



# New opportunities for genomic selection in sheep and goats

**Dr Joanne Conington**

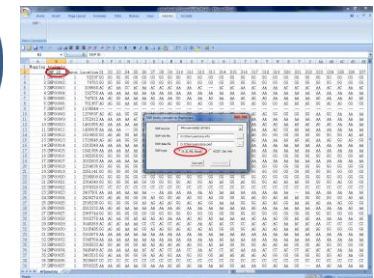
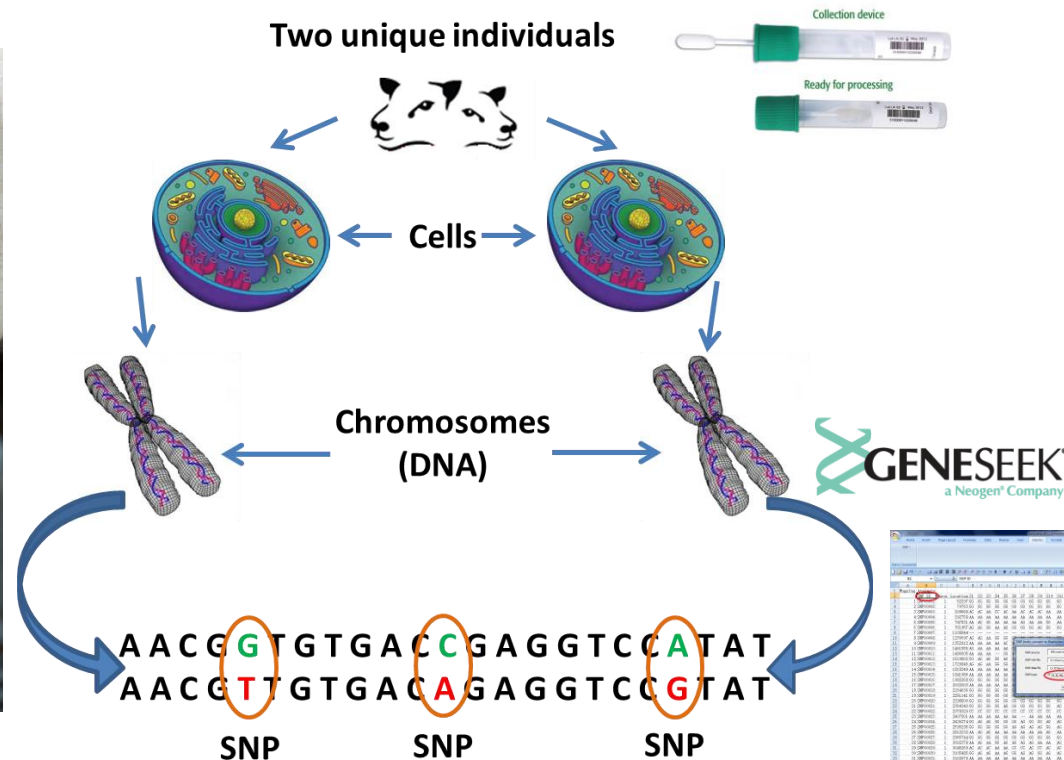


Scotland's Rural College (SRUC) Edinburgh, Scotland, UK

International Congress on the Breeding of Sheep and Goats, WCCB, Bonn, Germany 15 October 2020

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# Using genomic data in breeding programmes

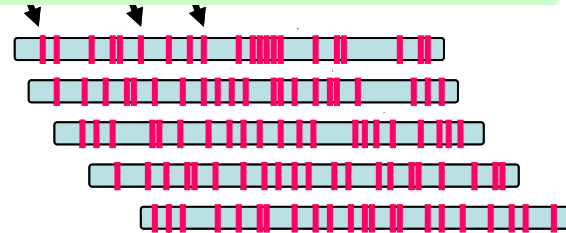


# Genomic selection

- Dense map of markers (SNPs) across the genome (50K)

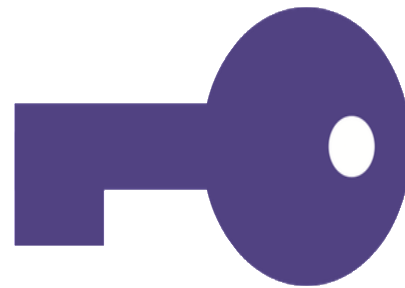


*Markers (SNPs) spaced across all chromosomes*



	SNP 1	SNP 2	SNP 3	SNP 4	SNP .....
Trait1	●	●	●	●	
Trait2	●	●	●	●	
Trait3	●	●	●	●	
Trait4	●	●	●	●	
Trait...	●	●	●	●	

