



SMALL RuminanTs breeding for Efficiency and Resilience

WP6

Practical Selection Tools to Benefit from International Cooperation Low density SNP chip

Jean-Michel Astruc (IDELE) & Donagh BERRY (TEAGASC)



Final meeting Toledo - Tue. 23rd May 2023



Task 6.1: Harmonisation of phenotypes, genotypes and pedigree to facilitate international evaluations

- Overview on selection programs and genetic/genomic evaluation (survey) – 19 countries x breeds
- 10 sharing agreements for pooling data signed → MS23
- File format for exchanging data (international ID, pedigree, phenotypes, genotypes) → D6.1
- Exchanged genomic data and completed research on allele frequency across country x breeds – 18 breeds, 5 countries → Panel SNP on SMARTER website
Focus (Donagh Berry)
- Recommendations/guidelines for recording novel traits → D6.3
Focus (J.M. Astruc)

Aim and objective

- The aim was to in silico develop a low density genotype panel for sheep that would be as informative as possible to a range of different breeds and populations represented in SMARTER
 - Identify the SNPs that are segregating across breeds* jurisdiction
 - Develop lower density panels 1000, 2000, 3000, 6000, 9000, 12000

High imputation accuracy from informative low-to-medium density single nucleotide polymorphism genotypes is achievable in sheep¹

Aine C. O'Brien,^{†‡} Michelle M. Judge,[†] Sean Fair,[‡] and Donagh P. Berry^{†,2}

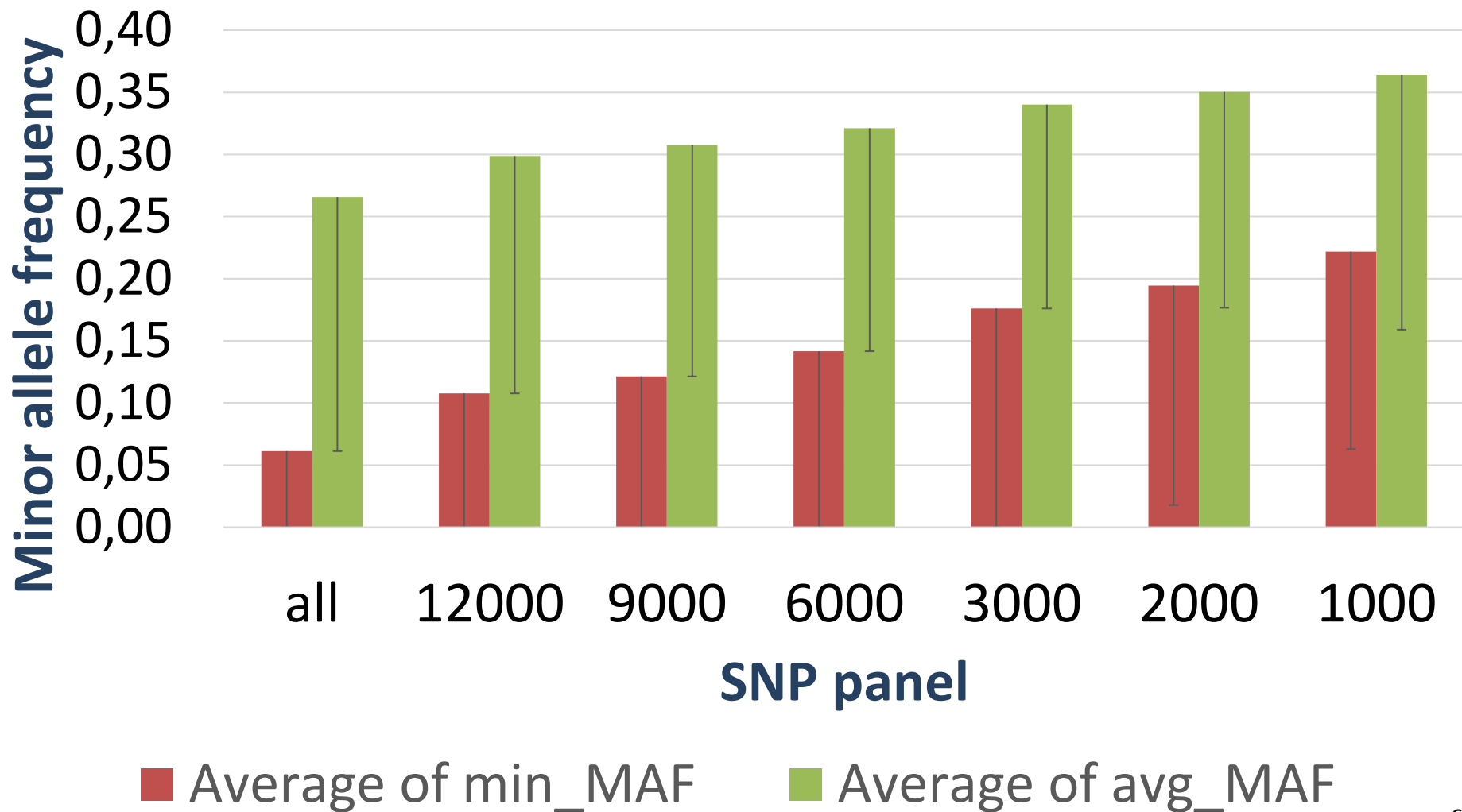
Data

- **Allele frequency per SNP**
- 5 meat sheep breeds from Ireland (44,040 SNPs)
 - Belclare, Charollais, Suffolk, Texel and Vendeen
- 2 meat sheep breeds from the UK (577,400 SNPs)
 - Scottish Blackface and Texel,
- 5 French dairy sheep breeds (48,059 SNPs)
 - Basco-Béarnaise, Black-faced Manech, Corse, Lacaune, and Red-faced Manech
- **38,883 SNPs in common**

Approach

- Average and minimum MAF per SNPs
- Genome broken into N blocks
- SNPs with greatest minimum MAF followed by greatest average MAF were chosen per block

Min and avg MAF across breeds



Conclusions

- Common SNP panels are possible with SNPs segregating in most populations

SMARTER PARTNERS



Thank you for your attention

www.smarterproject.eu