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#### ORIGINAL RESEARCH

The effect of antiviral therapy on patients with hepatitis B virus-related hepatocellular carcinoma after curative resection: a systematic review and meta-analysis

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**Background and aim:** Studies suggest that antiviral therapy performed after curative resection improves the postoperative prognosis of hepatitis B virus (HBV)-related hepatocellular carcinoma (HCC), but the evidence has been contradictory. The aim of this meta-analysis was to assess the effect of antiviral therapy with nucleoside analogs (NAs) after curative resection on the long-term postoperative survival of patients with HBV-related HCC.

**Materials and methods:** MEDLINE, PubMed, Embase, and Cochrane Library were systematically searched up to August 2017 with no limits. Outcome measures were the primary parameter of overall survival (OS) after radical resection of HBV-related HCC and the secondary parameter of postoperative recurrence-free survival (RFS).

**Results:** A total of 9,009 patients (2,546 of whom received antiviral therapy and 6,463 received no treatment) were included. The pooled analysis revealed that antiviral therapy was associated with significantly improved OS (hazard ratio [HR]: 0.58; 95% confidence interval [CI]: 0.51–0.67; P<0.00001) and RFS (HR: 0.68; 95% CI: 0.63–0.74; P<0.00001). Moderate heterogeneity among studies for both OS and RFS was observed, which disappeared or decreased after pooling studies using one type of NA as antiviral drug. In the subgroup analysis, antiviral therapy significantly prolonged both OS (HR: 0.69; 95% CI: 0.52–0.92; P=0.01) and RFS (HR: 0.58; 95% CI: 0.49–0.70; P<0.00001) in patients with high baseline HBV DNA level ( $\geq$ 20,000 IU/mL) with no heterogeneity, but not in patients with low baseline HBV DNA level (<20,000 IU/mL).

**Conclusion:** Antiviral therapy with NAs confers significant survival benefits in patients with HBV-related HCC after curative resection, especially in patients with high baseline HBV DNA level ( $\geq 20,000 \text{ IU/mL}$ ).

Keywords: hepatocellular carcinoma, hepatitis B virus, antiviral therapy, recurrence, survival

## Introduction

Hepatocellular carcinoma (HCC) is the third leading cause of cancer-related deaths in the modern world, and its incidence continues to increase.<sup>1</sup> Chronic hepatitis B virus (HBV) infection remains a major cause of HCC development (especially in Asia) either through direct transactivation, viral integration, or indirectly through inflammation, fibrosis, or cirrhosis.<sup>2,3</sup> Growing evidence has shown that antiviral therapy with nucleoside analogs (NAs) can reduce the risk of HCC development in patients with chronic HBV infection.<sup>4,5</sup> According to the current guidelines for the management of HCC, surgical resection should be considered as the first-line treatment for patients

OncoTargets and Therapy 2017:10 5363-5375

5363

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with resectable tumors and preserved liver function.<sup>6,7</sup> With advances in surveillance programs, early diagnosis, and surgical technologies, the long-term postoperative survival of patients with early-stage HCC has improved, but is still unsatisfactory due to the high recurrence rate. Therefore, how to decrease HCC recurrence after curative resection merits further attention.

In patients with resected HBV-related HCC, factors including high viral replication status, active inflammation, subsequent damage, and regeneration of hepatocytes are associated with an increased risk of recurrence and adverse long-term survival outcomes.8-10 Further, sustained low HBV load predicts good long-term recurrence-free survival (RFS) and overall survival (OS).<sup>11</sup> Recently, several pioneering studies used antiviral therapy with NAs (NA therapy) to treat patients with HBV-related HCC after curative resection and assessed the effect of such therapy on the long-term postoperative survival outcomes.12-36 Some trials reported significant postoperative survival benefits, but others failed to confirm such outcomes. Indeed, the previous metaanalyses on this issue did not exclude patients who received NA therapy before the diagnosis of HBV-related HCC, which may affect the natural course of HBV-related HCC.37 Compared to locoregional therapy including radiofrequency ablation (RFA), percutaneous ethanol injection (PEI), and transcatheter arterial embolization (TAE), surgical resection can usually achieve relatively complete elimination of HCC lesions and satisfied margin of normal liver tissue which is very important to avoid potential tumor tissue remnant and potentially benefits the prognosis of patients.<sup>38-40</sup> Patients who received locoregional treatments instead of curative resection for the initial treatment of HCC had not been excluded in the previous meta-analyses, which may have potentially biased the interpretation of survival outcomes.41-45 Otherwise, non-English articles were all excluded in previous meta-analyses, and this search strategy may not be sufficiently comprehensive since HBV infection is the main cause of HCC in Asia-Pacific region, and Chinese, Korean, and Japanese articles may contribute significantly to the meta-analysis. Therefore, to investigate this important issue, we performed a more comprehensive meta-analysis to evaluate the effect of NA therapy after curative resection on the long-term postoperative survival of patients with HBV-related HCC.

# Materials and methods Data sources and search strategy

A systematic literature search was performed up to August 2017 using MEDLINE, PubMed, Embase, and Cochrane

Library with no limits. The search strategy involved the Medical Subject Heading (MeSH) terms: "hepatitis B," "HBV," "antiviral," "nucleotide," "nucleotide analog," "lamivudine," "adefovir," "entecavir," "telbivudine," "hepatocellular carcinoma," "HCC," "liver cancer," "hepatic cancer," "liver resection," "surgical resection," "radical resection," "curative resection," "hepatic resection," and "hepatectomy," combined with free text words. The bibliographies of all retrieved review articles and primary studies were manually searched for more relevant studies. For studies with duplicate publications from the same cohort, the most recent comprehensive publication was included.

## Study selection

All clinical studies, including randomized controlled trials (RCTs) and prospective or retrospective cohort studies, were selected if they met the following criteria: 1) enrolled patients who were diagnosed with HBV-related HCC and underwent curative resection as the initial treatment; 2) enrolled patients underwent no other forms of antitumor therapy before curative resection, such as local ablation therapy, regional or systemic chemotherapy, molecular target therapy, or immunotherapy; 3) consisted of one or more groups treated with NA therapy and an untreated control group; 4) reported at least long-term results of OS or RFS for outcome measures; 5) had been published with full-text accessible.

Studies were excluded if they met one or more of the following criteria: 1) included patients who received NA therapy before the diagnosis of HBV-related HCC; 2) included patients with combined infection of other hepatitis viruses; 3) included patients with drug abuse or alcohol consumption; 4) nonhuman studies, abstracts, editorials, letters, case reports, reviews, and studies not clearly reporting the outcomes of interest.

#### Outcome measures

The primary analysis focused on OS of HBV-related HCC after curative resection, and postoperative RFS served as secondary outcome.

## Data extraction

Parameters regarding the following information were extracted in a standardized data extraction form: 1) study characteristics: reference, year of publication, country of origin, and study design; 2) patient characteristics: sample size, age, gender, hepatitis B e-antigen (HBeAg) status, and Child–Pugh classification; 3) tumor characteristics: tumor size and number; 4) outcomes of the antiviral therapy group and the control group: OS and RFS; 5) potential sources of heterogeneity. Any discrepancy in the extraction process was resolved by discussion and consensus.

# Quality assessment

The quality of each trial was assessed independently by two study investigators (X-XC and X-RY). The Jadad scale was used to score the methodological quality of RCTs based on the following items: randomization (0–2 points), blinding (0–2 points), and dropouts and withdrawals (0–1 point). A modified Newcastle–Ottawa Scale was used to assess non-RCTs across the following three factors: patient selection (0–2 stars), comparability of the cohort (0–2 stars), and outcome assessment (0–2 stars).<sup>46</sup>

# Statistical analysis

The meta-analysis was performed by using Review Manager (version 5.2), which was provided by the Cochrane Collaboration (The Nordic Cochrane Centre, Copenhagen, Denmark). Long-term outcomes (survival analysis) were analyzed by calculating the hazard ratio (HR) with 95% confidence interval (95% CI). HRs of the OS or RFS were calculated and combined using the data extracted from Kaplan–Meier curves; HR <1 represented survival benefits favoring the antiviral therapy group. A random-effect model (DerSimonian and Laird's method) was used to compare the overall effect estimates.

