Seroprevalence of hepatitis E virus infection in pregnant women: a systematic review and meta-analysis

Tauseef Ahmad, ^a Jin Hui, ^a Taha Hussain Musa, ^a Masoud Behzadifar, ^b Mukhtiar Baig^c

From the *Department of Epidemiology and Health Statistics, Southeast University, Nangjing, Jiangsu, China; *Hepatitis Research Center, Lorestan University of Medical Sciences, Khoram-Abad, Iran; *Department of Clinical Biochemistry/Medical Education, King Abdulaziz University, Jeddah, Saudi Arabia

Correspondence: Dr. Tauseef Ahmad Department of Epidemiology and Health Statistics, Southeast University, Nangjing, Jiangsu 210009, China hamdard_ hu@yahoo.com ORCID: https://orcid.org/0000-0001-8793-273X

Citation: Ahmad T, Hui J, Hussain M, Behzadifar M, Baig M. Seroprevalence of hepatitis E virus infection in pregnant women: a systematic review and meta-analysis. Ann Saudi Med 2020; 40(2): 136-146. DOI: 10.5144/0256-4947.2020.136

Received: July 27, 2019

Accepted: January 10, 2020

Published: April 2, 2020

Copyright: Copyright © 2020, Annals of Saudi Medicine, Saudi Arabia. This is an open access article under the Creative Commons Attribution-NonCommercial-NoDerivatives 4.0 International License (CC BY-NC-ND). The details of which can be accessed at http:// creativecommons. org/licenses/bync-nd/4 0/

Funding: This study was funded by Chinese National Natural Fund (81573258) and Jiangsu Provincial Six Talent Peak (WSN 002). The funder had no role in study design, data collection and writing. **BACKGROUND:** Hepatitis E virus (HEV) infection has emerged as a global public health problem that affects millions of people every year. **OBJECTIVE:** Systematically review data on the prevalence of HEV IgG antibody among pregnant women around the world.

DATA SOURCES: Potentially relevant studies were identified by a search of PubMed and ScienceDirect, and by a manual search of the reference lists of identified studies.

STUDY SELECTION: Observational studies in English with no age or area restriction. Reviews, duplicate, book chapters, and other irrelevant studies were excluded.

DATA EXTRACTION: Independent searching by two investigators (TA, THM).

DATA SYNTHESIS: In the 6137 retrieved studies, 15 studies met the inclusion criteria. The studies included 7160 pregnant subjects from 11 countries. Most studies were from Africa. Of the 7160 subjects, 1182 were positive to anti-HEV IgG antibody, and only 66 were anti-HEV IgM antibody positive. The highest seroprevalence of anti-HEV IgG antibody (61.29%) was reported in Sudan and the lowest (3.41%) was reported in Italy. The overall pooled prevalence was 16.51% (95% CI: 0.10-0.23). The heterogeneity level was I²= 98%; P≤.01.

CONCLUSION: The seroprevalence of anti-HEV IgG antibody among pregnant women differs by geographic location. Further studies are recommended to evaluate incidence, morbidity, and mortality in those areas where the disease is prevalent.

LIMITATIONS: Seroprevalence was only determined for the anti-HEV IgG antibody, which mostly indicates past infection. Heterogeneity was high among the studies in the analysis.

CONFLICT OF INTEREST: None.

nfection with the hepatitis E virus (HEV) is a significant public health problem that occurs in both developing and industrialized countries. Every year, an estimated 20 million cases of HEV are reported worldwide. In 2015, the World Health Organization (WHO) reported approximately 44000 deaths attributed to HEV infection worldwide, accounting for 3.3% of mortality due to viral hepatitis. The mortality rate ranges from 0.2 to 4%, but can be significantly higher in at-risk groups such as young children, those with pre-existing liver disease, and pregnant women. In pregnant women, the mortality rate reaches 10% to 25% and largely occurs in the third trimester.

HEV is an icosahedral, non-enveloped, singlestrand, positive-sense RNA virus with a 7.2kb genome.8,9 According to the 9th Report of the International Committee on the Taxonomy of Viruses (ICTV), HEV is classified in the family Hepeviridae, 10,11 and includes two genera, Orthohepevirus and Piscihepevirus.11 The genus Orthohepevirus, which infect mammals and birds,14 contains four species, A, B, C and D.11,12 Orthohepevirus A species are isolated from humans, pigs, deer, mongeese, rabbits, wild boars and camels; Orthohepevirus B is isolated from chicken; Orthohepevirus C is isolated from rats, Asian musk shrews, ferrets, greater bandicoots, and minks; and Orthohepevirus D is isolated from bats. 11,13 Species A contains eight genotypes (HEV1, HEV2, HEV3, HEV4, HEV5, HEV6, HEV7 and HEV8).14,15 Genotypes HEV1 and HEV2 are obligate human pathogens. 15,16 HEV3, HEV4 are endemic in several animal species, causing zoonotic infections in humans. 15,16 Genotypes HEV5 and HEV6 appear to be restricted to wild boars, and genotypes HEV7 and HEV8 have been isolated from dromedary and Bactrian camels. 15,16 In addition, a case of HEV7 has been reported in a human.¹⁷

The incubation period of HEV ranges from 2 to 10 weeks.² Symptoms of HEV infection include anorexia, fever, jaundice, myalgia, abdominal pain, back pain, rash, arthralgia, nausea, and vomiting.^{2,3,18} HEV infection is not clinically distinguishable from other types of acute viral hepatitis.² HEV infection is responsible for 30% to 70% cases of acute sporadic hepatitis,¹⁹ and is one of the major causes of acute liver failure.²⁰ HEV is mainly transmitted through the fecal-oral route due to contaminated drinking water,^{9,21} and zoonotic transmission.²¹ Foodborne transmission has also been documented.²² Other uncommon routes of HEV transmission have been documented such as vertical transmission,⁷ and blood-borne transmission.²³⁻²⁵

