Concept 14.4: Many human traits follow Mendelian patterns of inheritance

- Humans are not good subjects for genetic research
  - Generation time is too long
  - Parents produce relatively few offspring
  - Breeding experiments are unacceptable
- However, basic Mendelian genetics endures as the foundation of human genetics
Pedigree Analysis

- A **pedigree** is a family tree that describes the interrelationships of parents and children across generations
- Inheritance patterns of particular traits can be traced and described using pedigrees
• Pedigrees can also be used to make predictions about future offspring
• We can use the multiplication and addition rules to predict the probability of specific phenotypes
Sample Pedigree
<table>
<thead>
<tr>
<th>Term</th>
<th>Definition</th>
</tr>
</thead>
<tbody>
<tr>
<td>locus</td>
<td></td>
</tr>
<tr>
<td>codominant</td>
<td></td>
</tr>
<tr>
<td>compound heterozygote</td>
<td></td>
</tr>
<tr>
<td>allele</td>
<td>dominant</td>
</tr>
<tr>
<td>carrier (obligate heterozygote)</td>
<td></td>
</tr>
<tr>
<td>genotype</td>
<td>recessive</td>
</tr>
<tr>
<td>genetic heterogeneity</td>
<td></td>
</tr>
<tr>
<td>phenotype</td>
<td>homozygous</td>
</tr>
<tr>
<td>pleiotropy</td>
<td></td>
</tr>
<tr>
<td>autosomal</td>
<td>heterozygous</td>
</tr>
<tr>
<td>age of onset</td>
<td></td>
</tr>
<tr>
<td>X-linked</td>
<td>hemizygous</td>
</tr>
<tr>
<td>sex-limited</td>
<td></td>
</tr>
<tr>
<td>penetrance</td>
<td>expressivity</td>
</tr>
<tr>
<td>sex-influenced</td>
<td></td>
</tr>
<tr>
<td>pedigree</td>
<td>proband</td>
</tr>
<tr>
<td>imprinting</td>
<td></td>
</tr>
<tr>
<td>trinucleotide repeat</td>
<td></td>
</tr>
</tbody>
</table>
A **pedigree** is a concise summary of the medical family history; it is the symbolic language of clinical genetics and human genetics research.

- It is an easy, fast, and efficient means of recording a wealth of information about the family.

- Standardization of symbols is essential to facilitate communication - See Robin Bennett’s article referenced in resources at the end of the syllabus for more details if interested.

- Nomenclature is an evolving process.

- Several ethical and legal dilemmas - Potential for discrimination, issues of privacy raised, and need for guidelines.
Designation of generations and individuals

1. Each horizontal line is a generation
2. Place the oldest generation at the top
3. Use Roman numerals to identify generations
4. Use Arabic numbers to identify individuals within a generation
5. List siblings from oldest to youngest, from left to right
6. Male partner is usually placed to the left of the female partner
7. Record full name, current age and date of birth, or age at death for each individual
8. Record race and ethnic origin of each individual
9. Note health problems and/or cause of death for each individual
10. There are appropriate symbols to use for both adoption and assisted-reproductive technologies
• The proband is an affected individual coming to medical attention independently of other family members. The proband is designated with an arrow in the pedigree, and there may be more than one proband per family.
Medical status and results of genetic evaluation/testing of family members

1. Shading or fill (hatches, dots, etc.) is used to denote medical status or symptoms of individuals. A key/legend is used to define meaning.

2. Results of an evaluation (E) are recorded below the symbol and a key/legend defines the notations. Currently this is the least standardized pedigree nomenclature.
PEDIGREE NOMENCLATURE

Adapted from Bennett RL et al. (1995) AJHG 56:745-752.

- Male
- Female
- Sex Unspecified
- Number of children of sex indicated
- Affected
- Heterozygotes for autosomal trait
- Carrier of X-linked recessive trait
- Adopted out of a family
- Relationship
- Relationship that no longer exits
- Consanguineous mating
- Monozygotic twins
- Dizygotic twins
The Gene is the Unit of Inheritance

The location of a gene on a chromosome is its **locus**.

