

Fundamentals in R: Data Exploration









Module 4 - Data Expoloration

In this module we'll cover subsetting, summarizing and visualizing data. We'll cover how to remove duplicates, learning about loops and custom functions in the process. Lastly, we'll see how we can merge two data sets.

The objectives are

- Become comfortable in subsetting using [] and subset()
- Understand various ways to remove duplicates
- Be able to summarize data numerically and visually
- Be able to aggregate and merge data
- Have a basic understanding of how to write your own custom function



Load Your Work



load("data/Module3_songData.RData")





We can subset into the data frame and select specific rows & columns

```
songSubset <- songData[1:100, 1:10]
class(songSubset)
## [1] "data.frame"
rm(songSubset)</pre>
```





... alternatively

```
songSubset2 <- songData[, c(1, 3, 3, 9)]
rm(songSubset2)</pre>
```

See

?Extract



... we can *subset* using column names (or row names) instead of column numbers (or row numbers)

```
songSubset3 <- songData[, c("artist.name", "artist.hotttnesss", "artist
    "artist.location", "artist.latitude", "artist.longitude", "song.hot"
    "tempo", "duration", "term", "hfRatio")]</pre>
```

... or we can *subset* using a vector of logicals. The vector must be the same length as the number of columns (or rows, if subsetting rows)

```
songSubset3 <- songData[, c(FALSE, FALSE, FALSE, FALSE, TRUE, TRUE, TRUE
TRUE, TRUE, FALSE, TRUE, TRUE,
FALSE, FALSE, TRUE, FALSE, TRUE, TRUE)]</pre>
```



Instead of typing out the entire vector, we can create it

```
## [1] FALSE FALSE FALSE FALSE TRUE TRUE TRUE TRUE TRUE TRUE FALSE
## [12] FALSE TRUE
## [34] FALSE TRUE TRUE
```

songSubset3 <- songData[, colSelect]</pre>





Do we have any duplicate artists in our subset? If so, which are the 5 most duplicated artists?

summary(songSubset3)

```
##
                artist.name artist.latitude artist.longitude
   Mario Rosenstock
                         : 13
                                Min. :-41
                                              Min. :-162
   Aerosmith
                                              1st Qu.: -93
##
                         : 12 1st Qu.: 34
## Phil Collins
                         : 12 Median : 39
                                             Median: -80
   Sugar Minott
                         : 12 Mean : 37
##
                                              Mean : -64
   The Jackson Southernaires: 12
                                3rd Qu.: 44
                                              3rd Qu.: -9
```



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Create a new data frame called songAerosmith containing all rows of songSubset3 where the artist name is "Aerosmith". Your code should return a data frame with 12 rows and 11 columns.

Bonus: Take a look at the documentation for the subset() functions. See if you can do the same using subset()

How many missing observations of song.hottnesss does this subset have?



Solution

```
songAerosmith <- songSubset3[rowSelect, ]
dim(songAerosmith)

## [1] 12 11

# alternatively, using subset() instead of []
songAerosmith <- subset(songSubset3, subset = artist.name == "Aerosmith")</pre>
```

rowSelect <- songSubset3\$artist.name == "Aerosmith"</pre>



Solution

summary(songAerosmith) # identify number of NA's for song.hotttnesss

```
##
                                     artist.name artist.latitude
##
   Aerosmith
                                                :12 Min. : NA
   111
                                                : 0 1st Qu.: NA
##
  (hed) p.e.
                                                : 0 Median : NA
##
##
   :Blacks On :Blondes
                                                : 0 Mean :NaN
    [kaleidoskop]
                                                : 0 3rd Qu.: NA
##
. . .
```

summary(songAerosmith\$song.hotttnesss)

```
## Min. 1st Qu. Median Mean 3rd Qu. Max. NA's
## 0.415 0.434 0.505 0.530 0.588 0.731 6
```





Working with Missing Values

Note that song.hotttness variable in songAerosmith data frame has 6 missing values.

