

Human Papillomavirus Prevalence and Genotype Distribution Landscapes in Shannan City, Tibet Tibetan Autonomous Region, China

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Abstract

Background:

Data regarding human papillomavirus (HPV) prevalence and genotype distribution are limited in Shannan City, Tibet Tibetan Autonomous Region, China. The purpose of this study is to provide reliable data for guiding women in Shannan City in cervical cancer screening and HPV vaccine inoculation.

Methods:

HPV testing was performed on women aged 16 to 109 years (mean age 44.03 ± 9.25 years) from Shannan City in 2019 and 2020, which was implemented technically by gynecological examination, vaginal discharge smear microscopy, cytology, and HPV detection. The overall prevalence, age-specific prevalence, and genotype distribution were analyzed.

Results:

A total of 48,126 women received HPV testing, of which 3,929 were detected human papillomavirus. The HPV-positive rate was 8.16% (3,929/48,126), and the highest prevalence was in the ≤ 25 -year-old age group (12.68%). After the age of 25, the prevalence rate decreased rapidly, and then slowly increased from 7.49% in the 46-55 age group to 9.82% in the ≥ 66 age group, showing a “U-shaped” pattern. The positive prevalence of HPV 16 or 18-only was 1.43%, that of other HPV genotypes except HPV 16 or 18 was 6.39%, and mixed HPV infections including HPV 16 or 18 was 0.34%.

Conclusions:

The HPV infection rate in Shannan city is rather low, and the age-specific prevalence of HPV infection presents a “U” curve, suggesting the importance of screening among younger women and the necessity of detection among older women.

Background

Cervical cancer is one of the common malignant tumors in gynecology. According to the latest global cancer burden data in 2020 released by the international agency for research on cancer (IARC) of the World Health Organization, cervical cancer is the fourth most common female cancer in China and the fourth leading cause of female death[1]. Human papillomavirus (HPV) infection is the main cause of cervical cancer and intraepithelial neoplasia (CIN)[2][3], of which HPV 16 and 18 subtypes are the most common causative factors[4]. Prevention of cervical cancer starts with the prevention of HPV infection. Previous studies have shown that the HPV prevalence has obvious regional and age-specific distribution[5][6]. Understanding the characteristics of HPV infection plays an important role in the

prevention and treatment of cervical cancer. In this study, We collected basic information and HPV high-risk subtypes of patients from Shannan Maternal and Child Health Hospital from January 2019 to December 2020, and analyzed the HPV infection landscapes in different and continuous age groups to intervene early in cervical precancerous lesions and cervical cancer, thereby reducing the incidence rate of cervical cancer in Shannan city.

Shannan city is located in Southwest China, with an average altitude of about 3700 meters. It is a vast and remote area composed of Tibetan, Han, Hui, and other nationalities. Cervical cancer is still a major public health problem for women living in Shannan city due to the lack of Health Science Popularization for HPV detection and prevention, insufficient local health conditions, scattered population, and ethnic culture. However, data on HPV infection rates in Shannan city are badly limited.

Here, we examined the HPV infection rate of women in Shannan city and its relationship with age according to the data collected from female patients. These data will provide epidemiological evidence to support the development of a comprehensive and correct cervical cancer prevention program. We also aim to emphasize the need for complete, thorough and detailed data collection to increase our understanding of HPV infection rates.

Methods

Study population

A total of 48,126 females aged 16-109 years were enrolled in this study from January 2019 to December 2020. All participants were from Shannan Maternal and Child Health Hospital. Patients visited the hospital for various reasons, including physical examination, vaginitis, abnormal vaginal bleeding, Gynaecological tumors, etc. Clinical information was collected from the patients, and a molecular survey of HPVs was conducted. Eligible women were included in the study after signing an informed consent form.

