

IRISH CATTLE BREEDING FEDERATION

ICBF Dairy & Beef Genetic Evaluations Meetings.

Wednesday 21st July 2010. Maldron Hotel, Portlaoise.



Agenda 1. Dairy Traits & Dairy Breeding Programs.

- Agenda 1 (10 AM 11.30).
- 9.30 Tea & Coffee.
- 10.00 Index developments, including EBI, new labour sub-index, culling index and other traits - Andrew
- 10.20 Test-day evaluations; milk production traits
 John McCarthy.
- 10.40 Dairy genomics research Donagh.
- 11.00 Genomics operational Francis.



Agenda 2. Dairy & Beef Traits

- Agenda 2 (11.45 3.00)
- 11.45. Female fertility traits Ross.
- 12.30. Male fertility traits Ross.
- 1.00. Lunch.
- 1.30. Calving traits Francis.
- 2.15 Carcass cut data & other beef traits Thierry.
- 2.30 Al application process Pat.
- 2:45 Catalogues Brian W



Agenda 3. Beef Traits & Beef Breeding Programs.

- Agenda 3 (3.00 5.30).
- 3.00 Lessons from Grange research & BETTER farms programs – Pearce Kelly
- · 3.20 Linking Tully & beef industry data John Crowley.
- · 3.40 Beef linear traits Contemporary groups Ross.
- 4.10 Maternal weaning weight Ross.
- 4.30 Developments in €uro-Star evaluations Tim Byrne.
- 4.50 Beef genomics research Donagh.
- 5:00 Beef Specialist Brian W
- 5.15 Close of meeting.



Work-plan & recommendations.

- · Interbull official evaluations.
 - 23rd August 2010.
 - 13th December 2010.
- · Interbull test evaluations.
 - 7th September 2010.
- Trait improvements, e.g., fertility, calving, go through test run (7th September). If pass, then part of official evaluation (13th December).





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Index developments, including EBI, new labour sub-index, culling index and other traits

Andrew Cromie



EBI Developments 2009/2010 - Review.

- Beef Sub-Index.
 - Updating income/costs for rearing replacements and surplus stock.
 - Separation of live-weight into; (i) surplus beef value, and (ii) cow maintenance.
 - Lighter animals/breeds benefited (+/- €20 in EBI terms.
- Economic Values.
 - Update all incomes & costs.

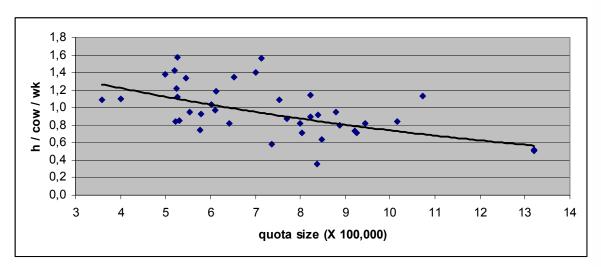


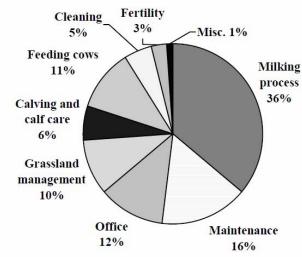
EBI Developments 2010/2011 - Work-plan.

- · Economic Values.
 - Update all incomes & costs.
 - Increased focus on economics of health traits.
- · New labour sub-index.
 - Currently EBI covers direct cost of health and fertility problems, e.g., cost of feed, cost of treatment, cost of withholding milk etc.
 - What about cost of labour? Time taken for; (i) milking duration, (ii) multiple serves, (iii) treatments, (iv) calving difficulty, (v) temperament etc.



New labour sub-index (i)





- Work from Netherlands (Zijlstra et. al., 2006) & Ireland (O'Brien, 2006) from dairy farms.
 - Larger dairy herds = less time for individual cows.
 - Large difference in time for individual cows (average 60 minutes per cow with SD of 30 minutes).
- With increasing scale and less labour, increasing demand for "easy-care" profitable cows.

