



ORIGINAL ARTICLE

Baseline prevalence and type distribution of *Human papillomavirus* in sexually active non-vaccinated adolescent girls from Argentina



Joaquín Víctor González^a, Gerardo Daniel Deluca^b, Domingo Javier Liotta^{c,d}, Rita Mariel Correa^a, Jorge Alejandro Basiletti^a, María Celeste Colucci^a, Nathalia Katz^e, Carla Vizzotti^e, María Alejandra Picconi^{a,*}, on behalf of the MALBRAN HPV Surveillance Study Group

^a Servicio Virus Oncogénicos, Instituto Nacional de Enfermedades Infecciosas - ANLIS "Dr. Malbrán", Av. Velez Sarsfield 563, C1282AFF, Buenos Aires, Argentina

^b Facultad de Medicina, Universidad Nacional del Nordeste, Mariano Moreno 1240, W3400ACX, Corrientes, Argentina

^c Laboratorio de Biología Molecular Aplicada, Facultad de Ciencias Exactas, Químicas y Naturales, Universidad Nacional de Misiones, Av. Mariano Moreno 1375, N3300, Posadas, Misiones, Argentina

^d Instituto Nacional de Medicina Tropical-ANLIS "Dr. Malbrán", Neuquén y Jujuy s/n, N3370, Puerto Iguazú, Misiones, Argentina

^e Dirección de Control de Enfermedades Inmunoprevenibles, Ministerio de Salud de la Nación, Rivadavia 875, C1002AAG, Buenos Aires, Argentina

Received 15 November 2019; accepted 17 June 2020

Available online 9 August 2020

KEYWORDS

Human papillomavirus;
Monitoring;
Vaccines;
HPV infection;
Adolescent women

Abstract In 2011, Argentina launched a government-funded national *Human papillomavirus* (HPV) immunization program incorporating a bivalent HPV vaccine, with a 0–1–6-month schedule, for girls 11 years of age, born after January 2000. Monitoring the changes of HPV infection prevalence among young women has been proposed as an endpoint for early assessment of HPV vaccination programs. However, the data on HPV prevalence at young ages are very limited. The aim of this work was to determine the prevalence of HPV infection and type-specific distribution in sexually active 15–17-year-old non-vaccinated girls. Cervical samples from 1073 adolescents were collected for HPV detection and genotyping using the BSGP5+/GP6+PCR-reverse line blot (RLB) assay. Out of 957 specimens analyzed, 56.3% were positive for any HPV type; 42.2% harbored at least one high-risk HPV (HR-HPV) type and 30.8% low-risk HPV (LR-HPV) types. Multiple and single infections were identified in 36.3% and 20.0% of the samples respectively. The 6 most common HR-HPV types were HPV16 (11.1%), HPV52 (10.8%), HPV56 (8.3%), HPV51 (7.4%), HPV58 (7.3%) and HPV31 (7.1%). The prevalence of HR-HPV-16/18 was 15.2%. In conclusion, results

* Corresponding author.

E-mail addresses: mapicconi@anlis.gov.ar, mapicconi@gmail.com (M.A. Picconi).

confirm that HPV (particularly HR-types) are very common among sexually active adolescents, and prevalence rises quickly after their sexual debut. Our HPV type-specific prevalence baseline may be used to monitor post-vaccinal longitudinal changes in Argentina.

© 2020 Asociación Argentina de Microbiología. Published by Elsevier España, S.L.U. This is an open access article under the CC BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).

PALABRAS CLAVE

Virus del papiloma humano;
Monitorización;
Vacunas;
Vigilancia;
Adolescentes

Prevalencia basal y distribución por tipo específica del virus del papiloma humano en mujeres adolescentes sexualmente activas y no vacunadas de Argentina

Resumen En 2011 se introdujo en el Calendario Nacional de Vacunación de Argentina una vacuna bivalente contra el virus del papiloma humano (HPV, por sus siglas en inglés) tipo 16 y 18, para ser aplicada en niñas de 11 años. Se propuso la monitorización de los cambios en la prevalencia de infección por este virus HPV en las adolescentes como un criterio de valoración temprano para evaluar programas de vacunación; sin embargo, los datos sobre prevalencia del HPV en la población joven son limitados. El objetivo de este estudio fue determinar la prevalencia de la infección por el HPV y su distribución tipo-específica en mujeres de 15 a 17 años, sexualmente activas y no vacunadas. Se colectaron 1.073 muestras cervicales para la detección y tipificación del HPV mediante la técnica de PCR BSGP5⁺/GP6⁻-hibridación reversa en línea (RLB). Sobre 957 muestras analizadas, la frecuencia global de infección fue del 56,3%. El 42,2% de los casos positivos presentó algún genotipo de alto riesgo oncogénico (HPV-AR) y un 30,8% al menos un genotipo de bajo riesgo oncogénico (HPV-BR). El 20,0 y el 36,3% fueron infecciones simples y múltiples, respectivamente. Los HPV-AR más prevalentes fueron HPV16 (11,1%), HPV52 (10,8%), HPV56 (8,3%), HPV51 (7,4%), HPV58 (7,3%) y HPV31 (7,1%). La prevalencia conjunta de HPV16/18 fue del 15,2%. En conclusión, los HPV (en especial los HPV-AR) son muy frecuentes en adolescentes jóvenes sexualmente activas, con un rápido aumento de la prevalencia tras el comienzo sexual. La prevalencia tipo-específica basal registrada podrá servir para monitorizar cambios en la prevalencia de los tipos vacunales y no vacunales en el tiempo.

© 2020 Asociación Argentina de Microbiología. Publicado por Elsevier España, S.L.U. Este es un artículo Open Access bajo la licencia CC BY-NC-ND (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).

Introduction

Genital *Human papillomavirus* (HPV) is the most common sexually transmitted viral infection²⁹. More than 200 HPV types have been identified and officially recognized by the International Human Papillomavirus Reference Center, located at the Karolinska Institute in Stockholm, Sweden¹⁹.

