

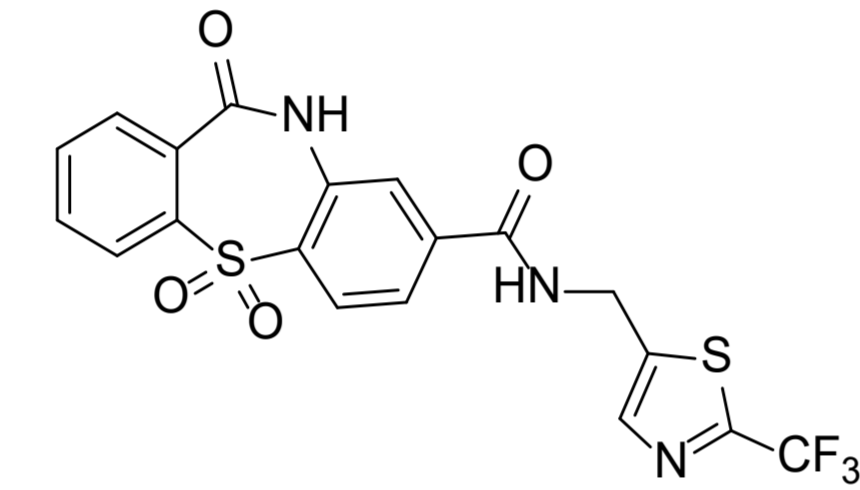
# Deeper virologic suppression with the addition of vebicorvir, a first-generation hepatitis B core inhibitor, to entecavir correlates with reduced inflammation and fibrosis-4 index in treatment-naïve patients with HBeAg positive chronic hepatitis B

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## Background

- Chronic hepatitis B virus infection (cHBV) is a significant global health problem
  - Worldwide, an estimated 296 million people have cHBV infection, resulting in approximately 887,000 deaths each year, mostly due to cirrhosis and hepatocellular carcinoma<sup>1-4</sup>
- A goal of nucleos(t)ide reverse transcriptase inhibitor (Nrti) therapy for cHBV is suppression of HBV DNA below the lower limit of quantification (LLOQ) for available assays
  - Nrtis are well tolerated, and while suppression of HBV DNA <LLOQ is achieved in most patients, low-level viremia remains, and treatment duration is indefinite<sup>5</sup>
  - Novel combination approaches incorporating agents with complementary mechanisms of action, such as core inhibitors, are likely required to further suppress viral replication, establish finite-duration regimens, and improve clinical outcomes
- Vebicorvir (VBR), a novel first-generation inhibitor of the HBV core protein (Figure 1), has previously demonstrated deeper reductions in HBV DNA and pregenomic RNA (pgRNA) and more rapid normalization of alanine aminotransferase (ALT) when added to entecavir (ETV)<sup>6,7</sup>
- In this report, we explore the associations between the observed deeper reductions in HBV parameters with changes in Fibrosis-4 (FIB-4) Index
  - FIB-4 is a validated prognostic marker of liver fibrosis used in the management of cHBV<sup>8,9</sup>
  - FIB-4 values tend to range between 0.2 and 10, with higher scores more strongly predicting the presence of fibrosis



**Figure 1. VBR is a novel, first-generation inhibitor of HBV core protein**

- Core inhibitors allosterically bind and interfere with the core protein by mechanisms distinct from Nrtis
  - Inhibition of pgRNA encapsidation, prevention of viral assembly/release, and disruption of viral capsids
- Broad in vitro antiviral activity<sup>10</sup>
  - Inhibits virion and pgRNA particle production (half-maximal effective concentration [EC<sub>50</sub>]=0.17–0.31 μM; half-maximal cytotoxic concentration [CC<sub>50</sub>] ≥20 μM)
  - Inhibits de novo formation of covalently closed circular DNA and downstream hepatitis B e antigen (HBeAg) and hepatitis B surface antigen production (EC<sub>50</sub>=2–7 μM)
  - Pangenotypic and fully active against Nrti-resistant HBV
- Orally administered as 300 mg once daily (QD) without regard to food, with a favorable clinical safety profile in Phase 2 studies<sup>11</sup>
- VBR+Nrti resulted in deeper on-treatment viral suppression in treatment-naïve and virologically-suppressed patients in Phase 2 studies<sup>6,12</sup>

