

# Homework 1

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## Part 1 - Distrubtion

**1. Give three reasons why a plant or animal might be patchily distributed like this. For each, describe whether you would expect the location of the high-density patches to be consistent from year to year (1 point).**

1. The landscape might be patched. Even at a small scales, landscapes may have distinct patches of vegetation or land cover that would cause populations to cluster in this pattern. An example might be a golf course, where there might exist small patches of unmanaged land or ponds between the managed greens. In this case, unless the land use of the area was altered, you would expect the patches to be consistent year to year. Another, more natural example is a tidepool landscape. Given the shifting nature of coastlines and intertidal communities, this may be less consistent year over year.
2. Access to light, or other limiting abiotic environmental factors may be unevenly distributed across the landscape. An example might be a forest with some very tall canopy trees that block lights in large patches, causing shorter species to congregate in specific areas. This may not be totally consistent year to year depending on the growth rate of the canopy species, as leaf cover and branches may shift as the species grows. However, given trees are not mobile some consistency could be expected.
3. Feeding behavior might also explain why certain species are distributed this way. Certain species of insects, fungi or plants may act as decomposers, attracting to areas where another animal has died. In this example, the high-density patches would not most likely not be consistent year to year.

## Part 2 - Eureka Dune grass

**2. Read in the data, and use a two-sample t-test (e.g., using `t.test()` in R) to assess whether and how mean abundance changed from 2009 to 2010. Choose a level for alpha, and justify your choice. On the basis of this analysis, what do you conclude about the change in the grass's abundance? (1 point)**

```
#explore the means
mean_2009 <- mean(swallenia$count_2009)

mean_2010 <- mean(swallenia$count_2010)

#run the t.test
t.test(swallenia$count_2009, swallenia$count_2010, conf.level = 0.95)
```

Using a confidence level of 95% ( $\alpha = 0.05$ ) it appears that there is not a significant difference between the mean abundance of swallenia between 2009 and 2010 ( $t_{10} = -0.81791$ ,  $p > 0.05$ ). This confidence level was used to minimize the risk of comitting a type 1 error, ie finding that there was a significant change in grass population when there actually was not. The greater management implications of a type 1 error in this case could be park managers interpreting a significant increase in population as a sign that current management is sufficient. This may overlook critical issues and prevent managers and scientists to do a more in depth analysis of the area and each site.

3. Because the plant was counted in the same plots each year, you can also do a paired t-test (R hint: use `paired = TRUE` in `t.test()`). On the basis of this analysis, what do you conclude about the change in the grass's abundance? (1 point)

```
#run paired t.test
t.test(swallenia$count_2009, swallenia$count_2010, paired = TRUE, conf.level = 0.95)
```

On the basis of this analysis, there is sufficient evidence that there is a difference in abundance of swallenia between the years 2009 and 2010 ( $t_{10} = -2.41$ ,  $p < 0.05$ ).

4. Which of these analyses is more appropriate? Why? (Hint: think about your answers to question 1, and whether any apply here) (1 point)

The second (paired t.test) analysis is more appropriate as it allows you to assess changes in abundance without confounds from any variance across sites. For example, there might be slightly different conditions at one cluster than another, therefore when comparing across timescales it is more informative to understand the change in growth given that the site and conditions are consistent. Furthermore, the dune grass population in a given year depends on the population at that site in the prior year, forming small sub populations at each site. It therefore makes sense to compare each subpopulation to its previous year rather than grouping the entire population in the area together.

5. Write a short paragraph to the park superintendent describing your finding about the changes (if any) to the dune grass population. (1 point)

Using a two-sided t.test across the entire population found no significant difference between Eureka Dune Grass abundance in 2009 and 2010. However, a paired t-test did find a significant difference. This suggests that there are some sites or subpopulations of dune grass that are doing better or worse than others, signalling that park management may want to do a site specific assessment to find out what ecological differences might be occurring at different sites. This will help inform the best management practices to keep all subpopulations and the overall population healthy.

