

## JARAMOGI OGINGA ODINGA UNIVERSITY OF SCIENCE AND TECHNOLOGY SCHOOL OF AGRICULTURAL AND FOOD SCIENCES

# SECOND YEAR FIRST SEMESTER UNIVERSITY EXAMINATION FOR THE DEGREE OF BACHELOR OF SCIENCE ANIMAL SCIENCE 2018/2019 ACADEMIC YEAR

### REGULAR

**COURSE CODE: AAS 3217** 

**COURSE TITLE: QUANTITATIVE GENETICS** 

**EXAM VENUE:** STREAM: BSc (Animal Science)

DATE: EXAM SESSION:

TIME: 2.00 HOURS

#### **Instructions:**

- 1. Answer ALL question in Section A (compulsory) and ANY other TWO questions in Section B.
- 2. Candidates are advised NOT to write on the question paper.
- 3. Candidates must hand in their answer booklets to the invigilator while in the examination room.

#### SECTION A [30 MARKS]

#### Answer ALL questions from this Section.

#### **Question 1**

Question 1 consists of 10 multiple choice questions. Only one of the four alternatives labelled (a), (b), (c) and (d) is correct. Write the letter corresponding to the correct answer in your booklet.

Q1.1 - 1.4 The data below are on egg weight in chickens. The values have been transformed to a base of 100.

| 1.1 | in the broad sense.  |      | Calculate | heritability (1 mark) |
|-----|----------------------|------|-----------|-----------------------|
|     | (a)                  | 0.35 |           |                       |
|     | (b)                  | 0.30 |           |                       |
|     | (c)                  | 0.65 |           |                       |
|     | (d)                  | 0.75 |           |                       |
|     |                      |      |           |                       |
| 1.2 | in the narrow sense. |      | Calculate | heritability (1 mark) |
|     | (a)                  | 0.35 |           |                       |
|     | (b)                  | 0.30 |           |                       |
|     | (c)                  | 0.65 |           |                       |
|     | (d)                  | 0.75 |           |                       |
|     |                      |      |           |                       |
| 1.3 |                      | Ca   | lculate r | epeatability.         |
|     | mark)                |      |           | (1                    |
|     | (a)                  | 0.35 |           |                       |
|     | (b)                  | 0.30 |           |                       |
|     | (c) 0.65             |      |           |                       |
|     | (d)                  | 0.75 |           |                       |

| 1.4 | environmental variance.<br>mark)                               |          | Cal      | culate  | ter     | mporary<br>(1 |
|-----|--|----------|----------|---------|---------|---------------|
|     | (a) 1  | 5        |          |         |         |               |
|     | (b) 2  | 5        |          |         |         |               |
|     | (c) 3  | 5        |          |         |         |               |
|     | (d) 6  | 5        |          |         |         |               |
| 1.5 | If additive genetic variance of a quantitative trait (1 mark)  | is zero  | , then   |         |         |               |
|     | (a) h  | eritabil | ity is a | at maxi | mum va  | alue.         |
|     | (b) dominance variance is zero.                                |          |          |         |         |               |
|     | (c) the environment.   | ne trait | is 1     | argely  | contro  | lled by       |
|     | (d) r  | one of   | the ab   | ove.    |         |               |
| 1.6 | Pleiotropy is a special situation in which mark)               |          |          |         |         | (1            |
|     | (a) many genes control a single trait.                         |          |          |         |         |               |
|     | (b) there is polygenic inheritance.                            |          |          |         |         |               |
|     | (c) there is no polygenic inheritance.                         |          |          |         |         |               |
|     | (d) same gene has different effects on different traits at the | e same   | time.    |         |         |               |
| 1.7 | Additive genetic effects are the most important because        | part o   | of the   | pheno   | typic v | ariation      |
|     |  |          |          |         | (1 r    | mark)         |
|     | (a) transmitted from one generation to the next.               | hey ar   | e the    | stable  | and re  | egularly      |

