

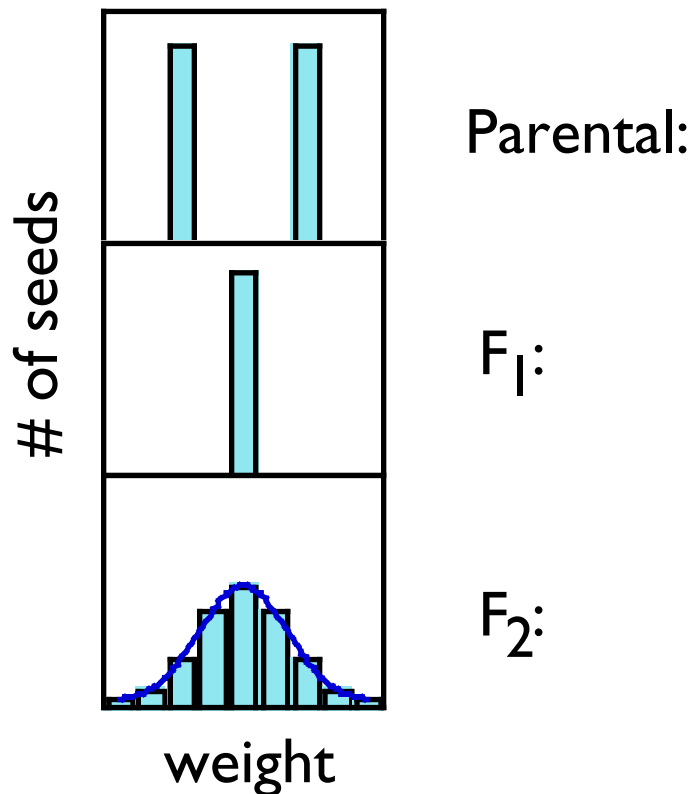
Quantitative genetics

Genetics 371B Lecture 32

30 Nov. 1999

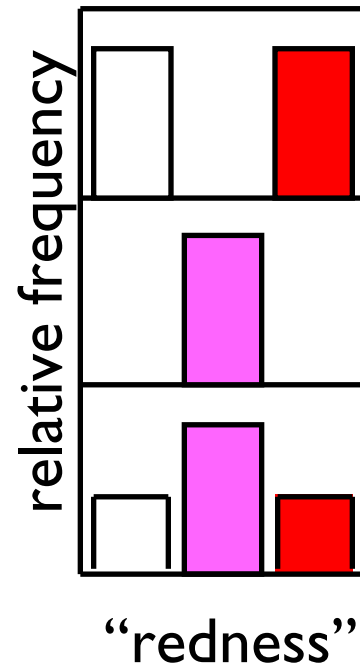
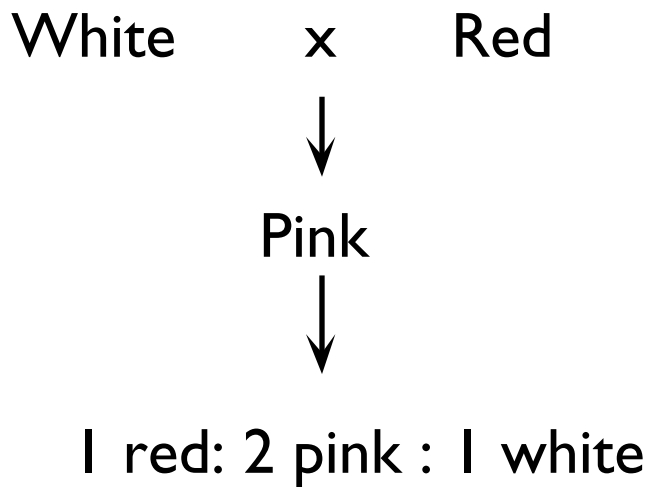
Many traits don't behave in a simple Mendelian fashion

e.g., seed weight



Explanation:

Reminder: Snapdragon flower color inheritance
(lecture 3)



Basal level:

One increment of color:

Two increments:

Additive or **contributing** allele:

Non-additive or **non-contributing** allele:

Suppose there are **two genes** contributing to color?
Locus **A/a** and locus **B/b**

How many possible genotypes?