### Part 3 - Grizzly Bears

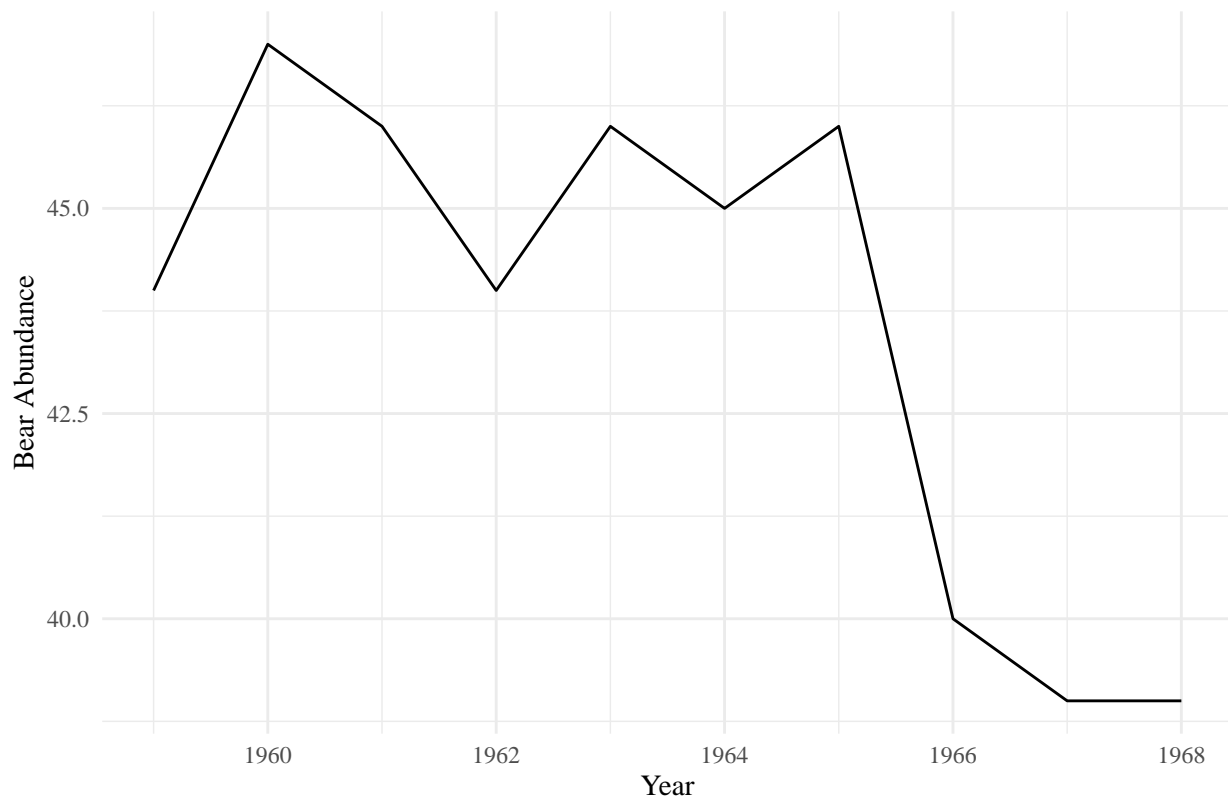
6. Open the file, and look at the data from 1959 to 1968. Did the grizzly bear population decline over this period? Support your conclusion with graphs, statistics, and logical reasoning. (2 points)

```
#graph
grizzly_early <- grizzly %>%
  filter(Year < 1969)

grizzly_plot_1 <- ggplot(grizzly_early, aes(x = Year, y = N)) +
  geom_line() +
  theme_minimal() +
  labs(x = "Year", y = "Bear Abundance", title = "Yearly Grizzly Bear population 1959 - 1968") +
  theme(text = element_text(family = "Times"))

grizzly_plot_1
```

Yearly Grizzly Bear population 1959 – 1968



```
#statistics - how has the population declined?

early_diff <- (grizzly_early$N[10]-grizzly_early$N[1])

early_pyr <- early_diff/10

#the difference is -1/2 a bear per year

early_perc <- early_diff/sum(grizzly_early$N)

#total decline of about 1%

early_model <- lm(grizzly_early$N ~ grizzly_early$Year)

summary(early_model)
```

