Contents lists available at ScienceDirect

Annals of Hepatology

journal homepage: www.elsevier.es/annalsofhepatology

Original article Global prevalence of occult HBV infection in children and adolescents: A systematic review and meta-analysis



Hepatology

Jiaying Wu^{a,b}, Jiayao He^c, Hongmei Xu^{a,b,*}

^a Department of Infectious Diseases, Children's Hospital of Chongging Medical University, National Clinical Research Center for Child Health and Disorders, Ministry of Education Key Laboratory of Child Development and Disorders, Chongqing Key Laboratory of Child Infection and Immunity, Chongqing, China ^b The First batch of key Disciplines On Public Health in Chongqing, Health Commission of Chongqing, Chongqing, China

^c Affiliated Hospital of Chengdu University, Chengdu, China

ARTICLE INFO

Article History: Received 28 June 2023 Accepted 11 September 2023 Available online 23 September 2023

Keywords: Occult hepatitis B Prevalence Children

ABSTRACT

Introduction and Objectives: Occult HBV infection (OBI) is a specific form of hepatitis B virus (HBV) infection and has the possibility of developing into hepatocellular carcinoma (HCC) in adults. This study aimed to estimate the global prevalence of occult HBV infection in children and adolescents.

Materials and Methods: We systematically searched PubMed, Embase, Web of Science, and Cochrane databases for relevant studies on the prevalence of OBI in children and adolescents. Meta-analysis was performed using STATA 16 software.

Results: Fifty studies were included. The overall prevalence of OBI in children and adolescents was 7.5% (95% CI: 0.050-0.103). In different risk populations, OBI prevalence was remarkably high in the HIVinfected population (24.2%, 95% CI: 0.000-0.788). The OBI prevalence was 0.8% (95% CI:0.000-0.029) in the healthy population, 3.8% (95% CI:0.012-0.074) in the general population, and 6.4% (95% CI: 0.021 -0.124) in children born to HBsAg-positive mothers. Based on different serological profiles, the prevalence of OBI in HBsAg-negative and anti-HBc-positive patients was 6.6% (95% CI: 0.016-0.136), 3.0% (95% CI: 0.009-0.059) in HBsAg-negative and anti-HBc-negative patients, 4.6% (95% CI: 0.015-0.088) in HBsAg-negative and anti-HBs-positive patients, and 3.7% (95% CI: 0.001-0.102) in HBsAg-negative and anti-HBs-negative patients.

Conclusions: Despite HBV vaccination and hepatitis B immunoglobulin (HBIG), OBI is common in children and adolescents in high-risk groups.

© 2023 Fundación Clínica Médica Sur, A.C. Published by Elsevier España, S.L.U. This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0/)

1. Introduction

Hepatitis B virus (HBV) is an enveloped DNA hepadnavirus that is responsible for hepatitis, liver cirrhosis, and even hepatocellular carcinoma (HCC) [1–2]. Despite expanded immunization and antiviral treatment, HBV infection is still a remarkable global health issue. According to the World Health Organization (WHO), approximately 296 million people had chronic HBV infection, and approximately 6 million children under five years old were infected with HBV worldwide in 2019 [3].

Occult HBV infection (OBI) is a specific form of HBV infection and is defined as the presence of HBV DNA in the liver or serum of individuals who test negative for HBsAg using currently available assays

Abbreviations: OBI, Occult HBV infection; HBV, Hepatitis B virus; HCC, Hepatocellular carcinoma; CI, Confidence interval; WHO, World Health Organization; MTCT, Motherto-child transmission; HAART, Highly active antiretroviral therapy

[4]. Many studies have demonstrated that mutations or deletions in the pre-S/S region of HBV may alter viral antigenicity and phenotype, which can lead to false-negative results of HBsAg [5-8]. OBI is classified into seropositive OBI (anti-HBc and/or anti-HBs positive) and seronegative OBI (anti-HBc and anti-HBs negative) based on the anti-HBc serostatus [4]. OBI has the pro-oncogenic properties of HBV and it has the possibility of developing into HCC in adults [9-11].

The incidence of OBI varies with the prevalence of HBV, and individuals from HBV hyper-endemic regions are more susceptible to occult HBV infection [12]. The majority of cases of HBV-infected children occur in the perinatal or early childhood period. In addition, HBV vaccination is not completely effective, and children born to mothers with OBI or HBV infection are likely to have occult HBV infections [13].

Some studies have reported the global prevalence of OBI in adults. However, no systematic review for OBI in children and adolescents has been conducted. Therefore, this meta-analysis aims to estimate the global prevalence of OBI in children and adolescents.

https://doi.org/10.1016/j.aohep.2023.101158

1665-2681/© 2023 Fundación Clínica Médica Sur, A.C. Published by Elsevier España, S.L.U. This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0/)



Corresponding author.

E-mail address: xuhongm0095@cqmu.edu.cn (H. Xu).

2. Materials and methods

2.1. Search strategy and study selection

For this meta-analysis, we systematically searched four databases (PubMed, Cochrane, Embase, and Web of Science) to collect relevant studies about the prevalence of OBI in children and adolescents published up to February 03, 2023.

We used the following terms to search for studies: "occult hepatitis B virus infection" and its synonyms, "children" and its synonyms (the full search strategy is available in the Supplementary Information).

All included studies were required to meet the following criteria: 1) defined OBI as the presence of HBV DNA in serum and/or liver tissue without detectable HBsAg, 2) assessed the prevalence of OBI, 3) sample size ≥ 10 , 4) included HBsAg-negative children and adolescents (age ≤ 18 years old), 5) written in English, and 6) could be retrieved in full-text. The exclusion criteria were as follows: 1) not pediatric population, 2) HBV-DNA was not tested, 3) data were duplicated or/and cannot be extracted, 4) case reports, editorial letters, conference abstracts, and reviews, and 5) studies were retracted.

Two independent reviewers (WJY and XHM) selected potentially eligible studies based on the inclusion and exclusion criteria by screening the title and abstract. The full texts of studies deemed eligible were reviewed thoroughly. Any discrepancies in study selection were resolved through discussion.

2.2. Data extraction

Two reviewers used an Excel form to record the following information from eligible studies: first author, year of publication, study type, study region, sample size, and participant characteristics (number of OBI patients, number of anti-HBc-positive patients, and population group).

2.3. Quality assessment

Two reviewers (WJY and XHM) independently assessed the quality of the eligible studies using the method described in a previous study [14]. This assessment method includes 7 items divided into three dimensions: sample size, laboratory methods, and external validity. Any disagreements were resolved by a third reviewer (HJY).

2.4. Statistical analysis

The pooled prevalence of OBI was calculated with a 95% confidence interval (CI). Heterogeneity of studies was assessed by the Cochrane Q test and quantified by I^2 values. A *P* value of the Q test < 0.1 or/and $I^2 \ge 75\%$ was identified as high heterogeneity [15]. A random-effect model was applied for statistical analysis. Subgroup analysis was performed to determine the source of heterogeneity. Otherwise, a fixed-effect model was performed. Publication bias was assessed by a funnel plot and Egger's test. If the funnel plot is asymmetric and the *P* value of Egger's test is less than 0.05, publication bias may exist. Statistical analysis was conducted by STATA version 16.0 software (Stata Corporation, College Station, TX, USA).

2.5. Ethical statements

This systematic review and meta-analysis was conducted following the Preferred Reporting Items for Systematic Reviews and Metaanalyses (PRISMA) reporting guidelines. The protocol was not registered.

3. Results

A total of 1275 studies were obtained from the initial search of four online databases: 421 from PubMed, 441 from Embase, 366 from Web of Science, and 47 from the Cochrane Library. A total of 456 studies were removed because of duplication, 514 studies were excluded after screening the titles and abstracts, and 302 studies were retrieved to read the full-text for further evaluation. Ultimately, 50 studies that met the inclusion criteria were included in this systematic review and meta-analysis. The details of the study selection process are shown in Fig. 1.

3.1. Characteristics of the included studies

The main characteristics of the 50 included studies are described in Table 1 [13,16–64]. Overall, a total of 12977 HBsAg-negative participants from 50 studies were included, including 743 OBI patients.



Fig. 1. Flow-chart of study selection.

Table 1	
Characteristic of the studies used in the systematic review and meta-analysis.	

