

Human papillomavirus prevalence in oral and oropharyngeal squamous cell carcinoma in South America: A systematic review and meta-analysis

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Abstract

Some studies have addressed the prevalence of human papillomavirus (HPV) in head and neck cancer in South America; however, no studies have systematically gathered prevalence and con-

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©Copyright: the Author(s), 2022 Licensee PAGEPress, Italy Oncology Reviews 2022; 16:552 doi:10.4081/oncol.2022.552 ducted a meta-analysis. This study aims to estimate the prevalence of HPV in oral and oropharyngeal squamous cell carcinomas in South America. We performed a systematic review and metaanalysis using the following databases: PubMed, Embase, Lilacs, Medline, Scopus, and Web of Science. Data were extracted and analyzed using random-effects models to estimate the pooled prevalence of HPV. We identified 209 nonduplicated studies, of which 38 were selected. The overall prevalence of HPV was 24.31% (95% CI 16.87-32.64; I² = 96%, pheterogeneity <0.001). HPV prevalence in oropharyngeal cancer was 17.9% (95% CI 7.6-31.4; $I^2 = 96\%$, $p_{\text{heterogeneity}} < 0.001$) and that in oral cavity cancer was 23.19% (95% CI 14.94–32.63; $I^2 = 94\%$, $p_{heterogeneity} < 0.001$). We found an overall prevalence of HPV in 24.31% of oral and oropharyngeal squamous cell carcinomas in South American patients. The prevalence of HPV was 17.9% for oropharyngeal cancer and 23.19% for oral cavity cancer.

Introduction

In 2020, the International Agency for Research on Cancer (IARC) registered 476,125 new cases of oropharyngeal and oral cavity cancer worldwide, of which 98,412 belonged to oropharynx and 377,713 belonged to the oral cavity.¹ In South America, with approximately 7292 new oropharyngeal cancer cases and 14,191 new oral cavity cancer cases, the combined incidence was estimated as 1.97% of all new cases recorded in 2020.¹

According to the National Cancer Institute (INCA), cancers of the oral cavity and oropharynx had an incidence in Brazil of 10.69 per 100,000 inhabitants in 2020 and represented the fifth highest incidence among men.²

Squamous cell carcinoma (SCC) is the most common histological type of oral cancer,^{3,4} and even with advances achieved in treatment regimes, mortality remains high after 5 years.⁵ Despite recognizing a multifactorial process, the risk of developing head and neck cancer (HNC) is 30 times greater for individuals who smoke and drink.⁶ Several studies have reported human papillomavirus (HPV) as a relevant risk factor for HNC development, particularly for oropharynx.^{7–9}

More than 170 types of HPV are known, and they are divided into two large groups: high-risk (hrHPV) and low-risk (lrHPV). The most common genotypes in head and neck cancers are HPV16 and HPV18.^{10,11} A meta-analysis published in 2014, with data from 44 countries, recorded a pooled prevalence of 29.5% for HPV DNA in HNCSCC; in the oral cavity and oropharynx, the pooled prevalences were 24.2% and 45.8%, respectively.¹¹ Moreover, HPV16 DNA could contribute to 82.2% of all cases of HPV-positive DNA.¹¹ A recent meta-analysis of Brazilian studies estimated an overall prevalence of 11.89% for different types of HPV in the oral cavity and 4.69% for hrHPVs.¹²

Methods for detecting and diagnosing HPV can target viral DNA or RNA, oncoproteins, cellular proteins, and serum antibodies specific to HPV.¹³ The detection of messenger RNA E6/E7 (mRNA) is considered as gold standard for identifying HPV.¹³ Among DNA detection techniques, the most cited are direct hybridization, such as Southern blotting, *in situ* hybridization (ISH), and polymerase chain reaction (PCR);^{13–15} whereas, among the biomarker detecting methods, the most common is immunohistochemical (IHC) detection of overexpression of p16^{INK4a}.^{13–15}

Furthermore, the literature results vary for the association between HPV and oral cavity and oropharyngeal cancers. Few studies have verified the prevalence of HPV in HNCC in South American countries. Considering the need for studies that collate the data recorded in this region, as well as the variation of data from Brazil and other South American countries concerning the rest of the world, the present study aimed to conduct a systematic review and meta-analysis to estimate the prevalence of HPV in oral and oropharyngeal squamous cell carcinoma in South America.

Materials and methods

We performed a systematic review and meta-analysis, where three authors carried out the initial searches on the following databases: Embase, Lilacs, Medline, Scopus, and Web of Science. We searched for articles in the following languages: English, Portuguese, and Spanish.

