Impact of Long-Term Tenofovir-Based Treatment on Hepatocellular Carcinoma Risk in Patients With Chronic Hepatitis B Using the reREACH-B Score

Young-Suk Lim¹, Harry LA Janssen²,³, Grace LH Wong⁴, Frida Abramov⁵, Dana Tedesco⁵, Hongyuan Wang⁵, Leland J Yee⁵, Wai-Kay Seto⁶, Kosh Agarwal७, Jinlin Hou⁶, Seng Gee Lim⁶, Jia-Horng Kao¹⁰, Maria Buti¹¹,¹²

¹Asan Medical Centre, University of Ulsan College of Medicine, Seoul, South Korea; ²Toronto Centre for Liver Disease, Toronto General Hospital Research Institute, University of Hong Kong, Shatin, Hong Kong; ⁵Gilead Sciences, Inc., Foster City, CA, USA; ⁶Department of Medicine, School of Clinical Medicine, The University of Hong Kong; ⁷Institute of Liver Studies, King's College Hospital NHS Foundation Trust, London, UK; ⁸Department of Infectious Diseases, State Key Laboratory of Organ Failure Research, Nanfang Hospital, Southern Medical University Hospital (NUH), Singapore; ¹⁰Graduate Institute of Clinical Medicine, National Taiwan University College of Medicine, Taiwan; ¹¹Hospital Universitario Vall d'Hebron, Barcelona, Spain; ¹²CIBEREHD del Instituto Carlos III, Madrid, Spain

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Conclusions

- In a cohort of >1600 patients with hepatitis B e antigen (HBeAg)-positive and HBeAg-negative chronic hepatitis B (CHB) enrolled in 2 large Phase 3 studies, antiviral treatment for up to 8 years demonstrated low rates of hepatocellular carcinoma (HCC) with tenofovir alafenamide (TAF) or tenofovir disoproxil fumarate (TDF) followed by TAF treatment (1.3% and 2.2%, respectively)
- Using Revised Risk Estimation for Hepatocellular Carcinoma in Chronic Hepatitis B (reREACH-B) scoring, long-term treatment with tenofovir-based therapies showed few patients at low risk at baseline shifted to higher risk, whereas many patients at higher risk shifted to lower risk on long-term tenofovir-based treatment
- These data highlight the importance of antiviral treatment to the attenuation of HCC risk in both low-risk and high-risk patients

Plain Language Summary

- We used a risk-assessment tool to predict liver cancer risk in patients treated with tenofovir-based treatments in clinical trials
- Over 8 years, few patients (26 of 1632, or 1.6%) developed hepatocellular carcinoma; many patients (52%) with higher risk at the start of the study shifted to lower risk with long-term antiviral treatment. Importantly, few patients (8%) at lower risk shifted to high risk by year 8

