

Analytical performance of the ScreenFire HPV RS Zebra BioDome assay on four different qPCR platforms

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Abstract

Objectives Cervical cancer is one of the most frequently diagnosed cancers and a leading cause of cancer-related deaths in women in low- and middle-income countries (LMICs), accounting for nearly 85% of the global cervical cancer burden. High-risk human papillomavirus (hrHPV) infection is the main cause of cervical cancer. Easy-to-use, rapid, scalable, high-throughput, and cost-effective HPV tests are urgently needed for low-resource settings. Atila Biosystems' clinically validated ScreenFire HPV Risk Stratification (RS) assay identifies 13 hrHPV in 4 groups based on their oncogenic risk (i.e., HPV16, HPV18/45, HPV31/33/35/52/58, and HPV51/59/39/56/68). While the current standard format is subject to laboratory contamination Atila has developed an innovative, contamination-preventive Zebra BioDome format. Recently we published the analytical performance of ScreenFire RS Zebra BioDome on the BioRad CFX-96 real-time PCR instrument. This current study evaluated its analytical performance on three additional qPCR platforms: Atila Portable iAMP-PS96, Atila Powergene9600 Plus, and Thermo Fisher Quantstudio-7.

Methods We tested 173 DNA samples from Nigerian women with cervical cancer. These samples were tested simultaneously using the ScreenFire HPV Zebra BioDome assay (M5FHPV-96) on four different real-time PCR machines (Atila portable iAMP-PS96, Atila Powergene9600 Plus, Thermo Fisher QuantStudio-7, and BioRad CFX-96). We used overall agreement rate and unweighted kappa values to compare different platforms.

Results The overall agreement for detection of hrHPV using Atila portable iAMP-PS96 was 96.5% with kappa value 0.95 (95% confidence interval: 0.91–0.99) compared to Thermo Fisher QuantStudio-7, and 97.1% with kappa value 0.96 (95% confidence interval: 0.92–0.99) compared to BioRad CFX-96. For genotype HPV16 and risk stratification (RS) genotype groups (HPV18/45, HPV31/33/35/52/58, and HPV51/59/39/56/68) agreement rates were all > 98.3%. For Atila Powergene9600 Plus the overall agreement was 98.8% with a kappa value of 0.98 (95% confidence interval: 0.96–1.0) compared to Thermo Fisher QuantStudio-7, and 96.5% with a kappa value of 0.96 (95% confidence

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interval: 0.94–0.99) compared to BioRad CFX-96. The agreements for the HPV16 and RS genotype groups (HPV18/45, HPV31/33/35/52/58, and HPV39/51/56/59/68) were at least 98.3%.

Conclusion The novel ScreenFire HPV Zebra BioDome format produced highly concordant hrHPV positivity and RS genotype results on all four qPCR platforms. The data suggests that this innovative technology has the potential to improve HPV testing uptake in low-resource settings without further investment in purchasing new equipment.

Keywords High-risk human papillomavirus (hr-HPV), ScreenFire HPV Zebra biodome assay, iAMP-PS96 personal station gPCR system, Powergene9600 plus Real-Time PCR system

Introduction

Globally, cervical cancer is one of the most frequently diagnosed cancers and a leading cause of cancer-related deaths in women in low- and middle-income countries (LMICs) which accounts for nearly 85% of the global cervical cancer burden [1, 2]. Cervical cancer is driven by the persistence of high-risk human papillomavirus (hrHPV) infection, the most significant risk factor for the development of cervical cancer [3, 4]. In 2022, the World Health Organization (WHO) released new guidance shifting primary cervical cancer screening recommendations to HPV DNA testing in all settings, and away from unaided visual inspection using acetic acid (VIA) [5]. Current HPV genotyping assays used in LMICs have a variety of limitations: they are time-consuming, labour-intensive, costly, lacking high-throughput capabilities, and at risk of laboratory contamination [6-8]. This latter risk is particularly common in PCR-based assays, and hinders the adoption of these assays for large-scale hrHPV testing in low-resource areas without a standard (i.e., costly) laboratory setup and available expert personnel.

