



## Sheep genetics research

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IFA Meeting, 2nd December 2021



#### Outline

1. Overview Irish breeding programme

2. Genomic Selection

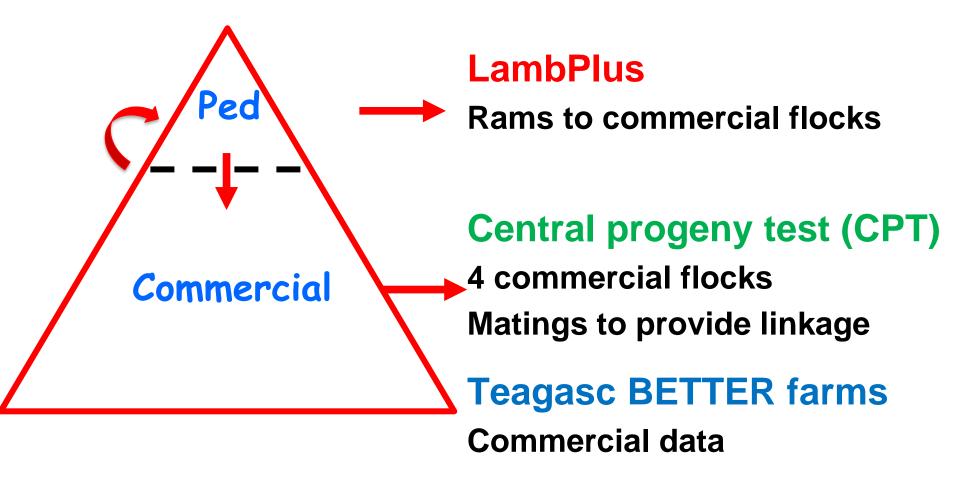
3. Future research priorities

4. Ireland's long term strategy



## Irish sheep breeding programme

**Data Sources** 





## Irish sheep breeding programme





Breeding Objectives



Lambing

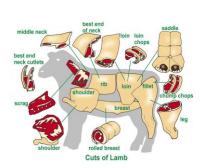


Growth



Days to slaughter

Meat



Carcass fat & conformation

Health



Lameness & dag score

Difficulty & Survival

## Irish sheep breeding programme



Breeding Objectives



Replacement

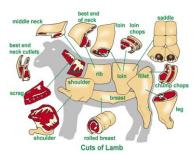
Ewe traits



Lambing



Meat



Health



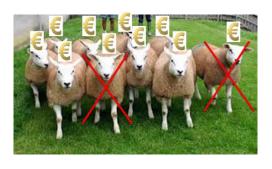
Milk Yield No. lambs born Ewe weight

Difficulty & Survival

Carcass fat & conformation

Lameness & dag score

## **Recent Improvements**







Scrapie Validation



Carcass Data



ACROSS-BREED GENOMIC EVALUATIONS

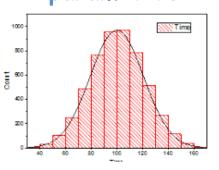
#### Across-Breed Model



#### New Traits

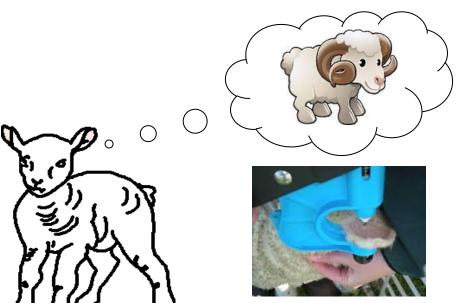
- Barren Rate
  - Lameness
  - Dag Scores

# Updating Genetic parameters

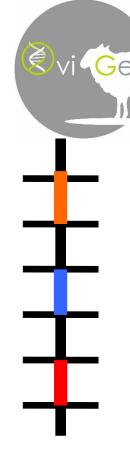




## **Genomic Selection**







- At birth we know parts of the lamb's DNA
- Lamb accuracy increase by 15 to 52%
  - » Equiv. of 6 (51) progeny for high (low) h<sup>2</sup>



## **Genomic Selection**

Applied to all genes & all traits



#### **Parentage**

Low cost option for hill and commercial flocks

## Major gene effects

Scrapie, lethal recessive genes....