To replace missing values with the average value across all non-missing observations

```
mean(songAerosmith$song.hotttnesss, na.rm = TRUE) # calculate replacem
```

```
## [1] 0.5304
```

rowIndex <- is.na(songAerosmith\$song.hotttnesss) # identify</pre>





Working with Missing Values

songAerosmith[rowIndex,] # subset

```
##
     artist.name artist.latitude artist.longitude artist.location
## 389
        Aerosmith
                               NΑ
                                               NΑ
                                                       Boston, MA
## 393
       Aerosmith
                               NΑ
                                               NΑ
                                                       Boston, MA
## 394
      Aerosmith
                               NA
                                               NA
                                                       Boston, MA
## 395 Aerosmith
                               NA
                                               NA
                                                       Boston, MA
## 396 Aerosmith
                               NA
                                               NA
                                                       Boston, MA
```

. . .





Working with Missing Values

```
songAerosmith$song.hotttnesss[rowIndex] <- mean(songAerosmith$song.hott
    na.rm = TRUE) # replace
songAerosmith$song.hotttnesss # verification 1
  [1] 0.5304 0.4150 0.4433 0.4314 0.5304 0.5304 0.5304 0.5304 0.5955 0.73
## [11] 0.5662 0.5304
mean(songAerosmith$song.hotttnesss) # verification
```



[1] 0.5304



Removing Duplicates: How can we ensure that we only have one entry for each artist? What will choose to do with the values in the other columns

songAerosmith

##	artist.name artist.latitude artist.longitude artist.location					
##	389	Aerosmith	NA	NA	Boston,	MA
##	390	Aerosmith	NA	NA	Boston,	MA
##	391	Aerosmith	NA	NA	Boston,	MA
##	392	Aerosmith	NA	NA	Boston,	MA
##	393	Aerosmith	NA	NA	Boston,	MA





We can remove duplicates, ignoring other columns... keep just 1 observation (first occurring) for values of other columns

```
songDedup <- songSubset3[!duplicated(songSubset3[1]), ]
summary(songDedup)</pre>
```



We lose some information here... what?

```
songDedup[songDedup$artist.name == "Aerosmith", ]
##
     artist.name artist.latitude artist.longitude artist.location
## 389
        Aerosmith
                               NA
                                               NΑ
                                                      Boston, MA
     artist.hotttnesss artist.familiarity tempo duration song.hotttnesss
##
                               0.8725 108.7
## 389
               0.6107
                                                425
                                                               NA
##
       term hfRatio
## 389 rock 0.6999
```

We have song hottnesss information about half of the observations corresponding to Aerosmith. However, our deduping exercise left us with just the observation missing a value for song hottnesss.



A better way may be to average song hottnesss across artists. Instead of obtaining 1 observation for each artist, we can return the average of all observations for that artist

We can write a for loop, we'll replace the value of song hotness obtained in the previous de-duping with the average song hotness

```
for (i in unique(songSubset3$artist.name)) {
    sSub <- subset(songSubset3, artist.name == i)
    meanHot <- mean(sSub$song.hotttnesss, na.rm = TRUE)
    songDedup[songDedup$artist.name == i, "song.hotttnesss"] <- meanHot
}</pre>
```



```
head(songDedup)
summary(songDedup)
songDedup[songDedup$artist.name == "Aerosmith", ]
     artist.name artist.latitude artist.longitude artist.location
##
## 389
        Aerosmith
                               NA
                                               NA
                                                      Boston, MA
     artist.hotttnesss artist.familiarity tempo duration song.hotttnesss
##
              0.6107
                               0.8725 108.7 425
                                                          0.5304
## 389
##
       term hfRatio
## 389 rock 0.6999
```

That took some time! There's a better way – the aggregate() function. See





R provides a multitude of tools for exploratory data analysis, summarization and visualization.

