ANSWERS TO Exam Questions from Final Exam – Human Genetics, Nondisjunction, and Cancer, and Cumulative Questions

- **1.** You are working on two different organisms -- the fruit fly *Drosophila* and the yeast *S. cerevisiae*
- (a) how to make this strain in yeast: mate C haploid mutants to E haploid mutants. how to make this strain in *Drosophila*: mate true-breeding C mutants to true-breeding E mutants

list **all** of the possible genetic tests for which this strain could be used: complementation, dominant/recessive, trans

(b) how to make this strain in yeast: mate C mutant haploids to D mutant haploids, induce sporulation of the diploids, look for NPD tetrads (4 spores, 2 normal, 2 double mutant)

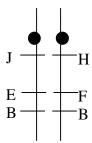
how to make this strain in *Drosophila*: can't be made, flies are diploid list **all** of the possible genetic tests for which this strain could be used: epistasis

- (c) complementation, dominant/recessive, cis, trans
- **2.** On the next page is a pedigree showing a couple that has a child with trisomy of chromosome 21.

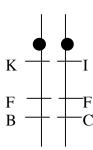
(a)

maternally inherited	С	В
allele(s) at SSR 53		_
paternally inherited	ВВ	В
allele(s) at SSR 53		
maternally inherited	F	F
allele(s) at SSR 78		
paternally inherited	FF	E
allele(s) at SSR 78		
maternally inherited	I	I
allele(s) at SSR 99		
paternally inherited	HJ	Н
allele(s) at SSR 99		

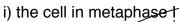
(b) Father (individual 1):

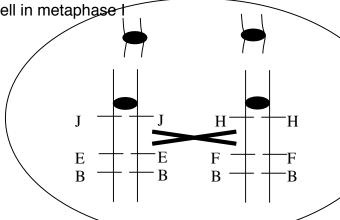


Mother (individual 2):

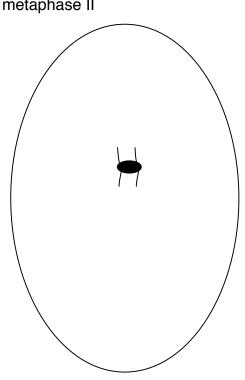


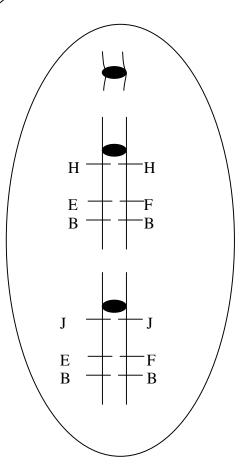
- (c) Individual 3
- (d) the dad (individual 1)
- (e) meiosis I
- **(f)**



