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Prevention of human papillomavirus (HPV) infection and cervical cancer in China: How does HPV vaccination bring about benefits to Chinese women?

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Summary

Cervical cancer is the second leading cause of cancer deaths among women in the world and more than 85% of cervical cancer cases occur in women living in developing countries. Human papillomavirus (HPV) infection is the major cause of cervical cancer. Since 2006, two prophylactic vaccines against the high-risk strains of HPV have been developed and approved in more than 100 countries around the world. However, in China, HPV vaccines are still under clinical trials for government approval. In this paper feasibility and justification of HPV vaccine introduction into China is examined by reviewing experiences in both developed and developing countries where the vaccination program has been implemented. The vaccination program has showed significant cost-effectiveness and great health and economic impacts on cervical cancer prevention and control in both high-income and middle- and low-income countries. On the other hand, based on the lessons from both developed and developing countries, secondary prevention alone cannot fully play a role to reduce the incidence and the disease burden, and neither does the vaccination program. The epidemiological characteristics in China suggest an urgent need to introduce the vaccines and the geographically diversified prevalence of oncogenic HPV types as well as socioeconomic status also highlight the importance of region-driven approaches for cervical cancer prevention and control by integration of a screening and vaccination program.

Keywords: Human papillomavirus (HPV), cervical cancer, vaccination, screening, China

1. Introduction

Cervical cancer is the second leading cause of cancer death among women in the world and more than 85% of cervical cancer cases occur in women living in developing countries, where approximately 529,000 new cases and 275,000 deaths occur every year (1). Human papillomavirus (HPV) infection acquired from sexual activities is the most common viral infection of the reproductive tract and the causal relationship between HPV infection and the cervix and cervical cancer was built by zur Hausen, the Nobelist in Physiology

and Medicine in 2008 for his epoch-making findings. The infection can be detected in more than 95% of carcinoma issues (2). So far, more than 100 different HPV genotypes have been detected, and among them type 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, and 68 are a high-risk for cervical carcinoma. The most common genotypes among infected women are 16, 18, 31, 58, and 52, accounting for approximately half of HPV infections (3).

Recently, there has been a milestone which prevents HPV infection and development of cervical cancer: two prophylactic vaccines against the high-risk strains of HPV, a quadrivalent vaccine *Gardasil* developed by *Merck* and a bivalent vaccine *Cervarix* by *GlaxoSmithKline* have been developed and approved in more than 100 countries around the world. The prophylactic vaccines mainly target HPV-16 and -18 types, which are the most prevalent genotypes globally

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and responsible for about 70% of cervical cancer worldwide (1). To fully play a role for the prevention of cervical cancer and diseases related to HPV infection, they need to be widely implemented in the appropriate target population, as both are only effective for those not infected with HPV, preferably prior to sexual debut. When girls begin having sex, the risk of HPV infection increases dramatically, weakening the effectiveness of the vaccination strategy. Table 1 summarizes basic information of the two vaccines (4,5). Both vaccines have shown excellent efficacy with minimal toxicity; on the other hand, numerous questions remain, such as delivery strategies, accessibility to vaccination for underserved populations, social acceptance, monitoring of safety and effectiveness post-licensure, and integration of current existing HPV screening in both developed and developing countries. Monitoring and evaluation of the long-term health and socioeconomic impacts including side effects is necessary not only for each country, but also for the global society.

Having characterized the geographical diversity, epidemiological characteristics of HPV infection in China is quite different by region (6). The disease burden of cervical cancer is high, particularly in the rural area. It is estimated that among Chinese women aged 30 to 50 years, the prevalence of infection with high-risk HPV is 15.0~20.8%, and the mortality of cervical cancer increases 4.1% per year (7). On the other hand, the

two prophylactic vaccines are still under clinical trials among the Chinese population for government approval. Although time is needed for the introduction of the HPV vaccines in China, it is expected to bring benefits to Chinese women with more effective prevention of HPV infection and cervical cancer in the long-term and the feasibility and justification should be examined by reviewing experiences in developed and developing countries where the vaccination program has been launched.

2. Strategies and impacts of HPV vaccination on the prevention of HPV infection and cervical cancer

The World Health Organization (WHO) strongly recommended an introduction and scaling-up of the HPV vaccination program (1). So far, there are more than 160 countries which have approved the prophylactic vaccines and have gradually introduced the vaccines into the national routine immunization program. Generally, the program in most countries targets pre-adolescent and adolescent girls whose age ranges from 9 to 13 years old by school-based or healthcare facility-based or mixed approaches with a catch-up group aged up to 26 years old. It provides 3 injections during a 6 month period, as the protective effect by the vaccines has been shown prior to exposure of the risk to HPV infection, *i.e.*, the sexual debut. In

Table 1. Basic information of two types of HPV vaccine

Items	<i>Gardasil (quadrivalent)^a</i>	<i>Cervarix (bivalent)^b</i>
Pharmaceutical company	Merck, Whitehouse Station, NJ, USA	GlaxoSmithKline, Rixensart, Belgium
HPV types	6, 11, 16, 18	16, 18
Prevention of diseases	Girls and women <ul style="list-style-type: none"> • Cervical, vulvar, vaginal and anal cancer caused by HPV types 16 and 18 • Genital warts (condyloma acuminata) caused by HPV types 6 and 11 • Precancerous or dysplastic lesions caused by HPV types 6, 11, 16, and 18 Boys and men <ul style="list-style-type: none"> • Anal cancer caused by HPV types 16 and 18 • Genital warts (condyloma acuminata) caused by HPV types 6 and 11 • Precancerous or dysplastic lesions caused by HPV types 6, 11, 16, and 18 	Girls and women <ul style="list-style-type: none"> • Following diseases caused by HPV types 16 and 18 • cervical cancer • cervical intraepithelial neoplasia (CIN) grade 2 or worse and adenocarcinoma <i>in situ</i>, and cervical intraepithelial neoplasia (CIN) grade 1
Target population	Females and males aged 9 to 26 years	Females aged 9 to 25 years
Administration	Intramuscular shoulder injection	Intramuscular shoulder injection
Efficacy against precancerous lesions	For females without infection of types 16 and 18: 98% (95% CI: 86-100%) For females with infection of types 16 and 18: 44% (95% CI: 26-58%)	For females without infection of types 16 and 18: 93% (95% CI 80-98%) For females with infection of types 16 and 18: less efficient
Cross-protection effects	Protection against types 31 was 46% (95% CI 15-66) for persistent infection and 57% (29-75) for any CIN or adenocarcinoma in situ	Protection against CIN2 or worse associated with types 31 and 33 in lesions with no co-infection with the vaccine types, and to a lesser extent against types 45 and 51
Safety	0.8% of individuals who received the vaccine and 1.0% of individuals who received the placebo reported a serious systemic adverse reaction	5.3% of individuals who received the vaccine and 5.9% of individuals who received the placebo reported at least one serious adverse event, without regard to causality

^a, Merck & Co. Gardasil (Human papillomavirus quadrivalent (types 6, 11, 16, and 18) vaccine, recombinant); ^b, GlaxoSmithKline. Cervarix (Human papillomavirus bivalent (types 16 and 18) vaccine, recombinant).

developed countries, during the past several years, the coverage of the vaccines has increased and caused a significant impact on reduction of the incidence of cervical and other HPV-associated cancers, including prevalence of HPV infection, genital warts and cervical lesions (8). With evidence that showed high efficacy for prevention of genital warts and anal precancerous lesions in males, *Gardasil* is also licensed and recommended for use in boys in some countries such as the United States since last year (9). In middle- and low-income countries, although several challenges such as affordability, infrastructure for delivery, coverage, and communication strategies to boost public acceptability still remain in the national cervical cancer prevention and control strategies (10), the vaccination program has been adopted by public financing, out-of-pocket payment or donations. The GAVI Alliance has subsidized low-income countries to increase financial feasibility for the introduction.

As summarized in Table 2 (11-42), the vaccination program has showed significant cost-effectiveness and great health and economic impacts on cervical cancer prevention and control in both high-income and middle- and low-income countries, even if the protocol of vaccination and coverage is diversified depending on the specific situation of each country. In middle- and low-income countries, the combination of vaccination and screening showed better cost-effectiveness than that of vaccination or screening alone. On the other hand, uncertainties for the cost-effectiveness of the different options of the prevention and control programs included coverage, price of vaccines, oncogenic and epidemiological characteristics of HPV infection, suggesting the issues should be carefully examined when introducing the vaccination program. Moreover, the vaccination program has also brought significant cross-protection impacts on prevention of HPV-related diseases, such as cervical intraepithelial neoplasia (CIN), adding further value to the preventive potential and clinical benefits of the vaccines (43).

3. Cervical cancer control in China: current status and challenges

3.1. Strategies and limitations of early detection of cervical cancer by the screening program

Recently in China, without official approval of the vaccines, the most essential strategy for cervical cancer prevention and control is the routine screening program targeting women at reproductive age for early detection and treatment of HPV infection, CIN (from which developing cervical cancer generally needs several years) and cervical cancer at an early stage. Based on previous large-scale studies, a systematic routine screening program can reduce the incidence of cervical cancer by at least 60% and has been recommended by

WHO, particularly in developing countries (44). The major methodologies of screening include: *i*) Pap smear cytology test; *ii*) liquid-based cytology test; *iii*) HPV DNA test (HC-II); *iv*) visual inspection with acetic acid (VIA); *v*) colposcopy. Table 3 briefly lists advantages and limitations of each methodology (45).

The Ministry of Health of China has launched the Guideline for Screening and Early Detection and Treatment of Cervical Cancer. According to the guideline, the target population is women older than 21 years old or with sexual intercourse experience for more than three years. The guideline also defined high-risk women as those with several sexual partners, too early sexual debut, HPV infection, low immunity, poor health knowledge and accessibility to healthcare. Depending on diversified geographical socioeconomic status and levels of exposure to the risks of the population, the guideline has recommended three protocols with a different combination of methodologies based on feasibility and a cost-effectiveness evaluation. The three protocols are: *i*) primary screening by liquid-based cytology test + HPV DNA test, an optimal one with best sensitivity and specificity but relevantly high cost, suitable for a screening program in developed regions and/or women with good economic status; *ii*) primary screening by Pap smear cytology test + HPV DNA test, suitable in moderately developed regions; *iii*) primary screening by VIA, a basic one as an alternative in low-resource settings. High-risk women are prioritized in the guideline, with a recommended more intensive frequency of screening and follow-up. Ideally, with such a strengthened HPV screening program, early detection and treatment of cervical cancer can be realized and incidence as well as disease burden can be reduced.

For women living in rural areas, among whom the incidence and the disease burden of cervical cancer is much higher, since 2009 the nationwide pilot campaign of free screening for cervical cancer and breast cancer has been implemented with a strong political commitment from the annual government report and the 12th Five-Year Plan for Healthcare System Reform. On the other hand, due to huge geographical diversity in socioeconomic development, the current screening program has limitations accomplishing its function. Although the national guideline recognized the huge gap in technical capacity and accessibility in different regions, the current recommended region-driven protocols for screening need to be re-examined, because incidence and disease burden in less developed regions tends to be higher and the difference of the protocol implementation may enlarge the gap. In reality, a relevant amount of high-risk cases from patients living in low resource regions may be missed due to low sensitivity of the screening protocols and bad performance of healthcare facilities at the primary level, in which the technical capacity tends to be poor and human resources are lacking. This remains an

Table 2. Strategy of HPV vaccination in some countries

Country/Region (Year introduced)	Protocol of vaccination	Estimated coverage	Impacts, effectiveness and economic evaluation	References
High-income				
The United States (2006)	3 doses targeting females aged 11-12 and catch-up ages 13-26, delivered at facilities	32%	Vaccination for HPV in combination with screening can be a cost-effective health intervention, but it depends on maintaining effectiveness during the ages of peak oncogenic HPV incidence.	Kulasingam <i>et al</i> , 2003; Markowitz <i>et al</i> , 2013
Canada (2007)	3 doses targeting females aged 12 and catch-up ages different by regions, delivered at schools	75% (estimated in the simulation model)	The bivalent vaccine had lower ICER compared to quadrivalent vaccine. It is associated with more reduced cervical cancer morbidity and mortality. Differences in these outcomes depend on the extent of cervical disease prevented by cross-protection and the burden of GW caused by HPV-6/11.	Kohli <i>et al</i> , 2012; Anonychuk <i>et al</i> , 2009
The United Kingdom (2008)	3 doses targeting females and males aged 12-13 and catch-up ages 13-17, delivered at schools	84%	The quadrivalent HPV vaccination programme that includes a catch-up strategy can reduce the incidence of cervical cancer, CIN and genital warts at a cost per QALY ratio within the range typically regarded as cost-effective. Vaccination with screening, compared to screening alone, was associated with an incremental cost-effectiveness ratio of 21,059 pound per QALY.	Dasbash <i>et al</i> , 2008; Kulasingam <i>et al</i> , 2008
France (2007)	3 doses targeting females aged 14 and catch-up ages 15-23, delivered at schools or facilities	80% (estimated in the simulation model)	The incremental cost per QALY for the introduction of HPV vaccination alongside the French cervical cancer screening program was €8,408, suggesting that a quadrivalent HPV vaccine in the current screening program in France is cost-effective.	Bergeron <i>et al</i> , 2008
Germany (2007)	3 doses targeting females aged 12-17, delivered at schools or facilities	-	The quadrivalent HPV vaccination programme of females ages 12 to 17 in Germany is cost-effective with an ICER of 5,525€/QALY. The current vaccination and cervical cancer screening programmes in Germany will substantially reduce the incidence of cervical cancer, CIN and genital warts.	Damm <i>et al</i> , 2009; Schobert <i>et al</i> , 2012
Italy (2008)	3 doses targeting females aged 11 and catch-up ages different by regions, delivered at schools or facilities	56%	The bivalent vaccine would prevent an additional reduction of 7976 abnormal pap smears, 601 CIN1, 1826 CIN2/3 and 295 CC cases compared to the quadrivalent vaccine while 25,848 genital wart cases would be prevented by the quadrivalent vaccine. The additional cost averted with the bivalent vaccine was estimated at €2,385,354 per year.	Capri <i>et al</i> , 2011; Mennini <i>et al</i> , 2009
Netherlands (2010)	3 doses targeting females aged 12 with catch-up ages 13-16, delivered at schools or facilities	50% (estimated in the simulation model)	The bivalent or quadrivalent vaccine reduces the cervical cancer incidence by 221 and 207 /100,000, corresponding to ICERs of €17,600 /QALY and €18,900 /QALY, respectively. The quadrivalent vaccine additionally prevents 4390 cases of genital warts, reducing the ICER to €16,300 /QALY.	Westra <i>et al</i> , 2013
Belgium (2007)	3 doses targeting females aged 12-18 with catch-up ages 13-18, delivered at schools or facilities	82%	The vaccine reduces the lifetime risk of cervical cancer from 0.94% to 0.34%, preventing 362 cases of cervical cancer and 131 related deaths in a cohort of 60,000 girls aged 12 years. The vaccination with current screening is at €10,546 /QALY.	Annemans <i>et al</i> , 2009
Ireland (2010)	3 doses targeting females aged 12-13, delivered at schools or facilities	-	ICER for quadrivalent vaccine would be 25,349 euros/QALY and 30,460 euros/QALY for the bivalent vaccine. At current prices, the bivalent vaccine would need to be 22% cheaper than the quadrivalent vaccine in order to have equivalent cost effectiveness.	Dee <i>et al</i> , 2010
Norway (2009)	3 doses targeting females aged 11-12 with catch-up ages 12-24, delivered at schools	63%	Implementation of a quadrivalent HPV vaccine national program in Norway could reduce the incidence of cervical cancer, cervical intraepithelial neoplasia and genital warts at a cost-effectiveness ratio.	Dasbash <i>et al</i> , 2008; Sander <i>et al</i> , 2012
Switzerland (2008)	3 doses targeting females aged 10-14 with catch-up ages 14-19, delivered at schools or facilities	80% (estimated in the simulation model)	Compared to screening only, adding a quadrivalent vaccine could prevent over lifetime 62% of cervical cancers and related deaths, 19% of Cervical Intraepithelial Neoplasia (CIN 1), 43% of CIN 2, 45% of CIN 3 and 66% of genital warts per cohort. ICER were estimated to be CHF45,008 /life-year and CHF26,005 /QALY.	Szucs <i>et al</i> , 2008
Australia (2007)	3 doses targeting females aged 12-13 with catch-up ages 13-26, delivered at schools	71%	Vaccination with screening compared with screening alone was associated with ICER of \$51,103 /life-year and \$18,735 /QALY.	Kulasingam <i>et al</i> , 2007
Japan (2009)	3 doses targeting females aged 12-16, delivered at schools or facilities	-	Vaccinating a 12-year-old cohort can reduce CC incidence and deaths from CC by 73%, associated with ICER of yen1.8 million per QALY gained. The vaccination program is more cost-effective to increase the coverage of the screening along with the universal administration of HPV vaccine.	Yamamoto <i>et al</i> , 2012; Konno <i>et al</i> , 2010

(continued on next page)

(Table 2. continued)

