

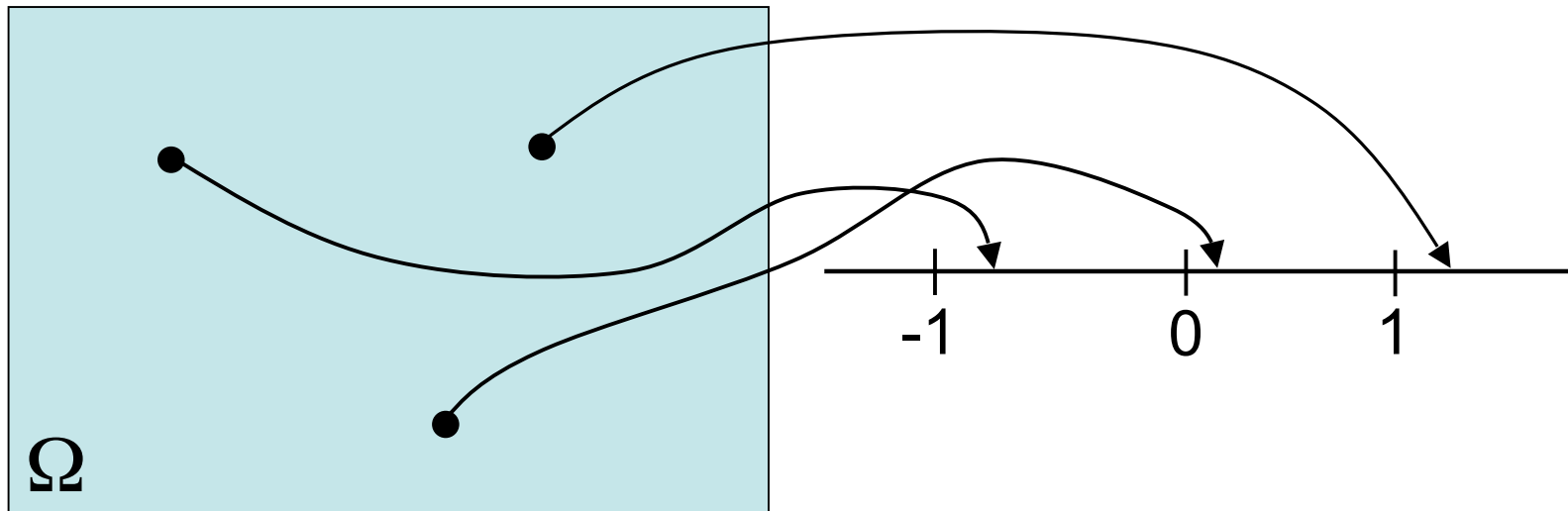
Lecture 4: Random Variables and Distributions

Goals

- Random Variables
- Overview of discrete and continuous distributions important in genetics/genomics
- Working with distributions in R

Random Variables

A rv is any rule (i.e., function) that associates a number with each outcome in the sample space



Two Types of Random Variables

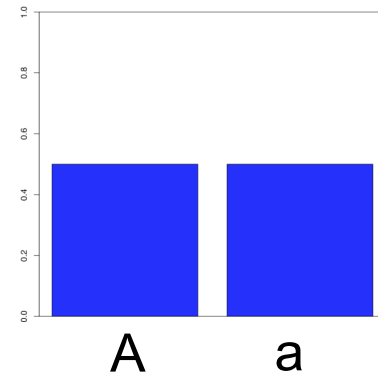
- A **discrete** random variable has a **countable** number of possible values
- A **continuous** random variable takes all values in an interval of numbers

Probability Distributions of RVs

Discrete

Let X be a discrete rv. Then the *probability mass function (pmf)*, $f(x)$, of X is:

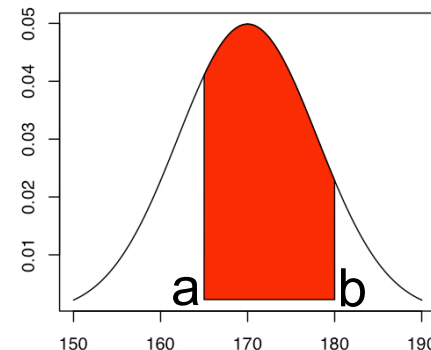
$$f(x) = \begin{cases} P(X = x), & x \in \Omega \\ 0, & x \notin \Omega \end{cases}$$



Continuous

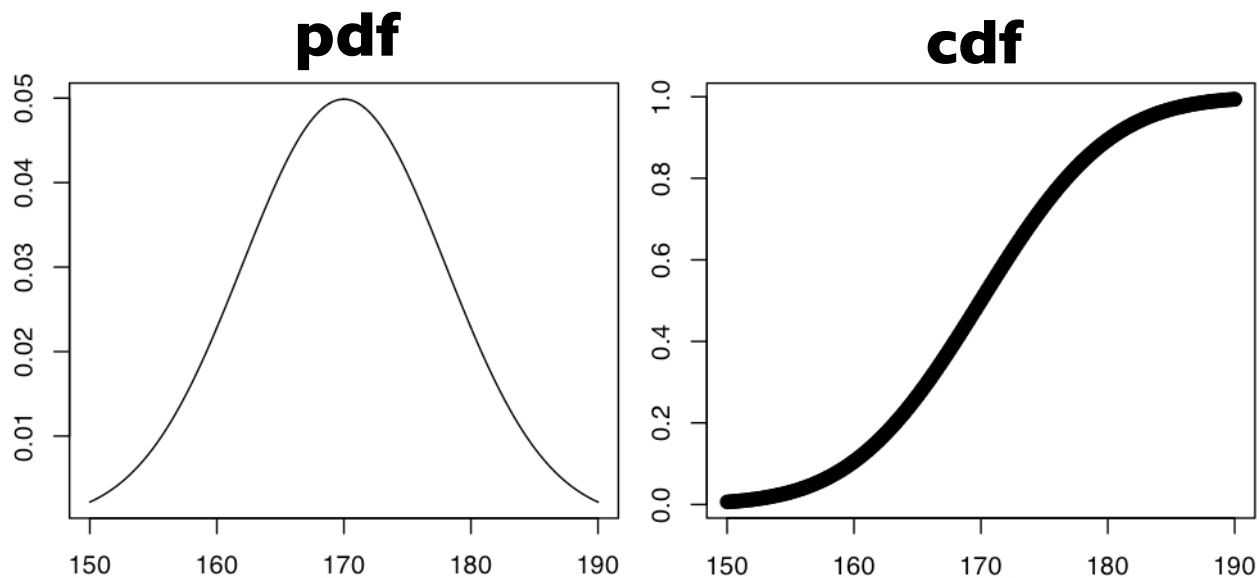
Let X be a continuous rv. Then the *probability density function (pdf)* of X is a function $f(x)$ such that for any two numbers a and b with $a \leq b$:

$$P(a \leq X \leq b) = \int_a^b f(x) dx$$



Using CDFs to Compute Probabilities

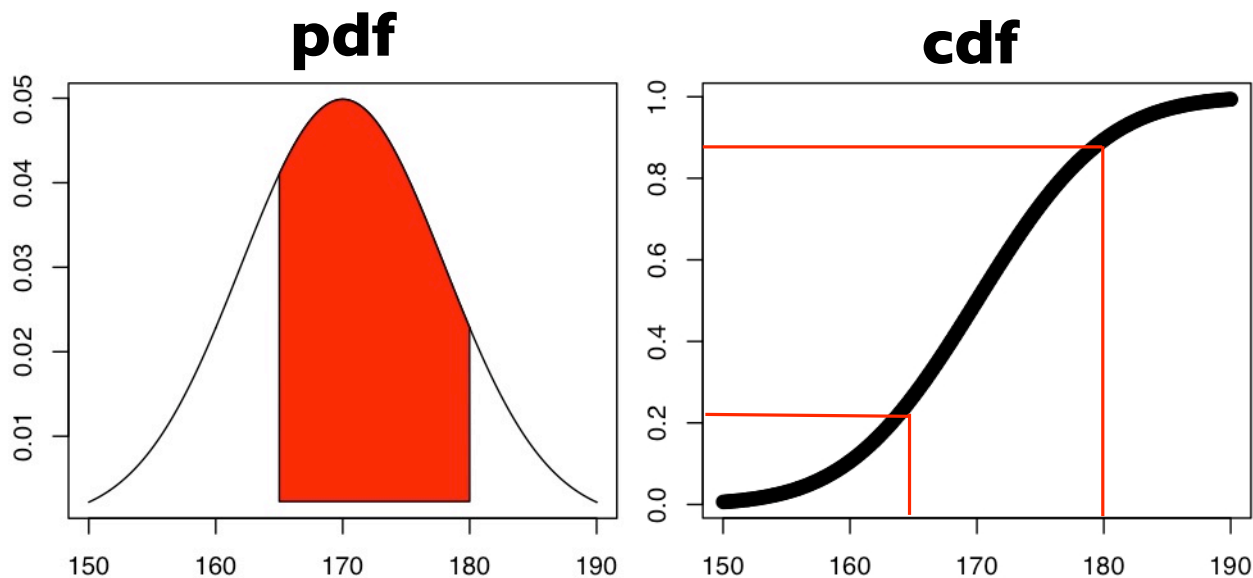
Continuous rv: $F(x) = P(X \leq x) = \int_{-\infty}^x f(y)dy$



$$P(a \leq X \leq b) = F(b) - F(a)$$

Using CDFs to Compute Probabilities

Continuous rv: $F(x) = P(X \leq x) = \int_{-\infty}^x f(y) dy$



$$P(a \leq X \leq b) = F(b) - F(a)$$

Expectation of Random Variables

Discrete

Let X be a discrete rv that takes on values in the set D and has a pmf $f(x)$. Then the expected or mean value of X is:

$$\mu_X = E[X] = \sum_{x \in D} x \cdot f(x)$$

Continuous

The expected or mean value of a continuous rv X with pdf $f(x)$ is:

$$\mu_X = E[X] = \int_{-\infty}^{\infty} x \cdot f(x) dx$$

Variance of Random Variables

Discrete

Let X be a discrete rv with pmf $f(x)$ and expected value μ . The variance of X is:

$$\sigma_X^2 = V[X] = \sum_{x \in D} (x - \mu)^2 = E[(X - \mu)^2]$$

Continuous

The variance of a continuous rv X with pdf $f(x)$ and mean μ is:

$$\sigma_X^2 = V[X] = \int_{-\infty}^{\infty} (x - \mu)^2 \cdot f(x) dx = E[(X - \mu)^2]$$

Example of Expectation and Variance

- Let L_1, L_2, \dots, L_n be a sequence of n nucleotides and define the rv X_i :

$$X_i \begin{cases} 1, & \text{if } L_i = A \\ 0, & \text{otherwise} \end{cases}$$

- pmf is then: $P(X_i = 1) = P(L_i = A) = p_A$

$$P(X_i = 0) = P(L_i = C \text{ or } G \text{ or } T) = 1 - p_A$$

- $E[X] = 1 \times p_A + 0 \times (1 - p_A) = p_A$

- $\text{Var}[X] = E[X - \mu]^2 = E[X^2] - \mu^2$
 $= [1^2 \times p_A + 0^2 \times (1 - p_A)] - p_A^2$
 $= p_A (1 - p_A)$

The Distributions We'll Study

1. Binomial Distribution
2. Hypergeometric Distribution
3. Poisson Distribution
4. Normal Distribution

Binomial Distribution

- **Experiment consists of n trials**
 - e.g., 15 tosses of a coin; 20 patients; 1000 people surveyed
- **Trials are identical and each can result in one of the same two outcomes**
 - e.g., head or tail in each toss of a coin
 - Generally called “success” and “failure”
 - Probability of success is p , probability of failure is $1 - p$
- **Trials are independent**
- **Constant probability for each observation**
 - e.g., Probability of getting a tail is the same each time we toss the coin