Useful functions for birds eye view summaries: structure using str() and basic statistical summary using summary()

```
str(songSubset3)
summary(songSubset3)
```





Cross tabulations using xtabs() returns to number of observations at each value of a given variable.

xtabs(~term, songSubset3)

```
## term
##
         ballad
                       blues
                                   chanson
                                               chill-out
                                                                country
             47
                          385
                                        208
                                                       90
                                                                     75
##
##
      dancehall
                        disco
                                       dub easy listening
                                                                   folk
##
             97
                           58
                                         19
                                                       65
                                                                    141
##
           funk
                                    hip hop
                                                    house
                                                                   jazz
                       grunge
. . .
```





Cross tabulations using xtabs can be used to return the number of observations at each combination of two or more variables.

```
songSubset3$song.hotttnesssCut <- cut(songSubset3$song.hotttnesss, break
na.rm = TRUE)[2:5], labels = c("low", "med", "high"))
xtabs(~term + song.hotttnesssCut, songSubset3)</pre>
```

```
##
                  song.hotttnesssCut
## term
                  low med high
##
    ballad
                   10
    blues
                   45 35 24
##
                   30 25 6
##
    chanson
    chill-out
                 18 12
                            14
##
```





prop.table() can be used to return percent instead of count. The margin argument specifies percent of what – total across rows, columns or both.

Which term has the highest proportion of songs rated *highly* in hotttnesss?

```
prop.table(xtabs(~term + song.hotttnesssCut, songSubset3), margin = 1)
```

```
##
                    song.hotttnesssCut
## term
                         low
                                 med
                                        high
                    0.47619 0.42857 0.09524
##
     ballad
##
     blues
                    0.43269 0.33654 0.23077
##
     chanson
                    0.49180 0.40984 0.09836
##
     chill-out
                    0.40909 0.27273 0.31818
```



. . .

Sorting Data

To better identify which terms have the highest proportion of songs rates .538 and higher, we reorder the proportion table just generated by the third column.

```
songTable <- prop.table(xtabs(~term + song.hotttnesssCut, songSubset3),</pre>
colIndex <- order(songTable[, 3], decreasing = TRUE)</pre>
songTable <- songTable[colIndex, ]</pre>
head(songTable, 3)
##
           song.hotttnesssCut
## term
                low
                        med
                              high
##
     punk 0.12057 0.2624 0.6170
##
     techno 0.06667 0.3333 0.6000
```





Use xtabs() and order() to identify the top 10 terms by observation count in songSubset3.

Poll: Which term is most frequently occuring in our dataset?

- pop
- rock
- punk
- blues
- techno
- disco
- hip hop







Solution

```
songTable2 <- xtabs(~term, songSubset3)</pre>
songTable2 <- songTable2[order(songTable2, decreasing = TRUE)]</pre>
songTable2[2:11] # rock is the most frequently occurring term
## term
##
    rock
           pop metal jazz blues hip hop
                                           house
                                                          punk
                                                    rap
    1619
            777
                  485
                      461 385
                                       364
                                             304
                                                    283
                                                           213
##
## chanson
##
      208
```



R ships with a variety of tools for visualizing data. Using plot() on a data frame with multiple (numeric!) columns returns a scatter plot matrix.

```
cols <- names(songSubset3)[sapply(songSubset3, is.numeric)]
cols

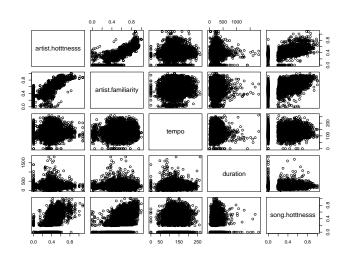
## [1] "artist.latitude" "artist.longitude" "artist.hotttnesss"
## [4] "artist.familiarity" "tempo" "duration"
## [7] "song.hotttnesss" "hfRatio"

plot(songData[, cols[3:7]])</pre>
```











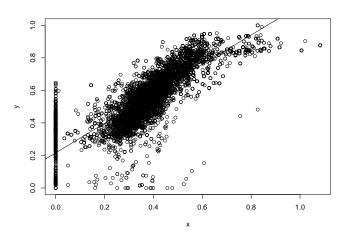


Using plot on two vectors (two arguments) creates a single scatter plot. We add an abline showing the results of linear regression on top.

```
x <- songSubset3$artist.hotttnesss
y <- songSubset3$artist.familiarity
plot(x, y)
abline(lm(y ~ x))</pre>
```











Summarizing the model returned by the linear regression

```
##
## Call:
## lm(formula = y ~ x)
##
## Residuals:
## Min 1Q Median 3Q Max
...
```



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Exercise

Create a new data frame that excludes observations of songSubset3 where artist.hotttnesss or artist familiarity is 0. Regenerate the previous plot & model based on this new data frame.

