Molecular Tools for Early Detection of Invasive Malaria Vector *Anopheles* stephensi Mosquitoes

Appendix

DNA Sequencing of Mosquito Samples Detected False Positive for *Anopheles* stephensi in Real-time PCR with Late Ct Value

Two pre-isolated DNA of mosquitoes, one each of *Aedes aegypti* and *Anopheles*. *subpictus*, were detected false-positive as *Anopheles stephensi* by real-time PCR with late Ct values (32.95 and 33.09, respectively). To check if false positivity is due to the contamination of DNA from *An. stephensi*, these samples were subjected to DNA sequencing following methods described in the main article under section "DNA sequencing strategy for the confirmation of PCR-based identification of *An. stephensi* in pooled samples" with modification where number of PCR cycles were increased to 45. Both samples showed ≈450 bp amplicon (Appendix Figure 3). The PCR products were sequenced from both directions of the strand. NCBI-blast search revealed 100% homology of all sequences with *An. stephensi*, confirming contamination of samples with DNA from *An. stephensi*.

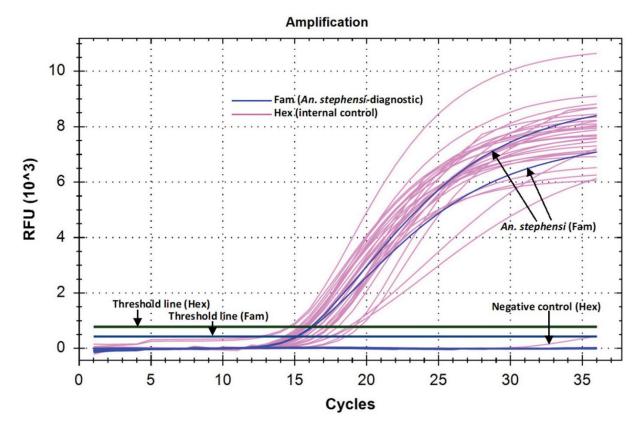
Appendix Table. List of biologic material used to evaluate specificity of An. stephensi-diagnostic PCR assays Description of species Source and nature of material (origin) Anopheline species An. albimanus, strain STECLA BEI Resources, USA, frozen adult (Santa Tecla, El Salvador) An. atroparvus, strain EBRO BEI Resources, USA, frozen adult (Tarragona, Amposta village, Ebro delta, Spain) An. dirus, strain WRAIR2 BEI Resources, USA, frozen adult (Thailand) An. farauti, strain FAR1 BEI Resources, USA, frozen adult (Papua, New Guinea) BEI Resources, USA, frozen adult (USA Marysville, California) An. freeborni, strain F-1 An. funestus, strain FUMOZ BEI Resources, USA, frozen adult (Matolo Province, southern Mozambique) BEI Resources, USA, frozen adult (Pimperena, Mali) An. gambiae, strain G3 An. merus, strain OPHANSI BEI Resources, USA, frozen adult (South Africa) An. quadrimaculatus, strain ORLANDO BEI Resources, USA, frozen adult (USA) An. stephensi. strain STE2 BEI Resources, USA, frozen adult (India) An. stephensi type form, strain DEL Lab colony, adults and larvae (Delhi) An. culicifacies species A, strain NIMR Lab colony, adults (Delhi) An. culicifacies species B Pre-isolated DNA from adults (Raipur, India) An. fluviatilis species S Pre-isolated DNA from adults (Sundergarh, India) Pre-isolated DNA from adults (Hardwar, India) An. fluviatilis species T An. subpictus molecular form A Wild caught adults (Delhi, India) An. subpictus molecular form A Pre-isolated DNA from adults (Raipur, India) An. subpictus molecular form B Pre-isolated DNA from adults (Chilka, Odisha, India) Pre-isolated DNA from adults (A&N Island, India) An. sundaicus cytoform D An. stephensi intermediate form Wild caught adults (Nuh, Haryana, India) An. stephensi var. mysorensis Wild caught adults (Nuh, Haryana, India) Culicines species Ae. aegypti, NIMR strain Lab colony, adults (Delhi) Pre-isolated DNA from adults (Bengaluru) Ae. aegypti Ae. albopictus Wild caught adults (Delhi, India) Cx. quinquefasciatus Wild caught adults (Delhi, India)

Pool of mixed species

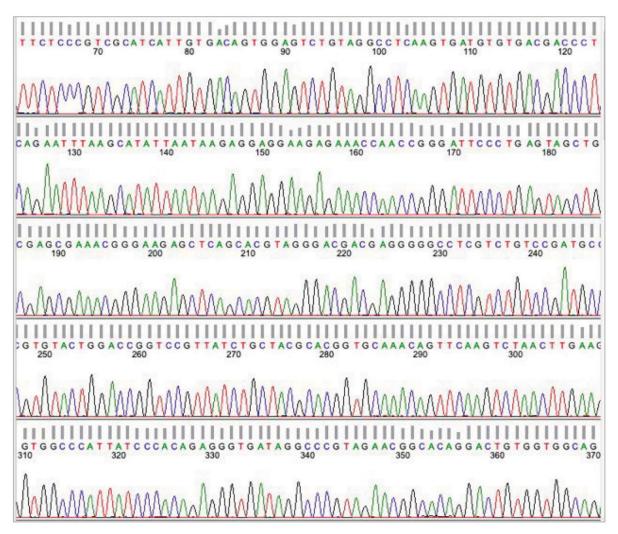
and Cx. quinquefasciatus)

(An. culicifacies s.l., An. subpictus s.l., An. fluviatilis s.l.

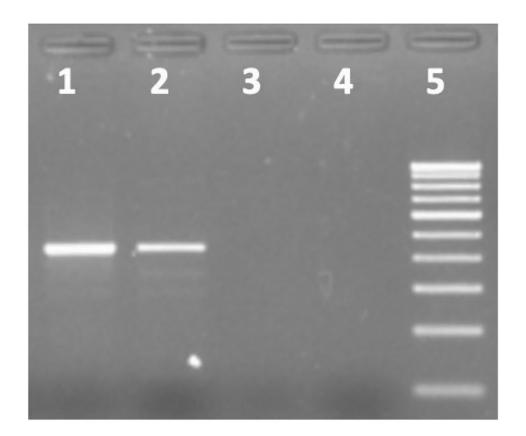
Wild caught adults (Dadri, Uttar Pradesh, India)



Appendix Figure 1. Amplification curve of real-time PCR on *An. stephensi* and other mosquitoes (anophelines and culicine) showing threshold lines for Fam and Hex as determine by the software. Amplification crossing threshold value for Fam was seen in *An. stephensi* only and for Hex was seen in all mosquitoes tested (*An. stephensi*, *An. albimanus*, *An. quadrimaculatus*, *An. dirus*, *An. farauti*, *An. gambiae*, *An. freeborni*, *An. funestus*, *An. atroparvus*, *An. culicifacies* complex, *An. subpictus* complex, *An. merus*, *An. fluviatilis* complex, *Cx. quinquefasciatus*).



Appendix Figure 2. DNA sequence chromatogram (partial ITS2 and 28S) of *An. stephensi* showing the quality of sequence derived from DNA isolated from a pool of 500 mosquitoes containing single *An. stephensi* and rest *An. culicifacies*.



Appendix Figure 3. Gel photograph showing amplification of *An. stephensi*-specific PCR product in DNA isolated from *Ae. aegypti* (lane 1) and *An. subpictus* (lane 2) which were detected false positive for *An. stephensi* in real-time PCR with late Ct values. Lanes 3 and 4 are negative controls; Lane 5: 100 bp DNA ladder.