Malaria Diagnosis across the International Centers of Excellence for Malaria Research: Platforms, Performance, and Standardization

Tamaki Kobayashi,* Dionicia Gamboa, Daouda Ndiaye, Liwang Cui, Patrick L. Sutton, and Joseph M. Vinetz*

Department of Epidemiology, Johns Hopkins Bloomberg School of Public Health, Baltimore, Maryland; Laboratorio de Malaria, Laboratorios de Investigación y Desarrollo, Facultad de Ciencias y Filosofía, Universidad Peruana Cayetano Heredia, Lima, Peru; Institute de Medicina Tropical, Universidad Peruana Cayetano Heredia, Lima, Peru; Department of Parasitology and Mycology, Faculty of Medicine, Pharmacy, and Odontology, Université Cheikh Anta Diop de Dakar, Dakar, Senegal; Department of Entomology, Pennsylvania State University, University Park, Pennsylvania; Center for Genomics and Systems Biology, New York University, New York, New York; Division of Infectious Diseases, Department of Medicine, University of California San Diego School of Medicine, La Jolla, California

Abstract. Diagnosis is "the act of identifying a disease, illness, or problem by examining someone or something." When an individual with acute fever presents for clinical attention, accurate diagnosis leading to specific, prompt treatment often saves lives. As applied to malaria, not only individual patient diagnosis is important but also assessing population-level malaria prevalence using appropriate diagnostic methods is essential for public health purposes. Similarly, identifying (diagnosing) fake antimalarial medications prevents the use of counterfeit drugs that can have disastrous effects. Therefore, accurate diagnosis in broad areas related to malaria is fundamental to improving health-care delivery, informing funding agencies of current malaria situations, and aiding in the prioritization of regional and national control efforts. The International Centers of Excellence for Malaria Research (ICEMR), supported by the U.S. National Institute of Allergy and Infectious Diseases, has collaborated on global efforts to improve malaria diagnostics by working to harmonize and systematize procedures across different regions where endemicity and financial resources vary. In this article, the different diagnostic methods used across each ICEMR are reviewed and challenges are discussed.

INTRODUCTION

Combating the global malaria burden begins with accurate diagnosis, which guides specific treatment and public health reporting. Reliable malaria diagnosis improves health-care delivery and informs funding agencies of current malaria situations, which is key for prioritization of regional- and national-level control efforts. Adequate malaria diagnostics contribute toward the long-term goal of malaria elimination, but accuracy in both testing and reporting depends on resource availability, health worker expertise, and formal reporting policies that vary among regions. Microscopy and rapid diagnostic tests (RDTs), which detect parasite antigen, are most commonly used to diagnose malaria; each method has its own strengths and weaknesses.¹ Current World health Organization (WHO) recommendations for malaria diagnosis focus on the identification of acute symptomatic malaria.² Therefore, individuals with subclinical malaria parasitemia who do not present at health facilities, and those with parasitemia below the detection limit of microcopy or RDTs, are missed.^{3,4} Individuals with sub-patent, subclinical parasitemia are increasingly recognized as key for maintaining regional malaria transmission, and as malaria transmission declines in some areas because of increased malaria control, such individual contributions to malaria endemicity become increasingly important.^{3,5,6} Because regional malaria transmission profiles will change as control strategies progress to elimination, diagnosing these changes will become necessary on a global level. Consequently, the National Institute of Allergy and Infectious Diseases established the Interna-

*Address correspondence to Tamaki Kobayashi, Department of Epidemiology, Johns Hopkins Bloomberg School of Public Health, 615 N. Wolfe Street, W4612, Baltimore, MD 21205, E-mail: tkobaya2@jhu.edu or Joseph M. Vinetz, Division of Infectious Diseases, Department of Medicine, University of California San Diego School of Medicine, 9500 Gilman Drive, La Jolla, CA 92093-0760, E-mail: jvinetz@ucsd.edu. tional Centers of Excellence for Malaria Research (ICEMR) program in 2010. As summarized in the foreword of this journal supplement, the ICEMR network includes 10 independent research programs representing all major malaria-endemic regions. Integral to each ICEMR is the inclusion of multiple, epidemiologically contrasting field sites and innovative multidisciplinary approaches and the use of robust diagnostic strategies to help assess the changing malaria situation in each endemic region. The goal of this integrated approach is to generate a knowledge base for improving clinical and field management of malaria. In this article, we discuss laboratorybased diagnostic tools used by ICEMRs, which is a major focus of their research programs.

Since 2010, WHO has recommended either RDT or microscopy confirmation of suspected malaria cases before treatment.² The use of RDTs increased globally from less than 200,000 in 2005 to more than 74 million in 2011.⁷ The increased availability and use of RDTs in the public sector has resulted in fewer cases of non-malaria acute febrile illness being empirically treated with antimalarial drugs, which helps to prevent development of drug resistance. Although both microscopy and RDTs are of great value in guiding appropriate malaria treatment, subclinical/sub-patent *Plasmodium* parasitemic individuals do not come to clinical attention and cannot be detected by these techniques because of limited sensitivity.^{3,5,6}

During the past decade, highly sensitive and specific nucleic acid amplification techniques have been developed to detect malaria parasites including polymerase chain reaction (PCR), real-time quantitative PCR (qPCR), and reverse transcriptase PCR (RT-PCR) (reviewed in references^{1.8}). Compared with light microscopy's limit of detection (about 30 parasites/µL) by the best microscopists) or RDTs (> 100 parasites/µL), nucleic acid amplification methods can detect fewer than 10 parasites/µL.^{1.8} PCR-based methods are extensively used by ICEMRs for the field of malaria epidemiology research. Highly sensitive qPCR methods were developed by either

increasing the volume of sample extracted and analyzed⁹ or by targeting multi-copy genes.¹⁰ However, such techniques often require sophisticated equipment and training and are significantly more expensive than microscopy and RDT. To make highly sensitive but technologically intense malaria diagnostics more readily available in the resource-limited setting, isothermal amplification techniques were adapted.¹¹ Loop-mediated isothermal amplification (LAMP), the best characterized of these techniques, is as sensitive and specific as conventional PCR without needing sophisticated equipment for extraction, amplification, or detection.^{12,13} Comparison of the detection limit between various PCR methods and isothermal detection methods has shown that the lower detection limit of LAMP is 5-10 parasites/µL, comparable to that of conventional PCR.¹¹ Other isothermal techniques such as nucleic acid sequence-based amplification (NASBA) can achieve detection limit of < 1 parasite/ μ L on blood sample sizes of at least 50–100 µL.^{14,15} Other molecular approaches can be used to detect malaria parasites (Table 1). For fieldbased epidemiology studies, a major focus area of ICEMR efforts, PCR-based methods are most commonly used to detect, quantify, and speciate low-density malaria parasitemia, whether asexual or gametocyte forms. Because of space limitations here, we primarily focused on PCR-based methods currently used by the ICEMR network.

Malaria control and elimination strategies depend on providing appropriate treatment of parasitologically confirmed clinical malaria cases as well as asymptomatic carriers. Such specific treatment depends not only on accurate diagnostics but also on effective (and pure or "valid"¹⁶) drugs.¹⁷ Throughout the malaria-endemic world, *Plasmodium falciparum* has developed some level of resistance to many current antimalarial drugs, and chloroquine-resistant *Plasmodium vivax* has been reported in some regions.¹⁸ To address multidrug resistance malaria parasites, WHO has recommended the use of artemisinin-based combination therapy (ACT) to treat *P. falciparum* in most malaria-endemic regions.¹⁹ In areas of *P. vivax* chloroquine resistance, ACTs are also recommended or deployed for treatment of vivax malaria.^{20,21} Because of the high demand and its market value, there has been an issue with counterfeit ACTs.^{22,23} A field deployable method to test drug potency is needed to help prevent the emergence of parasite drug resistance. Recent work by the southeast Asia ICEMR toward this end, to develop an assay to verify the quality of artemisinin class antimalarials, is discussed later in the section, RDT to check quality of artemisinin class antimalarials.

