

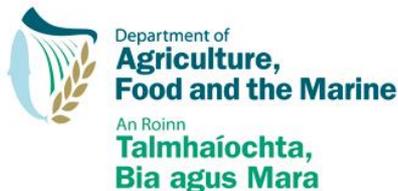


IRISH CATTLE BREEDING FEDERATION

The Irish Beef Genomics Scheme; Applying the latest DNA technology to address global challenges around GHG emissions and food security.



*Dr Andrew Cromie, Technical Director ICBF.*



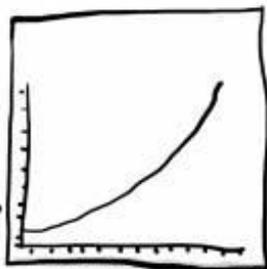
# The Global Challenge; GHG Emissions & Food Security.

## THE JEOPARDY OF PROGRESS

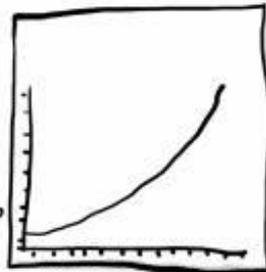
This graph shows the increase in crop yields due to improved agriculture.



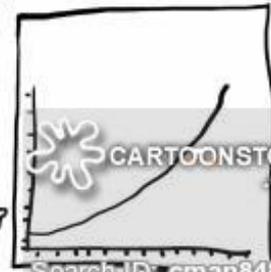
This graph shows the increase in the human population due to the increased crop yields.



This graph shows the increase in environmental degradation due to the increased population.



This graph shows the increase in despair as we realise that we can use the same graph to measure them all.



