# **Genetics Unit Exam**

#### Question 1

You are working with an ornamental fish that shows two color phenotypes, red or white. The color is controlled by a single gene. These fish are hermaphrodites – meaning they can either (1) self-fertilize or (2) mate with another fish. You have three fish: fish 1, fish 2, and fish 3.

a) You set up the mating experiment #1 below using these fish. From the following statements check all that might be true (that is, are consistent with the data, allowing for reasonable statistical fluctuations), given only the results of experiment #1.

	Number of progeny	with following phenotype
Experiment	Red	White
#1: Fish 2 (red) with Fish 3 (red)	100	0

O Fish 2 and f	fish 3 are	homozygotes.
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- O The red phenotype is dominant to the white phenotype.
- O The white phenotype is dominant to the red phenotype.
- O Both fish 2 and fish 3 are heterozygotes.
- O Fish 2 is a homozygote, and fish 3 is a heterozygote.
- O Fish 2 is a heterozygote, and fish 3 is a homozygote.

b) You set up mating experiments #1 (same as above) and #2 below using these fish. From the following statements check all that might be true (that is, are consistent with the data, allowing for reasonable statistical fluctuations), given both experiment #1 and experiment 2.

	Number of progeny	with following phenotype
Experiment	Red	White
#1: Fish 2 (red) with Fish 3 (red)	100	0
#2: Fish 2 (red) with Fish 2 (red)	70	30

- O Fish 2 and fish 3 are homozygotes.
- O The red phenotype is dominant to the white phenotype.
- O The white phenotype is dominant to the red phenotype.
- O Both fish 2 and fish 3 are heterozygotes.
- O Fish 2 is a homozygote, and fish 3 is a heterozygote.
- O Fish 2 is a heterozygote, and fish 3 is a homozygote.
- c) You perform experiment #3 and obtain 400 progeny. Note that fish 2 is the same fish in all three experiments. In the table below, list the phenotype or phenotypes you would see in the progeny. Then give the number of offspring expected with each phenotype.

Experiment
#3: Fish 1 (white) with Fish 2 (red)

Possible Phenotypes	total number showing this phenotype

## Question 1, continued

While working further with this fish, you discover one other mutation (in a different gene) that leads to the recessive phenotype of long dorsal fins. We now have two phenotypes:

Color: red or white (described in part a). Alleles denoted by R and r.

Dorsal Fins: long (recessive) or short (dominant). Alleles denoted by D and d

\*each trait is controlled by a single gene. For the questions below, use the upper case letter for the allele associated with the dominant phenotype. Use the lower case letter for the allele associated with the recessive phenotype.

In experiment A, you cross true-breeding white fish with short fins to true-breeding red fish with long fins and get all red fish with short fins. You then cross the F1 fish to true-breeding white fish with long fins.

Experiment A: white fish with short fins X red fish with long fins

(F1) red fish with short fins X white fish with long fins

F2: ?

d) You obtain 1000 F2 offspring. Assume that the color and fin loci are 5 cM apart. In the table below, list all the possible phenotypes seen in the F2 offspring. For each phenotype given, list all the possible genotypes seen in the F2 offspring. Finally, give the number of offspring expected with each phenotype.

Possible Phenotypes	Possible Genotypes	Total F2 showing this phenotype
red fish with short fins		
red fish with long fins		
white fish with long fins		
white fish with short fins		

You are studying a genetically inherited trait. The pedigree for a family with this trait is shown below. This trait shows complete penetrance. Assume no new mutations.

Unaffected male
Affected male
Unaffected female
Affected female
Note: carriers are not distinguished from non-carriers

Complete each table below ONLY if that mode of inheritance is possible.

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If Yes, give all the possible genotype or genotypes for the individuals listed in the table below. Use A for the allele associated with the dominant phenotype and a for the allele associated with the recessive phenotype.

Individual	possible genotype or genotypes
1	
2	
3	
4	
5	

h١	Could this	trait he	inherited i	in an	Autosomal	Recessive	manner?	
v,	Could tills	tiait be	IIIIICIIICU	III aii	Autosomai	INCCCSSIVE	manifer:	

If Yes, give all the possible genotype or genotypes for the individuals listed in the table below. Use A for the allele associated with the dominant phenotype and a for the allele associated with the recessive phenotype.

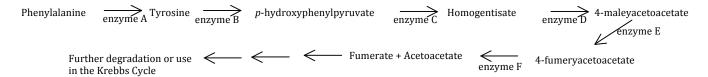
Individual	possible genotype or genotypes
1	
2	
3	
4	
5	

c)	Could this	trait be	inherited in	n an X-linked	Recessive	manner?

If Yes, give all the possible genotype or genotypes for the individuals listed in the table below. Use XA for the allele associated with the dominant phenotype and Xa for the allele associated with the recessive phenotype.

