

Comparison of Onclarity Human Papillomavirus (HPV) Assay with Hybrid Capture II HPV DNA Assay for Detection of Cervical Intraepithelial Neoplasia Grade 2 and 3 Lesions

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Analytical and clinical performance validation is essential before introduction of a new human papillomavirus (HPV) assay into clinical practice. This study compares the new BD Onclarity HPV assay, which detects E6/E7 DNA from 14 high-risk HPV types, to the Hybrid Capture II (HC2) HPV DNA test, to concurrent cytology and histology results, in order to evaluate its performance in detecting high-grade cervical lesions. A population of 567 women, including 325 with \geq ASCUS (where ASCUS stands for atypical cells of undetermined significance) and any HC2 result and 242 with both negative cytology and negative HC2 results, were prospectively enrolled for the study. The overall agreement between Onclarity and HC2 was 94.6% (95% confidence intervals [CI], 92.3% to 96.2%). In this population with a high prevalence of disease, the relative sensitivities (versus adjudicated cervical intraepithelial neoplasia grades 2 and 3 [CIN2+] histology endpoints) of the Onclarity and HC2 tests were 95.2% (95% CI, 90.7% to 97.5%) and 96.9% (95% CI, 92.9% to 98.7%), respectively, and the relative specificities were 50.3% (95% CI, 43.2% to 57.4%) for BD and 40.8% (95% CI, 33.9%, 48.1%) for HC2. These results indicate that the BD Onclarity HPV assay has sensitivity comparable to that of the HC2 assay, with a trend to an increased specificity. Moreover, as Onclarity gives the chance to discriminate between the different genotypes, we calculated the genotype prevalence and the absolute risk of CIN2+: HPV 16 was the most prevalent genotype (19.8%) with an absolute risk of CIN2+ of 77.1%.

uman papillomavirus (HPV) is the etiological agent for the development of cervical cancer and its precursor lesions (1): a persistent infection with high-risk (HR) HPV types has been established as a necessary step in the progression from precancerous to neoplastic disease (2). It is widely recognized that different HR HPV genotypes have different oncogenic potential, with genotype 16 being the most oncogenic of the cancer-causing HPV genotypes (3, 4).

For decades, the test of choice for the detection of cervical intraepithelial neoplasia (CIN) was the Pap smear (5). However, the Pap test has severe limitations: a relatively low sensitivity for detection of disease (range, 51 to 74%, depending on the study), limited reproducibility, interobserver variability, and finally the occurrence of equivocal results (6). The advent of HPV testing has introduced an epochal change in cervical cancer screening: in recent years, we observed a gradual shift in the use of HPV test from a "triage test," in case of equivocal cytology, or as an adjunct test to cytology in women older than 30 years, to a primary screening test. This introduction of molecular HPV testing is slowly changing the premise of cervical screening, promising better performance than cytology-based screening but also posing new challenges (7). Over the last 7 years, HPV testing has gained a foothold in a number of national screening programs, most notably in triage of equivocal cytology or as an adjunct test to cytology in women older than 30 years, as well as more recently as a primary screening modality. Four European randomized control trials (RCTs), as recently summarized by Ronco and colleagues, clearly demonstrate the benefit of HPV testing over the use of the Pap smear (8). In this scenario, the performance of the HPV testing system of choice is crucial: the key is for an HPV assay to have an optimal balance between sensitivity and specificity for detection of CIN grade 2 or 3 (CIN2+) lesions in order to identify virtually all women with immediate precursors or in rare cases cervical cancer, and at the same time not to result in too many screening false-positive samples (9, 10). The advantage of HPV testing relies on its high sensitivity, \geq 95% for CIN2+ lesions, which translates into a high negative predictive value (NPV) allowing for extended screening intervals without increased risk of cervical cancer: HPV-negative women have a reduced risk for CIN2/3 lesions and cancer in the subsequent screening round as demonstrated by the European RCTs (8).