First author	Publication year	Type of study	Location	Group	Occult HBV infection	Sample size	Anti-HBc+	Serological criteria to test for OBI	OBI prevalence (total OBI / number tested for HBV DNA)
H. Elrashidy [16]	2014	Cohort	Egypt	Healthy/diabetes population	0	170	0	HBsAg-/Anti-HBc-	0.0%
A. Youssef [17]	2013	Cross-sectional	Egypt	Acute hepatitis	7	24	0	HBsAg-	29.2%
C M Jaramillo [18]	2017	Cross-sectional	Colombia	General population	2	24	24	HBsAg-/Anti-HBc+	8 3%
G Gachara [19]	2017	Cross-sectional	Cameroon	HIV infected population	4	34	NR	HBsAg_	11.8%
	2017	Cross sectional	Taiwan	Ceneral population	3	683	0	HBsAg /Anti HBc	0.4%
T. I. HSU [20]	2017	Cobort	China	Born to UPsAg positive mothers	2	105	NP	HDSAg-/AIIU-HDC-	1.6%
Z. A. Chen [21]	2017	Conort	Unind	DOIN TO HDSAg-positive mothers	3	100	INK	IDSAg-	1.0%
E. Seremba [22]	2017	Cross-sectional	Uganda	Born to HIV of HBV infected /nealthy mother	0	20	INK	HBSAg-	0.0%
M. W. Lai [23]	2016	Cross-sectional	Taiwan	Full vaccinated population	36	675	0	HBsAg-/Anti-HBc-	5.3%
L. L. A. Rodriguez Lay [24]	2017	Cross-sectional	Cuba	Born to HBsAg-positive mothers	1	30	3	HBsAg-	3.3%
E. Amponsah-Dacosta [25]	2015	Cohort	South Africa	Non-HIV/HIV infected population with exposure to HBV	35	53	NR	HBsAg-	66.0%
Q. Q. Yao [26]	2013	Cohort	China	Born to HBI/Non-HBI mother	2	92	3	HBsAg-	2.2%
Z. N. A. Said [27]	2009	Case-control	Egypt	Population with haematological disorders and malignancies	21	55	NR	HBsAg-	38.2%
A. Ghaziasadi [28]	2020	Cross-sectional	Iran	General population	91	660	0	HBsAg-/Anti-HBc-	12.8%
N. S. Mohamed [29]	2020	Case-control	Egypt	Frequently Blood Transfused population	27	45	NR	HBsAg-	60%
M. Dapena [30]	2013	Cross-sectional	Spain	HIV-infected population	0	251	NR	HBsAg-	0
S. Shahmoradi [31]	2012	Cross-sectional	Iran	born to HBsAg-positive mothers	21	75	9	HBsAg-	28%
M. H. El-Save [32]	2021	Cohort	Egypt	Polytransfused Children with hematologic malignancy	26	58	8	HBsAg-	44.8%
I. S. Elefsiniotis [33]	2011	Cross-sectional	Greece.Russia.	Born to chronic HBV infected mother	0	44	32	HBsAg-	0
	2011		Bulgaria, Romania, and Serbia		Ū		52		
S. Barfi [34]	2019	Cohort	Iran	ASD/healthy popolation	1	254	7	HBsAg-	0.4%
S. Zhou [35]	2017	Cross-sectional	China	born to HBsAg positive mothers	28	74	NR	HBsAg-	37.8%
L. M. Villar [36]	2014	Cross-sectional	Brazil	General population	9	29	29	HBsAg-/Anti-HBc+	31.0%
X [in [37]	2016	Cross-sectional	China	General population	0	34	34	HBsAg-/Anti-HBc+	0
X Oi [38]	2023	Cross-sectional	China	General population	103	1679	NR	HBsAg_	61%
M R Aghasadeghi [39]	2020	Cross-sectional	Iran	Ceneral population	0	742	3	HBsAg_	0
O Shaker [40]	2020	Cohort	Faunt	Transfused population with thalassemia	26	80	NP	HBcAg	27.5%
0. Slidker [40]	2012	Conortional	Egypt	Deputation with chronic liver disease	20	00 4E	10	HBAAg Janti LIDal	52.5% 9.0%
A. SIIVaStava [41]	2015	CIOSS-Sectional	IIIUIa	Population with chronic liver disease	4	45	40	or anti-HBs+	0.9%
W. L. Hung [42]	2019	Cohort	Taiwan	Healthy/Non A to E hepatitis/ CHC population	22	422	90	HBsAg-	5.2%
G. Beykaso [43]	2022	Cross-sectional	Ethiopia	General population	1	12	12	HBsAg-/Anti-HBc+	8.3%
G. Y. Minuk [44]	2005	Cross-sectional	Canada	General population	6	119	NR	HBsAg-	5.0%
H. X. Su [45]	2013	Cross-sectional	China	Born to HBsAg positive mother	9	183	63	HBsAg-	4.92%
H. Foaud [46]	2015	Cohort	Egypt	born to HBsAg-positive mothers	1	63	0	HBsAg-	1.6%
S. C. Mu [47]	2009	Cross-sectional	Taiwan	HBV vaccinated children	5	46	3	HBsAg-	10.9%
K Yokoyama [48]	2017	Cross-sectional	lanan	horn to HBsAg-nositive mothers	2	158	NR	HBsAg_	0
	2021	Clinical trial	Taiwan	born to HBsAg_positive mothers	8	220	NR	HBsAg_	3.6%
	2021	Cohort	China	Born to HBI/Non HBI mother	3	328	NP	HBcAg	0.0%
H Su [51]	2021	Conort Cross soctional	China	Conoral population	15	1102	ND	HBcAg	1.2%
G. D. Zhugo [52]	2017	Ciuss-sectional	China	been to LIBA a positive perents	15	227	25	IIBAA	1.3%
5. K. Zhuge [52]	2020			Doin to HDSAg-positive patents	40	327	25	HDSAg-	14.1%
L. Yong [53]	2014	Cross-sectional	China	born to HBSAg-positive mothers	0	207	8	HBSAg-	0
S. J. Chen [54]	2012	Cross-sectional	China	Healthy population	9	1146	141	HBsAg-	0.8%
A. Marjani [55]	2022	Cross-sectional	Iran/Afghanistan	Working population	0	368	2	HBsAg-	0
T. Utsumi [56]	2010	Cross-sectional	Indonesia	General population	5	89	NR	HBsAg-/anti-HBs+ and/or anti- HBc+	5.6%
H. E. Raouf [57]	2015	Cohort	Egypt	HCV positive/negative cancer population	16	100	NR	HBsAg-	16%
A. Eilard [58]	2019	Cross-sectional	Sweden	born to HBsAg-positive mothers	3	44	NR	HBsAg-	6.8%
N. Weis [59]	2017	Cross -sectional	Denmark	born to CHB mothers	0	125	19	HBsAg-	0
A. Walz [13]	2009	Cross-sectional	Germany	born to Anti-HBc mothers	5	103	NR	HBsAg-	4.9%
C.Chakvetadze [60]	2011	Cohort	Mayotte, France	born to HBsAg-positive mothers	2	99	NR	HBsAg-	2.0%
C. Pande [61]	2013	Clinical trial	India	born to HBsAg-positive mothers	89	204	NR	HBsAg-	43.6%
A Y Li [62]	2020	Cohort	China	born to HBsAg and HBeAg	20	169	NR	HBsAg-/Anti-HBs+	11.8%
	2020			positive mothers	20	105	ND		22.4%
C. J. Hoffmann [63]	2014	Cohort	South Africa	born to CHB/Non-CHB mothers living with HIV	3	13	NK	HBSAg-	23.1%
H. Y. Hsu [64]	2015	cross-sectional	Taiwan	General population	23	1125	515	HBSAg-	2.0%