Among the existing assays, Atila Biosystems' clinically validated ScreenFire HPV RS assay (M5FHPV-100) has been specifically designed for use in LMIC cervical cancer screening programs [9–11]. It provides hierarchal risk stratification (RS) genotyping information by identifying 13 hrHPV genotypes in 4 groups based on their oncogenic risk (i.e., HPV16, HPV18/45, HPV31/33/35/52/58, and HPV51/59/39/56/68). The ScreenFire HPV RS assay uses isothermal amplification and reports the four hrHPV channels using fluorescent detection, with channel sensitivity designed according to hierarchical cancer risk. It does so at a low cost per test, with high throughput and a reported overall sensitivity of 94.7%.^{9,10} The ScreenFire HPV RS assay offers high capacity in standard 96-well plates with less than one hour of processing time. In addition, it is particularly suited for primary cervical cancer screening since collected specimens do not require DNA extraction and purification [11]. However, the current format of the ScreenFire HPV RS assay still relies on manual preparation of the master mix of reagents [12-15], which requires a standard molecular laboratory setup and well-trained laboratory personnel to minimize possible laboratory contamination.

To address these challenges, Atila Biosystems has developed a new format for the ScreenFire HPV RS assay: the ScreenFire HPV RS assay Zebra BioDome (M5FHPV-96). This novel format features pre-loaded reagents in either 8-well PCR strips or 96-well PCR plates, covered by a temperature-sensitive hydrophobic gel matrix. The hydrogel matrix in the reagent tube is highly stable and semi-solidified during transportation and storage before full implementation. This protects the reagents from leakage or spillage and contaminating the laboratory. The matrix liquefies and moves to the top of the liquid to seal the reaction wells during isothermal heating and re-solidifies after amplification and before disposal. Thus, in addition to the common features shared with the ScreenFire Standard format (e.g. easy to use, high-throughput, cost-appropriate in low-resource settings, and no requirements for DNA extraction) the contamination-prevention feature along with the lack of required reagent preparation makes this Zebra BioDome format highly desirable. It only requires the single step of adding the patient's sample before the remaining automated process, thus making it particularly suitable for large-scale cervical cancer screening via primary HPV screening.

In our recently published data, we reported that the Zebra BioDome format generated highly concordant results compared to the standard ScreenFire HPV RS assay when the BioRad CFX-96 PCR instrument was used [12]. The overall agreement for detection of hrHPV was 96.0%. The agreement rates between hrHPV genotype 16 and risk stratification genotype group (HPV18/45, HPV31/33/35/52/58, and HPV51/59/39/56/68) were all >97.5%. The U.S. National Cancer Institute has also reported that the Zebra BioDome performed similarly to the standard version of the ScreenFire HPV assay using Atila Powergene Instrument [7]. Specifically, Zebra Bio-Dome showed agreement with the standard version in the channel-specific analysis with positive percent agreement between 88.4% and 100% and negative percent agreement between 97.8% and 100%, as well as in hierarchical analysis with overall agreement 97.2%. This validation strengthened the case for wider adoption of the ScreenFire BioDome assay. However, many laboratories in LMICs may already have other PCR instruments (e.g.,

the common Thermo Fisher QuantStudio-7 and Bio-Rad CFX-96). For this study we evaluated the analytical performance of the ScreenFire RS hrHPV assay on four qPCR platforms: Atila iAMP-PS96, Atila Powergene9600 Plus, Thermo Fisher Quantstudio-7, and BioRad CFX-96.

Materials and methods

Study samples

This study is built upon our NCI-funded U54 consortium to study Epigenomic Biomarkers of HIV-Associated Cancers in Nigeria (U54CA221205). In total, the 173 cervical tissue samples used in this study were collected between 2018 and 2022 in Nigerian women diagnosed with cervical cancer [16]. The age range was 26 to 80 years. The use of these samples was covered under IRB approval at Northwestern University, Jos University, and Lagos University. The samples were obtained at the Jos University Teaching Hospital (JUTH) and Lagos University Teaching Hospital (LUTH) in Nigeria. The further HPV testing was allowed in the informed consent and the results available to all women who underwent testing. DNA was extracted from the cervical biopsies using the Qiagen QIAamp DNA Mini Kit and quantified using Qubit 4.0 fluorometer. DNA samples were stored at -80 C until shipment. All DNA samples were de-identified when shipped in dry ice to Northwestern University and stored at – 80 C.

