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**EVALUATION OF TYPE-SPECIFIC HPV PERSISTENCE AND
HIGH-RISK HPV VIRAL LOAD QUANTITATION IN HPV
POSITIVE WOMEN UNDER 30 WITH NORMAL CERVICAL
CYTOLOGY**

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Keywords:	Human Papillomavirus, Cervical cancer, viral load, genotyping, screening

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	HPV test	
	Initial (n=199)	1-year follow up (n=39)
Number of HPV types detected n (%)		
1	41 (20.6)	17 (43.6)
≥ 2	16 (8)	8 (20.5)
HPV types detected n (%)		
HPV16	22 (11.1)	15 (38.5)
HPV18	11 (5.5)	8 (20.5)
HPV31	6 (3)	2 (5.1)
HPV33	8 (4)	0 (0)
Other HR-HPV types*	5 (2.5)	1 (2.6)
LR-HPV types**	25 (12.6)	7 (17.9)
HPV viral load (n copies / 10 ⁶ cells) mean (SD)		
HPV16	8.5.10 ⁵ (2.04.10 ⁶)	5.27.10 ⁵ (1.08.10 ⁶)
HPV18	9.4.10 ⁵ (2.87. 10 ⁶)	3.66.10 ⁵ (9.24.10 ⁵)
HPV31	4.57.10 ⁶ (4.65.10 ⁶)	2.11.10 ⁸ (2.88.10 ⁸)
HPV33	2.37.10 ⁶ (3.85.10 ⁶)	-

Table 1. Results of initial and 1-year follow up HPV testing and typing.

HR-HPV: high-risk HPV; LR-HPV: low-risk HPV.

*Other high-risk HPV types initially detected: 4(2%) HPV58 and 1(0.5%) HPV82.

**Low-risk HPV types initially detected – 25 patients were infected by 30 LR-HPV types (because of multiple infections): 8(4%) HPV6, 6(3%) HPV53, 4(2%) HPV66, 2(1%) HPV61, 2(1%) HPV62, 2(1%) HPV 83, 1(0.5%) HPV1, 1(0.5%) HPV11, 1(0.5%) HPV54, 1(0.5%) HPV70, 1(0.5%) HPV81 and 1(0.5%) HPV84.

HPV type initially detected	Number of patients followed-up	1-year specific persistent HPV infection rate (%) [95% CI]	Cervical intraepithelial neoplasia			
			All grades		Grade 2 or 3	
			Incidence rate (%) [95% CI]	Women with 1-year specific persistent HPV infection n (%)	Incidence rate (%) [95% CI]	Women with 1-year specific persistent HPV infection n (%)
HPV16	12	92 [62-100]	33 [10-65]†	3 (75)	8 [0-39]	1 (100)
HPV18	7	57 [18-90]	14 [0-58]	0 (0)	0	-
HPV31	5	20 [1-72]	0	-	0	-
HPV33	3	0	0	-	0	-
Other HR-HPV types*	4	25 [1-81]	0	-	0	-
LR-HPV types**	20	15 [3-38]	5 [1-25]‡	0 (0)	0	-

Table 2. Rate of one-year HPV type persistence and cervical intraepithelial neoplasia incidence.

HR-HPV: high-risk HPV; LR-HPV: low-risk HPV.

* Other high risk HPV types: 3 HPV58 and 1 HPV82.