				%
study	events	total	ES (95% CI)	vveight
H. Elrashidy, 2014	0	170	♦ 0.000 (0.000, 0.021)	2.13
A. Youssef, 2013	7	24	0.292 (0.126, 0.511)	1.63
C. M. Jaramillo, 2017	2	24	0.083 (0.010, 0.270)	1.63
G. Gachara, 2017	4	34	0.118 (0.033, 0.275)	1.77
H. Y. Hsu, 2017	3	683	• 0.004 (0.001, 0.013)	2.22
Z. X. Chen, 2017	3	185	• 0.016 (0.003, 0.047)	2.14
E. Seremba, 2017	0	20	0.000 (0.000, 0.168)	1.55
M. W. Lai, 2016	36	675	0.053 (0.038, 0.073)	2.22
L. L. A. Rodríguez Lay, 2017	1	30	0.033 (0.001, 0.172)	1.72
E. Amponsah-Dacosta, 2015	35	53	0.660 (0.517, 0.785)	1.92
Q. Q. Yao, 2013	2	92	0.022 (0.003, 0.076)	2.04
Z. N. A. Said, 2009	21	55	0.382 (0.254, 0.523)	1.93
A. Ghaziasadi, 2020	91	660	0.138 (0.112, 0.167)	2.22
N. S. Mohamed, 2020	27	45	• 0.600 (0.443, 0.743)	1.87
S. Shahmoradi, 2012	21	75	0.280 (0.182, 0.396)	2.00
M. H. El-Sayed, 2021	26	58	0.448 (0.317, 0.585)	1.94
M. Dapena, 2013	0	251	• 0.000 (0.000, 0.015)	2.17
I. S. Elefsiniotis, 2011	0	44	• 0.000 (0.000, 0.080)	1.86
S. Zhou, 2017	28	74	0.378 (0.268, 0.499)	2.00
L. M. Villar, 2014	9	29	0.310 (0.153, 0.508)	1.71
S. Barfi, 2019	1	254	• I 0.004 (0.000, 0.022)	2.17
X. Lin, 2016	0	34	• <u>•</u> 0.000 (0.000, 0.103)	1.77
X. Qi, 2023	103	1679	▲ 0.061 (0.050, 0.074)	2.24
M. R. Aghasadeghi, 2020	0	742	• 0.000 (0.000, 0.005)	2.22
O. Shaker, 2012	26	80	0.325 (0.224, 0.439)	2.02
A. Srivastava, 2015	4	45	0.089 (0.025, 0.212)	1.87
W. L. Hung, 2019	22	422	0.052 (0.033, 0.078)	2.20
G. Beykaso, 2022	1	12	0.083 (0.002, 0.385)	1.29
G. Y. Minuk, 2005	6	119	0.050 (0.019, 0.107)	2.09
H. X. Su, 2013	9	183	0.049 (0.023, 0.091)	2.14
H. Foaud, 2015	1	63	• • • • • • • • • • • • • • • • • • •	1.96
S. C. Mu, 2009	5	46	0.109 (0.036, 0.236)	1.87
K.Yokoyama, 2017	2	158	• 0.013 (0.002, 0.045)	2.13
H. Y. Hsu, 2021	8	220	0.036 (0.016, 0.070)	2.16
A. Q. Hu, 2021	3	328	• 0.009 (0.002, 0.026)	2.19
H. Su, 2017	15	1192	• 0.013 (0.007, 0.021)	2.23
S. R. Zhuge, 2020	46	327	0.141 (0.105, 0.183)	2.19
L. Yong, 2014	0	207	◆ 0.000 (0.000, 0.018)	2.15
A. Marjani, 2022	0	368	◆ 0.000 (0.000, 0.010)	2.20
T. Utsumi, 2010	5	89	0.056 (0.018, 0.126)	2.04
H. E. Raouf, 2015	16	100	0.160 (0.094, 0.247)	2.06
S. J. Chen, 2012	9	1146	• <u>1</u> 0.008 (0.004, 0.015)	2.23
A. Eilard, 2019	3	44	0.068 (0.014, 0.187)	1.86
N. Weis, 2017	0	125	• 0.000 (0.000, 0.029)	2.09
A. Walz, 2009	5	103	••••	2.06
C.Chakvetadze, 2011	2	99	• 0.020 (0.002, 0.071)	2.06
C. Pande, 2013	89	204	0.436 (0.367, 0.507)	2.15
A. Y. Li, 2020	20	169	0.118 (0.074, 0.177)	2.13
C. J. Hoffmann, 2014	3	13	0.231 (0.050, 0.538)	1.33
H. Y. Hsu, 2015	23	1125	•	2.23
Overall (I ² = 96.389%, p = 0.0	000)		0.075 (0.050, 0.103)	100.00
		T		
		1	0 .1 .2 .3 .4 .5 .6 .7 .8	

Fig. 2. Overall prevalence of OBI.

The majority of included studies (31/50) were cross-sectional studies, 15 were cohort studies, 2 were clinical trials, and 2 were case-control studies. Of the 50 studies included, 27 studies were conducted in Asia, 14 in Africa, 5 in Europe, 2 in South America, and 2 in North America.

According to the risk of acquiring HBV infection, participants enrolled in the included studies were divided into four populations, including the healthy population in 4 studies, the general population in 12 studies, the population born to HBsAg-positive mothers in 19 studies, and the HIV-infected population in 3 studies.

3.2. Quality assessment

The details of the methodological quality assessment are illustrated in Supplement Table 1. 14 studies were considered at low risk of bias, 20 studies were considered at moderate risk of bias, and 16 studies were considered at high risk of bias.

3.3. Overall OBI prevalence in children

A random-effect model was performed to estimate the overall prevalence of occult HBV infection in children and adolescents because it has significantly high heterogeneity (l^2 = 96.389%, p < .1).

The overall pooled OBI prevalence in children and adolescents was 7.5% (95% CI: 0.050-0.103) (Fig. 2). Based on different serological criteria, the overall prevalence of OBI was as follows: 7.9% (95% CI: 0.050-0.114) for HBsAg-negative, 3.2% (95% CI: 0.000-0.108) for HBsAg-negative and anti-HBc-negative, 8.9% (95% CI: 0.006-0.232) for HBsAg-negative and anti-HBc-positive, and 7.3% (95% CI: 0.039-0.114) for HBsAg-negative and other criteria.

Healthy H. Elrashidy, 2014 S. Barfi, 2019	events	total	ES (95% CI)	vveiar
Healthy H. Elrashidy, 2014 S. Barfi, 2019				J
S. Barfi, 2019	0	107) 0.70
5. Dalli, 2019	0	147) 2.70
W I Hung 2010	16	201		2.04
VV. L. Hulig, 2019	0	1146) 2.97
S. J. Chen, 2012	9	1140		0 3.03
Subtotal (1 $^{\prime}$ 2 = 65.502%, p = 0.	.000)) 11.00
HIV infected				
G. Gachara, 2017	4	34	0.118 (0.033, 0.275) 2.29
E.Amponsah–Dacosta, 2015	11	12	• 0.917 (0.615, 0.998) 1.59
M. Dapena, 2013	0	251	• 0.000 (0.000, 0.015) 2.92
Subtotal (I ² = .%, p = .)			0.242 (0.000, 0.788	6.81
Born to HBsAg positive mother				
Z. X. Chen, 2017	3	185	• 0.016 (0.003, 0.04)) 2.88
L. L. A. Rodríguez Lay, 2017	1	30	0.033 (0.001, 0.172	2.22
Q. Q. Yao, 2013	0	33	0.000 (0.000, 0.100) 2.28
S. Shahmoradi, 2012	21	75	0.280 (0.182, 0.396) 2.65
I. S. Elefsiniotis, 2011	0	44	0.000 (0.000. 0.080) 2.43
S. Zhou, 2017	28	74	0.378 (0.268, 0.499) 2.65
H. X. Su, 2013	9	183	0.049 (0.023, 0.09) 2.88
H. Foaud, 2015	1	63	0.016 (0.000, 0.08	2.59
K Yokovama 2017	2	158	0.013 (0.002, 0.04	2 85
H Y Hsu 2021	8	220) 291
A O Hu 2021	3	53	0.057 (0.012, 0.157) 2.52
S B Zhuge 2020	24	172) 2.87
L Yong 2014	0	207		2.07
A Filard 2019	3	44) 243
N Weis 2017	0	125) 2.40
C Chakvetadze 2011	2	99) 2.00
C Pando 2013	20	204) 200
A V Li 2020	20	160	0.430 (0.307, 0.307) 2.90
C. I. Hoffmann 2014	20	109	0.200 (0.025, 0.55)) 2.00
C. J. Holimann, 2014	2	10) 1.40
Subtotal (1 $^{\prime}$ 2 = 95.251 $^{\prime}$ 6, p = 0.	.000)		0.004 (0.021, 0.124	49.01
General				
C. M. Jaramillo, 2017	2	24	0.083 (0.010, 0.270) 2.08
H. Y. Hsu, 2017	3	683	• 0.004 (0.001, 0.013) 3.01
A. Ghaziasadi, 2020	91	660	0.138 (0.112, 0.16) 3.01
L. M. Villar, 2014	9	29	0.310 (0.153, 0.508) 2.20
X. Lin, 2016	0	34	0.000 (0.000, 0.103) 2.29
X. QI, 2023	103	1679	0.061 (0.050, 0.074) 3.04
M. R. Aghasadeghi, 2020	0	742	• 1 0.000 (0.000, 0.008) 3.01
G. Beykaso, 2022	1	12	0.083 (0.002, 0.385) 1.59
G. Y. Minuk, 2005	6	119	0.050 (0.019, 0.10) 2.79
H. Su, 2017	15	1192	♦ 1 0.013 (0.007, 0.02 ⁻) 3.03
T. Utsumi, 2010	5	89	0.056 (0.018, 0.126	i) 2.71
H. Y. Hsu, 2015	23	1125	• I 0.020 (0.013, 0.03) 3.03
Subtotal (I ² = 96.384%, p = 0.	.000)		0.038 (0.012, 0.074) 31.78
Heterogeneity between aroups:	p = 0.021		i i i i i i i i i i i i i i i i i i i	
Overall (I^2 = 95.841%, p = 0.0	000);		O .051 (0.029, 0.07)) 100.0
		_ 1		
		1	۰.۱۰،۷۰،۷۰،۷۰،۷۰،۷۰،۷۰،۷۰،۷۰	

Fig. 3. OBI prevalence in different populations.

