



Global burden of vaccine-associated multiple sclerosis, 1967–2022: A comprehensive analysis of the international pharmacovigilance database

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Abstract

Vaccine-associated multiple sclerosis (MS) is rare, with insufficient evidence from case reports. Given the scarcity of large-scale data investigating the association between vaccine administration and adverse events, we investigated the global burden of vaccine-associated MS and potential related vaccines from 1967 to 2022. Reports on vaccine-associated MS between 1967 and 2022 were obtained from the World Health Organization International Pharmacovigilance Database (total number of reports = 120 715 116). We evaluated global reports, reporting odds ratio (ROR), and information components (IC) to investigate associations between 19 vaccines and vaccine-associated MS across 156 countries and territories. We identified 8288 reports of vaccine-associated MS among 132 980 cases of all-cause MS. The cumulative number of reports on vaccine-associated MS gradually increased over time, with a substantial increase after 2020, owing to COVID-19 mRNA vaccine-associated MS. Vaccine-associated MS develops more frequently in males and adolescents. Nine vaccines were significantly associated with higher MS reporting, and the highest disproportional associations were observed for hepatitis B vaccines

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(ROR 19.82; IC₀₂₅ 4.18), followed by encephalitis (ROR 7.42; IC₀₂₅ 2.59), hepatitis A (ROR 4.46; IC₀₂₅ 1.95), and papillomavirus vaccines (ROR 4.45; IC₀₂₅ 2.01). Additionally, MS showed a significantly disproportionate signal for COVID-19 mRNA vaccines (ROR 1.55; IC₀₂₅ 0.52). Fatal clinical outcomes were reported in only 0.3% (21/8288) of all cases of vaccine-associated MS. Although various vaccines are potentially associated with increased risk of MS, we should be cautious about the increased risk of MS following vaccination, particularly hepatitis B and COVID-19 mRNA vaccines, and should consider the risk factors associated with vaccine-associated MS.

KEYWORDS

global, vaccine-associated multiple sclerosis, vaccines, World Health Organization

1 | INTRODUCTION

Multiple sclerosis (MS) is the most common chronic inflammatory demyelinating disease of the central nervous system.¹ The neuro-inflammatory demyelination affecting the brain, spinal cord, and optic nerve leads to dysfunction of vision, sensory and motor disturbances, ataxia, and cognitive impairment.¹ These neurologic dysfunctions are reversible and predominantly present in individuals in early adulthood. However, after usually 10–20 years, progressive neurologic disability occurs in many affected individuals, and functional and financial burdens develop.²

The etiology and pathogenesis of MS are complex and remain uncertain, despite decades of research on both genetic and environmental factors.¹ Currently, genetically susceptible individuals and environmental triggers including smoking, obesity, vitamin D deficiency, autoimmune disorders, infections (e.g., Epstein-Barr virus, cytomegalovirus, and varicella-zoster virus), drugs, and alteration of the gut microbiota are known to aid in the development of MS.^{3,4} Previous studies have suggested that immunization with the recombinant hepatitis B vaccine is associated with an increased risk of MS.⁵ Furthermore, some reports suggest an association between influenza and human papillomavirus (HPV) vaccines with incident MS; however, it is important to recognize that some of these studies have been constrained by methodological limitations including limited sample sizes.^{6,7}

Vaccination is an important public health intervention for preventing many life-threatening infectious diseases and ultimately early mortality.⁸ The undeniable benefits of vaccines far outweigh the risks of potential adverse events (AEs).⁹ This has been particularly highlighted during the COVID-19 pandemic, emphasizing the paramount importance of vaccination.^{10–12} However, some AEs of vaccination, including MS, have been reported. As COVID-19 vaccines are urgently approved, MS associated with COVID-19 vaccines should be closely monitored.¹³

To better understand vaccine-associated MS on a global scale, further research is required to determine the burden, long-term

trends, and related factors associated with MS. Using VigiBase, the World Health Organization (WHO) global pharmacovigilance database, we aimed to explore the burden and long-term trends of vaccine-associated MS and its related factors on a global scale.

2 | METHODS

2.1 | Data sources

We conducted a comprehensive analysis of international pharmacovigilance studies focusing on vaccine-associated MS reported in VigiBase, which is the global postmarketing pharmacovigilance database maintained by the Swedish WHO Collaborating Centre for International Drug Monitoring, branded as the Uppsala Monitoring Centre (UMC), and has accumulated individual case safety reports (ICSR).^{14–16} Individual reports include basic health data of the patient, adverse effect data, and drug data. The database consists of ICSR from over 150 countries, involving 25 000 drugs and vaccines, and includes 120 715 116 reports submitted from national pharmacovigilance centers since 1967.¹⁷ Physicians, healthcare professionals, pharmacists, pharmaceutical companies, and patients spontaneously notify the ICSR database, and they are checked for quality of report, regularly reviewed, and analyzed based on predefined criteria.^{14,15} ICSR is coded into preferred terms using the Medical Dictionary for Regulatory Activities (MedDRA).¹⁸ The Institutional Review Boards of Kyung Hee University Medical Center (KHUH 2022-06-042) and the Uppsala Monitoring Centre (WHO Collaborating Center) approved the use of confidential and electronically processed patient data.

