Final Sanyika 2023-12-07 Python version: https://colab.research.google.com/drive/1H4mIVk9aYLchsxtm73B4AjsVX8lzlpzn?usp=sharing library(dplyr) ## ## Attaching package: 'dplyr' ## The following objects are masked from 'package:stats': ## ## filter, lag The following objects are masked from 'package:base': ## ## ## intersect, setdiff, setequal, union library(tidyr) setwd("~/Desktop") math <- read.csv("student math clean.csv")</pre> lang <- read.csv("student portuguese clean.csv")</pre> QUESTION 1 (a) Using the Portugese language data set for the guestion. Consider students only if they are at least 18 years or older. For these students, they each received a grade 1 and a grade 2. Create a variable for each student that is the larger of these two scores. Then find the average score of this maximum grouping by the combination of mother job and father job. Of these combinations (i.e. mother job and father job) with more than 5 cases, what combination of jobs had the highest average of the larger of their two test scores? #see column names in the lang data set colnames(lang) "school" [1] "student id" "sex" ## [4] "age" "address type" "family size" ## [7] "parent status" "mother education" "father education" ## ## [10] "mother job" "father job" "school choice reason" ## [13] "guardian" "travel time" "study time" ## [16] "class failures" "school support" "family support" ## [19] "extra paid classes" "activities" "nursery school" "internet access" ## [22] "higher ed" "romantic relationship" ## [25] "family relationship" "free time" "social" ## [28] "weekday alcohol" "health" "weekend alcohol" ## [31] "absences" grade 1" "grade 2" ## [34] "final grade" #filter for age > 18 lang %>% filter(age >= 18) %>% #add variable that calculates the max between 2 grades mutate(max grade = pmax(grade 1, grade 2)) %>% group by(father\_job, mother\_job) %>% #calculate ave score and count of each group summarize(ave score = mean(max grade), cases = n()) %>% #filter cases>5 and select top average score filter(cases>5) %>% arrange(-ave score) %>% head(1) ## `summarise()` has grouped output by 'father job'. You can override using the ## `.groups` argument. ## # A tibble: 1 × 4 ## # Groups: father job [1] father job mother job ave score cases ## ## <chr> <dbl> <int> <chr> 7 ## 1 services teacher 13.4 b. Using the math data set, find the mean age by sex and school. Then compute the mean final score by sex and school. Then present the results in a table where each row is a school and each column is an average of the particular variable for one sex (i.e. the first column is mean age females, the second column is mean age males, etc.). #see names of columns colnames(math) ## [1] "student id" "school" [4] "age" ## "address type" "family size" "mother education" "father education" [7] "parent status" ## ## [10] "mother job" "father job" "school choice reason" "travel time" ## [13] "guardian" "study time" ## [16] "class failures" "school support" "family support" ## [19] "extra paid classes" "nursery school" "activities" ## [22] "higher ed" "internet access" "romantic relationship" "free time" "social" ## [25] "family relationship" ## [28] "weekday alcohol" "health" "weekend alcohol" ## [31] "absences" "grade 1" "grade 2" ## [34] "final grade" #calculate average age by sex and school age mean <- math %>% group by(sex, school) %>% summarize(mean age = mean(age)) ## `summarise()` has grouped output by 'sex'. You can override using the `.groups` ## argument. #calculate average final grade by sex and school score mean <- math %>% group by(sex, school) %>% summarize(mean final = mean(final gra de)) ## `summarise()` has grouped output by 'sex'. You can override using the `.groups` ## argument. #join by sex and school and pivot wider score mean %>% left join(age mean, by = c("sex", "school")) %>% pivot wider(names from = "sex", values from = c("mean age", "mean final")) ## # A tibble:  $2 \times 5$ ## school mean age F mean age M mean final F mean final M <dbl> ## <dbl> <dbl> <dbl> ## 1 GP 16.6 16.5 9.97 11.1 ## 2 MS 17.8 18.2 9.92 9.76 c. Reproduce the provided plot. library(ggplot2) #select relevant columns and add column to state the subject lang timeGrade <- lang %>% select(sex, study time, final grade) %>% mutate(subject = "port") math timeGrade <- math %>% select(sex, study time, final grade) %>% mutate(subject = "math") #stack the two data frames on top of each other school <- rbind(lang timeGrade, math timeGrade)</pre> #set the order of study time for it to display on the x axis order time <- c("<2 hours", "2 to 5 hours", "5 to 10 hours", ">10 hours") school\$study time <- factor(school\$study time, levels = order time)</pre> #create plot with final grade as a response to study time and stratify by sex ggplot(data = school, aes( x = study time,y = final grade, color = sex) ) + #create box plot faceted by subject geom boxplot() + facet wrap( ~ subject) + xlab("Study Time") + ylab("Final Grade") math port 20 -15 -Final Grade sex 5 -0 -<2 hours 2 to 5 hours5 to 10 hours >10 hours <2 hours 2 to 5 hours5 to 10 hours >10 hours Study Time d. Using