Non-additive alleles: **a, b**

Basal level = no additive alleles =

One additive allele:

Two additive alleles:

Three additive alleles:

Four additive alleles:

Looking at a cross...

white x fully red

aabb x **AABB**

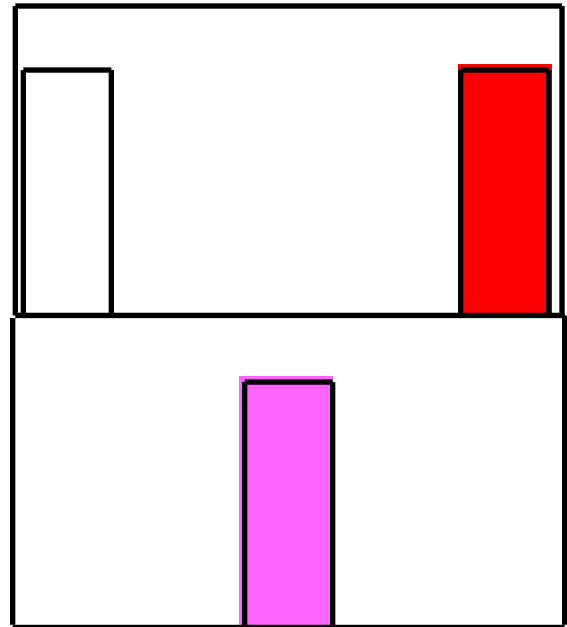


AaBb **Pink**

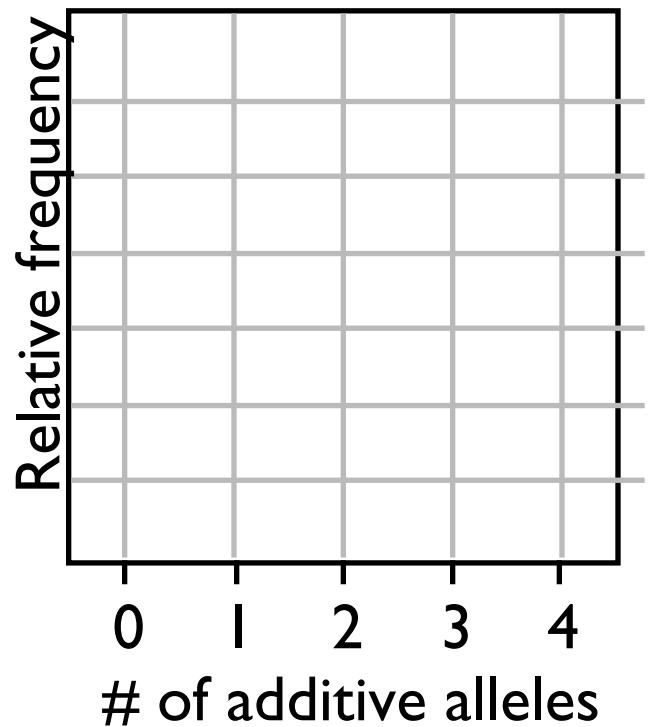
F₁ x F₁



Plot the number of additive alleles



	AB	Ab	aB	ab
AB				
Ab				
aB				
ab				



of genes = 2

of alleles = 4

of phenotypes =

distribution of additive allele frequencies:

fraction exhibiting extreme phenotype =

In general:

- ◆ # of genes:
- ◆ # of alleles
- ◆ # of phenotypes:
- ◆ distribution of additive allele frequencies:
- ◆ fraction exhibiting extreme phenotype:

Some assumptions:

Determining the number of polygenes (n):

1. Obtain true-breeders
2. Make F_1 . Phenotype:
3. Cross F_1 to generate F_2 . Phenotype:
4. Fraction of F_2 showing either extreme phenotype =

Why study quantitative genetics?

- ◇ Agriculture
- ◇ Human biology and health
- ◇ Studying evolution