



Diversity of sheep in the UK – do we have a problem or an opportunity?

Dr Joanne Conington



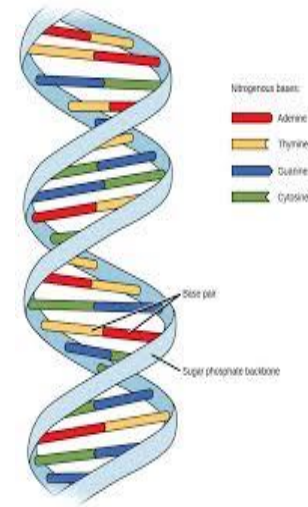
Scotland's Rural College (SRUC) Edinburgh, Scotland, UK

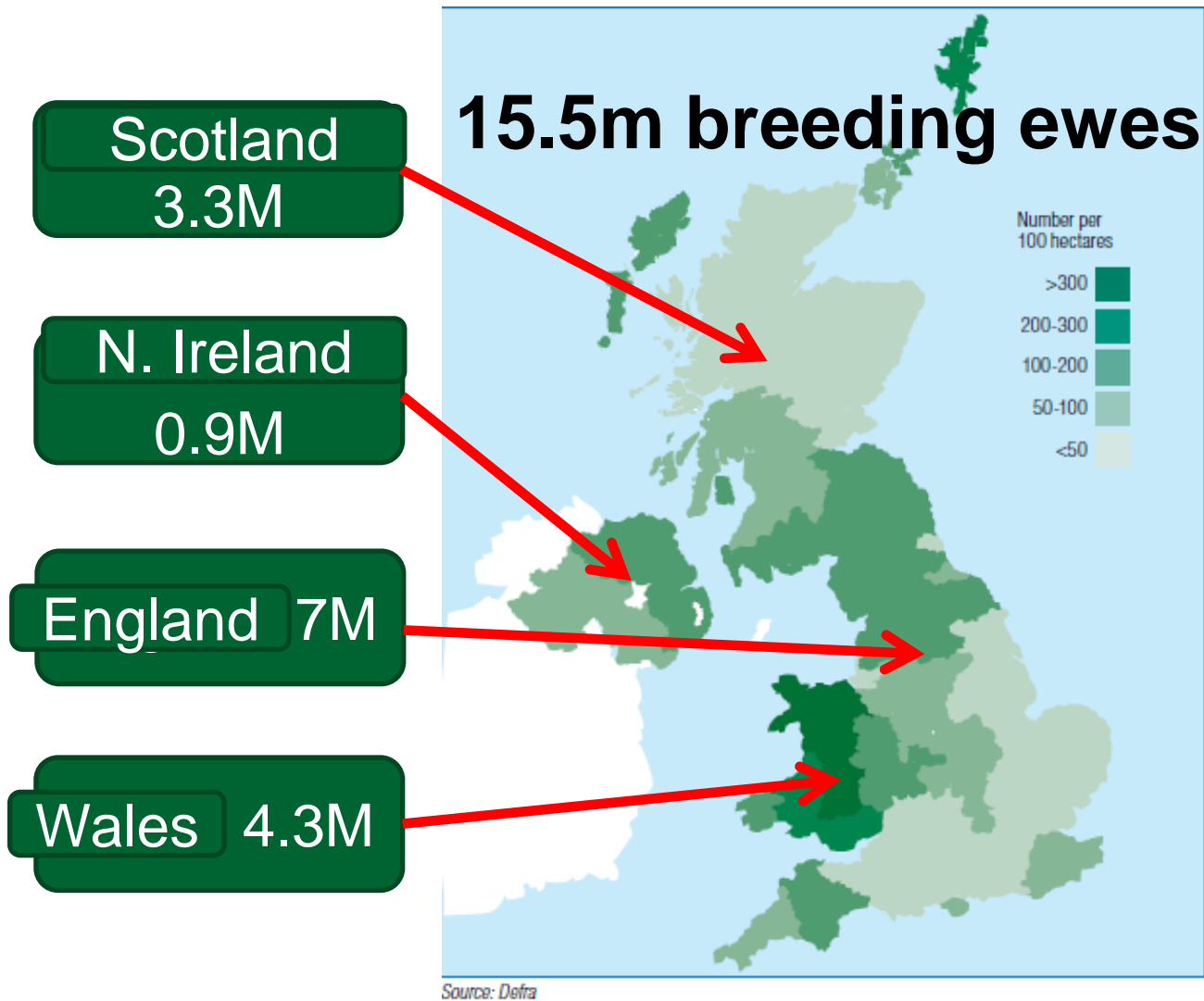
International Congress on the Breeding of Sheep and Goats, Bonn, Germany 15 October 2020