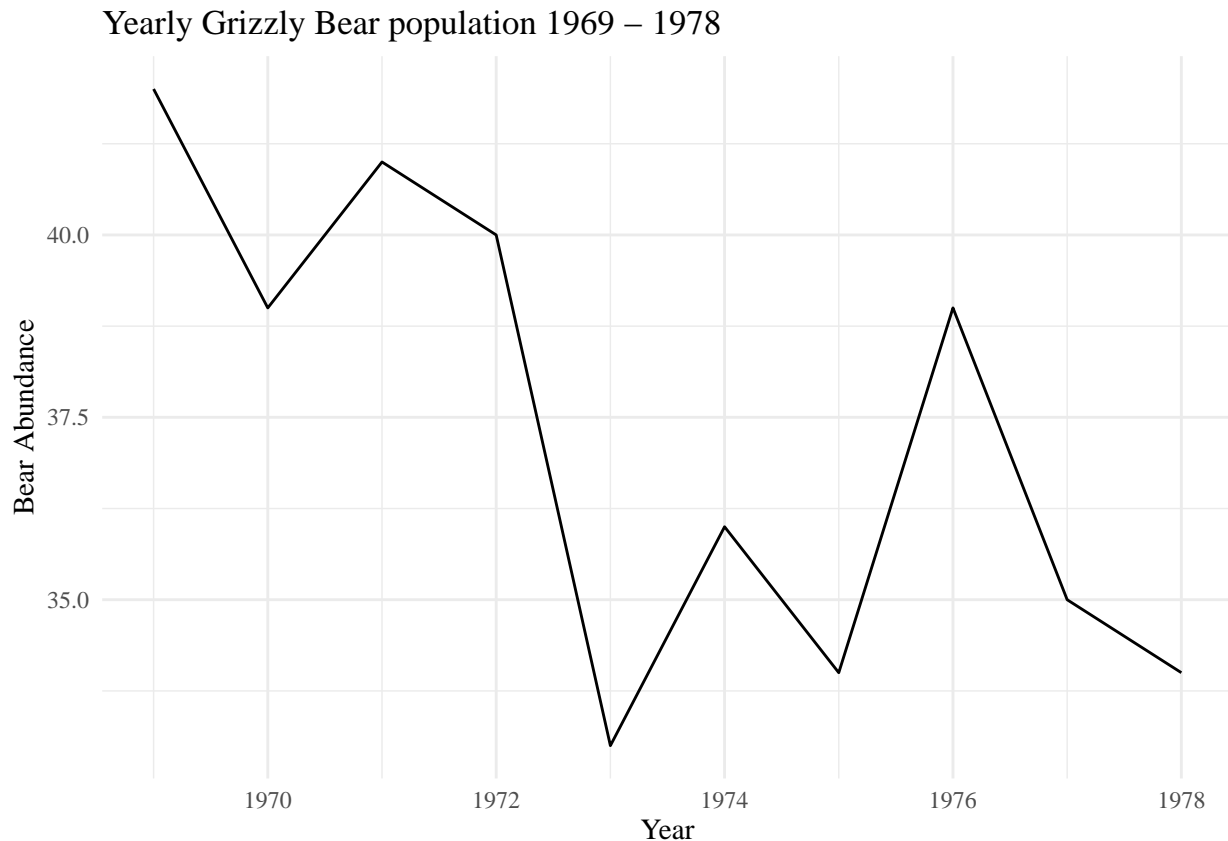
The graph suggests that the grizzly population did decline slightly from 1959 to 1968. Logically, this would make sense as a main food source (the dumps) was removed around this time, causing more human-bear conflict that can lead to bear deaths. Statistically, we find that there are 5 fewer bears reported in 1968 than in 1959 representing a total decline of about 1% during that time period.