HEV can be diagnosed by detecting anti-HEV antibodies (IgM and IgG) or RNA-based tests for the

detection of HEV RNA in biological specimens such as liver biopsy, serum, and stool. ^{26,27} In the past two decades, two recombinant vaccines have been developed by GlaxoSmithKline (Belgium) ²⁸ and Xiamen Innovax Biotech (China). ²⁹ The only licensed vaccine, HEV 239 Hecolin, has been approved by China but is not yet available commercially. ^{30,31} To reduce the number of cases of acute and chronic HEV infection, improvements in preventive measures and control strategies have to be made. ³² This meta-analysis aimed to scrutinize the burden and pooled prevalence of HEV IgG antibody in pregnant women around the world to inform researchers and policymakers.

SUBJECTS AND METHODS

Study protocol

During December 2018 and January 2019, we performed a systematic search of the published literature on HEV infection in pregnant women. Well-defined and clear criteria were set before conducting the search. This study was conducted according to the proposed protocol following the "Preferred Reporting Items for Systematic Reviews and Meta-analyses (PRISMA),"33 so that all the steps were conducted independently by two investigators and discrepancies were discussed and resolved by the third investigator.

Study selection

Observational studies published in 2015 to 2018 on HEV infection in pregnant women around the world were included in this study.³⁴ Only articles published in the English language were included. There was no age nor area restriction. Excluded were reviews, duplicate, book chapters, and other irrelevant studies. Studies with human immunodeficiency virus (HIV) patients or with co-infections, and studies whose required seroprevalence data were not accessible even after a request to the authors were also excluded.

Article search strategies

The search was carried out by two investigators (TA, THM) independently on PubMed and ScienceDirect. The following keywords; "Hepatitis E virus" AND "pregnant women" OR "pregnancy" were used (**Table 1**). We also reviewed and searched the relevant articles from the selected studies manually.

Outcome

Seroprevalence of anti-HEV IgG antibody in pregnant women was the outcome of interest. Country, and chronological time were investigated as predictors of

the outcome. The seroprevalence was calculated by the following formula.

Seroprevalence (%) = HEV-infected women

Total number of pregnant women

The operational definition of seroprevalence was 'the presence of anti-HEV antibody (IgG) in the plasma/ serum of pregnant women by the ELISA method'.

Source of data and extraction

Included studies on seroprevalence were from China (n=3), Iran (n=2), Ethiopia (n=2), Benin (n=1), Burkina Faso (n=1), Cameroon (n=1), Eritrea (n=1), Italy (n=1), Ghana (n=1), Nigeria (n=1) and Sudan (n=1) (**Figure 1**). The data from included studies were extracted by two investigators (TA, THM) independently, and disagreements and discrepancies were resolved via discussion with the third investigator. The following parameters were extracted for each selected study; author last name(s), country, publication year, study design, sample size, anti-HEV antibody (IgG and IgM), and anti-HEV antibody detection method. The data was entered into Microsoft Excel 2016.

Quality assessment

To assess the quality of the included studies, we used methodological guidelines from the Joanna Briggs Institute (JBI), University of Adelaide.³⁵ The JBI appraisal checklist is used for prevalence studies to assess the possibility of bias in study design, conduct, and analysis. The JBI appraisal checklist is composed of nine items, and the sources of bias are scored either

yes or no. Three investigators independently scored each study for quality assessment and disagreements were discussed between the investigators until a consensus was reached; however, the quality score was not used in the selection of studies for inclusion in the analysis. Studies that obtained scores from 0 to 3 are presented as high risk of bias, 4 to 6 are at moderate risk of bias, and 7 to 9 are at low risk of bias in study design, conduct, and analysis. Publication bias was calculated using a funnel plot and the Egger test.

Statistical analysis

The data were statistically analyzed by using R software for Windows, Version 3.6.0 (for overall pooled prevalence and subgroup analysis), and Stata software for Windows, Version 12.00 (for sensitivity analysis, meta-regression analysis by year of publication and sample size, and publication bias). The pooled prevalence and 95% CI were calculated based on a random effects model, taking into account the possibility of heterogeneity among the included studies. Heterogeneity was assessed using the I² statistic, with value ranges from 25% to 50%, 51% to 75%, and >75%, representing low, medium, and high heterogeneity between studies, respectively. Due to the high heterogeneity (P<.10) among the included studies, a random effect model was used. Meta-regression analysis was performed to examine the following factors; sample size (continuous variable), and year of study publication (continuous variable). In addition, a sensitivity analysis was also performed by excluding each study to examine whether a single study might influence the study results.36

Table 1. Search terms and history for studies on hepatitis E virus infection in pregnant women.

Database	Step 1	Yielded documents	Step 2	Assessed for eligibility
PubMed	Link: https://www.ncbi.nlm.nih. gov/pubmed/ Keys: (Hepatitis E virus[Title/ Abstract] AND pregnant women[Text Word]) AND pregnancy[Text Word])	163	(Hepatitis E virus[Title/Abstract] AND pregnant women[Text Word]) AND pregnancy[Text Word] AND ((Clinical Study[ptyp] OR Observational Study[ptyp] OR Letter[ptyp] OR Journal Article[ptyp]) AND ("2015/01/01"[PDAT]: "2018/12/31"[PDAT]) AND "humans"[MeSH Terms])	39
Science Direct	Link: https://www.sciencedirect. com/search/advanced Keys: *Hepatitis E virus* AND *Pregnant women* AND *Pregnancy*	5952	Filtered the documents	245
Manual search	Searched from the included studies references	22	Filtered the references of eligible studies	17

