Review Article

Antiviral Therapy in Lamivudine-Resistant Chronic Hepatitis B Patients: A Systematic Review and Network Meta-Analysis

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The relative efficacy of different strategies for chronic hepatitis B (CHB) patients with lamivudine resistance (LAM-R) has not yet been systematically studied. Clinical trials were searched in PUBMED, MEDLINE, EMBASE, and CNKI databases up to February 15, 2016. Nine trials including 764 patients met the entry criteria. In direct meta-analysis, TDF showed a stronger antiviral effect than any one of ETV, LAM/ADV, and ADV against LAM-R hepatitis B virus. LAM/ADV therapy was superior to ADV in suppressing viral replication. ETV achieved similar rate of HBV DNA undetectable compared to ADV or LAM/ADV. In network meta-analysis, TDF had higher rates of HBV DNA undetectable compared to ETV (OR, 24.69; 95% CrI: 5.36–113.66), ADV (OR, 37.28; 95% CrI: 9.73–142.92), or LAM/ADV (OR, 21.05; 95% CrI: 5.70–77.80). However, among ETV, ADV, and LAM/ADV, no drug was clearly superior to others in HBV DNA undetectable rate. Moreover, no significant difference in the rate of ALT normalization or HBeAg loss was observed compared the four rescue strategies with each other. TDF appears to be a more effective rescue therapy than LAM/ADV, ETV, or ADV. LAM plus ADV therapy was a better treatment option than ETV or ADV alone for patients with LAM-R.

1. Introduction

Chronic hepatitis B virus (HBV) infection remains a major public health problem and affects approximately 400 million people worldwide, especially in Asia [1]. Large cohort studies have demonstrated that the risk of liver disease progression in patients with chronic hepatitis B (CHB) is associated with elevated HBV DNA levels [2]. Therefore, the goals of therapy in HBV infected patients are to reverse progression of the disease by long-term suppression of HBV replication [3]. With the availability of potent nucleos(t)ide analogues (NAs), such as tenofovir disoproxil fumarate (TDF) and entecavir (ETV), suppression of serum HBV DNA to undetectable levels is achievable in most NA-naïve patients in the absence of drug-resistant HBV mutants [4, 5]. However, for most NAs such as lamivudine (LAM) and adefovir (ADV), one of the major limitations of these drugs is resistance development in long-term treatment, which can lead to a rebound in HBV replication and exacerbation of HBV-related disease [2].

LAM was prescribed extensively for the treatment of CHB in the era of the early generation NAs. However, resistance to LAM emerges in approximately 20% of patients after 1 year and in 70% of patients after 5 years of treatment [6]. Until TDF rescue therapy becomes available, switching to ADV or ETV monotherapy and adding on ADV were once suggested against LAM-R HBV in earlier international guidelines [3, 7]. Unfortunately, sequential ADV monorescue therapy for LAM-R induced high resistance to ADV [8, 9], and ETV monorescue therapy resulted in 50% of these patients developing ETV-resistance (ETV-R) after 5 years of treatment [10]. Combination of ADV and LAM therapy reduces the development of ADV resistance and has been a practical option for treatment of LAM resistance. However, LAM plus ADV therapy has limited antiviral efficacy in
LAM-R patients, and a substantial proportion of patients show a suboptimal virological response which may then result in selection of multidrug-resistant HBV variants and progression of liver disease [11]. As rescue therapies for LAM-R patients, TDF, an oral NA with the most potent activity against HBV and high genetic barrier [12], has demonstrated favourable virological outcomes. Some recent studies have revealed that TDF monotherapy was highly effective in LAM-R patients as well as NA-naive patients, and the presence of resistance mutations to LAM did not alter the response rates [13, 14]. Although some meta-analyses [24, 25] have compared the efficacy of LAM/ADV versus ETV or ADV for patients with LAM-R, the relative efficacy of various rescue strategies in LAM-R patients has not yet been systematically studied, as well as the efficacy of TDF versus LAM/ADV or ETV or ADV. Network meta-analysis can help assess comparative effectiveness of multiple interventions and synthesize evidence through simultaneous analysis of direct evidence and indirect evidence to calculate a mixed-effect estimate as the weighted average of the two. Such a technique may improve the precision of the estimate (compared with direct evidence alone) and also allows estimation of the comparative efficacy of two active treatments, even if no studies directly compare them [26]. Therefore, in this study, we conducted a network meta-analysis with updated evidence to evaluate effects of different rescue strategies including TDF, ETV, LAM/ADV, and ADV in the treatment of LAM-R patients.