2.2 | Selection of cases

The study evaluated vaccine-associated MS between 1967 and 2022, and vaccines were classified into 19 categories, which include (1) anthrax, (2) cholera, (3) diphtheria, tetanus, acellular pertussis, polio,

and *Haemophilus influenzae* type b (DTaP-IPV-Hib), (4) meningococcal, (5) pneumococcal, (6) tuberculosis, (7) typhoid, (8) encephalitis, (9) influenza, (10) hepatitis A, (11) hepatitis B, (12) rotavirus diarrhea, (13) measles, mumps, and rubella (MMR), (14) varicella zoster, (15) HPV, (16) COVID-19 mRNA, (17) adenovirus type-5 (Ad5)-vectored COVID-19, (18) inactivated whole virus COVID-19, and (19) other (brucellosis, plague, typhus, leptospirosis, rabies, yellow fever, smallpox, Ebola, and dengue) vaccines.

Vaccines were categorized using the Level 4 Anatomical Therapeutic Chemical Code in the WHO drug classifications. All the MS cases were identified using MedDRA at the 25.0 preferred terms level (Supporting Information S1: Table 1). According to WHO causality assessment recommendations, vaccines are considered as “suspected” when calculating disproportional association with MS.

2.3 | Data collection

The study collected ICSR data where there was a suspicion of vaccine-associated MS for further description. The ICSR consists of administrative information (reporting region, year, reporter qualification, and reporting type) and patient characteristics (sex and age). The ICSR also included vaccine information (vaccine class and single vaccine suspected), information about AEs such as the time to onset (TTO) of the AEs, nature and severity of the AEs, and fatal outcome. TTO was defined as the number of days between the administration date of the vaccine and the date when MS was diagnosed. All spontaneous reports included at least one vaccine suspected to be involved in the occurrence of AEs following immunization. According to the WHO definition, each AE is characterized as “non-serious” or “serious.”^{14,15} Fatal outcome was defined as fatal or death among serious AE. Physicians who submitted the ICSR determined the severity of the MS. Concomitant AEs were reported with vaccine-associated MS. These concomitant AEs were classified using MedDRA terms, which are listed in Supporting Information S1: Table 2.¹⁹ Given that ICSR was reported voluntarily based on anonymity and did not contradict patient consent, informed consent was not needed in this VigiBase study.

2.4 | Statistical analysis

The VigiBase data set was categorized into two groups (case and noncase), and disproportionality analysis was performed on each vaccine in the data set from 1967 to 2022 (Supporting Information S1: Tables 1 and 2). Disproportionality analysis involves comparison of the proportion of specific AEs reported for a single vaccine with the proportion of the same AEs for a control group of vaccines, which comprises the entire database of all vaccines.^{14,15} The total number of reported AEs for each vaccine group was used as the denominator in these analyses. If the proportion of cases of AEs associated with a specific vaccine is greater than that of patients not experiencing

them, a signal of disproportionality association (signal identification) between the vaccine and AE was identified.^{14,15}

To investigate which vaccines were significantly associated with MS, two common pharmacovigilance measures of disproportionate analysis were used: information components (IC) and reporting odds ratio (ROR). The IC was calculated for case-non-case analysis using a Bayesian confidence propagation neural network, developed and validated by the Uppsala Monitoring Centre as the indicator value for disproportionate reporting.²⁰ The IC has been validated and its effectiveness to identify signals has been confirmed in several studies.^{14,15} The statistical formula for calculating the IC is as follows: $IC = \log_2([N_{\text{observed}}+0.5]/[N_{\text{expected}}+0.5])$. N_{expected} is the number of cases expected for the combination of vaccine and AEs and is calculated as $[N_{\text{vaccine}} \times N_{\text{adverse event}}]/N_{\text{total}}$; N_{observed} refers to the number of case reports for a certain AE associated with a specific vaccine; N_{vaccine} refers to the number of case reports for a specific vaccine regardless of AEs; $N_{\text{adverse event}}$ refers to the number of case reports for a given AE regardless of the vaccines; and N_{total} represents the total number of case reports in the database. $IC_{0.25}$ is the lower limit of the 95% confidence interval (CI) for IC. A positive value of $IC_{0.25}$ is the conventional threshold used to detect statistical signals.^{18,21,22}