The following terms were used to search in the titles, abstracts, and keywords: ("HPV" OR "papillomavirus" OR "papillomaviridae") AND (("head" AND "neck") OR "oropharynx" OR "oropharyngeal" OR "tongue" OR "mouth" OR "oral" OR "oral cavity") AND ("cancer" OR "tumor" OR "neoplasms" OR ("carcinoma" AND "squamous" AND "cell")) AND ("Brazil" OR "Brazilian" OR "Argentina" OR "Bolivia" OR "Chile" OR "Colombia" OR "Ecuador" OR "Guyana" OR "Paraguay" OR "Peru" OR "Suriname" OR "Uruguay" OR "Venezuela").

Initially, the search in each database resulted in 494 records. After the first screening and exclusion of duplications, 209 studies were selected to read the titles and abstracts. Of these, 52 studies were selected from a review of titles and abstracts. After analyzing these 52 articles' references, another 2 were selected for detailed assessment. To control the data collection process, each of the 54 studies selected for detailed assessment received a sequential code ranging from A01 to A54. After reading the full texts, 38 articles were selected for qualitative and quantitative reviews.

The search and selection stages of the studies are presented in Appendix A and are illustrated in Figure 1.

Exclusion criteria

The following exclusion criteria were established: articles not written in English, Portuguese, or Spanish; articles in which the correlation between HPV and oral cavity squamous cell carcinoma (OSCC) or oropharynx squamous cell carcinoma (OPSCC) were not found; review articles with or without meta-analysis or case reports; animal studies; search results with incomplete information or those without access to full text; studies with incomplete or



inaccurate information that would make it impossible to extract the necessary information reliably; duplicate studies; studies that revealed strong evidence of sample overlap; and studies whose samples had less than five cases of OSCC or OPSCC per country.

Data extraction

In articles where the lesion subsites were not classified into the oropharynx and oral cavity, the following grouping was applied, according to the ICD-O classification:¹⁶ oropharynx (C01.9, 02.4, 09.0–10.9), oral cavity (C00.0–00.9, 02.0–02.3, 02.8–03.1, 03.9–04.1, 04.8–05.0, 05.8–06.2, 06.8–06.9).

Data collection was performed only for viral DNA detection techniques in studies where HPV diagnostic technique involved more than one detection method and the criterion for diagnosis did not include combination of results.

Data analysis

The collected data were tabulated using STATA IC version 16.¹⁷ The combined prevalence was estimated using the R metaprop package, version 3.6.1,¹⁸ via the Rstudio interface, version 1.2.1335.¹⁹ The following parameters were used for statistical analysis: inverse variance method, DerSimonian-Laird estimator for the variance between studies, Jackson's method for calculating confidence intervals, arcsine transformation for gross proportions,



Figure 1. Flow chart of study selection.



and Clopper–Pearson CI to estimate the confidence intervals (95% CI) for each of the studies. Heterogeneity tests (I^2) between studies were performed, and Cochran's Q test was used to establish the significance of heterogeneity. These parameters refer to the random-effects analysis, considering the heterogeneity of our sample (p<0.01).

Heterogeneity was analyzed using meta-regression and stratification. The explanatory variables selected for the meta-regression analysis included the country of origin of the studies, types of HPV investigated, type of material collected, and method for diagnosing HPV. Stratification included all subgroups of the metaregression and the patient's sex and profile (smokers and alcohol consumers). For the type of material collected, the analysis included in this document involves two groups: material fixed by freezing or inclusion in paraffin, and exfoliation off mucosa or swab. As for the diagnostic method, four subgroups were evaluated: HPV DNA detection techniques, HPV and IHC DNA alone, HPV and IHC DNA combined, and IHC. Meta-regression analysis was performed using the metareg package ¹⁸ via moderator test. Subgroup comparisons were calculated using the Q test for differences in subgroups of the metaprop package.¹⁸

The R codes, dataset, and the complete list of nonduplicate records with the respective reasons for exclusion can be found in Mendeley Data doi:10.17632/9ss9d42f74.1.

Results

We identified 209 unduplicated records. After reading the titles and abstracts, 52 studies were included. The references of these 52 articles were revised, and another two studies were added, adding up to 54 full texts selected for reading. After reading, 38 articles were selected for the qualitative and quantitative reviews. The selected studies were published between 1998 and August 2020 and included patients assessed between 1970 and 2019. Brazil has the largest number of studies (n=28),²⁰⁻⁴⁷ followed by Colombia (n=4),⁴⁸⁻⁵¹ Argentina (n=2),^{52,53} Chile (n=2),^{54,55} and Venezuela (n=2)^{56,57} presented only two studies each.

The patients' sample resulting from the 38 studies included 2788 individuals–1643 cases of OSCC and 1145 of OPSCC. Brazil had highest number of cases (2329 cases), followed by Colombia (190 cases), Chile (102 cases), and Argentina (101 cases), and Venezuela had the lowest number (66 cases) (Table 1).