2. Materials and Methods

2.1. Search Strategy. We searched PUBMED, MEDLINE, EMBASE, and CNKI databases up to February 15, 2016. The following keywords were used for the search to find relevant citations: chronic hepatitis B, lamivudine-resistant, tenofovir, adefovir, and entecavir. In addition, reference lists from retrieved documents were reviewed, and a manual search was conducted to supplement the computer search. The search results were downloaded to a reference database and were further screened by 2 authors (Hui-Lian Wang and Xudong Yang).

2.2. Inclusion and Exclusion Criteria. The following inclusion criteria were used for this meta-analysis: (1) CHB patients with LAM resistance which was defined as the presence of HBV variants with amino acid substitutions conferring resistance against LAM (rtM204V/I ± rtL180M); (2) intervention therapies: TDF, ETV, or ADV, or LAM plus ADV therapy. The following types of studies were excluded: (1) studies of patients with liver failure, HCC, coinfection with hepatitis C, hepatitis D or HIV, and previous liver transplant, (2) studies not reporting any efficacy measures or not conveying sufficient statistical information, and (3) studies that were not controlled trials.

2.3. Efficacy Measures. Efficacy was assessed by rates of undetectable HBV DNA (<400 copies/mL), ALT normalization (<40 IU/mL), HBeAg loss, and virological breakthrough for patients 24, 48, and 96 weeks after therapy.

2.4. Data Extraction. Data extraction was assessed independently by two reviewers (Hui-Lian Wang and Xudong Yang). Discrepancies among reviewers were resolved by discussions between the reviewers or by a third person (Nan Xu). Basic information obtained from each eligible trial included the study design, patient characteristics, numbers in each group, and treatments. Data were reviewed to eliminate duplicate reports of the same trial.

2.5. Quality Assessment. The risk of bias of included trials was assessed by Cochrane Collaboration's tool with the outcome shown in Sup. Figure 2 in Supplementary Material available online at http://dx.doi.org/10.1155/2016/3435965. The following factors were taken into consideration for the risk of bias: random sequence generation (selection bias), allocation concealment (selection bias), blinding of participants and personnel (performance bias), blinding of outcome assessment (detection bias), incomplete outcome data (attrition bias), selective reporting (reporting bias), and other bias. The risk of bias of each study was explicitly determined on those factors and classified as three different levels: low, high, or unclear. The percentages of low risk of selection bias, performance bias, and the detection bias were less than 50% according to the description of each study. The percentages of low risk of bias of incomplete outcome data, selective reporting, and other bias were all more than 50%. The outcome of risk of bias graph showed that there was low risk of bias in this meta-analysis.

2.6. Statistical Analysis. For direct meta-analysis, the data was conducted on continuous and dichotomous outcomes and assessed by Revman. The $\chi^2$ and $I^2$ tests were first calculated to assess the heterogeneity of the included trials. For $P$ values more than 0.1, the fixed-effects model was used because of the homogeneity; otherwise, data need to be dealt with the random-effects model. Pooled odds ratios (OR) with 95% confidence intervals (95% CI) were calculated using either the fixed-effects model (M-H methods) or random-effects model (D-L methods). A two-tailed $P$ value of less than 0.05 suggested statistical significance.

We used network meta-analysis methods to compare LAM-R patients with different incorporating evidence on both direct and indirect comparisons. Network meta-analysis was performed using R version 3.2.2 to calculate point estimates (OR) with 95% credibility intervals (CrI) and generate forest plots using random-effects models comparing the effect estimates of different therapies relative to comparator. Rank probabilities were generated to determine the rank of therapies in which the given treatment ranked first as the most effective therapy, second, and so on.