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Introduction

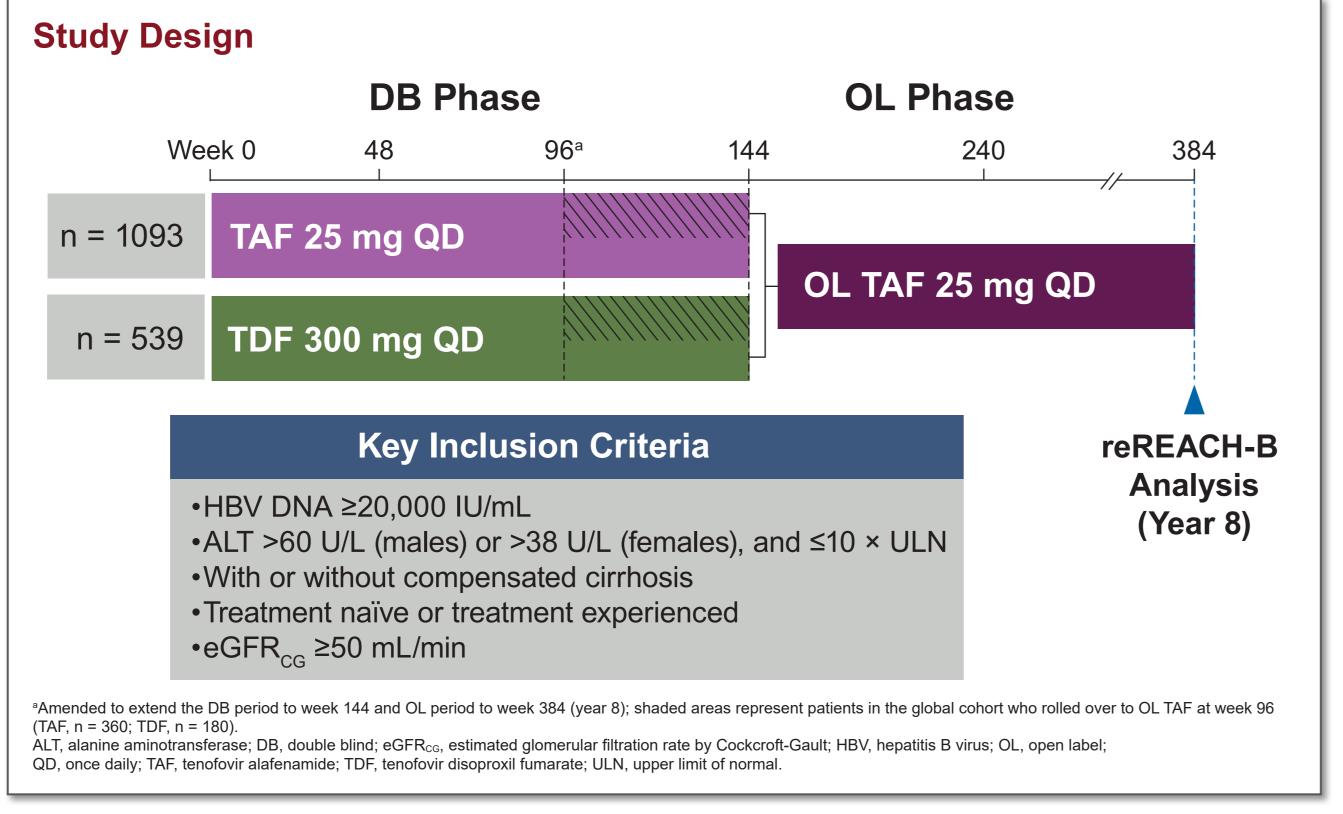
- CHB is a leading cause of HCC, accounting for over 50% of all HCC cases globally¹
- Oral antivirals can reduce but do not eliminate HCC risk^{2,3}
- Various scoring systems incorporating disease-specific factors, such as hepatitis B virus (HBV) DNA level and HBeAg status, have been developed to predict HCC risk in patients with CHB. These include nomograms, such as the REACH-B score⁴
- Recent reports indicate that the relationship between HBV DNA level and HCC risk follows a nonlinear, parabolic pattern⁵; thus, the REACH-B score was updated to incorporate age, sex, platelet count, HBV DNA level, alanine aminotransferase (ALT) level, and HBeAg status
- The reREACH-B model showed improved overall performance and offered greater clinical utility vs the REACH-B model⁶

Objective

To evaluate HCC incidence and risk based on the updated reREACH-B score in patients with HBeAg-positive and HBeAg-negative CHB enrolled in 2 Phase 3 studies evaluating tenofovir-based treatments

Methods

- This analysis used pooled data from 2 global Phase 3, randomised, double-blind, active-controlled trials
- Study 108 (N = 579): HBeAg-negative patients (global and China cohorts: NCT01940341 and NCT02836236, respectively)
- Study 110 (N = 1053): HBeAg-positive patients (global and China cohorts: NCT01940471 and NCT02836249, respectively)
- Patients were randomised 2:1, stratified by HBV DNA level and treatment status (naïve/experienced), and treated with one of the following:
- TAF 25 mg once daily (QD) for up to 144 weeks (3 years) followed by open-label (OL) TAF
 25 mg QD
- TDF 300 mg QD for up to 144 weeks (3 years) followed by OL TAF 25 mg QD
- Patients were assessed through 384 weeks (8 years)



- HCC was a predefined adverse event
- Screening, diagnosis, and treatment were per local standards of care
 - Hepatic ultrasonography (every 6 months) was added at week 96
- Predicted HCC risk was assessed at baseline and during treatment through year 8 (384 weeks)
 using reREACH-B scores
- Patients were stratified into low (<0.01) and high (≥0.01) HCC risk categories based on reREACH-B scores (minimum, 0; maximum, 1), reflecting an annual incidence rate of 0.2%, the cost-effectiveness threshold for HCC surveillance⁷
- Using the reREACH-B model, the standardised incidence ratios (SIRs) for HCC (observed cases vs model-predicted rates) with 95% CIs (calculated by Poisson regression) were determined

