TAs:_____

PRACTICE EXAM 2GENOME 371Autumn 2003

These questions are derived from exams given Winter 2002 and Autumn 2002.

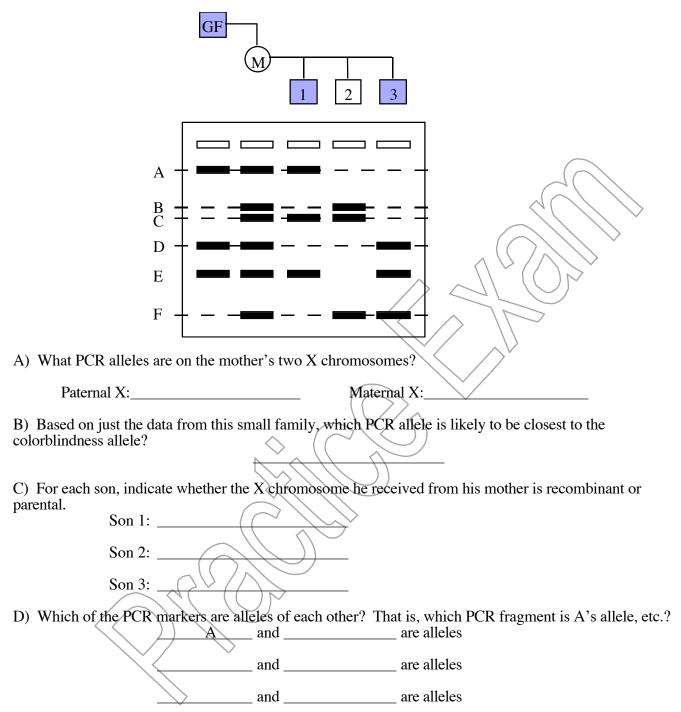
Take the exam in a quiet place and only when you are sure you will have time to complete the exam uninterrupted. Time yourself. This exam should require about 60 minutes to complete.

A key is posted separately on the web. Do NOT look up the answers until you have completed the exam.

Name:			_

TAs:_____

1) (21 pts) The gene for colorblindness is on the X chromosome, very near the end of the long arm. Three generations of a family in which colorblindness (shaded symbol) is segregating are shown below. Irrelevant individuals (the grandmother and the mother's husband) are omitted. Below each family member is the result of a PCR analysis in which three regions of the X chromosome have been amplified and combined in one lane of a gel.



E) Show the linkage arrangements and **one possible** relative order of PCR and colorblindness alleles on the mother's two X chromosomes.

Name:			
-			

TAs:___

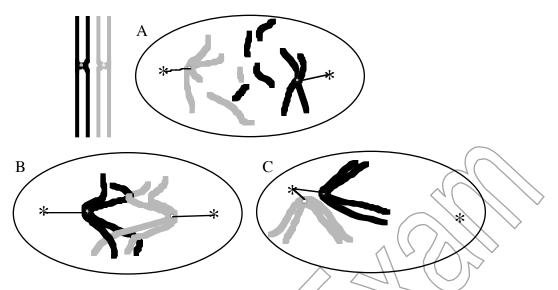
Maternal X	
Q	
Paternal X	
	$\wedge \wedge (O)$

F) On the chromosomes above, illustrate the cross-over(s) that gave rise to one of her sons; indicate son 1, 2 or 3 in your answer and circle the chromatid he received.

3

Name:

Practice Exam 2; Nov 1, 2003 TAs: 2) (12 pts) One pair of homologues from a diploid organism is illustrated below. Three cells that are attempting anaphase I of meiosis are shown in A, B, and C. None of the divisions is proceeding correctly. Some gene product was missing from each meiosis. (Assume that the spindle is normal. In the drawings only kinetochore microtubules are shown.)



A) Match the loss of function mutations listed below with the aberrant meioses.

Mutations in a gene for	could result in aberrant meiosis
ligase	
cohesin	
resolvase	
synaptonemal complex	
recBCD endonuctease	
	\rangle

B) Which of your answers to part A do you think we will mark wrong? (In other words, in which answer are you least confident?) Defend your answer as you would if you were turning this question in for a regrade!

