

L4:Simulation and probability

Learning objectives

- Understand Point Estimates and Sampling Variability
- Visualize and Interpret Sampling Distributions
- Calculate and Interpret Standard Error

Can we verify this theorem??

Central Limit Theorem

- Let X_1, \dots, X_n be iid random variables with finite mean μ and standard deviation σ . Then

$$\frac{\bar{X} - \mu}{\sigma/\sqrt{n}} \rightarrow Z \quad \text{as } n \rightarrow \infty$$

where Z is a standard normal random variable.

1 - Different shapes of distributions

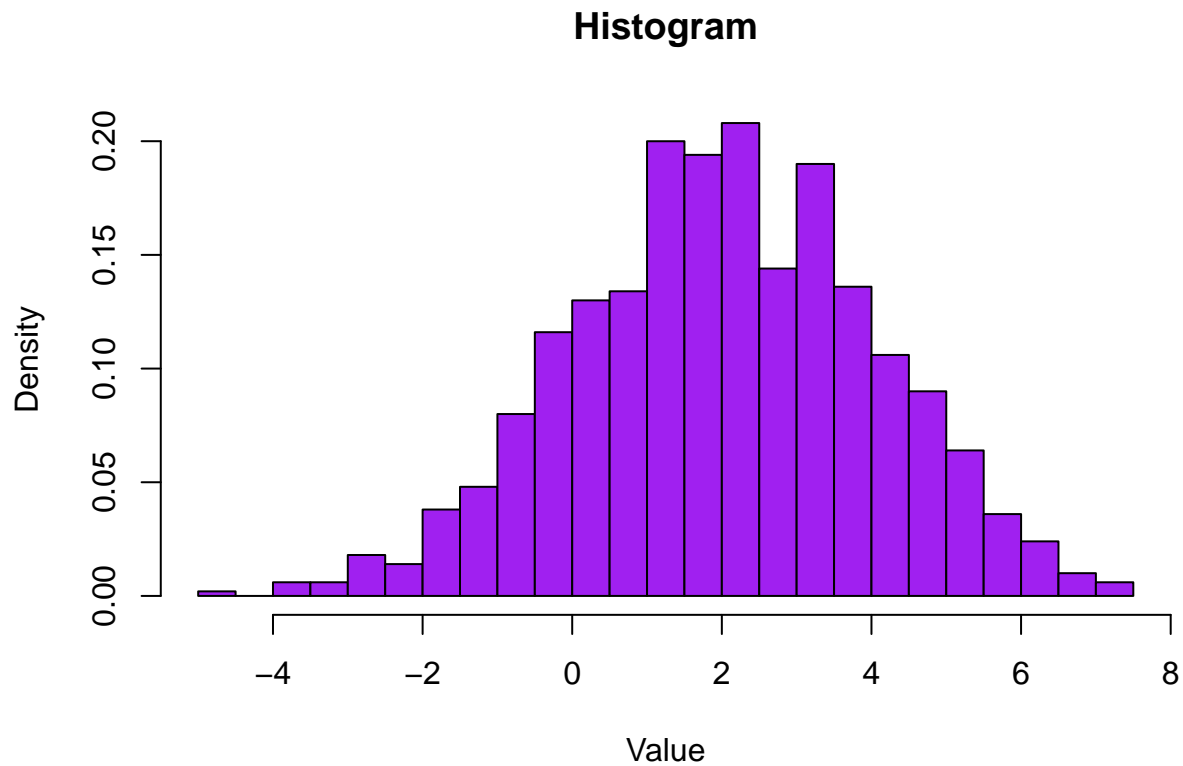
Some distribution functions in R.

- **Normal distribution** : `rnorm(n, mean, sd)`
 - **n**: Number of random values to generate
 - **mean**: Mean (center) of the distribution
 - **sd**: Standard deviation (spread) of the distribution
- **Exponential distribution** : `rexp(n, rate)`
 - **n**: Number of random values to generate
 - **rate**: Rate parameter (inverse of the mean)
 - The mean will be `1/rate`
- **Binomial distribution** : `rbinom(n, size, prob)`
 - **n**: Number of random values to generate
 - **size**: Number of trials
 - **prob**: Probability of success in each trial
 - The mean will be `n*prob`
- **Poisson distribution** : `rpois(n, lambda)`
 - **n**: Number of random values to generate
 - **lambda**: Expected number of occurrences in a fixed interval
 - The mean will be `lambda`
- **Gamma distribution** : `rgamma(n, shape, scale)`
 - **n**: Number of random values to generate

- `shape`: Shape parameter (determines the distribution's form)
- `scale`: Scale parameter (stretches or shrinks the distribution)
- The mean will be `shape*scale`

```
data <- rnorm(1000, mean=2, sd=2)
```