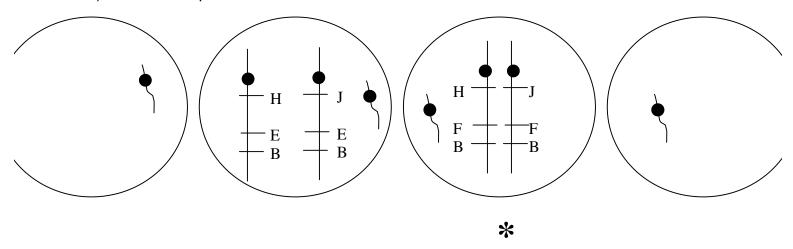


ii) the two cells in metaphase II

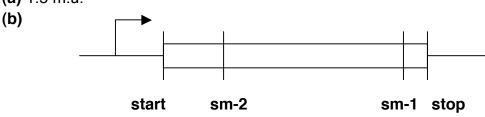




iii) the four final products of the meiosis



- **3.** After extensive genetic linkage studies, you map the locus for the ability to taste or not taste the compound PTC to a 2-centiMorgan (cM) region on human chromosome 7.
- (a) gene targeting
- (b) remove both copies of the Z gene
- (c) the Z gene disrupted by a gene that encodes antibiotic resistance. This construct would integrate at the Z locus by homologous recombination.
- (d) wild-type ES cell
- **(e)** Mate the chimeric heterozygote that results to wild-type to get a non-chimeric heterozygote. Then mate two non-chimeric heterozygotes together, and 1/4 of their progeny will be the mouse you want.
- (f) If the mouse will eat PTC-laced food, then gene Z is the locus for PTC tasting. If the mice won't eat the PTC-laced food, then gene Z is not the locus for PTC tasting.
- **(g)** Inject the wild-type human Z DNA as a transgene into the homozygous Z mutant fertilized mouse egg.
- 4. You have isolated three mutations in phage λ .
- (a) 1.5 m.u.



(c) 0.6 kb per 1% recombination



- 5. A ship carrying 7,000 passengers is about to land on an island that has 33,000 occupants.
- (a) q = (21/3500) = 0.006
- **(b)** 2pq = 0.00073
- (c) q = 0.00135
- **(d)** (1/2)q = 0.000675

6.

Genotype of strain	Will <i>cheX</i> be expressed	Will <i>cheX</i> be expressed
	when chemoattractants	when chemoattractants
	are absent?	are present?
wild-type	No	Yes
cheC1	Yes	Yes
cheB2 / F' P _X cheX+	Yes	Yes
cheA3 / F' cheC1	No	No
	A1	.
cheX+ cheB2 cheA3	No	No
P _X cheX+	No	Yes
/ F' cheC1 P _X + cheX+		

- **7.** You are studying cancer progression in mice.
- (a) constitutive
- (b) recessive
- (c) tumor suppressor
- (d) to inhibit cell division whenever nutrients are absent
- (e) constitutive
- (f) dominant
- (g) proto-oncogene

- (h) to promote cell division whenever nutrients are present.
- (i) yes because of loss of heterozygosity
- (j) one wild-type allele
- (k) two mutant alleles
- **8.** You are mapping a certain rare disorder that is caused by an allele at the N locus.

maternally inherited allele at SSR112	С	Α	Α	Α	С	С	Α
paternally inherited allele at SSR112	В	С	С	В	В	С	В

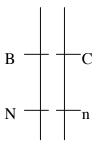
maternally inherited allele at the N locus	n	n	n	n	n	n	n
paternally inherited	N	n	N	N	N	n	n
allele at the N locus							

IF the condition is autosomal recessive [parts (a) and (b)] (Individual 2 is "nn")

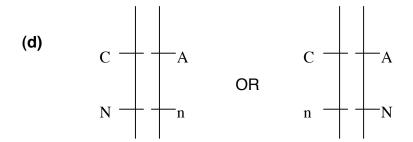
maternally inherited allele at the N locus	n	N	n	n	n	N	N
paternally inherited allele at the N locus	n	n	n	n	n	n	n

IF the condition is autosomal dominant [parts (c) - (e)] (Individual 2 is "Nn")

- (a) father individual one
- (b)



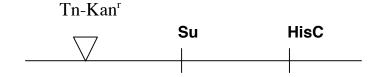
(c) mother - individual two



(e)
$$(1/2) (0.4)^3 (0.1)^4 + (1/2) (0.4)^4 (0.1)^3 = 0.262$$

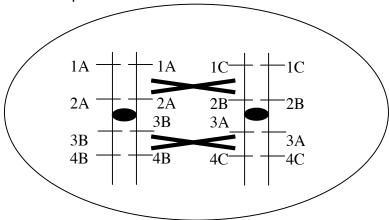
 $(0.25)^7$

- **9.** You have isolated an *E. coli* mutant that carries both an amber mutation in the HisC gene (HisC-am) and an amber suppressor mutation in a gene encoding a tRNA gene (Su+).
- (a) 20% is the distance between the Su locus and the transposon insertion
- (b)

