

Is cytomegalovirus infection related to inflammatory bowel disease, especially steroid-resistant inflammatory bowel disease? A meta-analysis

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Background: Human cytomegalovirus (HCMV) infection has been associated with inflammatory bowel disease (IBD). Numerous studies have been conducted to analyze the association between HCMV infection and risk of IBD and steroid-resistant IBD, but no clear consensus had been reached.

Objectives: The aim of this study was to confirm this relationship precisely by doing a systematic review and meta-analysis.

Study design: We identified relevant studies through a search of PubMed and Embase. Studies were eligible for inclusion if they 1) evaluated the association between HCMV infection and IBD disease; 2) evaluated the association between HCMV infection and steroid-resistant IBD disease; 3) were case-control studies or nested case-control studies; 4) provided the numbers (or percentage) of positivity for HCMV infection in cases and controls, respectively. Data were extracted and analyzed independently by two investigators.

Results and conclusion: A total of 18 studies including 1,168 patients and 951 health groups was identified, and HCMV infection was distinctly confirmed as a risk factor for the occurrence and development of IBD. When involving 17 studies including 1,306 IBD patients, a total of 52.9% of patients in the cytomegalovirus (CMV)-positive groups were observed to have steroid resistance, compared with 30.2% of patients in the CMV-negative groups. There was a significant difference in the risk of steroid resistance between people exposed to HCMV infection and those not exposed HCMV infection in IBD patients. This meta-analysis suggested that HCMV infection is associated with an increased risk for IBD and steroid-resistant IBD.

Keywords: cytomegalovirus, infection, inflammatory bowel disease, Crohn's disease, ulcerative colitis, meta-analysis

Background

Inflammatory bowel disease (IBD) is a group of chronic inflammatory diseases of the gastrointestinal tract with no specific etiology, and it includes ulcerative colitis and Crohn's disease, and other chronic, nonspecial inflammatory diseases of the gut with unknown etiology. The diagnosis is based on clinical, radiological, endoscopic, and histopathological studies. Several attempts have been made to uncover a possible infectious agent as a cause of IBD. But to date, such efforts have failed to prove any suspected pathogen.

Even though the etiology of IBD is unknown, recent studies have shown that the pathogenesis of IBD is related to susceptible genes, immune dysregulation, and

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microbiota of the gut.¹ IBD is characterized by alternating phases of exacerbation and remission, and cytomegalovirus (CMV) infection is one prominent cause of exacerbation. In addition, CMV infection is an emerging and controversial complication in patients with IBD.² There are many case reports which suggest that IBD patients are particularly vulnerable to CMV infection.

CMV is a β herpes virus with double-stranded DNA. Worldwide, the current infective rate ranges between 40% and 100%. CMV infection is a type of opportunistic infection, as the virus is reactivated and results in active CMV infection with characteristics of viremia and visceral injury (for example pneumonia or colitis) in immunocompromised patients. During the past few decades, abundant observational research has produced data on CMV prevalence in patients with IBD.³ Accordingly, previous studies recommended that CMV infection be suspected in IBD patients with resistance to steroid treatment.⁴ However, it is still uncertain whether CMV infection has an active role in IBD or whether it exists coincidentally.

The prevalence of CMV can be assessed in several ways. Serological CMV IgG testing provides data about latent infection, while IgM testing provides information on recent infection or, as is more likely in adults, reactivation. Other techniques used to demonstrate the virus include immunohistochemistry with antibodies to CMV early antigen or pp65, in situ hybridization to detect CMV mRNA, and polymerase chain reaction (PCR) for CMV DNA,^{5,6} and each has been used to attempt to quantify the infection.

The aims of this study were to clarify the association between CMV infection and IBD and to identify the relationship between CMV infection and steroid-resistant IBD.

Study design

Search strategy

We searched PubMed and Embase and for eligible studies to include in the present meta-analysis, using the terms “human cytomegalovirus” and “inflammation bowel disease” or “Crohn’s disease” or “ulcerative colitis.” An upper date limit of June 10, 2017 was applied; we used no lower date limit. Papers published in English were included. We reviewed citations in the retrieved papers to search for additional relevant studies. The retrieved studies were then read in their entirety to assess their appropriateness for the inclusion in this meta-analysis.