Sample collection

Doctors from Wuhan Union Hospital (Wuhan, China) and local clinics were trained in the collection of cervical cell samples for HPV testing. All participants were informed to avoid sex within 24 hours before collecting samples. With the help of a vaginal endoscope, the orificium externum isthmus was fully exposed and was scraped through clockwise 5 rotations gently using the HPV detection brush, so that the mucosa and secretions could adhere to the flat sides of the bristles. Then the tips of the cervical brush were put into a vial containing transport medium separately, which were stored at 2-8°C until HPV DNA extraction and genotyping could be carried out. The HPV genotype detection of samples was completed within 48 hours.

Statistical analysis

Analyses were done with SPSS (version 25.0) and Excel (version 2004). Descriptive statistical analysis was performed on the distribution of HPV genotypes using indicators such as frequency and prevalence.

For comparisons among different age groups, the categorical variables were compared by using the Chi-square test. All the reported P values were made based on two-sided tests with a significance level of 0.05.

Results

Prevalence and genotype distribution of human papillomavirus

From January 2019 to December 2020, a total of 48,126 human papillomavirus genotype samples were collected and detected in Shannan Maternal and Child Health Hospital. The subjects were aged 16-109 years, and the mean age was 44.03 ± 9.25 years. The genotype test showed that 3,929 samples were HPV positive, so the overall prevalence of all types of human papillomavirus in the study population was 8.16%.

The details of genotypes distribution were shown in Table 1 and 2. Notably, HPV 16-only infection rate was 0.52% (250/48,126). Correspondingly, HPV 16-total was 0.70% (339/48,126). HPV 18-only infection rate was 0.91% (440/48,126) and HPV 18-total was 1.09% (526/48,126), respectively. In addition, the HPV infection rate of other genotypes was 6.73% (3,239/48,126), accounting for 82.44% (3239/3929) of all HPV-positive samples.

Table 1
Frequency and prevalence of genotypes of HPV among women.

Genotype	Positive samples (n)	Proportion among 48,126 HPV-positive samples (%)	Proportion among 3,929 HPV-positive samples (%)
16	250	0.52	6.36
18	440	0.91	11.20
others	3077	6.39	78.32
16+others	76	0.16	1.93
18+others	73	0.15	1.86
16+18+others	13	0.03	0.33

Table 2
Frequency and prevalence of HPV16-18 and other genotypes among women.

Genotype	Positive samples (n)	Proportion among 48,126 HPV-positive samples (%)	Proportion among 3,929 HPV-positive samples (%)
16	339	0.70	8.63
18	526	1.09	13.39
others	3239	6.73	82.44

Prevalence of HPV grouped by age

Overall, 48,126 women (aged 16-109 years) were included in this study. All the participants were divided into six groups ranging from ≤ 25 years, 26–35 years, 36–45 years, 46–55 years, 56–66 years, and ≥ 66 years. There were significant differences in HPV infection rates among the six age groups ($P < 0.05$). Relevant data could be seen in Table 3.

The distribution of HPV infection rates in specific age groups presented a “U” curve. The first peak appeared in the youngest women (12.68%). However, the HPV infection rate decreased sharply after the first peak, and reached the bottom in the 46-55 age group (7.49%). On the contrary, it increased slowly, and finally reached the second peak at 9.82% in the oldest women group.

Table 3
Prevalence of HPV grouped by age groups.

Age group (years)	Mean age (years)	Total samples (n)	Positive samples (n)	Proportion among total samples (%)
≤ 25	23.32 \pm 1.80	891	113	12.68
26-35	31.99 \pm 2.79	8289	726	8.76
36-45	40.35 \pm 2.85	17640	1415	8.02
46-55	50.15 \pm 2.80	15438	1157	7.49
56-65	58.76 \pm 2.35	5705	502	8.80
≥ 66	69.57 \pm 6.31	163	16	9.82
Total	44.03 \pm 9.25	48126	3929	8.16
χ^2	41.542			
P	<0.05			

Region-Specific Prevalence of HPV Infection

The prevalence of HPV infection in Shannan city and in 19 different areas of China was compared separately (Table 4). The positive rate of HPV obtained in our study (Shannan City) was significantly different from those in other areas ($p < 0.05$). Moreover, the highest rate of HPV infection was in Henan Province (38.10%, 1,536/4,033)[7], whereas the lowest area was in Shannan City (8.16%, 3,929/48,126).

