

# Towards an improved estimation of the biological components of residual feed intake in growing cattle

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# Outline

Motivation

Opening the black box

Objectives

Data available

Preliminary evidences

Phenotypic Models (PM)

Results PM

Genetic Models (GM)

Results GM

Conclusions

# Motivation

Feed is expensive !  
Environmental footprint ...



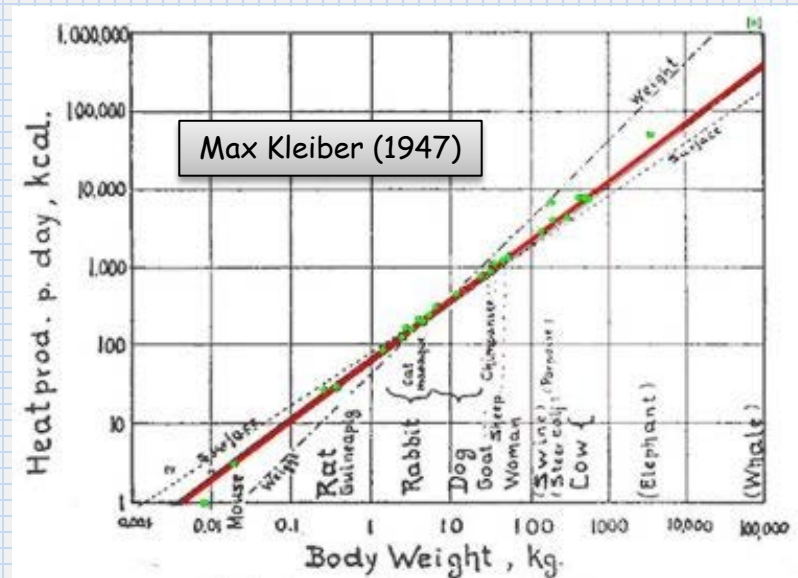
# Motivation

Many definitions...  
...ratio & residual traits



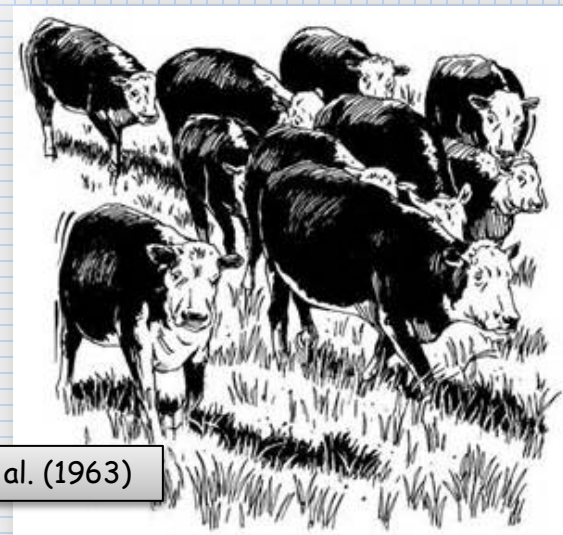
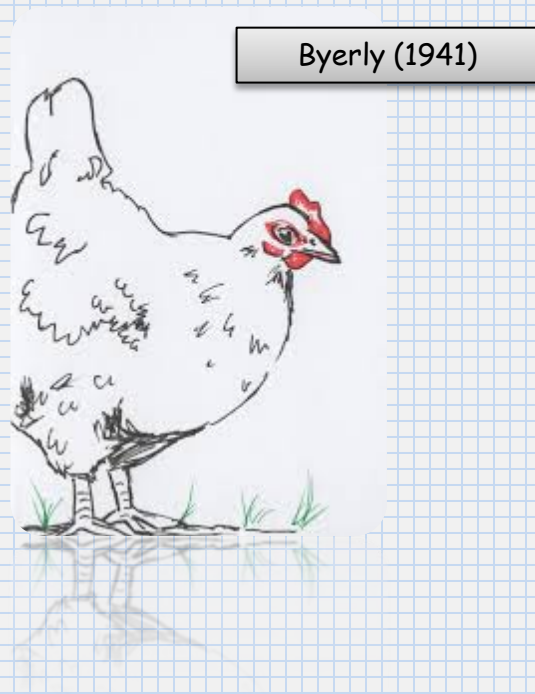
# Motivation

Feed conversion ratio  
is easy and simple, but...



