

LEAP-Agri: OPTIBOV-project

Genetic characterization of cattle populations for optimized performance in African ecosystems

13-09-2018, Dr. Richard Crooijmans (Richard.Crooijmans@wur.nl)



Why OPTIBOV

- Maintain traditional cattle breeds
- Capture genetic and genomic variation
- Use strength of these breed (adaptation)
- Use these breeds to find selective sweeps related to adaptation
- Use known variation in production to improve production (marker assistant selection)
- Train, educate and involve people to perform optimal breeding (longterm investment) (workshops, app, website)

Aim of the project: Adaptation!

- Improve production and survival of traditional breeds adapted to their local environment
 - Select local breeds
 - Collect phenotypes
 - Collect DNA
 - Estimate the amount of inbreeding
 - Find selection signals on the genome
 - Find genes and variant on the genome
 - How to improve adaptation and production

Partners (PI)

■ Europe:

- Prof Dr. J. Kantanen (FI:LUKE)
- Dr. C. Ginja (PT:CIBIO/ICETA)
- Dr. R. Crooijmans (NL: WUR) project leader

■ Africa:

- Dr. D. Kuganza (UG:MU-CAES)
- Dr. M. Makgahlela (ZA:ARC)
- Dr N. Ghanem (EG:CU)

Breeds

Eastern, northern and western Finn cattle

Groninger Whiteheaded Dutch belted, Tauros

Mirandesa, Barrosã and Mertolenga.

Menofi (Baladi) / Domiaty (Damietta) / Maryuti

Ankole, East African Zebu

Nguni, Drakenberger

EU: North
Finland

EU: mid
The Netherlands

EU: south
Portugal

Africa: North
Egypt

Africa: Mid
Uganda

Africa: South
South Africa



Potential traits:

-Climate, amount of rain, ecosystem

- Temperature-humidity index

Availability of water\food

- Food quality

- Amount of food available

- Type of food

- Disease recording

- Disease resistance

- Parasites infections (ticks, nematodes)

- Housing (outdoor-indoor)

- Production records

- Calving interval/total number of calves

- Age of first calf

- Treatments

- Biochemical measurements on blood

- Immune parameters (Ig)

- MHC haplotypes



WAGENINGEN
UNIVERSITY & RESEARCH



100years
1918 — 2018

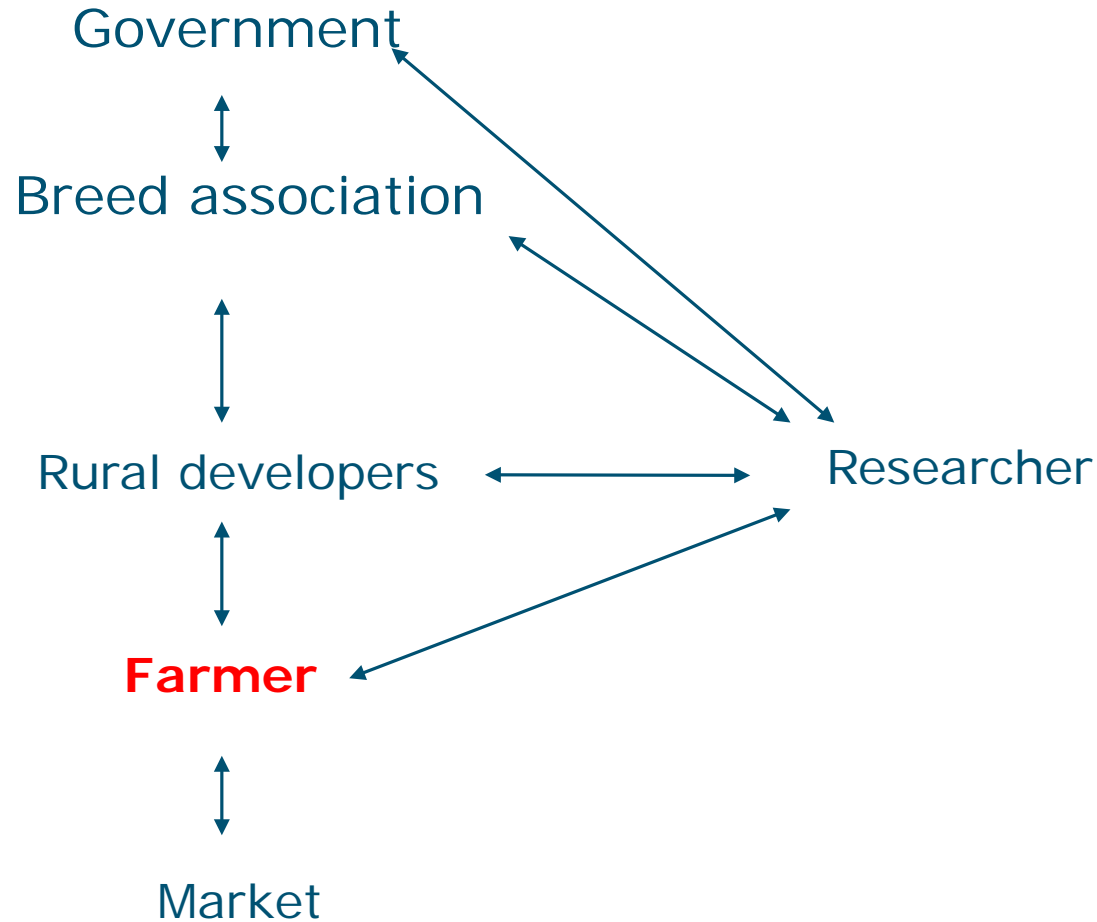
breeds



Recording phenotype

- Basis is the FAO list but expanded with new knowledge
- Record what is going wrong
- Easy access
 - App (info/pictures)
 - Website
 - Database

Information/data flow



Outcomes OPTIBOV

- Markers detected for selective adaptation traits
- Phenotype list
- African cattle breed variants
- Traditional/local breed characterisation
- New African SNP array
- Recording system (app, database, website)
- Training schools (phenotype and genotype analysis)

Thanks to

- LEAP Agri
- Local financiers