RESULTS

Search results and characteristics of included studies

Initially, we identified a total of 6137 articles, of which 5952 were identified from ScienceDirect, 163 from PubMed, and 22 articles were identified manually from the references of included studies. Of the total identified articles, 301 articles remained after screening for eligibility. After screening by title and abstract, we selected 15 studies for the meta-analysis.37-51 Articles were excluded on the basis of the following: review articles (n=251); book chapters (n=381); topics other than HEV infection in pregnant women such as HEV infection among blood donors, animal studies, duplicate, published before 2015, insufficient information (n=5193); and other languages (n=11) as described in Figure 2. The 15 studies reported on 8337 subjects, of which 1175 subjects were non-pregnant women and were subtracted from the total sample size. In addition, in another study 2 pregnant women were excluded from the analysis by the authors based on invalid results.⁵⁰ Finally, the analysis included data from 7160 pregnant women, and the sample size ranged from 93 to 1331 per study (Table 2).

Prevalence of HEV IgG antibody according to region

A study conducted in 2015 in Sudan reported a high seroprevalence of HEV IgG antibody (61.29%) in pregnant women, 40 while a study from Italy, in 2015, reported the lowest prevalence (3.41%), findings consistent with national data.⁴⁵ Another two studies from Ethiopia also reported a high prevalence of HEV IgG antibody among pregnant women (42.43%⁴⁹ and 31.61%⁴⁶), respectively, as compared to other included studies. In Eritrea, a study conducted in 2016, reported the prevalence of HEV IgG antibody was 26.8%.47 In Benin, a study conducted in 2011, 22.3% of total samples were HEV IgG positive, of which 45 (16.19%) were confirmed as positive by immunoblotting (if samples were positive, the test was repeated).41 Among the three reported studies from China, an increase was observed a reported 11.07% seroprevalence of HEV IgG antibody in 2009-2010,39 16.26% in 2011-2013,38 and 21.78% in 2013-2014.48 In Burkina Faso, a study conducted in 2014 reported that HEV IgG prevalence was 10.61%.44 In Iran, studies conducted in 2010-2011 and 2016-2017 reported 7.43%³⁷ and 6.24%,⁵¹ HEV IgG prevalence, respectively. In Cameroon, a study conducted in 2016, reported 9% seroprevalence of HEV IgG antibody.⁴² A study from Nigeria reported 9.89% HEV IgG prevalence.⁴³ In Ghana, HEV IgG prevalence was reported as 12.06%,50 (Table 3).

Overall pooled prevalence among pregnant

All the included studies were pooled for meta-analysis as presented on the forest plot (**Figure 3**). The estimated pooled prevalence was 16.51% (95% CI: 0.10-0.23). The heterogeneity level among the included studies was reported high. The I statistics (I²=98%) represents a high and statistically significant heterogeneity level (*P*<.01) (**Figure 3**). HEV IgG antibody seroprevalence ranged from 3.41% (95% CI: 0.02-0.06) to 61.29% (95% CI: 0.51-0.71) (**Table 3 and Figure 3**). Of the total includ-

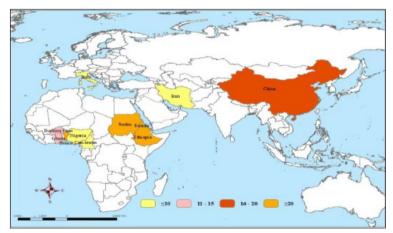


Figure 1. The prevalence of HEV IgG antibody among pregnant women around the world (≤10; 11-15%; 16-20%, ≥20%).

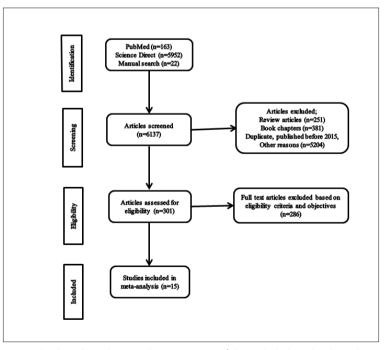


Figure 2. Flow chart showing the processing of the included studies based on PRISMA guidelines.

 Table 2.
 Characteristics of studies in the meta-analysis (n=15).

Year S	S	Year Sampling year	Country	Pregnant women (n)	Urban (n)	Rural (n)	Age (years)	lgG positive (n)	IgM positive (n)	Study type	Detection method	Ref.
2015		2010-2011	Iran	1050	725	325	27.2 (5.6)	78	0	Prospective cross-sectional	DiaPro	37
2015		2009-2010	China	497	138	359	<25 ->35	55	က	Cohort	Santa Cruz	39
2015		2011-2013	China	066	545	445	18-43	161	56	Case-control observational	Wantai	38
2016		2014	Burkina Faso	179	Ω	174	<25 -35	19	0	Cross-sectional	Creative Diagnostic	44
2016		2015	Sudan	93	16	41	15-45	27	0	Cross-sectional	Soronno	37
2016		2011	Benin	278	0	278	15-41	62	7	ΣΖ	DiaPro	41
2016		2016	Cameroon	200	200	0	16-41	18	0	Cross-sectional	Prestige Diagnostics	42
2016		2016	Nigeria	182	ΣΖ	Σ Z	18-45	18	0	ΣZ	Euroimmun	43
2017		2016	Eritrea	153	Σ Z	ΣΖ	15-44	41	0	Cross-sectional Descriptive	Euroimmun	47
2017		2015	Italy	352	352	0	18-≥36	12	0	Hospital based	DiaPro	45
2017		2014-2015	Ethiopia	386	386	0	16-40	122	2	Hospital based	Wantai	46
2018		2013-2014	China	225	Σ Z	ΣΖ	20-35	49	∞	Case-control observational	Wantai	48
2018		2014-2016	Ethiopia	846	356	490	18-50	359	œ	Cross-sectional	Wantai	49
2018		ΣZ	Ghana	398	341	27	28.0 (5.9)	48	—	Cross-sectional	Tangshan	20
2018		2016-2017	Iran	1331	1331	0	14-45	83	11	Descriptive cross-sectional	DiaPro	21

Age data are mean (SD) or range. NM: not mentioned...