Leading the way in Agriculture and Rural Research, Education and Consulting

My talk

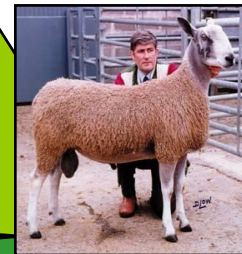
- UK sheep industry structure & phenotypic diversity
- Genetic diversity
- Genomic diversity





'Stratified' UK sheep industry

Hill and
Mountain



Uplands



Lowlands





Scottish Blackface





Welsh Mountain



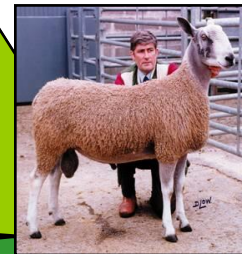


Swaledale



'Stratified' UK sheep industry

Hill and
Mountain



Uplands



Lowlands



Prolific ‘longwool’ breeds

Longwool Ram Breed Bluefaced Leicester



Longwool Ram Breed Border Leicester



F1 sheep
(cross between hill breeds and prolific breeds) are the most numerous breeding ewe in the UK

Crossbred Ewe Breed

e.g. Scotch Mule



(Bluefaced Leicester x Scottish Blackface)

Crossbred Ewe Breed

English mule / 'North Country mule'



Crossbred Ewe Breed

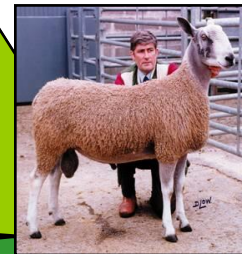
Welsh mule



Bluefaced Leicester x Welsh Mountain, Beulah or
Welsh Hill Speckled-face

'Stratified' UK sheep industry

Hill and
Mountain



Uplands



Lowlands



Meat sheep breeds

Suffolk



Texel



Charollais



Advantages of this structure



- Breeds and their crosses fit well to their location and system of production
- Use of specialised males and females – breed improvement occurs in purebred
- Crossing system maximises use of hybrid vigour

Disadvantages of this system

- Supermarket nightmare ! - Very diverse end product
 - Only ~50% of lambs meet EUROP (fat and conformation) spec
- Capture of the value of genetic improvement is complex
- Biosecurity compromised with purchase of animals from other farms / markets



UK sheep industry is very diverse



- 83 'Purebreeds'
- Many more crosses and composite 'breeds'

Only ~19 breeds have >10K

Native breeds

Jacob



Native breeds



Wensleydale



Dorset

Native breeds



Cute imports!

Valais Blacknose



Genetic variation – ewe & lamb traits

Core elements:-

Number of offspring /
year & survival
(low h^2) – but high σ_p^2

Rate of growth and
quality of carcass of
offspring
(Moderate h^2)