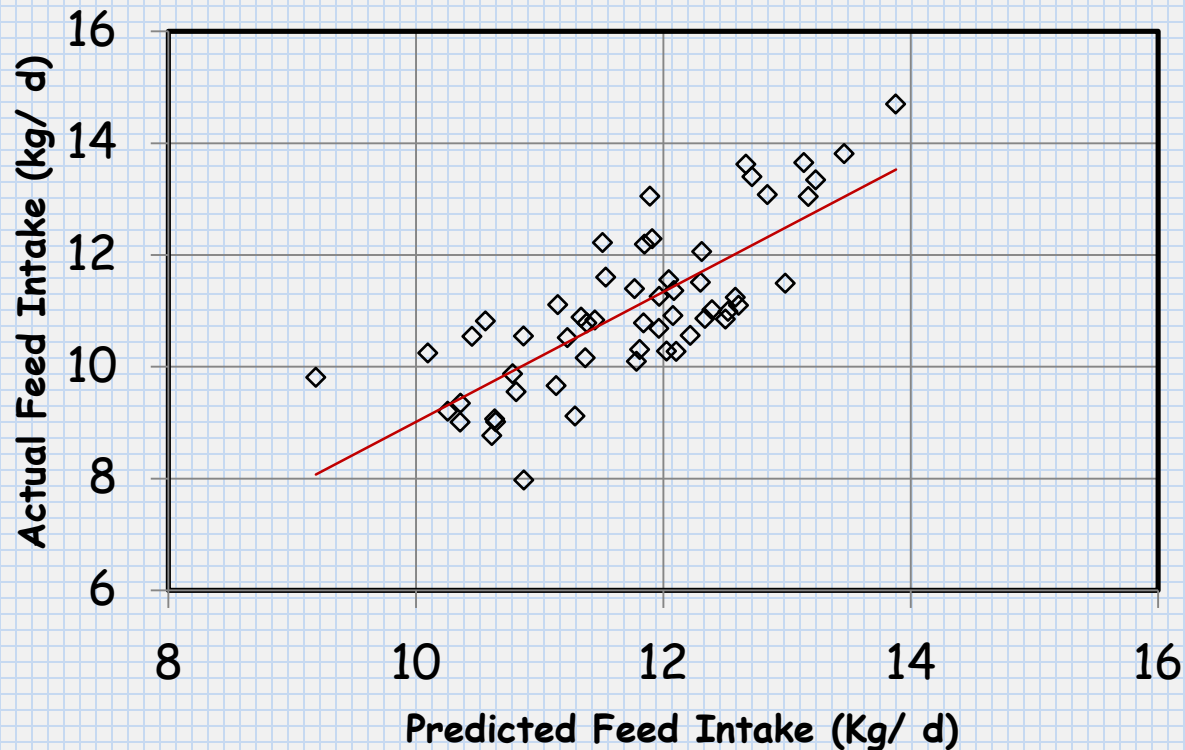
# Motivation

Residual feed intake  
has interesting properties, but...



