Name _	 Date

Pedigree Construction Notes

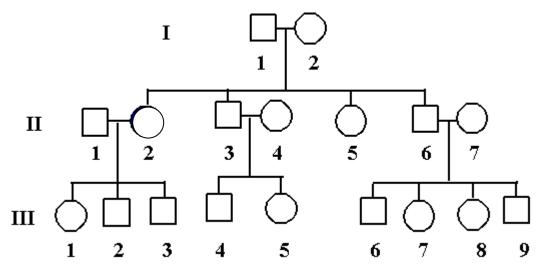
GO TO → Mendelian Inheritance (http://www.uic.edu/classes/bms/bms655/lesson3.html)

When human geneticists first began to publish family studies, they used a variety of symbols and conventions. Now there are agreed upon standards for the construction of pedigrees.

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Re	emember:		
1.	Males are always represented	by symbols, fem	ales withsymbols.
	A drawn between a so		
٦.	Two lines drawn between a so individuals are	sually second cousins or closer	relatives.
4.	When possible, the square sho of the mating line	ould be placed on the	and the circle on the
5.	Generations are connected by line to the next generation.		ending down from the mating

6.	Children of a mating are connected to a line, called the	,
7.	by short vertical lines. The children of a sibship are always listed in, the oldest being of	n
	the	
8.	Sometimes to simplify a pedigree only one parent is shown, the other is The neither signifies parthenogenic development nor does it signify divinely inspired concept	is tion
	it merely means the parent left out is not from the family being studied and is genotypica	
	for the trait being studied.	J
9.	Normal individuals are represented by an or, depending upon the gender, and affected individuals by a solid square or circle.	
10	D. Each generation is numbered to the of the sibship line with	
11.	. Individuals in each generation are numbered sequentially, beginning on the left, with	·
	For example the third individual in the second	
	generation would be identified as individual	
SC	CROLL DOWN TO "AUTOSOMAL DOMINANT INHERITANCE"	
	ead the short passage, shade the appropriate boxes and fill in the blanks below	
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	Pedigree 1. An idealized pedigree of a family with hypercholesterolemia, an autosomal dominant disease where the heterozygote has a	
	reduced number of functional low density lipoprotein receptors.	
	ne family represented by Pedigree 1 is a good example of how autosomal dominant disease	
app	ppear in a pedigree. Each of the four hallmarks of autosomal dominant inheritance are fulfil	lled.
•	Each affected individual has an there is no skipping of generat	ions
•	Each affected individual has an; there is no skipping of generat and are equally likely to be affected.	10110.
•	About 1/2 of the offspring of an affected individual are affected ().
•	(II-3) of affected individuals have all Low density lipoprotein receptors are structural proteins or polypeptides, not enzymes.	<u>_</u> .
	Low density lipoprotein receptors are structural proteins or polypeptides, not enzymes.	

ofl	, an affected female, were to produce a child that child would have a 1/2 chance being normal and a 1/2 chance of being affected. If her normal brother,, were produce a child that child would have a nearly 0 chance of being affected.
	D → AUTOSOMAL RECESSIVE INHERITANCE (www.uic.edu/classes/bms/bms655/lesson5.html)
	rst, and most important, thing to remember about autosomal recessive inheritance is that f not all, affected individuals have parents with normal phenotypes.
T	here are five hallmarks of autosomal recessive inheritance:
1.	Males and females are likely to be affected.
2.	On average, the recurrence risk to the unborn sibling of an affected individual is
3.	The trait is characteristically found in, not parents of affected or the offspring of affected.
4.	Parents of affected children may be The rarer the trait in the general population, the more likely a mating is involved.
5.	The trait may appear as an (sporadic) event in small sibships.

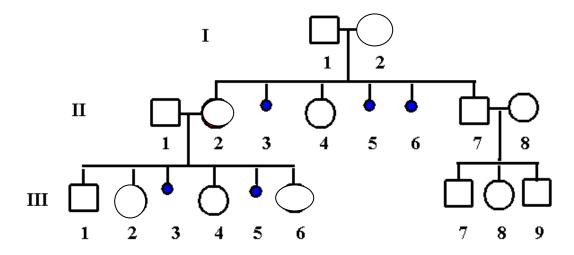


•	The above pedigree illustrates four of the five hallmarks of autosomal recessive
	inheritance and are unrelated, yet they produced an affected offspring
	<u> </u>
•	By chance, they both must have been carriers. Even though II-2 is affected, she produced
	no affected offspring ().