# The World needs to “wake-up”

The  
Economist

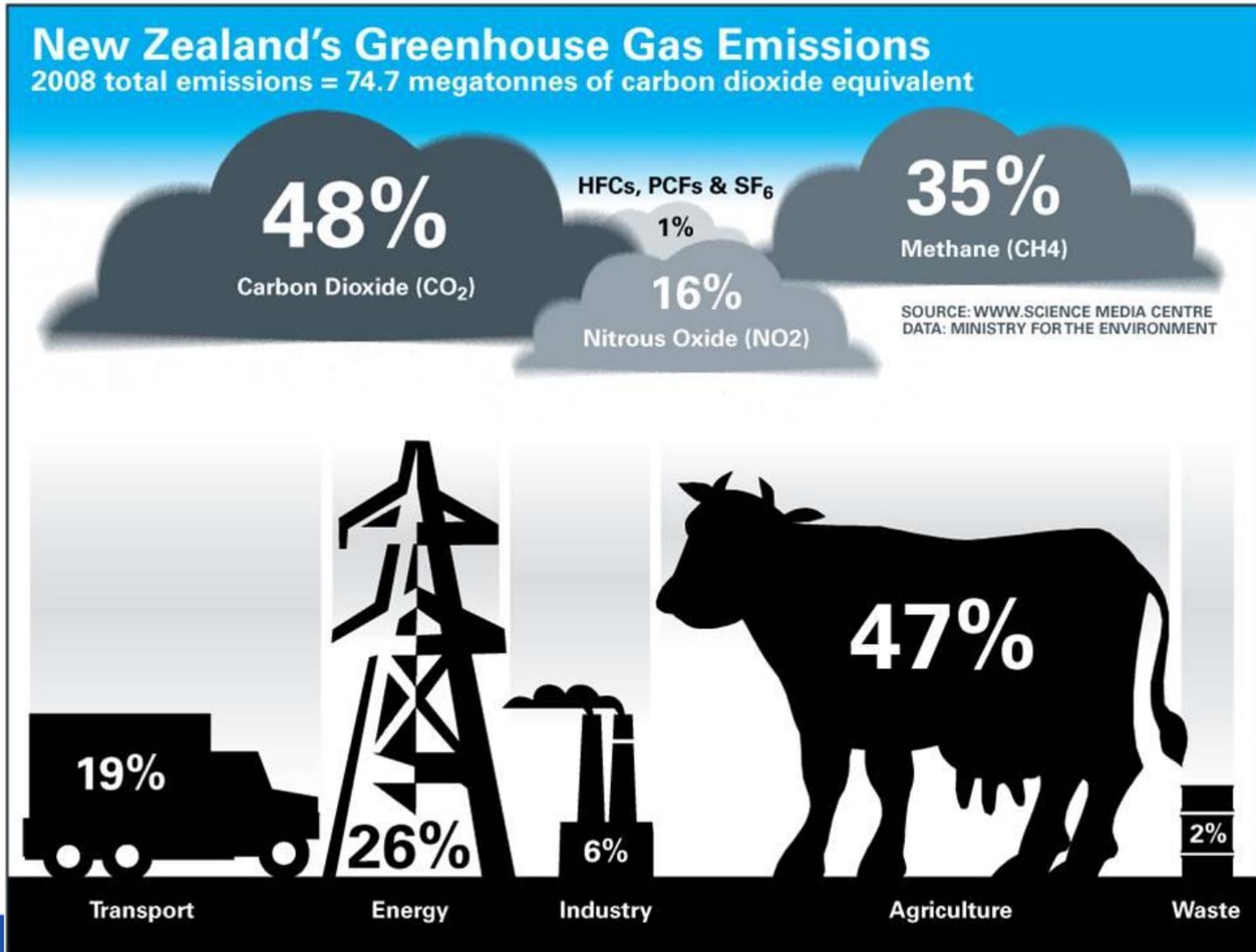
Events

## FEEDING THE WORLD



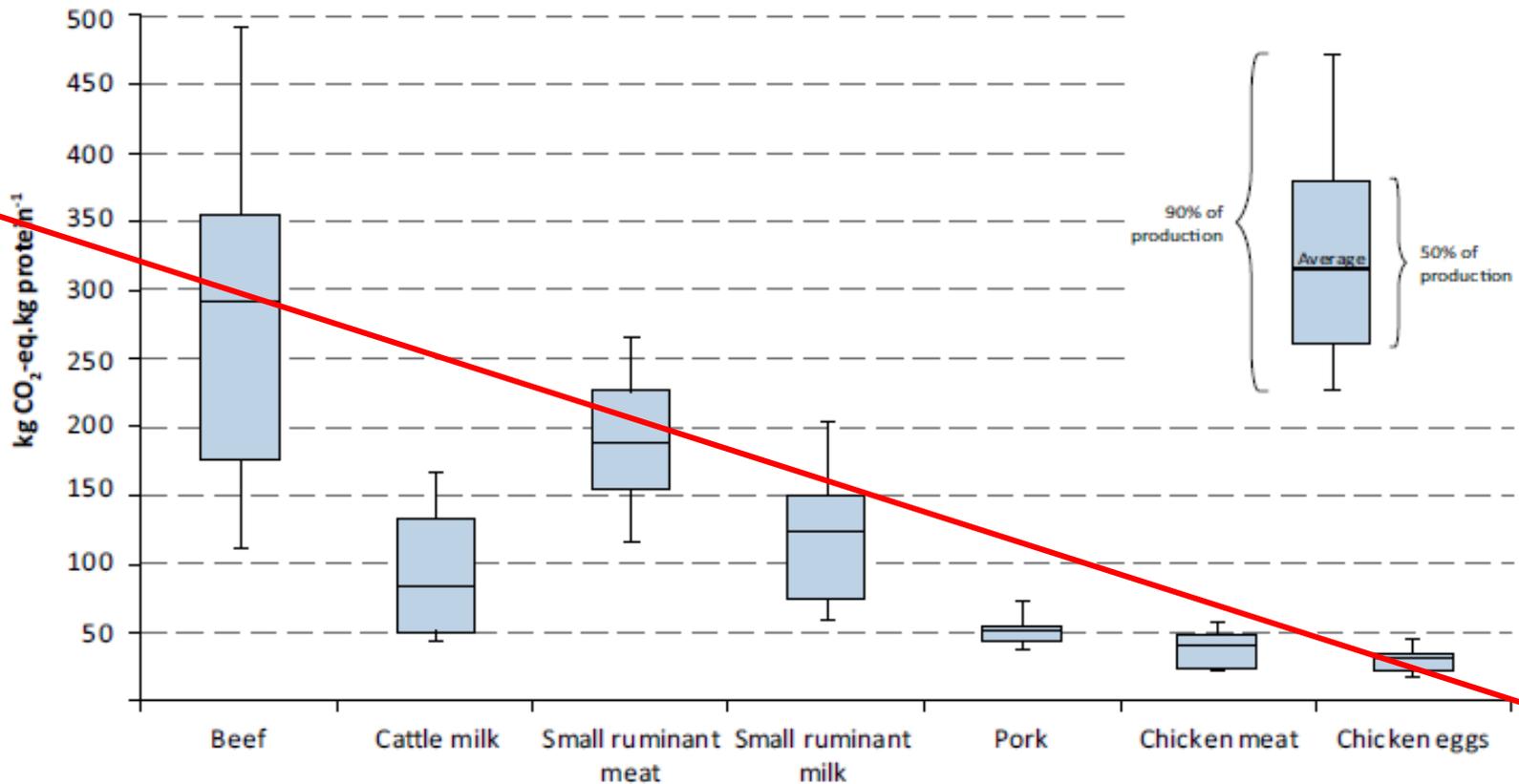
SUSTAINABLE SOLUTIONS FOR A GLOBAL CRISIS  
The world needs to wake up to the impending 2050 food security crisis

# Agriculture is a Problem!



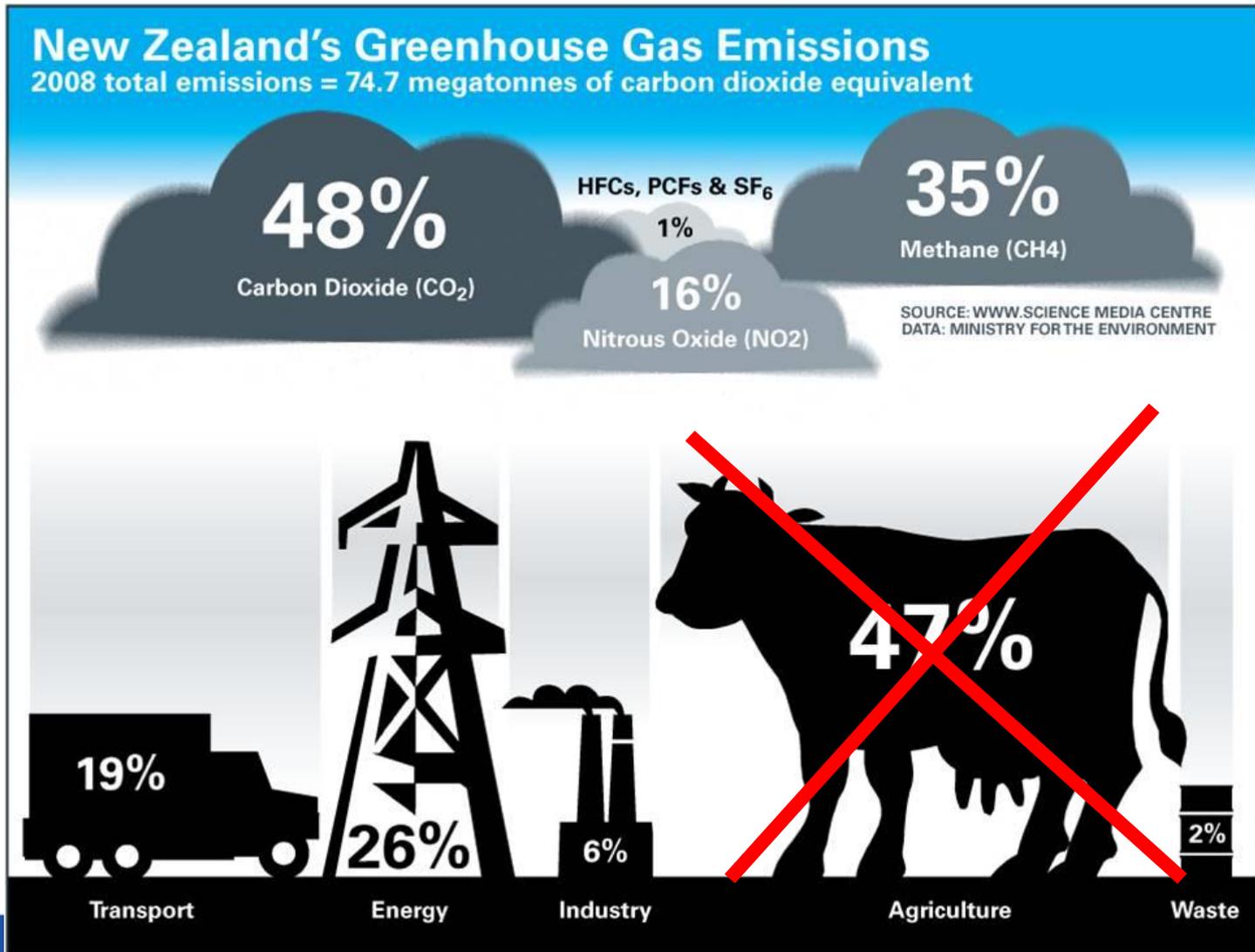
# Beef Cows are a particular Problem!!