3.4. OBI prevalence in different populations

There was a significant variation in different populations. The prevalence of OBI in the healthy population was estimated to be 0.8% (95% CI:0.000–0.029), with a total of 1791 participants from 4 prevalence datasets. The prevalence of OBI in the general population was estimated to be 3.8% (95% CI:0.012–0.074), with a total of 6388 participants from 12 prevalence datasets. The prevalence of OBI in the population with HIV infection was 24.2% (95% CI: 0.000–0.788), with a total of 297 participants from 3 prevalence datasets. The prevalence of OBI in children born to HBsAg-positive mothers was 6.4% (95% CI: 0.021–0.124), with a total of 2148 participants from 19 prevalence datasets (Fig. 3).

OBI prevalence varied with different methods of passive-active immunoprophylaxis. For the population born to HBsAg-positive

mothers, the prevalence of OBI was 3.0% (95% CI: 0.000–0.124) in children vaccinated with 3 doses of vaccine and HBIG, followed by 7.2% (95% CI: 0.000–0.309) in children vaccinated with 4 doses of vaccine and HBIG, 7.7% (95% CI: 0.000–0.476) in children vaccinated with 4 doses of vaccine, and 50% (95% CI: 0.313–0.687) in children vaccinated with 2 doses of vaccine and HBIG (Fig. 4).

3.5. OBI prevalence in different serological profiles of HBV infection

The prevalence of OBI varied with HBV serological status. The prevalence of OBI in HBsAg-negative and anti-HBc-positive participants was 6.6% (95% CI: 0.016–0.136), with a total of 1075 participants from 18 prevalence datasets. The prevalence of OBI in HBsAg-negative and anti-HBc-negative participants was 3.0% (95% CI: 0.009

Study	Events	Total	ES (95% CI)	1
4 dosos vaccinos			i i	
L A Bodríguez Lav 2017	0	30		
A Filard 2019	0	35		
C Pande 2013	44	104		
Subtotal $(1/2 = \% n =)$		104	0.077 (0.000, 0.476)	
3 doses vaccines+HBIG				
S. Shahmoradi, 2012	21	75	0.280 (0.182, 0.396)	
S. Zhou, 2017	13	44	0.295 (0.168, 0.452)	
H. Foaud, 2015	0	1	0.000 (0.000, 0.975)	
K.Yokoyama, 2017	2	158	♦ 1	
H. Y. Hsu, 2021	8	220	0.036 (0.016, 0.070)	
L. Yong, 2014	0	207	•	
A. Y. Li, 2020	20	167	0.120 (0.075, 0.179)	
Q. Q. Yao, 2013	0	33	0.000 (0.000, 0.106)	
Subtotal (I^2 = 93.510%, p =	0.000)		0.030 (0.000, 0.124)	
4 doses vaccines+HBIG				
. S. Elefsiniotis, 2011	0	44	• • • • 0.000 (0.000, 0.080)	
H. Foaud, 2015	1	62	0.016 (0.000, 0.087)	
A. Eilard, 2019	1	8	0.125 (0.003, 0.527)	
C.Chakvetadze, 2011	1	83	• 0.012 (0.000, 0.065)	
C. Pande, 2013	45	100	• 0.450 (0.350, 0.553)	
Subtotal (I^2 = 95.955%, p =	0.000)		0.072 (0.000, 0.309)	
2 doses vaccines+HBIG				
S. Zhou, 2017	15	30	• 0.500 (0.313, 0.687)	
Heterogeneity between group	s: p = 0.000			
Overall (I ² = 95.301%, p = 0	.000);		0.070 (0.009, 0.163)	
		1	0 1 2 3 4 5 6 7 8	

Fig. 4. OBI prevalence in different methods of immunoprophylaxis.

-0.059), with a total of 5775 participants from 21 prevalence datasets (Fig. 5).

For anti-HBs in serum, ≥ 10 mIU/ml was considered positive. Thus, the prevalence of OBI in HBsAg-negative and anti-HBs-positive participants was 4.6% (95% CI: 0.015–0.088), with a total of 2281 participants from 14 prevalence datasets. The prevalence of OBI in HBsAg-negative and anti-HBs-negative participants was 3.7% (95% CI: 0.001–0.102), with a total of 1342 participants from 13 prevalence datasets (Fig. 6).

3.6. Publication bias

Publication bias was assessed using Egger's test and a visual inspection of the funnel plot. The shape of the funnel plot is not symmetrical, indicating a potential bias (Fig. 7). The results of Egger's test (p value = 0.002) confirm the existence of publication bias.

4. Discussion

This is the first meta-analysis to estimate the global prevalence of OBI in children and adolescents. The prevalence of OBI varies in different populations. In this study, we found that the prevalence of OBI was 0.8% in the healthy population and 3.8% in the general population. The prevalence of OBI in children born to HBsAg-positive mothers was almost twice that in the general population. OBI prevalence was remarkably higher in the HIV-infected population than in the population born to HBsAg-positive mothers.

Mother-to-child transmission (MTCT) of HBV is a key route of transmission. WHO recommends that children born to HBsAg-positive mothers should receive HBIG and at least 3 doses of hepatitis B vaccine to prevent MTCT [65]. However, there is still a residual risk of HBV infection in children born to HBsAg-positive mothers despite neonatal passive-active immunoprophylaxis. In our study, we found that the usage of HBIG and 3 doses of hepatitis B vaccine was the most effective method to prevent children born to HBsAg-positive mothers from occult HBV infection. We need studies of large samples to confirm this result. In addition, maternal HBeAg status and HBV DNA load had positive correlations with the MTCT incidence of HBV [66]. However, Li et al [67] found that maternal age, HBsAg titer, HBeAg status, HBV DNA viral load, alanine aminotransferase level, child's sex, feeding pattern, HBIG dosage, birth weight, and anti-HBs level had no significant association with OBI incidence in 7-monthold infants. These controversies need further research to resolve. In addition, some studies have indicated that OBI in children born to HBsAg-positive mothers may be transient or become overt after

Study	Events	Total	ES (95% CI)	% Weight
anti-HBc -				
Heba Elrashidy,2014	0	170	• I 0.000 (0.000, 0.021)	3.23
A. Youssef,2013	7	24	0.292 (0.126, 0.511)	2.35
H. Y. Hsu, 2017	3	683	◆ I 0.004 (0.001, 0.013)	3.38
M. W. Lai, 2016	36	675	◆ 0.053 (0.038, 0.073)	3.38
L. L. A. Rodríguez Lay, 2017	1	27	0.037 (0.001, 0.190)	2.44
Q. Q. Yao, 2013	2	89	• 0.022 (0.003, 0.079)	3.05
A.Ghaziasad , 2020	91	660	0.138 (0.112, 0.167)	3.38
S. Shahmoradi, 2012	16	66	0.242 (0.145, 0.364)	2.94
S. Zhou, 2017	2	14	0.143 (0.018, 0.428)	1.93
S. Barfi, 2019	1	247	• I 0.004 (0.000, 0.022)	3.29
A. Srivastava, 2015	0	5	0.000 (0.000, 0.522)	1.13
H. X. Su, 2013	8	120	0.067 (0.029, 0.127)	3.14
H. Foaud, 2015	1	63	0.016 (0.000, 0.085)	2.92
S. C. Mu, 2009	5	43	0.116 (0.039, 0.251)	2.73
S. R. Zhuge, 2020	40	302	0.132 (0.096, 0.176)	3.32
L. Yong, 2014	0	199	0.000 (0.000, 0.018)	3.25
A. Marjani, 2022	0	366		3.34
S. J. Chen, 2012	0	1005		3.40
N. Weis 2017	0	106	0.000 (0.000, 0.034)	3.11
H. Y. Hsu, 2015	4	610	0.007 (0.002, 0.017)	3.38
W. L. Hung	8	301	0.027 (0.012, 0.052)	3.31
Subtotal $(1^2 = 95.519\%, p =$	0.000)		0.030 (0.009, 0.059)	62.41
anti HBa			l i i i i i i i i i i i i i i i i i i i	
	0	04		0.05
C. M. Jaramino,2017	2	24		2.35
C. C. Xao, 2012	0	3		0.01
S Shahmoradi 2012	5	0	0.556 (0.212, 0.863)	1.57
S Zhou 2017	26	60		2.00
5. 2100, 2017	20	20	0.310 (0.153 0.508)	2.90
S Barfi 2010	9	29		1 27
X Lin 2016	0	34		2.59
A Srivastava 2015	4	40		2.69
H X Su 2013	1	63		2.00
S C Mu 2009	0	3		0.81
S. B. Zhuge, 2020	6	25	0.000 (0.000, 0.100)	2.38
L. Yong. 2014	0	8	0.000 (0.000, 0.369)	1.48
A. Mariani, 2022	0	2		0.62
S. J. Chen. 2012	9	141		3.18
N. Weis 2017	0	19		2.18
H. Y. Hsu. 2015	19	515	•	3.37
W. L. Hung	8	90	0.089 (0.039, 0.168)	3.06
Subtotal (1/2 = 83.672%, p =	0.000)		0.066 (0.016. 0.136)	37.59
Heterogeneity between aroup	s: p = 0.02	24		
Overall (1/2 = 93.675%, p = 0	.000);		0.036 (0.013, 0.064)	100.00
,	,.		Ť	
		, I		
		1	0.1.2.3.4.5.6.7.8	

