# YEDOMON ANGE BOVYS **ZOCLANCLOUNON**

#### **Bioinformatician | Plant Genetics and Genomics**

As a newly graduated Doctor of Philosophy in Plant Genetics and Genomics, my fervor lies in the exploration of plant, fungi, fish, and mammalian genomes to uncover their evolutionary significance and advantageous genetic traits. My goal is to contribute towards promoting biodiversity preservation as well as enhancing advancements in molecular plant breeding.



#### **EDUCATION**

2020 2023

#### Jeonbuk National University

Ph.D. in Crop Science and Biotechnology

Jeonju, South Korea

Thesis: Genomic and phenotypic characterization of sesame germplasm for phylogenomics and breeding

2014 2016

#### **University Cheikh Anta Diop**

M.S. in Plant Pathology and Crop Protection

◆ Dakar, Senegal

Thesis: Characterization of downy mildew [Sclerospora graminicola (Sacc.) J. Schröeter] incidence and severity on 20 pearl millet [Pennisetum glaucum (L.) R. Br.] varieties and assessment of agronomic performance

2007 2012

### **University of Abomey-Calavi**

B.Sc. Agricultural Pest Management

Abomey-Calavi, Benin

Thesis: Toxicity of two plant powders as biopesticides in the management of Callosobruchus maculatus F. (Coleoptera: Chrysomelidae, Bruchinae) on two stored grain legumes

## RESEARCH EXPERIENCE

2018 2020

#### **Graduate Research Assistant**

Genomics Division, Department of Agricultural Biotechnology, National Institute of Agricultural Sciences

Jeonju, South-korea

- · Genome assembly annotation, and comparative analyses using Sesamum and Fusarium species.
- · Identification of peroxidase and laccase enzymes related to the biosynthesis of lignans from sesame seed.
- Construction of a sesame core collection and identification of accessions with high oil, and lignans content



Birth date: January 27, 1990 Native country: Benin

#### CONTACT INFO

- angez9914@gmail.com
- github.com/Yedomon
- twitter.com/AngeOmics
- https://yedomon-site.netlify.app/

#### **SKILLS**

#### Coding

R • Python • Bash • Shiny • Snakemake • Nextflow

#### Wet Lab

Plant and fungi DNA and RNA extraction • Sequencing library preparation • PCR • RT-PCR

#### **Dry Lab**

Genome assembly • Genome annotation • Comparative genomics • Pan-genome analysis • Population genomics • QTL • **GWAS** 

#### Languages

French: Native • English: Fluent •

Korean: Intermediate

2014 2015

#### **Graduate Research Assistant**

Centre d'Etudes Régional pour l'Amélioration de l'Adaptation à la Sècheresse (CERAAS)

Thies, Senegal

- · Evaluation and mapping pearl millet disease incidence within Senegal agro-ecological areas.
- · Identification of resistant genotypes as parental lines for millet breeding.

### **TEACHING EXPERIENCE**

2021

Introduction to R Language for GIS application - Raster data

Instructor of R and GIS data mining for the DAAD climapAfrica -Working group Climate change Animal and Plant Physiology and Stress. | Press release | Training material

Remote

2020

Introduction to R Language for GIS application - Shapefile data session

Instructor of R and GIS data mining for the DAAD climapAfrica -Working group Climate change Animal and Plant Physiology and Stress. | Press release | Training material

Remote

### A HONORS AND AWARDS

2014

In Country and in Region Scholarship for Master Science graduate Award

German Academic Exchange Service, Deutscher Akademischer Austausch Dienst (DAAD)



## SOCIETY MEMBERSHIP

2020 Present International Society for Computational Biology: ISCB Member



#### **VOLUNTEERING**

2021 Present African Biogenome Project (AfricaBP) - Pilot Committee Co-Chair of the Genome Annotation Sub-committee | Link

2022

African Biogenome Project (AfricaBP) - Pilot Committee Co-organizer of the AfricaBP Workshop on Biodiversity Genomics Technologies and Infrastructures | Link

### SELECTED PUBLICATIONS

Genomic and phenotypic characterization of sesame germplasm for phylogenomics and breeding 
PhD Thesis Document | Download
Zoclanclounon YAB

- Genome-wide development of intra- and inter-specific transferable SSR markers and construction of a dynamic web resource for yam molecular breeding: Y2MD Preprint bioRxiv | doi: https://doi.org/10.1101/2023.03.09.531889

  Moussa D, Zoclanclounon YAB, Mboup PA, Diouf D, Malédon E, Rivallan R, Chair H and Dossa K
- Haplotype-resolved telomere-to-telomere assembly of the African catfish (Clarias gariepinus) provides insights for terrestrial adaptation of air-breathing catfishes
   Preprint bioRxiv | doi: https://doi.org/10.1101/2023.03.23.533919

   Nguinkal JA, Zoclanclounon YAB, Brunner RM and Goldammer T
- Characterization of peroxidase and laccase gene families and in silico identification of potential genes involved in upstream steps of lignan formation in sesame

  MDPI Life | doi: https://doi.org/10.3390/life12081200

  Zoclanclounon YAB, Rostás M, Chung N-J, Mo Y, Karlovsky P and Dossa K
- A chromosome-level genome of the Korean cultivar

  Sesamum indicum var. Goenbaek

  Data Set | NCBI Project: PRJNA810203 | doi: https://doi.org/10.5281
  /zenodo.6350880

  Zoclanclounon YAB, Kim J-G, and Lee K
- A genome assembly of the sesame wilt disease pathogen

  Fusarium oxysporum f.sp. sesami

  Data Set | NCBI Project: PRJNA667197

  Zoclanclounon YAB, Jung H-J, Kim S-U, Yoon Y, Choi H-W, Kim TH and Lee K

#### REFEREES

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- Komivi Dossa, Ph.D.
  Researcher at CIRAD, UMR AGAP Institut, Guadeloupe, France. |
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