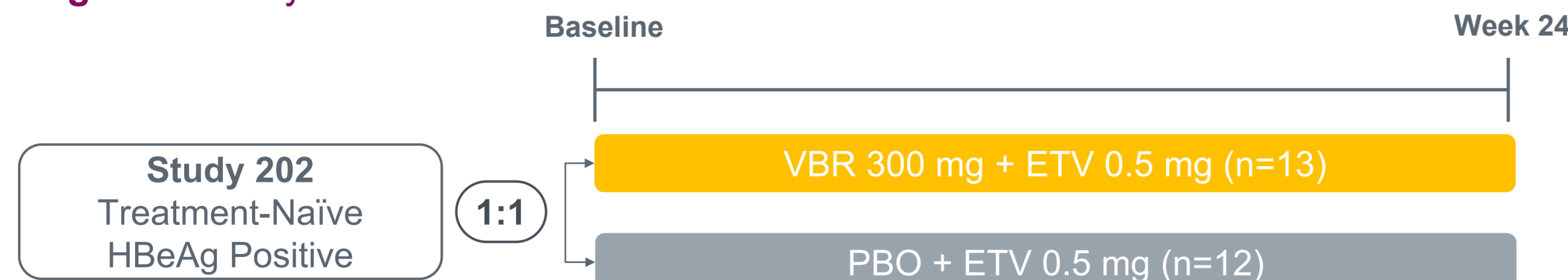
## Objective of this analysis

- This post hoc analysis from a Phase 2a study in treatment-naïve patients<sup>6,7</sup> explores correlations between the deeper level of viral suppression and more rapid normalization of ALT observed with the addition of VBR to ETV with FIB-4, a validated prognostic marker of liver fibrosis<sup>8,9</sup>

## Methods

- In this study (NCT03577171), 13 patients received VBR (300 mg) with ETV (0.5 mg) QD, and 12 patients received placebo (PBO) with ETV (0.5 mg) QD for 24 weeks (Figure 2)
- For this analysis, HBV DNA, pgRNA, and HBeAg were measured at Screening, Baseline (Day 1), and then every 4 weeks until Week 24. HBV DNA, pgRNA, and hepatitis B core antigen (HBcAg) were also measured at Week 2
  - Laboratory assessments (ALT, aspartate aminotransferase [AST], and platelets) were also assessed at the same timepoints as virological parameters
- FIB-4 was calculated according to the following formula: (age [years] × AST [U/L]) / (platelet count [10<sup>9</sup>/L] × √ALT [U/L])<sup>8</sup>
- Post hoc correlation analyses were performed, and Pearson's (for parametric analyses) and Spearman's (for nonparametric analysis) correlation coefficients were calculated where appropriate to describe associations between parameters and FIB-4
- Statistical analyses were based on a mixed-effects, repeated-measure model

**Figure 2. Study 202 overview**



## Key eligibility criteria

- Male or female, aged 18–70 years
- Treatment-naïve, HBeAg positive cHBV
- METAVIR F0–F2 or equivalent (no history of hepatic decompensation)
- HBV DNA >10<sup>5</sup> IU/mL; ALT <10× ULN; platelet count ≥100,000/mm<sup>3</sup>

ALT, alanine aminotransferase; cHBV, chronic hepatitis B virus; ETV, entecavir; HBeAg, hepatitis B e antigen; METAVIR, Meta-analysis of Histological Data in Viral Hepatitis; PBO, placebo; ULN, upper limit of normal; VBR, vebicorvir.