The pooled prevalence of HPV for OSCCs $(33 \text{ studies})^{20-23,25-27,30-40,42-57}$ and OPSCC (14 studies)^{20-22,24,25,28-30,34,37,38,40,41,51} was estimated at 24.31% (95% CI 16.87–32.64) (Figure 2). In Brazil, this prevalence was estimated to be 21.15% (95% CI, 12.83–30.90). The highest prevalence was observed in Venezuela (57.6% [95% CI 45.57–69.18]), Argentina (42.57% [95% CI 33.13–52.30]), and Colombia (30.27% [95% CI 16.14–46.63]). Chile had the lowest prevalence (10.77% [95% CI 5.53–17.48]) (Table 2). Nevertheless, notably, with the exception of Brazil, all other countries presented few studies for analysis and a reduced number of cases (Table 2).

The pooled prevalence for HPV16 in OSCC and OPSCC was 12.8% (95% CI 8.7–17.5) (Figure 3) (23 studies).^{20-25,27,30,34–40,46,48–50,52,53,55,56} In Brazil (16 studies),^{20-25,27,30,34–40,46} this prevalence was estimated at 9.6% (95% CI 6.1-13.9). Chile⁵⁵ (6.2% [95% CI 2.0–12.6]) and Venezuela⁵⁵ (44% [95% CI 30.5–57.8]) presented only one study each for this criterion. Argentina considered two studies^{52,53} (9.9% [95% CI 4.8–16.4]) and Colombia presented three studies⁴⁸⁻⁵⁰ (38.6% [95% CI 13.8–67.2]). In the metaregression analysis, studies from Colombia and Venezuela, using the data obtained from Argentinian studies as a reference, revealed

statistical significance for heterogeneity moderation in the pooled prevalence of HPV16 in OPSCCs and OSCCs, thereby contributing positively to the calculated percentage (p<0.05). Except for countries' stratification in the set of studies that investigated HPV16 infection, none of the other stratifications revealed significant results for moderating heterogeneity (p>0.05). This information is presented in Table 2.

The pooled prevalence of HPV in OSCCs was estimated at 23.19% (95% CI 14.94–32.63) (Figure 5) (33 studies).^{20,22,23,25-27,30–40,42–57} In Brazil (23 studies),^{20,22,23,25-27,30–40,42–47} this prevalence was 18.71% (95% CI 9.51–30.15) (Table 2). Argentina^{52,53} (42.57% [95% CI 33.13–52.30]), Chile^{54,55} (10.77% [95% CI 5.53–17.48]), and Venezuela^{56,57} (57.60% [45.57–69.18]) presented only two studies each. Colombia^{48–51} (31.80% [95% CI 17.61–48.97]) maintained the four initial studies (Table 2).

The subgroup analysis by type of material collected and HPV investigated did not reveal any statistical significance between the pooled prevalence (p>0.05). About 22.64% of the tissue samples fixed by frozen or paraffin embedding techniques (35 studies)²⁰⁻ 41,43-46,48-52,54-57 presented a positive diagnosis for some type of HPV (95% CI 15.13–31.17). In the samples of mucosa scraping or swab (3 studies).^{42,47,53} this percentage was 44.22% (95% CI 24.15-65.32). In studies that investigated only hrHPV (9 studies),^{20,24,25,27-30,35,50} the calculated prevalence was 16.45% (95% CI 05.03-32.75), those that investigated infection for both hrHPV and lrHPV (29 studies),^{21-23,26,31-34,36-49,51-57} the pooled prevalence was 27.14% (18.31-36.99). In studies where data collection considered only the results of molecular tests to detect HPV DNA (34 studies),21-23,25-27,30-57 the combined prevalence of HPV DNA was 24.67% (95% CI 17.13-33.10). Moreover, only two studies applied the combined methodology between PCR and IHC to diagnose HPV, considering only positive cases in which the p16 biomarker and viral DNA were present.^{20,24} The pooled prevalence of HPV was 4.15% (95% CI 1.37-8.34). In study A16,²⁹ detection of HPV involved viral DNA molecular detection techniques and IHC to detect the p16 biomarker; however, it was not specified which technique tested cases, and the pooled HPV prevalence in that study, in the random-effects model, was estimated at 59.07% (95% CI 52.18-65.71). Furthermore, Study A22²⁸ was the only one included in this review in which HPV was diagnosed only via IHC; the pooled prevalence of the cases in this study, in the randomeffects model, was 35.29% (95% CI 22.89-48.80). This information is detailed in Table 2.