Inclusion/exclusion criteria

Studies were eligible for inclusion if they 1) evaluated the association between human cytomegalovirus (HCMV)

infection and IBD; 2) evaluated the association between HCMV infection and steroid-resistant IBD; 3) were case–control studies or nested case–control studies; 4) and provided the numbers (or percentage) of positivity for HCMV infection in cases and controls, respectively. Conference abstracts, case reports, editorials, review articles, and letters were excluded.

Search selection and data extraction

Two authors (Ya-li Lv and Fei-fei Han) extracted data and reached a consensus on all of the eligibility items according to the aforementioned inclusion criteria. The following data were collected from each study: first author’s surname, the year of publication, source of cases, HCMV detection method, total numbers of cases and controls, and the positivity for HCMV infection in cases and controls or the incidence of steroid resistance in HCMV positive and negative patients, respectively.

All articles reporting data for HCMV prevalence based on enzyme-linked immunosorbent, PCR detection methods, and immunohistochemistry with antibodies to CMV early antigen or pp65 since April 12, 1975, were selected.

Statistical analysis

All statistical analyses were performed using Review Manager (Review Manager 5.0 software; Cochrane, London, England, UK) and Stata/MP 11.0. Cochran’s w^2 test and the inconsistency index (I^2) were used to evaluate heterogeneity across the included studies. Random-effects model was applied in all the analysis. Odds ratio (OR) and corresponding 95% confidence intervals (CIs) were estimated. Z-test was performed to determine the statistical significance of pooled OR, and was considered significant when $P < 0.05$. Statistical heterogeneity test was performed by using the χ^2 and I^2 statistics, and I^2 value of more than 50% was considered as substantial heterogeneity.⁷

Results

Search results

Our search identified 705 potentially relevant studies, 30 of which met the predefined study criteria and were included in our meta-analysis. Figure 1 summarizes the process of study identification, inclusion, and exclusion. There were 245 abstracts from the PubMed search and 460 articles from Embase search. After duplicates were removed, 667 records were assessed for eligibility, and 30 records were finally included in the analysis.^{1,3,8–35}

Characteristics of included studies

The characteristics of included studies and patients are presented in Table 1, including 18 studies, and Table 2 including

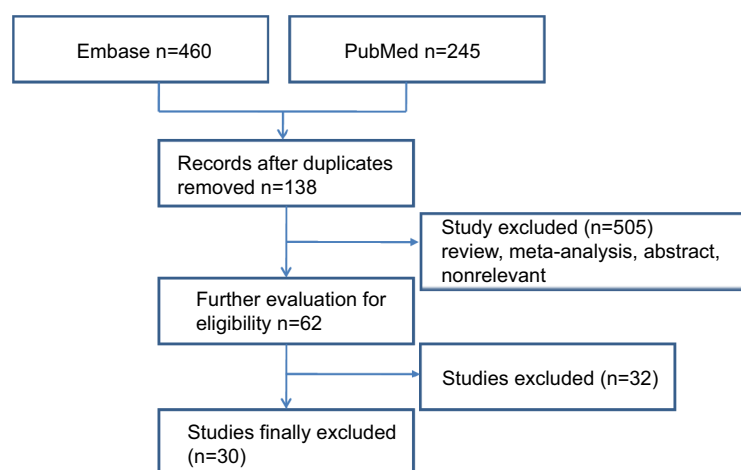


Figure 1 Flow diagram of selected studies.

