LEAP-Agri: OPTIBOV-project

Genetic characterization of cattle populations for optimized performance in African ecosystems

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Outline

- What is LEAP-Agri
- OPTIBOV- project
- Nagoya protocol





LEAP-Agri

LEAP-Agri: Long-term EU-Africa research and innovation Partnership on food and nutrition security and sustainable Agriculture

LEAP-Agri operates under the EU-Africa High Level Policy Dialogue on science, technology and innovation, which includes the implementation of the jointly funded EU-Africa Research and Innovation Partnership focusing on **food and nutrition security and sustainable agriculture** (endorsed by the EU-Africa Summit 2014).



This project has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 727715









30 partners in **9 African** and **10 European** countries: research funding institutions (like NWO and LNV in NL) and some research organisations (like WUR)

 Common call for proposals
National funding agencies pool their resources funds are provided and administrated in the respective country
Different national requirements for the funding -> for NL: involvement of (preferably) Dutch private sector business & close cooperation with African Union partner
✓ EC tops-up the national funds by variable amounts





LEAP-Agri funding –focus NL

Partner contributions: 18.5 million Euro EC top-up of the budget: 9.25 million Euro Total of 27.75 million Euro for three years of research projects approved

Funding details NL: •LNV and NWO: 1.5 million Euro TOTAL FUNDS Share of contributions 18.5 million Euro France Germany The Netherland Kenya Turkey Uganda Finland Ghana International Egypt Belgium Algeria South Africa Norway Cameroun Spain Portugal Senegal Burkina Faso15





Projects approved –start Sept 2018

Total number of LEAP-Agri projects approved: **27** (120 pre-proposals of which 87 full proposals submitted)

of which

14 projects with NL partners 8 projects with WUR as partner

OP IBOV

Genetic characterization of cattle populations for optimized performance in African ecosystems⁶





Partners (PI) within OPTIBOV



Europe:

- Prof Dr. J. Kantanen (LUKE)
- Dr. C. Ginja (PT)
- Dr. R. Crooijmans (NL) project leader

Africa:

- Dr. D. Kuganza (MU-CAES)
- Dr. M. Makgahlela (ARC)
- Dr N. Ghanem (CU)





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
 - Ecosystem (past- present- future): modelling
 - 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







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, nematotes, mosquito's)
- 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
- Milk components

Local breeds









ecosystems





equator

Adaptation started in the past

Modelling: climate and climate change; availability of water and food; diseases (transmitted by insects like ticks);





Recording phenotype (1)



- Basis is the FAO list but expanded with new knowledge
- Well defined phenotyping protocol
 - Blood collection
 - Whole blood, EDTA/Heparin blood, blood in RNA later
 - Milk collection
 - Hair collection
 - Feces collection
 - Mouth swaps





Recording phenotype (2)



Records of abbreviations of normal status

- In time
- Easy access
 - App (info/pictures)
 - Website
 - Database







Recording genotypes



WGS > 720 animals

- 10x coverage
- Illumina based
- Per breed at least 10 males and 10 females

New designed African SNP array (50K)

- Taurine and Indicine
- Traditional breed info





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 (Taurine-Indicine)
- Recording/communication systems (app, database, website)
- Training schools (phenotype and genotype analysis)





Nagoya protocol



What:

The Nagoya Protocol on Access to Genetic Resources and the Fair and Equitable Sharing of Benefits Arising from their Utilization (ABS) to the Convention on Biological Diversity is a supplementary agreement to the Convention on Biological Diversity.

Why:

The Nagoya Protocol will create greater legal certainty and transparency for both providers and users of genetic resources by:

- Establishing more predictable conditions for access to genetic resources.
- Helping to ensure benefit-sharing when genetic resources leave the country providing the genetic resources

By helping to ensure benefit-sharing, the Nagoya Protocol creates incentives to conserve and sustainably use genetic resources, and therefore enhances the contribution of biodiversity to development and human well-being.





Nagoya protocol: countries









Access and benefit sharing



Nagoya guidelines

- MAT agreement over 6 countries
- student started working on this topic
- Local guidelines importing animal material
 - National food security services/ national animal health services





Thanks





A Long term EU-Africa research and innovation Partnership on food and nutrition security and sustainable AGRIculture