|      | (b) phenotypic expression.  | the have the greatest effects on the |  |  |
|------|---|--------------------------------------|--|--|
|      | (c)   | they interact with the environment.  |  |  |
|      | (d)   | none of the above.                   |  |  |
| 1.8  | Genotype x Environment interaction occurs mark)   | (1                                   |  |  |
|      | (a) genotypes defies the environment in which they are k  | when relative performance of ept.    |  |  |
|      | (b)   | in Bos indicus breeds only.          |  |  |
|      | (c) genotypes changes with genotypes.   | when relative performance of         |  |  |
|      | (d) exposed to good environment.  | when certain genotypes are           |  |  |
| 1.9  | If there are records available on the performance of an animal, then the best esti of its future performance is mark) |                                      |  |  |
|      | (a)   | her first record.                    |  |  |
|      | (b) records.  | the mean of all her previous         |  |  |
|      | (c)   | her breeding value estimate          |  |  |
|      | (d) Ability estimate.   | her Most Probable Producing          |  |  |
| 1.10 | Mass selection is suitable for genetic improveme mark)  | ent of (1                            |  |  |
|      | (a)   | traits of high heritability.         |  |  |
|      | (b)   | reproductive traits.                 |  |  |
|      | (c)   | sex limited traits.                  |  |  |
|      | (d)   | traits of low heritability.          |  |  |

#### **Question 2**

| (a) e:      | xample    | Differentiate between the following. You may use illustrations.   | ons and /or  |
|-------------|-----------|---|--------------|
|             | i.        | Mass selection and pedigree selection.  | (2 marks)    |
|             | ii.<br>ma | Genotype x Environment interaction and Genotype – Environment corrks)   | relation. (2 |
| i           | iii.      | Breeding value and Most Probable Producing Ability.   | (2 marks)    |
| :           | iv.       | Pure breeding and crossbreeding.  | (2 marks)    |
|             | v.        | Heritability in the broad sense and repeatability.  | (2 marks)    |
| (b)         |           | Explain the following terms, giving examples where necessary.   |              |
|             | i.        | Pleiotropy.   | (2 marks)    |
|             | ii.       | Selection differential.   | (2 marks)    |
|             | iii.      | Correlated characters   | . (2 marks)  |
|             | iv.       | Heterosis.  | (2 marks)    |
|             | v.        | Inbreeding depression   | n. (2 marks) |
| <u>Oues</u> | stion 3   |   |              |
| whic        | h the a   | at selection is carried out separately for males and females in a herd of Bouverage daily gain was 0.22 kg. If the means of the selected males and female $\frac{1}{2}$ /g/day, respectively. |              |
| (a)         | Com       | apute the average selection differential when both males and females are select   | ted.         |
|             |           |   | (3 marks)    |
| (b)         | Repe      | eat (a) above when only males are selected.   | (3 marks)    |
| (c)         | By h      | ow much has the potential genetic gain changed by selecting only the males.   | (2 marks)    |
| (d)         | Expl      | ain the implication for the change in (c) above.  | (2 marks)    |

#### SECTION B [40 MARKS]

#### Answer ANY TWO questions from this Section.

#### **Question 4**

| What is family   | selection in the context of quantitative genetics?       | (3 marks) |  |  |
|--|--|-----------|--|--|
| State the two m  | nain family lines and explain how ech of them can arise. | (6 marks) |  |  |
| Briefly discuss the limitations of whole family selection and within family selection. |  |           |  |  |
| Indicate and explain the conditions under which family selection can be recommended.   |  |           |  |  |
| Question 5   |  |           |  |  |
| Write short notes on the following:  |  |           |  |  |
| (a)<br>marks)  | Maternal effects.  | (5        |  |  |
| (b)<br>marks)  | Correlated traits.                                       | (5        |  |  |

#### **Question 6**

marks)

marks)

(c)

(d)

Answer the following questions briefly and clearly. Explain your calculations.

Indirect selection.

Backcrossing.

Assume a repeatability of weaning weight, as trait of the dam, is 0.4, while heritability is 0.3. If herd average is 200 kg.

(5

(5

- (a) Compute the best estimate of the next record of a cow named Kitale which has first calf with a weaning weight of 220 kg. (3 marks)
- (b) Repeat (a) above when Kitale has had four calves with weaning weights of 220, 232, 245 and 250 kg. (7 marks)
- (c) Consider weaning weight as a trait of the dam, calculate the breeding value of Kitale using her first record only. (3 maks)

| (d) | Repeat (c) above using all records available on Kitale |  |  |
|-----|--|--|--|
|     |  |  |  |
|     |  |  |  |