Results

Baseline Characteristics by HCC Risk Low Risk High Risk n = 1084 n = 1631 n = 547 Age, years, mean (SD) 33 (9.7) 44 (11.2) 40 (11.8) Treatment group, n (%) 708 (65) 384 (70) 1092 (67) TDF→TAF 539 (33) 376 (35) 163 (30) 803 (74) 1062 (65) Male sex, n (%) 259 (47) 917 (85) 1353 (83) Asian, n (%) 436 (80) Cirrhosis, n (%) 6.9 (1.67) HBV DNA, log₁₀ IU/mL, mean (SD) 7.4 (1.91) Genotype, n (%) 29 (5) 56 (5) 85 (5) 371 (23) 120 (22) 251 (23) 578 (53) 823 (50) 245 (45) 146 (27) 184 (17) 6 (1) 13 (1) 2 (<1) 3 (<1) 1 (<1) Unknown 172 (46.2) 193 (57.4) Platelets, 10³/μL, mean (SD) 235 (54.8) ALT, U/L, mean (SD) 81 (50.5) 134 (138.8) 116 (119.5) HBeAg positive, n (%) 421 (77) 1038 (64) Albumin, g/dL, mean (SD) 43 (3.9) 43 (3.7) 43 (3.4) reREACH-B score, mean (SD) 0.00 (0.003) 0.11 (0.131) 0.07 (0.118) FibroTest score, mean (SD) 0.46 (0.229) 0.38 (0.232) 0.23 (0.153) reREACH-B. Revised Risk Estimation for Hepatocellular Carcinoma in Chronic Hepatitis B: TAF, tenofovir alafenamide: TDF, tenofovir disoproxil fumarate

Compared with patients at low risk, those at high risk were more likely to be older and male; were less likely to be HBeAg positive; had lower platelet counts, lower HBV DNA levels, and higher baseline ALT levels; and were more likely to have cirrhosis

P-value for HCC vs

No HCC

<.0001

Baseline Characteristics by HCC Development HCC n = 26Age, years, mean (SD) No HCC n = 160640 (11.7)