Network meta-analysis was conducted with R 3.2.2, addis and stata 13.0. Direct meta-analysis and figures of risk of bias were generated through Review Manager 5.3.
3. Results

3.1. Study Characteristics and Quality Assessment. After the selection procedure (Sup. Figure 1), a total of nine studies [15–23] met the inclusion criteria for this review, including 764 patients with LAM-R. Among nine studies, two studies compared TDF versus LAM/ADV [16, 17] and one study compared TDF versus ETV [16] or ADV [15], respectively, six studies compared ETV versus LAM/ETV [18–22], and two studies compared ADV versus LAM/ADV [22, 23]. All nine studies were published in English and in full-text form. The characteristics of each study are listed in Table 1.

3.2. Virological Response

3.2.1. Direct Meta-Analysis. At 24 weeks of treatment (Figure 1(a)), TDF had higher HBV DNA undetectable rate compared to ADV (50.0% versus 20.0%; OR, 4.00; 95% CI: 1.71–9.34). ADV had lower HBV DNA undetectable rate compared to LAM/ADV (30.2% versus 63.1%; OR, 0.25; 95% CI: 0.12–0.52). ETV had similar rates of HBV DNA undetectable compared to ADV (33.3% versus 27.3%; OR, 1.33; 95% CI: 0.45–3.92) or LAM/ADV (21.6% versus 40.7%; OR, 0.62; 95% CI: 0.14–2.79), respectively.

At 48 weeks of treatment (Figure 1(b)), TDF had higher rates of HBV DNA undetectable compared to ETV (88.9% versus 36.4%; OR, 14.00; 95% CI: 2.06–95.09), ADV (86.4% versus 21.5%; OR, 23.07; 95% CI: 8.12–65.57), or LAM/ADV (93.5% versus 14.8%; OR, 74.42; 95% CI: 20.01–276.70), respectively. ADV had lower HBV DNA undetectable rate compared to LAM/ADV (42.7% versus 75.0%; OR, 0.25; 95% CI: 0.13–0.47). ETV had similar rates of HBV DNA undetectable compared to ADV (54.2% versus 40.9%; OR, 1.71; 95% CI: 0.63–4.65) or LAM/ADV (32.3% versus 34.2%; OR, 0.90; 95% CI: 0.42–1.92), respectively.

At 96 weeks of treatment (Figure 1(c)), ADV had lower HBV DNA undetectable rate compared to LAM/ADV (50.0% versus 83.3%; OR, 0.20; 95% CI: 0.08–0.51). ETV had similar rates of HBV DNA undetectable compared to LAM/ADV (48.6% versus 41.2%; OR, 1.41; 95% CI: 0.53–3.75).

3.2.2. Network Meta-Analysis. On Bayesian network meta-analysis (Sup. Table 1), TDF had higher rates of HBV DNA undetectable as compared to ETV (OR, 24.67; 95% CrI: 5.36–113.66), ADV (OR, 37.28; 95% CrI: 9.73–142.92), or LAM/ADV (OR, 21.05; 95% CrI: 5.70–77.80) through 48 weeks of treatment. However, among ETV, ADV, and LAM/ADV, no drug was clearly superior to others in HBV DNA undetectable rate during the same period.

TDF, LAM/ADV, and ETV had the highest probability of being ranked first, second, and third for achieving HBV DNA undetectable after 48-week treatment, respectively, whereas ADV had had highest probability of being ranked fourth (Figure 3).

3.3. Biochemical Response

3.3.1. Direct Meta-Analysis. After 48-week treatment (Sup. Figure 3), no significant difference in the rate of ALT normalization was observed when comparing TDF to ADV (59.1% versus 69.2%; OR, 0.64; 95% CI: 0.29–1.43) or LAM/ADV (89.3% versus 67.7%; OR, 3.97; 95% CI: 0.96–16.33), respectively. Moreover, there was no significant difference in ALT normalization between ETV and LAM/ADV (77.7% versus 84.0%; OR, 0.75; 95% CI: 0.35–1.58) or ADV and LAM/ADV (71.2% versus 79.2%; OR, 0.65; 95% CI: 0.26–1.63).

3.3.2. Network Meta-Analysis. On Bayesian network meta-analysis, when comparing TDF to others, including ETV, ADV, and LAM/ADV, no drug was clearly superior to others (Sup. Table 2) in the rates of ALT normalization after 48 weeks of therapy.