Eventually, Table 2 details the stratification by sex and profile of the cases analyzed (smokers and alcoholics). In studies where cases were stratified by sex, the pooled prevalence of HPV in OPSCC and OSCC for men (19 studies)^{20,21,23,24,26,27,29-31,39-41,44,46,49,50,54,55,57} was 19.11% (95% CI 9.37–31.29), and for women (19 studies),^{<math>20,21,23,24,27,29-31,39,40,41,44,46,49,50,54-57} it was 22.35% (95% CI 11.85–35.04). Among smokers considering 947 cases (13 studies),^{<math>20,21,23,24,27-29,38,39,41,44,49,56} the pooled prevalence was 20.75% (95% CI 9.21–35.44) and for alcoholics, with 720 cases (10 studies),^{<math>20,21,23,24,27,29,41,44,49,56} the pooled prevalence of HPV was 22.23% (95% CI 8.23–40.62).</sup></sup></sup>

Discussion

This review included 38 studies that examined the prevalence of HPV DNA and p16^{INK4a} positivity, alone or in combination, in OPSCC and OSCC in patients from South American countries.

Based on 2788 cases, 1145 OPSCC and 1643 OSCC, we found that approximately 25% of the cases were associated with HPV infection. The highest pooled prevalence was observed in

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Table

Study by country	Code	Period	Collected	Diagnostic method	z	HPV	HPV16+ -		OSCC		DPSCC
			material	6		+		+V4H	-VAH	HPV+	HPV-
Argentina											
Gonzalez et al, 2007 ⁵²	A48	2007 ^a	FFPE	PCR, Dot blot	16	L	2	L	6		i
Mosmann et al, 2019 53	A38	2019 ^a	Swab	PCR	85	36	8	36	49	1	1
Brazil											
Kaminagakura et al, 2012 ³⁵	A37	1970-2006	FFPE	IHC, PCR, Dot blot	114	22°	22°	22°	92°	:	
De Cicco et al, 2018 ²⁹	A16	1984-2014	FFPE	IHC, PCR, Reverse blot	215	127				127	88
Betiol et al, 2016 ²²	A03	1991-2012	FFPE	IHC, PCR	120	27°	19°	23°	e9°	4°	24°
Demathe <i>et al</i> , 2010 ³²	A42	1991-2007	FFPE	nPCR	30	13	1	13	17	;	:
Simonato et al 2008 44	A45	1991-2005	FFPE	PCR	29	v		v	24		
Anantharaman et al. 2017 ²⁰	A01	2002-2015	FFPF	THC & DCR	278	L	7	0	107	7	164
$\Gamma_{raconari}$ of al 20133	V S N	1007 2005	EEDE			. v		o v	17		-
FIGUIDEL <i>et at</i> , 2012	DCA	CUU2-2661	TTT	LCN D	11	, i	: •	, i	71	÷	:
Oliveira et al, 2008 39	A17	1990-2002	FFPE	PCR	87	17	ŝ	17	70	1	:
Buexm et al, 2020 ²⁴	A05	1999-2010	FFPE	IHC & qPCR	346	21	21	;		21	325
López <i>et al</i> , 2014 ³⁷	A20	1998-2009	FFPE/Frozen	PCR	212	14	8	8	113	9	85
de Carvalho et al, 2020 ²⁸	A22	2009-2017	FFPE	IHC	51	18	1	;	:	18	33
Soares et al., 2002 ⁴⁵	A25	1992-1998	FFPE	ISI	10	0		0	10		
Hauck et al. 2015 ³⁴	A21	2007-2011	Frozen	THC. PCR	154	19°	110	6°	770	130	58°
de Abreil <i>et al</i> 2018 ²⁷	A07	2012-2015	FFPE/Frozen	PCR	06	"	"	"	87	1	1
Detito at al 2017 40	A 7 8	2005 2007	FEDF	DCB	62	10		01	06	E	37
$\mathbf{M}_{1} = \mathbf{M}_{1} $	07V	1002-0007	LI LI L	DCD DC4 PC6	70	17 17		- -	9 6	11	1 6
Nuguet et al, 1996 -	A14	0661-0661		FUK, DOI DIOL	17	1 1	4	- 1	70 ,	ŋ	C
Kibeiro <i>et al</i> , 2017^{42}	IcA	2013-2014	Cytobrush	PCK	9	0	1	0	-	I	l
Soares et al., 2007 ⁴⁶	A43	2005-2006	FFPE	PCR, Dot blot	75	18	1	18	57		1
Cortezzi et al, 2004 ²⁵	A52	2004 ^a	Frozen	PCR, Dot blot	32	4	4	1	10	ŝ	18
Vidal <i>et al.</i> , 2004 ⁴⁷	A29	2004 ^a	Cytobrush	Hybrid capture	40	П	ł	11	29	:	:
Rivero e Nunes, 2006 ⁴³	A31	2006 ^a	FFPE	ISH, PCR	40	0	i	0	40	÷	:
da Silva <i>et al</i> , 2007 ²⁶	A08	2007 ^a	Frozen	PCR	50	37	ł	37	13	÷	
de Spíndula-Filho <i>et al</i> , 2011 ³¹	A13	2010 ^a	FFPE	PCR	39	0		0	39	:	
Lira et al, 2010 ³⁶	A47	2010 ^a	FFPE	PCR	70	61	6	61	6	÷	ł
Brew et al, 2012 ²³	A04	2012 ^a	Frozen	PCR, Reverse blot	17	8	9	8	6	:	:
de Freitas Cordeiro-Silva <i>et al</i> 2012											
30	A23	2012 ^a	FFPE/Frozen	PCR	45	ŝ	ε	Э	38	0	4
Piña <i>et al</i> , 2016 ⁴¹	A35	2017 ^a	FFPE	IHC, ISH	13	0c	1	i	:	0°	13°
Barros-Filho et al, 2018 ²¹	A06	2018 ^a	Frozen	PCR	40	23	19	;	:	23	17
Chile											
Reyes et al., 2015 55	A34	2000-2014	FFPE	IHC, PCR	80	9c	5°	9°	71°	i	
Pennacchiotti et al, 2016 54	A44	2016 ^a	Frozen	PCR	22	2	l	2	20		
Colombia											
Quintero et al, 2013 ⁵¹	A32	1999-2008	FFPE	PCR, Reverse blot	112	22	:	16	51	9	39
Castillo et al, 2011 ⁴⁸	A49	1996-2001	FFPE	PCR	9	5	5	5	1	÷	
Mancilla <i>et al</i> , 2011 ⁵⁰	A19	2005-2007	FFPE	PCR	26	8	8	8	18		
Erira <i>et al</i> , 2015 ⁴⁹	A18	2013-2014	Fixed ^b	PCR	46	10	6	10	36	i	:
Venezuela											
Premoli-De-Percoco et al, 2001 56	A30	2001 ^a	FFPE	PCR, Southern blot	50	30	22	30	20	:	
Correnti et al, 2004 57	A39	2004 ^a	FFPE	PCR	16	8	1	8	8	:	1
aShidies did not mention the evect date of the recearch in the text of	so the date in the c	olumn "Period" refers t	o the date of publication of	the study bThe study does not snecify which	fivation techr	ione was adont	ed informing only th	hat the cenetic mate	orial was an All	Pren DNA/RNA/	ProteinOuiaden®
but the restriction of the restriction of the restriction of the restriction of the restriction 2788 . HPV + = HPV kit. Results for HPV DNA only. N = Total cases was 2788. HPV + = HPV	/ positive, the total	number of cases was 630	. HPV16+ = HPV genotype 1	6 positive, the total number of cases was 206	OSCC/HPV+ =	squamous cel	cut, intermined wing the or	ral the generic mun	-, the total num	ber of cases wa	s 388. OSCC/HPV-

