# Stat 5810, Section 003 Statistical Visualization II Spring 2019 Homework 1

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Stat 5810/6910 Statistical Visualization II

Spring 2019

Homework Assignment 1 (3/4/2019)

90 Points — Due Wednesday 3/20/2019 (via Canvas by 11:59pm)

(i) Visualization of Multivariate Data (60 Points):

In this question, you have to work with the *crabs* data set from the *MASS* package. See the *crabs* help page for further details.

(a) (3 Points) Load all required R packages to answer this question. Show your R code. Do not just blindly trust the information on the help page! How many observations and how many variables are included in this data set overall?

Use something like the following to incorporate results from your R code directly into your LaTeX text: "Apparently, there are 50 observations and 2 variables in the cars data set."

#### Answer:

- > setwd("C:/Users/Shaun/Desktop/StatVis/StatVis2/HW1")
- > library(ggplot2)
- > library(MASS)
- > library(lattice)
- > library(GGally)
- > library(dplyr)
- > library(PairViz)
- > library(RColorBrewer)
- > library(gplots)
- > data(crabs)

There are 200 observations and 8 variables in the crabs data set.

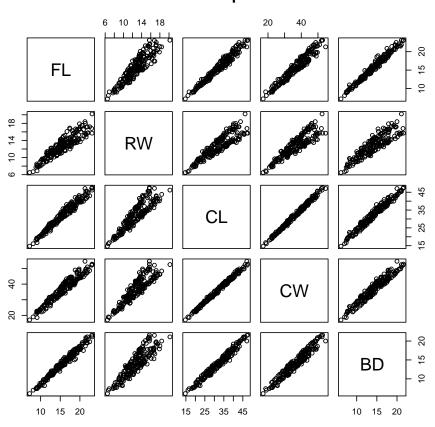
(b) (4 Points) Create a default scatterplot matrix of the five quantitative variables (omitting sp,  $sex \ & index$ ) via baseR. Do not optimize this scatterplot matrix and do not use any colors or symbols. Describe this scatterplot matrix. Which questions naturally come to your mind? What would you initially anticipate as the answers to your questions? Do not modify your answer here, even if it turns out later on that your anticipation was not correct! Hint: Think of which variables have been ignored in your scatterplot

matrix. Include your figure and your R code.

#### Answer:

- > pairs(crabs[,4:8],
- + main="Default Scatterplot Matrix")

### **Default Scatterplot Matrix**



There appears to be a strong linear correlation between the variables, so we might assume that this correlation might also hold between the variables sp and sex.

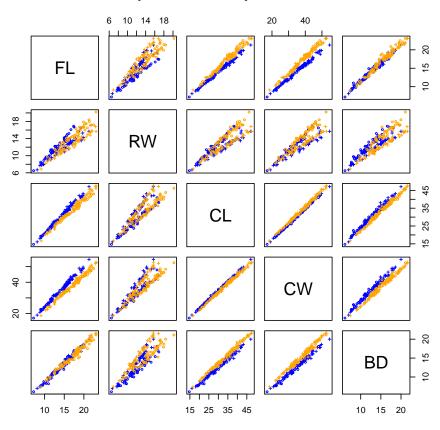
(c) (5 Points) Redo your scatterplot matrix from part (b) of the five quantitative variables (omitting sp, sex & index) via baseR. Now use different colors and symbols. Recall that "B" represents blue crabs and "O" represents orange crabs, so use these two colors. Use "o" to represent male crabs and "+" to represent female crabs. This is a perfect opportunity to make use of "ifelse" expressions in your R code! Also reduce the symbol size (cex) to 0.5. Describe this scatterplot matrix. Does this answer your previous questions from part (b)? Was your anticipation correct? Include your figure and your

