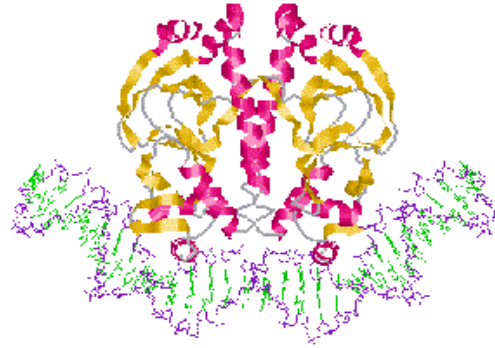




# Punnett Squares, Testcross, Monohybrid, dihybrid and trihybrid cross



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# Outline of course

- Punnett Squares
- Testcross
- Monohybrid
- Dihybrid and trihybrid cross
- Chi-Square Analysis Evaluates the Influence of Chance on Genetic Data

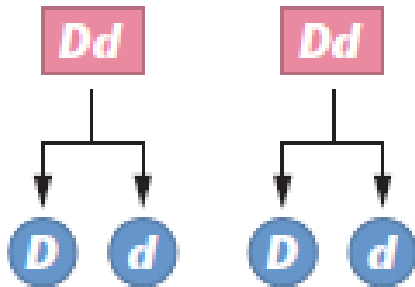
# Punnett Squares

F<sub>1</sub> cross

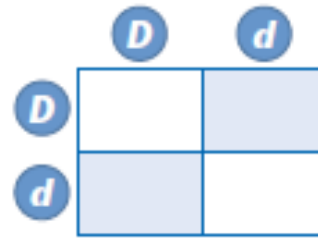
**Dd** × **Dd**  
tall × tall



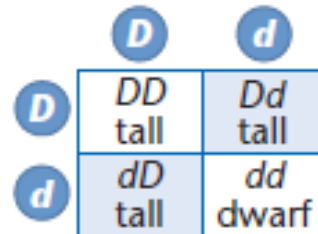
Gamete formation  
by F<sub>1</sub> generation



Setting up a  
Punnett square



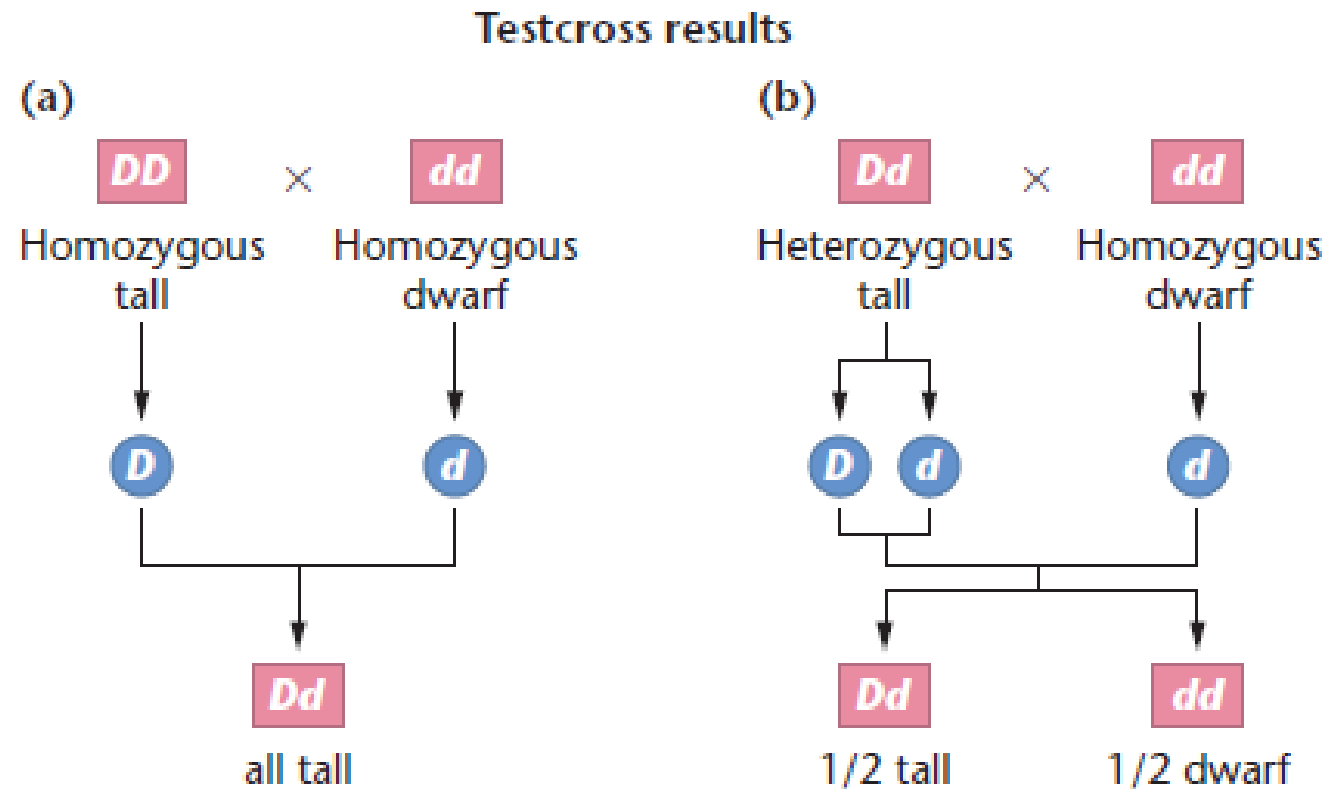
Filling out squares  
representing fertilization



