



HAL
open science

Modelisation of genomic selection in French dairy goats

Céline Carillier, Helene H. Larroque, Christèle Robert-Granié

► **To cite this version:**

Céline Carillier, Helene H. Larroque, Christèle Robert-Granié. Modelisation of genomic selection in French dairy goats. 16. Séminaire des thésards du Département de Génétique Animale, Apr 2013, Amboise, France. 2013. hal-02805824

HAL Id: hal-02805824

<https://hal.inrae.fr/hal-02805824>

Submitted on 6 Jun 2020

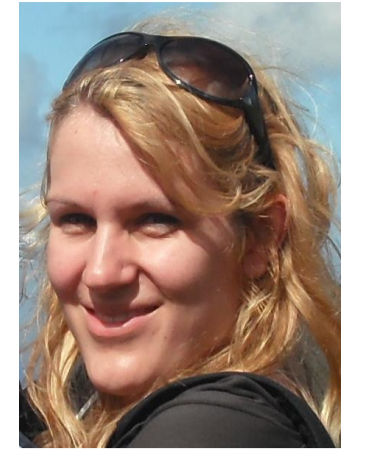
HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

Modelisation of genomic selection in French dairy goats

Céline Carillier, Hélène Larroque, Christèle Robert-Granié

INRA-SAGA, UR631, 24 Chemin de Borde Rouge, Auzeville, CS 52627, 31326 Castanet-Tolosan Cedex



Introduction

- Illumina 50K goat bead chip in 2011
- 2 **multi-breed** (Alpine and Saanen) populations with genotypes:
 - 1,985 **females** from commercial flocks born between 2008 and 2009 and their 20 sires
 - 657 artificial insemination **bucks** born between 1993 and 2009 and 148 young males born between 2010 and 2011
- Small generation interval (less than 4 years in sire-daughter pathway)
- High value of accuracy of young bucks at birth

Objectives and strategy

1

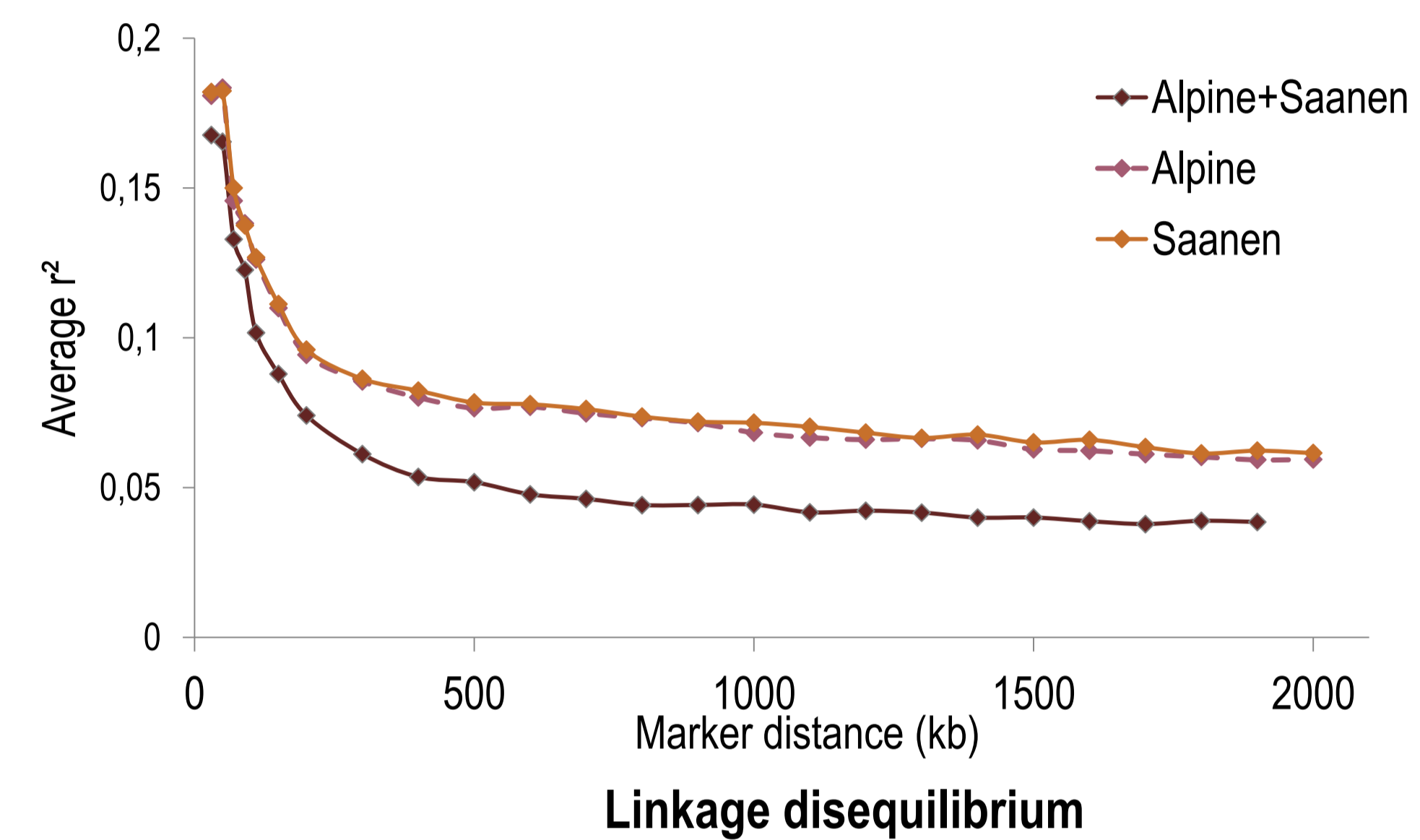
Characterization of reference population structure