Detection and RS genotyping of HrHPV

A total of 100ng purified DNA was prepared in 100µL of 1X lysis buffer and processed following procedures for hrHPV RS genotyping using the ScreenFire RS Zebra BioDome HPV test kits purchased from Atila BioSystems, Inc (Atila, Sunnyvale, CA). The prepared 10 µl DNA samples were added into the prepacked Zebra Bio-Dome reaction tubes. The capped reaction tubes were spun for 10 s to bring all the liquid down to the bottom. The strips were then loaded into four different real-time PCR machines (Atila iAMP-PS96, Atila Powergene9600 Plus, Thermo Fisher QuantStudio-7 or BioRad CFX-96) and the assays were carried out on the isothermal program mode run at 1 min per cycle at 60° C for 60 cycles. Fluorescence was obtained from CY5 (for HPV16), ROX (for HPV18/45), CY5.5 (for HPV31/33/35/52/58), FAM (for HPV39/51/56/59/68) and HEX (for human beta globin gene as internal control). A sample was considered positive for the corresponding HPV genotype if the signal was detected within 60 min in the channel, regardless of the signal in the HEX channel. If no signal was detected for any of the four HPV channels within 60 min, then a signal was required in the HEX channel for the batch run to be called a valid negative.

Statistical analysis

The performance of the ScreenFire HPV Zebra BioDome assay on Atila iAMP-PS96 and Atila Powergene9600 Plus was evaluated based on the consistency with Thermo Fisher QuantStudio-7 and BioRad CFX-96. The data was assessed according to positive, negative, and overall agreement; as well as the unweighted kappa values for the 13 HPV genotypes (HPV16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59 and 68) in the four detection groups. The groups (channels) were analysed hierarchically based on cervical cancer risk: HPV16 positive, else positive for HPV18/45, else positive for HPV 31/33/35/52/58, else positive for HPV51/59/39/56/68, or else negative.

Results

On the pairwise hierarchical analysis, Atila iAMP-PS96 showed an overall agreement of 96.5% (167/173) with a kappa value of 0.95 (95% confidence interval: 0.91-0.99) compared to the results from Thermo Fisher QuantStudio-7 (Table 1A). Similarly, when compared to the results from BioRad CFX-96, the overall agreement was 97.1% (168/173) with a kappa value of 0.96 (95% confidence interval: 0.92-0.99, Table 1B). The detailed agreement for HPV RS genotyping is shown in Table 2A. The respective agreement rates for the genotypes of HPV16, HPV18/45, HPV31/33/35/52/58, and HPV39/51/56/59/68 were 99.4%, 98.3%, 99.4%, and 98.8%; the corresponding kappa values were 0.98 (95% confidence interval 0.95-1.0), 0.94 (0.86-1.0), 0.97 (0.91-1.0), and 0.90 (0.77-1.0) respectively. The agreement rates for RS genotypes when comparing iAMP-PS96 vs. CFX-96 were at least 98.8% (Table 2B). Thus, the Atila iAMP-PS96 demonstrated highly consistent results when compared to well-established qPCR technologies, such as Thermo Fisher QuantStudio-7 and BioRad CFX-96, using the ScreenFire RS Zebra BioDome HPV assays.

The pairwise hierarchical analysis between the results on Atila Powergene9600 Plus and Thermo Fisher Quant-Studio-7 also showed high consistency with an overall agreement of 98.8% (171/173) with a kappa value of 0.98 (95% confidence interval: 0.96-1.0, Table 3A). When the results on Atila Powergene9600 Plus were compared to those from BioRad CFX-96, the overall agreement of 96.5% (167/173) with a kappa value of 0.96 (95% confidence interval: 0.94-0.99, Table 3B). As shown in Table 4A, the agreement rate for HPV16 genotype was 98.8% with a kappa value of 0.97 (95% confidence interval 0.92-1.0) when comparing Atila Powergene9600 Plus and Thermo Fisher QuantStudio-7. The agreement for RS genotypes of HPV18/45, HPV31/33/35/52/58, and HPV39/51/56/59/68 were all 100%. The agreement rates for respective RS genotypes were at least 98.3% when comparing Atila Powergene9600 Plus and BioRad CFX-96 (Table 4B). Thus, Atila Powergene9600 Plus also