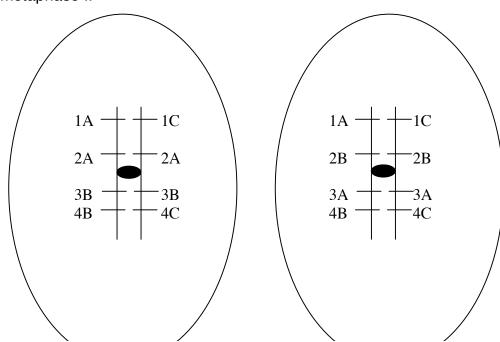


- **10.** Mendel's concept of the gene was first applied to a human trait in Archibald Garrod's landmark 1902 paper entitled "The Incidence of Alkaptonuria: A Study in Chemical Individuality."
- (a) autosomal recessive
- **(b)** q = 0.0014
- (c) $2pq = 2^* (0.9986)(0.0014) = 0.0028$
- **(d)** F = (1/16)
- (e) 2.3% of matings are between 1st cousins
- **11.** Consider the following mouse pedigree in which the indicated male exhibits a distinctive rare trait.
- **(a)** (1/3)
- **(b)** (1/4)
- **(c)** (1/4)
- (d) 45%

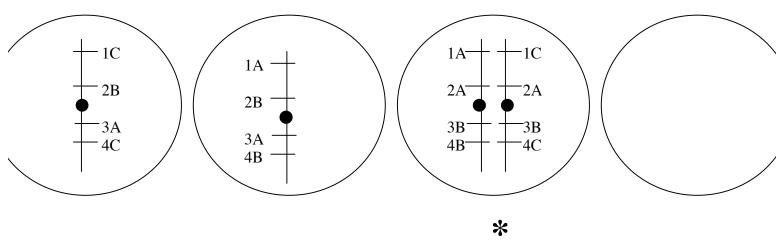
- **12.** The **cl** gene of phage lambda encodes a repressor protein that has a molecular weight of 24 kDa.
- (a) an amber nonsense mutation 2/3 of the way into the coding region
- (b) a +1 or −1 frameshift mutation that lies shortly before the nonsense mutation
- (c) 5'-TGG-3'
- (d) 5'-TAG-3'
- **13.** Your colleague, who is a medical geneticist, seeks your help in interpreting a patient: an XXY girl.
- (a) mother
- (b) meiosis II
- (c)
- i) the cell in metaphase I



ii) the two cells in metaphase II



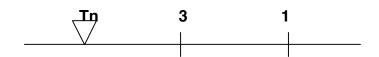
iii) the four final products of the meiosis



- (d) a translocation during the development of the sperm in the father, so that the Sry gene went to the X chromosome, and SSR1 went to the Y chromosome
- **14.** Your colleague seeks your advice regarding a family in which several individuals (filled circles or squares below) developed colon cancer in their 30's or 40's.
- (a) if you see SSR instability in the tumor cells, then it is most likely HNPCC
- **(b)** log $(0.5)^7 = 2.1$ $(0.25)^7$
- (c) yes, it becomes:

$$\log \frac{(0.5)^6}{(0.25)^6} = 1.8$$

- **(d)** yes
- (e) no, child #2 is a recombinant
- (f) yes
- **15.** Wild-type *E. coli* bacteria are motile (that is, they can swim around).
- (a) 70%
- (b) unlinked (more than 100kb away)
- (c)



- **16.** In order for yeast cells use the amino acid arginine as a nitrogen source, arginine is broken down by the enzyme arginase.
- (a) it is in a different gene than Arg1 and thus must act in trans
- (b) activator
- (c) yes, at 50cM
- (d) constitutive
- (e) Type Five
- (f) arginine → Arg2 --] Arg3 --] Arg1
- **17.** One in 20,000 human males is an (infertile) XX male (due to a translocation that moves the sex-determination gene Sry onto an X chromosome).
- (a) DZ = (1/2) (1/20,000)
- **(b)** MZ = 100%
- (c) DZ = 50%
- (d) MZ = 100%
- (e) DZ = (1/885)
- (f) MZ = 100%
- **18.** Consider a codominant blood antigen where individuals homozygous for one allele express only antigen M, individuals homozygous for the other antigen express only antigen N, and heterozygous individuals express both N and M antigens. **(a)**

