



## Improving carcass traits by selection in five beef cattle breeds

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# Study objective

## Why to improve carcass composition?

- Effective production of animal protein for human consumption
- Control of carcass quality

## Main beef breeds in Finland

British breeds: Hereford, Aberdeen Angus

Swiss breed: Simmental

French breeds: Charolais, Limousin

## Objectives

- 1) Estimate heritabilities and genetic correlations for carcass weight, carcass conformation and carcass fat for five main breeds
- 2) Compare alternative selection scenarios to quantify the way genetic correlations constrain breeding of the three traits



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# Data

Carcass weight, conformation and fat recorded in five slaughter houses

Carcass conformation and fat recorded with EUROP scoring:

- Conformation (1-15): 1 = poor and 15 = extensive muscularity
- Fat (1-5): 1 = low fat and 5 = extensive fat

**Breeding objective:** Increase weight and conformation, avoid extensive fat

| <b>Breed</b> | <b>Phenotyped animals</b> | <b><i>n</i> offspring per bull<br/>(mean; range)</b> | <b><i>n</i> herds per bull<br/>(mean; range)</b> |
|--------------|---------------------------|--|--|
| Hereford     | 19539                     | 14.1 (1-181)   | 2.8 (1-50)                                       |
| Ab. Angus    | 13598                     | 13.3 (1-99)  | 2.9 (1-55)                                       |
| Simmental    | 6879                      | 12.4 (1-151)   | 2.7 (1-34)                                       |
| Charolais    | 13611                     | 13.3 (1-134)   | 2.7 (1-45)                                       |
| Limousin     | 15072                     | 14.8 (1-290)   | 2.6 (1-50)                                       |

+ pedigree back to 60's

# Statistical model

For carcass weight, conformation and fat:

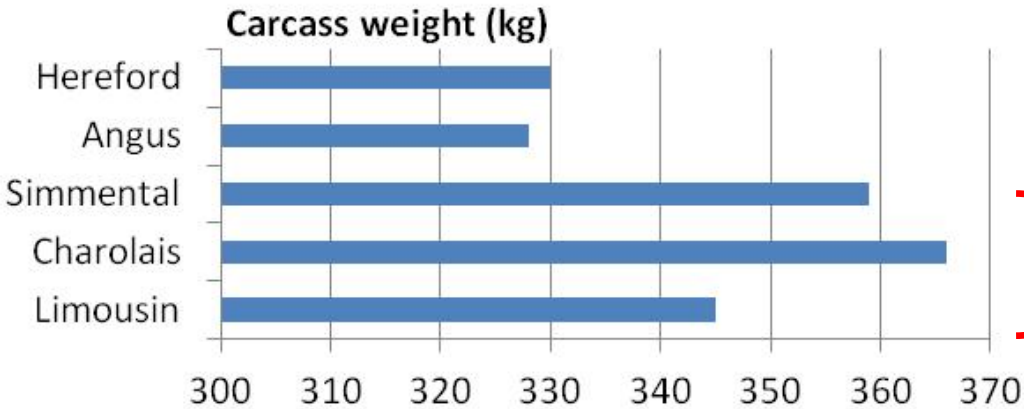
## Random effects

- $\text{Animal}_j$  random genetic effect of an animal  $j$  ( $j = 1$ -nro of animals)
- $\text{Error}_{ijklmno}$  random residual term