# R code.

#### Answer:

```
> # data(state)
> # tristate <- as.data.frame(state.x77[, c(3, 6, 5)])
> # illiteracy <- tristate[, 1]</pre>
> # grad <- tristate[, 2]</pre>
> # pairs(tristate, col = unclass(state.region))
> cols <- character(nrow(crabs))</pre>
> #cols[] <- "black"
> cols[crabs$sp == "B"] <- "blue"
> cols[crabs$sp == "0"] <- "orange"
> shapes <- character(nrow(crabs))</pre>
> shapes[] <- 0
> shapes[crabs$sex == "F"] <- '1'
> shapes[crabs$sex == "M"] <- '3'
> pairs(crabs[,4:8],
       main="Improved Scatterplot Matrix",
       col = unclass(cols), pch = c(1,3)[as.numeric(crabs$sex)], cex = 0.5)
```

# **Improved Scatterplot Matrix**



It would seem that my anticipation was correct, and that there is some correlation between gender and

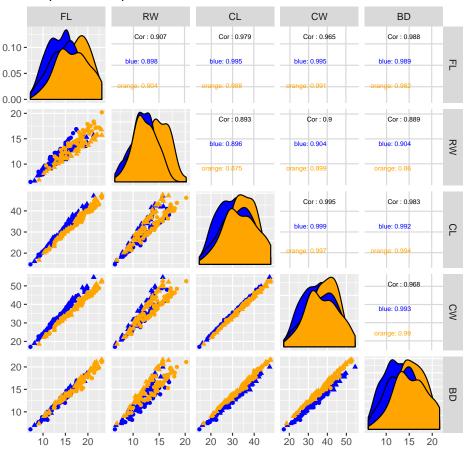
color as well.

(d) (4 Points) Redo your scatterplot matrix from part (c) of the five quantitative variables (omitting sp, sex & index) via ggpairs. Use the same colors and symbols as in part (c). For everything else, use the default settings and show the four densities on the diagonal and the correlations in the upper triangular matrix. Include your figure and your R code.

#### Answer:

```
> ggplot <- function(...) ggplot2::ggplot(...) +
+ scale_color_manual(values = c("blue","orange")) +
+ scale_fill_manual(values = c("blue","orange"))
> unlockBinding("ggplot",parent.env(asNamespace("GGally")))
> assign("ggplot",ggplot,parent.env(asNamespace("GGally")))
> ggpairs(crabs, columns = 4:8,
+ title = "GGplot Scatterplot Matrix",
+ aes(color = cols, shape = shapes),
+ upper = list(continuous = wrap("cor", size = 2.25)))
```

# **GGplot Scatterplot Matrix**



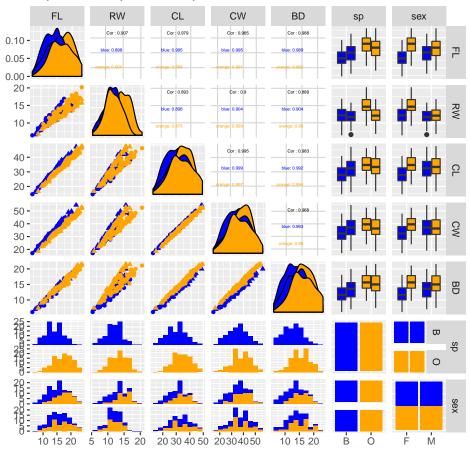
(e) (5 Points) Redo your scatterplot matrix from part (d) of the five quantitative

variables including  $sp \, \mathcal{E} \, sex$  (but still omitting index) via ggpairs. Place  $sp \, \mathcal{E} \, sex$  after the five quantitative variables so that the histograms and box plots appear on the bottom and on the right. Use the same colors and symbols as in part (d). Choose 10 bins for all histograms. For everything else, use the default settings and show the four densities on the diagonal and the correlations in the upper triangular matrix. Include your figure and your R code.