Table 1 Main characteristics of the studies included in the meta-analysis

Study	Source of cases	Study design type	Disease type	Definition of disease	Detection of HCMV	Case		Control	
						HCMV+	Total	HCMV+	Total
Rahbar et al, ⁸ 2003	Sweden	Case-control	UC and CD (IBD)	WHO	DNA	22	23	0	10
Wakefiels et al, ¹⁰ 1992	United Kingdom	Case-control	IBD	WHO	DNA	36	50	6	21
Verdonk et al, ⁹ 2006	The Netherlands	Case-control	IBD	WHO	pp65	12	31	17	53
Marszatek et al, ¹¹ 2011	Poland	Case-control	IBD	WHO	DNA	13	32	8	15
					IgG	9		5	
					IgM	0		0	
Nahar et al, ¹² 2015	Japan	Case-control	UC	WHO	DNA	26	71	6	188
Domenech et al, ¹³ 2008	Spain	Case-control	UC	WHO	IgG	66	94	19	25
Sipponen et al, ¹⁴ 2011	Finland	Case-control	IBD	WHO	pp65	64	79	6	15
Knosel et al, ¹⁵ 2009	Germany	Case-control	CD	WHO	DNA	2	56	0	10
Lavagna et al, ¹⁶ 2006	Italy	Case-control	UC	WHO	IE	1	24	0	20
					DNA	3			
Dimitroulia et al, ¹⁷ 2006	Greece	Case-control	IBD	WHO	pp65	10	85	0	42
					DNA	23		5	
Van Kruiningen et al, ¹⁸ 2007	USA	Case-control	CD	WHO	DNA	1	70	1	41
Kuwabara et al, ¹⁹ 2007	Japan	Case-control	IBD	WHO	IE	18	34	3	31
Yi et al, ¹ 2013	People's Republic of China	Case-control	IBD	WHO	DNA	190	226	173	290
					IgG	172		147	
					IgM	4		1	
Ciccocioppo et al, ²⁰ 2015	Italy	Case-control	IBD	WHO	IE	9	40	2	40
Ciccocioppo et al, ³⁴ 2016	Italy	Case-control	IBD	WHO	IE	51	64	7	25
Criscuoli et al, ²¹ 2015	Italy	Case-control	UC	WHO	DNA	11	24	2	24
					Pp65	8		2	
Taherkhani et al, ²² 2015	Iran	Case-control	UC	WHO	DNA	12	98	0	67
Thörn et al, ²³ 2016	Sweden	Case-control	IBD	WHO	IgM	14	67	1	34
					DNA	12		0	

Abbreviations: CD, Crohn's disease; HCMV, human cytomegalovirus; IBD, inflammatory bowel disease; IE, immediate-early; UC, ulcerative colitis; WHO, World Health Organization.

17 researches; there were 5 duplicates studies among these, so 30 records were included in the final analysis. Table 1 shows the characteristics and variables collected for the prevalence of CMV infection in IBD disease. And Table 2 displays the characteristics of the relationship between CMV infection and steroid-resistant IBD.

As shown in Table 1, data were available for a total of 1,168 patients and 951 healthy people; containing 14 studies from a Caucasian subgroup, 2 from an Asian subgroup, and 1 from a subgroup of other ethnicities. As can be seen in Table 2, data were available for a total of 1,306 patients, containing 11 studies from a Caucasian

Table 2 Main characteristics of the relationship between CMV infection and steroid-resistant IBD

Study	Source of cases	Study design type	Disease type	Definition of disease	Detection of HCMV	HCMV+		HCMV-	
						Steroid resistant/refractory	Total	Steroid resistant/refractory	Total
Lévêque et al, ²⁴ 2010	France	Case	IBD	WHO	DNA	0	5	8	29
Domènech et al, ¹³ 2008	Spain	Case	UC	WHO	IgG	13	29	6	14
Iida et al, ³ 2013	Japan	Case	UC	WHO	IgG	62	141	20	46
					IgM				
Wada et al, ²⁵ 2003	Japan	Case	UC	WHO	pp65	13	16	9	31
Kambham et al, ²⁶ 2004	USA	Case	UC	WHO	CMV inclusions	12	13	28	67
					IE				
Maconi et al, ²⁷ 2005	Italy	Case	UC	WHO	IE	15	17	40	60
Kuwabara et al, ¹⁹ 2007	Japan	Case	IBD	WHO	IE	13	14	2	8
Maher et al, ²⁸ 2009	Egypt	Case	IBD	WHO	IgM	8	9	15	63
					IE				
Criscuoli et al, ²⁹ 2011	Italy	Case	UC	WHO	CMV inclusions	6	28	11	57
					IE				
					DNA				
					Pp65				
Mccurdy et al, ³⁰ 2015	USA	Case	IBD	WHO	CMV inclusions	40	68	61	202
					IE				
Ciccocioppo et al, ²⁰ 2015	Italy	Case	IBD	WHO	IE	4	9	13	31
Ciccocioppo et al, ³⁴ 2016	Italy	Case	IBD	WHO	IE	25	51	0	13
Cottone et al, ³¹ 2001	Italy	Case	IBD	WHO	DNA	7	7	12	55
Criscuoli et al, ³¹ 2004	Italy	Case	IBD	WHO	pp65	4	9	8	33
					DNA				
Criscuoli et al, ²¹ 2015	Italy	Case	UC	WHO	CMV inclusions	5	11	7	13
					IE				
					DNA				
					pp65				
Kim et al, ³⁵ 2012	Korea	Case	UC	WHO	DNA	14	31	7	41
Ormeci et al, ³³ 2016	Turkey	Case	IBD	WHO	DNA	8	13	5	72

Abbreviations: HCMV, human cytomegalovirus; IBD, inflammatory bowel disease; IE, immediate-early; CMV, cytomegalovirus; UC, ulcerative colitis; WHO, World Health Organization.

subgroup, 4 from an Asian subgroup, and 2 from a subgroup of other ethnicities.

