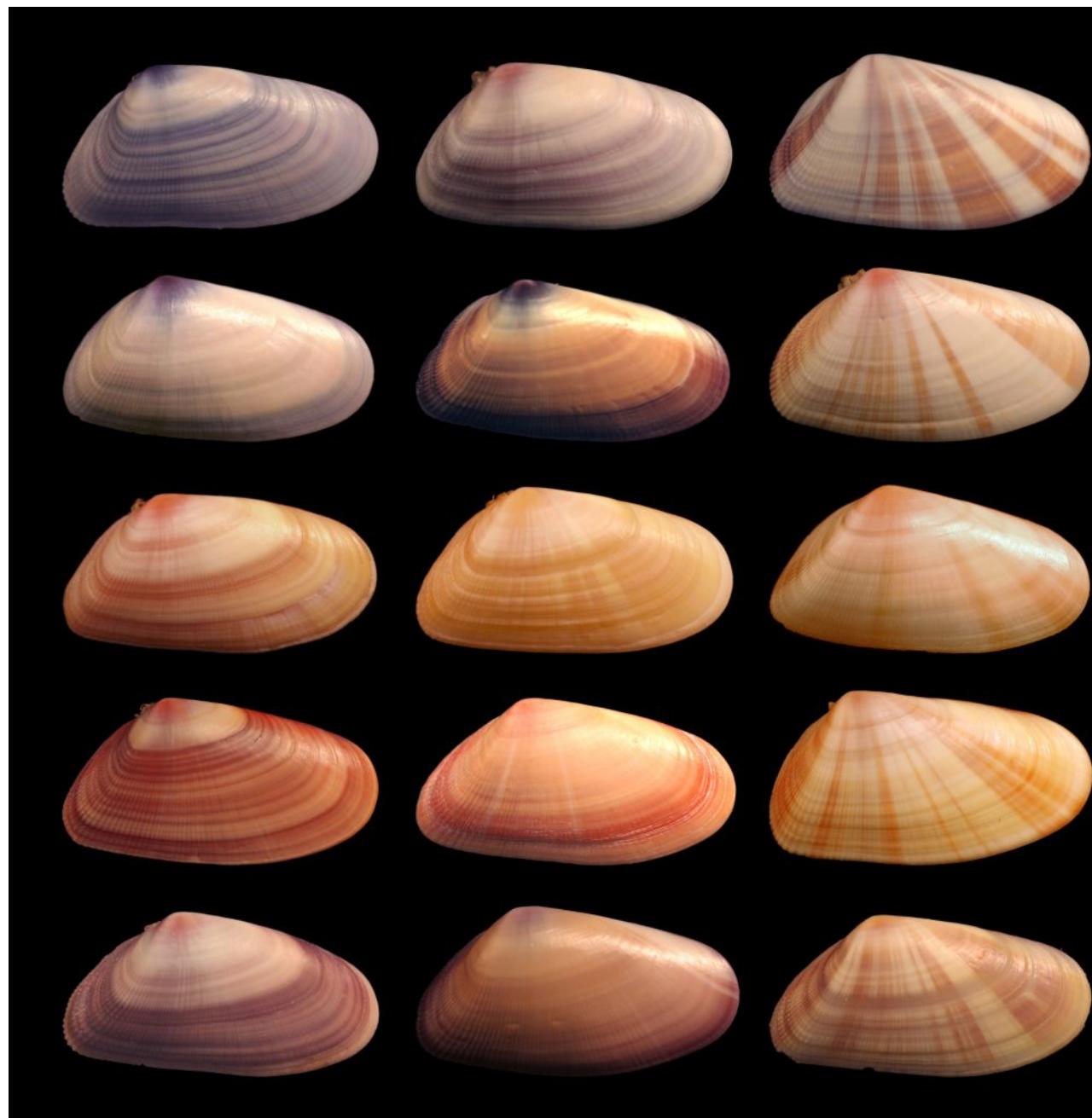


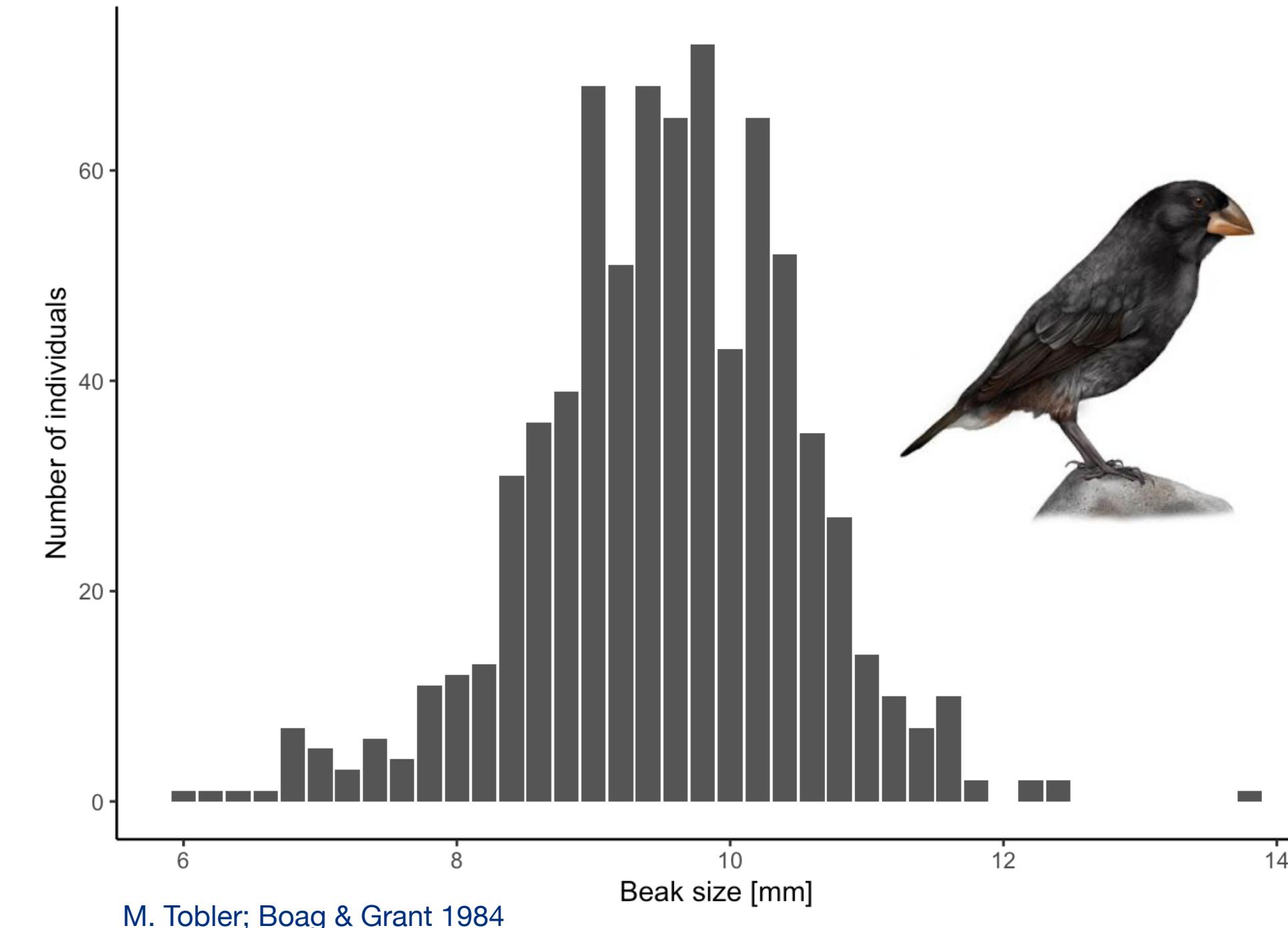
# Quantitative Traits in Nature: How do first principles of genetic inheritance explain their underlying statistical distributions?