We also evaluated the disproportionality association using the ROR, which is a measure of the association derived from the number of AEs and the contingency table of the vaccine, and compared the number of specific AEs occurring with a targeted vaccine to the probability of the same event occurring with all other vaccines in the database. The ROR was calculated using a contingency table of the number of AEs. If the lower 95% CI of the ROR is greater than 1, it was considered a significant association between the vaccine and a certain AE.^{23,24} The vaccine and all reported cases were compared using the unpaired Kruskal–Wallis test for continuous variables and the χ^2 test, or Fisher's exact test, for categorical variables.²⁵ A two-sided-value <0.05 was considered significant. All analyses were performed using SAS version 9.4 (SAS Inc.).^{26–30}

3 | RESULTS

3.1 | Clinical characteristics of vaccine-associated MS

The number of reports for vaccine-associated MS was 9511 out of 141 935 MS reports in the full database. After excluding demographic information missingness vaccine-associated MS was 8288 (Supporting Information S1: Figure 1).

The age of onset of MS was most prevalent between 18 and 64 years, with females representing 72.1% of the cases. Health professionals contributed to over 50.8% of the reported cases of MS. The median time to onset was 2.0 days for MS. Fatal clinical outcomes were reported in only 0.3% (21/8288) of all cases of vaccine-associated MS (Table 1).

TABLE 1 Patient characteristics of reports on vaccine-associated multiple sclerosis ($n = 8288$).

Variables		Number (%)
Region reporting	African Region	7 (0.1)
	Region of the Americas	3746 (45.2)
	South-East Asia Region	1 (0.0)
	European Region	4355 (52.6)
	Eastern Mediterranean Region	7 (0.1)
	Western Pacific Region	172 (2.1)
Reporting year	1967–1979	0 (0.0)
	1980–1989	8 (0.1)
	1990–1999	165 (2.0)
	2000–2009	689 (8.3)
	2010–2019	3146 (38.0)
	2020–2022	4280 (51.6)
	Reporter qualification	Health professional
Nonhealth professional		4076 (49.2)
Studies	Study related	7829 (94.5)
	Nonstudy related	459 (5.5)
Sex	Male	2104 (25.4)
	Female	5974 (72.1)
	Unknown	210 (2.5)
Age	12–17 years	554 (6.7)
	18–64 years	5366 (64.7)
	≥65 years	273 (3.3)
	Unknown	2095 (25.3)
Delay (TTO), days	Median days (IQR)	2.0 (2.0–2.0)
Vaccine class	Anthrax vaccines	39 (0.5)
	Cholera vaccines	5 (0.1)
	DTaP-IPV-Hib	299 (3.6)
	Meningococcal vaccines	67 (0.8)
	Pneumococcal vaccines	41 (0.5)
	Tuberculosis vaccines	6 (0.1)
	Typhoid vaccines	51 (0.6)
	Encephalitis vaccines	162 (2.0)
	Influenza vaccines	360 (4.3)
	Hepatitis A vaccines	296 (3.6)
	Hepatitis B vaccines	2301 (27.8)
	Rotavirus diarrhea vaccines	0 (0.0)
	MMR vaccines	116 (1.4)
	Zoster vaccines	97 (1.2)
	Papillomavirus vaccines	629 (7.6)

TABLE 1 (Continued)

Variables		Number (%)
	COVID-19 mRNA vaccines	3113 (37.6)
	Ad5-vectored COVID-19 vaccines	426 (5.1)
	Inactivated whole-virus COVID-19 vaccines	0 (0.0)
	Others	280 (3.4)
	Outcomes	Alive
	Fatal	21 (0.3)
	Unknown	3233 (39.0)
Single vaccine suspected		8288 (100.0)

Abbreviations: DTaP-IPV-Hib, diphtheria, tetanus toxoids, pertussis, polio, and Hemophilus influenza type b; IQR, interquartile range; MMR, measles, mumps, and rubella; TTO, time to onset; WHO, World Health Organization.

The temporal changes in the reported cases of vaccine-associated MS are illustrated (Figures 1 and 2). The number of vaccine-associated MS cases gradually increased over time. The proportion of vaccine-associated MS cases among all vaccine-related MS cases also increased. There was a sharp increase in the number and proportion of vaccine-associated MS cases since 2020. European regions had the highest number of reported cases of vaccine-associated MS, followed by America.

3.2 | Cumulative report analysis of vaccine-associated MS

The cumulative number of vaccine-associated MS cases evaluated for different vaccines is shown in Figure 3. Vaccine-associated MS has been reported for 17 vaccines. Before 2020, hepatitis B vaccines had the highest cumulative counts of MS, whereas after 2020, COVID-19 mRNA vaccines were most commonly reported for MS, followed by hepatitis B, HPV, influenza, DTaP-IPV-Hib, and hepatitis A vaccines.