Risk of CMV infection in patients with IBD

Meta-analysis results

A meta-analysis of 18 studies is reported in Table 1 and Figures 2–6, dealing with the association between HCMV infection and IBD disease. When involving 18 studies including 1,168 patients and 951 health groups, a total HCMV positive ratio of 69.6% in IBD patients (247/352) and 51.82% in control groups (171/330) based HCMV IgG tests, 5.5% in IBD patients (18/325) and 0.59% in control groups (28/895)

based HCMV IgM tests, 42.5% in IBD patients (351/826) and 26.4% in control groups (201/762) based HCMV DNA tests, 48.8% in IBD patients (79/162) and 10.3% in control groups (12/116) based on HCMV immediate-early (IE) tests, and 40.4% in IBD patients (92/228) and 6.6% in control groups (8/121) based on HCMV pp65 tests was observed.

Compared with the control groups, people exposed to HCMV infection had higher risk than those not exposed to CMV. The ORs of HCMV infection rate in IBD patients compared to the control groups were, respectively, 7.14 [95% CI =1.58–32.25, $p=0.01$] based on HCMV IgM tests, 4.99 [95% CI =2.40–10.40, $p<0.0001$] based on HCMV DNA tests, 8.43 [95% CI =4.08–17.42, $p<0.00001$] based on

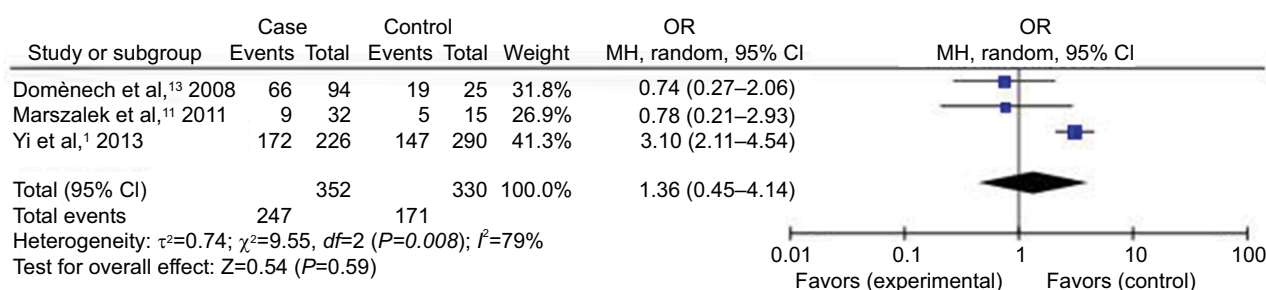


Figure 2 Forest plot (random-effect model) of the association between HCMV infection and IBD stratified by the study design based on HCMV IgG tests.

Abbreviations: CI, confidence interval; HCMV, human cytomegalovirus; IBD, inflammatory bowel disease; MH, Mantel–Haenszel; OR, odds ratio.

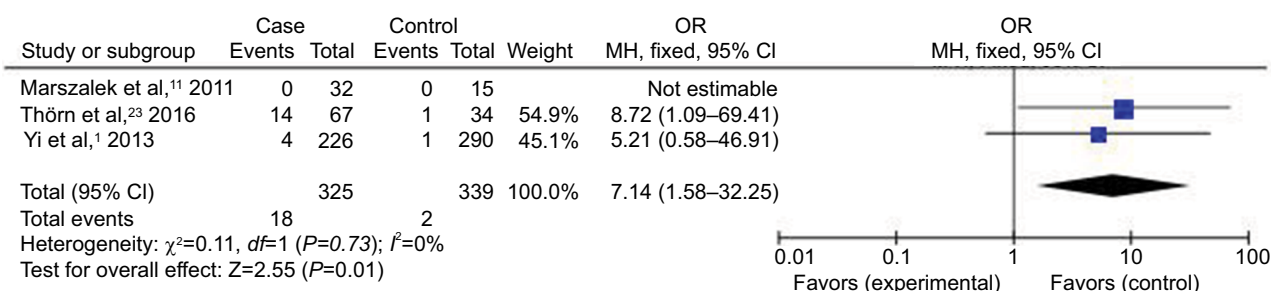


Figure 3 Forest plot (random-effect model) of the association between HCMV infection and IBD stratified by the study design based on HCMV IgM tests.

