


RESEARCH ARTICLE

Open Access



Low-cost HPV testing and the prevalence of cervical infection in asymptomatic populations in Guatemala

Hong Lou^{1†}, Eduardo Gharzouzi^{2†}, Sarita Polo Guerra^{2†}, Joël Fokom Domgue³, Julie Sawitzke¹, Guillermo Villagran², Lisa Garland¹, Joseph F. Boland¹, Sarah Wagner¹, Héctor Rosas⁵, Jami Troxler⁶, Heidi McMillen⁶, Bailey Kessing⁶, Enrique Alvarez⁷, Miriam Castillo², Hesler Morales², Victor Argueta⁸, Andert Rosingh⁹, Femke J. H. B. van Aerde-van Nunen⁹, Griselda Lopez⁸, Herbert M. Pinedo¹⁰, Mark Schiffman³, Michael Dean^{4*}  and Roberto Orozco⁸

Abstract

Background: A low cost and accurate method for detecting high-risk (HR) human papillomavirus (HPV) is important to permit HPV testing for cervical cancer prevention. We used a commercially available HPV method (H13, HybriBio) which was documented to function accurately in a reduced volume of cervical specimen to determine the most prevalent HPV types and the distribution of HPV infections in over 1795 cancer-free women in Guatemala undergoing primary screening for cervical cancer by cytology.

Methods: HR-HPV detection was attempted in cervical samples from 1795 cancer-free women receiving Pap smears using the HybriBio™ real-time PCR assay of 13 HR types. The test includes a globin gene internal control. HPV positive samples were sequenced to determine viral type. Age-specific prevalence of HPV was also assessed in the study population.

Results: A total of 13% (226/1717) of women tested HPV+, with 78 samples (4.3%) failing to amplify the internal control. The highest prevalence was found in younger women (< 30 years, 22%) and older ones (≥60 years, 15%). The six most common HR-HPV types among the 148 HPV+ typed were HPV16 (22%), HPV18 (11%), HPV39 (11%), HPV58 (10%), HPV52 (8%), and HPV45 (8%).

Conclusions: In this sample of cancer free women in Guatemala, HPV16 was the most prevalent HR type in Guatemala and the age-specific prevalence curve peaked in younger ages. Women in the 30-59-year age groups had a prevalence of HR-HPV of 8%, however, larger studies to better describe the epidemiology of HPV in Guatemala are needed.

Keywords: HPV, Cervical cancer, Guatemala, Prevalence, Screening, Real-time PCR

Background

Across the world, women living in poverty suffer disproportionately from cervical cancer (CC). In Guatemala, CC is a leading cause of cancer in women (1530 cases/year, Age-standardized rate (ASR) 31/100,000) resulting in an estimated 717 deaths) (Globocan 2012). In the Instituto de Cancerología (INCAN), the main adult oncology

hospital in Guatemala, over 40% of women diagnosed with malignancies have advanced CC, requiring costly management that often has a poor outcome [1]. Therefore, a focus on prevention is important.

There is overwhelming evidence that persistent infection with specific types of HPV is the main cause of CC [2, 3]. HPV types that are classified as established carcinogens are HPV16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, and possibly 68 [4]. While prophylactic vaccination of adolescents and possibly young adults is the ultimate preventive strategy, screening will remain important for decades to come. Cytology-based screening has been

* Correspondence: deanm@mail.nih.gov

†Equal contributors

⁴Laboratory of Translational Genomics, Division of Cancer Epidemiology and Genetics, National Cancer Institute, Gaithersburg, MD, USA

Full list of author information is available at the end of the article



associated with a major reduction in the incidence and mortality of the disease in developed countries [5]. However, cytology is either unavailable or poorly conducted in most low-income countries [6].

In most populations, HPV incidence peaks in women in their late teens or early 20s [7–9] following the average age of first sexual intercourse. Incidence and prevalence typically decline from age 30–45 [10]. A second peak is often observed in postmenopausal women, possibly related to immune senescence and escape from long-term cell-mediated immune control of infections acquired earlier in life [11, 12]. HPV DNA testing has been proposed as an alternative to cytology in women older than 25–30 years, when prevalence declines and predictive value (for signaling risk of precancer) of a positive HPV test increases. [13–15]. However, commercially available tests are typically expensive and require sophisticated equipment [16, 17]. The use of HPV assays targeting lower-resource settings would be useful for CC prevention in such settings, which contribute most of the world CC burden.