	Frequency	expressing	Allele fred	quencies	
Population	M only	N only	M	N	H-W equilib (yes or no)?
1	0.25	0.25	0.5	0.5	yes
2	0.36	0.16	0.6	0.4	yes
3	0.01	0.64	0.185	0.815	no

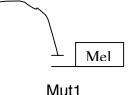
(b)
$$q^2 + 0.1q - 10^{-5} = 0$$

so
$$q = 10^{-4}$$

(c) Fq= 0.004

- **19.** HNPCC (Hereditary Non-Polyposis Colon Cancer) shows autosomal dominant inheritance in humans.
- (a) knockout of the mouse MSH2 gene
- (b) more quickly
- **20.** Your project is to genetically map the locus for color blindness, an X-linked recessive trait, with respect to SSR markers.
- (a) AB
- (b) ac
- (d) $\log \frac{(1/2)(0.4)^1(0.1)^9 + (1/2)(0.4)^9(0.1)^1}{(0.25)^{10}} = 1.14$
- (f) log $(1/2) (0.45)^{1} (0.05)^{9} + (1/2) (0.45)^{9} (0.05)^{1}$ = 1.30 $(0.25)^{10}$
- (g) the SSR72 would be in the middle

- **21.** The following three crosses involve mice from either true-breeding mutant strains or true-breeding wild-type strains.
- (a) X-linked recessive only
- (b) X-linked recessive and autosomal recessive
- (c) autosomal recessive only
- **22.** *E. coli* can utilize the sugar melibiose after induction of the enzyme melibiase.
- (a) yes, at 95%
- (b) it gives a dominant phenotype
- (c) It works in cis
- (d) melibiose --] repressor



- (e) the mutant repressor can't bind to the operator any more
- **23.** The sequence of the amber stop codon is ⁵'UAG³'.
- (a) the gln-tRNA gene and the trp-tRNA gene
- (b) gln: 5'-CAG-3' ← this strand is used as a template during transcription 3'-GTC-5'
- trp: 5'-TGG-3' ← this strand is used as a template during transcription 3'-ACC-5'
- (c) 5'-TAG-3' ← this strand is used as a template during transcription 3'-ATC-5'
- **24.** You are studying the yeast genes needed to metabolize organic phosphates.
- (a) an unlinked suppressor mutation that has no phenotype on its own for instance, a tRNA suppressor allele
- **(b)** this suppressor mutation is very close to the original mutation, and thus might be an intragenic suppressor mutation (and even may be a back mutation to revert the gene to its original sequence)

- **(c)** an unlinked suppressor mutation whose phenotype on its own is recessive and constitutive; it could be a mutation in a gene encoding a repressor protein that works downstream from Pho4 in the pathway that regulates the expression of phosphatase
- **25.** Shown below is a hypothetical scheme for the formation of eye pigment in *Drosophila*.
- (a) purple eyes
- (b) because the double mutant flies (bl pr mutants) have blue eyes
- (c)

|--|

Purple-eyed males: 42

Blue-eyed males: 50

Red-eyed males: 8

26. When setting out to determine the chromosomal location of a rare human disease gene by genetic linkage analysis (LOD scores), it is useful to calculate the theoretical maximum LOD score that a family of a given size and structure might contribute.

