

Johnbosco Tayebwa

BIOINFORMATICIAN

School of Applied Health Sciences, Kampala International University

□ +256 704 281904 | ☑ bioinfotayebwa@gmail.com | ② 0000-0002-2355-8475 | ② biocodebreaker | ⓓ Johnbosco

Tayebwa

Experienced bioinformatician with a strong background in analyzing high-throughput genomic data and developing bioinformatics pipelines. Skilled in R, Perl, and Linux environments with expertise in next-generation sequencing data analysis.

Education

Faculty of Medicine Lund University, Sweden

PHD CANDIDATE 2015

University of Skovde Sweden

MASTER OF SCIENCE IN BIOINFORMATICS 2011-2012

Makerere University

Uganda

Bachelor of Biomedical Laboratory Technology, Second Class Honours – Upper Division 2005-2008

Professional Experience

School of Applied Health Sciences, Kampala International University

Assistant Lecturer Current

· Current Position

Department of Clinical Genetics, Lund University Hospital

BIOINFORMATICIAN 2012-2014

Sweden

School of Biosciences, University of Exeter

VISITING POSTGRADUATE RESEARCHER 2010-2011

· United Kingdom

Skills_____

PROGRAMMING

• Advanced skills in Linux-Unix bash/shell, R statistical programming, and Perl

HIGH PERFORMANCE COMPUTING

• Experience with HPC clusters (UPPMAX, SNIC) and cloud computing (Amazon AWS, docker)

NGS DATA ANALYSIS

• 5 years of experience in management and analysis of high-throughput data from Next-Generation Sequencing Platforms, especially Illumina

MICROARRAY ANALYSIS

· Experience with gene expression array and microarray (SNP-array) data analysis

Research Experience _____

Department of Clinical Genetics, Lund University Hospital

BIOINFORMATICIAN 2012-2014

 Analyzed NGS data (WGS, WES, TruSeq, RNA-Seq) for bone and soft tissue tumor studies. Developed pipelines for variant calling, fusion gene detection, and differential expression analysis.

School of Biosciences, University of Exeter

VISITING POSTGRADUATE RESEARCHER

2010-2011

• Developed competence in Linux environment and Perl scripting. Assembled bacterial genomes and developed in-silico primers for Xanthomonas species.

Publications

For a complete list of publications, please visit: https://scholar.google.com/citations?user=PZ6u9ocAAAAJ

Professional Membership

Member of the International Society for Computational Biology (ISCB)

Awards_

Swedish Institute

SWEDISH INSTITUTE SCHOLARSHIP

2011-2012

NARO/WORLD BANK PROJECT

MILLENNIUM SCIENCE INITIATIVE RESEARCH GRANT

2010-2011

Muljibhai Madhvani Foundation

MULJIBHAI MADHVANI FOUNDATION SCHOLARSHIP

2006-2008

References_

Available upon request.