Binomial Distribution

pmf:

$$P\{X = x\} = \binom{n}{x} p^x (1 - p)^{n-x}$$

cdf:

$$P\{X \leq x\} = \sum_{y=0}^x \binom{n}{y} p^y (1 - p)^{n-y}$$

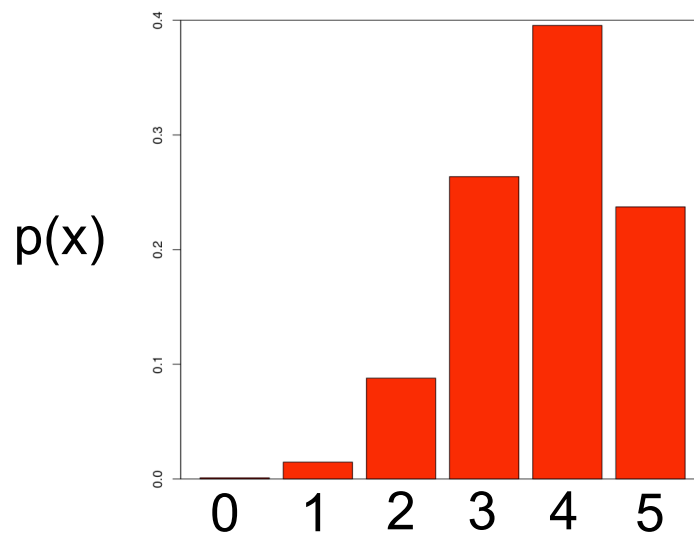
$$\mathbf{E(x) = np}$$

$$\mathbf{Var(x) = np(1-p)}$$

Binomial Distribution: Example 1

- A couple, who are both carriers for a recessive disease, wish to have 5 children. They want to know the probability that they will have four healthy kids

$$\begin{aligned}P\{X = 4\} &= \binom{5}{4} 0.75^4 \times 0.25^1 \\ &= 0.395\end{aligned}$$



Binomial Distribution: Example 2

- Wright-Fisher model: There are i copies of the A allele in a population of size $2N$ in generation t . What is the distribution of the number of A alleles in generation $t + 1$?

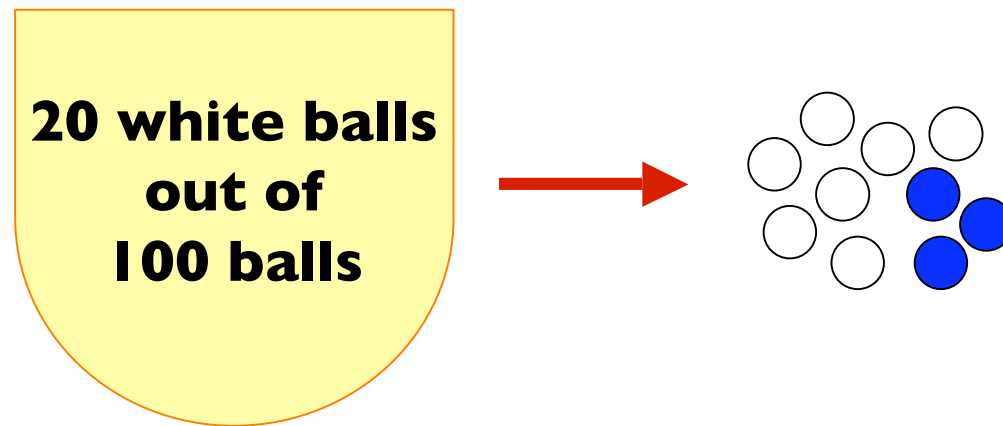
$$p_{ij} = \binom{2N}{j} \left(\frac{i}{2N}\right)^j \left(1 - \frac{i}{2N}\right)^{2N-j} \quad j = 0, 1, \dots, 2N$$

Hypergeometric Distribution

- **Population to be sampled consists of N finite individuals, objects, or elements**
- **Each individual can be characterized as a success or failure, m successes in the population**
- **A sample of size k is drawn and the rv of interest is $X =$ number of successes**

Hypergeometric Distribution

- Similar in spirit to Binomial distribution, but from a **finite** population **without** replacement



If we randomly sample 10 balls, what is the probability that 7 or more are white?