FIGURE 3. Global emission intensities by commodity



Source: GLEAM. FAO, 2013

# Solution – Get Rid of Cows



# We all need beef cows.



- Suckler cows & beef cattle are a key part of rural infrastructure, especially in Ireland
  - Small fragmented farms, marginal land etc.

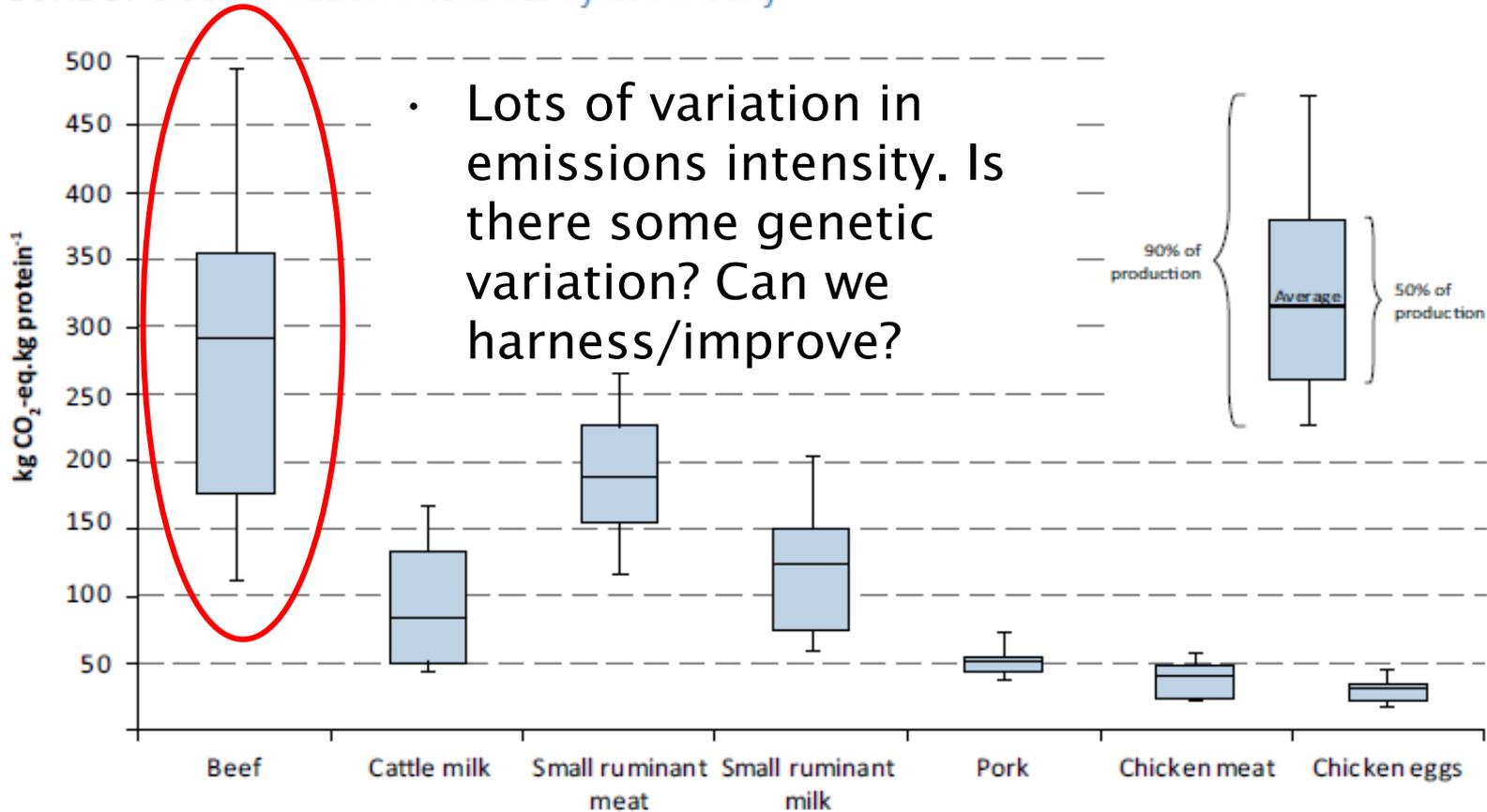
# Innovation; One Approach!



