

Unraveling the unparalleled 2024 epidemic of Dengue in the Americas

Desentrañando la incomparable epidemia de dengue de 2024 en Las Américas

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Document prepared by a panel of international experts convened by AJRM on the occasion of the reactivation and expansion of dengue in Latin America

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Conflicts of Interest. AJ Rodríguez-Morales has served as speaker and consult of Takeda and Valneva laboratories in relation with dengue and chikungunya vaccines, respectively. Rest of authors, no conflicts of interest declared.

Received: May 23th, 2024

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The year 2024 marked a pivotal moment in the history of public health in the Americas as the region grappled with an unprecedented surge in dengue cases^{1,2}. Dengue, primarily transmitted by *Aedes aegypti* and *A. albopictus*, now also has the risk of transmission by *A. vitattus*³, detected in Caribbean countries such as the Dominican Republic and Puerto Rico. Dengue has long been a concern in tropical and subtropical regions. However, the magnitude and severity of the outbreaks witnessed so far in 2024 surpassed anything previously recorded, posing immense challenges to healthcare systems, governments, and communities across the continent^{4,5}. Further, it must be regarded as the potential role of travellers in the risk of introducing dengue into areas where mosquito vectors may be present and where there is no indigenous circulation of the virus. Therefore, there is a need to educate on preventive measures for both the population living in endemic areas and travellers from non-endemic areas. Since Latin America is one of the most frequent destinations for travellers from Europe and North America, dengue vaccination should be widely considered as a prevention tool within the scope of national dengue control programs in endemic areas and in a perspective of traveler health^{6,7}.

In the Americas, which reported 80% of the cases worldwide in 2023 (<https://www.who.int/emergencies/disease-outbreak-news/item/2023-DON498>), more than 30 countries and territories report, weekly and systematically, dengue information to the Pan American Health Organization (PAHO) (www.paho.org). Additionally, the laboratories that integrate the Dengue Laboratory Network of the Americas provide periodic information regarding the circulating serotypes in the region⁸. Nevertheless, heterogeneity in diagnosis and report may occur among some countries⁹. Just during the first 20 epidemiological weeks of 2024, a total of 8,246,951 dengue cases have been reported in the Americas (Figure 1). Of these, 3,839,613 (46.6%) were confirmed. From the total, 8,084 cases (0.09%) were classified as severe dengue, with significant variation among countries (Figure 2). In 2024, 3,666 deaths have been reported, with a case fatality rate of 0.04% (4 deaths per 10,000 cases), also with high variation among countries (Figure 3). This burden, specifically in terms of Disability-Adjusted Life Years (DALYs), represents 194,022.35 DALYs, for a rate of 32.65 DALYs per 100,000 pop. in Latin America just in 2021, according to the data of the 2021 Global Burden of Study (GBS) (<https://vizhub.healthdata.org/gbd-results/>) (Figure 4).

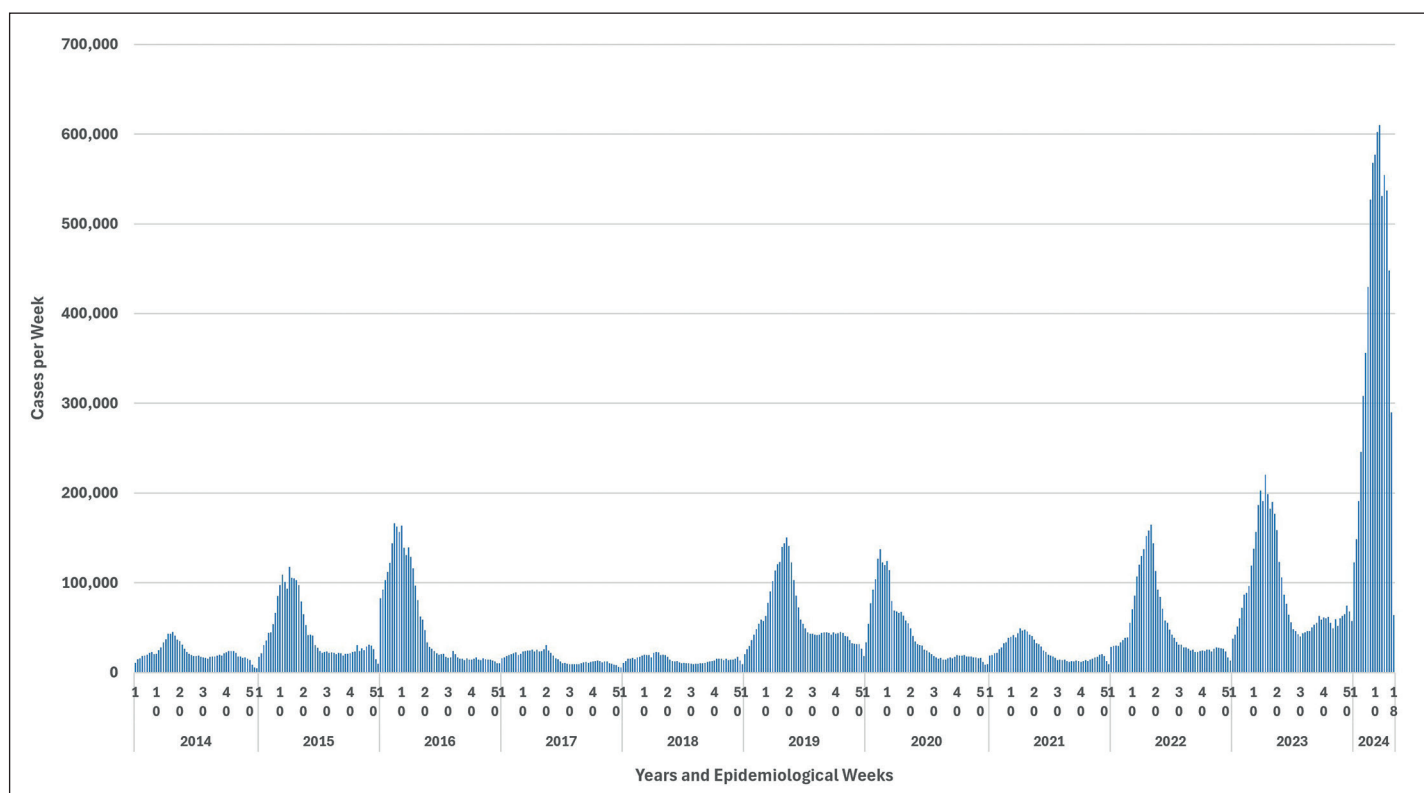


Figure 1. Dengue cases by epidemiological weeks, Americas, 2014-2024. It is modified from PAHO.