Abbreviations: CI, confidence interval; HCMV, human cytomegalovirus; IBD, inflammatory bowel disease; MH, Mantel–Haenszel; OR, odds ratio.

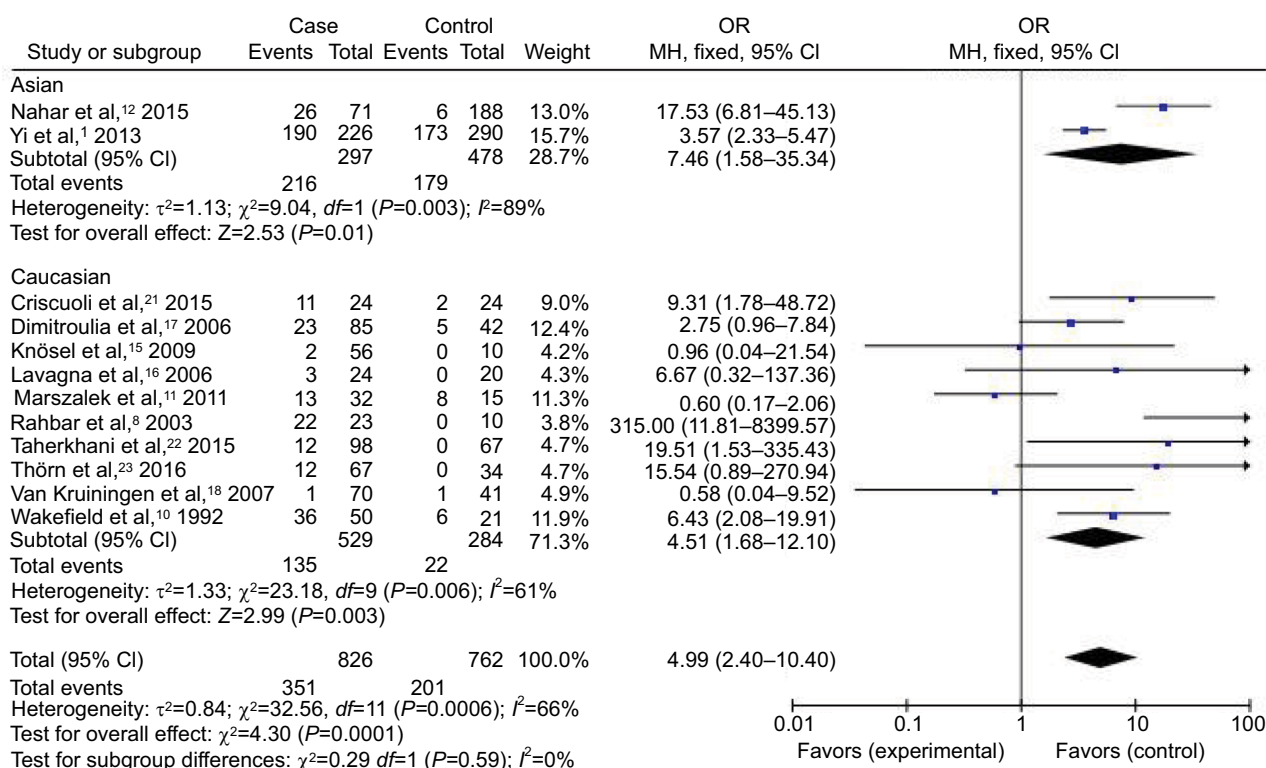


Figure 4 Forest plot (random-effect model) of the association between HCMV infection and IBD stratified by the study design based on HCMV DNA tests.

Abbreviations: CI, confidence interval; HCMV, human cytomegalovirus; IBD, inflammatory bowel disease; MH, Mantel–Haenszel; OR, odds ratio.