F<sub>2</sub> results

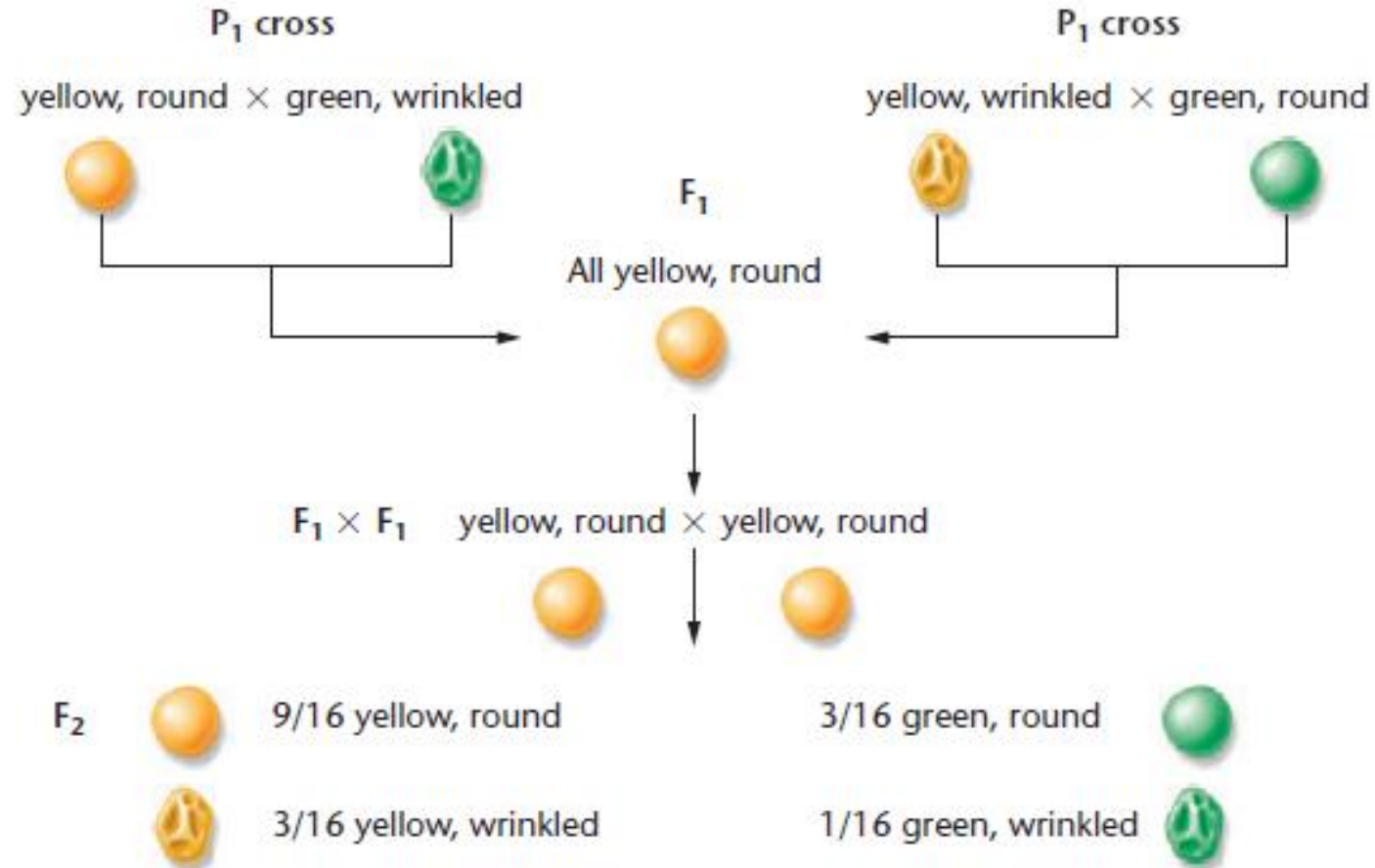
Genotype	Phenotype
1 DD	} 3/4 tall
2 Dd	
1 dd	} 1/4 dwarf
1:2:1	3:1

