Recent Developments in Selection for Docility in Ireland

National Seminar on Strategies for improving Safety with Cattle

Ross Evans ICBF





Importance of docility

- In 2005 612 on farm accidents due to temperamental beef cattle (McNamara et al., 2007)
 - 27 deaths between 1996 and 2007 due to cattle livestock incidents (Irish Health and Safety Authority - code of practice, 2007)
 - Average of 2.25/ year. 85% of these were attacks by bulls, cows, and weanlings (HSA, 2007)

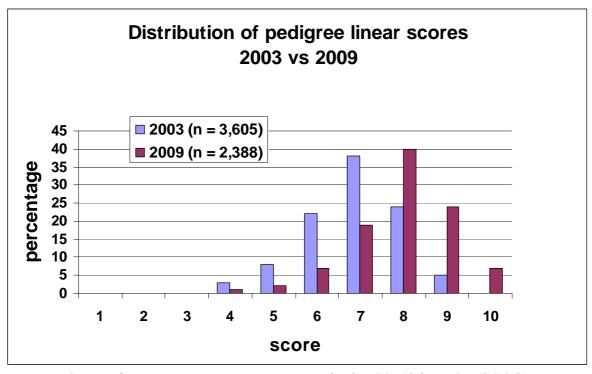


Evaluation pre -2009

- ICBF Trained Linear Scorers
- Limousine breed published figures based on pedigree animals
 - Breed average of 100, sd. of 10
- ~ 122,397 animals with docility
 - LM, CH, SI, HE, smaller numbers of other breeds including commercials
 - scale 1 to 10

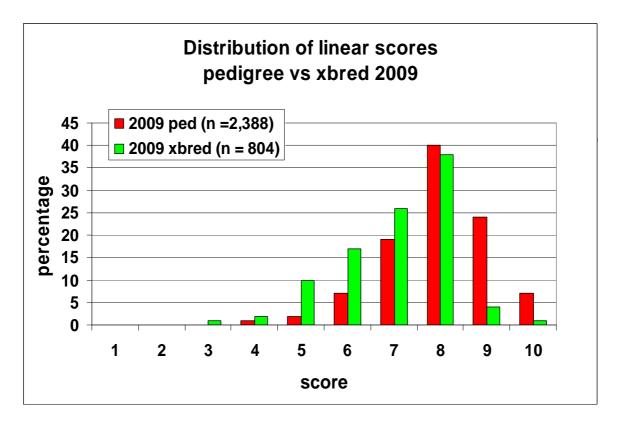
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~70% of scores recorded now 8, 9, 10 (28% in 2003)

But ~ less than 50% of population measured Genuine improvement or selective scoring



More variation when scoring crossbreds

(std = 1.24 for xbreds vs 1.13 for pedigrees)

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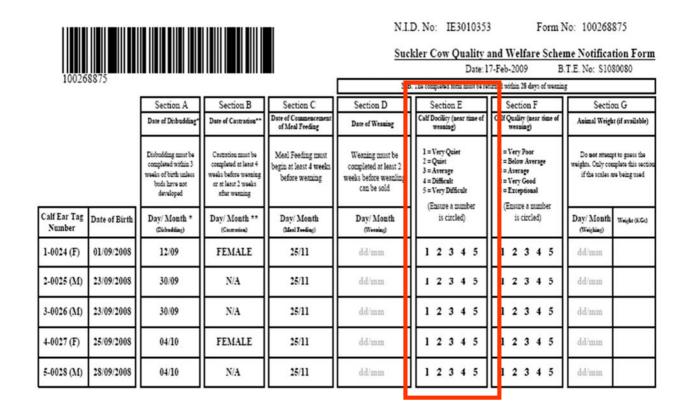
Animal welfare recording and breeding scheme



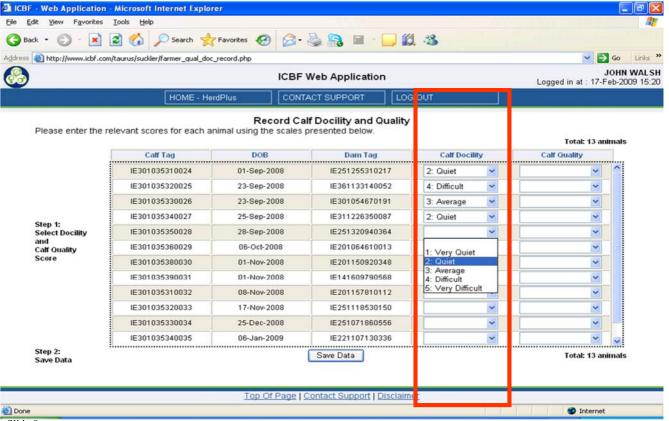
- Launched in January 2008 by Dept. of Agriculture
- ~ 60,000 herds, voluntary sign-up, monetary payment
- · Adhere to certain welfare measures, weaning measures
- Recording of sire, calving ease, disbudding, meal feeding, weaning, docility and calf quality