Fig. 5. OBI prevalence in different serological profiles of HBV infection.

several years [24,35,58]. Therefore, long-term follow-up is needed in children born to HBsAg-positive mothers.

Considering the overlap of transmission routes, HBV infection is common in people living with HIV. A meta-analysis study [68] indicated that the global OBI prevalence in HIV-infected patients was 16.26%. Kajogoo *et al* [69] reported that the prevalence of OBI was 12.4% among HIV-infected individuals in Africa. Xie *et al* [70] reported that 9% of patients living with HIV in Asia had occult HBV infection. While we found that the prevalence of OBI was 24.2% in the HIVinfected population in our study. The following reasons may be responsible for this: 1) HIV-infected children have inadequate expression of cytokines related to the immune response and inadequate production of anti-HBs, which may lead to HBV infection [71]. 2) HIV-infected patients have immune dysfunction, which may lead to HBV reactivation [72]. 3) HBV-resistance mutations may occur when HIV-infected patients receive lamivudine-based therapies [73]. 4) Low CD4 count is related to OBI incidence in HIV-infected patients [74]. However, one study had the opposite result, in which the occurrence of OBI was not concerned with CD4 count [75]. Therefore, detection of HBV DNA should be a routine examination in the HIVinfected population. Highly active antiretroviral therapy (HAART) with 2 anti-HBV nucleos(t)ide analogs is recommended for HIVinfected patients with OBI [76].

Anti-HBc is considered a sign when people have been exposed to HBV, and it can be detected before the appearance of HBsAg [77]. In this study, we found that the prevalence of OBI in the HBsAg-negative and anti-HBc-positive participants was significantly higher than that in the HBsAg-negative and anti-HBc-negative participants. This result was in line with two meta-analysis studies that reported 20.1% vs. 8% and 51% vs. 19%, respectively [14,78]. Thus, the detection of anti-HBc is recommended for screening OBI in underdeveloped regions. For anti-HBs, the prevalence of OBI in HBsAg-negative and anti-HBs-positive subjects was similar to that in HBsAg-negative and anti-HBs-negative subjects. However, among children born to HBsAg-positive

Anti-HBs <10 IU/L Heba Elrashidy,2014 C M. W. Lai, 2016 1 L. L. A. Rodríguez Lay, 2017 C A.Ghaziasad , 2020 6 M. H. El-Sayed, 2021 1 S. Zhou, 2017 1 S. Barfi, 2019 C H. X. Su, 2013 3	0 11 0 68 13 1 0	87 238 12 312 33 4	•	+								
Heba Elrashidy,2014 C M. W. Lai, 2016 1 L. L. A. Rodríguez Lay, 2017 C A.Ghaziasad , 2020 6 M. H. El–Sayed, 2021 1 S. Zhou, 2017 1 S. Barfi, 2019 C H. X. Su, 2013 3	0 11 0 68 13 1 0	87 238 12 312 33 4	*	ŀ								
M. W. Lai, 2016 1 L. L. A. Rodríguez Lay, 2017 0 A.Ghaziasad, 2020 6 M. H. El–Sayed, 2021 1 S. Zhou, 2017 1 S. Barfi, 2019 0 H. X. Su, 2013 3	11 0 68 13 1 0	238 12 312 33 4	•	-							0.000 (0.000, 0.042)	4.07
L. L. A. Rodríguez Lay, 2017 (A.Ghaziasad , 2020 6 M. H. El–Sayed, 2021 1 S. Zhou, 2017 1 S. Barfi, 2019 0 H. X. Su, 2013 3	0 68 13 1 0	12 312 33 4	•								0.046 (0.023, 0.081)	4.37
A.Ghaziasad , 2020 6 M. H. El–Sayed, 2021 1 S. Zhou, 2017 1 S. Barfi, 2019 0 H. X. Su, 2013 3	68 13 1 0	312 33 4									0.000 (0.000, 0.265)	2.48
M. H. El–Sayed, 2021 1 S. Zhou, 2017 1 S. Barfi, 2019 0 H. X. Su, 2013 3	13 1 0	33 4	1								0.218 (0.173, 0.268)	4.42
S. Zhou, 2017 5. Barfi, 2019 6. H. X. Su, 2013 3.	1 0	4			_		•		•		0.394 (0.229, 0.579)	3.47
S. Barfi, 2019 (H. X. Su, 2013 3	0				•						- 0.250 (0.006, 0.806)	1.37
H. X. Su, 2013 3		126	+								0.000 (0.000, 0.029)	4.21
11 5 1 0015	3	39		+	_						0.077 (0.016, 0.209)	3.60
H. Foaud, 2015	0	13	♦ 1								0.000 (0.000, 0.247)	2.56
S. C. Mu, 2009 2	2	23		•							0.087 (0.011, 0.280)	3.15
L. Yong, 2014 (0	43	+	_							0.000 (0.000, 0.082)	3.67
N. Weis, 2017 0	0	23	+								0.000 (0.000, 0.148)	3.15
H. Y. Hsu, 2015 4	4	389	◆ - !								0.010 (0.003, 0.026)	4.44
Subtotal (I^2 = 93.020%, p = 0.	.000)			>							0.037 (0.001, 0.102)	44.98
Anti-HBs >=10 IU/L												
Heba Elrashidy,2014 0	0	83	+	_							0.000 (0.000, 0.043)	4.05
M. W. Lai, 2016 2	25	437		-							0.057 (0.037, 0.083)	4.46
L. L. A. Rodríguez Lay, 2017 0	0	18	•		-						0.000 (0.000, 0.185)	2.91
A.Ghaziasad , 2020 2	23	257		•							0.089 (0.058, 0.131)	4.38
M. H. El-Sayed, 2021 1	13	25	i					•		_	0.520 (0.313, 0.722)	3.23
I. S. Elefsiniotis, 2011 0	0	44	+ +	_		_					0.000 (0.000, 0.080)	3.69
S. Zhou, 2017 2	27	70	i				•				0.386 (0.272, 0.510)	3.97
S. Barfi, 2019 1	1	128	+								0.008 (0.000, 0.043)	4.22
H. X. Su, 2013 6	6	144	-+	_							0.042 (0.015, 0.088)	4.25
H. Foaud, 2015 1	1	50	+ -								0.020 (0.001, 0.106)	3.78
S. C. Mu, 2009	3	23	<u>+</u>	•							0.130 (0.028, 0.336)	3.15
L. Yong, 2014 C	0	164	← i								0.000 (0.000, 0.022)	4.29
N. Weis 2017 0	0	102	• !								0.000 (0.000, 0.036)	4.14
H. Y. Hsu, 2015 1	19	736	+								0.026 (0.016, 0.040)	4.50
Subtotal (I^2 = 92.072%, p = 0.	.000)		\diamond	>							0.046 (0.015, 0.088)	55.02
11-1		10										
Heterogeneity between groups:	p = 0.9	49	-								0.044 /0.040 0.6= **	100.00
Overall (I ² = 92.333%, p = 0.0	000);		Ŷ	>							0.041 (0.016, 0.074)	100.00
		- 1	0	1	2	3	4	5	6	1	8	

Fig. 6. OBI prevalence in different serological profiles of HBV infection.

mothers, the OBI prevalence in infants with low anti-HBs (< 100 mIU/ mL) was higher than that in infants with high anti-HBs (\geq 100 mIU/ mL) [62]. Therefore, a high-risk population may need a booster vaccine and/or a higher dosage of vaccine.

