The 2nd International Electronic Conference on Entomology

19-21 May 2025 | Online





Plasmodium-mosquito midgut interactions: Role of peritrophins and digestive proteases

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INTRODUCTION AND AIM

- Malaria remains a major global health burden, with transmission intricately dependent on the interactions between *Plasmodium* parasites and the *Anopheles* midgut.
- Post-infected blood meal in 1st 24-hour period is a critical bottleneck, during which gametogenesis, fertilization, ookinete development, and digestion occurs
- Digestion is mediated by proteases and peritrophic containing peritrophin

To identify significantly peritrophins and digestive proteases during the early-stage infection

AIM

To characterize Per16, 44 and 48 & check its potential for transmission blocking

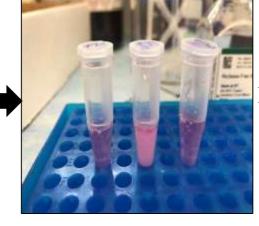
2 METHODS

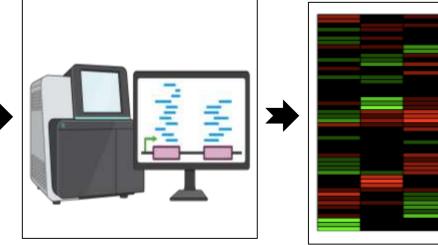


BF-IBF feeding/

Gut Isolation

Molecular Docking

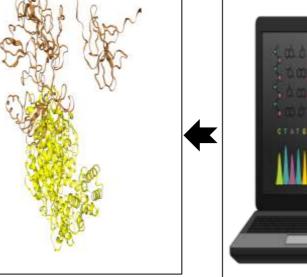


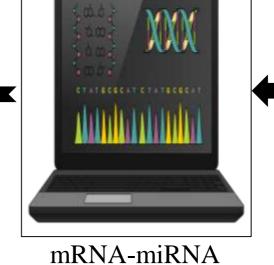


RNA isolation/ Seq Quantification Quar

Sequencing/ Quantification

Data analysis





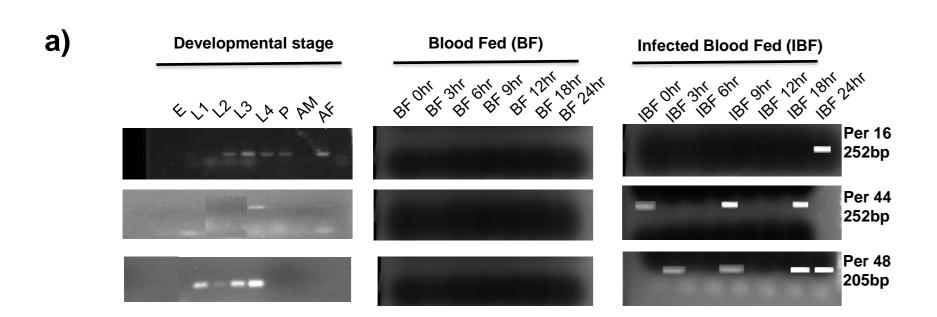
interaction

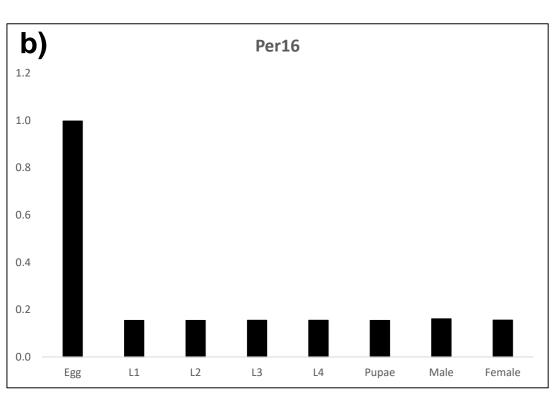
Que course

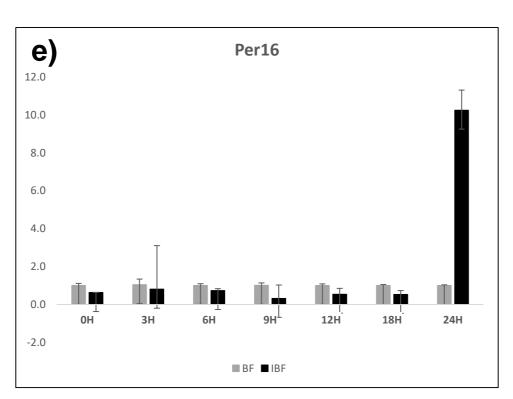
Cloning/ Sequencings

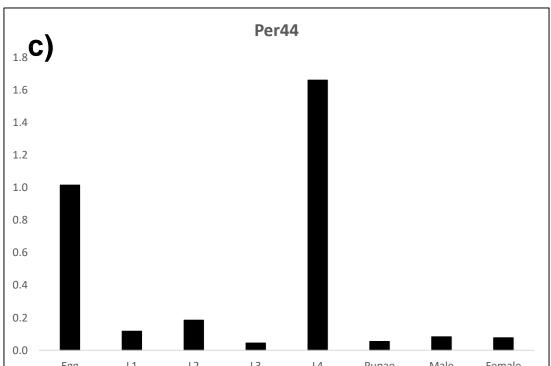
cDNA synthesis/ RT-PCR

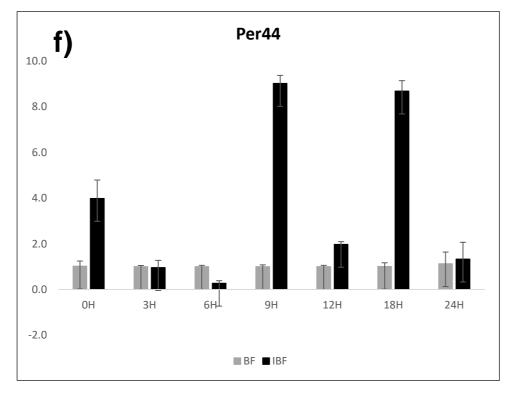
3 RESULTS

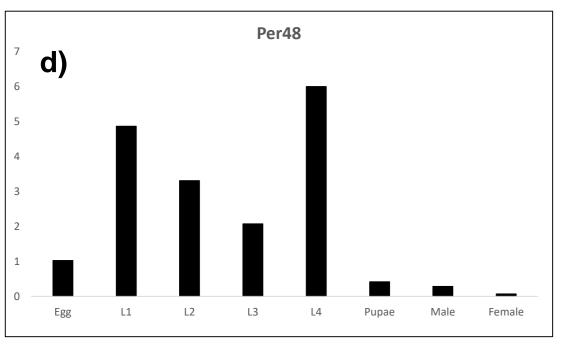












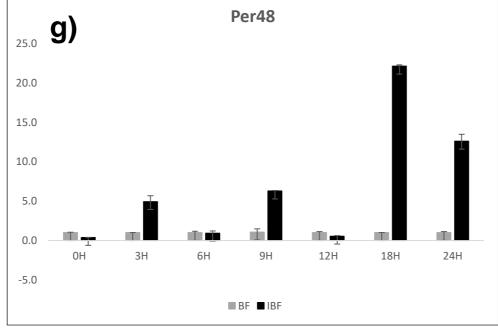


Fig 1: a) Semi-quantitative agarose gel (1.5%) of BF-IBF Per 16(252bp), 44(252bp), 48(205bp), b) Semi-quantitative agarose gel (1.5%) of Developmental stage Per 16-48, c) Developmental stage relative mRNA characterization of Per16, 44, and 48 c-e respectively, d) Bf-IBF relative mRNA characterization of Per16, 44, and 48 f-h respectively