Country/Region (Year introduced)	Protocol of vaccination	Estimated coverage	Impacts, effectiveness and economic evaluation	References
Singapore (2010)	3 doses targeting females aged 9-26, delivered at facilities	-	Comparing the bivalent to the quadrivalent vaccine, the ICER was \$12,488 per life-year saved. The quadrivalent vaccine dominates to the bivalent vaccine due to the additional QALY effect from reduction in genital warts. The overall outcomes were most sensitive to vaccine cost and coverage.	Lee <i>et al</i> , 2011
Taiwan (2006)	3 doses targeting females aged 9-26, delivered at facilities	-	An additional 768 QALY and 11.6 million new Taiwan dollars costs saved for the bivalent vaccine versus the quadrivalent vaccine after discounting.	Demarteau <i>et al</i> , 2012
Middle- and low- income				
Thailand (2010)	3 doses targeting females aged 9-12, delivered at schools or facilities	80% (estimated in the simulation model)	Pre-adolescent HPV vaccination alone was projected to reduce the lifetime risk of cervical cancer by 55%, which was greater than any strategy of screening alone. Pre-adolescent vaccination and HPV DNA testing five times per lifetime, starting at age 35 years, reduced the lifetime cervical cancer risk by 70%, and had a cost-effectiveness ratio less than Thailand's GDP per capita.	Sharma <i>et al</i> , 2012
Malaysia (2010)	3 doses targeting females before age 13 with catch-up ages 13-18, delivered at schools	-	Vaccination increase life expectancy with better QOL of women when cancer can be avoided. Cost effective strategies will include increasing the screening coverage to 70% or higher. Since feasibility and long term screening adherence is doubtful among Malaysian women, vaccination is a more cost effective strategy against cervical cancers.	Ezat <i>et al</i> , 2010
India (2011)	3 doses targeting females aged 11-12 with catch-up ages 13-26, delivered at facilities	70% (estimated in the simulation model)	If high coverage of pre-adolescent girls with a low-cost HPV vaccine that provides long-term protection is achievable, vaccination followed by screening three times per lifetime is expected to reduce cancer deaths by half, and be cost-effective.	Diaz <i>et al</i> , 2008
Hungary (2010)	3 doses targeting females aged 12 with catch-up ages 12-24, delivered at facilities	-	The ICER of adding bivalent vaccine to the current national cancer screening program was estimated to be 27 588 \$/QALY. By quadrivalent vaccine, the ICER of the routine vaccination targeting females aged 12 and the routine vaccination plus the catch-up group were €9,577 and €10,646 per QALY.	Voko <i>et al</i> , 2011; Dabash <i>et al</i> , 2010
Mexico (2008)	3 doses targeting females aged 9-12 with catch-up ages 12-24, delivered at schools or facilities	67%	The quadrivalent vaccine could reduce the probability of persistent HPV-16/18 infection by at least 60%, resulting in a near-proportional reduction in HPV-16/18-associated invasive cervical cancer and CIN 3. The most effective strategy therein was vaccination of 12-year-olds, plus a temporary 12-24-year-old catch-up program covering both sexes; whereby HPV 6/11/16/18-related cervical cancer, high-grade cervical pre-cancer and genital wart incidence was reduced by 84-98%.	Reynales-Shigematsu <i>et al</i> , 2009; Insinga <i>et al</i> , 2007; CDC, 2011
Peru (2011)	3 doses targeting females aged 10-13, delivered at schools	82%	Enhanced screening in adult women combined with preadolescent vaccination had incremental cost-effectiveness ratios lower than Peru's 2005 per capita GDP and considered to be cost-effective.	Goldie <i>et al</i> , 2012
Brazil (2006)	3 doses targeting females aged 9-12, delivered at schools or facilities	50% (estimated in the simulation model)	Vaccination in addition to the current screening programme is likely to save years of life and, depending on the cost of vaccination, may even save resources.	Vanni <i>et al</i> , 2012

obstacle for implementation and effective coverage of the regular screening program. Accessibility to regular screening is limited for quite a large number of high-risk women due to lack of knowledge and awareness as well, leading to a loss of opportunity for prevention of cancer development. Therefore, the technical capacity at the primary level and in less developed regions urgently needs to be strengthened; moreover, in the long-term, the introduction of the prophylactic vaccines is crucial to supplement the screening program, especially targeting high-risk women.

3.2. Factors potentially affecting the introduction of HPV vaccination

Although it showed great health and socioeconomic impacts in numerous countries, the worldwide introduction of the HPV vaccination program still has a short history with various questions remaining. In China, the government has showed a conservative attitude for approval of the current vaccines and has strictly required clinical trials targeting the Chinese population, even though it met with opposition from some scholars and the mass media. Because political commitment plays a core role in the introduction, solid evidence is necessary to persuade the policy makers. Table 4 summarizes factors potentially affecting the introduction based on results of studies having been implemented in China so far, including epidemiological characteristics, efficacy

Table 3. Major advantages and limitations of the screening methodologies for cervical cancer

Methodology	Advantages	Limitations
Pap smear cytology test	Regular screening tool for more than 50 years in settings where the cytology screening system has been established, with relevantly high sensitivity (50-80%) and specificity (85-90%)	The operation requires well trained personnel, mature laboratory technique, high financial costs, and three or more diagnostic follow-ups and treatment for the positive cases.
Liquid-based cytology test	Similar to Pap smear cytology test, while the operation improved collection efficacy accuracy of samples, with higher sensitivity (85%) and specificity (90%). In settings where the laboratory technical capacity for the cytology screening is weak, samples can be restored and sent to outside.	High financial costs (prolonged screening interval may reduce the costs)
HPV DNA test (HC-II)	It can explore the level of the risks and determine the screening interval with higher sensitivity than liquid-based cytology test for detecting significant precancerous lesions. Moreover, the processing of results can be automated, making the test more objective and requiring less training of personnel. It is especially suitable for large-scale screening in high-risk populations.	High financial costs (prolonged screening interval may reduce the costs)
Visual inspection with acetic acid (VIA)	Simple, less personnel and laboratory requirement, low financial costs, quick result and easy to operate, particularly suitable for low-resource setting with limited technical capacity of the cytology screening	Relatively low sensitivity (50-70%) and specificity (85%)
Colposcopy	As an essential supplementary tool for the early detection of cervical cancer and precancerous lesions, it is conducted for suspicious cases and positive results from the screening test. The combination of colposcopy with HPV DNA test and the cytology tests can further improve sensitivity and specificity.	Facility-based, intensive personnel and technical requirement and not suitable for a large-scale screening

Table 4. Literature review for factors affecting implementation of HPV vaccination program in China

Literature review
<p><u>Epidemiological characteristics of HPV infection</u></p> <ul style="list-style-type: none"> ● In rural Guangdong Province: HPV types 16 and 18 accounted for 28.52% of total infection while types 52 and 58 presented 48.24%. (Chen <i>et al</i>, 2012) ● HPV 16 (76.7%) and HPV 18 (7.8%) were the most common, together accounting for 84.5% of squamous cell carcinoma (SCC), followed by HPV 31 (3.2%), HPV 52 (2.2%), and HPV 58 (2.2%). Positive HPV in SCC did not differ notably by region. The potential impact of vaccines against oncogenic HPV types 16 and 18 is estimated to be high (84.5%) against total SCC. (Chen <i>et al</i>, 2009) ● In Western China: HPV-16 and -58 were the most prevalent types, with prevalence of 37.8% and 21.8%, respectively; HPV-18 and -45 were uncommon types. (Li <i>et al</i>, 2012) ● In Wufeng County: HPV 16, 52, and 58 are common genotypes. (Zhang <i>et al</i>, 2012) ● The most prevalent HPV are types 52 and 58 with positive rate of 42.5% in cervical cancer patients, greater than types 16 and 18 in Shanghai, in southern China the HPV 52- and 58-positive rate is greater than that in northern China. (Lo <i>et al</i>, 2002) ● The most prevalent types found were HPV16 (2.9 %), HPV52 (1.7 %), HPV58 (1.5 %), HPV33 (1 %), and HPV18 (0.8 %). Patterns of HPV prevalence differed by age, geographic region, and cytology findings. (Wu <i>et al</i>, 2013) <p><u>Efficacy, safety and immunogenicity</u></p> <ul style="list-style-type: none"> ● A double-blinded RCT in China showed the quadrivalent vaccine was generally well tolerated, with no vaccine-related serious adverse events. High antibody levels were observed for each of the four HPV types and sero-conversion was > 96%. (Li <i>et al</i>, 2010) <p><u>Cost-effectiveness of cervical cancer prevention and control strategies</u></p> <ul style="list-style-type: none"> ● Per-dose HPV vaccine cost of approximately < \$9-14 would be required for strategies involving vaccination to be cost-effective. Combined screening and vaccination approaches are required to maximize outcomes in rural China. (Canfell <i>et al</i>, 2011) ● Assuming a cost per vaccinated girl of \$25, the cost per DALY averted is \$1,360 in China, reflecting the greater number of girls that need to be vaccinated to prevent a death from cervical cancer in China. Vaccine price has an even greater effect on predicted affordability. (Goldie <i>et al</i>, 2008) ● Making an HPV16, 18 vaccine accessible to 70% of young adolescent girls in 72 countries including China could prevent the future deaths of more than four million women vaccinated over the next decade. Provided the cost per vaccinated girl is less than \$10-\$25, adolescent HPV16,18 vaccination would be cost-effective even in relatively poor countries. (Goldie <i>et al</i>, 2008) <p><u>Knowledge and attitude</u></p> <ul style="list-style-type: none"> ● Only 15.0% of women have ever heard of HPV, and this knowledge differs by rural (9.3%) and metropolitan areas (21.6%) and also by education. Most (84.6%) participants were willing to be vaccinated if HPV vaccine became available to them. (Li <i>et al</i>, 2009) ● Knowledge of HPV among the general female population was low; only 24% had heard of HPV. Less than 20% of healthcare providers recognized sexually naive women as the most appropriate population for HPV vaccination. There was high acceptance of the HPV vaccine for all categories of respondents. Only 6% of women were willing to pay more than US \$300 for the vaccine. (Zhao <i>et al</i>, 2012)

and safety of the vaccines, cost-effectiveness of cervical cancer prevention and control strategies Sexual behavior of the population, and knowledge and attitudes towards the vaccines and HPV-related diseases are also important (46-57).

In China, the prevalence of HPV 16 and 18, which are prevented by the current vaccines, is generally high, while there is a huge geographic diversity of the prevalence of oncogenic HPV types. Other prevalent genotypes include HPV 52 and 58, as well. Such epidemiological trends suggest introduction of the prophylactic vaccines and development of vaccines targeting other prevalent genotypes have potential clinical and social benefits. The clinical trials for the development are ongoing. Based on the current data, the quadrivalent vaccine showed great efficacy in the Chinese population and no vaccine-related serious adverse events were reported so far. Based on a mathematical simulation model, introduction of the vaccines and integrated vaccination and screening program will potentially be very cost-effective in China. Providing the program universally covers the high-risk and high-burden population and the price is low enough to ensure accessibility for the underserved. Like other developing countries, there is a concern about financial costs and affordability, highlighting the need for lowering vaccine prices, cost-efficient mechanisms for delivery of vaccinations to high-risk and high-burden populations, and creative sources of financing. A strong political commitment by the government is essential, because universal coverage is expected to be achieved by injection of public subsidies and adaptation of medical insurance, rather than out-of-pocket payment, particularly for the poor living in rural areas. An interesting finding shown in the previous studies is that although the overall related knowledge is lacking, people have a good willingness to receive the vaccines when they are available in China. Strengthening of health education on the prevention and control of cervical cancer and HPV-related disease is another important issue to improve potential coverage.

4. Prospects for the future

Although the disease burden of cervical cancer is relatively high in China, the two current prophylactic vaccines are currently not available due to ongoing clinical trials among Chinese women and multivalent vaccines that encompass additional oncogenic HPV strains are under development as well. The major strategy for cervical cancer prevention and control is screening of women at reproductive ages and secondary prevention. Time is still needed to introduce the prophylactic vaccines, currently, early detection and treatment by the universal coverage screening program is the core of a comprehensive strategy with region-driven approaches for cervical cancer prevention and

control.

So far, based on lessons from both developed and developing countries, secondary prevention alone cannot fully play a role to reduce the incidence and the disease burden, and neither does the vaccination program. As a preventive tool, both these vaccines prefer girls and women not yet exposed to the risk of infection, *e.g.*, prior to first sexual intercourse, and are not effective for those already infected. Moreover, because immunity to HPV is primarily type specific, protection by the current generation of vaccines with a limited number of HPV types cannot provide complete protection against all oncogenic HPV types. Therefore, the functional screening program is still necessary. Previous economic evaluations indicated that an integrated vaccination and screening program is the most effective tool for great cost-effectiveness and health impact. According to WHO, a HPV vaccination program combined with regular screening in women over age 30 for precancerous lesions followed by adequate treatment are key tools to prevent the 530,000 new cervical cancer cases diagnosed every year (1). Therefore, like other countries in the Asia Pacific area and the world, the short-term goal for cervical cancer control is to identify feasible and effective screening measures, and to find the most effective way to combine vaccination with sustainable screening programs (58,59).

The high prevalence of HPV 16 and 18 in the overall population in China suggests an urgent need to introduce the current vaccines (60). The geographically diversified prevalence of oncogenic HPV types as well as socioeconomic status also highlights the importance of region-driven approaches for cervical cancer prevention and control. With the profound tendency for cervical cancer epidemics and the tremendous task of control, in the long-term the introduction of the vaccines and the development of the new generation vaccines for additional oncogenic HPV types are crucial. Besides the current genotypes protected by the vaccines, cross-reactivity suggests that even better clinical benefits may be achieved by its wide application. Challenges for ensuring the benefits for Chinese women by the HPV vaccines include political commitment of the government, provision of solid evidence for policy making, monitoring of safety and side effects, health education, affordable prices and possible public subsidies for the poor and the vulnerable, and strengthened screening programs particularly for the high-risk population and for primary level healthcare facilities with poor technical capacity.

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Measures to combat H7N9 virus infection in China: Live poultry purchasing habits, poultry handling, and living conditions increase the risk of exposure to contaminated environments

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Summary

From March 31 to May 31, 2013, 132 cases of humans were infected with the H7N9 avian influenza virus, 39 of which resulted in death in China, which sparked global concerns about public health. Fortunately, no new case was reported in China since May 8, which seems like to make it step into a stable stage, and the emergency response to the event launched by Shanghai, Jiangsu, Zhejiang, Anhui, Jiangxi, Shandong, and Hu'nan of China have been terminated currently. However, on July 20 and August 10, two new cases were reported from two provinces - Hebei and Guangdong - where no case was reported during the period of spring of 2013. The emerged two new cases rung an alarm bell, thus, the continued public health response cannot let down its guard. Based on our before studies, we found that live poultry purchasing habits, poultry handling, and living conditions increase the risk of exposure to H7N9 virus contaminated environments in China. Due to the difficulty in changing live poultry purchasing habits and in thoroughly removing or closing live poultry markets in China, we suggest that enhanced regulation of poultry markets would be a more feasible and effective strategy to fight against H7N9 virus infection in China. Moreover, in view of the fact that frequent and inevitable contact between rural residents and poultry where rural residents lived also exists due to poultry handling and living conditions, the enhanced regulations on environmental health are also needed for free-range poultry, especially in rural areas.

Keywords: H7N9 virus, avian influenza, environmental exposure, live poultry market

1. Introduction

On July 20, 2013, a 61 years of age female living in Hebei was laboratory-confirmed with H7N9 virus infection and dead on August 11 (1), besides, on August 10, another 51 years of age female infected with H7N9 virus was also laboratory confirmed in Guangdong and now remains in hospital in a critical condition (2). Both

of the two new cases in China had a history of exposure to live animals in live poultry markets and were from the provinces where no case was reported during the period of spring of 2013, thus causing concerns in public that whether the virus will return in a more dangerous form and may spread to even more provinces in the near future?

From March 31 to May 31, 2013, 132 cases of human were infected with the H7N9 avian influenza virus in China, 39 of which were resulted in death (3,4), which sparked global concerns about public health. Fortunately, no new case was reported with H7N9 virus infection in China since May 8, and the emergency response to the event launched by Shanghai, Jiangsu, Zhejiang, Anhui, Jiangxi, Shandong, and Hu'nan of

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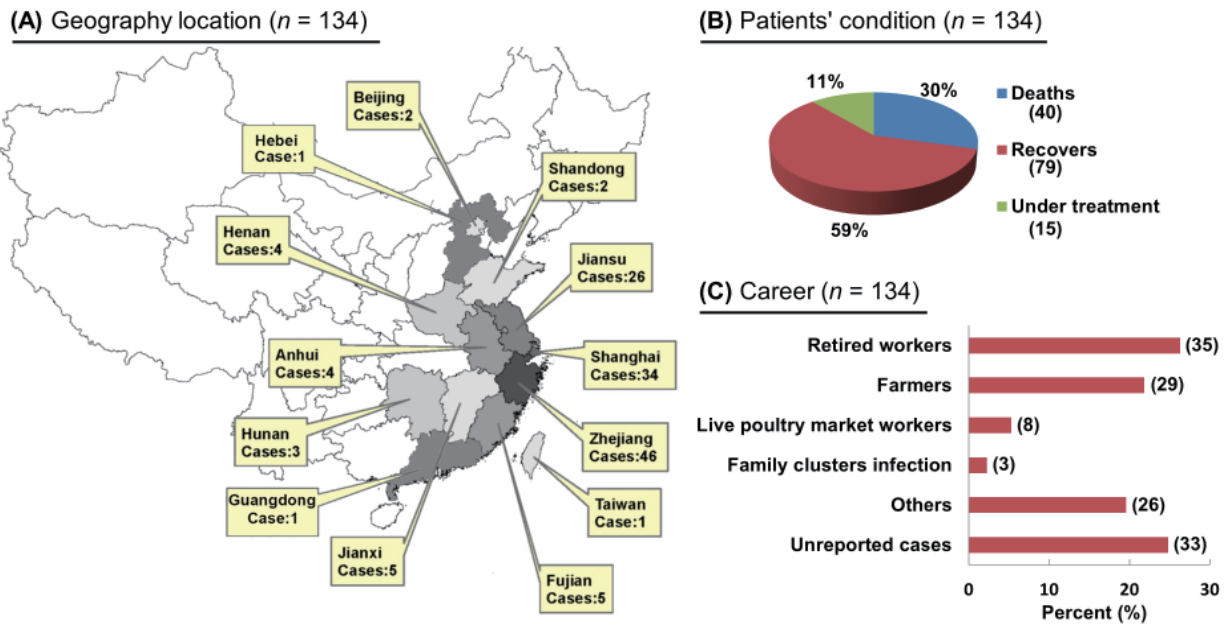


Figure 1. Epidemiologic characteristics of 134 confirmed cases of human infection with H7N9 virus in China. Data collection and analysis based on the information from weekly report on number of confirmed human cases of avian influenza A(H7N9) reported to WHO, the prevention and control work of human infection with avian influenza A(H7N9) by the National Health and Family Planning Commission (NHFPC) of China, and the epidemic reports from 13 provinces with infected patients in China during the period from March 31 to August 10, 2013.