This study sought to assess the performance of a validated low-cost HPV assay (H13, HybriBio) in the setting of a low and middle income country [18]. We determined the prevalence of HR-HPV infection in women attending screening clinics in Guatemala.

Methods

Study populations

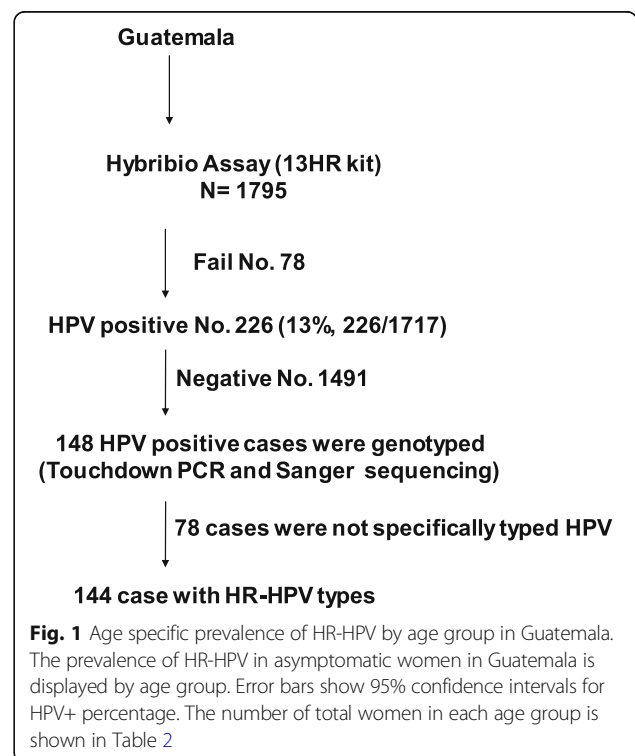
To explore the epidemiology of HPV in Guatemala using H13 (HybriBio Ltd., Hong Kong), a total of 1795 samples were obtained from asymptomatic sexually active women undergoing routine primary screening with Pap smear at hospital-based screening clinics in Guatemala, after obtaining informed written consent. The participants were recruited from women attending cervical cytology screening programs at public hospitals in two urban areas (Guatemala City, and the coastal city of Puerto Barrios), and were tested only one time. Most women live near the respective hospital and so this is a largely urban sample. A questionnaire on reproductive and socio-demographic characteristics was administered by trained personnel on the Guatemalan subjects [19]. The samples were collected by a medical practitioner using a Dacron swab placed in a tube containing 3 ml of Scope mouthwash, maintained at 4° C and transported at room temperature, and stored at -20oC [20].

HPV testing

The detection of HPV was performed using a HybriBio Assay that detects the 13 high-risk HPV types by real-time PCR. The performance of the assay was previously evaluated in a clinical validation study [18]. A cell lysate was prepared per the manufacturer’s instructions and

real-time PCR performed with 1 ul of lysate, 8.75ul of Master mix (containing probes and primers) and 0.25ul of Taq DNA polymerase. Reactions were incubated for 2 min at 50o, 10 min at 95o and 45 cycles of 10 s, 95o; 50 s 58o with data collection at each cycle during the 58o phase on an ABI 9700 or 7000. Each 96-well plate included four HPV+ controls: CC cell line DNA from HeLa (HPV18), CaSki (HPV16), MS751 (HPV45) DNA and positive control DNA from the HybriBio Assay kit; HPV- controls included C33A DNA and water. The kit contains an internal control (human globin gene), and samples that failed to amplify this internal control (78 samples, 4.3%) were excluded from further analysis.

To determine HPV type by sequencing, HPV positive samples were subjected to Touchdown PCR and DNA sequencing (Fig. 1). Samples in which the internal control (IC) did not amplify on two separate attempts were excluded (171 samples). The sensitivity of the Touchdown PCR was determined by a series of 10-fold dilutions of DNA from HPV+ and HPV- cell lines using Broad-Spectrum (BS) GP5+ and GP6+ Primers (BSGP5 +/6+) (150 bp) (Additional file 1: Figure S1) [21]. The HPV+ samples were amplified by using 400 nM BSGP5 +/6+. Briefly, 10 min denaturation step at 95 °C was followed by 40 cycles of amplification. Each cycle included denaturation at 94 °C for 20 s, annealing at 38 °C for 30 s, and elongation at 71 °C for 80 s. The final elongation step was 5 min. The ramping rates were



adjusted as described [22]; 1.8 °C/cycle from 74 °C to 38 °C in first 20 cycles. Each experiment included HPV+ and HPV- controls and a sample lacking template DNA (Additional file 1: Figure S2). The PCR products were subjected to Sanger Sequencing on an ABI3730XL. Sequences were analyzed by assembly and trimming in SeqMan (DNASTAR, Madison, WI) followed by BLAST search (NCBI). Samples with inconclusive Sanger sequence were repeated with a next-generation sequencing method (S. Wagner, and J. Boland, manuscript in preparation).