Given their oncogenic potential and clinical relevance, approximately 40 HPV types within the genus *Alphapapillomavirus* are roughly subdivided into high-risk (HR) and low-risk (LR) HPV types^{2,34,45,50}. According to the classification of the International Agency for Research on Cancer (IARC-WHO), HPV types 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58 and 59 are *carcinogenic to humans* (Group 1), being etiologically associated with more than 98% of cervical cancers (CCs), 70–90% of anal and vaginal cancers, 40% of vulvar cancers, 47% of penile cancers, and 25 to 30% of oropharyngeal cancers¹¹. HPV 68 is *probably carcinogenic* to humans (Group 2A). HPV types 26, 30, 34, 53, 66, 67, 69, 70, 73, 82, 85, and 97 are classified as *possibly carcinogenic* to humans (Group 2B). HPV types 6 and 11, the archetypes of LR-HPVs, are *not classifiable as to their carcinogenicity to humans* (Group 3),

being associated with more than 90% of anogenital warts and laryngeal papillomas (benign lesions)²⁵.

In Argentina, around 4500 new CC cases are diagnosed every year, and almost half of the women affected die²³.

The introduction of HPV vaccines offers a valuable tool to prevent CCs. Currently, three prophylactic HPV vaccines are available: a bivalent HPV16/18 vaccine (Cervarix; GlaxoSmithKline Biologicals, Belgium), a quadrivalent HPV6/11/16/18 vaccine (Gardasil/Silgard; Merck & Co., USA/Sanofi Pasteur MSD, France), and a nonavalent HPV6/11/16/18/31/33/45/52/58 vaccine (Gardasil9; Merck & Co.). After the first 13 years of routine use of HPV vaccines, comprehensive clinical trials and real-life data have confirmed their safety and effectiveness⁴⁷.

The World Health Organization (WHO) recommends the inclusion of the HPV vaccine in national immunization programs primarily targeting girls aged 9–14, prior to becoming sexually active, and as a secondary target, females aged ≥15, or males (only when feasible and cost-effective, and without diverting resources from the primary population vaccination program)⁴⁷.

To prevent CC, in October 2011 Argentina launched the most comprehensive government-funded national HPV

prevention program of Latin America, incorporating a bivalent HPV vaccine, with a 0–1–6-month schedule, for girls 11 years of age, born after January 2000¹⁴. This intervention also involved reinforcing CC screening in women aged 30–64 as a secondary prevention strategy, while gradually introducing HPV testing³. In 2014, the programme switched to the quadrivalent vaccine, and was extended to males and females aged 11 to 26 living with HIV, and transplanted individuals with a 3-dose schedule (0, 2 and 6 months). In 2015 the number of doses was reduced to two and two years later, vaccination was extended to boys aged 11, born after 2006¹⁴.

Since the first generation of HPV vaccines (bivalent and quadrivalent) protects against some, but not all the types capable of causing associated cancers, monitoring both the positive and negative effects of these vaccines on other HPV types is critical to evaluate their impact on the population. Data on type-specific HPV distribution may be useful to evaluate the additional benefits of vaccines, e.g. cross protection against non-vaccine types and herd immunity, and to monitor unlikely adverse phenomena such as type replacement.

Because the major benefit of HPV vaccination -the prevention of cervical and other less common HPV-associated cancers-will take decades to be seen, a number of previous outcomes are being monitored toward an earlier assessment of the impact of HPV vaccines, primarily in young adolescents^{7,49}. The short-term evidence of the effectiveness of the vaccine program is crucial to encourage government planners to sustain and improve HPV vaccination services.

Since no data existed on the HPV burden in adolescent women in Argentina, a study was conducted to determine the type-specific HPV prevalence in cervicovaginal samples from sexually active, unvaccinated girls aged 15–17 years. This data provides an important baseline to compare future changes in overall and type-specific HPV prevalence after HPV vaccination.

Materials and methods

Study design

From April 2014 to October 2015, a cross sectional study about HPV genotype circulation in unvaccinated girls was conducted by the National and Regional Reference HPV Laboratory (N&R-HPV Lab) (National Institute of Infectious Diseases, ANLIS Malbrán, Buenos Aires).

When girls sought medical advice, particularly about contraception, the gynecologists of the Adolescence Services invited them to join the study. Inclusion criteria was: no HPV vaccine, 15–17 years of age, and sexually active.

Fifteen years of age was established as the lower limit of the reference range since that is the average age at which adolescents start sexual activity in Argentina⁴².

Due to difficulties related to the target population, a convenience sampling method was designed and used.

For the adolescent population sample to be broad and representative, girls were recruited in six public hospitals from four cities of Argentina: *Hospital Evita Pueblo* (Berazategui, Province of Buenos Aires; east-central area of Argentina), *Hospital Madariaga* (Posadas, Province of

Misiones; northeastern area of Argentina), *Centro Integral de Salud La Banda* (La Banda, Province of Santiago del Estero, north-central area of Argentina; *Hospital Argerich*, *Hospital Rivadavia* and *Hospital Durand* (Buenos Aires city, Capital city, east-central area of Argentina).

Recruitment and sample collection

The girls who agreed to participate signed an informed consent and answered a short questionnaire that included basic information on age, date and place of birth, current address and age of sexual debut. These data are relevant to define certain basic characteristics of the group under investigation, and allow for a valid comparison with data to be obtained from vaccinated girls in the near future.

In order to standardize the collection, storage and transfer of the samples obtained in the different centers, a Standard Operating Procedure (SOP) was developed and shared in the training workshops carried out in each of the centers before launching the project. Audit visits were made to evaluate the progress of these processes.

Cervical samples were obtained from both the endocervix and the ectocervix, using a Cytobrush (hc2 DNA Collection Device, Qiagen) introduced in the cervical canal and rotated 360°, 5 times. The cellular material was placed in a vial containing specimen transport medium (STM, Qiagen) and stored at +4 °C until its shipment (within fifteen days) to the N&E-HPV Lab for processing.

DNA extraction from cervical samples

DNA was extracted from 200 µl of STM sample using commercial columns (Qiagen) on a robotic system (QIAcube system, Qiagen), following the manufacturer's instructions. DNAs were stored at –20 °C until use. For every ten samples, a tube containing only STM medium was included as a control for contamination.

