Truenat $^{\mathbb{R}}$ for *Plasmodium* sub-microscopic infections: Miles to go. . .



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Although Shankar and Kumar suggested Truenat[®] for *Plasmodium* sub-microscopic infections (SMIs)¹ but it is not backed by compelling evidence^{2,3} due to certain limitations:

- (I) **Single-copy target**: Truenat[®] relies on detection of *P. falciparum* (Pf) and *P. vivax* (Pv) single-copy Erythrocyte Binding Protein gene which might compromise comparable limit of detection (LoD) with nPCR based on multi-copy 18s rRNA. The chances of false-negative results could be higher with single-copy target due to spontaneous mutations in the target.
- (2) Comparison with nPCR: Nair and colleagues did not compare Truenat[®] with nPCR on patients' samples, but rather compared it with microscopy/RDT on febrile patients' samples.² They did compare Truenat[®]-tested samples with nPCR but the reference cited described RDTs instead of nPCR.
- (3) **LoD:** Truenat[®]'s performance was validated using only 10 each of Pf and Pv patients' samples for concurrence with nPCR but its species-specific LoD has not been estimated.³ Further, the amount of blood/DNA used for nPCR was not mentioned which imparts ambiguity while comparing concurrence.
- (4) Estimation of LoD: The LoD for *Plasmodium* by PCR can only be determined using accurate count of parasites/microliter of whole blood. 4·5 The published LoD of Truenat[®] is 4.7 parasites/μL and 10 genome-equivalents per PCR reaction for Pv.² It is unclear how exactly were the dilutions prepared, which diluent was used and how the LoD was estimated. Moreover, according to Truenat[®] packinsert sheet, the LoD of Pf/Pv was estimated using plasmid DNA. Using WBC-depleted blood/plasmid DNA for LoD and then extrapolating it on patients' blood might be erroneous as the majority of DNA from a malaria patient's blood is derived from the host WBCs rather than the parasite and is therefore, not comparable.

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(5) Mixed infections: Nair et al., did not estimate the LoD of mixed PfPv using Truenat[®] and nPCR. Because mixed *Plasmodium* infections pose significant diagnostic challenges, it is critical for any new technique to be at least equi-sensitive to mono- and mixed-infections.² Furthermore, Truenat[®] can only detect Pf and Pv against nPCR that can detect all species.

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Truenat[®] appears to be more sensitive than microscopy/RDT but when compared with nPCR, it seems that Truenat[®] has miles to go, particularly for screening SMIs. Further, Nair et al. had strong competing interests: 6/10 authors being employees of the technology developer which also funded the study.² Therefore, Truenat[®]'s performance must undergo unbiased evaluation on sufficient number of febrile patients and results replicated in varied malaria settings.

Contributors

ND: Original draft of the manuscript, Data analysis, Interpretation, Editing the manuscript critically; AS: Conceptualization, Data interpretation, Drafting the manuscript and revising it critically.

Declaration of interests

None.

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