

Comparative Performance of Anyplex II HPV28 and Cobas 4800 Human Papillomavirus (HPV) Assays for High-Risk HPV Detection in Self-collected Anal Samples

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26 ABSIKACI

28	We compared 2 HPV assays to detect the 14 hrHPV genotypes in self-collected anal
29	samples. We found a good agreement and similar performance to detect HPV16, HPV18 and
30	the 12 others hrHPV genotypes. The global performance to detect the 14 hrHPV genotypes
31	was not significantly different between the 2 assays.
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33	KEYWORDS
34	Anal lesions; self-sampling; HPV-testing; Anyplex II HPV28; Cobas 4800; HPV16; MSM;
35	TWG; Cambodia
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51 **BACKGROUND**

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Around 30 000 new cases of anal squamous cell carcinoma (ASCC) are diagnosed 53 each year worldwide [1]. Key-population, such as men having sex with men (MSM) and 54 people living with HIV (PLWHIV), have an increased risk of developing ASCC. For instance, 55 HIV-infected MSM has the highest incidence rate of ASCC with 85 new cases per 100 000 56 person-year [2]. Human Papillomavirus (HPV) is involved in roughly 90% of ASCC, of 57 58 whom 90% are HPV16 positive. ASCC is preceded by precancerous lesions, i.e. anal intraepithelial neoplasia 1, 2 and 3 (AIN1, AIN2 and AIN3), AIN1 being identify as low 59 grade anal lesions (LGAIN), AIN2 and AIN3 as high grade anal lesions (HGAIN). However 60 61 for now, high-risk HPV testing (hrHPV) is not recommended for the screening of anal precancerous and cancer lesions, while it has been clinically validated with multiple hrHPV 62 63 tests in cervical cancer screening [3,4]. Although anal HPV16 genotyping is not yet a widely 64 accepted molecular marker in anal cancer screening, it is by far the most carcinogenetic type 65 in the anus [5]. Indeed, (i) HPV16 prevalence increases across the anal lesion from normal to 66 precancerous and anal cancer [6] and (ii) in a recent prospective study, HGAIN HPV16positive are less likely to clear compared to those with other hrHPV genotypes [7]. 67

In this study, we aim to compare the agreement and performance of the Cobas 4800
HPV-test (Roche®) and the AnyplexTMII HPV28 Detection (Seegene Inc) to detect hrHPV in
self-sampling anal swabs from asymptomatic MSM and transgender women (TGW) in
Phnom Penh, Cambodia; especially for HPV16 detection which was the most prevalent HPV
genotype found in this population [8].

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76 MATERIALS AND METHODS

77 Study population

A cross-sectional retrospective study was conducted between October 1st, 2021 and 78 November 30th, 2021 in different clinics in Phnom Penh, Cambodia and under the scope of 79 80 activities of Men's Health Cambodia, a non-governmental and non-profit organization working on men, including men having sex with men (MSM) and transgender women (TGW) 81 issues in Cambodia. During consultation, a questionnaire was used to collect patient data and 82 83 the anal collection (FLOQSwabs, Copan) was self-performed using anal swabs discharged in Viral Transport Media (VTM). They were then stored at -80°C at the Institut Pasteur du 84 Cambodge (IPC) until further analysis. A comprehensive explanation of eligibility criteria and 85 procedures has been published before [8]. 86

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88 HPV-testing and methods' comparison

First analysis was performed at IPC on 400 µl of sample with the Cobas 4800 HPV-test 89 90 (Roche®) which allows the detection of 14 hrHPV types by PCR and nucleic acid hybridization. This test specifically identifies HPV16, HPV18 and 12 others hrHPV in the 91 same channel including HPV31-33-35-39-45-51-52-56-58-59-66-68. Then, 1ml of frozen 92 93 VTM was sent to the Virology Department of the Pitié-Salpêtrière hospital, Paris, France. DNA-extraction (from 200 µl of sample to 100 µl) was performed with the E-mag device 94 (Biomérieux) and both samples and extracted DNA were stored at -80°C. Then HPV-testing 95 was performed with the AnyplexTMII HPV28 Detection (Seegene Inc) which detects and 96 97 differentiates 28 distinct HPV-types including the 14 hrHPV detected by the Cobas 4800 98 HPV-test and 14 others HPV (not analyzed in this paper). Both HPV-testing also detect a 99 human gene as internal control (IC) to assess the quality of the sampling and the efficiency of 100 the PCR amplification. If IC was not amplified, samples were referred as invalid.

101 Statistical analysis was performed with the package *rstatix* from R software. McNemar and 102 kappa indices were calculated to compare the proportion of positive samples and the degree of 103 agreement between the two methods to detect the 14 hrHPV, respectively. A *p*-value of < 0.05104 in two-tailed were considered significant.

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106 **RESULTS**

107 Overall, 162 participants were included with a median age of 28.1 [24.4 – 31.8] years,
108 of whom 56% were MSM and 8% were HIV-infected. None of them received any HPV109 vaccine.

110 One hundred and sixty-two anal samples were tested with both the Roche and Seegene 111 methods and HPV-testing failed for 59 (36%) and 41 (25%) samples, respectively. The 112 prevalence of HPV16, HPV18 and others hrHPV types were not significantly different 113 between the two methods: 20% versus 25% (p=0.52), 16% versus 17% (p=0.86) and 50% 114 versus 59% (p=0.18), respectively (Table 1).