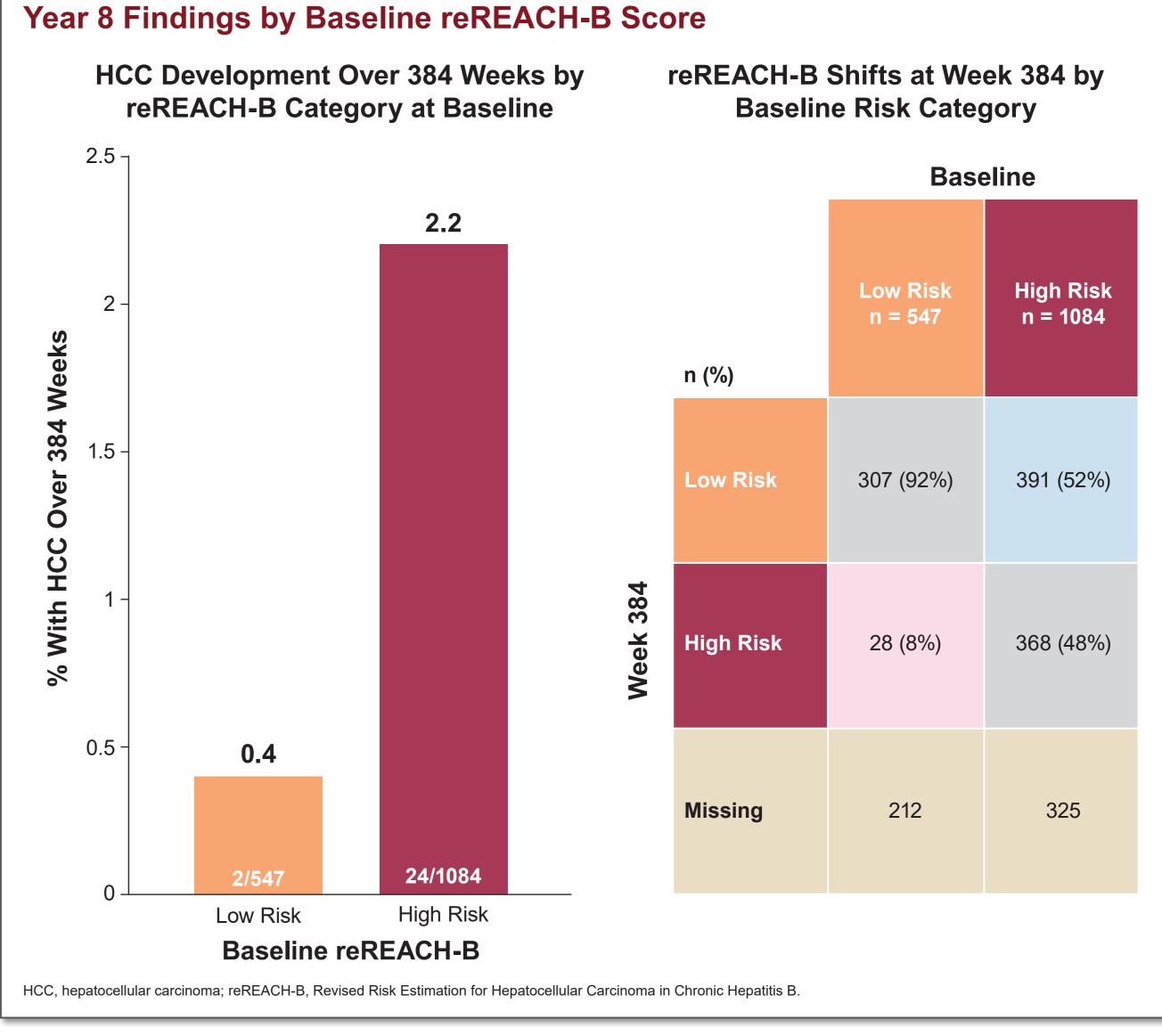
Treatment group, n (%)			
TAF	14 (54)	1079 (67)	.1515
TDF→TAF	12 (46)	527 (33)	
Male sex, n (%)	23 (88)	1040 (65)	.0119
Asian, n (%)	25 (96)	1329 (83)	.5135
Cirrhosis, n (%)	9 (35)	146 (9)	<.0001
HBV DNA, log ₁₀ IU/mL, mean (SD)	6.6 (1.10)	6.9 (1.68)	.1385
Genotype, n (%)			
A	0	85 (5)	.0705
В	2 (8)	369 (23)	
С	21 (81)	802 (50)	
D	3 (12)	328 (20)	
Other	0	19 (1)	
Unknown	0	3 (0)	
ALT, U/L, mean (SD)	84 (57.8)	117 (120.1)	.1272
HBeAg positive, n (%)	13 (50)	1026 (64)	.1443
Albumin, g/dL, mean (SD)	40.1 (4.26)	43.0 (3.67)	.0003
reREACH-B score, median (Q1, Q3)	0.21 (0.11, 0.36)	0.02 (0.01, 0.08)	<.0001
FibroTest score, mean (SD)	0.61 (0.219)	0.38 (0.230)	<.0001

Cirrhosis was defined as a FibroTest score ≥0.75 in Studies 108/110. *P*-values are from the CMH test and the 2-sided Wilcoxon rank sum test for categorical data and continuous data, respectively. Bold text indicates significant *P*-values.

