# GENETIC PARAMETERS FOR GROWTH, WOOL AND IN VIVO CARCASS TRAITS IN DOHNE MERINO SHEEP OF URUGUAY



Marques, C.B., De Barbieri, I., Montossi, F., Abella, I., Ramos, Z., Rovira, F., Monzalvo, C., Gimeno, D., Pintos, M.G., Ciappesoni, G.\*

gciappesoni@inia.org.uy

### INTRODUCTION

Dohne Merino (DM) is a synthetic breed that adaptates to different agroecological situations. Imported into Uruguay in 2002, this breed stands out for its fast growth, fine wool and potential use both as a pure breed and in crossbreeding.

The objective of this work was to estimate national genetic parameters for Birth weight (BWT), Weaning weight (WWT), Yearling weight (YWT), Scanning weight (SWT), Rib eye area (REA), Fat thickness (FAT), Greasy fleece weight (GFW), Clean fleece weight (CFW), Fibre diameter (FD) and Staple length (SL) of DM to develop a more precise evaluation and therefore a greater potential genetic progress of the breed.

#### MATERIALS AND METHODS

4331 )) animals

of **12** 

from 2

progenies

experimental stations

were

Data from

included in the fo

ESTIMATING
GENETIC
PARAMETERS

Table 1. Descriptive statistics of ages, growth, wool and in vivo carcass traits.

TRAIT	N	MEAN	STANDARD DEVIATION	RANGE
Weaning age (days)	2891	110	23.4	55-175
Scanning age (days)	1666	335	58.7	236-442
Yearling age (days)	2847	363	30.2	300-501
BWT (kg)	1828	5.0	1.11	2.00-8.70
WWT (kg)	2988	29,0	6.32	10.0-48.5
SWT (kg)	1735	54.4	14.53	22.5-95.0
YWT (kg)	3060	52.0	14.11	22.0-95.0
REA (cm²)	1735	10.8	3.32	3.1-22.99
FAT (mm)	1731	2.5	1.13	0.7–7.5
GFW (kg)	3060	3.04	0.894	0.97–9.9
CFW (kg)	2412	2.26	0.582	0.9-6.4
FD (µm)	2911	18.4	1.46	14.0-23.7
SL (cm)	2416	8.9	1.45	4.0–14.5

# RESULTS AND DISCUSSION

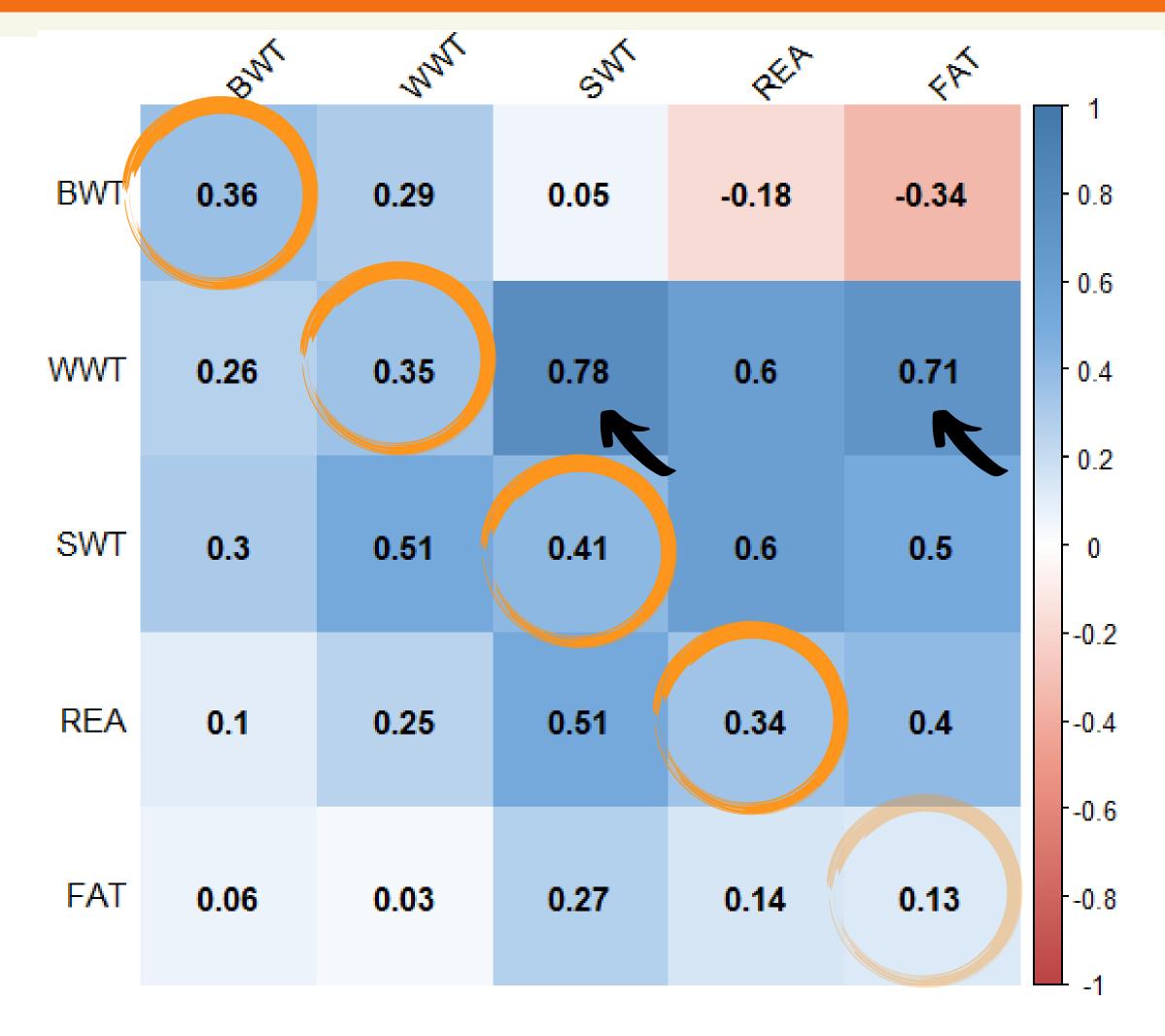


Figure 1 - Heritabilities (diagonal), genetic (above diagonal) and phenotypic correlations (below diagonal) of growth and carcass traits.