the math scores data set, for each grade on the first exam, find the average grade on the final exam for all the students who had the same score on exam one. That is, for all students who score, for example, a 10 on the first exam, what was the average final score for this group of students. Do this for all grades on the first exam. Repeat this for the second exam (i.e. For each grade on the second exam, find the average grade on the final exam for all the students who had the same score on exam two). Plot the exam score (first or second) on the x-axis and the average final score on the y-axis using color to indicate whether the point was the first or second exam. The final plot should look like the one provided. #calculate the mean of grade 1 and add a variable that states the exam number one <- math %>% group by(grade 1) %>% summarize(final = mean(final grade)) %>% mutate(type = "one") %>% #rename the variable in preparation for stacking rename(grade = grade 1) #calculate the mean of grade 2 and add a variable that states the exam number two <- math %>% group by(grade 2) %>% summarize(final = mean(final grade)) %>% mutate(type = "two") %>% #rename the variable in preparation for stacking rename(grade = grade 2) #stack the data frames on top of each other allgrade <- rbind(one, two) #create plot of final as a response to grade ggplot(data = allgrade, aes( x = grade, y = final,color = type ) + geom point() 20 -15 type final 10 one two 5 -0 -10 15 grade **QUESTION 2** spot <- read.csv("spotify songs.csv")</pre> a. Plot a line plot of the release year on the x-axis and the number of songs released in this year on the yaxis. library(lubridate) ## ## Attaching package: 'lubridate' ## The following objects are masked from 'package:base': ## ## date, intersect, setdiff, union release <- spot %>% #add column for date year mutate(release year = year(as.Date(track album release date))) %>% #count songs in each year group by (release year) %>% summarize(song num = n())ggplot(data = release, aes( x = release year,y = song num) + geom line() + xlab("Release Year") + ylab("Number of Songs") ## Warning: Removed 1 row containing missing values (`geom line()`). 7500 -Number of Songs 2500 -0 -2000 1960 1980 2020 Release Year b. Remove all the songs that only have the release year (as opposed to the full date of release). For each release month, find the 5 most danceable songs (higher scores are better). This will give you a list of 60 total songs (i.e. 12 months times 5 songs each month equals 60). Of these 60 songs, what is the name of the playlist that contains the song with the highest tempo? Of the same 60 songs, what is the name of the playlist that contains the song with the lowest tempo? Finally, what is the most common genre among these 60 songs, and how many songs belong to that genre? #filter for dates in the correct format topDance <- spot %>% filter(as.Date(track album release date) == format(as.Date(track album release date), format = "%Y-%m-%d")) %>% #add variable for release month mutate(release month = month(as.Date(track album release date))) %>% #remove any repeated track IDs distinct(track id, .keep all = TRUE) %>% #select songs with top 5 danceability scores for each month group by(release month) %>% slice max(order by = danceability, n=5) %>% #due to tie for 5th in scores for December, remove one of the tracks filter(!(track id %in% c("4ih3rak54fNYhweawmNZcj"))) high tempo <- topDance %>% arrange(-tempo) %>% select(playlist name, tempo) %>% head(1)## Adding missing grouping variables: `release month` print(high tempo) ## # A tibble: 1 × 3 ## # Groups: release month [1] release month playlist name ## tempo <dbl> <chr> <dbl> ## ## 1 2 Dirty South Rap Classics by DJ HOTSAUCE 139. low tempo <- topDance %>% #select slowest tempo arrange(tempo) %>% select(playlist name, tempo) %>% head(1)## Adding missing grouping variables: `release month` print(low\_tempo) ## # A tibble: 1 × 3 ## # Groups: release\_month [1] ## release\_month playlist\_name tempo <dbl> <chr> ## <dbl> ## 1 9 Zona Trap 103. comm genre <- topDance %>% #count songs in each genre group by(playlist genre) %>% summarize(num songs = n()) %>% #select genre with most songs arrange(-num songs) %>% head(1) print(comm genre) ## # A tibble: 1 × 2 playlist genre num songs ## ## <chr> <int> 27 ## 1 rap c. You are a wedding DJ and you are working with a night-mare couple and they have a lot of very specific requests. They want their first dance to be exactly 10 minutes long and consist of two back to back songs (they cannot be the same song!) by the artist Depeche Mode. Find the two Depeche Mode songs in this data set when played back to back the duration is as close as possible to 10 minutes (hint: a cross join may be useful here). (Note1: For loops are not allowed in your solution.) #filter for artist and select the relevant columns DM songs <- spot %>% filter(track artist == "Depeche Mode") %>% select(track name, dur ation ms) #create a duplicate DM songs2 <- DM songs #cross join the two data frames merge(DM songs, DM songs2, by=NULL) %>% #sum