recordsource	% of total
ICBF_papersheets	83%
online_DEPT_website	12%
online_farmsoftware	2%
online_www.ICBF.com	4%

Paper based recording



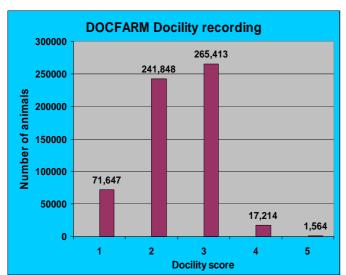
ICBF on-line recording



Two Forms of Docility

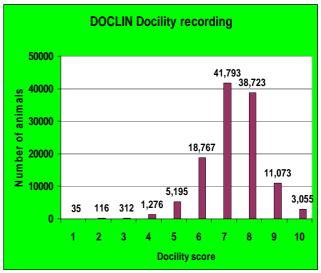
Farmer (DOCFARM)

n = 597,686



Technician (DOCLIN)

n = 122,397



Two Forms of Docility

Farmer (DOCFARM)

Breed	N-Obs	% com	% female
Angus	43,641	94	49
Aubrac	2,119	78	48
Blonde D'A	5,972	96	49
Belgian Blue	26,780	99	49
Charolais	246,811	97	49
Hereford	19,087	91	49
Limousine	190,973	97	49
Piemontese	920	89	48
Partenaise	997	91	49
Saler	6,332	92	50
Shorthorn	7,011	89	52
Simmental	39,096	95	49

Technician (DOCLIN)

Breed	N-Obs	% com	% female
Angus	2,570	34	51
Aubrac	1,031	12	64
Blonde D'A	1,033	46	51
Belgian Blue	2,874	92	50
Charolais	28,903	21	51
Hereford	3,040	7	49
Limousine	69,287	10	51
Piemontese	322	15	52
Partenaise	247	38	55
Saler	1,029	13	62
Shorthorn	356	49	71
Simmental	9,220	21	51

Genetic Parameter Estimation

Edits

- Only Animals recorded prior to sale DOCFARM
- Animals scored between 150-300 days for both traits
- Herds with variation in contemporary group (at least 3 scores)
- Variation in calf quality score and recorded same day
- Wean to docility recording <100 days
- CGs with at least 3 different sires (DOCFARM), 2+ different sires (DOCLIN)
- Only contemporary groups with >10 animals (DOCFARM), >5 animals (DOCLIN)
- Remaining *DOCFARM* = 73,827, *DOCLIN* =18,310

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Genetic Parameter Estimation

- 5 generation pedigree,

Animal model, no maternal effects

DOCFARM: CG of score date, sex, age, parity of dam, heterosis and recombination, meal feed

DOCLIN: CG of score date, sex, age, parity of dam, scorer, heterosis and recombination

Trait	N-Obs	N-Obs with both	-2LogL	h²	r _g	r _p
DOCFARM	73,827			0.44		
DOCLIN (150-300 day)	18,310	1,238	24449	0.25	-0.68	-0.19
DOCLIN (150-600 day)	72,228	2,304	72468	0.24	-0.40	-0.16

Genetic Parameter Estimation

Fixed		
Effect	DOCFARM	DOCLIN
Age	Older more docile	Older more docile
Sex	Females slightly less docile	Females slightly less docile
Heterosis	Crossbreds less docile	Crossbreds less docile
Meal Feeding	More docile with longer days on meal	
Parity	Animals from 1st parity cows less docile	No significant difference

Genetic Parameter Estimation

Maternal Effects: tested on smaller dataset

Model	-2LogL	h ² d	h ² _m	mat env (DOCLIN)	r _{gm}
Genetic effects only	32331	0.28			
Genetic & maternal envionment	32312	0.28		0.03	
Genetic & maternal genetic	32276	0.24	0.04		-0.47
Genetic & maternal genetic & maternal envionment	32255	0.28	0.03	0.03	-0.69