3.3 | Disproportionality analysis of vaccine-associated MS

Nine out of 17 vaccines were significantly associated with MS. Among them, the highest disproportional association with MS was hepatitis B vaccines (ROR 19.82, 95% CI 19.02–20.67; $IC_{0.25}$ 4.18). In the age-specific subgroup analysis, disproportionate signals of hepatitis B vaccine-associated MS were significant in individuals aged 12–64 years (Table 2). A significant disproportionality in MS reporting was also found for encephalitis vaccines (ROR 7.42, 95% CI 6.38–8.66; $IC_{0.25}$ 2.59), hepatitis A vaccines (ROR 4.46, 95% CI

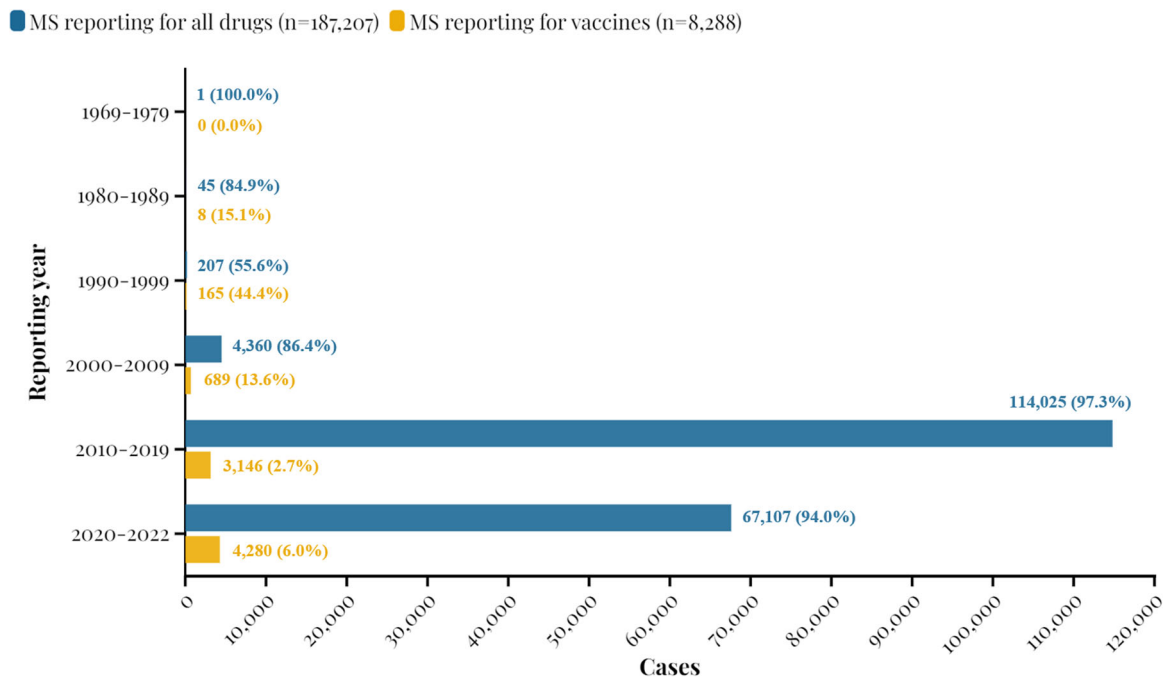


FIGURE 1 Temporal changes of reported cases of vaccine-associated multiple sclerosis. Absolute reported counts of multiple sclerosis according to the reporting year. The proportions of relevant adverse reactions among all drug- or vaccine-related adverse reactions are indicated.