#### Answer:

```
> ggpairs(crabs, columns = c(4:8,1:2),
+ title = "Expanded GGplot Scatterplot Matrix",
+ aes(color = cols, shape = shapes),
+ lower = list(combo = wrap("facethist", bins = 10)),
+ upper = list(continuous = wrap("cor", size = 1.5)))
```

#### **Expanded GGplot Scatterplot Matrix**



(f) (6 Points) Frankly, I have a problem with the interpretation of the (extended) scatterplot matrix from part (e). Can you easily identify which boxplot and histogram belongs to which species or which sex? Therefore, do the following: Add an "interaction variable" of *sp.sex* to the crabs data set. Recall

how we did this for "SexClassSurvived" for the Titanic data set. **Reorder** your factor levels as B.M, B.F, O.M, and O.F. Do not leave the factor levels in the order that was produced by R. We have done such reorderings of factor levels in *Statistical Visualization I*.

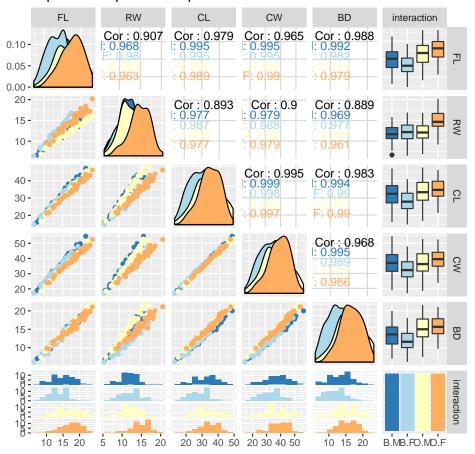
Redo your scatterplot matrix from part (e), but now with sp.sex as the only categorical variable. Use a diverging 4–class RdYlBu color scheme from the RColorBrewer R package where "blue" represents blue crabs and "red" represents orange crabs. Use the two darker (outside) colors for the male crabs and the two fainter (inside) colors for the female crabs. Warning: Make sure that you correctly map these colors to the species and sex. If you are not certain, compare with your scatterplot matrix from baseR.

As before, choose 10 bins for all histograms. For everything else, use the default settings and show the four densities on the diagonal and the correlations in the upper triangular matrix. Reduce alpha to 0.5 address the overplotting problem. Likely, your font size will be too large for the values of the correlations. Also check and adjust the font sizes on the axes if necessary. Recall that this has to look good in your final pdf version and not in the Plots window in RStudio. If necessary, google for a solution how to adjust the font sizes in *qapairs*. Include your figure and your R code.

#### Answer:

```
> ggplot <- function(...) ggplot2::ggplot(...) +
+ scale_color_manual(values = rev(brewer.pal(5,"RdYlBu"))) +
+ scale_fill_manual(values = rev(brewer.pal(5,"RdYlBu")))
> unlockBinding("ggplot",parent.env(asNamespace("GGally")))
> assign("ggplot",ggplot,parent.env(asNamespace("GGally")))
> crabs.ordered <- crabs
> crabs.ordered <- crabs
> crabs.ordered$interaction <- with(crabs, interaction(sp, sex))
> crabs.ordered$interaction <- factor(crabs.ordered$interaction,
+ levels = c("B.M", "B.F", "O.M", "O.F"))
> ggpairs(crabs.ordered, columns = 4:9,
+ title = "Improved GGplot Scatterplot Matrix",
+ aes(color = crabs.ordered$interaction),
+ lower = list(combo = wrap("facethist", bins = 10)))
```

# Improved GGplot Scatterplot Matrix

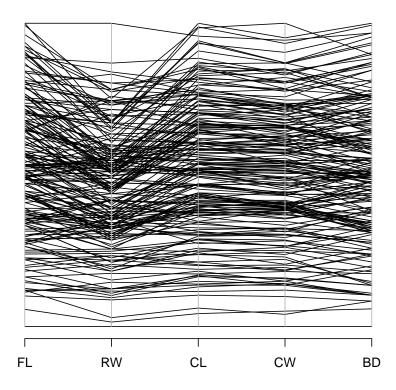


(g) (4 Points) Create a default parallel coordinates plot (PCP) of the five quantitative variables (omitting sp, sex & index) via baseR. Do not optimize this PCP and do not use any colors. Describe this PCP. Are there any variables that allow to separate one of the species or sexes? Include your figure and your R code.

