Future of Malaria Research Symposium
Friday 13 November 2020
Johns Hopkins Malaria Research Institute

Call for Abstracts

Instructions & Format
The submission deadline for short (3 min) or long (10 min) talk consideration is 12 October 2020. Please use the below format and upload your abstract in the following link:
https://forms.gle/TEgfyDPePf3xVjSw5

Please provide title, authors, institution, and a brief abstract of your research (300 words maximum).

Audience: Both researchers and lay persons.
Font: Times New Roman, Type size: 12 pt, Format: MS Word
Title: Bold, capitalizing first word and proper nouns, as well as italicizing when appropriate
Authors: Please use full names, (first name last name), and use numeric symbols to designate institutions
Institution(s): Please order alphabetically and designate with numeric symbols

EXAMPLE
Charaterizing the Anopheles gambiae late-phase immune responses that limit Plasmodium oocyst survival

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Mosquitoes of the genus Anopheles serve as the obligate vectors of the malaria parasite Plasmodium. During its development within the mosquito host, several factors and developmental bottlenecks limit parasite success, including two distinct phases of the mosquito innate immune response. Emerging evidence suggests that parasite numbers are largely influenced by an “early-phase” that targets ookinetes as they reach the basal lamina of the mosquito midgut through the exposure to complement-like components of the hemolymph and a “late-phase” response that limits oocyst survival through the production of nitric oxide mediated by the STAT pathway. Recently, we have identified a novel LITAF-like transcription factor (LL3) in Anopheles gambiae that is an important component of the mosquito response to Plasmodium. Following LL3-silencing, oocyst numbers are significantly increased and preliminary evidence suggests that LL3 is involved in the “late-phase” response by limiting oocyst survival.

Experiments to determine the relationship between the “late-phase” response for LL3 and those previously described for the STAT pathway imply that the LL3 and STAT phenotypes occur through independent, yet closely related mechanisms involving hemocyte function. Current experiments aim to further dissect LL3 function to address a critical gap in our knowledge of the mosquito late-phase immune response and the mechanisms that limit oocyst survival in its mosquito host.

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