To help the environment farmer Mick converted his tractor to gas

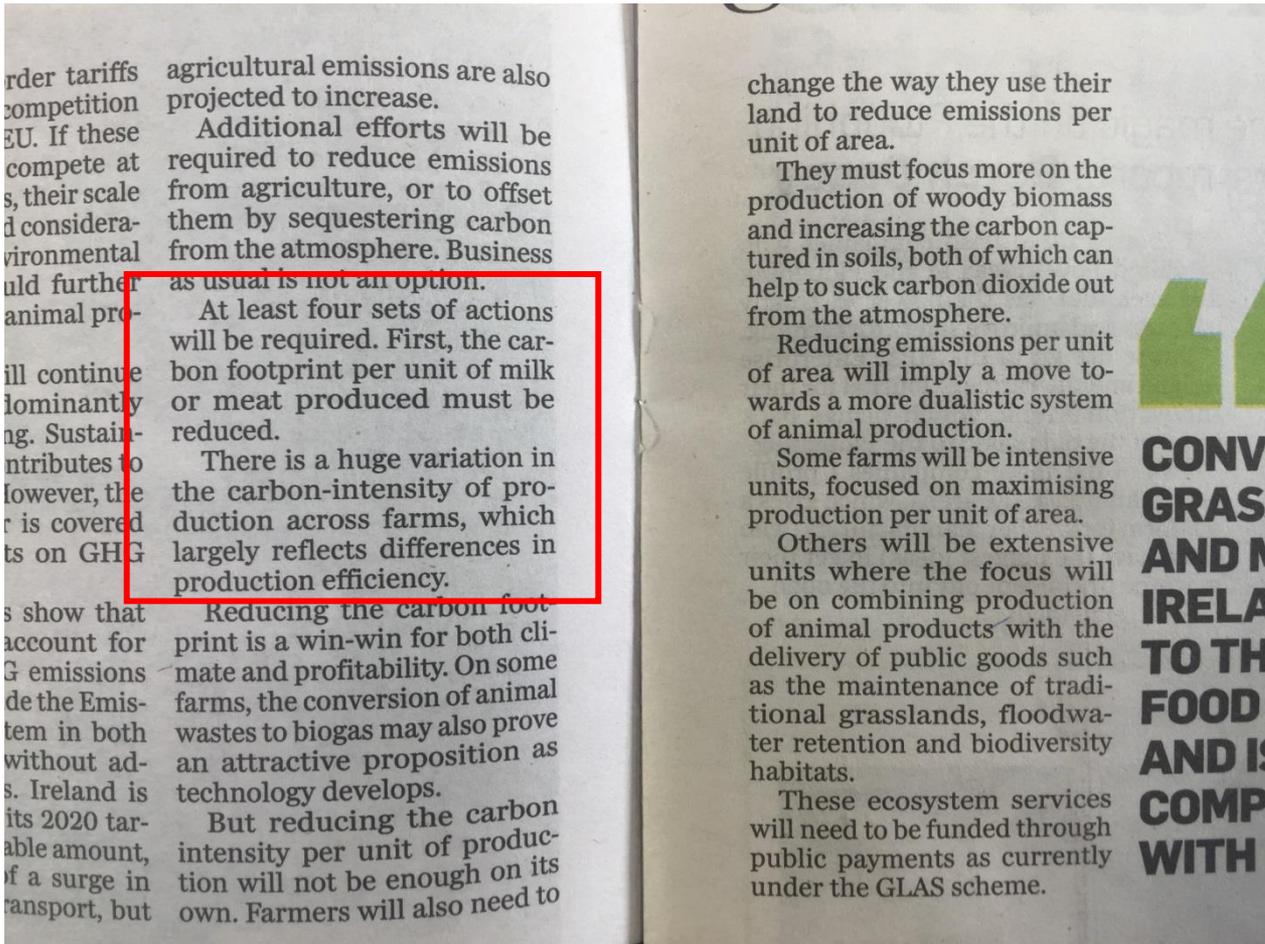
# Innovation; Another Approach!

FIGURE 3. Global emission intensities by commodity



Source: GLEAM. FAO, 2013

# Creating a “Win:Win”.



- Alan Matthews, Professor of European Agricultural Policy, TCD.
- Key climate change advisor for IRL and EU

# Genetic parameters in Australian Beef Cattle (de Haas et al., JAM 2016)

Trait	$\sigma^2_a$	$\sigma^2_p$	MeP	RPM	RGM	DMI	WT
MeP	49.7	166.9	<b>0.30</b> ( <b>0.06</b> )	0.65 (0.11)	0.55 (0.14)	<b>0.83</b> (0.05)	<b>0.80</b> (0.06)
RPM	12.9	84.7	0.71 (0.02)	<b>0.19</b> ( <b>0.05</b> )	0.98 (0.02)	0.04 (0.17)	-0.01 (0.17)
RGM	11.8	96.7	0.62 (0.02)	0.94 (0.00)	<b>0.15</b> ( <b>0.05</b> )	0.00 (0.18)	0.00 (0.18)
DMI	0.1	0.2	0.70 (0.02)	0.00 (0.04)	-0.10 (0.03)	<b>0.39</b> ( <b>0.06</b> )	0.98 (0.01)
WT	415.4	1010.6	0.67 (0.02)	0.00 (0.04)	0.03 (0.03)	0.93 (0.01)	<b>0.41</b> ( <b>0.06</b> )

\* Heritabilities on diagonal, genetic correlations above diagonal, phenotypic correlations below diagonal

- *Smaller cows => eat less feed => produce less methane.*

# €uro-Star Replacement Index.

Trait	Goal	Relative wt
Calving ease	More	16%
Feed efficiency	More	15%
Carcase	More	15%
Maturity	More	15%
Female fertility	More	25%
Docility	More	4%

**Emphasis:**

**Cow traits 71%**

**Calf traits 29%**

# 5 star cows are more profitable & more carbon efficient.

Stars	Repl Index	Cow Wt	Calf wean weight.	Calving Interval	Progeny carcass wt	Co2 Output
5 Stars	€108	669 kg	336 kg	403 days	358 kg	3,355 kg
4 stars	€86	680 kg	324 kg	407 days	356 kg	3,432 kg
3 Stars	€60	690 kg	319 kg	411 days	356 kg	3,475 kg
2 stars	€43	691 kg	315 kg	416 days	357 kg	3,502 kg
1 Star	€12	739 kg	309 kg	423 days	357 kg	3,552 kg

- Compared to 1 star cows, 5 stars are;
  - more profitable, sustainable & carbon efficient (+€100/cow).
  - Cows that will produce more from less.