China have been terminated, which seems like to make it step into a stable stage. However, the two new laboratory-confirmed cases emerged on July 20 and August 10 rung an alarm bell, thus, the continued public health response cannot let down its guard.

2. The public health response to combat H7N9 virus infection in China

Until now, 134 cases of H7N9 infection were reported in 43 cities of 13 provinces in China; most were in Zhejiang (46 cases, 7 deaths), Shanghai (34 cases, 13 deaths), and Jiansu (26 cases, 8 deaths) (Figure 1A and 1B). Based on experience fighting SARS, H5N1, and H1N1 outbreaks over the past ten years, China's public health response to H7N9 virus infection is faster and more effective in terms of transparency in reporting, surveillance, screening, and stockpiling of antimicrobials (5,6). Moreover, important information such as viral gene sequence data and patient information were shared in a timely manner within China and with the international community via the WHO under International Health Regulations (7).

After the outbreak of infection with the H7N9 virus, China promptly initiated a survey of the virus' prevalence. In total, 84,444 specimens were collected from 473 live poultry markets, 32 poultry slaughterhouses, 896 poultry farms, 79 wild bird habitats, 36 pig slaughterhouses, and 137 environmental sampling sites. Results were announced by the Ministry of Agriculture of China on April 17 (8). This

results and the update information involving a total of 899,758 specimens announced on May 23 showed that Avian influenza (H7N9) viruses were only isolated in samples from "wet" poultry markets and wild pigeons but not from poultry farms, wild bird habitats, or slaughterhouses (8,9). From April 18-24, a joint team of international and Chinese influenza experts convened by the WHO visited poultry markets and neighborhoods in Shanghai where infections were reported to assess the prevalence of H7N9 in China (10). Results indicated that the risk of H7N9 infection stemmed from live poultry markets and that people could be infected through contact with virus-carrying birds or exposure to contaminated environments.

On April 29, a joint report on the environmental exposure risk at poultry farms from the Ministry of Agriculture and World Organization for Animal Health indicated that there were previously no cases of human or animal H7N9 infections. And virological data have indicated that the new H7N9 virus is a recombination of H7N3, H9N2, and the original H7N9 (11,12). Genetic recombination and key gene mutations have presumably occurred in poultry markets and in transit but not at poultry farms. Thus, there may be no pressing need for large-scale culling at henneries and sampling and inspection of poultry farms may prove adequate.

In this condition, focus on live poultry markets, active public health responses have been implemented in China, including closure of affected markets, halting the sale of fowl, thorough disinfection, poultry culling, and prohibiting the entry of exotic live poultry that

could benefit in preventing more people infected with H7N9 virus. Currently, closing live poultry markets is regarded as a major prevention measure in China.

3. Administration of live poultry markets in management of H7N9 virus infection: thorough closure or enhanced regulation?

It is expected to avoid H7N9 virus infection at large if the live poultry markets could be closed thoroughly in the nationwide. However, it should be noted that nationwide thoroughly closing live poultry markets faces huge challenge in China. Currently, the measures of closing live poultry markets usually start to enforce after finding H7N9 virus positive samples in the affected market, just like the measures that have been implemented in affected areas such as Shanghai, and the measure are implementing in Hebei and Guangdong since the new cases were reported. Actually, as early as 2006, the Chinese Government have issued a regulation to direct live poultry markets to move out of densely populated areas in large and medium-sized cities and encouraging large cities to take the lead in prohibiting the sale and slaughter of live poultry in live poultry markets (13). But until now, a large amount of live poultry are still sold and slaughtered in markets, even in Shanghai, Jiangsu, and Zhejiang where most of the cases of H7N9 infection were reported.

Several studies have indicated that the greater risk factor for infection in urban residents is contact with virus-carrying birds or exposure to contaminated environments in live poultry markets. This risk is closely related to the habit of purchasing freshly slaughtered poultry. Changing live poultry purchasing habits will be difficult and take a long time. Correspondingly, thoroughly removing or closing live poultry markets face challenge and need a long time to solve it. So, due to the difficulty in changing live poultry purchasing habits and in thoroughly removing or closing live poultry markets in China, enhanced regulation of poultry markets will be a more feasible and effective strategy expected to be implemented. Such regulation would be expected to establish quarantine areas at production sites, establish fixed locations for slaughtering and quarantine, implement regular spot checks, implement bio-safety level disinfection, and establish better traceability of frozen poultry.

4. Other environments involving close contact with poultry: poultry handling, and living conditions in rural areas

Beside the environmental exposure risk at live poultry markets and poultry farms, the environments involving close contact with poultry also exist in rural areas. According to available information on 134 laboratory-confirmed cases of H7N9 virus infection, 35 cases

involved urban retirees and 29 involved farmers (Figure 1C); most affected individuals had a history of exposure to live animals, and especially chickens.

In order to ascertain the relationship between poultry handling and avian influenza in rural areas, we conducted a cross-sectional study involving 1,379 participants in Shandong Province, Anhui Province, and the Inner Mongolia Autonomous Region from September 2007 to January 2008 (14). Results showed that the risk of environmental exposure to avian influenza is closely related to Chinese living conditions that influence poultry handling: of 1,379 participants, half had contact with wild birds ("often" and "occasionally"); nearly 50% have a semi-closed or open yard that poultry could pass through, 51% kept poultry in their homes. Some respondents had special chicken coops while others did not. Respondents who did not have special chicken coops instead had poultry that were raised in a common room, an extra room, or a restroom. Poultry feces were often found in the yard, living room, restroom, or kitchen of most homes with poultry and even in those with special coops. Thus, enhanced regulations on environmental health are needed for free-range poultry, especially in rural areas.

5. Conclusion

The emerged two new laboratory-confirmed cases infected with H7N9 virus on July 20 and August 10 rung an alarm bell, thus, the continued public health response cannot let down its guard. Based on our before studies, we found that live poultry purchasing habits increase the risk of exposure to contaminated environments. Due to the difficulty in changing live poultry purchasing habits and in thoroughly removing or closing live poultry markets in China, we suggest that enhanced regulation of poultry markets would be a more feasible and effective strategy to fight against H7N9 virus infection in China. Such regulation would be expected to establish quarantine areas at production sites, establish fixed locations for slaughtering and quarantine, implement regular spot checks, implement bio-safety level disinfection, and establish better traceability of frozen poultry. Moreover, in view of the fact that frequent and inevitable contact between rural residents and poultry where rural residents lived also exists due to poultry handling and living conditions, the enhanced regulations on environmental health are also needed for free-range poultry, especially in rural areas.

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Advances in the study of biomarkers of idiopathic pulmonary fibrosis in Japan

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Summary

Idiopathic pulmonary fibrosis is an intractable disease with a median survival time of 2 to 3 years. Serum levels of Krebs von den Lungen-6 (KL-6), surfactant protein A (SP-A), and surfactant protein D (SP-D) are useful biomarkers for idiopathic pulmonary fibrosis and they are widely used in Japan. Based on clinical use in Japan, a combination of KL-6, SP-A, and SP-D is useful at diagnosing interstitial lung diseases and predicting the prognoses for patients with these diseases. However, the differential diagnosis of idiopathic pulmonary fibrosis from other interstitial lung diseases is still challenging. Several other biomarkers have been identified and are being studied in Japan.

Keywords: Interstitial lung disease, biomarker, KL-6, SP-A, SP-D

1. Introduction

Idiopathic pulmonary fibrosis (IPF) is a specific form of chronic, progressive, fibrosing interstitial pneumonia of unknown etiology (1). Although its exact epidemiology is still unknown, IPF is reported to have no distinct geographical distribution, a greater prevalence in males, and mostly affect individuals who smoke (2). IPF has the characteristic appearance of usual interstitial pneumonia (UIP), which is usually limited to the lungs. It is the most common and severe form of idiopathic interstitial pneumonia (IIP). It is progressive, irreversible, and associated with an extremely poor prognosis. No effective pharmacological therapies have been identified to date. The median survival time for patients with IPF is 2 to 3 years from the time of diagnosis (3).

The prevalence of IPF in Japan is reported to be 11.8 cases per 100,000 population. IPF has been designated as an intractable disease by the Ministry of Health, Labor, and Welfare of Japan, and patients diagnosed with the condition receive full insurance coverage (4). In 1989, Kohno first reported Krebs von den Lungen-6 (KL-6) as a new serum indicator for

IPF (5). Since then, several types of biomarkers were identified and have been studied extensively mostly in Japanese. Several types of biomarkers for IPF have been used clinically in Japan. The current article briefly reviews advances in the study of biomarkers for IPF in Japan.

2. Search strategy

A search of the Medline database was done using a combination of the keywords "idiopathic pulmonary fibrosis" and "biomarker" or a combination of the keywords "idiopathic pulmonary fibrosis" and "marker". All articles related to Japan were reviewed. Searches of the databases of the Japan Science and Technology Agency (J-GLOBAL) and Japan Medical Abstracts Society (JMAS) were also done using a combination of the keyword "idiopathic pulmonary fibrosis" or "idiopathic interstitial pneumonia" and "biomarker." All abstracts were reviewed and the full text of relevant articles was reviewed.

3. Biomarkers used in clinic in Japan

The diagnosis of IPF was once a major challenge to clinicians. According to the American Thoracic Society's (ATS) 2000 guidelines on diagnosis and treatment of IPF (6), the golden standard for diagnosing IPF is by pathological findings. However, a lung biopsy

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is not feasible for many patients suspected of having IPF, so the identification of diagnostic biomarkers for IPF would help both clinicians and patients, particularly in cases where a lung biopsy cannot be obtained. With the development of high-resolution computed tomography (HRCT), the ATS had revised its criteria so that patients with UIP patterns according to HRCT can be diagnosed as having IPF without the need for a lung biopsy (1). This sounds as though the value of biomarkers for IPF has decreased. Because of the potential side effects of many types of therapies for IPF, however, IPF should be carefully be diagnosed, and biomarkers may help in this regard. Biomarkers also have important value in predicting the progression and activity of IPF as well as patient response and prognosis.

3.1. KL-6

KL-6 is the earliest and most intensively studied biomarker of interstitial lung diseases (ILDs) in Japan. KL-6 is a high-molecular-weight glycoprotein classified as a polymorphic epithelial mucin. It is expressed on the surface of various epithelial cells and highly expressed by regenerating type II pneumocytes. It was first investigated as a serum tumor biomarker, but it resulted in a very high rate of false positives in patients with pulmonary fibrosis (7). A later study (5) identified it as a biomarker for ILDs.

KL-6's value in diagnosing patients with ILDs compared to healthy controls has been demonstrated by several studies (8,9). The clinical cut-off value for distinguishing patients with ILDs from healthy subjects and patients with lung diseases other than ILDs had been set at the level of 500 U/mL (10). That said, serum levels of KL-6 were also significantly higher in patients with active pulmonary tuberculosis, with a false positive rate of 28% (11). A study (12) found that serum levels of KL-6 were elevated in 70-100% of patients with ILDs and therefore cannot be used to differentiate IPF from other interstitial pneumonias. Data from a study (13) with a large Japanese sample were used to create Figure 1 to show the different rates at which KL-6 tests positive based on a cut-off value of 500 U/mL; data are from 225 patients with various lung diseases and 200 healthy individuals.

Serum levels of KL-6 are also related to therapeutic efficacy. A study treated 14 Japanese patients with rapidly progressive IPF (14) with high-dose corticosteroid pulse therapy and it followed all of the patients for at least 3 weeks. Patients whose serum levels of KL-6 significantly decreased had a better response and better long-time survival. Several studies (15,16) found that serum levels of KL-6 were related to the extent and activity of IPF.

Serum levels of KL-6 are also related to the prognosis for a patient with IPF. Elevated serum levels

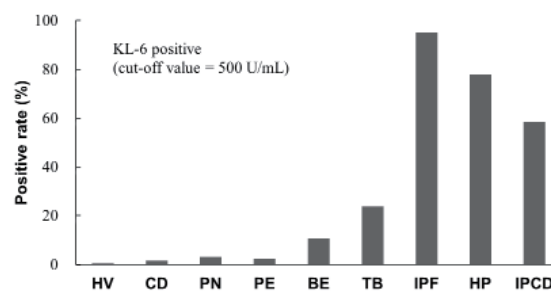


Figure 1. Different rates at which KL-6 tests positive in different lung diseases (Ref. 13). Abbreviations: HV, health volunteer; CD, collagen disease; PN, ordinary pneumonia; PE, pulmonary emphysema; BE, bronchiectasis; TB, pulmonary tuberculosis; HP, hypersensitivity pneumonia; IPCD, interstitial pneumonia with collagen disease.

of KL-6, and especially levels over 1,000 U/mL, upon initial measurement are reported to be related to increased mortality (17). The progression of IPF is significantly faster in patients with KL-6 levels of 1,000 U/mL or higher upon initial measurement than in patients with KL-6 levels below 1,000 U/mL (18).

3.2. Surfactant proteins

Surfactant protein A (SP-A) and surfactant protein D (SP-D) are both water-soluble lung-specific proteins. They are also collectins, a subgroup of the C-type lectin family. They are mainly produced by alveolar epithelial type II pneumocytes and Clara cells within the lungs. Elevated serum levels of SP-A in patients with IPF were first noted in 1993 (19). Serum levels of SP-A in patients with IPF and pulmonary alveolar proteinosis (PAP) were found to be significantly higher than those in healthy volunteers and patients with other non-interstitial lung diseases. A study found that serum levels of SP-A indicate the activity of IPF (20). Another study (12) found that SP-A was better at distinguishing IPF from other ILDs. That study found that serum levels of SP-A in patients with UIP were significantly higher than in patients with non-specific interstitial pneumonia (NSIP).

Elevated serum levels of SP-D were first found to be a possible marker for IPF in 1995 (21). In that study, serum levels of SP-D were significantly higher in patients with IPF, IPCD, and PAP. Serum levels of SP-D were significantly related to the activity of IPF and IPCD. They also closely indicated the severity of PAP.

Serum levels of SP-A and SP-D were also reported to indicate the prognosis for patients with IPF (22). In that study, serum levels of SP-A and SP-D were significantly correlated with the extent of alveolitis. Serum levels of SP-D were also related to the extent of parenchymal collapse on CT and deterioration in pulmonary function. Patients with higher levels of SP-A

Table 1. Rates at which SP-A and SP-D test positive in different lung diseases (Ref. 23)

Rate of testing positive (%)	SP-A	SP-D
70-100	IPF	IPF, IPCD, PAP
50-70	PAP	
30-50	IPCD	
10-30	SAR, PN, DPB, TB, PC	SAR, PN, DPB, TB, PE
0-10	BA, PE	BA

Abbreviations: IPF, idiopathic pulmonary fibrosis; IPCD, interstitial pneumonia with collagen disease; PAP, pulmonary alveolar proteinosis; SAR, pulmonary sarcoidosis; PN, ordinary pneumonia; DPB, diffuse panbronchiolitis; TB, pulmonary tuberculosis; PE, pulmonary emphysema; BA, bronchial asthma.

and SP-D had a worse 3-year survival rate.

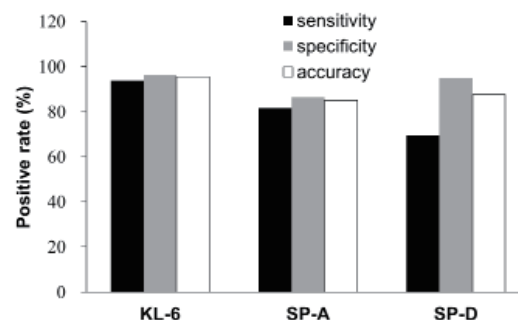
A study (23) of serum levels of SP-A and SP-D in a large Japanese sample set the cut-off value of serum SP-A and SP-D at 48.3 ng/mL and 109 ng/mL, respectively. The study described serum levels of SP-A and SP-D in 87 patients with IPF, 303 patients with HV, and 215 patients with other lung diseases (Table 1).

Many studies have investigated the levels of KL-6, SP-A, and SP-D in bronchoalveolar lavage fluid (BALF). Levels of KL-6 in BALF were reported to be consistent with those in serum (5,8,9), whereas the levels of SP-A and SP-D in BALF were reported to be inversely correlated with levels in serum (19-21). Although the levels of KL-6, SP-A and SP-D in BALF are potential biomarkers for IPF, the inconvenience of obtaining BALF has resulted in their limited use clinically.

KL-6, SP-A, and SP-D have been recognized by Japan's National Health Insurance System as diagnostic markers for ILDs in 1999. KL-6 has been used in clinical practice for more than 10 years in Japan, and KL-6 levels are examined in more than 2,000,000 samples per year in Japan (24). Kits to measure KL-6, SP-A, and SP-D have been successfully developed by reagent manufacturers (Picolumi KL-6: Sanko Junyaku Co., Tokyo, Japan; SP-A test-F: Kokusai Shiyaku Co., Hyogo, Japan; SP-D kit Yamasa; YamasaShoyu Co., Tokyo, Japan). KL-6, SP-A, and SP-D have been included a fourth edition of clinical diagnostic criteria for IPF from a Study Group on IPF of the Ministry of Health, Labor, and Welfare, and all three markers are widely used in many hospitals in Japan.

