

COMPARATIVE INVESTIGATION OF *PLASMODIUM VIVAX* IN BONE MARROW AND PERIPHERAL BLOOD SAMPLES FROM INDIVIDUALS NATURALLY EXPOSED TO MALARIA *VIVAX* IN BRAZILIAN AMAZON

Jéssica RS Alves^{1#}, João L Silva-Filho^{2#}, Anne CG Almeida⁴, João PC Tavares¹, Juliana A Leite¹, Marcelo Brito⁴, Italo FA Antunes⁴, Adriana PB Lopes⁴, Cindy BS Rodrigues⁴, Daniel GQ Ribeiro⁴, Vitória GG de Siqueira⁴, Marcia VG Vallejos⁴, Pedro AC Costa¹, Letusa Albrecht³, Marcus VG Lacerda⁴, Erich V de Paula⁵, Gisely C Melo⁴, Stefanie CP Lopes⁷, Matthias Marti^{2,6*} and Fabio TM Costa^{1*}.

¹Laboratory of Tropical Diseases – Prof. Dr. Luiz Jacintho da Silva, Department of Genetics, Evolution, Microbiology and Immunology, Institute of Biology, University of Campinas, Campinas, Brazil; ²Wellcome Centre for Integrative Parasitology, Institute of Infection, Immunity & Inflammation, University of Glasgow, Glasgow, United Kingdom; ³Research laboratory of Apicomplexa, Fiocruz, Paraná, Brazil; ⁴Tropical Medicine Foundation Dr. Heitor Vieira Dourado, Manaus, Brazil; ⁵Department of Clinical Pathology, School of Medical Sciences, University of Campinas, Campinas, Brazil; ⁶Institute of Parasitology Zurich (IPZ), University of Zurich.

⁷Institute Leônidas & Maria Deane, Fiocruz, Manaus, Brazil;

Contributed to this work equally

*Coordinated this work equally

Corresponding authors: jessica.rsalves@hotmail.com and joao.dasilvafilho@glasgow.ac.uk

Background: *Plasmodium vivax* (Pv) has a unique tropism to invade immature reticulocytes. Recent studies revealed significant enrichment of parasite biomass in the bone marrow (BM) and spleen when compared to the peripheral blood (PB). The effects behind this cell tropism in disease progression are poorly understood, hindering progress in understanding Pv immunopathogenesis.

Objective: The present study hypothesizes phenotypic and functional differences between parasite populations developing in the BM versus those in the PB.

Methods: To test this hypothesis, we aim to integrate transcriptome data analyzes and functional assays performed *ex-vivo* with Pv isolated from BM vs. PB reticulocytes. BM aspirates and PB matched samples were collected from uncomplicated Pv cases admitted at FMT-HVD (Manaus-AM).

Results: To date, 26 patients have consented to the collection of matched BM and PB samples. Whole blood, plasma, and mononuclear cells were isolated and cryopreserved. Analysis of BM and blood smears showed no differences in total parasitemia (% of infected red blood cells) and stage distribution between BM and PB. However, analysis of transcriptomic signatures by bulk-RNA-seq, indicated enrichment of sexually committed schizonts (AP2-G+ parasites), rings and immature gametocytes in the BM, compared to blood, during *P. vivax* infection. Upregulation of genes associated with different metabolic pathways, including PC biosynthesis and reticulocyte invasion in BM parasites were also observed. Additionally, functional analysis of rosette formation in 9 patients demonstrated a higher density in BM samples when compared to PB, particularly in those collected during malaria recurrence episode. **Conclusion:** These findings represent a pioneering study to better understand the unique adaptation of *P. vivax* in the BM niche, with implications to malaria diagnosis and treatment as well as strategies towards the malaria elimination agenda.

Keywords: *Plasmodium vivax*, bone marrow and pathogenesis. Financial Support: FAPESP (Processos 2021/04632-8 e 2017/18611-7), Fundação Bill & Melinda Gates (Processo 5597) and

Conselho Nacional de Desenvolvimento Científico e Tecnológico – CNPq (Processo 442946/2019-8).