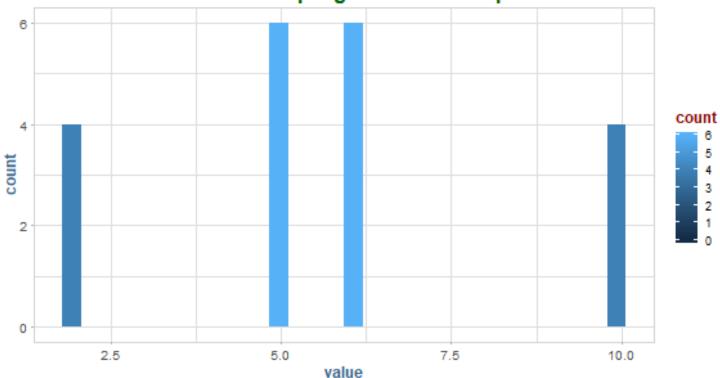
Consider the population {1, 2, 5, 6, 10, 12}.

Find (and plot) the sampling distribution of medians for samples of size 3 without replacement.

```
p <- c(1, 2, 5, 6, 10, 12)
c <- combinations(v = p, n = 6, r = 3)
t <- apply(c, 1, median)

ggplot(data.table(value = t), aes(value, fill = ..count..)) +
    geom_histogram(bins = 30) +
    labs(title = "Median Sampling Distribution of p")</pre>
```





Compare the median of the population to the mean of the medians.

Median of p = 5.5. Mean of Medians of p = 5.7

2.5

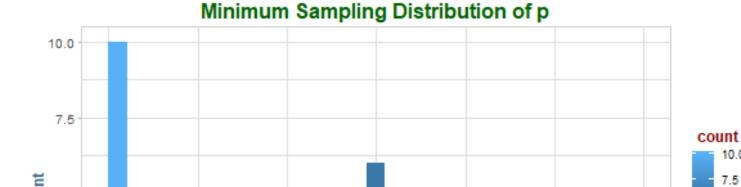
0.0

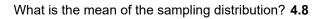
4.2

Consider the population {3, 6, 7, 9, 11, 14}.

For samples of size 3 without replacement, find (and plot) the sampling distribution for the minimum.

```
p \leftarrow c(3, 6, 7, 9, 11, 14)
c \leftarrow combinations(v = p, n = 6, r = 3)
t <- apply(c, 1, min)
ggplot(data.table(value = t), aes(value, fill = ..count..)) +
   geom_histogram(bins = 30) +
   labs(title = "Minimum Sampling Distribution of p")
```





The statistic is an estimate of some parameter - what is the value of that parameter?

This is an estimation of the minimum, which is: 3

6

value

8

10.0 7.5

5.0 2.5 0.0

Let A denote the population {1, 3, 4, 5} and B the population {5, 7, 9}.

```
A \leftarrow c(1, 3, 4, 5)

B \leftarrow c(5, 7, 9)
```

Let *X* be a random value from *A*, and *Y* and random value from *B*.

a.) Find the sampling distribution of X + Y.

```
result = numeric(12)
index <- 1
for(j in 1:length(A))
{
   for(k in 1:length(B))
   {
      result[index] <- A[j] + B[k]
      index <- index + 1
   }
}</pre>
sort(result)
```

```
[1] 6 8 8 9 10 10 10 11 12 12 13 14
```

b.) In this example, does the sampling distribution depend on whether you sample with or without replacement?

No.

Why or why not?

Because 5 in is both sets.

c.) Compute the mean of the values for each of A and B and the values in the sampling distribution of X + Y.

Mean of A: 3.25. Mean of B: 7.

Mean of A + B: 10.25

How are the means related?

mean(A) + mean(B) = mean(A + B).

d.) Suppose you draw a random value from A and a random value from B.

```
prob <- sum(result >= 13) / length(result)
```

What is the probability that the sum is 13 or larger? **16.666667%**

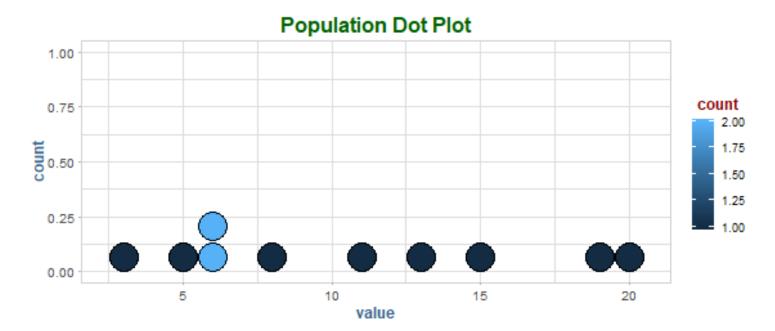
Consider the population {3, 5, 6, 6, 8, 11, 13, 15, 19, 20}.