Alternative forms of a gene at a particular locus are referred to as **alleles**.

An individual’s **genotype** (genetic composition) at a particular locus is defined by the nature of the alleles at that locus.

If both alleles are identical, then the individual is **homozygous** at the locus. Homozygosity may refer to the presence of two normal or two mutant alleles.

If the alleles differ, then the individual is **heterozygous** at the locus. If two different mutant alleles are present, then the individual is a **compound heterozygote**.
A   A               A    a              a    a              a
1
2

homozygote     heterozygote      homozygote         compound
A allele                                        a allele             heterozygote

A A
homzygote
A allele

A a
heterozygote

a a
homozygote
a allele

a1 a2
compound
heterozygote
The genotype at a particular locus and the environment in which it is expressed determines the phenotype or observed characteristics of an individual.

Traits that are determined by loci on one of the 22 autosomes are **autosomal**. Traits determined by loci on the X chromosome are **X-linked**, and those determined by loci on the Y chromosome are **Y-linked**.
Gregor Mendel’s Laws of Inheritance

– Law of Unit Inheritance - parental characteristics do not blend because there is a unit of inheritance. Mendel’s “units” are now known as genes or alleles.

– Law of Segregation - the two alleles at a particular locus segregate into different gametes.

– Law of Independent Assortment - alleles at different loci are transmitted independently of each other. Linkage is an exception to this rule.
Dominant and Recessive Inheritance

- Nomenclature: For dominant traits the capital letter (e.g. A) represents the mutant allele and the small letter (e.g. a) represents the normal allele. For recessive traits, the small letter (e.g. a) represents the mutant allele and the capital letter (e.g. A) represents the normal allele.

- **Autosomal dominant traits** are those traits in which the phenotype of the heterozygote and the homozygote for the dominant allele are the same, i.e., Aa and AA have the same phenotype where A=dominant allele. These traits are expressed when only one copy of the dominant allele is present. In practice, if the heterozygote expresses the trait, then the trait is classified as dominant, even if the phenotype of the homozygote (AA) and heterozygote (Aa) are different.

- **Autosomal recessive traits** are those traits in which the phenotype is expressed only if homozygous for the recessive allele, i.e., aa where a=recessive allele. Two copies of the recessive allele are necessary for expression.
Dominant and Recessive Inheritance

- If the heterozygote (AB) has a different phenotype than either of the homozygotes (AA or BB), then the alleles are said to be **codominant**.

- **X-linked dominant traits** are those expressed when either males or females have one copy of the dominant allele, i.e., \( X^A Y \) or \( X^A X^a \) where \( A = \) dominant allele.

- **X-linked recessive traits** are those expressed in males who carry one copy of the recessive allele (i.e., are hemizygous, \( X^a Y \) where \( a = \) recessive allele). Two copies of the recessive allele are generally required for females to express the trait, i.e., \( X^a X^a \).
Types of Genetic Disease

- Chromosomal
- Single gene (Mendelian)
- Multifactorial
- Teratogenic
Examples and Features of Autosomal Dominant Inheritance