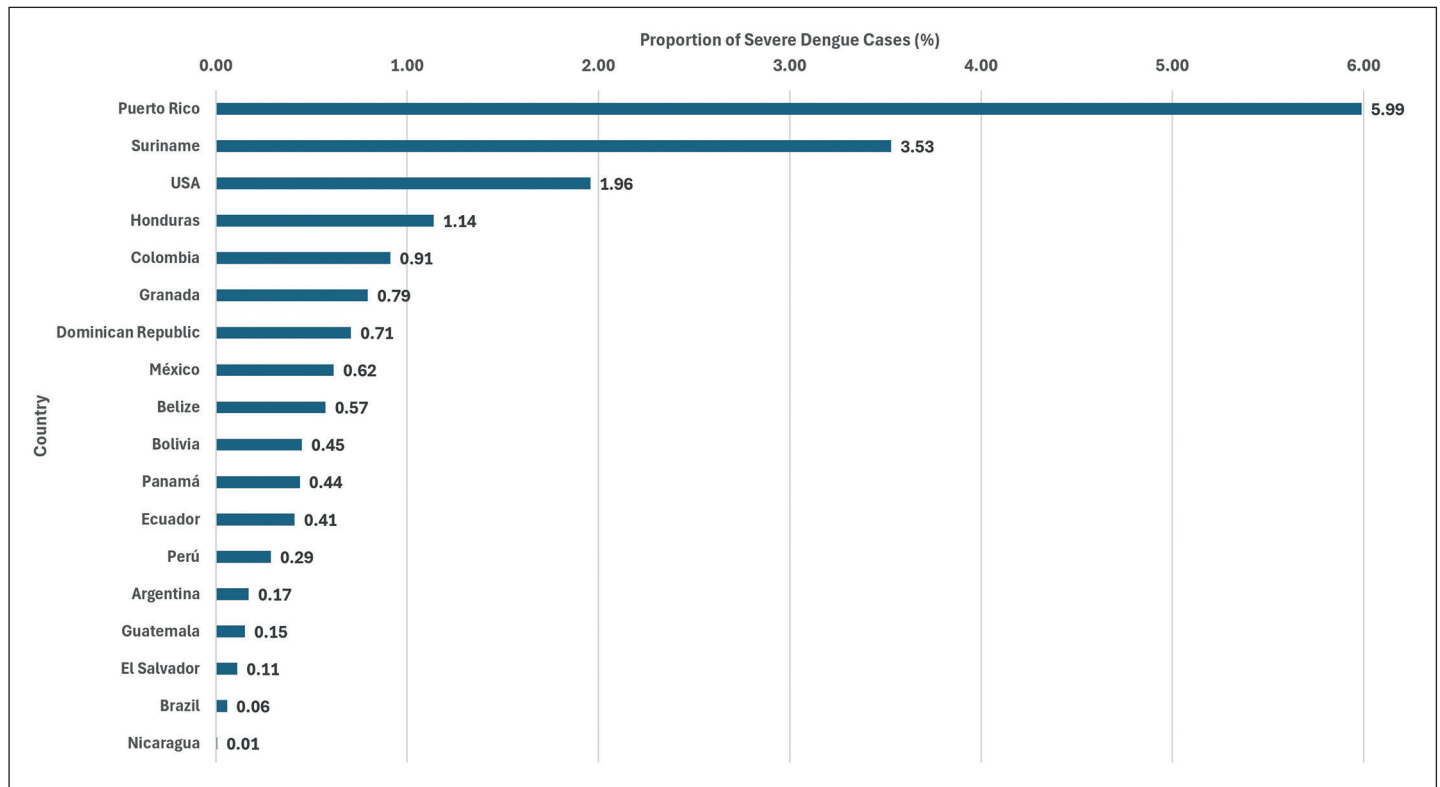


Figure 2. Proportion of Severe Dengue cases by countries, Americas, 2024. Modified from PAHO.

