## **CONCEPT: MATHMATICAL MEASRUMENTS**

- Common statistical measurements are used in genetics to \_\_\_\_\_\_ phenotypes
  - ☐ The **mean** is an average of values
    - A **population** is all individuals within the group you're measuring
    - A **sample** is a representative subset of individuals in a population

**EXAMPLE:** Mean calculation

$$M = \frac{\Sigma(X)}{N}$$

Where  $\Sigma = \text{Sum of}$ 

X = Individual data points

N = Sample size (number of data points)

- ☐ The **variance** measures how far a set of values is from the mean
  - Covariance measures how much variation is common to 2+ traits

**EXAMPLE:** Variance calculation

$$S^2 = \frac{\Sigma (X-M)^2}{n-1}$$

Where  $\Sigma = \text{Sum of}$ 

X = Individual score

M = Mean of all scores

N =Sample size (number of scores)

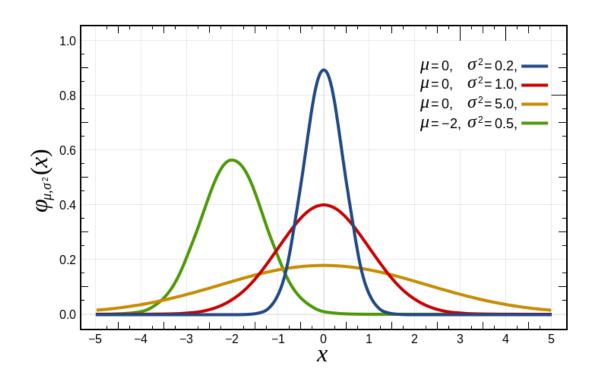
- ☐ The **standard deviation** measures the amount of variation that exists within a set of data
  - Standard error measures the accuracy of the sample mean

**EXAMPLE:** Standard deviation calculation

$$s = \sqrt{\frac{\sum (x - \bar{x})^2}{n - 1}}$$

- □ A **normal distribution** is the "bell curve" and visualizes the range of variation of a phenotype
  - Sometimes called a "frequency histogram" as it measures frequency of the trait on y axis

## **EXAMPLE**:



## PRACTICE:

Bristle Number	Number of Individuals
1	2,
2	3,
3	9
4	29,
5	55,
6	18,
7	4,

- 1. The table shows a distribution of bristle numbers in a *Drosophila* population. What is the mean bristle number?
  - a. 4.7
  - b. 80
  - c. 562
  - d. 5.0

Bristle Number	Number of Individuals	(X-M) <sup>2</sup>	Sum (X-M) <sup>2</sup>
1	2	13.69	27.38
2	3	7.29	21.87
3	9	2.89	26.01
4	29	0.49	14.21
5	55	0.09	4.95
6	18	1.69	30.42
7	4	5.29	21.16

- 2. The table shows a distribution of bristle numbers in a *Drosophila* population. What is the variance?
  - a. 1.0
  - b. 1.2
  - c. 5.5
  - d. 3.0

- 3. Using the variance calculated in problem #2, what is the standard deviation?

  - a. 1.0 b. 1.1 c. 2.3
  - d. 1.7