A = mutant allele
a = normal allele

Affected individual

Unaffected individual
Examples

- familial hypercholesterolemia
- Huntington disease
- neurofibromatosis type I (NF1)
- myotonic dystrophy
- Marfan syndrome
- achondroplasia
<table>
<thead>
<tr>
<th>DISEASE</th>
<th>CLINICAL FEATURES</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Autosomal Dominant</strong></td>
<td>Note: Key aspects of phenotypic expression or inheritance features are bolded</td>
</tr>
<tr>
<td><strong>HUNTINGTON DISEASE</strong></td>
<td>Progressive loss of brain neurons, dementia, loss of motor control Affects 1/20,000 persons of European descent Late onset, typically between 30-40 years, but may be earlier (See lecture on unstable trinucleotide repeats.)</td>
</tr>
<tr>
<td><strong>MYOTONIC DYSTROPHY</strong></td>
<td>Facial weakness Cataracts Progressive muscular weakness Variable onset Variable expressivity</td>
</tr>
<tr>
<td><strong>NEUROFIBROMATOSIS TYPE I (NFI)</strong></td>
<td>Cafe-au-lait spots (hyperpigmented skin) Lisch nodules (benign growths on the iris) Peripheral nerve tumors Variable expressivity High mutation rate</td>
</tr>
<tr>
<td><strong>FAMILIAL HYPERCHOLESTEROLEMIA,</strong></td>
<td>Arteriosclerosis, xanthomas Heterozygotes: Increased LDL coronary heart disease in middle age Homozygotes: childhood coronary heart disease</td>
</tr>
<tr>
<td><strong>MARFAN SYNDROME (Connective tissue disorder)</strong></td>
<td>Tall stature with long limbs Narrow facies with high, narrow palate Dislocated lenses &amp; myopia Cardiac manifestations, i.e., aortic aneurysm Variable expressivity Pleiotropy</td>
</tr>
<tr>
<td><strong>ACHONDROPLASIA</strong></td>
<td>Short-limbed dwarfism Megaloecephaly Lordosis &amp; Kyphosis 80% new mutations Increased mutations with increasing paternal age</td>
</tr>
</tbody>
</table>


Dominantly Inherited Disorders

- Some human disorders are caused by dominant alleles
- Dominant alleles that cause a lethal disease are rare and arise by mutation
- *Achondroplasia* is a form of dwarfism caused by a rare dominant allele
Figure 14.17

Parents

Dwarf

\[ Dd \]

\[ \times \]

Normal

\[ dd \]

Sperm

\[ D \]

\[ d \]

Eggs

\[ d \]

\[ Dd \]

Dwarf

\[ dd \]

Normal

\[ Dd \]

Dwarf

\[ dd \]

Normal

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Achondroplasia

From www.hopkinsmedicine.org

From www.sciencemuseum.org.uk
Neurofibromatosis Type 1
Neurofibromatosis Type 1
Neurofibromatosis Type 1

Fig. 121-3. — Recklinghausen neurofibromatosis.
Features of Autosomal Dominant Inheritance

1. Vertical transmission – direct transmission from grandparent to parent to child without skipping generations
2. Both sexes affected in 1:1 ratio
3. Both sexes may transmit the trait
4. Heterozygotes much more common than homozygotes
5. May see variable expressivity and variable age of onset
6. Homozygotes usually more seriously affected than heterozygotes
7. May be due to new mutation
8. Gene product is usually a structural (non-enzymatic) protein
Autosomal Dominant Inheritance
(Affected Father)

Parental Gametes

A

a

Maternal Gametes

a

Aa

aa

1Aa: 1aa
A = mutant, a = normal
Transmission probabilities and the use of the Punnett square

1. If one parent has the disorder (assumed to be Aa) and the other does not (aa) then there is a 50% chance that the child will inherit the disorder and a 50% chance that they will not.

2. If both parents have the disorder (assumed to be Aa x Aa) then there is a 75% chance that their children will inherit the disorder, and a 25% chance that they will not.
Examples and Features of Autosomal Recessive Inheritance
Recessively Inherited Disorders

• Many genetic disorders are inherited in a recessive manner
• These range from relatively mild to life-threatening
Examples

- cystic fibrosis
- sickle cell anemia
- Tay-Sachs disease
- Phenylketonuria
- most inborn errors of metabolism
The Behavior of Recessive Alleles

