

Evaluation of the Clinical Proficiency of RDTs, Microscopy and Nested PCR in the Diagnosis of Symptomatic Malaria in Ilorin, North-Central, Nigeria

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Abstract

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BACKGROUND: Accurate laboratory diagnosis of suspected malaria is the hallmark to the control of the disease.

AIM: The clinical proficiency of commercial Rapid Diagnostic test kits (RDTs) using nested PCR as quality control was evaluated among patients attending two public healthcare providing institutions in Ilorin, Kwara state, North-Central, Nigeria.

METHOD: A cross-sectional evaluation of finger prick blood samples of volunteer patients were accessed for malaria parasites with pLDH, HRP2, Pf, Pf/PAN and nested PCR molecular assays. The data derived were analysed using standard formulae for diagnostic accuracy, and the obtained predictive values were subjected to a comparison with one-way analysis of variance (ANOVA).

RESULT: Three hundred and sixty-eight (368) patients comprising 203 (55%) females and 165 (45%) males participated in this study. Routine microscopy revealed that 54 (32.7%) males and 80 (39.4%) was infected with Plasmodium falciparum. SD Bioline (pLDH) 47.4%; Carestart Malaria (HRP2) 49.8% recorded low sensitivities. Micropoint (pfPAN) 82.8% and Micropoint (Mal. Pf) 64.4% recorded a high sensitivity. SD Bioline (pLDH) 67.4%; Carestart Malaria (HRP2) 85.9%; Micropoint (Pf/PAN) 62.2% and Micropoint (Mal. Pf) 86.7% had high specificities. The positive predictive value (PPV) ranged from 67.7% to 85.94%, while the negative predictive values (NPV) of 64.4% for SD Bioline (pLDH); 86.7% for Carestart Malaria (HRP2); 89.3% for Micropoint (pfPAN) and 58.5% for Micropoint (Mal. Pf). Agarose gel analysis of *P. falciparum* *ssrRNA* gene (206 bp) for 28 specimens containing 10% concordant and discordant samples showed that all 12 negative specimens for RDTs and routine microscopy were truly negative for nPCR. However, the remaining 16 specimens were positive for nPCR and showed discrepancies with routine microscopy and RDTs. Cohen's interrater diagnostic measure analysis revealed that the weighted kappa for the RDTs was moderate 0.417 ($p=0.027$), 95%CI (0.756, 0.078) and good for nPCR 0.720 ($p < 0.001$), 95%CI (0.963, 0.477). The area under the curve (AUC) specify that nPCR has been more effective than the RDTs (nPCRAUC = 0.875; $p < 0.001$ and RDTsAUC = 0.708; $p = 0.063$).

CONCLUSION: A thorough large-scale quality control is advocated on all commercial RDTs being used in most sub-Saharan African countries. This is to avoid double jeopardy consequent upon misdiagnosis on unidentified positive cases serving as pool reservoir for the insect vector and cyclical infection and re-infection of the populace.

Introduction

In spite of the continued efforts at eradicating malaria in sub-Saharan Africa, it has remained a major public health concern in the region. One of the remarkable milestones left to be achieved for the complete eradication of malaria is the development of adequate biological and clinical diagnostic tools [1] [2]

[3] [4]. Microscopy has been the usual clinical practice for the diagnosis of malaria in endemic regions, but this method of diagnosis is faced with so many issues bothering on expertise, quality of field microscopy, epileptic electricity supply and poor reagents to mention a few [5] [6] [7] [8]. Although, detection thresholds of 4–20 parasites per microlitre are achievable using a Giemsa-stained thick blood film in controlled laboratory situations [9], thresholds of 100–200 parasites per microlitre are more common in field

