Stat. 651 Homework 1

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```
library(pacman)
p_load(tidyverse, macleish, nasaweather, palmerpenguins, mdsr)
```

Problem 4 (Medium):

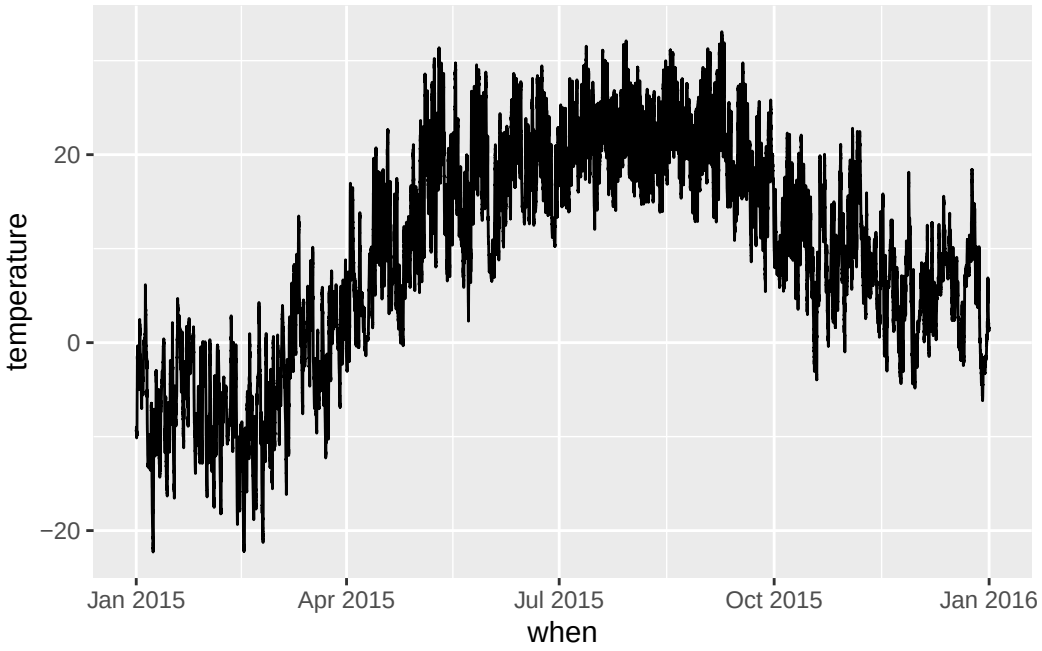
The `macleish` package contains weather data collected every 10 minutes in 2015 from two weather stations in Whately, MA.

```
head(whately_2015)
```

```
# A tibble: 6 x 8
  when                temperat~1 wind_~2 wind_~3 rel_h~4 press~5 solar~6 rainf~7
  <dttm>                   <dbl>   <dbl>   <dbl>   <dbl>   <int>   <dbl>   <dbl>
1 2015-01-01 00:00:00      -9.32    1.40    225.    54.6     985       0       0
2 2015-01-01 00:10:00      -9.46    1.51    248.    55.4     985       0       0
3 2015-01-01 00:20:00      -9.44    1.62    258.    56.2     985       0       0
4 2015-01-01 00:30:00      -9.3     1.14    244.    56.4     985       0       0
5 2015-01-01 00:40:00      -9.32    1.22    238.    56.9     984       0       0
6 2015-01-01 00:50:00      -9.34    1.09    242.    57.2     984       0       0
# ... with abbreviated variable names 1: temperature, 2: wind_speed,
#   3: wind_dir, 4: rel_humidity, 5: pressure, 6: solar_radiation, 7: rainfall
```

Using `ggplot2`, create a data graphic that displays the average temperature over each 10-minute interval (`temperature`) as a function of time (`when`).