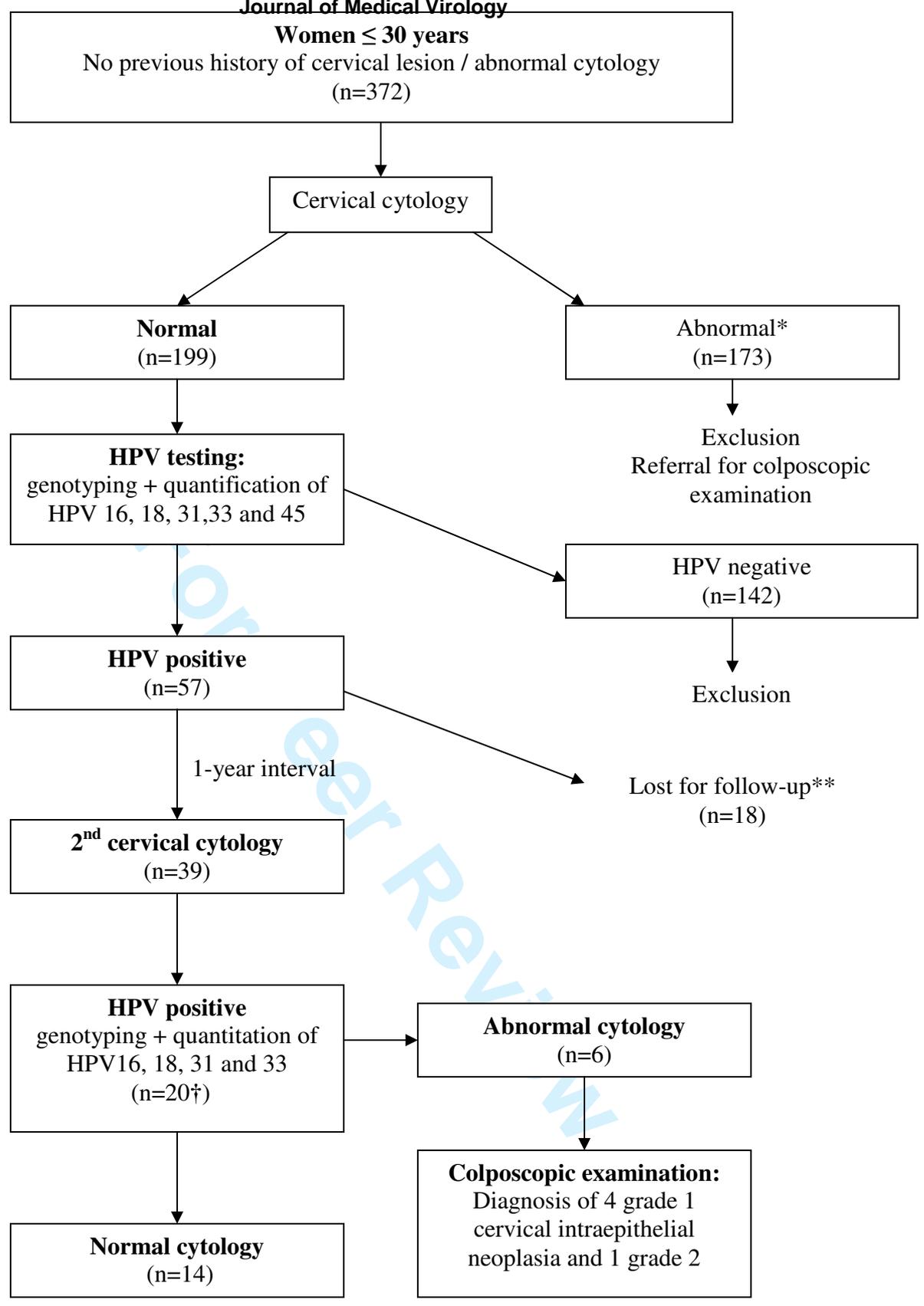
** Low risk HPV types: 6 HPV53, 4 HPV6, 2 HPV62, 2 HPV66, 2 HPV83, 1 HPV11, 1 HPV61, 1 HPV 81 and 1 HPV84.

† 3 patients were diagnosed with grade 1 Cervical intraepithelial neoplasia and one with grade 2 cervical intraepithelial neoplasia.

‡ 1 patient was diagnosed with grade 1 cervical intraepithelial neoplasia.

Bold style indicates statistical significance when comparing with other HPV types.

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3 **Figure 1.** Study design
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5 * Smear was considered abnormal if borderline nuclear abnormalities (BNA) or more severe
6 abnormalities were identified.
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9 ** 1 patient got pregnant and thus did not attend to the 2nd cervical smear.
10

11 † 20 cases of type-specific HPV persistent infection were identified: 11 HPV16, 4 HPV 18, 1
12 HPV 31, 0 HPV 33, 3 LR-HPV, and 1 HR-HPV different from types 16, 18, 31 and 33.
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For Peer Review