Table 4
Comparison of HPV Infection Rates in Different Areas in China

Regions	Total samples (n)	Positive samples (n)	Prevalence (%)	P-value
Shannan City	48,126	3,929	8.16	-
Beijing City[8]	29,436	3,586	12.18	<0.001
Shanghai City[9]	23,724	3,816	16.08	<0.001
Wuhan City[10]	13,775	2,436	17.68	<0.001
Xinjiang Province[11]	37,722	5,287	14.02	<0.001
Jiangxi Province[12]	71,435	16,065	22.49	<0.001
Sichuan Province [13]	14,185	3,382	23.84	<0.001
Yunnan Province[14]	17,898	3,681	12.94	<0.001
Zhejiang Province[11]	77,443	17,270	22.30	<0.001
Guangdong Province[16]	33,328	3,526	10.58	<0.001
Jiangsu Province[17]	62,317	16,775	26.92	<0.001
Shaanxi Province[18]	17,341	4,559	26.30	<0.001
Shandong Province[19]	94,489	26,839	28.40	<0.001
Henan Province[7]	4,033	1,536	38.10	<0.001
Shanxi Province[20]	7,640	1,441	18.86	<0.001
Liuzhou City[21]	2,300	522	22.70	<0.001
Jilin Province[22]	20,648	7,095	34.40	<0.001
Heilongjiang Province[23]	18,522	5,011	27.10	<0.001
Liaoning Province[24]	6,479	667	10.30	<0.001
Inner Mongolia[25]	2,345	844	36.00	<0.001

Discussion

HPV screening is very important for the prevention and detection of cervical cancer[26]. However, cervical cancer screening started late in China, and there is still a big gap between the current promotion and

developed countries. In addition, the data of HPV infection rate and genotype distribution in different regions are not comprehensive, especially in the remote areas in Western and Northern China. There is a lack of relevant data to guide the health education of cervical cancer. Therefore, collecting and analyzing the epidemiological evidence of local HPV infection can provide a reliable scientific basis for the prevention, treatment, and elimination of human papillomavirus infection-related diseases.

Shannan city of Tibet Tibetan Autonomous Region was located in Northwest China. Its economy was underdeveloped, and the prevention and screening of cervical cancer were also quite backward. Therefore, improving the screening of HPV in Tibet was particularly important for the primary prevention of cervical cancer in Tibet. This is not only conducive to women's health but also can save a lot of medical expenses for the country. Studying the prevalence and genotype distribution of human papillomavirus in different regions and periods is highly important for cervical cancer screening and evaluating the effectiveness of the human papillomavirus vaccine for women. For the last 20 years, the world has made great efforts to generate epidemiological data on cervical HPV-DNA. In China, although certain studies have been performed to assess the prevalence and incidence rate of human papillomavirus genotypes in Tibet, they are based on small samples[27][28]. Our study is the first large-scale sample study in Tibet.

Due to the differences in regional, population, living environment and lifestyle, and human papillomavirus DNA test methods, the reported results of global human papillomavirus distribution vary from study to study. A meta-analysis[29] of a total of 1,016,719 screening people included in 194 studies around the world showed that the adjusted infection rate of HPV in the global population with normal cytology was 11.7%, of which the infection rate was the highest in sub-Saharan Africa (24.0%), Eastern Europe (21.4%) and Latin America (16.1%) and the lowest in Western Asia (1.7%). According to research, the overall prevalence of human papillomavirus infection in China is 15.54%[30]. In our study, the rate of human papillomavirus infection among women in Shannan City, Tibet was 8.16%, lower than the global average, lower than the Chinese average, and lower than many other cities or regions in China. Some researchers speculated that the variability of HPV prevalence in China was due to China's large population and territory[30]. At the same time, the level of economic development and population migration also led to the differences in the distribution landscapes of human papillomavirus among regions. As shown in Table 4, both Beijing and Shanghai were economically developed cities, but the HPV infection rate varied greatly[8][9]. The reason may be that Beijing, as China's political, cultural, and economic center, had good medical conditions and protection strategies. Otherwise, Shanghai's economic development level was also very high, but due to a large number of foreigners and migrant population, the city was expanding and the population composition was complex, resulting in a high rate of human papillomavirus infection. In Shannan City, although the economy and medical treatment were relatively backward, the HPV infection rate was the lowest. The reason behind it may be the simple folk customs, the majority of the people who believe in Tibetan Buddhism and the conservative traditional concept of sex, which was similar to Xinjiang Province. Another study on the HPV infection rate in Tibet which was 9.19% (279/3,036), which supports this conclusion[27].