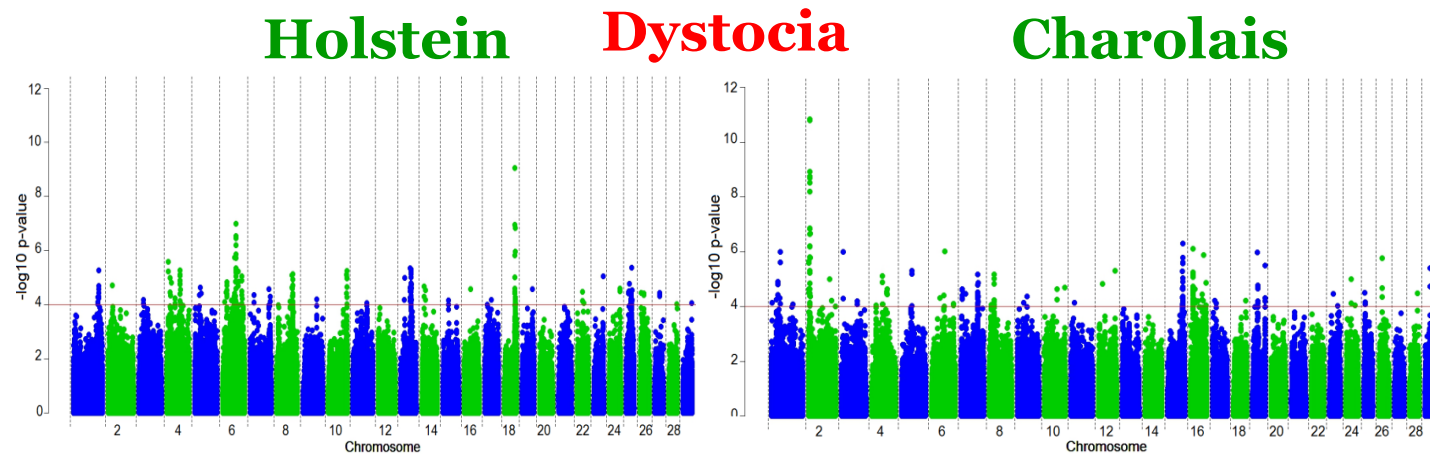




Genomic  
Estimated  
Breeding  
Values

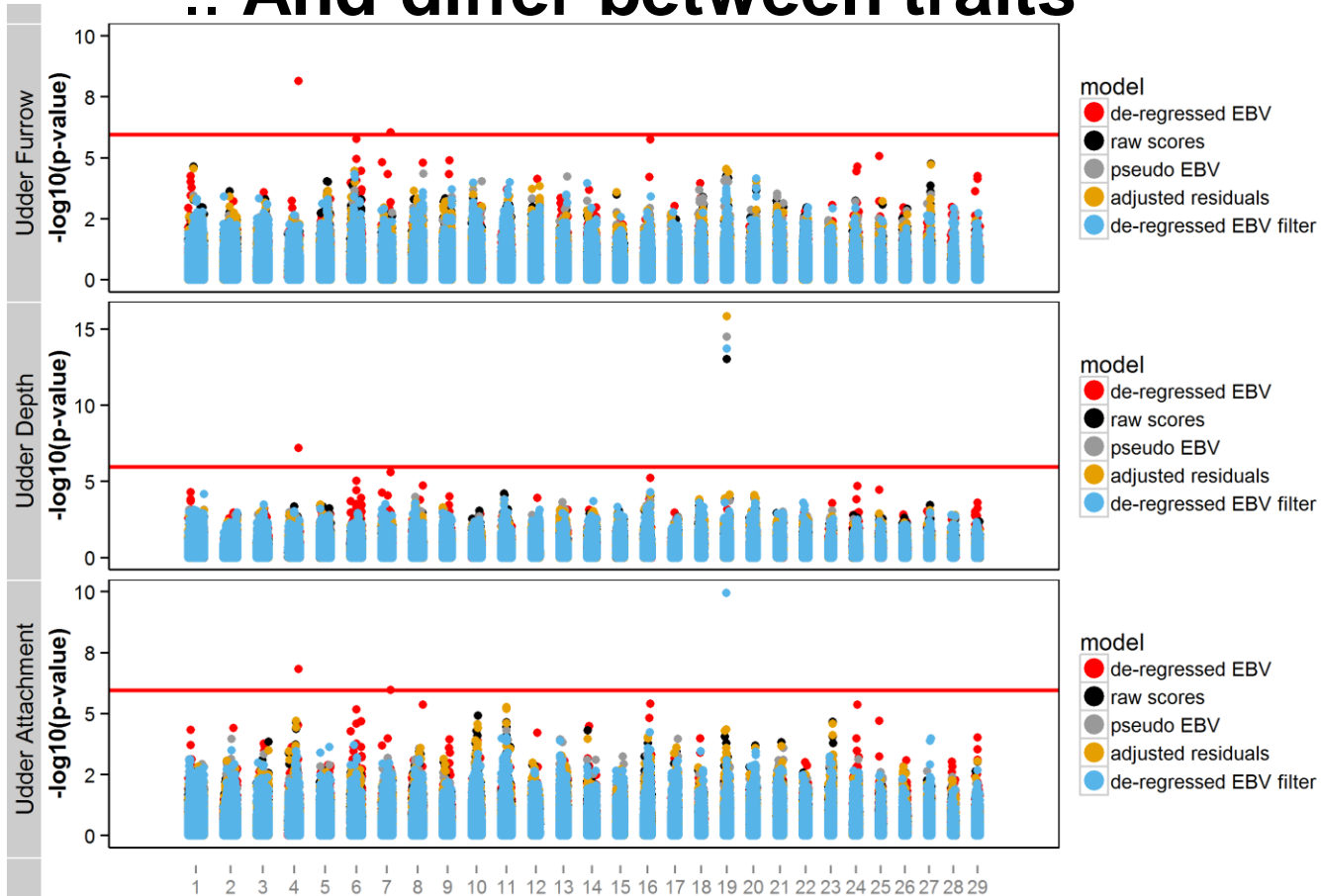
# SNP 'patterns'

DNA effects can differ between breeds



Purfield et al., (2014)

## .. And differ between traits



Mucha, Mrode, Coffey, Kizilaslan, Desire, Conington J. 2018 J. Dairy Sci.101(3):2213-2225



# Hard to measure / expensive / measured later in life traits

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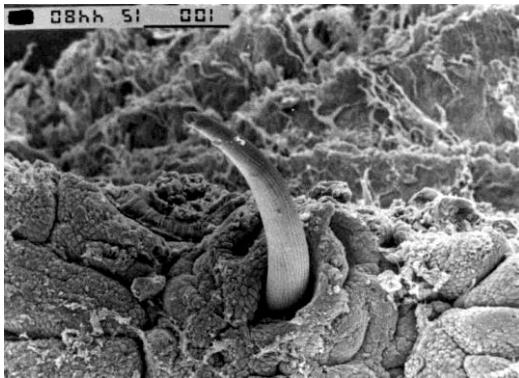
Maternal traits  
Longevity