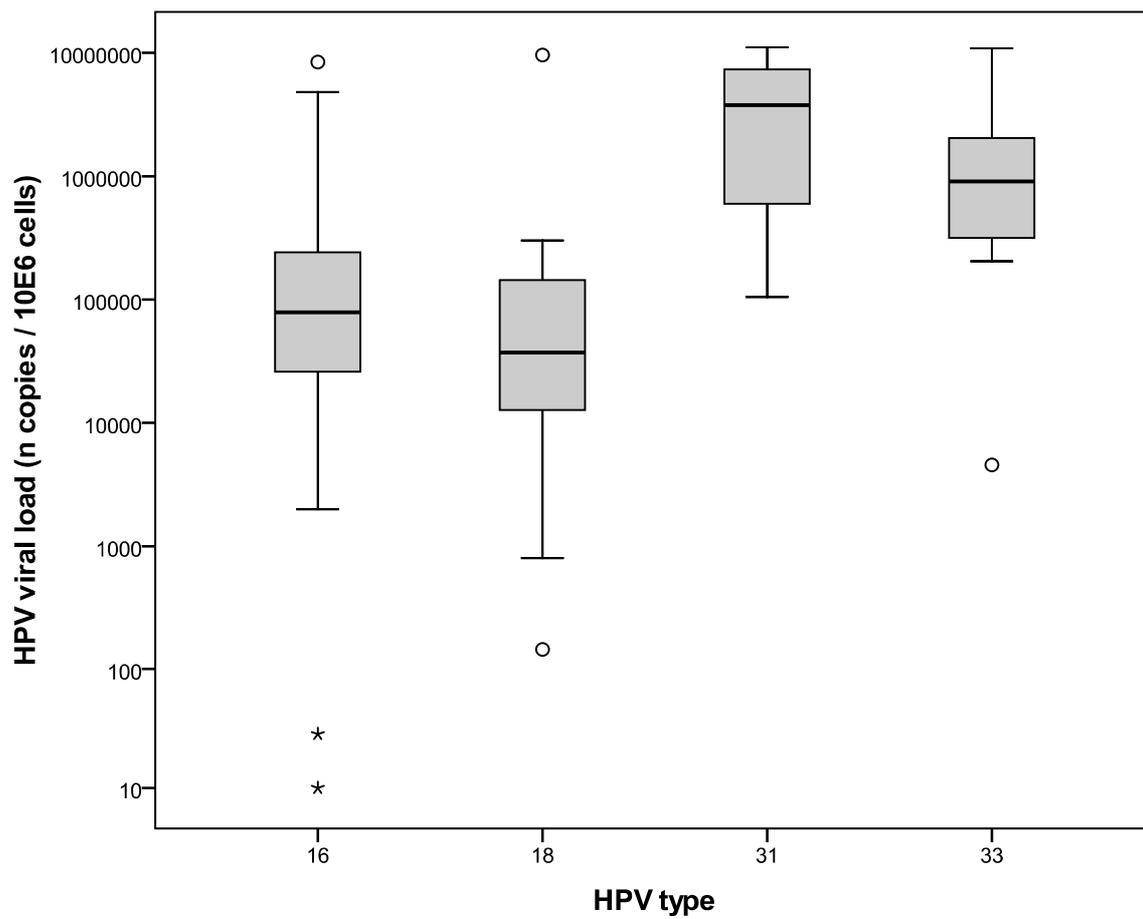


Figure 2. Distribution of HPV16, 18, 31 and 33 viral load in women under 30 with normal smear at the time of inclusion (logarithmic scale). HPV31 viral load was statistically higher than HPV16 ($p=0.015$) and HPV18 viral load ($p=0.019$).

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3 1 **EVALUATION OF TYPE-SPECIFIC HPV PERSISTENCE AND HIGH-RISK HPV**
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5 2 **VIRAL LOAD QUANTITATION IN HPV POSITIVE WOMEN UNDER 30 WITH**
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7 3 **NORMAL CERVICAL CYTOLOGY**
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11
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7 **Shortened title:** HPV type and viral load in young women with normal smear