Table 1 Pairwise comparison between the HPV detection results of Atila iAMP-PS96 and (A) Thermo Fisher QuantStudio-7 or (B)BioRad CFX-96 using Screenfire HPV Zebra BioDome assay categorized hierarchically according to HPV RS genotypes

(A) iAMP-PS96 vs. QuantSt	tudio-7									
iAMP-PS96	QuantStudio-7									
	HPV16	HPV18/45	HPV31/33 /35/52/58	HPV39/51 /56/59/68	Negative	Total				
HPV16	36	0	0	0	1	37				
Row %	97.3	0	0	0	2.7	100				
Column %	100	0	0	0	1.2	21.4				
HPV18/45	0	26	1	0	1	28				
Row %	0	92.8	3.6	0	3.6	100				
Column %	0	96.3	5	0	1.2	16.2				
HPV31/33/35 /52/58	0	0	19	0	0	19				
Row %	0	0	100	0	0	100				
Column %	0	0	95	0	0	11				
HPV39/51/56 /59/68	0	0	0	10	2	12				
Row %	0	0	0	83.3	16.7	100				
Column %	0	0	0	100	2.6	6.9				
Negative	0	1	0	0	76	77				
Row %	0	1.3	0	0	98.7	100				
Column %	0	3.7	0	0	95	44.5				
Total	36	27	20	10	80	173				
Row %	20.8	15.6	11.6	5.8	46.2	100				
Column %	100	100	100	100	100	100				

Overall agreement rate = 96.5% (167/173); Unweighted kappa (95% CI) = 0.95 (0.91, 0.99)

(B) iAMP-PS96 vs. CFX-96

iAMP-PS96	CFX-96								
	HPV16	HPV18/45	HPV31/33 /35/52/58	HPV39/51 /56/59/68	Negative	Total			
HPV16	36	0	0	0	1	37			
Row %	97.3	0	0	0	2.7	100			
Column %	97.3	0	0	0	1.3	21.4			
HPV18/45	0	27	1	0	0	28			
Row %	0	96.4	3.6	0	0	100			
Column %	0	96.4	5	0	0	16.2			
HPV31/33/35 /52/58	0	0	19	0	0	19			
Row %	0	0	100	0	0	100			
Column %	0	0	95	0	0	11			
HPV39/51/56 /59/68	0	0	0	11	1	12			
Row %	0	0	0	91.7	8.3	100			
Column %	0	0	0	100	1.3	6.9			
Negative	1	1	0	0	75	77			
Row %	1.3	1.3	0	0	97.4	100			
Column %	2.7	3.6	0	0	97.4	44.5			
Total	37	28	20	11	77	173			
Row %	21.4	16.2	11.6	6.4	44.5	100			
Column %	100	100	100	100	100	100			
Overall agreement rate = 9	97.1% (168/173); U	Inweighted kappa (95	6% CI) = 0.96 (0.92, 0.99	9)					

 Table 2
 Agreement between Atila iAMP-PS96 and (A) Thermo Fisher QuantStudio-7 or (B) BioRad CFX-96 using Screenfire HPV Zebra