Several in-house HPV DNA detection methods have been successfully used in research laboratories worldwide for more than 2 decades, and some have been employed for diagnostic purposes in countries where "nonapproved" tests for routine clinical testing are allowed (11). Recently, new tests using microarrays or real-time PCRs have been developed: the majority of them have been designed to

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Laboratory test. After vortexing, each sample was aliquoted for cytological analysis, HC2, LA, and Onclarity assays.

HC2. The Qiagen HC2 test is a sandwich capture molecular hybridization assay: it is a signal amplification detection method based on chemiluminescence that detects 13 HR HPV types (HPV 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, and 68). The DNA-RNA hybrids are captured on a microplate, and the emitted light is measured in a luminometer as relative light units (RLU). Samples were considered positive if the RLU/cutoff ratio was >1.0 (equivalent to 1.0 pg HPV DNA/ml). All samples with RLU between 1 and 2.5 should be retested as requested in the package insert

RESULTS We prospectively enrolled 567 women. The median age was 39

years (range, 18 to 81 years). The majority of women were found in the category of \geq 30 years (78.1%) (Fig. 1).

Subsequent histology was available for 352 women, with 46.9% of them showing samples determined as \geq CIN2 and 25% with \geq CIN3. Women with no histological evaluation were considered

FIG 2 BD Onclarity HPV assay design. The assay is comprised of 3 wells, designated G1 to G3, each of which has a beta-globin internal control (IC) on one channel. The 14 high-risk HPV types are read out on the other 3 channels of each well as an individual genotype or as a group of 2 or 3 pooled HPV types as shown.

detect the entire genome or a region thereof (usually within L1 region) with different grades of automation. Some of these tests are also able to provide genotyping (single or pooled), possibly allowing a stratification of the risk of cervical high-grade lesions (12).

Becton Dickinson (BD) has recently developed a new assay for the detection of HR HPV: the BD Onclarity HPV assay. It is based on type-specific E6/E7 DNA target detection using the BD Viper LT platform (BD Diagnostics, Sparks, MD, USA). The assay simultaneously detects 13 high-risk genotypes and one possibly carcinogenic HPV genotype (HPV66) (13); it provides genotyping information on six individual genotypes (16, 18, 31, 45, 51, and 52) and reports the remaining eight HPV genotypes in three distinct groups (33/58, 56/59/66, and 35/39/68). The assay has an internal human beta globin (HBB) gene control for sample sufficiency and assay performance.

This study was aimed at an explorative analytical evaluation of the Onclarity assay in a referral colposcopy center, with a high prevalence of cervical high-grade lesions, to evaluate the relative sensitivity and specificity of the assay in this setting. We compared the results of the BD Onclarity TM HPV assay (i) to those of the Qiagen HC2 HPV DNA test (HC2; Qiagen GmbH, Hilden, Germany) and (ii) to reference cytology and histology results.

MATERIALS AND METHODS

Population. Women (n = 567, ranging in age from 18 to 81 years) (Fig. 1) attending the European Institute of Oncology in Milan referred for cervical conization, for colposcopy, or for screening purposes were enrolled in the study. For each woman a ThinPrep PreservCyt (Hologic, Inc. Bedford, MA) cervical sample was collected for Onclarity, HC2, and the Roche Diagnostics linear array (LA) test for genotyping and cytology. Cytological diagnoses were reported according to the 2001 Bethesda system terminology (14). Histology was performed for all women with positive cytology (\geq ASCUS, where ASCUS stands for atypical cells of undetermined significance) and/or a positive HPV test. Histology was performed either from a specimen obtained by colposcopy with biopsy or from the evaluation of a conization specimen, and in the case of high-grade lesions a conservative surgical procedure was performed with histology evaluation of the tissue. Informed consent was obtained from all study women according to local ethical approval (IEO S689/212 study approved 28 March 2012).

instructions, but the residual material volume was not sufficient: these samples were discarded from the analysis.