Statistical heterogeneity was explored by the  $\chi^2$  and  $I^2$  statistics.  $I^2 < 25\%$  was considered to reflect low heterogeneity, an  $I^2$  value between 25% and 50% was considered to reflect moderate heterogeneity, and  $I^2 > 50\%$  was considered to reflect high heterogeneity. Heterogeneity was considered statistically significant when the Cochrane Q test P < 0.10. Two-sided value of P < 0.05 was considered statistically significant. A funnel plot was conducted to screen for potential publication bias.

In addition, a sensitivity analysis was performed to assess the effect of individual studies on the pooled estimates. To anticipate potential heterogeneity among the included studies, subgroup analysis was performed for the following study-related variables if the necessary data were provided: 1) NA type; 2) viral load (baseline HBV DNA level  $\geq$ 20,000 IU/mL versus <20,000 IU/mL); 3) fully preserved hepatic function (Child class A).

# Results

## Characteristics of identified studies

A total of 360 potentially relevant studies were identified through database searching and other sources. After detailed

screening, based on the inclusion and exclusion criteria, 25 references involving 26 studies (two RCTs and 24 non-RCTs) were included for the final meta-analysis.<sup>12–36</sup> One study was a two-stage longitudinal clinical study, which included a first-stage non-RCT to assess the effect of NA therapy on the postoperative prognosis of HBV-related HCC and a second-stage RCT to validate the initial result.<sup>20</sup> Furthermore, propensity score matching (PSM) analysis was performed in one retrospective study to reduce patient selection bias, and we only extracted the data after PSM for the current meta-analysis.<sup>15</sup> The detailed study screening and selection process is shown in Figure 1.

A total of 9,009 patients with resected HBV-related HCC were included in the analysis, 2,546 of whom received NA therapy (antiviral therapy group), whereas the other 6,463 patients received no treatment (control group). Table 1 lists the baseline characteristics of the included studies and the main features of the enrolled patients.

# Antiviral therapy and virological response

In the 26 included studies, lamivudine was the most commonly used antiviral drug, followed by entecavir and adefovir. When lamivudine resistance occurred, adefovir was added or entecavir was used instead. Three studies reported HBV DNA suppression rates, and NA therapy was associated with significantly higher HBV DNA suppression rate at 1, 2, 3, and 5 years.<sup>13,14,16,26,29,36</sup> In the antiviral therapy group, the HBV DNA suppression rate ranged from 51.3% to 87.2% at 1 year, from 62.7% to 98.0% at 2 years, from 67.2% to 91.7% at 3 years, and 92.8% at 5 years of NA therapy. The cumulative HBeAg seroconversion rate at 1 year ranged from 12.0% to 57.2% in the antiviral therapy group.<sup>14,16</sup>

# Primary outcome: OS

A total of 17 studies reported the comparative data for postoperative OS. Meta-analysis of these studies revealed that the NA therapy was significantly associated with higher OS (HR: 0.58; 95% CI: 0.51–0.67; P<0.00001; Figure 2A). Moderate heterogeneity was detected in the analysis (P=0.04,  $I^2$ =40%). However, after pooling studies using one type of NA, no heterogeneity was observed within subgroups of studies (Figure 3).

While the subgroup analysis of patients with high baseline HBV DNA level ( $\geq$ 20,000 IU/mL) showed stable results (HR: 0.69; 95% CI: 0.52–0.92; *P*=0.01) with no heterogeneity among studies, the subgroup analysis of patients with low baseline HBV DNA level (<20,000 IU/mL) indicated no significant difference between the antiviral therapy group



Figure I Flowchart of search strategy for meta-analysis study selection.

Abbreviations: HBV, hepatitis B virus; HCC, hepatocellular carcinoma; HDV, hepatitis D virus.

and the control group (HR: 0.80; 95% CI: 0.38–1.68; *P*=0.56; Figure 4A). Stratified meta-analysis of fully preserved hepatic function (Child class A) showed stable results with no heterogeneity (Figure 5A).

# Secondary outcome: RFS

The selected 25 studies reported the comparative data for postoperative RFS. Meta-analysis of these studies showed that patients with NA therapy had a significantly increased RFS after surgery (HR: 0.68; 95% CI: 0.63–0.74; P < 0.00001). Moderate heterogeneity was detected in the analysis (P=0.08,  $I^2=30\%$ ; Figure 2B). However, after pooling studies using one type of NA, only a little heterogeneity was observed within subgroups of studies (Figure 6).

According to subgroup analysis, patients with high baseline HBV DNA level showed stable results (HR: 0.58; 95% CI: 0.49–0.70; P<0.00001) with on heterogeneity among studies, whereas results from patients with low baseline HBV DNA level showed no significant difference between the antiviral therapy group and the control group (HR: 0.86; 95% CI: 0.51–1.46; P=0.57) with moderate heterogeneity among studies (Figure 4B). Stratified meta-analysis of fully preserved hepatic function (Child class A) showed stable results with high heterogeneity (Figure 5B).

# Liver function reserve at HCC recurrence and subsequent treatment for recurrence

Two studies reported significantly better liver function in the antiviral therapy group compared to the control group at the time of HCC recurrence.<sup>13,15</sup> Similarly, several studies reported significantly improved liver function in the antiviral therapy group compared to the control group at 6 months after surgery;<sup>20,25,26,28</sup> Li et al's<sup>16</sup> study also reported a significant residual liver volume improvement in the antiviral therapy group compared to the control group at 6 months after surgery. Two studies reported a significantly higher amenability rate of radical retreatment (eg, surgical resection or local ablation therapy) for HCC recurrence in the antiviral therapy group due to better liver function reserve than that of the control group.<sup>13,15</sup>

# Publication bias

Figure 7 illustrates a funnel plot of the included studies comparing postoperative RFS in HBV-related HCC patients with or without NA therapy. Visual inspection of the funnel plot revealed asymmetry, and Begg's test for publication bias showed statistically significant results which indicated