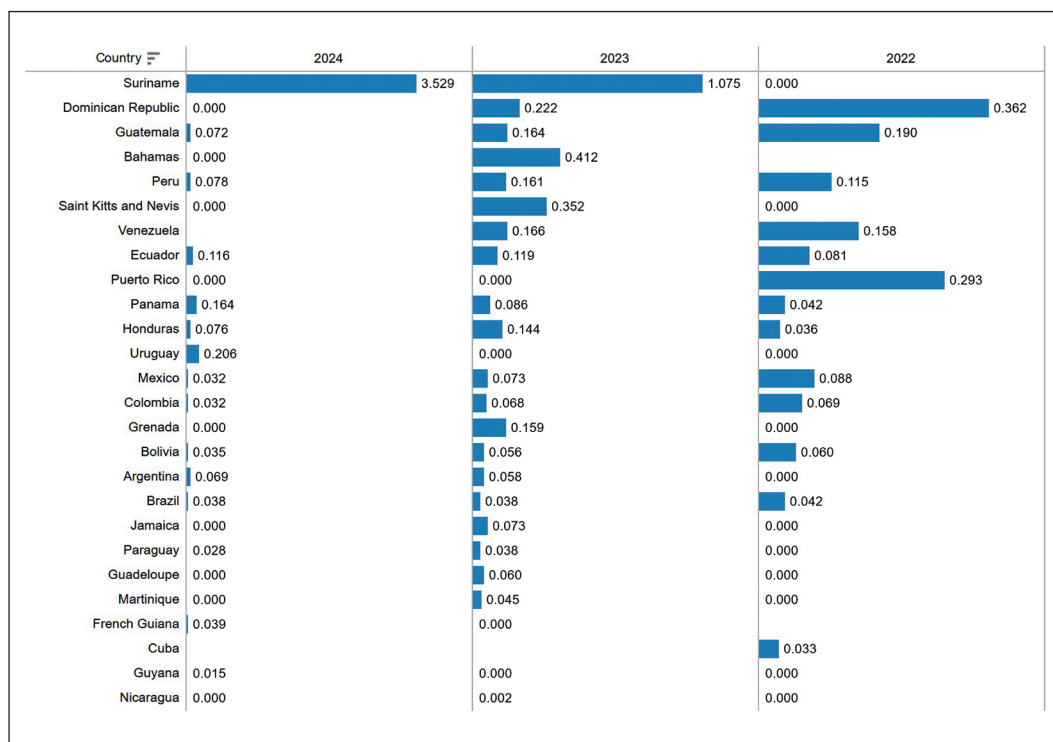


Figure 3. Case fatality rate (%CFR) associated with Dengue cases by countries, Americas, 2022-2024. From PAHO.

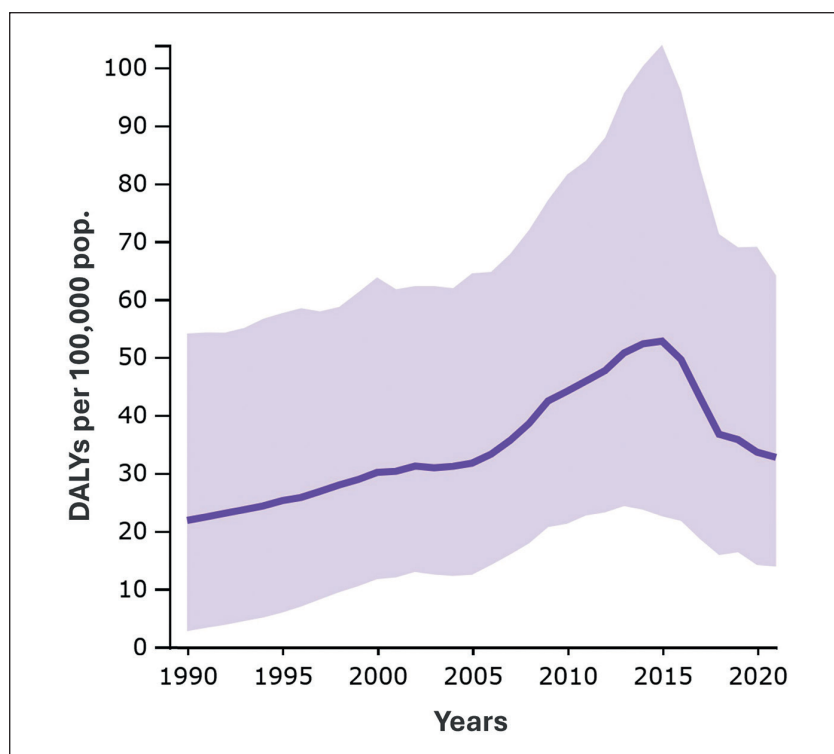


Figure 4. DALYs associated to Dengue in Latin America and the Caribbean, 1990-2021. Modified from 2021 GBS.

Regarding severe dengue, it is essential to consider the associated risk of antibody-dependent enhancement (ADE) of infection, which occurs when preexisting antibodies present in the body from a primary (first) dengue virus (DENV) infection bind to an infecting virus particle during a subsequent infection with a different dengue serotype^{10,11}. Antigenic distance between primary and secondary dengue infections correlates with disease risk¹². However, severe dengue may occur as a consequence of primary infection. Recent findings question the widely held belief that severe dengue is associated predominantly with secondary infections and emphasize the importance of developing vaccines or treatments to protect dengue-naïve populations and seropositive subjects¹³.

The four distinct dengue virus serotypes (DENV-1, DENV-2, DENV-3, and DENV-4), are circulating in the Americas. Several factors converged to fuel the unparalleled dengue epidemic in the Americas in 2024. Climate change, urbanization, globalization, co-circulation of different serotypes in areas with a large proportion of the population exposed in previous epidemics, and inadequate mosquito control efforts created a perfect storm for the proliferation of *Aedes* mosquitoes and the massive transmission of DENV^{14,15}. Rising temperatures and changing

rainfall patterns created conducive environments for mosquito breeding. At the same time, rapid urbanization led to densely populated areas with inadequate sanitation and water management systems, providing ample breeding grounds for mosquitoes¹⁶⁻¹⁸. Multiple other arboviral diseases are concurrently circulating in the Americas, especially Chikungunya and Zika, as well as Mayaro, Oropouche and Yellow Fever¹⁹⁻³². In addition, outbreaks of South American endemic hemorrhagic fever (HF) viruses such as Junin (Argentinian HF), Machupo and Chapare (Bolivian HF), Sabia (Brazilian HF), and Guanarito (Venezuelan HF)³³⁻³⁷ were also reported.

Given the arboviral cocirculation, the Global Arbovirus Initiative, launched on 31 March 2022, is an essential cross-cutting effort initiated across the World Health Organization's Health Emergencies Programme, the Department of Control of Neglected Tropical Diseases, and the Immunization, Vaccines and Biologicals Department in collaboration with a growing body of multisectoral international partners. The pillars of the initiative provide a framework for objectives and priority activities to tackle emerging and re-emerging arboviruses with epidemic and pandemic potential focusing on monitoring risk, pandemic prevention, preparedness, detection and response, and building a coalition of partners (<https://www.who.int/initiatives/global-arbovirus-initiative>).

Globalization facilitated the rapid spread of dengue, as increased travel and trade facilitated the movement of infected individuals and vectors across borders. Moreover, the challenges posed by the COVID-19 pandemic, which strained healthcare systems and diverted resources away from mosquito control programs, further exacerbated the situation³⁸⁻⁴². Hospitals and clinics were overwhelmed with patients presenting with dengue symptoms, leading to overcrowding, shortages of medical supplies, and increased pressure on healthcare workers. The dual burden of dengue and COVID-19 stretched resources to their limits, highlighting the need for robust and resilient healthcare systems capable of responding to multiple concurrent crises^{43,44}.