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Table 2. Pooled prevalence of human papillomavirus in squamous cell carcinomas of the oral cavity (OSCC) and oropharynx (OPSCC) in South American countries by stratification variables.

						P-value	H	leterogeneity
	Studies	Cases	HFV positive cases	Pooled Prevalence (95% CI)	Test for subgroup differences	Test of Moderators	12	P-value
ODSCC and OSCC hy soundary					~0 0001	0 7644		
	¢	101	2		1000.0~	0.20	700	50100
Argentina	7	101	45	42.5/% (55.15-52.50)	:	rel.'	0%0	C/ 16.0
Brazil	28	2329	493	21.15% (12.83-30.90)	:	0.2689	96.4%	<0.0001
Chile	2	102	11	10.77% (5.53-17.48)	:	0.1816	0%0	0.7664
Colombia	4	190	45	30.27% (16.14-46.63)	:	0.7388	74.4%	0.0084
Venezuela	2	99	38	57.60% (45.57-69.18)	:	0.6747	%0	0.4833
OPSCC by country ^a				r.	0.5475	0.8270		
Brasil	13	1100	236	18.31% (7.33-32.82)	:	ref. ^b	96.2%	<0.0001
Colombia	1	45	9	13.33% (5.10-24.68)	i	0.8270	:	i
OSCC by country					<0.0001	0.2341		
Argentina	2	101	43	42.57% (33.13-52.30)	:	ref. ^b	%0	0.9175
Brazil	23	1229	257	18.71% (9.51-30.15)	i	0.2383	95.4%	<0.0001
Chile	2	102	11	10.77% (5.53-17.48)	:	0.2041	0%0	0.7664
Colombia	4	145	39	31.80% (17.61-47.97)	:	0.8028	69.9%	0.0189
Venezuela	2	66	38	57.60% (45.57-69.18)	:	0.6905	0%0	0.4833
HPV16 by country					<0.0001	0.0027		
Argentina	2	101	10	9.88% (4.85-16.42)	:	ref. ^b	0%0	0.7161
Brazil	16	1789	147	9.61% (6.06-13.88)	:	0.8969	86.3%	<0.0001
Chile	1	80	5	6.25% (2.03-12.56)	:	0.6533	:	:
Colombia	3	78	22	38.65% (13.84-67.17)	:	0.0433	80.7%	0.0057
Venezuela	1	50	22	44.00% (30.65-57.81)	:	0.0263		
Collected material					0.0530	0.1364		
Fixed	35	2657	578	22.64% (15.13-31.17)	:	ref. ^b	95.8%	<0.0001
Exfoliated or swab mucosa	33	131	52	44.22% (24.15-65.32)	:	0.1364	75.6%	0.0165
HPV investigation					0.2433	0.2356		
HR	6	1591	417	16.45% (5.03-32.75)	:	ref. ^b	97.6%	<0.0001
LR & HR	29	1197	213	27.14% (18.31-36.99)	:	0.2356	94.2%	<0.0001
Diagnostic method					<0.0001	0.1296		
HPV DNA and IHC	2	624	28	4.15% (1.37-08.34)	:	ref. ^b	79.8%	0.0261
HPV DNA	34	1898	457	24.67% (17.13-33.10)	:	0.0775	93.7%	< 0.0001
HPV DNA or IHC	1	215	127	59.07% (52.18-65.71)	:	0.0249	1	1
IHC	1	51	18	35.29% (22.89-48.80)	:	0.1579	:	1
Profile								
Smoker	13	947	169	20.75% (9.21-35.44)	:	:	95.5%	<0.0001
Drinker	10	720	122	22.23% (8.23-40.62)	:	:	95.7%	<0.0001
Sex								
Male	19	1213	270	19.11% (9.37-31.29)	:	:	95.5%	<0.0001
Female	19	326	87	22.35% (11.85-35.04)		:	82.7%	< 0.0001