Study	Study	NA type	Arms	Cases	Age	Gender	HBV DNA level	HBeAg (±)	Tumor	No of	Tumor stage $^{\mathrm{b}}$	Child-Pugh	Follow-up	Quality
	type				(years) <sup>a</sup>	(M/F)	(log copies/mL) <sup>ª</sup>		size (cm) <sup>4</sup>	tumors (S/M)		(A/B/C)	duration (months) <sup>a</sup>	score
Chan et al (2011) <sup>12</sup>	Pro-retro	LAM, ETV	Treatment	42	57	31/11	N/A	N/A	9.3	N/A	11/14/16/0 (AICC)	42/0/0	N/A	4
			Control	94	55	74/20	N/A	N/A	9.0	N/A	28/18/48/0	84/10/0	N/A	
Chen et al	Retro	LAM, ETV	Treatment	192	47	172/20	N/A	48/144	4.7	176/16	(5)(4)(0 (TNM)	192/0/0	74 <sup>d</sup>	4∘
(2016) <sup>22</sup>			Control	253	49	218/35	N/A	35/218	5.4	232/21	215/38/0 (TNM)	253/0/0		
Chen	Pro	LAM	Treatment	45	52.2	39/6	N/A	N/A	N/A	37/8	38/5/2/0 (AJCC)	41/4/0	N/A	•4
(2015) <sup>24</sup>			Control	40	51.9	36/4	N/A	N/A	N/A	35/5	34/5/1/0 (AJCC)	38/2/0	N/A	
Cheng et al	Retro	LAM	Treatment	50	45.3	38/12	5.56 (5.53–5.64)	N/A	N/A	N/A	N/A	N/A	N/A	4°
(2011) <sup>35</sup>			Control	43	42.2	32/11	5.68 (5.50–5.83)	N/A	N/A	N/A	N/A	N/A	N/A	
Chong et al (2015) <sup>13</sup>	Pro-retro	LAM, ETV	Treatment	254	55	222/32	N/A	42/212	3.5	198/56	168/50/34/2 (AJCC)	243/10/1	39.2 (0.2–163.9)	ũ
		ADV, LdT	Control	150	56	125/25	N/A	12/138	3.8	118/32	97/26/27/0	145/4/1	43.3 (0.1–151.2)	
			1	i	ļ						(AJCC)			
Ding et al	Retro	ETV	I reatment	74	47.2	56/18 20/0	5.21	N/A	N/A	N/A	N/A	54/20/0	N/A	4
(2014) <sup>38</sup>			Control	45 7	46.5	30/9	20.5	N/A	A/A	N/A	N/A	31/8/0	N/A	
Fang et al	Retro	LAM	l reatment	26	48.9	20/6	6.38	N/A	N/A	N/A	N/A	N/A	41	4
(2012) <sup>25</sup>			Control	30	50.0	23/7	6.55	N/A	N/A	N/A	N/A	N/A	20	
Huang et al (2015) <sup>14</sup>	RCT	ADV	Treatment	8	50.6	01/06	N/A	51/49	4.9	0/001	8/60/0 (BCLC [0/A/B])	0/0/001	60 (4−70)⁴	ň
			Control	001	50.5	89/11	N/A	50/50	5.1	0/001	8/59/0 (BCLC	0/0/001		
			F	Ļ						<b>V</b> 11 <b>V</b>	[0/A/B])	41,010		, r
пиалу ес аг (2016) <sup>26</sup>	Verto		Control	6 m	A/N	A/N	4.75	A/N	A/N	A/N		33/0/0	-(+c-c)	ł-
Kaatol Kaatol	Ratro	Μ	Treatment	141	48.9	C1/6C1	4 97	15/176	4.5	95/001	11/20/201	141/0/0	74 (7-65)	ŭ
(2013) <sup>15</sup>		-	ווכמתווכוור	Ē	202	711/71		07 101	2		(BCLC [A/B/C])		(20-2) 12	r
			Control	4	49.7	127/14	4.78	16/125	5.0	107/34	105/26/10 (BCLC_FA/B/C1)	141/0/0	23 (1–73)	
Li et al	Pro	LAM, ADV	Treatment	43	46	34/9	6.49 (4.04–7.38)	38/5	7.1	N/A	(FUL) (TNM)	28/14/1	12	4
(2010)16				36	45	30/6	7.27 (3.45–8.32)	27/9	8.5	N/A	4/10/22 (TNM)	21/12/3	12	
Lin et al	Retro	ETV	Treatment	35	54.2	26/9	7.48	N/A	N/A	N/A	N/A	28/7/0	54	4∘
(2016) <sup>27</sup>			Control	25	2.5	20/5	6.94	N/A	N/A	N/A	N/A	20/5/0	38	
Qian et al	Retro	LAM, ADV	Treatment	70	N/A	58/12	6.70	N/A	6.5	N/A	N/A	70/0/0	N/A	•4
(2016) <sup>28</sup>			Control	65	N/A	52/13	6.71	N/A	6.6	N/A	N/A	65/0/0	N/A	
Su et al (2013) <sup>17</sup>	Retro	LAM, ETV	Treatment	62	52	56/6	5.89 (4.83–7.15)	9/53	2.7	48/14	40/15/5 (BCLC [A/B/C])	N/A	N/A	4

Study	Study type	NA type	Arms	Cases	Age (years) <sup>a</sup>	Gender (M/F)	HBV DNA level (log copies/mL) <sup>a</sup>	HBeAg (±)	Tumor size (cm)ª	No of tumors	Tumor stage <sup>b</sup>	Child-Pugh (A/B/C)	Follow-up duration	Quality score
			Control	271	58	232/39	5.48 (4.09–7.83)	28/243	4.2	(M/c) 144/127	142/83/44	N/A	(monens) <sup>-</sup> N/A	
											(BCLC [A/B/C])			
Tian et al	Retro	LAM, ETV	Treatment	29	57.8	18/11	4.7	N/A	2.8	81/11	N/A	25/4/0	28.6 <sup>d</sup>	4∘
(2015) <sup>29</sup>		ADV	Control	21	58.3	14/7	6.1	N/A	3.4	8/13	N/A	18/3/0		
Wang et al	Retro	LAM, ETV	Treatment	76	53.7	58/18	5.46	N/A	6.1	64/12	35/30/11 (TNM)	N/A	30.3	4∘
$(2015)^{30}$		ADV	Control	80	51.3	57/23	5.40	N/A	5.5	71/9	32/28/10 (TNM)	N/A	18.4	
Wei et al	Retro	LAM, ADV	Treatment	86	50.8	79/7	6.9 (2.0–8.8)	14/72	8.0	27/59	0/36/50 (TNM)	82/4/0	30.6 (3.3–73.2) <sup>d</sup>	4∘
(2016) <sup>23</sup>			Control	40	50.0	35/5	5.7 (2.0–6.7)	6/34	8.9	15/25	0/22/18 (TNM)	39/1/0		
Wu et al	Retro	LAM, ETV	Treatment	518	54.4	435/83	N/A	N/A	N/A	N/A	N/A	N/A	32	4∘
(2012) <sup>18</sup>		LdT	Control	4,051	54.6	3,335/716	N/A	N/A	N/A	N/A	N/A	N/A	26	
Xu et al	Retro	N/A	Treatment	29	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	4∘
(2016) <sup>31</sup>			Control	82	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	
Yang et al	Retro	LAM	Treatment	30	22/8	N/A	N/A	N/A	N/A	N/A	27/2/1 (TNM)	29/1/0	26.2 <sup>d</sup>	4∘
(2010) <sup>32</sup>			Control	30	23/7	N/A	N/A	N/A	N/A	N/A	21/7/2 (TNM)	26/4/0		
Yang et al	Pro	LAM, ETV	Treatment	142	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	49.1 <sup>d</sup>	4
(2012) <sup>19</sup>		ADV	Control	I 88	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A		
rao et al	Retro	LAM, ADV	Treatment	76	53.9	54/22	5.43	N/A	N/A	N/A	N/A	N/A	30.4	4
(2016) <sup>33</sup>			Control	80	52.3	55/25	5.40	N/A	N/A	N/A	N/A	N/A	18.5	
*Yin et al	Pro	LAM, ETV	Treatment	215	50.1	194/21	4.51	80/135	N/A	184/31	6/159/50 (BCLC	209/6/0	24 <sup>d</sup>	ñ
(2013a) <sup>20</sup>		ADV	Control	402	50.3	336/66	3.87	100/302	N/A	351/51	[0/A/B]) 6/296/98 (BCLC	392/10/0		
		-									[0/A/B])			
*Yin et al	RCT	LAM, ETV	Treatment	8	47.9	74/7	4.88	41/40	N/A	21/10	4/67/10 (BCLC	80/1/0	40 <sup>d</sup>	3e
(2013b) <sup>20</sup>		ADV	Control	82	49.4	20/12	4.57	26/56	N/A	64/18	[0/A/B]) 2/58/22 (BCLC	82/0/0		
											[0/A/B])			
Zhang	Pro	LAM	Treatment	45	52.2	39/6	N/A	N/A	N/A	37/8	38/5/2 (TNM)	41/4/0	N/A	4∘
(2015) <sup>34</sup>			Control	40	51.9	36/4	N/A	N/A	N/A	35/5	34/5/1 (TNM)	38/3/0	N/A	
Zhang et al	Retro	ETV	Treatment	40	N/A	26/14	N/A	N/A	4.6	40/0	N/A	33/6/1	N/A	ŭ
(2014) <sup>21</sup>			Control	47	N/A	31/16	N/A	N/A	4.8	47/0	N/A	37/10/0	N/A	