# Food Wise; Smart & Green.

- Apply the latest DNA technology to support an important indigenous industry
- Simultaneously addressing global challenges around GHG and food security



 Department of  
Agriculture,  
Food and the Marine  
An Fórsa  
Talmhaíochta,  
Bia agus Mara

LOCAL ROOTS GLOBAL REACH  
**Food Wise 2025**  
A 10-year vision for the Irish agri-food industry

# The Irish Beef Genomics Scheme.

- €300m total funding 6 years (2015-2020)
  - Farmers paid ~€90/cow/year to complete key actions re: the scheme, e.g., data recording, genotyping & targets for 4/5 star cows & bulls.
  - 24k farms & 600k cows.
  - ~1.2 m animals genotyped to-date. ~2.5m animals in total will be genotyped during period of scheme.
  - Current cost of genomic service is €22/animal.
- Subsidy or investment?
  - How much would it cost to collect this data through team technicians => must think differently.

# Key Project partners.

- **DAFM/EU**; Scheme “owners” and responsible for scheme delivery.
- **ICBF**; Data collection, genotyping, delivery of genetic/genomic evaluations & reports.
  - Role of Scientific Advisory Committee (Amer, Garrick, Mantysaari, Meuwissen & Veerkamp).
- **Teagasc**; Research, extension & training.
- **Illumina**; Delivery of IDB 54k cust chip.
- **Weatherby’s/Eurofins**; Lab services.
- **Bord Bia**; Carbon Navigator.

# Beef performance evaluations in a multi-layered and mainly crossbred population

R.D. Evans\*, J.F. Kearney\*, J.McCarthy\*, A. Cromie\* and T. Pabiou\*

\*Irish Cattle Breeding Federation, Highfield House, Bandon, Cork, Ireland



Department of Agriculture database

Calf registration, animal movements; 2.3 m calves per year (1.4m dairy & 900k beef)

Feed intake, live-weights, linear classification, carcass, meat eating quality; 400 animals/year



Progeny test

Sire ancestry, calving scores, live-weights, calf quality scores, docility, fertility; 1.5m

Live-weights and sale prices /year; 1,8m/year



50 Cattle Auctions

60k commercial Beef & 18k Dairy Farms

Ancestry, live-weights, linear scores, fertility; 30k calves/yr

Carcass data; 1.6 m/year



40 Abattoirs

Pedigree beef herds 14 herd societies

Genotyping companies

Genotypes; 350k/yr

500k/year

Inseminations; 400k/year

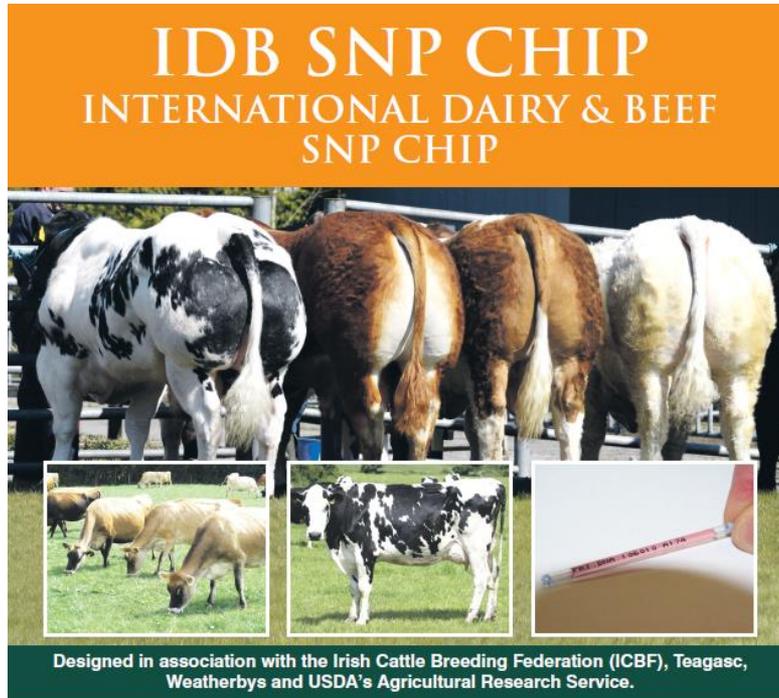


6 AI Companies

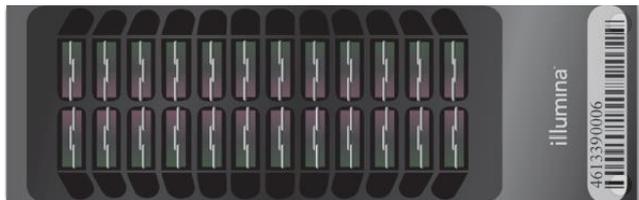
Beef Genomics Scheme; farmer recorded data.



# IDB Chip – The database in 54k SNP's!



- The International Dairy & Beef Chip.
- Developed in Ireland, with Illumina. Currently on v3.
- 54k SNP's.
  - 40k core, 6k for better imputation, 7k for “regions of interest” & 1k for major genes/defects.
- 160 Major genes/defect.
  - Database will drive this.
- See Jen McClure poster.



# Benefits of farmers recorded data.

Form 2: Beef Data & Genetics Programme - Requirement 2: Ewe and Lambing Data

Form ID: 01204507 (1/2016) Date: 24-Jan-2016

The following tables are missing a valid site and/or lambing year. Please review and correct these details and return this page in the separate envelope provided.

Notes: Contact information on site and lambing year must be given as part of your requirements under the Beef Data & Genetics Programme.

Only the Data not listed in the table below may be recorded.