Variation in maternal behaviour



Within-breed estimates of heritability for maternal behaviour

0.13 Lambe, et al (2001). Animal Science **72**, p415-425

0.20 Brown et al (2015). Anim.Prod. Sci. 56:4 767-774

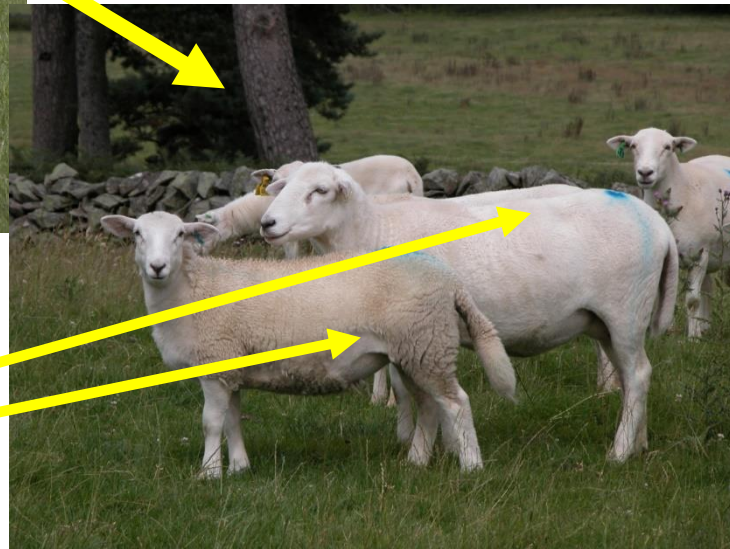
Wool shedding



Photos courtesy of Sandy Welsh

Shedding wool in Spring
(May)

August
Ewe shedding complete
Lamb shedding coat



Exploiting natural genetic variation

Genetic propensity to shed fleece



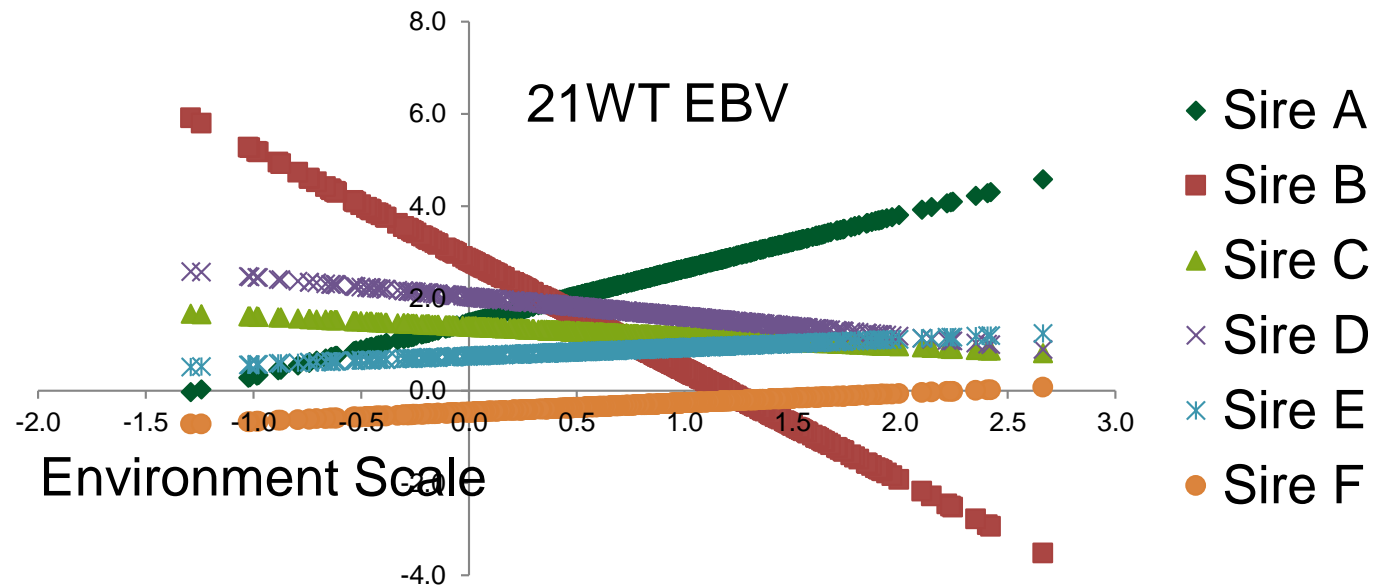
Heritability 0.65-0.8 Matika et al 2013. Anim.Genet 44(6):742-9;
0.54 Pollot, 2011 J Anim Sci 89:2316-25