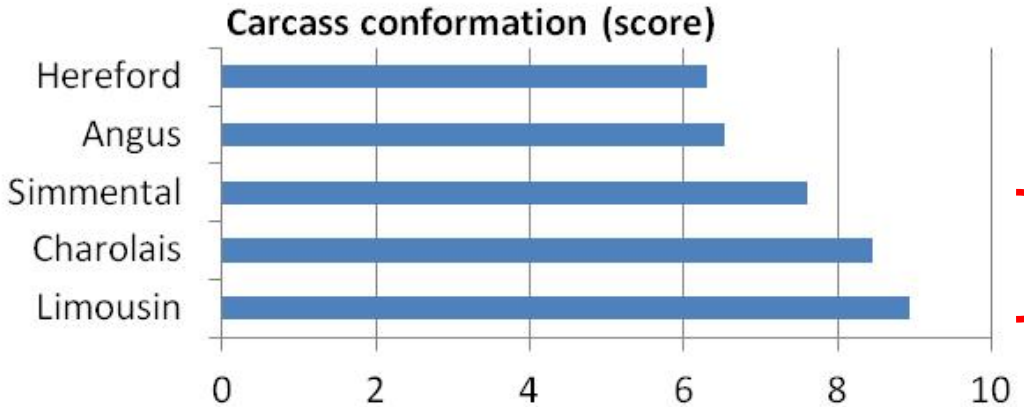
## Fixed effects

- $\text{HerdYear}_k$  herd-year interaction ( $k = 1$ -number of combinations)
- $\text{DamAge}_l$  age class of a dam of an individual  $j$  ( $l = 1$ -7)
- $\text{BirthSeason}_m$  season of the birth date ( $m = 1$ -5)
- $\text{Twin}_n$  twin or as a singleton birth ( $n = 1$ -2)
- $\text{Gender}_o$  gender ( $o = 1$ -2)
- $b_{\text{Age1}}(\text{Gender})$  linear regression coefficient of age for each gender
- $b_{\text{Age2}}(\text{Gender})$  quadratic regression coefficient of age for each gender