3.3. The comparative diagnostic value of KL-6, SP-A, and SP-D

A study (25) compared the usefulness of KL-6, SP-A, and SP-D in clinically diagnosing ILDs (Figure 2). KL-6 had the highest specificity and sensitivity for ILDs, but SP-A had the best ability to differentiate IPF from other ILDs (12). To date, KL-6, SP-A, and SP-D have proven value at diagnosing ILDs, but the

**Figure 2. Different diagnostic values of KL-6, SP-A, and SP-D for ILDs (Ref. 25).**

differential diagnosis of ILDs is still difficult. Thus, a combination of KL-6, SP-A, and SP-D is needed to diagnose IPF.

4. Potential biomarkers being studied in Japan

4.1. Vascular endothelial growth factor (VEGF)

Since many interstitial lung diseases are associated with aberrant angiogenesis, the key angiogenesis regulator VEGF has been investigated as a potential biomarker for IPF. A study in the UK (26) found that serum levels of VEGF were not related to IPF. A study in Japan (27) grouped patients depending on the alveolar-arterial difference in oxygen (AaDO₂) and the study found that there were significant differences in the serum levels of VEGF in different groups with a certain AaDO₂ and healthy volunteers. IPF patients with serum levels of VEGF above the median tended to have a worse survival than those with serum levels of VEGF below the median.

4.2. YKL-40

YKL-40, also known as human cartilage glycoprotein 39, is a chitinase-like protein. Its roles in tissue remodeling and fibrosis have been investigated. A study in Japan (28) found that YKL-40 levels in serum and BALF were significantly higher in patients with IPF than in healthy controls. Serum levels of YKL-40 are significantly related to serum levels of KL-6. This means that YKL-40 may be a potential biomarker for IPF.

4.3. Osteopontin

Osteopontin is a type of glycoprotein involved in immune response and tissue repair. Its role in promoting the migration, adhesion, and proliferation of fibroblasts has been demonstrated in a mouse model of bleomycin-induced pulmonary fibrosis (29). A study in 2005 (30) found that levels of osteopontin in plasma were significantly higher in patients with ILDs than in those

with sarcoidosis or in healthy controls. No significant differences in the plasma concentration of osteopontin were noted in patients with IPF or other ILDs.

4.4. Periostin

Periostin is a type of protein known for its roles in the maintenance and development of bones, teeth, and the heart. A recent study (31) found that serum levels of periostin were significantly higher in patients with IPF than in healthy volunteers and patients with cryptogenic organizing pneumonia (COP). Furthermore, periostin levels were inversely correlated with patients' pulmonary function.

4.5. Napsin A

Napsin A, which is expressed in type II pneumocytes and alveolar macrophages, is a type of aspartic proteinase. A recent study (32) found that serum levels of napsin A were significantly higher in patients with IPF than in healthy controls. Serum levels of napsin A are also correlated with the severity of disease.

4.6. Connective tissue growth factor (CCN2)

CCN2 (also known as CTGF) is a type of secreted peptide that mainly serves to produce extracellular matrix and perform other profibrotic activities. Plasma levels of CCN2 were reported to be significantly higher in patients with IPF than in patients with non-IPF IIPs and healthy controls (33). That study also reported a correlation between plasma CCN2 and changes in the 6-month forced vital capacity (FVC).

4.7. S100A9

S100A9 is a type of calcium-binding protein also known as calgranulin B. It is reported to be a useful marker for non-infectious inflammatory diseases. A study of BALF (34) found that S100A9 levels were significantly higher in patients IPF than in patients with other IPDs and healthy volunteers.

4.8. Heat shock protein 47 (HSP47)

HSP47 is a type of collagen-specific molecular chaperone that mainly functions in biosynthesis and the secretion of collagen. Serum levels of HSP47 in cases of acute exacerbation of IPF (AE-IPF) are reported to be significantly higher than those in cases of stable IPF (35). Details are shown in Table 2.

5. Conclusion

This review briefly presented advances in the study of biomarkers for IPF in Japan. Although no specific

Table 2. Details on other biomarkers being studied in Japan

Biomarker	Location	Difference between ILDs and HV	Difference between IPF and other ILDs	Correlation with disease activity or severity	Correlation with KL-6	Correlation with SP-A or SP-D	Number of HV	Number of IPF	Other IIPs	Ref.
VEGF	serum	Yes*	--	HRCT	--	--	43	41	--	(27)
YKL-40	serum	Yes	--	DLco and PaO ₂	Yes	--	41	63	--	(28)
	BALF	Yes	--	--	--	No	16	18	--	(28)
Osteopontin	plasma	Yes	No	PaO ₂	Yes	--	20	8	8	(29,30)
Periostin	serum	Yes	No	Pulmonary function	--	--	--	51	41	(31)
Napsin A	serum	Yes	--	Forced vital capacity	--	--	20	20	--	(32)
CCN2	plasma	Yes	Yes	Forced vital capacity	--	--	101	33	14	(33)
S100A9	BALF	Yes	Yes	--	--	--	23	28	70	(34)
	serum	No	No	Exacerbation vs. stable	--	--	23	28	70	(34)
HSP47	serum	--	--	--	--	--	--	53	--	(35)

Abbreviations: DLco, diffusing capacity for carbon monoxide; PaO₂, partial pressure of oxygen. *Differences were significant for the subgroup with a certain alveolar-arterial difference in oxygen.

biomarkers for IPF had been identified to date, KL-6, SP-A, and SP-D have shown good specificity and sensitivity at diagnosing ILDs. Of the three, KL-6 has the best specificity and sensitivity at diagnosing ILDs while SP-A is specifically able to differentiate IPF from other ILDs. Based on clinical use in Japan, the combination of KL-6, SP-A, and SP-D provides certain value at diagnosing IPF and predicting the prognosis for patients with the condition.

That said, the differentiation of IPF from other ILDs is still challenging. Biomarkers should help to overcome this challenge. Several other biomarkers have also been identified, but further research and evaluation are still needed.

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Factors affecting childhood immunization in Lao People's Democratic Republic: A cross-sectional study from nationwide, population-based, multistage cluster sampling

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Summary

Vaccines are one of the most important achievements in public health, and a major contributor to this success is the Expanded Programme on Immunization (EPI). The effective vaccination series of the EPI should be used by its target population. Various factors influence the utilization of the EPI vaccination series. In Lao People's Democratic Republic (Lao PDR), immunization coverage was lower than the regional average. This study evaluates risk factors affecting immunization underutilization among children five to nine years of age. It is a cross-sectional study from nationwide, population-based, multistage cluster sampling. The children who have received 'standard six' antigens and those who have been partially immunized are compared. In a bivariate analysis, household occupation, maternal age, means of transportation, time to the nearest health facilities, the child's birthplace, birth attended by medical staff, and notification of vaccination date by medical staff, village authority, or megaphone were associated with vaccination status. The final multivariate logistic regression model revealed that maternal age and notification of vaccination date by the village authority increased the odds of full vaccination, while notification of vaccination date by megaphone had decreased those odds. Further detailed qualitative research may be needed to discover how maternal sociodemographic factors influence the utilization of these services. Future research needs to target younger children and must include health care provider factors related to vaccination services.

Keywords: Expanded programme on immunization (EPI), full vaccination, childhood vaccination

1. Introduction

Vaccines have substantially reduced the global burden of infectious diseases. They are considered one of the most important achievements in public health and one of the most cost-effective preventive services for children (1-7). The major contributor to this success is the Expanded Programme on Immunization (EPI) of the World Health Organization (WHO), United Nations Children's Fund (UNICEF) and Global Alliance Vaccine Initiative (GAVI) (6). The EPI was launched in 1974 as a worldwide

alliance of collaborating nations whose goal was to expand immunization services and coverage (6).

The success of EPI does not only depend on effective vaccination series, but also on achieving optimal use by its target population and high immunization coverage (4,8). Pinpointing non-vaccination factors is important for achieving the EPI targets (9,10). The utilization of vaccination services depends on numerous factors such as provision of EPI services including outreach services, accessibility of these services, number of health workers, availability of safe needles, syringes, and cold chain, as well as health education and knowledge and attitude of mothers (5,11,12). Once a child enters the vaccination system, completion of the series is determined by the mother's educational level, employment status, experience with vaccination services, adequate schedule information, immigration status, and overall socioeconomic status (13-15). Various factors

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are important for the initiation and completion of the vaccination series.

The EPI was initiated in Lao People's Democratic Republic (Lao PDR) in 1979 (16). This programme greatly contributed to the progress of basic immunization coverage through the support of international partners and the government of Japan (17). However, immunization coverage in Laos PDR became stagnant after the WHO Western Pacific Region achieved regional polio eradication in 2000 (17). Immunization coverage was lower than regional average: measles immunization coverage among 1-year-olds was 64%, diphtheria tetanus toxoid and pertussis (DTP3) immunization coverage among 1-year-olds was 74%, hepatitis B (HepB) immunization coverage among 1-year-olds was 74%, tuberculosis (Bacille Calmette-Guérin vaccine: BCG) immunization coverage among 1-year-olds was 72% and polio (OPV3) immunization coverage among 1-year-olds was 76% (18,22). The risk factors for non-vaccination need to be studied to achieve optimal use of the vaccination services in Lao PDR. Therefore, the aim of this study is to evaluate risk factors affecting underutilization of childhood immunization among children five to nine years of age.

2. Material and Methods

2.1. Study location and population

Lao PDR is located in Southeast Asia and bordered by five countries: Burma, China, Vietnam, Cambodia, and Thailand. In 2010, its population was approximately 6.2 million and its under-five mortality rate was 54 out of 1,000 live births (19).

2.2. Sampling and sampling frame

This analysis has been done in the sub-population of the nation-wide survey for Hepatitis B sero-prevalence. All 143 districts in Lao PDR were stratified into two strata according to their immunization coverage. Twelve districts were randomly selected from each stratum, and two villages were selected from each district *via* probability of population proportional to size sampling. After randomly selecting 21 children (five to nine years old) and their mothers (15 to 45 years old) from the selected villages, questionnaires were administered.

One thousand and eight pairs of mothers and children were recruited and assessed for eligibility for this study. Forty-three pairs were excluded either because mothers were younger than 15 years of age or older than 45 years of age, or children were younger than five years of age or older than nine years of age. Four hundred and sixty-five pairs were excluded because they did not have vaccination certifications such as yellow cards or mother and child handbooks. Three pairs were excluded because possession of the vaccination certification was unknown.

Among the 497 pairs with the vaccination certification, 284 pairs were excluded because they could not show their certification on the day of the survey. From this, 213 pairs were included in this study.

In Lao PDR, the yellow cards or mother and child handbooks are provided to children who received any vaccination regulated by the Laos PDR Ministry of Health. Vaccination dates were transcribed from those cards, and any vaccination documentation was considered sufficient evidence (20). Each child's immunization record was checked against the EPI immunization schedule recommended by the WHO (21,22). The following categories were used: fully immunized, if the "standard six" antigens – BCG, DTP3 (3 doses), OPV3 (3 doses), and measles vaccines – have been received on the day of interview; and partially immunized, if at least one recommended vaccine dose was not given (23,24). This study compared factors between children who completed their standard vaccinations (full vaccination) and children who had not completed their vaccinations at the time of the survey (partial vaccination).

2.3. Collection of data

The survey, which used a face-to-face, interview-based questionnaire, was conducted from 25 January 2012 to 4 February 2012 by a survey team. A pilot study was conducted prior to the survey to check for clarity and consistency of the questionnaire. The survey team collected demographic information, vaccination status, and other relevant information. Each survey team had two surveyors. These surveyors received two days of training, which consisted of an overview of research methods, interview strategies, and ethical considerations. Before each interview, written consent was obtained from the child's mothers and vaccination dates were transcribed from yellow cards or mother and child handbooks.

2.4. Statistical analysis

The proportions of children who received all the recommended routine vaccination according to the policy of the Expanded Programme of Immunization (EPI) in Lao PDR and its 95% confidence interval (CI) were calculated. Bivariate analysis was performed to assess the relationship between vaccination status and its risk factors. A chi-square test was used for categorical variables and Student's *t* test was used for continuous variables. Crude odds ratios (ORs) and 95% CIs were calculated. Based on the results of the bivariate analysis, all significant factors and ethnicity were entered into the multivariate logistic regression analysis. Ethnicity was added to the final multivariate model because its diversity presented a major challenge in health service delivery due to cultural and language barriers. However, the variable "birthplace of the children" was excluded

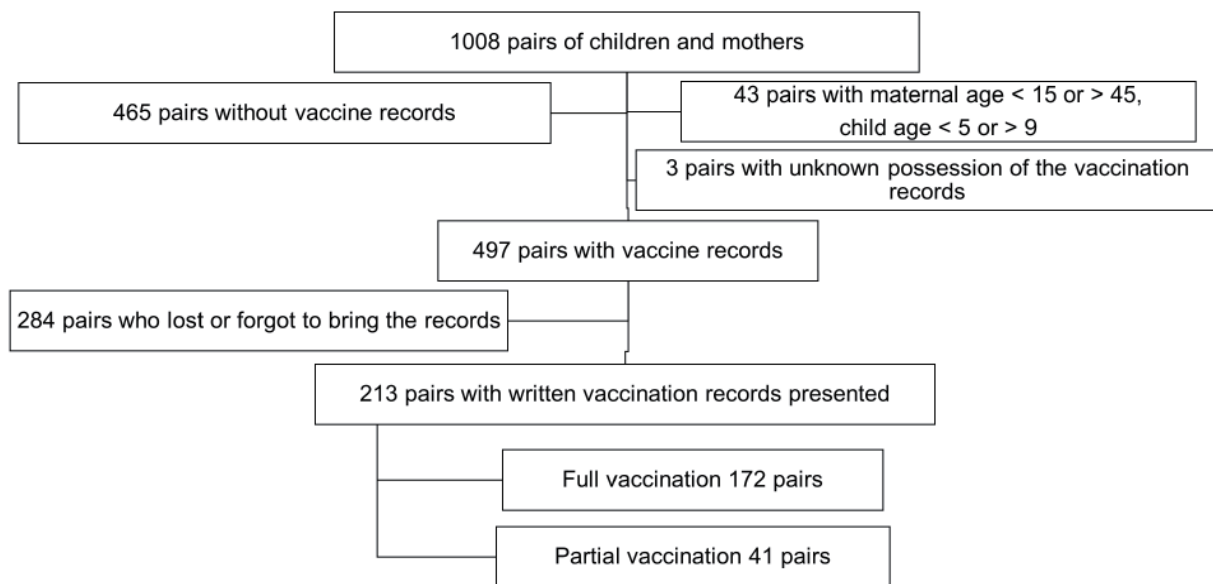


Figure 1. Study profile.

from the final multivariate logistic regression analysis since it showed the highest multicollinearity among the variables in the model. Multicollinearity was detected by using the variance inflation factor (VIF). Stata version 11.0 was used to perform all statistical analyses.

2.5. Ethical considerations

The survey was reviewed and approved by the Ethical Committee of the Ministry of Health, Lao PDR, and the institutional review board of the National Center for Global Health and Medicine, Japan (NCGM-G-001130-00). Access to selected households was granted by the Ministry of Health and the provincial and district government authorities.

3. Results

3.1. Study profile of all subjects

Figure 1 shows the profile of research subjects. 213 pairs were eligible. The case (full vaccination) group had 172 children and the control (partial vaccination) group had 41 children. The proportion of children, aged five to nine, who hold certificates of vaccination (either a yellow card, a mother and child handbook, or both) was 49.3% of the sample, while the proportion of children with an eligible record of vaccination was 21.1% of the sample (data not shown).

3.2. Proportion of children vaccinated by antigen

The statistics of children vaccinated with each antigen are presented in Table 1. All vaccination rates were over 80% except for the HepB birth dose, which

Table 1. Proportion of children having received the various EPI antigens according to vaccination cards

EPI antigen	Total (%), (n = 213)	95%CI
HepB0	19.2	13.9-24.6
BCG	97.2	94.9-99.4
OPV1	97.7	95.6-99.7
OPV2	94.8	91.8-97.8
OPV3	93.4	90.1-96.8
DPT1	98.6	97.0-100
DPT2	95.8	93.1-98.5
DPT3	93.4	90.1-96.8
MV1	83.6	78.6-88.6
Full vaccination***	80.8	75.4-86.1

***Full vaccination – BCG, diphtheria-tetanus pertussis (DTP) (3 doses), polio (3 doses), and measles vaccines

was 19.2%. Among those 19.2%, only five children received the birth dose on the day of birth or the next day. The dates for the HepB birth dose for the rest of the children (80.8%) were not accurate. This low rate and record confusion may be due to the fact that the HepB vaccination for newborns was integrated into the EPI in 2004, which gradually expanded from central hospitals to rural areas. Most (80.8%) of the children were fully immunized between the ages of five and nine, according to the recommended EPI schedule.