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ABSTRACT

The persistence of high-risk HPV (HR-HPV) infection is necessary for the development of cervical intraepithelial neoplasia. The aim of this study was to evaluate if HR-HPV typing and HPV16, 18, 31 and 33 quantitation are predictive for type-specific infection persistence and/or the development of CIN in women under 30 with normal cervical cytology. Young women (under 30) attending a family planning clinic who were HPV positive with normal cervical cytology were included. HPV genotyping was assessed by MY09/ MY11 PCR, sequencing, phylogenetic analysis, and cloning when necessary. HR-HPV viral load was quantified using duplex real-time PCR. Study patients were offered for a second smear and HR-HPV detection and quantitation after twelve months. HR-HPV was identified in 43 (21.9%) of the 199 included women. Of these, 39 patients had a second cervical sample taken within a mean interval of 11.7 months (8.8-18.3 months). The mean HR-HPV 16, 18, 31 and 33 initial viral load was 1.9×10^6 copies per millions cells. The level of viral load did not reveal any significant association with type-specific HR-HPV persistence or the subsequent development of cervical intraepithelial neoplasia. Only HPV16 infection was significantly more likely to persist (91.7% vs. 33.1%, $p=0.001$) and to develop CIN (33.3% vs. 3.7%, $p=0.025$). In women under 30 with normal cytology, HR-HPV viral load is common and is not predictive of HPV persistence or the development of cervical intraepithelial neoplasia. HPV16 positive women are significantly more likely to have persistent infection and to develop cervical intraepithelial neoplasia.

248 words

Key words: Human Papillomavirus, cervical cancer, viral load, genotyping, screening.

1 INTRODUCTION

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1 High-risk or oncogenic, human papillomavirus (HR-HPV) infection is the major risk factor
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3 for the development of cervical cancer and cervical intraepithelial neoplasia. In France, 97%
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5 of cervical cancers are HPV positive [Pretet et al., 2007]. Among HR-HPV types, HPV16, is
6
7 by far the most prevalent type and is identified in 73% of cervical cancers, followed by HPV
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9 18 (19 %), 31 (7 %) and 33 (4 %) [Pretet et al., 2007]. For this reason HPV testing has been
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11 considered as an alternative method of cervical pre-cancer screening. Compared to cervical
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13 cytology, HPV testing is more sensitive but less specific in the detection of grade 2 or more
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15 cervical intraepithelial neoplasia [Arbyn et al., 2006; Cuzick et al., 2006; Koliopoulos et al.,
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17 2007]. The Superiority of HPV testing over cytology or visual inspection techniques has also
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19 been demonstrated in low resource setting countries [Sankaranarayanan et al., 2009]. In this
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21 study screening with HPV and subsequent management of screen positive women has been
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23 shown to result clinically important and statistically significant reduction in the number of
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25 advanced stages cervical cancers and of overall cervical cancer related mortality
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27 Age has a major impact on the clinical utility of HPV testing being very much more specific
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29 in women over 30 years [Koliopoulos et al., 2007; Mayrand et al., 2007]. HPV infection is
30
31 extremely common in young women and in the great majority of cases it is both transient and
32
33 ultimately harmless. Following initial exposure to HR-HPV, the great majority of women will
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35 clear their infection in 8 to 10 months without the development of a cervical lesion [Dalstein
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37 et al., 2003; Ho et al., 1998; Woodman et al., 2001]. Thus, persistence of HPV infection only
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39 affects a small minority of women [Bosch and Munoz, 2002; Franco et al., 1999; Ho et al.,
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41 1998]. More than HR-HPV infection itself, it is the persistence of the same HR-HPV
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43 infection that has been demonstrated to be the most important risk factor for cervical
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45 intraepithelial neoplasia [Dalstein et al., 2003; Nobbenhuis et al., 1999; Schlecht et al., 2001].

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3 1 Thus, the positive predictive value of a single HPV test in young women is poor and only a
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5 2 minority of these women will develop a cervical lesion [Cuzick et al., 2003]. Factors that
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7 3 influence HR-HPV persistence remain unknown and predictive factors for HR-HPV infection
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9 4 persistence would be clinically valuable.

10 5 With the increased use of HPV testing in routine practice, physicians must consider how to
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12 6 manage HPV positive young women who have normal cytology. This presents a clinical
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14 7 dilemma and may easily result in unnecessary treatment with the potential for subsequent
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16 8 pregnancy related morbidity and perinatal mortality [Arbyn et al., 2008; Kyrgiou et al., 2006].
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18 9 Determination of HR-HPV viral load would be useful if it could predict HR-HPV persistence
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20 10 and Cervical Intraepithelial Neoplasia [Bae et al., 2009; Dalstein et al., 2003; Lai et al., 2008;
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22 11 Munoz et al., 2009; van Duin et al., 2002]. The predictive value of HR-HPV detection could
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24 12 be HPV type-dependant with perhaps a higher risk for women with high HPV 16 viral load
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26 13 [Gravitt et al., 2007; Moberg et al., 2004; Moberg et al., 2005].