New labour sub-index (ii)

- Project initiated with Animal Sciences Group, Wageningen (Roel Veerkamp).
- Two key questions regarding work-plan?
 - Which "labour" traits are most important?
 - · Milking speed, temperament, number serves, calf health, mastitis incidence, lameness incidence...
 - How important are these labour traits relative to other profit traits in the EBI?
- Undertake survey of farmers (August/Sept)
 - Web-based survey & a meeting with "targeted farmers.
- Establish genetic differences between bulls in "labour required" for daughters.
- Present update at next industry meeting.



New traits – Mastitis & lameness (i).

- · Health sub-index.
 - We use SCC with an assumed correlation of 0.70 to indicate mastitis.
 - We use locomotion with an assumed correlation of -0.40 to indicate lameness.
- Now starting to receive "incidence" data from commercial farmers.
 - ~100k incidence records from; (i) G€N€
 IR€LAND herds and (ii) Discussion group herds
 - dairy efficiency program.

New traits - Mastitis & lameness (ii).

- Work from Donagh (Nov 09) indicated;
 - Incidence of 10% for mastitis, 5% heritability and rg of ~0.5 with SCC
 - Incidence of 10% for lameness, 4% heritability and rg of -0.35 with locomotion score.
 - Results compare well with international estimates.
- · Develop new health evaluation system.
 - Mastitis + lameness + predictor traits.
- Update at next industry meeting.



New Culling Index

- New PhD project involving ICBF, Teagasc and Abacus Bio. Walsh fellowship secured.
- Looking at 2 stage implementation.
 - Stage 1. New fertility evaluations + output from test day evaluations + economic value work from Abacus Bio.
 - Stage 2. Refinements based on work of PhD student.
- Update at next industry meeting.





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Test Day Models for Milk Production Traits John McCarthy.



Background

- Uses 305 day values
- Operated on contract by CRV Holland for ICBF
- Genomic data is incorporated separately to main evaluation in postblending process
- Uses variation of tools



Background(ctd)

- 305 day model uses 1 305 day figure for milk/fat/prot/scc which summarises whole lactation
- The 305d figures are calculated using "lactation curves" software
- Change from 305 day model to test day model where all individual recordings are included in evaluation
- Use this opportunity to examine "Irishspecific" issues to include in new test-day model



Why

- More accurate estimation of environmental effects from including the influence of particular days of recording
- Optimal use of information from all test days
- Better use of records in progress
- Remove necessity of predicting 305d
- Method of choice for many dairy evaluations internationally (NZ, HOL, CAN, ...)



What

- Change from 305 day model to test day model
- Examine whether variance component structure needs to incorporate Season Calving
- Estimate Initial Genetic Parameters
- Examine Breed Effects
- Examine effect of recording method (eDIY, Technician)



What(ctd)

- Examine Heterogeneity of Variance –
 i.e. some herds have more variance
 than others
- Incorporate into overall model
- International experience has shown that there will be re-ranking, particularly on cows and young bulls



Raw Milk Production Data

6,064,769	Num lactations in Production file
2,297,342	Num animals in Production file
35,572,153	Total number of animal tests
25,030	Number of Herds

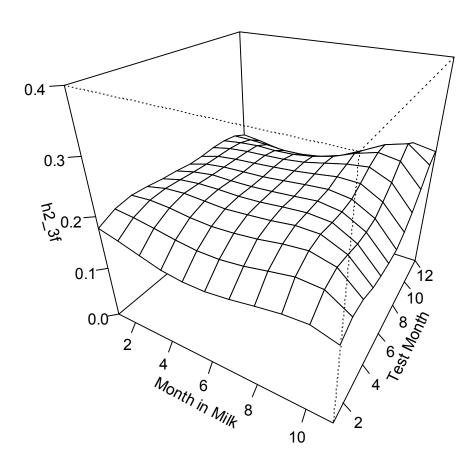
Current Status

- Analysis of data to determine if variance component structure needs to incorporate Seasonal Calving
 - Nearing Completion it appears there is a need for this step
- Estimate Initial Parameters
 - Follows from above



Current Status(ctd)

Heritability of milk yield





Current Status(ctd)

- Breed Analysis
 - Initial investigation of existing work by John Hickey
- Examine effect of milk recording method
 - To do



