

## UNIVERSITY OF TORONTO

### Microbiology & Infectious Diseases Research Days

Monday, June 3rd, 2019 – Trainee Day (Selected from Abstracts)

Tuesday, June 4th, 2019 – Invited Lectures & Poster Session

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**Talks in Medical Sciences Building, Room 2170**

**Posters & Lunch in Medical Sciences Building,  
Room 2171 (C. David Naylor Student Commons)**

**Website:** <http://microbeto.ca/mid-2019/>

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#### Monday, June 3<sup>rd</sup>, 2019

##### **9:30 - 9:40 WELCOME ADDRESS**

##### **9:45 – 10:00: Avid Mohammadi**

Characterizing the impact of penile-vaginal sex on HIV-susceptible CD4<sup>+</sup> T cell subsets in the female genital tract

##### **10:05 - 10:20: Erin O. Y. Wong**

Developing defined microbiota to model inflammation in the mouse gut

##### **10:25 - 10:40: Nora Mellouk**

An ATG16L1-dependent pathway promotes plasma membrane repair and limits *Listeria monocytogenes* cell-to-cell spread

##### **10:45 - 11:15: COFFEE BREAK**

##### **11:20 - 11:35: Jean-Paul R. Soucy**

Joint modelling of resistance to six antimicrobials in urinary *Escherichia coli* isolates in Quebec, Canada

##### **11:40 – 11:55: Sarah Birstonas**

EHEC utilizes two-component systems to modulate expression of major flagellar subunit protein, FliC, in response to host intestinal cues

##### **12:00 - 12:15: Nathaniel Winsor**

NLRP6 regulates the colonic mucus layer during *Trichomonas* infection

##### **12:35 – 1:30: LUNCH**

##### **1:35 - 12:50: Samuel Salamun**

Epstein-Barr Virus Protein BMRF1 Modulates Cellular SUMO and DNA Damage Response Pathways by Binding the Cellular NuRD Complex

##### **1:55 - 2:10: Nicola Case**

Elucidating the mechanism of *Candida albicans* morphogenesis in response to phagocytosis by macrophages

##### **2:15 - 2:30: Sarah Kronheim**

A small molecule anti-phage defense mechanism in *Streptomyces*

##### **2.30 - 3:00: COFFEE BREAK**

##### **3:05 - 3:20: Alexandra Willis**

Understanding inherited immunity using a *C. elegans* model of microsporidia infection

##### **3:25 - 3:40: Genevieve Mailhot**

Differentiating between protective and pathogenic neutrophil responses during *Neisseria gonorrhoeae* infection

##### **3:45 – 4:00: Tiffany Fitzpatrick**

Successes of anti-RSV prophylaxis among infants in Ontario: results from a multi-decade, population-based controlled interrupted time series analysis using health administrative data

# Poster Presentations

## 80) Intercurrent Flaviviral Viremia in Ill Returned Travelers with *Plasmodium vivax* Malaria

Ruwandi Kariyawasam, Katherine Tan, Min Qun Chen, Ruben Cuidamat, Aisha Khatib, Rachel Lau, Filip Ralevski, Andrea K. Boggild

Institute of Medical Sciences, University of Toronto; Human Biology, University of Toronto; Public Health Ontario Laboratories; Tropical Disease Unit, UHN-Toronto General Hospital; Department of Medicine, University of Toronto

**Background:** Similar epidemiology and clinical presentations of arboviral infections and malaria coupled with the typically sequential approach to diagnostic testing, where malaria is confirmed or excluded urgently in febrile returned travelers, may mask the true epidemiology of co-infections. Flaviviruses are known to trigger relapsing forms of malaria, including *Plasmodium vivax*, long after primary malaria infection, and this may delay the diagnosis of malaria.

**Objective:** We aim to understand the incidence of intercurrent flaviviral infection in confirmed *Plasmodium vivax* infection.

**Method:** DNA and RNA from biobanked isolates of *P. vivax* detected in whole blood at the Public Health Ontario Laboratory between 2006 and 2019 were extracted and screened for intercurrent flaviviral infections using previously validated real-time PCR (qPCR) assays targeting multiple flaviviruses (pan-FLAV) and, specifically, dengue virus types 1-4 (DEN1, DEN2, DEN3, DEN4).

**Results:** Five hundred and two unique isolates of *P. vivax* were identified, of which 175 have been tested to date. Males accounted for 69.1% (n=121/175) of *P. vivax* cases, while females accounted for 28.6% (n=50/175), and sex was unassigned in 2.3% (4/175). Median age of *P. vivax* cases was 34.2 years (range 3.7 years – 87.6 years; IQR 24.0 – 51.9 years). Median parasitemia was 0.1% (range < 0.01% - 1.1%). Sixty-eight (38.9%) *P. vivax* cases had documented travel history exclusively to South Asia, with India as the most common source country (34/175 [19.4%]). Pan-FLAV assay yielded a 0.6% (1/175) positivity rate. DENV assay yielded a 0.6% (1/175) positivity rate. Type-specific real-time PCR revealed DEN1, which was also detected on both Pan-FLAV and pan-DENV assays.

**Conclusion:** Intercurrent flaviviral viremia, was noted in at least 0.6%, which may suggest that primary flaviviral infection, in this case, DEN1, triggered a relapse of *P. vivax*. Alternatively, such co-occurrence may suggest primary infection with both organisms known to cause fever in returning travelers. Consideration of flaviviral coinfection should be given to the *P. vivax* patient with deep thrombocytopenia, lymphopenia, and high-yield arboviral symptomatology such as rash and retro-orbital headache.