Linear array test. The Roche Diagnostics LA test uses biotinylated PGMY09/11 consensus primers to amplify a 450-bp region of the L1 gene: it is capable of detecting 37 HPV genotypes (6, 11, 16, 18, 26, 31, 33, 35, 39, 40, 42, 45, 51, 52, 53, 54, 55, 56, 58, 59, 61, 62, 64, 66, 67, 68, 69, 70, 71, 72, 73 [MM9], 81, 82 [MM4], 83 [MM7], 84 [MM8], IS39, and CP6108). The denatured PCR products were then hybridized to an array strip containing immobilized oligonucleotide probes. The results were visually interpreted by using the provided reference guide according to the manufacturer's protocol by two independent operators, and then the results were compared to reach a consensus.

BD Onclarity. The BD Onclarity HPV assay detects 14 HPV genotypes and coamplifies a beta-globin internal control (IC), which acts as a processing control. The primers for the 14 HPV genotypes are designed to target a region of 79 to 137 bases in the E6/E7 genome, whereas the IC primers amplify a 75-base region in the human beta-globin gene. The assay consists of three PCR assay tubes (G1, G2, and G3) and four optical channels for the detection of the 14 HPV genotypes (16, 18, 31, 45, 51, 52 as single infections and the remaining eight genotypes in three groups [33/58, 56/59/66, and 35/39/68]) and the IC (Fig. 2).

Cytology and histology. Pap tests were all liquid based and prepared with the ThinPrep 5000 slide processor (Hologic, Inc., Bedford, MA). The final diagnosis of women with abnormal cytology (>ASCUS) was made by histological evaluation of colposcopically directed biopsy specimens (at least 2 punches of the transformation zone alone or with endocervical curettage [ECC]) or on surgical tissue. Histology was adjudicated by two independent pathologists. Discrepancies were further adjudicated by a third independent pathologist.

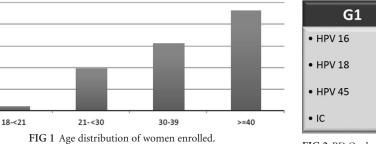
Data analysis and statistics. The data were recorded on case report forms and analyzed using standard statistical techniques.

We compared the results of Onclarity HPV assay with those of the HC2 HPV assay using the McNemar test.

Sensitivity and specificity were calculated for these two tests in relationship to cytology category. Positive predictive value and negative predictive value were evaluated. The results of HC2, Linear Array (Roche), and LBC Onclarity HPV assays were summarized and cross-tabulated.

All tests were considered statistically significant at α values of <0.05. All statistical analyses were performed using SAS 9.3 (SAS Institute Inc., Cary, NC).

G2 G3 • HPV 16 HPV 33_58 • HPV 51 HPV 18 HPV 31 HPV 52 HPV 45 • HPV 56 59 66 HPV 35_39_68 • IC • IC • IC



50

40

30

20 10

%

Cytology category	Onclarity HPV assay result	No. of samples with indicated HC2 HPV assay result		Total no.		% agreement (95% CI)			
		Positive	Negative	NA^b	of samples	P value	Positive	Negative	Overall
NILM	Positive	15	5	4	24		71.4 (50.0-86.2)	98.0 (95.3–99.1)	95.9 (92.8–97.7)
	Negative	6	242	6	254	0.76			
	Total	21	247	10	278				
ASCUS	Positive	13	1	1	15		92.9 (68.5–98.7)	95.2 (77.3–99.2)	94.3 (81.4–98.4)
	Negative	1	20	0	21	1.00			
	Total	14	21	1	36				
>ASCUS	Positive	219	3	1	223		94.0 (90.2–96.4)	81.3 (57.0-93.4)	93.2 (89.3–95.7)
	Negative	14	13	3	30	0.008			
	Total	233	16	4	253				
Overall	Positive	247	9	6	262		92.2% (88.3–94.8)	96.8 (94.1–98.3)	94.6 (92.3–96.2)
	Negative	21	275	9	305	0.03	. /	. ,	. ,
	Total	268	284	15	567				

TABLE 1 Percent agreement of BD HPV and Qiagen HC2 HPV assays by each cytology category (NILM, ASCUS, >ASCUS) and all categories combined for PreservCyt samples^a

^a The McNemar test was used to compare results.

