

# Applied Genetic Evaluation - Exercise 2

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*2020-04-27*

## Problem 1: Analysis of Variance

Estimate the variance component for the sire effect using an analysis of variance. The data is available from [https://charlotte-ngs.github.io/GELASMSS2020/ex/w10/data\\_sire\\_w10.csv](https://charlotte-ngs.github.io/GELASMSS2020/ex/w10/data_sire_w10.csv). Because the data contains just female animals, the fixed effect of the sex can no longer be estimated.

### Hint

- Use the functions `aov()` to do the analysis of variance and the function `summary()` on the ANOVA result to get the relevant parts of the variance components.

## Problem 2: Variance Components Estimation Using REML

Use the same data set as for Problem 1 and a sire model to estimate the same sire variance  $\sigma_s^2$ . The sire model is the linear mixed effects model that contains sire effects as random component. The model can be specified as

$$y = Xb + Zs + e$$

where  $y$  is the vector of observations,  $b$  is the vector of fixed effects which are the same as in Problem 1,  $s$  is the vector of random sire effects and  $e$  is the vector of random error terms.

### Hint

- Use the package `pedigreemm` to get a REML estimate for the sire variance component  $\sigma_s^2$ .
- We assume that the sires are not related. Hence variance-covariance matrix  $var(s)$  of the sire components are  $var(s) = I * \sigma_s^2$ .

## Additional Problem: Variance Components Estimation Using an Animal Model

We are given the dataset with the response variable `carcass weight` (CW) and the predictor variables that resulted from the model selection process from Exercise 1. These consisted of

- sex (`sex`)
- slaughterhouse (`slh`)
- herd (`hrd`)
- age at slaughter (`age`)

The data is available from [https://charlotte-ngs.github.io/GELASMSS2020/ex/w10/data\\_bp\\_w10.csv](https://charlotte-ngs.github.io/GELASMSS2020/ex/w10/data_bp_w10.csv).

We use a mixed linear effects model to estimate the variance components for the random effects in the model.

$$y = Xb + Za + e \tag{1}$$

where  $y$  is a vector of observations,  $b$  is a vector of fixed effects found to be relevant in Exercise 1,  $a$  is a vector of random breeding values and  $e$  is a vector of random errors.

**Hint**

- Use the package `pedigreemm` to get an estimate of the variance components

**Your Task**

- Estimate the variance components  $\sigma_a^2$  and  $\sigma_e^2$  for the two random component  $a$  and  $e$ , respectively.