Chen et al

5368

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Study or subgroup	Log (HR)	Antiviral SE	therapy Total	Control Total	Weight (%)	HR IV, random, 95% CI	HR IV, random, 9	95% CI
Chan et al (2011) <sup>12</sup>	-0.5798	0.3537	42	94	3.1	0.56 (0.28–1.12)		
Chen et al (2016)22	-0.4463	0.1575	192	253	8.8	0.64 (0.47-0.87)	-	
Chong et al (2015)13	-0.5604	0.1678	254	150	8.3	0.57 (0.41-0.79)	-	
Ding et al (2014)36	-0.6733	0.3638	74	39	2.9	0.51 (0.25-1.04)		
Huang et al (2015)14	-0.6675	0.2084	100	100	6.5	0.51 (0.34-0.77)		
Ke et al (2013)15	-0.6733	0.254	141	141	5.0	0.51 (0.31–0.84)		
Li et al (2010) <sup>16</sup>	-0.3857	0.2221	43	36	6.0	0.68 (0.44-1.05)		
Su et al (2013)17	-1.3863	0.3336	62	271	3.4	0.25 (0.13-0.48)		
Tian et al (2015)29	-0.5351	0.3317	29	21	3.4	0.59 (0.31-1.12)		
Wang et al (2015)30	-0.8062	0.255	76	80	5.0	0.45 (0.27-0.74)		
Wei et al (2016)23	-1.0966	0.324	86	40	3.5	0.33 (0.18-0.63)	_ <b>_</b>	
Wu et al (2012) <sup>18</sup>	-0.3567	0.0959	518	4,051	12.5	0.70 (0.58-0.84)	+	
Yang et al (2012)19	-0.2485	0.1783	142	188	7.8	0.78 (0.55–1.11)		
Yao et al (2016)33	-0.4691	0.1851	76	80	7.5	0.63 (0.44-0.90)		
*Yin et al (2013a)20	-0.2357	0.1319	215	402	10.3	0.79 (0.61-1.02)	-	
*Yin et al (2013b)20	-1.3093	0.3729	81	82	2.8	0.27 (0.13-0.56)		
Zhang et al (2014) <sup>21</sup>	-0.2877	0.3469	40	47	3.2	0.75 (0.38–1.48)	-+-	
Total (95% CI)			2,171	6,075	100	0.58 (0.51–0.67)	•	
Heterogeneity: $\tau^2=0.03$		· · · ·	; /²=40%					
Test for overall effect: 2	Z=7.84 (P<0.00	0001)					0.01 0.1 1	10
							Favors	Favors control

Study or subgroup	Log (HR)	Antiviral SE	therapy Total	Control Total	Weight (%)	HR IV, random, 95% CI	HR IV, random, 95% CI
Chan et al (2011) <sup>12</sup>	-0.4155	0.2306	42	94	2.9	0.66 (0.42-1.04)	
Chen et al (2016)22	-0.0305	0.1245	192	253	6.8	0.97 (0.76–1.24)	+
Chen (2015) <sup>24</sup>	-0.297	0.3001	45	40	1.8	0.74 (0.41–1.34)	
Cheng and Lv (2011) <sup>35</sup>	-0.4082	0.219	50	43	3.1	0.66 (0.43–1.02)	
Chong et al (2015)13	-0.3147	0.1262	254	150	6.7	0.73 (0.57–0.93)	-
Ding et al (2014) <sup>36</sup>	-0.5987	0.3155	74	39	1.7	0.55 (0.30-1.02)	
Fang et al (2012) <sup>25</sup>	-0.9086	0.3812	26	30	1.2	0.40 (0.19–0.85)	
Huang et al (2015)14	-0.399	0.1827	100	100	4.1	0.67 (0.47-0.96)	
Huang et al (2016) <sup>26</sup>	-0.5086	0.2205	45	33	3.1	0.60 (0.39-0.93)	
Ke et al (2013) <sup>15</sup>	-0.1054	0.146	141	141	5.6	0.90 (0.68–1.20)	-+
Li et al (2010)16	-0.2107	0.1793	43	36	4.2	0.81 (0.57–1.15)	
Lin et al (2016)27	-1.1841	0.3932	35	25	1.1	0.31 (0.14-0.66)	
Qian et al (2016)28	-0.685	0.2324	70	65	2.8	0.50 (0.32-0.79)	
Su et al (2013)17	-0.5798	0.1846	62	271	4.0	0.56 (0.39-0.80)	
Tian et al (2015)29	-0.5351	0.3317	29	21	1.5	0.59 (0.31–1.12)	
Wang et al (2015)30	-0.4291	0.1767	76	80	4.3	0.65 (0.46-0.92)	-
Wei et al (2016)23	-0.4005	0.1809	86	40	4.2	0.67 (0.47-0.96)	
Wu et al (2012)18	-0.2744	0.072	518	4,051	10.8	0.76 (0.66–0.88)	-
Xu et al (2016) <sup>31</sup>	-0.6038	0.4061	29	82	1.1	0.55 (0.25–1.21)	
Yang et al (2012)19	-0.6349	0.1436	142	188	5.7	0.53 (0.40-0.70)	-
Yao et al (2016)33	-0.3479	0.1574	76	80	5.1	0.71 (0.52-0.96)	
*Yin et al (2013a) <sup>20</sup>	-0.2485	0.0852	215	402	9.6	0.78 (0.66–0.92)	-
*Yin et al (2013b) <sup>20</sup>	-0.734	0.1759	81	82	4.3	0.48 (0.34–0.68)	<b>-</b>
Zhang (2015) <sup>34</sup>	-0.2942	0.2752	45	40	2.1	0.75 (0.43–1.28)	
Zhang et al (2014) <sup>21</sup>	-0.5108	0.2606	40	47	2.3	0.60 (0.36–1.00)	
Total (95% CI)			2,516	6,433	100	0.68 (0.63–0.74)	
. ,				0,400		0.00 (0.00-0.14)	· · · · · · · · · · · · · · · · · · ·
Heterogeneity: $\tau^2$ =0.01; Test for overall effect: Z			; /-=30%				0.01 0.1 1 10
	·						Favors Favors antiviral control

Figure 2 Forest plots for postoperative survival outcomes.

Notes: (A) Meta-analysis of OS. (B) Meta-analysis of RFS. \*Study of Yin et al (2013)<sup>20</sup> was a two-stage longitudinal clinical study which included a first stage pro study and a second stage RCT. Therefore, the first stage pro study was defined as Yin et al (2013a)<sup>20</sup> and the second stage RCT was defined as Yin et al (2013b)<sup>20</sup>. Abbreviations: CI, confidence interval; HR, hazard ratio; IV, inverse variance; OS, overall survival; RFS, recurrence-free survival; SE, standard error.

possibility of publication bias (z=2.73, P=0.006). Subsequently, trim and fill method was used to correct and identify whether the asymmetry funnel plot was caused by publication bias. The results showed that no study was missing in the iterative

algorithm and the effect size of the meta-analysis after trimming and filling was exactly same as the primary results, which indicated that there was no publication bias and the asymmetry funnel plot may not be caused by publication bias.

