

1. c

5. c (same value as A and T are complementary bases)

2. a

6. b

3. b

7. b (in a typical heterozygous cross 25% change of homozygous recessive)

4. b

9. male has recessive phenotype & homozygous recessive (rr)  
female is heterozygous (Rr)

	r	r
R	Rr	Rr
r	rr	rr

genotype = 50% Rr, 50% rr

phenotype = 50% red coat (Rr), 50% silver-black coat (rr)

10. Sex-linked (XX + XY). The question assumes colour blindness is recessive (mum is a carrier for colour blindness)

male - has recessive phenotype  $X_b Y$

mum - is a carrier (heterozygous)  $X_B X_b$

	$X_b$	Y
$X_B$	$X_B X_b$	$X_B Y$
$X_b$	$X_b X_b$	$X_b Y$

boys: 50%  $X_B Y$ , 50%  $X_b Y$   
"normal", colour blind

girls - 50%  $X_B X_b$ , 50%  $X_b X_b$   
"normal", colour blind

(a) 50% - see above

(b) 50% - see above

- 8.
- |            |                 |
|------------|-----------------|
| a. asexual | f. recessive    |
| b. sexual  | g. dominant     |
| c. gene    | h. heterozygous |
| d. 46      | i. homozygous   |
| e. 23      |                 |

11. a) Recessive. In the pedigree, 2 "normal" parents have had an affected child. This is only possible if the trait is caused by a recessive allele. This occurs twice for parents 1+2, and once for parents 10+11.

ALTERNATIVELY - Recessive. In the pedigree, the trait skipped a generation and went from grand<sup>(2+4)</sup>parents to grand<sup>(12+13)</sup>children without being in parents<sup>(10,11)</sup>. This is only possible if recessive

b) 1. Aa    4. aa    10. Aa    11. Aa.

12. a) Three

b) II<sub>2</sub> - normal tooth enamel, II<sub>4</sub> - normal, II<sub>8</sub> - normal enamel

c) (brother-sister) siblings

d) Assumption made that this trait is caused by a dominant allele (see answer to e)

II<sub>2</sub> - Ff, II<sub>2</sub> - ff, II<sub>4</sub> - ff, II<sub>6</sub> - Ff, II<sub>8</sub> - ff.

III<sub>2</sub> - Ff, III<sub>4</sub> - ff, III<sub>6</sub> - ff, III<sub>8</sub> - ff, III<sub>10</sub> - Ff

e) Actually could be either, no proof it is recessive and both assumptions work for this pedigree. Because it did NOT skip a generation I have assumed it is a dominant allele, but this is NOT proof.

f) Assuming it is dominantly inherited, I<sub>1</sub> would be homozygous recessive.

Assuming it is recessively inherited, I<sub>1</sub> would be heterozygous

13. Darwin's theory of evolution would explain future change by

- Penguins will continue to have more chicks than can survive
- The chicks will continue to be different colours including white & grey
- The environmental change would mean that non-white chicks would be better adapted and more likely to survive to adulthood and have offspring
- This trait (non-white) would be inherited by the next generation
- The next generation of penguins (after the climate change) would mainly be non-white (grey)