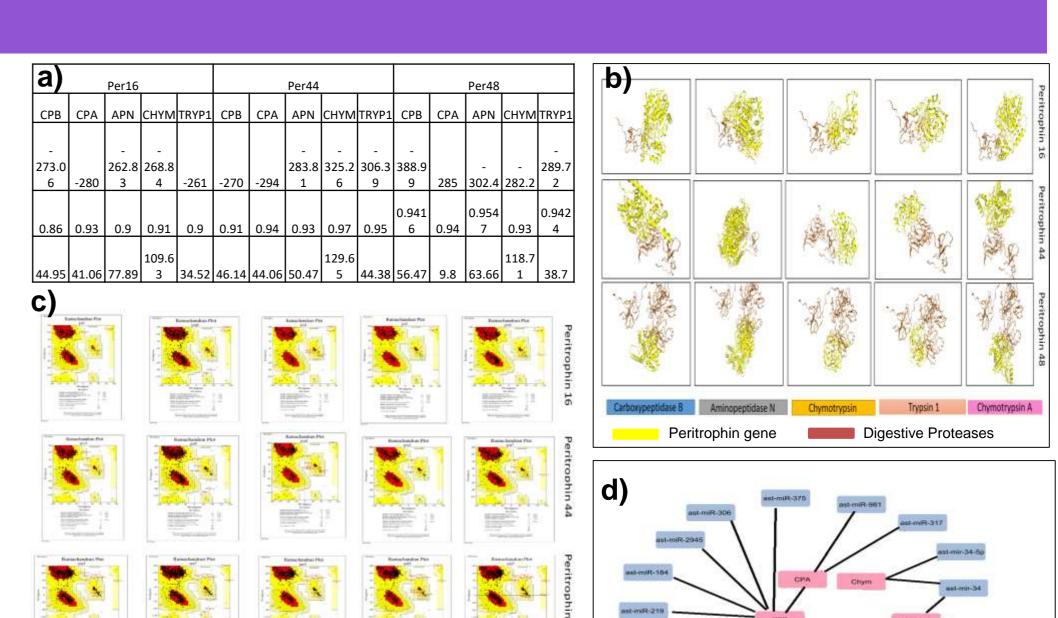


Fig 2: a) Docking of Per16, 44, and 48 with CPB, CPA, Chymotrypsin, Trypsin and APN a-c, b) (d) miRNA-mRNA targeting analysis,

4 CONCLUSION

- Expression of *Per* genes in later larval stages, BF, and IBF stages indicates their potential functional significance across mosquito developmental phases and role in Plasmodium transmission
- Molecular interaction signifies the complexity and specificity of these binding events. The observed variations in binding affinities and structural stabilities suggest distinct roles for different peritrophin proteins in their interactions with digestive enzymes, potentially reflecting their diverse biological functions and potential role in transmission-blocking

5 FUTURE WORKS/ REFERENCES

siRNA and subsequent investigations will be conducted to examine the impact of *per*-gene silencing on *CPB* and viceversa, as well as to evaluate the influence of gene silencing on the peritrophic matrix, vector proficiency, and the transmission of Plasmodium.

References

- Singh, M., Singh, G., Dubey, A., & Chaitanya, R. K. (2021). Plasmodium's journey through the Anopheles mosquito: A comprehensive review. *Biochimie*, 181, 176-190.
- Rao, V. V., Kolli, S. K., Bargava, S., & Chaitanya, R. K. (2017). Modulation of midgut peritrophins' expression during Plasmodium infection in Anopheles stephensi (Diptera: Culicidae). Current Science, 154-160.
- Dong, Y., Aguilar, R., Xi, Z., Warr, E., Mongin, E., & Dimopoulos, G. (2006). Anopheles gambiae immune responses to human and rodent Plasmodium parasite species. *PLoS pathogens*, 2(6), e52.