Carcass &  
Meat quality



# Hard to measure / expensive / measured later in life traits



**Disease**



**Feed intake**

(photo courtesy of H.Wishart)



# Number of Texels with new disease phenotypes



Year	No. ewes
2015*	3,339
2016*	3,482
2017	1,712
2018	1,652

**> 10,000  
records**

**8,764  
genotypes**

\*ewes scored twice @ 38 & 115 weeks of age

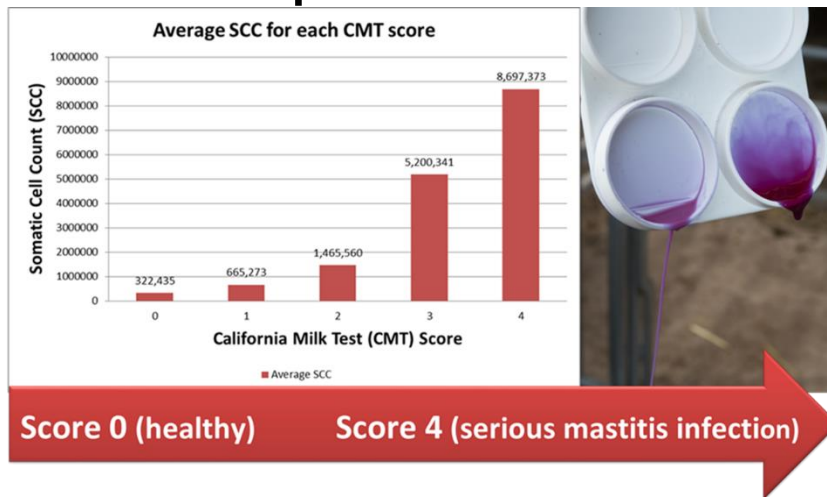
# Mastitis 'phenotyping'



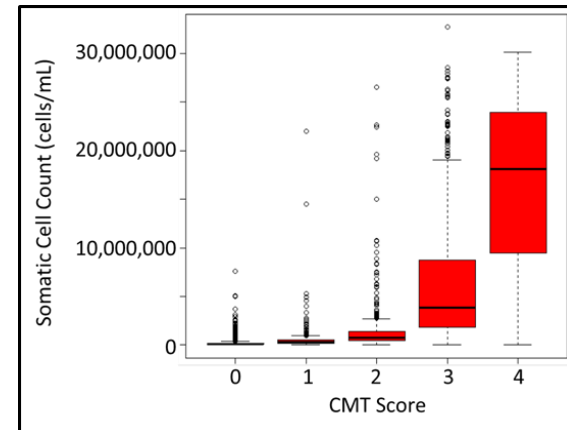
McLaren, A., Kaseja, K., Yates, J., Mucha, S., Lambe, N and Conington J. 2018 Animal 12:1-10

# Somatic Cell Count vs CMT?

- Good predictor ✓

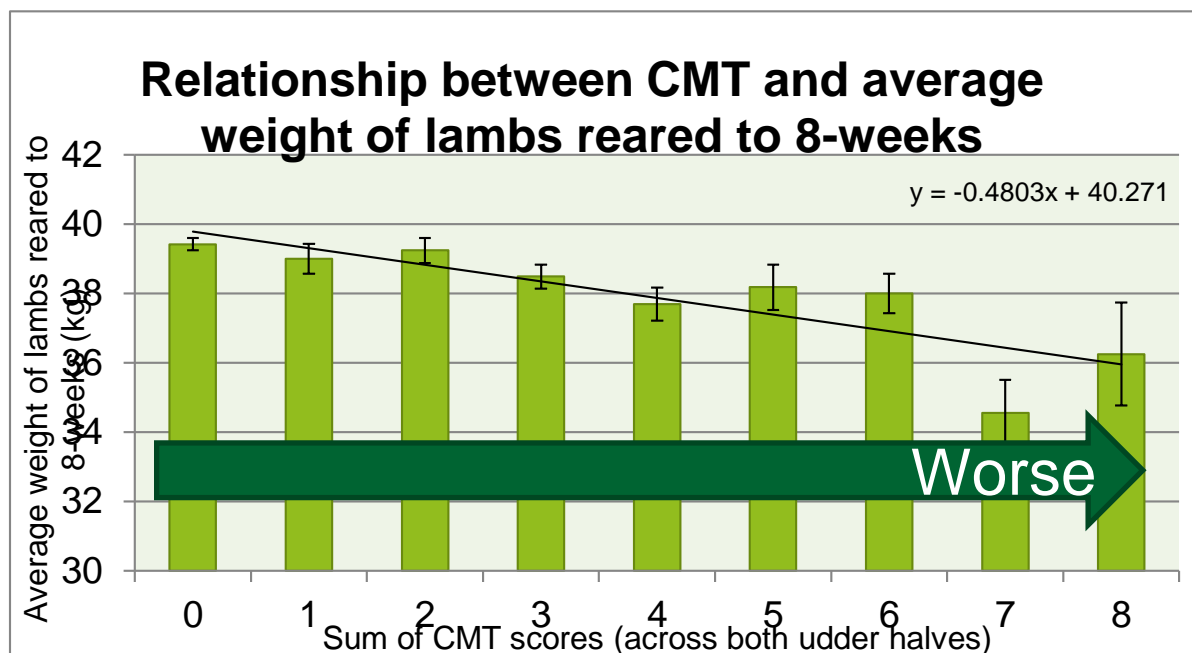


➡ Same (ish) genes



Trait	Heritability
CMT	0.14 (0.08)
SCC	0.23 (0.08)

# CMT and lamb live weights



Difference between a ewe scored 0 and scored 8:

- 3.84 kg average lamb live weight reared
- £6.30 a lamb (live weight price of £1.64 per kilo)