Breed Composition of Dairy Replacements born in 2009

Sire Breed

		FR	JE	MO	MY	NR	RB	SR	Total
	FR	250,717	5,807	1,930	241	2,052	1,169	210	262,126
Da	JE	2,497	1,131	15		117	22	2	3,784
3	MO	3,802	162	2,570	23	132	95	15	6,799
W	MY	490	11	38	388	7	54		988
re	NR	443	144	3		155	6	10	761
ed	RB	1,682	81	74	33	78	772	14	2,734
	SR	319	80	3		20	5	41	468
	Total	259,950	7,416	4,633	685	2,561	2,123	292	277,660

~10% non Black and White



Breed Composition of cows milk recorded in 2009

Sire Breed

		FR	JE	МО	MY	NR	OTH	RB	SR	Total
	FR	420,170	3,924	5,536	592	823	4,683	2,304	674	438,706
Da	JE	1,483	1,518	21	1	16	43	8	20	3,110
3	MO	2,678	99	3,024	17	30	154	105	34	6,141
₩	MY	441	4	30	465		48	34	5	1,027
Te e	NR	214	11	5		529	2	12	24	797
ed	OTH	2,116	100	176	20	18	1,376	55	15	3,876
	RB	1,110	65	36	6	3	36	1,121	22	2,399
	SR	151	49	9		12	10	4	89	324
	Total	428,363	5,770	8,837	1,101	1,431	6,352	3,643	883	456,380

~10% non Black and White



Next Steps

- Complete investigation of variance component structure Seasonal Calving
- Begin to examine in detail breed effects and what constitutes different genetic groups



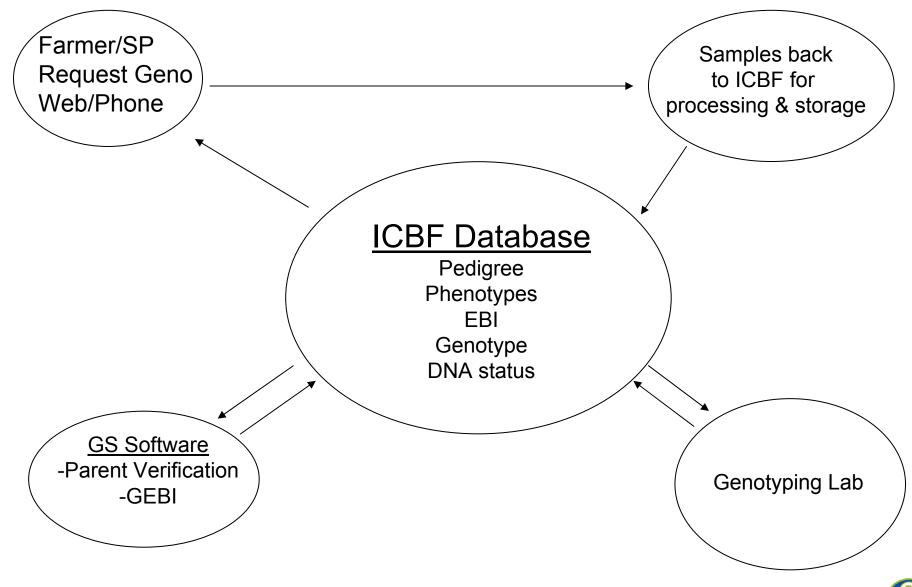


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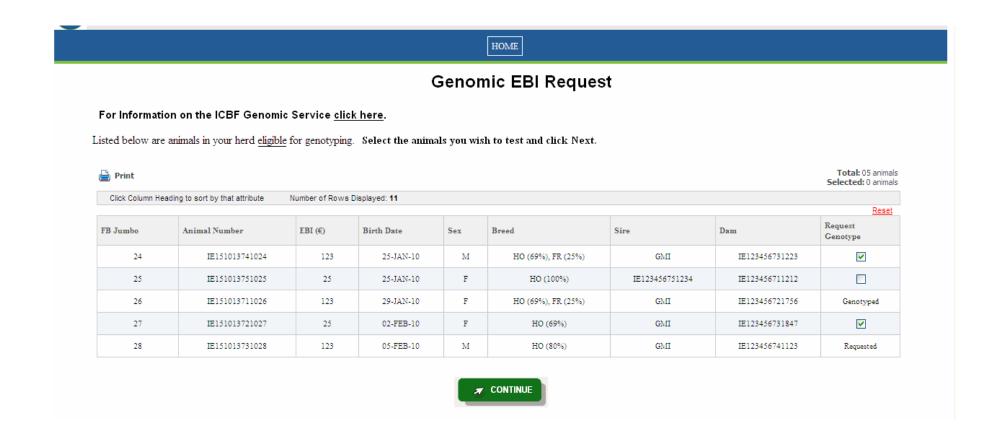
Genomic Service