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28 14 The aim of this study was to evaluate HPV type-specific prevalence and persistence in a
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30 15 population of women under 30 with normal cervical cytology. In that specific population,
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32 16 HPV 16, 18, 31 and 33 viral load, referred to as “oncogenic” HPV viral load, was evaluated
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34 17 as a predictive factor of type-specific HPV infection persistence and of colposcopically
35
36 18 documented cervical intraepithelial neoplasia. The impact of HPV genotyping was also
37
38 19 evaluated.

39 20

40 21 **MATERIALS AND METHODS**

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42 23 **Patients and design of the study**

43 24 The study design is summarized in Figure 1. All women under 30 with no previous history of
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45 25 cervical lesion or of abnormal cervical cytology attending an Irish Family Planning

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3 1 Association (IFPA) clinic in Tallaght, (Dublin, Ireland) for routine screening cervical
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5 2 cytology were invited to participate to the study. All had a liquid-based cervical cytology
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7 3 performed using the ThinPrep® pap test (Hologic Inc, Nasdaq: Holx) and dedicated cervical
8
9 4 brush. Women with abnormal cervical cytology were referred for colposcopic examination or
10
11 5 repeat cervical cytology according to British Society of Colposcopy and Cervical Pathology
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13 6 (BSCCP) guidelines and were then excluded from the study. All women with normal cervical
14
15 7 cytology had a HPV test performed directly on the remaining suspension of the ThinPrep®
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17 8 pap test vial. HPV negative women were excluded from the study and referred for routine
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19 9 screening cytology i.e. three yearly. HPV positive women were asked to return in one-year for
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21 10 a repeat cervical cytology and HPV test. Subsequent cervical smears were performed at the
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23 11 Colposcopy Clinic of the Coombe Women and Infants University Hospital. All were tested
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25 12 for HPV with systematic genotyping of HPV positive sample and quantitation of HPV 16, 18,
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27 13 31 and 33 viral load. Women with a second abnormal smear were referred for colposcopic
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29 14 examination. Colposcopic examinations were blinded to the results of HPV genotyping and
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31 15 viral load quantitation.
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33 16 Study protocol received approval from Ethics Committees of both the Coombe Women and
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35 17 Infants University Hospital and the Adelaide and Meath Hospital in Tallaght, Dublin.
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37 18 Informed consent was obtained from all the women participating to the study.
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48 **Cervical cytology analysis**

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50 21 All cervical smears were analyzed at the Pathology department of the Coombe Women and
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52 22 Infants University Hospital (Dublin, Ireland). Analysis were performed using ThinPrep®
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54 23 Imaging system (Hologic Inc, Nasdaq: Holx). All slides were first screened using the Review
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56 24 Scopes® followed by manual secondary screening. Negative cases were signed out by the
57
58 25 cytotechnologists and all abnormal cases (Borderline Nuclear Abnormality or greater) were
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1 signed out by the consultant Pathologist. Cervical cytology analysis were blinded to the result
2 of HPV genotyping and viral load quantitation.

3

4 **Virological analysis**

5 HPV testing was performed at the department of Virology of the Timone's Hospital
6 (Marseilles, France). All HPV tests were performed using the remaining fluid in the liquid
7 based cytology vials. HPV genotyping was assessed by PCR (MY09/ MY11 primers),
8 sequencing, phylogenetic analysis and cloning if necessary, as described previously [Tamalet
9 et al., 2010]. Quantitation of HPV 16, 18, 31 and 33 positive samples were performed using
10 quantitative duplex real-time PCR method, as reported previously [Carcopino et al., 2006].
11 HPV viral load was expressed as the number of HPV copies per million cells.

