Personal Data	
Name:	Duncan N. Ndegwa, PhD
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## Bio

Duncan N. Ndegwa, holds a PhD from the University of Tuebingen in Germany where he studied the biology of the apicoplast during erythrocytic schizogony of *P. falciparum*. He then joined Embu University as a Lecturer in Biochemistry and molecular cell biology during which he secured a Postdoctoral fellowship with Julian Rayner at the Welcome Sanger Institute, UK, focusing on antigen screening for blood stage vaccine candidates in *Plasmodium vivax* using genetics, molecular and cellular biology. He later worked as a Post-Doc associate with Dr. Amy Bei lab at Yale University, US, where he researched on the mechanisms behind the persistence of specific parasite genotypes and the role of naturally arising polymorphisms in blood stage vaccine candidate antigens on receptor binding and immune evasion. He is now working as a Lecturer at Embu University. His interests are both in basic and applied research using experimental genetics, proteomics and bioinformatics to study host-parasite interaction and to apply this for development of intervention strategies against malaria and other neglected tropical diseases.

Career	
01/2022 – To Date	Lecturer in the Department of Biological Sciences University of Embu, Kenya
	<ul> <li>Duties and responsibilities: Lecturing and training of both undergraduate and Post graduate students</li> </ul>
10/2019-12/2021	<ul> <li>Post-Doctoral Scientist         Yale University New Haven CT.         <ul> <li>Research Project:</li> <li>Mechanisms behind the persistence of specific parasite genotypes and</li> <li>The role of naturally arising polymorphisms in blood stage vaccine candidate antigens on receptor binding and immune evasion</li> </ul> </li> </ul>
06/2017- 09/2019	Post-Doctoral Scientist Welcome Sanger Institute, Hinxton, Cambridge, UK • Research Project:
	<ul> <li>Antigen screening of <i>Plasmodium vivax</i> blood-stage malaria vaccine targets using <i>P</i>.</li> </ul>
Education and Trair	ning
10/2012 - 07/2016	Doctor of Science in Biochemistry (Grade: Magna cum laude)
	Tübingen University and Magdeburg University, Germany
	• Thesis:
	<ul> <li>Study of the apicoplast biology in <i>Plasmodium falciparum</i> during erythrocytic schizogony</li> </ul>
08/2009 – 12/2011	Masters of Science in Biochemistry (Grade: A)
	Heidelberg University, Germany and Edgerton University, Kenya
	• Thesis:
	<ul> <li>Study of resistance to artemisinin in <i>Plasmodium falciparum</i> isolates from Kilifi County, Kenya</li> </ul>
08/2005 – 06/2009	Bachelor of Science (Grade: First-class honors) Egerton University, Kenya
Other experiences	
01/2017 - 06/2017	Lecturer in the Department of Biological Sciences University of Embu, Kenya
	<ul> <li>Duties and responsibilities: Lecturing undergraduate and Post graduate students</li> </ul>
04/2015 – 06/2016	<ul> <li>Development of new malaria diagnosis methods</li> <li>Carl Zeiss AG / Institute for tropical medicine, Tübingen University, Germany</li> <li>Cell culture, cell-based assays, sample preparation, evaluation of assays and equipments, report writing, training of colleagues, project management</li> </ul>
04/2014 – 05/2016	Volunteer Microscopist Institute for tropical medicine, Tübingen University, Tübingen Germany

	<ul> <li>Malaria diagnosis during malaria clinical trials</li> </ul>
08/2011– 12/2011 Skills	<ul> <li>Volunteer Intern</li> <li>Wellcome Trust Kemri-Kilifi, Kilifi, Kenya</li> <li>Sample processing of field isolates for sequencing and preservation (parasite, freeze down), carrying out research experiments, data collection, database management, project managements</li> </ul>
Wet-lah skills	Genetic modification: CRISPR-case DNA and RNA Isolation CDNA- synthesis PCR aPCP cloning
Wet-lab Skills	transfection construct design and generation for gene targeting, unconditional and conditional gene knock-out, transfection, agarose electrophoresis <b>Cell culture:</b> <i>P. falciparum, P. knowlesi,</i> HEK293, THP1
	Inhibition assays: Invasion/growth inhibition assays, drug inhibition assays, flow cytometry Immuno-assays: Polyclonal and monoclonal antibody purification, protein microarray, western blot, ELISA, immunofluorescence assay(IFA)
	<b>Cellular Imaging:</b> Fluorescence microscopy, light microscopy <b>Others</b> : Parasite synchronisation, merozoite isolation, protein expression, protein purification, SDS-Page electrophoresis, NGS library preparation
Computational-skills	<b>Data analysis:</b> Protein microarray analysis, Python, Bash (Basic user), R, Stata and Prism GraphPad <b>Bioinformatics:</b> NGS Whole genome sequence analysis, Bash, Python, Benchling, DNASTAR Navigator <b>Others:</b> Flow jo
Additional Informatio	n
Languages	English (Fluent), German (Intermediate level), and Swahili (Native speaker)
Awards	<ol> <li>2012 German Academic Exchange Services PhD fellowship</li> <li>2011 German Academic Exchange Services Masters fellowship</li> <li>2010 German Academic Exchange Services in country scholarship-Kenya</li> </ol>
Conferences/Workshops	<ol> <li>Aspiring leaders in science, Sanger Institute, Hinxton, UK</li> <li>7th Short Course for Young Parasitologists, Hamburg, Germany</li> <li>Innovation workshop on malaria diagnosis, Carl Zeiss, Jena Germany</li> <li>RNAseq data analysis, University of Glasgow, Scotland, UK</li> </ol>
References	
Dr. Amy Bei Yale Un	iversity. Yale School of Public Health. 60 College Street. New Haven CT 06520-8034
Prof. Julian Rayner Ca	ambridge University. CIMR, The Keith Peters Building, Hills R, Cambridge, CB2 0XY. UK
Publications	

Published:

Ndegwa DN, Kundu P, Hostetler JB, Marin-Menendez A, Sanderson T, Mwikali K, et al. (2021) Using *Plasmodium knowlesi* as a model for screening *Plasmodium vivax* blood-stage malaria vaccine targets reveals new candidates. PLoS Pathog 17(7): e1008864. https://doi.org/10.1371/journal.ppat.1008864

## Preprints:

- Ndegwa DN, Tuju J, Chepsat E, Mwai R, Mwai K, Nyamako L, Mosobo M, Deme AB, Dieye B, Ndiaye PI, Diedhiou Y, Mbaye AM, Wirth D, Ndiaye D, Osier F, Bei AK. High-throughput profiling of natural acquired humoral immune response to a large panel of known and novel full length ectodomain P. falciparum merozoite vaccine candidates under reduced malaria transmission. doi: <a href="https://doi.org/10.1101/2022.02.15.479108">https://doi.org/10.1101/2022.02.15.479108</a>
- Khadidiatou Mangou, Adam J. Moore, Laty Gaye Thiam2, Aboubacar Ba, Alessandra da Silva Orfano, Ife Desamours, Duncan Ndungu Ndegwa, Justin Goodwin, Yicheng Guo, Zizhang Sheng, Saurabh D. Patel, Fatoumata Diallo, Seynabou Diouf Sene, Mariama Nicole Pouye, Awa Thioub Faye, Bacary Djilocalisse Sadio, Lawrence Shapiro, Ousmane Faye, Alassane Mbengue, Amy K. Bei1. Identification of novel variants in the vaccine candidate PfRh5 in a highly endemic region of Senegal. doi:https://doi.org/10.1101/2022.03.17.22271950