TDF, ADV, and LAM/ADV had the highest probability of being ranked first, second, and third for improving biochemical response after 48-week treatment, respectively, whereas ETV had had highest probability of being ranked fourth (Sup. Figure 5).

3.4. Serological Response

3.4.1. Direct Meta-Analysis. After 48-week treatment (Sup. Figure 4(A)), no significant difference in the rate of HBeAg loss was observed when comparing TDF to ADV (91.9% versus 4.6%; OR, 2.07; 95% CI: 0.44–9.73) or LAM/ADV (3.6% versus 0; OR, 3.44; 95% CI: 0.13–87.85), respectively. Moreover, there was no significant difference in HBeAg loss between ETV and LAM/ADV (3.6% versus 9.4%; OR, 0.41; 95% CI: 0.09–1.89) or ADV and LAM/ADV (48.1% versus 66.7%; OR, 0.46; 95% CI: 0.21–1.04).

After 96-week treatment (Sup. Figure 4(B)), no significant difference in the rate of HBeAg loss was observed when comparing ETV to ADV (33.3% versus 50.0%; OR, 0.50; 95% CI: 0.18–1.41) or LAM/ADV (21.6% versus 32.8%; OR, 0.48; 95% CI: 0.17–1.31), respectively. Moreover, there was no significant difference in HBeAg loss between ADV and LAM/ADV (54.2% versus 64.3%; OR, 0.66; 95% CI: 0.36–1.22).

3.4.2. Network Meta-Analysis. On Bayesian network meta-analysis, when comparing TDF to others, including ETV, ADV, and LAM/ADV, no drug was clearly superior to others (Sup. Table 3) in the rates of HBeAg loss after 48 weeks of therapy.

TDF, LAM/ADV, and ADV had the highest probability of being ranked first, second, and third for achieving HBeAg loss after 48-week treatment, respectively, whereas ETV had highest probability of being ranked fourth (Sup. Figure 6).

3.5. Viral Breakthrough and Genotypic Resistance

3.5.1. Direct Meta-Analysis. After 48-week treatment (Figure 2(a)), no significant difference in the rate of virological breakthrough and genotypic resistance was observed when comparing TDF to LAM/ADV (0 versus 6.5%; OR, 0.21; 95% CI: 0.01–4.50). However, ETV had higher rate of virological breakthrough and genotypic resistance compared to LAM/ADV (12.4% versus 0.79%; OR, 5.84; 95% CI: 1.15–29.78).
Table 1: Characteristics of the included studies.