115 Further statistical analyses for method comparison were performed on 97 samples, 65 samples 116 being referred as invalid in one or the two methods. The concordance of the results for 117 HPV16, HPV18 and others hrHPV types detection were 97%, 100% and 88%, respectively 118 (Table2). Fifteen samples were discordant: (i) 3 for HPV16 positivity, of whom 2 HPV16 119 missed by Roche and 1 missed by Seegene, (ii) 12 for others hrHPV positivity, of whom 8 120 hrHPV-types missed by Roche and 4 by Seegene (Supplementary Table 1). HPV52 was the 121 most common hrHPV type missed by Roche (6/10 cases), followed by HPV16 and HPV56 122 (2/10 cases each one). For most of the hrHPV missed by Roche, they were detected tardily by Seegene, after 40 cycles of amplification. 123

The agreement (Cohen's Kappa test) for HPV16, HPV18 and others hrHPV detection
were 0.91 [0.81-1.00], 1.00 [1.00-1.00] and 0.75 [0.62-0.88], respectively. The performance

to detect HPV16, HPV18 and others hrHPV positivity were not significantly different between the 2 tests (McNemar test, p>0.05) (Table 2 and Supplementary Table 2). In total, the global performance to detect any of the 14 hrHPV types was not significantly different (McNemar test, p=0.44).

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131 DISCUSSION

To our knowledge, this is the first study conducted in self-collected anal samples
which compared the Cobas 4800 HPV-test (Roche®) and the AnyplexTMII HPV28 Detection
(Seegene Inc) tests for hrHPV detection.

First, comparing the 2 methods on 97 samples revealed a good agreement between 135 them and comparable worldwide ability to identify any of the 14 hrHPV genotypes currently 136 recommended for the cervical cancer screening. This result was also consistent for HPV16 137 138 alone, HPV18 alone and the others hrHPV genotypes detected with the 2 methods. However, 139 compared to the Seegene approach, the Cobas test missed more anal hrHPV infection. In term 140 of hrHPV prevalence in normal cervical tissue, a difference was also identified between the 141 Cobas method and GP5+/6+ PCR testing followed by genotyping with Luminex [4]. 142 However, equal frequency was showed in samples with abnormal cytology. In our work, we 143 dealt with anal samples and we did not assess the association with cytology results, thus these 144 two factors may have an impact on the difference between the 2 procedures.

The prevalence of HPV16 was the highest in our high-risk population as expected and the 2 methods showed high agreement and performance to detect the most oncogenic genotype in anal site. This is of importance as HPV16 testing could be one of the biomarkers used as anal cancer screening in key-population in the near future. For instance in France, new guidelines propose to use HPV16 testing in anal cytology specimen collected by physician as anal cancer screening marker in MSM living with HIV [9]. If positive, a reflex cytology is performed onthe same sample and patient is referred to a proctologist.

Similarly to self-collected vaginal swabs for cervical cancer screening [10], self-collected anal samples could be of interest to reach patients outside the healthcare system [11]. In our work, 41 (25%) and 59 (36%) samples failed to be amplified by the AnyplexTMII HPV28 Detection and Cobas 4800 HPV-test, respectively. Low-cellularity due to inappropriate sampling and misunderstanding of the participants may explain this high rate of invalid results. If implemented in this population, self-collection instructions should be improved.

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In conclusion, both the AnyplexTMII HPV28 Detection and the Cobas 4800 HPV-test are suitable to detect the 14 hrHPV genotypes in self-collected anal samples, as well as more specifically the HPV16 genotype.

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Authors contributions. Conception and design of the study: AJ, SC and GD; acquisition of
the clinical data: SW, SP and KV; execution of the experiments: GAM; analysis of the data:
AJ, SM and GD; drafting of significant portion of the manuscript or figures: AJ, VC, AGM
and GD. All the authors red, corrected and approved the final version of the manuscript.

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180 Transparency declarations. The authors declare that the research was conducted in the 181 absence of any commercial or financial relationships that could be construed as a potential 182 conflict of interest.

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184 Ethics. Permission for the study was obtained from the National Ethics Committee for Health
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	Seegene N= 121 ¹	Roche N= 103²	<i>p-value</i> ³
HPV16, n (%)			
Negative	91 (75%)	82 (80%)	0.52
Positive	30 (25%)	21 (20%)	
HPV18, n (%)			
Negative	101 (83%)	87 (84%)	0.86
Positive	20 (17%)	16 (16%)	
hrHPV others, n (%)			
Negative	50 (41%)	52 (50%)	0.18
Positive	71 (59%)	51 (50%)	

- 234 ¹41 samples being referred as invalid with Seegene HPV-test
- 235 ² 59 samples being referred as invalid with Roche HPV-test
- **236** ³Fisher's exact test
- hrHPV others: 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66 and 68
- 238 HPV: human papillomavirus; hrHPV: high-risk human papillomavirus

Table 2: Agreement and performance of the two methods to detect HPV16, HPV18 and the

256 others hrHPV types

	$N = 97^{1}$	Cohen's Kappa test	McNemar test
HPV16, n (%)			
Concordant	94 (97%)	0.91 [0.81 - 1.00]	1.00
Discordant	3 (3%)		
HPV18, n (%)			
Concordant	97 (100%)	1.00 [1.00 - 1.00]	NA
Discordant	0 (0%)		
hrHPV others, n (%)			
Concordant	85 (88%)	0.75 [0.62-0.88]	0.39
Discordant	12 (12%)		
All hrHPV, n(%)			
Concordant	90 (93%)	0.85 [0.74–0.96]	0.44
Discordant	7 (7%)		

259 ¹65 samples being referred as invalid in one or both HPV-test

260 Cohen's Kappa test and McNemar test were performed to compare the agreement and the performance of the

two methods to detect HPV16, HPV18, the others hrHPV and the 14 hrHPV as a whole.

hrHPV others: HPV31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66 and 68

263 HPV: human papillomavirus; hrHPV: high-risk human papillomavirus