



Genetic Parameters for Beef Carcass Cuts Yield in Ireland

**T. Pabiou,^{a,*} F. Fikse^b, A. Näsholm^c, A. Cromie^a, M.D. Drennan^d,
M.G. Keane^d, and D. Berry^e**

^a *The Irish Cattle Breeding Federation, Shinagh House, Bandon, Co. Cork, Ireland.*

^b *Swedish University of Agricultural Sciences, Interbull Centre, Department of Animal Breeding and Genetics, Uppsala, Sweden.*

^c *Swedish University of Agricultural Sciences, Department of Animal Breeding and Genetics, Uppsala, Sweden.*

^d *Grange Research Center, Teagasc, Dunsany, Co. Meath, Ireland.*

^e *Moorepark Dairy Production Research Center, Teagasc, Fermoy, Co. Cork, Ireland.*





Introduction: Yield & Value?

- Different € value cuts in a carcass

Sirloin = €€€€

Shoulder = €€

- Using dissections results

→ genetic analysis to detect
possible benefits for beef farmers



16-18 June 2008

Interbull Meeting – Niagara Falls - USA



Two Raw Datafiles

■ Commercial data

- ◆ From an Irish supermarket
- ◆ 11 000 animals
 - ◆ *slaughtered from 1996 to 2005*
 - ◆ *1 carcass => up to 16 different cuts*

■ Experimental data

- ◆ From Teagasc beef research center
- ◆ 500 animals
 - ◆ *slaughtered from 2000 to 2008*
 - ◆ *1 carcass => 23 different cuts*



16-18 June 2008

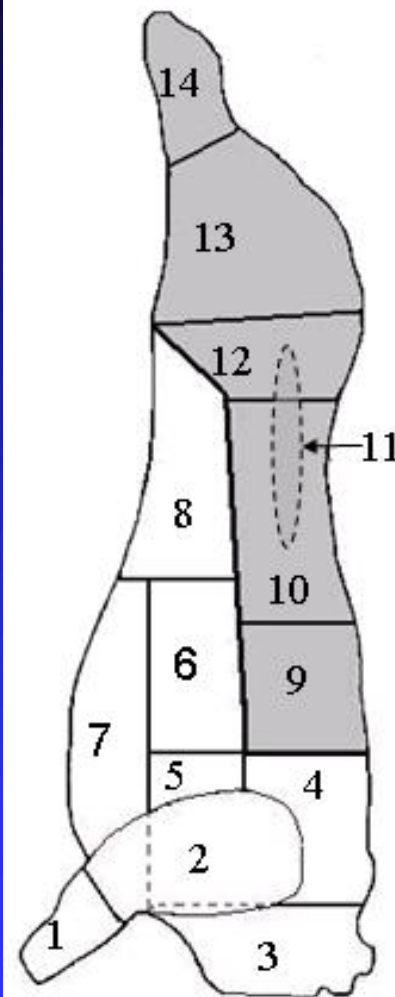
Interbull Meeting – Niagara Falls - USA



The Different Cuts



- Higher & Lower value cuts
 - ◆ Graph based on the experimental dissections



Forequarter

1. Front shin
2. Shoulder
3. Neck
4. Chuck
5. Ribs 1 to 5
6. Ribs 6 to 13
7. Brisket
8. Flank

Hindquarter

9. Rib Roast
10. Strip-loin
11. Fillet (inside)
12. Sirloin
13. Round
14. Back shin



The Models

	Commercial	Experimental
Model	Animal	Animal
Pedigree (breed grp)	6 250 (6)	8 300 (7)
Fixed effects (levels)	Sex (3) C. Groups (83)	Dam age (4) C Groups (11)
Covariates	Slaughter age Sex x Slaug. age	Heterosis Slaughter age Sex x Slaug. age
Packages	SAS DMU ASREML	SAS ASREML



Results: Heritability



	Commercial		Experimental	
	N	h^2 (s.e.)	N	h^2 (s.e.)
Carc. Weight (kg)	635	0.59 (0.200)	373	0.42 (0.216)
Carc. Conf. (1-15)	345	0.78 (0.265)	373	0.44 (0.224)
Carc. Fat (1-15)	345	0.63 (0.264)	373	0.31 (0.189)
Shoulder (kg)	635	0.61 (0.195)	373	0.75 (0.238)
Strip-loin (kg)	523	0.41 (0.217)	373	0.45 (0.231)
Sirloin (kg)	635	0.55 (0.199)	373	0.61 (0.227)
Round (kg)	635	0.42 (0.293)	373	0.84 (0.247)
Fillet (kg)	520	0.62 (0.204)	373	0.32 (0.214)



Results: Correlations

■ Experimental dataset

	Correlation
	≤ 0.0
	$>0.0 \ \& \ \leq 0.2$
	$>0.2 \ \& \ \leq 0.5$
	$>0.5 \ \& \ \leq 0.7$
	> 0.7



Genetic Phenotypic	C. Weight	C. Conf.	C. Fat	Shoul- der	Strip- loin	Sirloin	Round	Fillet
C. Weight								
C. Conf.								
C. Fat								
Shoulder								
Strip-loin								
Sirloin								
Round								
Fillet								



16-18 June 2008

Interbull Meeting – Niagara Falls - USA



Conclusions

- Genetic parameters
 - ◆ Moderate to high heritability & correlations
 - ◆ Stronger estimates in the hindquarter *vs* the forequarter
- EUROP grading for conformation *vs* genetic potential of cuts : room for improvements
- Selection for carcass weight will increase high value cuts weight





What's next?

■ Use of digital images

- ◆ VBS2000 (E+V, Germany)
- ◆ 2 pictures / carcass
- ◆ Around 90 variables calculated to produce the EUROP grading for conformation & fat.





What's next?

■ Creation of prediction equations

- ◆ *Images variables* → *main cut yields*
- ◆ *Genetic analysis of the predicted yields*

■ First steps

- ◆ +5 million images stored since 2004
 - ◆ + 1 million animals match the ICBF database
 - + 600 000 animals with known sire.



Thank you.



16-18 June 2008

Interbull Meeting – Niagara Falls - USA