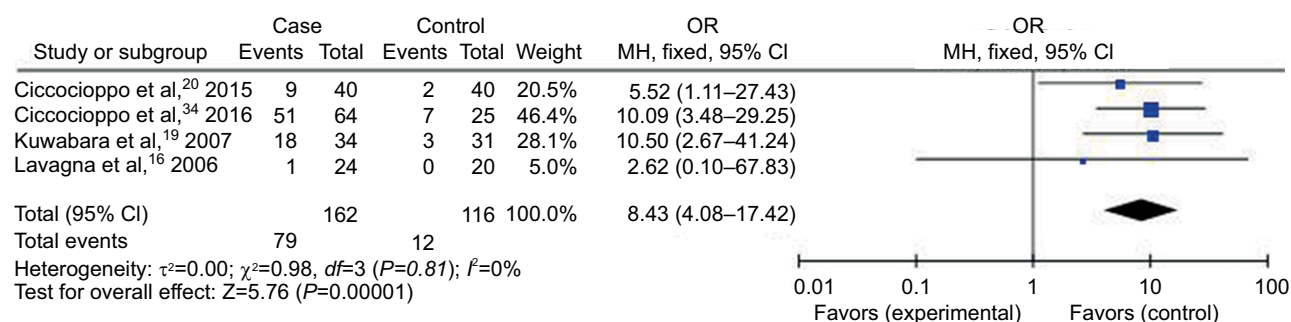


Figure 5 Forest plot (random-effect model) of the association between HCMV infection and IBD stratified by the study design based on HCMV IE tests.

Abbreviations: CI, confidence interval; HCMV, human cytomegalovirus; IBD, inflammatory bowel disease; IE, immediate-early; MH, Mantel–Haenszel; OR, odds ratio.

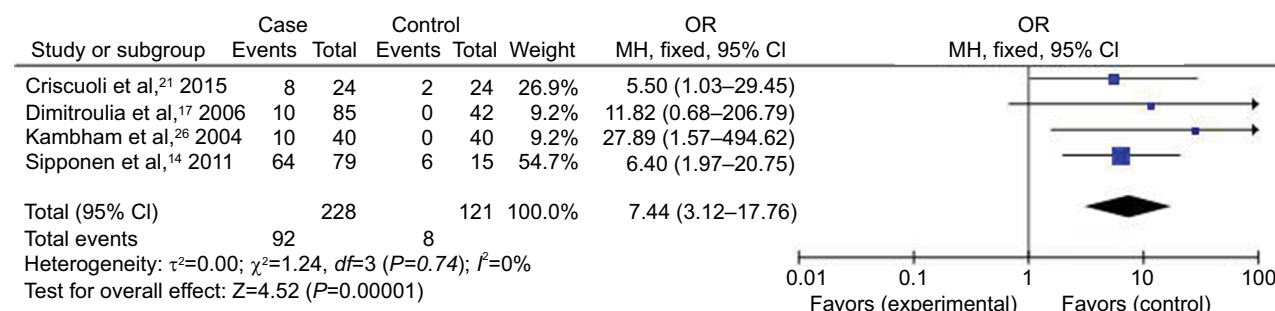


Figure 6 Forest plot (random-effect model) of the association between HCMV infection and IBD stratified by the study design based on HCMV pp65 tests.

Abbreviations: CI, confidence interval; HCMV, human cytomegalovirus; IBD, inflammatory bowel disease; MH, Mantel–Haenszel; OR, odds ratio.

HCMV IE tests, and 7.44 [95% CI=3.12–17.76, $p<0.00001$] based on HCMV pp65 tests. However, there was no difference in the risk of HCMV infection rate in IBD patients compared to the control groups based on HCMV IgG tests [OR=1.36, 95% CI=0.45–4.14, $p=0.59$].

Subgroup analysis

In the stratified analysis by ethnicity, through the experimental results (Figure 4), HCMV infection was distinctly confirmed as a risk factor for the occurrence and development of IBD in Asian and Caucasian groups based on HCMV DNA tests, with ORs of 7.46 [95% CI=1.58–35.34, $p=0.01$] and 4.51 [95% CI=1.68–12.10, $p=0.003$], respectively.

Risk of CMV infection in patients with steroid-resistant IBD

Meta-analysis results

When involving 17 studies including 1,306 IBD patients, a total of 52.9% of patients in the CMV-positive groups were observed to have steroid resistance, compared with 30.2% of patients in the CMV-negative groups. There was a significant difference in the risk of steroid resistance between people exposed to HCMV infection and those not exposed HCMV infection for IBD patients [OR=3.63, 95% CI=1.99–6.62, $p<0.0001$].