 BioDome assay for HPV RS genotyping

	+/+ n (%)	-/+ n(%)	+/- n(%)	-/- n(%)	Positive agreement % (95% CI)	Negative agree- ment % (95% CI)	Overall agree- ment % (95% Cl)	Unweight- ed kappa (95% Cl)
HPV16	36 (20.8)	0 (0)	1 (0.6)	136 (78.6)	100 (90.3–100)	99.3 (96–100)	99.4 (96.8–100)	0.98 (0.95-1)
HPV18/45	26 (15)	1 (0.6)	2 (1.2)	144 (83.2)	96.3 (81-99.9)	98.6 (95.1–99.8)	98.3 (95-99.6)	0.94 (0.86-1)
HPV31/33 /35/52/58	19 (11)	1 (0.6)	0 (0)	153 (88.4)	95 (75.1–99.9)	100 (97.6–100)	99.4 (96.8–100)	0.97 (0.91-1)
HPV39/51 /56/59/68	10 (5.8)	0 (0)	2 (1.2)	161 (93.1)	100 (69.2–100)	98.8 (95.6–99.9)	98.8 (95.9–99.9)	0.90 (0.77-1)
(B) iAMP-PS9	6 vs. CFX-96							
	+/+ n(%)	-/+ n(%)	+/- n(%)	-/- n(%)	Positive agreement % (95% CI)	Negative agree- ment % (95% Cl)	Overall agree- ment % (95% Cl)	Unweight- ed kappa (95% Cl)
HPV16	36 (20.8)	1 (0.6)	1 (0.6)	135 (78)	97.3 (85.8–99.9)	99.3 (96–100)	98.8 (95.9–99.9)	0.97 (0.92-1)
HPV18/45	27 (15.6)	1 (0.6)	1 (0.6)	144 (83.2)	96.4 (81.7–99.9)	99.3 (96.2–100)	98.8 (95.9–99.9)	0.96 (0.9-1)
HPV31/33 /35/52/58	19 (11)	0 (0)	1 (0.6)	153 (88.4)	100 (82.4–100)	99.4 (96.4–100)	99.4 (96.8–100)	0.97 (0.91-1)
HPV39/51 /56/59/68	11 (6.4)	1 (0.6)	0 (0)	161 (93.1)	91.7 (61.5–99.8)	100 (97.7–100)	99.4 (96.8–100)	0.95 (0.86-1)

demonstrated highly consistent results when compared to Thermo Fisher QuantStudio-7 and BioRad CFX-96 using the ScreenFire RS Zebra BioDome HPV assay.

Discussion

Understanding the analytical performance of the Screen-Fire HPV RS Zebra BioDome assay on commonly used qPCR instruments, not only from Atila but also from other manufacturers, will provide important information for the future use of this innovative HPV DNA test for cervical cancer screening. As the widely used clinically validated HPV assays on the market are mostly closed systems that require dedicated platforms, such as GeneXpert (cartridge-based), Roche Cobas, BD Onclarity and Seegene Allplex, so far none of the clinical validation studies has been done on the lab existing qPCR equipment. In contrast, the Atila ScreenFire HPV assay is an open-platform solution, allowing for broader compatibility. It has been evaluated and compared to other qPCR-based HPV genotyping assays on specific devices separately [9–15]. To our knowledge this is the first study to evaluate Zebra BioDome format across multiple existing platforms simultaneously. This study verified Screen-Fire Zebra BioDome HPV assays are compatible with widely used platforms existed in the lab.

The ScreenFire Zebra BioDome HPV assays not only offers the compatibility capability but also helps to address many other hurdles or limitations for implementing cervical cancer screening in low resource constrained settings, such as the needs for trained laboratory personnel and a dedicated laboratory setting to prevent contamination. The ScreenFire HPV RS Zebra BioDome format greatly reduced the number of processing steps, which now only involves adding lysed samples to pre-loaded Zebra BioDome reaction tube strips. The resulting process is fast, easy-to-use and high throughput, besides virtually eliminating the risk of laboratory contamination. In addition, the Zebra BioDome assay is also resistant to environmental fluctuations, as its performance remains unaffected by environmental temperature variations and other external conditions.

This study provides the first scientific data to show that the Zebra BioDome assays can be used in several commonly used qPCR platforms for HPV DNA testing. This cross-platform applicability has high public health significance in order to remove financial burden of capital investment to purchase new equipment for implementation cervical cancer screening. Our study in Mali and Nigeria has demonstrated that a robust and sustained implementation of a community-based ScreenFire Zebra BioDome HPV detection system using self-collected samples and operated by minimum-laboratory trained persons is feasible in LMIC settings.