Results: similar to Beckman et al., (2007) on US Limousines

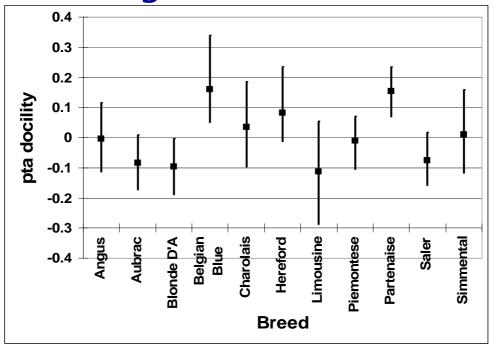
Follow up study conclusion: bias in herds due to selective recording, not all poor phenotypes recorded

Breeding Value Estimation

- Relaxed restriction on contemporary groups with at least 3 different sires represented
- % Crossbreds DOCFARM (96%) DOCLIN (39%)

Trait	N-obs	Animals with both	Average score	S.D.
AWRBS	302,065	3,816	2.4	0.76
linear scores	23,721	3,010	7.3	1.14

Average PTA Breed effects

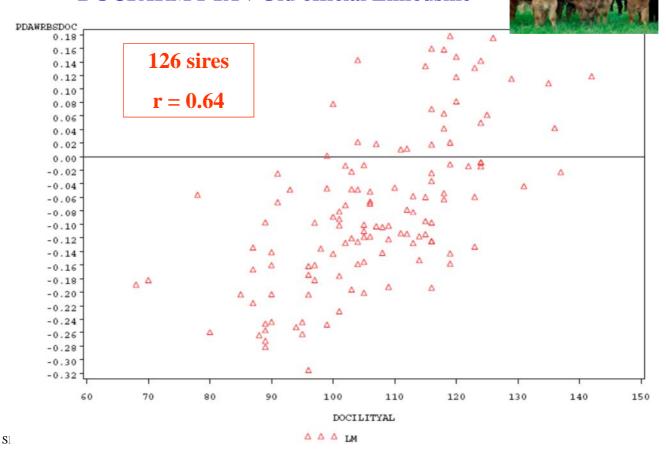


Belgian Blue highest average docility, Limousine lowest

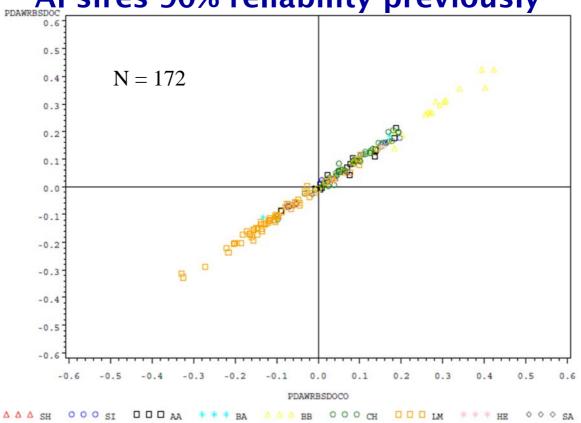
But 2 breed highest within breed deviation in PTA