Francis Kearney





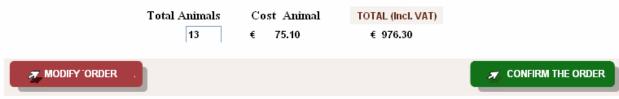
Order Process





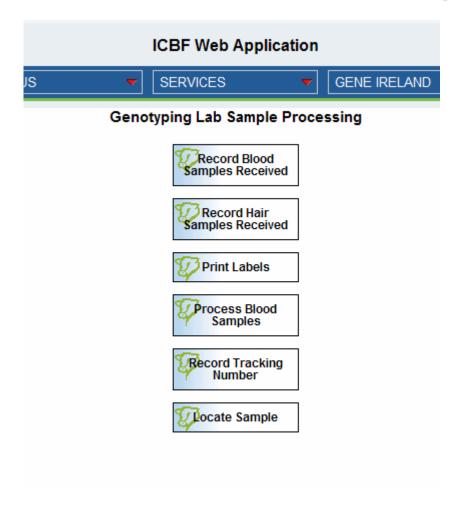
Order Process

Jumbo	Animal Number	DOB	Sex	Breed	Name	Sire	MGS	EBI€			
855	IE151013740855	20-MAR-05	F	HO (94%) FR (3%)	PARKDUV IRIS 25	HRZ	ELC	118	×		
1135	IE151013741135	15-APR-08	F	HO (91%) FR (9%)	PARKDUV SNOWDROP 19	HFL	ESZ	95	×		
1136	IE151013751136	17-APR-08	М	HO (97%) FR (3%)		SAO	RMB	45	× .		
1137	IE151013761137	20-APR-08	F	HO (97%) FR (3%)	PARKDUV NANNYBREEZE 41	KES	ESZ	80	x		
1138	IE151013771138	18-SEP-08	М	HO (100%)		RUU	GMI	112	×		
1139	IE151013781139	22-SEP-08	F	HO (91%) FR (9%)	PARKDUV NANNYBREEZE 43	KWA	AHD	119	×		
1140	IE151013711140	24-SEP-08	F	HO (97%) FR (3%)		RUU	FAL	92	×		
1141	IE151013721141	24-SEP-08	F	HO (97%) FR (3%)	PARKDUV ROSE 13	KWA	RMB	117	×		
1142	IE151013731142	02-OCT-08	F	HO (100%)	PARKDUV DAISY 15	KWA	QUR	95	×		
1143	IE151013741143	06-OCT-08	М	HO (97%) FR (3%)		OJI	RUU	192	×		
1144	IE151013751144	06-OCT-08	F	HO (97%) FR (3%)	PARKDUV ROSE 12	OJI	RUU	198	×		
1145	IE151013761145	07-OCT-08	М	HO (94%) FR (3%)		OJI	RUU	205	× :		
1146	IE151013771146	08-OCT-08	F	HO (97%) FR (3%)	PARKDUV MARIA 33	RUU	GMI	109	× :		
	Total Animals Cost Animal TOTAL (Incl. VAT)										