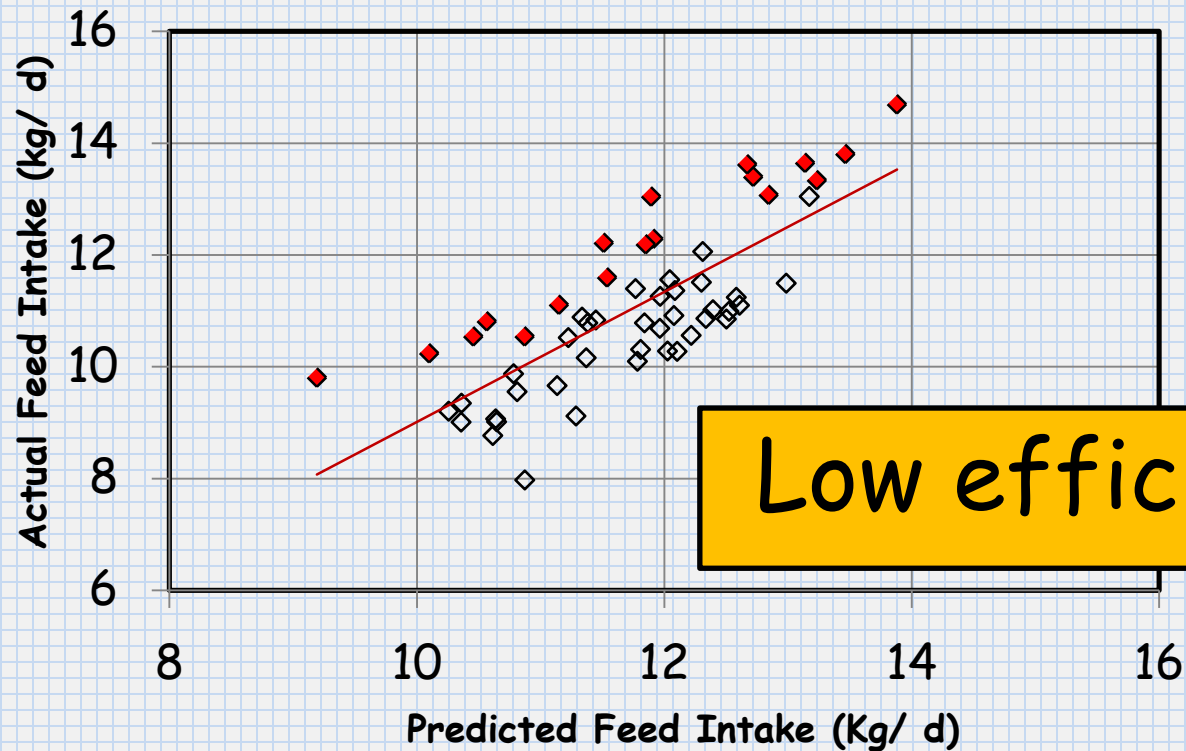
# Opening the black box

$$FI_i = \mu + b_0 \cdot BW_i^{0.75} + b_1 \cdot ADG_i + e_i$$