Table 3. Prevalence rate and quality score of the studies in the meta-analysis by prevalence (n=15).

Study	Sample size (n)	HEV IgG positive (n)	Prevalence (%)	Quality score (average)	Ref.
Musa	93	57	61.29	7	37
Niguse	846	359	42.43	8	49
Abebe	386	122	31.61	8	46
Tekeste	153	41	26.80	6	47
De Paschale	278	62	22.30	7	41
Rui	225	49	21.78	7	48
Cong	990	161	16.26	7	38
Obiri-Yeboah	398	48	12.06	8	50
Gu	497	55	11.07	7	39
Florence	179	19	10.61	6	44
Alkali	182	18	9.89	6	43
Noufensi	200	18	9.00	6	42
Mamani	1050	78	7.43	8	37
Farshadpour	1331	83	6.24	8	51
La Fauci	352	12	3.41	6	45

ed studies, only 7 studies reported IgM antibody, and the pooled rate was 1.33% (95% CI: 0.01-0.02), I²=77% (**Additional file: Figure S1**).

Subgroup analysis

HEV IgG seroprevalence among pregnant women by

region was high in Africa 22% (95% CI: 0.13-0.34) as compared to other regions (**Figure 4**). The prevalence rate was increased by publication year, 11% (95% CI: 0.08-0.16) in 2015 and 17% (95% CI: 0.08-0.33) in 2018 (**Figure 5**). This increase may be attributed due to different factors, such as studies from different geo-

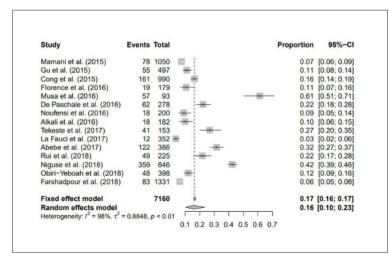


Figure 3. Forest plot meta-analysis of HEV IgG antibody prevalence in pregnant women around the world.

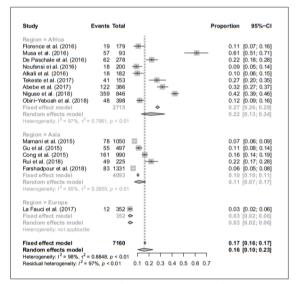


Figure 4. Forest plot of subgroup analysis of HEV IgG antibody prevalence by region.

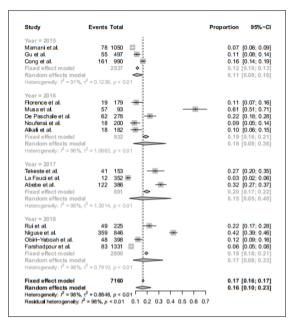


Figure 5. Forest plot of subgroup analysis of HEV IgG antibody prevalence by publication year.

graphical regions, different sample sizes, and different detection methods. The pooled HEV IgG antibody seroprevalence by different ELISA assays showed high variability among included studies ranging from 8% (95% CI: 0.04-0.15) to 27% (95% CI: 0.18-0.38). Dia Pro and Wantai were frequently used commercial assays (**Figure 6**). The seroprevalence of HEV IgG antibody varied with sample size (**Figure 7**). The methodological quality scores of the included studies are presented in **Table 3**. Meta-regression by year of publication (*P*=.582) and sample size (*P*=.003) were calculated (**Additional file 1: Figure S2 and S3**).

Publication bias

The publication bias was determined using the funnel plot, which showed a dispersed distribution suggesting the possibility of publication bias (**Figure 8**) as indicated by the Egger test (P=.007).

Sensitivity analysis

We also performed a sensitivity analysis by removing one study with a large sample size.⁵¹ The overall HEV IgG antibody pooled rate was 17% (95% CI: 0.11-0.25), with I^2 =98%, $P \le .01$ (Additional file 1: Figure S4 and S5).

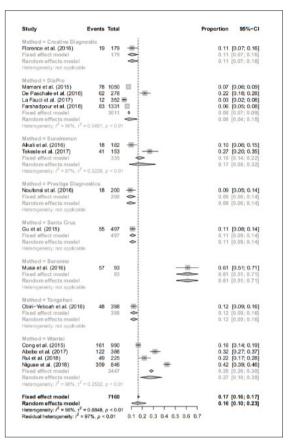


Figure 6. Forest plot of subgroup analysis of HEV IgG antibody prevalence by commercial assay method.

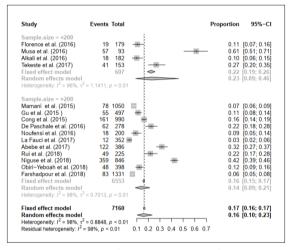


Figure 7. Forest plot of subgroup analysis of HEV IgG antibody prevalence by sample size.

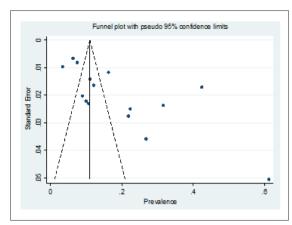


Figure 8. Funnel plot to assess publication bias.

DISCUSSION

HEV is an old and underestimated infection, due to inappropriate diagnosis and lack of awareness among physicians. ^{52,53} In recent years, both in developing and developed countries, surveillance and seroprevalence-based research has gained more attention. However, there are significant gaps in the published literature. Therefore, this study aimed to estimate the burden and pooled prevalence of HEV IgG antibody in pregnant women around the world. The information in this report may help researchers and policymakers to better understand the disease distribution in order to launch effective control strategies and preventive measures.

