

The use of seemingly unrelated regression to predict in vivo the carcass composition of lambs

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Outline

- 1 Introduction
- 2 Materials and methods
- 3 Results
- 4 Conclusions

Carcass quality

Good quality carcasses

- Reduced amount of fat
- But enough fat to guarantee: Good presentation and Protect the meat

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Carcasses value

- A carcass with an optimum composition: Highest price
- Composition moves away from the optimum: Value should suffer depreciation

Carcass classification

Aims of classification

- Base for fair payments
- Communication with consumers'

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Classification systems

- Accurate
- Based on simple models
- Predict muscle proportion (**not muscle weight**)

Published work

Common feature of published work

- Predict **lean meat weight** or **lean meat proportion**
- Several independent equations are **estimated separately by ordinary least squares (OLS)**
- Assumption of **independence** is not supported
 - Biological knowledge
 - Carcass compositional traits are correlated both phenotypically and genotypically
 - The equations for predicting carcass compositional traits are interrelated
 - Single-equation approach is inefficient from a statistical point of view

Multiple equations models

Set of equations

- 1 Set of equations **sharing a common error structure** with non-zero covariance is said to be **contemporaneously correlated**
- 2 **Seemingly Unrelated Regression (SUR) estimator**
 - Accounts for these contemporaneous correlations
 - Allows the **p dependent variables** to have **different sets of explanatory variables**
 - SUR method estimates the parameters of all equations simultaneously
 - Greater efficiency of the parameter estimates

Objectives

- 1 To compare alternative models for simultaneously predicting carcass compositional traits

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- 1 To compare alternative models for simultaneously predicting carcass compositional traits
- 2 To compare the efficiency of the OLS and SUR estimators

Animals

- 125 lambs - Churra Galega Bragançana
 - 82 males
 - 43 females

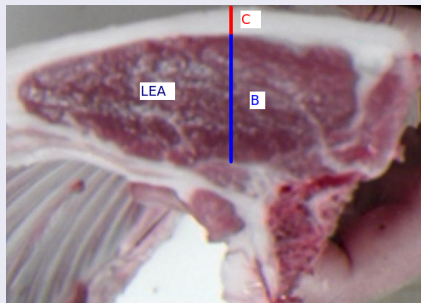
Slaughtering

- Carcasses weighted - HCW
- Carcasses were cooled at 4°C for 24h
- Left side dissected
 - Lean meat proportion (LMP)
 - Subcutaneous fat proportion (SFP)
 - Intermuscular fat proportion (IFP)
 - Bone plus remainder proportion (BP)
 - Kidney and knob channel fat proportion (KCFP)

Tissues thickness measurements

Subcutaneous fat thickness

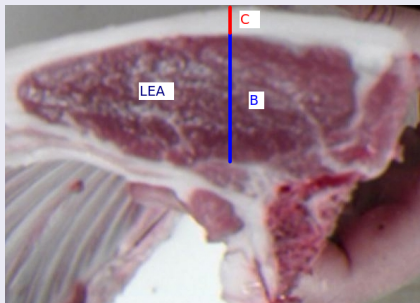
- C12: 12th and 13th ribs



Tissues thickness measurements

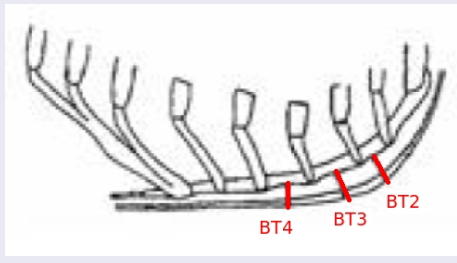
Subcutaneous fat thickness

- C12: 12th and 13th ribs



Breast bone tissues thickness

- E2: middle of the 2nd sternebrae



Statistical analysis

Models

- Three **multiple equations models** were developed to **simultaneously predict**: LMP, SFP, IFP, BP and KCFP
- All statistical analyses were undertaken using the **software R**
- **Add-on package systemfit**
- After estimating the **full models** by OLS and SUR
 - All explanatory variables that had a parameter with a marginal level of significance (P value) larger than 0.20 were removed.

Statistical analysis

Models fitting quality

- Coefficients of determination of estimation (R_e^2)
- standard error of the estimate (SEE)
- Standard errors (SE) of the estimated parameters

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Models validation

- Leaving-one-out **cross-validation procedure**
- Predicted residuals sum of squares (**PRESS**) statistic
- Coefficient of determination of prediction (R_p^2)

Statistical analysis

The base model: C12+E2

$$LMP = \alpha_0 + \alpha_1 HCW + \alpha_2 C12 + \alpha_3 E2 + \varepsilon_1 \quad (1)$$

$$SFP = \beta_0 + \beta_1 HCW + \beta_2 C12 + \beta_3 E2 + \varepsilon_2 \quad (2)$$

$$IFP = \gamma_0 + \gamma_1 HCW + \gamma_2 C12 + \gamma_3 E2 + \varepsilon_3 \quad (3)$$

$$BP = \delta_0 + \delta_1 HCW + \delta_2 C12 + \delta_3 E2 + \varepsilon_4 \quad (4)$$

$$KCFP = \theta_0 + \theta_1 HCW + \theta_2 C12 + \theta_3 E2 + \varepsilon_5 \quad (5)$$