HPV detection and genotyping

HPV detection and genotyping was performed by PCR using Broad-Spectrum General Primers (BSGP) 5+/6+ biotin-labeled designed to amplify a highly conserved 140 bp fragment of the HPV-L1 gene^{41,44}. DNA amplification was combined with a reverse line hybridization (RLB) which identifies 36 HPV genotypes (6, 11, 16, 18, 26, 31, 33, 34, 35, 39, 40, 42, 43, 44, 45, 51, 52, 53, 54, 55, 56, 57, 58, 59, 61, 66, 68, 70, 71, 72, 73, 81, 82, 83, 84 and 89), including all the HR-HPV established to date.

Briefly, the denatured biotinylated amplicons were hybridized with genotype-specific oligonucleotide probes immobilized as parallel lines on a Biodyne C membrane (Life Science) using the Miniblotter MN45. The hybrids were treated with alkaline phosphatase-streptavidin conjugate (GE HealthCare) and substrate (ECL Detection Reagents, GE HealthCare), resulting in a chemiluminescent product subsequently detected by exposure to autoradiography film. The β-globin gene (208 bp fragment) was co-amplified and its relative abundance detected for each sample to serve as internal control⁴¹.

Every year the N&R-HPV Lab participates in the external control of HPV genotyping (HPV DNA typing proficiency panel), provided by the International Papillomavirus Reference Center, which serves as Quality Control of the results obtained in this study¹⁶.

Statistical analyses

Data were entered and analyzed in an especially designed database, and subsequently processed using the SPSS Statistics 23.0.0.3 (IBM) statistical software. The association between proportions was assessed by the chi-square test and the 95% confidence interval (CI) was considered in all cases. For all the data analyses conducted, *p* values <0.05 were considered statistically significant.

Ethical and legal considerations

The present study was approved by the ANLIS Malbrán Research Ethics Board and the Ethics Committee of each participating hospital to ensure confidentiality and anonymity. A written informed consent was obtained before enrolling the study participants.

Under the Argentine legal system, adolescents over 14 years of age have the right to be assisted and receive health-related care unaccompanied by an adult or guardian; they may also decide to participate in research projects and sign their consents.

Because the teenage girls invited to participate in the study were not within the age range for CC screening, special care was taken to report that it was only a research study; the opportunity was taken to teach basic concepts about HPV infection and health care which included the age at which screening should begin to avoid generating alarm and/or an inappropriately early screening onset.

In some provinces, social workers gave talks in secondary schools informing about the study within the framework of health care counseling seminars.

Results

The study enrolled 1073 unvaccinated, sexually active girls who provided cervical cell samples; 116 were excluded because their ages were not within the established range. The remaining 957 samples were amplifiable for beta-globin, thus being adequate for the PCR analysis; therefore, this analysis refers to them. The distribution of the samples collected in the four cities was as follows: 272 from Berazategui, 172 from Posadas; 215 from La Banda, and 298 from Buenos Aires.

The prevalence of HPV infection of any type was 56.3% (95% CI; 53.2–59.5), 42.2% (95% CI; 39.1–45.3) were infected with at least one HR-HPV genotype and 7.9% (95% CI; 6.2–9.7) with LR-HPV genotypes only.

The frequencies of the different genotypes in descending order are presented in Figure 1. The 6 most common HR types were HPV16 [11.1%; (95% CI, 9.1–13.1)], HPV52 [10.8%; (95% CI, 8.8–12.7)], HPV56 [8.3%; (95% CI, 6.5–10.0)], HPV51 [7.4%; (95% CI, 5.8–9.1)], HPV58 [7.3%; (95% CI, 5.7–9.0)] and HPV31 [7.1%; (95% CI, 5.5–8.7)].

Among the LR types, HPV42 had the highest prevalence [11.5% (95% CI, 9.5–13.5)], followed by HPV6 [6.7%; (95% CI, 5.1–8.3)], HPV40 and HPV81 [5.0%; (95% CI, 3.6–6.4)] and HPV11 [3.1%; (95% CI, 2.0–4.2)].

The prevalence of types closely related to the HPV vaccine, types against which cross-protection has been reported in clinical trials, and types according to IARC criteria are shown in Table 1.

Data about HPV prevalence broken down by city of sample collection are shown in Table 2. Significant differences were observed for the prevalence of HPV infection of any type only between Berazategui and the rest of the cities (*p* < 0.000); the prevalence of HPV16/18 and 6/11/16/18 differed significantly only between Berazategui and Posadas (*p* = 0.001 and *p* = 0.004, respectively).

In 36.3% (95% CI: 33.3–39.4) multiple infections were identified and up to 9 different HPV types were found in one sample (Table 3).

Discussion

Several studies have shown cervical HPV prevalence and type distribution in women older than 18 years of age in different study designs worldwide^{10,13,22,30,33,36,38}; however, there is very scarce information for young adolescents. The present work reports HPV prevalence and genotype distribution among 957 sexually active non-vaccinated girls from Argentina, aged 15–17 years.

In this study, the overall HPV prevalence (56.3%) was similar to previous local data from the subgroup of 15–23-year-old (y) women from Buenos Aires and Santa Fe (Argentina) (46.7% and 64.2%, respectively)⁹, and also close to that obtained for comparable age groups in other regions like Sweden (60%, subgroup 16 y)³⁷, USA (49% for 11–19 y¹⁵; 71% for 14–17 y¹⁷), Australia (66.3%; subgroup under 21 y)⁴³, and Colombia (60.3% for 18–25 y³⁵; and 49% for 12–19 y¹²), but higher than Africa where 36% was reported in non-vaccinated women aged ≤16 years⁴⁶. Results confirm that this infection is very common among sexually active adolescents, and rises quickly after their sexual debut^{21,48}.

The present study showed a prevalence of 42.2% for HR-HPV types (Group 1 and 2A IARC), which is consistent with previous surveys from Colombia (48% for 12–19 y)¹², Denmark (40% for 15–19 y)²⁸, Sweden (over 50% for 16 y)³⁷ and France (35.71%; 15–18 y)⁴.

The prevalence of HPV infection of any type was similarly high in all the cities of sample collection, with the exception of Berazategui, where a significant smaller value was recorded (Table 2). Although Berazategui is located in the same geographical region as Buenos Aires city, it has very different urban and demographic characteristics since it is a small town (170 000 inhabitants). Anyhow, this observation is difficult to explain because we have no information about the participants' variables related to sexual behavior such as the number of sexual partners. Furthermore, we cannot rule out that these estimates may be unstable due to the limited sample size by city.