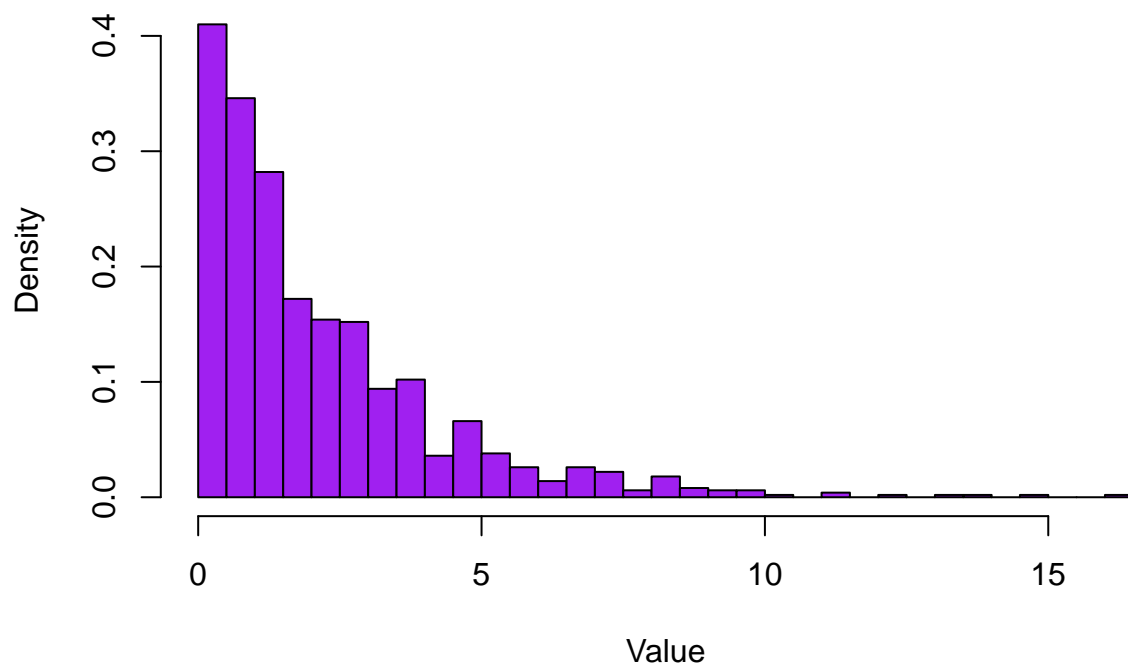
```
hist(data, breaks = 30, col = "purple", main = "Histogram", xlab = "Value", probability = TRUE)
```



```
data <- rexp(1000, rate = 1/2)
```

```
hist(data, breaks = 30, col = "purple", main = "Histogram", xlab = "Value", probability = TRUE)
```