settings [7] [8]. The use of rapid diagnostic tests as an alternative to microscopy in remote areas has however been advocated [1]. RDT was developed to improve the timeless sensitivity, and objectivity of malaria diagnosis through less reliance on expert microscopy [10]. These kits come in strips and cassettes embedded with immuno-chromatographic cork material that has antibodies targeted for *Plasmodium* parasite antigens. RDTs work on the detection of Histidine-Rich Protein2 (HRP-2) from *Plasmodium falciparum* and Parasite-Specific Lactate Dehydrogenase (pLDH) or *Plasmodium* aldolase from the parasite glycolytic pathway found in all species [11] [13]. RDTs are increasingly being used for malaria diagnosis because they are rapid and easier to use especially in limited resource settings and do not require trained personnel or special equipment [1] [8] [14] [15] [16]. One major pitfall in the use of RDTs in clinical situations is the issue of false-positive/false negative results which may lead to misdiagnosis and over administration of antimalarial drugs which often culminate into increased financial costs, side effects, and selection pressure for development of resistance in malaria parasite populations [17] [18] [19]. Inadvertently, false-negative results may lead to excess morbidity and mortality and further transmission. Thus, quality diagnostics are essential [20]. There are many commercial RDTs in the market today, but their efficacy remains a subject of concern. Therefore, it is not clear which RDT is more appropriate for different epidemiological settings [5] [21] [22]. It has been estimated that a diagnostic test with 95% sensitivity and 95% specificity requiring minimal infrastructure would avert more than 100,000 deaths and about 400 million unnecessary treatments (Long, 2009). The ominous disparity in RDTs and routine microscopy is gradually becoming contentious, hence the need for a more appropriate clinical diagnostic technique. A new diagnostic technique undergoing clinical trials becomes evident. Several molecular detection methods (MDM) are increasingly being researched in clinical practice [11] [23] [24] [25] [26]. It operates on the basis of small nuclear subunit ribosomal (SSU) rRNA genes which are targets, extensively used for the molecular detection of human malaria parasites [25]. These genes are known to have highly conserved regions and their copy numbers ranging from 4 to 8. These characteristics ensure that they are suitable genes for phylogenetic studies and molecular detection of *Plasmodium* parasites [27].

There are two basic approaches for species detection, single polymerase chain reaction and nested polymerase chain reaction (PCR). In general, nested-PCR is more sensitive than single PCR because they detect malaria parasites at very low-level malaria parasitemia [28]. The nested PCR is highly sensitive and has been widely used for diagnosis, confirmation of diagnosis, epidemiological studies, drug efficacy assessment and to measure the accuracy of microscopy [23] [29]. The nested PCR is

used as a quality control laboratory technique to evaluate the clinical proficiency of established RDTs for this study. There is a dearth of published information on the assessment of the diagnostic efficiency of malaria in Ilorin, north-central Nigeria, thus the need for the present study. This study, therefore, seeks to comparatively evaluate the diagnostic efficiency of the various malaria diagnostic methods in Ilorin, north-central Nigeria.

Materials and Methods

A prospective study was conducted from January to March 2016 in Ilorin, Kwara State. A total of 368 subjects (203 males and 165 females) were enrolled at the following comprehensive health centres: University of Ilorin clinic and Children Specialist Hospital (Center Igboro) in Ilorin metropolis. The above areas were used to avoid bias in the sample collection, and so both children and adults were included in the study. The patients were screened for *Plasmodium falciparum* parasites using RDTs, Microscopy and nested PCR techniques.

Peripheral blood samples were collected from voluntary donors at the hospitals mentioned earlier. Examined samples were categorized into 2 groups; concordant and discordant samples after the methods of [24] with slight modifications. The concordant samples were those samples that tested simultaneously positive or negative with PLDH, HRP2 (RDTs) and Microscopy. The discordant group1 were samples that tested positive for PLDH, HRP2 but negative with Microscopy. The discordant groups 2 were samples that tested negative with PLDH, HRP2 but positive with microscopy. Dried blood spot (DBS) from all blood samples with discordant results and 5% of randomly selected samples with concordant results were analysed using nested PCR technique.

