

# Genetic parameters for live animal ultrasound measures, scrotal circumference, carcass and growth traits in Aberdeen Angus

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# OBJECTIVES

- Genetic parameters for
  - Live animal ultrasound measures
- Genetic correlations with
  - Carcass traits (SEUROP)
  - Growth traits

# DATA

## Live animal ultrasound measures

- Collected in 2019, 2020 and 2022
- Accredited ultrasound scanning technician
- Age 250 – 500 days
- 1,949 animals (1,021 bulls)



## Carcass traits (SEUROP classification)

- 2006 - 2021
- Commercial abattoirs
- Age 250 – 1080 days
- 7,672 animals



## Growth traits

- 2006 - 2022
- Collected by inspectors
- 36,958 animals with weaning weight
- 15,572 animal with yearling weight



## Ultrasound

Body weight at scanning (SCW)

Scrotal circumference (SC)

Rump fat thickness (P8FT)

Rib fat thickness (RBFT)

Eye muscle area (EMA)

Intramuscular fat content (IMF)

## Carcass

Carcass weight (CW)

Carcass conformation (CC)

Carcass fatness (CF)

## Growth

Adjusted weaning weight (AWW)

Adjusted yearling weight (AYW)

# MODEL EQUATIONS

	HYS	CLASS		REGRESSION		GENETIC ANIMAL
		SEX*TWIN	AGE OF DAM	AGE + AGE <sup>2</sup>	HETEROSIS	
Body weight at scanning (SCW)	✓	✓	✓	✓		✓
Scrotal circumference (SC)	✓		✓	✓		✓
Rump fat thickness (P8FT)	✓	✓	✓	✓		✓
Rib fat thickness (RBFT)	✓	✓	✓	✓		✓
Eye muscle area (EMA)	✓	✓	✓	✓		✓
Intramuscular fat content (IMF)	✓	✓	✓	✓		✓
Carcass weight (CW)	✓	✓		✓	✓	✓
Carcass conformation (CC)	✓	✓		✓	✓	✓
Carcass fatness (CF)	✓	✓		✓	✓	✓
Adjusted weaning weight (AWW)	✓	✓	✓		✓	✓
Adjusted yearling weight (AYW)	✓	✓	✓		✓	✓

# VARIANCE COMPONENT ESTIMATION

- AIREMLF90 (Ignacy Misztal et al.)
- Animal model

$$y_i = X_i \beta_i + Z_i u_i + e_i$$

$$\text{Var}(u) = G \times A$$

$$\text{Var}(e) = \begin{bmatrix} R_U \times I & 0 & R_{UG} \times I \\ 0 & R_C \times I & R_{CG} \times I \\ R_{UG} \times I & R_{CG} \times I & R_G \times I \end{bmatrix}$$

← ultrasound  
← carcass  
← growth

# VARIANCE COMPONENT ESTIMATION

	LIVE ANIMAL ULTRASOUND MEASURES						CARCASS TRAITS			GROWTH TRAITS	
	SCW	SC	P8FT	RBFT	EMA	IMF	CW	CC	CF	AWW	AYW
SCW	MULTITRAIT MODEL						TWO-TRAIT MODEL			TWO-TRAIT MODEL	
SC											
P8FT											
RBFT											
EMA											
IMF											
CW							MULTITRAIT MODEL			TWO-TRAIT MODEL	
CC											
CF											
AWW										MULTITRAIT MODEL	
AYW											

SCW - body weight at scanning, SC - scrotal circumference, P8FT - rump fat thickness, RBFT - rib fat thickness, EMA - eye muscle area, IMF - intramuscular fat content

CW - carcass weight, CC - carcass conformation, CF - carcass fatness

AWW - adjusted weaning weight, AYW - adjusted yearling weight

# HERITABILITIES (SD)

	LIVE ANIMAL ULTRASOUND MEASURES						CARCASS TRAITS			GROWTH TRAITS	
	SCW	SC	P8FT	RBFT	EMA	IMF	CW	CC	CF	AWW	AYW
SCW	0.69 (0.05)										
SC		0.77 (0.06)									
P8FT			0.49 (0.08)								
RBFT				0.44 (0.07)							
EMA					0.71 (0.01)						
IMF						0.25 (0.03)					
CW							0.57 (0.10)				
CC								0.37 (0.12)			
CF									0.33 (0.13)		
AWW										0.45 (0.01)	
AYW											0.38 (0.02)

SCW - body weight at scanning, SC - scrotal circumference, P8FT - rump fat thickness, RBFT - rib fat thickness, EMA - eye muscle area, IMF - intramuscular fat content