Statistical analyses

Statistical analyses on the samples were performed to determine age-specific HPV prevalence, comparing the age groups with the Pearson Chi-square test using GraphPad Prism version 7 for Windows. $P < 0.05$ was

regarded as statistically significant. We performed analyses of association between HPV infection and age.

Results

To evaluate the minimum required assay volume of the H13 test we tested the standard 20 ul and 10ul and determined that a 10ul real-time PCR volume gave equivalent results (Table 1); whereas 5ul did not (data not shown). In a separate study, we determined that the H13 test give high concordance with the Qiagen HC2 and Onclarity assays, and has good clinical accuracy compared to histologic diagnosis [18].

To apply the H13 test to samples from a low-middle income country, we used the test to determine the prevalence and distribution of HPV types in Guatemala. We recruited asymptomatic women from the general population at hospital clinics performing primary screening for

Table 1 Comparison of the H13 Assay 10 µl and 20 µl volumes

Sample	Total Volume (µl)	Input DNA (5 ng/µl)		13HR HPV Types (FAM) Ct	Internal Control (JOE) Ct	HPV Status
		Volume (µl)	Input DNA (ng)			
HPV positive controls						
CaSki (HPV16)	20	2	10	15.74	27.26	+
	10	1	5	16.15	28.61	+
HeLa (HPV18)	20	2	10	22.26	26.23	+
	10	1	5	23.03	26.26	+
MS751 (HPV45)	20	2	10	26.90	29.79	+
	10	1	5	27.96	29.91	+
ME180 (HPV39/18)	20	2	10	27.61	25.99	+
	10	1	5	27.72	25.96	+
Unknown samples from cell lysate						
S1 DD015959	20	2		ND ^a	24.05	-
	10	1		ND	24.80	-
S2 DD015960	20	2		17.60	27.59	+
	10	1		18.02	28.10	+
S3 DD015963	20	2		16.85	20.78	+
	10	1		17.20	21.30	+
S6 DD015964	20	2		16.01	22.05	+
	10	1		16.80	22.70	+
HPV negative control (C33A)	20	2	10	ND	25.66	-
	10	1	5	ND	25.04	-
Controls from kit						
Positive control (Kit)	20	2		23.88	23.39	+
	10	1		25.07	25.68	+
Negative control (Kit) ^a	20	2		ND	ND	-
	10	1		ND	ND	-

DNA samples from positive and negative controls along with four unknown samples were assayed using the H13 kit using both 10 and 20ul volumes. The Ct values are shown as well as the interpretation of HPV status

ND not detected

^athe negative control is DNase-free distilled water from the kit

cervical cancer by cytology. The women with available data on age ranged in age from 17 to 79 years attending clinics in Guatemala City and the city of Puerto Barrios. To determine the prevalence of HPV infection, 1717 subjects were tested (Fig. 1). The overall prevalence of HPV infection was 13% (226/1717) (Table 3). To understand the age specific prevalence of HPV infection, the women were divided into 6 age groups (Table 3). HPV prevalence ranged from 21% in the <30 group to 7.0% to 9.0% in the 30-59 age groups and 14% in the ≥60 group. Age specific prevalence for HPV was significantly higher in the younger age group (<30) than the 30-59-year age groups (Table 2 and Fig. 2).

HPV typing was successful in 148 out of 226 HPV+ samples and the 13 HR-HPV types 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 68 were detected in 143 subjects (Table 3.) (143/226). Of the 13 HR types, HPV16 (22%, 32/148), HPV18 (11%, 16/148) accounted for 32% (48/148) and HPV39, 58, 45, 52 combined accounted for 37%, 55/148). Types considered to convey very low risk (Unk), HPV67 and 74 were detected in 1% (Table 3).