Study or		Antiviral		Control	Weight	HR	HR
subgroup	Log (HR)	SE	Total	Total	(%)	IV, random, 95% (	CI IV, random, 95% CI
Studies using lamivu	dine as antivi	ral drug					
Ke et al (2013)15	-0.6733	0.254	141	141	5.0	0.51 (0.31–0.84)	
Subtotal (95% CI)			141	141	5.0	0.51 (0.31–0.84)	•
Heterogeneity: not app							
Test for overall effect:	Z=2.65 (P=0.00	08)					
Studies using enteca	vir as antivira	l drua					
Ding et al (2014) <sup>36</sup>	-0.6733	0.3638	74	39	2.9	0.51 (0.25-1.04)	
Zhang et al (2014)21	-0.2877	0.3469	40	47	3.2	0.75 (0.38-1.48)	
Subtotal (95% CI)			114	86	6.1	0.62 (0.38-1.02)	•
Heterogeneity: $\tau^2=0.00$	); χ²=0.59, <i>df</i> =1	(P=0.44); l <sup>2</sup>	=0%			· · · ·	•
Test for overall effect:	Z=1.88 (P=0.00	6)					
Studies using adefov	vir as antiviral	drua					
Huang et al (2015) <sup>14</sup>	-0.6675	0.2084	100	100	6.5	0.51 (0.34-0.77)	
Subtotal (95% CI)			100	100	6.5	0.51 (0.34-0.77)	•
Heterogeneity: not app	olicable					. ,	
Test for overall effect:	Z=3.20 (P=0.00	01)					
Studies using more t	han one kind (	of NAs as an	utiviral druc				
Chan et al (2011) <sup>12</sup>	-0.5798	0.3537	42	94	3.1	0.56 (0.28-1.12)	
Chen et al (2016)22	-0.4463	0.1575	192	253	8.8	0.64 (0.47–0.87)	-
Chong et al (2015)13	-0.5604	0.1678	254	150	8.3	0.57 (0.41-0.79)	-
Li et al (2010)16	-0.3857	0.2221	43	36	6.0	0.68 (0.44–1.05)	
Su et al (2013)17	-1.3863	0.3336	62	271	3.4	0.25 (0.13-0.48)	
Tian et al (2015)29	-0.5351	0.3317	29	21	3.4	0.59 (0.31–1.12)	
Wang et al (2015)30	-0.8062	0.255	76	80	5.0	0.45 (0.27-0.74)	
Wei et al (2016) <sup>23</sup>	-1.0966	0.324	86	40	3.5	0.33 (0.18–0.63)	
Wu et al (2012) <sup>18</sup>	-0.3567	0.0959	518	4,051	12.5	0.70 (0.58-0.84)	-
Yang et al (2012)19	-0.2485	0.1783	142	188	7.8	0.78 (0.55–1.11)	-
Yao et al (2016) <sup>33</sup>	-0.4691	0.1851	76	80	7.5	0.63 (0.44–0.90)	-
*Yin et al (2013a) <sup>20</sup>	-0.2357	0.1319	215	402	10.3	0.79 (0.61–1.02)	-
*Yin et al (2013b) <sup>20</sup>	-1.3093	0.3729	81	82	2.8	0.27 (0.13–0.56)	
Subtotal (95% CI)			1,816	5,748	82.4	0.59 (0.50-0.69)	♦
Heterogeneity: $r^2=0.04$	$\chi^2 = 24.70$ . df=	12 (P=0.02)		-,			•
Test for overall effect:		· · · · · ·					
Total (95% CI)			2,171	6,075	100	0.58 (0.51–0.67)	•
Heterogeneity: $\tau^2=0.03$	3; χ <sup>2</sup> =26.89, df	=16 ( <i>P</i> =0.04)	; <i>l</i> ²=40%	-		. ,	· · · ·
Test for overall effect: 2	· ·	,					0.01 0.1 1 10 10
Test for subgroup diffe	rences: $\chi^2 = 0.6$	7, df=3 (P=0	.88); /²=0%				Favors Favors
							antiviral control

Figure 3 Stratified meta-analysis of OS according to the type of NAs.

Note: \*Study of Yin et al (2013)<sup>20</sup> was a two-stage longitudinal clinical study which included a first stage pro study and a second stage RCT. Therefore, the first stage pro study was defined as Yin et al (2013a)<sup>20</sup> and the second stage RCT was defined as Yin et al (2013b)<sup>20</sup>.

Abbreviations: CI, confidence interval; HR, hazard ratio; IV, inverse variance; NAs, nucleoside analogs; OS, overall survival; SE, standard error.

# Discussion

Tumor recurrence is the most common cause of mortality for HCC patients after curative resection.<sup>47</sup> Despite the advances in surveillance programs, surgical technologies, and multidisciplinary treatments, there are still no adjuvant therapy options that effectively prevent HCC recurrence after curative resection. Most of the well-known risk factors for HCC recurrence, such as tumor characteristics, liver cirrhosis, and alpha fetoprotein level, are irreversible.<sup>48</sup> However, HBV status is an important risk factor for HCC recurrence that can be reversed by NA therapy. Thus, the exact effect of NA therapy on patients with HBV-related HCC after curative resection becomes a subject of great interest to hepatobiliary surgeons or physicians, and several studies have been performed recently, but the results are inconsistent. Otherwise, there is still not enough convincing evidence to support this issue because of the potential bias mentioned earlier in the previous studies or meta-analyses. Therefore, we conducted the current study, a more comprehensive meta-analysis, to assess the exact effect of NA therapy after curative resection on the long-term survival of patients with HBV-related HCC as far as possible.

The current meta-analysis demonstrated that NA therapy significantly improved the RFS of patients after surgical resection for HBV-related HCC, which suggested that NA therapy can prevent or delay the recurrence of HBV-related HCC. Recent studies have demonstrated that sustained viremia may impair tumor immune surveillance and favor the development of hepatocellular carcinogenesis,<sup>49,50</sup>