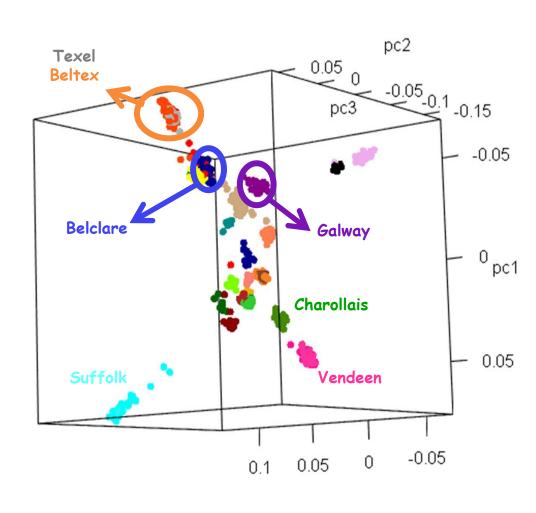
## **Breed mapping**

How closely related are breeds??



# Number of genotyped sheep

Breed	Total
Belclare	4,145
Charollais	4,843
Suffolk	4,860
Texel	8,472
Vendeen	1,956
Other - pedigree	1,337
Other - crossbred	29,808
Total	55,421





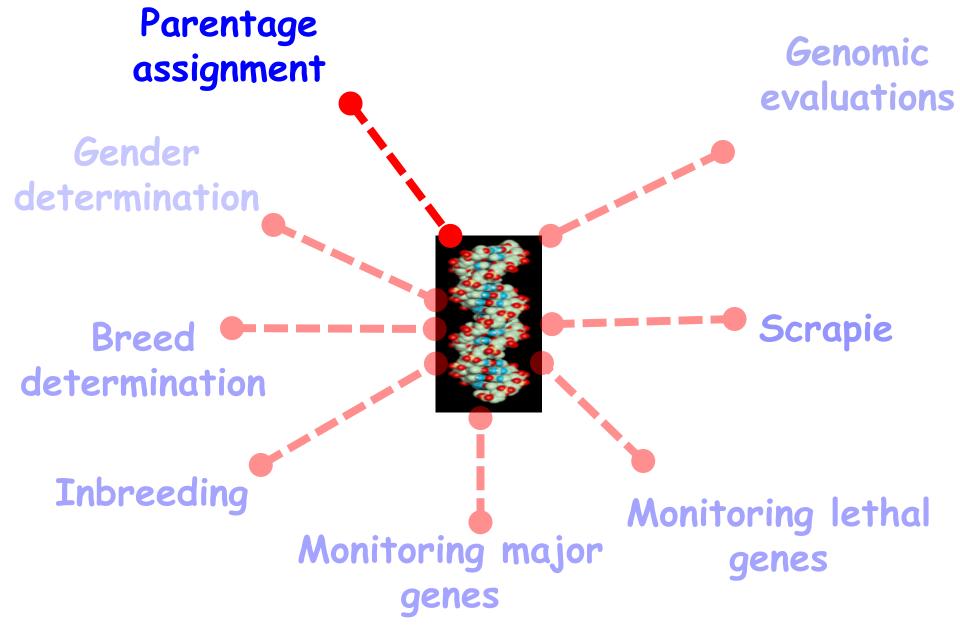






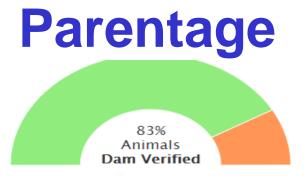










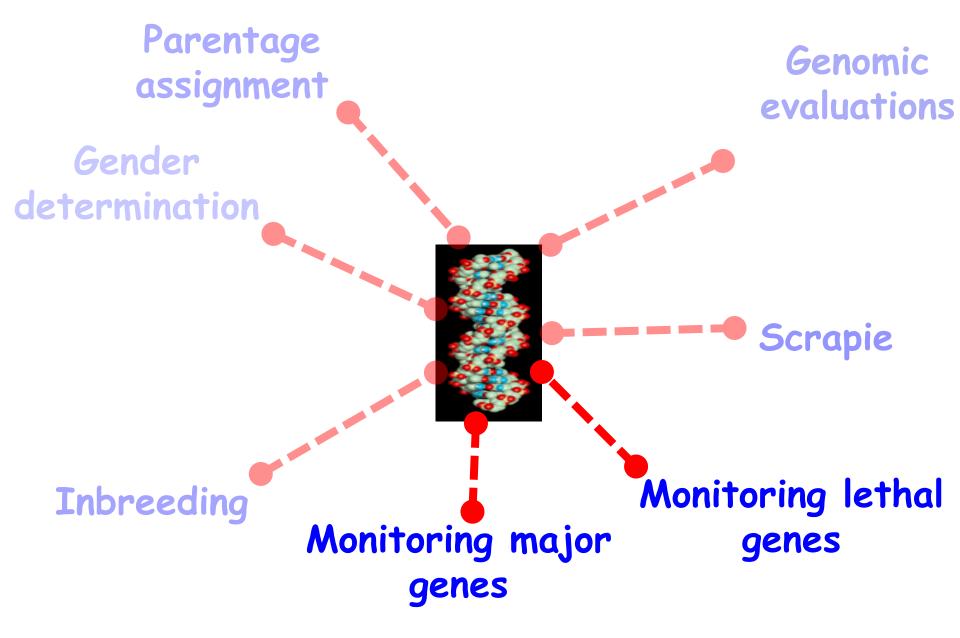


Search

 $\bigcirc$  Invalidated Sample  $\bigcirc$  Resample Needed  $\bigcirc$  Sire Mismatch  $\bigcirc$  Dam Mismatch  $\bigcirc$  All

18 animals of 521 animals found, 18 displayed

Animal	Sex	DOB	Breed	Animal Result	Tested Sire	Sire Result	Tested Dam	Dam Result
<b>IE0424133285</b> 25H	F	16-MAR-2020	SU	Resample Needed				