3.3. Risk factors of childhood immunization

Among the sociodemographic factors of the family, household occupation, maternal age, means of transportation, and time to the nearest health facilities were associated with full vaccination of the children (Table 2). Birthplace of the children and birth attended by medical staff were associated with vaccination status (Table 3). Notification of vaccination date by

Table 2. Risk factors of childhood immunization (sociodemographic factors)

Items	Full vaccination n = 172, (%)	Partial vaccination n = 41, (%)	Crude OR (95%CI)	p-value
Ethnicity			Reference	
Lowland Lao	102 (59.3)	25 (61.0)		
Midland Lao	61 (35.5)	14 (34.2)	1.07 (0.52-2.21)	0.86
Highland Lao	9 (5.2)	32 (4.9)	1.10 (0.22-5.43)	0.90
Household occupation			Reference	
farmer	126 (73.3)	35 (89.7)		
not farmers	46 (26.7)	4 (10.3)	0.31 (0.10-0.94)	0.03
Mean maternal age (years)	32.7 (CI; 31.8-33.7)	29.3 (CI; 27.5-31.2)		0.00
Maternal education			Reference	
no education	68 (39.5)	16 (39.0)		0.32
primary school	55 (32.0)	19 (46.3)	0.68 (0.32-1.45)	0.25
junior high school	34 (19.8)	4 (9.8)	2.00 (0.62-6.45)	0.43
high school	10 (5.8)	1 (2.4)	2.35 (0.28-19.73)	0.89
university	5 (2.9)	1 (2.4)	1.18 (0.13-10.78)	
Sex of children			Reference	
boy	74 (43.0)	21 (51.2)	0.69 (0.34-1.38)	0.29
girl	97 (56.4)	41 (46.3)		
unknown	1 (0.6)	1 (2.4)		
Mean number of children	3.4 (CI; 3.1-3.7)	3.2 (CI; 2.6-3.9)		0.71
Transportation			Reference	
walk	65 (37.8)	13 (31.7)		
bicycle	3 (1.7)	0 (0)		
bike	44 (25.6)	18 (43.9)	0.49 (0.22-1.10)	0.08
car	27 (15.7)	0 (0)		
tractor	33 (19.2)	5 (12.2)	1.32 (0.43-4.02)	0.63
boat	0 (0)	3 (7.3)		
N/A	0 (0)	2 (4.9)		
Mean time to the nearest health facilities (minutes)	29.5 (CI; 25.0-33.9), n = 152	48.1 (CI; 20.5-75.7), n = 37		0.02
Decision-maker for vaccination			Reference	
Father	36 (20.9)	9 (22.0)		
Mother	129 (75.0)	29 (70.7)	1.11 (0.48-2.56)	0.80
Grandparents	4 (2.3)	3 (7.3)	0.33 (0.06-1.76)	0.20
Village leader	2 (1.2)	0 (0)		
Others	1 (0.6)	0 (0)		

N/A; Non applicable, CI; 95% confidence interval.

Table 3. Risk factors of childhood immunization (delivery history)

Items	Full vaccination n = 172, (%)	Partial vaccination n = 41, (%)	Crude OR (95%CI)	p-value
Birth place of children			Reference	
provincial hospital	38 (22.1)	3 (7.3)		
district hospital	24 (14.0)	9 (16.9)	0.21 (0.05-0.86)	0.03
health centre	4 (2.3)	1 (2.4)	0.32 (0.03-3.80)	0.36
private clinic	2 (1.2)	1 (2.4)	0.16 (0.01-2.29)	0.18
home	72 (41.9)	26 (63.4)	0.22 (0.06-0.77)	0.02
bush near house	31 (18.0)	0 (0)		
N/A	1 (0.6)	1 (2.4)		
Birth attended by the medical staff			Reference	
yes	94 (54.7)	14 (34.2)	2.27 (1.09-4.69)	0.02
no	77 (44.8)	26 (63.4)		
N/A	1 (0.6)	1 (2.4)		
Birth attended by the village health volunteer			Reference	
yes	32 (18.6)	3 (7.3)	2.88 (0.83-10.10)	0.08
no	137 (79.7)	37 (90.2)		
N/A	3 (1.7)	1 (2.4)		
Birth attended by the traditional birth attendant			Reference	
yes	33 (19.2)	9 (22.0)	0.81 (0.35-1.87)	0.62
no	136 (79.1)	30 (73.2)		
N/A	3 (1.7)	2 (4.9)		
Birth attended by the family member			Reference	
yes	75 (43.6)	19 (46.3)	0.79 (0.39-1.60)	0.51
no	95 (55.2)	19 (46.3)		
N/A	2 (1.2)	3 (7.3)		
Birth attended by nobody			Reference	
yes	4 (2.3)	2 (4.8)	0.46 (0.08-2.60)	0.36
no	167 (97.1)	38 (92.7)		
N/A	1 (0.6)	1 (2.4)		

N/A; Non applicable, CI; 95% confidence interval.

Table 4. Risk factors of childhood immunization (source of information of vaccination)

Items	Full vaccination <i>n</i> = 172, (%)	Partial vaccination <i>n</i> = 41, (%)	Crude OR (95%CI)	<i>p</i> -value
Source of information of vaccination (medical staff)				
yes	129 (75.0)	32 (78.1)	0.66 (0.27-1.60)	0.35
no	43 (25.0)	7 (17.1)	Reference	
N/A	0 (0)	2 (4.9)		
Source of information of vaccination (information on the vaccination cards)				
yes	51 (24.8)	9 (22.0)	1.45 (0.64-3.29)	0.37
no	117 (68.0)	30 (73.2)	Reference	
N/A	4 (1.3)	2 (4.9)		
Source of information of vaccination (family member or friends)				
yes	40 (23.3)	14 (34.2)	0.58 (0.27-1.21)	0.14
no	129 (75.0)	26 (63.4)	Reference	
N/A	3 (1.7)	1 (2.4)		
Source of information of vaccination (TV)				
yes	67 (39.0)	13 (31.7)	1.36 (0.66-2.84)	0.40
no	102 (59.3)	27 (65.9)	Reference	
N/A	3 (1.7)	1 (2.4)		
Source of information of vaccination (local authority)				
yes	97 (56.4)	28 (68.3)	0.58 (0.27-1.22)	0.14
no	72 (41.9)	12 (29.3)	Reference	
N/A	3 (1.7)	1 (2.4)		
Source of information of vaccination date (medical staff)				
1. yes	33 (19.2)	18 (43.9)	0.31 (0.14- 0.64)	0.00
2. no	138 (80.2)	23 (56.1)	Reference	
N/A	1 (0.6)	0		
Source of information of vaccination date (village health volunteer)				
yes	62 (36.1)	24 (58.5)	1.16 (0.57-2.36)	0.68
no	108 (62.8)	16 (39.0)	Reference	
N/A	2 (1.2)	1 (2.4)		
Source of information of vaccination date (local authority)				
yes	170 (98.8)	38 (92.7)	6.71 (1.05-42.7)	0.02
no	2 (1.2)	3 (7.3)	Reference	
Source of information of vaccination date (woman's union)				
yes	78 (45.4)	19 (46.3)	0.96 (0.48-1.90)	0.91
no	94 (54.7)	22 (53.7)	Reference	
Source of information of vaccination date (official letter from district governor)				
yes	27 (15.7)	6 (14.6)	1.09 (0.42-2.86)	0.85
no	144 (83.7)	35 (85.4)	Reference	
N/A	1 (0.6)	0		
Source of information of vaccination date (megaphone)				
yes	40 (23.3)	20 (48.8)	0.32 (0.15-0.66)	0.00
no	132 (76.7)	21 (51.2)	Reference	
Source of information of vaccination date (poster)				
yes	27 (15.7)	5 (12.2)	0.57 (0.48-3.74)	0.57
no	145 (84.3)	36 (87.8)	Reference	

N/A; Non applicable, CI; 95% confidence interval.

medical staff, village authority, or megaphone was also associated with full vaccination (Table 4).

Based on the bivariate analysis all variables significantly associated with full vaccination status and ethnicity were included in the multivariate logistic regression model. The results revealed that maternal age and obtaining information of the vaccination date by the village authority increased the odds of full vaccination. However, obtaining information on the vaccination date by megaphone had decreased the odds

of full vaccination (Table 5).

4. Discussion

This study highlights the factors associated with the vaccination status of children aged five to nine in Lao PDR. Multivariate analysis of the risk factors for childhood immunization showed that maternal age and notification of the vaccination date by the village authority were positively associated with full

Table 5. Factors associated with full vaccination of children (logistic regression); occupation of household head-grouped

Factors	Adjusted odds ratio	95%CI	p-value
Sociodemographic factors			
Ethnicity	1.321	0.575-3.036	0.512
Household occupation	0.269	0.067-1.085	0.065
Maternal age	1.087	1.008-1.172	0.031
Transportation to the health facility	1.174	0.852-1.618	0.327
Time to the nearest health facility	0.989	0.977-1.001	0.076
Delivery history			
Birth attended by medical staff	2.617	0.936-7.317	0.067
Source of EPI information			
Source of information of vaccination date by medical staff	0.422	0.150-1.185	0.102
Source of information of vaccination date by local authority	17.430	1.827-166.280	0.013
Source of information of vaccination date by megaphone	0.204	0.065-0.637	0.006

immunization. Notification of the vaccination date by megaphone was negatively associated with full immunization.

4.1. Sociodemographic factors

According to the bivariate analysis, maternal age was associated with vaccination status. Furthermore, there was a linear trend that showed that full vaccination increases with maternal age. However, the relationship between childhood immunization and maternal age varies in the literature. There are some studies that show that maternal age was not associated with childhood immunization (6). One study from Africa showed the influence of younger maternal age on the utilisation of medical care (8).

This study did not show that maternal education was associated with vaccination status, which may be due to the contextual effects of maternal education on children's immunization in Lao PDR (25). Many studies have shown that maternal education or literacy is positively associated with the vaccination status of the children (1,6,8,9,13,25). In addition, some studies showed that the mother's knowledge of specific immunizations was associated with full vaccination (3,5,17). However, in Mali, Koumaré *et al.* reported no difference associated with parental knowledge about EPI diseases and full vaccination (3). In India, Parashar showed that literate women in a village may influence other women's capacity to seek and take advantage of state-provided healthcare, and even children of uneducated mothers may have better health knowledge due to residential or employment proximity to literate women through social influence (25). Further detailed research is needed to determine why maternal education was not associated with childhood immunization in this context.

Maekawa *et al.* showed that distance to health facilities in a rural region of Lao PDR was associated with vaccination status (17). This study focused on time rather than distance to health facilities since access depends on not only the distance but also the accessibility of the road and the availability of

transportation. Therefore, mean time to the nearest health facilities was associated with full vaccination status in the bivariate analysis; however, it was not significant in the logistic regression model, which may be due to the established nationwide outreach service of EPI (26). Further research is needed to explore the association of vaccination status and the actual accessibility to the vaccination sites, combining precise information on distance, time to the vaccination sites, and availability of transportation.

4.2. Delivery history

Birth attended by medical staff was associated with full vaccination of the children; however, it was not significant in the multivariate analysis. According to Maekawa *et al.*, bivariate analysis showed that immunization status was associated with whether mothers obtained information on immunisation before delivery; however, this factor was not significant in the multivariate analysis (17). Maekawa *et al.* also revealed that household visits and receipt of information before delivery influenced the number of fully immunized children. Further analysis revealed that household visits contributed to higher full-vaccination rates, especially among illiterate mothers. Therefore, it may not be sufficient to give information only once before or after the delivery; it might be better to provide information over several household visits.

4.3. Source of EPI information

The notification of the vaccination date by village authority had higher odds of full vaccination. On the other hand, notification of the vaccination date by megaphone was negatively associated with vaccination status. These results show that comprehensive and appropriate information dissemination may be key for vaccinating children against the EPI diseases (3,6).

There are several strengths in this study. This information has been collected through a nationwide survey using multistage cluster sampling. Geographically, the survey dissemination covered a wide portion of

Lao PDR and the ethnic minorities. A sampling frame allowed sampling bias to be a minimum. Only written records were used to avoid inaccuracies in vaccination history. Yellow cards and mother and child handbooks were the sole source of immunization information in this study. Parent's recall was not included in the interview. Rodewald *et al.* stated that the gold standard for measuring vaccination status, other than serological testing to detect immunity, is a parent-linked and provider-validated immunization status measure (2). Parent-linked means that parents name all immunization providers and provider-validated means obtaining the immunization records for each child and combining those data into a single record. Multiple studies have documented the inaccuracy of parents as the sole source of immunization status. The single-provider record check method, used in this study, lies between the parental recall and the parent-linked, provider-verified measure. The validity of this method depends on the frequency and reliability with which immunizations are included in the medical records (2). However, this study has several limitations. Firstly, this analysis has been done in the sub-population of the nation-wide survey for Hepatitis B sero-prevalence and only included the children and mothers who possessed any certifications of vaccination; therefore, the number of children in full and partial vaccination is skewed and no immunization or risk factor information was included from the children without the vaccination certifications. Second, the target of this study was children five to nine years old, which is older than the normal target population for a risk factor-based study. Thirdly, this study did not focus on health care provider factors, which may influence the vaccination status. Utilization of vaccination services is dependent on sociodemographic factors of the target population, as well as the number of health workers and availability of safe needles and syringes (5). Two studies show that children missed immunizations due to the provider's reluctance to vaccinate children while sick or when visiting health care centres for other purposes (9,28). This study did not focus on these provider risk factors. Lastly, this is a cross-sectional study; therefore, it is difficult to generate cause-and-effect relationships of childhood immunization and risk factors due to the study's design.

In conclusion, this study reinforces the importance of appropriate means of notification or provision of information on vaccination services in order to ensure full immunization (13). In the policy and programme level in Lao PDR, it may be necessary to implement appropriate information system on vaccination services at the community level such as mobilizing the village authorities. Older mothers showed higher odds of having fully immunized children, however, other sociodemographic factors such as maternal education was not associated with vaccination status. Further

detailed qualitative research may be needed to discover how this factor influences vaccination services, along with other maternal sociodemographic factors such as education. Future research needs to target younger children to accurately collect information on vaccination records and other sociodemographic factors. Future studies also have to include health care provider factors that affect vaccination services.

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An iTRAQ approach to quantitative proteome analysis of cerebrospinal fluid from patients with tuberculous meningitis

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Summary

To study the cerebrospinal fluid (CSF) protein profiles of tuberculous meningitis (TBM) and discover potential biomarkers for TBM, differential expression of proteins in the CSF of patients with TBM, patients with cryptococcal meningitis, and a control group were compared using isobaric tags for relative and absolute quantitation labelling (iTRAQ) coupled with 2-dimensional liquid chromatography-tandem mass spectrometry (LC-MS). As a result, a total of 208 unique proteins with a molecular weight ranging from 10 KD to 135 KD were identified and quantified in CSF samples from patients with TBM. Of the proteins, 9 were expressed at levels differing 2.0 fold, 6 were up-regulated, and 3 were down-regulated. These proteins appear to be involved in calcium ion binding, lipoprotein metabolism, immune response, and signal conduction. Two differentially expressed proteins were identified using ELISA. The present study represents the successful use of iTRAQ to examine CSF from patients with TBM. The differentially expressed proteins identified may be potential diagnostic biomarkers and provide valuable insight into the underlying mechanisms of TBM. This study also demonstrated that the differential protein profiles of diseases can be quickly determined using iTRAQ-LC-MS, a potential method for quantitative comparative proteomics.

Keywords: Tuberculous meningitis (TBM), iTRAQ, LC-MS, biomarker

1. Introduction

Tuberculous meningitis (TBM) is an infection of the meninges caused by *Mycobacterium tuberculosis* (MTB) and is associated with high mortality rates and severe neurological sequelae. Its prognosis largely depends on early diagnosis as well as timely treatment (1,2). Currently, however, the diagnosis of TBM remains a major challenge due to inadequate diagnostic methods and poor sensitivity and/or specificity of existing markers. Thus, the search for more potential biomarkers to facilitate diagnosis and predict prognosis is of great importance.

Central nervous system (CNS) diseases can cause

changes in protein expression in cerebrospinal fluid (CSF) (3). At present, differential proteomics provides a powerful approach to screen for variations in protein levels and posttranslational modifications associated with disease, culminating in the identification of many potential therapeutic targets and disease-related biomarkers. Numerous researchers have discovered potential biomarkers of disease using proteomics (4-6). Kumar *et al.* performed quantitative protein expression profiling of the brains of patients with TBM and normal individuals using isobaric tags for relative and absolute quantitation labelling (iTRAQ) labelling and LC-MS/MS, and they found several potential diagnostic biomarkers for tuberculous (7). However, the global proteomic profiling of CSF from individuals with TBM has proceeded very slowly, and changes in the CSF proteomes of patients with TBM are rarely reported.

Although 2-dimensional gel electrophoresis is commonly used in conventional proteomics, its drawbacks have been noted; these include low sensitivity,

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taking an excessive amount of time, and failure to detect low-abundance proteins (8). A new labelling method that uses isobaric tags for relative and absolute quantitation labelling (iTRAQ) avoids these drawbacks and offers several advantages (9). iTRAQ reagents have been developed for the relative and/or absolute quantification of complex protein mixtures by labelling tryptic fragments of protein mixtures with reporter tags at the N-terminus and lysine residues (10). Quantification is achieved by comparing reporter ion abundance (*i.e.*, m/z 114, 115, 116, and 117) in the MS/MS spectra. Isotope-coded affinity tag (ICAT) reagents react with cysteine residues, but iTRAQ labels the unblocked N-termini of tryptic peptides and lysines, thus improving protein coverage.

In the present study, iTRAQ and liquid chromatography-tandem mass spectrometry (LC-MS) were used to identify differential proteins in patients with TBM. Nine proteins that were differentially expressed in CSF from patients with TBM were identified. These proteins may be potential diagnostic protein markers for tuberculous. Two of these potential biomarkers were identified using ELISA.

2. Material and Methods

2.1. Subjects

Subjects were patients with TBM and patients with cryptococcal meningitis seen by the Department of Infectious Diseases of Shanghai Public Health Clinical Center from January 2009 to December 2010 and healthy volunteers. *i*) Patients with TBM: twenty patients definitively diagnosed with TBM (having the disease no longer than 2 weeks) consisting of 13 males and 7 females. Their ages ranged from 18 to 52 years, with an average age of 32.6 years. All patients had typical clinical manifestations of TBM and CSF that tested positive for *M. tuberculosis* DNA according to acid-fast bacillus smears, cultures, or PCR results. The patients were yet to receive treatment. *ii*) Patients with cryptococcal meningitis: twenty patients consisting of 12 males and 8 females. Their ages ranged from 18 to 65 years, with an average age of 35.8 years. These patients had typical clinical manifestations, positive smear or culture results, and were yet to receive treatment. *iii*) Normal control group: twenty healthy volunteers consisting of 13 males and 7 females. Their ages ranged from 16 to 55 years, with an average age of 38.4 years. Individuals in this group were free of CNS infections and had normal CSF biochemical and routine test results.