A high prevalence of HR-HPV infections in young women does not necessarily mean a high rate of persistent infection, a pre-requisite for the development of precancerous and cancerous lesions, since most of these infections are

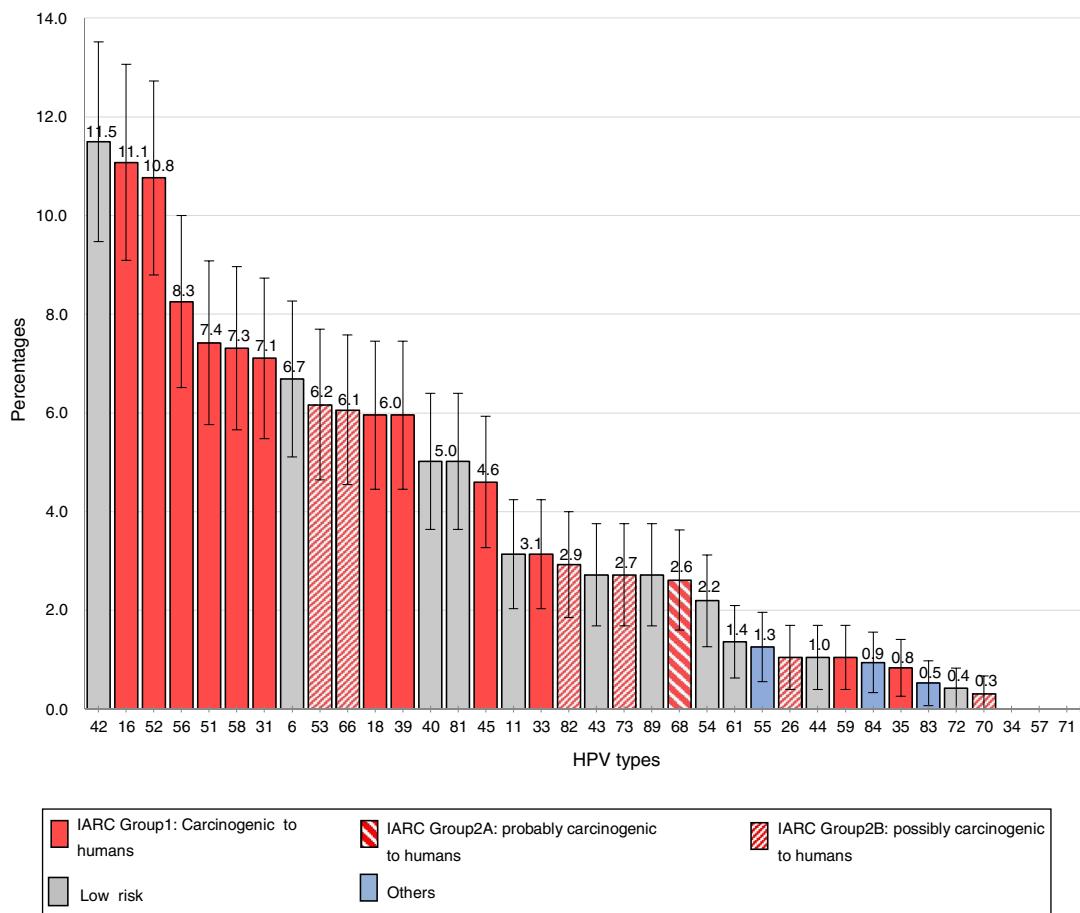


Figure 1 Prevalence, with 95% CI, of all identified HPV types in non-vaccinated sexually active 15–17-year old girls from Argentina.

transient and will be cleared in less than two years²⁴; this is the main reason for CC screening to be discouraged under 25 years of age⁴⁰. In cross-sectional prevalence studies such as this one, it is not possible to differentiate transient infections from those that will persist; however, premature contact with HR-HPV at the onset of adolescence could increase the risk of cervical disease at an earlier age if a persistent infection is established.

Estimates of type-specific HPV prevalence in cervical samples vary greatly worldwide among different populations according to age range, cytological status, laboratory method used, and the biological interaction between different viruses and the host's immunogenetic factors¹⁰. In this study, it was found that 15.2% of the girls were infected with HPV 16/18, the HR types included in the bivalent and quadrivalent vaccines. This rate is close to that reported in Denmark (13.6%) for women aged 15–19 y²⁸, England (11.5%) for 13–15 y²² and USA (14.3%) for 11–19 y¹⁵.

It is well known that HPV vaccines do not impact on infections that are present at the time of immunization²⁰. Under the public health plan, the Argentine regular immunization program introduced HPV vaccination in 11-year-old girls to get the maximum benefit. The fact that 15% of the girls participating in this study were already infected by HPV16/18 supports the decision of local sanitary authorities to vaccinate at an early age, before their sexual debut⁴⁷. However,

the vaccines were also approved for older women who were already sexually active; since it is unlikely that they were infected with all the vaccine HPV types, they could benefit from vaccination and speed up the public health impact of vaccination and increase short-term benefits⁵.

The 6 most common HR types identified in this work were HPV16 (11.1%), HPV52 (10.8%), HPV56 (8.3%), HPV51 (7.4%), HPV58 (7.3%) and HPV31 (7.1%), 4 of which were included in the nonavalent vaccine formula; this could represent a beneficial future scenario if the nonavalent vaccine were used.

In the case of the bivalent and quadrivalent HPV vaccines, the possibility of cross-protection against other HPV types is an extremely important aspect since it could increase the number of prevented cervical cancers. To date, there is evidence on the bivalent and quadrivalent vaccines cross-protection against HPV types 31, 33 and 45⁴⁷. It was shown that the efficacy in preventing 6-month persistent infection in HPV naive women vaccinated with the current vaccines does not exceed 30% for non-vaccine genotypes, namely HPV 31, 33, 45, 52, 58^{8,31}. However, a more recent report from Scotland revealed that immunization with the bivalent vaccine has led to a startling reduction in vaccine and cross-protective HPV types seven years after vaccination; there is also evidence of herd protection against the vaccine-specific and cross-protective types (HPV types 31, 33 and 45)^{6,26}.