```
ggplot(data = whately_2015, mapping = aes(x = when, y = temperature)) +
  geom_line()
```



Problem 8 (Medium):

Using data from the `nasaweather` package, use the `geom_path` function to plot the path of each tropical storm in the `storms` data table. Use color to distinguish the storms from one another, and use faceting to plot each `year` in its own panel.

```
head(storms)
```

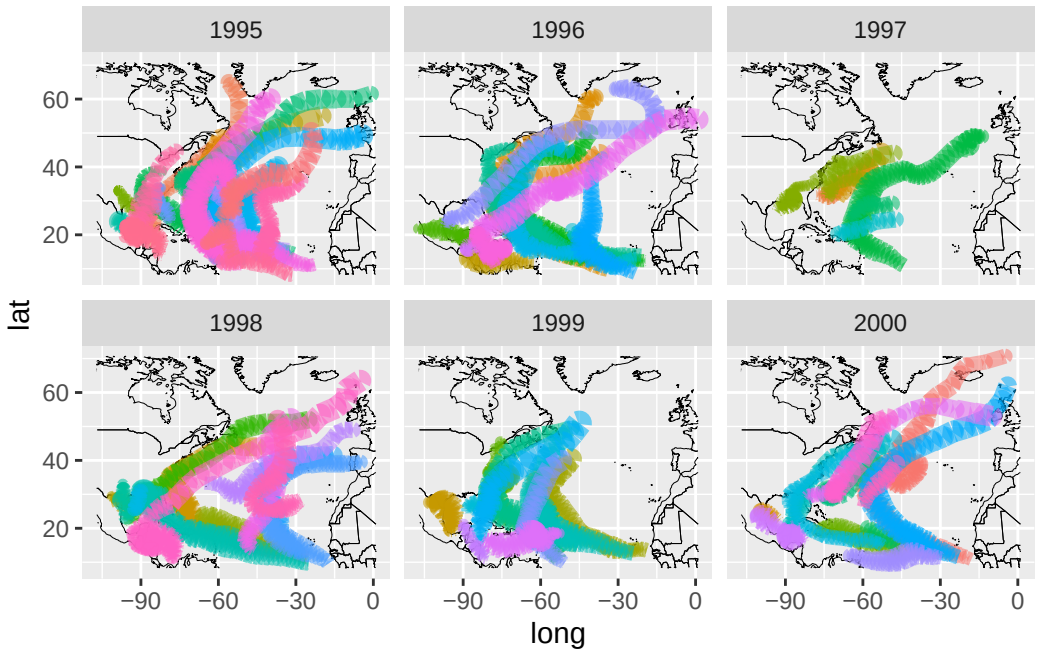
```
# A tibble: 6 x 11
  name     year month   day  hour   lat  long pressure  wind type        seasday
  <chr>   <int> <int> <int> <int> <dbl> <dbl>    <int> <int> <chr>         <int>
1 Allison  1995     6     3     0  17.4 -84.3     1005    30 Tropical D~       3
2 Allison  1995     6     3     6  18.3 -84.9     1004    30 Tropical D~       3
3 Allison  1995     6     3    12  19.3 -85.7     1003    35 Tropical S~       3
```