a.) Compute the mean and standard deviation and create a dot plot of its distribution.

```
p <- c(3, 5, 6, 6, 8, 11, 13, 15, 19, 20)

mu <- mean(p)
sigma <- sd(p)

ggplot(data.table(value = p)) +
    geom_dotplot(aes(value, fill = ..count..), binwidth = 1) +
    labs(title = "Population Dot Plot")</pre>
```



```
\mu = 10.6, \sigma = 5.9851668
```

b.) Simulate the sampling distribution of \bar{X} by taking random samples of size 4 and plot your results.

```
N <- 10e2
results <- numeric(N)

for( i in 1:N)
{
   index <- sample(length(p), size = 4, replace = F)
   results[i] <- mean( p[index] )
}</pre>
```

```
ggplot(data.table(value = results)) +
  geom_histogram(aes(value, fill = ..count..), bins = 30) +
  labs(title = "Sample Means")
```



```
xbar <- mean(results)
se <- sd(results) / sqrt(N)</pre>
```

Compute the mean and standard error, and compare to the population mean and standard deviation.

mean: 10.644, standard error: 0.0738408

c.) Use the simulation to find $P(\bar{X} < 11)$.

$$P(\bar{X} < 11) = 53.7\%$$

Consider two populations A = {3, 5, 7, 9, 10, 16}, B = {8, 10, 11, 15, 18, 25, 28}.

```
A <- c(3, 5, 7, 9, 10, 16)
B <- c(8, 10, 11, 15, 18, 25, 28)
```

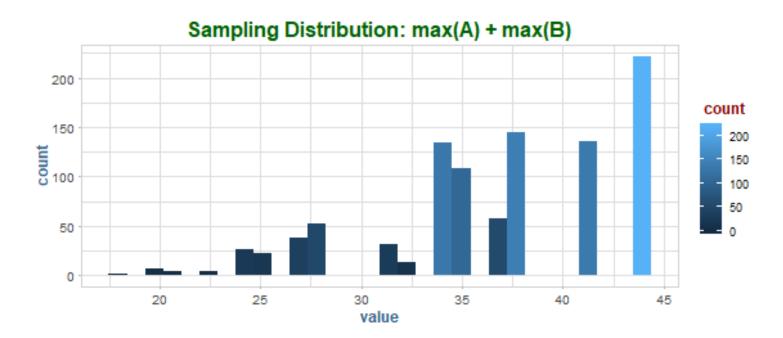
a.) Using R, draw random samples (without replacement) of size 3 from each population, and simulate the sampling distribution of the sum of their maximums.

```
N <- 10e2
results <- numeric(N)

for(i in 1:N)
{
    samp.a <- sample(A, 3, replace = F)
    samp.b <- sample(B, 3, replace = F)

    results[i] <- max(samp.a) + max(samp.b)
}

ggplot(data.table(value = results)[, index := .I]) +
    geom_histogram(aes(value, fill = ..count..), bins = 30) +
    labs(title = "Sampling Distribution: max(A) + max(B)")</pre>
```



b.) Use your simulation to estimate the probability that the sum of the maximums is less than 20.

```
prob <- mean(results < 20)</pre>
```

Probability: 0.1%

c.) Draw random samples of size 3 from each population, and find the maximum of the union of these two sets.

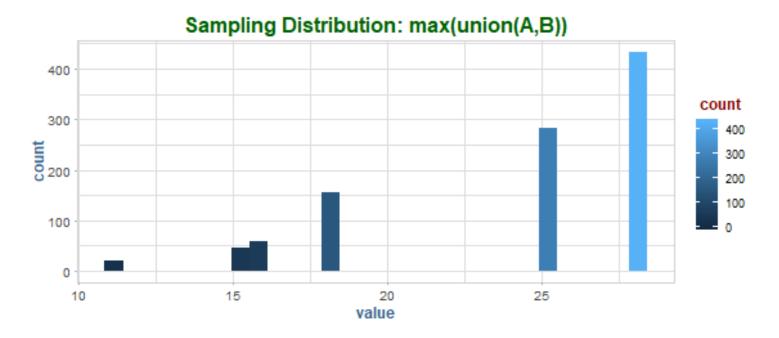
Simulate the sampling distribution of the maximums of this union.

```
results <- numeric(N)

for(i in 1:N)
{
    samp.a <- sample(A, 3, replace = F)
    samp.b <- sample(B, 3, replace = F)

    results[i] <- max(union(samp.a, samp.b))
}

ggplot(data.table(value = results)[, index := .I]) +
    geom_histogram(aes(value, fill = ..count..), bins = 30) +
    labs(title = "Sampling Distribution: max(union(A,B))")</pre>
```



d.) Use simulation to find the probability that the maximum of the union is less than 20.

```
prob <- mean(results < 20)</pre>
```

Probability: 28.2%