THE DIAGNOSTIC DILEMMA AND CURRENTLY AVAILABLE PLATFORMS

Accurate diagnosis is essential for treating suspected malaria cases, but in many malaria-endemic regions fever is commonly presumed to be malaria without confirmation.²⁴ A diagnostic test for malaria must be specific to identify the infecting Plasmodium species, readily available with a rapid turnaround time, and inexpensive. Even though malaria testing in Africa focuses on P. falciparum, effective diagnostics for P. vivax are important elsewhere because parasitemia is typically lower. Identifying very low parasitemia individuals is not feasible with RDTs, but requires highly competent microscopy complemented by molecular assays. Currently, it is not possible to diagnose asymptomatic P. vivax (and Plasmodium ovale) hypnozoite carriers, yet developing methods to do so is important because the biology of P. vivax makes elimination of this parasite more difficult.^{18,25-27} Even though diagnostic strategies in some countries have started to use combination RDTs more often, RDTs are not useful in elimination settings, but only for diagnosis of acute disease. Further such strategies do not take the diagnosis of P. ovale or Plasmodium malariae into account, particularly in Africa, or the zoonotic Plasmodium knowlesi in Asia. It is important to identify infections due to both single and more than one Plasmodium species to ensure proper antimalarial drug

	Curi	rently available tools to detect malaria parasites
Platform	Target	Description and potential use
Microscopy	Whole parasite	Detects asexual and sexual blood stage parasites of all species under microscope Reliable readings require skilled microscopists
RDT	Antigen	Detects malaria antigen by immunochromatographic assay with monoclonal antibodies to target parasite antigen
		Detect parasite antigens (HRP, aldolase, or pLDH) circulating in the blood
PCR (conventional and real time)	DNA	Amplifies target parasite DNA. Depending on the target gene, genus- and species-level diagnoses are available. The result of conventional PCR is qualitative whereas qPCR is quantitative
RT-PCR	RNA	Detects mRNA expressed at specific life cycle of the parasite. The test can be used to measure the transmissibility of the infection by quantifying the presence of mosquito-infective sexual stages
NASBA	RNA	Amplifies target RNA in a single step isothermal condition
LAMP	DNA	Detecting infection by a turbidity meter after amplifying parasite DNA; DNA extraction methods are key
Microarrays	DNA	Use extract parasite DNA on a hybridization platform to quantify parasitemia by fluorescence-based detection
Mass spectrometry	Heme	Detects infection based on identification of heme by laser desorption mass spectrometry
Flow cytometry	Hemozoin	Detects infection based on hemozoin concentration
Automated blood cell counter	Hemozoin	Detects infection based on malarial pigment in activated monocytes
Serological tests	Malaria parasite specific antibody	Detects and measures antibodies against malaria parasites as an indicator of recent and/or past exposure to parasites

TABLE 1

HRP = histidine-rich protein; LAMP = loop-mediated isothermal amplification; NASBA = nucleic acid sequence-based amplification; pLDH = *Plasmodium* lactate dehydrogenase; RDT = rapid diagnostic test; qPCR = real-time quantitative PCR; RT-PCR = reverse transcriptase PCR.

treatment.² P. vivax and P. ovale form dormant hypnozoites that should be treated with an 8-aminoquinoline (primaquine; tafenoquine in clinical trials), despite challenges related to using these medications on population scales because of the potential for adverse events due to the presence of even small percentages of people having some form of glucose-6-phosphate dehydrogenase deficiency.^{18,28–33} On a population and geographic level, antimalarial treatment carried out without regard to specificity of diagnosis is a major risk for drug tolerance and resistance as well as potential clinical complications, all of which are concerns throughout the malariaendemic world. With regard to population-level reservoirs of continuing transmission, a specific diagnostic test must be able to identify sub-patent cases, defined as infections with parasite levels below the limit of microscopic or RDT detection.⁵ Subpatent parasitemia cases likely contribute to transmission in endemic regions.^{3,6} Furthermore, sub-patent parasitemia cases are often subclinical, and inaccurate detection of subclinical cases, either due to insufficiently effective diagnostics or policies that impede the diagnosis and treatment of subclinical cases, interferes with malaria control efforts in regions, particularly those nearing elimination. The superior sensitivity of molecular diagnostics could ameliorate both of these problems, and provides opportunity for researchers to investigate other important epidemiological questions, including exposure frequency and the transmissibility of malaria parasites based on the quantification of sexual stages.

Currently, there are many molecular diagnostics platforms being used in the ICEMR network (Table 1). Even though their performance has many advantages, most molecular diagnostics are either unable to be sustained in the field because of cost/resource availability or too laborious to be effective for point-of-care diagnosis, hence reducing effectiveness. RDTs have been useful for point-of-care diagnosis, but the issue of false positive results due to the residual antigenemia after the clearance of the parasite remains problematic.^{34–36} Because of low sensitivity, RDTs are not suitable for diagnosis of very low parasitemias. The limitations of molecular diagnostics and RDTs make the continued use of light microscopy important for accurate diagnosis.

CROSS-ICEMR COMPARISON

Diagnostic methods. The ICEMR program was designed to include a wide range of malaria transmission intensities in contrasting epidemiological settings. Studies were required to be designed to identify and quantify the Plasmodium species causing malaria toward a more complete understanding of the complex interactions among human hosts, malaria parasites, and mosquito vectors in diverse ecological niches taking into account expected and unexpected changes occur over space and time. All 10 ICEMRs implemented light microscopy-based diagnosis as the primary quantification metric. Nine study sites used microscopy plus RDTs; Amazonia ICEMR did not use RDTs for several reasons: microscopy is generally available and accurate; the dominance of P. vivax compared with P. falciparum malaria, combined with the lactate dehydrogenase (LDH)-based RDTs that were not considered sufficiently sensitive; and national policy requires microscopy-based diagnosis. In Peru, one country in which the Amazonia ICEMR is based, pfhrp2and/or pfhrp3-lacking P. falciparum were first reported, which made the histidine-rich protein 2/3 (HRP2/3)-based RDT inadequate for use as described later in the section, HRP2/3 deletion among the circulating parasites. Among ICEMRs, the choice of different RDTs has been largely dependent on availability and national policy-based recommendations (Table 2). ICEMRs reporting the use of RDT were aware of their limitation. There, RDTs are meant to be used for clinically apparent cases, primarily where P. falciparum is the major parasite and where parasitemia is sufficiently high to enable RDT detection. The sensitivity of RDTs is not sufficient for screening of subclinical cases where the parasitemia is generally very low, a focus of most ICEMRs. All 10 ICEMRs implemented molecular diagnostic methods either by conventional PCR, qPCR, or RT-PCR (Table 2). In addition, half of the ICEMRs have also implemented gametocyte detection by sexual stage (Pfs25, Pvs25)-specific RT-PCR (Table 2). Two of the 10 ICEMRs use nucleic acid amplification techniques as a point-of-care diagnosis (i.e., southern Asia ICEMR and west Africa ICEMR), whereas the majority of the ICEMRs use nucleic acid amplification techniques only in the research setting (Table 2). The detection limit reported by the majority of ICEMRs was 11-50, 51-200, and < 10 parasites/µL for microscopy, RDTs, and molecular diagnostics (i.e., LAMP, conventional PCR, qPCR, and gametocyte-specific RT-PCR), respectively.

For the detection of malaria parasite DNA, the majority of ICEMRs reported the use of the 18S rRNA as a target gene either by conventional PCR or qPCR to identify malaria species (Table 2).^{37–39} Other target genes such as *pfr364*, *cytb*, and *pfldh* were also used, but less frequently (Table 2).^{40,41}

Factors affecting the sensitivity of different diagnostics. Microscopy and RDTs are extensively used to diagnose malaria at health facilities. Because of extensive experience, quality assurance/quality control (QA/QC) protocols are well established to assure reliable and comparable results. For light microscopy, double reading of the slides with a third reader for discrepancies is standard among ICEMRs. WHO recommends that parasite quantification be performed against 200 white blood cells (WBCs), and if the parasite count is less than 100 after counting 200 WBCs, the microscopist should continue to 500 WBCs.⁴² The west Africa ICEMR reported implementation of minimum count of 300 WBCs per specimen can improve the sensitivity of the slide reading (D. Krogstad, unpublished data). Increasing the WBC count ensures that low-parasitemia cases are not missed, but requires a substantial increase in microscopy time.

