

# Topic 10 – data-based questions

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- coloured, starchy both dominant traits Cc; white, waxy recessive traits Ss; F<sub>1</sub> are all CcSs; so F<sub>1</sub> × F<sub>1</sub>; CcSs × CcSs produces typical dihybrid ratio of 9 coloured starchy: 3 coloured waxy: 3 white starchy: 1 white waxy in F<sub>2</sub>;
- **2.** the actual frequencies do not follow the 9:3:3:1 ratio and so the genes must be linked as they differ from the theoretical ratio for dihybrid crosses;
- **3.** coloured, shrunken CCnn; white, non-shrinken ccNN; F<sub>1</sub> coloured, non-shrunken is CcNn are test-crossed with homozygous recessive: ccnn; CcNn × ccnn; typical ratio of 1 coloured non-shrunken: 1 coloured shrunken: 1 white non-shrunken: 1 white shrunken
- **4.** actual frequencies frequencies differ from typical ratio of 1:1:1:1, so genes must be linked;
- **5.** if starchy/waxy and non-shrunken/shrunken are both linked to colour, then they must also be linked to each other;

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#### 1 and 2.

|          | White Crested | Non-white, Non<br>crested | Non-white<br>Crested | White<br>Non-crested | Total |
|----------|---------------|---------------------------|----------------------|----------------------|-------|
| observed | 337           | 337                       | 34                   | 46                   | 754   |
| expected | 188.5         | 188.5                     | 188.5                | 188.5                | 754   |

3. 3 degrees of freedom expected;

**4.** critical value for 3 df = 7.815;

- 5.  $X^2 >> 7.815;$
- 6.  $H_0$  the traits are not linked and differences between observed and expected are due to sampling error;  $H_1$  the traits are linked and differences between observed and expected are not due to sampling error;  $X^2 >> 7.815$ , therefore p<<0.05; reject  $H_0$  and accept  $H_1$ ;

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- a) negative correlation / mean length declining with time;
- **b)** the longer the horns, the more likely the sheep will be shot; advantage to having short horns; long horns removed from reproductive pool; mean length becomes shorter with time; shorter horn alleles become more common in population with time; this is directional selection;
- **c)** long horns more likely to win in courtship battles and become more common in reproductive pool; long horns more likely to be hunted and removed from the pool; the latter seems to be the most relevant factor;

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- a) any value from 3.25 to 3.49 kg;
- **b)** any value from 3.50 to 3.74 kg;
- **c)** initially as birth mass increases up to 3.5 kg, survival increases, hencemortality decreases; then, as birth mass further increases beyond 3.5 kg, survival decreases andmortality increases; further from mode the higher the mortality, the highest survival and lowest mortality nearest to mode value;
- d) birth mass shows variation; selection against very low / very high birth weights;

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- a) (i) sneaking approximately 80 cm
  - (ii) fighting approximately 200 cm





**b)** (i) >60 cm body size for fighting

(ii) 25-29 for sneaking;

- **c)** 45–49 / 40–44;
- d) extreme size forms reproduce; intermediate size forms are selected against;

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- **1.** *C. lucasina* has uniform loudness whereas the *C. mediterranea* grows louder and then softer; the *C. lucasina* song lasts longer; individual notes would be more audible in *C. lucasina*;
- **2.** lacewings with certain songs will attract some mates, but not others; over time, gene pools become isolated within the population; this would lead to sympatric reproductive isolation;
- **3. a)** allopatric speciation is reproductive isolation due to geographic barriers; founder populations may differ in allele frequencies; the source and founder populations are subjected to different selection pressures; leading to increasing differences between the two;
  - **b)** sympatric speciation is reproductive isolation within the same geographic area; it could be caused by behavioural or temporal isolation; variables prevent interbreeding of sub-populations; the same selection pressures present in the habitat may affect the two sub-populations differently.