HEV infection is one of the leading causes of fetal and maternal death.⁵⁴ In pregnant women with HEV infection, cytokine gene polymorphisms are associated with adverse pregnancy outcomes.⁵⁵ During the second and third trimester of pregnancy, HEV infection leads to hepatic failure and increases the risk of mortality 30% to 100%.⁷ However, the mechanism of liver injury has not been clarified. Additional research is needed to explore the exact mechanism and determine preventive strategies. A possible reason may relate to the interplay of immunological and hormonal changes that along with high viral load render women more vulnerable to HEV infection.⁵⁶

According to our findings, the prevalence of HEV IgG antibody ranged from 3.4%⁴⁵ to 61.2%.⁴⁰ The observed discrepancies of HEV seroprevalence may be attributable to different assay methods and different geographic regions. Keeping the above factors in mind, we conducted a subgroup analysis by year of publica-

tion, study region, sample size, and assay method. The meta-analysis of the included studies showed that the estimated pooled prevalence of HEV IgG antibody among pregnant women around the world was 16.51%. These results suggest that HEV IgG antibody is endemic in pregnant women.

In Chinese blood donors, the estimated pooled prevalence of HEV IgM antibody was 30%.57 However, the finding of our study is higher than a systematic review conducted by Iranian authors reported 5.4% seroprevalence of HEV.58 Another meta-analysis conducted among the general population of Iran reported the overall pooled prevalence of HEV was 9.7%.⁵⁹ The estimated seroprevalence of HEV ranged from 0.6% to 52.5% in European countries. 60 Another systematic review of reports of the Brazilian population reported an overall seroprevalence of HEV infection of 6.0%.61 Moreover, the pooled prevalence in our study is also higher than that in some primary studies conducted in different countries among pregnant women, such as in Tunisia of 5.1%,62 Mexico of 5.7%,63 France of 7.7%,64 Pakistan of 8.86%,65 Sudan of 10.3%,66 and in Serbian blood donors of 15.0%.67 On the other hand, a high prevalence of HEV was reported among pregnant women in Egypt of 83.4%, 68 and India of 60.0%. 69 In addition, some other studies reported a high prevalence of HEV IgG antibody in Iran, with 46.1% in the adult population.70 In a study conducted in German blood donors, only 3 of 13 HEV RNA positive results had detectable IgM antibody titers.⁷¹ Spread of HEV through contaminated blood products remains unknown.⁷¹

This meta-analysis has a relatively large sample size but several limitations. Most of the studies used different ELISA kits for the detection of anti- HEV IgG antibody with different specificity and sensitivity, which may affect the reliability and accuracy of the test. Seroprevalence was only determined by the anti-HEV IgG antibody level, which shows mostly after infection. Furthermore, the studies included in the analysis were observational and varied ub baseline characteristics, sample size, and year of sampling.

The consequences of HEV infection— severe liver injury, and high morbidity and mortality rate, especially in pregnant women, have gained the attention of more scientists and researchers. As a result, more focus has been paid to the pathophysiology and immunology of HEV interaction during pregnancy. But it is also important to study genetic and environmental factors, espe-

cially in HEV endemic areas. Immunological research and preventive and management strategies need to be improved to control and stop the disease in the near future. Also, cost-effective vaccination programs are needed in those countries where the disease is endemic.

In conclusion, the seroprevalence of anti-HEV IgG antibody among pregnant women differs by geographic location and is highest in African countries. Further studies are recommended to evaluate the incidence, morbidity, and mortality in those areas where the disease is prevalent. However, necessary actions should be taken to control and prevent HEV infection in gen-

eral and particularly in pregnant women. Special care must be taken in traveling to endemic areas, especially in strictly following drinking water (only bottled water) and food precautions.

Acknowledgments

The authors are very grateful to the US National Library of Medicine and National Institute of Health for open access to PubMed, and we also thanked the ScienceDirect database for free access to the articles. We are also thankful to the reviewers for their valuable comments and suggestions.

REFERENCES

- **1.** Murrison LB, Sherman KE. The Enigma of Hepatitis E Virus. J Gastroenterology Hepatol. 2017; 13:484-91.
- 2. World Health Organization. Hepatitis E: fact sheet. https://www.who.int/news-room/fact-sheets/detail/hepatitis-e. Updated 8 July 2019. Accessed 30 October, 2019.
- **3.** Kamar N, Bendall R, Legrand-Abravanel F, Xia N, Ijaz S, Izopet J, et al. Hepatitis E. Lancet 2012; 379:2477-88.
- **4.** Sharapov MB, Favorov MO, Yashina TL, Brown MS, Onischenko GG, Margolis HS, et al. Acute viral hepatitis morbidity and mortality associated with hepatitis E virus infection: Uzbekistan surveillance data. BMC Infect Dis. 2009:9:35.
- **5.** Teshale EH, Howard CM, Grytdal SP, Handzel TR, Barry V, Kamili S, et al. Hepatitis E epidemic, Uganda. Emerg Infect Dis. 2010;16:126-9.
- **6.** Acharya SK, Sharma PK, Singh R, Mohanty SK, Madan K, Jha JK, et al. Hepatitis E virus (HEV) infection in patients with cirrhosis is associated with rapid decompensation and death. J Hepatol. 2007; 46:387-94.
- 7. Kumar A, Beniwal M, Kar P, Sharma JB, Murthy NS. Hepatitis E in pregnancy. Int J Gynaecol Obstet 2004; 85:240-4.
- **8.** Debing Y, Moradpour D, Neyts J, Gouttenoire J. Update on hepatitis E virology: implications for clinical practice. J Hepatol. 2016; 65:200-12.
- **9.** Yugo DM, Meng XJ. Hepatitis E virus: foodborne, waterborne and zoonotic transmission. Int J Environ Res Public Health. 2013; 10:4507-33.
- **10.** Meng XJ, Anderson DA, Arankalle VA, Emerson SU, Harrison TJ, Jameel S, et al. Hepeviridae. In: King AMQ, Adams MJ, Carstens EB, Lefkowitz EJ, eds. Virus Taxonomy, 9th Report of the ICTV. London: Elsevier Academic Press 2012; 1021-1028.
- 11. Smith DB, Simmonds P, Jameel S, Emerson SU, Harrison TJ, Meng XJ, et al, International Committee on the Taxonomy of Viruses Hepeviridae Study Group. Consensus proposals for classification of the family Hepeviridae. J Gen Virol. 2014; 95(Pt 10):2223.
- **12.** Sridhar S, Teng J, Chiu TH, Lau S, Woo P. Hepatitis E virus genotypes and evolution: emergence of camel hepatitis E variants. Int J Mol Sci. 2017;18.
- **13.** Pérez?Gracia MT, Suay?García B, Mateos?Lindemann ML. Hepatitis E and