Besides the unique features that have been described above, the ScreenFire Zebra BioDome HPV detection system is one of the most affordable HPV tests on the market compared to other commercially available HPV screening options. The ScreenFire HPV RS assay with

 Table 3
 Pairwise comparison between the HPV detection results of Atila PowerGene9600 plus and (A) Thermo Fisher QuantStudio-7 or (B) BioRad CFX-96 using Screenfire HPV Zebra BioDome assay categorized hierarchically according to HPV RS genotypes

 (A) PowerGene9600 plus vs
 QuantStudio-7

PowerGene9600 Plus	QuantStudie	o-7				
	HPV16	HPV18/45	HPV31/33 /35/52/58	HPV39/51 /56/59/68	Negative	Tota
HPV16	36	0	0	0	2	38
Row %	94.7	0	0	0	5.3	100
Column %	100	0	0	0	2.5	22
HPV18/45	0	27	0	0	0	27
Row %	0	100	0	0	0	100
Column %	0	100	0	0	0	15.6
HPV31/33/35	0	0	20	0	0	20
/52/58						
Row %	0	0	100	0	0	100
Column %	0	0	100	0	0	11.6
HPV39/51/56 /59/68	0	0	0	10	0	10
Row %	0	0	0	100	0	100
Column %	0	0	0	100	0	5.8
Negative	0	0	0	0	78	78
Row %	0	0	0	0	100	100
Column %	0	0	0	0	97.5	45.1
Total	36	27	20	10	80	173
Row %	20.8	15.6	11.6	5.8	46.2	100
Column %	100	100	100	100	100	100
Overall agreement rate = 9	8.8% (171/173); U	nweighted kappa (95	% CI) = 0.98 (0.96, 1)			

(B) PowerGene9600 Plus vs. CFX-96

PowerGene9600 Plus CFX-96 HPV16 HPV18/45 HPV31/33 HPV39/51 Negative Total /35/52/58 /56/59/68 HPV16 36 0 0 0 37 1 Row % 97.3 0 0 0 2.7 100 100 0 0 21.4 Column % 0 1.2 HPV18/45 0 26 0 28 1 1 Row % 0 92.8 3.6 0 3.6 100 0 0 Column % 96.3 5 1.2 16.2 HPV31/33/35 0 0 19 0 0 19 /52/58 Row % 0 0 100 0 0 100 Column % 0 0 0 0 11 95 HPV39/51/56 0 0 0 10 2 12 /59/68 Row % 0 0 0 83.3 16.7 100 0 0 0 100 Column % 2.6 6.9 Negative 0 1 0 0 76 77 Row % 0 1.3 0 0 98.7 100 Column % 0 3.7 0 0 95 44.5 Total 36 27 20 10 80 173 Row % 20.8 15.6 11.6 5.8 46.2 100 Column % 100 100 100 100 100 100 Overall agreement rate = 96.5% (167/173); Unweighted kappa (95% CI) = 0.96 (0.94, 0.99)

 Table 4
 Agreement between Atila PowerGene9600 plus and (A) Thermo Fisher QuantStudio-7 or (B) BioRad CFX-96 using Screenfire

 HPV Zebra BioDome assay for HPV RS genotyping

	+/+ n(%)	-/+ n(%)	+/- n(%)	-/- n(%)	Positive agreement % (95% CI)	Negative agree- ment % (95% CI)	Overall agree- ment % (95% Cl)	Unweight- ed kappa (95% Cl)
HPV16	36 (20.8)	0 (0)	2 (1.2)	135 (78)	100 (90.3–100)	98.5 (94.8–99.8)	98.8 (95.9–99.9)	0.97 (0.92-1)
HPV18/45	27 (15.6)	0 (0)	0 (0)	146 (84.4)	100 (87.2–100)	100 (97.5–100)	100 (97.9–100)	1 (1-1)
HPV31/33 /35/52/58	20 (11.6)	0 (0)	0 (0)	153 (88.4)	100 (83.2–100)	100 (97.6–100)	100 (97.9–100)	1 (1-1)
HPV39/51 /56/59/68	10 (5.8)	0 (0)	0 (0)	163 (94.2)	100 (69.2–100)	100 (97.8–100)	100 (97.9–100)	1 (1-1)