<table>
<thead>
<tr>
<th>Author</th>
<th>(year)</th>
<th>Interventions</th>
<th>Group (n)*</th>
<th>Country</th>
<th>Age (y)</th>
<th>Baseline HBV DNA level (log10 IU/mL)</th>
<th>Baseline ALT level (U/L)</th>
<th>HBeAg+ (%)</th>
<th>Study design</th>
</tr>
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<tbody>
<tr>
<td>Hann et al. [15]</td>
<td>(2008)</td>
<td>TDF (NA)</td>
<td>TDF (44)</td>
<td>USA</td>
<td>46 (11)</td>
<td>6.4 (1.6)</td>
<td>91 (161)</td>
<td>78</td>
<td>nRCT</td>
</tr>
<tr>
<td></td>
<td></td>
<td>ADV (NA)</td>
<td>ADV (65)</td>
<td></td>
<td></td>
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<td></td>
<td></td>
</tr>
<tr>
<td>Ong et al. [16]</td>
<td>(2011)</td>
<td>TDF (NA)</td>
<td>TDF (18)</td>
<td>Hong Kong, China</td>
<td>48.1 ± 11.4</td>
<td>5.78 ± 1.67</td>
<td>89 (44–210)</td>
<td>50</td>
<td>nRCT</td>
</tr>
<tr>
<td></td>
<td></td>
<td>ETV (NA)</td>
<td>ETV (11)</td>
<td></td>
<td></td>
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<tr>
<td>Maklad et al. [18]</td>
<td>(2014)</td>
<td>LAM/ADV*</td>
<td>LAM/ADV (25)</td>
<td>Egypt</td>
<td>38.5 ± 11.5</td>
<td>5.57 ± 4.41</td>
<td>69.75 ± 31.5</td>
<td>41.4</td>
<td>nRCT</td>
</tr>
<tr>
<td></td>
<td></td>
<td>ETV*</td>
<td>ETV (13)</td>
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<td></td>
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<tr>
<td>Ze et al. [19]</td>
<td>(2014)</td>
<td>LAM/ADV*</td>
<td>LAM/ADV (36)</td>
<td>Korea</td>
<td>51.85 ± 10.45</td>
<td>6.85 ± 1.1</td>
<td>172 ± 155</td>
<td>63.45</td>
<td>nRCT</td>
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<td></td>
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<td>ETV*</td>
<td>ETV (24)</td>
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<td>Chung et al. [20]</td>
<td>(2011)</td>
<td>LAM/ADV*</td>
<td>LAM/ADV (32)</td>
<td>Korea</td>
<td>44 (21–64)</td>
<td>7.36 (5.31–9.48)</td>
<td>122 (22–887)</td>
<td>88.15</td>
<td>RCT</td>
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<td></td>
<td></td>
<td>ETV*</td>
<td>ETV (44)</td>
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<tr>
<td>Ryu et al. [21]</td>
<td>(2010)</td>
<td>LAM/ADV*</td>
<td>LAM/ADV (47)</td>
<td>Korea</td>
<td>45.3 ± 9.57</td>
<td>6.69 ± 1.46</td>
<td>165.37 ± 185.5</td>
<td>69.3</td>
<td>nRCT</td>
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<tr>
<td></td>
<td></td>
<td>ETV*</td>
<td>ETV (45)</td>
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<td></td>
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<td></td>
<td></td>
</tr>
<tr>
<td>Kim et al. [22]</td>
<td>(2010)</td>
<td>LAM/ADV*</td>
<td>LAM/ADV (36)</td>
<td>Korea</td>
<td>42.8 ± 10.15</td>
<td>5.1 ± 1.12</td>
<td>100.3 ± 38.45</td>
<td>84.05</td>
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<tr>
<td></td>
<td></td>
<td>ADV*</td>
<td>ADV (44)</td>
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</tr>
<tr>
<td>Wang et al. [23]</td>
<td>(2014)</td>
<td>LAM/ADV*</td>
<td>LAM/ADV (48)</td>
<td>China</td>
<td>42.8 ± 10.15</td>
<td>5.1 ± 1.12</td>
<td>100.3 ± 38.45</td>
<td>84.05</td>
<td>nRCT</td>
</tr>
</tbody>
</table>

*CHB patients with LAM-R treated with tenofovir disoproxil fumarate (TDF), lamivudine plus adefovir (LAM/ADV), entecavir (ETV), or adefovir (ADV), respectively.

TDF$: patients treated with 300g/d TDF; LAM/ADV*: patients treated with 100g/d LAM and 10mg/d ADV; ETV*: patients treated with 1mg/d ETV; ADV*: patients treated with 10 mg/d ADV; NA: data not available.
1.1.1 TDF-LAM-R versus ADV-LAM-R

Hann et al., 2008 22 44 13 65 100.0% 4.00 [1.71, 9.34]
Subtotal (95% CI) 22 44 13 65 100.0% 4.00 [1.71, 9.34]
Total events Heterogeneity: not applicable
Test for overall effect: Z = 3.21 (P = 0.001)

1.1.2 ETV-LAM-R versus ADV-LAM-R

Kim et al., 2010 8 24 12 44 100.0% 1.33 [0.45, 3.92]
Subtotal (95% CI) 8 24 12 44 100.0% 1.33 [0.45, 3.92]
Total events Heterogeneity: not applicable
Test for overall effect: Z = 0.52 (P = 0.60)

1.1.3 ETV-LAM-R versus (LAM-ADV)-LAM-R

Chong et al., 2011 4 51 5 31 32.1% 4.00 [1.71, 9.34]
Subtotal (95% CI) 4 51 5 31 32.1% 4.00 [1.71, 9.34]
Total events 19
Heterogeneity: r^2 = 1.31; χ^2 = 7.99, df = 2 (P = 0.02); I^2 = 75%
Test for overall effect: Z = 2.70 (P = 0.007)