Individual	possible genotype or genotypes		
1			
2			
3			
4			
5			

The pathway for the breakdown of phenylalanine is as follows. There are six enzymes involved in this pathway. Enzyme A is encoded by gene A, enzyme B is encoded by gene B, etc. The genes involved in this pathway all lie on different chromosomes and are thus unlinked to each other.



- a) PKU is a human autosomal recessive genetic disorder where the inability to breakdown the amino acid phenylalanine and the resulting accumulation can cause neurological problems. What would be the genotype of an individual with PKU? Consider only the relevant gene and use the upper case letter for the allele associated with the dominant phenotype, the lower case letter for the allele associated with the recessive phenotype.
- b) Alkaptonuria is a human autosomal recessive genetic disorder where the individual is missing the function of enzyme D. This results in the accumulation of a byproduct that is excreted in the urine. What byproduct accumulates in these individuals?
- c) A woman with PKU who is also a carrier for Alkaptonuria marries a man with Alkaptonuria who is also a carrier for PKU.
  - i) What would be the genotype of the woman? What would be the genotype of the man?
  - ii) What are the chances that their child will appear phenotypically normal? Explain your answer or show your work.
  - iii) What are the chances that their child will show a both the PKU and the Alkaptonuria phenotype? Explain your answer or show your work.

You are studying development in a new species of fruit fly, Drosophila mit-ensis, and you find a fly that has curly wings as seen here:

Curly Wings

**Normal Wings** 

- a) The Curly wings phenotype is due to a mutation at a single locus. Assume that you have flies that are true-breeding for the curly wing phenotype (cur/cur) and flies that are true-breeding for normal wings (+/+).
  - Propose a genetic cross to determine whether the mutant phenotype is dominant or recessive, and list the genotype(s) of the F1 offspring you would obtain from this cross. Use cur for the allele associated with curly wings and + for the allele associated with normal wings.
  - What phenotype(s) would you expect to see in the F1 generation if the mutant phenotype is dominant? In what proportion would you see these phenotypes?
  - What phenotype(s) would you expect to see in the F1 generation if the mutant phenotype is recessive? In what proportion would you see these phenotypes?

You perform the experiment and find that the curly wing phenotype is recessive. A friend of yours has been studying a mutation that causes short antenna. The short antenna phenotype is recessive to the wild-type long antenna phenotype. You designate the mutant allele (ant) and the wild-type allele (+).

You set up the following crosses:

Curly wings, long antenna Normal wings, short antenna cross 1: cur + + ant cur + Χ + ant F1: Normal wings, long antenna cross 2: Normal wings, long antenna Curly wings, short antenna Χ (F1 from cross 1)

- b) What type of cross is cross 2?
- c) If the two genes are unlinked...
  - i) What phenotypic classes do you expect to see from cross 2 and in what ratios?
  - ii) Could these two genes be located on the same chromosome? Explain.

# Question 4, continued

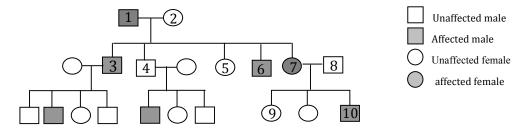
d) You perform cross 2, examine 1000 progeny and see that the genes are linked. In the table below, give the genotypes and phenotypes for the parental and recombinant progeny.

Please note that you must use the following nomenclature. Use cur and + for the alleles of the wing gene. Use ant and + for the alleles of the antennae gene.

	Number of progeny	Genotype	Phenotype
Parental	431 + 429		
Recombinant	69 + 71		

e) What is the recombination frequency between the wing and the antennae genes?

You are studying a genetically inherited disease. The pedigree for a family with this disease is shown below. Assume 1) the disease phenotype shows complete penetrance and 2) the disease allele is present at a high frequency in the population.



a) Circle all possible modes of inheritance that are consistent with this disease and assumptions 1 and 2.

Autosomal dominant Autosomal recessive X-linked dominant

X-linked recessive Y-linked

\*Now assume that individual 8 does not carry any disease alleles. Complete the remainder of this question based on this information.

Use the genotypes +/-, +/+, or -/- for autosomal traits.

Use the genotypes X<sup>+</sup>Y, X<sup>-</sup>Y, X<sup>+</sup>X<sup>-</sup>, X<sup>-</sup>X<sup>-</sup> or X<sup>+</sup>X<sup>+</sup> for X-linked traits.

Use the genotypes XY<sup>+</sup> or XY<sup>-</sup> for Y-linked traits.

\*In all cases, + indicates the wild-type allele and – the mutant allele.

- b) Give all possible genotypes for the following individuals below.
  - individual 2
  - · individual 4

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