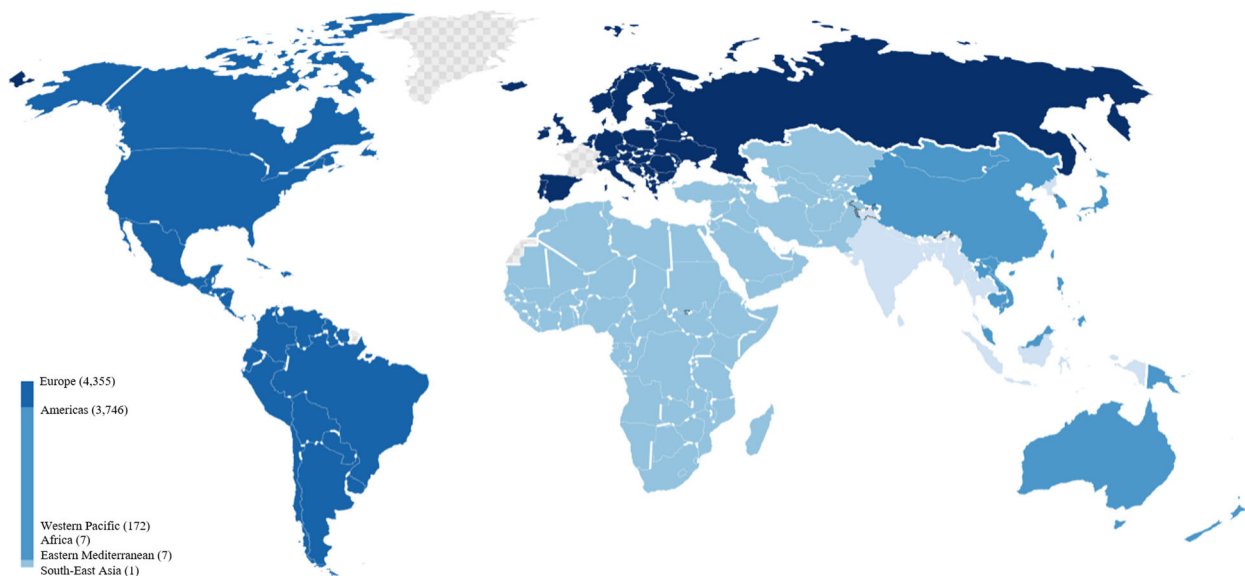


FIGURE 2 World map of reported cases of vaccine-associated multiple sclerosis by continent. Global reported counts based on the continent.

3.98–5.00; IC_{025} 1.95), HPV vaccines (ROR 4.45, 95% CI 4.11–4.81; IC_{025} 2.01), anthrax vaccines (ROR 3.58, 95% CI 2.61–4.90; IC_{025} 1.26), typhoid vaccines (ROR 2.80, 95% CI 2.13–3.68; IC_{025} 0.99), COVID-19 mRNA vaccines (ROR 1.55, 95% CI 1.49–1.60; IC_{025} 0.52), other vaccines (ROR 1.42, 95% CI 1.26–1.60; IC_{025} 0.31), and MMR vaccines (ROR 1.33, 95% CI 1.11–1.60; IC_{025} 0.11).

From an age-specific perspective, the 12–17 years age group had the highest risk of developing vaccine-associated MS (IC_{025} 7.33). Regarding

individual vaccines, age-specific top disproportionality was found in DTaP-IPV-Hib, encephalitis, hepatitis A, and HPV vaccines for 12–64 years, anthrax and typhoid vaccines for 18–64 years, and COVID-19 mRNA and Ad5-vectored COVID-19 vaccine for ≥ 65 years (Table 2). Concurrent AEs in vaccines with significantly disproportionate signals of vaccine-associated MS showed similar patterns. Neurological events were the most frequently reported concurrent manifestations, followed by muscular and psychiatric events (Supporting Information S1: Table 3).