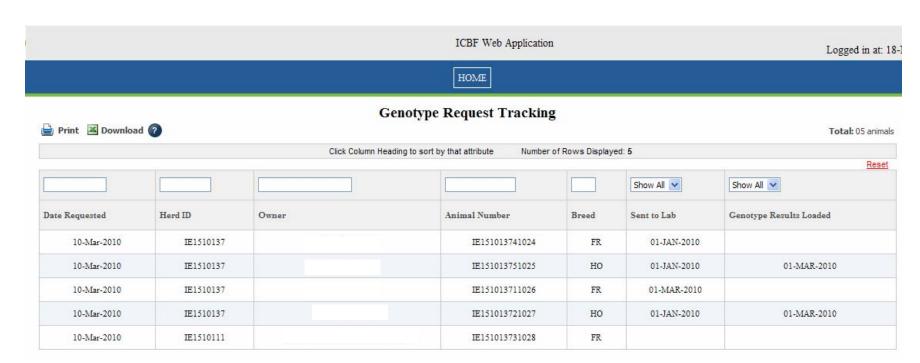


ICBF Processing





Tracking





Results - Profile

											Milk							
Jumbo Animal Number	DOB	Sex	Lact Sire	EBI€	Milk€	Fert€	Calv€	Beef€	Maint€	Health€	Kg	FatKg	ProtKg	Fat%	Prot%	Calv Int	Surv%	Genomic
200 IESNYH0200Q	21-OCT-97	F	9 ASI	109	40.8	68	13	-25.8	10.4	2.8	95	5.5	7	0.04	0.07	-4.4	1.3	Υ
441 IE151013730441	29-MAR-00	F	7 AHD	46	70.1	-17.3	3.9	-5.7	-4	-1.2	387	14	14.5	-0.01	0.03	2.2	0.7	N
467 IE151013740467	09-OCT-00	F	7 MFX	110	69.8	31.8	13.4	-13.6	5.5	3.5	63	6.5	11	0.08	0.17	-1.5	1.2	N
487 IE151013780487	19-NOV-00	F	6 QUR	134	95.8	8.4	32.3	-7.4	7.6	-2.5	142	14.5	15	0.17	0.19	-0.8	-0.1	N
504 IE151013780504	10-FEB-01	F	8 AHD	80	6	65.6	8.5	7.2	-3.6	-4.1	-113	-1	-0.5	0.07	0.07	-3.8	1.7	N
550 IE151604920550	25-MAR-06	F	3 RUU	131	48.6	77.6	14.9	1.1	-11.8	0.6	23	13	6	0.23	0.1	-4.7	1.8	N
562 IE151013790562	17-NOV-01	F	5 MFX	24	43.7	-20.5	4.8	-2.6	-1.4	-0.4	28	8.5	6	0.14	0.1	2.3	0.5	N
566 IE151013740566	09-DEC-01	F	5 MFX	88	76.9	0	15.8	-10.5	5.5	0.8	144	11.5	12.5	0.11	0.14	0.5	0.5	N
568 IE151013760568	26-JAN-02	F	6 FAL	73	63	0.2	8.6	-0.8	5.7	-3.9	226	5	12.5	-0.07	0.09	0.4	0.4	N
569 IE151013770569	26-JAN-02	F	6 FAL	48	29.5	10.8	12.5	-3.2	3.8	-5.1	95	3.5	5.5	0	0.05	-0.6	0.3	N
590 IE151013740590	03-MAR-02	F	6 ESZ	76	56.8	17.5	15.1	-10.2	-4.7	1.7	271	6	12	-0.08	0.06	-0.5	1	N
605 IE151013710605	06-SEP-02	F	6 MFX	42	67.3	-21.4	3.1	-8.6	-0.5	2.4	403	9.5	15	-0.1	0.03	2.4	0.5	N
633 IE151013750633	29-OCT-02	F	4 HRZ	65	68.1	-10.7	8.9	-9.5	3.8	4.5	259	13	12.5	0.06	0.07	2.4	1.5	N
651 IE151013770651	15-JAN-03	F	5 ESZ	90	58.6	34.9	8.6	-11.5	0.2	-0.4	215	9	11	0.02	0.07	-1.9	1	N
673 IE151013740673	12-FEB-03	F	4 GMI	93	54.1	48.2	10.5	-33.6	15.8	-1.5	137	13	8.5	0.15	0.08	-1.7	2.4	N
675 IE151013760675	16-FEB-03	F	5 GMI	91	65.6	7.8	28.2	-10.3	1.9	-2.1	212	9.5	12	0.03	0.09	0.5	1.1	N
690 IE151013750690	02-APR-03	F	6 LLO	89	56.6	33.5	11.6	-10.1	-2.5	0	238	6	11.5	-0.06	0.07	-1.3	1.5	N
699 IE341460380699	28-JAN-06	F	3 RUU	130	21.9	82.6	28.6	6.5	-12.6	2.8	20	8	2.5	0.14	0.04	-4.3	2.6	N
702 IE151013780702	07-OCT-03	F	4 RGK	-7	49.4	-36.7	-3.2	-13.4	2.9	-6.2	45	9.5	7	0.15	0.11	1.6	-1.5	N
706 IE151013730706	09-OCT-03	F	4 ESZ	44	59.1	-8.8	7.9	-17.4	3.4	0	222	7	11.5	-0.03	0.08	0.7	0	N
721 IE151013720721	16-NOV-03	F	5 ESZ	81	45.8	44.2	1.8	-7.3	-0.8	-2.3	111	2.5	8.5	-0.03	0.09	-2.7	1	N
728 IE151013790728	24-DEC-03	F	5 AHD	79	67.8	9.1	5.7	4.6	-4.9	-3.4	385	8.5	15	-0.1	0.04	-0.3	0.5	N
752 IE151013790752	02-FEB-04	F	4 AAP	35	-4	40.6	16.3	-11.2	-0.5	-6.7	1	-7	0.5	-0.14	0.01	-2.4	1	N
761 IE151013710761	13-FEB-04	F	5 AHD	112	65.5	36.1	14.5	0.5	-2.6	-2.1	143	12.5	10.5	0.13	0.11	-2.4	0.6	N
764 IE151013740764	18-FEB-04	F	4 ESZ	74	49.3	20.4	7.7	-0.1	-0.9	-2.6	231	10.5	9.5	0.03	0.04	-0.7	1	N
769 IE151013790769	25-FEB-04	F	5 AAP	123	42	66.7	29.2	-20.1	8.5	-3	198	6.5	8.5	-0.02	0.04	-4	1.6	N
772 IE151013740772	28-FEB-04	F	4 GMI	69	47.1	11.3	15.4	-6.7	4.4	-2.3	223	4.5	10	-0.07	0.05	0.3	1.3	N
781 IE151013750781	05-APR-04	F	3 DCD	68	52.6	9.1	19.6	-1.5	-8.6	-3.4	205	11.5	9.5	0.07	0.05	-0.9	-0.2	N
784 IE151013780784	22-SEP-04	F	4 RMB	77	72.8	9.4	9.6	-6	-8.3	0	196	9	13	0.03	0.12	-0.9	-0.1	N
785 IE151013790785	25-SEP-04	F	3 RMB	42	43.2	6	5.3	3	-9.5	-5.7	202	8	8.5	0.01	0.04	0.3	0.8	N
789 IE151013740789	01-OCT-04	F	4 RMB	44	55.5	-0.5	-2.3	-2.7	-6.7	1.1	-33	5.5	7.5	0.13	0.17	0.7	0.6	N
791 IE151013770791	03-OCT-04	F	4 QUR	38	40.3	-7.4	14.3	2.4	-8	-3.4	304	5	10	-0.11	0	0.4	-0.2	N



