

# Accelerate genetic progress to reduce methane emission – **Africa working group**

Chairs: Yvette de Haas & Raphael Mrode, May 23, 2025



# Aim working groups

- Introduce Global Methane Genetics & running projects
- Bring people together and build network
- Share knowledge, experience, hurdles & issues
- Collaborate on key topics
  - E.g. Smaller groups, task force
- 2 meetings a year

# Agenda

- Welcome & Aim
- Global Methane Genetics initiative (Birgit)
- ICAR Feed&Gas guidelines & wiki (Birgit)
- GMG Africa project (Raphael)
- Any other project to mention in Africa?
- Discussion:
  - Gaps & Needs, ideas, opportunities
- AOB

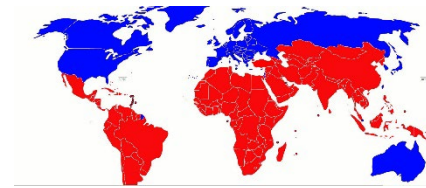
# Global Methane Genetics (GMG)

Accelerating Genetic Progress to reduce  
methane in ruminants



Coordinators: Roel Veerkamp & Birgit Gredler-Grandl (WUR)

# GMG: Why? How? What?



- Genetic progress can make a **permanent** and **impressive contribution** to reducing methane output from livestock systems **globally**
- We aim to accelerate genetic progress and to implement breeding strategies for reduced methane emissions in ruminants in the **global North and South** by supporting
  - **sharing of protocols and data,**
  - **expanding phenotyping,** breeding program design
  - **genetic evaluations**
  - development of **Global Livestock Genetics and Genomics Programs**

# Global Methane Genetics initiative (GMG)

Accelerate genetic progress for low CH<sub>4</sub> emitting ruminants

**Investment of 27M US\$**

**25 countries, 50 partners, 25 breeds**

**Methane pheno- & genotypes ~110k cattle & sheep, ~20k microbiome**

## **Dairy:**

Holstein (~42k)  
Jersey (~8k)  
(Nordic) Red Breeds  
(~7.3k)  
Brown Swiss (~3.3k)

World-wide sharing  
Develop protocols  
Phenotyping for  
reference populations  
Genetic evaluation  
Impact of genetics

## **Africa**

Dairy & crossbreeds (~1.5k)

## **South America**

Beef & indigenous (~7k)

## **Microbiome:**

World-wide reference  
population  
(~20k samples)

## **Beef:**

North America (~6k)  
Australia, Ireland,  
UK, NZ (~18.5k)

## **Sheep: world-wide reference population**

Australia & New Zealand  
UK & Ireland  
Uruguay (~17k)

# Global Methane Genetics (GMG) initiative



# Key areas – current activities

- Projects get started
- Share data – data base for external and GMG internal data sharing
- Fair share principle
- SOP for recording & harmonisation of data pipelines across partners
- Build facility to test methane recordings of many individual cows





THE GLOBAL STANDARD  
FOR LIVESTOCK DATA

Network. Guidelines. Certification.

# ICAR Feed&Gas working group

## ICAR Sheep, Goats & Camelids

### ICAR Wiki-page

**Birgit Gredler-Grandl & Chantal van Gemert**

**Wageningen University & Research**

- Group Objectives


- Update, promote and extend guidelines for phenotype recording – methane and feed intake in ruminants
- Guidelines for full range of traits in sheep and goats
- Harmonisation & standardisation
- Conduct surveys, reports for recording schemes
- Provide a forum and foster knowledge exchange
- Facilitate and coordinate international collaboration

<https://www.icar.org/group/working-group-sheep-goats-and-camelids/>

<https://www.icar.org/group/working-group-feed-and-gas/>

- [ICAR Wiki](https://wiki.icar.org/index.php/Guidelines): <https://wiki.icar.org/index.php/Guidelines>
- Section 20: Methane emissions for genetic evaluations

ICAR Wiki



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Section 20 – Methane Emission for Genetic Evaluation

 Page

 Discussion

 View source

 History

This is the approved revision of this page; it is not the most recent. [View the most recent revision.](#)

**NOTE: This version of Section 20 has been approved by the working group's Chair. Please be aware that further revisions may occur before final review and approval by the Board and ICAR members per the [Approval of Page Process](#).**

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- ICAR Wiki: <https://wiki.icar.org/index.php/Guidelines>
- Section 20: Methane emissions for genetic evaluations

