

Introduction to Bioinformatics course: IBT

Introduction Part 4 Why IBT?







Why IBT?



The Need Basic Bioinformatics training for Molecular Biologists

'need for basic bioinformatics training for individuals entering the discipline, or for those who need a basic foundational understanding of bioinformatics before moving on to more complex areas' – H3ABioNet Education & Training Working Group







Audience



- molecular biologists
- Little to no bioinformatics experience





Skills-based curriculum



Learning outcomes:

- measurable skills gained
- what the participant will learn

 Important for participants to be able to go out and perform the analyses on their own after the course -> web based applications







Modules and primary trainers



Resources and databases (16 hrs)



Shaun Aron, University of the Witwatersrand, South Africa



Nicola Mulder, University of Cape Town, South Africa

• **Linux** (12 hrs)



Amel Ghouila, Institute Pasteur de Tunis, Tunisia

Molecular evolution
 and Phylogenetics (16 hrs)



Jean-Baka Domelevo Entfellner, University of the Western Cape, South Africa

 Sequence alignment theory & applications (12 hrs)



Jonathan Kayondo, Uganda Virus Research Insitute, Uganda



Sonal Henson, International Centre of Insection Physiology and Ecology, Kenya

Multiple sequence alignment (8 hrs)



Ahmed Mansour Alzohairy, Zagazig University, Egypt

Genomics (16 hrs)



Colleen Saunders,
University of the
Western Cape, South Africa



Fatma Guerfali, Institute Pasteur de Tunis, Tunisia

H3ABioNet
Pan African Bioinformatics Network for H3Africa





Next



watch video labeled: Part 5