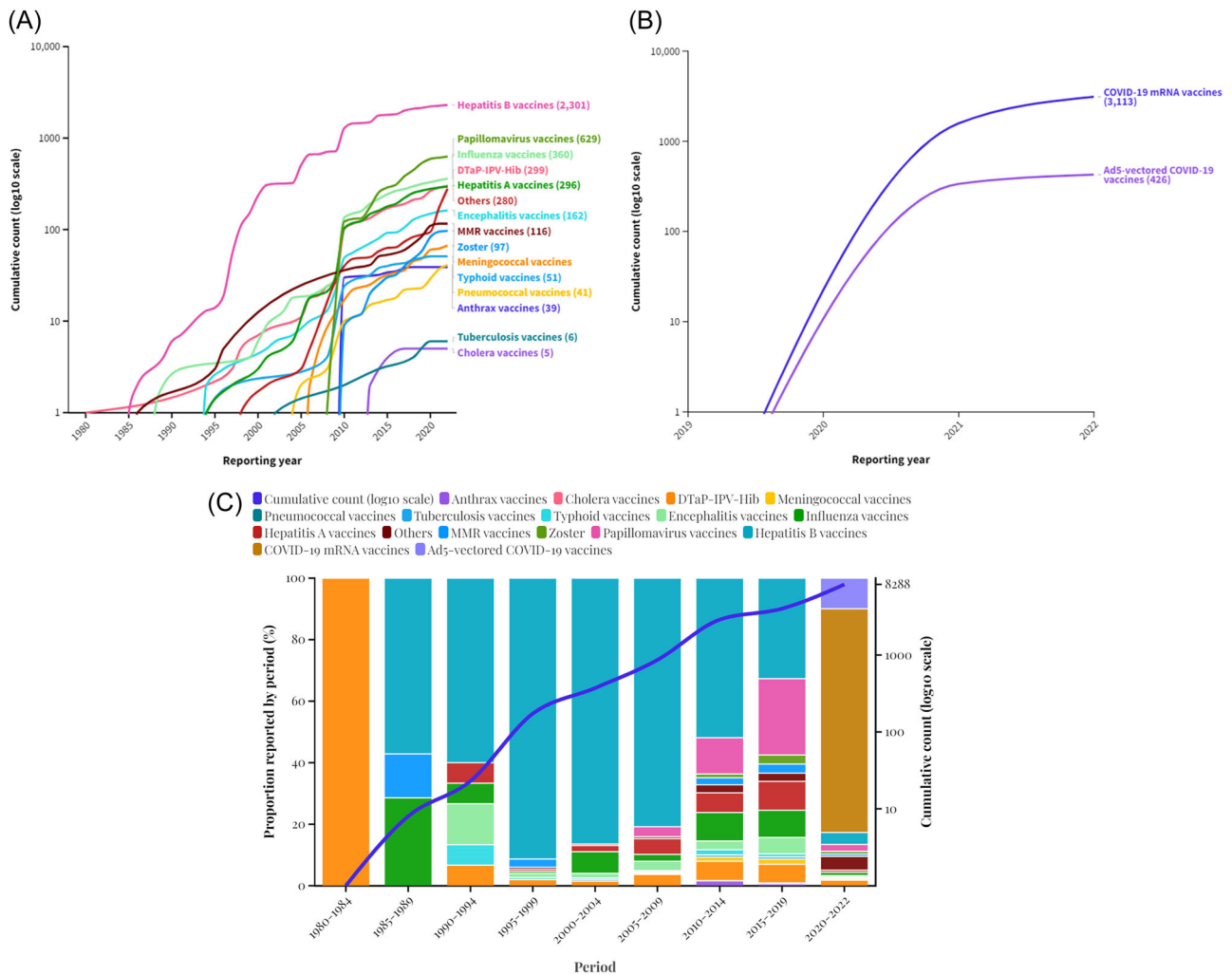


FIGURE 3 Cumulative number of reports of multiple sclerosis per year in association with different vaccines. (A–C) Cumulative counts of multiple sclerosis according to the reporting year. Other vaccines included brucellosis, plague, typhus, leptospirosis, rabies, yellow fever, smallpox, Ebola, and dengue vaccines. Ad5, adenovirus type-5; COVID, coronavirus disease 2019; DTaP-IPV-Hib, diphtheria, tetanus, acellular pertussis, polio, and *Hemophilus influenzae* type b; HPV, human papillomavirus; MMR, measles, mumps, and rubella.

4 | DISCUSSION

This global pharmacovigilance data set, consisting of almost 120 million reports between 1967 and 2022, was analyzed for a comprehensive investigation of vaccine-associated MS. We observed a gradual increase in the number of cases of vaccine-associated MS over the years, with a substantial increase observed after 2020 due to COVID-19 mRNA vaccine-associated MS. Vaccine-associated MS develops more frequently in males and adolescents. Nine vaccines were significantly associated with higher MS reporting, and the highest disproportional associations were observed for hepatitis B vaccines, followed by encephalitis, hepatitis A, and papillomavirus vaccines. In addition, MS showed a significantly disproportional signal for COVID-19 mRNA vaccines (ROR 1.55; IC₀₂₅ 0.52). These results imply that vaccines are emerging as an important potential cause of MS and suggest the importance of continuous pharmacovigilance monitoring for vaccine AEs.^{31,32}

Our analysis revealed nine vaccines with significant disproportionality in MS reporting. Our study clearly suggests that COVID-19 mRNA vaccines had the highest number of vaccine-associated MS reports, which was significantly disproportionate to vaccine-associated MS reporting. COVID-19 vaccines were granted with emergency use authorization, and several case reports of vaccine-associated MS pointed out safety concerns.^{6,33–38} Moreover, Ad5-vectored COVID-19 vaccines and inactivated whole-virus COVID-19 vaccines did not exhibit disproportional MS signals among the various vaccines analyzed.³⁹ It is also unclear how COVID-19 mRNA vaccines could trigger the immunologic processes that lead to MS. COVID-19 mRNA vaccine-associated MS may develop via an overactive immune system response, including a self-reactive T-cell response, through exposure of dendritic cells to exogenous mRNA.^{37,38,40,41} However, these findings suggest that COVID-19 mRNA vaccines have the greatest potential for vaccine-associated MS. Additionally, the risk of MS in the context of COVID-19 mRNA

TABLE 2 Disproportionality analysis of vaccine-associated multiple sclerosis and relevant subgroups analysis.