	+/+ n(%)	-/+ n(%)	+/- n(%)	-/- n(%)	Positive agreement % (95% Cl)	Negative agree- ment % (95% CI)	Overall agree- ment % (95% Cl)	Unweight- ed kappa (95% CI)
HPV16	36 (20.8)	2 (1.2)	1 (0.6)	134 (77.5)	94.7 (82.3–99.4)	99.3 (95.9–100)	98.3 (95-99.6)	0.95 (0.89-1)
HPV18/45	26 (15)	1 (0.6)	2 (1.2)	144 (83.2)	96.3 (81-99.9)	98.6 (95.1–99.8)	98.3 (95-99.6)	0.94 (0.86-1)
HPV31/33 /35/52/58	20 (11.6)	0 (0)	0 (0)	153 (88.4)	100 (83.2–100)	100 (97.6–100)	100 (97.9–100)	1 (1-1)
HPV39/51 /56/59/68	10 (5.8)	0 (0)	1 (0.6)	162 (93.6)	100 (69.2–100)	99.4 (96.6–100)	99.4 (96.8–100)	0.95 (0.85-1)

Zebra BioDome technology is currently priced at \$5.95 per test and the iAMP-PS96 device costs \$13,500, about one-quarter the cost of the most common existing platforms. The low cost per test combined with the portable and battery-operated nature of the iAMP-PS96 platform makes it a highly cost-effective solution for large-scale cervical cancer screening programs, particularly in low-resource settings. This study shows the value of Screen-Fire HPV technology to make the WHO's goal to screen 70% women in the world closer to a reality.

Acknowledgements

Not applicable.

Author contributions

JW performed the assays, interpreted the experimental data, prepared the manuscript. GI, ASA, JM and RA performed recruitment and biopsy collection. YZ analyzed and interpreted the data. BJ, IA, IOM, JB, MM and DBG interpreted the experimental data. ASS, FTO, RLM and LH supervised the study. All authors read and approved the final manuscript.

Funding

Research findings reported in this manuscript was supported by the National Cancer Institute of the National Institutes of Health under award number U54CA221205, D43CA260658, and U01CA275129, and by the Fogarty International Center of the National Institutes of Health under award number D43TW009575. The content is solely the responsibility of the authors and does not necessarily represent the official views of the National Institutes of Health.

Data availability

No datasets were generated or analysed during the current study.

Declarations

Ethics approval and consent to participate

This study is covered under IRB approval at Northwestern University (STU00207051), Jos University (JUTH/DCS/ADM/127/XXVII/630) and Lagos University (CMUL/HREC/01/18/327).

Consent for publication

Not applicable.

Competing interests

The authors declare no competing interests.

