

RAGE

Rabies Accessible Genomic Epidemiology

TRAINING WORKSHOP

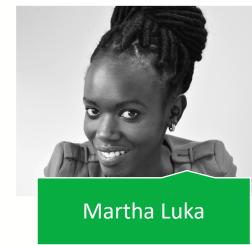
> NCDC, Abuja 12-16th Feb 2024



Who are we?















Workshop etiquette

- Treat each other with respect
- General lab rules apply
- Turn off your cellphones
- For demonstrator attention: raise a hand
- Participants can ask questions at any time following a raised hand
- Everyone may take a moment to step out of the room discreetly without explanation at any time

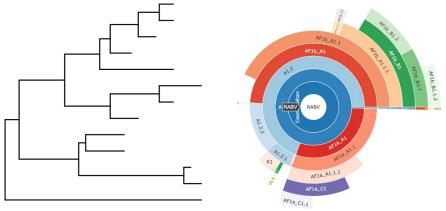


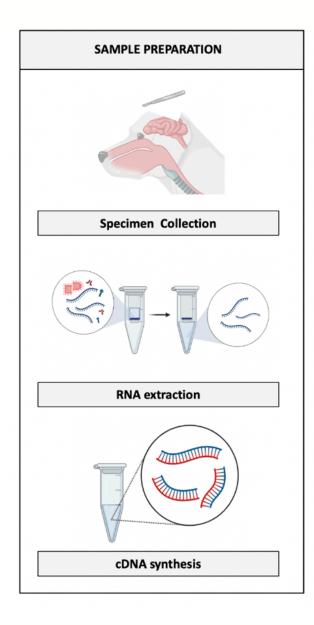


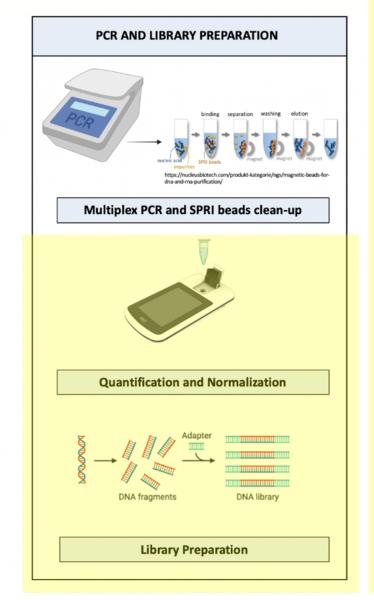
Workshop content

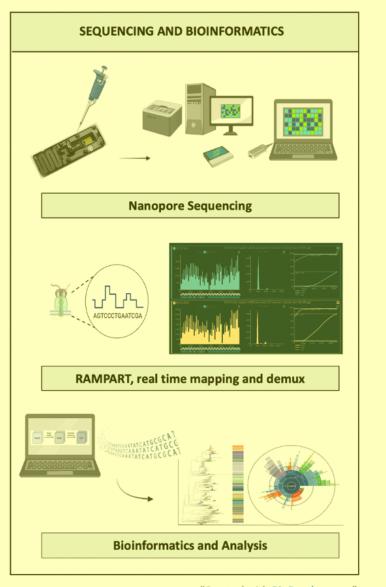
- An introduction to genomic surveillance for rabies virus
 - Wet lab
 - Sequencing library preparation
 - Nanopore sequencing
 - Monitoring the run
 - Dry lab
 - Processing raw data
 - Building trees
 - Visualising & interpreting your data











"Created with BioRender.com."

Bautista et al., JoVE

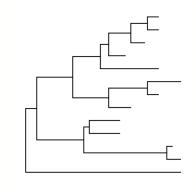


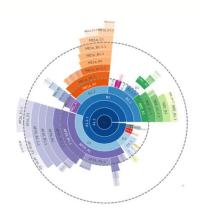
Schedule

MONDAY	TUESDAY	WEDNESDAY	THURSDAY	FRIDAY
OPENING WET LAB STARTER	WET LAB	WET LAB/DRY LAB	DRY LAB	DRY LAB
Background & lab intro	Masterclasses & Nanopore library prep I	Nanopore library prep II & sequencing Intro to Bioinformatics	Bioinformatics pipeline Alignments & Phylogenies	Advanced analysis, interpretation & visualisation











Intended learning outcomes

- An understanding of an end-to-end workflow for sequencing rabies virus
- Background theoretical knowledge
- Mastering key techniques
- How to apply "light-touch" analytical tools





Who are you?



Quick poll

Join by Web PollEv.com/kirstynbrunker994





Icebreaker groups

- 1. Introduce yourself (name, position, where you are from)
- 2. What is your past experience (if any) with sequencing?
- 3. What are you hoping to gain from the workshop?