Adapting to climate change



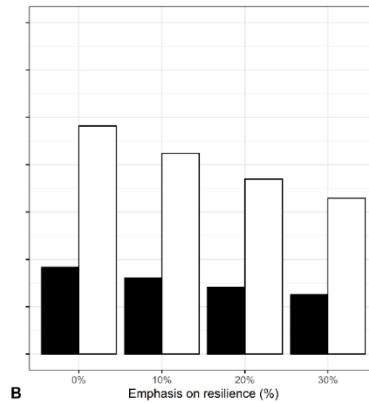
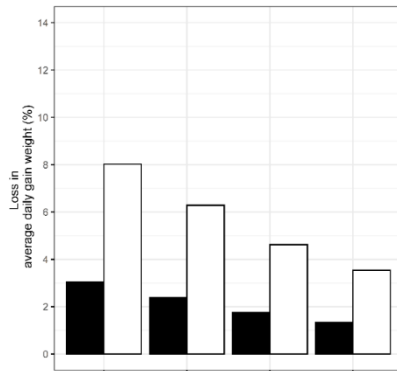
Sheep – evidence of G x E



- Scaling and re-ranking observed.
- ‘Robust’ sires, and sires suited to specific environments identified.

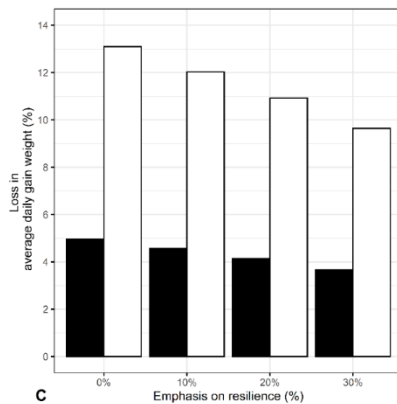
McLaren et al. 2015, Animal 9:10 1624-1634