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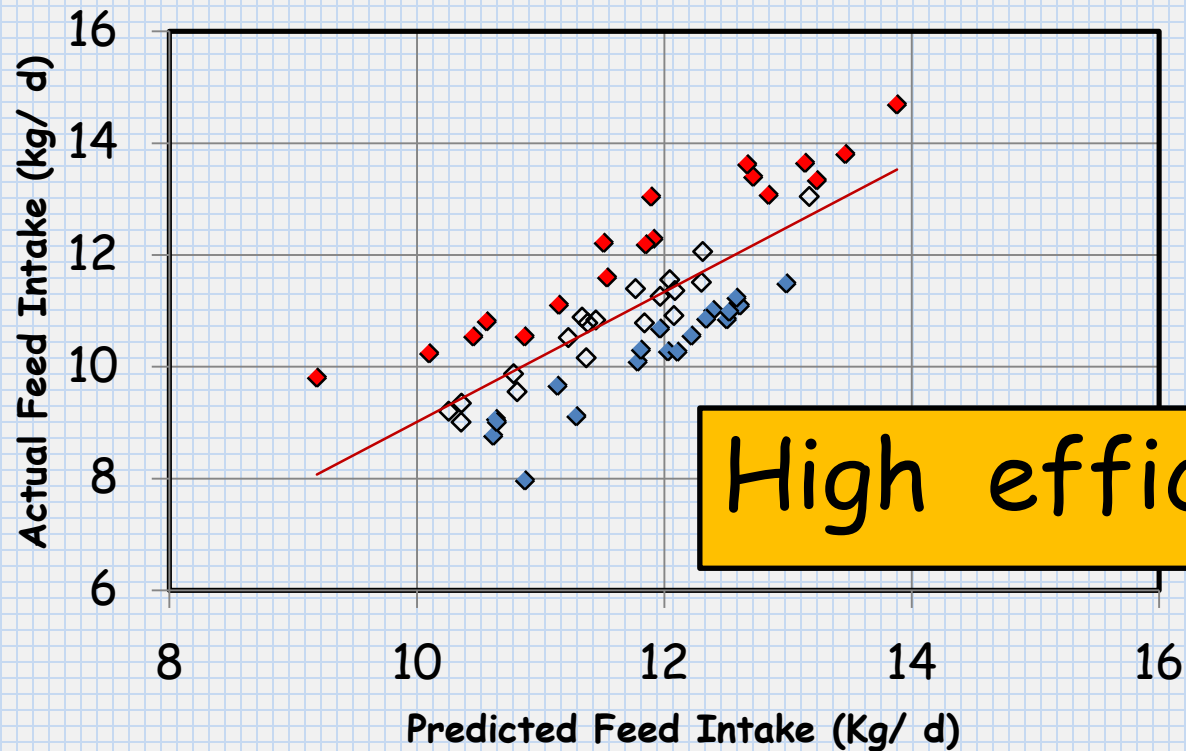
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