- pregnancy: current state. Rev Med Virol. 2017; e1929.
- **14.** Purdy MA, Harrison TJ, Jameel S, Meng XJ, Okamoto H, Van der Poel WHM, et al, ICTV Report Consortium. ICTV Virus Taxonomy Profile: Hepeviridae. J Gen Virol. 2017; 98:2645-6.
- **15.** Webb GW, Dalton HR. Hepatitis E: an underestimated emerging threat. Ther Adv Infect Dis. 2019; 6:1-18.
- **16.** Primadharsini PP, Nagashima S, Okamot H. Genetic Variability and Evolution of Hepatitis E Virus. Viruses 2019; 11:E456.
- 17. Lee GH, Tan BH, Teo EC, Lim SG, Dan YY, Wee A, et al. Chronic infection with camelid hepatitis E virus in a liver transplant recipient who regularly consumes camel meat and milk. Gastroenterology 2016; 150:355-7.

 18. Dalton HR, Stableforth W, Thurairajah
- **18.** Dalton HR, Stableforth W, Thurairajah P, Hazeldine S, Remnarace R, Usama W, et al. Autochthonous hepatitis E in Southwest England: natural history, complications and seasonal variation, and hepatitis E virus IgG seroprevalence in blood donors, the elderly and patients with chronic liver disease. Eur J Gastroenterol Hepatol. 2008; 20:784-90.
- **19.** Panda SK, Thakral D, Rehman S. Hepatitis E virus. Rev Med Virol. 2007; 17:151-80.
- **20.** Acharya SK, Panda SK, Saxena A, Gupta SD. Acute hepatic failure in India: a perspective from the East. J Gastroenterol Hepatol. 2000; 15:473-9.
- **21.** Meng XJ. Hepatitis E virus: Animal reservoirs and zoonotic risk. Vet Microbiol. 2010; 140:256-65.
- **22.** Meng XJ. Zoonotic and foodborne transmission of Hepatitis E virus. Semin Liver Dis. 2013; 33: 41-9.
- **23.** Fukuda S, Ishikawa M, Ochiai N, Suzuki Y, Sunaga J, Shinohara N, et al. Unchanged high prevalence of antibodies to Hepatitis E virus (HEV) and HEV RNA among blood donors with an elevated alanine aminotransferase level in Japan during 1991–2006. Arch Virol. 2007; 152:1623-35.
- **24.** Sakata H, Matsubayashi K, Takeda H, Sato S, Kato T, Hino S, et al. A nationwide survey for Hepatitis E virus prevalence in Japanese blood donors with elevated alamine aminotransferase. Transfusion. 2008; 48:2568-76.
- **25.** Adlhoch C, Kaiser M, Pauli G, Koch J, Meisel H. Indigenous Hepatitis E virus infection of a plasma donor in Germany. Vox

- Sang. 2009; 97:303-8.
- **26.** Mirazo S, Ramos N, Mainardi V, Gerona S, Arbiza J. Transmission, diagnosis, and management of Hepatitis E: an update. Hepat Med. 2014;6:45-59.
- **27.** Gupta E, Agarwala P. Hepatitis E virus infection: An old virus with a new story!. Indian J Med Microbiol. 2018; 36:317-23.
- **28.** Shrestha MP, Scott RM, Joshi DM, Mammen Jr MP, Thapa GB, Thapa N, et al. Safety and efficacy of a recombinant hepatitis E vaccine. New Engl J Med. 2007; 356:895-9.
- 29. Zhu FC, Zhang J, Zhang XF, Zhou C, Wang ZZ, Huang SJ, et al. Efficacy and safety of a recombinant hepatitis E vaccine in healthy adults: a large-scale, randomised, double-blind placebo-controlled, phase 3 trial. Lancet. 2010; 376:895-902.
- **30.** Wu T, Li SW, Zhang J, Ng MH, Xia NS, Zhao Q. Hepatitis E vaccine development: a 14 year odyssey. Hum Vaccin Immunother. 2012; 8:823-7.
- **31.** Zhang J, Shih JW, Wu T, Li SW, Xia NS. Development of the hepatitis E vaccine: from bench to field. Semin Liver Dis. 2013; 33:79-88
- **32.** Melgaço JG, Gardinali NR, Mello VD, Leal M, Lewis-Ximenez LL, Pinto MA. Hepatitis e: Update on prevention and control. BioMed Res Int. 2018; 5769201.
- **33.** Moher D, Liberati A, Tetzlaff J, Altman DG, The PRISMA Group. Preferred Reporting Items for Systematic Reviews and Metanalyses: The PRISMA Statement. PLoS Med. 2009; 6: e1000097.
- **34.** Mann CJ. Observational research methods. Research design II: cohort, cross sectional, and case-control studies. Emerg Med J 2003; 20:54-60.
- **35.** Munn Z, Moola S, Lisy K, Riitano D, Tufanaru C. Methodological guidance for systematic reviews of observational epidemiological studies reporting prevalence and incidence data. Int J Evid Based Healthc 2015; 13:147-53.
- **36.** Copas J, Shi JQ. Meta-analysis, funnel plots and sensitivity analysis. Biostatistics 2000; 1: 247-62.
- **37.** Mamani M, Zamani M, Hashemi SH, Keramat F. Seroprevalence of antibodies to Hepatitis E virus among pregnant women. Avicenna J Clin Microbiol Infect. 2015; 2: e25339
- 38. Cong W, Sui JC, Zhang XY, Qian AD,