# Foot rot

## FOOTROT

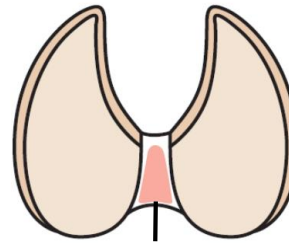


Mucha et al Genet.Sel'n.Evol'n 2015



# Foot scoring method

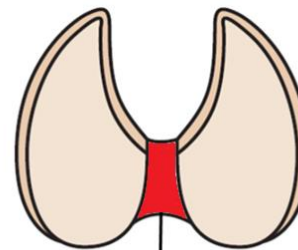
- 1: Non-specific inflammation of the interdigital skin



Interdigital  
dermatitis



- 2: Necrotic, severe in



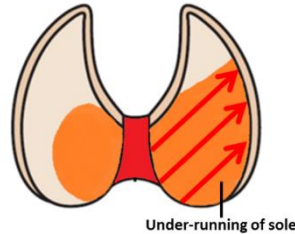
Severe  
inflammation



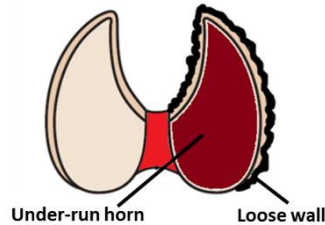


# Foot scoring method

- 3: Under-run of the sole, restricted to soft horn of the heel



- 4: Under-run of the sole of the foot including the hard horn of the toe and wall