#### Answer:

> parcoord(crabs.ordered[,4:8], main = "Parallel Coordinate Plot")

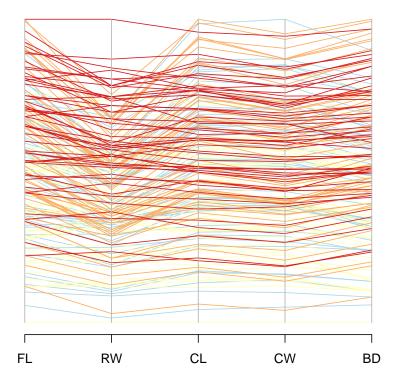
#### **Parallel Coordinate Plot**



It would appear that there is at least two patterned groups that would be a valid show of gender's influence on the other factors. The one that sticks out most prominently is the variable RW.

#### Answer:

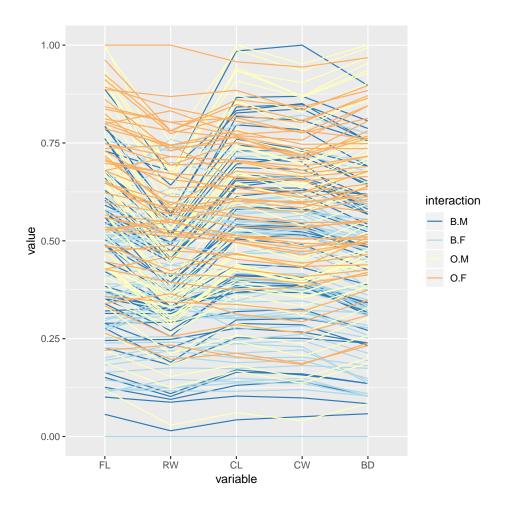
> parcoord(crabs.ordered[,4:8], col = rev(brewer.pal(5,"RdYlBu")[unclass(crabs.ordered\$interaction)]))



(i) (5 Points) Redo your PCP from part (h) of the five quantitative variables (omitting sp, sex & index) via the ggparcoord function from the GGally R package. Use the same colors as in part (f). Use a 0–1–scale for all parallel axes. Hint: The legend of this graph should reveal whether you correctly use the colors as specified in part (f). If something goes wrong here, likely you use the colors incorrectly in some of your previous parts as well. Include your figure and your R code.

#### Answer:

<sup>&</sup>gt; ggparcoord(crabs.ordered, columns = 4:8, groupColumn = 9, scale = "uniminmax")



(j) (6 Points) Let's try something: First install the *BiocManager* R package from CRAN. Then install the *graph* R package from Bioconductor as follows:

BiocManager::install("graph", version = "3.8")

Finally install the *PairViz* R package from CRAN. Read the PCP vignette at https://cran.r-project.org/web/packages/PairViz/vignettes/pcp.html. Adapt the example code to create a "Weighted Eulerian with Correlation Guide," placing the bars for the correlations below the PCP. Do not colorcode the lines in the PCP. Describe and interpret this version of the PCP

code the lines in the PCP. Describe and interpret this version of the PCP (recall the made—up examples from class). It will also be helpful to revisit the correlations in the scatterplot matrix in part (f). Include your figure and your R code.

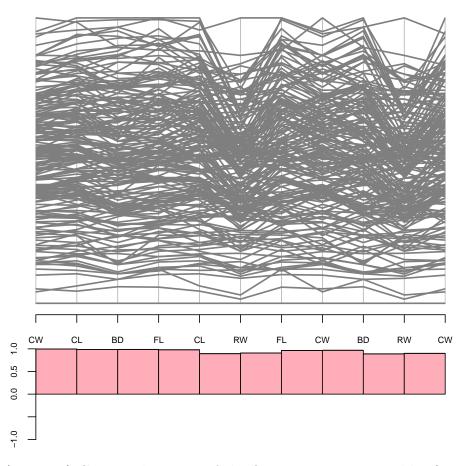
#### Answer:

> data <- crabs[4:8]