Subgroup analysis

In the stratified analysis by ethnicity and CMV detection method, HCMV infection was distinctly confirmed as a risk factor for the occurrence of steroid-resistant IBD through the experimental results (Table 3).

Sensitivity analysis

The purpose of sensitivity analysis is to estimate the stability of the results. When any single study was omitted, the pooled results showed a lack of change. This means that in the overall meta-analysis, no single study substantially affected the stability of the results. Therefore, the results of this meta-analysis were stable and reliable.

Publication bias

The publication bias of the included studies was investigated with the Begg's funnel plot and Egger's test. The results are presented in Figures 7 and 8. The shapes of the funnel plots showed some dissymmetry. Egger's test was used to provide statistical evidence of funnel plot symmetry, and there was no evidence of publication bias.

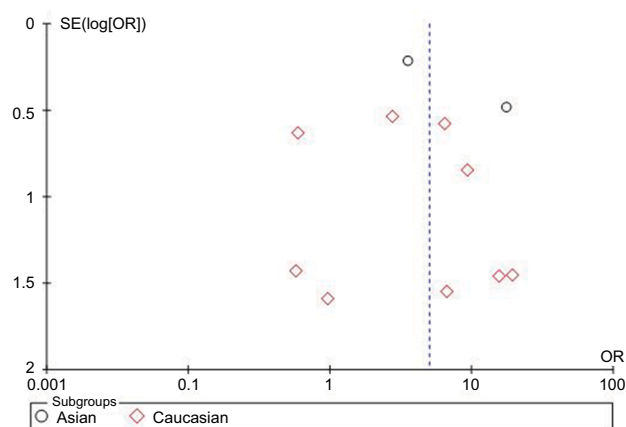
Discussion

CMV is an opportunistic pathogenic microorganism that was discovered in immunocompetent individuals in the 1960s. As

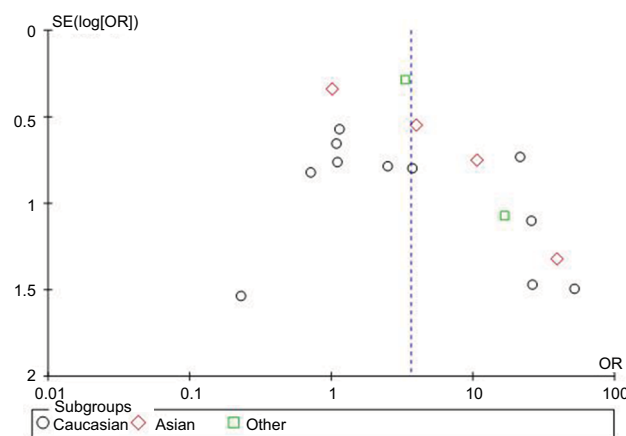
Table 3 Main characteristics of the studies included in the meta-analysis

Characteristics	Subgroup	N (Study)	N (Subject)	OR	95% CI	I ² (%)
All		17	1,306	3.84	2.12–6.97	67
ethnicity	Caucasian	11	220	3.47	1.43–8.41	65
	Asian	4	140	4.75	1.16–19.52	80
	Other	2	141	5.43	1.22–24.17	55
CMV detection	IgG	1	43	1.08	0.30–3.92	Not applicable
	DNA	4	253	6.40	1.13–36.33	71
	IE	4	203	5.86	1.17–29.37	61
	pp65	1	47	10.59	2.42–46.33	Not applicable
	CMV inclusion + IE	2	350	5.43	1.22–24.17	55
	IgG + IgM	1	187	1.02	0.52–2.00	Not applicable
	IgM + IE	1	72	25.60	2.96–221.58	Not applicable
	pp65 + DNA	1	42	2.50	0.54–11.63	Not applicable
	CMV inclusion + IE + DNA + pp65	2	109	0.98	0.39–2.45	0

Abbreviations: CI, confidence interval; CMV, cytomegalovirus; IE, immediate-early; OR, odds ratio.

**Figure 7** Begg's funnel plot of the association between HCMV infection and IBD based on HCMV DNA tests.

Abbreviations: HCMV, human cytomegalovirus; IBD, inflammatory bowel disease; OR, odds ratio; SE, standard error of the mean.