Clay Small

Data Science Department - University of Oregon



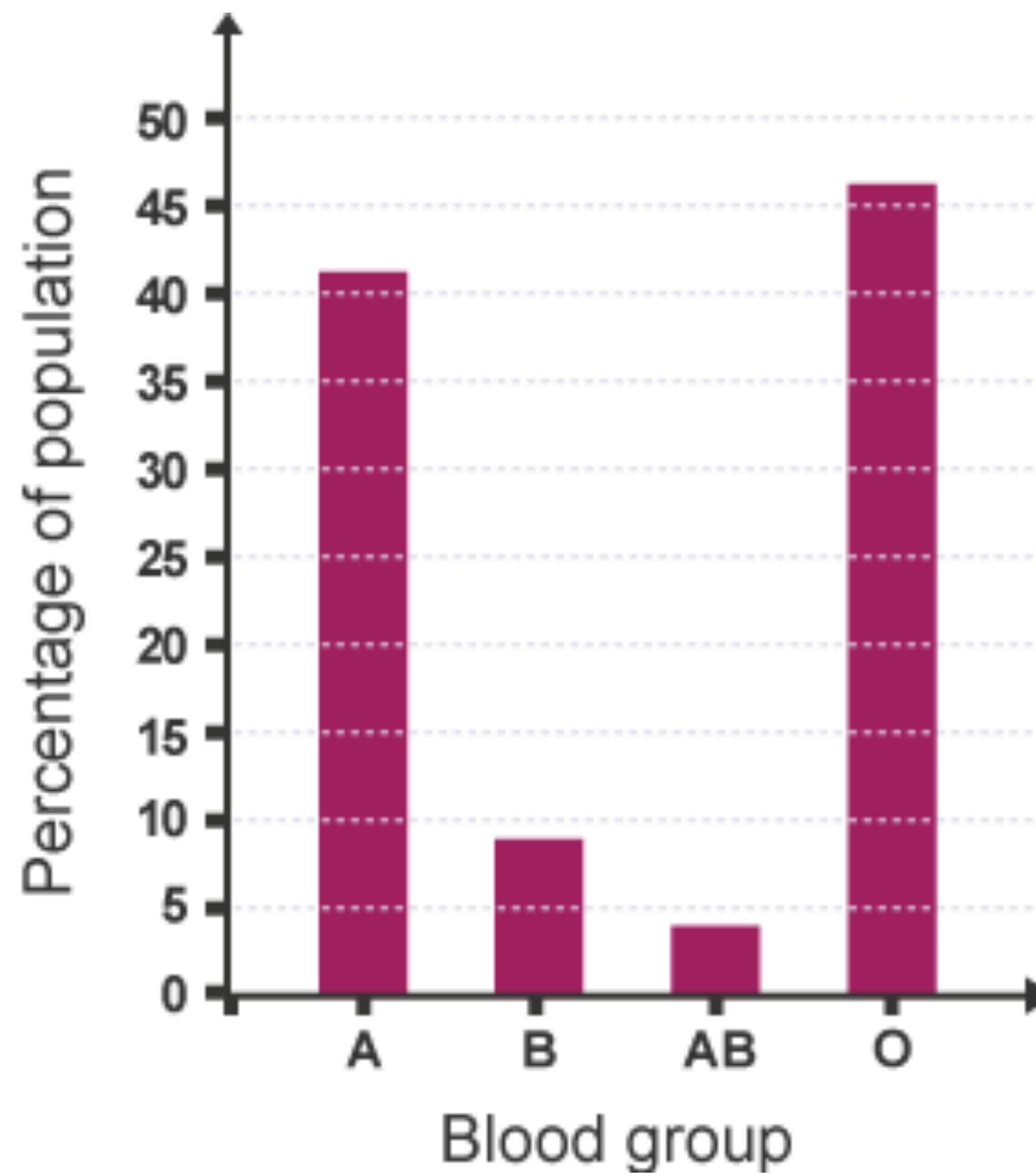
B. de Bivort



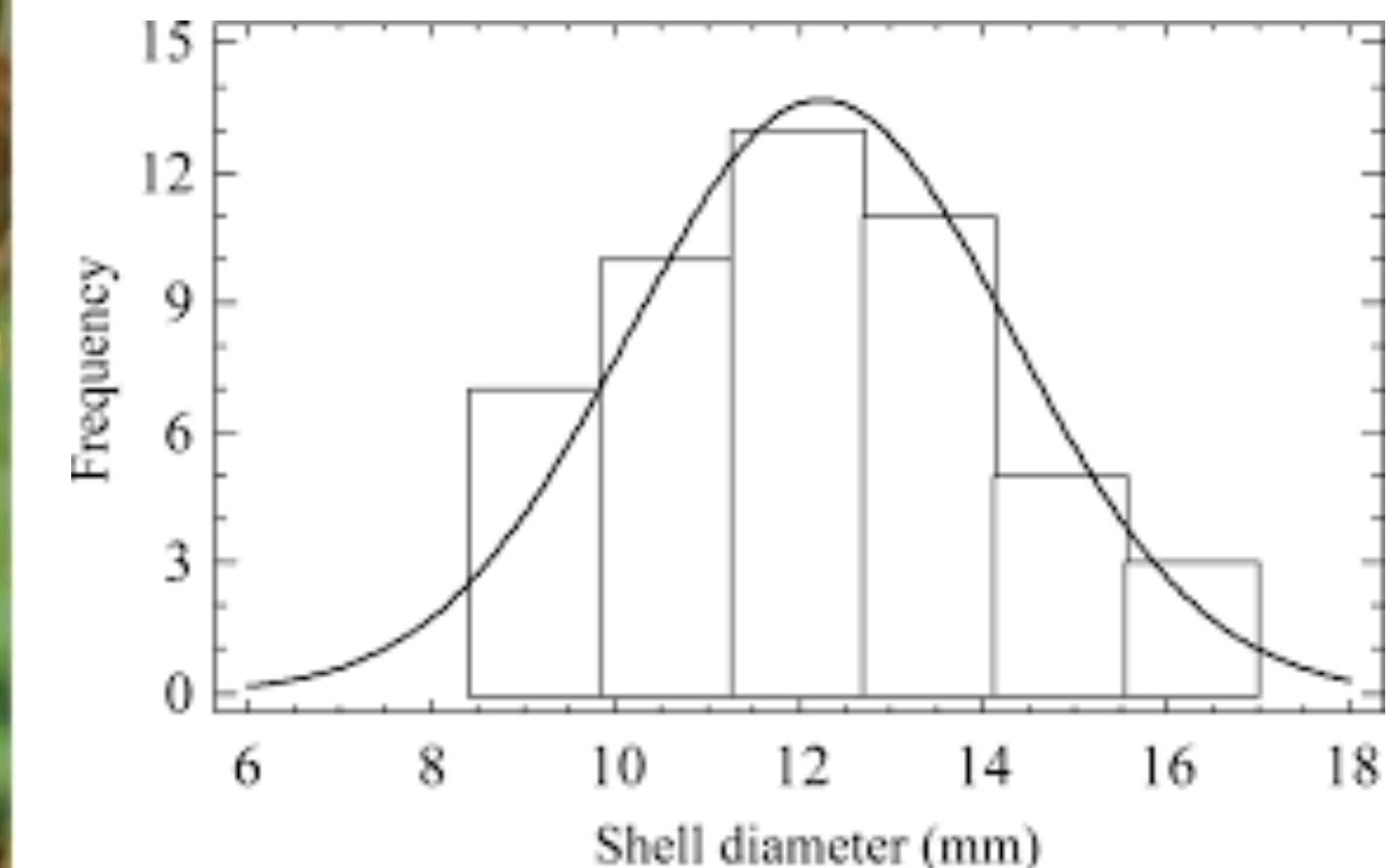
M. Tobler; Boag & Grant 1984

# Measuring and quantifying traits

## Discrete