Take a look at the documentation for plot. Add axis titles (xlab, ylab) and a main title (main).

Take a look at the documentation for abline. Change the color of the line to "blue" (col), line width to "3" (lwd) and the line type to "twodash" (lty)

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Solution

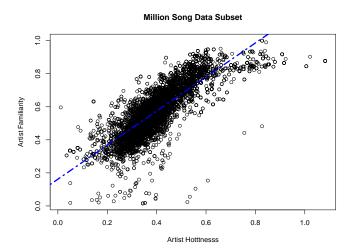


(()





Solution





Note that, by default...

```
* `plot([2+ column dataframe])` returns a scatter plot matrix.
```

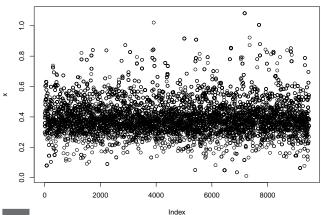
- * `plot([two vectors])` returns one scatter plot.
- * `plot([one vector])` returns one scatter plot.
- * `plot([lm object])` returns 4 plots that can be used to evaluate the
- * `plot([density object])` returns one scatter plot.
- * `hist([one vector])` returns a histogram.





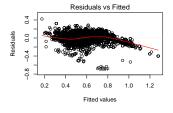


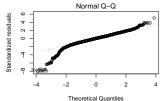
plot(x)

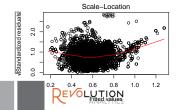


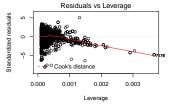






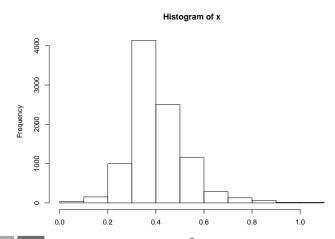








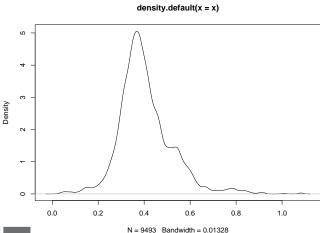
hist(x)







plot(density(x))







Plot is flexible function: the output depends on the input, the input can be a variety of different R objects.

For each valid input object class, a method is defined. When plot(data.frame) is run, plot.data.frame gets called which, in turn, calls pairs(). You can see plot.data.frame specific arguments by looking at the help file.

methods(plot)
help(plot.data.frame)





Additional Basic plots.

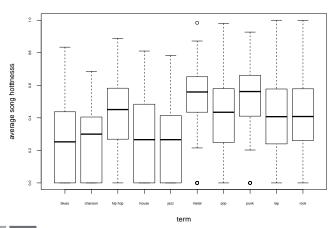
```
load("data/songTermHot.RData")
```

```
boxplot(song.hotttnesss ~ term, data = songTermHot, cex.axis = 0.5, xla
ylab = "average song hotttnesss")
```





Additional Basic plots.

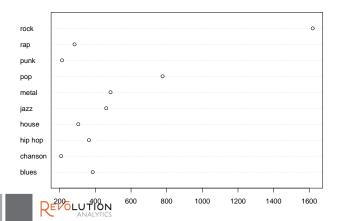






Additional Basic plots.

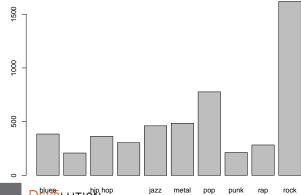
dotchart(xtabs(~songTermHot\$term))





Additional Basic plots.

barplot(xtabs(~songTermHot\$term))



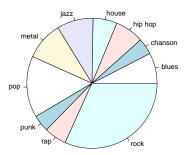






Additional Basic plots.

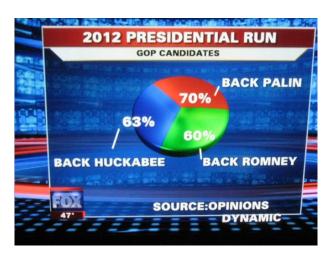
pie(xtabs(~songTermHot\$term))







Pie Charts are Bad



http://www.graphgraph.com/2011/12/pie-charts-are-terrible/



Factor Variables

Why did we load in a separate data set to generate plots for the top 10 most popular terms in our data set?

