Andree Valle Campos, MSc(c)

♥ avallecam.github.io ■ avallecam@gmail.com Pron: Él/He/His \bigcirc (+51) 950 951 722 \triangle Lima - Peru https://orcid.org/0000-0002-7779-481X Interests Systems developmental biology. Quantitative and Bioengineering approaches. Reproducibility. Open science. Best-practices of teaching. Causal questions for hypothetical interventions. 2018-2018 Master of Science in Epidemiological Research **EDUCATION** Universidad Peruana Cayetano Heredia (UPCH), Lima-Peru Rank: Top fifth among 28 students. 2011-2015 Bachelor of Science in Genetics and Biotechnology Universidad Nacional Mayor de San Marcos (UNMSM), Lima-Peru Rank: Top third at the 4th term among 30 students. 2019-2020. National Center for Epidemiology (CDC Peru) Researcher Affiliations Epidemiological Research and Surveillance Group, Ministry of Health. 2017-2019 Universidad Peruana Cayetano Heredia (UPCH), Peru. Intern Emerge, Emergent Diseases and Climate Change Research Unit. 2016-2017 Universidad Nacional de la Amazonía Peruana (UNAP), Peru. Consultant Fundación para el Desarrollo Sostenible de la Amazonía Baja. 2015-2016 U.S. Naval Medical Research Unit Six (NAMRU-6), Peru. Intern Dept. of Parasitology, Div. of Immunology and Vaccine Development. Peer-reviewed (n=6) **PUBLICATIONS** -Reyes-Vega MF, Soto-Cabezas MG, Cárdenas F, Martel KS, Valle A, et al. "SARS-CoV-2 prevalence (N=8)associated to low socioeconomic status and overcrowding in an LMIC megacity: A population-based seroepidemiological survey in Lima, Peru". EClinical Medicine. doi: 10.1016/j.eclinm.2021.100801 & . -Gunderson AK, Kumar RE, Recalde-Coronel C, Vasco LE, Valle-Campos A, et al. "Malaria Transmission and Spillover across the Peru-Ecuador Border: A Spatiotemporal Analysis". Int. J. Environ. Res. Public Health 2020, 17, 7434. doi: 10.3390/ijerph17207434 & . -Quispe AM, Pinto DF, Huamán MR, Bueno GM, & Valle-Campos A. ["Quantiative Methodologies: Sample size calculation with STATA and R."] Revista del Cuerpo Médico del HNAAA, 2020, 13(1), 78-83. doi: 10.35434/rcmhnaaa.2020.131.627 ピ. -Munayco CV, Tariq A, Rothenberg R, Soto-Cabezas MG, Reyes MF, Valle A., et al. "Early transmission dynamics of COVID-19 in a southern hemisphere setting: Lima-Peru: February 29th-March 30th, 2020.". Infectious Disease Modelling, 2020, 5, 338 - 345. doi: 10.1016/j.idm.2020.05.001 \(\mathref{\pi}\). -Loyola S., Valle A., Montero S. and Carrasco-Escobar G. ["Recomendations to properly describe a COVID-19 epidemic curve."] Revista Peruana de Medicina Experimental y Salud Pública, 2020, 37(2). doi: $10.17843/\text{rpmesp.}2020.372.5461 \ \ \ \ \ \ \ \ \ \ \$ -Saavedra-Langer R., Marapara J., Valle-Campos A., et al. "IgG subclass responses to excretedsecreted antigens of Plasmodium falciparum in a low transmission malaria community of the Peruvian Amazon". Malaria journal, 2018, 17(1), 328. doi: $10.1186/s12936-018-2471-6 \ \mathbb{Z}$. At Newsletters (n=2) -[Opinion] Carrasco-Escobar G, Incio J, Valle A., Martínez JJ, Prochazka M, Ugarte C. ["Data and Transparency to fight against to coronavirus."] Ojo Público, 2020. url: ojo-publico.com & . -[Editorial] Valle-Campos A. ["Health Data Science: Applications at the Peruvian Center for Epidemiology, Prevention and Disease Control, CDC-Peru." Boletín Epidemiológico del Perú, 2019, 18(49), 1245. doi: 10.5281/zenodo.4014211 2. Scholarship. Emerge Training Grant NIH/FIC TG D43 TW007393 2018 USD 10.644 Grants, Awards Grant. Undergraduate research: Camelid Reproduction Group - UNMSM 2014 **USD 500** AND RECOGNITIONS Ranked 1st. UPCH X Summer Course on Molecular Biology (40 students) 2013 Ranked 1st. UNMSM Admission Test to Basic Sciences (1000 applicants) 2011

R (developer of work-in-progress packages like serosurvey).