	Multiple sclerosis		IC (IC ₀₂₅) based on age, years				
	Total	Observed	ROR (95% CI)	IC (IC ₀₂₅)	12–17 years	18–64 years	≥65 years
Total							
IC (IC ₀₂₅)					7.48 (7.33)	1.22 (1.18)	2.91 (2.71)
ROR (95% CI)					8.08 (7.13–9.15)	1.07 (1.04–1.10)	1.59 (1.41–1.80)
Sex difference							
Male	2 533 435	2104	1.19 (1.14–1.25)	0.24 (0.17)	1.89 (1.52)	0.46 (0.37)	1.01 (0.64)
Female	4 283 879	5974	1.01 (0.98–1.04)	0.01 (–0.03)	2.18 (2.02)	–0.04 (–0.10)	0.46 (0.22)
Anthrax vaccines	9923	39	3.58 (2.61–4.90)	1.79 (1.26)	–0.01 (–10.34)	1.30 (0.72)	–0.01 (–10.33)
Cholera vaccines	2310	5	1.97 (0.82–4.73)	0.85 (–0.71)	–0.10 (–10.42)	0.77 (–1.00)	–0.06 (–10.39)
DTaP-IPV-Hib vaccines	777 222	299	0.35 (0.31–0.39)	–1.52 (–1.71)	0.88 (0.13)	0.85 (0.61)	1.36 (–0.06)
Meningococcal vaccines	144 492	67	0.42 (0.33–0.53)	–1.24 (–1.65)	0.42 (–0.40)	0.36 (–0.27)	–0.29 (–10.61)
Pneumococcal vaccines	264 284	41	0.14 (0.10–0.19)	–2.81 (–3.33)	0.35 (–3.43)	–0.91 (–1.57)	–1.07 (–2.64)
Tuberculosis vaccines	33 415	6	0.16 (0.07–0.36)	–2.52 (–3.94)	–1.49 (–11.81)	0.89 (–0.87)	–0.52 (–10.85)
Typhoid vaccines	16 578	51	2.80 (2.13–3.68)	1.46 (0.99)	–0.62 (–10.95)	1.36 (0.84)	–0.32 (–10.65)
Encephalitis vaccines	19 976	162	7.42 (6.36–8.66)	2.85 (2.59)	3.16 (1.95)	3.16 (2.85)	–0.60 (–10.92)
Influenza vaccines	346 453	360	0.94 (0.85–1.05)	–0.08 (–0.26)	0.16 (–1.26)	0.18 (–0.02)	0.48 (–0.25)
Hepatitis A vaccines	60 558	296	4.46 (3.98–5.00)	2.14 (1.95)	2.27 (1.37)	2.79 (2.56)	–0.62 (–10.94)
Hepatitis B vaccines	109 304	2301	19.82 (19.02–20.67)	4.25 (4.18)	5.09 (4.82)	4.77 (4.69)	–0.58 (–10.90)
MMR vaccines	78 971	116	1.33 (1.11–1.60)	0.41 (0.11)	–3.01 (–13.33)	–6.08 (–16.41)	–0.17 (–10.49)
Zoster vaccines	203 900	97	0.43 (0.35–0.53)	–1.21 (–1.54)	1.14 (–0.17)	–0.67 (–1.18)	0.83 (–0.01)
Papillomavirus vaccines	129 318	629	4.45 (4.11–4.81)	2.14 (2.01)	3.11 (2.89)	2.46 (2.23)	–0.04 (–10.37)
COVID-19 mRNA vaccines	3 230 266	3113	1.55 (1.49–1.60)	0.58 (0.52)	–0.47 (–1.14)	–0.62 (–0.70)	0.72 (0.46)
Ad5-vectored COVID-19 vaccines	1 073 625	426	0.61 (0.56–0.67)	–0.70 (–0.86)	–1.62 (–11.94)	–1.81 (–1.99)	0.76 (0.30)
Others ^a	178 902	280	1.42 (1.26–1.60)	0.51 (0.31)	–0.41 (–4.20)	0.07 (–0.17)	0.35 (–1.21)

Note: Bold style indicates when the value of IC₀₂₅ is greater than 0.0 or the lower end of the ROR 95% CI is greater than 1.0. This means it is statistically significant. Numbers in bold indicate statistical significance.

Abbreviations: CI, confidence interval; DTaP-IPV-Hib, diphtheria, tetanus toxoids, pertussis, polio, and Hemophilus influenza type b; IC, information component; MMR, measles, mumps, and rubella; ROR, reported odds ratio.