Hypergeometric Distribution

- pmf of a hypergeometric rv:

$$P\{X = i \mid n, m, k\} = \frac{\binom{m}{i} \binom{n}{k-i}}{\binom{m+n}{k}} \quad \text{For } i = 0, 1, 2, 3, \dots$$

Where,

k = Number of balls selected

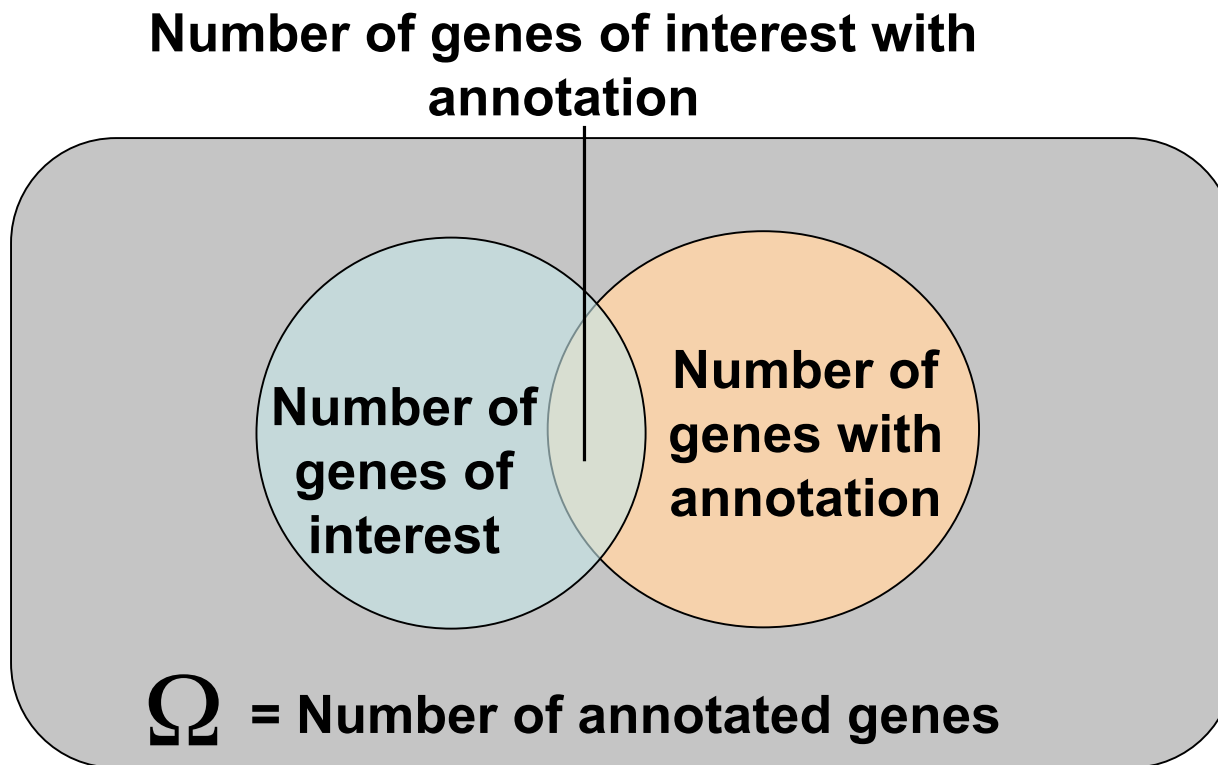
m = Number of balls in urn considered “success”

n = Number of balls in urn considered “failure”

m + n = Total number of balls in urn

Hypergeometric Distribution

- Extensively used in genomics to test for “enrichment”:



Poisson Distribution

- **Useful in studying rare events**
- **Poisson distribution also used in situations where “events” happen at certain points in time**
- **Poisson distribution approximates the binomial distribution when n is large and p is small**