Study or subgroup	Log (HR)	Antiviral SE	therapy Total	Control Total	Weight (%)	HR IV, random, 95% CI	HR IV, random, 9	5% CI	
HBV DNA level ≥20,000	IU/mL								
Chan et al (2011) <sup>12</sup>	-0.5798	0.3537	42	94	14.8	0.56 (0.28-1.12)			
Ding et al (2014) <sup>36</sup>	-0.6733	0.3638	74	39	14.0	0.51 (0.25-1.04)			
Yang et al (2012)19	-0.2485	0.1783	142	188	58.3	0.78 (0.55–1.11)	-		
Subtotal (95% CI)			258	321	87.1	0.69 (0.52-0.92)	•		
Heterogeneity: $\tau^2=0.00$ ;	χ <sup>2</sup> =1.51, df=2 (F	P=0.47); I <sup>2</sup> =0	%						
Test for overall effect: Z=	2.56 ( <i>P</i> =0.01)								
HBV DNA level <20,000	IU/mL								
Chen et al (2016)22	-0.2231	0.3798	51	154	12.9	0.80 (0.38-1.68)	_ <b>_</b>		
Subtotal (95% CI)			51	154	12.9	0.80 (0.38-1.68)	•		
Heterogeneity: not applic	cable					. ,	•		
Test for overall effect: Z=									
Total (95% CI)			309	475	100	0.70 (0.54–0.92)	•		
Heterogeneity: $\tau^2=0.00$ ;	χ <sup>2</sup> =1.65, df=3 (F	P=0.65); /2=0	%			` ´ ⊢			
Test for overall effect: Z=	2.60 (P=0.009)					0.01	I 0.1 1	10	10
Test for subgroup differe	nces: $\gamma^2 = 0.14$	df=1 (P=0.71	); /²=0%				Favors	Favors	
loot for bubgroup amore									
loot for cabgroup amore	, τ	, i i i i i i i i i i i i i i i i i i i					antiviral	control	
		,	therapy	Control	Weight	HR		control	
	Log (HR)	Antiviral SE	therapy Total	Control Total	Weight (%)	HR IV, random, 95% Cl	antiviral		
Study or	Log (HR)	Antiviral					antiviral HR		
Study or subgroup	Log (HR)	Antiviral					antiviral HR		
Study or subgroup HBV DNA level ≥20,000	Log (HR)	Antiviral SE	Total	Total	(%)	IV, random, 95% CI	antiviral HR		
Study or subgroup HBV DNA level ≥20,000 Chan et al (2011) <sup>12</sup>	Log (HR) IU/mL -0.4155	Antiviral SE 0.2306	Total 42	<b>Total</b> 94	<b>(%)</b> 12.8	IV, random, 95% CI 0.66 (0.42–1.04)	antiviral HR		
Study or subgroup HBV DNA level ≥20,000 Chan et al (2011) <sup>12</sup> Cheng and Lv (2011) <sup>35</sup>	Log (HR) IU/mL -0.4155 -0.4082	<b>Antiviral</b> <b>SE</b> 0.2306 0.219	<b>Total</b> 42 50	<b>Total</b> 94 43	(%) 12.8 14.1	IV, random, 95% CI 0.66 (0.42–1.04) 0.66 (0.43–1.02)	antiviral HR		
Study or subgroup HBV DNA level ≥20,000 Chan et al (2011) <sup>12</sup> Cheng and Lv (2011) <sup>35</sup> Ding et al (2014) <sup>36</sup>	Log (HR) IU/mL -0.4155 -0.4082 -0.5987	Antiviral SE 0.2306 0.219 0.3155	<b>Total</b> 42 50 74	<b>Total</b> 94 43 39	(%) 12.8 14.1 6.9	IV, random, 95% CI 0.66 (0.42–1.04) 0.66 (0.43–1.02) 0.55 (0.30–1.02)	antiviral HR		
Study or subgroup           HBV DNA level ≥20,000           Chan et al (2011) <sup>12</sup> Cheng and Lv (2011) <sup>35</sup> Ding et al (2014) <sup>36</sup> Su et al (2013) <sup>17</sup>	Log (HR) -0.4155 -0.4082 -0.5987 -0.5323	Antiviral SE 0.2306 0.219 0.3155 0.2205	<b>Total</b> 42 50 74 44	<b>Total</b> 94 43 39 156	(%) 12.8 14.1 6.9 14.0	IV, random, 95% CI 0.66 (0.42–1.04) 0.66 (0.43–1.02) 0.55 (0.30–1.02) 0.59 (0.38–0.90)	antiviral HR		
Study or subgroup           HBV DNA level ≥20,000           Chan et al (2011) <sup>12</sup> Cheng and Lv (2011) <sup>35</sup> Ding et al (2014) <sup>36</sup> Su et al (2013) <sup>17</sup> Yang et al (2010) <sup>32</sup>	Log (HR) -0.4155 -0.4082 -0.5987 -0.5323 -0.5027	Antiviral SE 0.2306 0.219 0.3155 0.2205 0.4713	<b>Total</b> 42 50 74 44 15	<b>Total</b> 94 43 39 156 15	(%) 12.8 14.1 6.9 14.0 3.1	IV, random, 95% CI 0.66 (0.42–1.04) 0.66 (0.43–1.02) 0.55 (0.30–1.02) 0.59 (0.38–0.90) 0.60 (0.24–1.52)	antiviral HR IV, random, 9		
Study or subgroup           HBV DNA level ≥20,000           Chan et al (2011) <sup>12</sup> Cheng and Lv (2011) <sup>35</sup> Ding et al (2014) <sup>36</sup> Su et al (2013) <sup>17</sup> Yang et al (2010) <sup>32</sup> Yang et al (2012) <sup>19</sup>	Log (HR) IU/mL -0.4155 -0.4082 -0.5987 -0.5323 -0.5027 -0.6349	Antiviral SE 0.2306 0.219 0.3155 0.2205 0.4713 0.1436	<b>Total</b> 42 50 74 44 15 142 <b>367</b>	<b>Total</b> 94 43 39 156 15 15 188	(%) 12.8 14.1 6.9 14.0 3.1 31.7	IV, random, 95% CI 0.66 (0.42–1.04) 0.66 (0.43–1.02) 0.55 (0.30–1.02) 0.59 (0.38–0.90) 0.60 (0.24–1.52) 0.53 (0.40–0.70)	antiviral HR IV, random, 9		
Study or subgroup           HBV DNA level ≥20,000           Chan et al (2011) <sup>12</sup> Cheng and Lv (2011) <sup>35</sup> Ding et al (2014) <sup>36</sup> Su et al (2013) <sup>17</sup> Yang et al (2010) <sup>32</sup> Yang et al (2012) <sup>19</sup> Subtotal (95% CI)	Log (HR) IU/mL -0.4155 -0.4082 -0.5987 -0.5323 -0.5027 -0.6349 χ <sup>2</sup> =1.13, df=5 (F	Antiviral SE 0.2306 0.219 0.3155 0.2205 0.4713 0.1436 P=0.95); /²=0	<b>Total</b> 42 50 74 44 15 142 <b>367</b>	<b>Total</b> 94 43 39 156 15 15 188	(%) 12.8 14.1 6.9 14.0 3.1 31.7	IV, random, 95% CI 0.66 (0.42–1.04) 0.66 (0.43–1.02) 0.55 (0.30–1.02) 0.59 (0.38–0.90) 0.60 (0.24–1.52) 0.53 (0.40–0.70)	antiviral HR IV, random, 9		
Study or subgroup           HBV DNA level ≥20,000           Chan et al (2011) <sup>12</sup> Cheng and Lv (2011) <sup>35</sup> Ding et al (2014) <sup>36</sup> Su et al (2013) <sup>17</sup> Yang et al (2010) <sup>32</sup> Yang et al (2012) <sup>19</sup> Subtotal (95% CI)           Heterogeneity: τ <sup>2</sup> =0.00;	Log (HR) IU/mL -0.4155 -0.4082 -0.5987 -0.5323 -0.5027 -0.6349 $\chi^2$ =1.13, <i>df</i> =5 ( <i>F</i> -5.98 ( <i>P</i> <0.0000	Antiviral SE 0.2306 0.219 0.3155 0.2205 0.4713 0.1436 P=0.95); /²=0	<b>Total</b> 42 50 74 44 15 142 <b>367</b>	<b>Total</b> 94 43 39 156 15 15 188	(%) 12.8 14.1 6.9 14.0 3.1 31.7	IV, random, 95% CI 0.66 (0.42–1.04) 0.66 (0.43–1.02) 0.55 (0.30–1.02) 0.59 (0.38–0.90) 0.60 (0.24–1.52) 0.53 (0.40–0.70)	antiviral HR IV, random, 9		
Study or subgroup HBV DNA level ≥20,000 Chan et al (2011) <sup>12</sup> Cheng and Lv (2011) <sup>35</sup> Ding et al (2014) <sup>36</sup> Su et al (2013) <sup>17</sup> Yang et al (2010) <sup>32</sup> Yang et al (2012) <sup>19</sup> Subtotal (95% CI) Heterogeneity: $r^2$ =0.00; Test for overall effect: Z=	Log (HR) IU/mL -0.4155 -0.4082 -0.5987 -0.5323 -0.5027 -0.6349 $\chi^2$ =1.13, <i>df</i> =5 ( <i>F</i> -5.98 ( <i>P</i> <0.0000	Antiviral SE 0.2306 0.219 0.3155 0.2205 0.4713 0.1436 P=0.95); /²=0	<b>Total</b> 42 50 74 44 15 142 <b>367</b>	<b>Total</b> 94 43 39 156 15 15 188	(%) 12.8 14.1 6.9 14.0 3.1 31.7	IV, random, 95% CI 0.66 (0.42–1.04) 0.66 (0.43–1.02) 0.55 (0.30–1.02) 0.59 (0.38–0.90) 0.60 (0.24–1.52) 0.53 (0.40–0.70)	antiviral HR IV, random, 9		
Study or subgroup HBV DNA level ≥20,000 Chan et al (2011) <sup>12</sup> Cheng and Lv (2011) <sup>35</sup> Ding et al (2014) <sup>36</sup> Su et al (2013) <sup>17</sup> Yang et al (2010) <sup>32</sup> Yang et al (2012) <sup>19</sup> Subtotal (95% Cl) Heterogeneity: $\tau^2$ =0.00; Test for overall effect: Z= HBV DNA level <20,000	Log (HR) IU/mL -0.4155 -0.4082 -0.5987 -0.5323 -0.5027 -0.6349 χ <sup>2</sup> =1.13, df=5 (F -5.98 (P<0.0000) IU/mL	Antiviral SE 0.2306 0.219 0.3155 0.2205 0.4713 0.1436 P=0.95); /²=0 01)	<b>Total</b> 42 50 74 44 15 142 <b>367</b> %	94         43         39         156         15         188         535	(%) 12.8 14.1 6.9 14.0 3.1 31.7 <b>82.6</b>	IV, random, 95% CI 0.66 (0.42–1.04) 0.66 (0.43–1.02) 0.55 (0.30–1.02) 0.59 (0.38–0.90) 0.60 (0.24–1.52) 0.53 (0.40–0.70) 0.58 (0.49–0.70)	antiviral HR IV, random, 9		
Study or subgroup HBV DNA level ≥20,000 Chan et al (2011) <sup>12</sup> Cheng and Lv (2011) <sup>35</sup> Ding et al (2014) <sup>36</sup> Su et al (2013) <sup>17</sup> Yang et al (2012) <sup>19</sup> Subtotal (95% Cl) Heterogeneity: $\tau^2$ =0.00; Test for overall effect: Z= HBV DNA level <20,000 Chen et al (2016) <sup>22</sup>	Log (HR) IU/mL -0.4155 -0.4082 -0.5987 -0.5323 -0.5027 -0.6349 χ <sup>2</sup> =1.13, df=5 (F -5.98 (P<0.0000) IU/mL 0.131	Antiviral SE 0.2306 0.219 0.3155 0.2205 0.4713 0.1436 P=0.95); /²=0 01) 0.2416	<b>Total</b> 42 50 74 44 15 142 <b>367</b> %	<b>Total</b> 94 43 39 156 15 188 <b>535</b> 154	(%) 12.8 14.1 6.9 14.0 3.1 31.7 <b>82.6</b> 11.7	IV, random, 95% CI 0.66 (0.42–1.04) 0.66 (0.43–1.02) 0.55 (0.30–1.02) 0.59 (0.38–0.90) 0.60 (0.24–1.52) 0.53 (0.40–0.70) 0.58 (0.49–0.70) 1.14 (0.71–1.83)	antiviral HR IV, random, 9		
Study or subgroup HBV DNA level ≥20,000 Chan et al (2011) <sup>12</sup> Cheng and Lv (2011) <sup>35</sup> Ding et al (2014) <sup>36</sup> Su et al (2013) <sup>17</sup> Yang et al (2013) <sup>17</sup> Yang et al (2012) <sup>19</sup> Subtotal (95% CI) Heterogeneity: $r^2$ =0.00; Test for overall effect: Z= HBV DNA level <20,000 Chen et al (2016) <sup>22</sup> Su et al (2013) <sup>17</sup>	Log (HR) IU/mL -0.4155 -0.4082 -0.5987 -0.5323 -0.5027 -0.6349 χ <sup>2</sup> =1.13, df=5 (F -5.98 (P<0.0000 IU/mL 0.131 -0.6255	Antiviral SE 0.2306 0.219 0.3155 0.2205 0.4713 0.1436 D=0.95); / <sup>2</sup> =0 01) 0.2416 0.3811	<b>Total</b> 42 50 74 44 15 142 <b>367</b> % 51 17	<b>Total</b> 94 43 39 156 15 188 <b>535</b> 154 104	(%) 12.8 14.1 6.9 14.0 3.1 31.7 <b>82.6</b> 11.7 4.8	IV, random, 95% CI 0.66 (0.42–1.04) 0.66 (0.43–1.02) 0.55 (0.30–1.02) 0.59 (0.38–0.90) 0.60 (0.24–1.52) 0.53 (0.40–0.70) 0.58 (0.49–0.70) 1.14 (0.71–1.83) 0.53 (0.25–1.13)	antiviral HR IV, random, 9		
Study or subgroup HBV DNA level ≥20,000 Chan et al (2011) <sup>12</sup> Cheng and Lv (2011) <sup>35</sup> Ding et al (2014) <sup>36</sup> Su et al (2013) <sup>17</sup> Yang et al (2010) <sup>32</sup> Yang et al (2012) <sup>19</sup> Subtotal (95% CI) Heterogeneity: $\tau^2$ =0.00; Test for overall effect: Z= HBV DNA level <20,000 Chen et al (2016) <sup>22</sup> Su et al (2013) <sup>17</sup> Yang et al (2010) <sup>32</sup>	Log (HR) IU/mL −0.4155 −0.4082 −0.5987 −0.5323 −0.5027 −0.6349 χ <sup>2</sup> =1.13, df=5 (F :5.98 (P<0.0000) IU/mL 0.131 −0.6255 −0.1664	Antiviral SE 0.2306 0.219 0.3155 0.2205 0.4713 0.1436 P=0.95); /²=0 01) 0.2416 0.3811 0.8497	Total 42 50 74 44 15 142 <b>367</b> % 51 17 15 <b>83</b>	Total 94 43 39 156 15 188 <b>535</b> 154 104 15	(%) 12.8 14.1 6.9 14.0 3.1 31.7 82.6 11.7 4.8 1.0	IV, random, 95% CI 0.66 (0.42–1.04) 0.66 (0.43–1.02) 0.55 (0.30–1.02) 0.59 (0.38–0.90) 0.60 (0.24–1.52) 0.53 (0.40–0.70) 0.58 (0.49–0.70) 1.14 (0.71–1.83) 0.53 (0.25–1.13) 0.85 (0.16–4.48)	antiviral HR IV, random, 9		
Study or subgroup HBV DNA level ≥20,000 Chan et al (2011) <sup>12</sup> Cheng and Lv (2011) <sup>35</sup> Ding et al (2014) <sup>36</sup> Su et al (2013) <sup>17</sup> Yang et al (2010) <sup>32</sup> Yang et al (2012) <sup>19</sup> Subtotal (95% CI) Heterogeneity: $r^2$ =0.00; Test for overall effect: Z= HBV DNA level <20,000 Chen et al (2016) <sup>22</sup> Su et al (2013) <sup>17</sup> Yang et al (2010) <sup>32</sup> Subtotal (95% CI)	Log (HR) IU/mL -0.4155 -0.4082 -0.5987 -0.5323 -0.5027 -0.6349 $\chi^2=1.13, df=5$ ( <i>F</i> $z^2=1.13, df=5$ ( <i>F</i> 0.131 -0.6255 -0.1664 $\chi^2=2.82, df=2$ ( <i>F</i>	Antiviral SE 0.2306 0.219 0.3155 0.2205 0.4713 0.1436 P=0.95); /²=0 01) 0.2416 0.3811 0.8497	Total 42 50 74 44 15 142 <b>367</b> % 51 17 15 <b>83</b>	Total 94 43 39 156 15 188 <b>535</b> 154 104 15	(%) 12.8 14.1 6.9 14.0 3.1 31.7 82.6 11.7 4.8 1.0	IV, random, 95% CI 0.66 (0.42–1.04) 0.66 (0.43–1.02) 0.55 (0.30–1.02) 0.59 (0.38–0.90) 0.60 (0.24–1.52) 0.53 (0.40–0.70) 0.58 (0.49–0.70) 1.14 (0.71–1.83) 0.53 (0.25–1.13) 0.85 (0.16–4.48)	antiviral HR IV, random, 9		