By far the most probable genotype for an individual from III-1, III-2 and III-3 are all	(II-1) is
(heterozygotes), since they are not affected but could only have inherited to	the recessive
gene from II-2 II-3, II-5, and II-6 each have a 2/3 chance of being a carrier	
chance of being homozygous normal. They are not affected, but they come	
• II-4 and II-7 have a high probability of being sin from outside the family. III-4, III-5, III-6, III-7, III-8, and III-9 all have a being and a 2/3 chance of being	
• One parent of each is probably, the other has a 2 being a carrier and a 1 in 2 chance of passing on the recessive allele if they	/3 chance of y were a carrier.
GO TO → X-LINKED INHERITANCE (http://www.uic.edu/classes/bms/bms655/lesson6.html)	
When the locus for a gene for a particular trait or disease lies on the X chromo	osome the
disease is said to be The inheritance pattern for X-linked inh	
from autosomal inheritance only because the X chromosome has	errance anners
in the male, the male has an X and a Y chromosome. Very few genes have been d	
the Y chromosome.	
The inheritance pattern follows the pattern of segregation of the X and Y chro	
and A male child always gets his X from or	ne of his
and his Y chromosome from his X-linked g	
passed from A child always gets the father's I	
and one of the two X's of the mother. An affected female must have are always hemizygous for X linked traits, that is, they can never be heterozygose	
homozygotes. They are never A single dose of a mutant allele w	
mutant phenotype in the male, whether the mutation is dominant or recessiv	
hand, females must be either homozygous for the normal allele, heterozygous, or	
for the mutant allele, just as they are for autosomal loci.	, ,
When an X-linked gene is said to express inheritance, it means the	
of the mutant allele will affect the phenotype of the female. A X-links	ed gene requires
two doses of the mutant allele to affect the female phenotype. The following are	the hallmarks of
X-linked dominant inheritance:	
• The trait is never passed from to	
All of an affected male and a normal female are affected male are affected male and a normal female are affected male and a normal female are affected male affected male and a normal female are affected male are affected male are affected male are	ected. All sons
of an affected male and a normal female are	

 Matings of affected females and the daughters affect 	d normal males produce the sons affected and ted.
• Males are usually morein males.	affected than females. The trait may be
• In the general population, if the disease is not lethal in ma	are more likely to be affected than males, even lles.
producing the non-functioning producing the non-functioning product at all. Affected female has two X chromosome as a tadeleterious in males than it is in fem	rected than females because in each affected female there is g a normal gene product and oneallele duct, while in each affected male there is only the stioning product and the, no normal gene are more prevalent in the general population because the ther of which could carry the mutant allele, while the male reget for the mutant allele. When the disease is no more nales, are about twice as likely to be a Pedigree 5 below, X-linked dominant inheritance has a
I	
II 2 3	
IV $\left(\begin{array}{c} 1 \\ 2 \\ 3 \end{array}\right)$	
Pedigree 5. X-linked dominant	inheritance.
The key for determining if a domination	ant trait is X-linked or autosomal is to look at the
If the affectedhas an affected	d son, then the disease is
All of his must also be affect	ted if the disease is In Pedigree 5,
both of these conditions are met.	

What happens when	_ are so	severely	affected	that they	can't reprod	duce'?
Suppose they are so severely affected t	hey never	survive to	term, the	en what ha	ppens? This	is not
uncommon in X-linked	diseases.	There are	no affect	ed males to	test for X-l	inked
dominant inheritance to see if the prod	uce all affe	ected		and no	affected	
Pedigree 6 shows the effects of such a	disease in	a family.	There			,
only affected females, in the populatio	n. Living _			outnu	mber living 1	nales
two to one when the mother is affe	cted. The	ratio in	the offspi	ring of af	fected female	es is:
affected female: normal fe	emale:	norma	l male.			



Pedigree 6.

You v	will note that in Pedi	gree 6 there have also been sev	reral spontaneous
in the offsp	oring of affected	Normally, in the genera	al population of us normal couples,
one in	recognized preg	nancies results in a	abortion. Here the ratio is
much	Presuma	ably many of the spontaneous	abortions shown in Pedigree 6 are
	_ that would have b	een affected had they survived	to term.