## Continuous



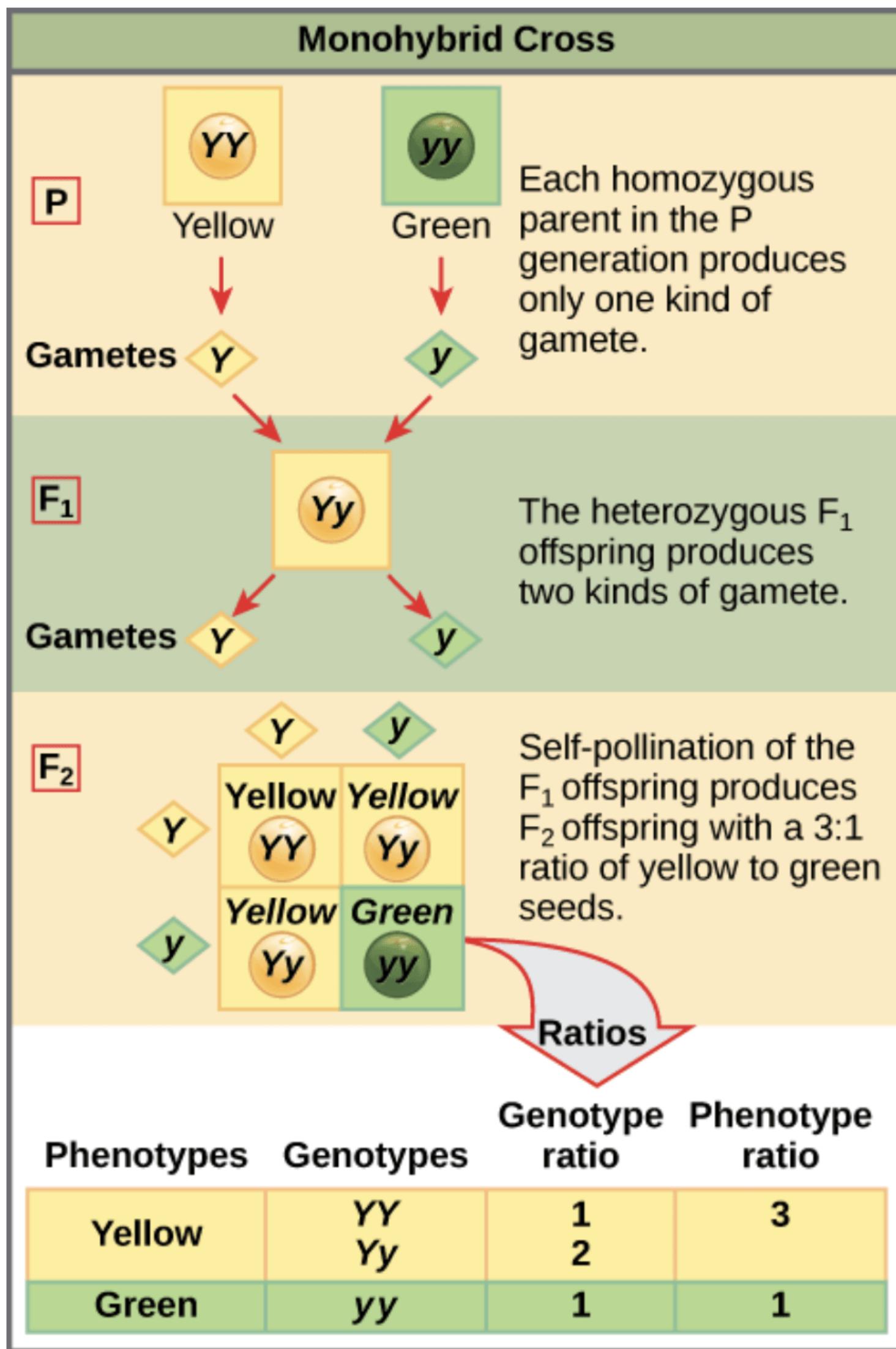
# Measuring and quantifying traits

## **Importance of understanding quantitative trait variation**

- Plant and animal breeding
- Predicting risk for human disease traits