^aOhly Brazil and Colombia presented eligible studies. ^aReference value for the moderator test. HR = high-risk, LR = low-risk. HPV = human papillomavirus. HPV DNA = HPV DNA detection techniques. IHC = immunohistochemical detection of overexpression of pl6. HPV DNA or IHC = HPV DNA or IHC = non-combined techniques. DNA and IHC, embined techniques.



	n/N	Prevalence (95% CI)	
Gonzalez et al., 2007 ⁵²	7/16		3.75% (19.75-70.12)
Mosmann et al., 2019 ⁵³	36/85		2.35% (31.70-53.55)
Kaminagakura et al., 2012 ³⁵	22/114		9.30% (12.51-27.75)
De Cicco et al., 2018 ²⁹	127/215	— + — 5	9.07% (52.18-65.71)
Oliveira et al., 2008 ³⁹	17/87		9.54% (11.81-29.43)
Simonato et al., 200844	5/29	— · —	17.24% (5.85-35.77)
Demathe et al., 2010 ³²	13/30		3.33% (25.46-62.57)
Betiol et al., 2016 ²²	27/120	2	2.50% (15.38-31.02)
Soares et al., 2002 ⁴⁵	0/10	P	0.00% (0.00-30.85)
Fregonezi et al., 2012 ³³	5/17		9.41% (10.31-55.96)
Miguel et al., 1998 ³⁸	4/27		14.81% (4.19-33.73)
López et al., 2014 ³⁷	14/212	*	6.60% (3.66-10.83)
Buexm et al., 2020 ²⁴	21/346	+-	6.07% (3.80-9.13)
Anantharaman et al., 2017 ²⁰	7/278	+-	2.52% (1.02-5.12)
Cortezzi et al., 2004 ²⁵	4/32		12.50% (3.51-28.99)
Vidal et al., 2004 ⁴⁷	11/40		7.50% (14.60-43.89)
Soares et al., 2007 ⁴⁶	18/75	2	4.00% (14.89-35.25)
Petito et al., 2017 ⁴⁰	21/82		5.61% (16.60-36.44)
Rivero e Nunes, 200643	0/40	H	0.00% (0.00-8.81)
Hauck et al., 2015 ³⁴	19/154		12.34% (7.59-18.59)
da Silva et al., 2007 ²⁶	37/50	• • 7	4.00% (59.66-85.37)
de Carvalho et al., 2020 ²⁸	18/51	3	5.29% (22.43-49.93)
de Spíndula-Filho et al., 2011 ³¹	0/39	H	0.00% (0.00-9.03)
Lira et al., 2010 ³⁶	61/70	8	7.14% (76.99-93.95)
de Abreu et al., 2018 ²⁷	3/90	+	3.33% (0.69-9.43)
Brew et al., 2012 ²³	8/17	4	7.06% (22.98-72.19)
de Freitas Cordeiro-Silva et al., 2012 ³⁰	3/45		6.67% (1.40-18.27)
Ribeiro et al., 2017 ⁴²	5/6	8	3.33% (35.88-99.58)
Piña et al., 2016 ⁴¹	0/13	H	0.00% (0.00-24.71)
Barros-Filho et al., 2018 ²¹	23/40	5	7.50% (40.89-72.96)
Reyes et al., 201555	9/80		11.25% (5.28-20.28)
Pennacchiotti et al., 201654	2/22		9.09% (1.12-29.16)
Castillo et al., 2011 ⁴⁸	5/6	· · · · · · · · · · · · · · · · · · ·	3.33% (35.88-99.58)
Quintero et al., 2013 ⁵¹	22/112		9.64% (12.74-28.22)
Mancilla et al., 2011 ⁵⁰	8/26	3	0.77% (14.33-51.79)
Erira et al., 2015 ⁴⁹	10/46	2	1.74% (10.95-36.36)
Premoli-De-Percoco et al., 2001 ⁵⁶	30/50	· 6	0.00% (45.18-73.59)
Correnti et al., 200457	8/16	• • • • •	0.00% (24.65-75.35)
Pooled Prevalence	630/2788	2	4.31% (16.87-32.64)
Heterogeneity I ² =95.6%; p<0.0001		I I I I I I 0 20 40 60 80 100	