> corw <- as.dist(cor(data))</pre>

```
> o <- eulerian(-corw)
> par(cex.axis=.7)
> corw <- dist2edge(corw)
> edgew <- cbind(corw*(corw>0), corw*(corw<0))
> par(cex.axis=.7)
> guided_pcp(data,edgew, path=o,pcp.col="grey50",lwd=2,
+ main="Weighted eulerian with correlation guide",
+ bar.col = c("lightpink1","lightcyan1"),
+ bar.ylim=c(-1,1),bar.axes=TRUE)
```

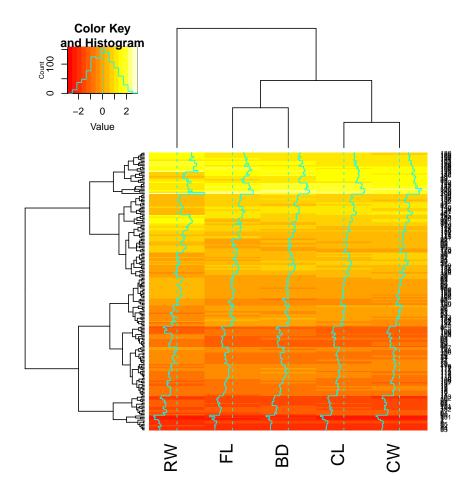
# Weighted eulerian with correlation guide



(k) (5 Points) Create a heatmap of the five quantitative variables (omitting sp,  $sex \ &endowned index$ ) via the heatmap.2 function from the gplots R package. Make sure to scale your variables! What can we learn from this heatmap (if anything at all)? Include your figure and your R code.

#### Answer:

```
> crabs48 <- as.matrix(crabs[4:8])
> heatmap.2(scale(crabs48))
```



It appears as though there does exist strong correlation between the factor levels moving from smallest value to highest value.

(l) (8 Points) Start with a basic description of the data set and its variables. Then provide a detailed interpretation of your graphical results. Point out differences and similarities for the two species and the two sexes. Also comment on similarities of the variables. Do this for the univariate graphical summaries of the five quantitative variables, the pairwise scatterplots of these variables, the PCPs, and the heatmap. Compare and combine results obtained from the various graphs. Ideally, different graphs should result in similar interpretations, but, if necessary, point out when different graphs result in contradictory interpretations. Be specific and refer back to specific question parts and do not just say "the PCP or the scatterplot matrix shows shows . . . ".

This summary should be at least 1 page in length.

Answer: I believe that each of these plots convey the fundamental correlation between each of the factor levels, including *species* and *sex*. I believe that the Matrix plots from parts (b)-(f) hammer home this point, with particular emphasis on part (f) which included information on distributions, correlation, and box plots for

#### (ii) From Tables to Graphs (30 Points):

You have to find a suitable table and translate this table into a graph. This must be a "multivariate" (non-trivial) table with several variables, possibly with different estimates, confidence intervals, etc. A table that could just be translated into several columns of dotplots is not acceptable.

(a) (5 Points) Present your proposed table to me after class, during office hours, or via e-mail by **Sunday 3/10/2019**, **11:59pm** to get my approval before you start translating it into a graph.

The table should have been published in the last four years (2016 to 2019). It can originate from a journal or conference proceedings article, textbook, or from the Web. If you can find a suitable table related to your MS or PhD research (or from your main adviser), that would be perfect. Include the reference where you located the table.

Table is included below. Primary report was generated by Lieutenant Mikelshan Bartschi on March 10, 2019 citing the number of 'Calls to Service' performed by the Cache County Sheriff's Office in the previous year by weekday and hour.

(b) (15 Points) You should review the section titled *From Tables to Plots* in our lecture notes before you start with your conversion. Also, take a closer look at the references cited in that section and under *Further Reading*. Examples 1 and 2 from that section might provide you with useful R commands. The barley example from *Statistical Visualization I* may also provide you with some useful design suggestions and R commands.

Moreover, take a look at the examples in the section titled *Linked Micromap Plots*. Even if your table is not related to any geographic data, the other components from linked micromap plots may still be useful for your conversion. For example, see how we have displayed confidence bounds (or minima and maxima), population means (or medians), annual changes, and even time series in micromaps. The same principles may be suitable for your conversion.