Bottom 20% of sires across breeds: 8% increase in unfavourable scores

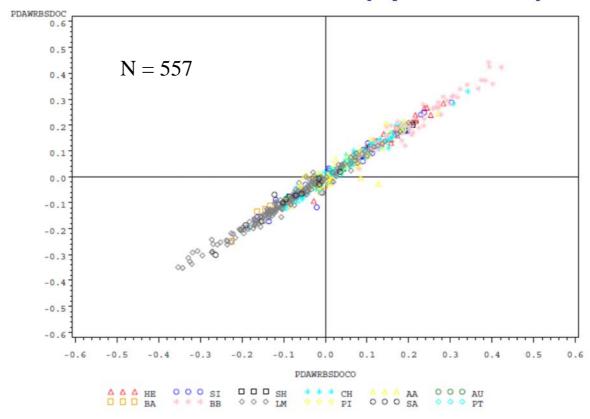
DOCFARM PTA v Old official Limousine







Al sires 70% reliability previously

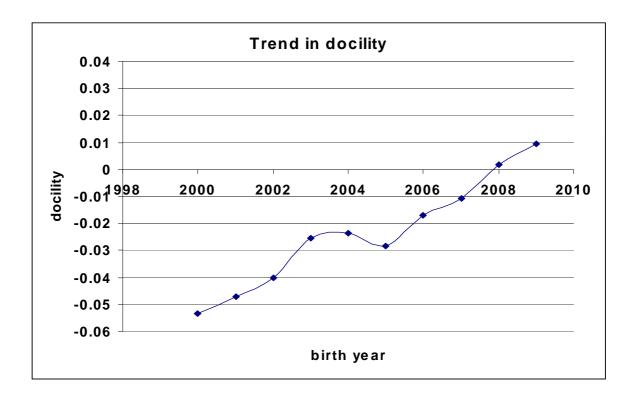


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Comparison of sires >90%

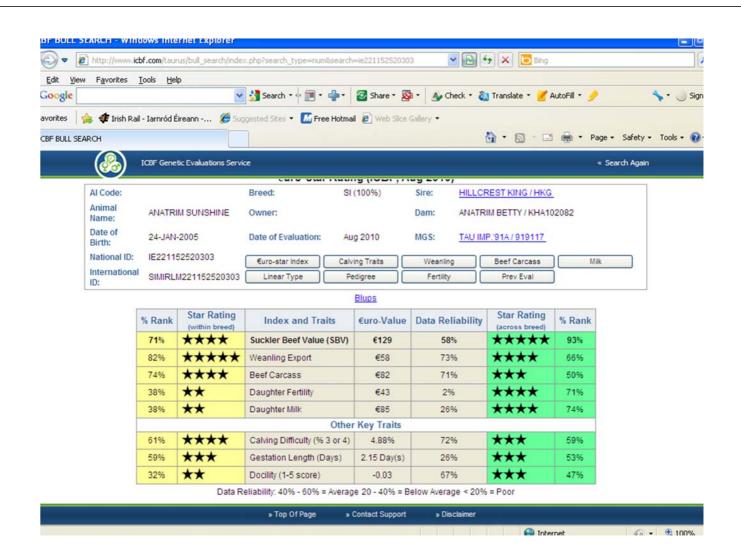
	Average no. progeny	Average no. of herd		Average progeny	Average herd mate	Average % of progeny
	with	mates	Average	DOCFARM	DOCFARM	scored 4 or
RATING	DOCFARM	with	reliability	score	score	5
across breed: *****	191	1,211	95	2.2	2.4	5
across breed: ***	160	1,101	96	2.4	2.4	9
across breed: *	182	1,128	95	2.6	2.4	13

Compared against their herdmates
Clear differences appearing



Slight trend towards improved docility

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Further work

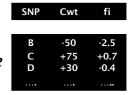
- Investigate maternal effects further with repeated observations on cows
- · Economic value of docility?
- · €17.87 per unit increase in weanlings
- · €33.20 per unit increase in weanlings
- · International collaboration
- Mature stock bull docility same trait?
 - Need system to collect and measure, then analyse associations
 - Use as correlated trait
 - Would include dairy herd docility
- · Genomic selection

Using the DNA to select animals called "Genomic selection"

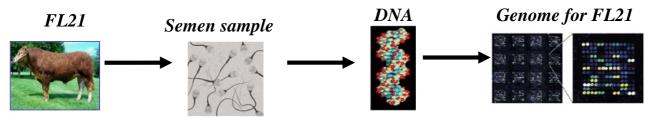
Collect DNA and get a genome for all well proven AI bulls

Why AI bulls?

So we can compare their genotype versus their EBI through their well proven daughter performance

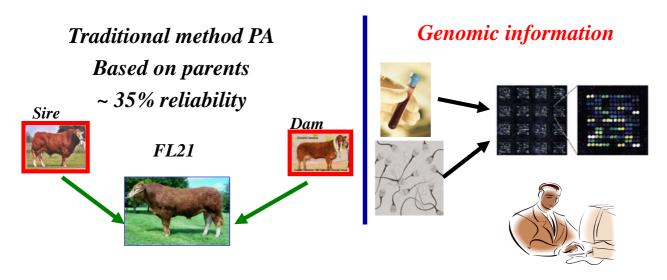


Best source to get DNA is a blood sample Most well proven AI sires are dead so Semen, hair



About 1,000 proven bulls with genomic information at ICBF

Genomic Selection in beef cattle



<u>Proof potentially based on average of sire and dam but also</u> genomic information ~ 50% reliability











~ equivalent of 15-20 progeny

records



Current status

- Across breed DOCFARM proofs now official for Active AI sires, natural service sires and females
- Across breed nature evaluation unique in the world
- 200,888 Farmer scores, 99,204 linear and increasing rapidly
- Will help to reduce cattle related injuries on farm if recorded properly