- Phenotypic evolution in natural populations

**Genes -> Phenotypes**

# A historical controversy



## The “Mendelians”

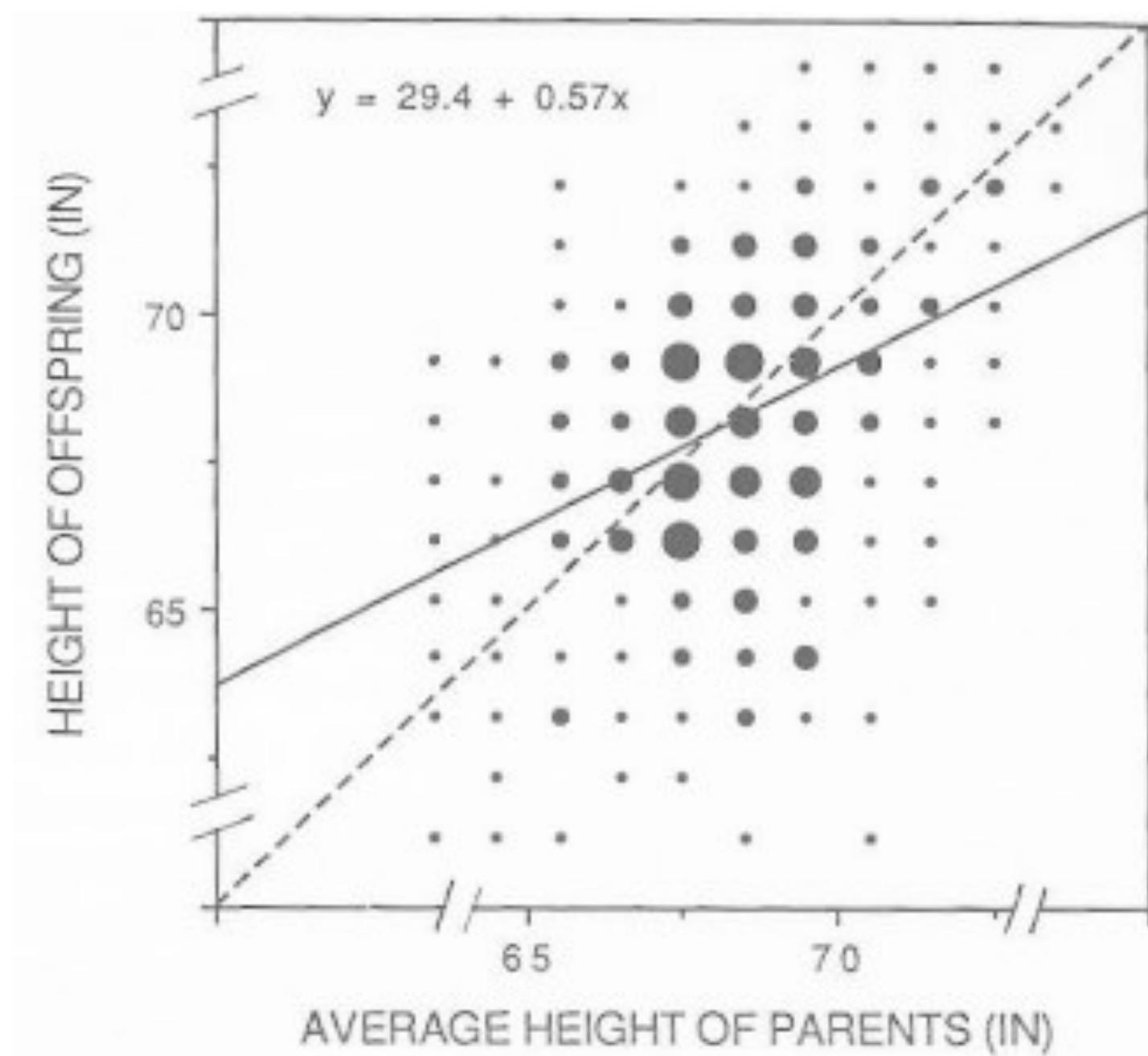
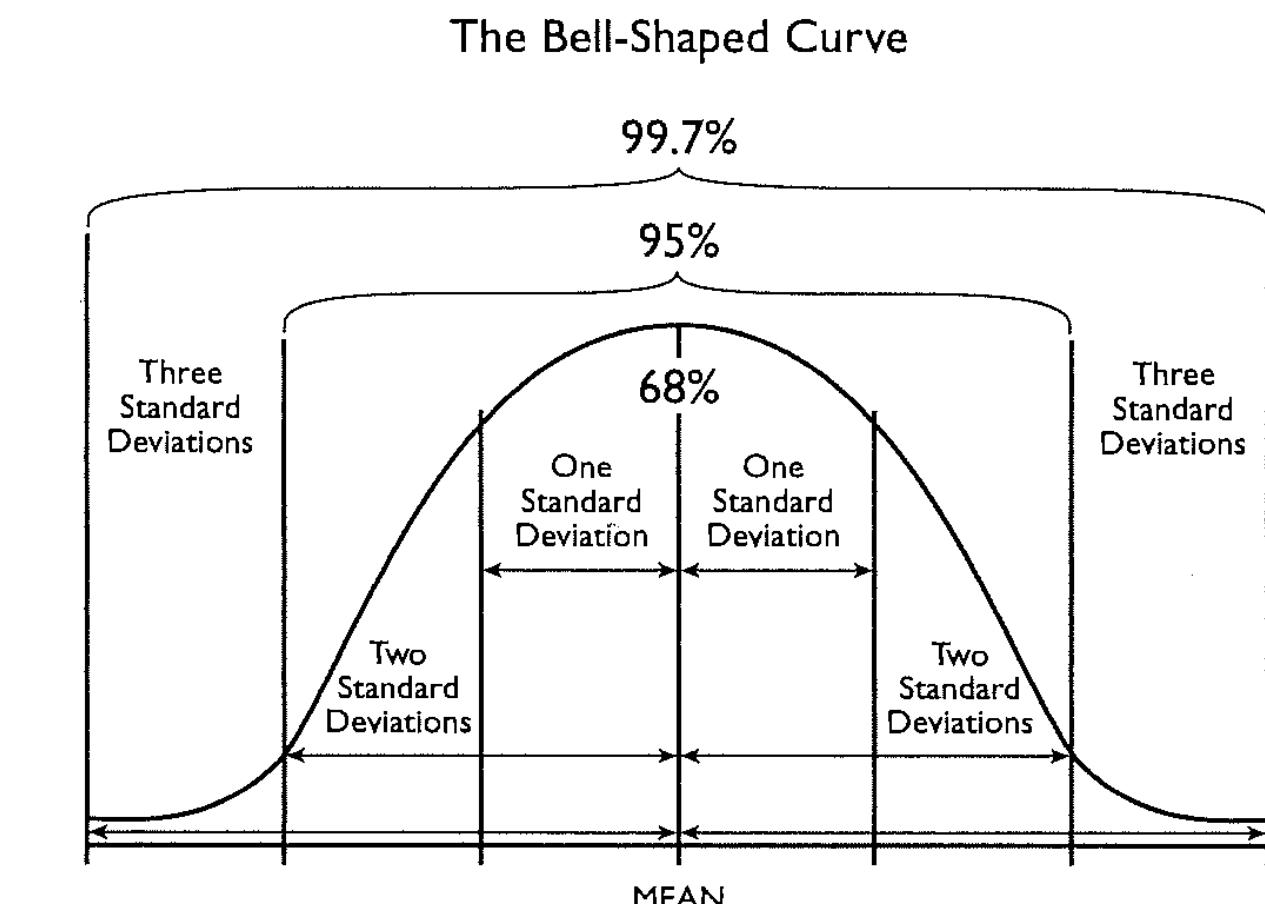
discrete genetic factors ->  
discrete phenotypes