^aOthers include brucellosis, plague, typhus, leptospirosis, rabies, yellow fever, smallpox, Ebola, and dengue vaccines.

vaccinations should be emphasized, and importance of monitoring and managing these risks should be stressed.

We also confirmed several significant signals between different vaccines and higher reporting of MS. Previous pooled analyses suggested that there is no evidence that vaccination with tuberculosis, hepatitis B, influenza, MMR, DTaP-IPV-Hib, or typhoid vaccines is associated with an increased risk of developing MS.^{42,43} However, in the present study, hepatitis B vaccines exhibited the strongest disproportionate signal for MS among the various vaccines analyzed. These were the commonly reported vaccine-associated MS cases before 2020. The potential link between hepatitis B vaccines and an increased risk of MS has been evaluated in several previous studies.^{44–46} In addition to previous studies,^{42,46,47} our findings

showed significant disproportionality for vaccine-associated MS in recipients of encephalitis, HPV, hepatitis A, anthrax, typhoid, and MMR vaccines. Our findings provide valuable insights into associations between specific vaccines and MS occurrence in different age subgroups.

To the best of our knowledge, this is the first comprehensive study to investigate the global trends of vaccine-associated MS and identify which of them have significant signals. Despite this, our study has several limitations. First, the database (VigiBase) lacks clinical information, including neurological symptoms or signs, laboratory findings, or radiological features needed to diagnose MS. Second, the ICSR of suspected vaccine-associated MS uses a passive reporting method with the potential for reporting bias, such as

underreporting, and the sources of reports are also heterogeneous. Third, we conducted our analysis using MedDRA codes, which have limitations in appropriately mapping to ICD codes. This could affect the reproducibility. Nevertheless, VigiBase collects ICSR data from more than 150 countries spanning 50 years, which guarantees the generalizability of our findings. Third, the exact denominator of recipients exposed to the vaccines could not be determined. In this study, the total number of ICSRs was used as a denominator for the disproportionality analysis, which has been validated in pharmacovigilance studies of signals. Finally, a causal relationship between COVID-19 mRNA vaccinations and the risk of MS cannot be determined. Therefore, it is important that further research be conducted to provide the necessary evidence to support this result.

In conclusion, using the global pharmacovigilance database, we found that the number of vaccine-associated MS cases substantially increased after 2020. Vaccine-associated MS develops more frequently in males and adolescents. A total of nine vaccines were significantly associated with higher MS reporting, and the highest disproportionate associations were observed for hepatitis B vaccines, followed by encephalitis, hepatitis A, papillomavirus, anthrax, typhoid, COVID-19 mRNA, and MMR vaccines. We should be cautious about the increased risk of MS following vaccination and its risk factors, and further studies are needed to inform the early diagnosis and treatment of vaccine-associated MS.

AUTHOR CONTRIBUTIONS

Study concept and design: Ho Geol Woo, Hyeon Jin Kim, Jaeyu Park, and Dong Keon Yon. *Acquisition, analysis, or interpretation of data:* Ho Geol Woo, Hyeon Jin Kim, Jaeyu Park, and Dong Keon Yon. *Drafting of the manuscript:* Ho Geol Woo, Hyeon Jin Kim, Jaeyu Park, and Dong Keon Yon. *Critical revision of the manuscript for important intellectual content:* Ho Geol Woo, Hyeon Jin Kim, Jaeyu Park, Jinseok Lee, Hayeon Lee, Min Seo Kim, Ai Koyanagi, Lee Smith, Masoud Rahmati, Seung Geun Yeo, Dong Keon Yon. *Statistical analysis:* Ho Geol Woo, Hyeon Jin Kim, Jaeyu Park, and Dong Keon Yon. *Study supervision:* Ho Geol Woo, Hyeon Jin Kim, Jaeyu Park, and Dong Keon Yon. Ho Geol Woo, Seung Geun Yeo, and Dong Keon Yon contributed equally as corresponding authors. Ho Geol Woo, Hyeon Jin Kim, and Jaeyu Park contributed equally as first authors. Dong Keon Yon supervised the study and were guarantors. The corresponding authors attest that all listed authors meet the authorship criteria and that no others meeting the criteria have been omitted. Ho Geol Woo and Dong Keon Yon had full access to all the data in the study and take responsibility for the integrity of the data and the accuracy of the data analysis. All authors have approved the final version of the manuscript before submission.

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CONFLICT OF INTEREST STATEMENT

The authors declare no conflict of interest.

DATA AVAILABILITY STATEMENT

Data are available on reasonable request. Study protocol, statistical code: available from DKY (email: yonkkang@gmail.com). Data set: available from the Uppsala Monitoring Centre or World Health Organization through a data use agreement.

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SUPPORTING INFORMATION

Additional supporting information can be found online in the Supporting Information section at the end of this article.

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