WHO collaborates with the Foundation for Innovative New Diagnostics to evaluate and standardize the use of RDTs.⁴³ For molecular diagnostics, standardization has not been established. The decision to choose a protocol and target genes to use largely depends on the research hypothesis. Nine of the ICEMRs use conventional PCR or qPCR to detect, quantify, and speciate submicroscopic parasitemia. RT-PCR coupled with qPCR is used to detect and quantify the gametocyte stage of the parasite with high sensitivity.⁵

Sample preparation. All 10 ICEMRs currently use at least one of the molecular diagnostic methods such as conventional PCR, qPCR, and LAMP (Table 2). Suitable sample preparation (i.e., DNA extraction) enables optimal sensitivity and is probably the most important variable to affect field deployability. Various DNA extraction methods, which ultimately depend on starting material, have been reported

AntiolIndex basesRadioIndexIndexRadioRadioSubment fractionSubment fractionSubment fractionNum dyterion100 NumWHOLNENENENENENENENENENENENet Addition100 NumWHOLWHOLNENENENENENENENENENet Addition100 NumWHOLWHOLNENENENENENENENENENet AdditionNet					Detail	l of diagnostic tool	I ADLE 2 Is and the usage at	t each ICEMR				
one Use and between the second s			Amazonia	East Africa	India	Latin America	Malawi	Southeast Asia	Southern Africa	Southern Asia	Southwest Pacific	West Africa
Using the field in the field	copic	Use as a diagnostic	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes
Question and and and and and and 		Used as point- of-care	Yes	Yes	Yes	Yes	Yes	Yes	No	Yes	No	Yes
Intensitie Neurality consistent Neurality consisten		diagnosis Quantification metric	WHO guidelines	WHO guidelines	WHO guidelines	WHO guidelines	WHO guidelines	WHO guidelines; percent	WHO guidelines	WHO guidelines; percent	WHO guidelines	WHO guidelines
ONC Dubbit Desktranding <t< td=""><td></td><td>Differentiates asexual and gametocyte stages</td><td>Asexual (all together); sexual (all together)</td><td>Asexual (all together)</td><td>Asexual (all together); sexual (all together)</td><td>Asexual (all together); sexual (all together)</td><td>Asexual (all together); sexual (all together)</td><td>parasitemia Asexual (all together); sexual (all together)</td><td>Asexual (all together); sexual (all together)</td><td>parasitemia Asexual stages (all together); merozoites, trophozoites,</td><td>Asexual (all together); sexual (all together);</td><td>Asexual (all together); trophozoites (separately);</td></t<>		Differentiates asexual and gametocyte stages	Asexual (all together); sexual (all together)	Asexual (all together)	Asexual (all together); sexual (all together)	Asexual (all together); sexual (all together)	Asexual (all together); sexual (all together)	parasitemia Asexual (all together); sexual (all together)	Asexual (all together); sexual (all together)	parasitemia Asexual stages (all together); merozoites, trophozoites,	Asexual (all together); sexual (all together);	Asexual (all together); trophozoites (separately);
CMOC Double reading integration (or solid stepration (or solid stepration stepration (or solid stepration (or solid stepration (or solid stepration (or solid stepration (or solid stepration (or solid stepration (or										semizonts (separately); sexual stages (all together); male and female gametocytes	mate and remate gametocytes (separately)	sexual (all together)
Use as one of diagnostic CR and diagnostic break diagnostic di dia		QA/QC practices	Double reading; independent reading of 10% of slides in reference laboratory	Double reading: triple for discrepancies	Double reading: triple for discrepancies; diagnostic PCR	Double reading	Double reading: triple for discrepancies	Double reading	Double reading: triple for discrepancies	(separately) Double reading: triple for discrepancies; diagnostic PCR	Double reading; triple for discrepancies; independent reading of 10% of slides in reference laboratory;	Double reading: triple for discrepancies
$ \begin{array}{cccccc} Ucda a point. Not applicable for constant of the sectional sectinal sectional sectional sectional sectional secti$	nosis	Use as a diagnostic	No	Yes	Yes	Yes	Yes	Yes	Yes	Yes	diagnostic PCR Yes	Yes
Unsponse unsponseNot applicableParachcek, P^{TM} Falcivar FalcivarSD Bioline AlereSD Bioline (Zambia); ICTFalcivar ApoorCareStart ^M , ApoorParachcek, First ApoorDetectionNot applicableHRP2:HRP2:HRP2:HRP2:HRP2:HRP2:RP2:Uppe $P_{asmodium}$ $P_{falciparum}$		Used as point- of-care	Not applicable	Yes, cross- sectional	Yes, cross- sectional	No	Yes, based on availability	Yes, based on availability	Yes	Yes	Yes	Yes
$ \begin{array}{c cccc} Detection & Not applicable & HRP2: & HRP2$		diagnosis RDT(s)	Not applicable	studies Paracheck- <i>Pf</i> TM	FalciVax	SD Bioline	SD One-Step	BinaxNOW [®] , Alere	SD Bioline (Zambia); ICT (Zimbabwe)	FalciVax	CareStart TM , Apacor	Paracheck; First Response; SD Rioline
SpeciesNot applicableP falciparumP falciparumP falciparumP falciparumP falciparumP falciparumP falciparumspecificityand P. vivaxP. vivaxP. vivaxP. vivaxP. vivaxP. vivaxP. vivaxspecificityR. vivaxP. vivaxP. vivaxP. vivaxP. vivaxP. vivaxP. vivaxDeterminingNot applicableGovernmentBivalentiGovernmentBivalentiGovernmentGovernmentfactor foravailabilityrecommendation:efficacyrecommendation:efficacyrecommendation:availabilityrecommendation:availabilityRDT(s)RDT(s)RDT(s)RVRVRVRVRVRVRVRDT(s)RDT(s)RDT(s)RVRVRVRVRVRVRV		Detection type	Not applicable	HRP2: Plasmodium falciparum	HRP2: P. falciparum; pLDH: Plasmodium	HRP2: P. falciparum; pLDH: P. vivax	pLDH: <i>P. falciparum</i> and Pan	HRP2: <i>P. falciparum</i> ; pLDH: Pan	HRP2: P. falciparum	HRP2: P. falciparum; pLDH: P. vivax	HRP2: <i>P. falciparum</i> ; pLDH: Pan	HRP2: P. falciparum
DeterminingNot applicableGovernmentBivalent; GovernmentGovernmentBivalent; GovernmentGovernmentGovernmentfactor forrecommendation; recommendation; efficacyefficacy recommendation; availabilityGovernmentBivalent; recommendation; recommendation; recommendation; recommendation; availabilityGovernment recommendation; <b< td=""><td></td><td>Specificity</td><td>Not applicable</td><td>P. falciparum</td><td>P. falciparum and P. vivax</td><td>P. falciparum and P. vivax</td><td>P. falciparum, P. vivax, Plasmodium malariae, and Plasmodium</td><td>P. falciparum, P. vivax, P. malariae, and P. ovale</td><td>P. falciparum</td><td>P. falciparum and P. vivax</td><td>P. falciparum, P. vivax, P. malariae, and P. ovale</td><td>P. falciparum</td></b<>		Specificity	Not applicable	P. falciparum	P. falciparum and P. vivax	P. falciparum and P. vivax	P. falciparum, P. vivax, Plasmodium malariae, and Plasmodium	P. falciparum, P. vivax, P. malariae, and P. ovale	P. falciparum	P. falciparum and P. vivax	P. falciparum, P. vivax, P. malariae, and P. ovale	P. falciparum
		Determining factor for selection of RDT(s)	Not applicable	Government recommendation; availability	Bivalent; efficacy results; availability	Government recommendation	Government recommendation; availability	Bivalent; availability	Government recommendation	Bivalent; availability	Government recommendation; sensitivity to <i>P. vivax</i>	Government recommendation; availability

TABLE 2

102

KOBAYASHI AND OTHERS

		West Africa	Yes	HRP2 deletions	Yes	Yes	Conventional PCR; qPCR	Asexual	18S rRNA, Pfr364	Centralized location in Mali (Senegal)
		Southwest Pacific	No	Ŷ	Yes	No	Conventional PCR; qPCR; RT-PCR	Asexual and sexual	18S rRNA, pfs25 and pvs25	Onsite in Papua New Guinea; offsite at the Walter and Eliza Hall Institute and the Swiss Tropical and Public Health Institute
		Southern Asia	Yes	°Z	Yes	Yes (LAMP)	Conventional PCR; LAMP	Asexual	18S rRNA	Onsite in India
		Southern Africa	No	HRP2 deletions	Yes	No	Conventional PCR; qPCR; RT-PCR	Asexual and sexual	PfCytb, Pfs25	Onsite in Macha and Ndola (Zambia) and Harare (Zimbabwe)
		Southeast Asia	No	°Z	Yes	No	Conventional PCR	Asexual	18S rRNA	Centralized location in host countries
TABLE 2	Continued	Malawi	No	HRP2 deletions	Yes	No	qPCR, RT-PCR	Asexual and sexual	pfLDH	Centralized location in Blantyre
	U	Latin America	Yes	°Z	Yes	No	qPCR, RT-PCR, LAMP	Asexual and sexual	18S rRNA	Real-time PCR in centralized laboratory, LAMP in field
		India	Yes	°Z	Yes	No	Conventional PCR	Asexual	18S rRNA	Onsite in Raurkela (Odisha), Chennai (Tamil Nadu), Nadiad (Gujarat); QA/OC in centralized location in Dwarka, New Delhi
		East Africa	No	°Z	Yes	No	Conventional PCR	Asexual	18S rRNA	Onsite in Kampala
		Amazonia	Not applicable	Not applicable	No	No	Conventional PCR; qPCR; RT-PCR	Asexual and sexual	18S rRNA	Onsite in Iquitos (Loreto) and Lima (Lima Province)
			Performed RDT efficacy study	Genotyping of parasite population for mutations that may impede accurate RDT diagnosis	Use as a diagnostic tool	Used as point- of-care diagnosis	Molecular diagnostic(s) used	Morphological stage-specific molecular diagnostic(s)	Target	Assays performed on-site or in centralized location
					Molecular diagnosis					