While dengue fever is endemic to many countries in the Americas, the burden of the 2024 epidemic was not evenly distributed^{45,46}. Countries with weaker healthcare systems and limited resources faced disproportionate challenges in responding to the outbreak, as is probably the case for some Caribbean islands and low-income countries (Figures 2 and 3). Rural areas with limited access to healthcare and sanitation were particularly vulnerable, as were marginalized communities lacking basic infrastructure and public health services⁴⁷.

Furthermore, socioeconomic factors such as poverty, overcrowding, poor environmental management of solid waste, and inadequate housing exacerbated the risk of dengue transmission in vulnerable populations. The

lack of access to mosquito repellents, proper sanitation, and healthcare services hindered prevention and control efforts, allowing the virus to spread unrestrictedly in many communities⁴⁸⁻⁵⁰. It should be emphasized that the One Health perspective should be included as a multidisciplinary approach to better understanding the course of emergencies and the re-emergence of dengue, which can trigger epidemics worldwide^{51,52}. The pressure of climate change, global human migration, and growing interactions between wild fields and human implications in that field^{1,2}. Conversely, certain regions in South America facilitate the accumulation of waste, including containers, vehicle tires, and other items that collect rainwater, thereby serving as artificial breeding grounds for mosquitoes. Compounding the challenge of dengue control, *Aedes* mosquitoes have developed resistance to repellents and insecticides^{53,54}.

The economic consequences of the 2024 dengue epidemic are profound, affecting individuals, communities, and entire countries. The direct and indirect costs associated with medical treatment, lost productivity, and vector control efforts burden already strained economies. Businesses suffered as employees fell ill, and tourism, a significant source of revenue for many countries in the Americas, was impacted by travel advisories and concerns about dengue transmission⁵⁵⁻⁵⁷.

Moreover, the long-term economic impact of dengue fever⁵⁸, including its effects on workforce productivity and healthcare expenditures, threatened to exacerbate existing inequalities and hinder economic development in the region⁵⁹.

In the face of this unprecedented dengue epidemic, governments and public health authorities in the Americas implemented a range of response and mitigation efforts to contain the spread of DENV and reduce its impact. These efforts included intensified mosquito control measures, such as insecticide spraying, larval source reduction, and community mobilization campaigns to promote awareness and prevention⁶⁰⁻⁶².

Healthcare systems ramped up their capacity to diagnose and treat dengue cases. At the same time, efforts were made to improve the infrastructure for laboratory diagnosis, surveillance and reporting to understand the dengue epidemic's scope, dynamics and circulation of other arboviruses. Collaboration and coordination among countries and international organizations are crucial in

sharing information, expertise, and resources to support the response efforts⁶³⁻⁶⁵.

Innovations in vector control, such as Wolbachia-infected mosquitoes and novel insecticides, are promissory to control mosquito populations and reduce DENV transmission^{1,66-70}. The inclusion of spatio-temporal modelling and Artificial Intelligence (AI) can be used for promising predictive purposes in the fight against dengue^{1,51,52,71,72}. The integrated management strategy for preventing and controlling arboviral diseases in the Americas is key for the region (<https://iris.paho.org/handle/10665.2/51787>). Additionally, increase production capacity and access to the vaccine already available, in addition to research into new dengue vaccines and therapeutics continued, offering hope for future prevention and treatment strategies⁷³⁻⁷⁸. Furthermore, the role of the current introduction of a second dengue vaccine in the region, in countries such as Argentina and Brazil, needs further assessment, especially regarding effectiveness and safety in the upcoming months and years.

The unparalleled epidemic of dengue fever in the Americas in 2024 served as a stark reminder of the complex interplay of factors driving infectious disease outbreaks in an increasingly interconnected world. Climate change, urbanization of new areas without adequate planning in basic public services, globalization, and socioeconomic disparities all contributed to the emergence and spread of dengue, highlighting the need for a multifaceted and coordinated approach to disease prevention and control⁷⁹⁻⁸².

While the challenges posed by the 2024 epidemic were immense, they also spurred innovation, collaboration, and a renewed commitment to strengthening public health systems, addressing the root causes of disease transmission, and delving into innovative methods in epidemiological surveillance and vector control. As the region continues to grapple with the threat of dengue and other infectious diseases, the lessons learned from the 2024 epidemic will be invaluable in shaping future efforts to protect the health and well-being of populations across the Americas. As all these preventive measures have failed to control dengue in the region for many decades, the approval and introduction of new dengue vaccines seem to be an urgent public health intervention and probably one of the best strategies to face this disease.