# The Testcross: One Character

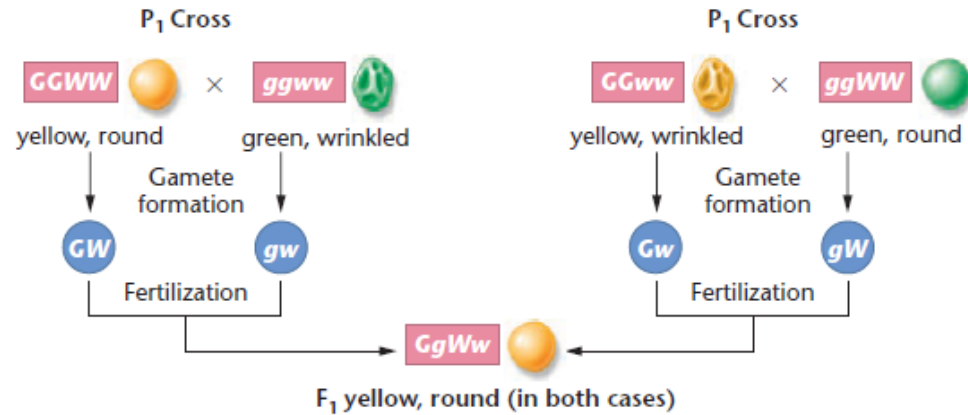


**FIGURE 3.4** Testcross of a single character. In (a), the tall parent is homozygous, but in (b), the tall parent is heterozygous. The genotype of each tall  $P_1$  plant can be determined by examining the offspring when each is crossed with the homozygous recessive dwarf plant.

