

## OR072: Midterm Quiz Solutions

- polymerase
1.  
A. matches mRNA codons to tRNA anticodons  
B. matches tRNA anticodons to amino acids  
 C. makes a complementary copy of DNA or RNA  
D. algebraic expression for a genotype
- intron
- A. area of DNA that has genetic material  
 B. area of DNA that has no genetic material  
C. one of a pair of genes determining a particular trait  
D. computer chip manufacturer
- Tjio&Levan
- A. discovered number of human chromosomes  
B. showed that DNA contains genetic information  
C. found gene for Huntington's disease  
D. world-renown juggling team
- Linus Pauling
- A. established structure of proteins  
B. "Father of American Genetics"  
C. discovered first genetic disease  
D. found mapping from DNA triples to amino acids
- Erwin Schrödinger
- A. head of laboratory where DNA was discovered  
 B. famous physicist who asked "What is Life"  
C. invented the first computer  
D. determined the crystallographic structure of DNA
- Oswald Avery
- A. established that proteins are responsible for bodily functions  
 B. established that DNA contained hereditary properties  
C. discovered the structure of proteins  
D. discovered the structure of DNA
- Myrna Gopnyk
- A. discovered first genetic disease  
B. established the genetic nature of Huntington's disease  
C. used X-ray crystallography to determine the structure of DNA  
 D. established that grammar is hereditary
- A rooted tree with at most 3 children per node can have *at most* \_\_\_\_ leaves of depth 3.
- A. 9  
B. 16  
 C. 27  
D. 41

2. A.  $(.40)(.40)\frac{1}{16}(n+n)(H+h)(n+n)(H+h)$   
 $+ (.60)(.60)\frac{1}{16}(N+N)(H+h)(N+N)(H+h)$   
 $+ 2(.40)(.60)\frac{1}{16}(n+n)(H+h)(N+N)(H+h)$   
 $= .04nnHH + .08nnHh + .04nnhh + .09NNHH + .18NNHh$   
 $+ .09NNhh + .12nNHH + .24nNhh + .12nNhh$
- B.  $(.90)(.80)(.40)(.60)\frac{1}{16}(n+n)(H+h)(N+N)(H+h)$   
 $= .0864nNHH + .1728nNhh + .0864nNhh$
- C.  $\frac{1}{4}(nn + Hh)(NN + Hh) = .25nnNN + .25nnHh + .25HhNN + .25HhHh$
- D.  $(.40)(.60)\frac{1}{16}(n+n)(H+h)(N+N)(H+h)$   
 $= .12nNHH + .24nNhh + .12nNhh$
- E.  $\frac{1}{16}(n+n)(H+h)(N+N)(H+h) = .25nNHH + .5nNhh + .25nNhh$
- F.  $.04(1)(1) + .08(1)(1) + .04(1)(.9) + .09(.8)(1) + .18(.8)(1) + .09(.8)(.9) + .12(.8)(1) + .24(.8)(1) + .12(.8)(.9)$   
 $= .04 + .08 + .036 + .072 + .144 + .0648 + .096 + .192 + .0864 = .8112$

A particular species of bird has two pairs of traits, nearsightedness( $N$ )/farsightedness( $n$ ), and hard-shelled( $H$ )/soft-shelled( $h$ ) eggs, with  $N$  and  $H$  the dominant traits.

- (a) Suppose a pure far-sighted, half hard-shelled bird ( $nnHh$ ) mated with a pure nearsighted, half hard-shelled bird ( $NNHh$ ). What would be the percentage of nearsighted soft-shelled offspring in the following generation?

**Expression E:** Coefficient of  $nNhh$  term = 25%

- (b) Suppose you have a population of these two species with 40%  $nnHh$  and 60%  $NNHh$ . Now what would be the percentage of nearsighted soft-shelled offspring in the following generation?

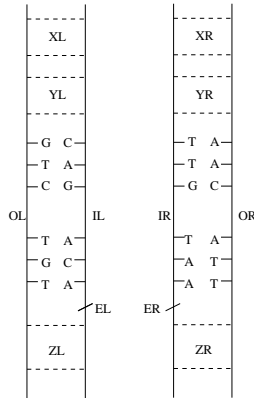
**Expression A:** Add coefficients of  $NNhh$  and  $nNhh$  terms =  $.09 + .12 = 21\%$

- (c) Finally, suppose that 10% of all soft-shelled eggs are destroyed before they hatch, and that 20% of all hatched nearsighted birds are killed before they mate? (Assume that eggs/birds are not destroyed by any other means.) Starting with the mating-age population given in (b), what would be the percentage nearsighted soft-shelled offspring reaching mating age in the following generation?

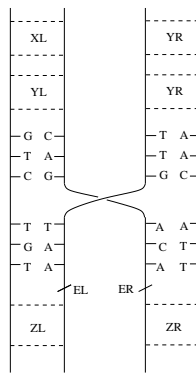
**Expression F:** Add terms corresponding to  $NNhh$  and  $nNhh$ , and then divide by the sum of all the terms

$$\frac{.0648 + .0864}{.8112} = 18.64\%$$

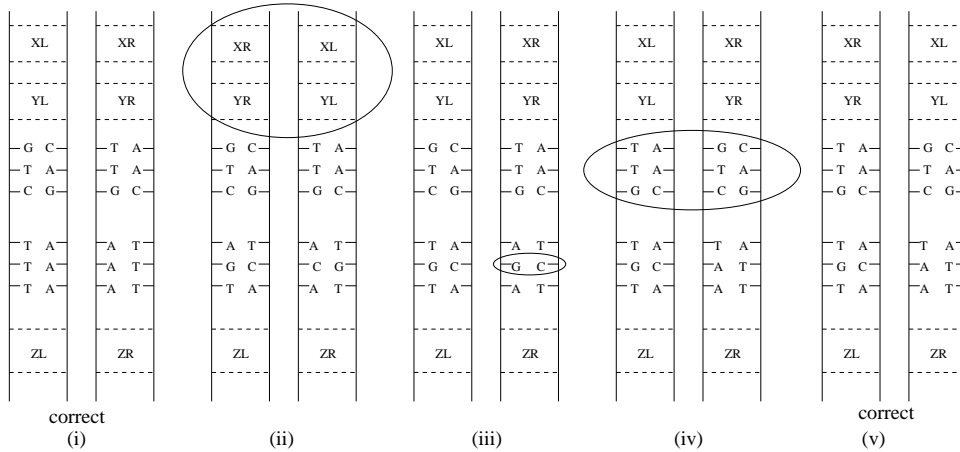
3. The initial picture:



(a) Draw a picture of the situation after the crossing occurs.



(b) Which of the following pairs could be valid results of a crossing between these pairs? If the pair is invalid, circle the parts that are wrong.



(c) How many possible results of a crossing of these chromatids can there be?

There are 4 mismatched pairs, and two ways of cutting the strands, for a total of  $2 \times 2 \times 2 \times 2 \times 2 = 32$  possible results of the crossing.