<b>IE0424133285</b> 57F	F	17-MAR-2020	SU	Genotyped	IE042413326083G	Mismatch New Sire IE043538902911J	IE042413324615E	Match
<b>IE0424133285</b> 51C	F	17-MAR-2020	TX 属	Genotyped		Not Recorded on Database New Sire MURPHYS CHURCHILL	IE042413321911J	Match
<b>IE0424133285</b> 83A	F	17-MAR-2020	BR 📕	Genotyped	TEAGASC BEAUTY	Match	IE042413323348G	No Result Available
IE0424133285	F	17-MAR-2020	BR	Genotyped	COOLKELLURE DUSTIN	Match	IE042413324881H	No Result Available
IE0424133285	F	17-MAR-2020	BR	Genotyped	ALLOON DANE	Match	IE042413326568G	No Result Available
<b>IE0424133285</b> 31A	F	16-MAR-2020	BR	Genotyped	FORAS EASTON	Match	IE042413326537B	No Result Available
								No Result





# Major genes

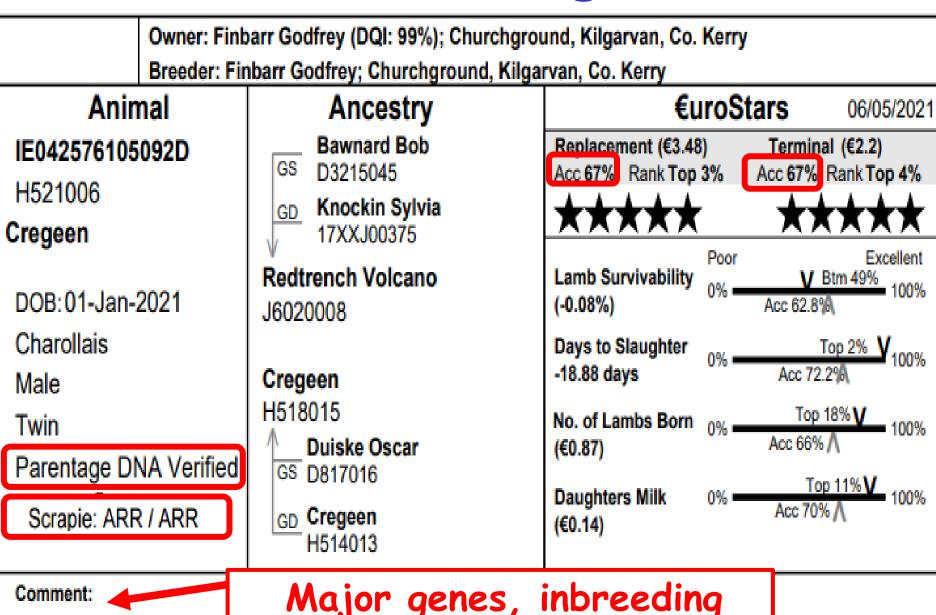
- Prolificacy genes
- BMP15 Xb → associated with Belclare breed
- 1 copy increased ovulation rate
- 2 copies sterile