the durations and convert to minutes by dividing by 60000 mutate(minutes = (duration ms.x+duration ms.y)/60000) %>% #filter for row where minutes are closest to 10 filter(minutes == minutes[which.min(abs(minutes - 10))]) ## track name.x duration ms.x ## 1 Enjoy The Silence - 2006 Remastered Version 372813 ## 2 Strangelove - 7" Version 227413 ## track\_name.y duration\_ms.y minutes ## 1 Strangelove - 7" Version 227413 10.00377 ## 2 Enjoy The Silence - 2006 Remastered Version 372813 10.00377 'Enjoy The Silence - 2006 Remastered Version' and 'Strangelove - 7" Version' by Depeche Mode are the songs I'd play. QUESTION 3 (a) Lets say that you have a population that follows a normal distribution with a mean of 10 and a variance of 20. Generate a simple random sample from this population with sample size n = 25. From this sample, compute the sample mean (i.e.  $x^{-}$ ). Repeat this process (i.e. generate a sample of size n = 25 and compute x<sup>-</sup>) a large number of times (say 5000) and store the value of x<sup>-</sup> each time. Compute the standard deviation of this collection of x<sup>-</sup>'s. This is a simulated approximation of the standard error of x<sup>-</sup>. What is the value that you obtain? set.seed(1234)mean < -10variance <- 20 sd <- sqrt(variance)</pre> n < -25#number of simulations nsim < -5000sample\_means <- numeric(nsim)</pre> for (i in 1:nsim){ sample\_data <- rnorm(n, mean, sd)</pre> sample means[i] <- mean(sample data)</pre> } mean error25 = sd(sample means)mean error25 ## [1] 0.8899491 b. Repeat the previous question but use the sample median instead of the mean. What is the approximate value of the standard error of the sample median? Is this bigger, smaller, or the same as the standard error of the mean? sample medians <- numeric(nsim)</pre> for (i in 1:nsim){ sample data <- rnorm(n, mean, sd)</pre> sample medians[i] <- median(sample data)</pre> } median error25 = sd(sample medians) median\_error25 ## [1] 1.116712 This is bigger than the standard error of the mean. c. Now repeat both of the previous questions with values of n equal to 10, 25 (you already have this one), 50, 100, 250, 500, 1000. (Note: n is the sample size and it is not the same as the number of simulations!) Now plot n, the sample size on the x-axis versus the standard error on the y-axis as a line plot with points at each of the values of n that were used. There should be one line for median and one line for mean you should use color to distinguish between them. #n=10n=10sample means <- numeric(nsim)</pre> sample medians <- numeric(nsim)</pre> for (i in 1:nsim){ sample\_data <- rnorm(n, mean, sd)</pre> sample means[i] <- mean(sample data)</pre> sample\_medians[i] <- median(sample\_data)</pre> } mean error10 = sd(sample means) median\_error10 = sd(sample\_medians) #n=50n=50sample means <- numeric(nsim)</pre> sample medians <- numeric(nsim)</pre> for (i in 1:nsim){ sample\_data <- rnorm(n, mean, sd)</pre> sample means[i] <- mean(sample data)</pre> sample\_medians[i] <- median(sample\_data)</pre> } mean error50 = sd(sample means) median\_error50 = sd(sample\_medians) #n=100n = 100sample means <- numeric(nsim)</pre> sample medians <- numeric(nsim)</pre> for (i in 1:nsim){ sample\_data <- rnorm(n, mean, sd)</pre> sample means[i] <- mean(sample data)</pre> sample\_medians[i] <- median(sample data)</pre> mean error100 = sd(sample means) median\_error100 = sd(sample\_medians) #n=250n = 250sample means <- numeric(nsim)</pre> sample medians <- numeric(nsim)</pre> for (i in 1:nsim){ sample data <- rnorm(n, mean, sd)</pre> sample\_means[i] <- mean(sample\_data)</pre> sample\_medians[i] <- median(sample\_data)</pre> } mean\_error250 = sd(sample\_means) median\_error250 = sd(sample\_medians) #n=500n = 500sample\_means <- numeric(nsim)</pre> sample\_medians <- numeric(nsim)</pre> for (i in 1:nsim){ sample\_data <- rnorm(n, mean, sd)</pre> sample\_means[i] <- mean(sample\_data)</pre> sample\_medians[i] <- median(sample\_data)</pre> } mean\_error500 = sd(sample\_means) median\_error500 = sd(sample\_medians) #n=1000n=1000sample\_means <- numeric(nsim)</pre> sample medians <- numeric(nsim)</pre> for (i in 1:nsim){ sample\_data <- rnorm(n, mean, sd)</pre> sample\_means[i] <- mean(sample\_data)</pre> sample\_medians[i] <- median(sample\_data)</pre> } mean\_error1000 = sd(sample\_means) median\_error1000 = sd(sample\_medians) #create data frame errors = data.frame(N = c(10, 25, 50, 100, 250, 500, 1000), error\_type = c("mean\_error", "mean\_error", "mean\_error" "mean\_error", "mean\_error", "mean\_error", "mean\_error", "median\_error", "median\_error", "median\_error", "median\_error", "median\_error", "median\_error", "median\_error"), error\_value= c(mean\_error10, mean\_error25, mean\_error50, mean\_error100, mean\_error250, mean\_error500, mean\_error1000, median\_error10, median\_error25, median\_error50, median\_error100, median\_error250, median\_error500, median\_error1000) ) ggplot(data = errors, aes(x=N, y=error\_value, color=error\_type))+ geom\_line() 1.5 error\_type mean\_error median\_error 0.5 -250 500 750 1000 Ν