# Mendel's Dihybrid Cross Generated a Unique F<sub>2</sub> Ratio



# Mendel's Dihybrid Cross Generated a Unique F2 Ratio



F<sub>2</sub> Genotypic ratio

1/16 GGWW  
 2/16 GGWw  
 2/16 GgWW  
 4/16 GgWw

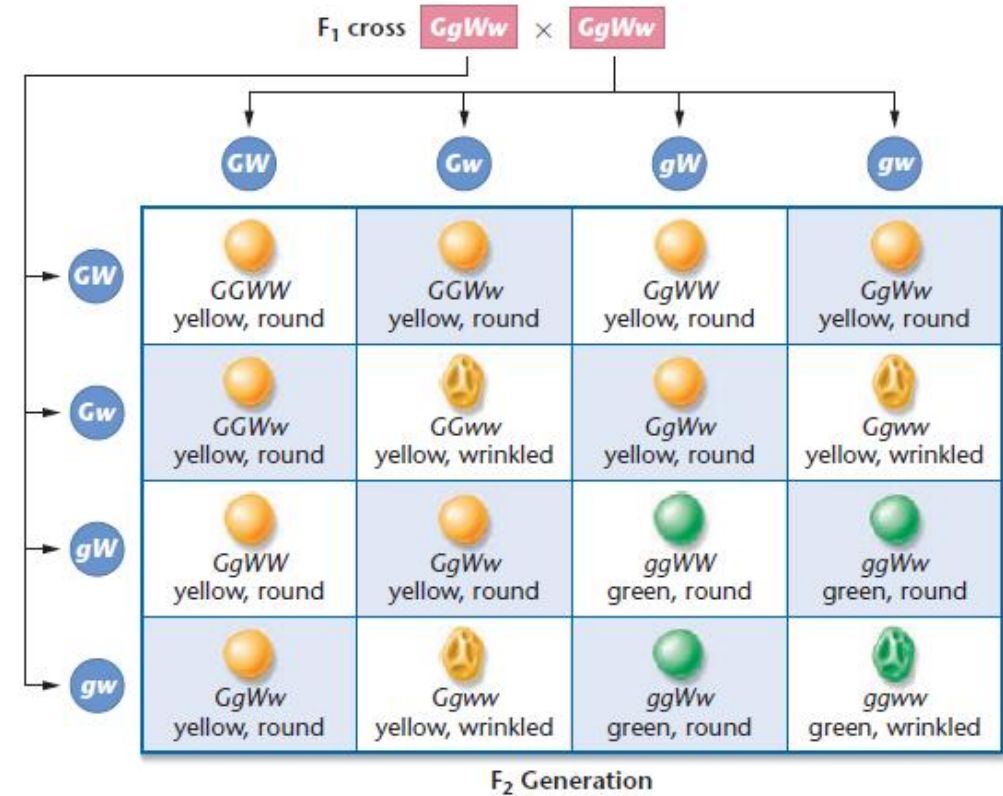
F<sub>2</sub> Phenotypic ratio

9/16 yellow, round

3/16 yellow, wrinkled

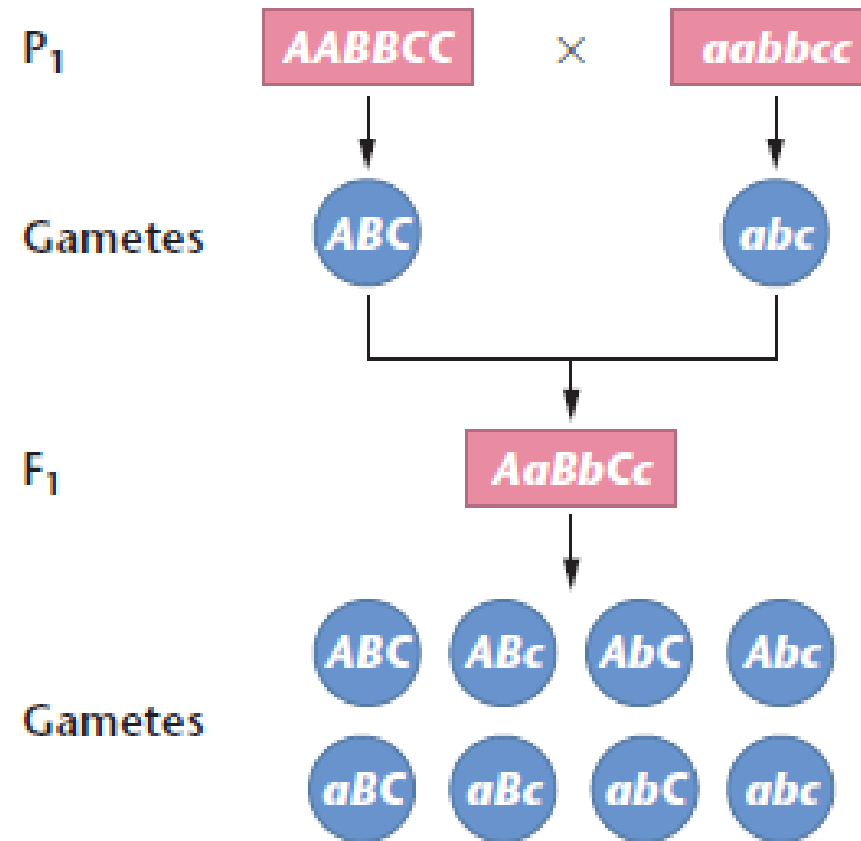
3/16 green, round

1/16 green, wrinkled



# Triybrid gamete formation

## Trihybrid gamete formation



**FIGURE 3.8** Formation of P<sub>1</sub> and F<sub>1</sub> gametes in a trihybrid cross.

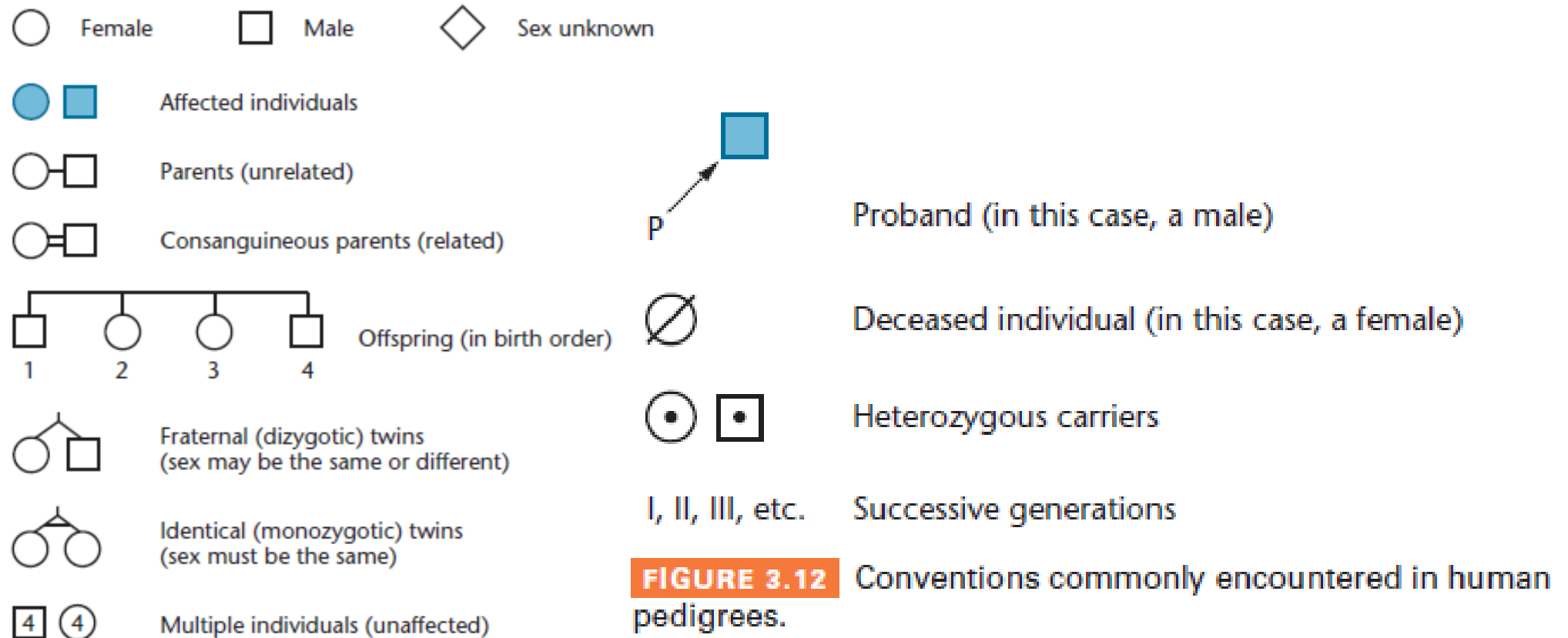