**Figure 8** Begg's funnel plot of the association between HCMV infection and steroid-resistant IBD.

Abbreviations: HCMV, human cytomegalovirus; IBD, inflammatory bowel disease; OR, odds ratio; SE, standard error of the mean.

a β -herpesvirus, it is an important pathogen in many patients and widely exists in the majority of the population. After primary infection, the virus persists in the body and is located mainly in white blood cells and endothelial cells. CMV also has wide tissue tropism and is able to infect multiple organs, including the intestine.³⁶ CMV was first found to be related with IBD in 1961.³⁷ Since then, research on the effect of CMV infection on IBD activation has attracted broad attention and interest among clinicians. In IBD patients, immunosuppressive therapy, impaired absorption of nutrients, and dysfunction of the immune system render them susceptible to CMV infection.³⁸ However, the role of CMV infection in patients with IBD is not confirmed yet. This meta-analysis sought to address unresolved debate regarding the relationship between CMV infection and IBD, especially CMV infection and steroid-resistant IBD.

When analyzing the relationship between CMV infection and IBD, our meta-analysis involved 19 studies including 1,218 patients and 972 healthy groups. Compared with the control groups, people exposed to HCMV infection had higher risk than those not exposed to CMV. Among 19 included studies, anti-CMV IgG, IgM, DNA, IE, and pp65 were used to detect CMV infection in IBD patients and healthy controls. Compared to the control groups, anti-CMV IE and pp65 were significantly higher in IBD patients, and the ORs of HCMV infection rate in IBD patients were, respectively, 8.43 [95% CI =4.08–17.42, $p<0.00001$] and 7.44 [95% CI =3.12–17.76, $p<0.00001$]. This suggested that CMV infection might be nearly 7–8 times higher in IBD patients than in control groups, thereby indicating the association between IBD and CMV. Our results are in

agreement with the previous reports^{39,40} suggesting that immunosuppressive medications might be an important risk factor for CMV infection. Significant heterogeneity of HCMV prevalence was noted when analyzed by ethnic subgroup. Our findings indicated that HCMV infection in Asia was the highest. Positive rates in Asian studies were 72.7% in the IBD population and 37.4% in the control population based on HCMV DNA tests, which might be the cause of the increased risk of IBD in the Asian population. However, there are limited included studies and insufficient participants in our research, so it needs further study in order to validate the relationship between ethnic difference and HCMV infection in IBD risk.

When analyzing the relationship between CMV infection and steroid resistance in IBD patients, our meta-analysis involved 17 studies including 1,306 IBD patients. Compared to CMV-negative groups, the OR in CMV-positive groups was 3.63 [95% CI = 1.99–6.62, $p < 0.0001$], which suggested that CMV infection might cause an approximately fourfold risk of steroid resistance in IBD patients. When analyzed by ethnic subgrouping, the percentage of steroid-resistant patients in the CMV-positive groups was higher than in the CMV-negative groups regardless of the ethnicity. And when analyzed by CMV detection method, the percentage of steroid resistance in CMV-positive groups was also higher than that in CMV-negative groups for the majority of the CMV detection methods. These results thus showed a positive correlation between CMV infection and steroid resistance in IBD patients. Previous studies have also reported that local reactivation of CMV is closely related with severe and steroid-refractory IBD in patients.^{3,41,42}

Previous studies had linked CMV infection with IBD and its resistance to steroids; however, those researches failed to provide a strong and systematic relationship between CMV infection and IBD.^{1,34,42} In this research, we carried out a systematic meta-analysis assessing the effect of CMV infection on IBD and its resistance to steroids. Our results showed that there exists a significant association between CMV infection with IBD and its resistance to steroids. However, our study has several limitations. First, the number of articles included in this research is relatively small, ie, there were only 33 studies. Second, there have been some differences in the method of CMV detection and protocol, this may be because we only included published studies and did not include unpublished ones. An additional limitation in this research may be that there is a potential publication bias. In order to provide a more precise estimation on the basis

of adjustment for confounders, more well-designed studies should be performed. Further prospective controlled trials will help to confirm these findings and identify subgroups of patients who might suffer from CMV infection.

In summary, our meta-analysis strongly suggests that CMV infection might be a probable cause of IBD and its resistance to steroids.

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Author contributions

All authors contributed toward data analysis, drafting and critically revising the paper and agree to be accountable for all aspects of the work.

Disclosure

The authors report no conflicts of interest in this work.

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