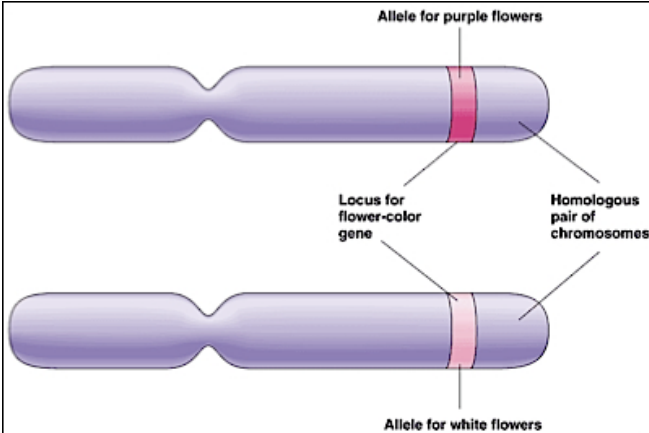


**Mendelian Genetics (Inheritance)**

keywords/questions	notes	extra info	i get it
	<b><u>I. Background</u></b>		<input type="checkbox"/>
	<b>A. Gregor Mendel (1822-1884. He was a monk.</b>		<input type="checkbox"/>
	1. Discoveries in 1865 --> credit around 1900		<input type="checkbox"/>
	2. worked with pea plants		<input type="checkbox"/>
	3. distinguished varieties of peas		<input type="checkbox"/>
	4. established "true breeding" strains		<input type="checkbox"/>
	5. worked on cross-breeding AKA hybridization		<input type="checkbox"/>
	<b><u>II. Mendel's Experiments - many on pea traits</u></b>		<input type="checkbox"/>
	<b>A. example of experiment -</b>		<input type="checkbox"/>
	1. Cross: Parental (P) generation: Round seed X wrinkled seed		<input type="checkbox"/>
	2. results: First filial generation (F1) --> all round! ohsh!		<input type="checkbox"/>
	3. then Mendel self fertilized some F1 plants. (F1 X F1)		<input type="checkbox"/>
	4. Results : F2 generation, 5474 round; 1850 wrinkled , a 2.96:1 ratio.		<input type="checkbox"/>
	5. the crazy part is that Mendel repeated many many true breeding crosses, then F1 crosses and got a generalizable 3:1 of "dominant" to "recessive" in the F2 gen.		<input type="checkbox"/>
	<b>B. Mendel's explanation for his results.</b>		<input type="checkbox"/>
	1. for each trait, each organism has 2 genes (this is not always true, especially in animals like humans; new discovery 11/2006 that shows that some humans have multiple copies of each certain genes, and other humans have none. Very unexpected...)		<input type="checkbox"/>

keywords/questions	notes	extra info	i get it
	<p>2. Alternative forms of the same gene are called <b>alleles</b></p> 		<input type="checkbox"/>
	<p>a. For example: gene for pea color, then a green allele and yellow allele.</p>		<input type="checkbox"/>
	<p>3. If alleles are the same in both chromosomes --&gt; "<b>homozygous</b>" for trait, (green/green)</p>		<input type="checkbox"/>
	<p>4. If alleles are diff. --&gt; "<b>heterozygous</b>" (green/yellow)</p>		<input type="checkbox"/>
	<p>5. If organism is heterozygous, the allele that "appears" or that is expressed is called the <b>dominant</b> allele. Usually represented by capital letter symbol (expressed = phenotype)</p>		<input type="checkbox"/>
	<p>6. Allele that is hidden is called <b>recessive</b> allele</p>		<input type="checkbox"/>
	<p>7. What alleles code for = genotype</p>		<input type="checkbox"/>
	<p>8. What alleles show up = phenotype</p>		<input type="checkbox"/>
	<p><b>III. When organism makes gametes, alleles separate, which is called segregation</b></p>		<input type="checkbox"/>
	<p>A. only one allele in each gamete (remember <math>2n \rightarrow n</math> in meiosis!)</p>		<input type="checkbox"/>
	<p>B. From a heterozygote, 50% of gametes will be dom. and 50% will be recessive.</p>		<input type="checkbox"/>
	<p>C. Hey, then why is there a 3:1 ratio in the F2 of a heterozygote cross?</p>		<input type="checkbox"/>
	<p>1. Use a punnet square to figure it out:</p>		<input type="checkbox"/>
	<p>a. How to use punnett squares:</p>		<input type="checkbox"/>
	<p>i. write out genotypes of parents</p>		<input type="checkbox"/>

