Andree Valle Campos, MSc(c)

♥ avallecam.github.io ■ avallecam@gmail.com Pron: Él/He/His (9 (+51) 950 951 722 ♣ Lima - Peru https://orcid.org/0000-0002-7779-481X Interests Quantitative and systems developmental biology. Causal questions for hypothetical interventions. Reproducible and open science. Biostatistics. Bioengineering approaches. Best-practices of teaching. Master of Science in Epidemiological Research 2018-2018 **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. Researcher 2019-2020. National Center for Epidemiology (CDC Peru) 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 - 2016U.S. Naval Medical Research Unit Six (NAMRU-6), Peru. Intern Dept. of Parasitology, Div. of Immunology and Vaccine Development. **PUBLICATIONS** Peer-reviewed (n=6) -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 \(\mathref{C}\). -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 \Box . -Munayco CV, Tariq A, Rothenberg R, Soto-Cabezas MG, Reves 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 \(\mathred{\texts} \). -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: $10.17843/\text{rpmesp.}2020.372.5461 \ \text{\square}$. -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: $10.1186/s12936-018-2471-6 \ \text{C}$. 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 ♂. Scholarship. Emerge Training Grant NIH/FIC TG D43 TW007393 2018 USD 10.644 Grants, Awards Grant. Undergraduate research: Camelid Reproduction Group - UNMSM USD 500 2014 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: serosurvey, and more).

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 Misclassification Elevator pitch at the useR! Conference. \bigcirc	on. 2021
	[Epidemiological analysis of the epidemic of Guillain Barré Syndrome in Peru.] Poster presentation at the INS International Scientific Congress. Ω	2019
	Human mobility and malaria history in a periurban community in Iquitos, Peru Poster presentation at the ASTMH Annual Meeting. Ω	2019
Workshop Instructor	Outbreak Analytics and Modelling for Public Health	2021
	[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 student	2021
	[Epidemiological analysis using R] Ω 4 hours Applications to case-control, cohort and time to event study designs. 30 students.	2019
	[Introduction to Inferential Statistics for biologist] O 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
TEACHER AND LECTURER	[Data analysis in epidemiological surveillance II: spatial analysis] 2 hours Generate maps with risk estimates using R. 35 grad students.	2021
	[Data analysis in epidemiological surveillance I: time, space, person] 2 hours Descriptive and statistical analysis of outbreaks. 35 grad students.	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 year In charge of practical sessions, monthly reviews, and test correction. 48 grad students.	2019
	[Gene Regulatory Networks: Topology and Dynamics] 3 hours Applications from Graph Theory and Finite Automata. 10 undergrad students.	2015-18
	[On #tardigate and Horizontal Gene Transfer bioinformatics] 2 2 hours Review of the controversy around the first tardigrade genome. 5 undergrad students.	2016
Talks	[Introduction to spatial analysis.] 2 30 participants.	2021
	[Analysis of #multiple epidemics and prevalences with R and purrr.] \circlearrowleft 50 part.	2020
	[Hypothesis testing with nonparametric statistical methods.] \circlearrowleft 30 participants.	2020
	[How to use R in Epidemiology?] $\@ifnextchar[{\@model{C}}{\mathcal{C}}$ 25 participants.	2019
TRAININGS WITH SCHOLARSHIP	[How to teach programming online] $\ \ \ \ \ \ \ \ \ \ \ \ \ $	2021
	Outbreak Analytics and Modelling for Public Health O 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] 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
	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). Biomedical Research - Basic/Refresher: CITI program. Expiration date: 04 M. Responsible Conduct in Research: OUIPI program - Peru. Completition date: 05 M.	May 2021

Responsible Conduct in Research: QUIPU program - Peru.

Completition date: 05 May 2018