Our study had some limitations. First, we used a homemade assessment tool to evaluate the quality of articles, and the majority of included studies were considered at moderate risk. Second, there was significant heterogeneity that we could not explain, although we performed subgroup analyses. Third, we cannot evaluate the prevalence of OBI in the HCV-infected population and transfused population, because there were not enough studies. Fourth, the studies included children born to HBsAg-positive mothers were from different countries with different methods of HBV prophylaxis at birth. Last, the sensitivity and specificity of the HBsAg assay was improved gradually, and the methods of HBV DNA detection were different in the included studies. No gold standards can be followed to estimate OBI prevalence between the studied periods now.

Despite the above limitations, the main strength of our study was that the prevalence of OBI in children born to HBsAg-positive mothers was first estimated. We also accounted the prevalence of OBI according to the anti-HBc/anti-HBs serostatus.



Fig. 7. Funnel plot.

5. Conclusions

This review first summarized the global prevalence of OBI in children and adolescents. With the popularity of the HBV vaccine and HBIG, children still have the chance to be infected with occult HBV. OBI prevalence varies with different populations. In high-risk groups, OBI prevalence is remarkably high. We should pay more attention to OBI in children and adolescents.

Funding

This research did not receive any specific grant from funding agencies in the public, commercial, or not-for-profit sectors.

Author contributions

Jiaying Wu developed the study protocol and wrote the manuscript. Jiaying Wu and Hongmei Xu performed the search, screened the articles and data analysis. Jiaying Wu, Hongmei Xu, and Jiayao He performed quality assessment. All authors reviewed and agreed with the final version of the manuscript.

Data availability statement

Data are available upon request, please contact author for data requests.

Declaration of interests

None.

Supplementary materials

Supplementary material associated with this article can be found in the online version at doi:10.1016/j.aohep.2023.101158.

References

- Hu J, Seeger C. Hepadnavirus genome replication and persistence. Cold Spring Harb Perspect Med. 2015;5(7):a021386. https://doi.org/10.1101/cshperspect. a021386.
- [2] McMahon BJ. The natural history of chronic hepatitis B virus infection. Hepatology 2009;49(5 Suppl):S45–55. https://doi.org/10.1002/hep.22898.
- [3] WHO global progress report on HIV, viral hepatitis and sexually transmitted infections. https://www.who.int/publications/i/item/9789240027077; 2021 [accessed 15 July 2021].
- [4] Raimondo G, Locarnini S, Pollicino T, Levrero M, Zoulim F, Lok AS, et al. Update of the statements on biology and clinical impact of occult hepatitis B virus infection. J Hepatol 2019;71(2):397–408. https://doi.org/10.1016/j.jhep.2019.03.034.
- [5] Huang CH, Yuan Q, Chen PJ, Zhang YL, Chen CR, Zheng QB, et al. Influence of mutations in hepatitis B virus surface protein on viral antigenicity and phenotype in occult HBV strains from blood donors. J Hepatol 2012;57(4):720–9. https://doi. org/10.1016/j.jhep.2012.05.009.
- [6] Wang H, Wang M, Huang J, Xu R, Liao Q, Shan Z, et al. Novel hepatitis B virus surface antigen mutations associated with occult genotype B hepatitis B virus infection affect HBsAg detection. J Viral Hepat 2020;27(9):915–21. https://doi.org/ 10.1111/jvh.13309.
- [7] Wang J, Liu Y, Liao H, Liu L, Chen R, Si L, et al. The sK122R mutation of hepatitis B virus (HBV) is associated with occult HBV infection: Analysis of a large cohort of Chinese patients. J Clin Virol 2020;130:104564. https://doi.org/10.1016/j. jcv.2020.104564.
- [8] Liao H, Liu Y, Chen J, Ding W, Li X, Xu Z, et al. Characterization of hepatitis B virus (HBV) preS/S gene mutations in blood donors with occult HBV infection in the Baoji area of North China. Transfusion 2017;57(3pt2):857–66. https://doi.org/ 10.1111/trf.14046.
- [9] Wong DK, Huang FY, Lai CL, Poon RT, Seto WK, Fung J, et al. Occult hepatitis B infection and HBV replicative activity in patients with cryptogenic cause of hepatocellular carcinoma. Hepatology 2011;54(3):829–36. https://doi.org/10.1002/ hep.24551.
- [10] Chemin I, Zoulim F, Merle P, Arkhis A, Chevallier M, Kay A, et al. High incidence of hepatitis B infections among chronic hepatitis cases of unknown aetiology. J Hepatol 2001;34(3):447–54. https://doi.org/10.1016/s0168-8278(00)00100-8.
- [11] Pollicino T, Squadrito G, Cerenzia G, Cacciola I, Raffa G, Craxi A, et al. Hepatitis B virus maintains its pro-oncogenic properties in the case of occult HBV infection. Gastroenterology 2004;126(1):102–10. https://doi.org/10.1053/j. gastro.2003.10.048.