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References

- Sung H, Ferlay J, Siegel RL, Laversanne M, Soerjomataram I, Jemal A, et al. Global cancer statistics 2020: GLOBOCAN estimates of incidence and mortality worldwide for 36 cancers in 185 countries. CA Cancer J Clin. 2021;71(3):209–49.
- Arbyn M, Weiderpass E, Bruni L, de Sanjosé S, Saraiya M, Ferlay J, Bray F. Estimates of incidence and mortality of cervical cancer in 2018: a worldwide analysis. Lancet Glob Health. 2020;8:e191–203.
- Plummer M, de Martel C, Vignat J, Ferlay J, Bray F, Franceschi S. Global burden of cancers attributable to infections in 2012: a synthetic analysis. Lancet Global Health. 2016;4:e609–16.
- Small W Jr., Bacon MA, Bajaj A, Chuang LT, Fisher BJ, Harkenrider MM, Jhingran A, Kitchener HC, Mileshkin LR, Viswanathan AN, Gaffney DK. Cervical cancer: A global health crisis. Cancer. 2017;123:2404–12.
- World Health organization. New recommendations for screening and treatment to prevent cervical cancer: https://www.who.int/news/item/06-07-202 1-new-recommendations-for-screening-and-treatment-to-prevent-cervical-c ancer (Last Time Accessed, June 2024).
- Siqueira JD, Alves BM, Castelo Branco ABC, Duque KCD, Bustamante-Teixeira MT, Soares EA, Levi JE, Azevedo E, Silva G, Soares MA. Comparison of four different human papillomavirus genotyping methods in cervical samples: addressing method-specific advantages and limitations. Heliyon. 2024;10(3):e25474.
- Desai KT, Ajenifuja KO, Adepiti CA, Inturrisi F, Dagnall C, Hoffman AC, Egemen D, Gage JC, Wentzensen N, de Sanjose S, Schiffman M. Validation of a simplified HPV genotyping assay designed for cervical screening in low-resource settings. J Clin Microbiol 2024 Dec 31:e0163924. https://doi.org/10.1128/jcm. 01639-24. Epub ahead of print.
- de Sanjosé S, Perkins RB, Campos N, Inturrisi F, Egemen D, Befano B, Rodriguez AC, Jerónimo J, Cheung LC, Desai K, Han P, Novetsky AP, Ukwuani A, Marcus J, Ahmed SR, Wentzensen N, Kalpathy-Cramer J, Schiffman M, PAVE Study Group. Design of the HPV-automated visual evaluation (PAVE) study: validating a novel cervical screening strategy. Elife. 2024;12:RP91469.
- Desai KT, Adepiti CA, Schiffman M, et al. Redesign of a rapid, Lowcost HPV typing assay to support risk-based cervical screening and management. Int J Cancer. 2022;151(7):1142–9.

- Inturrisi F, de Sanjosé S, Desai KT, Dagnall C, Egemen D, Befano B, et al. A rapid HPV typing assay to support global cervical cancer screening and risk-based management: A cross-sectional study. Int J Cancer. 2024;154(2):241–50.
- Hou J, Belinson JL, Du H, Li C, Zhang W, Zhang L, Zhang Y, Qu X, Wu R. AmpFire HPV and screenfire RS HPV validation trial. Am J Clin Pathol. 2024;161(6):535–42.
- Wang J, Imade G, Akanmu AS, Musa J, Anorlu R, Zheng Y, Garcia-Bedoya O, Sanchez GI, Belinson J, Kim K, Maiga M, Gursel DB, Sagay AS, Ogunsola FT, Murphy RL, Hou L. Analytic performance of screenfire HPV RS assay zebra biodome format and its potential for large-scale population HPV screening. Infect Agents Cancer. 2024;19(1):59.
- Zhang W, Du H, Huang X, Wang C, Duan X, Liu Y, Shi B, Zhang W, Qu X, Wei L, Schiffman M, Belinson JL, Wu R. Evaluation of an isothermal amplification HPV detection assay for primary cervical cancer screening. Infect Agent Cancer. 2020; 23;15:65.
- 14. Moyo S, Ramogola-Masire D, Moraka NO, Tawe L, Noubary F, Motsumi K, Manowe G, Zuze B, Radibe B, Hungwe FTT, Mohammed T, Maphorisa C, Shapiro R, Gaseitsiwe S, Luckett R. Comparison of the AmpFire multiplex HPV assay to the Xpert* HPV assay for detection of human papillomavirus and cervical disease in women with human immunodeficiency virus: a pragmatic performance evaluation. Infect Agent Cancer. 2023;18(1):29.
- Mungo C, Guliam A, Chinula L, Inturrisi F, Msowoya L, Mkochi T, Jawadu S, de Sanjosé S, Schiffman M, Tang JH, Smith JS. Comparison of the screenfire and Xpert HPV assays for the detection of human papillomavirus and cervical precancer among women living with HIV in Malawi. Infect Agent Cancer. 2024;19(1):24.
- Musa J, Kim K, Zheng Y, Qu Y, Joyce BT, Wang J, Nannini DR, Gursel DB, Silas O, Abdulkareem FB, Imade G, Akanmu AS, Wei J-J, Kocherginsky M, Kim K-YA, Wehbe F, Achenbach CJ, Anorlu R, Simon MA, Sagay A, Ogunsola FT, Murphy RL, Hou L. Accelerated epigenetic age among women with invasive cervical cancer and HIV-Infection in Nigeria. Front Public Health. 2022;10:834800.

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