Calf Tag	Calving Date	Valid Box A1 Code as per Box Tag	Sex	Retention
8-8818	2015-01-11	1 2 3 4	<input type="checkbox"/>	<input type="checkbox"/>
2-8818	2015-01-11	1 2 3 4	<input type="checkbox"/>	<input type="checkbox"/>
4-8818	2015-01-11	1 2 3 4	<input type="checkbox"/>	<input type="checkbox"/>
5-8818	2015-01-11	1 2 3 4	<input type="checkbox"/>	<input type="checkbox"/>
6-8818	2015-01-11	1 2 3 4	<input type="checkbox"/>	<input type="checkbox"/>
7-8818	2015-01-11	1 2 3 4	<input type="checkbox"/>	<input type="checkbox"/>
8-8818	2015-01-11	1 2 3 4	<input type="checkbox"/>	<input type="checkbox"/>
9-8818	2015-01-11	1 2 3 4	<input type="checkbox"/>	<input type="checkbox"/>
10-8818	2015-01-11	1 2 3 4	<input type="checkbox"/>	<input type="checkbox"/>

Signature of herd owner: \_\_\_\_\_ Date: 11 12 13 14

Form 3: Beef Data & Genetics Programme - Requirement 2: Annual Survey

Form ID: 01204507 (1/2016) Date: 24-Jan-2016

The calves listed below were alive and in your herd at 5 months of age. Calf quality and calf destiny must be recorded on all animals in this section.

Calf Tag	Birth Date	Calf Quality	Calf Destiny
81234567891011	01-Jan-2014	WG D A P NP	WG D A F WF
81234567891012	08-Jan-2014	WG D A P NP	WG D A F WF
81234567891013	15-Feb-2014	WG D A P NP	WG D A F WF
81234567891014	01-Jan-2014	WG D A P NP	WG D A F WF
81234567891015	07-Jan-2014	WG D A P NP	WG D A F WF
81234567891016	08-Jan-2014	WG D A P NP	WG D A F WF
81234567891017	10-Feb-2014	WG D A P NP	WG D A F WF
81234567891018	01-Jan-2014	WG D A P NP	WG D A F WF
81234567891019	01-Jan-2014	WG D A P NP	WG D A F WF
81234567891020	08-Jan-2014	WG D A P NP	WG D A F WF
81234567891021	10-Feb-2014	WG D A P NP	WG D A F WF
81234567891022	01-Jan-2014	WG D A P NP	WG D A F WF
81234567891023	10-Feb-2014	WG D A P NP	WG D A F WF
81234567891024	01-Jan-2014	WG D A P NP	WG D A F WF

Signature of herd owner: \_\_\_\_\_ Date: 11 12 13 14

Form 4: Beef Data & Genetics Programme - Requirement 2: Slaughter Reasons

Form ID: 01204507 Date: 24-Jan-2016

The animals that have left your herd, since the start of your 2014 lam season, are listed below. Mark as 10 if the most recent reason for the animal leaving your herd is at all. For a more extensive list of reasons, report this information on page 02 of this form.

All slaughter reasons must be recorded as 10.

Cow Tag	Separation Date	Reasons for Leaving the Herd
81234567891011	11-Feb-2014	10 11 12 13 14 15 16 17 18 19 20 21 22 23 24
81234567891012	11-Feb-2014	10 11 12 13 14 15 16 17 18 19 20 21 22 23 24
81234567891013	11-Feb-2014	10 11 12 13 14 15 16 17 18 19 20 21 22 23 24
81234567891014	11-Feb-2014	10 11 12 13 14 15 16 17 18 19 20 21 22 23 24
81234567891015	11-Feb-2014	10 11 12 13 14 15 16 17 18 19 20 21 22 23 24
81234567891016	11-Feb-2014	10 11 12 13 14 15 16 17 18 19 20 21 22 23 24
81234567891017	11-Feb-2014	10 11 12 13 14 15 16 17 18 19 20 21 22 23 24
81234567891018	11-Feb-2014	10 11 12 13 14 15 16 17 18 19 20 21 22 23 24

Signature of herd owner: \_\_\_\_\_ Date: 11 12 13 14

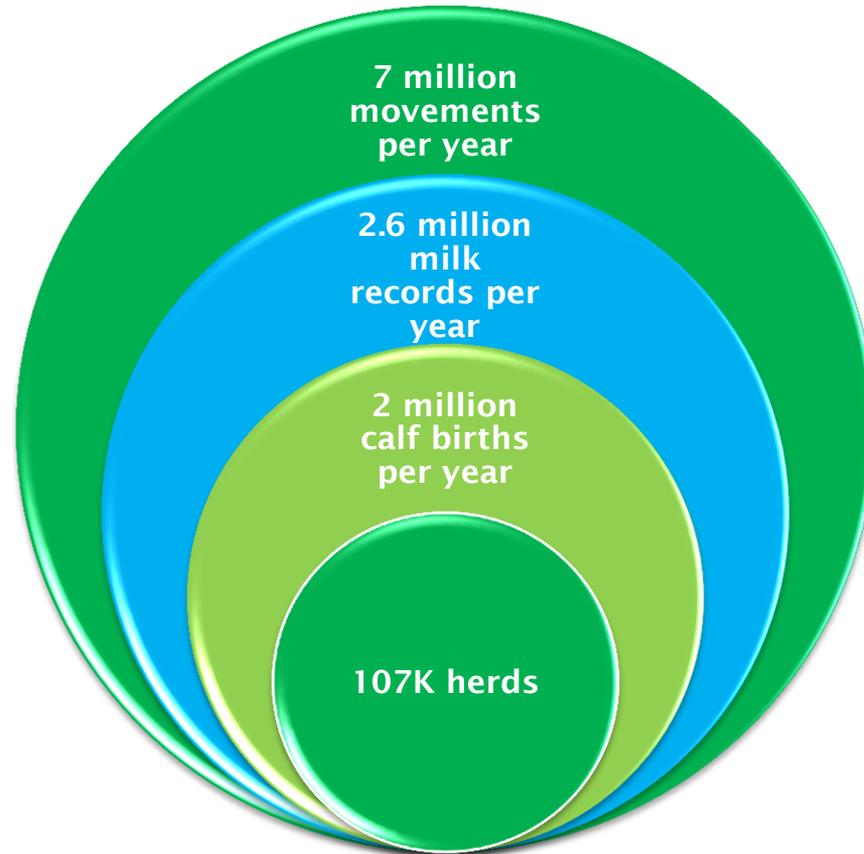
## Information Collected

- Calf
  - Sire
  - Calving Ease
  - Birth Size
  - Vigor
  - (at 5 months)
    - Quality
    - Docility
    - Scour
    - Pneumonia
- Dam
  - Docility
  - Milk-ability
  - Departure Reason
- Stock Bull
  - Docility
  - Functionality
  - Departure Reason