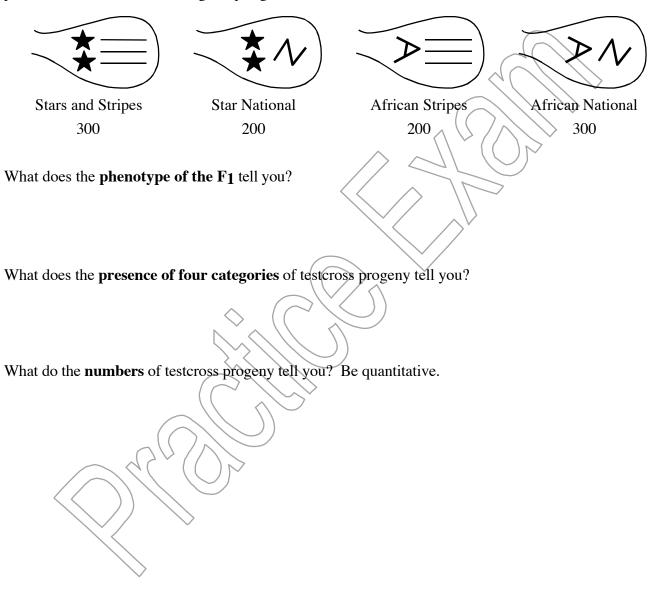
	Name: TAs:			
Eye color	Females	Males	Total	
Red	50	50	100	
White	50	50	100	
	white-ey	ed sons		
Recessive?	,	\frown		
Autosomal	?			
crossed: Defin	ne your notatio	n. <u>(</u>)		
wild ty	pe female	$\bigcirc \bigcirc \bigcirc \bigcirc$	\searrow	
B) Celeste then took white-eyed sons from				
Eye color	Females	Males	Total	
Red	1000	1	1000	
White	1000	2000	3000	
)		II	
herate this seco	nd set of offsp	ring?		
Morgan's whi	te-eyed female	es		
C) What is the genotype of the single red-eyed male (in the offspring of the second cross)? Remember that in flies, sex is determined by the X:Autosome ratio, not by the presence/absence of a Y.				
D) How did this single male arise? Be specific.				
	Red White Recessive? Autosomal crossed: Defin wild ty Eye color Red White herate this seco Morgan's whi herate (in the offs)	Eye color Females Red 50 White 50 white-eye Recessive? Autosomal? crossed: Define your notation	TAs: Eye color Females Males Red 50 50 White 50 50 white-eyed sons Recessive? Autosomal?	

Name:		

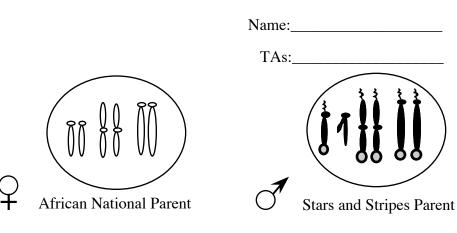
TAs:_____

4) (25 pts) You discover a new species of fruit fly with beautiful patterns on the wings. You name the species *Flaga pterae*, because the patterns remind you of various countries' flags.

A) You mate pure-breeding flies with the African National pattern to pure-breeding flies with the Stars and Stripes pattern. All the F₁ have the African National pattern. You cross the F₁ progeny to testcross parents and obtain the following offspring:



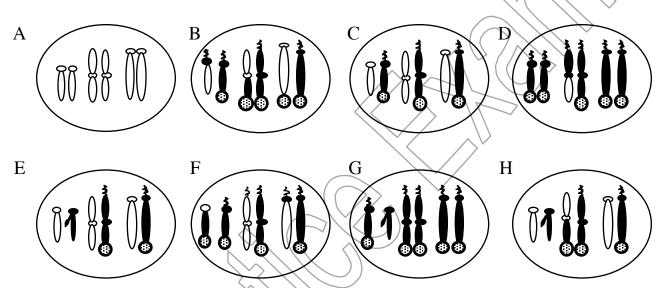
B) You isolate cells from each pure-breeding strain and analyze the chromosomes. Typical karyotypes are shown below.



Shown below are eight potential karyotypes for the F1 progeny.

Which karyotype best fits the expected chromosomes of the F1 female?

Which karyotype best fits the expected chromosomes of the F1 male?



C) You now wish to map the chromosomal location of the information for wing patterning.

When you examine the karyotype for two of the Star National testcross progeny, you see that they look like B and H above.

What would be the karyotype of the African Stripes testcross progeny? Choose a letter or letters that best fit(s) the expected karyotypes.

\mathbf{X}	
	/

Star National

African Stripes

D. Draw the pair of homologues from the F1 hybrid female that carry the wing patterning loci. Show the arrangement of alleles before she undergoes any crossing over in meiosis.

dead colony _

Name:____ TAs: 5) (14 pts) The yeast genome contains 8 different loci that encode the tyrosine-tRNAs that recognize UAC codons in mRNA. These 8 wild type genes are called sup2, 3, 4, 5, 6, 7, 8 complete plate and 11. A mutation in any of these loci that changes the anticodon loop so that it now recognizes a UAG codon is called a "suppressor". SUP3 is one such dominant mutation; SUP11 \frown \bigcirc is another. Haploid cells with either of these mutations are sick \bigcirc \bigcirc 0 \bigcirc and make small colonies on complete plates. \bigcirc 0 0 0 \bigcirc The sup3 gene and the sup11 gene are not linked. 0 \bigcirc 0 С A в A. When a strain that contains a SUP3 mutation is crossed to a strain that contains a SUP11 mutation, the diploid survives, but it is also sick. The tetrads produced by sporulating the diploid are of three basic types illustrated on the right. What type of tetrads are class A? What type of tetrads are class B? What type of tetrads are class C? B. What are the genotypes of the four spores that make up the class A tetrad? large colony and small colonies

C. What is your hypothesis for why some of the spores in type A and C tetrads are dead? Explain.