MALARIA DIAGNOSIS ACROSS THE ICEMR

HRP2 = histidine-rich protein 2: LAMP = loop-mediated isothermal amplification; PLDH = *Plusmodium* lactate dehydrogenase; QA = quality assurance; QC = quality control; qPCR = quantitative polymerase chain reaction; RDT = rapid diagnostic test; RT-PCR = reverse transcriptase polymerase chain reaction; WHO = World Health Organization.

KOBAYASHI AND OTHERS

	DNA extraction methods for various starting materials					
Starting material	Method	Reference				
Whole blood	GTC preparation with subsequent phenol:chloroform extraction	44				
	Rapid boiling method	45				
	PURE method (followed by LAMP)	46				
	Commercial kits (<i>Oiagen DNeasy Blood and Tissue Kit, OIA Amp Kit</i> , bioMérieux easyMag [®] , Abbott m2000)	47,48				
Dried blood spot	Chelex boiling					
1	Commercial kits (<i>Oiagen DNeasy Blood and Tissue Kit, OIA Amp Kit</i> , bioMérieux easyMag [®])	53,54				
	Tris-EDTA protocol	55				
	PURE method (followed by LAMP)	46				
Thick blood smear	Qiagen mini kit with minor modification	56				
RDT cassette	Boiling nitrocellulose component of the RDT strip in molecular grade water	57				
Urine/saliva	Qiagen DNeasy Kit	58				

TABLE 3

EDTA = ethylenediaminetetraacetic acid; GTC = guanidine isothiocyanate; LAMP = loop-mediated isothermal amplification; RDT = rapid diagnostic test. Method reported to be in use by ICEMRs are given in italics.

(Table 3). The methods that were reported to be used by ICEMRs are shown in italics in Table 3. The factors influencing the sensitivity of molecular diagnostics, particularly for conventional PCR, are reviewed in brief, but individual ICEMR protocols are not detailed here.

Dried blood spot (DBS) sampling is commonly used to preserve whole blood samples because it is an easy, durable method with an efficient use of space and optimal transport conditions. In research setting, DBSs have been used as starting material for PCR, RT-PCR, and serology.⁵⁹⁻⁶¹ Four ICEMRs have collected DBSs and whole blood as packed red cells, five ICEMRs collected only DBS, and one ICEMR collected only whole blood as packed red cells but not DBSs. DNA extraction differed among ICEMRs. Three of the 10 ICEMRs used a Chelex-based extraction method⁴⁹; the remaining seven ICEMRs used commercial DNA extraction kits, particularly individual spin column-type kits from Qiagen (Hilden, Germany) (Table 3). When DBSs were used as starting material for DNA extraction, the number/volume of blood spots varied between ICEMRs and the reported amount varied between using a partial blood spot to using more than one blood spot. One full blood spot is thought to typically correspond to approximately 50 µL blood. The available sample volume is affected by how blood is collected, whether by finger prick or venipuncture. With venipuncture, a higher volume of blood is typically available, therefore the starting sample volume is more standardized.

HRP2/3 deletion among the circulating parasites. HRP2 is a protein found only in P. falciparum asexual stages and young gametocyte stages. Hence HRP2 has been used in RDTs to detect P. falciparum.⁶² Plasmodium LDH (pLDH), from the glycolytic pathway, is found in all malaria species and is the second most commonly used RDT antigen.⁶² A third target antigen for RDTs is aldolase, another glycolytic enzyme.⁶² Several studies have compared RDTs based on these antigens, with different results in terms of sensitivity, specificity, and positive and negative predictive values.^{63,64} The majority of RDT experience in the field setting is in the detection of *P. falciparum*.

RDTs are particularly useful where microscopy is not available, whether because of financial constraints or lack of equipment or sufficiently trained personnel. RDT performance in the field is influenced by many factors such as manufacturing quality, handling and storage conditions, user interpretation, parasite density and limits of detection (which partly depend on parasite biology, e.g., P. vivax parasitemia is characteristically low), and the recent discovery of some region-specific field isolates lacking the HRP2 protein. In 2010, P. falciparum lacking pfhrp2 and/or pfhrp3 genes were first isolated from infected human subjects in the Amazon region of Peru.⁶⁵ Other endemic regions also reported false negative results using RDTs based on PfHRP2 due to pfhrp2 and/or pfhrp3 gene deletions.^{66–68} The presence of pfhrp2/3 deletion in P. falciparum strains has serious implications for diagnosis especially for countries and regions where the antimalarial treatment is primarily based on RDT diagnosis. Therefore, it is important to monitor the prevalence of pfhrp2 and/or pfhrp3 gene deletions in regions where RDTs based on PfHRP2 are the primary mode of malaria diagnosis, given the potential need to base procurement decisions on specific RDT detection proteins.

Peru was the first country where pfhrp2- and/or pfhrp3lacking P. falciparum was identified in infected patients.65 These data indicated that the deletion was widespread throughout the Peruvian Amazon region, which had major implications for miscalculated endemicity based solely on RDT detection.⁶⁵ The *pfhrp3* deletion was the most prevalent (70%) compared to pfhrp2 (41%), and 22% of isolates had both genes deleted.⁶⁵ Peru is not likely the only country affected, as the Amazon River basin runs through other malaria-endemic countries, including Colombia, Ecuador, Brazil, Venezuela, Guyana, Suriname, and Bolivia. Recent reports from these areas showed variable results with regard to the presence/absence of pfhrp2 and/or pfhrp3.69-71 Plasmodium falciparum with the double deletion of pfhrp2 and pfhrp3 was found in a patient who returned to Europe from Brazil and misdiagnosed using an RDT based on aldolase and PfHRP2; this patient was treated as having P. vivax.⁶⁹ Colombia also reported two P. falciparum isolates from clinically infected subjects; however, it was not possible to amplify exon 2 for both pfhrp2 and pfhrp3.⁷⁰ In a recent survey in communities around Iquitos-Nauta road in the Peruvian Amazon region, it was difficult to find P. falciparum PfHRP2-positive individuals even though they were diagnosed positive by light microscopy and pLDH (D. Gamboa, A. Llanos, and J. M. Vinetz, unpublished data). Thus far, Guyana is the only country in South America where all P. falciparum isolates have been found to carry pfhrp2; therefore, to date, the performance of PfHRP2based RDTs here has not yet been affected.⁷¹

RDT TO CHECK OUALITY OF ARTEMISININ CLASS ANTIMALARIALS

Because they are relatively expensive and widely used, ACTs have become a target of counterfeiters.^{22,23} Counterfeit artemisinins have the potential to be a public health and clinical threat because they contain little to no active ingredient hence providing inadequate treatment.⁷² Substandard drugs not only compromise the expected therapeutic effects, but also facilitate the development of resistance.⁷³ Although counterfeit artemisinins are known to be a problem in the Greater Mekong subregion of southeast Asia,74-80 the Worldwide Antimalarial Resistance Network database has reported that poor-quality artemisinins are also a growing concern in Africa.⁸¹⁻⁸⁵ Counterfeit and substandard antimalarials are an immediate and urgent threat to the current momentum of global malaria control and elimination efforts. Artemisinin counterfeiting is becoming more sophisticated. Thus far, as many as 14 different formulations of fake artesunate have been identified.^{72,86,87} To combat the entry of both counterfeit and substandard artemisinin compounds, both national and international regulatory authorities need to strengthen drug quality monitoring to deter the introduction and circulation of poor-quality antimalarials. Until recently, the accurate determination of artemisinin contents in commercial drugs has required sophisticated instrumentation and expertise. The recent development of simple and fieldapplicable tests to determine the quality of artemisinin drugs is being investigated.