^b NA, not available.

negative if they fulfilled the combination of being cytology negative, HPV negative, and cytology negative at the previous round of screening.

Overall, we collected 567 cervical samples: 15 samples were not analyzed because they gave an HC2 result with an RLU between 1.0 to 2.5 and the volume was insufficient for retesting. Concordance between the BD Onclarity HPV and the Qiagen HC2 assays was 94.6% (522/552), with 30 discordant samples (5.4%). Percent agreements subcategorized into cytology results and combined results are shown in Table 1. Onclarity and HC2 results were significantly different for overall cytology categories (P = 0.03); however, these differences occurred only for samples with a concurrent cytology showing >ASCUS (P = 0.008, Table 1). Nine samples were found to be HC2 negative and Onclarity positive; of these, 4 tested HR HPV positive with LA while 5 were LA negative (Table 2). Resulting histology was available in 4 cases, and 1 ≥CIN3 sample was detected. Twenty-one samples were HC2 positive and Onclarity negative; 11 were concordantly HR HPV negative with LA, and the resulting histology showed 1 sample to be ≥CIN3 and 10 samples to be <CIN2. Ten patients showed discrepant genotype findings between Onclarity and LA whether in

single or multiple infections. Histology of these patients revealed that 2 were \geq CIN2 and 8 were < CIN2 (Table 3).

We analyzed the data coming from genotyping and calculated the prevalence and the absolute risk of CIN2+: HPV16 was the most prevalent genotype (19.8%) with an absolute risk of CIN2+ of 77.1%. The second-most-frequent genotype was HPV31 (10.6%) with an associated absolute risk of CIN2+ of 66.7%. HPV18 was found in 4.2%, with an absolute risk of 52.2%.

A comparison of the performance of BD Onclarity HPV assay with that of HC2 stratified according to histology results (\geq CIN2 versus <CIN2) by each cytology category (negative for intraepithelial lesion or malignancy [NILM], ASCUS, or >ASCUS) is shown in Table 3. Overall sensitivity was 95.2% (95% confidence intervals [CI], 90.7% to 97.5%) for BD and 96.9% (95% CI, 92.9% to 98.7%) for HC2 HPV, and overall specificity was 50.3% (95% CI, 43.2% to 57.4%) for BD and 40.8 (95% CI, 33.9% to 48.1%) for HC2 HPV (Table 4).

DISCUSSION

The aim of the current study was to evaluate the analytical performance of the BD Onclarity HPV assay against that of the HC2

Subject ID	Result by:			Onclarity HPV	LA result or HPV genotype(s)		
	Cytology	Lab histology	HC2	Onclarity	genotype(s)	HR HPV	Other HPV
MIL1091	NILM	NEG	NEG	POS	31	31	NEG
MIL1372	NILM	NA^{a}	NEG	POS	59/56/66	NEG	NEG
MIL1375	NILM	NA	NEG	POS	16	NEG	NEG
MIL1617	NILM	NA	NEG	POS	33/58	NEG	NEG
MIL1563	>ASCUS	NA	NEG	POS	16	NEG	NEG
MIL1326	NILM	CIN1	NEG	POS	16	16	NEG
MIL1125	ASCUS	CIN1	NEG	POS	16	16	NEG
MIL1128	>ASCUS	CIN2	NEG	POS	52	NEG	NEG
MIL1248	>ASCUS	≥CIN3	NEG	POS	52	52	6, 53

TABLE 2 Discordant results between HC2 assay (negative results) and Onclarity assay (positive results) compared to cytology, histology, and linear array

^{*a*} NA, not available.