Genetic variation in resilience to weather



1 deg = white bar

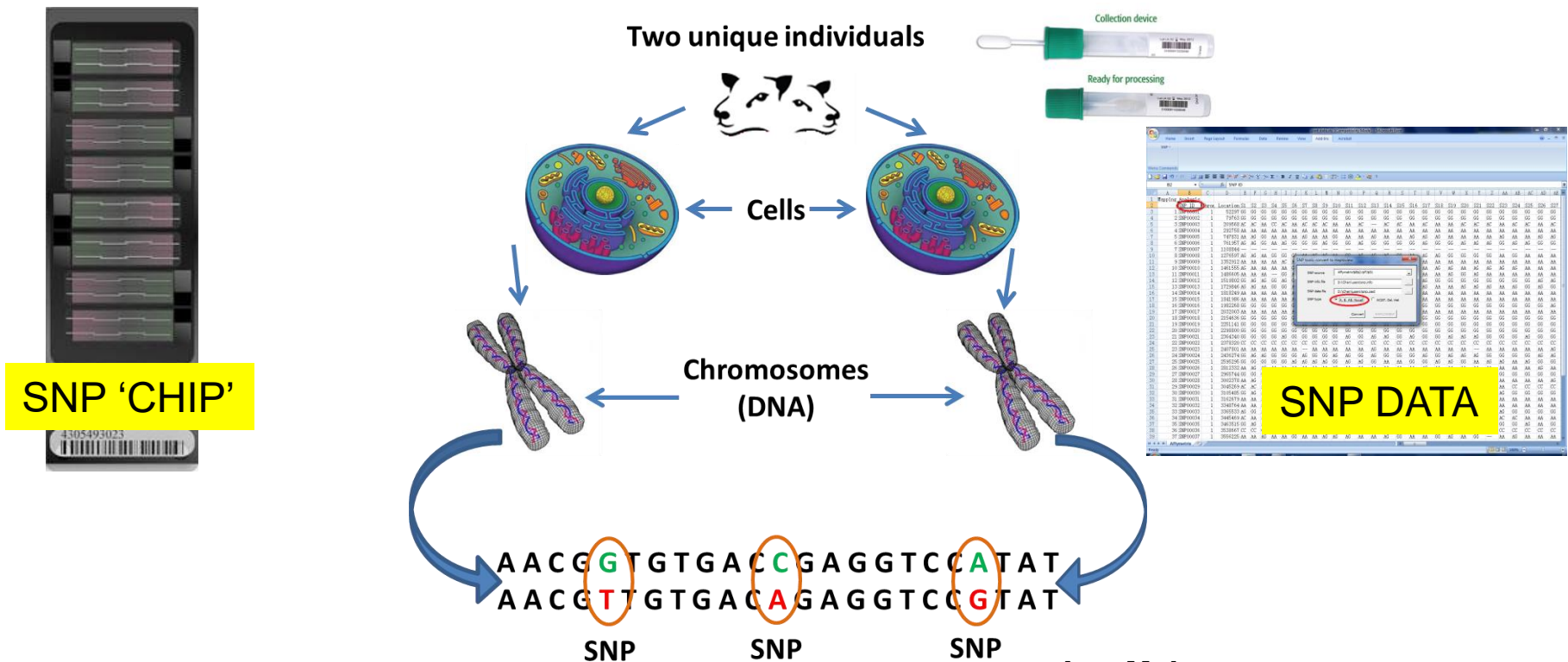
2.6 deg = black bar



Sanchez-Molano et al., 2020. In press, BMC Genetics.

Breeding strategies for animal resilience to weather variation in meat sheep

Using genomic information



Ann McLaren

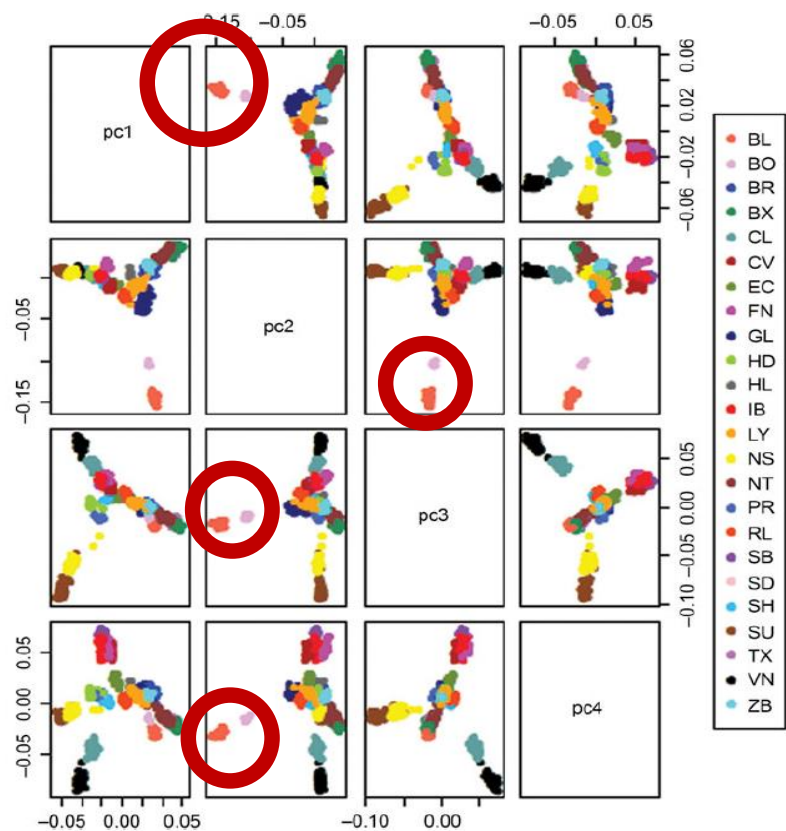
Knowledge of population structure & breed composition - SNP data



Important for:-

- Design of optimal (cross)breeding strategies
 - maximise non-additive genetic effects
-
- Maintaining flockbook integrity
 - authenticating animals being registered
- QA for the genotyping process – amongst other things..