There were no statistically significant differences between the patients and healthy individuals in terms of age and gender ($p > 0.05$). The present study was approved by the Ethics Committee of Shanghai Public Health Clinical Center. All subjects provided written

informed consent for the collection of samples and subsequent analysis.

2.2. Preparation of CSF samples

CSF was extracted *via* a lumbar puncture. Pooled TBM, cryptococcal meningitis, and control samples were created by combining 800 μ L of each individual CSF sample. Proteins were precipitated by mixing 10 volumes of cold acetone containing 10% trichloroacetic acid (TCA) with the CSF samples. The mixtures were then incubated overnight at -20°C , followed by centrifugation at 8,000 rpm/min for 15 min at 4°C . The concentrations of pooled samples were measured using the Bradford method (11).

2.3. iTRAQ labelling and strong cation exchange (SCX) chromatography

About 100 μ g of each of three pooled samples was labelled. iTRAQ labelling was performed in accordance with a kit protocol (Applied Biosystem Inc, Foster City, CA, USA) and a previous report (12). After trypsin digestion, iTRAQ reagents (114, 115, and 117) in 70 μ L ethanol were added separately to each tube and incubated at room temperature for 1 h (114 for patients with TBM, 115 for patients with cryptococcal meningitis, and 117 for the control group). All labelled peptide solutions were then pooled in a new vial and dried using a rotary vacuum concentrator (Christ RVC 2-25, Christ, Germany). Pooled iTRAQ-labelled peptide samples were desalted prior to SCX chromatographic fractionation and LC-MS/MS analysis.

iTRAQ-labelled mixed peptides were fractionated using SCX chromatography in a 20 AD HPLC system (Shimadzu, Kyoto, Japan) using a polysulfoethyl column (2.1 mm \times 100 mm, 5 μ m, 200 \AA , The Nest Group, Southborough, MA, USA). The peptide mixture was reconstituted in Buffer A [10 mM KH_2PO_4 in 25% ACN (Fisher Scientific, Fair Lawn, New Jersey), pH 2.6] and loaded onto the column. The peptides were separated at a flow rate of 200 μ L/min for 60 min with a gradient of 0% to 80% Buffer B (Buffer A containing 350 mM KCl) in Buffer A. Fractions were collected in 3-min increments and vacuum-dried.

2.4. Protein identification by MS

Fractions were vacuum-dried and resuspended in 50 μ L HPLC Buffer A (5% ACN, 0.1% formic acid (TEDIA, Fairfield, USA)), loaded onto the RPLC column (ZORBAX 300 SB-C18 column, 5 μ m, 300 \AA , 0.1 mm \times 15 mm, Microm, Auburn, CA, USA) and analyzed on a QSTAR XL System (Applied Biosystem, Inc.) and a 20AD HPLC system (Shimadzu). MS data were acquired automatically using Analyst QS 1.1 Service Pack 8 software (ABI/MDS SCIEX, Concord, Canada).

Survey scans were acquired from m/z 400 to 1,800, with up to 6 precursors selected for MS/MS from m/z 100 to m/z 2,000.

Ratios of 114.1, 115.1, and 117.1 amu signature mass tags were generated from MS/MS fragmentation of iTRAQ-labelled tryptic peptides and calculated using Protein Pilot (version 3.0, Applied Biosystem, Inc.). MS and MS/MS tolerances were set to 0.2 Da. The International Protein Index (IPI) (version 3.45, HUMAN) database was used to search for peptides. All proteins identified must have $\geq 95\%$ confidence, and the protein confidence threshold cut-off was set at 1.3 (unused) with at least more than two peptides above the 95% confidence level. Fold-changes > 2.0 or < 0.5 were set as cut-off values to designate significant protein expression changes. The subcellular location and function of identified proteins were determined according to the UniProt knowledgebase (Swiss-Prot/UniProtKB) and Gene Ontology (GO) database.

2.5. ELISA analysis of individual samples

Differential proteins were measured in an independent sample set of subjects (25 patients with TBM, 25 patients with cryptococcal meningitis and 25 healthy volunteers) by commercially available ELISA in accordance with the manufacturer's instructions. Data were expressed as mean \pm S.D. and statistical analysis was performed with SPSS 13.0 software (SPSS, Chicago, IL). $p < 0.05$ was considered statistically significant.

3. Results

3.1. CSF proteomes in patients with TBM

After LC-MS/MS analysis, 208 proteins of CSF in TBM were identified and quantified. According to the GO database, most of the proteins identified were located outside cells or in the membrane, cytoplasm, or cytoskeleton, and most function through binding, catalytic activity, enzyme regulator activity, and transporter activity. Most of the proteins identified may be involved in biological regulation, cellular processes, response to stimulus, metabolic process, or the like (Figure 1).

3.2. Identification of differentially expressed proteins

Nine differentially expressed proteins were identified: 6 proteins were up-regulated and 3 proteins were down-regulated. Representative MS/MS spectra for one peptide, identified from the APOB antigen, are shown in Figure 2.

Up-regulated and down-regulated proteins are showed in Tables 1 and 2. These proteins were related to cholesterol metabolism, response to stimulus, calcium ion binding, and prostaglandin-D synthase activity.

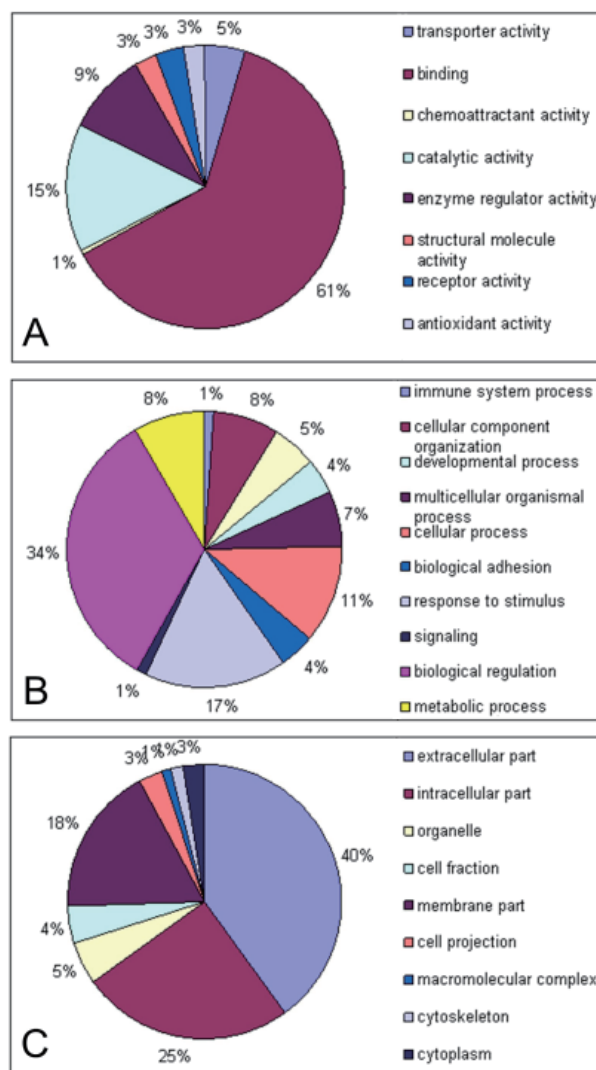


Figure 1. Gene ontology (GO) analysis of 208 proteins in CSF from patients with TBM. (A) molecular function; (B) biological process; (C) cellular component.

3.3. Differential expression of S100A8 and APOB according to ELISA

The levels of S100A8 and APOB expression in patients with TBM, patients with cryptococcal meningitis, and the control group were examined with ELISA to further ascertain differences in protein expression observed using iTRAQ. Differences between patients with TBM and patients with cryptococcal meningitis and the control group ($p < 0.05$) were noted. This result validates the current findings regarding the proteomic probing of differential CSF proteins (Table 3).

4. Discussion

The iTRAQ approach was first described in 2004. The method uses new isobaric reagents that consist of a charged reporter group, a peptide-reactive group, and a neutral balance portion, and the method can identify more proteins than other proteomic methods. It has

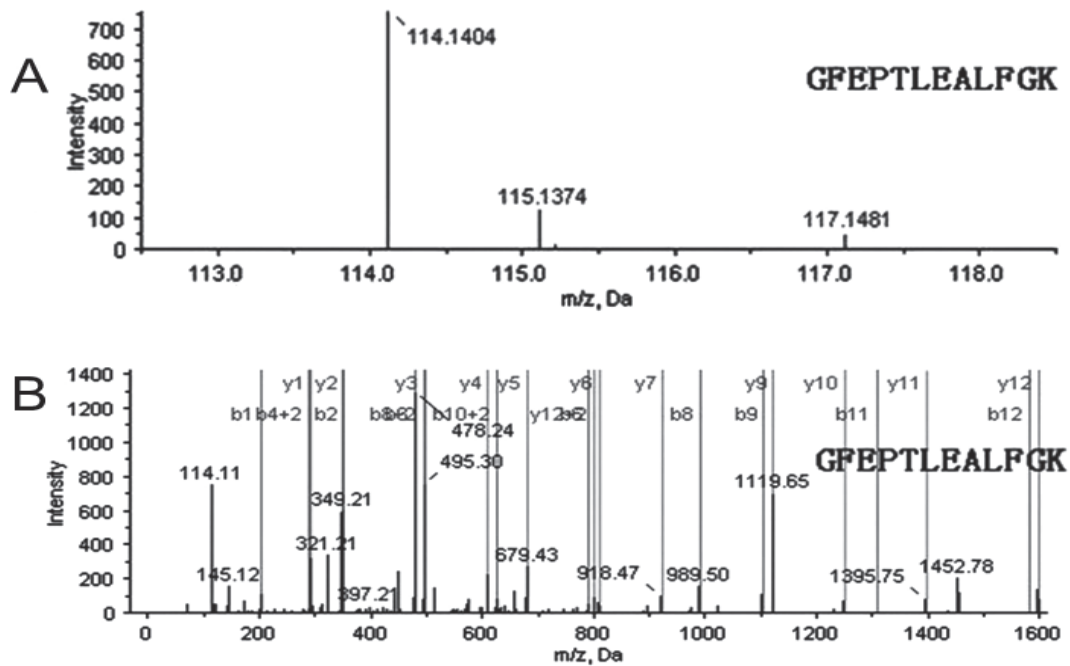


Figure 2. APOB Precursor MS region (GFEPTLEALFGK). The ion assignments were as follows: 114 for patients with TBM, 115 for patients with cryptococcal meningitis, and 117 for the control group. The intensity of reporter ions from precursor peptides indicates protein expression levels (A). The MS/MS spectra revealed peptide sequences with GFEPTLEALFGK, leading to the identification of APOB (B).

Table 1. Up-regulated proteins in CSF from individuals with TBM

IPI	Protein name	MW (D)	Peptide (95%)	Ratio 114:117	Ratio 114:115	Subcellular Location	Molecular function
IPI00847179.1	Apolipoprotein A-I (APOA4)	45372	8	4.4	2.2	secreted	cholesterol transporter activity, antioxidant activity, and copper ion binding
IPI00884926.1	Orosomuroid 1 (ORM1)	23540	20	9.8	2.1	secreted	regulation of immune system processes and protein binding
IPI00292532.6	Cathelicidin antimicrobial peptide (CAMP)	19591	2	2.9	7.4	secreted	antibacterial activity and antibiotic activity
IPI00007047.1	S100 Calcium-binding Protein A8 (S100A8)	10835	4	3.9	9.9	cytoplasm; cell membrane	calcium ion binding and protein binding
IPI00022229.1	Apolipoprotein B-100 (APOB)	515563	7	13.4	3.9	secreted	cholesterol transporter activity, enzyme binding, and low-density lipoprotein receptor binding
IPI00847635.1	a1-antichymotrypsin (SERPINA3)	47652	17	3.1	2.3	secreted	serine-type endopeptidase inhibitor activity and protein binding

CSF: cerebrospinal fluid; TBM: tuberculous meningitis; IPI: international protein index; MW: molecular weight; D: daltons.

Table 2. Down-regulated proteins in CSF from individuals with TBM

IPI	Protein name	MW (D)	Peptide (95%)	Ratio 114:117	Ratio 114:115	Subcellular location	Molecular function
IPI00514285.2	Prostaglandin D2 synthase (PTGDS)	21080	86	0.35	0.42	nucleus membrane	fatty acid binding, transporter activity, and prostaglandin-D synthase activity
IPI00021842.1	Apolipoprotein E (APOE)	36154	31	0.44	0.36	secreted	antioxidant activity, low-density lipoprotein particle receptor binding, and beta-amyloid binding
IPI00607600.1	Calsyntenin-1 (CLSTN1)	108643	6	0.11	0.48	postsynaptic cell membrane	calcium ion binding

CSF: cerebrospinal fluid; TBM: tuberculous meningitis; IPI: international protein index; MW: molecular weight; D: daltons.

Table 3. Differential expression of S100A8 and APOB in CSF

Protein	Patients with TBM (ng/mL)	Patients with cryptococcal meningitis (ng/mL)	Control group (ng/mL)
S100A8	48.9 ± 9.1 ^{*#}	16.9 ± 5.2	13.9 ± 6.7
APOB	480.1 ± 215.2 ^{*#}	166.5 ± 241.9	125.6 ± 131.2

Compared to patients with cryptococcal meningitis, [#] $p < 0.05$; Compared to the control group, ^{*} $p < 0.05$.

been extensively used to identify biomarkers in various disease contexts (13,14).

The present study successfully used iTRAQ-LC-MS/MS to analyze the expression of proteins in CSF from individuals with TBM. A total of 208 proteins with a molecular weight ranging from 10 KD to 135 KD were identified through LC-MS/MS and a database search. GO analysis indicated that these proteins are involved in transport, biological regulation, cellular processes, immune response, metabolic process, and the like. Nine of the 208 proteins were found to be differentially expressed in patients with TBM compared to patients with cryptococcal meningitis and the control group. Six of the 9 proteins were up-regulated and three were down-regulated. Two differential proteins were identified using ELISA. The 9 differentially expressed proteins may be involved in the host response to TBM at the molecular level and may be potential diagnostic biomarkers for TBM.

S100A8 belongs to the S100 protein family and originates from neutrophil granulocytes and macrophages. This protein favors cell growth and restrains proliferation, has the ability to mediate apoptosis and enzymatic activity, promotes a dynamic cytoskeleton, maintains calcium ion stability, and stimulates the development of cancer (15,16). In recent years, S100A8 has been found to be able to regulate inflammatory activity and S100A8 is overexpressed in many infectious diseases (17). In the present study, patients with TBM had higher levels of S100A8 expression in CSF than did patients with cryptococcal meningitis and the control group, which implies that S100A8 may be related to the pathological and biological progression of TBM. CLSTN1 is allocated in the postsynaptic membrane of CNS synapses and is a proteolytically processed protein of the postsynaptic membrane with a calcium-binding cytoplasmic domain (18). It is a dynamic modulator of postsynaptic calcium via extracellular proteolysis and may modulate local Ca transport either beneath the postsynaptic membrane or around intracellular Ca stores. In the present study, CLSTN1 levels in the CSF of patients with TBM were lower than those in patients with cryptococcal meningitis and the control group. Symptoms such as loss of memory and impaired cognition are presumed to be related to the abnormal expression of CLSTN1. However, further research is needed to determine the exact mechanism of this process.

APOB, a component of low-density lipoprotein, can participate in endogenous cholesterol and triacylglycerol transport (19). Upregulation of APOB can cause arterial atherosclerosis, which is responsible for coronary disease and arteriosclerosis (20). APOA4 plays a key role in human cholesterol homeostasis, secretion of chylomicron particles, and antioxidant activity (21). The present study found that levels of APOB and APOA4 expression were higher in patients with TBM than in patients with cryptococcal meningitis and the control group, which suggests that the onset and development of TBM is related to abnormal cholesterol metabolism. Further studies are needed to prove this hypothesis. APOE in human brains is mainly synthesized and secreted by astrocytes, oligodendrocytes, neurons, and activated neuroglia cells (22). It modulates cholesterol and phospholipid homeostasis and adjusts the mobilization and redistribution of cholesterol and phospholipids during remodeling of the neurilemma to mediate the maintenance of synapse flexibility and repair of impaired nerve cells (23). The *APOE* gene can affect memory and facilitate neuron maintenance and repair. Research has shown that *APOE*-knockout mice have learning and memory deficits compared to normal controls (24). In the present study, *APOE* levels in the CSF of patients with TBM were lower than those in patients with cryptococcal meningitis and the control group, presumably because astrocyte cells or neurons that have been attacked exhibit impeded composition and secretion of *APOE*. The lack of *APOE* degrades the repair function of nerve cells. An insufficient supply of acetylcholine leads to nervous system sequelae, such as memory loss and dementia. *APOE* gene polymorphism has been associated with Alzheimer's disease, ischemic stroke, hemorrhagic stroke, multiple sclerosis, Parkinson's disease, schizophrenia, and other diseases (25-27). Further studies are needed to determine the effects of *APOE* gene polymorphism on TBM.

ORM1, SERPINA3, and CAMP are proteins known to be related to acute inflammatory reactions. When MTB attacks meninges, it causes a severe inflammatory reaction. Large amounts of cell factors and mediators of inflammation are released and cause an acute reaction, resulting in an increase in ORM1 and SERPINA3 secretion. Several studies have found that SERPINA3 is overexpressed in CSF from patients with Alzheimer's disease and that it plays an important role in the pathology and pathogenesis of the disease (28,29). Presumably there are common factors involved in the pathogenesis of both TBM and Alzheimer's disease. CAMP plays an important part in natural immune systems. It functions as an antimicrobial, regulating the immune system and the release of inflammation mediators and accelerating revascularization (30,31). The present study found that MTB can cause high levels of CAMP expression, indicating that CAMP serves as an immediate and effective frontline defense.