Patients of all age groups, presenting with signs and symptoms of malaria infections were included in the study. Demographic information, clinical details and basic information regarding prevention measures of the patients were recorded using questionnaires to establish inclusion and exclusion criteria. Venous blood samples were collected in EDTA and Plain bottles from the patients with suspected malaria cases. A drop was spotted on filter paper (Whatman no. 1). Each filter paper was dried at room temperature and stored carefully in a plastic container to avoid cross-contamination at -20°C.

The diagnostic accuracies of the following RDTs were assessed, viz; pLDH (SD Bioline), HRP2 (Carestart Malaria), Pf/pan (Micropoint) and Mal. Pf (Micropoint). Fresh blood samples were transferred directly from the EDTA bottle to RDTs sample pads

with 2 drops of buffer solution (according to manufacturers' instruction). The buffer solution allows the blood to migrate towards the diagnostics and control line.

Thin and thick blood films were prepared following the methods of Gilles [30]. The slides were fixed in methanol in order to allow lysis of red blood cells. The slides were then stained with 10% Giemsa solution for 30 minutes, and thereafter washed, and screened under oil immersion (X 1000 Mg) Olympus microscope for *Plasmodium spp.* Parasite density was determined as the number of parasites per 200 leukocytes (WBC).

Dried blood spot (DBS) from all blood samples with discordant results and 5% of randomly selected samples with concordant results were analysed using molecular (nested PCR) technique. The various techniques were carried out in separate rooms.

Two methods were used for the DNA extraction according to Berczky *et al.*, [31]. Briefly:

Tris-EDTA buffer-based extraction: Each filter paper punch was placed in Eppendorf tube, soaked in 65 μ l of TE buffer and incubated at 50°C for 15 minutes. The punches were then pressed gently at the bottom of the tube several times, using pipette tip for each punch and heated at 97°C for 15 minutes to elute the DNA templates. The liquid condensing on the lid and the wall of the tubes were removed by short centrifugation. DNA extract was kept at 4°C for use within a few hours or stored at 20°C.

Chelate extraction: Each filter paper punch was incubated overnight at 4°C in 1ml of saponin in phosphate buffered saline (PBS). The punches were washed for 30 minutes in PBS at 4°C, transferred into new tubes containing 25 μ l of stock solution (20% chelex and 75 μ l of distilled water) and vortexed for 30 seconds. The tubes were heated at 99°C for 15 minutes to elute DNA templates.

A Master Mix containing all the reagents is prepared and aliquoted into the reaction tubes and overlaid with mineral oil. DNA templates generated from each sample by the two respective methods were added last into the master mix. Two amplification reactions were carried out. In the first amplification reaction (nest 1), a pair of oligonucleotide primers, which hybridised to a sequence in ssRNA gene of *Plasmodium falciparum*, were used. The product of the first reaction is then used as DNA template for a second amplification reaction (nest 2). The second amplification reaction involves the use of genus-specific and species-specific primers. The genus-specific and species-specific primers (oligonucleotide primers) was used to indicate the presence of malaria parasites and the *Plasmodium* species in the samples.

The PCR products from the amplification reaction were subjected to agarose gel electrophoresis. The electrophoresed products (result) were then interpreted by molecular detection specialist.

Data was entered and analysed using Statistical package for social sciences (SPSS 16.0) to determine the discrepancies between RDTs and nested PCR. The Sensitivity (sn), specificity (sp), positive predictive values (ppv) and negative predictive value (npv) were calculated using standard formulae, and their differences were analysed by comparing their mean values with One way ANOVA. The level of significance was estimated at $p < 0.05$. An interrater reliability analysis using the Kappa statistic was performed to determine consistency among raters. Cohen's interrater statistics was used to validate a measure of agreement between the diagnostic outcomes among the laboratory diagnostic tools.

Results

A Total of 368 patients; 203 (55%) females and 165 (45%) males were randomly enrolled and screened for malaria parasites using microscopy and RDT test kits. Four commercial RDTs were employed for this study, viz; Malaria Ag Pf/Pan by Standard Diagnostics Inc. Hagandong, Korea; Malaria HRP2 (Pf) by Carestart™ by Access Bio, Inc., New Jersey, USA; Malaria PF by Micropoint Inc. The USA.