# Generation of F<sub>2</sub> trihybrid phenotypes

## Generation of F<sub>2</sub> trihybrid phenotypes

A or a	B or b	C or c	Combined proportion
3/4 A	3/4 B	3/4 C →	(3/4)(3/4)(3/4) ABC = 27/64 ABC
		1/4 c →	(3/4)(3/4)(1/4) ABc = 9/64 ABc
	1/4 b	3/4 C →	(3/4)(1/4)(3/4) AbC = 9/64 AbC
		1/4 c →	(3/4)(1/4)(1/4) Abc = 3/64 Abc
1/4 a	3/4 B	3/4 C →	(1/4)(3/4)(3/4) aBC = 9/64 aBC
		1/4 c →	(1/4)(3/4)(1/4) aBc = 3/64 aBc
	1/4 b	3/4 C →	(1/4)(1/4)(3/4) abC = 3/64 abC
		1/4 c →	(1/4)(1/4)(1/4) abc = 1/64 abc



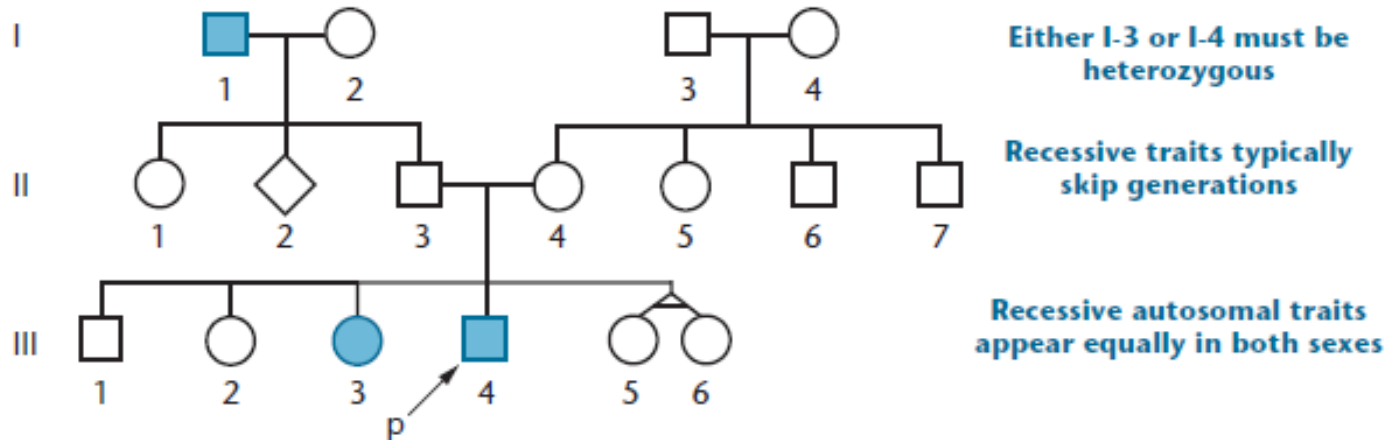
# Pedigrees Reveal Patterns of Inheritance of Human Traits