keywords/questions	notes	extra info	i get it
	ii. write out gametes		<input type="checkbox"/>
	iii. combine gametes		<input type="checkbox"/>
	iv. analyze!		<input type="checkbox"/>
	<b>IV. Types of Mendelian crosses</b>		<input type="checkbox"/>
	<b>A. monohybrid cross - one trait, one gene pair. (P=purple; p= white)</b>		<input type="checkbox"/>
	1. Can use monohybrid cross if you don't know a genotype of individual with dom phenotype. Breed it with a homozygous rec indiv. and look at the offspring results. Called a test cross.		<input type="checkbox"/>
	<b>B. Dihybrid cross - 2 traits, 2 genes, 4 possible alleles.</b>		<input type="checkbox"/>
	1. F1 heterozygote cross results in F2 gen with 9:3:3:1 ratio. Dom/Dom:Dom/Rec: Rec/Dom : Rec/Rec		<input type="checkbox"/>
	2. Why does this crazy ratio happen? The answer is the principle of <b>independent assortment</b> --> One allele does not affect the others.		<input type="checkbox"/>
	<b>C. Using a testcross to find unknown genotypes.</b>		<input type="checkbox"/>
	1. only works for things that can be mated with each other.		<input type="checkbox"/>
	2. requires a full generation to get results.		<input type="checkbox"/>
	3. How to do it:		<input type="checkbox"/>
	a. get unkown genotype and mate it with known homozygous recessive individual (known because recent ancestors were also homo. rec.)		<input type="checkbox"/>
	b. look at offspring:		<input type="checkbox"/>
	i. if phenotype is homo. dom., then offspring should be _____		<input type="checkbox"/>
	ii. if phenotype is hetero., then offspring should be _____		<input type="checkbox"/>
	<b>D. Mendelian Genetics and Probability</b>		<input type="checkbox"/>
	1. probability is the chance that something occurs.		<input type="checkbox"/>

keywords/questions	notes	extra info	i get it
	2. <b>rule of multiplication</b> - used when two events are independent from each other. like sex of child, or coin flips.		<input type="checkbox"/>
	3. <b>rule of addition</b> - used when there is more than one way to get the same result. usually when heterozygotes and homo. dom. are possible. It is the sum of different probabilities.		<input type="checkbox"/>
	4. <b>How to solve specific stuff with out punnet squares!</b>		<input type="checkbox"/>
	a. Multiply probabilities!!!!!!		<input type="checkbox"/>
	i. e.g. What is the probability you get AaBBCCDd from AaBbCcDd X AaBbCcDd ?		<input type="checkbox"/>
	I prob of Aa = 1/2		<input type="checkbox"/>
	II prob of BB = 1/4		<input type="checkbox"/>
	III prob of CC = 1/4		<input type="checkbox"/>
	IV prob of Dd = 1/2		<input type="checkbox"/>
	ii. so, to solve 1/2 times 1/4 times 1/4 times 1/2 equals 1/64! it is not like wow, it is like whoa!		<input type="checkbox"/>
	<b><u>V. Pedigrees track traits across generations.</u></b>		<input type="checkbox"/>
	A. Rules for pedigrees. carrier = heterozygote.		<input type="checkbox"/>

keywords/questions	notes	extra info	i get it
	<b><u>VI. Single gene Autosomal traits in humans</u></b>		<input type="checkbox"/>
	<b>A. recessive</b>		<input type="checkbox"/>
	1. albinism		<input type="checkbox"/>
	2. cystic fibrosis		<input type="checkbox"/>
	3. galactosemia		<input type="checkbox"/>
	4. phenylketonuria		<input type="checkbox"/>
	5. sickle cell disease (aka sickle cell anemia)		<input type="checkbox"/>
	6. Tay-Sachs		<input type="checkbox"/>
	<b>B. Dominant</b>		<input type="checkbox"/>
	1. achondroplasia		<input type="checkbox"/>
	2. huntington's disease		<input type="checkbox"/>
	<b>C. hypercholesterolemia</b>		<input type="checkbox"/>
	<b><u>VII. NON-Dominant/Recessive Genetics</u></b>		<input type="checkbox"/>
	<b>A. Incomplete Dominance</b>		<input type="checkbox"/>
	1. Hybrids have phenotype that is between parents phenotypes.		<input type="checkbox"/>
	a. red + white = pink		<input type="checkbox"/>
	b. in humans - hypercholesterolemia		<input type="checkbox"/>
	<b>B. Codominance - both show up.</b>		<input type="checkbox"/>
	1. in humans - Blood groups		<input type="checkbox"/>

keywords/questions	notes	extra info	i get it
	<b>C. Pleiotropy - when one gene affects many traits.</b>		<input type="checkbox"/>
	1. sickle cell gene		<input type="checkbox"/>
	<b>D. Polygenic traits - many genes, one phenotype</b>		<input type="checkbox"/>
	1. skin color		<input type="checkbox"/>
	<b>E. Linked Genes - genes on same chromosome. Do not assort independently</b>		<input type="checkbox"/>
	1. linkage maps.		<input type="checkbox"/>
	2. Morgan's Drosophila		<input type="checkbox"/>
	<b>F. Sex Linkage - genes on X chromosome. A sex-linked trait has a gene on the X chromosome.</b>		<input type="checkbox"/>
	1. red - green colorblindness		<input type="checkbox"/>
	2. hemophilia		<input type="checkbox"/>
	3. Duchenne Muscular Dystrophy		<input type="checkbox"/>