BIODIVERSITY — DAY 6			
Time	Activity Description	Intended Learning Outcomes	Relevance
		<i>After completion, trainees will (be able to):</i>	Why this is important for you as:
			Bioinformaticians should understand bacterial strain
			nomenclature from genomic data.
1400- 1530	Bacterial strain nomenclature and epidemiological tracking (Sylvain Brisse)	Learn about the cgMLST-based Life Identification numbers (LIN) codes concept Explore the practical applications of bacterial strain nomenclature and LIN code in epidemiological surveillance, outbreak investigation, and public health response.	Microbiologists should get familiar with LIN codes and how to accurately communicate strain information within the scientific community and collaborate effectively in epidemiological investigations. Epidemiologists should understand the LIN code example of bacterial strain nomenclature and how to interpret nomenclatural data to analyze transmission patterns.

## Details

## Bacterial strain nomenclature and epidemiological tracking

This course will provide participants with insights into the cgMLST-based Life Identification numbers (LIN) codes concept, a novel approach to bacterial strain taxonomy. Participants will explore how bacterial strain nomenclature and LIN codes can be used in epidemiological surveillance, outbreak investigation, and public health response.