Max.
                                          :0.6500
                                                    Max. :1.1300
##
   Whole weight
                   Shucked weight
                                   Viscera weight
                                                     Shell weight
         :0.0020
                   Min. :0.0010
                                   Min. :0.0005
                                                    Min.
                                                          :0.0015
## Min.
                   1st Qu.:0.1860
                                                    1st Qu.:0.1300
  1st Qu.:0.4415
                                   1st Qu.:0.0935
  Median :0.7995
                   Median :0.3360
                                   Median :0.1710
                                                    Median :0.2340
  Mean :0.8287
                   Mean
                         :0.3594
                                   Mean :0.1806
                                                    Mean
                                                         :0.2388
##
   3rd Qu.:1.1530
                   3rd Qu.:0.5020
                                   3rd Qu.:0.2530
                                                    3rd Qu.:0.3290
##
  Max.
         :2.8255
                   Max. :1.4880
                                   Max. :0.7600
                                                    Max. :1.0050
##
       Rings
## Min. : 1.000
  1st Qu.: 8.000
##
## Median: 9.000
## Mean : 9.934
## 3rd Qu.:11.000
## Max. :29.000
```