Information on the distribution of human papillomavirus infection in different age groups is extremely important for the design of human papillomavirus preventive vaccines in specific age groups[31]. What many studies have in common is that the first peak occurs in the younger age group (just after the beginning of sexual relations). In some areas, the second peak can be observed at the age of > 45 or > 55 or > 65, while in some other areas, no second peak can be observed. In conclusion, age-specific HPV distribution is either shown as a bimodal curve (including "U" curve)[32][33] or a left inclined unimodal distribution[34]. In this study, age-specific HPV distribution showed a "U" curve. The first peak of human papillomavirus infection occurred in the age group ≤ 25 years old (12.68%), then decreased gradually, reached the lowest in the age group 46-55 years old, and then increased gradually. The reason for this trend may be that young women were sensitive to human papillomavirus soon after sexual activity due to immature immune protection[35]. With the stimulation of immune response, a large part of primary HPV infection would be temporary and would be cleared spontaneously[36], so the HPV infection rate decreased gradually. The immune ability of elderly women decreased with age, especially in the premenopausal and postmenopausal women, which resulted their ability to eliminate previous and new HPV infections being weakened. Furthermore, as past (latent) infections reappeared, both factors led to a higher HPV infection rate of elderly women[37]. Based on these findings, the earlier young women were vaccinated with HPV vaccine, the higher the antibody titer and the better the protection[38]. Once HPV infection was detected, HPV viral load should be continuously monitored and cervical biopsy should be performed regularly.

Although a variety of studies provided large-scale information on the recent HPV prevalence and genotype distribution in Shannan City, Tibet, China, there were still some limitations. First, most cervical screening tests received by patients did not carry out detailed HPV typing, nor did they be combined with cytology. Women included in some studies were unable to obtain cervical cytological or histological results. Therefore, it is impossible for doctors or researchers to associate HPV infection and genotype distribution with different cervical abnormalities. Secondly, the collection of personal information of patients was not complete. Tibet had a high altitude and a wide area. There was a variety of distinctive information about the nationality, living habits, reproductive history, climatic conditions cultural, and other backgrounds of the population in this area. Unfortunately, it was not recorded in this study so that we couldn't specify the impact of these different backgrounds on the rate of HPV infection. In addition, some studies had shown that human papillomavirus infection may lead not only to cervical cancer but also to oropharyngeal cancer and head and neck cancer[39]. Therefore, it was suggested to also focus on the relationship between HPV infection and oropharyngeal cancer, and head and neck cancer in Tibetan women in future research. In addition, obtaining data on human papillomavirus infection in Tibetan men could be regarded as a potential direction for future research.

After all, our study revealed the HPV infection rate and genotype distribution of women from Shannan City, Tibet, China. This information may provide guidance and suggestions for the prevention of cervical cancer for women in this area. According to the current results, young women should be vaccinated with HPV vaccine as soon as possible, and elderly women should focus on crevice screening.

Declarations

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Authors' contributions

XWZ and HBW designed and supervised the study. DLF and JC collected patient clinical information. ZCY analyzed and interpreted the data. STW prepared the manuscript. All authors read and approved the final manuscript.