- [12] Morales-Romero J, Vargas G, García-Román R. Occult HBV infection: a faceless enemy in liver cancer development. Viruses 2014;6(4):1590–611. https://doi.org/ 10.3390/v6041590.
- [13] Walz A, Wirth S, Hucke J, Gerner P. Vertical transmission of hepatitis B virus (HBV) from mothers negative for HBV surface antigen and positive for antibody to HBV core antigen. J Infect Dis 2009;200(8):1227–31. https://doi.org/10.1086/ 605698.
- [14] Pisaturo M, Onorato L, Russo A, Chiodini P, Coppola N. An estimation of the prevalence of occult HBV infection in Western Europe and in Northern America: a meta-analysis. J Viral Hepat 2020;27(4):415–27. https://doi.org/10.1111/ jvh.13248.
- [15] Higgins JP, Thompson SG, Deeks JJ, Altman DG. Measuring inconsistency in metaanalyses. BMJ 2003;327(7414):557–60. https://doi.org/10.1136/bmj.327.7414.557.
- [16] Elrashidy H, El-Didamony G, Elbahrawy A, Hashim A, Alashker A, Morsy MH, et al. Absence of occult hepatitis B virus infection in sera of diabetic children and adolescents following hepatitis B vaccination. Hum Vaccin Immunother 2014;10 (8):2336–41. https://doi.org/10.4161/hv.29521.
- [17] Youssef A, Yano Y, El-Sayed Zaki M, Utsumi T, Hayashi Y. Characteristics of hepatitis viruses among Egyptian children with acute hepatitis. Int J Oncol 2013;42 (4):1459–65. https://doi.org/10.3892/ijo.2013.1822.
- [18] Jaramillo CM, de La Hoz F, Porras A, di Filippo D, Choconta-Piraquive LA, Payares E, et al. Characterization of hepatitis B virus in Amerindian children and mothers from Amazonas State, Colombia. PLoS One 2017;12(10):e0181643. https://doi. org/10.1371/journal.pone.0181643.
- [19] Gachara G, Magoro T, Mavhandu L, Lum E, Kimbi HK, Ndip RN, et al. Characterization of occult hepatitis B virus infection among HIV positive patients in Cameroon. AIDS Res Ther 2017;14(1):11. https://doi.org/10.1186/s12981-017-0136-0.
- [20] Hsu HY, Chang MH, Ni YH, Chiang CL, Wu JF, Chen HL, et al. Chronologic changes in serum hepatitis B virus DNA, genotypes, surface antigen mutants and reverse transcriptase mutants during 25-year nationwide immunization in Taiwan. J Viral Hepat 2017;24(8):645–53. https://doi.org/10.1111/jvh.12687.
- [21] Chen ZX, Gu GF, Bian ZL, Cai WH, Shen Y, Hao YL, Zhang S, et al. Clinical course and perinatal transmission of chronic hepatitis B during pregnancy: A real-world prospective cohort study. J Infect 2017;75(2):146–54. https://doi.org/10.1016/j. jinf.2017.05.012.
- [22] Seremba E, Van Geertruyden JP, Ssenyonga R, Opio CK, Kaducu JM, Sempa JB, et al. Early childhood transmission of hepatitis B prior to the first hepatitis B vaccine dose is rare among babies born to HIV-infected and non-HIV infected mothers in Gulu, Uganda. Vaccine 2017;35(22):2937–42. https://doi.org/10.1016/j.vaccine.2017.04.020.
- [23] Lai MW, Lin TY, Liang KH, Lin WR, Yeh CT. Hepatitis B viremia in completely immunized individuals negative for anti-hepatitis B core antibody. Medicine (Baltimore) 2016;95(49):e5625. https://doi.org/10.1097/MD.00000000005625.
- [24] Rodríguez Lay LLA, Bello Corredor M, Montalvo Villalba MC, Chibás Ojeda AG, Sariego Frómeta S, Diaz González M, et al. Hepatitis B virus infection assessed 3 to 18 years after vaccination in Cuban children and adolescents born to HBsAg-positive mothers. Arch Virol 2017;162(8):2393–6. https://doi.org/10.1007/s00705-017-3365-6.
- [25] Amponsah-Dacosta E, Lebelo RL, Rakgole JN, Selabe SG, Gededzha MP, Mayaphi SH, et al. Hepatitis B virus infection in post-vaccination South Africa: occult HBV infection and circulating surface gene variants. J Clin Virol 2015;63:12–7. https:// doi.org/10.1016/j.jcv.2014.11.032.
- [26] Yao QQ, Dong XL, Wang XC, Ge SX, Hu AQ, Liu HY, et al. Hepatitis B virus surface antigen (HBsAg)-positive and HBsAg-negative hepatitis B virus infection among mother-teenager pairs 13 years after neonatal hepatitis B virus vaccination. Clin Vaccine Immunol 2013;20(2):269–75. https://doi.org/10.1128/CVI.00539-12.
- [27] Said ZN, El-Sayed MH, El-Bishbishi IA, El-Fouhil DF, Abdel-Rheem SE, El-Abedin MZ, et al. High prevalence of occult hepatitis B in hepatitis C-infected Egyptian children with haematological disorders and malignancies. Liver Int 2009;29 (4):518–24. https://doi.org/10.1111/j.1478-3231.2009.01975.x.
- [28] Ghaziasadi A, Fakhari Z, Aghcheli B, Poortahmasebi V, Farahmand M, Norouzi M, et al. High prevalence of occult hepatitis B infection (OBI) among healthy children and their parents in Alborz province, Iran; vertical OBI, myth or truth? Liver Int 2020;40(1):92–100. https://doi.org/10.1111/liv.14252.
- [29] Nageh SM, Hend MA. High prevalence of occult hepatitis B virus infection among frequently blood transfused children: a single Egyptian center experience. Int J Pediatr 2020;8(12):12523-32. https://doi.org/10.22038/ijp.2020.52174.4147.
- [30] Dapena M, Figueras C, Noguera-Julian A, Fortuny C, de José MI, Mellado MJ, et al. Implementation of occult hepatitis screening in the Spanish cohort of HIVinfected pediatric patients. Pediatr Infect Dis J 2013;32(9):e377–9. https://doi. org/10.1097/INF.0b013e31828e9b99.
- [31] Shahmoradi S, Yahyapour Y, Mahmoodi M, Alavian SM, Fazeli Z, Jazayeri SM. High prevalence of occult hepatitis B virus infection in children born to HBsAg-positive mothers despite prophylaxis with hepatitis B vaccination and HBIG. J Hepatol 2012;57(3):515–21. https://doi.org/10.1016/j.jhep.2012.04.021.
- [32] El-Sayed MH, Said ZNA, Abo-Elmagd EK, Ebeid FSE, Salama II. High risk of HBV infection among vaccinated polytransfused children with malignancy. J Pediatr Hematol Oncol 2021;43(1):e45–50. https://doi.org/10.1097/ MPH.000000000001887.
- [33] Elefsiniotis IS, Tsoumakas K, Papadakis M, Vlachos G, Saroglou G, Antsaklis A. Importance of maternal and cord blood viremia in pregnant women with chronic hepatitis B virus infection. Eur J Intern Med 2011;22(2):182–6. https://doi.org/ 10.1016/j.ejim.2010.12.005.
- [34] Barfi S, Narges C, Pouretemad HR, Poortahmasebi V, Norouzi M, Farahmand M, et al. Measurement of serum hepatitis B surface antibody levels in Iranian autistic children and evaluation of immunological memory after booster dose injection in

comparison with controls. J Med Virol 2019;91(7):1272-8. https://doi.org/ 10.1002/jmv.25429.

- [35] Zhou S, Li T, Allain JP, Zhou B, Zhang Y, Zhong M, et al. Low occurrence of HBsAg but high frequency of transient occult HBV infection in vaccinated and HBIGadministered infants born to HBsAg positive mothers. J Med Virol 2017;89 (12):2130-7. https://doi.org/10.1002/jmv.24861.
- [36] Villar LM, Amado LA, de Almeida AJ, de Paula VS, Lewis-Ximenez LL, Lampe E. Low prevalence of hepatitis B and C virus markers among children and adolescents. Biomed Res Int 2014;2014:324638. https://doi.org/10.1155/2014/324638.
- [37] Lin X, Yang J, Lu H, Zhou Y, Zhou G, Wu H, et al. Minimization of hepatitis B infection among children in Jiangsu, China, 12years after integration of hepatitis B vaccine into the expanded program on immunization. Vaccine 2016;34(51):6458– 63. https://doi.org/10.1016/j.vaccine.2016.11.022.
- [38] Qi X, Dai J, Wang X, Wang M, Wang Y. Molecular evolutionary characteristics of OBI virus S gene among the adolescent population in rural and pastoral areas of Xinjiang Province. Infect Genet Evol 2023;107:105395. https://doi.org/10.1016/j. meegid.2022.105395.
- [39] Aghasadeghi MR, Aghakhani A, Mamishi S, Bidari-Zerehpoosh F, Haghi Ashtiani MT, Sabeti S, et al. No evidence of occult HBV infection in population born after mass vaccination. Wien Med Wochenschr 2020;170(9-10):218–23. https://doi. org/10.1007/s10354-020-00748-z.
- [40] Shaker O, Ahmed A, Abdel Satar I, El Ahl H, Shousha W, Doss W. Occult hepatitis B in Egyptian thalassemic children. J Infect Dev Ctries 2012;6(4):340–6. https://doi. org/10.3855/jidc.1706.
- [41] Srivastava A, Mathias A, Yachha SK, Aggarwal R. Occult hepatitis B infection in children with chronic liver disease. Eur J Gastroenterol Hepatol 2015;27(4):375– 7. https://doi.org/10.1097/MEG.0000000000294.
- [42] Hung WL, Wu JF, Ni YH, Chen HL, Chiang CL, Chang MH, et al. Occult hepatitis B virus and surface antigen mutant infection in healthy vaccinated cohorts and children with various forms of hepatitis and multiple transfusions. Liver Int 2019;39 (6):1052–61. https://doi.org/10.1111/liv.14076.
- [43] Beykaso G, Mulu A, Giday M, Berhe N, Selamu M, Hailu D, et al. Occult hepatitis B virus infection and its risks of cryptic transmission in Southern Ethiopia. Infect Drug Resist 2022;15:619–30. https://doi.org/10.2147/IDR.S344668.
- [44] Minuk GY, Sun DF, Uhanova J, Zhang M, Caouette S, Nicolle LE, et al. Occult hepatitis B virus infection in a North American community-based population. J Hepatol 2005;42(4):480–5. https://doi.org/10.1016/j.jhep.2004.11.037.
- [45] Su H, Zhang Y, Xu D, Wang B, Zhang L, Li D, et al. Occult hepatitis B virus infection in anti-HBs-positive infants born to HBsAg-positive mothers in China. PLoS One 2013;8(8):e70768. https://doi.org/10.1371/journal.pone.0070768.
- [46] Foaud H, Maklad S, Mahmoud F, El-Karaksy H. Occult hepatitis B virus infection in children born to HBsAg-positive mothers after neonatal passive-active immunoprophylaxis. Infection 2015;43(3):307–14. https://doi.org/10.1007/s15010-015-0733-6.
- [47] Mu SC, Lin YM, Jow GM, Chen BF. Occult hepatitis B virus infection in hepatitis B vaccinated children in Taiwan. J Hepatol 2009;50(2):264–72. https://doi.org/ 10.1016/j.jhep.2008.09.017.
- [48] Yokoyama K, Kumagai H, Takahashi M, Nagashima S, Okamoto H, Yamagata T. Occult hepatitis B virus infection in immunized children born to carrier mothers. Pediatr Int 2017;59(9):1010–6. https://doi.org/10.1111/ped.13352.
- [49] Hsu HY, Chen HL, Wu JF, Ni YH, Chang KC, Chiang CL, et al. Occult hepatitis B virus infection in immunized infants born to untreated and tenofovir-treated highly viremic mothers. Clin Gastroenterol Hepatol 2021;19(7):1494–6. https://doi.org/ 10.1016/j.cgh.2020.07.041.
- [50] Hu AQ, Cai QY, Zhang M, Liu HY, Wang TL, Han WH, et al. Overt and occult hepatitis B infection after neonatal vaccination: mother-to-infant transmission and HBV vaccine effectiveness. Int J Infect Dis 2021;104:601–9. https://doi.org/10.1016/j. ijid.2021.01.045.
- [51] Su H, Shao Z, Pu Z, Wang Y, Zhang L, Zhang W, et al. Overt and occult hepatitis B virus infection among community children in Northwest China. J Viral Hepat 2017;24(9):797–803. https://doi.org/10.1111/jvh.12709.
- [52] Zhuge S, Ge C, Yang Y, Cui Y, Yue X, Zhang Z, et al. The prevalence of occult HBV infection in immunized children with HBsAg-positive parents: a hospital-based analysis. Hepatol Int 2020;14(4):503–12. https://doi.org/10.1007/s12072-020-10055-9.
- [53] Liu Y, Wen J, Chen J, Xu C, Hu Y, Zhou YH. Rare detection of occult hepatitis B virus infection in children of mothers with positive hepatitis B surface antigen. PLoS One 2014;9(11):e112803. https://doi.org/10.1371/journal.pone.0112803.
- [54] Chen SJ, Zhao YX, Fang Y, Xu WZ, Ma YX, Song ZW, et al. Viral deletions among healthy young Chinese adults with occult hepatitis B virus infection. Virus Res 2012;163(1):197–201. https://doi.org/10.1016/j.virusres.2011.09.029.
 [55] Marjani A, Garshasbi S, Khanaliha K, Kahyesh-Esfandiary R, Dehghani-Dehej F,
- [55] Marjani A, Garshasbi S, Khanaliha K, Kahyesh-Esfandiary R, Dehghani-Dehej F, Babaei R, et al. Screening of occult hepatitis B and C virus infection in working children, Tehran, Iran. Arch Pediatr Infect Dis 2022;10(4):e118763. https://doi. org/10.5812/pedinfect-118763.
- [56] Utsumi T, Yano Y, Lusida MI, Amin M, Soetjipto, Hotta H, et al. Serologic and molecular characteristics of hepatitis B virus among school children in East Java, Indonesia. Am J Trop Med Hyg 2010;83(1):189–93. https://doi.org/10.4269/ ajtmh.2010.09-0589.