From the PLDH (SD BIOLINE test kit), 145 (39%) were infected consisting of 72 (49.7%) males and 73 (50.3%) females. Carestart (HRP2) showed that 146 (39.7%) with male (72, 49.3%) and female (74, 50.7%). Micropoint (pfPAN) reported 167 (45.4%) with male (87, 52.1%) and female (80, 47.9%). Results from microscopy indicated that 234 (63.6%) were infected, consisting of 111 (47.4%) males and 123 (52.6%) females. However, only Micropoint (pfPAN) and (*Mal. Pf*) were statistically significant as regards sex at $P > 0.05$ (Table 1).

Table 1: Diagnostic detection of *P. falciparum* based on gender

Diagnostics	Sex	Prevalence of Malaria infections		p-value
		Infected n (%)	Uninfected n (%)	
SD Bioline (pLDH)	Male	72 (49.7)	93 (41.7)	0.082
	Female	73 (50.3)	130 (58.3)	
Carestart Malaria (HRP2)	Male	72 (49.3)	93 (41.9)	0.098
	Female	74 (50.7)	129 (50.1)	
Micropoint (pfPAN)	Male	87 (52.1)	78 (38.8)	0.007
	Female	80 (47.9)	123 (61.2)	
Micropoint (<i>Mal. Pf</i>)	Male	72 (46.8)	93 (43.5)	0.028
	Female	82 (53.2)	121 (56.5)	
Microscopy	Male	111 (47.4)	54 (32.7)	0.112
	Female	123 (52.6)	80 (39.4)	

The diagnostic detection of *P. falciparum* across age-groups in the study revealed that there were discrepancies in the diagnostic accuracy of the four RDTs when compared to routine microscopy. For instance at age ≤ 5yrs; routine microscopy identified varying parasitaemia with all the subjects at that age group being infected (40, 100%). However, SD Bioline identified only two (2, 5.0%) as positive for *P. falciparum*. At age-group 6-15yrs, the results were not comparable as slight differences occurred with all the diagnostic methods. There was a change in trend at age 16-25 yrs where it was observed microscopy detected more subjects as malaria positive (56, 62.9%) when compared with the other four RDTs (SD Bioline 15, 38.5%; Carestart 13, 33.3%; Micropoint pfPAN 24, 27.0%; Micropoint Pf 26, 29.2%). The diagnostic accuracy of the RDTs appear to positively pick up with increasing age of the subjects, for instance at age 36-45 yrs, there was no appreciable differences in the diagnostic outcome of the subjects (SD Bioline 31, 34.8%; Carestart 23, 25.8%; Micropoint pfPAN 23, 59.0%; Micropoint Pf 23, 59.0%; Microscopy 17, 43.6%). Malaria positive subjects declined with routine microscopy at age ≥ 46 yrs (9, 34.6%) while the RDTs showed more positive results within this age group. However, the diagnostic tests were all statistically significant at p < 0.001 (Table 2).