Discussion

Few studies of HPV prevalence have been described in Guatemala, even though this country has one of the highest incidences of HPV related diseases and mortality in Latin America [1, 23, 24]. However, our study has the advantage of collecting samples from a screening population using a low-cost HPV screening method. The overall prevalence of HR HPV infection in this study was 13% for Guatemala, similar to rural Costa Rica [25] and other countries (25–29%) [7, 26, 27].

HPV16 is the most common type found in women with cervical cancer worldwide, and is among the most common high-risk type in cancer-free women [26, 28–32], and the same is true in Guatemala. The combined prevalence of HPV16, and HPV18 was highest in the youngest age groups in this country. A similar prevalence of HPV16 and 18 was reported in other studies [33] as well as in Guatemalans with CC [19]. In addition to HPV16 and 18, 15 other HPV types were observed frequently in our study, most notably HPV39, 58, 45, and 52. The relatively small number of women <30 years of age had

the highest prevalence, while the HPV prevalence decreased markedly with increasing age, up to age 60. This trend has been observed in Costa Rica and other countries [7–9, 30].

Numerous studies support HPV testing as the most sensitive primary screening method for CC. However, there are few HPV tests that are affordable for LMICs. The HybriBio H13 test is a sensitive test that costs \$6/assay at the recommended 20ul volume. Commercial tests in Guatemala cost between \$100-210, out of the range of practical use. Mexico carried out a large screening program with the hybrid capture assay (HC2, Qiagen, Germantown, MD) at an approximate cost of \$11 per test, but many poor and rural areas remain unscreened. A method has been developed with support from donor and non-profit foundations, called careHPV (Qiagen). This system has been rigorously tested in China, India and in pilot programs in other areas [34] and is under further evaluation in several Latin American countries. It is unclear whether it can be scaled up to cover an entire whole population. Thus, there is still considerable discussion on the most effective strategies for managing HPV+ women in different economic and cultural settings [35–37].

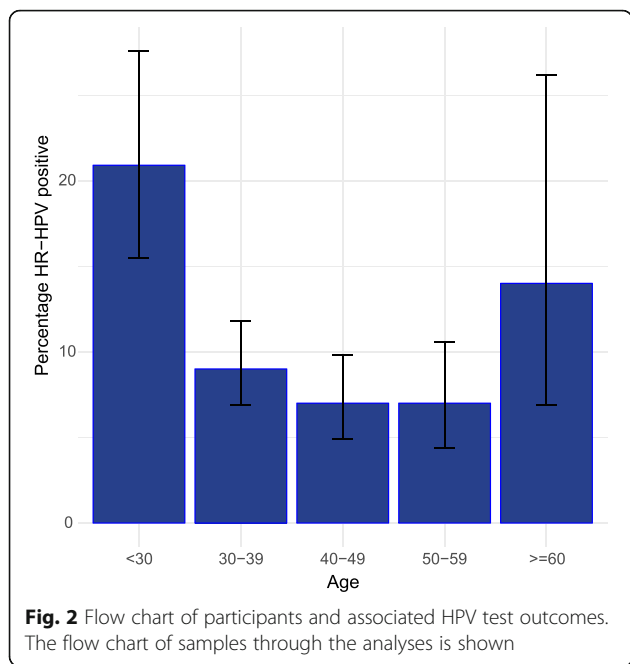
We sought to establish a method that would be cost effective, and use only equipment available in many molecular biology laboratories. We employed a validated storage buffer, mouthwash containing 15% ethanol, which costs \$0.01 per sample with fewer shipping requirements than methanol based buffers [20]. The HybriBio Assay under the conditions we employed (10ul reaction volume), costs about \$3/assay (Table 1). While HybriBio requires a real-time PCR instrument, we have purchased used ABI7000 instruments for under \$1000, and found that some Latin American hospitals and clinical laboratories have real-time PCR instrumentation.

Our study has several limitations that might affect our conclusions. We used a HybriBio assay that detects only 13 HR types. In addition, we attempted to sequence all positive samples to determine type. In a portion of samples (4.3%), we had a failure to detect the IC indicating a failure in sample collection, preservation or storage. Most of these samples were negative using both HybriBio and a touchdown PCR method (data not shown).