Results - Individual

	Ge	nomic Evalu	ation Re	port		
Jumbo	802			Lact. No	0	
Tag	IE12345678980	2		Sex	F	
Name	ICBF			Sire	GMI (€100)	
Date of Birth)y 1m		Dam	IE123456788111	(€95)
Breed	HO 100%			Dam's Sire	OJI (€246)	
Date of Evaluation	28-Jun-10					
				Change in		
	Official		% Weight	Evaluation	Reliability	Direct
	Genomic		on	from Parent	Increase with	Genomic
Index	Evaluation	Reliability %	Genomics	Average	Genomics %	Value
gEBI €	Evaluation	remaining 70	Genomics	Trierage	Genomics 70	, arac
Milk Sub Index €						
Fertility Sub Index €						
Calving Sub Index €						
Beef Sub Index €						
Maintenance Sub Index €						
Health Sub Index €						
Milk Sub Index						
Milk (Kg)						
Fat (Kg)						
Prot (Kg)						
Fat (%)						
Protein(%)						
Fertility Sub Index						
Calv Int (Days)						
Survival (%)						
Calving Sub Index						
Dir. Calv Diff (%)						
Mat. Calv Diff (%)						
Gest Len (Days)						
Calf Mort (%)						
Beef Sub Index						
Cull Cow Weight (Kg)						
Carcass Weight (Kg)						
Carc Conf (Grade)						
Carcass Fat (%) Maintenance Sub Index						
Cull Cow Weight (kg)						
Health Sub Index						
Lameness (Locomotion)						
Udder(SCC)						
)09	!	2	1	!		