Statistical analysis

The multiple equation model can be written as:

$$\begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_p \end{bmatrix} = \begin{bmatrix} X_1 & 0 & 0 & 0 \\ 0 & X_2 & 0 & 0 \\ 0 & 0 & \ddots & 0 \\ 0 & 0 & 0 & X_p \end{bmatrix} \begin{bmatrix} \beta_1 \\ \beta_2 \\ \vdots \\ \beta_p \end{bmatrix} + \begin{bmatrix} \varepsilon_1 \\ \varepsilon_2 \\ \vdots \\ \varepsilon_p \end{bmatrix} \quad (6)$$

Statistical analysis

Sub-models

- Sub-model: C12
with $\alpha_3 = \beta_3 = \gamma_3 = \delta_3 = \theta_3 = 0$
i.e. without the explanatory variable E2

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with $\alpha_2 = \beta_2 = \gamma_2 = \delta_2 = \theta_2 = 0$
i.e. without the explanatory variable C12.

Statistical analysis

OLS estimates of the entire system of equations:

$$\beta^{OLS} = (X'X)^{-1} X'y$$

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SUR estimates of the entire system of equations:

$$\beta^{SUR} = [X'\Omega^{-1}X]^{-1} [X'\Omega^{-1}Y]$$

- Ω^{-1} is a **weighting matrix** based on the covariance matrix of the error terms Σ .
- true error terms ε are unknown
 - replaced by observed residuals obtained from OLS estimates
 - i.e. $\hat{\varepsilon}_i = y_i - X_i\beta_i^{OLS}$
- Thus, a SUR model is an application of the **generalized least squares (GLS)** approach and the unknown residual covariance matrix is estimated from the data

Descriptive statistics

Table: Correlations among HCW, C12, E2, LMP, SFP, IFP, BP, and KCFP

Variable	C12	E2	LMP	SFP	IFP	BP	KCFP
HCW, kg	0.13	0.52	-0.14	0.20	0.47	-0.55	0.31
C12, mm	1	0.51	-0.36	0.63	0.25	-0.41	0.42
E2, mm		1	-0.61	0.77	0.52	-0.52	0.75
LMP, %			1	-0.84	-0.69	0.14	-0.73
SFP, %				1	0.56	-0.44	0.82
IFP, %					1	-0.63	0.55
BP, %						1	-0.48
KCFP, %							1

Three basic models

Table: Fitting quality for three basic models

Dep. var.	LMP			SFP		
Model	C12	E2	C12+E2	C12	E2	C12+E2
R_e^2	0.140	0.425	0.425	0.415	0.658	0.716
R_p^2	0.051	0.375	0.341	0.351	0.622	0.669
SEE	2.72	2.23	2.25	1.36	1.04	0.961

Three basic models

Table: Fitting quality for three basic models (Cont.)

Dep. var.	IFP			KCFP		
Model	C12	E2	C12+E2	C12	E2	C12+E2
R_e^2	0.252	0.323	0.323	0.238	0.576	0.576
R_p^2	0.164	0.239	0.220	0.164	0.544	0.523
SEE	1.64	1.56	1.58	0.452	0.337	0.342

Three basic models

Table: Fitting quality for three basic models

Dep. var.	BP		
Model	C12	E2	C12+E2
R_e^2	0.411	0.373	0.425
R_p^2	0.346	0.303	0.338
SEE	1.401	1.45	1.404

Residuals correlations

Table: Residuals correlations for model E2 estimated by ordinary least squares method (OLS)

	LMP	SFP	IFP	KCFP	BP
LMP	1	-0.709	-0.670	-0.505	-0.189
SFP		1	0.445	0.569	-0.242
IFP			1	0.337	-0.446
KCFP				1	-0.229
BP					1

Model E2a

Table: Fitting quality of model E2a

Dep. var.	LMP		SFP		IFP	
	OLS	SUR	OLS	SUR	OLS	SUR
R_e^2	0.425	0.413	0.657	0.649	0.323	0.317
R_p^2	0.375	0.382	0.636	0.631	0.239	0.242
SEE	2.23	2.25	1.03	1.04	1.56	1.56

Model E2a

Table: Fitting quality of model E2a

Dep. var.	KCFP		BP	
	OLS	SUR	OLS	SUR
R_e^2	0.533	0.533	0.373	0.371
R_p^2	0.534	0.534	0.303	0.302
SEE	0.344	0.344	1.445	1.448

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Conclusions

- 1 Novel approach to simultaneously predict carcass components using the SUR technique
- 2 The SUR estimator provides the lowest standard errors of the estimated parameters
- 3 SUR is a robust methodology for predicting the carcass composition
- 4 Results are relevant for implementing objective carcass classification systems to simultaneously predict carcass components
- 5 Our findings can have a positive effect on the meat industry, since the methodology **can improve decision support systems** by using **all carcass tissues** to determine the **price** as well as the **optimal processing** (e.g., refrigeration conditions) of the carcasses

THE END!

Thank You for Your Attention

Obrigado pela Atenção

Hvala na Pozornosti