Running boxplot() on just a subset of our data pertaining to the top 10 most popular terms:

```
topTerms <- sort(xtabs(~term, songSubset3), decreasing = TRUE)
topTerms <- names(topTerms[2:11])

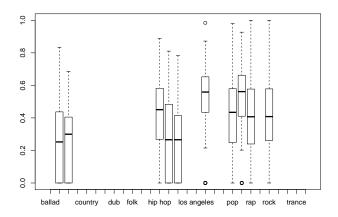
boxplot(song.hotttnesss ~ term, data = songSubset3[songSubset3$term %in ])</pre>
```



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Extras





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Extras

We have correctly subset our data to include only observations corresponding to the top 10 terms. However, term is a factor variables whose levels were defined upon import (read.csv()) to include all unique values of term, not just the top 10. Even though out data subset does not include any other observations but those corresponding to the top 10 terms, place holders for all terms exist.

```
str(songSubset3$term[songSubset3$term %in% topTerms])
```

```
## Factor w/ 29 levels "ballad", "blues",...: 25 25 22 22 22 22 2 2 2 2 5 ...
```





We can redefine term as a factor variable with levels corresponding to only those unique values in our subset.

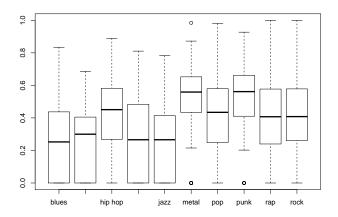
```
songTermHot <- songSubset3[songSubset3$term %in% topTerms, ]
songTermHot$term <- factor(songTermHot$term) # this is key
boxplot(song.hotttnesss ~ term, data = songTermHot)
# save(songTermHot, file='songTermHot.RData')</pre>
```



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Extras







Facts about Factor Variables: * for categorical data (numeric or string) * can be more memory efficient than strings * explicit levels/labels mean better graphs & tables * modeling of categorical variables in, say, Im()

... more information: http:
//www.ats.ucla.edu/stat/r/modules/factor_variables.htm



Custom Functions

One way to extend the functionality of R is to install and load additional packages and their containing functions. Another is to write your own functions.

Given a vector of values, we can normalize the values (ensure they have a mean of 0 and sd of 1) by, for each value, subtracting the mean and dividing the result by the sd.

```
x <- songData$song.hotttnesss
x <- (x - mean(x, na.rm = TRUE))/sd(x, na.rm = TRUE)
head(x)</pre>
```

```
## [1] NA NA NA -1.387 NA NA
```







Instead of recalculating x using these steps for each variable we'd like to normalize, we can define a custom function and apply it to a given variable.

```
normfunct <- function(x = songData$song.hotttnesss) {
    (x - mean(x, na.rm = TRUE))/sd(x, na.rm = TRUE)
}

x <- normfunct()
y <- normfunct(songData$duration)</pre>
```

Exercise

Use songData to create the perfect running playlist for efficient stride of 90 steps/minute/leg (180 or 90 bpm).

(http://gizmodo.com/5906815/the-most-mathematically-perfect-playlist-for-running/all)[http://gizmodo.com/5906815/the-most-mathematically-perfect-playlist-for-running/all]

tempo: tempo in BPM according to The Echo Nest

```
head(songData$tempo, 20)
```

```
## [1] 127.51 171.99 149.03 130.16 193.36 149.79 45.43 135.93 136.82 112.6 ## [11] 107.45 189.82 158.02 115.09 208.17 114.93 145.88 135.83 109.10 179.
```



Exercise

- Create a new variable tempDec which rounds tempo to the nearest 10 bpm. Hint: use round()
- Create a Subset of songData to identify songs where this rounded bpm/"tempo" is 180 or 90.
- Identify the top 10 terms by observation count. Pick three terms for your playlist.
- Subset the dataset again to only include the terms you chose and songs. We will randomly choose 10 songs from this term from our playlist.