Further studies are needed to clarify the effect of high levels of ORM1, SERPINA3, and CAMP expression on the onset of TBM and assessment of prognosis.

PTGDS is a member of the prostaglandin synthetases and a transporter of retinoids and thyroxine. PTGDS is expressed in tissues of the blood-brain barrier and secreted into the CSF. Changes in PTGDS expression may influence signal transmission and the activities of vitamin A and thyroxine (32). PTGDS may be involved in blood-brain development and maintenance. It may play an important role in the maturation and maintenance of both the CNS and the male reproductive system. A retrospective analysis of published data that quantified relative amounts of PTGDS in patients with multiple sclerosis, schizophrenia, and Parkinson's disease compared to controls revealed significant dysregulation (33). In the present study, levels of PTGDS in CSF from patients with TBM were lower than those in CSF from patients with cryptococcal meningitis and the control group. This implies that the pathogenesis of TBM may be related to the abnormal expression of PTGDS, although the exact mechanism of this process remains unclear.

In conclusion, 9 differentially expressed proteins associated with TBM were identified using an iTRAQ-based quantitative proteomic approach. Some proteins such as PTGDS, CLSTN1, and S100A8 were found to be expressed at high levels in TBM for the very first time. These promising potential markers warrant further study and evaluation in patients with TBM to determine their clinical utility. The present study also demonstrated that the differential protein profiles of diseases can be quickly determined by iTRAQ-LC-MS/MS, a commercially available method of quantitative comparative proteomics.

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Sub-acute exposure to the herbicide atrazine suppresses cell immune functions in adolescent mice

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Summary

Atrazine (ATR), one of the most widely used herbicides worldwide, has caused a series of toxicological and environmental problems. This study sought to investigate the effects of ATR on the immune system of mice. Four-week-old female C57B I/6 mice were treated with 5, 25, and 125 mg/kg ATR for 28 days. On day 29, blood was collected and the spleen was harvested to detect lymphocyte transformation, natural killer (NK) cell activity, cellular phenotypes, and cytokines. Results indicated that the thymus and spleen weights decreased after ATR treatment, and the spleen was found to be more sensitive to ATR than the thymus. Decreases in lymphocyte transformation and NK cell activity were also observed in mice treated with 25 mg/kg ATR and 125 mg/kg ATR compared to the control group. In addition, there were also alterations of lymphocyte phenotypes in the spleen, and the percentages of CD3⁺ and CD4⁺ cells decreased in mice treated with 25 mg/kg ATR and 125 mg/kg ATR compared to the control group. Moreover, serum interleukin-4 level decreased significantly after treatment with 25 mg/kg and 125 mg/kg ATR, but ATR did not affect the expression of interleukin-2, interferon- γ , and tumor necrosis factor- α . These results suggest that ATR may have induced damage in spleen cells. As ATR is an environmental contaminant, its immunosuppressive action raises concerns that it may potentiate clinical conditions such as tumors, inflammation, and infections. Thus, it needs to be carefully monitored and studied.

Keywords: Atrazine, sub-acute exposure, immunotoxic potential, adolescent mice

1. Introduction

Atrazine (2-chloro-4-ethylamino-6-isopropylamino-1,3,5-triazine, or ATR), is registered for the control of broadleaf weeds and certain grassy weeds. It has been used mostly to prevent weeds from growing in corn, sorghum, coffee, wheat, and sugarcane fields, conifer forests, Christmas tree farms, golf courses, and residential lawns (1). Since it first came on the market in 1958 in Switzerland, ATR has been one of the most commonly used herbicides worldwide (2). As a result, ATR has been found regularly in soil and surface moisture, where it tends to persist for months and migrate great distances

from where it is used. ATR has even been found above the Arctic Circle, albeit at low concentrations (3-5). An estimated 2-3 million people who use groundwater as their primary drinking water are exposed to 0.2 ppb ATR.

The extensive use of ATR and its persistence underline the importance of understanding its general impacts on the environment. Many epidemiological studies have been performed, and some have indicated a possible correlation between atrazine exposure and an increased incidence of neoplastic diseases (6). Other studies have found that atrazine may act as an endocrine-disrupting compound (EDC) with effects on the central nervous system (7), endocrine system (8), and immune system (9).

Laboratory studies have revealed that exposure to environmental contaminants may suppress the amphibian immune system, rendering animals more susceptible to infection (10-13). Fatima found that herbicides at concentrations present in water in Europe caused

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immune suppression in goldfish (11). A meta-analysis reveals that ATR exposure consistently compromises the immune function of fish and amphibians (5). Brodtkin *et al.* reported that in frogs ATR exposure suppressed the thioglycollate-stimulated recruitment of white blood cells to the peritoneal cavity compared to the background levels and also decreased the phagocytic activity of these cells (10). Langerveld *et al.* reported that ATR altered the expression of genes associated with growth and metabolism, proteolysis, fibrinogen complex formation, and immune regulation (14). A significant decrease in the number of intracelomatic cells and a significant decrease in the phagocytic index are observed after ATR exposure. In addition, silver catfish fingerlings exposed to ATR are more susceptible to intracelomatic challenge with the pathogen *Aeromonas hydrophila* (15).

Some studies have examined the immunotoxic potential of acute ATR exposure in adult animals. Exposure to ATR for 14 days can induce a significant decrease in the number of hematopoietic progenitors in murine bone marrow (16), cause a transient suppression of IgM production and T cell proliferation (17), decrease the CD4+ lymphocytes and MHC-II+ cells (18), and increase the mixed lymphocyte reaction (MLR) and cytotoxic T lymphocyte (CTL) response (19). Rajkovic *et al.* reported that ATR increased the number of degranulated mast cells of rats (20). Đikić *et al.* detected apoptotic cells in the thymus and lymph nodes of rats exposed to ATR (21).

Prenatal ATR exposure often causes substantially different effects in adult offspring from those noted in adult animals that are exposed (22,23). In one study, Rooney *et al.* orally administered ATR to Sprague-Dawley rats from day 10 of gestation to day 23 postnatally when pups were weaned. Two weeks after weaning, male pups that were inoculated with sheep red blood cells (SRBC) had a delayed hypersensitivity response that decreased in magnitude and a decreased IgM antibody response (24). Rowe *et al.* used time-release pellets to subcutaneously expose pregnant Balb/c dams to 700 µg/day of ATR for 21 days from day 10 and day 12 of pregnancy. At 3 months of age, male, but not female, mouse pups had increases in T cell proliferation, cytolytic activity, and the number of IgM-secreting B cells in the spleen, but there were no marked changes in the body weight, organ-to-body weight ratio of the spleen, and the numbers of CD8+ T cells, CD4+ T cells, and B220+ B cells in the spleen (25).

Previous studies described the toxic effects of atrazine on the adult and prenatal immune systems of animals. However, the immune system is still developing in the early postnatal period. The developing immune system is more sensitive to xenobiotic exposure than the adult immune system (26-28). A continuous remodeling of the primary and secondary immune organs occurs in mice throughout their lives, *i.e.*, a decline in the relative weight of the thymus,

spleen, and lymph nodes occurs before puberty, while only the thymus continues to involute after puberty (29). One of the proposed windows of vulnerability for the developing rodent immune system occurs postnatally, which is from day 30 to sexual maturity (30). Combined with the limited immunotoxicity data in terms of age and with the possible impact and relevance of such data for immunotoxicological risk assessment, the current study sub-acutely exposed 4-week-old C57B 1/6 mice to ATR for 28 days in order to assess the immunotoxic potential of ATR.

2. Materials and Methods

2.1. Chemicals and reagents

ATR (99% purity) and concanavalin A (ConA) were obtained from Sigma Chemical Company (St Louis, Missouri, USA). RPMI-1640 medium and fetal calf serum (FBS) were purchased from Gibco Laboratory (Gaithersburg, Maryland, USA). Interleukin-2 (IL-2), interleukin-4 (IL-4), interferon γ (IFN- γ), and tumor necrosis factor α (TNF- α) enzyme-linked immunosorbent assay (ELISA) kits were from R&D Systems (Minneapolis, Minnesota, USA). CD3, CD4, and CD8 fluorescence-conjugated antibody were from Ebioscience (San Diego, California, USA). ATR solutions (0.5 mg/mL, 2.5 mg/mL, and 12.5 mg/mL) were prepared by dissolving ATR in corn oil. These solutions were kept at 4°C for a maximum of 1 week.

2.2. Animals and treatment

Male mice housed three or more per cage often fight, and this may cause unpredictable stress responses that affect the immunological parameters being studied. Thus, 4-week-old pathogen-free female C57B 1/6 mice were used in this study. Animals were purchased from the Experimental Animal Center of Norman Bethune Medical College, Jilin University (Changchun, China). This study was conducted in accordance with internationally recognized guidelines and approved by the Animal Research Committee of Norman Bethune College of Medicine, Jilin University. Animals were housed in groups of five in a temperature- and humidity-controlled environment and given access to food and water *ad libitum*. These animals were randomly divided into 4 groups by body weight (ten/group) and were orally given 0, 5, 25, or 125 mg/kg ATR for 28 consecutive days. Blood and the spleen and thymus were collected 24 h after the final round of ATR treatment.

2.3. Body weight and organ weight

Mice were weighed once every 7 days starting on day 1 after ATR treatment. The organs were collected aseptically and weighed on day 29. The relative weights

of organs were calculated as organ weight (mg)/body weight (g).

2.4. Pathological examination

The spleen was collected, fixed in 10% buffered formaldehyde, embedded in paraffin, and then sectioned. The sections were stained with hematoxylin-eosin for histological assessment.

2.5. Preparation of splenocytes

Suspensions of single cells from the spleen were prepared by gently passing the spleen through a nylon mesh filter. Cellular debris was removed with a 400- μ m stainless steel mesh. Red blood cells were lysed with a hypotonic buffered solution and lymphocytes were washed with PBS and re-suspended in RPMI-1640 medium supplemented with 10% FBS, 100 U/mL penicillin, 100 μ g/mL streptomycin, and 2 mM L-glutamine. Viable cells were counted using the trypan blue exclusion method with a hemocytometer.

2.6. Lymphocyte transformation test (LTT)

Freshly isolated splenocytes were suspended in complete RPMI-1640 medium supplemented with 10% FBS. The splenocytes were seeded into 6 wells in a 96-well plate (each well containing 2×10^4 cells). Cells in three wells were treated with PBS and served as controls, while cells in the remaining three wells were treated with 20 μ g/mL ConA. After culturing in an incubator containing 5% CO₂ at 37°C for 48 h, cell counting was done using a 3-(4,5-dimethylthiazol-2-yl) -2,5-diphenyltetrazolium bromide (MTT) assay. The optical density (OD) was quantified using a microplate reader. The following formula was used to calculate the stimulation index (SI): SI = OD of ConA-treated cells/OD of control cells.

2.7. Natural killer (NK) cell cytotoxicity assay

NK cells from mouse spleens were prepared as described above and used as effector cells. Mouse lymphoma YAC-1 cells, which are sensitive to NK cells, were used as target cells. Briefly, 1×10^4 YAC-1 cells and effector cells were separately added to a 96-well plate. After incubation for 48 h, NK cytotoxicity was assessed using the MTT assay described above. Cytotoxic activity was calculated using the following formula: NK cytotoxicity (%) = $[1 - (OD_{E+T} - OD_E)/OD_T \times 100\%]$. (E + T = a mixture of effector cells and target cells, E = effector cells, T = target cells).

2.8. Flow cytometry

For cell phenotype analysis, fluorescence-conjugated antibody was used in accordance with the manufacturer's

instructions. Briefly, splenocytes were re-suspended in 100 μ L of binding buffer followed by addition of 10 μ L of anti-CD3-PE-Cy5, anti-CD4-FITC, or anti-CD8-PE. After incubation for 15 min at 37°C, cells were washed thrice and then subjected to flow cytometry on the FAC Scan (Becton Dickinson Immuno-cytometry, San Jose, CA), in which 1×10^5 cells were counted. Analysis was performed using the CELL Quest software package (BDIS) in list mode and the lymphocyte gate as defined by forward/side scatter characteristics.

2.9. Enzyme-linked immunosorbent assay (ELISA) of IL-2, IL-4, IFN- γ , and TNF- α

After treatment with ATR for 28 days, animals were sacrificed and blood was collected. The serum was obtained by centrifugation at 3,000 g \times for 10 min. The serum levels of murine IL-2, IL-4, IFN- γ , and TNF- α were detected with corresponding ELISA kits in accordance with the manufacturer's instructions.

2.10. Statistical analysis

Data were analyzed using the Statistical Package for Social Sciences (SPSS) version 10. Data were expressed as means \pm standard deviations (S.D.). Comparisons among multiple groups were done using two-way analysis of variance (ANOVA). A value of $p < 0.05$ was considered statistically significant.

3. Results

3.1. General state and body weight

All animals survived to the end of study. There were no overt changes in appearance and/or behaviors and no significant differences in body weight (Figure 1A) after ATR exposure.

3.2. Toxic effects of ATR on the spleen

The mean weights of the spleen and thymus decreased in mice treated with 25 mg/kg ATR and 125 mg/kg ATR ($p < 0.05$) compared to the control group (Figure 1B). Histopathological examination of the spleen revealed degenerative changes. The spleens appeared atrophic in mice treated with 25 mg/kg ATR and 125 mg/kg ATR and were characterized by effacement of germinal centers, diminution of white pulp, and congestion of red pulp (Figure 1C). There were no significant changes in mice treated with 5 mg/kg ATR in comparison to the control group.

3.3. Lymphocyte transformation

An MTT assay was performed to evaluate splenic lymphocyte proliferation in mice treated with different

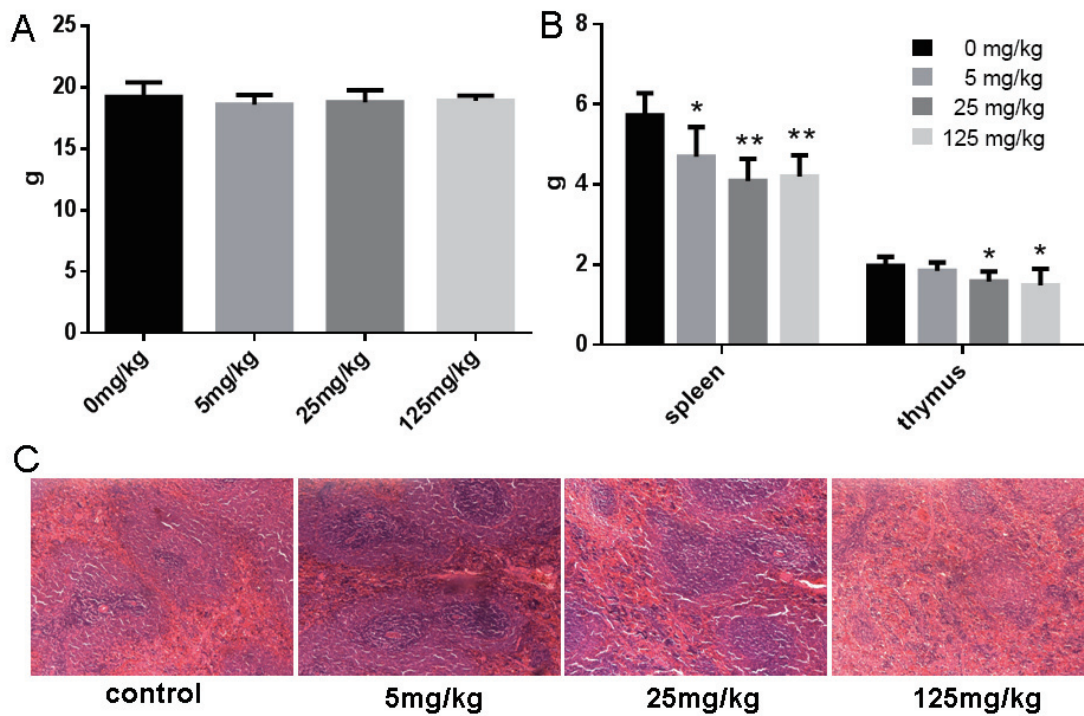


Figure 1. Toxic effects of ATR on the spleen. (A) Body weight was comparable among the 4 groups treated with ATR. (B) The mean weight of the spleen and thymus decreased in mice treated with 25 mg/kg ATR and mice treated with 125 mg/kg ATR (** $p < 0.01$, * $p < 0.05$). (C) Histopathological examinations of the spleen revealed degenerative changes in mice treated with 25 mg/kg ATR and mice treated with 125 mg/kg ATR.

concentrations of ATR. The SI was 2.69 ± 0.21 in mice treated with 25 mg/kg ATR and 1.64 ± 0.67 in mice treated with 125 mg/kg ATR, reflecting a significant decrease compared to the SI in the control group (3.19 ± 0.10) ($p < 0.05$, Figures 2A). The SI did not differ significantly in mice treated with 5 mg/kg ATR (2.66 ± 0.72) and the control group. These results suggest that ATR exposure reduced lymphocyte proliferation in mice.

3.4. NK cell cytotoxicity assay

To determine the impact of ATR on NK cell function, the activity of NK cells was measured *via* a cytotoxicity assay. Results indicated that NK cell cytotoxic activity was $71.16 \pm 2.29\%$ in mice treated with 125 mg/kg ATR, reflecting a significant decrease compared to the control group ($85.08 \pm 6.43\%$) ($p < 0.05$) (Figures 2B). Compared to the control group, there were no significant differences in the NK cell cytotoxic activity in mice treated with 25 mg/kg ATR ($90.54 \pm 3.23\%$) and 5 mg/kg ATR (84.04 ± 12.53).