Genotyped	1 сору	
All population	0.07%	
Belclare	9.78%	





# Sales Catalogue

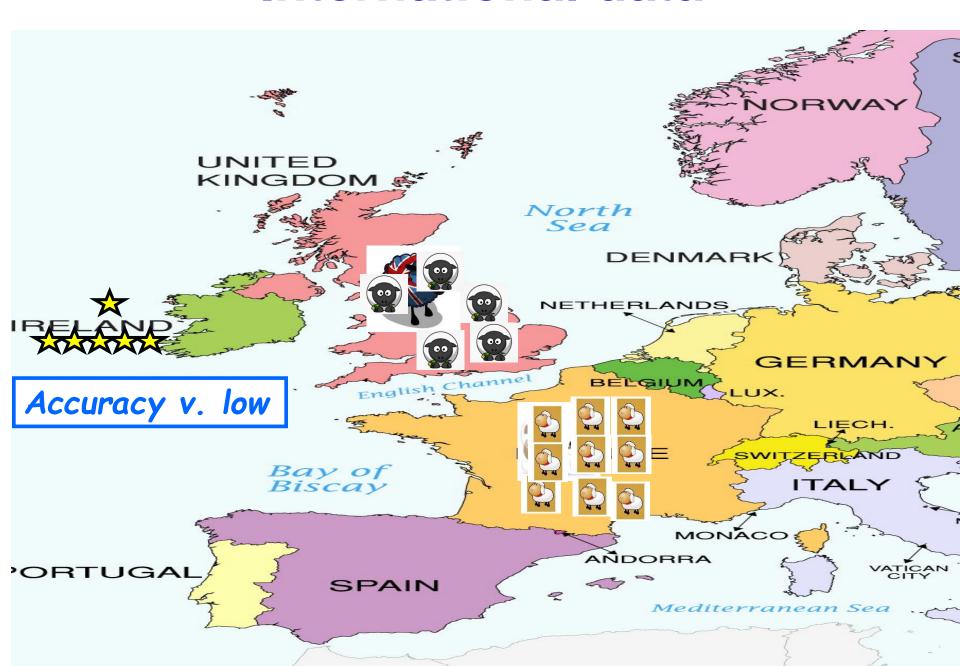


#### **Future Research**

- Across country evaluations
- Health
  - FEC, mastitis
- Methane emissions
- Survival to Weaning
- Ewe longevity/Stay-ability



## **International data**





#### Ireland versus UK



- Case-study → Texel breed
- Large number of common animals
- Strong genetic correlations between traits

Current rate of genetic gain (%)					
Pre weaning	1.81	2.37			
Post weaning	5.31	3.38			
Muscle	2.42	1.49			