References

- 1.- Cabrera M, Leake J, Naranjo-Torres J, Valero N, Cabrera JC, Rodríguez-Morales AJ. Dengue Prediction in Latin America using machine learning and the one health perspective: a literature review. *Trop Med Infect Dis* 2022;7. doi: 10.3390/tropicalmed7100322.
- 2.- Rodríguez-Morales AJ, Villamil-Gómez WE, Franco-Paredes C. The arboviral burden of disease caused by co-circulation and co-infection of dengue, chikungunya and Zika in the Americas. *Travel Med Infect Dis* 2016;14:177-9. doi: 10.1016/j.tmaid.2016.05.004.
- 3.- Alarcón-Elbal PM, Rodríguez-Sosa MA, Newman BC, Sutton WB. The first record of *Aedes vittatus* (Diptera: Culicidae) in the Dominican Republic: Public health implications of a potential invasive mosquito species in the Americas. *J Med Entomol* 2020;57:2016-21. doi: 10.1093/jme/tjaa128.
- 4.- Paz-Bailey G, Adams LE, Deen J, Anderson KB, Katzelnick LC. Dengue. *Lancet* 2024;403:667-82. doi:https://doi.org/10.1016/S0140-6736(23)02576-X
- 5.- Long H, Zhang C, Chen C, Tang J, Zhang B, Wang Y, et al. Assessment of the global circulation and endemicity of dengue. *Transbound Emerg Dis* 2022;69:2148-55. doi: 10.1111/tbed.14211.
- 6.- Huits R, Angelo KM, Amatya B, Barkati S, Barnett ED, Bottieau E, et al. Clinical characteristics and outcomes among travelers with severe dengue : a geosentinel analysis. *Ann Intern Med* 2023;176:940-8. doi: 10.7326/M23-0721
- 7.- Angelin M, Sjölin J, Kahn F, Hedberg AL, Rosdahl A, Skorup P, et al. Qdenga® - A promising dengue fever vaccine; can it be recommended to non-immune travelers? *Travel Med Infect Dis* 2023;54:102598. doi: 10.1016/j.tmaid.2023.102598.
- 8.- Perez F, Llau A, Gutierrez G, Bezerra H, Coelho G, Ault S, et al. The decline of dengue in the Americas in 2017: discussion of multiple hypotheses. *Trop Med Int Health* 2019;24:442-53. doi: 10.1111/tmi.13200
- 9.- Blacksell SD, Jarman RG, Bailey MS, Tanganuchitcharnchai A, Jenjaroen K, Gibbons RV, et al. Evaluation of six commercial point-of-care tests for diagnosis of acute dengue infections: the need for combining NS1 antigen and IgM/IgG antibody detection to achieve acceptable levels of accuracy. *Clin Vaccine Immunol* 2011;18:2095-101. doi: 10.1128/CVI.05285-11.
- 10.- Clarke T. Dengue virus: break-bone fever. *Nature* 2002;416:672-4. doi: 10.1038/416672a
- 11.- Katzelnick LC, Gresh L, Halloran ME, Mercado JC, Kuan G, Gordon A, et al. Antibody-dependent enhancement of severe dengue disease in humans. *Science* 2017;358:929-32. doi: 10.1126/science.aan6836
- 12.- Wang L, Huang AT, Katzelnick LC, Lefrancq M, Coello Escoto A, Duret L, et al. Antigenic distance between primary and secondary dengue infections correlates with disease risk. *Sci Transl Med* 2024;16:eade3259. doi: 10.1126/scitranslmed.adk3259
- 13.- Aggarwal C, Ahmed H, Sharma P, Reddy ES, Nayak K, Singla M, et al. Severe disease during both primary and secondary dengue virus infections in pediatric populations. *Nat Med* 2024;30:670-4. doi: 10.1038/s41591-024-02798-x
- 14.- Garcia-Rejon JE, Navarro JC, Cigarroa-Toledo N, Baak-Baak CM. An updated review of the invasive *Aedes albopictus* in the Americas; geographical distribution, host feeding patterns, arbovirus infection, and the potential for vertical transmission of dengue virus. *Insects* 2021;12. doi: 10.3390/insects12110967
- 15.- Bargielowski IE, Lounibos LP, Shin D, Smartt CT, Carrasquilla MC, Henry A, et al. Widespread evidence for interspecific mating between *Aedes aegypti* and *Aedes albopictus* (Diptera: Culicidae) in nature. *Infect Genet Evol* 2015;36:456-61. doi: 10.1016/j.meegid.2015.08.016
- 16.- Lühken R, Brattig N, Becker N. Introduction of invasive mosquito species into Europe and prospects for arbovirus transmission and vector control in an era of globalization. *Infect Dis Poverty* 2023;12:109. doi: 10.1186/s40249-023-01167-z.
- 17.- Silva SOF, de Mello CF, Campos J, Leite PJ, Sabino R, Alencar J. Report of mosquito vectors of arboviruses from a federal conservation unit in the atlantic forest, Rio de Janeiro State, Brazil. *Life (Basel)* 2022;12. doi: 10.3390/life12101597.
- 18.- Akanda AS, Johnson K, Ginsberg HS, Couret J. Prioritizing water security in the management of vector-borne diseases: lessons from Oaxaca, Mexico. *Geohealth* 2020;4:e2019GH000201. doi: 10.1029/2019GH000201
- 19.- Musso D, Rodríguez-Morales AJ, Levi JE, Cao-Lormeau VM, Gubler DJ. Unexpected outbreaks of arbovirus infections: lessons learned from the Pacific and tropical America. *Lancet Infect Dis* 2018;18:e355-e61. doi: 10.1016/S1473-3099(18)30269-X.
- 20.- Culquichicon C, Cardona-Ospina JA, Patiño-Barbosa AM, Rodríguez-Morales AJ. Bibliometric analysis of Oropouche research: impact on the surveillance of emerging arboviruses in Latin America. *F1000Res* 2017;6:194. doi: 10.12688/f1000research.10936.2
- 21.- Rodríguez-Morales AJ, Paniz-Mondolfi AE, Villamil-Gomez WE, Navarro JC. Mayaro, Oropouche and Venezuelan Equine Encephalitis viruses: following in the footsteps of Zika? *Travel Med Infect Dis* 2017;15:72-3. doi: 10.1016/j.tmaid.2016.11.001
- 22.- Arenivar C, Rodriguez Y, Rodríguez-Morales AJ, Anaya JM. Osteoarticular manifestations of Mayaro virus infection. *Curr Opin Rheumatol* 2019;31:512-6. doi: 10.1097/BOR.0000000000000635
- 23.- Patiño-Barbosa AM, Bedoya-Arias JE, Cardona-Ospina JA, Rodríguez-Morales AJ. Bibliometric assessment of the scientific production of literature regarding Mayaro. *J Infect Public Health* 2016;9:532-4. doi: 10.1016/j.jiph.2015.10.001
- 24.- Pezzi L, Rodríguez-Morales AJ, Reusken CB, Ribeiro GS, LaBeaud AD, Lourenco de Oliveira R, et al. GloPID-R report on chikungunya, o'nyong-nyong and Mayaro virus, part 3: epidemiological distribution of Mayaro virus. *Antiviral Res* 2019;172:104610. doi: 10.1016/j.antiviral.2019.104610.
- 25.- Forato J, Meira CA, Claro IM, Amorim MR; de Souza GF, Murano SP, et al. Molecular epidemiology of Mayaro virus among febrile patients, Roraima State, Brazil, 2018-2021. *Emerg Infect Dis* 2024;30:1013-6. doi: 10.3201/eid3005.231406
- 26.- Grisales-Nieto JC, Luz SLB, Alves do Nascimento V, Gomez Naveca F, Murcia-Montañón LM, et al. First case of Oropouche fever detected in the international border region of the Colombian Amazon: clinical characteristics and molecular diagnosis. *Mem Inst Oswaldo Cruz* 2024;119:e230221. doi: 10.1590/0074-02760230221.
- 27.- Acosta-España JD, Dueñas-Espín I, Grijalva Narvaez DF, Altamirano-Jara JB, Gómez-Jaramillo AM, Rodríguez-Morales AJ. Analysis of inpatient data on dengue fever, malaria and leishmaniasis in Ecuador: A cross-sectional national study, 2015-2022. *New Microbes and New Infections* 2024;101421. doi:10.1016/j.nmni.2024.101421
- 28.- Souza Chaves TS, Orduna T, Lepetic A, Macchi A, Verbanaz S, Risquez A, et al. Yellow fever in Brazil: Epidemiological aspects and implications for travelers. *Travel Med Infect Dis* 2018;23:1-3. doi: 10.1016/j.tmaid.2018.05.001.
- 29.- Ortiz-Martínez Y, Patiño-Barbosa AM, Rodríguez-Morales AJ. Yellow fever in the Americas: the growing concern about new epidemics. *F1000Res* 2017;6:398. doi: 10.12688/f1000research.11280.2
- 30.- Reno E, Quan NG, Franco-Paredes C, Chastain DB, Chauhan L, Rodríguez-Morales AJ, et al. Prevention of yellow fever in travellers: an update. *Lancet Infect Dis* 2020;20:e129-e37. doi: 10.1016/S1473-3099(20)30170-5
- 31.- Rifakis PM, Benitez JA, De-la-Paz-Pineda J, Rodríguez-Morales AJ. Epizootics of