- Chen J, Zhu XQ. Seroprevalence of Hepatitis E virus among pregnant women and control subjects in China. J Med Virol. 2015; 87: 446-50
- **39.** Gu G, Huang H, Zhang L, Bi Y, Hu Y, Zhou YH. Hepatitis E virus seroprevalence in pregnant women in Jiangsu, China, and postpartum evolution during six years. BMC Infect Dis 2015: 15: 560.
- **40.** Musa AO, Osman OH, Jaffer A, Ali AA, Ibrahim MEA, Abuzeid N. Seroprevalence of HEV infection and risk factors among Sudanese pregnant women in Khartoum state. Mediterr J Biosci 2016;1: 83-91.
- **41.** De Paschale M, Ceriani C, Romanò L, Cerulli T, Cagnin D, Cavallari S, Ndayake J, Zaongo D, Diombo K, Priuli G, Viganò P. Epidemiology of hepatitis E virus infection during pregnancy in Benin. Trop Med Int Health 2016; 21:108-13.
- **42.** Noufensi MRY, Kengne M, Njukeng PA, Anong DN, Masebe TM, Tamoufe U, Echelibe H, Goon D, Nwobegahay JM. Seroprevalence and risk factors of Hepatitis E virus infection among pregnant women at the Yaounde central hospital, Cameroon. Microbiol Res Int 2016; 4:50-4.
- **43.** Alkali BR, Bello M, Kabiru M, Shu'aibu AB, A'isha BI, Firdausi A, Bunza NM, Ashcroft OF. Seroprevalence of Hepatitis E virus (HEV) infection in pregnant women in Sokoto state, Nigeria. J Adv Microbiol. 2016; 1:1-5.
- **44.** Florence K, Djeneba O, Charlemagne G, Florencia DW, Dorcas O, Rebeca CT, et al. Hepatitis E in pregnant women at the Saint Camille Hospital of Ouagadougou in Burkina Faso: Prevalence and infection risk factors. Int J Recent Adv Multidiscip Res. 2016; 3:1885-8
- **45.** La Fauci V, Facciolà A, Riso R, Calimeri S, Giudice DL, Squeri R. Seroprevalence of HEV antibodies in a sample of pregnant women in the city of Messina. Ann Ig. 2017; 29: 232-8. **46.** Abebe M, Ali I, Ayele S, Overbo J, Aseffa
- **46.** Abebe M, Ali I, Ayele S, Overbo J, Aseffa A, Mihret A. Seroprevalence and risk factors of Hepatitis E virus infection among pregnant women in Addis Ababa, Ethiopia. PLoS ONE 2017: 12: e0180078.
- 47. Tekeste T. G. Dawoud AA, Abdelbagi TA, Mohamedahmed KA. Seroprevalence of Hepatitis E amongst Pregnant Women in Asmara, Eritrea. Eur Acad Res. 2017; 5:607-17.
 48. Rui Z, Jiang L, Wang X, Yang Y, Peng Y, Ling Y, Zhang W, Fu X, Zhou C, Yang S, Shen Q. Prevalence of hepatitis E virus infection among pregnant women in Zhenjiang, China. Front Lab Med. 2018; 2:116-9.

- **49.** Niguse S, Hailekiros H, Buruh G, Dejene T, Berhe N, Asmelash T. Seroprevalence and risk factors of hepatitis E virus infection among pregnant women attending antenatal care in health facilities of Tigray, northern Ethiopia. J Med Virol. 2018; 90:1364-9.
- **50.** Obiri-Yeboah D, Asante Awuku Y, Adu J, Pappoe F, Obboh E, Nsiah P, Amoako-Sakyi D, Simpore J. Seroprevalence and risk factors for hepatitis E virus infection among pregnant women in the Cape Coast Metropolis, Ghana. PLoS ONE 2018; 13: e0191685.
- **51.** Farshadpour F, Taherkhani R, Ravanbod MR, Eghbali SS, Taherkhani S, Mahdavi E. Prevalence, risk factors and molecular evaluation of hepatitis E virus infection among pregnant women resident in the northern shores of Persian Gulf, Iran. PLoS ONE 2018; 13: e0191090.
- **52.** Taherkhani R, Farshadpour F. Epidemiology of hepatitis E in pregnant women and children in Iran: a general overview. J Clin Trans Hepato. 2016; 4:269-76.
- **53.** Taherkhani R, Farshadpour F. Epidemiology of hepatitis E virus in Iran. World J Gastroenterol. 2016; 22:5143-53.
- **54.** Farshadpour F, Taherkhani, Taherkhani R. Hepatitis E virus infection during pregnancy: the overlooked cause of maternal and fetal mortality. Infect Disord Drug Targets. 2019; 19:334-6.
- **55.** Devi SG, Kumar A, Kar P, Husain SA, Sharma S. Association of pregnancy outcome with cytokine gene polymorphisms in HEV infection during pregnancy. J Med Virol. 2014; 86:1366-76.
- **56.** Fiore S, Savasi V. Treatment of viral hepatitis in pregnancy. Expert Opin Pharmacother 2009;10:2801-9.
- **57.** Wang M, Fu P, Yin Y, He M, Liu Y. Acute, recent and past HEV infection among voluntary blood donors in China: a systematic review and Meta-analysis. PLoS ONE 2016; 11(9):e0161089.
- **58.** Behzadifar M, Lankarani KB, Abdi S, Mirghaed MT, Beyranvand G, Keshavarzi A, Ghoreishinia G, Rezapour A, Behzadifar M. Seroprevalence of hepatitis E virus in Iran: a systematic review and Meta-analysis. Middle East J Dig Dis. 2016; 8:189-200.
- **59.** Hadifar S, Sedighi M, Mostafaei S, Miri A, Amiri H, Abiri R, Babaei F, Kabir K & Moghoofei M. Prevalence of hepatitis E infection in the general population of Iran: a systematic review and meta-analysis. Future Virol. 2017; 12: 227-36.