Solution

```
songData$tempoDec <- round(songData$tempo, -1)</pre>
songTemp <- songData[songData$tempoDec == 180 | songData$tempoDec == 90</pre>
songTable <- xtabs(~songData$term)</pre>
head(songTable[order(songTable, decreasing = TRUE)], 11)
## songData$term
##
    other rock pop metal jazz blues hip hop house
                                                             rap
     3423 1619
                   777
                          485
                                 461
                                        385
                                               364
                                                      304
##
                                                             283
##
    punk chanson
```



213

208

##

Solution

```
# hip hop, rock and house
songTemp <- songTemp[songTemp$term == "rock" | songTemp$term == "house"</pre>
    "hip hop", ]
songTemp[sample(nrow(songTemp), 10), c("artist.name", "title", "release
##
            artist.name
                                      title
## 9719 Les Sexareenos
                                    Ruby D.
## 2331 Jedi Mind Tricks Brute Force II
## 7703 Rachel Portman
                                    Bridget
## 4344
             Sly Dunbar Casava Piece Riddim
                            Final Flash
## 1815
                Hatiras
```



R comes with multiple functions built for aggregating data by applying a function (such as mean) over observations subset by one or multiple variables. Unlike other programming languages, this functionality in R doesn't require you to specify the dimensions or element names of your resulting object.

We use the aggregate function here to return all columns averaged across artist names.





Our goal, benchmark

```
subset(songDedup, subset = artist.name == "Aerosmith")
     artist.name artist.latitude artist.longitude artist.location
##
                                                      Boston, MA
## 389
        Aerosmith
                              NΑ
                                              NΑ
     artist.hotttnesss artist.familiarity tempo duration song.hotttnesss
## 389
              0.6107
                              0.8725 108.7 425
                                                          0.5304
       term hfRatio
##
## 389 rock 0.6999
```





```
artistAgg <- aggregate(. ~ artist.name, data = songSubset3, mean)
subset(artistAgg, subset = artist.name == "Aerosmith")

## [1] artist.name artist.latitude artist.longitude
## [4] artist.location artist.hotttnesss artist.familiarity
## [7] tempo duration song.hotttnesss
## [10] term hfRatio song.hotttnesssCut
## <0 rows> (or 0-length row.names)
```

What happened to Aerosmith?



Problem: mean() defaults to NA if *any* observations meaned are NA and aggregate(), by default, removes any rows with NA/NaN. Aerosmith disappears.

Solution: na.rm=TRUE is an argument for mean(), it allows us to calculate the mean of each var, excluding any NA values

Note: mean of variables where all values are missing will still default to NaN. aggregate() argument na.action="na.pass" leaves in NA containing rows



```
artistAgg2 <- aggregate(. ~ artist.name, data = songSubset3, mean, na.a
    na.rm = TRUE)
subset(artistAgg2, subset = artist.name == "Aerosmith")
    artist.name artist.latitude artist.longitude artist.location
       Aerosmith
                                                           109
## 76
                            NaN
                                            NaN
    artist.hotttnesss artist.familiarity tempo duration song.hotttnesss
             0.6108
                            0.8725 121.1 312.3
## 76
                                                     0.5304
     term hfRatio song.hotttnesssCut
## 76 25 0.7
                                 2.5
```







Like xtabs(): we can use aggregate() to obtain contingency tables

The number of observations at each value of term. songSubset3 contains 47 songs labeled ballad.

```
aggregate(artist.name ~ term, data = songSubset3, length)
```

We can use any variable in our dataset in place of artist.name here. The key is that aggregate returns the length of artist.name (the count of the number of observations of artist.name) at each value of term.

##		term	artist.name
##	1	ballad	47
##	2	blues	385
##	3	chanson	208
##	4	chill-out	90
##	5	country	75
##	6	BEKEFAILD	97



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Extras

Apply functions

There exist many other functions like aggregate. Aggregate works on a variety of R objects and returns a data frame. Other functions are object specific in both their input and output.

- apply()
- lapply()
- sapply()
- tapply()
- mapply()
- by()





The plyr package provides similar functionality.