Conington et al., Vet Res Comm 2008

# Resistance to Footrot is under low to moderate genetic control

Heritability 0.18

## Breeding resistance to footrot

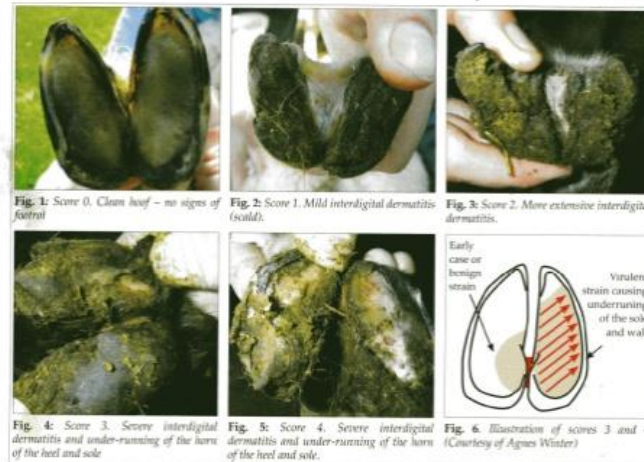
65

- Foot scoring useful tool to differentiate footrot & other hoof lesions

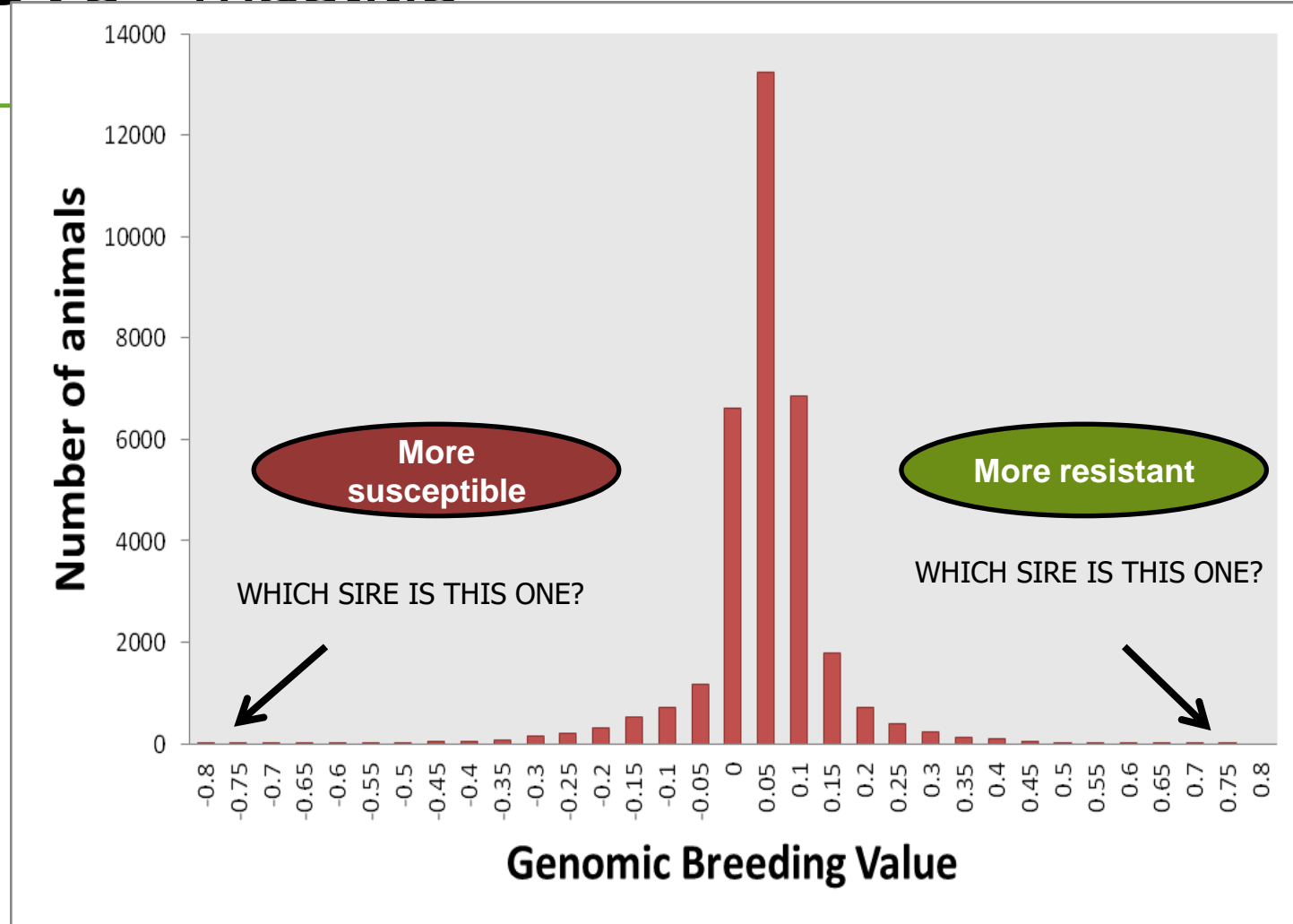
- 5-point scale,  $h^2 \sim 0.2$



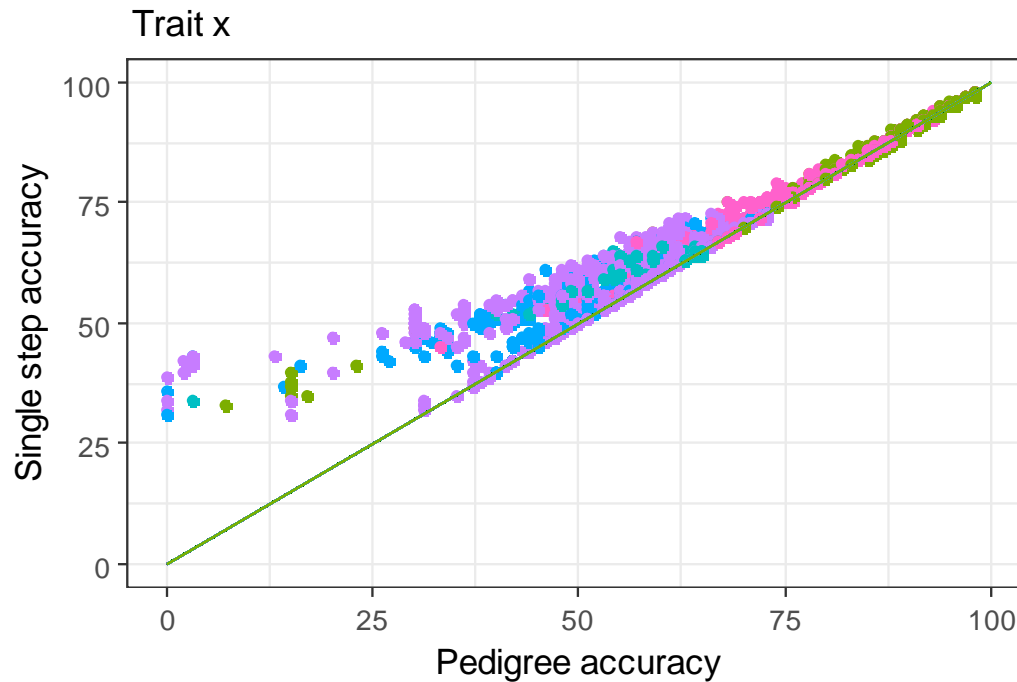
Conington et al., 2008 *Vet Res Comm.*  
Nieuwhof et al., 2008 *Animal*



# GEBVs - mastitis

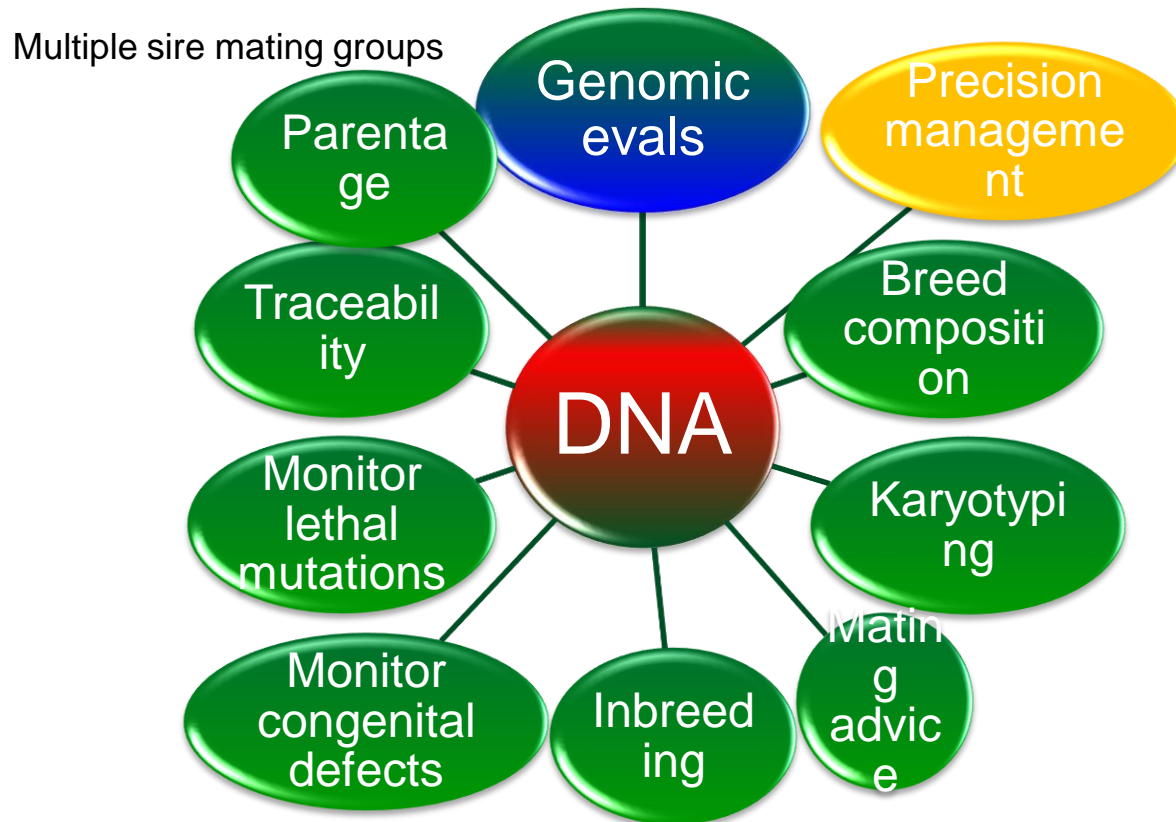


# Genomic selection most beneficial when EBV accuracy is low



D. Brown, pers. comm 2018

## What benefits ?







# Example

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- Dairy goat farm in UK
  - 2014 genomic selection
  - 14K genotyped goats (male & female\*)
  - Earlier & more accurate identification of elite animals
  - Shorter generation interval
    - Pick replacements younger
  - Higher rate of genetic gain
    - 10- ~30%