Given that workers in malaria-endemic populations are very familiar with using RDTs for malaria diagnosis, the southeast Asia ICEMR chose to develop a lateral flow dipstick as the point-of-care tests for qualitative and semiquantitative detection of artemisinins in antimalarial drugs. Specific antibodies are needed for such dipstick assays, hence new monoclonal antibodies have been made for such efforts, which include the differentiation of commonly used artemisinin derivatives including artemether, artesunate, and dihydroartemisinin. The first approach was to immunize mice with artesunate conjugated to bovine serum albumin at the succinate group, which yielded a monoclonal antibody (mAb) that reacted with artemisinin, artesunate, and dihydroartemisinin, but with limited reactivity to artemether, 88,89 and a separate mAb that was specific for only artesunate (L. Cui, unpublished data). To achieve specificity for different artemisinin derivatives, which differ in R-groups at position 12, conjugation of the artemisinin derivatives to a carrier protein at the opposite position is needed. For artemether, this is achieved through microbial fermentation of artemether and purification of 9-hydroxylartemether.⁹⁰ A prototype dipstick was designed based on one selected mAb that was found to have high avidity and broad reactivity for artemisinins, with sensitivity as low as 100-200 and 200-500 ng/mL for artesunate and dihydroartemisinin, respectively.⁹¹ Testing these new tools for monitoring quality of artemisinin drugs is underway. Having specific mAbs in hand for most artemisinin derivatives also allows for enzyme-linked immunosorbent assay quantification.

QA/QC SYSTEM FOR DIAGNOSTIC SYSTEMS AND IMPLEMENTATION IN ICEMR

All aspects of malaria control and elimination must have robust quality management and control standards. The concept of quality management for malaria diagnostics covers the entire process including establishment of standard operating procedures for each method and technique used in diagnosis, especially sample tracking and handling, to achieve accurate and reliable test results and reporting. QA aims at improving and standardizing each component of this complicated process to minimize or avoid unreliable results. QC emphasizes test accuracy and precision.⁹² The goal of QC in diagnosis is to detect, identify, evaluate, and correct errors due to technical failure, environmental conditions, and/or human error.

ICEMR sites have established several quality control strategies, most of which aim to comply with WHO guidelines (Table 2).^{42,93} In all ICEMR sites, slides are read by two microscopists and a third reader in case of discrepancy (Table 2). Discrepancies in slide reading are usually due to differences in parasite detection limits, species identification, and parasite quantification. In most cases, if there is a disagreement in parasite counts of more than 20% between the two readers, a third reader is consulted to resolve discrepancies.⁴² In some areas, more stringent rules may apply; for example in the Peruvian ICEMR the acceptable discrepancy is $\leq 5\%$.

To maintain microscopy quality, refresher training programs for microscopists using reference slides as recommended by WHO were reported.^{42,93} For any malaria diagnostic laboratory, one way to promote standardization is to request that slide readers pass the WHO accreditation course and to implement regular refresher training and continuing assessments for microscopists. Another QC tactic to help assure microscopy result quality is to have slides reexamined by microscopists at an external reference laboratory or a collaborating institution.

For molecular diagnosis, the inclusion of well-characterized negative and positive controls (including serial dilutions of positive controls for qPCR) was reported from all ICEMRs. Protocols to freshly extract positive and negative controls to assure efficiency and lack of contamination in processed samples remain to be standardized. WHO provides international standards for *P. falciparum* DNA,⁹⁴ and the Malaria Research and Reference Reagent Resource Center also provides useful reagents and other resources. To verify PCR results, some ICEMRs reported sending a fraction of the samples to be reanalyzed at a reference laboratory or a collaborating institution (i.e., Latin America ICEMR and Amazonia ICEMR). Specificity may be increased by performing more than one PCR assay for the identification of sub-patent infection. Although a formal external QA/QC program is not yet available, various attempts to establish such processes have been made, especially for clinical trials and eradication surveil-lance.^{95,96} As qPCR and other more advanced technologies are increasingly used in malaria epidemiology, there is a need to ensure reproducibility of data. A recent article compared different protocols using the WHO International Standard for P. falciparum DNA,⁹⁷ leading to a guideline for the minimum information for publication of real-time qPCR experiment. In such a way, the reproducibility of the protocols across different laboratories can be improved.98

OVERCOMING KEY GAPS

Although WHO has published standardized protocols for microscopy and RDTs, no such protocols exist for nucleic acid amplification-based diagnostic tests such as PCR. Each ICEMR site has established conventional PCR; variation in details of sample preparation (starting material volume, nucleic acid extraction methods) remains a challenge for effective comparisons of diagnostic and treatment outcomes data. Other molecular diagnostic tools, such as qPCR, LAMP, and other sensitive techniques, are either not yet available in all the ICEMR regions or were not selected as part of the research design.

International standardization of diagnostic platforms in settings and research networks such as the ICEMR will be tedious. However, this process will include routine technical training for technicians, monitoring of diagnostic accuracy by external QC procedures, outreach training, reference laboratory support, and regular proficiency testing to ensure comparable results among laboratories. An external QA plan is important to assure valid diagnostic results and accurate reporting. Harmonization of PCR procedures more recently has taken on higher priority given the increasing importance of molecular diagnostics in the ICEMR network and elsewhere in infectious diseases epidemiology, clinical care and research. Access to reference laboratories to assure accurate and available QA/QC in both malaria microscopy and PCR should be readily available, as a matter of public health policy within malaria-endemic regions.

Although this article focused on different diagnostic platforms and protocol standardization, being able to harmonize data across platforms is essential, in the case of the ICEMRs, enabling comparable case definitions across the ICEMRs for which diagnostics lead to interventions, whether for disease or surveillance.

Received January 2, 2015. Accepted for publication June 23, 2015.

Published online August 10, 2015.

Acknowledgments: We appreciate the critical reading of the manuscript by Andres Vallejo, Sócrates Herrera, and Don Krogstad. We thank Trevor A. Thompson for information on PCR sensitivity and slide reading quality assurance practice at the West Africa ICEMR. We also thank all the ICEMRs for contributing information and acknowledge Neena Valecha for her guidance regarding the RDTs used in India and India ICEMR project.

Financial support: This work was supported by the following Cooperative Agreements from the United States Public Health Service, National Institute of Allergy and Infectious Diseases: U19AI089672, U19AI089674, 5U19AI089676, U19AI089680, U19AI089681, U19AI089683, U19AI089686, U19AI089686, U19AI089696, and U19AI089702.

Authors' addresses: Tamaki Kobayashi, Department of Epidemiology, Johns Hopkins Bloomberg School of Public Health, Baltimore, MD, E-mail: tkobaya2@jhu.edu. Dionicia Gamboa, Universidad Peruana Cayetano Heredia, Lima, Peru, E-mail: dionigamboa@yahoo.com. Daouda Ndiaye, Department of Parasitology and Mycology, Université Cheikh Anta Diop de Dakar, Dakar, Senegal, E-mail: dndiaye@hsph .harvard.edu. Liwang Cui, Department of Entomology, Pennsylvania State University, University Park, PA, E-mail: luc2@psu.edu. Patrick Sutton, Acsel Health, New York, NY, E-mail: suttop02@gmail.com. Joseph M. Vinetz, Division of Infectious Diseases, Department of Medicine, University of California San Diego School of Medicine, La Jolla, CA, and Universidad Peruana Cayetano Heredia, Lima, Peru, E-mails: jvinetz@ucsd.edu or joseph.vinetz@upch.pe.

This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

REFERENCES

 Murphy SC, Shott JP, Parikh S, Etter P, Prescott WR, Stewart VA, 2013. Malaria diagnostics in clinical trials. *Am J Trop Med Hyg 89*: 824–839.