Table 2: Diagnostic detection of *P. falciparum* across age-group

Age-grp.	Diagnostics	Prevalence of Malaria infections		p-value
		Infected n (%)	Uninfected n (%)	
≤ 5 yrs	SD Bioline (pLDH)	2 (5.0)	38 (95.0)	
	Carestart Malaria (HRP2)	4 (10.0)	36 (90.0)	
	Micropoint (pfPAN)	9 (22.5)	31 (77.5)	
	Micropoint (Mal. Pf)	9 (22.5)	31 (77.5)	
	Microscopy	40 (100)	0	
6-15 yrs	SD Bioline (pLDH)	57 (65.5)	30 (34.5)	
	Carestart Malaria (HRP2)	65 (74.7)	22 (25.3)	
	Micropoint (pfPAN)	65 (74.7)	22 (25.3)	
	Micropoint (Mal. Pf)	62 (71.3)	25 (28.7)	
	Microscopy	78 (89.7)	9 (10.3)	
16-25 yrs	SD Bioline (pLDH)	31 (34.8)	58 (65.2)	
	Carestart Malaria (HRP2)	23 (25.8)	66 (74.2)	
	Micropoint (pfPAN)	24 (27.0)	65 (73.0)	
	Micropoint (Mal. Pf)	26 (29.2)	63 (70.8)	
	Microscopy	56 (62.9)	33 (37.1)	
26-35 yrs	SD Bioline (pLDH)	28 (32.2)	59 (67.8)	
	Carestart Malaria (HRP2)	29 (33.3)	58 (66.7)	
	Micropoint (pfPAN)	27 (31.0)	60 (69.0)	
	Micropoint (Mal. Pf)	27 (31.0)	60 (69.0)	
	Microscopy	34 (39.1)	53 (60.9)	
36-45 yrs	SD Bioline (pLDH)	15 (38.5)	24 (61.5)	
	Carestart Malaria (HRP2)	13 (33.3)	26 (66.7)	
	Micropoint (pfPAN)	23 (59.0)	16 (41.0)	
	Micropoint (Mal. Pf)	23 (59.0)	16 (41.0)	
	Microscopy	17 (43.6)	22 (56.4)	
≥46 yrs	SD Bioline (pLDH)	12 (46.2)	14 (53.8)	
	Carestart Malaria (HRP2)	12 (46.2)	14 (53.8)	
	Micropoint (pfPAN)	19 (73.1)	7 (26.9)	
	Micropoint (Mal. Pf)	19 (73.1)	7 (26.9)	
	Microscopy	9 (34.6)	17 (65.4)	
Total	SD Bioline (pLDH)	145 (39.4)	223 (60.6)	<0.001
	Carestart Malaria (HRP2)	146 (39.7)	222 (60.3)	<0.001
	Micropoint (pfPAN)	167 (45.4)	201 (54.6)	<0.001
	Micropoint (Mal. Pf)	166 (45.1)	202 (54.9)	<0.001
	Microscopy	234 (63.6)	134 (36.4)	<0.001

Routine microscopy was used as a gold standard against the four RDTs. There was low sensitivity for the first two diagnostic RDTs, viz; SD

Bioline (pLDH) 47.4%; Carestart Malaria (HRP2) 49.8%. However, Micropoint (pfPAN) recorded a high sensitivity of 82.8%, and Micropoint (Mal. Pf) had 64.4%. The probability that a diagnostic will indicate the absence of malaria parasite among those without the disease is called specificity and in the present study, the specificity of SD Bioline (pLDH) 67.4%; Carestart Malaria (HRP2) 85.9%; Micropoint (pfPAN) 62.2% and Micropoint (Mal. Pf) 86.7%. PPV has determined the probability that the positive results were positive. In the following RDTs, the ppv ranged from 67.7% to 85.94%. The probability that the subjects truly does not have malaria reflected with NPV values of 64.3% for SD Bioline (pLDH); 86.7% for Carestart Malaria (HRP2); 89.3% for Micropoint (pfPAN) and 58.5% for Micropoint (Mal. Pf) (Table 3).

Table 3: Performance of RDTs using routine microscopy as the gold standard

Diagnostics	Sensitivity	Specificity	PPV	NPV
SD Bioline (pLDH)	0.4739	0.6739	0.7078	0.6438
Carestart Malaria (HRP2)	0.4979	0.8593	0.8593	0.8667
Micropoint (pfPAN)	0.8278	0.6223	0.6773	0.8929
Micropoint (Mal. Pf)	0.6438	0.8667	0.7905	0.5850

Keys: PPV=positive predictive value; NPV=negative predictive value.