1.1.4 ADV-LAM-R versus (LAM-ADV)-LAM-R

Kim et al., 2010 12 44 25 36 35.9% 1.33 [0.45, 3.92]
Subtotal (95% CI) 12 44 25 36 35.9% 1.33 [0.45, 3.92]
Total events 29
Heterogeneity: r^2 = 0.07; χ^2 = 1.32, df = 1 (P = 0.25); I^2 = 24%
Test for overall effect: Z = 3.75 (P = 0.0002)

Test for subgroup differences: χ^2 = 24.60, df = 3 (P < 0.0001); I^2 = 87.8%

(a)

(b) Figure 1: Continued.
After 96-week treatment (Figure 2(b)), there was no significant difference in virological breakthrough and genotypic resistance between ETV and ADV (25% versus 13.6%; OR, 2.11; 95% CI: 0.60–7.46). However, ETV (22.9% versus 4.2%; OR, 5.98; 95% CI: 1.70–21.10) or ADV (26% versus 10.7%; OR, 3.22; 95% CI: 1.37–7.55) had higher rate of virological breakthrough and genotypic resistance compared to LAM/ADV, respectively.

3.5.2. Network Meta-Analysis. On Bayesian network meta-analysis, ETV had higher virological breakthrough and genotypic resistance as compared to TDF (OR, 31.37; 95% CrI: 1.62–605.81) or LAM/ADV (OR, 6.49; 95% CrI: 1.48–28.54) through 48 weeks of treatment (Sup. Table 4).

3.6. Sensitivity Analysis. Additionally, in direct meta-analysis, we only found there was clinical heterogeneity in 24-week ($I^2 = 75\%$) and 48-week ($I^2 = 59\%$) undetectable HBV DNA rate between ETV group and LAM plus ADV group. In 24-week virological response, the heterogeneity was brought by Maklad et al. 2014 study [18]; ETV had lower rate of virological response compared to LAM/ADV (OR, 0.29; 95% CI: 0.12–0.68; $P = 0.004$; $I^2 = 0\%$; Sup. Figure 7(A)) after removing this study. In 48-week virological response, the heterogeneity was brought by Ong et al’s 2011 study [16]. Because we did network meta-analysis of 48 weeks’ virological response, heterogeneity in 48 weeks’ virological response should be discussed to find its influence on the synthetic results of network meta-analysis.

When the heterogeneity was removed by excluding one study [16], there was still no significant difference in the direct meta-analysis of 48 weeks’ virological response between ETV and LAM/ADV (OR, 0.63; 95% CI: 0.38–1.03; $P = 0.064$; $I^2 = 3.3\%$; Sup. Figure 7(B)). In the network meta-analysis, TDF still had higher rates of HBV DNA undetectable as compared to ETV (OR, 25.58; 95% CrI: 4.67–140.24), ADV (OR, 42.43; 95% CrI: 11.25–160.06), or LAM/ADV (OR, 16.55; 95% CrI: 3.74–73.29) through 48 weeks of treatment. No drug was superior to others among ETV, ADV, and LAM/ADV. Also no change was made in rank probability of 48 weeks’ HBV DNA undetectable rate. Therefore, even though heterogeneity was found in the direct meta-analysis of 48 weeks’ virological response between ETV and LAM/ADV, it left no statistical influence on the synthetic results of network meta-analysis.

4. Discussion

This Bayesian network meta-analysis is the first study to evaluate the relative efficacy of TDF, ETV, LAM plus ADV, and ADV compared with each other for patients with LAM-R, in terms of rates of undetectable HBV DNA, ALT normalization, HBeAg loss, and virological breakthrough.