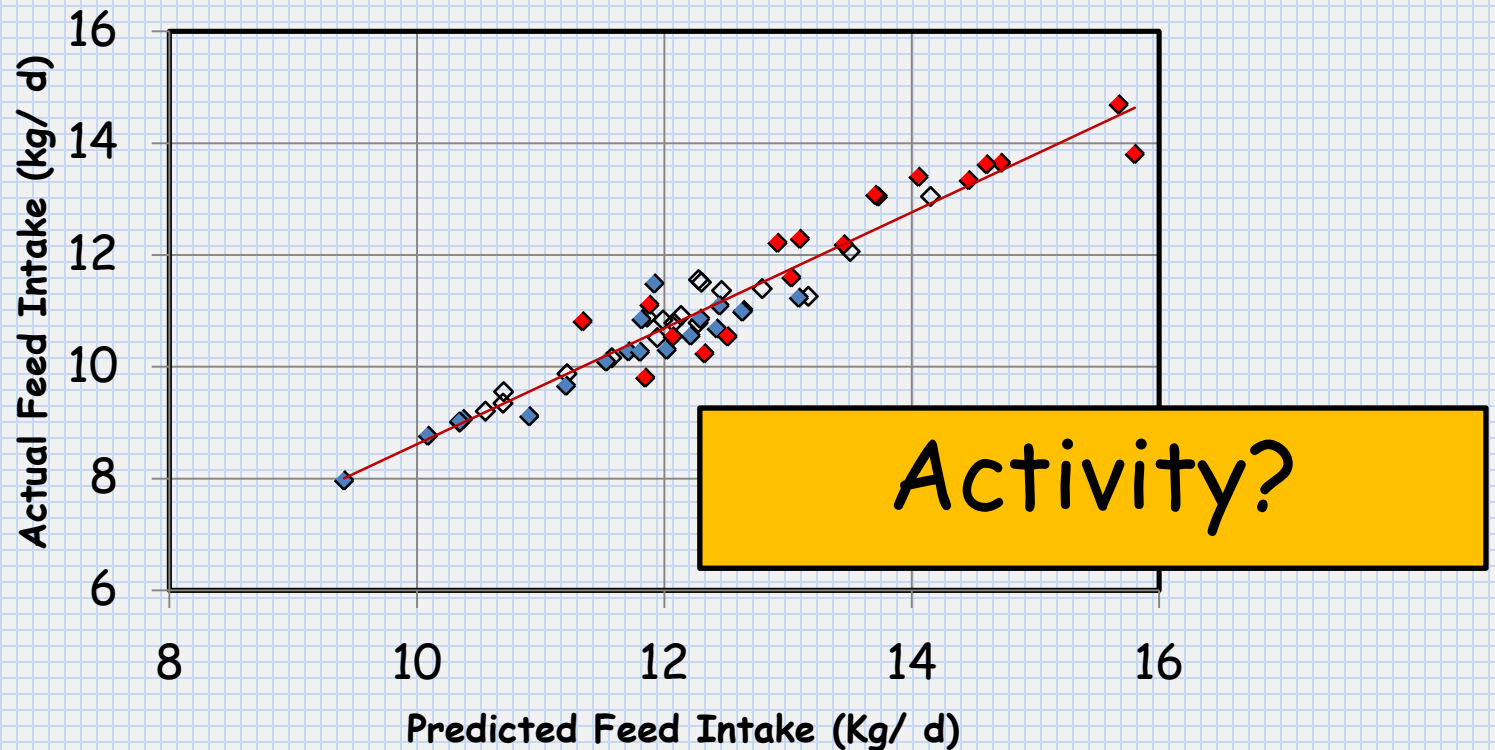
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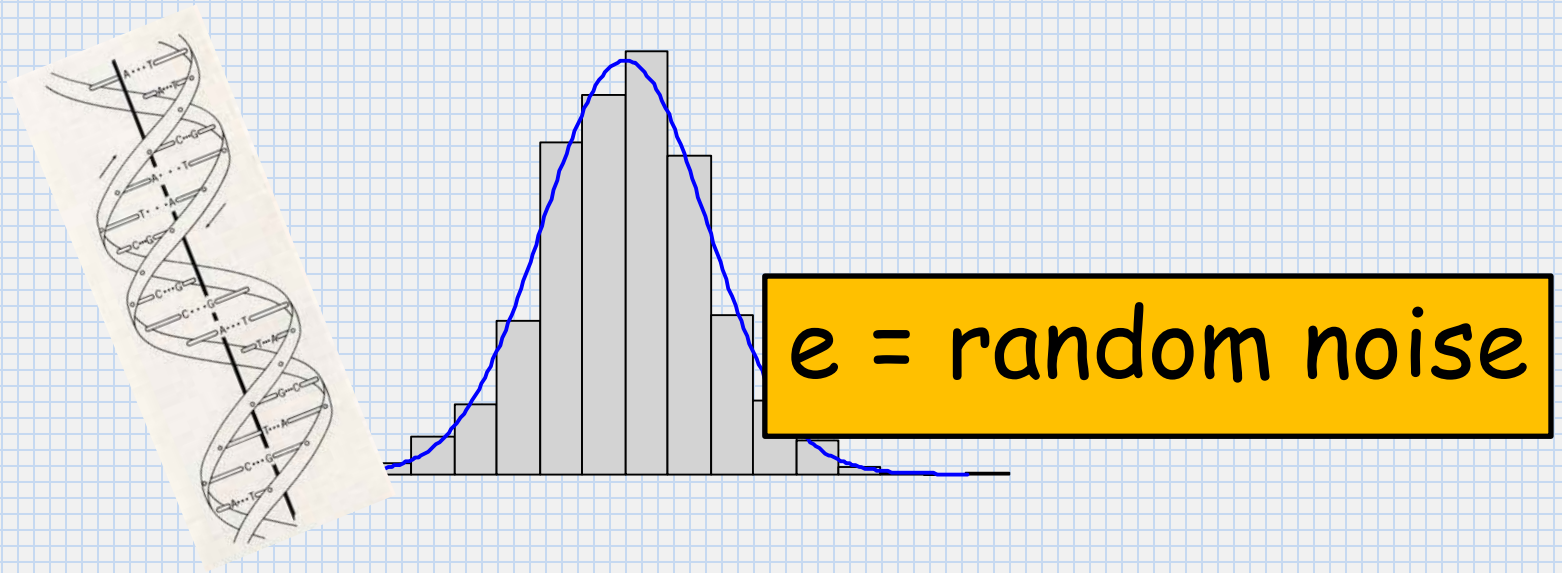
# Opening the black box