HPV = human papillomavirus. n = number of HPV-positive cases. N = total number of cases.

Figure 2. Pooled prevalence of human papillomavirus in squamous cell carcinomas of the oral cavity and oropharynx in South American countries.



Venezuela (57.6%), and Chile had the lowest percentage (10.77%). Nevertheless, notably, the number of cases registered for countries, except for Brazil, was reasonably low, making the sample more subject to random variations.

In Europe, in 2014, the pooled prevalence of HPV in HNCs was estimated at 40%, and the highest and lowest percentages were observed in one of the oropharynx and oral cavity subsites, that is tonsil (66.4%) and tongue (25.7%), respectively.⁵⁸ In another meta-analysis that analyzed records from 43 countries, the prevalence of HPV in non-oropharyngeal cancers was 21.8%.59 Although they did not specifically mention the term oral cavity, we can infer that the prevalence observed in the present study (24.2%) was relatively closer to the averages recorded in our study.

Article

in our meta-analysis was low. A recent study investigating the global prevalence of HPV in OPSCC patients reported a pooled percentage of 44.8%. Among the countries analyzed, Brazil was the only representative of South America, with the lowest prevalence (11.1%):⁶⁰ however, notably, the aforementioned study analyzed only 15 cases worldwide, and data from Brazil were collected from a single article with 63 cases.³⁴ In 2015, a meta-analysis was published that demonstrated the increased prevalence of HPV in OPSCCs over four periods in 23 countries: i. pre 1995: 32.3%: 1995–1999: 37%; 2000–2004: 51.8%; and 2005–2014: 52.9%.9 Another meta-analysis with the same theme, published in 2013, calculated a global prevalence of 47.7% for HPV infection in



HPV16 = human papillomavirus genotype 16. \mathbf{n} = number of HPV16-positive cases. \mathbf{N} = total number of cases.

Figure 3. Pooled prevalence of human papillomavirus genotype 16 in squamous cell carcinomas of the oral.



OPSCCs.⁵⁹ Data from the South American countries were not included.

A recently published meta-analysis revealed that the IHC tests for p16^{INK4a} and PCR for detecting HPV DNA are highly sensitive (94% and 98%, respectively), but with moderate specificity (83% and 84%, respectively), and that the combination of these two revealed high sensitivity (98%) and high specificity (96%) for HPV diagnosis in OPSCC.¹⁵ In the present study, the cases tested by IHC or HPV DNA had the highest prevalence of HPV (59.07%) in the random-effects model, followed by cases tested by only IHC (35.29%); however, it is relevant to note that these two situations only presented one study (A16²⁹ and A22²⁸). The lowest prevalence was observed in studies in which HPV was diagnosed using a combination of DNA and IHC techniques (4.15%), but only two studies presented this diagnostic methodology (A01²⁰ and A05²⁴).