evolution: large steps

? genetic factors ->  
continuous phenotypes

evolution: gradual

## The “Biometricians”



# A historical controversy (reconciled)

## Moving toward synthesis: “The multiple-factor hypothesis”

1 (diallelic) locus -> **3 genotypes:** AA Aa aa

10 (diallelic) loci ->  $3^{10} \approx 60,000$  **genotypes**

Also: Non-genetic (environmental) factors add continuous variation



# Simulating the "multiple-factor hypothesis" in R

Can we simulate a  $\approx$  Normal distribution for 500 Idaho giant salamander tail length measurements, from discrete genotypes at 5 loci?

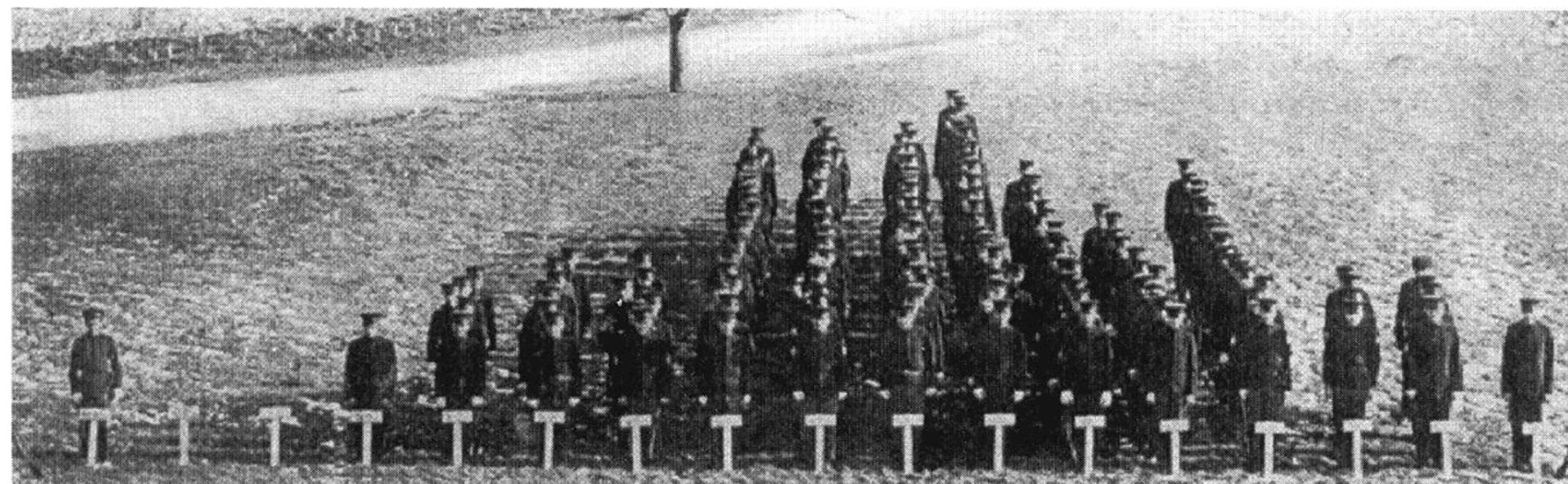
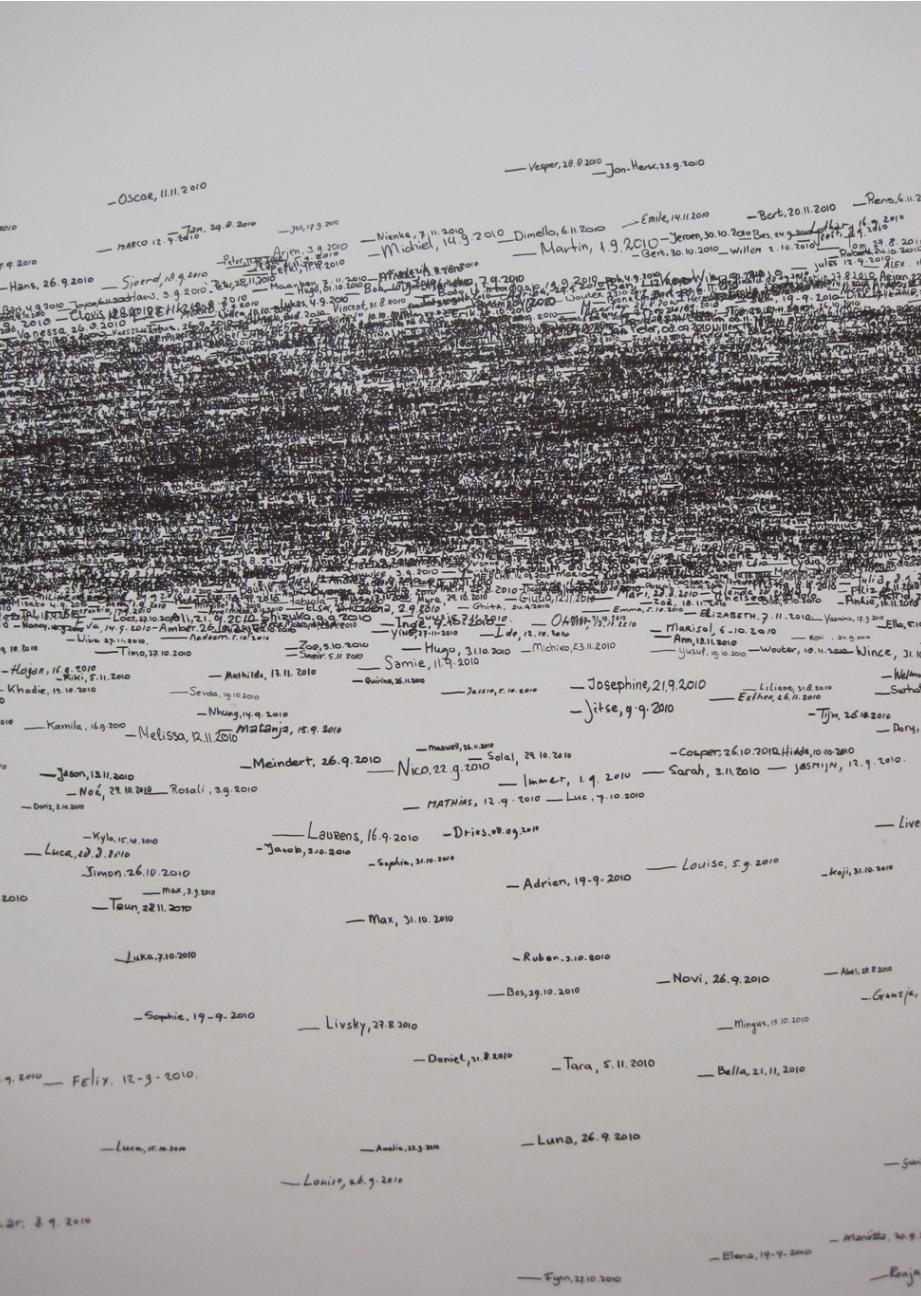