Histogram



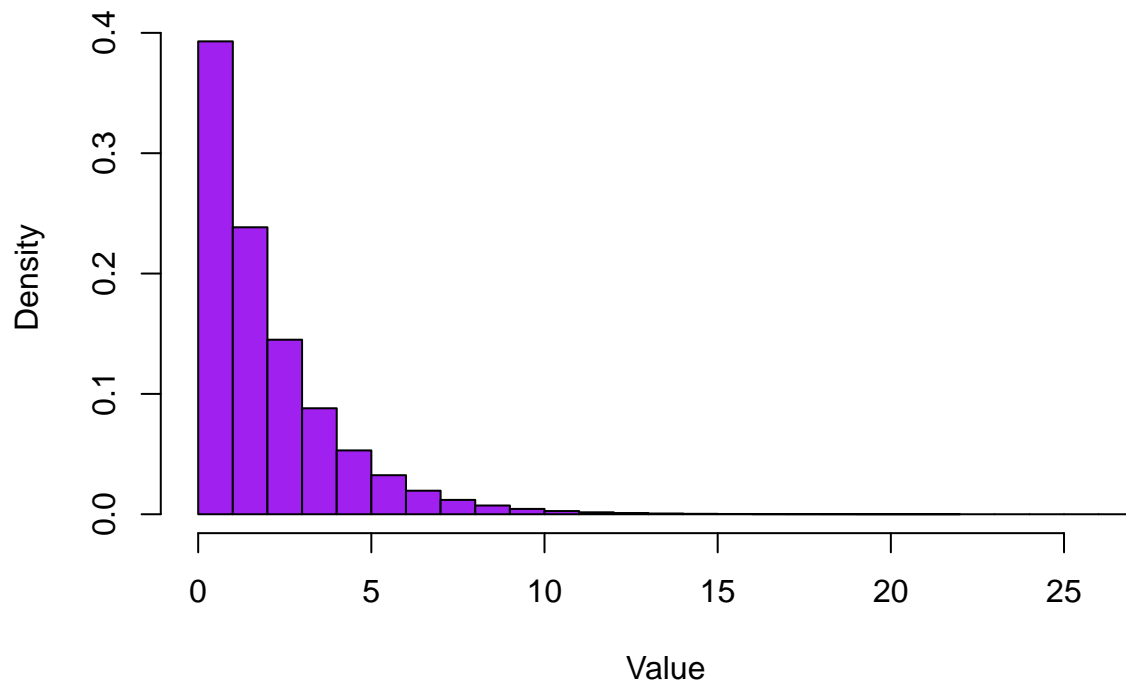
3 - Procedure

- 1. Obtain population data with some distribution.
- 2. Show the shape of the whole population.
- 3. Obtain \bar{X}_i for $i = 1 \dots n$ by repeating step 1.
- 4. Show the shape of \bar{X}_i for $i = 1 \dots n$.
- Change sample size n and do the whole step from 1 to 4.
- What can you conclude?

```
mean = 2
pop_data <- rexp(1000000, rate = 1/mean)

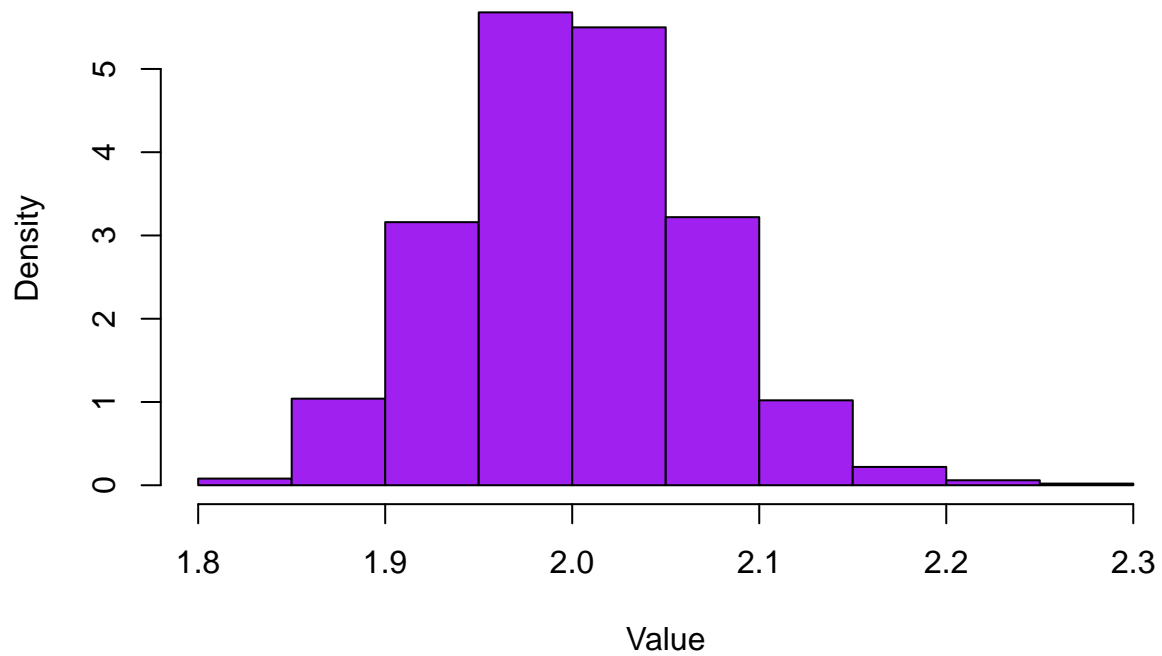
hist(pop_data, col = "purple", main = "Histogram of population data", xlab = "Value", probability = TRUE)
```

Histogram of population data



```
sample_and_get_xbar <- function(n, pop_data){  
  sample_data <- sample(pop_data, size=n, replace=TRUE)  
  xbar = mean(sample_data)  
  return(xbar)  
}  
  
n=1000  
# Xbar1 = sample_and_get_xbar(n, pop_data)  
Xbar_data <- replicate(n, sample_and_get_xbar(n, pop_data ))  
head(Xbar_data)  
  
## [1] 2.065205 1.899304 2.067580 2.030113 1.919196 1.926914  
hist(Xbar_data, col = "purple", main = "Histogram of Xbar", xlab = "Value", probability = TRUE)
```

Histogram of Xbar



```
# better version
ggplot(data = data.frame(Xbar_data),
       aes(x = Xbar_data)) +
  geom_histogram(fill = "blue", alpha = 0.5,
                 color = "black") +
  labs(title = "Histogram of Sample Mean",
       x = "Xbar",
       y = "Frequency") +
  theme_minimal()
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