In our meta-analysis, the pooled prevalence of HPV among smokers was 20.75% and 22.23% for alcoholics. The consumption of beverages and smoking increases the risk of developing cancers of the oropharynx and oral cavity.⁶ In addition, evidence suggests an association between smoking and HPV carcinogenesis and a worse prognosis in smokers' overall survival with HNC HPV positive cases.^{61,62}

The main limitation of our study is the heterogeneity of the sample. Nevertheless, even if more factors were stratified, hetero-

geneity would undoubtedly be maintained. Despite a broad research tactic, the initial number of 209 records without duplication, of which only 52 complete studies were selected for reading, is reasonably low if we consider that no filter was included to limit the period of publication. Another factor to be highlighted as limiting a more effective analysis was the deficiency of information presented in a considerable part of the analyzed studies, of 38 studies, 15 did not mention the period corresponding to the collection of the samples, making any analysis impossible about the evolution of prevalence over time. Data that could support heterogeneity analysis could not be extracted from many of the studies analyzed because the data presented did not provide sufficient detail to stratify the information. Cofactors such as sex, age, smoking, alcoholism, and risky sexual behavior have not been reported in many reviewed articles. Even when approached, they did not present sufficient details for the quantitative analysis. Indeed, the simple provision of raw data with metadata would be sufficient to improve the information collected. However, only three of the studies analyzed provided supplementary material for consultation.

In conclusion, we found an overall prevalence of HPV in 24.31% of oral and oropharyngeal squamous cell carcinomas in South American patients. The prevalence of HPV was 17.9% for oropharyngeal cancer and 23.19% for oral cavity cancer in South American patients.



HPV = human papillomavirus. n = number of HPV-positive cases. N = total number of cases.

Figure 4. Pooled prevalence of human papillomavirus in squamous cell carcinomas of the oropharynx in South American countries.



	n/N	Prevalence (95% CI)	
Gonzalez et al., 2007 ⁵²	7/16		43.75% (19.75-70.12)
Mosmann et al., 2019 ⁵³	36/85		42.35% (31.70-53.55)
Kaminagakura et al., 2012 ³⁵	22/114		19.30% (12.51-27.75)
Oliveira et al., 2008 ³⁹	17/87	· _	19.54% (11.81-29.43)
Simonato et al., 200844	5/29	— ,	17.24% (5.85-35.77)
Demathe et al., 2010^{32}	13/30		43.33% (25.46-62.57)
Betiol et al., 2016 ²²	23/92		25.00% (16.55-35.11)
Soares et al., 2002 ⁴⁵	0/10		0.00% (0.00-30.85)
Fregonezi et al., 2012 ³³	5/17		29.41% (10.31-55.96)
Miguel et al., 1998 ³⁸	1/21		4.76% (0.12-23.82)
López et al., 2014 ³⁷	8/121		6.61% (2.90-12.61)
Anantharaman et al., 2017 ²⁰	0/107	F	0.00% (0.00-3.39)
Cortezzi et al., 2004 ²⁵	1/11		9.09% (0.23-41.28)
Vidal et al., 200447	11/40		27.50% (14.60-43.89)
Soares et al., 2007 ⁴⁶	18/75		24.00% (14.89-35.25)
Petito et al., 2017 ⁴⁰	10/39		25.64% (13.04-42.13)
Rivero e Nunes, 200643	0/40	H	0.00% (0.00-8.81)
Hauck et al., 2015 ³⁴	6/83		7.23% (2.70-15.07)
da Silva et al., 2007 ²⁶	37/50	.	74.00% (59.66-85.37)
de Spíndula-Filho et al., 2011 ³¹	0/39	F	0.00% (0.00-9.03)
Lira et al., 2010 ³⁶	61/70	· _·	87.14% (76.99-93.95)
de Abreu et al., 2018 ²⁷	3/90	+	3.33% (0.69-9.43)
Brew et al., 2012 ²³	8/17		47.06% (22.98-72.19)
de Freitas Cordeiro-Silva et al., 2012 ³⁰	3/41		7.32% (1.54-19.92)
Ribeiro et al., 2017 ⁴²	5/6		— 83.33% (35.88-99.58)
Reyes et al., 201555	9/80		11.25% (5.28-20.28)
Pennacchiotti et al., 201654	2/22		9.09% (1.12-29.16)
Castillo et al., 2011 ⁴⁸	5/6		— 83.33% (35.88-99.58)
Quintero et al., 2013 ⁵¹	16/67	,	23.88% (14.31-35.86)
Mancilla et al., 2011 ⁵⁰	8/26		30.77% (14.33-51.79)
Erira et al., 2015 ⁴⁹	10/46		21.74% (10.95-36.36)
Premoli-De-Percoco et al., 2001 ⁵⁶	30/50		60.00% (45.18-73.59)
Correnti et al., 2004 ⁵⁷	8/16		50.00% (24.65-75.35)
Pooled Prevalence	388/1643	\sim	23.19% (14.94-32.63)
Heterogeneity I ² =94.5%; p<0.0001		I I I I I 0 20 40 60 80	l 100

HPV = human papillomavirus. n = number of HPV-positive cases. N = total number of cases.

Figure 5. Pooled prevalence of human papillomavirus in squamous cell carcinomas of the oral cavity in South American countries.



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