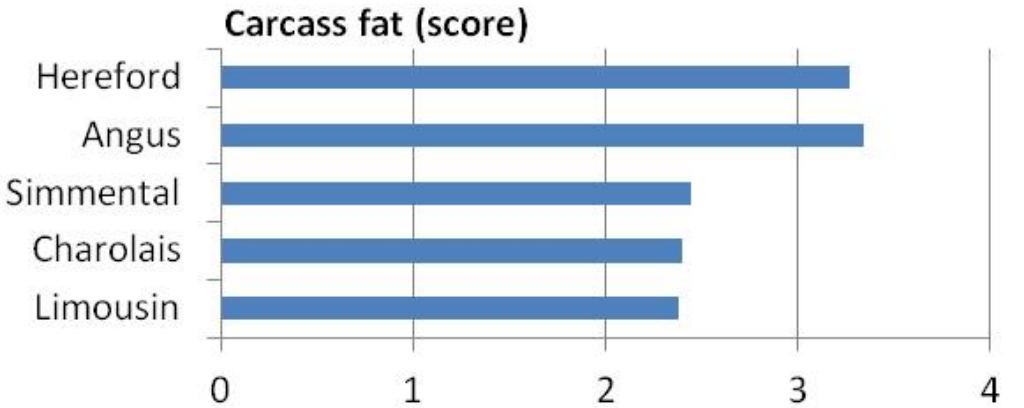
# Trait means



High carcass weight

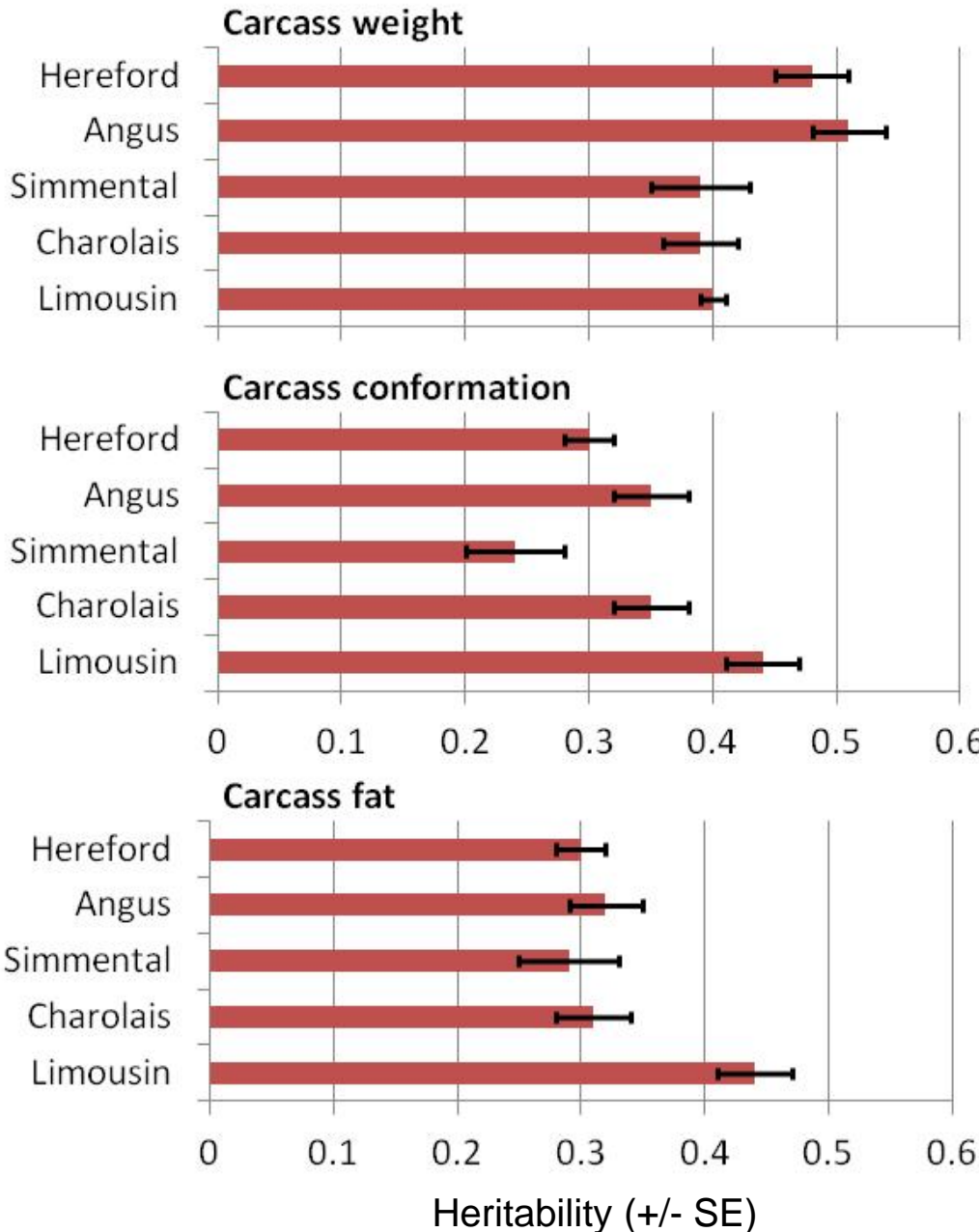


High muscularity



Low fat

# Trait heritability



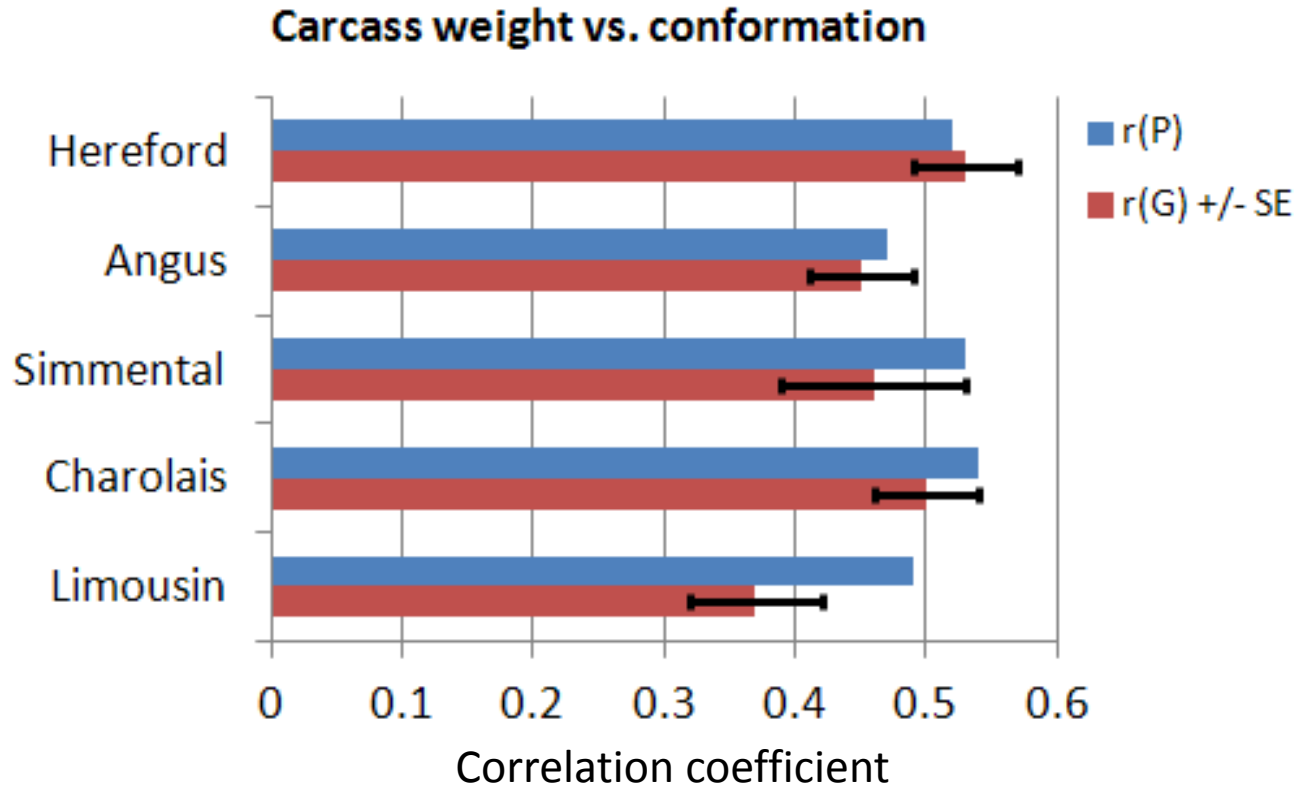
Our data, average  $h^2$  of 0.43 (weight), 0.34 (conformation), and 0.33 (fat)

In literature, average  $h^2$  of 0.31, 0.23, and 0.21

(More O'Ferrall et al. 1989; Dijkstra et al. 1990; Hirooka et al. 1998; van der Werf et al. 1998; Liinamo et al. 1999; Parkkonen et al. 2000; Eriksson et al. 2003; Hickey et al. 2011; Vesela et al. 2011)