Variation between populations



Bluefaced Leicester



Border Leicester



O'Brien et al 2020 *Animal* 14:3, pp 464–474

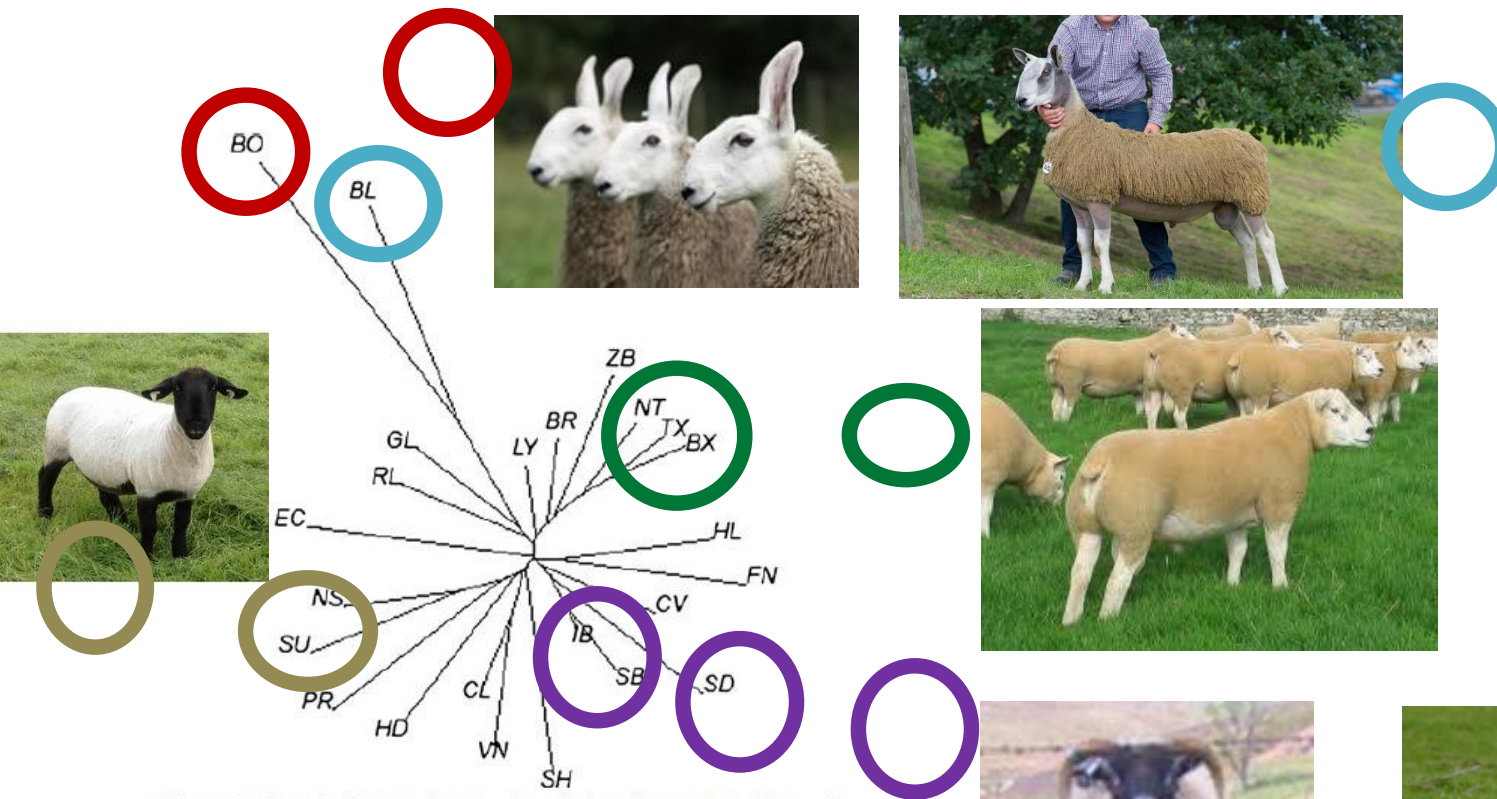
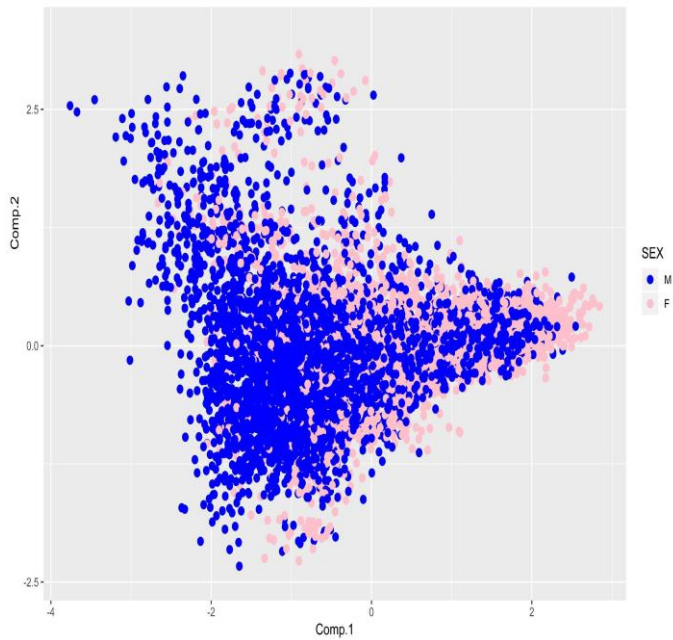