**UK's largest Goat Milk  
producer  
High-health Goats on 2  
sites in East Yorkshire.**

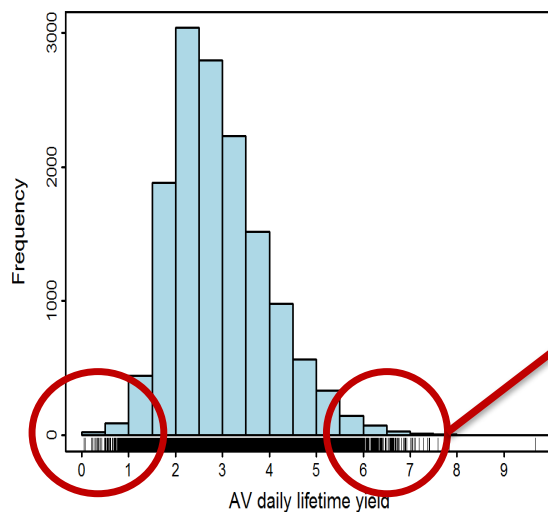
**+SRUC (2011+)**

**+ ££**

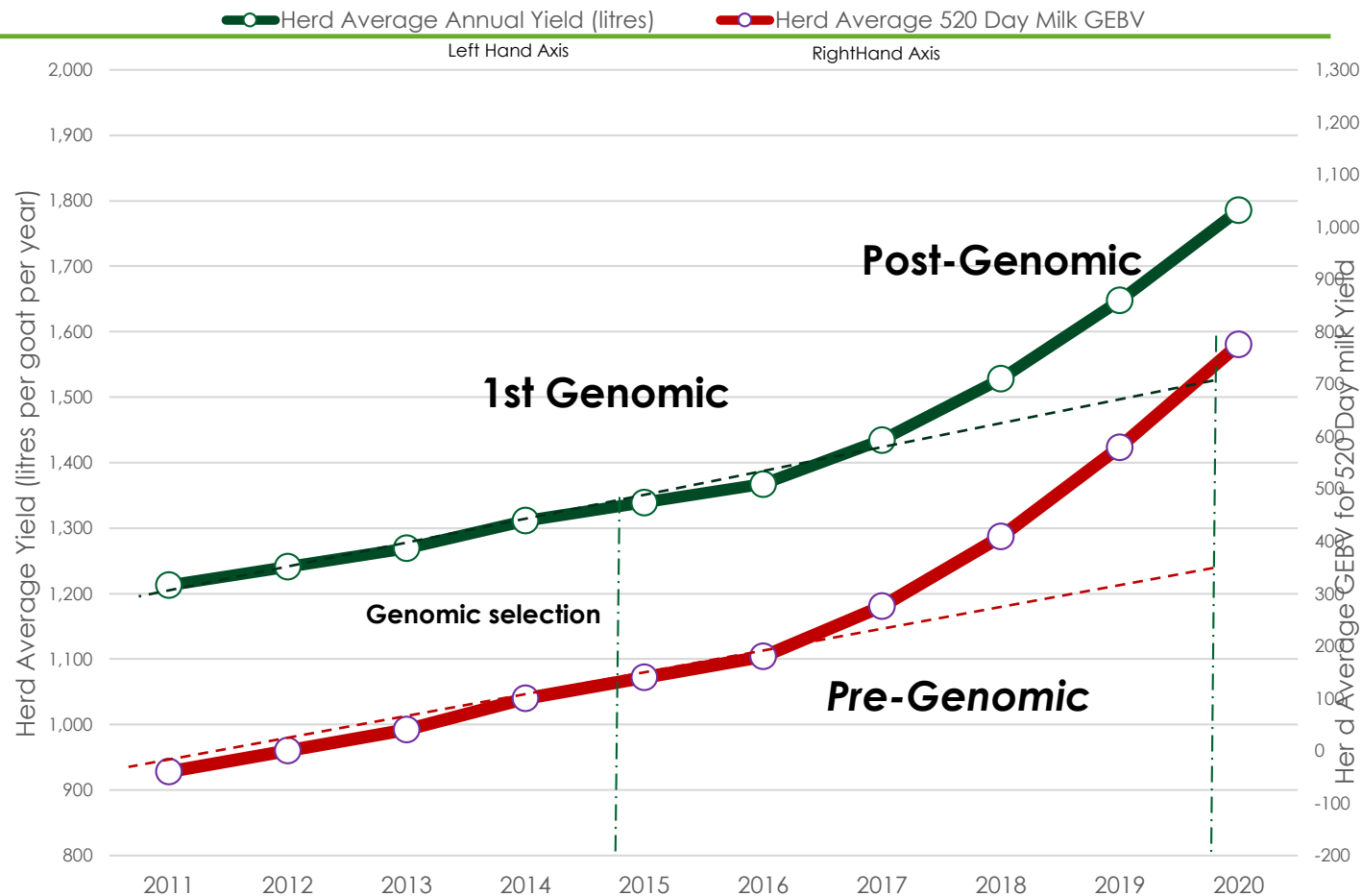
**Unique genomic  
capability  
Export & semen  
collection facility**