Malaria sensitive groups were divided into the concordant groups and the discordant groups. The concordant group was further divided into Group 1: all True positives (i.e. positive for pLDH, HRP2 and microscopy) and Group 2: True negatives (i.e. negative for pLDH, HRP2 and microscopy). The discordant groups were also divided into two: Discordant group false positives (i.e. positive for pLDH and HRP2 and negative for microscopy), and the discordant group 2: false negatives (i.e. negative for pLDH and HRP2 and positive for microscopy). The clinical accuracy of the diagnostics was evaluated with nPCR by taking 10% each of Groups 1 & 2 and randomly picking 4 samples each from Groups I & II (Figure 1).

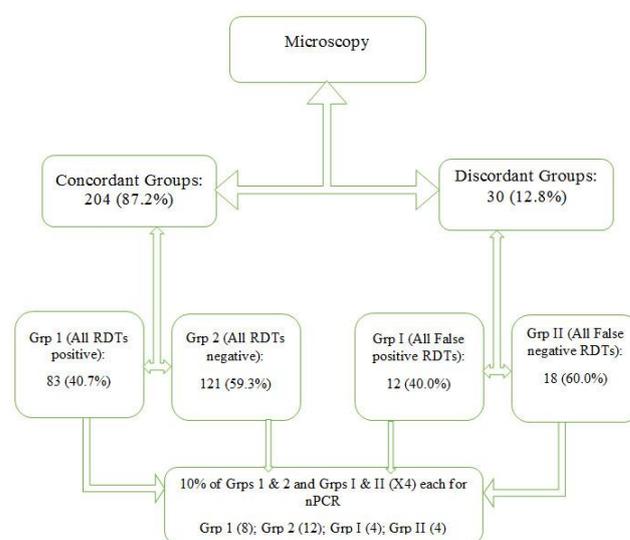


Figure 1: Flowchart describing the proportion of RDTs Sensitivity and samples subjected to nested PCR Analysis

The concordant group: Group 1: all True positives (i.e. positive for pLDH, HRP2 and microscopy) and Group 2: True negatives (i.e. negative for pLDH, HRP2 and microscopy). The discordant groups: Group 1: False positives (i.e. positive for pLDH and HRP2 and negative for microscopy), and Group 2: false negatives (i.e. negative for pLDH and HRP2 and positive for microscopy). The clinical accuracy of the diagnostics was evaluated with nPCR by taking 10% each of Groups 1 & 2 and 4 samples each from Groups 1 & 2 was randomly picked.

Agarose gel photograph of *P. falciparum* *ssrRNA* gene (206 bp) resolved on 1.2% Agarose gel from 28 clinical samples of patients with malaria suspected cases provided the following results: Lanes 3, 5, 6, 8, 11, 13, 15, 19, 22, 24, 27 and 31 were PCR negative for *P. falciparum*. All the 12 specimens diagnosed and grouped as negative for RDTs and routine microscopy was truly negative for nPCR. This further shows that the RDTs specificity can be relied upon. Lane 1 and 17 = DNA Ladder while Lane 2 and 18 = 3D7 Positive control (Figure 2). However, lanes 4, 7, 9, 10, 12, 14, 16, 20, 21, 23, 25, 26, 28, 29, 30, 32 showed *P. falciparum* positive cases meaning that the RDTs had low sensitivities as they initially reported non-malaria infections. To revalidate the present fallouts, the assays used as negative controls were subjected to nPCR and there was no amplification.