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Availability of data and materials

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Ethics approval and consent to participate

This study followed the ethical guidelines of the Declaration of Helsinki and was approved by Ethics Committee of the Tongji Medical college of Huazhong University of Science and Technology.

Consent to publication

All of the patients signed an informed consent form.

Competing interests

The authors declare that they have no competing interests.

References

1. Sung, H., et al., Global Cancer Statistics 2020: GLOBOCAN Estimates of Incidence and Mortality Worldwide for 36 Cancers in 185 Countries. *CA Cancer J Clin*, 2021. 71(3): p. 209–249.
2. Schiffman, M., et al., Human papillomavirus and cervical cancer. *Lancet*, 2007. 370(9590): p. 890–907.
3. Crosbie, E.J., et al., Human papillomavirus and cervical cancer. *Lancet*, 2013. 382(9895): p. 889–99.
4. Walboomers, J.M.M., et al., Human papillomavirus is a necessary cause of invasive cervical cancer worldwide. *The Journal of Pathology*, 1999. 189(1): p. 12–19.

5. de Sanjose, S., et al., Human papillomavirus genotype attribution in invasive cervical cancer: a retrospective cross-sectional worldwide study. *Lancet Oncol*, 2010. 11(11): p. 1048–56.
6. Hammer, A., et al., Age-specific prevalence of HPV16/18 genotypes in cervical cancer: A systematic review and meta-analysis. *Int J Cancer*, 2016. 138(12): p. 2795–803.
7. Zhao, J.W., et al., Prevalence and genotype distribution of human papillomavirus: implications for cancer screening and vaccination in Henan province, China. *Rev Soc Bras Med Trop*, 2016. 49(2): p. 237–40.
8. Yu, H., et al., Prevalence and Genotype Distribution of Human Papillomavirus Among Healthy Females in Beijing, China, 2016-2019. *Infection and drug resistance*, 2021. 14: p. 4173–4182.
9. Li, P., et al., Characteristics of human papillomavirus infection among women with cervical cytological abnormalities in the Zhoupu District, Shanghai City, China, 2014-2019. *Viol J*, 2021. 18(1): p. 51.
10. Xiang, F., et al., Distribution characteristics of different human papillomavirus genotypes in women in Wuhan, China. *J Clin Lab Anal*, 2018. 32(8): p. e22581.
11. Pan, Z., et al., Screening for HPV infection in exfoliated cervical cells of women from different ethnic groups in Yili, Xinjiang, China. *Sci Rep*, 2019. 9(1): p. 3468.
12. Zhong, T.Y., et al., Prevalence of human papillomavirus infection among 71,435 women in Jiangxi Province, China. *J Infect Public Health*, 2017. 10(6): p. 783–788.
13. Luo, Q., et al., Prevalence and genotype distribution of HPV and cervical pathological results in Sichuan Province, China: a three years surveys prior to mass HPV vaccination. *Viol J*, 2020. 17(1): p. 100.
14. Baloch, Z., et al., Epidemiologic characterization of human papillomavirus (HPV) infection in various regions of Yunnan Province of China. *BMC Infect Dis*, 2016. 16: p. 228.
15. Li, M., et al., Incidence, persistence and clearance of cervical human papillomavirus among women in Guangdong, China 2007-2018: A retrospective cohort study. *J Infect Public Health*, 2021. 14(1): p. 42–49.
16. Zhang, C., et al., Distribution of human papillomavirus infection: a population-based study of cervical samples from Jiangsu Province. *Viol J*, 2019. 16(1): p. 67.
17. Jiang, L., et al., HPV prevalence and genotype distribution among women in Shandong Province, China: Analysis of 94,489 HPV genotyping results from Shandong's largest independent pathology laboratory. *PLoS One*, 2019. 14(1): p. e0210311.
18. Li, J., et al., [Analysis of human papillomavirus infection and typing in Shanxi province]. *Zhonghua Yu Fang Yi Xue Za Zhi*, 2014. 48(3): p. 192–6.
19. Wu, X., et al., [Prevalence of type-specific human papillomavirus infection among 18-45 year-old women from the general population in Liuzhou, Guangxi Zhuang Autonomous Region: a cross-sectional study]. *Zhonghua Liu Xing Bing Xue Za Zhi*, 2017. 38(4): p. 467–471.