$$FI_i = \mu + b_0 \cdot BW_i^{0.75} + b_1 \cdot ADG_i + b_2 \cdot FAT_i + e_i$$



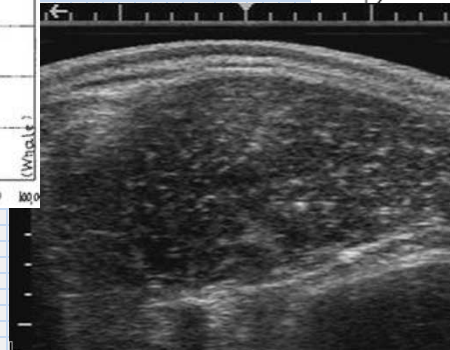
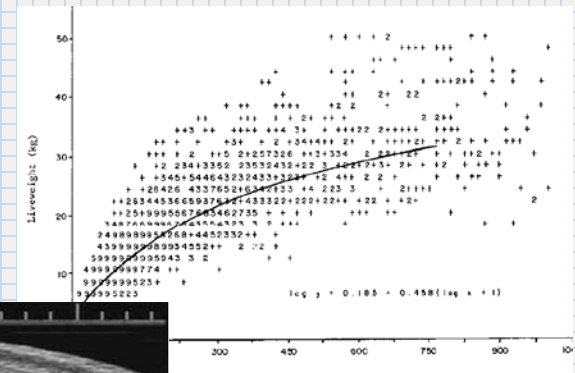
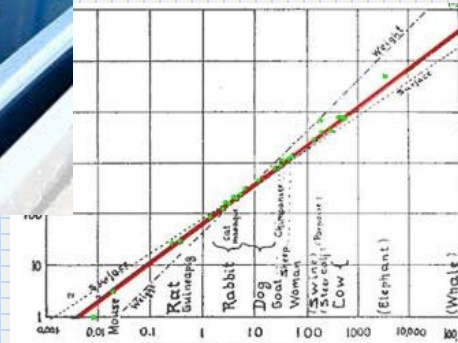
# Opening the black box

Correct model &  
variables are error free



# Objectives

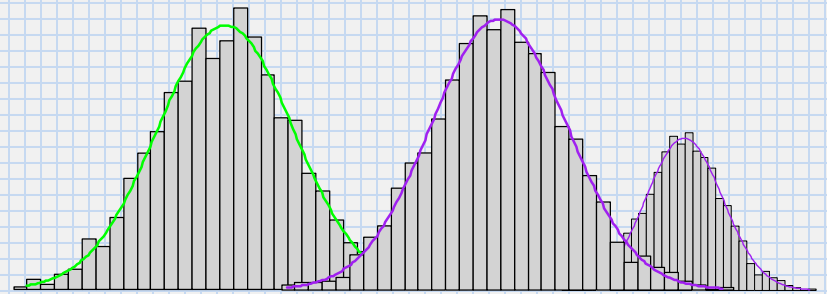
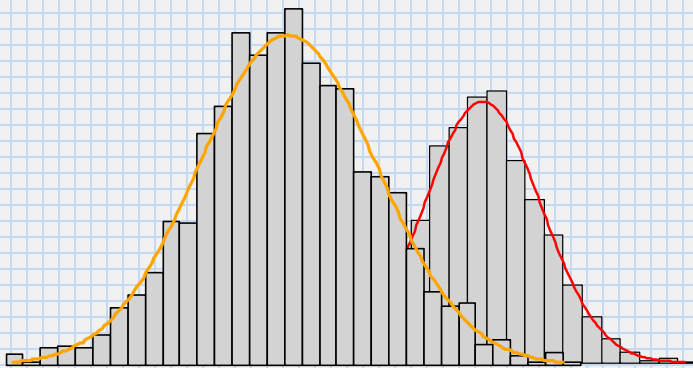
Precise estimate feed intake  
from limited information



# Objectives

Residuals are random noise

$$FI_i = \mu \cdot I_i (b_0 + b_1) \cdot BW_i^{0.75} + (b_2 + b_3) \cdot ADG_i + e_i$$