	Result by:				LA result or HPV genotype(s)		
Subject ID	Cytology	Lab histology	HC2	Onclarity	HR HPV	Other HPV	
MIL1084	NILM	NEG	POS	Negative	Negative	Negative	
MIL1232	NILM	NEG	POS	Negative	Negative	Negative	
MIL1561	NILM	NEG	POS	Negative	Negative	Negative	
MIL1043	NILM	NEG	POS	Negative	HPV58	HPV53	
MIL1068	NILM	NEG	POS	Negative	HPV58	HPV42	
MIL1235	NILM	NEG	POS	Negative	HPV16	Negative	
MIL1030	ASCUS	CIN1	POS	Negative	Negative	HPV70	
MIL1118	>ASCUS	NEG	POS	Negative	HPV51	HPV42	
MIL1049	>ASCUS	NEG	POS	Negative	HPV45, HPV51	HPV42, HPV53, HPV70, HPV83	
MIL1152	>ASCUS	NEG	POS	Negative	HPV45, HPV51, HPV66	HPV42, HPV67	
MIL1024	>ASCUS	CIN1	POS	Negative	Negative	Negative	
MIL1361	>ASCUS	CIN1	POS	Negative	Negative	HPV53	
MIL1485	>ASCUS	CIN1	POS	Negative	Negative	HPV70	
MIL1192	>ASCUS	CIN1	POS	Negative	Negative	HPV54, HPV70	
MIL1116	>ASCUS	CIN1	POS	Negative	Negative	HPV53, HPV54, HPV84, CP6108	
MIL1079	>ASCUS	CIN1	POS	Negative	Negative	HPV53, HPV67	
MIL1051	>ASCUS	CIN1	POS	Negative	HPV16, HPV59	Negative	
MIL1041	>ASCUS	CIN1	POS	Negative	HPV31, HPV45, HPV51	HPV61	
MIL1106	>ASCUS	CIN2	POS	Negative	HPV33, HPV39	Negative	
MIL1087	>ASCUS	≥CIN3	POS	Negative	Negative	HPV67	
MIL1006	>ASCUS	≥CIN3	POS	Negative	HPV16	HPV53, HPV61	

TABLE 3 Discordant results between HC2 assay (positive results) and Onclarity assay (negative results) compared to cytology, histology, and linear array

assay for the detection of HR HPV in cervical samples from a referral population attending our institution. Overall, the two assays showed good agreement using ThinPrep stored samples (94.6%), with similar sensitivity and specificity between the assays in this setting with high prevalence of disease.

HPV testing is becoming more and more recognized as the test of choice for primary screening and management of cervical disease, and recently released guidelines propose HPV testing as the primary screening test, replacing cytology (8, 9, 15). A shift to primary HPV screening can be implemented only if the test used has undergone a fully analytical and clinical validation. Yet a number of different systems are available and used today (12, 16), with large differences in analytical and clinical performance. This represents a serious challenge, as decisions

are taken by the clinicians on the basis of a result that may not be completely correct from a clinical perspective. In fact, at present only few systems have been fully validated.

In this study, we focused primarily on the evaluation of Onclarity in a referral population, associating the Onclarity HPV result with underlying clinical disease and comparing the outcomes in a split-sample fashion to results obtained with the HC2 test. The agreement was found to be very high, even though the Onclarity test, with its extended genotype readout, represents a more complex diagnostic system than does the HC2. These findings are in line with the conclusions previously reported by Wright et al. and Ejegod et al. (17, 18). Onclarity offers substantial advantages over "yes/no" assays like HC2 in terms of automation, reduced hands-on time, and the presence of an internal control based on a

TABLE 4 Performance comparison of BD Onclarity HPV and Qiagen HC2 assays with histology results (≥CIN2 versus <cin2) for="" preservcyt<="" th=""></cin2)>
samples by each cytology category (NILM, ASCUS, >ASCUS) and all categories combined