Table 1 Prevalence of selected HPV types and groups of types among sexually active non-vaccinated 15–17-year-old girls from Argentina.

HPV type	Tested adolescents (n = 957)	
	n (%)	95% CI (%)
Vaccine types		
Bivalent/Tetravalent types		
HPV 16	106 (11.1)	(9.1–13.1)
HPV 18	57 (6.0)	(4.4–7.5)
HPV 16, 18	145 (15.2)	(12.9–17.4)
HPV 6	64 (6.7)	(5.1–8.3)
HPV 11	30 (3.1)	(2.0–4.2)
HPV 6, 11	91 (9.5)	(7.6–11.4)
HPV 6, 11, 16, 18	215 (22.5)	(19.8–25.1)
Nonavalent types		
HPV 6, 11, 16, 18, 31, 33, 45, 52, 58	368 (38.5)	(35.4–41.5)
HR vaccine and closely related types		
HPV 16, 18, 31, 33, 45	237 (24.8)	(22.0–27.5)
HPV 31, 33, 45	130 (13.6)	(11.4–15.8)
Alpha 9 species (HPV 16, 31, 33, 35, 52, 58)	278 (29.0)	(26.2–32.0)
Alpha 9 species (without HPV16)	172 (18.0)	(15.5–20.4)
Alpha7 species (HPV 18, 39, 45, 59, 68, 70)	169 (17.7)	(15.2–20.1)
Alpha 7 species (without HPV 18)	112 (11.7)	(9.7–13.7)
HR types according to IARC criteria		
HR-HPV13 types (Group 1 and 2A) ^a	404 (42.2)	(39.1–45.3)
HR-HPV 20 types (Group 1, 2A and 2B) ^a	455 (47.5)	(44.4–50.7)
HR HPV 18 types (Group 1, 2A and 2B without HPV 16,18)	310 (32.4)	(29.4–35.4)
LR types included and not included in the vaccine		
HPV LR	295 (30.8)	(27.9–33.7)
HPV LR (without HPV 6,11)	191 (20.0)	(17.4–22.5)

HR-types: High risk HPV types; LR: low risk HPV types.

^a Group 1 IARC (carcinogenic to humans): HPV 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58 and 59. Group 2A IARC (probably carcinogenic to humans): HPV68. Group 2B IARC (possibly carcinogenic to humans): HPV26, 30, 34, 53, 66, 67, 69, 70, 73, 82, 85 and 97.

Table 2 Prevalence of HPV infection of any type, HPV 16/18, and HPV 6/11/16/18 by city of recruitment, among sexually active non-vaccinated 15–17-year-old girls from Argentina.

Sample collection city	n (%)	HPV Positive		HPV16/18		HPV6/11/16/18	
		n (%)	95% CI (%)	n (%)	95% CI (%)	n (%)	95% CI (%)
Berazategui, Province of Buenos Aires	272 (28.4)	117 (43.0)	37.1–48.9	25 (9.2)	5.8–12.6	43 (15.8)	11.5–20.2
Buenos Aires city	298 (31.1)	180 (60.4)	54.8–65.9	47 (15.8)	11.6–19.9	70 (23.5)	18.7–28.3
Posadas, Province of Misiones	172 (18.0)	106 (61.6)	54.3–68.9	36 (20.9)	14.8–27.0	47 (27.3)	20.7–34.0
La Banda, Province of Santiago del Estero	215 (22.5)	136 (63.3)	56.8–69.7	37 (17.2)	12.2–22.3	55 (25.6)	19.7–31.4

The data is relevant considering that Argentina started the national immunization program using the bivalent vaccine; local studies on the prevalence and risk of non-vaccine HPV types are important to better assess the effectiveness of vaccination.

In almost 30% of the samples at least one LR-HPV was identified, in line with data from Denmark that show 25% prevalence of these viruses among women aged 15–19 y²⁸. In this study, the prevalence of HPV6/11, the components of the quadrivalent and nonavalent vaccines is 9.5%; a similar amount was previously detected in Sweden (10.1%)³⁷,

and a higher percentage in Denmark (16.1%)²⁸; in the three cohorts, the rate of HPV11 prevalence was about one half of that obtained for HPV6, confirming the predominance of the latter.

Notably, HPV42 was the most common type in this group of adolescents (11.5%). In Sweden, HPV42 (6.4%) was the most frequent type among the LR-HPVs; however, the total type-specific prevalence ranking was led by HPV16³⁷. In several studies on cervical samples from healthy women in Denmark and the Netherlands, the prevalence of HPV42 was lower than that of HPV6 and roughly equal to that

Table 3 Single and multiple human papillomavirus infections among sexually active non-vaccinated 15–17-year-old girls from Argentina.

Single and Multiple infections	Adolescents Tested (n = 957)	
	n (%)	95% CI (%)
1 type	191 (20.0)	(17.4–22.5)
2 types	137 (14.3)	(12.1–16.5)
3 types	97 (10.1)	(8.2–12.0)
4 types	49 (5.1)	(3.7–6.5)
5 or more types	65 (6.8)	(5.2–8.4)

of HPV11^{28,32}. However, reports from Southern Europe and Africa show that the prevalence of HPV42 exceeds that of HPV6 and HPV11^{1,27}. At first, the high prevalence of HPV42 raised the question of whether it would be advantageous to include HPV42 in a vaccine. Nevertheless, a study on HPV types in external genital warts in a group of Argentine women showed that HPV6 and/or 11 were present in 93.3% of the cases, indicating that most of the lesions could be prevented by the quadrivalent HPV vaccine¹⁸.

The results of the present study also show that infection with multiple HPV types is frequent in adolescents (36.3%), in accordance with previously published data from Sweden (38%) in a cohort of 15–23-year-old non-vaccinated women³⁷ and Colombia (29%) in 12–19-year-old girls¹²; while in African girls, the value rises more than 50% for sexually active young women (<25 years of age)⁴⁶. Multiple cervical HPV infections have been extensively associated with young age, which may be due to a lack of natural immunity to HPV during the first years of sexual activity, but also to the number and characteristics of sexual partners³⁹.