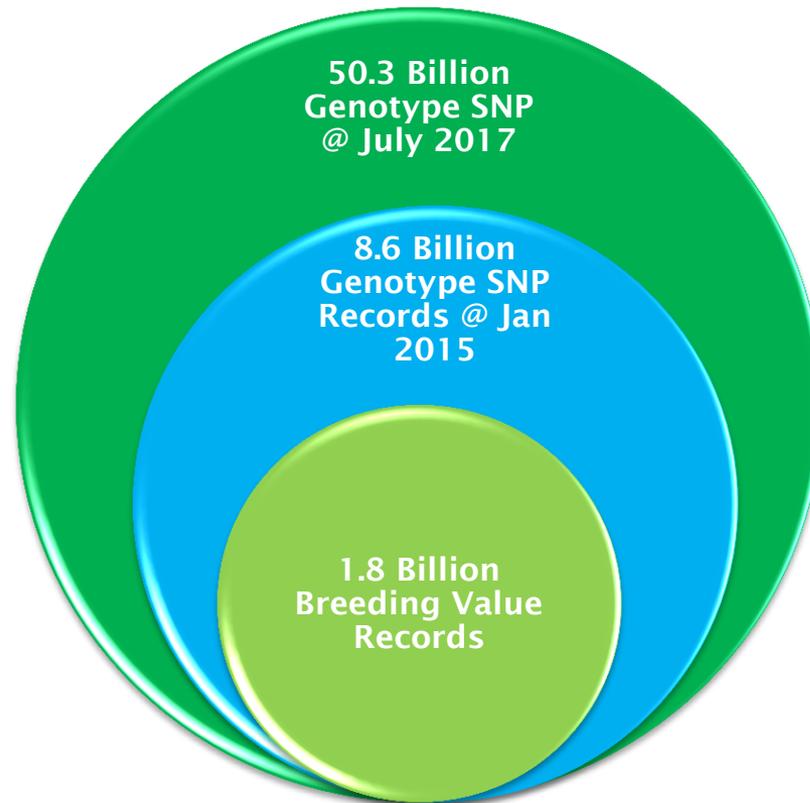
- ~600k records/trait/year (cow & calf).
- Excellent heritability estimates, e.g., docility @ 30% with rg of 0.85 with data from “expert” scorers.



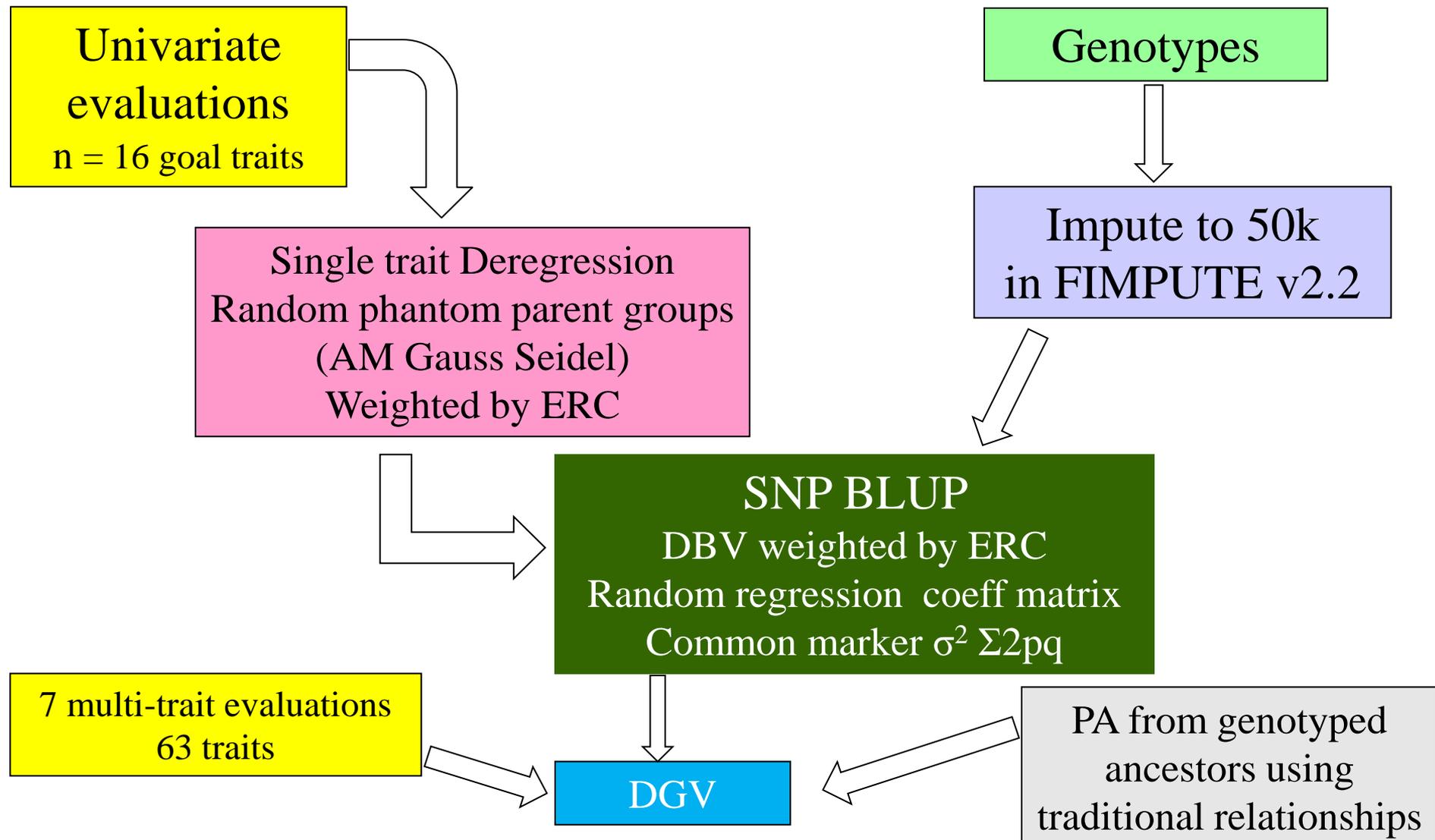
# Regular Data Set – What we used to think of as ‘Big Data’



# Big Data => Genotype Datasets.



# 2-Step Genomic Evaluation (Mix99)



- Blending using selection index methodology, (Van Raden et al. 2009)
- Research underway to move to “One-Step” (2018+).