Figure 1 presents a moderate magnitude heritability (h²) for BWT, WWT, SWT and REA, while FAT presented a low magnitude h². Highly positive genetic correlations between WWT-SWT and WWT-FAT. Moderate genetic correlations between WWT-BWT, WWT-REA, SWT-REA and SWT-FAT, and negative genetic correlations between BWT-REA and BWT-FAT were found.

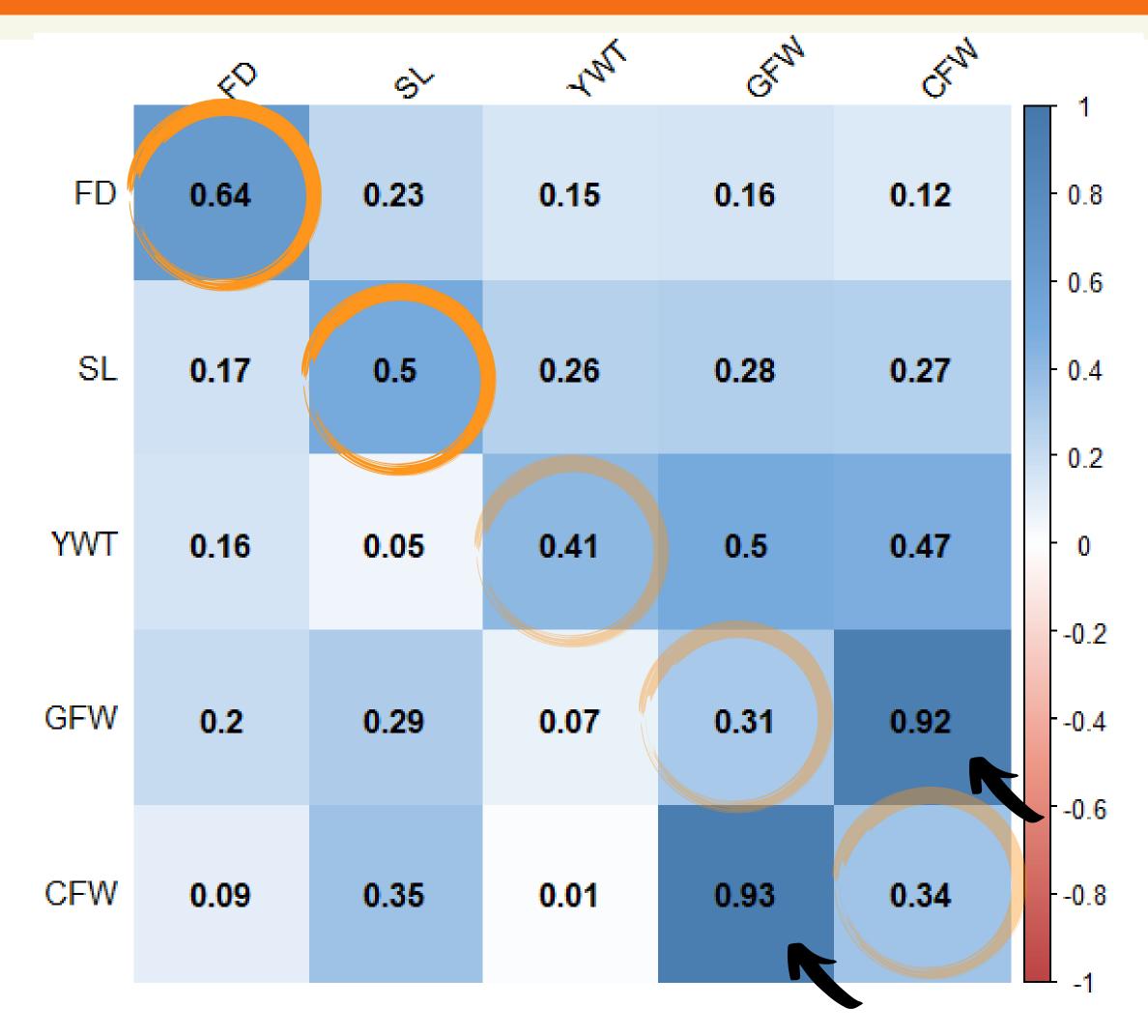


Figure 2 - Heritabilities (diagonal), genetic (above diagonal) and phenotypic correlations (below diagonal) of wool and YWT traits.

In **Figure 2**, a high h² for **FD** and **SL** was observed. This population also presented a moderate magnitude h² for **YWT**, **GFW**, and **CFW**, similarly to the reported in literature. There was a **very high** genetic and phenotypic correlation between **CFW** and **GFW**, poitive genetic correlations between **YWT-GFW**, **YWT-CFW**. Genetic correlations among the other wool traits were lower than mentioned above.



## CONCLUSION

Data evaluation in different geographic locations is a very important way to obtain assertiveness in research of a particular breed and location.

Our estimation of genetic parameters for growth, wool and carcass traits in DM sheep of Uruguay will help in developing an Uruguayan genetic evaluation of the breed.