- **60.** Hartl J, Otto B, Madden RG, Webb G, Woolson KL, Kriston L, et al. Hepatitis E seroprevalence in Europe: a meta-analysis. Viruses. 2016; 8:211.
- **61.** Tengan FM, Figueiredo GM, Nunes AK, Manchiero C, Dantas BP, Magri MC, et al. Seroprevalence of hepatitis E in adults in Brazil: a systematic review and meta-analysis. Infect Dis Poverty 2019; 8:3
- **62.** Neffatti H, Lebraud P, Hottelet C, Gharbi J, Challouf T, Roque-Afonso A-M. Southern Tunisia: A still high endemicity area for hepatitis A. PLoS ONE 2017; 12: e0175887.
- **63.** Alvarado-Esquivel C, Sánchez-Anguiano LF, Hernández-Tinoco J. Hepatitis E virus exposure in pregnant women in rural Durango, Mexico. Ann Hepatol. 2015;13:510-17.
- **64.** Renou C, Gobert V, Locher C, Moumen A, Timbely O, Savary J, Roque-Afonso AM, Association Nationale des Hépato-Gastroentérologues des Hôpitaux Généraux (ANGH). Prospective study of Hepatitis E Virus infection among pregnant women in France. Virol J. 2014; 11:68.
- **65.** Khaskheli MN, Baloch S, Sheeba A, Baloch S. Hepatitis E–A preventable health issue–endangering pregnant women's life and foetal outcomes. J Pak Med Assoc. 2015; 65: 655-9.
- **66.** Elduma AH, Osman WM. Dengue and hepatitis E virus infection in pregnant women in Eastern Sudan, a challenge for diagnosis in an endemic area. Pan Afr Med J. 2014;19:391.
- **67.** Petrovi? T, Lupulovi? D, de Oya NJ, Vojvodi? S, Blázquez AB, Escribano-Romero E, et al. Prevalence of hepatitis E virus (HEV) antibodies in Serbian blood donors. J Infect Dey Ctries. 2014: 8:1322-32.
- **68.** Stoszek SK, Abdel-Hamid M, Saleh DA, Kafrawy SE, Narooz S, Hawash Y, et al High prevalence of hepatitis E antibodies in pregnant Egyptian women. Trans R Soc Trop Med Hyg. 2006;100:95-101.
- 69. Patra S, Kumar A, Trivedi SS, Puri M, Sarin SK. Maternal and fetal outcomes in pregnant women with acute Hepatitis E virus infection. Ann Intern Med. 2007;147:28-33.
- M. Prevalence of hepatitis E virus among adults in south-west of Iran. Hepat Res Treatment. 2015; 759589.
- 71. Vollmer T, Diekmann J, Johne R, Eberhardt M, Knabbe C, Dreier J. Novel approach for detection of hepatitis E virus infection in German blood donors. J Clin Microbiol. 2012; 50:2708-13.

SUPPLEMENT FIGURES:

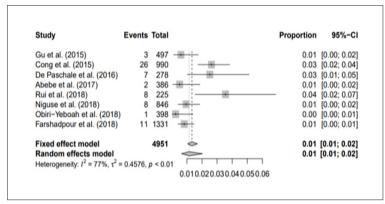


Figure \$1. Forest plot of HEV IgM antibody among pregnant women around the world

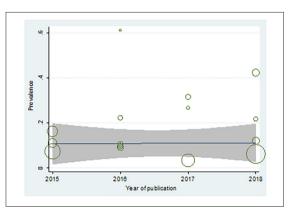


Figure S2. Meta-regression analysis by year of publication (P=.582)

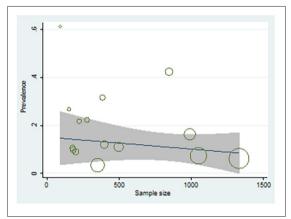


Figure S3. Meta-regression analysis by sample size (P=.003)

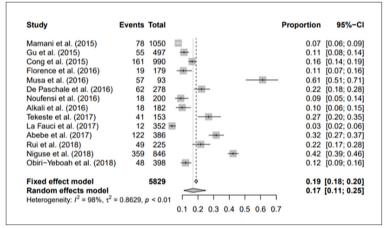


Figure S4. Forest plot of sensitivity analysis.

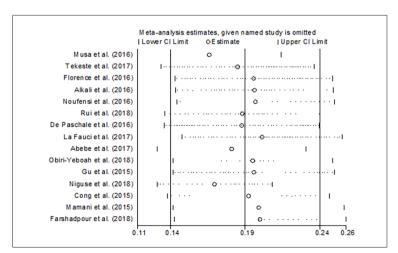


Figure S5. Sensitivity analysis.