Applying core competencies in Africa: the H3ABioNet experience

Nicola Mulder University of Cape Town





What is H3Africa?

 "The Human Heredity and Health in Africa (H3Africa) **Initiative** aims to facilitate a contemporary research approach to the study of genomics and environmental determinants of common diseases with the goal of improving the health of African populations. To accomplish this, the H3Africa Initiative aims to contribute to the development of the necessary expertise among African scientists, and to establish networks of African investigators."

www.h3africa.org





What is H3ABioNet?

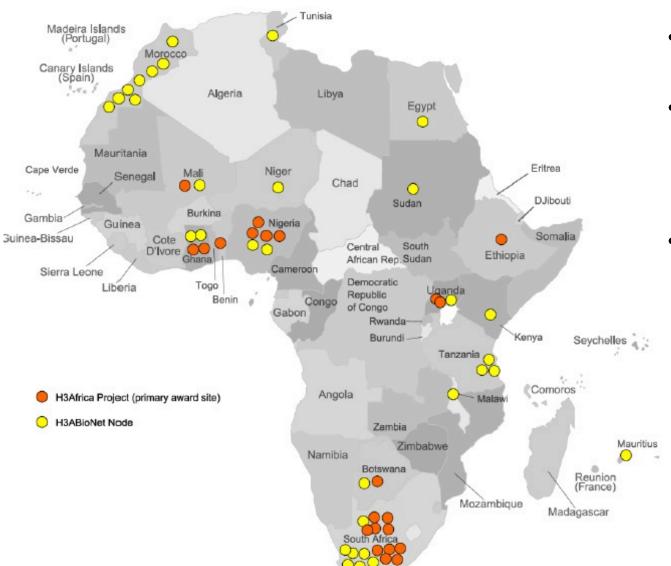
- H3ABioNet is a pan-African bioinformatics network that aims to develop sustainable African bioinformatics capacity (intellectual and physical infra-structure) to cope with the on going "omics" revolution and ensure African science is not left behind.
- The "omics" revolution for human populations is coming to Africa via a series of H3Africa funded projects.
- H3ABioNet has 32 institutions in 15 African countries, 1 in USA and 1 in UK, led by Computational Biology Group at the University of Cape Town.

www.h3abionet.org





H3ABioNet distribution in relation to H3Africa primary award sites



- 32 Nodes within Africa
- Nodes at different levels of bioinformatics expertise
- Geographically
 placed to help
 develop regional
 bioinformatics
 capacity for the
 Node and H3Africa
 project in the
 region



Need for MScs in Bioinformatics

- Only a handful of MSc in Bix outside SA
- Many institutions wishing to establish programs
- Held Bioinformatics curriculum development workshop in Botswana
- Established African Bioinformatics Education Committee (ABEC)





ABEC

- http://www.h3abionet.org/training-andeducation/african-bioinformatics-educationcommittee
- Curriculum development task force
- Set up website with guideline documents:
 - Considerations –key steps for starting a program
 - University processes for developing new programs
 - Existing programs and curricula
 - Curriculum development





H3ABioNet Bioinformatics curriculum

- Set of core and elective modules for a bioinformatics program using the ISCB guidelines have been defined
- Based on discussion, needs of individual institutions and existing curricula
- Trainers determined the content and contact hours for these modules
- Have included suggested lecturers from Africa
- http://training.h3abionet.org/ curriculum development wg/





H3ABioNet Bioinformatics curriculum

H3ABioNet Curriculum Development Taskforce

H3ABioNet taskforce to develop a curriculum of bioinformatics to be used throughout the continent

