Discussion 4

- 1. Consider a large population which has true mean μ and true standard deviation σ . We take a sample of size 3 from this population, thinking of the sample as the RVs X_1, X_2, X_3 where X_i can be considered iid (independent identically distributed). We are interested in estimating μ .
 - a. Consider the estimator $\hat{\mu}_1 = X_1 + X_2 X_3$. What is the mean of this estimator?
 - b. Find the variance of $\hat{\mu_1}$.
 - c. Consider the estimator $\hat{\mu_2} = \frac{X_1 + X_2 + X_3}{3}$. Is this estimator biased?
 - d. Find the variance of $\hat{\mu_2}$.
 - e. Now, consider the estimator $\hat{\mu}_3 = \frac{X_1 + 2X_2 + 3X_3}{6}$. What is the mean of this estimator?
 - f. Find the variance of $\hat{\mu_3}$.
 - g. Which of these three estimators is preferable? Why?
- 2. A packing plant fills bags with cement. The weight X kg of a bag of cement can be modeled by a normal distribution with mean 50kg and standard deviation 0.7kg.
 - a. Find P(X > 51)
 - b. Three bags are selected randomly. Find the probability that at least two weigh more than 51kg.
- 3. Weights of female cats are well approximated by a normal distribution with mean 4.1kg and standard deviation of $0.6 \text{kg } X \sim N(4.1, 0.6^2)$.
 - a. What proportion of female cats have weights between 3.7 and 4.4kg?
 - b. A certain female cat has a weight that is 0.5 standard deviations above the mean. What proportion of female cats are heavier than this one?
 - c. How heavy is a female cat whose weight is on the 80th percentile?
 - d. A female cat is chosen at random. What is the probability that she weighs more than 4.5 kg?
- 4. Take a look at the R-code below. [Note: You can also find this as a separate .Rmd file on Canvas. Everything after a pound symbol (#) is a comment, and NOT part of the R code.]
 - a. Walk through it, one line at a time, and try to make sense of it.
 - b. Run all the code, **EXCEPT FOR THE LAST LINE**. This should generate a bunch of QQ-plots. Half of them are from a normal distribution, the other half an exponential distribution. Can you tell which correspond to samples from the normal distribution, and which correspond to samples from the exponential distribution?
 - c. Repeat with sample_size <- 100. Does this change anything?

```
library(tidyverse)
library(distributions3)

## Create two random variables -- one Normal and one Exponential
X <- Normal(mu = 1, sigma = 1)
Y <- Exponential(rate = 1)</pre>
```

```
sample_size <- 10</pre>
## Create a vector of i's to use for scrambling
is <- sample(1:16)
normal_samples <- data.frame(i = is[1:8]) %>%
 mutate(distribution = "Normal",
                                                    # Create column that just says "Normal".
                                                    # For book keeping.
                                                    # map runs the function "random" with
         sample = map(i, random,
                      d = X, n = sample_size)) %% # arguments d = X, and n = sample_size
                                                    # for each i. I.e. it creates 8 random
                                                    # samples of size sample_size from the
                                                    # distribution X.
                                                    # The result is this weird list thing...
  unnest_longer(col = sample) # This qet's rid of the list and simply gives us a
                              # data.frame that we can work with...
exponential_samples <- data.frame(i = is[9:16]) %>%
 mutate(distribution = "Exponential",
         sample = map(i, random, d = Y, n = sample_size)) %>%
  unnest_longer(col = sample)
# Here we just stack the two data.frame on top of each other (i.e. bind the rows together)
all_samples <- bind_rows(</pre>
 normal_samples,
  exponential_samples
# Create QQ-plots
ggplot(all_samples,
       aes(sample = sample)) +
  geom_qq() +
 facet_wrap(~i) +
 geom_abline(aes(slope = sd(sample), intercept = mean(sample)))
## To reveal the truth behind the i's, run this line:
all_samples %>% select(i, distribution) %>% unique()
```