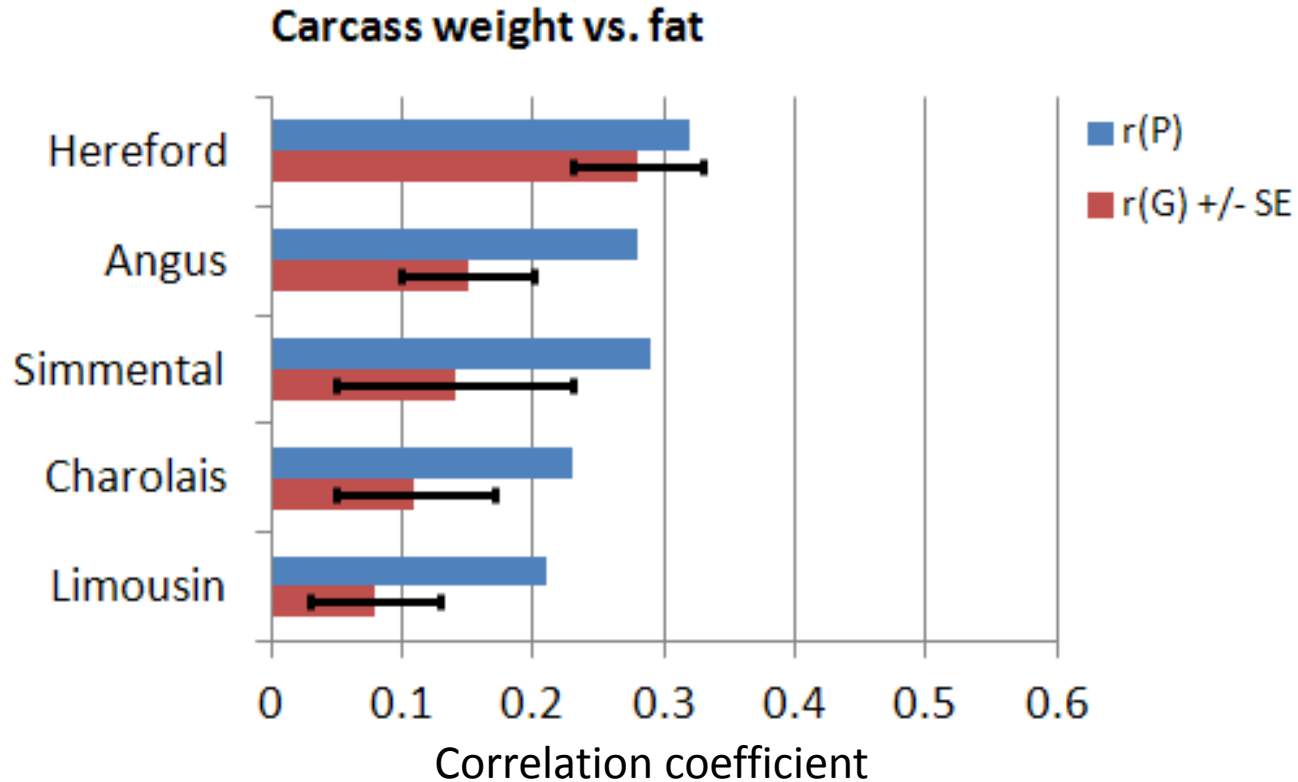
➡ Pattern the same, overall level higher in our data