Specificity: **Males and females**

Multi-breed

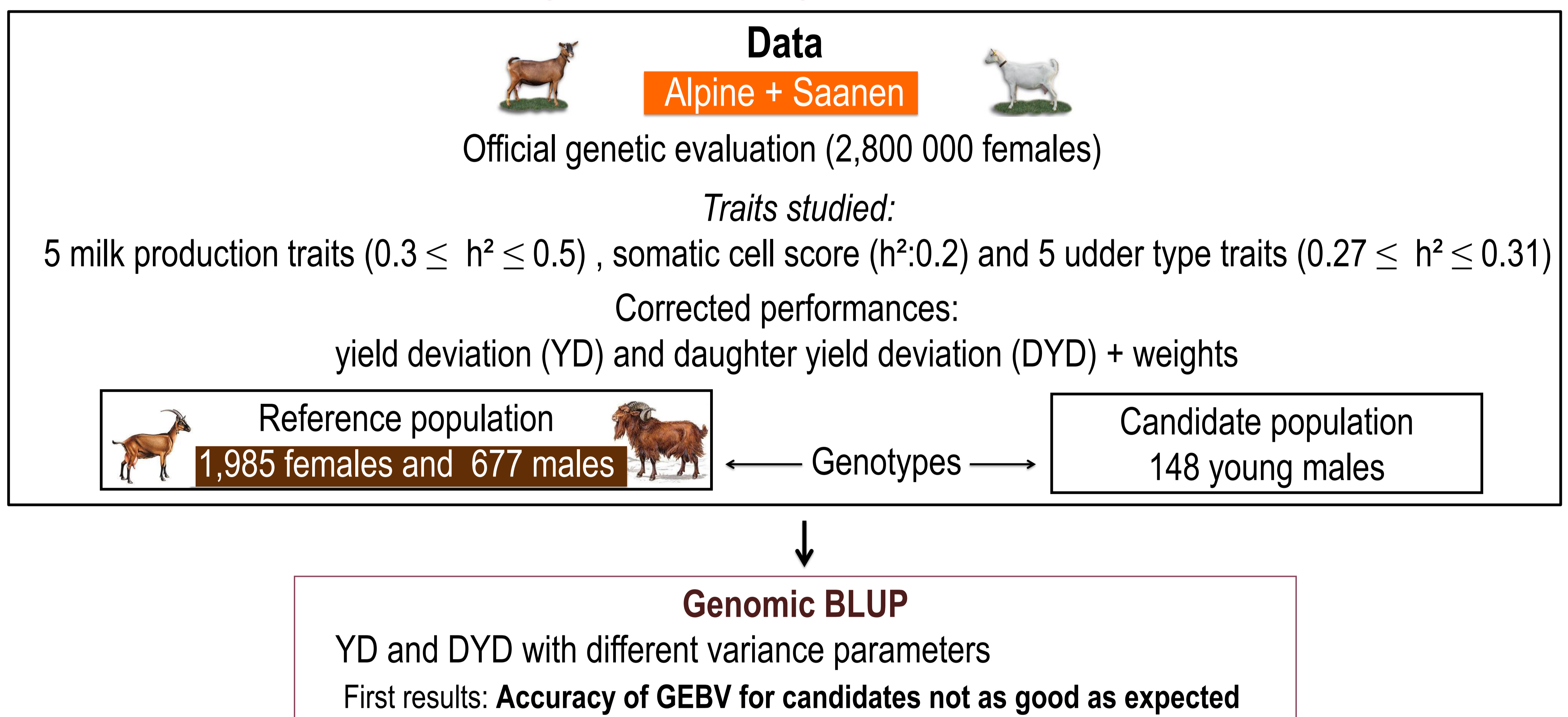
- Linkage disequilibrium
- Relationship (pedigree, genomic)

→ First results: **Not optimal for genomic selection**

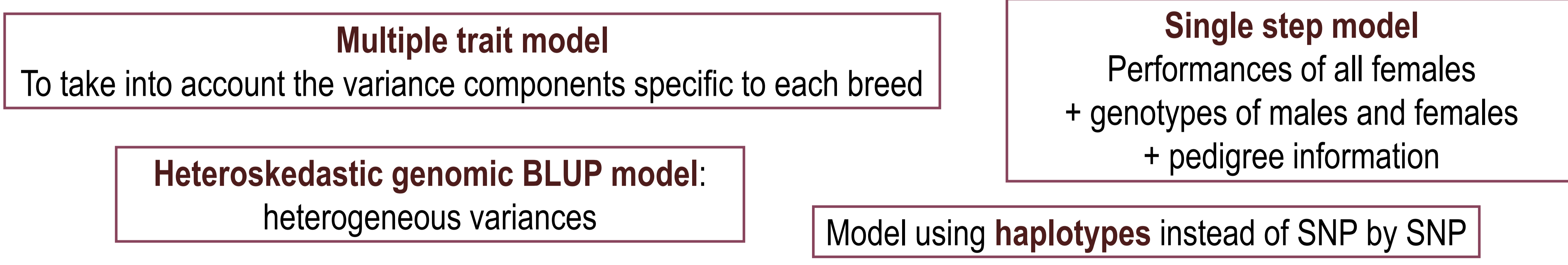


2

Working on models for genomic evaluation



To be done



3

Advices and strategies for future

Identify the best suited genomic model, which animals should be genotyped, SNP with largest effects



Funding

Midi Pyrénées region
INRA with selGen program

Background

Master degree in animal breeding and genetics (AgroParis Tech)
Master degree in agronomic engineering (AgroSup Dijon)

■ 16^e Séminaire des thésards du Département de Génétique Animale ■ Amboise, 2 & 3 avril 2013 ■