20. Hao, S., et al., HPV genotypic spectrum in Jilin province, China, where non-vaccine-covered HPV53 and 51 are prevalent, exhibits a bimodal age-specific pattern. *PLoS One*, 2020. 15(3): p. e0230640.
21. Liu, J., et al., Epidemiology and persistence of cervical human papillomavirus infection among outpatient women in Heilongjiang province: A retrospective cohort study. *J Med Virol*, 2020.
22. Xue, H., et al., Prevalence and genotype distribution of human papillomavirus infection in asymptomatic women in Liaoning province, China. *J Med Virol*, 2015. 87(7): p. 1248–53.
23. Ji, Y., et al., The Burden of Human Papillomavirus and Chlamydia trachomatis Coinfection in Women: A Large Cohort Study in Inner Mongolia, China. *J Infect Dis*, 2019. 219(2): p. 206–214.
24. Goodman, A., HPV testing as a screen for cervical cancer. *Bmj*, 2015. 350: p. h2372.
25. Jin, Q., et al., [Prevalence of human papillomavirus infection in women in Tibet Autonomous Region of China.]. *Zhonghua Fu Chan Ke Za Zhi*, 2009. 44(12): p. 898–902.
26. Chen, L., et al., The genomic distribution map of human papillomavirus in Western China. *Epidemiol Infect*, 2021. 149: p. e135.
27. Bruni, L., et al., Cervical Human Papillomavirus Prevalence in 5 Continents: Meta-Analysis of 1 Million Women with Normal Cytological Findings. *The Journal of Infectious Diseases*, 2010. 202(12): p. 1789–1799.
28. Zhu, B., et al., The prevalence, trends, and geographical distribution of human papillomavirus infection in China: The pooled analysis of 1.7 million women. *Cancer Med*, 2019. 8(11): p. 5373–5385.
29. Skinner, S.R., et al., Efficacy, safety, and immunogenicity of the human papillomavirus 16/18 AS04-adjuvanted vaccine in women older than 25 years: 4-year interim follow-up of the phase 3, double-blind, randomised controlled VIVIANE study. *Lancet*, 2014. 384(9961): p. 2213–27.
30. Xue, Y., et al., “U” shape of age-specific prevalence of high-risk human papillomavirus infection in women attending hospitals in Shanghai, China. *European Journal of Obstetrics & Gynecology and Reproductive Biology*, 2009. 145(2): p. 214–218.
31. Liao, G., et al., Multi-Infection Patterns and Co-infection Preference of 27 Human Papillomavirus Types Among 137,943 Gynecological Outpatients Across China. *Front Oncol*, 2020. 10: p. 449.
32. Chen, L., et al., The genomic distribution map of human papillomavirus in Western China. *Epidemiol Infect*, 2021. 149: p. e135.
33. Alder, S., et al., Acceptance of human papillomavirus (HPV) vaccination among young women in a country with a high prevalence of HPV infection. *Int J Oncol*, 2013. 43(4): p. 1310–8.
34. Rodríguez, A.C., et al., Rapid clearance of human papillomavirus and implications for clinical focus on persistent infections. *J Natl Cancer Inst*, 2008. 100(7): p. 513–7.
35. González, P., et al., Behavioral/lifestyle and immunologic factors associated with HPV infection among women older than 45 years. *Cancer Epidemiol Biomarkers Prev*, 2010. 19(12): p. 3044–54.
36. Spinner, C., et al., Human Papillomavirus Vaccine Effectiveness and Herd Protection in Young Women. *Pediatrics*, 2019. 143(2).