12 For quantification, we used a plasmid which contained the five target sequences of interest:
13 HPV 16, HPV 31, HPV 33, HPV 45 (each on E6 gene), HPV 18 (on E7 gene), and human
14 albumin gene (on exon12). This method allowed HPV 16 and 18, 31, 33, and 45 and albumin
15 gene copy number to be quantified in the same assay. So, HPV viral load could be expressed
16 as HPV copies per cell or per million cells. For the purpose of intra-assay reproducibility, the
17 standard curve was tested 10 times on the same plate and serial dilutions of positive samples
18 (from 10⁷ copies to 1 copy per assay) each with a high HPV 16, 18, 31, 33, or 45 value were
19 tested in duplicate to establish the sensitivity and the reproducibility of the method. We
20 obtained a lower detection limit of 5 copies per 5 μ l of sample as well for HPV 16, 18, 31, 33,
21 45 as for albumin. The linearity was excellent from the highest value to 1 copy per 5 μ l of
22 sample. The data revealed an excellent reproducibility of the assays as well for high, median
23 or low values. The coefficient of correlation varied from 98.9% for HPV 18 to 99.7% for
24 HPV 16. For the purpose of inter-assay reproducibility, the coefficient of variation (CV) was

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3 1 determined for each reaction. Mean CV for each point of the quantification curve in HPV 16,
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5 2 18, 31, 33, 45 and albumin was <3%.
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10 **Statistical analysis**

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12 5 Statistical analysis was performed using SPSS 15.0 software for Windows (SPSS Inc,
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14 6 Chicago, Illinois). Groups were compared using the Chi square test or the Fisher exact test for
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16 7 qualitative characteristics, and using the Mann-Whitney test for continuous ones. A two-sided
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18 8 P value of less than 0.05 was considered statistically significant.
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10 **RESULTS**

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12 12 Recruitment ran from September 2007 to the end of March 2009, during which 372 patients
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14 13 participated in the study. Of these, 199 had normal cytology. 173 had an abnormal cervical
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16 14 smear, were excluded from the study and were referred for colposcopic examination or repeat
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18 15 cervical cytology. All of the 199 patients whose cervical cytology was normal were tested for
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20 16 HPV. The overall HPV prevalence was 28.6% (57/199) (Table 1). HR-HPV was identified in
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22 17 43 (21.6%) patients. The most prevalent HPV genotypes were HPV 16 (11.1%) and HPV 18
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24 18 (5.5%) followed by HPV 33 (4%), HPV 6 (4%), HPV 31 (3%), HPV 53, 58, 66, 61, 62 and 83
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26 19 (3-1%). Single HPV infection was observed in 41 (20.6%) while 16 (8%) patients had at least
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28 20 two different HPV types. The mean HR-HPV viral load for HPV 16, 18, 31 and 33 was
29
30 21 1.5×10^6 copies / million cells (Standard Deviation (SD): 3.05×10^6). The mean HPV 31 viral
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32 22 load was statistically higher than the mean HPV 16 ($p=0.015$) or HPV 18 viral loads
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34 23 ($p=0.019$) (Figure 2).
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36 24 After the exclusion of all HPV negative women, 57 remained for analysis. Of these, 39
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38 25 (68.4%) patients had a second cervical sample taken within a mean interval of 11.7 months
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1 (min-max: 8.8-18.3 months). One patient conceived and declined a second cervical sample.
2 Seventeen patients were lost to follow up despite several reminder invitations by mail and
3 telephone (Figure 1). The mean patient age was 24.8 years (min-max: 18-30 years). The
4 mean interval between smears was comparable for HPV 16, 18, 31 and 33 positive patients as
5 well as for patients who were positive for other HPV types.

6 All of the 39 patients were tested for HPV. Type-specific persistent HPV infection was
7 identified in 20 (51.3%). The mean age of patients with persistent infection and of those who
8 cleared their infection was comparable: 24.7 years vs. 24.5, respectively. HPV 16 was
9 significantly more likely to persist than were all other types: 11 (91.7%) vs. 9 (33.3%),
10 respectively ($p < 0.001$).

11 Compared to HR-HPV types, low-risk HPV (LR-HPV) were significantly less likely to
12 persist: 17 (54.8%) vs. 3 (15.0%), respectively ($p = 0.004$) (Table 2). In HPV 16, 18, 31 and
13 33 positive patients the initial overall viral load was not predictive of type specific HPV
14 infection. Mean initial HPV viral load of women with persistent HPV 16, 18, 31 or 33
15 infection was 1.2×10^6 copies per million cells whereas viral load of women who had cleared
16 their infection was 5.3×10^6 copies per million cells ($p = 0.173$).