- yellow fever in Venezuela (2004-2005): an emerging zoonotic disease. *Ann N Y Acad Sci* 2006;1081:57-60. doi: 10.1196/annals.1373.005
- 32.- Rodríguez-Morales AJ, Bonilla-Aldana DK, Suarez JA, Franco-Paredes C, Forero-Peña DA, Mattar S, et al. Yellow fever reemergence in Venezuela - Implications for international travelers and Latin American countries during the COVID-19 pandemic. *Travel Med Infect Dis* 2021;44:102192. doi: 10.1016/j.tmaid.2021.102192.
- 33.- Silva-Ramos CR, Montoya-Ruiz C, Faccini-Martínez Á A, Rodas JD. An updated review and current challenges of Guanarito virus infection, Venezuelan hemorrhagic fever. *Arch Virol* 2022;167:1727-38. doi: 10.1007/s00705-022-05453-3.
- 34.- Silva-Ramos CR, Faccini-Martínez Á A, Calixto OJ, Hidalgo M. Bolivian hemorrhagic fever: a narrative review. *Travel Med Infect Dis* 2021;40:102001. doi: 10.1007/s00705-022-05453-3.
- 35.- Rodríguez-Morales AJ, Bonilla-Aldana DK, Rísquez A, Paniz-Mondolfi A, Suárez JA. Should we be concerned about Venezuelan hemorrhagic fever? - A reflection on its current situation in Venezuela and potential impact in Latin America amid the migration crisis. *New Microbes New Infect* 2021;44:100945. doi: 10.1016/j.nmni.2021.100945
- 36.- Nastri AC, Duarte-Neto AN, Casadio LVB, Maciel de Souza W, Claro IM, Manuli ER, et al. Understanding Sabiá virus infections (Brazilian mammarenavirus). *Travel Med Infect Dis* 2022;48:102351. doi: 10.1016/j.tmaid.2022.102351.
- 37.- Sarute N, Ross SR. New World arenavirus biology. *Annu Rev Virol* 2017;4:141-58. doi: 10.1146/annurev-virology-101416-042001.
- 38.- Cardona-Ospina JA, Arteaga-Livias K, Villamil-Gomez WE, Pérez-Díaz CE, Bonilla-Aldana DK, Mondragón-Aldona A, et al. Dengue and COVID-19, overlapping epidemics? An analysis from Colombia. *J Med Virol* 2021;93:522-7. doi: 10.1002/jmv.26194
- 39.- Leon-Figueroa DA, Abanto-Urbano S, Olarte-Durand M, Nuñez-Lupaca JN, Barboza JJ, Bonilla-Aldana DK, et al. COVID-19 and dengue coinfection in Latin America: A systematic review. *New Microbes New Infect* 2022;49:101041. doi: 10.1016/j.nmni.2022.101041.
- 40.- Navarro JC, Arrivillaga-Henriquez J, Salazar-Loor J, Rodríguez-Morales AJ. COVID-19 and dengue, co-epidemics in Ecuador and other countries in Latin America: Pushing strained health care systems over the edge. *Travel Med Infect Dis* 2020;37:101656. doi: 10.1016/j.tmaid.2020.101656.
- 41.- Patwary MM, Haque MZ, Bardhan M, Rodríguez-Morales AJ. COVID-19 and Dengue co-epidemic during the second wave of the pandemic in Bangladesh: a double blow for an overburdened health-care system. *Disaster Med Public Health Prep* 2022;1-3. doi: 10.1017/dmp.2022.105
- 42.- Plasencia-Duenas R, Failoc-Rojas VE, Rodríguez-Morales AJ. Impact of the COVID-19 pandemic on the incidence of dengue fever in Peru. *J Med Virol* 2022;94:393-8. doi: 10.1002/jmv.27298
- 43.- Dyatkin B. COVID-19 pandemic highlights need for US policies that increase supply chain resilience. *MRS Bull* 2020;45:794-6. doi: 10.1557/mrs.2020.258
- 44.- Islam Z, Mohanan P, Bilal W, Hashmi T, Rahmat Z, Abdi I, et al. Dengue virus cases surge amidst COVID-19 in Pakistan: Challenges, efforts and recommendations. *Infect Drug Resist* 2022;15:367-71. doi: 10.2147/IDR.S347571
- 45.- Tian N, Zheng JX, Guo ZY, Li LH, Xia S, LV S, et al. Dengue Incidence trends and its burden in major endemic regions from 1990 to 2019. *Trop Med Infect Dis* 2022;7. doi: 10.3390/tropicalmed7080180
- 46.- Malavige GN, Sjö P, Singh K, Piedagnel JM, Mowbray C, Estani S, et al. Facing the escalating burden of dengue: challenges and perspectives. *PLOS Glob Public Health* 2023;3:e0002598. doi: 10.1371/journal.pgph.0002598.