# Trait correlations



 High weight – High muscularity  
Favourable for all breeds

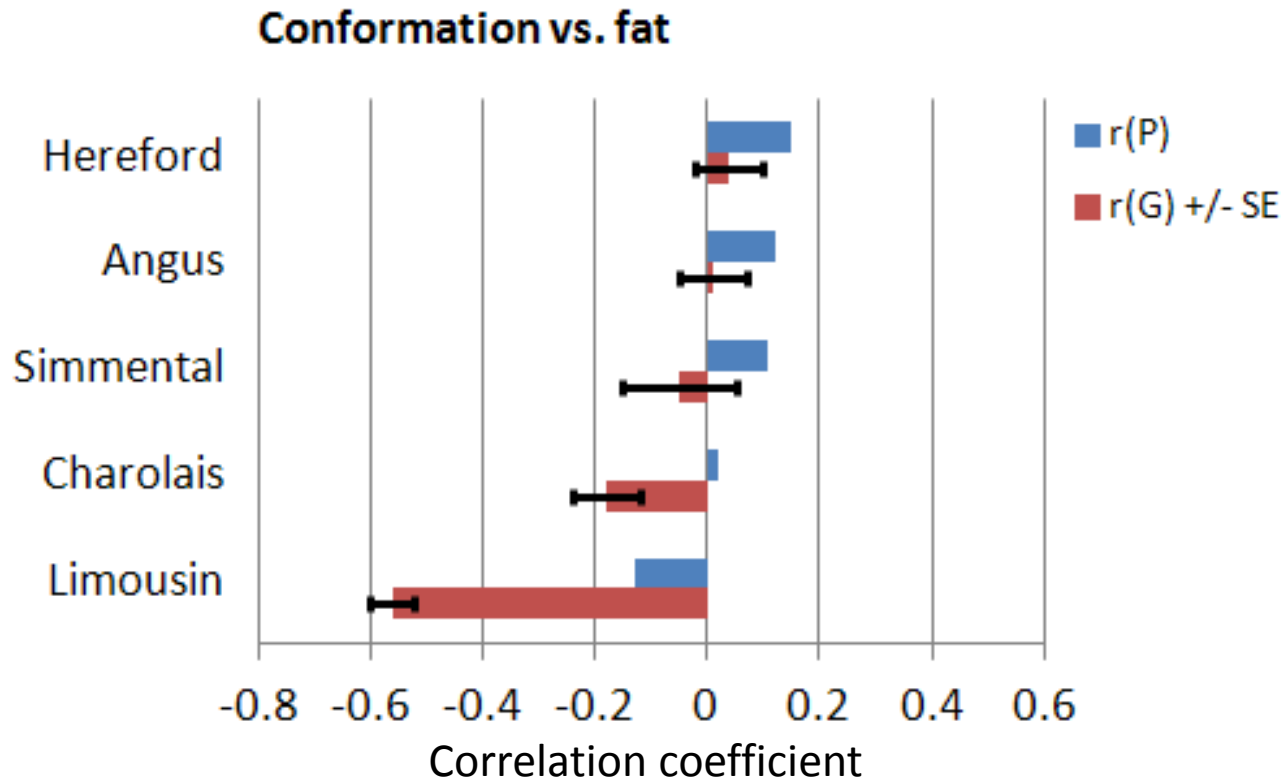
# Trait correlations



➔ High weight – High fat  
Unfavourable especially in British breeds



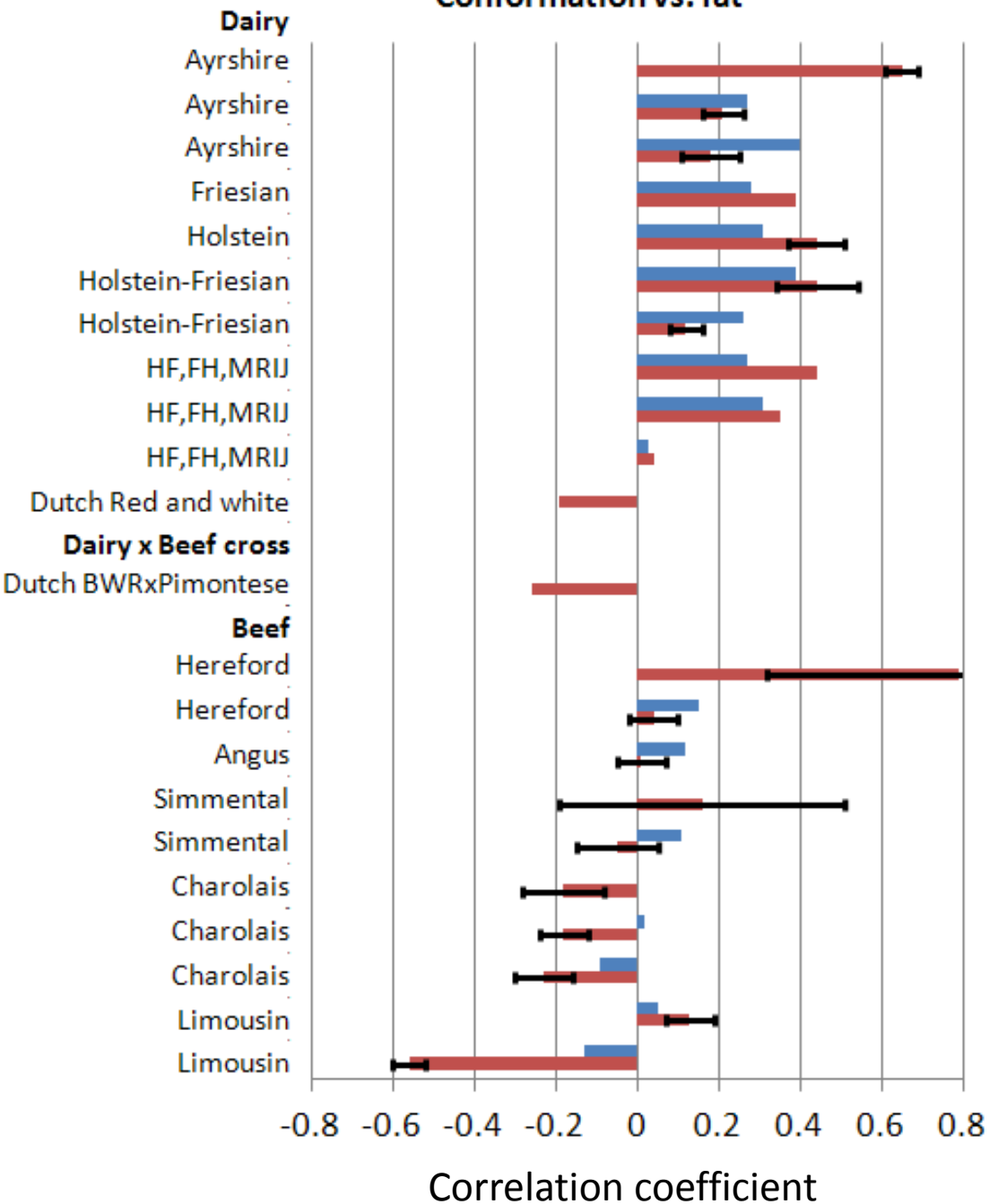
# Trait correlations



➔ More favourable in French breeds (High muscularity and Low fat - relationship)

## Conformation vs. fat

## Breed differences



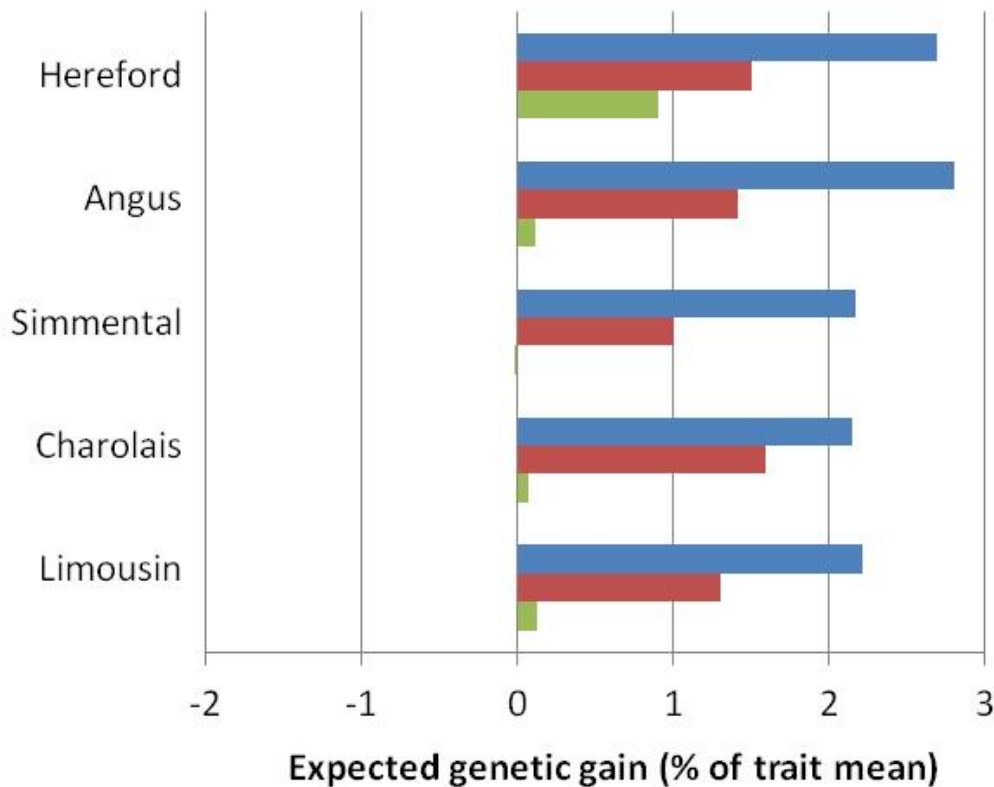
■ r(P)  
■ r(G) +/- SE

➔ In dairy, unfavourable 'high conformation-high fat' relationship

➔ In large body-sized beef, favourable 'high conformation-low fat' relationship

References (More O'Ferrall et al. 1989; Dijkstra et al. 1990; Hirooka et al. 1998; van der Werf et al. 1998; Fouilloux et al. 1999; Liinamo et al. 1999; Parkkonen et al. 2000; Eriksson et al. 2003; Hickey et al. 2011; Kause et al. EAAP)