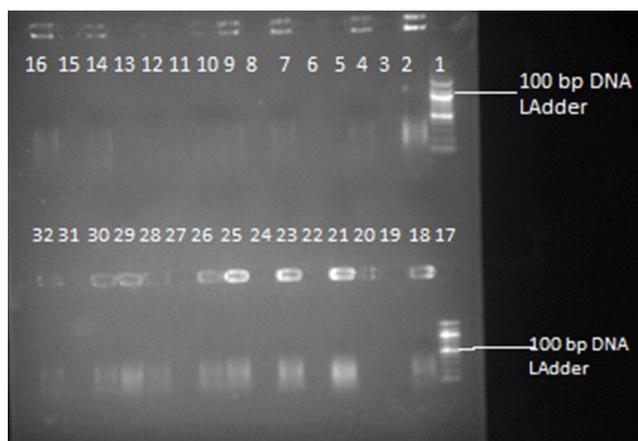


Figure 2: Schematic representation of Agarose gel photograph of *P. falciparum* *ssrRNA* gene (206 bp) from 28 clinical samples using species-specific oligonucleotide pairs for *P. falciparum*. Lanes 3, 5, 6, 8, 11, 13, 15, 19, 22, 24, 27 and 31 were PCR negative for *P. falciparum*. Lane 1 and 17 = DNA Ladder, while lane 2 and 18 = 3D7 Positive control. Lanes 4, 7, 9, 10, 12, 14, 16, 20, 21, 23, 25, 26, 28, 29, 30, 32 *P. falciparum* positive cases

Cohen's interrater diagnostic measure analysis revealed that the weighted kappa was estimated to assess the extent to which the sensitivity and specificity of RDTs and nPCR correspond with the routine microscopy in the diagnosis of malaria among a sampled population of the subjects. The kappa value for the RDTs was moderate 0.417 ($p = 0.027$), 95%CI (0.756, 0.078) and good for nPCR

0.720 ($p < 0.001$), 95%CI (0.963, 0.477). The area under the curve (AUC) is the percentage of randomly drawn pairs for which malaria is truly positive. The effectiveness of the RDTs and nPCR was also evaluated with AUC. The nPCR was found to be more effective than the RDTs (nPCR^{AUC} = 0.875; $p < 0.001$ and RDTs^{AUC} = 0.708; $p = 0.063$) (Table 4).

Table 4: Cohen's Interrater diagnostic measure for RDTs and nPCR with routine microscopy

Diagnostics	AUC	p-value	Kappa	95% CI	
				Upper Bound	Lower Bound
<i>Population for Quality control</i>					
RDTs	0.708	0.027	0.417	0.756	0.078
nPCR	0.875	<0.001	0.720	0.963	0.477

Key: AUC: area under curve -0.5: worthless, to 1, a perfect test; Kappa < 0.20: poor, 0.41-0.60: moderate, 0.61-0.80: good and 0.81-1: very good; P is significant < 0.05. 95%CI: Estimate \pm 1.96 SE.

Discussion

The diagnostic accuracy of malaria laboratory test reagents and kits has been a major pitfall in the efforts towards complete eradication of the disease in sub-Saharan Africa where presumptive evaluation has remained the norm [15] [16] [20]. The present study identified some incongruities between the diagnostic reliability of the sampled RDTs cum routine microscopy and nPCR. The diagnostic detectability of SD Bioline (pLDH) and Carestart Malaria

HRP2 were not gender significant ($p > 0.05$) while that of Micropoint (pfPAN) and (*Mal. Pf*) were statistically significant ($p < 0.05$). Concerning the stratified age groups in the study; a retinue of diagnostic discrepancies were also observed especially at age ≤ 5 yrs, the RDTs failed to detect *Plasmodium falciparum* antigens adequately. If the RDT results obtained for this age group were clinically relied upon; then diagnosis will be misleading the clinician. The low sensitivity recorded in the present situation among the ≤ 5 yrs group is in contrast with what was obtained in other studies [32] [33] [34] [35]. In previous studies, it was reported that RDTs failure to adequately indicate febrile illness usually occurs due to handling and storage [36] [37]. The RDTs on arrival in developing countries face challenges of immediate inspection and dispatch to health facilities and consequently are abandoned in the sun [38] [39].