Current Status

- Work almost complete on Lab screens
- Development of other screens ongoing
- Implement imputation
- Turnaround time ~4/5 days



Issues

- Results sent to person who pays for genotype in advance
- Results made official at next evaluation (weekly, bi-weekly, monthly, 3 times/year)?
- How to handle duplication?
 - Not an issue where contract exists
 - Dropped calves request from multiple sources



Recommendation

- Proceed with development work
- Update at next Industry meeting



Imputing genotypes from lowcost, less dense genotype arrays

Donagh Berry Teagasc, Moorepark

donagh.berry@teagasc.ie



Motivation

- Currently costs ~€150 per genotype with 50,000 SNPs
- · Genomics will dictate selection decisions
 - Heifers as replacements
 - Bulls as test bulls or stock bulls
- Parentage testing is a form of genotyping but in current form is not useful for genomic selection
- > reduce the cost of genomic selection



State of the art in genotyping platforms

Currently 54,001 SNPs

1,661 discarded 2,269 new

New version 54,609 SNPs

50k chip



2,900 SNPs *3k chip* ~790,000 SNPs HD chip High Density chip

```
Sire
```

....TCACCGCTGAG.....

.....CAGATAGGATT.....

....??<mark>G</mark>?????A??....



```
Sire
......CAGATAGGATT......
.....CAGATAGGATT......
.....??T??????T??.....
Offspring
```



```
Sire

.....TCACCGCTGAG.......AGTACATCTAG.....

.....CAGATAGGATT......CAGATGGATTG.....

.....??G??????A??....

.....??T??????T??....

Offspring
```

```
Sire

.....TCACCGCTGAG.......AGTACATCTAG.....

.....CAGATAGGATT......CAGATGGATTG.....

.....CAGATAGGATT.....

.....AGTACATCTAG.....

Offspring
```

Sire

....TCACCGCTGAG.....

.....CAGATAGGATT.....

Dam

....AGTACATCTAG.....

.....CAGATGGATTG.....

....CAGATAGGATT.....
....AGTACATCTAG.....



Population

MG-Sire

....AGTACATCTAG.....AGTACATCTAG.....

.....CAGATGGATTG.......CAGATGGATTG.....

....AGTCGTGACTG.....

Sire

....TCACCGCTGAG.....

.....CAGATAGGATT.....

Dam

....???????????

????????????



....??T?????T??....



Population

MG-Sire

....AGTACATCTAG.....AGTACATCTAG....

.....CAGATGGATTG......CAGATGGATTG.....

....AGTCGTGACTG.....

Sire

....TCACCGCTGAG.....

.....CAGATAGGATT.....

Dam

.....????????????.....

....???????????



....AGTACATCTAG.....