# Data available

1,963 bulls

69 Contemporaneous Group

314 d old & 456 kg

70 d on test

CI, BW\* and FAT

\*MTW & ADG



# Preliminary evidences

Table 1 - Summary statistics by breeds

<b>Traits</b>	<b>AN</b> n = 183	<b>CH</b> n = 485	<b>HE</b> n = 100	<b>LI</b> n = 821	<b>SI</b> n = 374
<b>CI</b> (kg/ d)	11.71 <sup>c</sup>	11.38 <sup>b</sup>	11.56 <sup>bc</sup>	10.19 <sup>a</sup>	12.24 <sup>d</sup>
<b>MTW</b> (kg <sup>0.75</sup> )	108.4 <sup>a</sup>	117.2 <sup>b</sup>	107.8 <sup>a</sup>	108.3 <sup>a</sup>	118.1 <sup>b</sup>
<b>ADG</b> (kg/ d)	1.63 <sup>a</sup>	1.75 <sup>b</sup>	1.74 <sup>b</sup>	1.60 <sup>a</sup>	1.74 <sup>b</sup>
<b>FAT</b> (cm)	0.552 <sup>c</sup>	0.256 <sup>a</sup>	0.559 <sup>c</sup>	0.249 <sup>a</sup>	0.316 <sup>b</sup>
<b>RFI</b> (kg/ d)	0.332 <sup>c</sup>	-0.124 <sup>ab</sup>	0.025 <sup>b</sup>	-0.205 <sup>a</sup>	0.443 <sup>c</sup>

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# Phenotypic Models (PM)

## Basic models

$$[P1] \text{CI}_i = b_0 \cdot \text{MTW}_i + e_i$$

$$[P2] \text{CI}_i = b_0 \cdot \text{MTW}_i + b_1 \cdot \text{ADG}_i + e_i$$

$$[P3] \text{CI}_i = b_0 \cdot \text{MTW}_i + b_1 \cdot \text{ADG}_i + b_2 \cdot \text{FAT}_i + e_i$$

# Phenotypic Models (PM)

No great expectations

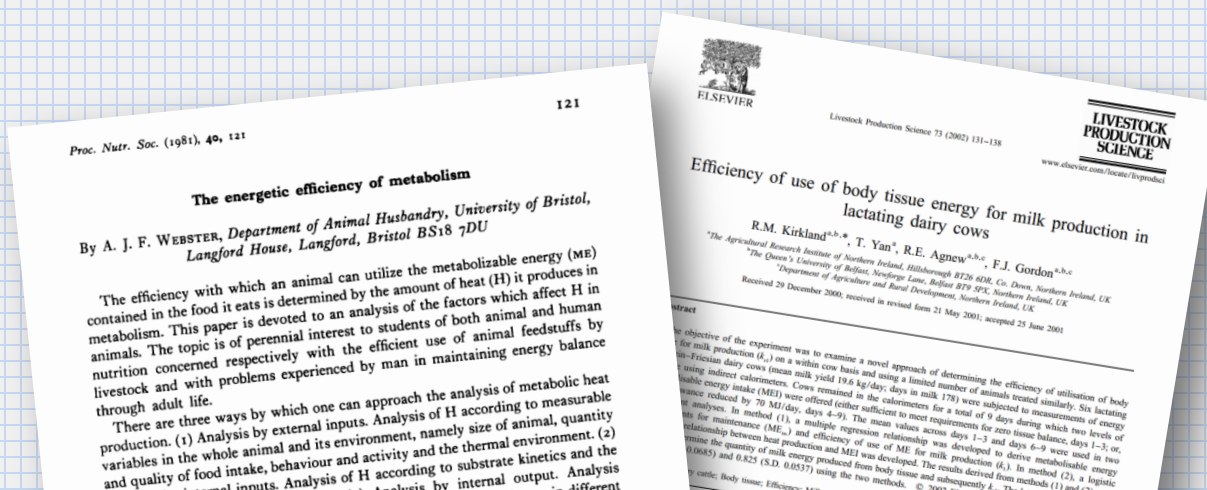
$$[P5] \text{ CI}_i = [P3] + b_3 \cdot \text{MTW} \times \text{ADG}_i + e_i$$