(a)
$$\log \frac{(1/2)(0.5)^{10}}{(0.25)^{10}} = 2.7$$

(b)
$$\log \frac{(0.5)^{10}}{(0.25)^{10}} = 3.01$$

(c)
$$\log \frac{(0.5)^{10}}{(0.25)^{10}} = 3.01$$

(d)
$$\log \frac{(1/2)(0.5)^2}{(0.25)^2} = 0.301$$

(e)
$$\log \frac{(1/2)(0.5)^1}{(0.25)^1} = 0$$

(f) log
$$(0.5)^{1}$$
 = 0.301 $(0.25)^{1}$

(g) log
$$(1/2) (0.5)^1 = 0$$

 $(0.25)^1$

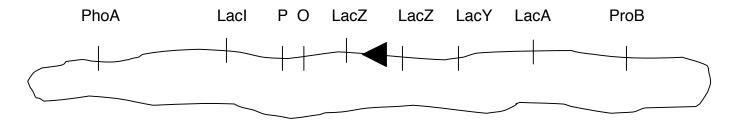
- **27.** You discover a frame-shift mutation in an X-linked gene called SPG in a man who is infertile because of poor sperm production.
- (a) Yes it is possible, because the mutation can be passed on by females indefinitely.
- (b) i) pronuclear injection
 - ii) wt human SPG
 - iii) fertilized egg
 - iv) a fertilized egg that is XY and carries a mutant form of SPG on the X chrom
 - v) randomly
 - vi) no
 - vii) none
- viii) if the male mouse is fertile, then the human and mouse SPG genes are interchangeable. If the male mouse is sterile, then the human and mouse genes are not interchangeable.
- **28.** You have isolated two different X-linked mutations in *Drosophila* that affect eye color.
- (a) orange
- (b) same gene
- (c) 1 cM
- (d)



29. You have constructed an **F**' plasmid that carries the **LacZ** gene.



(b) the Hfr:



- (c) no; it won't express LacY because the gene is disjointed from its promoter
- (d) late
- **30.** The **PyrG** gene is found to lie about 40 kb away from the group of **Lac** genes on the *E. coli* chromosome.

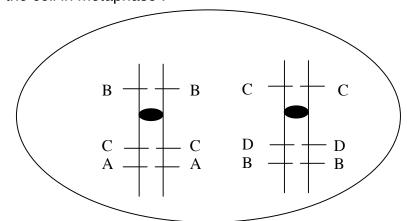
Draw a diagram of this region of the chromosome that shows where the **PyrG** gene maps relative to **LacZ** and **LacI**.

ANSWER:

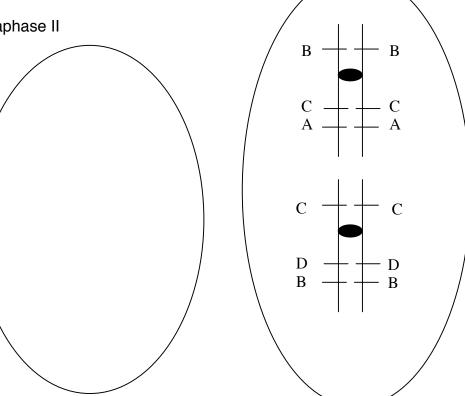


- **31.** Only a small fraction of human fetuses with trisomy 18 survive to birth, and most of those surviving to birth die in infancy.
- (a) before
- (b) mother
- (c) meiosis I