# Computer requirements for ICBF evaluations



iPhone 5  
1 GB RAM

Cerus x 2  
Ram 6,000 GB  
Disk 40,000 GB

Igen2  
Ram 760 GB  
Disk 4,000 GB

198  
Ram 356 GB  
Disk 1,500 GB

151  
Ram 120GB  
Disk 1,800GB

143  
Ram 64GB  
Disk 698GB

109  
Ram 16GB  
Disk 279GB

163  
Ram 2GB  
Disk 80GB

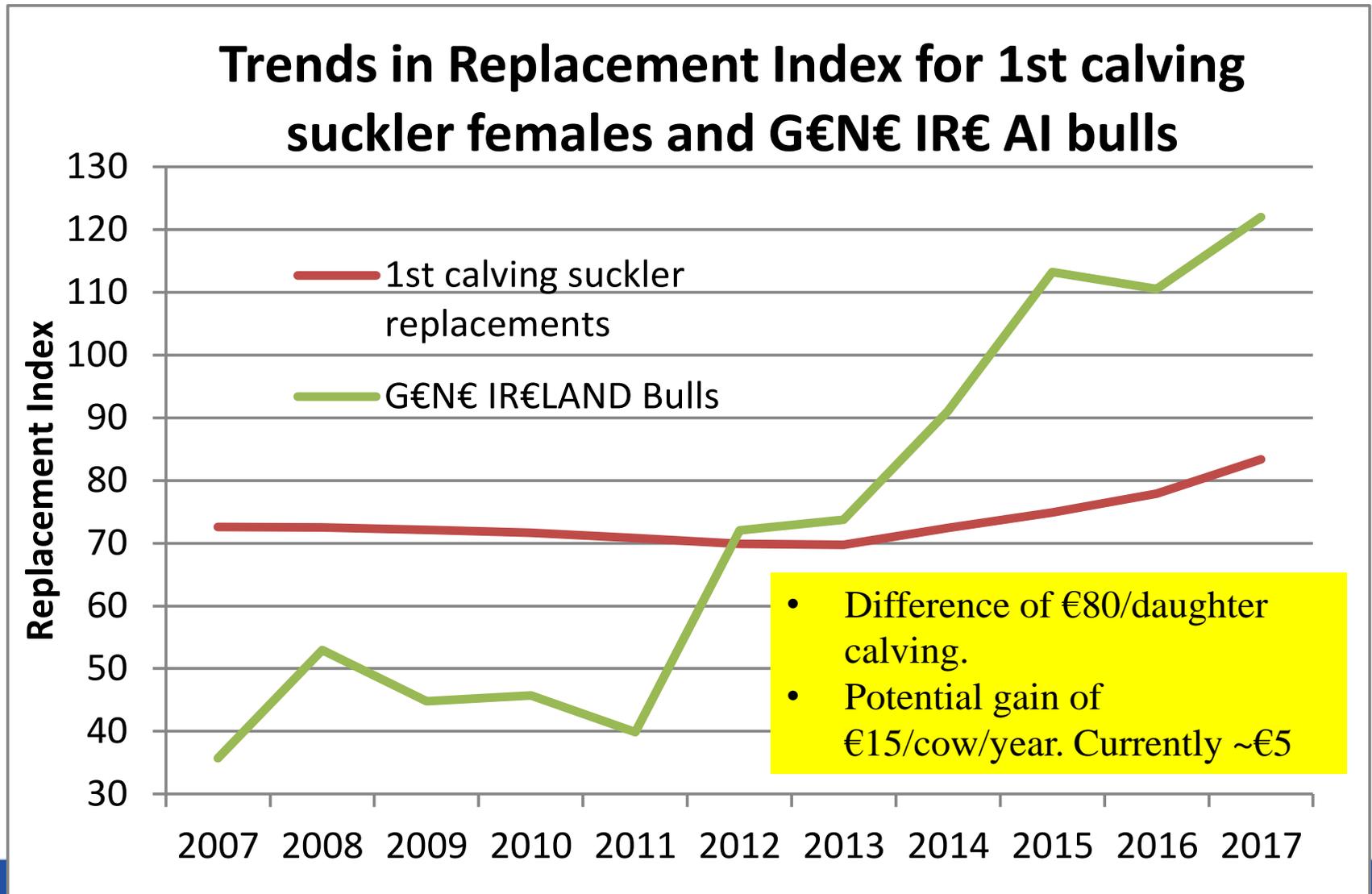
2005	2007	2010	2013	2015
------	------	------	------	------

# Validation carcass wt AI sires

- AI sires (n = 524) with first progeny born in 2012 had all their progeny phenotypes omitted (21,428 progeny)

	Validation category		
	Holstein	ALL Beef breeds	Charolais
Current Reliability	>95%	>95%	>85%
N	36	49	16
	Correlation with current deregressed univariate ebv		
EBV uni validation	0.750	0.891	0.590
EBV multi validation	0.719	0.899	0.584
DGV validation	0.790	0.910	0.628
GEV validation	0.788	0.893	0.674
% of bulls whose sires were in SNP BLUP	100%	49%	56%

# Progress has started – and will increase!



# Visit to Tully Tomorrow.



- Live-weight & feed intake; The correlation is not 1!
- Which animal ate the most?!
- Visit to Tully tomorrow. Interested?

# Summary.

- Genomics will have a major role in addressing global challenges around environment and food security in the future.
- The Irish beef genomics scheme has had a huge impact on the uptake/interest in genetic gain in Ireland.
- Strongly encourage other countries to consider such programs for their countries in the future => opportunities for collaboration.