Table 2 Analysis of age as related to 13HR HPV detection by the H13 assay

Age group	HPV-No.	HPV+ No.	HPV+ (%) (95% CI)	OR	95% CI of OR	P value
< 30	136	36	21 (16-28)	Ref		
30-39	474	47	9 (7-12)	0.43	0.27 to 0.69	0.0004
40-49	401	30	7 (5- 10)	0.33	0.20 to 0.56	< 0.0001
50-59	244	18	7 (4-11)	0.33	0.18 to 0.60	0.0003
≥60	43	7	14 (7-26)	ns		

Subjects with available data on age (N=1298) were used to determine the effect of age group on HR-HPV infection using the under 30 age group as a reference. The 95% CI of HPV+ (%) was calculated with the Binomial (Clopper-Person) exact method based on the beta distribution
 Ref reference, ns not significant



This could indicate a limitation to using Scope mouthwash as a preservative. We have limited histopathology or other outcome data on the women and have not demonstrated that HybriBio Assay is effective in CC prevention in Guatemala. However, the manufacturer reports data on a comparison with Qiagen (with >95% agreement with an FDA approved test), the test passed two

Table 3 Prevalence of HR-HPV types in Guatemalan women

HPV type	No.	Percentage in 13HR (%)
13 HR types	32	22%
16	16	11%
18	4	3%
31	2	1%
33	6	4%
35	16	11%
39	12	8%
45	5	3%
51	12	8%
52	6	4%
56	15	10%
58	10	7%
59	7	5%
68		
Total	143	

HPV type was determined from Sanger sequence after touch-down PCR or next-generation sequencing. Also detected were individual samples with HPV30, HPV42, HPV52, HPV55, HPV66, HPV67, HPV70, HPV73, HPV86, HPV114 and two samples each with HPV53, HPV71 and HPV74. Multiple infections could only be assessed in 27 samples positive by the next-generation sequencing assay; 6/27 samples (22%) had multiple infections but only 1/27 (4%) had multiple HR infections

WHO proficiency trials and has been used in a study of 48,559 women in China [11].

Conclusions

In conclusion, we have used the HybriBio H13 test, an affordable alternative for HPV screening to determine that the high-risk HPV prevalence is 13% in a population of asymptomatic women in Guatemala. The distribution of HPV types is typical of other countries and the highest HPV prevalence is in youngest age groups (< 30). This low-cost approach to detect HPV could be employed in other countries planning to introduce HPV screening to reduce the burden of CC.

Additional files

Additional file 1: Figure S1. Amplification of HPV positive and negative cell line genomic DNA by Touchdown PCR using BS GP5+/6+ primers. A dilution series of an HPV positive cell line (HeLa), HPV negative (HEK293) and HPV-status unknown cell lines were amplified by touchdown PCR with both HPV primer sets and a globin primer pair, and run on 1.5% agarose gels. **Figure S2.** Touchdown PCR from swab cell lysate, (A): Lane 1.- Lane 4 HPV positive control with input DNA 1 ng, 0.1 ng, 0.01 ng and 0.001 ng; Lane 5. HPV negative control (cell line); Lane 6. no template control. Lane 7 to Lane 16 indicate unknown swab samples. M indicate the DNA ladder marker. Touchdown PCR from HPV positive swabs by HybriBio Assay (B): Lane 1. - Lane 18 are swab cell lysate; Lane 19. HPV positive control (cell line); Lane 20. HPV negative control (cell line); Lane 21. no template control. The samples were amplified with the BSGP5+/6+ primers. (DOCX 2548 kb)

Abbreviations

CC: Cervical cancer; HPV: Human papilloma virus; HR: High-risk; INCAN: Instituto de Cancerología; PCR: Polymerase chain reaction

Acknowledgements

The authors would like to thank the staff and health professionals from the Instituto de Cancerología, Guatemala City, Guatemala, Hospital San Juan de Dios, Guatemala City, Guatemala and Medical Laboratory Services, Willemstad, Curacao, as well as Fundashon Prevencion, Curacao, and the Liga Nacional Contra Cancer, Guatemala; Edmundo Torres Gonzalez for data analysis, Patricia Zaid, Adolfo Santizo, Esther Avila and Lineth Boror for sample and data collection and shipping; Russ Hanson for approvals, Qi Yang and Shenchow Alfred Li for statistical advice and the Cancer Genomics Research Laboratory and BSP-CCR Genetics Core for technical support. The content of this publication does not necessarily reflect the views or policies of the Department of Health and Human Services, nor does mention of trade names, commercial products, or organizations imply endorsement by the U.S. government.

Funding

Supported in part by the Intramural Research Program of the National Institutes of Health, National Cancer Institute, Center for Cancer Research, and Division of Cancer Epidemiology and Genetics from Leidos-Frederick under contract # HHSN261200800001E. The funding body had no role in the design of the study and collection, analysis, and interpretation of data and in writing the manuscript.

Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

HL, LG, JB, SW, JT, HM, FJHBvA performed experimental assays, JS, LG, BK, EG, RO, VA, AR managed samples and data, SPG, GV, HR, HM, EA, HMP, MC, GL

recruited and examined patients, HL, JFD, MS, MD performed data analysis, HL, JFD, MS, MD wrote the manuscript. All authors read and approved the final manuscript.

Ethics approval and consent to participate

This study was approved by the ethics committee of the Hospital General San Juan de Dios, and the Instituto de Cancerología, Guatemala and testing in the US laboratory was judged exempt by the NIH Office of Human Studies Research. All subjects signed an approved informed consent form.

Competing interests

The authors declare that they have no competing interests.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Author details

¹Laboratory of Translational Genomics, Division of Cancer Epidemiology and Genetics, Leidos Biomedical Research, Inc., National Laboratory for Cancer Research, Gaithersburg, MD, USA. ²Instituto de Cancerología, 6^a Avenida 6-58, Zona 11, Guatemala City, Guatemala. ³Cancer Genetics Branch, Gaithersburg, MD, USA. ⁴Laboratory of Translational Genomics, Division of Cancer Epidemiology and Genetics, National Cancer Institute, Gaithersburg, MD, USA. ⁵Department of Gynecology and Obstetrics, Hospital General San Juan de Dios, Guatemala City, Guatemala. ⁶Cancer Research Technology Program, Leidos Biomedical Research, Inc., National Cancer Institute, 8560 Progress Drive, Frederick, MD 21701, USA. ⁷Hospital Central Universitario "Dr. Antonio M Pineda", Lara State, Barquisimeto, Venezuela. ⁸Department of Pathology, Hospital General San Juan de Dios, Guatemala City, Guatemala. ⁹Medical Laboratory Services, Willemstad, Curaçao. ¹⁰Fundashon Prevencionshon, Willemstad, Curaçao.