In addition to the graph types mentioned above, some tables may be translated into parallel coordinate plots, time series plots, or graphs that are based on the small multiples principle. Ultimately, it is your decision which design to choose for your graph.

One suggestion: Before you code your graph in R, manually design the layout of your graph on paper. Recall from  $Statistical\ Visualization\ I$  that even a tiny data set allows us to draw many different graphs. It is likely that you need to consider several possible designs for your graph first before you come up with a design that best highlights the information from the table.

Now go ahead and translate your table into a graph! Include the original table, the final version of your graph, and the R code that was used to create your graph as an answer to this question.

# Original Table:

03/10/19 19:37			Cache County Sheriff's Office Law Incident Analysis, Time Reported					1033
Hour	Monday	Tuesday	Wednesday	Thursday	Friday	Saturday	Sunday	Total
00:00-00:59		154	148	159	182			
01:00-01:59		99	126	120				
02:00-02:59		76	61	86	100			
03:00-03:59		53	51	54	67			
04:00-04:59		55	74	59	55		59	
05:00-05:59		71	75	65	91		56	
06:00-06:59		115	135	141	105	84		
07:00-07:59		206	199	203	236			
08:00-08:59		367	393	364	382			
09:00-09:59		498	501	480		314	185	
10:00-10:59		568	579	594	599			
11:00-11:59		556	510	567	525		238	
12:00-12:59		568	539	540		401		
13:00-13:59		648	547	544	614			
14:00-14:59		560	510	581	598		279	
15:00-15:59		574	641	660	579		263	
16:00-16:59		601	588	625		348	289	
17:00-17:59		484	471	532		397		
18:00-18:59		437	406	410	478			2871
19:00-19:59		350	378	330	390			
20:00-20:59		341	321	368	383		316	
21:00-21:59		273	323	316	416			
22:00-22:59		233	277	282		390		
23:00-23:59		213	226	200	294	351	173	1629
Total by Day	8066	8100	8079	8280	8680	6841	5094	53140

```
Report Includes:
All dates greater than `00:00:00 03/10/18`
All agencies
All offenses observed
All offenses reported
All offense codes
All nature of incidents
All location codes
```

\*\*\* End of Report /tmp/rptJrAK5e-rplwia.r1 3 \*\*\*

#### Code and Final graphical Visualization:

```
> ggplot <- function(...) ggplot2::ggplot(...)</pre>
```

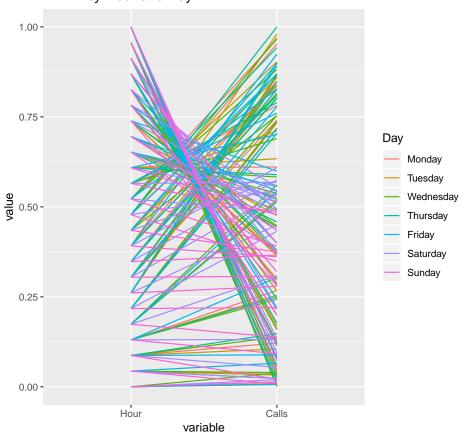
```
> unlockBinding("ggplot", parent.env(asNamespace("GGally")))
> assign("ggplot", ggplot, parent.env(asNamespace("GGally")))
> temp <- read.table("allCalls.csv", header=TRUE,
                           sep=",", stringsAsFactors = FALSE)
> all.Calls <- data.frame(Hour = rep(temp$Hour,7),</pre>
                     Day = c(rep("Monday", 24),
                              rep("Tuesday",24),
                              rep("Wednesday", 24),
                              rep("Thursday",24),
                              rep("Friday", 24),
                              rep("Saturday",24),
                              rep("Sunday",24)
                              ),
                      Calls = c(temp$Monday, temp$Tuesday,
                               temp$Wednesday, temp$Thursday,
                               temp$Friday, temp$Saturday,
                               temp$Sunday))
> all.Calls$Day <- factor(all.Calls$Day,
                           levels = c("Monday",
                                      "Tuesday",
                                      "Wednesday",
                                      "Thursday",
                                      "Friday",
                                      "Saturday",
                                      "Sunday"))
> all.Calls$Hour <- factor(all.Calls$Hour,
                            levels = c("04:00-04:59","05:00-05:59",
                                        "06:00-06:59","07:00-07:59",
                                        "08:00-08:59", "09:00-09:59",
                                        "10:00-10:59", "11:00-11:59",
                                        "12:00-12:59", "13:00-13:59",
                                        "14:00-14:59", "15:00-15:59",
                                        "16:00-16:59", "17:00-17:59",
                                        "18:00-18:59", "19:00-19:59",
                                        "20:00-20:59","21:00-21:59",
                                        "22:00-22:59","23:00-23:59",
```

```
# "00:00-00:59","01:00-01:59",
# "02:00-02:59","03:00-03:59"))

# basic line plot for personal reference
# ggplot(data = all.Calls, aes(x=Hour,y=Calls,group=Day)) +
# geom_line(aes(color=Day)) +
# theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
# scale_color_brewer(palette = "Spectral")

# ggparcoord(all.Calls, columns = c(1,3), groupColumn = 2,
# scale = "uniminmax",
# title = "Annual Calls to Service Breakdown
# by Hour and Day")
```