$$\text{MAYBE} = \frac{\text{YES}}{\text{NO}} + \frac{\text{NO}}{\text{YES}}$$

# Phenotypic Models (PM)

Composition affects maintenance

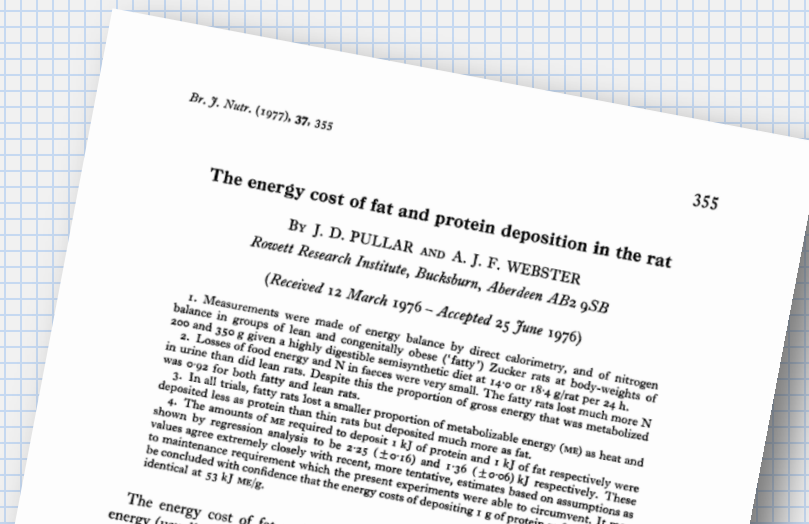
$$[P4] CI_i = [P3] + b_3 \cdot MTW \times FAT_i + e_i$$



# Phenotypic Models (PM)

Composition of gain has different cost

$$[P6] CI_i = [P3] + b_3 \cdot ADG \times FAT_i + e_i$$



# Results PM

Table 2 - Partial regression coefficients

Model	MTW	ADG	FAT	MTW×ADG	MTW×FAT	ADG×FAT	R <sup>2</sup>
P1	0.093	.	.	.	.	.	0.691
P2	0.081	1.887	.	.	.	.	0.760
P3	0.080	1.899	1.132	.	.	.	0.764
P4	0.104	3.471	1.167	-0.014	.	.	0.765
P5	0.080	1.900	1.254 <sup>NS</sup>	.	-0.001 <sup>NS</sup>	.	0.764
P6	0.080	1.831	0.758 <sup>NS</sup>	.	.	0.213 <sup>NS</sup>	0.764

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# Genetic Models (GM)

P3 was the base model

$$[G1] CI_i = [P3] + a_i + e_i$$

$$[G2] CI_i = (b_0 + b_{0a}) \cdot MTW_i + \dots + a_i + e_i$$

$$[G3] CI_i = (b_1 + b_{1a}) \cdot ADG_i + \dots + a_i + e_i$$

...

$$[G8] CI_i = (b_0 + b_{0a}) \cdot MTW_i + (b_1 + b_{1a}) \cdot ADG_i + (b_2 + b_{2a}) \cdot FAT_i + a_i + e_i$$

# Results GM

Table 3 - Additive and residual variances

Model	$a_i$	$b_{0a} \cdot \text{MTW}$	$b_{1a} \cdot \text{ADG}$	$b_{2a} \cdot \text{FAT}$	$e_i$	$h^2$	AIC
G1	0.317	.	.	.	0.315	0.50	1174.6
G2	0	0.00004	.	.	0.178	0.72	1137.3
G3	0.222	.	0.035	.	0.306	0.52	1172.8
G4	0.299	.	.	0.212	0.308	0.51	1175.2
G5	0	0.00004	0	.	0.178	0.72	1139.3
G6	0	0.00004	.	0.173	0.170	0.73	1138.1
G7	0.218	.	0.032	0.137	0.302	0.52	1174.2
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# Conclusions

## Limited information limits modelling

- Costs related to composition
  - Animals close to maturity
  - Dairy cows

## Random regression

- Differences in CI related to MTW
  - Similar to findings on gene expression
  - Probably less complex to understand