CW - carcass weight, CC - carcass conformation, CF - carcass fatness

AWW - adjusted weaning weight, AYW - adjusted yearling weight

# GENETIC CORRELATIONS (SD)

	LIVE ANIMAL ULTRASOUND MEASURES						CARCASS TRAITS			GROWTH TRAITS	
	SCW	SC	P8FT	RBFT	EMA	IMF	CW	CC	CF	AWW	AYW
SCW	0.69 (0.05)	0.38 (0.09)	0.54 (0.08)	0.61 (0.07)	0.90 (0.05)	0.70 (0.05)					
SC		0.77 (0.06)	0.19 (0.16)	0.20 (0.15)	0.42 (0.07)	0.60 (0.11)					
P8FT			0.49 (0.08)	0.99 (0.01)	0.35 (0.07)	0.87 (0.04)					
RBFT				0.44 (0.07)	0.43 (0.06)	0.88 (0.04)					
EMA					0.71 (0.01)	0.50 (0.06)					
IMF						0.25 (0.03)					
CW							0.57 (0.10)	0.41 (0.15)	0.92 (0.03)		
CC								0.37 (0.12)	0.19 (0.24)		
CF									0.33 (0.13)		
AWW										0.45 (0.01)	0.89 (0.01)
AYW											0.38 (0.02)

SCW - body weight at scanning, SC - scrotal circumference, P8FT - rump fat thickness, RBFT - rib fat thickness, EMA - eye muscle area, IMF - intramuscular fat content

CW - carcass weight, CC - carcass conformation, CF - carcass fatness

AWW - adjusted weaning weight, AYW - adjusted yearling weight

# GENETIC CORRELATIONS (SD)

	LIVE ANIMAL ULTRASOUND MEASURES						CARCASS TRAITS			GROWTH TRAITS	
	SCW	SC	P8FT	RBFT	EMA	IMF	CW	CC	CF	AWW	AYW
SCW	0.69 (0.05)	0.38 (0.09)	0.54 (0.08)	0.61 (0.07)	0.90 (0.05)	0.70 (0.05)	0.99 (0.01)	0.44 (0.19)	0.24 (0.22)	0.94 (0.01)	0.99 (0.01)
SC		0.77 (0.06)	0.19 (0.16)	0.20 (0.15)	0.42 (0.07)	0.60 (0.11)	0.99 (0.01)	0.97 (0.01)	0.85 (0.08)	-0.16 (0.14)	0.57 (0.22)
P8FT			0.49 (0.08)	0.99 (0.01)	0.35 (0.07)	0.87 (0.04)	0.61 (0.13)	0.28 (0.27)	0.99 (0.01)	0.16 (0.06)	-0.07 (0.11)
RBFT				0.44 (0.07)	0.43 (0.06)	0.88 (0.04)	0.80 (0.07)	0.78 (0.12)	0.99 (0.01)	0.24 (0.07)	0.02 (0.12)
EMA					0.71 (0.01)	0.50 (0.06)	0.99 (0.01)	0.98 (0.01)	0.03 (0.30)	0.47 (0.05)	0.43 (0.08)
IMF						0.25 (0.03)	-0.17 (0.01)	0.23 (0.02)	0.60 (0.06)	-0.51 (0.17)	-0.05 (0.05)
CW							0.57 (0.10)	0.41 (0.15)	0.92 (0.03)	0.55 (0.04)	0.70 (0.06)
CC								0.37 (0.12)	0.19 (0.24)	0.20 (0.01)	0.09 (0.01)
CF									0.33 (0.13)	0.27 (0.01)	0.15 (0.01)
AWW										0.45 (0.01)	0.89 (0.01)
AYW											0.38 (0.02)

SCW - body weight at scanning, SC - scrotal circumference, P8FT - rump fat thickness, RBFT - rib fat thickness, EMA - eye muscle area, IMF - intramuscular fat content

CW - carcass weight, CC - carcass conformation, CF - carcass fatness

AWW - adjusted weaning weight, AYW - adjusted yearling weight

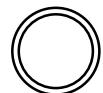
# CONCLUSIONS

The results support the potential value of live animal ultrasound measurements in Czech Aberdeen Angus breeding program to generate indicator traits for carcass quality.

Negative genetic correlation between intramuscular fat and adjusted weaning weight indicates that selection on high AWW could lead to lower IMF in the Aberdeen Angus.

- First results obtained on the basis of three years of measurements
- Expectation of improvement with more data
- Next step – incorporation of genomic information

THANK YOU FOR  
ATTENTION



ČESKÝ SVAZ CHOVATELŮ  
MASNÉHO SKOTU