```
4 Allison  1995     6     3    18  20.6 -85.8     1001    40 Tropical S~       3
5 Allison  1995     6     4     0  22   -86        997    50 Tropical S~       4
6 Allison  1995     6     4     6  23.3 -86.3      995    60 Tropical S~       4
```

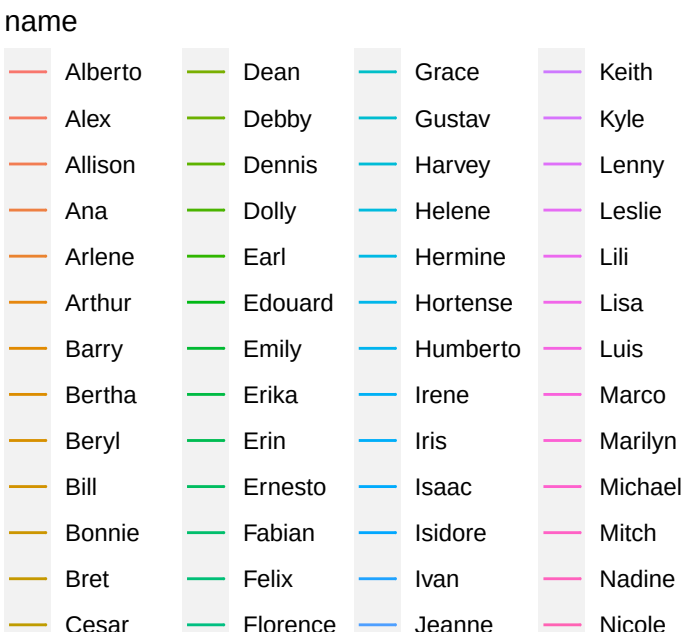
```
bbox <- storms %>%
  select(lat, long) %>%
  map_df(range)          # using the purrr R package

bbox
```

```
# A tibble: 2 x 2
    lat  long
  <dbl> <dbl>
1   8.3 -107.
2  70.7    1
```

```
base_map <- map_data("world") %>% ggplot( aes(x = long, y = lat)) +
  geom_path(aes(group = group), color = "black", size = 0.1) +
  lims(x = bbox$long, y = bbox$lat)

storms <- storms %>%
  unite("the_date", c(year, month, day), sep="-", remove="FALSE") %>%
  mutate(the_date = lubridate::ymd(the_date))

base_map <- base_map +
  geom_path(data = storms,
            aes(color = name, alpha = 0.01, size = wind, show.legend = FALSE),
            arrow = arrow(length = unit(0.005, "inches"))) +
  facet_wrap(~year)

base_map + theme(legend.position = "none")
```




```
legend<-cowplot::get_legend(base_map)
cowplot::plot_grid(legend)
```



Problem 9 (Medium):

Using the `penguins` data set from the `palmerpenguins` package:

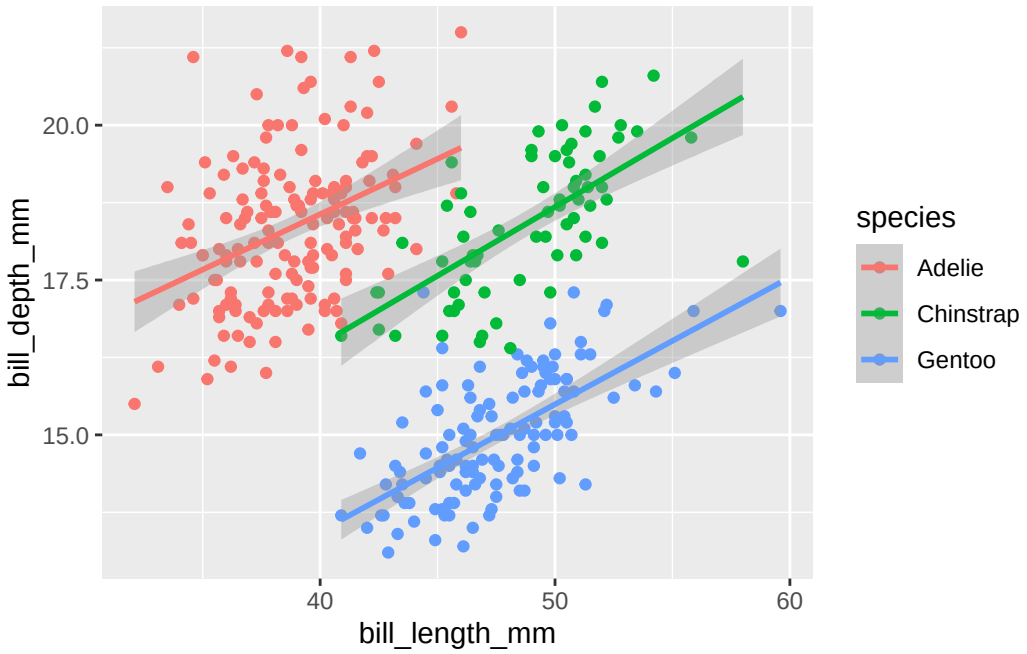
(a) Create a scatterplot of `bill_length_mm` against `bill_depth_mm` where individual species are colored and a regression line is added to each species. Add regression lines to all of your facets. What do you observe about the association of bill depth and bill length?