Test for subgroup differences:  $\chi^2$ =1.85, df=1 (P=0.17);  $I^2$ =45.9% **Favors** Favors antiviral control

Figure 4 Stratified meta-analysis of postoperative survival outcomes according to viral load (baseline HBV DNA level ≥20,000 IU/mL versus <20,000 IU/mL). Notes: (A) Meta-analysis of OS. (B) Meta-analysis of RFS.

Abbreviations: CI, confidence interval; HBV, hepatitis B virus; HR, hazard ratio; IV, inverse variance; OS, overall survival; RFS, recurrence-free survival; SE, standard error.

and chronic hepatitis activity and liver inflammation induced by immune response were associated with HCC recurrence after radical resection.<sup>51,52</sup> Similarly, other studies found that high HBV load and HBV mutation promote the growth and metastasis of HCC,<sup>17,53,54</sup> and the HBV X protein promotes the invasive and metastatic potential of HCC.<sup>20,55–57</sup> Therefore, the beneficial effect of NA therapy on HCC recurrence may be associated with the inhibition of these viral factors. Due to moderate heterogeneity that was detected among eligible studies, subgroup analyses were performed. After pooling studies using one type of NAs, only a little heterogeneity was observed. That may be partly

Heterogeneity: τ<sup>2</sup>=0.00; χ<sup>2</sup>=8.18, df=8 (P=0.42); l<sup>2</sup>=2%

Test for overall effect: Z=5.50 (P<0.00001)

attributed to the different viral suppression effect of different types of NAs which can further affect the tumor recurrence of HBV-related HCC. In the subgroup analyses of patients with different viral load, NA therapy can significantly prolong RFS in patients with high baseline HBV DNA level ( $\geq$ 20,000 IU/mL) with no heterogeneity, but not in patients with low baseline HBV DNA level (<20,000 IU/mL) with moderate heterogeneity. The results indicated that HCC patients with low baseline HBV DNA level may not significantly benefit from NA therapy as patients with high baseline HBV DNA level did. However, as there was moderate heterogeneity in the subgroup analysis of patients

0.01

0.1

10

100



Figure 5 Subgroup analysis on patients with fully preserved hepatic function (Child class A).

Notes: (A) Meta-analysis of OS. (B) Meta-analysis of RFS.

Abbreviations: CI, confidence interval; HR, hazard ratio; IV, inverse variance; OS, overall survival; RFS, recurrence-free survival; SE, standard error.

with low baseline HBV DNA level (<20,000 IU/mL), the findings are not conclusive and further high-quality studies are needed.