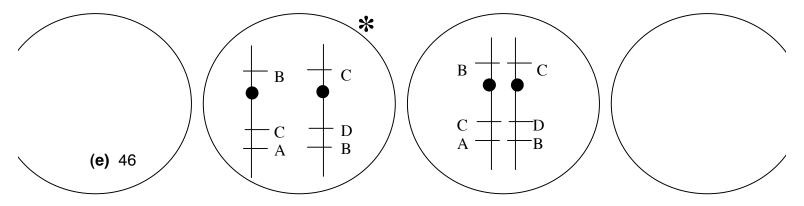
(d) i) the cell in metaphase I



ii) the two cells in metaphase II



iii) the four final products of the meiosis



32. In some families, breast cancer displays autosomal dominant inheritance.

(a) log
$$(1/2) (0.05)^6 (0.45)^1 + (1/2) (0.05)^1 (0.45)^6 = 0.53$$

 $(0.25)^7$

(b)
$$\log \frac{(0.05)^1 (0.45)^6}{(0.25)^7} = 0.83$$

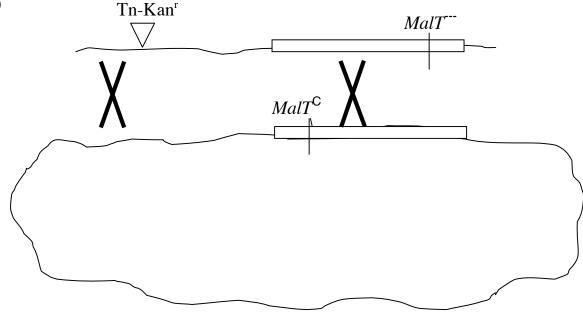
- (c) the cell that develops into a tumor needs to lose the 2nd copy of the gene by loss of heterozygosity
- (d) the phenotype caused by a homozygous loss of BRCA is lethality. Perhaps BRCA is necessary for normal fetal development to occur.
- **33.** You are genetically mapping the locus that determines a rare skin disease that shows autosomal dominant inheritance.
- (a) the father
- **(b)** A

(d)
$$\log \frac{(0.45)^5}{(0.25)^5} = 1.27$$

(e) three families

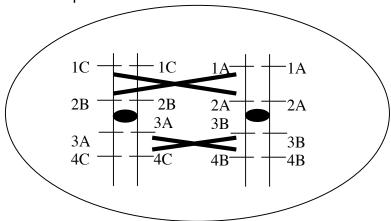
- **34.** Consider an autosomal gene at which a rare allele (call it allele **a**) results in homozygotes (**aa**) having only 20% of the number of offspring as average individuals in the population.
- (a) S = 0.8
- **(b)** h = 0.002
- (c) f(a) = 0.0025
- (d) 2pq = 0.005
- (e) new q = 0.00225
- **35.** You have isolated a Tn5 insertion in an otherwise wild-type *E. coli* strain that is linked to the gene encoding the MalT activator protein.
- (a) 80%

(b)

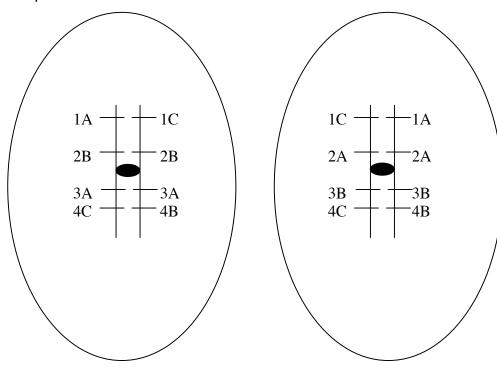


- (c) an ochre nonsense mutation
- (d) uninducible
- **36.** You are called by your family physician to provide an expert genetic opinion on an unusual patient: an XXX boy.

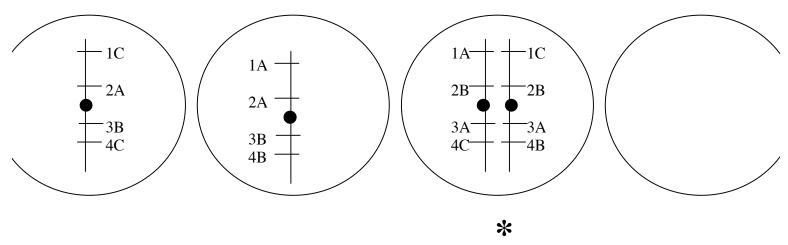
- (a) mother
- (b) meiosis II
- (c)
- i) the cell in metaphase I



ii) the two cells in metaphase II



iii) the four final products of the meiosis



- (d) the father must have passed on an X chromosome which mistakenly carried the Sry gene that causes maleness
- (e) a translocation occurred during the development of that sperm such that SSR1 went to the Y chromosome in exchange for the Sry gene a