Received: 25 September 2017 Accepted: 26 April 2018

Published online: 15 May 2018

References

- Valles X, Murga GB, Hernandez G, Sabido M, Chuy A, Lloveras B, Alameda F, de San Jose S, Bosch FX, Pedroza I, et al. High prevalence of human papillomavirus infection in the female population of Guatemala. *Int J Cancer*. 2009;125(5):1161–7.
- Ngelangel C, Munoz N, Bosch FX, Limson GM, Festin MR, Deacon J, Jacobs MV, Santamaria M, Meijer CJ, Walboomers JM. Causes of cervical cancer in the Philippines: a case-control study. *J Natl Cancer Inst*. 1998;90(1):43–9.
- Smith JS, Lindsay L, Hoots B, Keys J, Franceschi S, Winer R, Clifford GM. Human papillomavirus type distribution in invasive cervical cancer and high-grade cervical lesions: a meta-analysis update. *Int J Cancer*. 2007;121(3):621–32.
- Bouvard V, Baan R, Straif K, Grosse Y, Secretan B, El Ghissassi F, Benbrahim-Tallaa L, Guha N, Freeman C, Galichet L, et al. A review of human carcinogens—part B: biological agents. *Lancet Oncol*. 2009;10(4):321–2.
- Arbyn M, Raifu AO, Weiderpass E, Bray F, Anttila A. Trends of cervical cancer mortality in the member states of the European Union. *Eur J Cancer*. 2009;45(15):2640–8.
- Gakidou E, Nordhagen S, Obermeyer Z. Coverage of cervical cancer screening in 57 countries: low average levels and large inequalities. *PLoS Med*. 2008;5(6):e132.
- de Ona M, Alvarez-Arguelles ME, Torrents M, Villa L, Rodriguez-Feijoo A, Palacio A, Boga JA, Tamargo A, Melon S. Prevalence, evolution, and features of infection with human papillomavirus: a 15-year longitudinal study of routine screening of a women population in the north of Spain. *J Med Virol*. 2010;82(4):597–604.
- Leinonen MK, Anttila A, Malila N, Dillner J, Forslund O, Nieminen P. Type- and age-specific distribution of human papillomavirus in women attending cervical cancer screening in Finland. *Br J Cancer*. 2013;109(11):2941–50.
- Ley C, Bauer HM, Reingold A, Schiffman MH, Chambers JC, Tashiro CJ, Manos MM. Determinants of genital human papillomavirus infection in young women. *J Natl Cancer Inst*. 1991;83(14):997–1003.
- Schiffman M, Doorbar J, Wentzensen N, de Sanjose S, Fakhry C, Monk BJ, Stanley MA, Franceschi S. Carcinogenic human papillomavirus infection. *Nat Rev Dis Primers*. 2016;2:16086.
- Chen Q, Xie LX, Qing ZR, Li LJ, Luo ZY, Lin M, Zhang SM, Chen WZ, Lin BZ, Lin QL, et al. Epidemiologic characterization of human papillomavirus infection in rural Chaozhou, eastern Guangdong Province of China. *PLoS One*. 2012;7(2):e32149.
- Syrjanen K. New concepts on risk factors of HPV and novel screening strategies for cervical cancer precursors. *Eur J Gynaecol Oncol*. 2008;29(3):205–21.
- Mayrand MH, Duarte-Franco E, Rodrigues I, Walter SD, Hanley J, Ferenczy A, Ratnam S, Coutlee F, Franco EL. Human papillomavirus DNA versus Papanicolaou screening tests for cervical cancer. *N Engl J Med*. 2007;357(16):1579–88.
- Naucler P, Ryd W, Tornberg S, Strand A, Wadell G, Elfgrén K, Radberg T, Strander B, Johansson B, Forslund O, et al. Human papillomavirus and Papanicolaou tests to screen for cervical cancer. *N Engl J Med*. 2007;357(16):1589–97.
- Rijkaart DC, Berkhof J, Rozendaal L, van Kemenade FJ, Bulkman NW, Heideman DA, Kenter GG, Cuzick J, Snijders PJ, Meijer CJ. Human papillomavirus testing for the detection of high-grade cervical intraepithelial neoplasia and cancer: final results of the POBASCAM randomised controlled trial. *Lancet Oncol*. 2012;13(1):78–88.
- Barski A, Cuddapah S, Cui K, Roh TY, Schones DE, Wang Z, Wei G, Chepelev I, Zhao K. High-resolution profiling of histone methylations in the human genome. *Cell*. 2007;129(4):823–37.
- Green RE, Krause J, Ptak SE, Briggs AW, Ronan MT, Simons JF, Du L, Egholm M, Rothberg JM, Paunovic M, et al. Analysis of one million base pairs of Neanderthal DNA. *Nature*. 2006;444(7117):330–6.
- Fokom Domgue J, Schiffman M, Wentzensen NH, Gage JC, Castle PE, Raine-Bennett TR, Fetterman B, Lorey T, Poitras NE, Befano B, et al. Assessment of a new lower-cost real-time PCR assay for detection of high-risk human papillomavirus: useful for cervical screening in limited-resource settings? *J Clin Microbiol*. 2017;55(8):2348–55.
- Lou H, Villagran G, Boland JF, Im KM, Polo S, Zhou W, Odey U, Juarez-Torres E, Medina-Martinez I, Roman-Basaure E, et al. Genome analysis of Latin American cervical cancer: frequent activation of the PIK3CA pathway. *Clin Cancer Res*. 2015;21(23):5360–70.
- Castle PE, Sadorra M, Garcia FA, Cullen AP, Lorincz AT, Mitchell AL, Whitby D, Chuke R, Kornegay JR. Mouthwash as a low-cost and safe specimen transport medium for human papillomavirus DNA testing of cervicovaginal specimens. *Cancer Epidemiol Biomark Prev*. 2007;16(4):840–3.
- Schmitt M, Dondog B, Waterboer T, Pawlita M. Homogeneous amplification of genital human alpha papillomaviruses by PCR using novel broad-spectrum GP5+ and GP6+ primers. *J Clin Microbiol*. 2008;46(3):1050–9.
- Snijders PJ, van den Brule AJ, Jacobs MV, Pol RP, Meijer CJ. HPV DNA detection and typing in cervical scrapes. *Methods Mol Med*. 2005;119:101–14.
- Fokom Domgue J, Schiffman M, Wentzensen NH, Gage JC, Castle PE, Raine-Bennett TR, Fetterman B, Lorey T, Poitras NE, Befano B, et al. Assessment of a new lower-cost real-time PCR detection assay for high-risk HPV: useful for cervical screening in limited resource settings? *J Clin Microbiol*. 2017;55:2348.
- Nunez-Troconis J, Delgado M, Gonzalez J, Mendiola R, Velasquez J, Conde B, Whitby D, Munroe DJ. Prevalence and risk factors of human papillomavirus infection in asymptomatic women in a Venezuelan urban area. *Investig Clin*. 2009;50(2):203–12.
- Herrero R, Hildesheim A, Bratti C, Sherman ME, Hutchinson M, Morales J, Balmaceda I, Greenberg MD, Alfaro M, Burk RD, et al. Population-based study of human papillomavirus infection and cervical neoplasia in rural Costa Rica. *J Natl Cancer Inst*. 2000;92(6):464–74.
- de Sanjose S, Diaz M, Castellsague X, Clifford G, Bruni L, Munoz N, Bosch FX. Worldwide prevalence and genotype distribution of cervical human papillomavirus DNA in women with normal cytology: a meta-analysis. *Lancet Infect Dis*. 2007;7(7):453–9.
- Dunne EF, Unger ER, Sternberg M, McQuillan G, Swan DC, Patel SS, Markowitz LE. Prevalence of HPV infection among females in the United States. *JAMA*. 2007;297(8):813–9.
- Alibegashvili T, Clifford GM, Vaccarella S, Baidoshvili A, Gogiasvili L, Tsagareli Z, Kureli I, Snijders PJ, Heideman DA, van Kemenade FJ, et al. Human papillomavirus infection in women with and without cervical cancer in Tbilisi, Georgia. *Cancer Epidemiol*. 2011;35(5):465–70.
- Zhao R, Zhang WY, Wu MH, Zhang SW, Pan J, Zhu L, Zhang YP, Li H, Gu YS, Liu XZ. Human papillomavirus infection in Beijing, People's Republic of China: a population-based study. *Br J Cancer*. 2009;101(9):1635–40.