Figure 3 Genetic distance between breeds based on pairwise F_{st} estimates. Breeds included in the analysis were Bluefaced Leicester (BL), Border Leicester (BO), Belclare (BR), Beltex (BX), Charollais (CL), Cheviot (CV), EasyCare (EC), Finn (FN), Galway (GL), Hampshire Down (HD), Highlander (HL), Irish Blackface (IB), Lleyn (LY), New Zealand Suffolk (NS), New Zealand Texel (NT), Primera (PR), Rouge de l'Ouest (RL), Scottish Blackface (SB), Swaledale (SD), Shropshire (SH), Suffolk (SU), Texel (TX), Vendeen (VN), and Zwartble (ZB).

O'Brien et al 2020 *Animal* 14:3, pp 464–474

Genomic selection - testing population structures




Texel PCA (UK)
Kaseja 2019



Variation within populations

Inbreeding measurement(s)



- Standard F statistic
 - % homozygosity
 - 'Van Raden method'
- 
- A green bracket is positioned to the right of the last two items in the list, grouping them together.

VanRaden 2007. Genomic measures of relationship and inbreeding. Interbull Bull. 37: 33-36.

VanRaden 2008. Efficient methods to compute genomic predictions. J. Dairy Sci. 91: 4414-4423

VanRaden et al. 2011. Genomic inbreeding and relationships among Holsteins, Jerseys, and Brown Swiss. J. Dairy Sci. 94: 5673-5682.

Inbreeding measurement – example of Scottish Blackface n=1801

50K SNP used

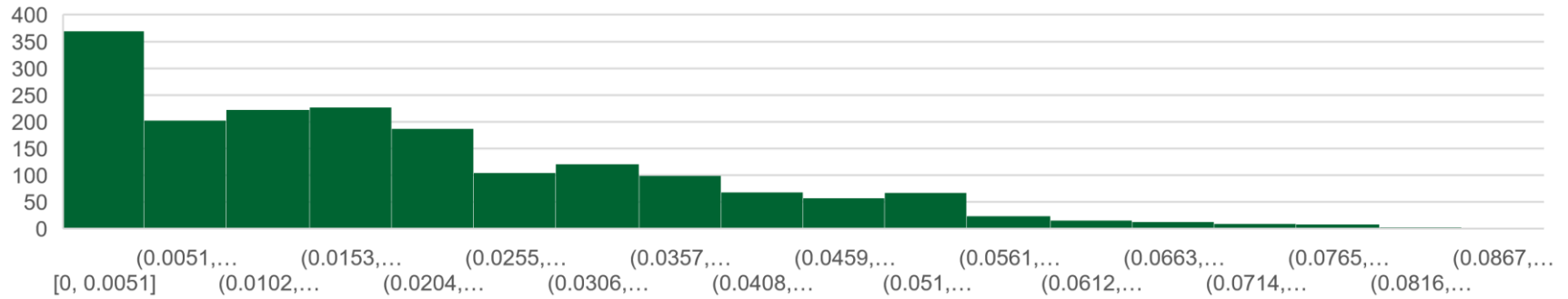
	F statistic	Homozy %	Van Raden
Average	0.02	0.64	0.07
Min-max	0 – 0.09	0.60 - 0.68	-0.03-0.18
Corr with ped	1	0.56	0.56