```
head(penguins)
```

```
# A tibble: 6 x 8
  species island    bill_length_mm bill_depth_mm flipper_l~1 body_~2 sex    year
  <fct>   <fct>              <dbl>         <dbl>       <int>   <int> <fct> <int>
1 Adelie  Torgersen           39.1          18.7         181    3750 male   2007
2 Adelie  Torgersen           39.5          17.4         186    3800 fema~  2007
3 Adelie  Torgersen           40.3          18           195    3250 fema~  2007
4 Adelie  Torgersen           NA            NA            NA      NA <NA>   2007
5 Adelie  Torgersen           36.7          19.3         193    3450 fema~  2007
6 Adelie  Torgersen           39.3          20.6         190    3650 male   2007
# ... with abbreviated variable names 1: flipper_length_mm, 2: body_mass_g
```

```
p1 <- penguins %>%
  ggplot(aes(x = bill_length_mm, # set aesthetics for x
             y = bill_depth_mm, # set aesthetics for y
             color = species)) + # color by species
  geom_point() + # create scatter plot
  geom_smooth(method = 'lm') # add regression line for each species


p1
```

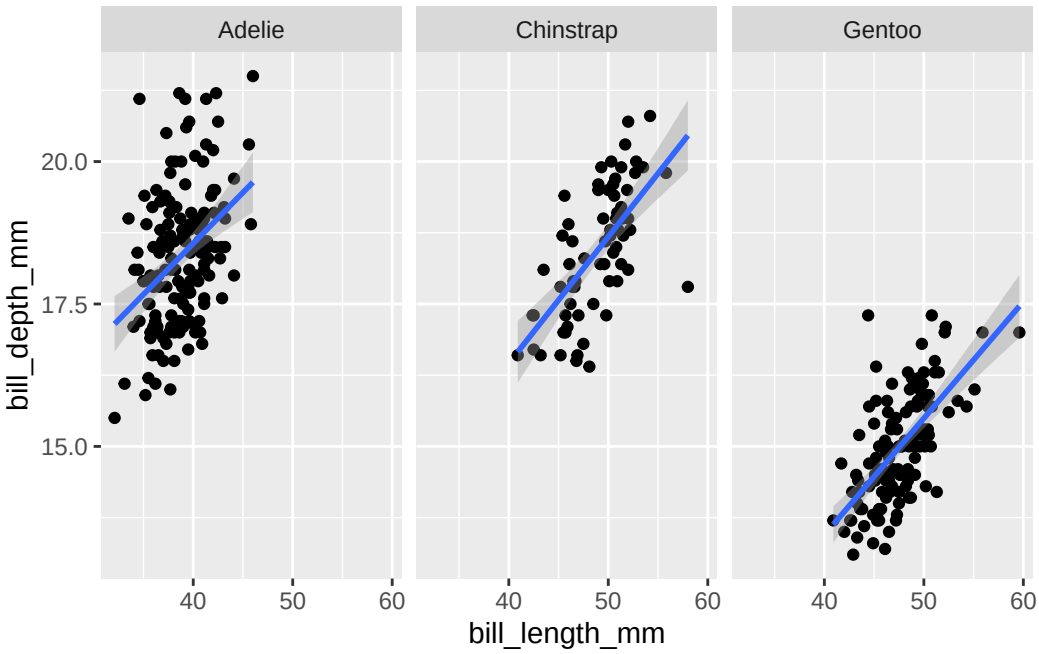
```
`geom_smooth()` using formula = 'y ~ x'
```



(b) Repeat the same scatterplot but now separate your plot into facets by `species`. How would you summarize the association between bill depth and bill length.

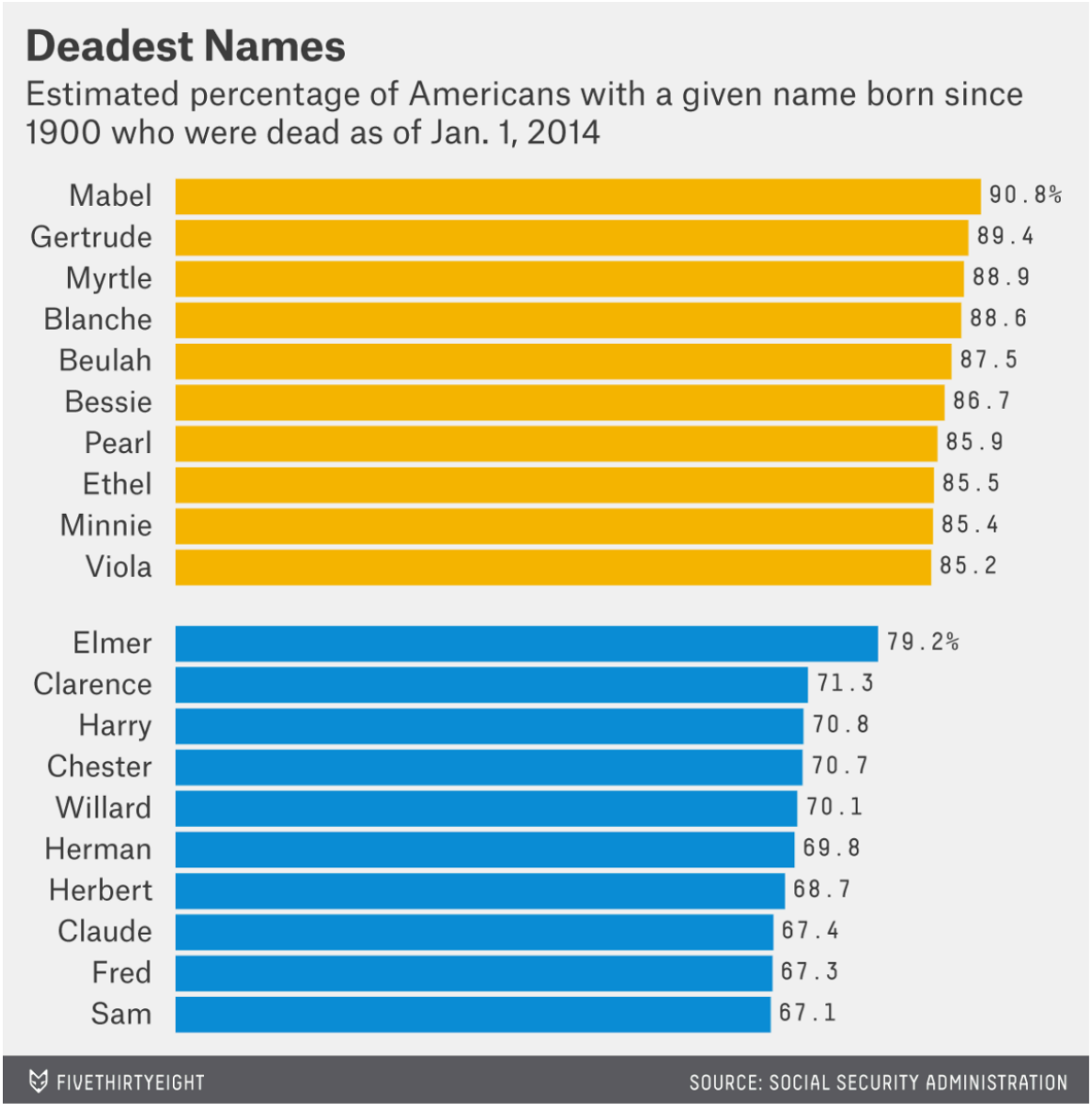
```
p2 <- penguins %>%
  ggplot(aes(x = bill_length_mm, y = bill_depth_mm)) + #set aesthetics
  geom_point() + #create scatterplot
  geom_smooth(method = 'lm') + #add regression line
  facet_wrap( ~ species) #facet by species