Resource for

apply functions & plyr: http://stackoverflow.com/questions/3505701/

r-grouping-functions-sapply-vs-lapply-vs-apply-vs-tapply-vs-by-vs 7141669#7141669



Using the function apply() to apply the custom normalizing function we wrote earlier to each column of a songData subset:

```
normfunct <- function(x = songData$song.hotttnesss) {
     (x - mean(x, na.rm = TRUE))/sd(x, na.rm = TRUE)
}

colNameSub <- c("artist.hotttnesss", "artist.familiarity", "tempo", "dur"
     "loudness", "song.hotttnesss")

x <- apply(songData[, colNameSub], 2, normfunct)</pre>
```



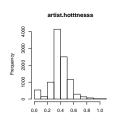
We can create a histogram plot for each normalized variable by using the function sapply() to apply the hist() function to each column of x.

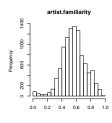
```
par(mfrow = c(2, 3))
sapply(names(songData[, colNameSub]), function(x) hist(songData[, x], maximum xlab = ""))
```

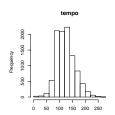
(()) I

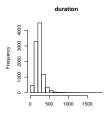


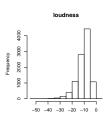
Extras

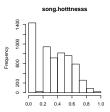












artist.hotttnesss artist.familiarity tempo

Bracketing Review

Poll: Which of the following can we use for subsetting or extracting elements of a data frame? Select all that apply.

- Ψ
- **(**)
- [[]]
- {}
- .
- subset()

Bracketing Review

Poll: Which of the following can we use to aid in specifying function arguments? Select all that apply.

- **\$**
- **(**)
- **[**[]]
- {}
- .
- subset()



(())



Merging Data

The artist location data is neither standardized nor complete. We do have latitude and longitude coordinates! The dataset revGeocodeDF.RData used ggmap2 package's revgeocode() function to obtain state information for these coordinates.

Recall we loaded this RData file which created an object called 'd' in our workspace.

load("data/revGeocodeDF.RData")

Exercise



Explore this data set. What information is common between this data set and songData? What additional information does this data set provide? How can we augment g with this information?

(())



Solution

```
summary(d)
summary(songData)
head(d)
head(songData)
```

View(d)

View(songData)

Exercise



- Use subset() on *songSubset3* to identify the observation where (artist.latitude, artist.longitude) is (34.23294, -102.41020). What is the artist.location?
- Use subset() on d to identify the observation where (artist.latitude, artist.longitude) is (34.23294, -102.41020). What is the geoCountry and geoSate?



Solution

##

subset(songSubset3, artist.latitude == 34.23294 & artist.longitude == -

```
##
    artist.name artist.latitude artist.longitude artist.location
## 35
            BT
                       34.23
                                    -102.4
                                                  Earth
## 36
           BT
                      34.23
                                    -102.4
                                                 Earth
## 37
          BT
                     34.23
                                  -102.4
                                                 Earth
## 38
          BT
                     34.23
                                  -102.4
                                               Earth
## 39 BT feat. JES
                     34.23
                                  -102.4
                                                 Earth
. . .
```

subset(d, artist.latitude == 34.23294 & artist.longitude == -102.4102)

artist.longitude artist.latitude geoCountry geoState ## <u>1</u>343 -102.434.23 United States Texas





Merging Data

We will merge the two based on their shared variables: artist.latitude and artist.longitude to obtain more complete and standardized location information than artist.location provides.

```
songSubset3 <- merge(d, songSubset3, by = c("artist.latitude", "artist.")</pre>
subset(songSubset3, artist.latitude == 34.23294 & artist.longitude == -
##
     artist.latitude artist.longitude geoCountry geoState artist.name
                         -102.4 United States Texas
## 1116
             34.23
                                                          BT
## 1117
             34.23
                        -102.4 United States Texas
                                                          BT
            34.23
                                                          BT
## 1118
                       -102.4 United States Texas
             34.23
## 1119
                        -102.4 United States Texas BT feat, JES
## 1120
             34.23
                        -102.4 United States Texas
                                                          BT
```



Save Your Work



```
save(songSubset3, file = "data/Module4_songSubset3.RData")
```





Thank you

Revolution Analytics is the leading commercial provider of software and support for the popular open source R statistics language.

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