HOME BIOSTATISTICS I DATABASES I DATABASES II ETHICS **EVOLUTION AND PHYLOGENETICS**

GENOMICS AND COMPARATIVE GENOMICS HIGH-THROUGHPUT SEQUENCING INTRODUCTION TO LINUX AND SHELL SCRIPTING

INTRODUCTION TO PROGRAMMING FOR BIOLOGISTS MACHINE LEARNING METABOLOMICS MOLECULAR BIOLOGY FOR PROGRAMMERS

POPULATION GENETICS AND GWAS STATISTICS II PROGRAMMING I PROGRAMMING II PROTEOMICS SEQUENCE ANALYSIS

STRUCTURAL BIOINFORMATICS WRITING & PRESENTATION SKILLS WG MEETING MINUTES

H3ABioNet Curriculum Development Taskforce

In the wake of the Gaborone Workshop (H3ABioNet / University of Botswana, March 11th-12th 2014), H3ABioNet decided to build a taskforce to develop a standard curriculum in bioinformatics, at the MSc level.

This site is the repository for the documents produced by this Curriculum Development Task-Force (CDTF), and primarily for the module outlines as proposed by the CDTF members.

Please find here the different modules proposed and the list of volunteers for these. The menu on top of this page provides links to the different outlines.

The meeting minutes are to be found on this website, here (tab "WG Meeting Minutes" on top of



RECENT POSTS

H3ABioNet Curriculum Development Taskforce

RECENT COMMENTS

Benjamin Kumwenda on Databases I Amel Ghouila on Sequence Analysis





Example module page

Module outline by the H3ABioNet Curriculum Development Taskforce

Biostatistics I

2014-03-06 07:03:32 Jean-Baka Domelevo Entfellner

Prepared by: Jean-Baka Domelevo Entfeliner

Possible Lecturers: Jean-Baka Domelevo Entfellner, or any bioinformatician with a strong background in mathematics and statistics, ideally from his/her primary education.

Contact hours: For consistency reasons, each contact hour is fixed at 45min. Theory (23), Practicals (30)

SPECIFIC OUTCOMES ADDRESSED

- 1. Generally speaking: develop an understanding of stochastic experiments
- Understand and be able to build the framework of a statistical test

BACKGROUND KNOWLEDGE REQUIRED

Basic general-purpose scientific knowledge, basic arithmetic skills, and some familiarity with basic linear algebra.

BOOKS & OTHER SOURCES USED

- Fundamentals of Biostatistics, 7th edition, by Bernard Rosner (Cengage Learning, 2011)
- 2. Biostatistics with R An introduction to statistics through biological data, by Babak Shahbaba (Springer, 2012)

COURSE CONTENT

(A) Theory lectures

- I. Probability theory:
- Atomic and complex events, probabilities as a measure on sets



Example module page

Module outline by the H3AI Curriculum Development Ta

Biostatistics I

2014-03-06 07:03:32 Jean-Baka Domelevo Entfellner

Prepared by: Jean-Baka Domelevo Entfeliner

Possible Lecturers: Jean-Baka Domelevo Entfellner, or any with a strong background in mathematics and statistics, ide primary education.

Contact hours: For consistency reasons, each contact hour Theory (23), Practicals (30)

SPECIFIC OUTCOMES ADDRESSED

- Generally speaking: develop an understanding of stochas
- 2. Understand and be able to build the framework of a stat

BACKGROUND KNOWLEDGE REQUIRED

Basic general-purpose scientific knowledge, basic arithmet familiarity with basic linear algebra.

BOOKS & OTHER SOURCES USED

- Fundamentals of Biostatistics, 7th edition, by Bernard Ro Learning, 2011)
- Biostatistics with R An introduction to statistics through Babak Shahbaba (Springer, 2012)

COURSE CONTENT

(A) Theory lectures

- I. Probability theory:
- 1. Atomic and complex events, probabilities as a measure

- Enumerative combinatorics: counting permutations, combinations and partitions. Binomial coefficients.
- 4. Some common discrete probability distributions: Bernoulli, binomial, Poisson. Behaviour of a binomial when the number of trials tends to infinity. Concepts: probability mass, expectation of a discrete distribution.
- 5. First continuous probability distributions: uniform, exponential
- Central limit theorem and the normal distributions
- 7. Other continuous distributions: Student's t and chi-square distributions.
- II. Statistical hypothesis testing
- III. Analysis of variance and regression models
- IV. Multidimensional dataset analysis: Principal Component Analysis

(B) Practical component

We suggest the use of Rstudio throughout the course, as an integrated development environment to work with R. Being the fundamental statistical software in use across various research areas, it is essential that the students develop mastery over R during this course.