p2
```

```
`geom_smooth()` using formula = 'y ~ x'
```


Adelie
Chinstrap
Gentoo
bill_depth_mm
20.0
17.5
15.0
40
50
60
bill_length_mm

Problem 10 (Hard):

Use the `make_babynames_dist()` function in the `mdsr` package to recreate the "Deadest Names" graphic from FiveThirtyEight (https://fivethirtyeight.com/features/how-to-tell-someones-age-when-all-you-know-is-her-name).



```r
babynames_dist <- make_babynames_dist()
head(babynames_dist)
```

```
# A tibble: 6 x 9
   year sex   name          n   prop alive_prob count_thousands age_to~1 est_a~2
  <dbl> <chr> <chr>     <int>  <dbl>      <dbl>           <dbl>    <dbl>   <dbl>
1  1900 F     Mary      16706 0.0526          0            16.7      114       0
2  1900 F     Helen      6343 0.0200          0             6.34     114       0
3  1900 F     Anna       6114 0.0192          0             6.11     114       0
4  1900 F     Margaret   5304 0.0167          0             5.30     114       0
5  1900 F     Ruth       4765 0.0150          0             4.76     114       0
6  1900 F     Elizabeth  4096 0.0129          0             4.10     114       0
# ... with abbreviated variable names 1: age_today, 2: est_alive_today
```

```r
deadest <- babynames_dist %>%
  filter(year >= 1900) %>% #filter by years greater than or equal to 1900
  group_by(name, sex) %>% # group by name and sex
  summarise(N = n(), # count observations
            total_est_alive_today = sum(est_alive_today), #create column of total estimate
            total = sum(n)) %>%
  mutate(percent_dead = 1 - (total_est_alive_today / total)) %>% #create column of percent
  filter(total > 50000) %>% #filter out rows less than or equal to 50000
  arrange(desc(percent_dead)) %>% #arrange in descending order by percentage dead
  group_by(sex) %>% #group by sex
  top_n(10) #
```

```
`summarise()` has grouped output by 'name'. You can override using the
`.groups` argument.
Selecting by percent_dead
```

```r
head(deadest)
```

```
# A tibble: 6 x 6
# Groups:   sex [1]
  name     sex       N total_est_alive_today  total percent_dead
  <chr>    <chr> <int>                 <dbl>  <int>        <dbl>
1 Mabel    F       111                20238.  96044        0.789
2 Gertrude F       111                31365. 145703        0.785
3 Myrtle   F        99                25492. 108943        0.766
```

```
4 Blanche  F       111              16511.  69526       0.763
5 Beulah   F       111              15647.  63367       0.753
6 Opal     F       111              17471.  65823       0.735
```

```
ggplot(deadest, aes(reorder(name, percent_dead), percent_dead, fill = sex)) +
  geom_bar(stat = "identity") +
  geom_text(aes(y = percent_dead + 0.05), label = paste(round(deadest$percent_dead * 100,
  coord_flip() +
  ggtitle("Deadest Names", subtitle = "Estimated % of Americans with a given name born sin
scale_x_discrete(NULL) + scale_y_continuous(NULL) +
scale_fill_manual(values = c("#f6b900", "#008fd5"))
```