## 4 Sub-sections

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Definition and Terminology

Methane determining factors

Methane measurements methods

Sniffer SOP  
GreenFeed SOP  
PAC  
LMD

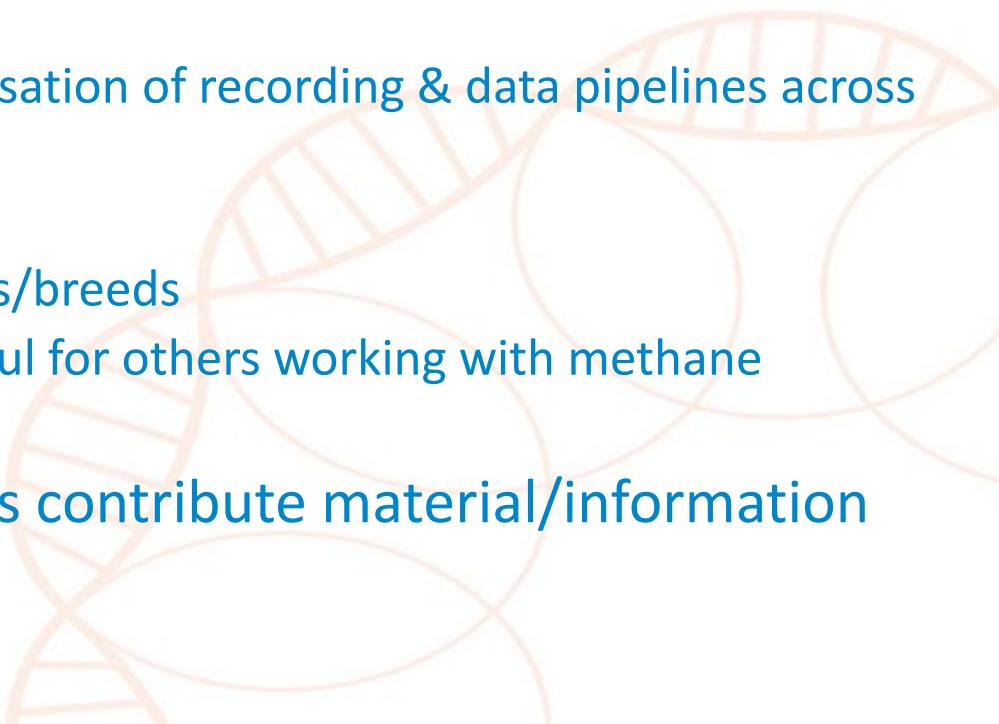
Proxies

Proxies discussion

Merging and sharing data in genetic evaluations

Ongoing activities


Projects  
Initiatives  
Course material  
...

- GMG projects/partners valuable source of existing & newly developed information
    - Recording devices
    - Protocols & SOPs – harmonisation of recording & data pipelines across countries
    - Do's and don'ts
    - Differences between species/breeds
    - Anything that might be useful for others working with methane emissions
  - Expect GMG partners & others contribute material/information
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- **Feedback is welcome! Start a discussion!**
  - If you have information you would like to add to a specific page
  - If you notice information that is wrong, no longer up to date or incomplete
- Click on the discussion button at the top of the page

## Section 20 – Methane Emission for Genetic Evaluation

 Page

 Discussion

- Fill in a subject and description and add the topic
  - The writers of the pages and everyone else can comment on these topics
- Would like to add information, feel free to open a discussion on the main page



*better lives through livestock*

# Global Methane Genetics Initiative: Accelerating reduced emissions in indigenous breeds in Africa

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*R. Mrode -- East Africa Team : ILRI, Nairobi, Kenya*

*West Africa Team: Habibou ASSOUMA (CIRAD) and Luc Hippolyte Dossa (UAC)*

*South Africa Team: Giel Scholtz (ARC)*



# Dimensions of the project

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- Project implemented in three regions of Africa
- West Africa:
  - Mohamed Habibou ASSOUMA (CIRAD) and Luc Hippolyte Dossa (UAC)
  - CIRAD, CIDES, Université d'Abomey-calavi (UAC)
  - Burkaso Faso and Benin
- South Africa:
  - Giel Scholtz
  - Agricultural Research Council, South Africa (ARC),
- East Africa: Kenya and Ethiopia
  - ILRI - Raphael Mrode





# Four Work packages

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- 1: Phenotyping for CH<sub>4</sub>, production traits and genotyping of animals
- 2: Microbiome profiling, fecal sampling and sequencing of microbiome.
- 3: Design genetic models for joint data analysis dairy cattle data from different regions of Africa; analysis of microbiome sequence data and mapping to CH<sub>4</sub> and productive traits.
- 4: Working with government national breeding centers and private breeding organizations to feed outcomes into existing breeding programs for improved productivity and efficiency



# WP1: Phenotyping

- West Africa
  - Two experimental stations
  - UAC (Benin) : 20 cows measured from 4 breeds - Girolando (exotic), White Fulani, Goudali, Borgou (LMD and Greenfeed)
  - CIRDES ( Burkina Faso) : 20 cows - Sudanese Zebu Peul and Azawak (LMD and GreenFeed)
  - Benin - 75 cows in government farm - LMD
- South Africa
  - 20 bull calves each, from the Afrikaner, Bonsmara and Nguni beef breeds over 3 years (LMD and GreenFeed)





# WP1: Phenotyping

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- East Africa
  - Kenya and Ethiopia
  - 1800 cows measured in smallholder farms in each country over 3 years (LMD)
  - 180 cows in medium /large farms in each country ( LMD and Greenfeed)



## WP2: Microbiome profiling, fecal sampling and sequencing of microbiome

- Microbiome sampling will be collected on 1000 tropical cows using the oral stomach tubing method
- West Africa : 40
- South Africa : 180
- East Africa : 680
- Fecal samples also available for the East Africa samples



## Update on Status

- Order placed with C-Lock for 2 Greenfeed units with Grid systems
- Order placed for 6 units of LMD units
- CRA with ARC, South Africa is being processed
- CRA with West Africa initiated
- Two farms for first location of Greenfeed units identified
- Work plans are being drafted- Ethiopia and Kenya
- Ethical approval (IREC and IACUC) at ILRI commencing
- Registered for overall launch of project in Innsbruck



## Some concerns

- Microbiome sampling will smallholder farms could be a challenge
  - May need some incentive
  - Permission from Veterinary Department
- Limited experience with utilizing GreenFeed







Thank you for  
your attention