The choice of HPV genotyping method is crucial, particularly due to the extensive variation in available tests; they mainly differ in the viral types that they are able to identify, the limit of detection (LOD) for each genotype, and their cost. The test should exhibit high analytical sensitivity to detect a broad spectrum of HPV types and to have a more accurate picture of their relative prevalence. The BSGP 5+/6+ PCR-RLB method applied in this study was developed and validated for the sensitive detection of 36 HPV types in single as well as multiple infections; it provides a powerful, highly flexible and sensitive tool for a more homogeneous amplification and HPV genotyping, at a considerably lower cost than commercial tests with a similar methodological basis⁴¹. This assay allowed to identify all HR-types (IARC groups 1, 2A and 2B), with the only exception of HPV30, 67, 69, 85 and 97, showing a remarkable ability for viral identification. It is important to emphasize that our results are supported by the correct resolution of the Proficiency Panel distributed on an annual basis by the Global HPV LabNet¹⁶; it guarantees that at least 5 copies of vaccine viruses are detected (in single and multiple infections) and that all negative controls are correct. This adds strength and confidence to these data.

Although this study was not population-based, data are particularly relevant because they provide information about HPV frequency in a group rarely assessable or included

in worldwide epidemiological surveys, mainly due to recruitment difficulties, which adds value to our work.

In conclusion, this is the first report about cervical HPV genotype distribution that involves sexually active 15–17-year-old non-vaccinated girls from Argentina, and one of the few existing in the world focusing exclusively on this group.

It may be used to inform on vaccination policies, as a baseline against which to measure the impact of the national HPV immunization program on the prevalence of vaccine-type and non-vaccine-type HPV infections. We are conducting the same study in vaccinated girls; assessments of the overall reduction in vaccine type infections, changes in related alpha types, and differential reductions based on the number of doses received would be the main points to evaluate and thus help to sustain and improve prevention strategies.

Funding

This study was financially supported by Salud Investiga (Carrillo-Oñativia fellowships), Dirección Nacional de Control de Enfermedades Inmunoprevenibles (Grant 2014–2015) and Instituto Nacional de Enfermedades Infecciosas - ANLIS Malbrán.

Conflict of interest

The authors declare that they have no conflicts of interest

Acknowledgments

The authors sincerely thank all hospitals and institutions involved in the study from the following cities: Berazategui (Province of Buenos Aires), Posadas (Province of Misiones), La Banda (Province of Santiago del Estero) and Buenos Aires, and the girls who participated in this study for their generosity.

The authors are particularly grateful to the members of the Health Ministry of Santiago del Estero Province: Minister, Dr. Luis Martínez; Undersecretary, Dr. Cesar Monti; Executive Director CISB, Dra. Liliana Garnica; Director of Maternity, Childhood and Adolescence, Dr. Pedro Carrizo; Director of Immunizations, Dra. Florencia Coronel; Coordinator of the Cervical Cancer Program, Dra. Yolanda Martínez; APS Director, Dra. Teresa Santillán and *Unidades de ProntaAtención* (UPAs); USM Director, José Alzogaray; *Departamento de Docencia e Investigación*, Dr. Carlos Marrodán, as well as to Dr. Walter Villalba (Director of the *Hospital Escuela de Agudos Ramón Madariaga*, Posadas, Misiones), and Delia Dominga Motta (nurse), for their support and commitment in patient recruitment.

We are also indebted to Dr Enrique Lamuedra and Marcela Grosso (*Centro Nacional de Red de Laboratorios*, ANLIS - Malbrán), for their assistance in sample transfer.

Appendix A. MALBRAN HPV Surveillance Study Group

Alejandra Giurgiović (Hospital Evita Pueblo, Berazategui); Gabriela Alzogaray, Ricardo Aboslaiman (Centro Integral

de Salud La Banda, Santiago del Estero); Cecilia Chami (Ministerio de Salud de Santiago del Estero); Juan José Carmona, Néstor Tappari, Andrea Morgenstern (Hospital Madariaga, Misiones); María Elina Totaro (Universidad Nacional de Misiones); Enrique Berner, Viviana Cramer, Sandra Vázquez, Paula Real (Hospital Argerich, Buenos Aires), Carlota Lopez Kaufman, Gabriela Kosoy, Lucía Katabian (Hospital Rivadavia, Buenos Aires); María Silvia Severino (Hospital Durand, Buenos Aires).