## How did we start?



**Innovate UK**  
Technology Strategy Board



# Sheep?

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- Australia ✓
- NZ ✓
- France (dairy sheep, within-breed) ✓
- Ireland ✓
- UK (Texel) routine planned for 2020 ~✓
  - COVID delay

# France – sorting out the reference population

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Situation in 2018	Genotyped rams	With daughters	Reference population depth	1 <sup>er</sup> complete year of birth
Lacaune	13212	5368	1996-2017	2003
RF Manech	3007	2238	1998-2017	2000
Basco-Béarnaise	945	667	1999-2017	1999
BF Manech	643	510	1996-2017	1996
Corse	746	243	2001-2017	2005

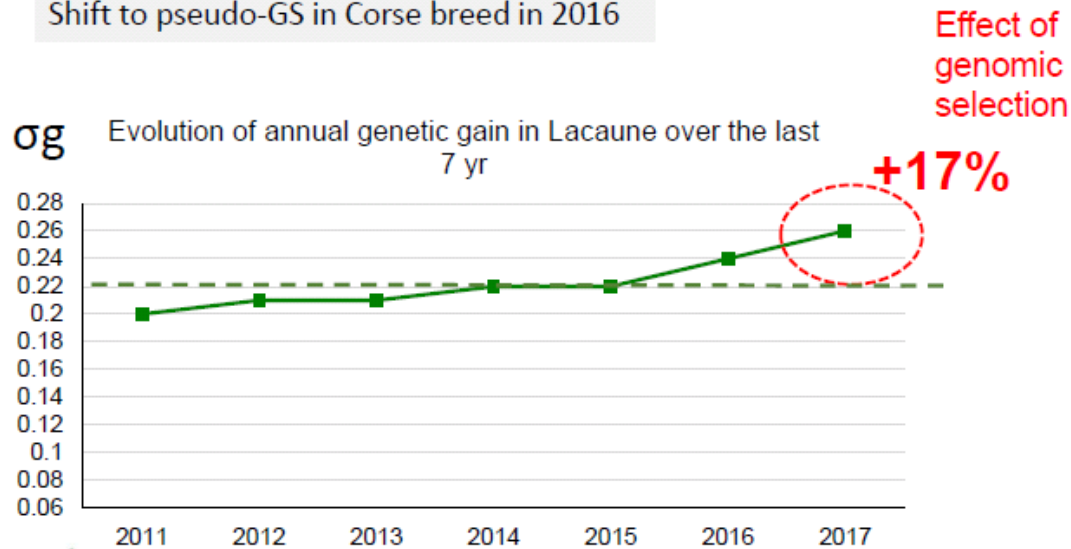
Courtesy of J.M. Astruc, 2018



# Proof it works - France

## Genomic selection: first results observed in Lacaune breed

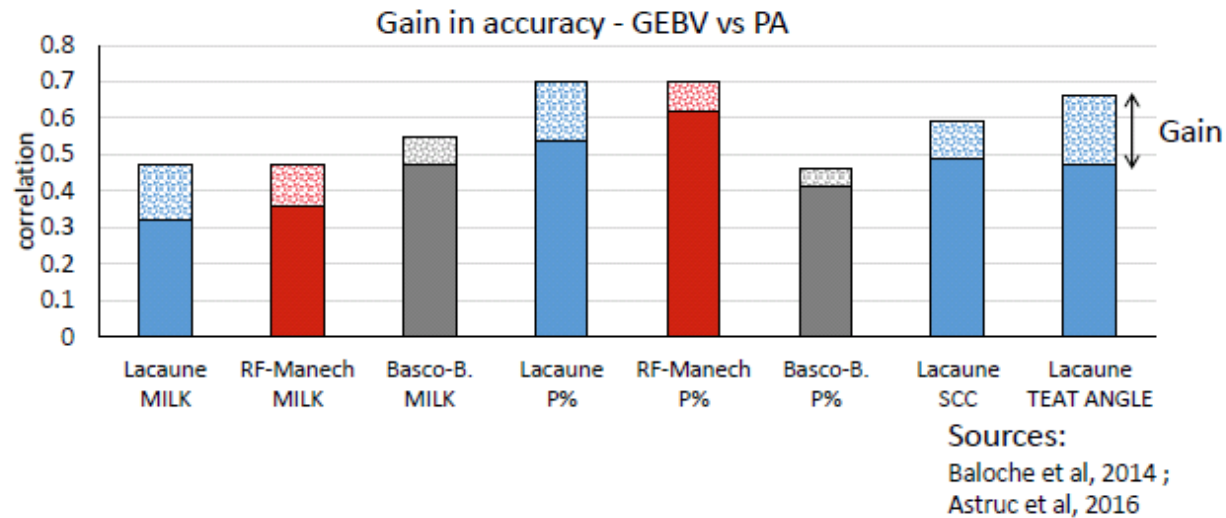
Shift to GS in Lacaune breed in 2015  
Shift to GS in Pyrenean breeds in 2017  
Shift to pseudo-GS in Corse breed in 2016