Alternatively, if computing resources are extremely scarce, use an interactive R interpreter to demonstrate the concepts, plus a simple text editor later on, once the students start writing functions.

This section "practical component" follows the same structure as the previous section "Theory lectures": practicals just aim at having the students manipulate the concepts seen in the lectures, right after they were introduced to them.

ASSESSMENT ACTIVITIES AND THEIR WEIGHTS

We would suggest two written exams during the course of the module (total weight = $\frac{1}{2}$), and a final programming exam (weight = $\frac{1}{2}$). Of course, practicals can also be for marks all along the module, but our advice is not to make each and every practical for marks, not to put too much counterproductive stress on the students. Practicals are the privileged moments when students actually understand the concepts as they put them into play.

Implementations

- First MSc in Bioinformatics program launched at the University of Bamako this year, lecturers local and distant
- New progam in Malawi about to be submitted for approval
- Other MSc in Bioinformatics programs such as in Kenya, Tunisia are being developed using the defined modules
- Using some as a base for Introduction to Bioinformatics course









Training evaluation framework



Centralized Evaluations

- Workshops
- Webinars
- Mentorship
- Internships
- Placements

Inputs &

Activities

Fellowships

How are the training activities impacting outputs & trainees' work?

Outcomes

How are trainees' careers changing?

Assess challenges, successes and needs

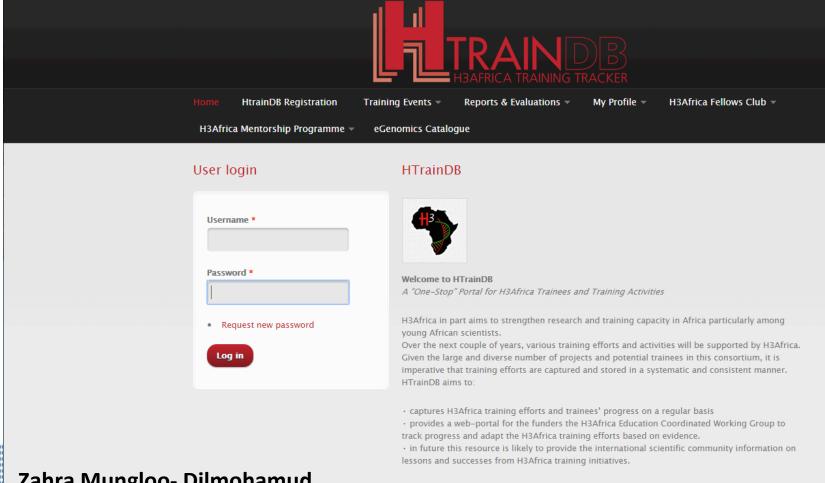
Evaluation results used to adjust training activities





Tracking Training and Education

 Developing a system to track, monitor and evaluate the training development of students and staff







Why HTrainDB?

- A major goal of H3Africa Consortium is to train the next generation of "genomics" scientists
- Ongoing training is widespread, un-trackable, and therefore not easy to monitor/evaluate
- Need to establish a "one-stop" center for training tracking, info & opportunities for trainees
- A tool to facilitate reflections & planning: where we are, where we have come from, plan next steps
- See Poster





First survey

- 498 trainees
- 10% response rate to the survey
 - 76% participated in H3Africa funded training
 - 94% of trainees who attended H3Africa funded training shared their knowledge/information with at least 3 people

"The trainings have been very informative and my request is for advanced courses for those who have gone through the introductions and are using the tools"

"Systems admin workshop: i can now confidently setup and administer Linux servers even of other departments"

Other impact:

- Set up new collaborations
- Organised their own workshop