# Annual Calls to Service Breakdown by Hour and Day



(c) (10 Points) In addition to translating the table into a graph, you have to explain and motivate your resulting graph! Why did you choose that particular graph design? What sorting did you use for the rows and columns of the

original table? How (and why) did you combine data from the table, e.g., means and standard deviations from the table into means and confidence intervals in your graph? How (and why) did you choose specific colors and glyphs in your graph, etc.?

Also emphasize what are the most important features that can be seen in your graph. Keep in mind that the general audience may have never seen such a graph type before. So, you may have to explain some of the basic design features of your graph. Motivation:

I chose this particular graph design because I believe that it quickly allows an individual to see that the number of calls is much higher in the middle of the day than in the morning or night. As is, this graph would still need to be explained to the law enforcement audience, as they would need to be told how to interpret the hours (since I modified the order to start at 4am and end at 3 am). This could be clarified by better labeling, although it could very well croud the graph with text. Ultimately, the intent of this graph is to make it quickly apparent that there are trends between the hour of the day and the calls, and upon closer inspection, also the day of the week that it occured. I would likely pair this plot with the more basic line plot that I created as well to increase informational awareness.

# General Instructions

- (i) Create a single pdf document, using R Markdown, Sweave, or knitr. When you take this course at the 6000–level, you have to use LaTeX in combination with Sweave or knitr. You only have to submit this one document.
- (ii) Include a title page that contains your name, your A-number, the number of the assignment, the submission date, and any other relevant information.
- (iii) Start your answers to each main question on a new page (continuing with the next part of a question on the same page is fine). Clearly label each question and question part.
- (iv) Show your R code for each question part!
- (v) Before you submit your homework, check that you follow all recommendations from Google's R Style Guide (see https://google.github.io/styleguide/Rguide.xml). Moreover, make sure that your R code is consistent, i.e., that you use the same type of assignments and the same type of quotes throughout your entire homework.
- (vi) Give credit to external sources, such as stackoverflow or help pages. Be specific and include the full URL where you found the help (or from which help page you got the information). Consider R code from such sources as "legacy code or third–party code" that does not have to be adjusted to Google's R Style (even though it would be nice, in particular if you only used a brief code segment).

# (vii) Not following the general instructions outlined above will result in point deductions!

- (viii) For general questions related to this homework, please use the corresponding discussion board in Canvas! I will try to reply as quickly as possible. Moreover, if one of you knows an answer, please post it. It is fine to refer to web pages and R commands, but do not provide the exact R command with all required arguments or which of the suggestions from a stackoverflow web page eventually worked for you! This will be the task for each individual student!
  - (ix) Submit your single pdf file via Canvas by the submission deadline. Late submissions will result in point deductions as outlined on the syllabus.