

Ruwandi Kariyawasam¹, Katherine Tan², Min Qun Chen³, Ruben Cuidamat³, Rachel Lau³, Filip Ralevski³, Andrea K. Boggild^{1,3,4,5}

Introduction

- Flaviviruses: transmitted to humans through infected bites of *Culex* and *Aedes* mosquitoes¹
- Plasmodium ovale*: spread by the bite of *Anopheles* mosquito²
- Flaviviral infection could precipitate a *P. ovale* relapse³
- Given overlap of epidemiological and clinical presentations of both flaviviral and malaria infections, diagnostic testing where malaria is confirmed or excluded, without subsequent flaviviral testing may mask true epidemiology of co-infections⁴

Objective: We aim to understand the incidence of intercurrent flaviviral infection in confirmed *Plasmodium ovale* infection in whole blood specimens from ill returned travelers

Methods

- DNA extracted from whole blood specimens and tested for malaria by microscopy and rapid diagnostic test (RDT) between 2006 and 2019 at Public Health Ontario Laboratory-Toronto^{5,6}
- RNA extracted from *P. ovale* positive whole blood specimens and examined by real-time PCR (qPCR) for the following targets: flaviviruses (pan-FLAV) and dengue virus types 1-4 (DEN1, DEN2, DEN3, DEN4)⁷⁻⁹

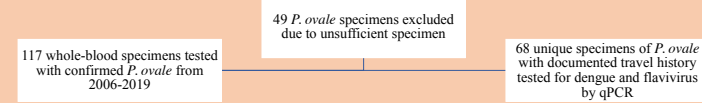


Figure 1: Workflow highlighting *P. ovale* confirmed diagnostic testing for intercurrent flaviviral infection using qPCR.

Results

Table 1: Clinical and parasitological characteristics of *P. ovale* cases

| | Total [n=68 (%)] |
|-------------------------------|-----------------------------------|
| Median Age, years (range) | 27.4 years (22 months – 72 years) |
| Median Parasitemia, % (range) | <0.01 % (< 0.01 % - 0.8 %) |
| Sex | |
| Male | 37 (54.4 %) |
| Female | 30 (44.1 %) |
| Unknown | 3 (4.4%) |
| Travel History | |
| Yes | 39 (57.4%) |
| Unknown | 29 (42.6%) |

Table 2: Top countries of travel

| | Total [n=68 (%)] |
|---------------------|------------------|
| Nigeria | 23 (33.8%) |
| Africa, unspecified | 5 (7.4%) |
| Congo | 3 (4.8%) |
| Tanzania | 3 (4.8%) |

Table 3: DENV and Flavivirus qPCR positive results

| DENV qPCR | Flavivirus qPCR |
|-----------|----------------------------|
| 0/68 (0%) | 1/68 (1.5 %) untypeable |

Discussion & Conclusions

- Pan-FLAV yielded a 1.5% positivity rate (1/68), while the DENV assays did not yield a positive result (Table 3)
- P. ovale* infections are commonly imported to Ontario from West Africa, especially from Nigeria which was our top country of travel (Table 2)
- Intercurrent flaviviral viremia was noted in 1.5% of specimens, suggesting that primary flaviviral infection could have triggered a relapse of *P. ovale*
- Alternatively, co-infection may suggest primary infection with both organisms given the overlap of vector populations in these endemic areas
- Consideration of flaviviral co-infection should be given to *P. ovale* patients to appropriately manage clinical manifestations including deep thrombocytopenia, lymphopenia, and high yield arboviral symptomatology including rash and retro-orbital headache³

References

- Huang YJS, Higgs S, Horne KM, Vanlandingham D. Flavivirus-Mosquito Interactions. *Viruses* 2014; 6(11): 4703-4730.
- Anantabotla VM, Antony HA, Parija SC, Rajkumari N, Kini JR, Manipura R, et al. Polymorphisms in genes associated with drug resistance of *Plasmodium vivax* in India. *Parasitol Int*. 2019; 70: 92-97
- Kariyawasam R, Lecce C, Tan K, Boggild AK. Don't forget co-infections! A case of intercurrent *Plasmodium vivax* and dengue infection in a traveler to India and the Caribbean. *Travel Med Infect Dis*. 2019.
- Lupi O, Ridolfi F, da Silva S, Zanini GM, Lavigne A, Nogueira RM, da Cruz Mde F, Daniel-Ribeiro CT, Brasil P. Dengue infection as a potential trigger of an imported *Plasmodium ovale* malaria relapse or a long incubation period in a non-endemic malaria region. *Int J Infect Dis* 2016; 44: 20-4.
- Phuong M, Lau R, Ralevski F, Boggild AK. Sequence-based optimization of a quantitative real-time PCR assay for detection of *Plasmodium ovale* and *Plasmodium malariae*. *J Clin Microbiol*. 2014;52:1068-73. <http://dx.doi.org/10.1128/JCM.03477-13.20>
- Phuong M, Lau R, Ralevski F, Boggild AK. Survival analysis of diagnostic assay in *Plasmodium falciparum* malaria. *Malar J* 2015; 14: 350.
- Kariyawasam R, Lau R, Eshaghi A, Patel SN, Sider D, Gubbay JB, Boggild AK. Spectrum of Viral Pathogens in Blood of Malaria-Free Ill Travelers Returning to Canada. *Emerg Infect Dis* 2016; 22(5): 854-861.
- Balm MN, Lee CK, Lee HK, Chiu L, Koay ES, Tang JW. A diagnostic polymerase chain reaction assay for Zika virus. *J Med Virol* 2012; 84(9): 1501-5.
- Pongsiri P, Praiantantavorn K, Theamboonlers A, Payungporn S, Poovorawan Y. Multiplex real-time RT-PCR for detecting chikungunya virus and dengue virus. *Asian Pac J Trop Med* 2012; 5(5): 345-6.
- Price RN, Tjitra E, Guerra CA, Yeung S, White NJ, Anstey NM. *Vivax Malaria: Neglected and Not Benign*. *Am J Trop Med Hyg*. 2007;77:6.