



Given $x_j \in X$ and $x_i^* \in X^*$, let x_1^*, \dots, x_n^* be n independent trials, each resulting in either success or failure, with success defined as $x_i^* = x_j$. The probability of k success is a binomial random variable with the probability of success for any trial $p = \frac{1}{n}$. Therefore, by Theorem 3.2.1 (Larsen & Marx (2018), p. 104):

$$P(k \text{ successes}) = \binom{n}{k} \left(\frac{1}{n}\right)^k \left(\frac{n-1}{n}\right)^{n-k}$$

We can see this in R by calculating the theoretical probability and creating a bootstrap sample for

$$\hat{\theta}_i^* = \sum_{i=1}^n I_{(x_j^* = x_i)}$$

Where $I(\phi)$ is a Binomial random variable with success probability ϕ

```
## Define nchoosek (Larsen & Marx (2018), p. 84).
nchoosek <- function(n,k){
  factorial(n)/((factorial(k))*(factorial(n-k)))
}

## Define full function (Larsen & Marx (2018), p. 104)
binom.nk <- function(n,k){
  nchoosek(n,k) * ((1/n)^k) * ((n-1)/n)^(n-k)
}

## Get a base X of size n
vctrzdData <- c(1,2,3.5,4,7.3,8.6,12.4,13.8)

## Set parameters
n <- length(vctrzdData)
k <- 2
B <- 1000

## Boot Strap
count <- c()
for (i in 1:B){
  ## Get B samples from X
  btstrpSmp1 = sample(vctrzdData,length(vctrzdData), replace = TRUE)
  ## Count how many times x_i shows up in the sample
  count[i] <- sum(btstrpSmp1 == vctrzdData[3])
}

## Probability
(sum(count == k))/B
```

```
## [1] 0.211
```

```
## Compare to theoretical probability
binom.nk(length(vctrzdData),k)
```