- 2. WHO, 2010. *Guidelines for the Treatment of Malaria*, 2nd edition. Geneva, Switzerland: World Health Organization.
- Okell LC, Bousema T, Griffin JT, Ouedraogo AL, Ghani AC, Drakeley CJ, 2012. Factors determining the occurrence of submicroscopic malaria infections and their relevance for control. *Nat Commun 3*: 1237.
- Tietje K, Hawkins K, Clerk C, Ebels K, McGray S, Crudder C, Okell L, LaBarre P, 2014. The essential role of infectiondetection technologies for malaria elimination and eradication. *Trends Parasitol 30:* 259–266.
- Bousema T, Okell L, Felger I, Drakeley C, 2014. Asymptomatic malaria infections: detectability, transmissibility and public health relevance. *Nat Rev Microbiol* 12: 833–840.
- Lin JT, Saunders DL, Meshnick SR, 2014. The role of submicroscopic parasitemia in malaria transmission: what is the evidence? *Trends Parasitol 30*: 183–190.
- Alonso PL, Tanner M, 2013. Public health challenges and prospects for malaria control and elimination. *Nat Med 19:* 150–155.
- Vasoo S, Pritt BS, 2013. Molecular diagnostics and parasitic disease. *Clin Lab Med 33*: 461–503.
- Imwong M, Hanchana S, Malleret B, Renia L, Day NP, Dondorp A, Nosten F, Snounou G, White NJ, 2014. Highthroughput ultrasensitive molecular techniques for quantifying low-density malaria parasitemias. *J Clin Microbiol* 52: 3303–3309.
- Hofmann N, Mwingira F, Shekalaghe S, Robinson LJ, Mueller I, Felger I, 2015. Ultra-sensitive detection of *Plasmodium falciparum* by amplification of multi-copy subtelomeric targets. *PLoS Med* 12: e1001788.
- Oriero EC, Jacobs J, Van Geertruyden JP, Nwakanma D, D'Alessandro U, 2015. Molecular-based isothermal tests for field diagnosis of malaria and their potential contribution to malaria elimination. J Antimicrob Chemother 70: 2–13.
- Paris DH, Imwong M, Faiz AM, Hasan M, Yunus EB, Silamut K, Lee SJ, Day NP, Dondorp AM, 2007. Loop-mediated isothermal PCR (LAMP) for the diagnosis of falciparum malaria. *Am J Trop Med Hyg* 77: 972–976.
- Abdul-Ghani R, Al-Mekhlafi AM, Karanis P, 2012. Loopmediated isothermal amplification (LAMP) for malarial parasites of humans: would it come to clinical reality as a pointof-care test? *Acta Trop 122*: 233–240.
- Schoone GJ, Oskam L, Kroon NC, Schallig HD, Omar SA, 2000. Detection and quantification of *Plasmodium falciparum* in blood samples using quantitative nucleic acid sequencebased amplification. *J Clin Microbiol* 38: 4072–4075.
- Mens PF, Schoone GJ, Kager PA, Schallig HD, 2006. Detection and identification of human *Plasmodium* species with realtime quantitative nucleic acid sequence-based amplification. *Malar J 5:* 80.
- Nayyar GM, Breman JG, Herrington JE, 2015. The global pandemic of falsified medicines: laboratory and field innovations and policy perspectives. *Am J Trop Med Hyg* 92: 2–7.
- 17. White NJ, Pukrittayakamee S, Hien TT, Faiz MA, Mokuolu OA, Dondorp AM, 2014. Malaria. *Lancet 383:* 723–735.
- Price RN, von Seidlein L, Valecha N, Nosten F, Baird JK, White NJ, 2014. Global extent of chloroquine-resistant *Plasmodium vivax*: a systematic review and meta-analysis. *Lancet Infect Dis* 14: 982–991.
- 19. WHO, 2012. World Malaria Report 2012. Geneva, Switzerland: World Health Organization.
- Baird JK, 2011. Resistance to chloroquine unhinges vivax malaria therapeutics. Antimicrob Agents Chemother 55: 1827–1830.
- Douglas NM, Anstey NM, Angus BJ, Nosten F, Price RN, 2010. Artemisinin combination therapy for vivax malaria. *Lancet Infect Dis* 10: 405–416.
- Nayyar GM, Breman JG, Newton PN, Herrington J, 2012. Poorquality antimalarial drugs in southeast Asia and sub-Saharan Africa. *Lancet Infect Dis* 12: 488–496.
- Karunamoorthi K, 2014. The counterfeit anti-malarial is a crime against humanity: a systematic review of the scientific evidence. *Malar J* 13: 209.
- 24. Koram KA, Molyneux ME, 2007. When is "malaria" malaria? The different burdens of malaria infection, malaria disease, and malaria-like illnesses. *Am J Trop Med Hyg* 77: 1–5.

- Shanks GD, 2012. Control and elimination of *Plasmodium vivax*. Adv Parasitol 80: 301–341.
- Baird JK, 2008. Real-world therapies and the problem of vivax malaria. N Engl J Med 359: 2601–2603.
- 27. Ingram RJ, Crenna-Darusallam C, Soebianto S, Noviyanti R, Baird JK, 2014. The clinical and public health problem of relapse despite primaquine therapy: case review of repeated relapses of *Plasmodium vivax* acquired in Papua New Guinea. *Malar J 13*: 488.
- 28. Llanos-Cuentas A, Lacerda MV, Rueangweerayut R, Krudsood S, Gupta SK, Kochar SK, Arthur P, Chuenchom N, Mohrle JJ, Duparc S, Ugwuegbulam C, Kleim JP, Carter N, Green JA, Kellam L, 2014. Tafenoquine plus chloroquine for the treatment and relapse prevention of *Plasmodium vivax* malaria (DETECTIVE): a multicentre, double-blind, randomised, phase 2b dose-selection study. *Lancet* 383: 1049–1058.
- Lell B, Faucher JF, Missinou MA, Borrmann S, Dangelmaier O, Horton J, Kremsner PG, 2000. Malaria chemoprophylaxis with tafenoquine: a randomised study. *Lancet 355:* 2041–2045.
- Kondrashin A, Baranova AM, Ashley EA, Recht J, White NJ, Sergiev VP, 2014. Mass primaquine treatment to eliminate vivax malaria: lessons from the past. *Malar J* 13: 51.
- Poirot E, Skarbinski J, Sinclair D, Kachur SP, Slutsker L, Hwang J, 2013. Mass drug administration for malaria. *Cochrane Database Syst Rev 12:* CD008846.
- Galappaththy GN, Tharyan P, Kirubakaran R, 2013. Primaquine for preventing relapse in people with *Plasmodium vivax* malaria treated with chloroquine. *Cochrane Database Syst Rev* 10: CD004389.
- 33. von Seidlein L, Auburn S, Espino F, Shanks D, Cheng Q, McCarthy J, Baird K, Moyes C, Howes R, Menard D, Bancone G, Winasti-Satyahraha A, Vestergaard LS, Green J, Domingo G, Yeung S, Price R, 2013. Review of key knowledge gaps in glucose-6-phosphate dehydrogenase deficiency detection with regard to the safe clinical deployment of 8-aminoquinoline treatment regimens: a workshop report. *Malar J 12*: 112.
- Wongsrichanalai C, Barcus MJ, Muth S, Sutamihardja A, Wernsdorfer WH, 2007. A review of malaria diagnostic tools: microscopy and rapid diagnostic test (RDT). Am J Trop Med Hyg 77: 119–127.
- 35. Mayxay M, Pukrittayakamee S, Chotivanich K, Looareesuwan S, White NJ, 2001. Persistence of *Plasmodium falciparum* HRP-2 in successfully treated acute falciparum malaria. *Trans R Soc Trop Med Hyg 95*: 179–182.
- 36. Iqbal J, Siddique A, Jameel M, Hira PR, 2004. Persistent histidinerich protein 2, parasite lactate dehydrogenase, and panmalarial antigen reactivity after clearance of *Plasmodium falciparum* monoinfection. J Clin Microbiol 42: 4237–4241.
- 37. Snounou G, Viriyakosol S, Jarra W, Thaithong S, Brown KN, 1993. Identification of the four human malaria parasite species in field samples by the polymerase chain reaction and detection of a high prevalence of mixed infections. *Mol Biochem Parasitol 58:* 283–292.
- Mahajan B, Zheng H, Pham PT, Sedegah MY, Majam VF, Akolkar N, Rios M, Ankrah I, Madjitey P, Amoah G, Addison E, Quakyi IA, Kumar S, 2012. Polymerase chain reactionbased tests for pan-species and species-specific detection of human *Plasmodium* parasites. *Transfusion* 52: 1949–1956.
- 39. Mangold KA, Manson RU, Koay ES, Stephens L, Regner M, Thomson RB Jr, Peterson LR, Kaul KL, 2005. Real-time PCR for detection and identification of *Plasmodium* spp. *J Clin Microbiol* 43: 2435–2440.
- 40. Demas A, Oberstaller J, DeBarry J, Lucchi NW, Srinivasamoorthy G, Sumari D, Kabanywanyi AM, Villegas L, Escalante AA, Kachur SP, Barnwell JW, Peterson DS, Udhayakumar V, Kissinger JC, 2011. Applied genomics: data mining reveals species-specific malaria diagnostic targets more sensitive than 18S rRNA. J Clin Microbiol 49: 2411–2418.
- 41. Rantala AM, Taylor SM, Trottman PA, Luntamo M, Mbewe B, Maleta K, Kulmala T, Ashorn P, Meshnick SR, 2010. Comparison of real-time PCR and microscopy for malaria parasite detection in Malawian pregnant women. *Malar J* 9: 269.
- WHO, 2009. Malaria Microscopy Quality Assurance Manual– Version 1. Geneva, Switzerland: World Health Organization.