17 The result of a second smear test was abnormal in 6 of the 39 patients. Three women were
18 reported as having borderline nuclear abnormality (BNA), 2 low-grade squamous
19 intraepithelial lesion (LSIL) and 1 high-grade squamous intraepithelial lesion (HSIL). After
20 colposcopic examination, one patient was diagnosed with normal cervix. Grade 1 cervical
21 intraepithelial neoplasia was diagnosed in 4 patients who were advised to have a follow up
22 colposcopy and smear within a year. One patient was diagnosed with grade 2 cervical
23 intraepithelial neoplasia, confirmed at LLETZ (large loop excision of the transformation
24 zone).

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3 1 Patients diagnosed with cervical intraepithelial neoplasia (any grade) were not significantly
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5 2 more likely to have persistent HPV infection: 3 (60%) vs. 17 (50%), respectively, though
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7 3 these numbers do not allow accurate assessment of risk. Nor was the initial HPV 16, 18, 31
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9 4 or 33 viral load predictive of cervical intraepithelial neoplasia. Initial mean HPV viral load in
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11 5 patients who were diagnosed with cervical intraepithelial neoplasia was 5.2×10^4 copies per
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13 6 million cells compared to 2.9×10^6 copies per million cells in women with a normal cervix
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15 7 ($p=0.263$). Also the mean age was comparable in both groups: 23.7 years vs. 24.7 years
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17 8 ($p=0.515$). Finally, compared to other HPV types, only HPV 16 infection was significantly
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19 9 associated with a higher probability of developing cervical intraepithelial neoplasia: 1 (3.7%)
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21 10 vs. 4 (33.3%), respectively ($p=0.025$) (Table 2). Despite the fact that the only patient
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23 11 diagnosed with grade 2 cervical intraepithelial neoplasia was HPV 16 positive, HPV 16
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25 12 infection was not significantly associated with higher probability of Grade 2 or 3 cervical
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27 13 intraepithelial neoplasia: 1 (8.33%) vs. 0 (0%), respectively ($p=0.3$).
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36 15 **DISCUSSION**

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41 17 With a HPV and HR-HPV prevalence of 28.6% and 21.6%, respectively, the present study
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43 18 confirms that HPV, and particularly HR-HPV infection, is very common in women under 30
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45 19 years old with normal cervical cytology. Such finding concurs with the results of other
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47 20 studies. In women who reported recent coitarche and who reported just one sexual partner, the
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49 21 1-year cumulative incidence rate of HPV infection is 28.5 % (IC à 95 % : 20,6-38,6 %) and
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51 22 reaches almost 50% after 3 years [Winer et al., 2008]. The overall cumulative incidence rate
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53 23 of HPV infection in 25 year old women has been estimated to be 80 % [Winer et al., 2006].
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55 24 Thus, the prevalence of HPV infection in young women is very high, with a peak of
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57 25 prevalence of 19.6% for women between 25 and 29 years. This appears to decline to 4.3% in
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Carcopino et al.

1 women over 30 [Jacobs et al., 2000]. Although the rates for HR-HPV are generally lower, the
2 prevalence and natural history is comparable to overall HPV. The best evidence suggests that
3 in 20-24 and 25-29 years old women, HR-HPV prevalence is 13% and 17% respectively and
4 will decrease to 2.5 and 3.9 % in women over 30 [Cibas et al., 2007; Jacobs et al., 2000].
5 Interestingly, despite the limited number of 39 women in the key core study group, it should
6 be noted that 20 women (51.3%) presented persistent HR-HPV infection. Of these 20 women,
7 six had abnormal cytology, while 14 had sustained normal cytology after a mean interval time
8 period of 11.7 months. Thus, since the critical point is actually to know the future of this
9 persistent HPV infection in these 14 young women it will be of highest importance to monitor
10 their cytological and virological follow up. What is more, given the consistent association
11 between persistent infection and increased risk of cervical precancer and cancer, persistent
12 HPV infection was the virological endpoint proposed as a relevant clinical marker in the
13 Cervarix vaccine trial [Paavonen et al., 2009](Paavonen, Lancet 2009, 374:301-14). Finally
14 the high incidence of transient HPV infection in young women could also explain the higher
15 probability of abnormal smear in young women as the cytopathic effects of HPV can be
16 detected on cytology [ACOG, 2009]. However, explanation for higher probability of
17 abnormal cytology, as observed in the present study, is probably multifactorial and includes
18 the low socioeconomic level of women attending to the IFPA clinic in Tallaght [Cotton et al.,
19 2007].