J. Cossel Jr.

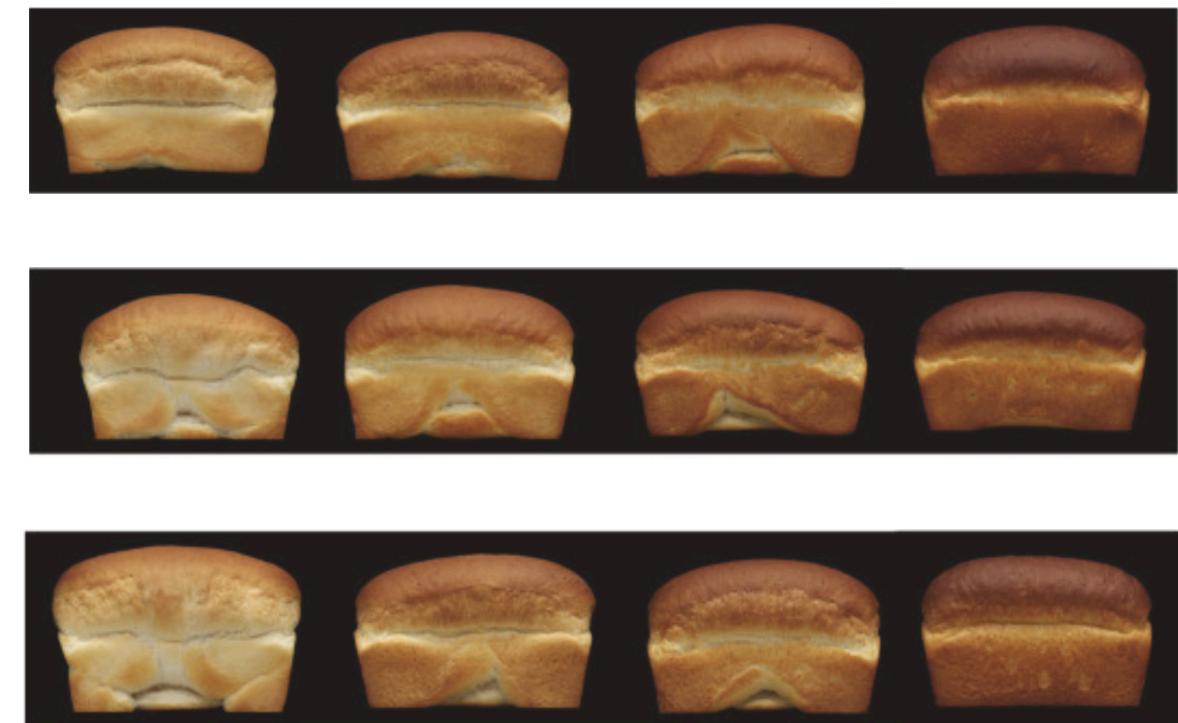
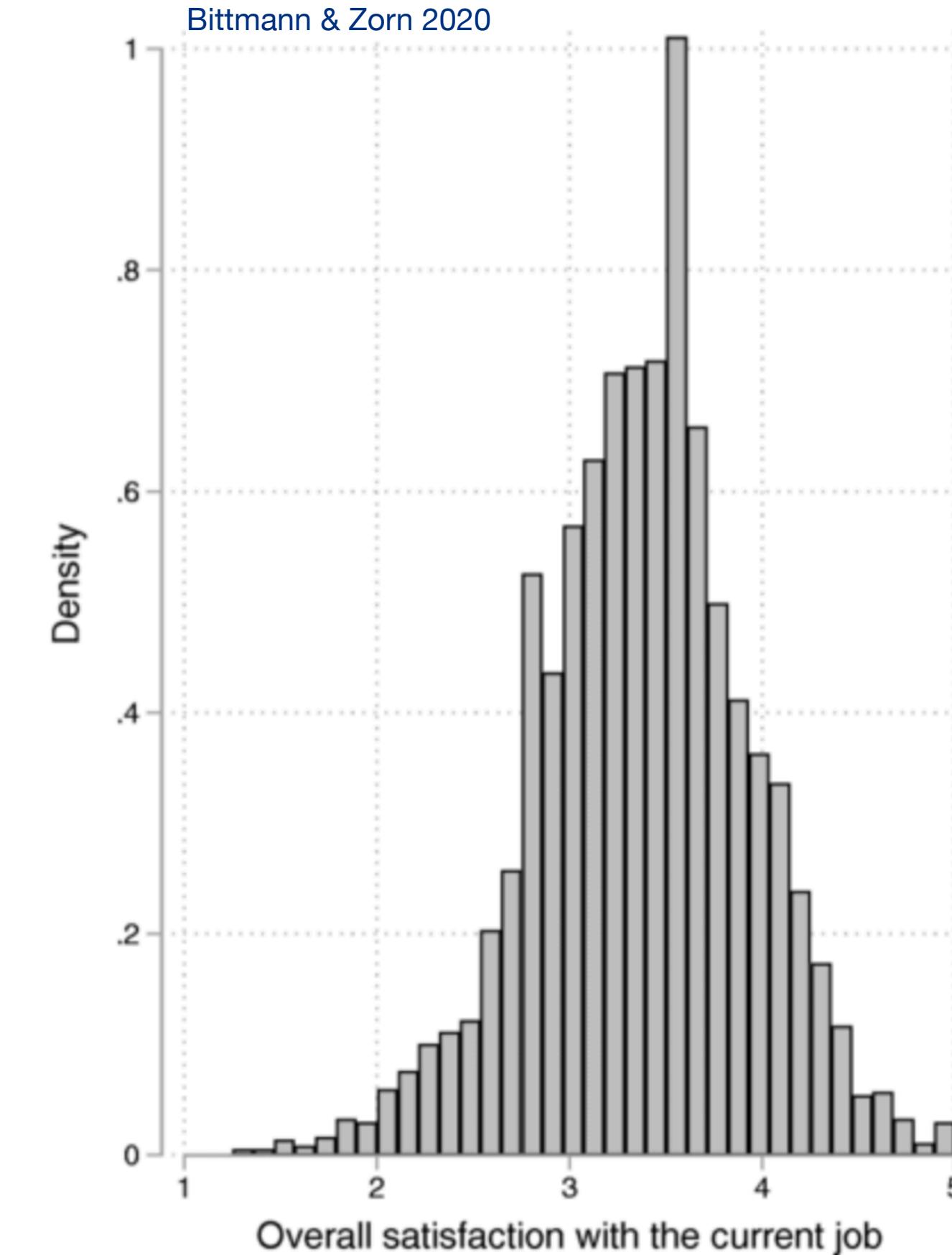
What “ingredients” do we need for our simulation?