- 43. FIND, 2013. Malaria Rapid Diagnostic Test Performance— Results of WHO Product Testing of Malaria RDTs: Round 5. Geneva, Switzerland: World Health Organization.
- Henning L, Felger I, Beck HP, 1999. Rapid DNA extraction for molecular epidemiological studies of malaria. Acta Trop 72: 149–155.
- 45. Foley M, Ranford-Cartwright LC, Babiker HA, 1992. Rapid and simple method for isolating malaria DNA from fingerprick samples of blood. *Mol Biochem Parasitol* 53: 241–244.
- FIND, 2012. Manual of Standard Operating Procedures for Malaria LAMP. Available at: http://www.finddiagnostics.org/export/ sites/default/programs/malaria-afs/docs/SOPs_LAMP_Malaria_ AUG12.pdf.
- Pillet S, Bourlet T, Pozzetto B, 2012. Comparative evaluation of the QIAsymphony RGQ system with the easyMAG/R-gene combination for the quantitation of cytomegalovirus DNA load in whole blood. *Virol J 9*: 231.
- Gomes P, Carvalho AP, Diogo I, Goncalves F, Costa I, Cabanas J, Camacho RJ, 2013. Comparison of the NucliSENS EasyQ HIV-1 v2.0 with Abbott m2000rt RealTime HIV-1 assay for plasma RNA quantitation in different HIV-1 subtypes. J Virol Methods 193: 18–22.
- 49. Kain KC, Lanar DE, 1991. Determination of genetic variation within *Plasmodium falciparum* by using enzymatically amplified DNA from filter paper disks impregnated with whole blood. *J Clin Microbiol 29*: 1171–1174.
- Wooden J, Kyes S, Sibley CH, 1993. PCR and strain identification in *Plasmodium falciparum. Parasitol Today 9*: 303–305.
- Cox-Singh J, Mahayet S, Abdullah MS, Singh B, 1997. Increased sensitivity of malaria detection by nested polymerase chain reaction using simple sampling and DNA extraction. *Int J Parasitol 27:* 1575–1577.
- 52. Chaorattanakawee S, Natalang O, Hananantachai H, Nacher M, Brockman A, Krudsood S, Looareesuwan S, Patarapotikul J, 2003. Storage duration and polymerase chain reaction detection of *Plasmodium falciparum* from blood spots on filter paper. *Am J Trop Med Hyg 69*: 42–44.
- 53. Mercier-Delarue S, Vray M, Plantier JC, Maillard T, Adjout Z, de Olivera F, Schnepf N, Maylin S, Simon F, Delaugerre C, 2014. Higher specificity of nucleic acid sequence-based amplification isothermal technology than of real-time PCR for quantification of HIV-1 RNA on dried blood spots. *J Clin Microbiol 52:* 52–56.
- 54. Lofgren SM, Morrissey AB, Chevallier CC, Malabeja AI, Edmonds S, Amos B, Sifuna DJ, von Seidlein L, Schimana W, Stevens WS, Bartlett JA, Crump JA, 2009. Evaluation of a dried blood spot HIV-1 RNA program for early infant diagnosis and viral load monitoring at rural and remote healthcare facilities. *AIDS 23*: 2459–2466.
- 55. Bereczky S, Martensson A, Gil JP, Farnert A, 2005. Short report: rapid DNA extraction from archive blood spots on filter paper for genotyping of *Plasmodium falciparum*. *Am J Trop Med Hyg* 72: 249–251.
- Cnops L, Van Esbroeck M, Bottieau E, Jacobs J, 2010. Giemsastained thick blood films as a source of DNA for *Plasmodium* species-specific real-time PCR. *Malar J 9*: 370.
- Cnops L, Boderie M, Gillet P, Van Esbroeck M, Jacobs J, 2011. Rapid diagnostic tests as a source of DNA for *Plasmodium* species-specific real-time PCR. *Malar J 10*: 67.
- Mharakurwa S, Simoloka C, Thuma PE, Shiff CJ, Sullivan DJ, 2006. PCR detection of *Plasmodium falciparum* in human urine and saliva samples. *Malar J 5*: 103.
- 59. Plowe CV, Djimde A, Bouare M, Doumbo O, Wellems TE, 1995. Pyrimethamine and proguanil resistance-conferring mutations in *Plasmodium falciparum* dihydrofolate reductase: polymerase chain reaction methods for surveillance in Africa. *Am J Trop Med Hyg 52*: 565–568.
- 60. Mlambo G, Vasquez Y, LeBlanc R, Sullivan D, Kumar N, 2008. A filter paper method for the detection of *Plasmodium falciparum* gametocytes by reverse transcription polymerase chain reaction. *Am J Trop Med Hyg 78*: 114–116.
- Corran PH, Cook J, Lynch C, Leendertse H, Manjurano A, Griffin J, Cox J, Abeku T, Bousema T, Ghani AC, Drakeley C, Riley E, 2008. Dried blood spots as a source of anti-malarial antibodies for epidemiological studies. *Malar J* 7: 195.

- Moody A, 2002. Rapid diagnostic tests for malaria parasites. Clin Microbiol Rev 15: 66–78.
- Mouatcho JC, Goldring JP, 2013. Malaria rapid diagnostic tests: challenges and prospects. J Med Microbiol 62: 1491–1505.
- 64. Maltha J, Guiraud I, Lompo P, Kabore B, Gillet P, Van Geet C, Tinto H, Jacobs J, 2014. Accuracy of PfHRP2 versus Pf-pLDH antigen detection by malaria rapid diagnostic tests in hospitalized children in a seasonal hyperendemic malaria transmission area in Burkina Faso. *Malar J 13*: 20.
- 65. Gamboa D, Ho MF, Bendezu J, Torres K, Chiodini PL, Barnwell JW, Incardona S, Perkins M, Bell D, McCarthy J, Cheng Q, 2010. A large proportion of *P. falciparum* isolates in the Amazon region of Peru lack *pfhrp2* and *pfhrp3*: implications for malaria rapid diagnostic tests. *PLoS One* 5: e8091.
- 66. Koita OA, Doumbo OK, Ouattara A, Tall LK, Konare A, Diakite M, Diallo M, Sagara I, Masinde GL, Doumbo SN, Dolo A, Tounkara A, Traore I, Krogstad DJ, 2012. Falsenegative rapid diagnostic tests for malaria and deletion of the histidine-rich repeat region of the *hrp2* gene. *Am J Trop Med Hyg* 86: 194–198.
- 67. Kumar N, Pande V, Bhatt RM, Shah NK, Mishra N, Srivastava B, Valecha N, Anvikar AR, 2013. Genetic deletion of HRP2 and HRP3 in Indian *Plasmodium falciparum* population and false negative malaria rapid diagnostic test. *Acta Trop 125:* 119–121.
- 68. Wurtz N, Fall B, Bui K, Pascual A, Fall M, Camara C, Diatta B, Fall KB, Mbaye PS, Dieme Y, Bercion R, Wade B, Briolant S, Pradines B, 2013. *Pfhrp2* and *pfhrp3* polymorphisms in *Plasmodium falciparum* isolates from Dakar, Senegal: impact on rapid malaria diagnostic tests. *Malar J 12*: 34.
- 69. Houze S, Hubert V, Le Pessec G, Le Bras J, Clain J, 2011. Combined deletions of *pfhrp2* and *pfhrp3* genes result in *Plasmodium falciparum* malaria false-negative rapid diagnostic test. J Clin Microbiol 49: 2694–2696.
- Mendoza NM, Cucunuba ZM, Aponte S, Gonzalez NE, Bernal SD, 2013. Field evaluation for diagnostic accuracy of the rapid test SD Bioline Malaria Antigen Pf/Pv(R) in Colombia. *Biomedica* 33: 587–597.
- 71. Trouvay M, Palazon G, Berger F, Volney B, Blanchet D, Faway E, Donato D, Legrand E, Carme B, Musset L, 2013. High performance of histidine-rich protein 2 based rapid diagnostic tests in French Guiana are explained by the absence of *pfhrp2* gene deletion in *P. falciparum. PLoS One 8:* e74269.
- 72. Newton PN, McGready R, Fernandez F, Green MD, Sunjio M, Bruneton C, Phanouvong S, Millet P, Whitty CJ, Talisuna AO, Proux S, Christophel EM, Malenga G, Singhasivanon P, Bojang K, Kaur H, Palmer K, Day NP, Greenwood BM, Nosten F, White NJ, 2006. Manslaughter by fake artesunate in Asia—will Africa be next? *PLoS Med 3*: e197.
- 73. Keoluangkhot V, Green MD, Nyadong L, Fernandez FM, Mayxay M, Newton PN, 2008. Impaired clinical response in a patient with uncomplicated falciparum malaria who received poor-quality and underdosed intramuscular artemether. Am J Trop Med Hyg 78: 552–555.
- 74. Dondorp AM, Newton PN, Mayxay M, Van Damme W, Smithuis FM, Yeung S, Petit A, Lynam AJ, Johnson A, Hien TT, McGready R, Farrar JJ, Looareesuwan S, Day NP, Green MD, White NJ, 2004. Fake antimalarials in southeast Asia are a major impediment to malaria control: multinational cross-sectional survey on the prevalence of fake antimalarials. *Trop Med Int Health 9:* 1241–1246.
- 75. Vijaykadga S, Cholpol S, Sitthimongkol S, Pawaphutanan A, Pinyoratanachot A, Rojanawatsirivet C, Kovithvattanapong R, Thimasarn K, 2006. Strengthening of national capacity in implementation of antimalarial drug quality assurance in Thailand. Southeast Asian J Trop Med Public Health 37 (Suppl 3): 5–10.
- 76. Lon CT, Tsuyuoka R, Phanouvong S, Nivanna N, Socheat D, Sokhan C, Blum N, Christophel EM, Smine A, 2006. Counterfeit and substandard antimalarial drugs in Cambodia. *Trans R* Soc Trop Med Hyg 100: 1019–1024.
- Rozendaal J, 2001. Fake antimalaria drugs in Cambodia. Lancet 357: 890.