		BD Onclarity HPV a	ssay	Qiagen HC2 HPV as	say
Cytology category	Performance measure	Value % ^a	95% CI	Value %	95% CI
NILM	Sensitivity	50.0 (2/4)	15.0-85.0	50.0 (1/2)	9.5-90.5
	Specificity	79.1 (53/67)	67.9–87.1	72.6 (45/62)	60.4-82.1
ASCUS	Sensitivity	100.0 (7/7)	64.6-100.0	100.0 (7/7)	64.6-100.0
	Specificity	72.0 (18/25)	52.4-85.7	75.0 (18/24)	55.1-88.0
>ASCUS	Sensitivity	96.1 (148/154)	91.8-98.2	97.4 (148/152)	93.4–99.0
	Specificity	24.2 (23/95)	16.7–33.7	10.8 (10/93)	5.9-18.7
Overall	Sensitivity	95.2 (157/165)	90.7–97.5	96.9 (156/161)	92.9–98.7
	Specificity	50.3 (94/187)	43.2-57.4	40.8 (73/179)	33.9-48.1
	Positive predictive value	62.8 (157/250)	59.5-66.4	59.5 (156/262)	56.7-62.7
	Negative predictive value	92.2 (94/102)	85.8-95.9	93.6 (73/78)	86.3–97.2

^{*a*} Values are percentages relatives to the CIN2+ endpoints determined by histology; fractions in parentheses represent the number of samples with results agreeing with histology/ total number of samples.

human housekeeping gene, the positivity of which confirms the adequacy of the specimen.

Looking at the discordant samples, HC2-positive/Onclaritynegative results dominated, constituting 70% of the discordant samples. The HR HPV result of HC2 was not confirmed by LA genotyping in one half of these samples, while for the other half the LA detected the presence of different low-risk genotypes. We found only 3 of 21 patients HC2 positive/Onclarity negative with a lesion \geq CIN2. The remaining HC2-positive/Onclarity-discordant samples showed mainly LR HPV types, thereby representing a "false-positive" HC2 test result, using LA HR HPV negativity as the adjudicator. On the other hand, 9 patients were HC2 negative/ Onclarity positive; 4 patients were positive for HR HPV by LA, and 1 patient was negative for any HPV. Histology was available for only 6 of these patients, with 2 patients bearing a \geq CIN2, both with HPV genotype 52. It should be noted that LA does not have an established clinical cutoff and would be expected to detect some analytically positive samples that are below the clinical cutoff in the Onclarity assay. Overall, our data corroborate already published data showing that a substantial cross-reactivity to LR HPV types damages the HC2 specificity (19, 20, 21). In conclusion, these data show that Onclarity not only performs as well as HC2 from a clinical sensitivity perspective but also offers some advantage in terms of specificity, although the comparison, though showing a trend, did not reach a statistical significance overall.

One interesting point is that, differently from other recently introduced systems, Onclarity is a PCR-based assay in which the target region is within the E6 and E7 regions of the HR HPV genome. This could be of help in the case of cervical cancer, in which the L1 part of the genome has been disrupted and therefore these lesions may give a negative HPV test result when systems based on L1 consensus primers PCR are used (22, 23). Moreover, the test is designed so that different genotypes can be distinguished: HPV genotypes 16, 18, 45, 31, 51, 52, 33, 58, 56, 59, 66, 35, 39, and 68 can be identified in single or multiple infections. This can be of help, both for the risk stratification of the patients, as HPV16 detection can refer the women to a different clinical approach, and for the follow-up of patients with a CIN lesion, after treatment, to evaluate the clearance or the persistence of a specific HPV type (4). One strength of this study is that women were prospectively enrolled, samples were stored and tested in batches with Onclarity, and we had a high percentage of specimens taken from patients with a cervical lesion. On the other hand, one limitation related to the inclusion criteria is that we could not have a histological diagnosis in case of both HC2 negativity and cytology negativity, in particular for those cases testing positive with Onclarity.

In conclusion, the BD Onclarity assay is an accurate HPV assay for the detection and genotyping of HR HPV infections. Further studies are ongoing to validate its clinical performance outside high-prevalence, high-risk referral populations like those used in this study.

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