LETTER TO THE EDITOR

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Molecular mass screening of mosquitoes for filarial parasites in Germany – re-interpretation of PCR xenomonitoring results would be required

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Abstract

Comments concerning interpretation of the PCR xenomonitoring results in the article "Molecular detection of *Setaria tundra* (Nematoda: Filarioidea) and an unidentified filarial species in mosquitoes in Germany" Parasites & Vectors 2012, 5:14.

Keywords: Xenomonitoring, PCR, Filaria, Dirofilaria, Infection rate

Dear Editor,

In the context of an evaluation of the use of xenomonitoring (sensu [1]) for detecting filarial parasites in Germany we would like to point out that the article by Czajka and colleagues [2] suffers from partially incorrect interpretation, and incomplete description, of PCR results. The performance of D. repens DNA detection in mosquitoes using the primers applied in the second round PCRs [3-5], was described neither by Czajka et al. [2] nor such performance tests were described in papers published prior to the one by Czajka et al. [2] (to the best of our knowledge). Furthermore, only one [6] of the primer pairs used in second round PCRs [3-5] perfectly matches D. repens sequences deposited in GenBank [6] - the primer pair 16SOvC and 16SOvB [4, 5]. We asked the authors which primers were used for cytochrome oxidase (COI) gene amplification and the authors informed us that they used, for confirmatory PCRs (second round PCRs), self-designed primers Co1-F and Co1-R that were not mentioned in their article [2].

The authors interpreted every positive result of the screening real time PCR performed on DNA obtained from mosquito pools as positive result of filaria detection. However, the positive results of real time PCR

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could be confirmed as filaria positive results by second round PCR, yet they use the number of screening PCR positive results to calculate minimum infection rate of mosquitoes in Table one [2]. Only the samples confirmed to contain filaria by second round PCR and/or sequencing should have been used for minimum infection rate calculations. Furthermore, the authors should publish information which PCR assays were actually used as second round PCRs for COI gene detection [2]. It would be interesting to see the data reanalyzed by the authors which would help to understand better what did molecular mass screening of mosquitoes for filarial parasites in Germany reveal?

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

RS and AM analyzed data. AM wrote the letter following the discussion with RS. Both authors approved the final version of the manuscript.

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