Poisson Distribution

- A rv X follows a Poisson distribution if the pmf of X is:

$$P\{X = i\} = e^{-\lambda} \frac{\lambda^i}{i!} \quad \text{For } i = 0, 1, 2, 3, \dots$$

- λ is frequently a rate per unit time:

$$\lambda = \alpha t = \text{expected number of events per unit time } t$$

- Safely approximates a binomial experiment when $n > 100$, $p < 0.01$, $np = \lambda < 20$)
- $E(X) = \text{Var}(X) = \lambda$

Poisson RV: Example 1

- The number of crossovers, X , between two markers is $X \sim \text{poisson}(\lambda=d)$

$$P\{X = i\} = e^{-d} \frac{d^i}{i!}$$

$$P\{X = 0\} = e^{-d}$$

$$P\{X \geq 1\} = 1 - e^{-d}$$

Poisson RV: Example 2

- Recent work in *Drosophila* suggests the spontaneous rate of deleterious mutations is ~ 1.2 per diploid genome. Thus, let's tentatively assume $X \sim \text{poisson}(\lambda = 1.2)$ for humans. What is the probability that an individual has 12 or more spontaneous deleterious mutations?

$$P\{X \geq 12\} = 1 - \sum_{i=0}^{11} e^{-1.2} \frac{1.2^i}{i!}$$

$$= 6.17 \times 10^{-9}$$

Poisson RV: Example 3

- Suppose that a rare disease has an incidence of 1 in 1000 people per year. Assuming that members of the population are affected independently, find the probability of k cases in a population of 10,000 (followed over 1 year) for $k=0,1,2$.

The expected value (mean) = $\lambda = .001*10,000 = 10$

$$P(X = 0) = \frac{(10)^0 e^{-(10)}}{0!} = .0000454$$

$$P(X = 1) = \frac{(10)^1 e^{-(10)}}{1!} = .000454$$

$$P(X = 2) = \frac{(10)^2 e^{-(10)}}{2!} = .00227$$

Normal Distribution

- **“Most important” probability distribution**
- **Many rv’s are approximately normally distributed**
- **Even when they aren’t, their sums and averages often are (CLT)**

Normal Distribution

- pdf of normal distribution:

$$f(x; \mu, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma}} e^{-(x-\mu)^2 / 2\sigma^2}$$

- standard normal distribution ($\mu = 0, \sigma^2 = 1$):

$$f(z; 0, 1) = \frac{1}{\sqrt{2\pi\sigma}} e^{-z^2 / 2}$$

- cdf of Z:

$$P(Z \leq z) = \int_{-\infty}^z f(y; 0, 1) dy$$

Standardizing Normal RV

- If X has a normal distribution with mean μ and standard deviation σ , we can standardize to a standard normal rv:

$$Z = \frac{X - \mu}{\sigma}$$

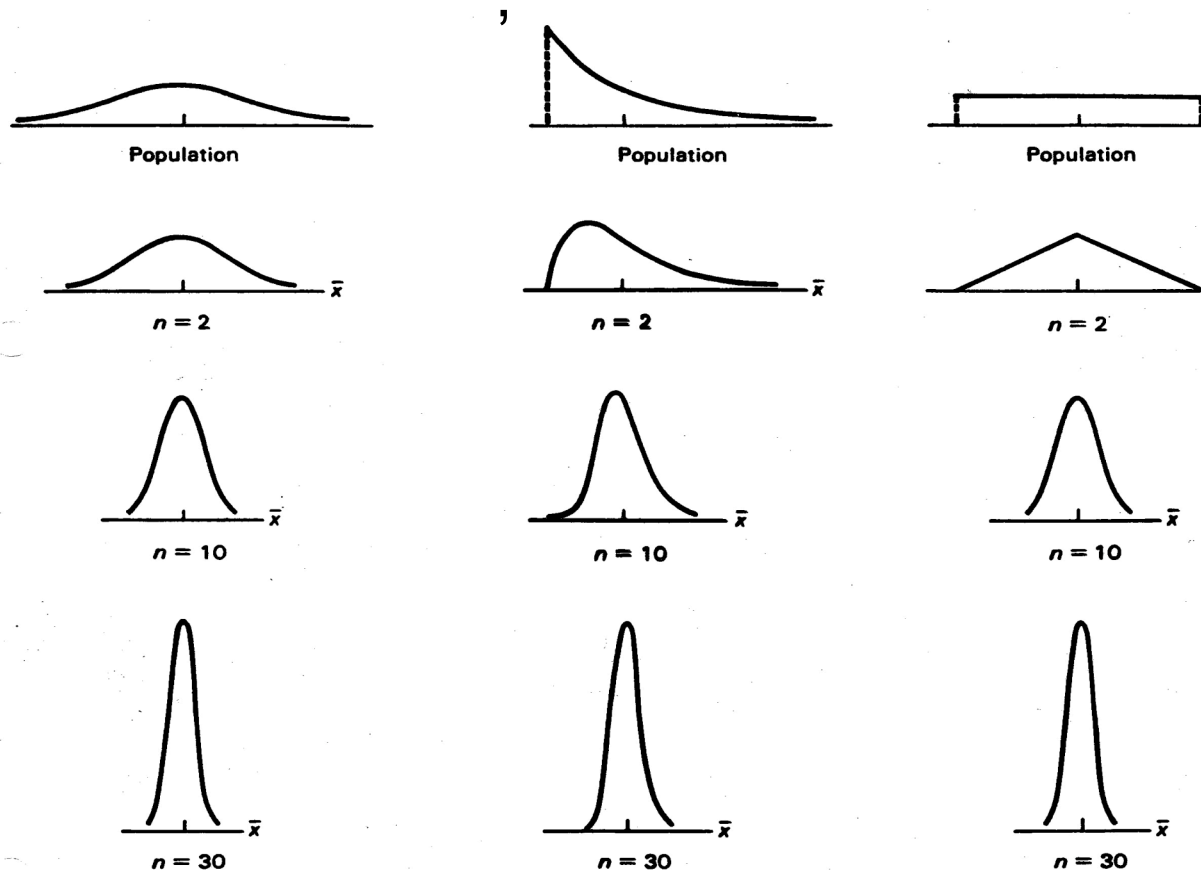
I Digress: Sampling Distributions

- Before data is collected, we regard observations as random variables (X_1, X_2, \dots, X_n)
- This implies that until data is collected, any function (statistic) of the observations (mean, sd, etc.) is also a random variable
- Thus, any statistic, because it is a random variable, has a probability distribution - referred to as a **sampling distribution**
- Let's focus on the sampling distribution of the mean, \bar{X}

Behold The Power of the CLT

- Let X_1, X_2, \dots, X_n be an iid random sample from a distribution with mean μ and standard deviation σ . If n is sufficiently large:

$$\bar{X} \sim N\left(\mu, \frac{\sigma}{\sqrt{n}}\right)$$



Example

- If the mean and standard deviation of serum iron values from healthy men are 120 and 15 mgs per 100ml, respectively, what is the probability that a random sample of 50 normal men will yield a mean between 115 and 125 mgs per 100ml?

First, calculate mean and sd to normalize (120 and $15/\sqrt{50}$)

$$\begin{aligned} p(115 \leq \bar{x} \leq 125) &= p\left(\frac{115 - 120}{2.12} \leq \bar{x} \leq \frac{125 - 120}{2.12}\right) \\ &= p(-2.36 \leq z \leq 2.36) \\ &= p(z \leq 2.36) - p(z \leq -2.36) \\ &= 0.9909 - 0.0091 \\ &= 0.9818 \end{aligned}$$

R

- **Understand how to calculate probabilities from probability distributions**
 - Normal: `dnorm` and `pnorm`
 - Poisson: `dpois` and `ppois`
 - Binomial: `dbinom` and `pbinom`
 - Hypergeometric: `dhyper` and `phyper`
- **Exploring relationships among distributions**