- Newton P, Proux S, Green M, Smithuis F, Rozendaal J, Prakongpan S, Chotivanich K, Mayxay M, Looareesuwan S, Farrar J, Nosten F, White NJ, 2001. Fake artesunate in southeast Asia. *Lancet* 357: 1948–1950.
- Newton PN, Dondorp A, Green M, Mayxay M, White NJ, 2003. Counterfeit artesunate antimalarials in southeast Asia. *Lancet* 362: 169.
- Hall KA, Newton PN, Green MD, De Veij M, Vandenabeele P, Pizzanelli D, Mayxay M, Dondorp A, Fernandez FM, 2006. Characterization of counterfeit artesunate antimalarial tablets from southeast Asia. *Am J Trop Med Hyg* 75: 804–811.
- Bate R, Coticelli P, Tren R, Attaran A, 2008. Antimalarial drug quality in the most severely malarious parts of Africa—a six country study. *PLoS One 3:* e2132.
- Atemnkeng MA, De Cock K, Plaizier-Vercammen J, 2007. Quality control of active ingredients in artemisinin-derivative antimalarials within Kenya and DR Congo. *Trop Med Int Health 12:* 68–74.
- Amin AA, Kokwaro GO, 2007. Antimalarial drug quality in Africa. J Clin Pharm Ther 32: 429–440.
- 84. Newton PN, Green MD, Mildenhall DC, Plancon A, Nettey H, Nyadong L, Hostetler DM, Swamidoss I, Harris GA, Powell K, Timmermans AE, Amin AA, Opuni SK, Barbereau S, Faurant C, Soong RC, Faure K, Thevanayagam J, Fernandes P, Kaur H, Angus B, Stepniewska K, Guerin PJ, Fernandez FM, 2011. Poor quality vital anti-malarials in Africa—an urgent neglected public health priority. *Malar J 10*: 352.
- Ambroise-Thomas P, 2012. The tragedy caused by fake antimalarial drugs. *Mediterr J Hematol Infect Dis 4*: e2012027.
- 86. Newton PN, Fernandez FM, Plancon A, Mildenhall DC, Green MD, Ziyong L, Christophel EM, Phanouvong S, Howells S, McIntosh E, Laurin P, Blum N, Hampton CY, Faure K, Nyadong L, Soong CW, Santoso B, Zhiguang W, Newton J, Palmer K, 2008. A collaborative epidemiological investigation into the criminal fake artesunate trade in south east Asia. *PLoS Med 5:* e32.
- Newton PN, Green MD, Fernandez F, 2007. Counterfeit artemisinin derivatives and Africa: update from authors. *PLoS Med 4:* e139.
- 88. He SP, Tan GY, Li G, Tan WM, Nan TG, Wang BM, Li ZH, Li QX, 2009. Development of a sensitive monoclonal antibody-based enzyme-linked immunosorbent assay for the antimalaria active ingredient artemisinin in the Chinese herb Artemisia annua L. Anal Bioanal Chem 393: 1297–1303.
- Wang M, Cui Y, Zhou G, Yan G, Cui L, Wang B, 2013. Validation of ELISA for quantitation of artemisinin-based antimalarial drugs. *Am J Trop Med Hyg 89*: 1122–1128.
- 90. Guo S, Cui Y, He L, Zhang L, Cao Z, Zhang W, Zhang R, Tan G, Wang B, Cui L, 2013. Development of a specific monoclonal antibody-based ELISA to measure the artemether content of antimalarial drugs. *PLoS One 8*: e79154.
- 91. He L, Nan T, Cui Y, Guo S, Zhang W, Zhang R, Tan G, Wang B, Cui L, 2014. Development of a colloidal gold-based lateral flow dipstick immunoassay for rapid qualitative and semi-quantitative analysis of artesunate and dihydroartemisinin. *Malar J 13*: 127.
- 92. Lourens C, Watkins WM, Barnes KI, Sibley CH, Guerin PJ, White NJ, Lindegardh N, 2010. Implementation of a reference standard and proficiency testing programme by the World Wide Antimalarial Resistance Network (WWARN). *Malar J* 9: 375.
- WHO, 2014. Technical Consultation to Update the WHO Malaria Microscopy Quality Assurance Manual. Geneva, Switzerland: World Health Organization.
- 94. Padley DJ, Heath AB, Sutherland C, Chiodini PL, Baylis SA, 2008. Establishment of the 1st World Health Organization International Standard for *Plasmodium falciparum* DNA for nucleic acid amplification technique (NAT)-based assays. *Malar J* 7: 139.
- 95. Taylor SM, Mayor A, Mombo-Ngoma G, Kenguele HM, Ouedraogo S, Ndam NT, Mkali H, Mwangoka G, Valecha N, Singh JP, Clark MA, Verweij JJ, Adegnika AA, Severini C, Menegon M, Macete E, Menendez C, Cistero P, Njie F, Affara M, Otieno K, Kariuki S, ter Kuile FO, Meshnick SR,

2014. A quality control program within a clinical trial consortium for PCR protocols to detect *Plasmodium* species. *J Clin Microbiol 52*: 2144–2149.

- 96. Murphy SC, Hermsen CC, Douglas AD, Edwards NJ, Petersen I, Fahle GA, Adams M, Berry AA, Billman ZP, Gilbert SC, Laurens MB, Leroy O, Lyke KE, Plowe CV, Seilie AM, Strauss KA, Teelen K, Hill AV, Sauerwein RW, 2014. External quality assurance of malaria nucleic acid testing for clinical trials and eradication surveillance. *PLoS One 9:* e97398.
- 97. Alemayehu S, Feghali KC, Cowden J, Komisar J, Ockenhouse CF, Kamau E, 2013. Comparative evaluation of published real-time PCR assays for the detection of malaria following MIQE guidelines. *Malar J 12:* 277.
- 98. Bustin SA, Benes V, Garson JA, Hellemans J, Huggett J, Kubista M, Mueller R, Nolan T, Pfaffl MW, Shipley GL, Vandesompele J, Wittwer CT, 2009. The MIQE guidelines: minimum information for publication of quantitative real-time PCR experiments. *Clin Chem* 55: 611–622.