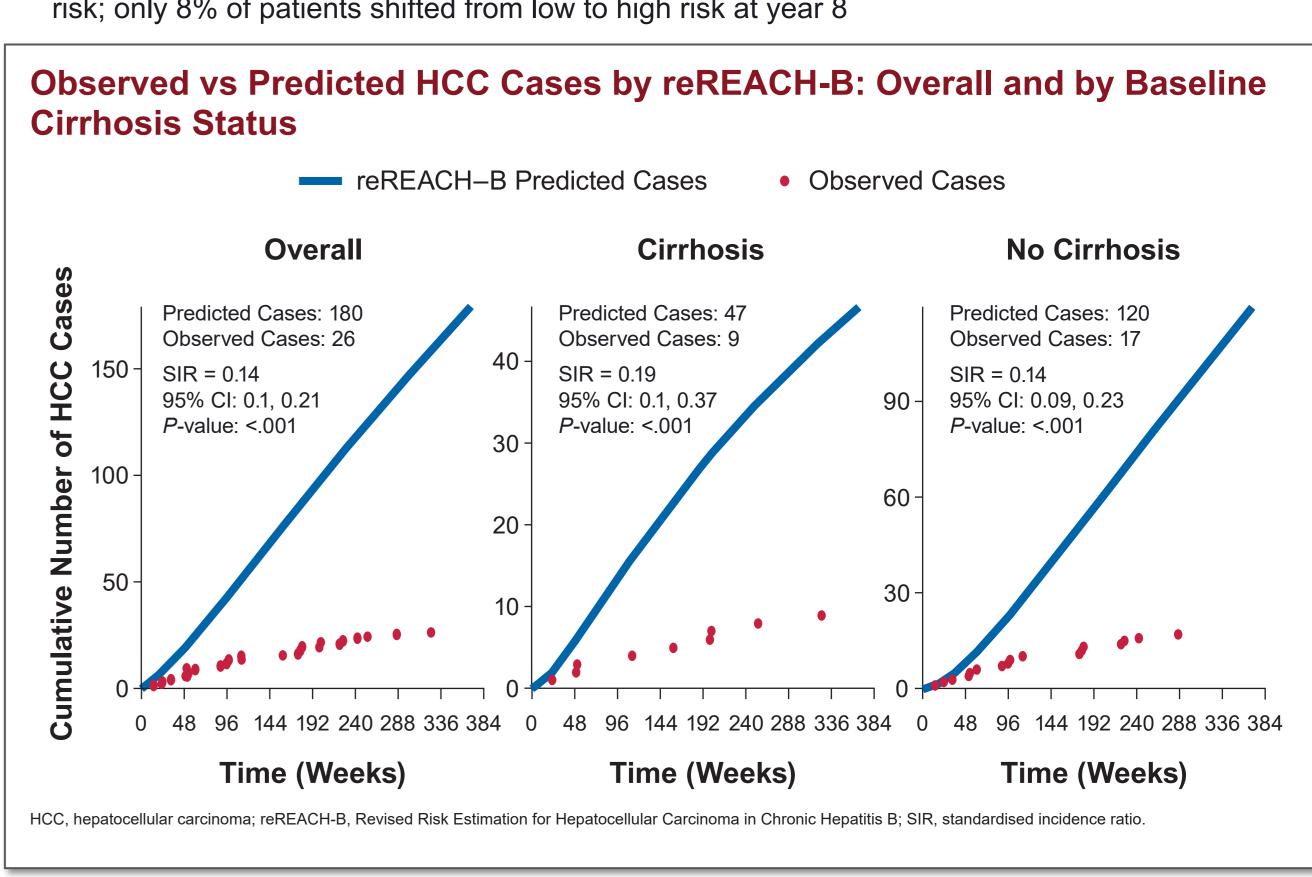
ALT, alanine aminotransferase; CMH, Cochran-Mantel-Haenszel; HBeAg, hepatitis B e antigen; HBV, hepatitis B virus; HCC, hepatocellular carcinoma; Q, quartile; reREACH-B, Revised Risk Estimation for Hepatocellular Carcinoma in Chronic Hepatitis B; TAF, tenofovir alafenamide; TDF, tenofovir disoproxil fumarate.

- Through 8 years of follow-up, HCC occurred in 26 of 1632 (1.6%) patients (TAF, 14 of 1093 [1.3%]; TDF, 12 of 539 [2.2%])
 - Of the 26 patients with HCC, 9 (35%) had cirrhosis at baseline

 Patients who developed HCC were significantly older, more likely to be male, had I
- Patients who developed HCC were significantly older, more likely to be male, had lower mean albumin, and more likely to have cirrhosis at baseline



- Among individuals characterised as high or low risk at baseline, 24 of 1084 (2.2%) and 2 of 547 (0.4%), respectively, developed HCC over 8 years
- At year 8, the mean (range) change from baseline in reREACH-B score was −0.1 (−1.0 to 0.1)
- Notably, 52% of patients deemed high risk at baseline with available data at year 8 shifted to low risk; only 8% of patients shifted from low to high risk at year 8



- Based on the SIR, the HCC incidence seen with treatment overall was 86% lower compared with the incidence predicted by the reREACH-B model
- Similar significant reductions in HCC incidence were observed with long-term antiviral treatment in patients with and without cirrhosis at baseline