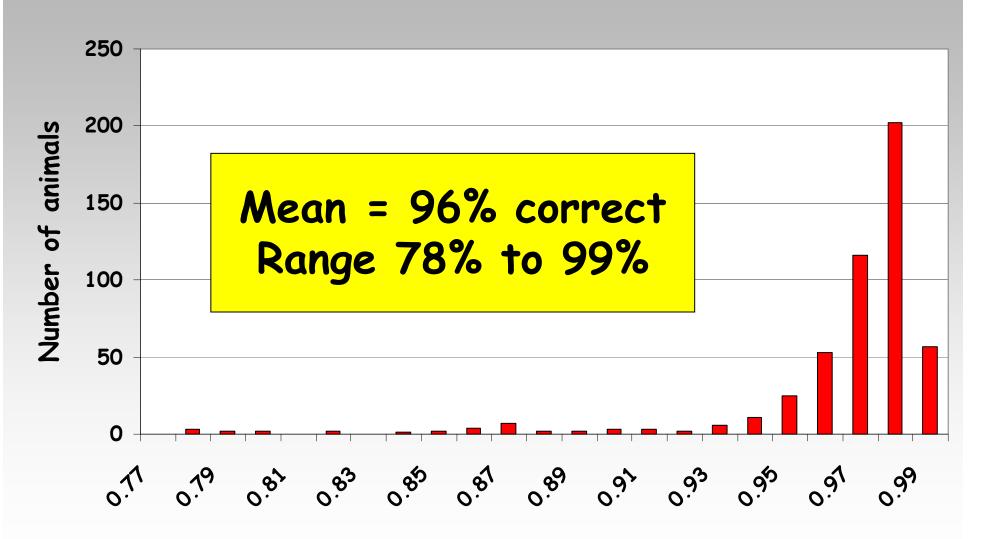


Testing

- 5,732 animals with genotypes on 54,000 SNPs
- 505 animals born since 2007 assumed to be genotyped only on smaller 3,000 SNP chip
 - Had their 54,000 genotypes so knew the "answer"
- · Accuracy
 - Percentage correct alleles (A,B)



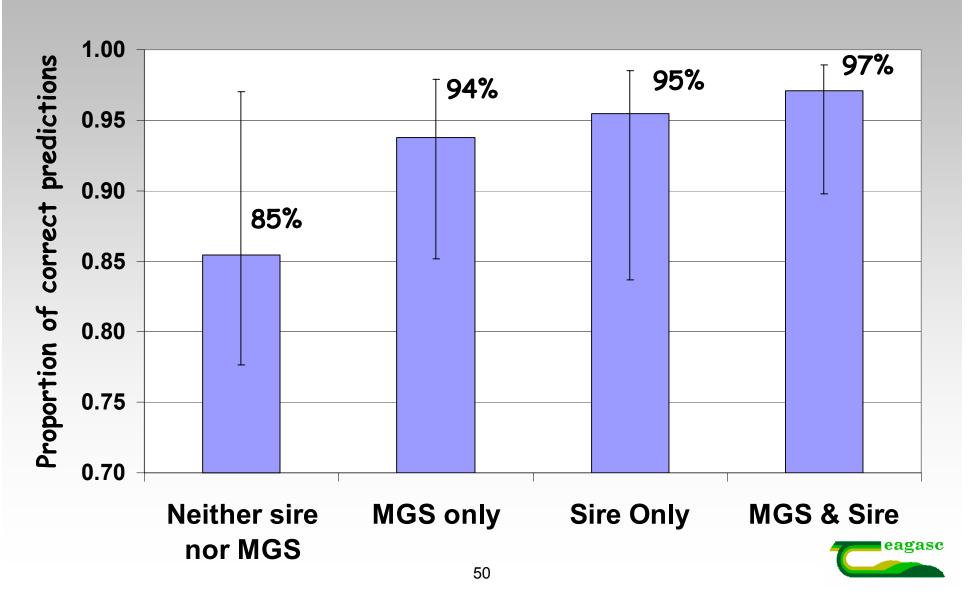
Accuracy per animal



Proportion of correct alleles



Accuracy by pedigree genotyped



Implementation

- 3k chip can also be used for parentage testing
- Computing time: ~24 hrs for imputation



Genomics and other breeds

- Key for successful genomic prediction is representation of the DNA signature in the training population
 - Breed (HO, FR, JER, MO, NO...) must be genotyped
 - Animals must have a traditional proof
- · Also vital for imputation



Conclusions & recommendations

- Accuracy of imputation is high particularly with back pedigree is genotyped
- Only impute where sire and MGS genotypes are available
- Bulls used in AI must have 54,000 SNPs genotyped
- · Continue research in imputation



Other genomic research

- Breeding programs (Noirin)
- Multi-breed genomic evaluation (Holsteins versus Friesians)
- Cows/stock bulls in training population
- Sharing of genotypes