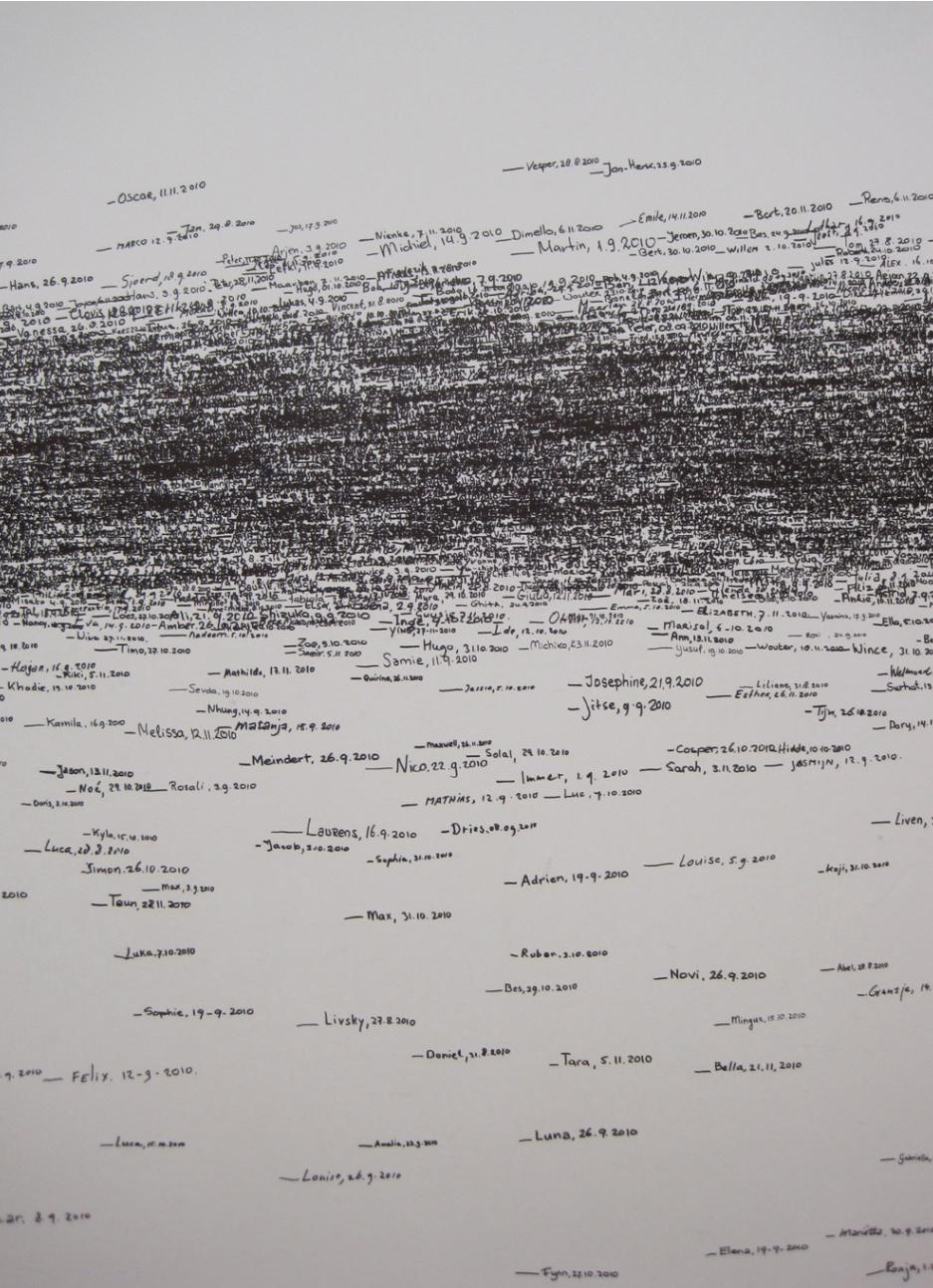
# Why are quantitative traits often normally distributed?



**Non-biological variables are, too!**



# Why are quantitative traits often normally distributed?



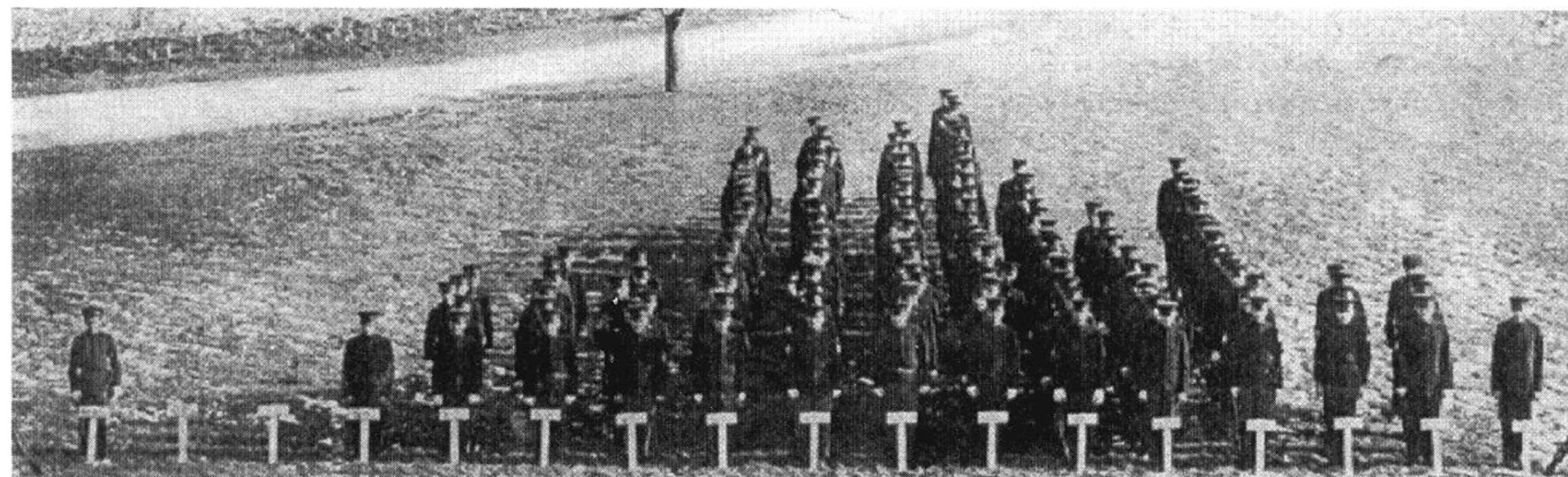
“fuzzy analogue” of the **central limit theorem**:

Variables which are, themselves, influenced by the sum of many, independent underlying variables tend to be normally distributed.

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

$\mu$  (“mu”) = mean  
 $\sigma$  (“sigma”) = standard deviation

$$\boxed{V_P} = V_G + V_E + V_{GxE}$$
$$\quad \quad \quad \underbrace{V_A + V_D + V_I}$$



4:10 4:11 5:0 5:1 5:2 5:3 5:4 5:5 5:6 5:7 5:8 5:9 5:10 5:11 6:0 6:1 6:2

# Summary and questions for further consideration

## **Quantitative Traits | Inheritance | Simulation of CLT**

- What assumptions does our simulation make?
- How might we introduce environmental contributions?
- How might we include dominance in our simulation?