- Recessively inherited disorders show up only in individuals homozygous for the allele.
- **Carriers** are heterozygous individuals who carry the recessive allele but are phenotypically normal; most individuals with recessive disorders are born to carrier parents.
- **Albinism** is a recessive condition characterized by a lack of pigmentation in skin and hair and eyes.
<table>
<thead>
<tr>
<th>Condition</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>CYSTIC FIBROSIS</td>
<td>Chronic, progressive pulmonary disease&lt;br&gt;Pancreatic endocrine insufficiency&lt;br&gt;Elevated sweat chloride&lt;br&gt;&lt;b&gt;Higher frequency in European Caucasians&lt;/b&gt;</td>
</tr>
<tr>
<td>TAY-SACHS DISEASE</td>
<td>Progressive neurological abnormalities&lt;br&gt;Retinal cherry-red spot&lt;br&gt;&lt;b&gt;Higher frequency in the Ashkenazi Jewish and French Canadian populations&lt;/b&gt;&lt;br&gt;Reduced serum hexosaminidase A&lt;br&gt;Usually fatal in early childhood</td>
</tr>
<tr>
<td>SICKLE CELL ANEMIA</td>
<td>Failure to thrive&lt;br&gt;Chronic anemia&lt;br&gt;&lt;s&gt;Vasoocclusive crisis (pain)&lt;/s&gt;&lt;br&gt;Increased risk for infection&lt;br&gt;&lt;b&gt;Higher frequency in those of African descent&lt;/b&gt;&lt;br&gt;Heterozygote advantage</td>
</tr>
</tbody>
</table>
Cystic Fibrosis

- **Cystic fibrosis** is the most common lethal genetic disease in the United States, striking one out of every 2,500 people of European descent.
- The cystic fibrosis allele results in defective or absent chloride transport channels in plasma membranes leading to a buildup of chloride ions outside the cell.
- Symptoms include *mucus buildup* in some internal organs and abnormal absorption of nutrients in the small intestine.
Cystic fibrosis (CF)

Cystic fibrosis is a hereditary disorder characterized by lung congestion and infection and malabsorption of nutrients by the pancreas.
A Organs affected by cystic fibrosis

Sinuses: sinusitis (infection)

Lungs: thick, sticky mucus buildup, bacterial infection, and widened airways

Skin: sweat glands produce salty sweat.

Liver: blocked biliary ducts

Pancreas: blocked pancreatic ducts

Intestines: cannot fully absorb nutrients

Reproductive organs: (male and female) complications

B Normal airway
Airway wall
Airway lined with a thin layer of mucus
(Airway in cross-section)

C Airway with cystic fibrosis
Thick, sticky mucus blocks airway
Widened airway
Blood in mucus
Bacterial infection
Figure 14.16

Parents

Normal
Aa

×

Normal
Aa

Sperm

A

a

Eggs

A

a

<table>
<thead>
<tr>
<th>AA Normal</th>
<th>Aa Normal (carrier)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Normal</td>
<td>Normal (carrier)</td>
</tr>
<tr>
<td></td>
<td>Albino</td>
</tr>
</tbody>
</table>

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• If a recessive allele that causes a disease is rare, then the chance of two carriers meeting and mating is low

• **Consanguineous matings** (i.e., matings between close relatives) increase the chance of mating between two carriers of the same rare allele

• Most societies and cultures have laws or taboos against marriages between close relatives
Sickle-Cell Disease: A Genetic Disorder with Evolutionary Implications

- **Sickle-cell disease** affects one out of 400 African-Americans
- The disease is caused by the substitution of a single amino acid in the hemoglobin protein in red blood cells
- In homozygous individuals, all hemoglobin is abnormal (sickle-cell)
- Symptoms include physical weakness, pain, organ damage, and even paralysis
Heterozygotes (said to have sickle-cell trait) are usually healthy but may suffer some symptoms.

About one out of ten African Americans has sickle cell trait, an unusually high frequency of an allele with detrimental effects in homozygotes.

Heterozygotes are less susceptible to the malaria parasite, so there is an advantage to being heterozygous.
Sickle Cell Anemia
PKU is an inherited disorder that increases the levels of phenylalanine in the blood.

Due to defective hepatic enzyme phenylalanine hydroxylase (PAH).

Necessary to metabolize the amino acid phenylalanine ('Phe') to the amino acid tyrosine.
**Symptoms**

- Elevated phenylalanine, phenylpyruvate, phenyllactate and phenylacetate in blood and urine *(musty odor of urine)*.

- **Neurological problems** (mental retardation, seizures, tremors, microcephaly etc) due to reduced production of catecholamines.

- **Hypopigmentation** (light skin, hair, blue eyes) due to reduced melatonin production. NO COMPLETE LOSS OF PIGMENT B/C WILL STILL HAVE SOME TYROSINE FROM DET