20 In the present study the initial HPV 16, 18, 31 and 33 viral load was not found to be
21 predictive of type-specific HPV persistence or the development of cervical intraepithelial
22 neoplasia although in other studies HPV viral load has been demonstrated to be correlated
23 with HPV persistence and subsequent cervical pre-malignancy [Bae et al., 2009; Dalstein et
24 al., 2003; Lai et al., 2008; Munoz et al., 2009; van Duin et al., 2002]. This correlation may
25 vary with HPV type and to be most reliable for HPV 16 [Gravitt et al., 2007; Moberg et al.,

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3 1 2004; Moberg et al., 2005]. It seems that no previous reports have evaluated such a
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6 2 correlation in young women with normal smears.

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8 3 Interestingly, mean HR-HPV viral load was relatively high. Indeed, the mean initial HR-HPV
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10 4 viral load measured in the present study was close to the HPV 16 viral load cut-off ($>2.2 \times 10^7$
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12 5 copies/million cells) identified by Saunier et al. to allow the identification of women with
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14 6 prevalent grade 2 or 3 cervical intraepithelial neoplasia with a high specificity [Saunier et al.,
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16 7 2008]. Thus, regardless of age, it has been shown that HR-HPV viral load increases
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18 8 significantly with the severity of cervical lesions [Saunier et al., 2008]. But, in a previous
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20 9 study, such significant association was only found in patients over 30 while mean HPV 16
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22 10 and 18 viral load was already particularly high in women under 30 [Carcopino et al., 2006].
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25 11 These results suggest that HR-HPV viral load is commonly high in young women, possibly
26
27 12 related to the recent occurrence of HPV infection, and would therefore not be of any clinical
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29 13 significance.
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34 14 In this study, only HPV 16 was found to persist and associated with an increased risk of
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36 15 cervical intraepithelial neoplasia. In comparison, LR-HPV types were significantly less likely
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38 16 to persist. This finding illustrates the high oncogenicity of HPV 16 and points to the potential
39
40 17 benefit of HPV genotyping in clinical practice. Indeed, HPV 16 has been already shown to
41
42 18 persist over a 8-16 months period [Ralston Howe et al., 2009]. The usefulness of HPV 16
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44 19 testing in the management of HPV positive women with normal cervical cytology has been
45
46 20 already demonstrated [Khan et al., 2005]. In their study the detection of HPV 16 was
47
48 21 associated with a 17.2% risk of grade 3 or more cervical intraepithelial neoplasia over a 10
49
50 22 years period (95% CI: 11,5-22,9). In the present study, the absence of significant association
51
52 23 between HPV 16 infection and grade 2 or more cervical intraepithelial neoplasia occurrence
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54 24 may be explained by the small number of HPV positive patients and because the study was
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56 25 limited to one-year's follow-up.
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3 1 The Management of HPV positive young women with normal cytology remains a clinical
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5 2 challenge. The balance between not overtreating women with a minimal risk of progression
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7 3 to cancer and the risk of treatment related morbidity is a fine line [Arbyn et al., 2008; Kyrgiou
8
9 4 et al., 2006]. These results suggest that the use of HPV genotyping and particularly HPV 16
10
11 5 identification may be valuable. Because of the risk of HPV 16 persistence and the
12
13 6 development of cervical intraepithelial neoplasia, young HPV 16 positive patients with
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15 7 normal smear should be monitored carefully.
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22 9 **CONCLUSION**

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27 11 This study extends the understanding of the natural history of HPV in women under 30 with
28
29 12 normal cervical cytology. In this specific population, HPV type 16, 18, 31 and 33 viral load is
30
31 13 often relatively high. It does not appear to predict type-specific HPV infection persistence or
32
33 14 the development of cervical intraepithelial neoplasia. However, in the present study, HPV 16
34
35 15 infection was more likely to persist and HPV 16 positive patients were also more likely to
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37 16 develop cervical intraepithelial neoplasia. The use of HPV typing may be valuable in women
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39 17 under 30.
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For Peer Review

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