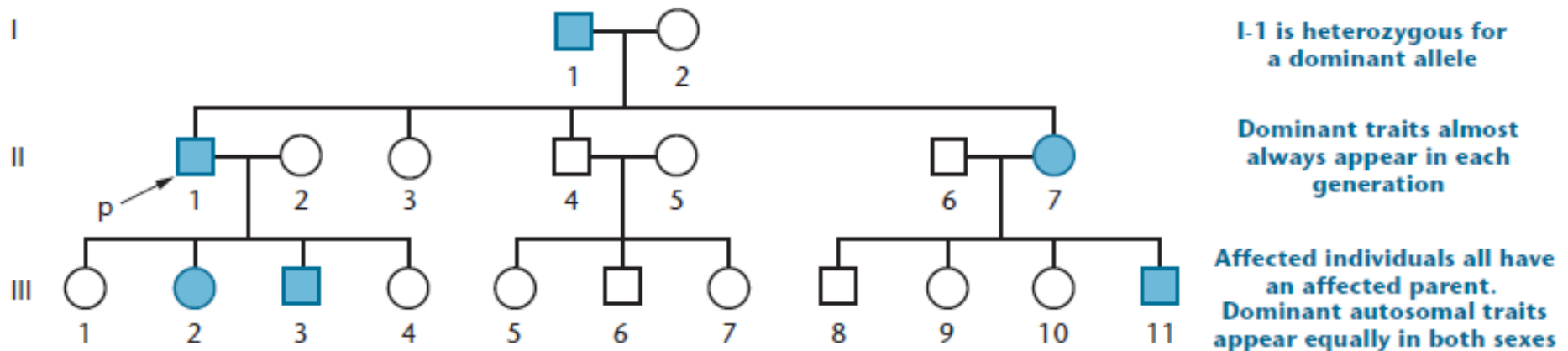
**FIGURE 3.12** Conventions commonly encountered in human pedigrees.

# Pedigrees Reveal Patterns of Inheritance of Human Traits

(a) Autosomal Recessive Trait



(b) Autosomal Dominant Trait



**FIGURE 3.13** Representative pedigrees for two characteristics, each followed through three generations.



# Chi-Square Analysis

**TABLE 3.1** Chi-Square Analysis

(a) Monohybrid					
Cross Expected Ratio	Observed ( <i>o</i> )	Expected ( <i>e</i> )	Deviation ( <i>o</i> - <i>e</i> = <i>d</i> )	Deviation <sup>2</sup>	<i>d</i> <sup>2</sup> / <i>e</i>
3/4	740	3/4(1000) = 750	740 - 750 = -10	(-10) <sup>2</sup> = 100	100/750 = 0.13
1/4	<u>260</u>	1/4(1000) = 250	260 - 250 = +10	(+10) <sup>2</sup> = 100	100/250 = <u>0.40</u>
	Total = 1000				$\chi^2 = 0.53$ <i>p</i> = 0.48
(b) Dihybrid					
Cross Expected Ratio	Observed ( <i>o</i> )	Expected ( <i>e</i> )	Deviation ( <i>o</i> - <i>e</i> = <i>d</i> )	Deviation <sup>2</sup>	<i>d</i> <sup>2</sup> / <i>e</i>
9/16	587	567	+20	400	0.71
3/16	197	189	+8	64	0.34
3/16	168	189	-21	441	2.33
1/16	<u>56</u>	63	-7	49	<u>0.78</u>
	Total = 1008				$\chi^2 = 4.16$ <i>p</i> = 0.26

# Chi-Square Analysis