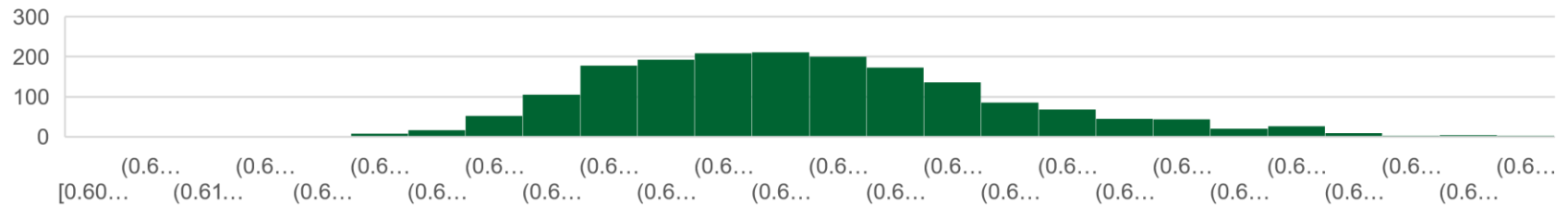
SBF= most numerous maternal purebred in UK

Banos 2020, pers. comm

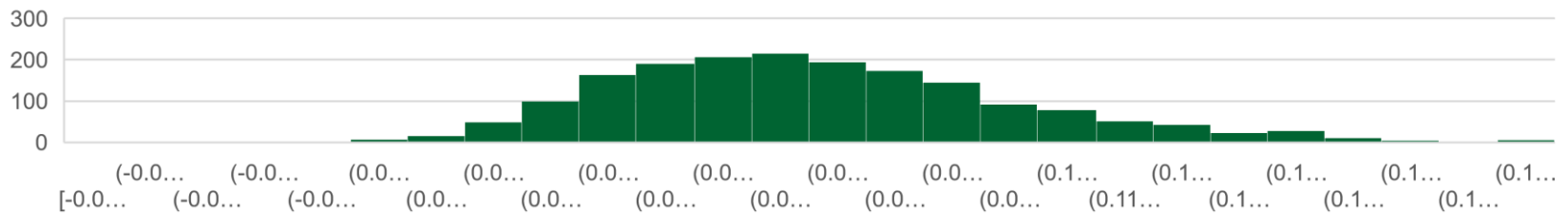
F stat



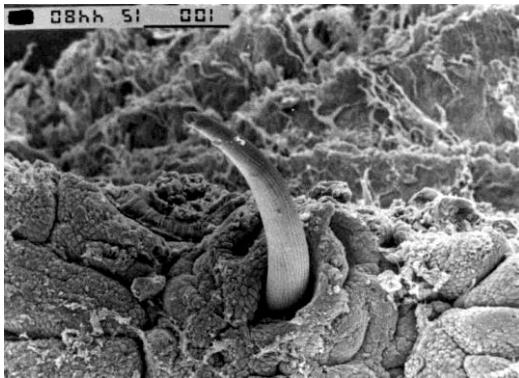
Homozygosity %



Van Raden



Hard to measure / expensive / measured later in life traits



Disease



Feed intake

(photo courtesy of H.Wishart)

Diversity of sheep in the UK – do we have a problem or an opportunity?



- Phenotypically and genetically diverse
 - Exploit diversity to meet new breeding goals – adapt to economic changes
- New methodologies to identify more robust animals
 - Resilience to changes in weather etc.
- Genomic tools – balancing genetic gain and inbreeding

.. UK sheep industry biggest problem...



Acknowledgements



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SRUC

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