The diagnostic accuracy picked up with increasing age in such a way as to believe that there exist a wider differences in addition to a substantial none overlap between the test methods. This assumption has also been reported by Endeshaw et al., [40]; Ojurongbe et al., [15]; Osei-Yeboah et al., [16]. Although positive malaria subjects declined with routine microscopy (≥ 46 yrs) which appears to be the normal occurrence in malaria endemic regions [41] [42] the results obtained with the sampled RDTs

indicated an array of false negatives. The sensitivity of the RDTs (Pf/HRP2 and Pf PLDH) were low and fell below the threshold approved by WHO [1]. The observed low sensitivity in this study is similar to reports in other field studies [4] [43] [44] [45] [46]. A probable reason for these might be because some of the RDT kits rely on PfHRP2 antigen for the detection of *P. falciparum*, and PfHRP2 gene deletions often lead to false-negative results [43] [47] [48]. On the other hand, Baker *et al.*, [49] conducted studies on PfHRP2 DNA obtained from isolates from African and South American countries with extreme sequence variation but concluded that diversity of the protein was not a major cause of the varying sensitivities of RDTs. Other possible explanations for the poor RDTs sensitivity, despite significant parasitemia, may include substandard diagnostics [50] [51], defects in device membrane [29], anti RDTs antibodies in human [7] and genetic diversity of parasite antigen PfHRP2 and PfPLDH [52].

Routine microscopy results were equivalent to those obtained by nPCR, except for few microscopically negative, but positive for nPCR as also reported by these authors [15] [22] [43] [53]. From a clinical perspective, failure to establish high parasite density from blood samples of febrile malaria patients can pose grave fatality [12] [20]. Diagnostic output with low specificity is less serious when compared with low sensitivity because presentations with low specificity only translate to overdiagnosis and an over the treatment of non- malaria cases [7] [44]. Faint colour bands formation on RDTs may also contribute to underreporting as subjective interpretation may contribute to poor performance [54]. This study used the nPCR techniques as a quality control technique between routine microscopy and available commercial RDTs. While appreciating that nPCR techniques provide astounding results even detecting parasitaemia at low microlitre, the high cost, complexity, materials and laborious exercise required may not make it a candidate diagnostic tool in many resource-poor communities [22] [55]. We determine the ability of a nested PCR assay to detect *Plasmodium* DNA in stored dried blood samples. The authors only choose the nPCR to ascertain the clinical diagnostic accuracy of the use of routine microscopy and RDTs in our settings. Often, malaria presentations at primary health providing centres in our communities are usually misleading, and the diagnostic failure of the RDTs may contribute over or/and under therapy.

Moreover, this has far-reaching implications such as treating patients on clinical suspicion, the threat of drug resistance, as well as possibility of missing a malaria case and consequent complications and mortalities [5] [36]. The WHO minimum selection criteria for malaria RDTs, i.e. panel detection score (PDS) against *P. falciparum* samples should be at least 75% at 200 parasites/ μ L [56]. WHO recommends that RDTs must demonstrate high sensitivity and specificity [1].

In conclusion, the results obtained from this study has indeed proven that DNA amplification technique provides a better diagnostic outcome as compared with microscopy and antigen-antibody complex detection. However, some of the shortcomings of the DNA amplification include; time, cost, sample quality and probably the volume of the sample available for PCR analysis. Furthermore, the ambiguity and complexity associated with the suitable choice of adequate RDTs to meet up with the prompt and immediate diagnosis are major challenges, especially now that the clinical approach to management, control, elimination and patient's well-being are paramount. Several factors may be militating against the reliability of any of these RDTs, some of these include genetic diversity of parasite antigen, parasite detection threshold, parasite target and level of malaria endemicity. A regular and thorough large-scale quality control is advised on all commercial RDTs imported into most sub-Saharan African countries, this is imperative to avoid the double jeopardy consequent upon misdiagnosis on unidentified positive cases serving as pool reservoir for *Anopheles* mosquitoes leading to cyclical infection and re-infection among the inhabitants.

Ethics approval and consent to participate

This study was part of a study approved by the University of Ilorin Ethical Consideration Committee on 10th December 2012 with protocol approval number: UERC/ASN/2012/221. The consent form was administered, approved by the volunteers and documented before commencing the study.

Authors' Contributions

OSH designed the protocols, performed the statistical analysis and wrote the final draft, OAO collected literature and wrote the first manuscript, OBO and OF provided volunteers for the research and obtained consent, OA carried out the nPCR analysis, OSH and OAO performed the routine microscopy, RDTs and administered the questionnaire.

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