- 47.- Patikorn C, Cho JY, Higashi J, Huang XX, Chaikayunapruk N. Financial hardship among patients suffering from neglected tropical diseases: A systematic review and meta-analysis of global literature. *PLoS Negl Trop Dis* 2024;18:e0012086. doi: 10.1371/journal.pntd.0012086
- 48.- Yitbarek S, Chen K, Celestin M, McCary M. Urban mosquito distributions are modulated by socioeconomic status and environmental traits in the USA. *Ecol Appl* 2023;33:e2869. doi: 10.1002/eap.2869
- 49.- Mulligan K, Dixon J, Sinn CL, Elliott SJ. Is dengue a disease of poverty? A systematic review. *Pathog Glob Health* 2015;109:10-8. doi: 10.1179/2047773214Y.0000000168.
- 50.- Telle O, Nikolay B, Kumar V, Binkimoun S, Pal R, Nagpal BN, et al. Social and environmental risk factors for dengue in Delhi city: a retrospective study. *PLoS Negl Trop Dis* 2021;15:e0009024. doi: 10.1371/journal.pntd.0009024
- 51.- Hossan J, Hasan S, Patwary MM, Bardhan M, Imran SA, Rodríguez-Morales AJ. Mapping hotspots and cluster analysis of the current dengue outbreak in Bangladesh. *New Microbes New Infect* 2023;55:101190. doi: 10.1016/j.nmni.2023.101190
- 52.- Zambrano LI, Sierra M, Lara B, Rodríguez-Núñez I, Medina MT, Lozada- Riascos CO, et al. Estimating and mapping the incidence of dengue and chikungunya in Honduras during 2015 using Geographic Information Systems (GIS). *J Infect Public Health* 2017;10:446-56. doi: 10.1016/j.jiph.2016.08.003.
- 53.- de Sousa SC, Carneiro M, Eiras AE, Bezerra JMT, Barbosa DS. Factors associated with the occurrence of dengue epidemics in Brazil: a systematic review. *Rev Panam Salud Publica* 2021;45:e84. doi: 10.26633/RPSP.2021.84.
- 54.- Yang L, Norris EJ, Jiang S, Bernier UR, Linthicum KJ, Bloomquist JR. Reduced effectiveness of repellents in a pyrethroid-resistant strain of *Aedes aegypti* (Diptera: culicidae) and its correlation with olfactory sensitivity. *Pest Manag Sci* 2020;76:118-24. doi: 10.1002/ps.5562.
- 55.- Semenza JC, Suk JE. Vector-borne diseases and climate change: a European perspective. *FEMS Microbiol Lett* 2018;365. doi: 10.1093/femsle/fnx244
- 56.- Fonseca SNS. Changing epidemiology of dengue fever in children in South America. *Curr Opin Pediatr* 2023;35:147-54. doi: 10.1097/MOP.0000000000001220
- 57.- Osman S, Preet R. Dengue, chikungunya and Zika in GeoSentinel surveillance of international travellers: a literature review from 1995 to 2020. *J Travel Med* 2020;27. doi: 10.1093/jtm/taaa222.
- 58.- Zambrano LI, Fuentes-Barahona IC, Portillo-Pineda R, Aguilar-Ponce M, Murillo-Padilla JC, Suazo-Menocal M, et al. Assessment of post-dengue rheumatic symptoms using the WOMAC and DAS-28 questionnaires in a Honduran population after a four-month follow-up. *Trop Med Infect Dis* 2022;7. doi: 10.3390/tropicalmed7120394
- 59.- Childs ML, Lyberger K, Harris M, Burke M, Mordecai EA. Climate warming is expanding dengue burden in the Americas and Asia. *medRxiv* 2024. doi: 10.1101/2024.01.08.24301015
- 60.- Ladner J, Rodrigues M, Davis B, Besson MH, Audureau E, Saba J. Societal impact of dengue outbreaks: Stakeholder perceptions and related implications. A qualitative study in Brazil, 2015. *PLoS Negl Trop Dis* 2017;11:e0005366. doi: 10.1371/journal.pntd.0005366
- 61.- Bonna AS, Pavel SR, Mehjabin T, Ali M. Dengue in Bangladesh. *Int J Infect Dis* 2023. doi: 10.1016/j.ijid.2023.06.020.
- 62.- Hladish TJ, Pearson CAB, Rojas DP, Gomez-Dantes H, Halloran ME, Vasquez-Prokopec GM, et al. Forecasting the effectiveness of indoor residual spraying for reducing dengue burden. *PLoS Negl Trop Dis* 2018;12:e0006570. doi: 10.1371/journal.pntd.0006570
- 63.- Agbo S, Gbaguidi L, Biliyar C, Sylla S, Fahnbulleh M, Dogba J, et al. Establishing national multisectoral coordination and collaboration mechanisms to prevent, detect, and respond to public health threats in Guinea,