In direct meta-analysis, TDF showed a stronger antiviral effect than any one of ETV, LAM/ADV, and ADV against LAM-R hepatitis B virus. When comparing TDF to ETV, LAM/ADV, or ADV, the proportion of patients with serum HBV DNA levels < 400 copies/mL was 93.5% versus 14.8%, 88.9% versus 36.4%, or 86.4% versus 21.5% at 48 weeks, respectively. Meanwhile, LAM/ADV combination therapy was superior to ADV in suppressing viral replication for patients with LAM-R, and the proportion of patients with undetectable HBV DNA was 63.1% versus 30.2%, 75.0% versus 42.7%, and 83.3% versus 50.0% when comparing LAM/ADV to ADV at 24, 48, and 96 weeks, respectively. However, ETV had similar rates of HBV DNA undetectable compared to ADV or LAM/ADV. The rates of HBV DNA undetectable for ADV were 21.6% versus 40.7%, 32.3% versus 34.2%, and 48.6% versus 41.2% when comparing LAM/ADV through 24, 48, and 96 weeks’ therapy, respectively. The rates of undetectable HBV DNA were 33.3% versus 27.3% and 54.2% versus 40.9% when comparing ETV to ADV at 24, 48, weeks, respectively. In the network meta-analysis, the results were basically similar to the results of the direct meta-analysis. Of note, on comparative effectiveness network meta-analysis among TDF, ETV, LAM/ADV, and ADV for patients with LAM-R by undetectable HBV DNA rate at 48 weeks, TDF and LAM/ADV had the highest probability of ranking first and second, respectively, whereas ETV and ADV had highest probability of being ranked third and fourth, respectively.
ALT normalization and HBeAg loss are usually associated with improved clinical outcomes. In this study, among TDF, ETV, LAM/ADV, and ADV, the results of direct and indirect meta-analysis revealed that no drug was clearly superior to others in the rates of ALT normalization and HBeAg loss compared with each other after 48 weeks of therapy. However, on comparative effectiveness network meta-analysis, TDF, ADV, and LAM/ADV had the highest probability of being ranked first, second, and third for improving biochemical response, respectively, whereas ETV had the worst probability of being ranked fourth. In addition, TDF, LAM/ADV, ADV, and ETV had the highest probability of being ranked first, second, third, and fourth for achieving HBeAg loss after 48-week treatment, respectively. Of note, ETV or ADV had a higher rate of virological breakthrough and genotypic resistance compared to LAM/ADV (22.9% versus 4.2%).

<table>
<thead>
<tr>
<th>Study or subgroup</th>
<th>Experimental Events Total</th>
<th>Control Events Total</th>
<th>Weight</th>
<th>Odds ratio M-H, random, 95% CI</th>
<th>Odds ratio M-H, fixed, 95% CI</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.1.1 TDF-LAM-R versus (LAM-ADV)-LAM-R</td>
<td>0 28 2 31 100.0% 0.21 [0.01, 4.50]</td>
<td></td>
<td></td>
<td></td>
<td></td>
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<tr>
<td>Yang et al., 2015</td>
<td>28 31</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Subtotal (95% CI)</td>
<td>0 2</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Total events</td>
<td>28 31</td>
<td></td>
<td></td>
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<tr>
<td>Heterogeneity: not applicable</td>
<td></td>
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<tr>
<td>Test for overall effect: Z = 1.00 (P = 0.32)</td>
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</tbody>
</table>

1.1.2 ETV-LAM-R versus (LAM-ADV)-LAM-R

<table>
<thead>
<tr>
<th>Study or subgroup</th>
<th>Experimental Events Total</th>
<th>Control Events Total</th>
<th>Weight</th>
<th>Odds ratio M-H, random, 95% CI</th>
<th>Odds ratio M-H, fixed, 95% CI</th>
</tr>
</thead>
<tbody>
<tr>
<td>Chung et al., 2011</td>
<td>12 52 0 44 31.8% 27.47 [1.58, 478.97]</td>
<td></td>
<td></td>
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<tr>
<td>Han, 2010</td>
<td>2 45 1 47 43.4% 2.14 [0.19, 24.45]</td>
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<td></td>
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<tr>
<td>Ze et al., 2014</td>
<td>1 24 0 36 24.8% 4.66 [0.18, 119.23]</td>
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<tr>
<td>Subtotal (95% CI)</td>
<td>121 127 100.0% 5.84 [1.15, 29.78]</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Total events</td>
<td>15 1</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Heterogeneity: $\chi^2 = 0.05; \chi^2 = 2.94, df = 2 (P = 0.36); I^2 = 2%$</td>
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<tr>
<td>Test for overall effect: Z = 2.12 (P = 0.03)</td>
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</tbody>
</table>