# Selection response: Selection for weight



■ Carcass weight  
■ Conformation  
■ Fat

Index theory calculations for genetic gain (Hazel 1943)

Mass selection with intensity = 0.5

Economic values used:

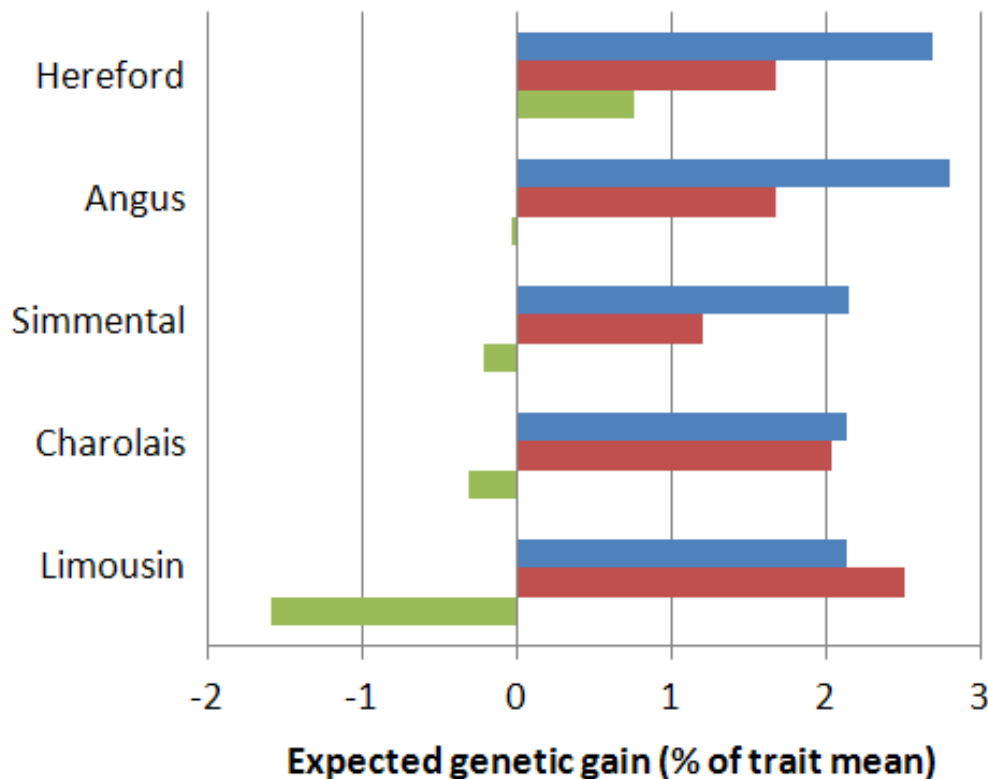
$a_{\text{Weight}} = 1 \text{ €}$

$a_{\text{Conformation}} = 0 \text{ €}$

$a_{\text{Fat}} = 0 \text{ €}$

➡ Consistent increase in weight and conformation (modest)

# Selection response: Economic values



■ Carcass weight  
■ Conformation  
■ Fat

Index theory calculations for genetic gain (Hazel 1943)

Mass selection with intensity = 0.5

Economic values used for Irish production system (Amer et al. 2001; Evans et al. 2012)

$a_{\text{Weight}} = 2.95 \text{ €}$

$a_{\text{Conformation}} = 14.77 \text{ €}$

$a_{\text{Fat}} = -7.86 \text{ €}$

➔ Breed-specific responses with carcass fat decreasing in continental breeds

# Conclusions

Genetic variation exists for carcass weight, carcass conformation and carcass fat

Simultaneous improvement of carcass weight and conformation easy in all breeds

For fat, unfavourable correlation with carcass weight especially in Hereford (British breed)

Breed difference - Correlations of fat more favourable in large-body sized and muscular continental breeds