3.5. Lymphocyte phenotypes in the spleen

Flow cytometry was used to detect the surface markers of lymphocytes. Results revealed that the percentage of CD3⁺ lymphocytes in the spleen decreased in mice treated with 25 mg/kg ATR ($36.22 \pm 2.62\%$) and 125 mg/kg ATR ($34.80 \pm 4.22\%$) compared to the control group ($45.22 \pm 0.55\%$) ($p < 0.05$) (Figure 3A). The percentage of CD4⁺ lymphocytes in the spleen decreased

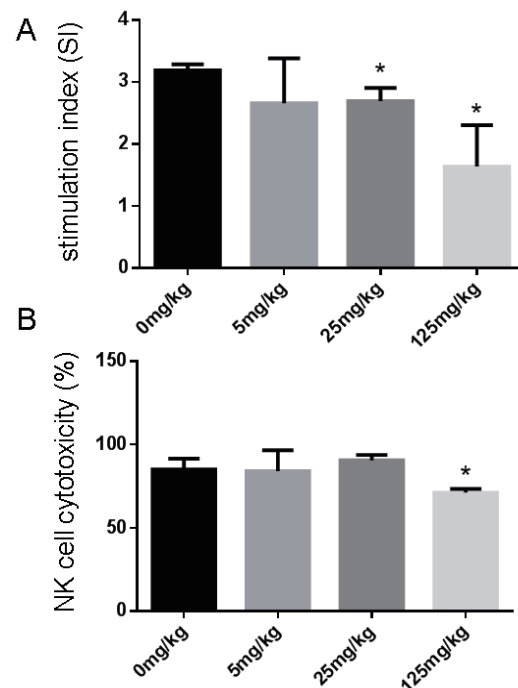


Figure 2. Toxic effects of ATR on splenocyte function. (A) Splenic lymphocyte proliferation decreased significantly ($p < 0.05$) in mice treated with 25 mg/kg ATR and mice treated with 125 mg/kg ATR compared to the control group. (B) NK cell cytotoxic activity decreased significantly ($p < 0.05$) in mice treated with 125 mg/kg ATR compared to the control group ($p < 0.05$).

significantly in mice treated with 25 mg/kg ATR ($8.39 \pm 0.56\%$) and 125 mg/kg ATR ($7.14 \pm 0.88\%$) compared to the control group ($11.08 \pm 0.49\%$) ($p < 0.05$) (Figure 3B).

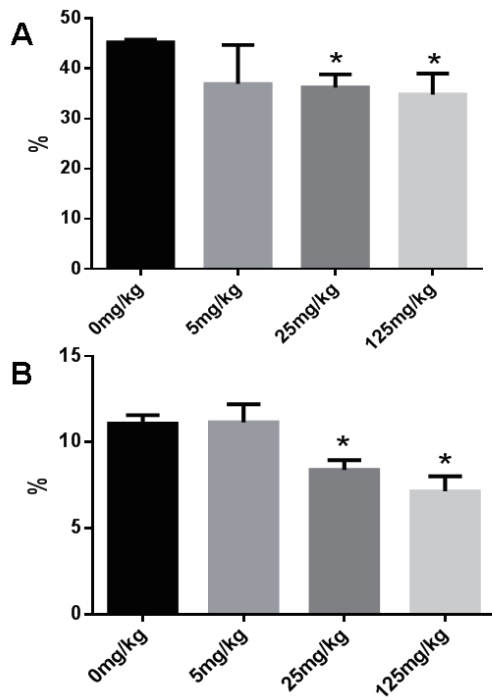


Figure 3. Effect of ATR on percentage of splenic lymphocyte phenotypes. (A) The percentage of CD3+ splenic lymphocytes decreased in mice treated with 25 mg/kg ATR and mice treated with 125 mg/kg ATR compared to the control group ($p < 0.05$). (B) The percentage of CD4+ splenic lymphocytes decreased significantly in mice treated with 25 mg/kg ATR and mice treated with 125 mg/kg ATR compared to the control group ($p < 0.05$). There were no significant changes in the percentage of CD8+ splenic lymphocytes in the four groups treated with ATR ($p > 0.05$).

Neither CD3+ ($36.94 \pm 7.79\%$) nor CD4+ ($11.14 \pm 1.16\%$) lymphocytes changed significantly in mice treated with 5 mg/kg ATR. There were no significant changes in the percentage of CD8+ lymphocytes in the groups treated with ATR and the control group. In fact, ATR treatment reduced the splenic CD4+/CD8+ ratio.

3.6. IL-2, IL-4, IFN- γ , and TNF- α

To further investigate the mechanisms by which ATR inhibits NK activation and decreases T lymphocytes, the serum levels of pro-inflammatory and immunosuppressive cytokines such as IL-2, IL-4, IFN- γ , and TNF- α were detected. As shown in Figure 4A, IL-2 levels in mice treated with ATR were comparable to those in the control group ($p > 0.05$), and the serum level of IL-2 was 330.88 ± 50.40 in mice treated with 0 mg/kg ATR, 307.24 ± 58.35 in mice treated with 5 mg/kg ATR, 322.70 ± 53.38 in mice treated with 25 mg/kg ATR, and 330.42 ± 53.05 in mice treated with 125 mg/kg ATR. As shown in Figure 4B, the IL-4 levels in mice treated with 25 mg/kg ATR (16.93 ± 1.14 ng/mL) and 125 mg/kg ATR (17.25 ± 1.98 ng/mL) had decreased about 15% from levels in the control group (20.45 ± 2.56 ng/mL) ($p < 0.05$), while there were no significant differences in those levels in mice treated with 5 mg/kg ATR (18.03 ± 2.50 ng/mL) and the control group. Figure 4C shows that IFN- γ levels in mice treated with ATR were similar to those in the control group ($p > 0.05$). The serum level of IFN- γ was 131.58 ± 40.95

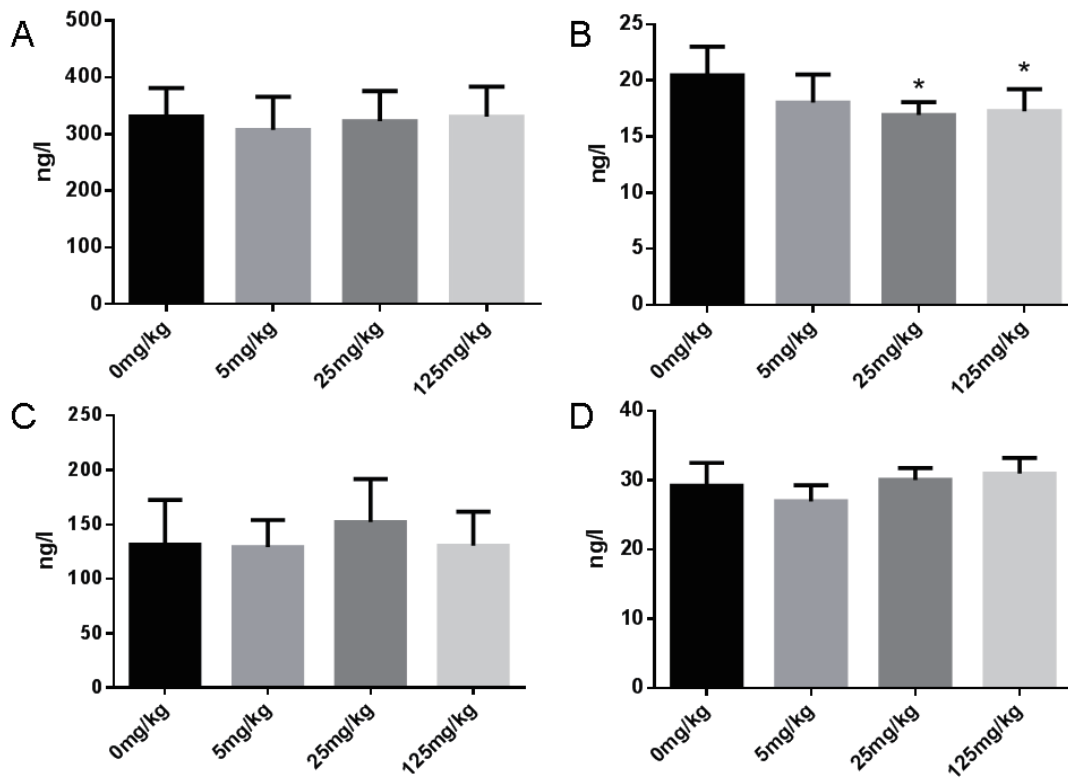


Figure 4. Effects of ATR on serum cytokine levels. (A) Serum IL-2 levels in groups of mice treated with ATR were comparable to those in the control group ($p > 0.05$). (B) Serum IL-4 levels decreased significantly ($p < 0.05$) in mice treated with 25 mg/kg ATR and mice treated with 125 mg/kg ATR. (C) The serum IFN- γ level in groups treated with ATR was similar to that in the control group ($p > 0.05$). (D) There were no significant changes in TNF- α in the four groups treated with ATR ($p > 0.05$).

in mice treated with 0 mg/kg ATR, 129.29 ± 24.77 in mice treated with 5 mg/kg ATR, 152.21 ± 39.59 , in mice treated with 25 mg/kg ATR, and 130.54 ± 31.26 in mice treated with 125 mg/kg ATR. As shown in Figure 4D, there were no significant differences in TNF- α among the four groups treated with ATR ($p > 0.05$). The serum level of TNF- α was 29.45 ± 3.30 in mice treated with 0 mg/kg ATR, 26.99 ± 2.33 in mice treated with 5 mg/kg ATR, 30.10 ± 1.70 in mice treated with 25 mg/kg ATR, and 30.97 ± 2.28 in mice treated with 125 mg/kg ATR.

4. Discussion

The immune system is a complex network of interacting regulatory genes, hormones, and cells that has evolved in multicellular organisms for the purpose of maintaining homeostasis against a dynamic battery of foreign environmental agents and/or pathogens. Environmental pollutants can interfere with the normal operation of the immune system, leading to a broad range of disorders, including endocrine dysfunction, tumorigenesis, increased rates of inflammatory infections, and autoimmune diseases. Widespread use of the herbicide ATR increases the chance for and rate of exposure to other foreign agents that the immune system must contend with, and yet the immunotoxicological potential of ATR has not been studied extensively. The present study sought to examine the effects of ATR on selected immune parameters in juvenile mice. ATR (up to 125 mg/kg per day) was orally administered to 4-week-old female C57BL/6 mice for 28 days. One day after the final round of ATR treatment, mice were sacrificed and their serum and spleens were collected.

During the experiment, overt changes in the appearance and behaviors were not observed in mice treated with ATR, and no significant difference in body weight was noted after ATR exposure. However, relative spleen and thymus weights decreased significantly in mice treated with 25 mg/kg ATR and 125 mg/kg ATR compared to the control group, but there were no significant changes in those weights in mice treated with 5 mg/kg ATR. Since abnormal weights of the spleen and thymus are important indicators used to investigate immunotoxic potential (9), the current results indicate that ATR was likely to be immunotoxic to juvenile mice. The decrease in immune organ weight may be associated with the inhibition of lymphocyte proliferation and/or the increase in lymphocyte death in the spleen and thymus (9). In the current study, the histopathological features of the spleen were detected by light microscopy, and results revealed degenerative changes, compared to the control group, in the spleens of mice treated with 25 mg/kg ATR and 125 mg/kg ATR but not in mice treated with 5 mg/kg ATR. These results suggested that ATR exposure had an adverse impact on the immunity of the spleen in mice. This conclusion is supported by other studies reporting the adverse effects

of ATR on the immune system (18).

T cells play a central role in organizing the immune defense and are practically involved in all types of immune reactions either by orchestrating the type of immune response or as effectors themselves (31). ConA is one of the biologically relevant stimuli applied to activate lymphocytes (32). In order to understand the effects of ATR induced T-cell response, the ability of ConA to stimulate the proliferation of T lymphocytes was investigated 1 day after the final round of ATR treatment. Results indicated that 5 mg/kg ATR induced similar T-cell proliferation in the control group while the SI of lymphocytes decreased significantly in mice treated with 25 mg/kg ATR and 125 mg/kg ATR in comparison to the control group. These findings suggest that ATR exposure for 28 days could lead to the inhibition of T-cell proliferation. NK cells play a central role in the immune defense against tumor development and viral infections. Thus, any agent that interferes with the ability of NK cells to lyse their targets may increase the risk for tumorigenesis and/or viral infections (1). To determine the impact of ATR on NK cell function, the antitumor activity of NK cells was detected *via* a cytotoxicity assay. Results showed the NK cell cytotoxic activity in mice treated with 125 mg/kg ATR was significantly lower than that in the control group, but 5 mg/kg ATR and 25 mg/kg ATR did not significantly affect NK cell cytotoxicity. These results suggest that treatment with 125 mg/kg ATR for 28 days may induce NK cell dysfunction.

A lymphoproliferative assay provides information only about overall proliferative responses without detailing the specific cell subset involved in these responses. Expression of activation molecules prior to proliferation, on the other hand, offers a useful method to predict lymphocyte proliferative activity (33). Exposure to foreign substances induces or stimulates a specific immune system. Some immune responses are directed to specific cells or cytokines that contribute to the immune response (34). To better understand the immune environment after ATR treatment, whether the cell phenotypes obtained in flow cytometry are modulated or whether they provide a true reflection of these cells must be determined.

Lymphocyte subsets are major cellular components of the adaptive immune response (35). CD3 is a signaling component of the TCR (T-cell receptor) complex, almost all T-lymphocytes express CD3, and therefore the CD3+ antibody is used as a genetic marker of T-lymphocytes. In addition, cytotoxic T lymphocytes express CD8 and recognize endogenous antigens by binding to MHC class I molecules; helper T-lymphocytes express CD4 and require the processing and presentation of antigens in association with MHC class II molecules by antigen-presenting cells and helper T-lymphocytes coordinate and assist other immune cells (36). In order to determine if ATR decreased the

yields of CD3+, CD4+, and CD8+ T lymphocytes, flow cytometry was used to detect surface markers of T lymphocyte subsets in this study. Interestingly, results indicated a significant decrease in the number of CD3+ T lymphocytes and a reduction in the ratio of CD4+/CD8+ T lymphocytes in the spleens of both mice treated with 25 mg/kg ATR and mice treated with 125 mg/kg ATR. A possible explanation for this is that ATR may affect T lymphocytes by decreasing CD4+ T lymphocytes mainly by affecting macrophage activation and production of cytokines (37).

CD4+ T lymphocytes can be further subdivided by the patterns of cytokine expression. Th1 CD4+ T lymphocytes, involved in cellular immunity, are known to produce IL-2, IFN- γ , and TNF- α and enhance macrophage function, cellular immunity, and synthesis of opsonizing and complement-fixing antibodies. Th2 CD4+ T lymphocytes involved in humoral responses are known to enhance the immunoglobulin G-mediated response and activation of eosinophils through the actions of IL-4, IL-5, and IL-13 (36). Numerous studies have found that the immune responses in mice and humans are critically influenced by the balance between Th1 and Th2 cytokines (38). Any absence in the cytokine network would cause an imbalance between Th1 and Th2 cytokines and potentially cause inflammation-related diseases. Inflammation-related cytokines have been regarded as either pro-inflammatory or anti-inflammatory according to their contribution to the inflammatory response. Th1 cells express mainly pro-inflammatory cytokines, and Th2 cells primarily secrete anti-inflammatory cytokines. The current results indicated that 25 mg/kg ATR and 125 mg/kg ATR reduced serum IL-4 levels while the serum levels of IFN- γ , IL-2, and TNF- α remained unchanged. IL-4, produced particularly during allergic, cellular, and humoral responses to selected pathogen infections, may modulate other lymphoid cell activities such as regulation of the differentiation of antigen-stimulated T lymphocytes and control of immunoglobulin class switching in B lymphocytes (39). Turnbull *et al.* found that IL-4 could inhibit macrophage cytokine production and restrain macrophage activation (40). The current results demonstrated that ATR at a concentration higher than 25 mg/kg could alter the production of inflammation-related cytokines in serum. As a result, the balance between Th1 and Th2 immune responses might be disrupted.

The immunotoxicity of ATR manifests as a decrease in immune response capacity, including the suppression of immune cell function and atrophy of immune organs (10,18), which were also observed in the present study. The ATR concentrations in this study ranged from 5 mg/kg to 125 mg/kg, which were determined based on findings in previous studies (21,41) and higher than what is expected in nature. However, these amounts ensured that the effects of ATR were noticeable in this

study. In California, potential occupational exposure to ATR was assessed during mixing, loading, and application of ATR on field corn, and the absorbed daily dosage (ADD) of ATR for a mixer/loader/applicator was estimated to be 1.8-6.1 $\mu\text{g}/\text{kg}/\text{day}$ in the event of short-term exposure (15-21 days) (9, 21). However, the ADD of ATR is expected to be higher for commercial applicators and farmers in developing countries due to inappropriate personal protective equipment and unintentional overspraying. In the surface water in the East Liaohe River Basin of Jinlin Province, the average ATR content was 9.7 $\mu\text{g}/\text{L}$ in regions with agricultural plots and 8.854 $\mu\text{g}/\text{L}$ in regions without agricultural plots; ATR content was as high as 18.93 $\mu\text{g}/\text{L}$ in July (42). These data indicate that the daily ATR exposure for ordinary residents in the watershed is about 0.37-0.79 $\mu\text{g}/\text{kg}/\text{day}$, and this value is higher for farmers. ATR is not removed from the body within 24 h and its metabolites can still be detected in the urine 48 h after exposure (9). Thus, some effects may occur after repeated exposures and result in accumulation of substances above a critical threshold.

5. Conclusions

This study found that ATR interferes with immune function by inducing degenerative changes in the spleen, inhibiting the proliferation of T lymphocytes and NK cell cytotoxic activity, decreasing the percentage of CD3+ and CD4+ T lymphocytes, and reducing the serum levels of IL-4. Results indicated that ATR may be associated with inflammation, infections, and other diseases. However, the many molecular mechanisms involved in this process are still unclear. Therefore, further studies are needed to elucidate the exact mechanisms underlying the deleterious effects of ATR on the immune system and to investigate the effects of prolonged exposure to ATR under natural conditions.

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