Test for subgroup differences: $\chi^2 = 3.53, df = 1 (P = 0.06); I^2 = 71.7\%$

Figure 2: Forest plot of direct meta-analysis of viral breakthrough and genotypic resistance. (a) 48 weeks; (b) 96 weeks.
Figure 3: Rank probability and network plot of 48 weeks of undetectable HBV DNA rate. (a) rank probability. Different color indicates different rank showed in the picture; for example, the lightest color means rank 4. The highest bar in the chart of one intervention represents its rank probability. Rank 1 means the best efficiency, and rank 4 means the worst. The bar which means rank 1 in TDF treatment group is the highest bar; therefore, TDF treatment had the highest probability for achieving HBV DNA undetectable after 48-week treatment; (b) network plot. The node size indicates the sample size in the treatment group that the node stands for; the thickness of the link represents the sample size of the direct comparisons.

or (26% versus 10.7%) through 96-week therapy; however, no patient included in our meta-analysis had virological breakthrough during TDF treatment.

TDF and ETV are highly potent antivirals with high genetic barriers in NA-naïve patients [27, 28]; however, direct evidence in this study demonstrated superiority of TDF over ETV for suppression of HBV DNA replication in patients with LAM-R. Although the finding was based on a single trial, our network meta-analysis by comparing TDF versus ETV, LAM/ADV versus ETV, and ADV versus ETV provided a comprehensive efficacy of ETV in patients with ADV-R, which revealed that ETV was less effective in LAM-R patients compared to TDF or LAM/ADV. Meanwhile, this study provided a comparison of the antiviral efficacy between TDF and LAM-ADV or ADV for patients with LAM-R and showed that TDF alone exerted greater viral suppression than LAM plus ADV or ADV alone therapy for these patients. Therefore, TDF mono-rescue therapy is superior to not only ETV or ADV alone but also the combination of LAM and ADV therapy in effectively suppressed viral activity in patients with LAM-R. In addition, our results showed that LAM plus ADV therapy for up to 96 weeks achieved better rates of virological suppression and resistance than ADV alone, which was consistent with previous meta-analyses by Chen et al. [25]. Although ETV alone had similar efficacy in viral suppression compared to LAM plus ADV therapy through 96 week of treatment, ETV-treated patients showed relatively high probabilities of virological breakthrough and resistance mutation compared to those of LAM plus ADV therapy. Since preventing the development of multidrug resistance is a key concern for evaluating the efficacy of rescue therapy for these patients with genotypic resistance [4, 5], therefore, taking into account all outcomes in the study, our results suggest that TDF appears to be a more effective rescue therapy than LAM/ADV, ETV, or ADV for patients with LAM-R, and ETV or ADV alone is not a reasonable therapeutic option for these patients. In addition, our network meta-analysis also presented the probabilities of ranking for all these treatment strategies by using the Bayesian approach. The results of the probabilities of ranking could help the clinicians to choose better decisions for treatment.

Several limitations regarding of our systematic review require comment. Firstly, some studies had a small sample size and some of the reports’ experimental controls were not very balanced. Secondly, it has been reported that geographic, ethnic, or disease status (such as baseline disease characteristics, genotype, pervious treatment history, etc.) differences are possibly associated with agent efficacy. Thirdly, long-term outcomes of TDF in treatment of LAM-R patients were not adequately assessed owing to limited published studies in this area.

In conclusion, TDF monotherapy appears to be a more effective rescue therapy than LAM/ADV, ETV, or ADV for patients with LAM-R. LAM and ADV combination therapy
was a better treatment option than ETV or ADV alone. ETV or ADV monotherapy is not a reasonable therapeutic option for CHB patients with LAM-R.

**Competing Interests**

No potential conflict of interests relevant to this article was reported.

**Authors’ Contributions**

Hui-Lian Wang and Xi Lu contributed equally to this study, sharing first authorship as co-first authors.

**Acknowledgments**

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**References**