References

1. Ammatuna P, Giovannelli L, Matranga D, Ciriminna S, Perino A. Prevalence of genital *Human papillomavirus* infection and genotypes among young women in Sicily, South Italy. *Cancer Epidemiol Biomark Prev.* 2008;17:2002–6.
2. Arbyn M, Depuydt TM, Dillner CJ. Are 20 *Human papillomavirus* types causing cervical cancer? *J Pathol.* 2014;234:431–5.
3. Arrossi S. The impact of the HPV test in screening programs in Latin America: the case of Argentina. *Salud Publica Mex.* 2019;61:86–94.
4. Baudu A, Prétet J, Riethmuller D, Chotard M, Mougin C, Mercier M. Prevalence and risk factors of *Human papillomavirus* infection types 16/18/45 in a cohort of French females aged 15–23 years. *J Epidemiol Glob Health.* 2014;4:35–43.
5. Bosch FX, Robles C, Díaz M, Arbyn M, Baussano I, Clavel C, Ronco G, Dillner J, Lehtinen M, Petry KU, Poljak M, Kjaer SK, Meijer CJ, Garland SM, Salmerón J, Castellsagué X, Bruni L, de Sanjósé S, Cuzick J. HPV-FASTER: broadening the scope for prevention of HPV-related cancer. *Nat Rev Clin Oncol.* 2016;13:119–32.
6. Brotherton JML. Impact of HPV vaccination: achievements and future challenges. *Papillomavirus Res.* 2019;7:138–40.
7. Brotherton JM, Kaldor J, Garland SM. Monitoring the control of *Human papillomavirus* (HPV) infection and related diseases in Australia: towards a national HPV surveillance strategy. *Sex Health.* 2010;7:310–9.
8. Brown DR, Kjaer S, Sigurdsson K, Iversen OE, Hernandez-Avila M, Wheeler CM, Perez G, Koutsky LA, Tay EH, Garcia P, Ault KA, Garland SM, Leodolter S, Olsson SE, Tang GW, Ferris DG, Paavonen J, Steben M, Bosch FX, Dillner J, Joura EA, Kurman RJ, Majewski S, Muñoz N, Myers ER, Villa LL, Taddeo FJ, Roberts C, Tadesse A, Bryan J, Lupinacci LC, Giacoletti KE, Sings HL, James M, Hesley TM, Barr E. The impact of quadrivalent *Human papillomavirus* (HPV; types 6, 11, 16, and 18) L1 virus-like particle vaccine on infection and disease due to oncogenic nonvaccine HPV types in generally HPV-naive women aged 16–26 years. *J Infect Dis.* 2009;199:926–35.
9. Chouhy D, Mamprín D'Andrea R, Iglesias M, Messina A, Ivancovich JJ, Cerdá B, Galimberti D, Bottai H, Giri AA. Prevalence of *Human papillomavirus* infection in Argentinean women attending two different hospitals prior to the implementation of the National vaccination program. *J Med Virol.* 2013;85:655–66.
10. Clifford G, Franceschi S, Diaz M, Muñoz N, Villa LL. Chapter 3: HPV type-distribution in women with and without cervical neoplastic diseases. *Vaccine.* 2006;24 Suppl. 3. S3/26–34.
11. de Martel C, Plummer M, Vignat J, Franceschi S. Worldwide burden of cancer attributable to HPV by site, country and HPV type. *Int J Cancer.* 2017;141:664–70.
12. Del Río-Ospina L, Soto-De León SC, Camargo M, Sánchez R, Mancilla CL, Patarroyo ME, Patarroyo MA. The prevalence of high-risk HPV types and factors determining infection in female Colombian adolescents. *PLOS ONE.* 2016;11:11.
13. Deluca GD, Marin MH, Blanco NS, Basiletti JA, González JV, Merino AL, Picconi MA. Distribution of *Human papillomavirus* genotypes in women with cervical alterations from north Argentina. *Indian J Med Microbiol.* 2013;31:138–41.
14. Dirección de Control de Enfermedades Inmunoprevenibles, Argentina. Available online at: <https://www.argentina.gob.ar/salud/inmunoprevenibles/manuales-lineamientos> [accessed October 2019].
15. Dunne EF, Klein N, Naleway AL, Baxter R, Weinmann S, Riedlinger K, Fetterman B, Steinau M, Scarbrough MZ, Gee J, Markowitz LE, Unger ER. Prevalence of HPV types in cervical specimens from an integrated healthcare delivery system: baseline assessment to measure HPV vaccine impact. *Cancer Causes Control.* 2013;24:403–7.
16. Eklund C, Forslund O, Wallin KL, Dillner J. Continuing global improvement in *Human papillomavirus* DNA genotyping services: the 2013 and 2014 HPV LabNet international proficiency studies. *J Clin Virol.* 2018;101:74–85.
17. Ermel AC, Shew ML, Weaver BA, Qadri B, Denski C, Tu W, Tong Y, Fortenberry JD, Brown DR. DNA detection and seroprevalence of *Human papillomavirus* in a cohort of adolescent women. *Sex Transm Infect.* 2014;90:64–9.
18. Fleider LA, Tatti S, Suzuki AV, Maldonado VA, Díaz LB, Chiesa IJ, Perez MS. *Human papillomavirus* types involved in external genital warts in a group of Argentinian Women in Buenos Aires. *J Low Genit Tract Dis.* 2016;20:365–6.
19. Global HPV Reference Center. Available online at: <http://www.hpvcenter.se/html/refclones> [accessed September 2019].
20. Hildesheim A, Herrero R, Wacholder S, Rodriguez AC, Solomon D, Bratti MC, Schiller JT, Gonzalez P, Dubin G, Porras C, Jimenez SE, Lowy DR. Costa Rican HPV Vaccine Trial Group Effect of *Human papillomavirus* 16/18 L1 virus like particle vaccine among young women with preexisting infection: a randomized trial. *JAMA.* 2007;298:743–53.
21. Houlihan CF, Baisley K, Bravo IG, Kapiga S, de Sanjósé S, Changalucha J, Ross DA, Hayes RJ, Watson-Jones D. Rapid acquisition of HPV around the time of sexual debut in adolescent girls in Tanzania. *Int J Epidemiol.* 2016;45:762–73.
22. Howell-Jones R, Bailey A, Beddows S, Sargent A, de Silva N, Wilson G, Anton J, Nichols T, Soldan K, Kitchener H, Study Group Collaborators. Multi-site study of HPV type-specific prevalence in women with cervical cancer, intraepithelial neoplasia and normal cytology, in England. *Br J Cancer.* 2010;103:209–16.
23. ICO-IARC Information Centre on HPV and Cancer. *Human papillomavirus* and related diseases report, Argentina; 2018. Available online at: www.hpvcentre.net [accessed October 2019].
24. Insinga RP, Perez P, Wheeler CM, Koutsky LA, Garland SM, Leodolter S, Joura EA, Ferris DG, Steben M, Hernandez-Avila M, Brown DR, Elbasha E, Muñoz N, Paavonen J, Haupt RM. FUTURE I Investigators Incident cervical HPV infections in young women: transition probabilities for CIN and infection clearance. *Cancer Epidemiol Biomark Prev.* 2011;20:287–96.
25. International Agency for Research on Cancer. IARC monographs on the evaluation of carcinogenic risks to humans-human papillomaviruses, vol. 100B. Geneva: World Health Organization; 2012. p. 255–313.
26. Kavanagh K, Pollock K, Cuschieri K, Palmer T, Cameron RL, Watt C, Bhatia R, Moore C, Cubie H, Cruickshank M, Robertson C. Changes in the prevalence of human papillomavirus following a national bivalent *Human papillomavirus* vaccination programme in Scotland: a 7-year cross-sectional study. *Lancet Infect Dis.* 2017;17:1293–302.
27. Keita N, Clifford G, Koulibaly M, Douno K, Kabba I, Haba M, Sylla BS, van Kemenade FJ, Snijders PJ, Meijer CJ, Franceschi S. HPV infection in women with and without cervical cancer in Conakry, Guinea. *Br J Cancer.* 2009;101:202–8.
28. Kjaer SK, Breugelmans G, Munk C, Junge J, Watson M, Iftner T. Population-based prevalence, type- and age-specific distribution of HPV in women before introduction of an HPV-vaccination program in Denmark. *Int J Cancer.* 2008;123:1864–70.