- [57] Raouf HE, Yassin AS, Megahed SA, Ashour MS, Mansour TM. Seroprevalence of occult hepatitis B among Egyptian paediatric hepatitis C cancer patients. J Viral Hepat 2015;22(2):103–11. https://doi.org/10.1111/jvh.12260.
- [58] Eilard A, Andersson M, Ringlander J, Wejstål R, Norkrans G, Lindh M. Vertically acquired occult hepatitis B virus infection may become overt after several years. J Infect 2019;78(3):226–31. https://doi.org/10.1016/j.jinf.2019.01.002.
- [59] Weis N, Cowan S, Hallager S, Dröse S, Kristensen LH, Grønbæk K, et al. Vertical transmission of hepatitis B virus during pregnancy and delivery in Denmark. Scand J Gastroenterol 2017;52(2):178–84. https://doi.org/10.1080/ 00365521.2016.1244704.
- [60] Chakvetadze C, Roussin C, Roux J, Mallet V, Petinelli ME, Pol S. Efficacy of hepatitis B sero-vaccination in newborns of African HBsAg positive mothers. Vaccine 2011;29(16):2846–9. https://doi.org/10.1016/j.vaccine. 2011.01.101.
- [61] Pande C, Sarin SK, Patra S, Kumar A, Mishra S, Srivastava S, et al. Hepatitis B vaccination with or without hepatitis B immunoglobulin at birth to babies born of HBsAg-positive mothers prevents overt HBV transmission but may not prevent occult HBV infection in babies: a randomized controlled trial. J Viral Hepat 2013;20(11):801–10. https://doi.org/10.1111/jvh.12102.
- [62] Li AY, Liu Z, Song Y, Xiao Y, Jiang J, Li L, et al. Reduction of the occurrence of occult HBV infection in infants by increasing the dose of hepatitis B vaccine: a large prospective cohort study. Emerg Microbes Infect 2020;9(1):1881–91. https://doi.org/ 10.1080/22221751.2020.1808533.
- [63] Hoffmann CJ, Mashabela F, Cohn S, Hoffmann JD, Lala S, Martinson NA. et al Maternal hepatitis B and infant infection among pregnant women living with HIV in South Africa. J Int AIDS Soc 2014;17(1):18871. https://doi.org/10.7448/ IAS.17.1.18871.
- [64] Hsu HY, Chang MH, Ni YH, Chiang CL, Wu JF, Chen HL. Universal infant immunization and occult hepatitis B virus infection in children and adolescents: a population-based study. Hepatology 2015;61(4):1183–91. https://doi.org/10.1002/ hep.27650.
- [65] World Health Organization. Hepatitis B vaccines: WHO position paper, July 2017 recommendations. Vaccine 2019;37(2):223–5. https://doi.org/10.1016/j.vaccine.2017.07.046.
- [66] Johannessen A, Mekasha B, Desalegn H, Aberra H, Stene-Johansen K, Berhe N. Mother-to-child transmission of hepatitis B virus in Ethiopia. Vaccines (Basel) 2021;9(5):430. https://doi.org/10.3390/vaccines9050430.
- [67] Li Y, Li L, Song Y, Liu M, Zhai X, Duan Z, et al. Booster vaccination in infancy reduces the incidence of occult HBV infection in maternal HBsAg-positive children. J Clin Transl Hepatol 2023;11(3):661–9. https://doi.org/10.14218/ ICTH.2022.00213.
- [68] Ji DZ, Pang XY, Shen DT, Liu SN, Goyal H, Xu HG. Global prevalence of occult hepatitis B: a systematic review and meta-analysis. J Viral Hepat 2022;29(5):317–29. https://doi.org/10.1111/jvh.13660.
- [69] Kajogoo VD, Swai SS, Gurung S. Prevalence of occult hepatitis B among HIV-positive individuals in Africa: a systematic review and meta-analysis. SAGE Open Med 2022;10:20503121211072748. https://doi.org/10.1177/ 20503121211072748.
- [70] Xie WY, Sun C, He H, Deng C, Sheng Y. Estimates of the prevalence of occult HBV infection in Asia: a systematic review and meta-analysis. Infect Dis (Lond) 2022;54(12):881–96. https://doi.org/10.1080/23744235.2022.2115126.
- [71] Su XL, Luo HJ, Yao J, Xia DY, Hao MQ, Chen WX, et al. HBV infections among HIVexposed infants in China. Int J Virol 2017;24(1):26–9. https://doi.org/10.3760/ cma.j.issn.1673-4092.2017.01.007.
- [72] Sarmati L, Malagnino V. HBV infection in HIV-driven immune suppression. Viruses 2019;11(11):1077. https://doi.org/10.3390/v11111077.
- [73] Lukhwareni A, Gededzha MP, Amponsah-Dacosta E, Blackard JT, Burnett RJ, Selabe SG, et al. Impact of lamivudine-based antiretroviral treatment on hepatitis B Viremia in HIV-coinfected South Africans. Viruses 2020;12(6):634. https://doi. org/10.3390/v12060634.
- [74] Cohen Stuart JW, Velema M, Schuurman R, Boucher CA, Hoepelman AI. Occult hepatitis B in persons infected with HIV is associated with low CD4 counts and resolves during antiretroviral therapy. J Med Virol 2009;81(3):441–5. https://doi. org/10.1002/jmv.21422.
- [75] Mitsumoto-Kaseida F, Murata M, Takayama K, Toyoda K, Ogawa E, Furusyo N, et al. Prevalence and characteristics of occult hepatitis B virus infection in Japanese human immunodeficiency virus-infected patients. J Infect Chemother 2020;26(1):28–32. https://doi.org/10.1016/j.jiac.2019.06.003.
- [76] Sagnelli E, Pisaturo M, Martini S, Filippini P, Sagnelli C, Coppola N. Clinical impact of occult hepatitis B virus infection in immunosuppressed patients. World J Hepatol 2014;6(6):384–93. https://doi.org/10.4254/wjh. v6.i6.384.
- [77] Urbani S, Fagnoni F, Missale G, Franchini M. The role of anti-core antibody response in the detection of occult hepatitis B virus infection. Clin Chem Lab Med 2010;48(1):23–9. https://doi.org/10.1515/CCLM.2010.002.
- [78] Ondigui JLN, Kenmoe S, Kengne-Ndé C, Ebogo-Belobo JT, Takuissu GR, Kenfack-Momo R, et al. Epidemiology of occult hepatitis B and C in Africa: a systematic review and meta-analysis. J Infect Public Health 2022;15(12):1436–45. https:// doi.org/10.1016/j.jiph.2022.11.008.