GNU/Linux (Ubuntu). LATEX, R Markdown. SublimeText. Git.

Bash (Unix shell, fluent), Python (basic), Stata (fluent).

Statistical programming:

Programming Language:

OS, Text editor, & more:

Computational

SKILLS

Conference Participant	serosurvey: Serological Surveys and Prevalence Estimation Under Misclassiff Elevator pitch at the useR! Conference. Online. \bigcirc	ication. 2021	
	[Epidemiological analysis of the epidemic of Guillain Barré Syndrome in Pe Poster presentation at the INS International Scientific Congress. Lima, Peru. • • • • • • • • • • • • • • • • • • •	•	
	Human mobility and malaria history in a periurban community in Iquitos, Poster presentation at the ASTMH Annual Meeting. Maryland, USA. • C	amunity in Iquitos, Peru. 2019	
	In vitro effect of ELF Magnetic Field on the sperm motility of Alpacas Poster, Annual Meeting of the Bioelectromagnetics Society, BioEM2015. Monterey, US	2015 A. ೧ ♂	
WORKSHOP INSTRUCTOR	Outbreak Analytics and Modelling for Public Health	2021 ats.	
	[Basic R applied to disease surveillance and outbreak analysis] O 6 hours Introduction to R projects and ggplot2 graphics for Ministry of Health personel. 30 str	2021 $udents.$	
	[Epidemiological analysis using R] \bigcirc 4 hours Applications to case-control, cohort and time to event study designs. 30 students.	2019	
	[Introduction to Inferential Statistics for biologist] • 6 hours Introduction to R, Linear models and Multiple comparison. 40 students.	2019	
	[Reproducible science and Microarray analysis] \bigcirc 8 hours Designs, statistics and visualizations with Bioconductor and Tidyverse. 50/20 students	2017/19 nts.	
LECTURER AND TEACHING	[Data analysis in epidemiological surveillance I: time, space, person] 2 2 how Descriptive and statistical analysis of outbreaks. 35 grad students.	urs 2021	
	[Visualizing public health and field epidemiology data] ? 2 hours Dashboards as tools for decision making in public health. 30 grad students.	2021	
	Teacher Assistant . At the Master's of Science in Epidemiological Research. 2 1 yea. In charge of practical sessions, monthly reviews, and test correction. 48 grad students.	r 2019	
	[Gene Regulatory Networks: Topology and Dynamics] 2 3 hours Applications from Graph Theory and Finite Automata. 10 undergrad students.	2015-18	
	[On #tardigate and Horizontal Gene Transfer bioinformatics] 2 hours Review of the controversy around the first tardigrade genome. 5 undergrad students.	2016	
TALKS	[Analysis of #multiple epidemics and prevalences with R and purrr.] $ ilde{\sc c}$ 50 $ ilde{\sc b}$	part. 2020	
	[Hypothesis testing with nonparametric statistical methods.] 2 30 participan	ts. 2020	
	[How to use R for Epidemiology at CDC Peru?] 25 participants.	2019	
Trainings with scholarship	[How to teach programming online] [2] Techniques to design programming courses and evaluate students. One day.	2021	
	Outbreak Analytics and Modelling for Public Health $\ \ \ \ \ \ \ \ \ \ $ Dynamic modeling in response to outbreaks and interventions. One week.	2019	
	CODATA-RDA Research Data Science School 2017 and 2020 Data management, open science, machine learning and infrastructure. Two weeks.	2017/20	
	[Minicourse on Spatio-Temporal Models in Epidemiology] C Theory and practice of areal data, point pattern analysis and geostatistics. Two days.	2017	
	Working with Parasite Database Resources © Genomic, proteomic, metabolomic applications of eupathdb.org. One week.	2016	
	School on Physics Applications in Biology © Game theory, non-linear dynamics and statistical physics. Three week.	2016	
	V Southern-Summer School on Mathematical Biology ♂ Population dynamics modeling in ecology and epidemiology. One week.	2016	
CERTIFICATIONS	English: TOEFL score 88 (21/23/20/24). Test date	Test date: 14 Dic 2020	

CERTIFICATIONS

English: TOEFL score 88 (21/23/20/24).

Biomedical Research - Basic/Refresher: CITI program.

Responsible Conduct in Research: QUIPU program - Peru.

Completition date: 05 May 2018