29. Koutsky L. Epidemiology of genital *Human papillomavirus* infection. *Am J Med.* 1997;102:3–8.
30. Lazcano-Ponce E, Herrero R, Muñoz N, Cruz A, Shah KV, Alonso P, Hernández P, Salmerón J, Hernández M. Epidemiology of HPV infection among Mexican women with normal cervical cytology. *Int J Cancer.* 2001;91:412–20.
31. Lehtinen M, Paavonen J, Wheeler CM, Jaisamrarn U, Garland SM, Castellsagué X, Skinner SR, Apter D, Naud P, Salmerón J, Chow SN, Kitchener H, Teixeira JC, Hedrick J, Limson G, Szarewski A, Romanowski B, Aoki FY, Schwarz TF, Poppe WA, De Carvalho NS, German MJ, Peters K, Mindel A, De Sutter P, Bosch FX, David MP, Descamps D, Struyf F, Dubin G, HPV PATRICIA Study Group. Overall efficacy of HPV-16/18 AS04-adjuvanted vaccine against grade 3 or greater cervical intraepithelial neoplasia: 4-year end-of-study analysis of the randomised, double-blind PATRICIA trial. *Lancet Oncol.* 2012;13:89–99.
32. Lenselink CH, Melchers W, Quint WG, Hoebers AM, Hendriks JC, Massuger LF, Bekkers RL. Sexual behaviour and HPV infections in 18 to 29 year old women in the pre-vaccine era in the Netherlands. *PLoS One.* 2008;13:e0206219. <https://journals.plos.org/> [online].
33. Mendoza LP, Paez M, Kasamatsu E, Castro A, Giménez G, Basiletti J, Gonzalez J, Mongelós P, Picconi MA. Distribution of *Human papillomavirus* genotypes in Paraguayan women according to the severity of the cervical lesion. *J Med Virol.* 2011;83:351–7.
34. Munoz N, Castellsague X, de Gonzalez AB, Gissmann L. Chapter 1: HPV in the etiology of human cancer. *Vaccine.* 2006;24 Suppl. 3. S3/1–10.
35. Puerto D, Reyes V, Lozano C, Buitrago L, Garcia D, Murillo RH, Muñoz N, Hernandez GA, Sanchez L, Wiesner C, Combita AL. Detection and genotyping of HPV DNA in a group of unvaccinated young women from Colombia: Baseline Measures Prior to Future Monitoring Program. *Cancer Prev Res (Phila).* 2018;11: 581–92.
36. Ramas V, Mirazo S, Bonilla S, Mendoza L, Lago O, Basiletti J, González J, Picconi MA, Arbiza J. *Human papillomavirus* genotypes distribution in cervical samples from Uruguayan women. *J Med Virol.* 2013;85:845–51.
37. Ramqvist T, Du J, Lundén M, Ahrlund-Richter S, Ferreira J, Mariions L, Sparén P, Andersson S, Dalianis T. Pre-vaccination prevalence of *Human papillomavirus* types in the genital tract of 15–23-year-old women attending a youth health clinic in Stockholm, Sweden. *Scand J Infect Dis.* 2010;43:115–21.
38. Revzina NV, Diclemente R. Prevalence and incidence of *Human papillomavirus* infection in women in the USA: a systematic review. *Int J STD AIDS.* 2005;16:528–37.
39. Sargent A, Bailey A, Almonte M, Turner A, Thomson C, Peto J, Desai M, Mather J, Moss S, Roberts C, Kitchener HC, ARTISTIC Study Group. Prevalence of type-specific HPV infection by age and grade of cervical cytology: data from the ARTISTIC trial. *Br J Cancer.* 2008;98:1704–9.
40. Sasieni P, Castanon A, Cuzick J. Effectiveness of cervical screening with age: population based case-control study of prospectively recorded data. *BMJ.* 2009;339:b2968. www.bmjjournals.com [online].
41. 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;46:1050–9.
42. Sociedad Argentina de Ginecología Infanto Juvenil. Available online at: <http://www.conders.org.ar/pdf/sagij.pdf> [accessed May 2019].
43. Tabrizi SN, Brotherton J, Stevens MP, Condon JR, McIntyre P, Smith D, Garland SM. WHINURS group HPV genotype prevalence in Australian women undergoing routine cervical screening by cytology status prior to implementation of an HPV vaccination program. *J Clin Virol.* 2014;60:250–6.
44. van den Brule AJ, Pol R, Fransen-Daalmeijer N, Schouls LM, Meijer CJ, Snijders PJ. GP5+/6+PCR followed by reverse line blot analysis enables rapid and high-throughput identification of human papillomavirus genotypes. *J Clin Microbiol.* 2002;40:779–87.
45. Walboomers JM, Jacobs M, Manos MM, Bosch FX, Kummer JA, Shah KV, Snijders PJ, Peto J, Meijer CJLM, Muñoz N. *Human papillomavirus* is a necessary cause of invasive cervical cancer worldwide. *J Pathol.* 1999;189:12–9.
46. Watson-Jones D, Baisley K, Brown J, Kavishe B, Andreasen A, Changalucha J, Mayaud P, Kapiga S, Gumodoka B, Hayes RJ, de Sanjosé S. High prevalence and incidence of *Human papillomavirus* in a cohort of healthy young African female subjects. *Sex Transm Infect.* 2013;89:358–65.
47. WHO position paper. Human papillomavirus vaccines. *Wkly Epidemiol Rec.* 2017;92:241–68.
48. Winer RL, Feng Q, Hughes JP, O'Reilly S, Kiviat NB, Koutsy LA. Risk of female *Human papillomavirus* acquisition associated with first male sex partner. *J Infect Dis.* 2008;197:279–82.
49. World Health Organization. Summary of HPV surveillance and monitoring meeting. 20 July 2009. Available online at: http://www.who.int/nutri/hpv/HPV_Monitoring_Meeting_Report_May_2009.pdf [accessed May 2019].
50. Zur Hausen H. Papillomaviruses causing cancer: evasion from host-cell control in early events in carcinogenesis. *J Natl Cancer Inst.* 2000;92:690–8.