Courtesy of J.M. Astruc, 2018

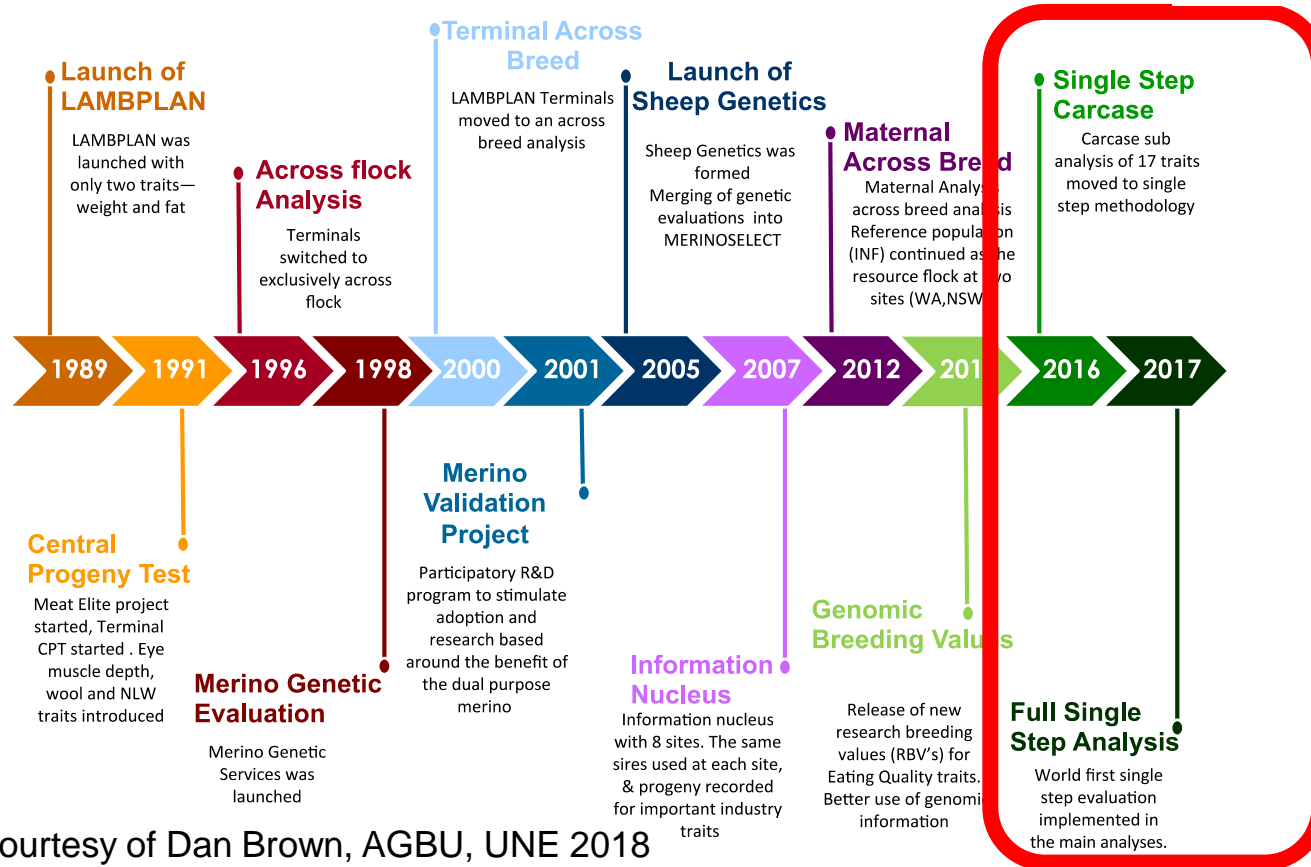
# More proof it works - France

## Single-Step GBLUP



Courtesy of J.M. Astruc, 2018

# Australia GS timeline



Courtesy of Dan Brown, AGBU, UNE 2018

# Reference population design

- Eight sites across Australia
- Mostly Merino ewe base
- 100 sires mated annually
- Comprehensive phenotyping of progeny
- 50K SNP genotypes  
+ (15K, HD, Seq)



Courtesy of D. Brown, AGBU, UNE 2018

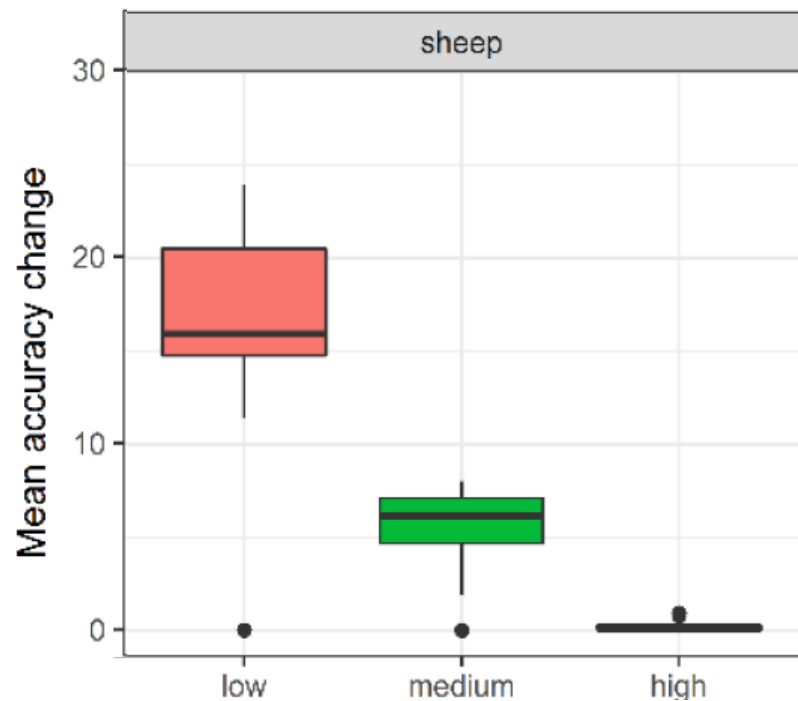
# The Genomic Reference Population

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Courtesy of Dan Brown, AGBU, UNE 2018

**Improvement in accuracy of using genomic selection vs ABLUP accuracies within bands of ABLUP accuracy from low (<30), medium (30 – 50) and high (>50)**



Australian EBV accuracies

Li et al 2018 AAABG



# Conclusions

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- Genomic selection is now operating in sheep and goat populations around the world
- Hard to measure / expensive traits ideal targets
  - Cheap & cheerful phenotypes – lots needed
- Key benefit is increase in accuracy
  - Reduces risk of selection
- Associated benefits of using genomic info
  - Healthier, higher welfare

# Acknowledgements



Ann McLaren  
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# SRUC

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