**7. Select the data from 1969 to 1978. Did the population continue to decline? Was the decline faster than the period prior to the dump closures? (1 point)**

```
#plot
grizzly_mid <- grizzly %>%
  filter( Year > 1968 & Year < 1979)
```

```
grizzly_plot_2 <- ggplot(grizzly_mid, aes(x = Year, y = N)) +
  geom_line()+
  theme_minimal()+
  labs(x = "Year", y = "Bear Abundance", title = "Yearly Grizzly Bear population 1969 - 1978")+
  theme(text = element_text(family = "Times"))
```

```
grizzly_plot_2
```



```
#statistics - how has the population declined?

mid_diff <- (grizzly_mid$N[10]-grizzly_mid$N[1])

mid_pyr <- mid_diff/10

#the total percent decline

mid_perc <- mid_diff/sum(grizzly_early$N)

#use linear regression to compare slopes - is the slope the rate of decline?

mid_model <-lm(grizzly_mid$N ~ grizzly_mid$Year)

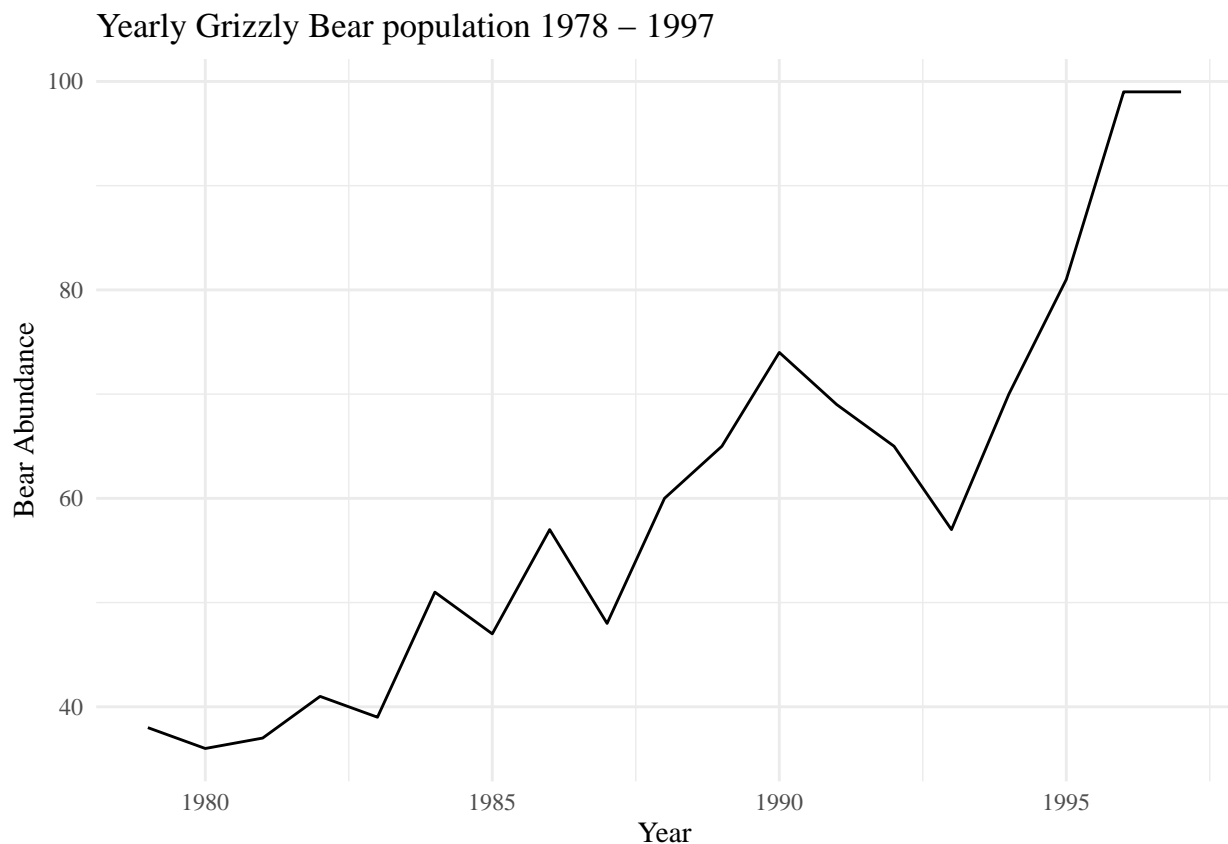
summary(mid_model)
```

The grizzly population continued to decline during this period. Between 1969 and 1978 the population decreased by 8 bears, representing close to a 2% decline overall. The rate of decline was similar to the

previous 10 years. Using a simple linear regression we find that from 1959 to 1968, the average rate of decline was -0.763 bears per year, while from 1969 to 1978 the population declined on average -0.757 bears per year.

**8. Finally look at the data after 1978. Did population size continue to change? What was the direction and magnitude of the trend? (1 point)**

```
grizzly_late <- grizzly %>%  
  filter( Year > 1978)  
  
grizzly_plot_3 <- ggplot(grizzly_late, aes(x = Year, y = N)) +  
  geom_line()+  
  theme_minimal()+  
  labs(x = "Year", y = "Bear Abundance", title = "Yearly Grizzly Bear population 1978 - 1997")+  
  theme(text = element_text(family = "Times"))  
  
grizzly_plot_3
```



```
#statistics - how has the population declined?  
late_diff <- (grizzly_late$N[10]-grizzly_late$N[1])  
late_pyr <- late_diff/10  
  
#the total percent decline  
late_perc <- late_diff/sum(grizzly_early$N)  
  
#use linear regression to compare slopes - is the slope the rate of decline?
```

```
late_model <-lm(grizzly_late$N ~ grizzly_late$Year)

summary(late_model)
```

For the last 10 years represented in the data set, there is an increase in bear abundance, with 22 more bears in 1979 than in 1968. Using linear regression, this amounts to an on average increase of 3.15 bears per year.

**9. Write a short paragraph to the park superintendent describing your conclusions on the effects (if any) of the dump closures on the grizzly bear population. (1 point)**

While the dump closures appeared to coincide with a decline in the grizzly bear population, our analysis suggests that it actually had very little short term effect. The bear population was declining prior to the dump closures at about -.76 bears per year, and this rate remained approximately the same for the 10 year period after. Longer term, the dump closures may have had a positive effect on the grizzly population, as bear abundance increased between 1979 and 1997 at an average rate of approximately 3 bears per year. These findings suggest that while dump closures may have at first increased human-bear conflict, in the long term the decision helped keep bears away from the more densely populated areas of the park and help to decrease conflicts. Another possibility is that the dump closures caused bears to rely more on their natural diets, helping individual bears to be healthier and strengthening the population overall.