Figures

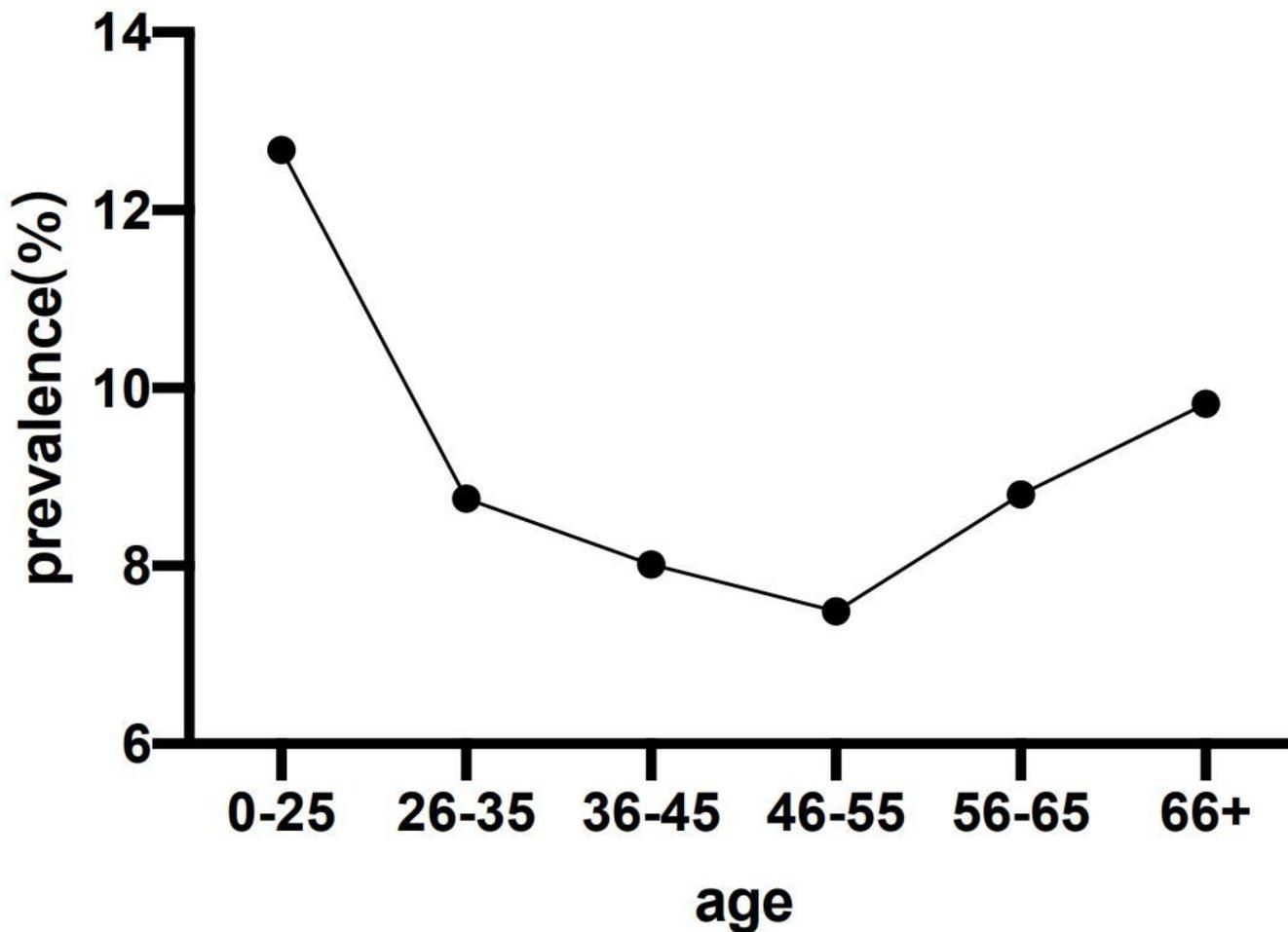


Figure 1

Prevalence of HPV infection in different age intervals. The highest HPV infection rates was in women aged ≤ 25 years (12.68%), and the lowest was in the 46-55 age group (7.49%). The overall performance was a “U” shape.