Increase accuracy and genetic gain by sharing data



## Ireland versus New Zealand

#### CON

Conservative breeders



#### **PRO**

Progressive breeders who never use foreign genetics



#### **PROFOR**

Progressive breeders who use foreign genetics





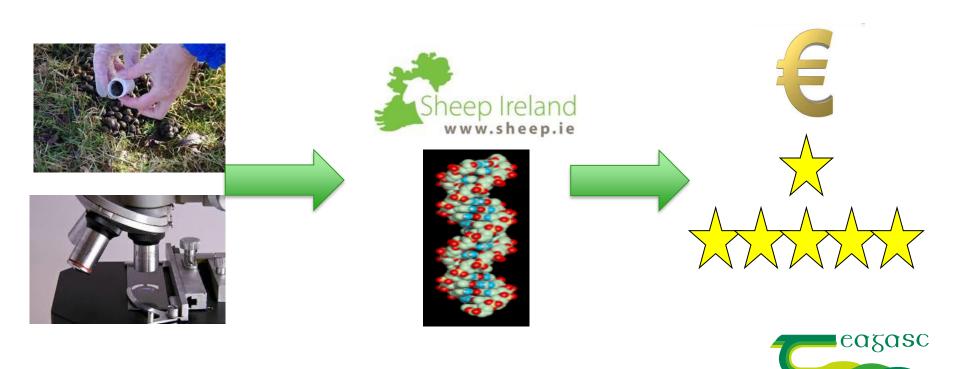
#### **FOR**

Foreign breeders who supply semen and/or live rams

	Benefit after 20 years (€m)
Current	33.598
Progressive	134.723
Foreign	135.601

# **OviFec**DAFM Stimulus Funded

Developing genetic index for resistance to gastrointestinal nematode infections



AGRICULTURE AND FOOD DEVELOPMENT AUTHORITY

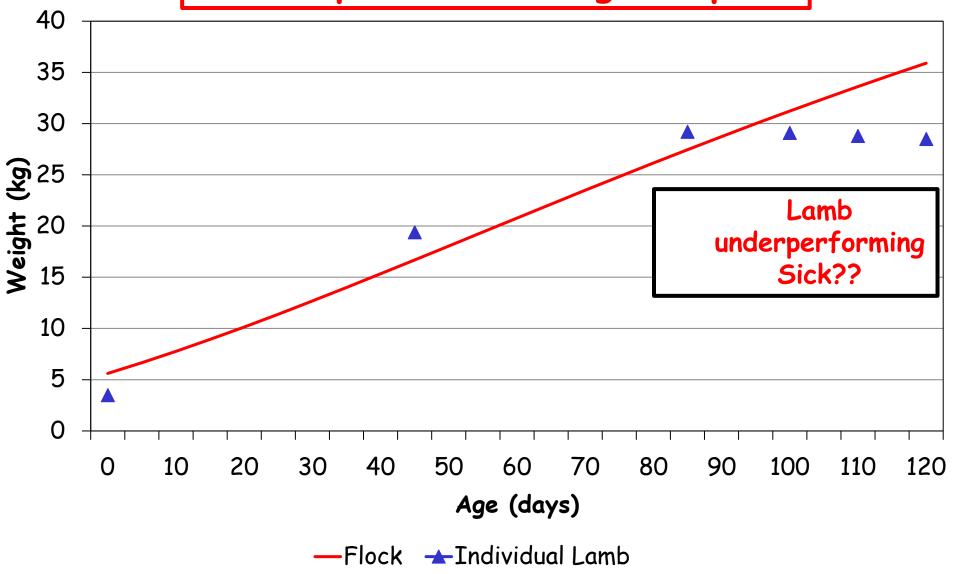
## Long term strategy

## 1. Decision Support tools



#### **Decision Support tools**

Example - flock weight report



PROBLEMENT AND E OUD EPEVELOPMENT PROTECTION !

## Long term strategy

#### 1. Decision Support tools

- 2. Animal welfare
  - Linked to health, lamb and ewe survival
- 3. Meat eating quality
- 4. Ram functionality
  - Difficulty to measure
- 5. More commercial/hill data
  - Genomics to provide parentage



#### **Conclusions**

- Large potential to increase genetic gain
- Short term
  - 1. Improve genetic evaluations
  - 2. Genomic selection & index validation
- Long term
  - Difficult to measure traits
- Integrated industry approach

# High Index Sheep Produce Less GHG per Kilo of Carcass

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<sup>1</sup>Sheep Ireland, <sup>2</sup>Teagasc





## Flocks Modelled



#### 1 Star Flock











#### 1 Star Flock

257 ewes

20 HA

Weaning 1.60 lambs/ewe

DTS: 200

#### 5 Star Flock











#### **5 Star Flock**

257 ewes

20 HA

Weaning 1.74 lambs/ewe

DTS: 192

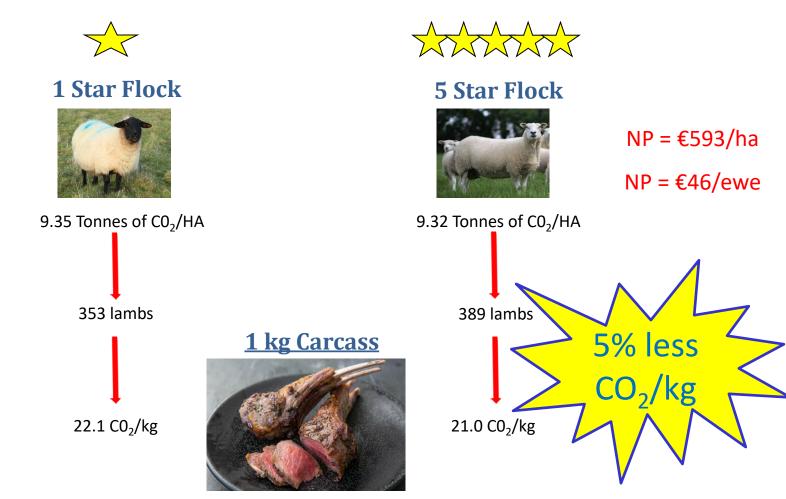
Using CPT commercial data

easasc

## **Economic and environmental output**

NP = €439/ha

NP = €34/ewe







#### **SMARTER PARTNERS**























































#### Thank you for your attention

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