- Liberia, and Sierra Leone 2016-2018. One Health Outlook 2019;1:4. doi: 10.1186/s42522-019-0004-z.
- 64.- Lawson B, Neimanis A, Lavazza A, López-Olvera JR, Tavernier P, Billinis C, et al. How to start up a National Wildlife Health Surveillance Programme. Animals (Basel) 2021;11. doi: 10.3390/ani11092543.
 - 65.- Srisawat N, Thisyakorn U, Ismail Z, Rafiq K, Gubler DJ. World Dengue Day: A call for action. PLoS Negl Trop Dis 2022;16:e0010586. doi: 10.1371/journal.pntd.0010586.
 - 66.- Aliota MT, Peinado SA, Velez ID, Osorio JE. The wMel strain of Wolbachia reduces transmission of Zika virus by *Aedes aegypti*. Sci Rep 2016;6:28792. doi: 10.1038/srep28792.
 - 67.- Aliota MT, Walker EC, Uribe Yepes A, Velez ID, Christensen BM, Osorio JE. The wMel strain of Wolbachia reduces transmission of chikungunya virus in *Aedes aegypti*. PLoS Negl Trop Dis 2016;10:e0004677. doi: 10.1371/journal.pntd.0004677
 - 68.- Velez ID, Santacruz E, Kutcher SC, Duque SL, Uribe A, Barajas J, et al. The impact of city-wide deployment of Wolbachia-carrying mosquitoes on arboviral disease incidence in Medellín and Bello, Colombia: study protocol for an interrupted time-series analysis and a test-negative design study. F1000Res 2019;8:1327. doi: 10.12688/f1000research.19858.2.
 - 69.- Velez ID, Tanamas SK, Arbelaez MP, Kutcher SC, Duque SL, Uribe A, et al. Reduced dengue incidence following city-wide wMel Wolbachia mosquito releases throughout three Colombian cities: Interrupted time series analysis and a prospective case-control study. PLoS Negl Trop Dis 2023;17:e0011713. doi: 10.1371/journal.pntd.0011713
 - 70.- Velez ID, Uribe A, Barajas J, Uribe S, Ángel S, Suazo-Vasco JD, et al. Large-scale releases and establishment of wMel Wolbachia in *Aedes aegypti* mosquitoes throughout the Cities of Bello, Medellín and Itagüí, Colombia. PLoS Negl Trop Dis 2023;17:e0011642. doi: 10.1371/journal.pntd.0011642.
 - 71.- Cabrera M, Taylor G. Modelling spatio-temporal data of dengue fever using generalized additive mixed models. Spat Spatiotemporal Epidemiol 2019;28:1-13. doi: 10.1016/j.sste.2018.11.006
 - 72.- Mamenun, Koesmaryono Y, Sopaheluwakan A, Hidayati R, Dasanto BD, Aryati R. Spatiotemporal characterization of dengue incidence and its correlation to climate parameters in Indonesia. Insects 2024;15:366. <https://doi.org/10.3390/insects15050366>
 - 73.- Coronel-Martínez DL, Park J, López-Medina E, Capeding MR, Cadena Bonfanti AA, Montalbán MC, et al. Immunogenicity and safety of booster CYD-TDV dengue vaccine after alternative primary vaccination schedules in healthy individuals aged 9-50 years: a randomised, controlled, phase 2, non-inferiority study. Lancet Infect Dis 2022;22:901-11. doi: 10.1016/S1473-3099(21)00706-4
 - 74.- López-Medina E, Biswal S, Saez-Llorens X, Borja-Tabora C, Bravo L, Sirivichayakul C, et al. Efficacy of a dengue vaccine candidate (TAK-003) in healthy children and adolescents 2 years after Vaccination. J Infect Dis 2022;225:1521-32. doi: 10.1093/infdis/jiaa761.
 - 75.- Rivera L, Biswal S, Saez-Llorens X, López-Medina E, Borja-Tabora C, Bravo L, et al. Three-year efficacy and safety of Takeda's dengue vaccine candidate (TAK-003). Clin Infect Dis 2022;75:107-17. doi: 10.1093/cid/ciab864.
 - 76.- Sirivichayakul C, Biswal S, Saez-Llorens X, et al. Efficacy and safety of a tetravalent dengue vaccine (TAK-003) in children with prior Japanese Encephalitis or Yellow Fever vaccination. J Infect Dis 2024. doi: 10.1093/infdis/jiae222
 - 77.- Tricou V, Yu D, Reynales H, Biswal S, Saez-Llorens X, Sirivichayakul. Long-term efficacy and safety of a tetravalent dengue vaccine (TAK-003): 4.5-year results from a phase 3, randomised, double-blind, placebo-controlled trial. Lancet Glob Health 2024;12:e257-e70. doi: 10.1016/S2214-109X(23)00522-3.
 - 78.- Ylade M, Crisostomo MV, Daag JV, Agrupis KA, López AL. Effect of single-dose, live, attenuated dengue vaccine in children with or without previous dengue on risk of subsequent, virologically confirmed dengue in Cebu, the Philippines: a longitudinal, prospective, population-based cohort study. Lancet Infect Dis 2024. doi:[https://doi.org/10.1016/S1473-3099\(24\)00099-9](https://doi.org/10.1016/S1473-3099(24)00099-9)
 - 79.- Herrera-Martínez AD, Rodríguez-Morales AJ. Potential influence of climate variability on dengue incidence registered in a western pediatric Hospital of Venezuela. Trop Biomed 2010;27:280-6. PMID: 20962726
 - 80.- Mattar S, Morales V, Cassab A, Rodríguez-Morales AJ. Effect of climate variables on dengue incidence in a tropical Caribbean municipality of Colombia, Cereté, 2003-2008. Int J Infect Dis 2013;17:e358-9. doi: 10.1016/j.ijid.2012.11.021.
 - 81.- Ritu MR, Sikder D, Patwary MM, Tamim AR, Rodríguez-Morales AJ. Climate change, urbanization and resurgence of dengue in Bangladesh. New Microbes New Infect 2024;59:101414. doi: 10.1016/j.nmni.2024.101414.
 - 82.- Zambrano LI, Sevilla C, Reyes-García SZ, Kafati R, Rodríguez-Morales AJ, Mattar S. Potential impacts of climate variability on dengue hemorrhagic fever in Honduras, 2010. Trop Biomed 2012;29: 499-507. <https://citeseerx.ist.psu.edu/document?repid=rep1&type=pdf&doi=122c7177f2099bccc71492de41b4548d48effbe1>