30. Guan P, Howell-Jones R, Li N, Bruni L, de Sanjose S, Franceschi S, Clifford GM. Human papillomavirus types in 115,789 HPV-positive women: a meta-analysis from cervical infection to cancer. *Int J Cancer*. 2012;131(10):2349–59.
31. Lo KW, Wong YF, Chan MK, Li JC, Poon JS, Wang VW, Zhu SN, Zhang TM, He ZG, Wu QL, et al. Prevalence of human papillomavirus in cervical cancer: a multicenter study in China. *Int J Cancer*. 2002;100(3):327–31.
32. Bosch FX, Manos MM, Munoz N, Sherman M, Jansen AM, Peto J, Schiffman MH, Moreno V, Kurman R, Shah KV. Prevalence of human papillomavirus in cervical cancer: a worldwide perspective. International biological study on cervical cancer (IBSCC) study group. *J Natl Cancer Inst*. 1995;87(11):796–802.
33. Haguenoer K, Giraudeau B, Gaudy-Graffin C, de Pinieux I, Dubois F, Trignol-Viguiet N, Viguiet J, Marret H, Goudeau A. Accuracy of dry vaginal self-sampling for detecting high-risk human papillomavirus infection in cervical cancer screening: a cross-sectional study. *Gynecol Oncol*. 2014;134(2):302–8.
34. Trope LA, Chumworathayi B, Blumenthal PD. Preventing cervical cancer: stakeholder attitudes toward CareHPV-focused screening programs in Roi-et Province, Thailand. *Int J Gynecol Cancer*. 2009;19(8):1432–8.
35. Gottschlich A, Rivera-Andrade A, Grajeda E, Alvarez C, Montano M, Meza R. Acceptability of human papillomavirus self-sampling for cervical cancer screening in an indigenous community in Guatemala. *J Glob Oncol*. 2017; <https://doi.org/10.1200/JGO.2016.005629>.
36. Jeronimo J, Holme F, Slavkovsky R, Camel C. Implementation of HPV testing in Latin America. *J Clin Virol*. 2016;76(Suppl 1):S69–73.
37. Schiffman M, Wentzensen N, Wacholder S, Kinney W, Gage JC, Castle PE. Human papillomavirus testing in the prevention of cervical cancer. *J Natl Cancer Inst*. 2011;103(5):368–83.

Ready to submit your research? Choose BMC and benefit from:

- fast, convenient online submission
- thorough peer review by experienced researchers in your field
- rapid publication on acceptance
- support for research data, including large and complex data types
- gold Open Access which fosters wider collaboration and increased citations
- maximum visibility for your research: over 100M website views per year

At BMC, research is always in progress.

Learn more biomedcentral.com/submissions