In this study, we also found that NA therapy can significantly improve the OS of patients with HBV-related HCC after surgical resection. The beneficial effect of NA therapy in preventing or delaying HCC recurrence contributes to better OS. Further, several studies have reported that NA therapy is effective in suppressing viral replication, modulating liver function, and increasing residual liver volume after radical resection.13,15,16,20 These effects may not only affect survival directly but also significantly improve the tolerance of patients to receive subsequent therapy (especially repeat surgical resection) after HCC recurrence which leads to a significant improvement in OS. Subgroup analyses were also conducted because of the moderate heterogeneity among studies. After pooling studies using one type of NA, no heterogeneity was observed among studies. In the subgroup analyses of patients with different viral load, NA therapy can significantly prolong OS in patients with high baseline HBV DNA level ( $\geq 20,000$  IU/mL) with no heterogeneity, but not in patients with low baseline HBV DNA level (<20,000 IU/mL) with moderate heterogeneity. Smaller size was one factor that led to nonsignificant groups of patients with low baseline HBV DNA level (<20,000 IU/mL). Moreover, it has been identified that HCC patients with persistently low HBV DNA levels had better survival results compared to those with high HBV DNA levels. After antiviral therapy, groups of HCC patients with high HBV DNA levels always could achieve a more substantial reduction in HBV DNA load, as compared to groups of HCC patients with low HBV DNA levels. Therefore, HCC patients with low HBV DNA levels may not significantly benefit from antiviral therapy compared to patients with high HBV DNA levels. However, as only one study was included in the subgroup analysis of patients with low baseline HBV DNA level (<20,000 IU/mL), more quality studies are needed to draw a definitive conclusion.

Nonetheless, there are several limitations to the current meta-analysis. First, most of the included studies were non-RCTs; the potential confounding factors in these studies may decrease the reliability of the results, even for the well-analyzed cohort studies. Second, several indirect data acquisition methods were used in the meta-analysis, which may have effects on the outcomes. Finally, moderate heterogeneity existed in the meta-analysis; the variation in HBV status, type of NA therapy, Child–Pugh class, and tumor stage may be responsible for the heterogeneity.

## Conclusion

The current meta-analysis suggests that antiviral therapy with NAs significantly improves the survival outcomes of patients with HBV-related HCC after curative resection, especially for patients with high HBV DNA level. To further

Study or	Log (HR)	Antiviral SE	therapy Total	Control Total	Weight	HR	<b>CI</b>	HR IV, random	95% 01	
subgroup	- · ·		Total	Total	(%)	IV, random, 95%		IV, random	, 95% CI	-
Studies using lamivudi		•								
Chen (2015) <sup>24</sup>	-0.297	0.3001	45	40	1.8	0.74 (0.41–1.34)		-+		
Cheng and Lv (2011) <sup>35</sup>	-0.4082	0.219	50	43	3.1	0.66 (0.43–1.02)		-		
ang et al (2012) <sup>25</sup>	-0.9086	0.3812	26	30	1.2	0.40 (0.19–0.85)				
Ke et al (2013) <sup>15</sup>	-0.1054	0.146	141	141	5.6	0.90 (0.68–1.20)		-		
Zhang (2015) <sup>34</sup>	-0.2942	0.2752	45	40	2.1	0.75 (0.43–1.28)		-+		
Subtotal (95% CI)			307	294	13.8	0.75 (0.60-0.93)		•		
Heterogeneity: $\tau^2$ =0.01; $\gamma$ Test for overall effect: Z=			=11%							
	,	,								
Studies using entecavi		-								
Ding et al (2014)36	-0.5987	0.3155	74	39	1.7	0.55 (0.30–1.02)				
₋in et al (2016) <sup>27</sup>	-1.1841	0.3932	35	25	1.1	0.31 (0.14–0.66)				
Zhang et al (2014) <sup>21</sup>	-0.5108	0.2606	40	47	2.3	0.60 (0.36–1.00)				
Subtotal (95% CI)			149	111	5.1	0.51 (0.35–0.73)		•		
Heterogeneity: $\tau^2$ =0.01; $\gamma$			=6%							
Test for overall effect: Z=	3.68 (P=0.00	02)								
Studies using adefovir	as antiviral o	drug								
Huang et al (2015)14	-0.399	0.1827	100	100	4.1	0.67 (0.47–0.96)				
Subtotal (95% CI)			100	100	4.1	0.67 (0.47-0.96)		•		
leterogeneity: not applic	able									
Test for overall effect: Z=	2.18 (P=0.03	)								
Studies using more that	in one kind o	of NAs as ar	ntiviral dru	ıg						
Chan et al (2011) <sup>12</sup>	-0.4155	0.2306	42	94	2.9	0.66 (0.42-1.04)		-		
Chen et al (2016)22	-0.0305	0.1245	192	253	6.8	0.97 (0.76–1.24)		+		
Chong et al (2015) <sup>13</sup>	-0.3147	0.1262	254	150	6.7	0.73 (0.57–0.93)		-		
-luang et al (2016) <sup>26</sup>	-0.5086	0.2205	45	33	3.1	0.60 (0.39–0.93)				
_i et al (2010) <sup>16</sup>	-0.2107	0.1793	43	36	4.2	0.81 (0.57–1.15)		-+		
Qian et al (2016) <sup>28</sup>	-0.685	0.2324	70	65	2.8	0.50 (0.32-0.79)				
Su et al (2013) <sup>17</sup>	-0.5798	0.1846	62	271	4.0	0.56 (0.39–0.80)		-		
Fian et al (2015) <sup>29</sup>	-0.5351	0.3317	29	21	1.5	0.59 (0.31–1.12)				
Nang et al (2015)30	-0.4291	0.1767	76	80	4.3	0.65 (0.46-0.92)				
Nei et al (2016)23	-0.4005	0.1809	86	40	4.2	0.67 (0.47-0.96)				
Nu et al (2012) <sup>18</sup>	-0.2744	0.072	518	4,051	10.8	0.76 (0.66-0.88)		-		
Ku et al (2016) <sup>31</sup>	-0.6038	0.4061	29	82	1.1	0.55 (0.25–1.21)				
Yang et al (2012)19	-0.6349	0.1436	142	188	5.7	0.53 (0.40-0.70)		-		
Yao et al (2016)33	-0.3479	0.1574	76	80	5.1	0.71 (0.52-0.96)		-		
Yin et al (2013a)20	-0.2485	0.0852	215	402	9.6	0.78 (0.66-0.92)		-		
Yin et al (2013b)20	-0.734	0.1759	81	82	4.3	0.48 (0.34-0.68)		-		
Subtotal (95% CI)			1,960	5,928	76.9	0.68 (0.62-0.75)		•		
Heterogeneity: $\tau^2 = 0.01$ ; Fest for overall effect: Z=			; /2=37%							
	1.02 (1 >0.00	001)								
			2,516	6,433	100	0.68 (0.63-0.74)		•		
Fotal (95% CI)	2-31 37 df-	24 (0-0 00)				. ,		· · · · · · ·		
Heterogeneity: $\tau^2$ =0.01;				-,		. ,	0 01	01 1	10	1
	8.87 (P<0.00	001)	; /2=30%	-			0.01	0.1 1 Favors	10 Favors	1

Figure 6 Stratified meta-analysis of RFS according to the type of NAs.

Note: \*Study of Yin et al (2013)<sup>20</sup> was a two-stage longitudinal clinical study which included a first stage pro study and a second stage RCT. Therefore, the first stage pro study was defined as Yin et al (2013a)<sup>20</sup> and the second stage RCT was defined as Yin et al (2013b)<sup>20</sup>.

Abbreviations: Cl, confidence interval; HR, hazard ratio; IV, inverse variance; NAs, nucleoside analogs; RFS, recurrence-free survival; SE, standard error.



**Figure 7** Funnel plot for the results from included studies comparing RFS in HBVrelated HCC patients who received antiviral therapy or no treatment. **Abbreviations:** HBV, hepatitis B virus; HCC, hepatocellular carcinoma; HR, hazard ratio; RFS, recurrence-free survival; SE, standard error. investigate the promising effects of antiviral therapy with NAs on patients with low HBV DNA level, high-quality studies are needed.

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# Disclosure

The authors report no conflicts of interest in this work.

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