## Results

**Table 1. Demographics and Baseline characteristics**

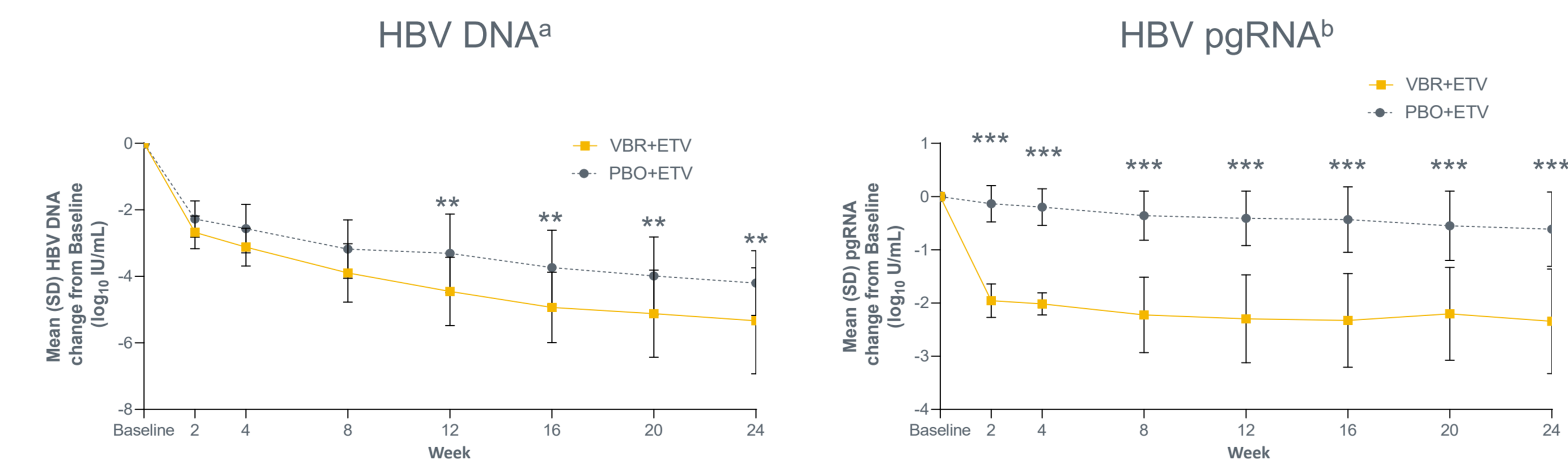
Study 202 Baseline	VBR+ETV (n=13)	PBO+ETV (n=12)	Total (N=25)
Age, years; median (min, max)	32 (20, 66)	32 (20, 57)	32 (20, 66)
Sex, female; n (%)	10 (77)	7 (58)	17 (68)
Race, Asian; n (%)	13 (100)	11 (92)	24 (96)
BMI, kg/m <sup>2</sup> ; median (min, max)	22.4 (18.4, 28.3)	21.7 (19.5, 32.3)	22.1 (18.4, 32.3)
HBV genotype; n (%)			
A	0	2 (17)	2 (8)
B	7 (54)	4 (33)	11 (44)
C	5 (38)	6 (50)	11 (44)
Unable to genotype	1 (8)	0	1 (4)
HBV DNA <sup>a</sup> , log <sub>10</sub> IU/mL; mean (SD)	7.9 (0.89)	8.0 (1.0)	8.0 (0.93)
HBV pgRNA <sup>b</sup> , log <sub>10</sub> U/mL; mean (SD)	7.1 (0.95)	7.4 (1.14)	7.2 (1.03)
HBeAg <sup>c</sup> , log <sub>10</sub> IU/mL; mean (SD)	2.5 (0.81)	2.5 (1.24)	2.5 (1.02)
HBsAg <sup>d</sup> , log <sub>10</sub> IU/mL; mean (SD)	4.5 (0.51)	4.7 (0.44)	4.6 (0.48)
HBcAg <sup>e</sup> , log <sub>10</sub> kU/mL; mean (SD)	5.5 (0.69)	5.4 (0.97)	5.4 (0.82)
ALT, U/L; mean (SD)	65.9 (87.11)	46.8 (31.64)	56.7 (65.94)
AST, U/L; mean (SD)	46.0 (42.62)	36.3 (26.28)	41.3 (35.35)
Platelets, K/μL; mean (SD)	228.2 (57.53)	227.3 (69.70)	227.8 (62.30)
INR; mean (SD)	1.0 (0.05)	1.0 (0.08)	1.0 (0.06)
FIB-4; mean score (SD)	0.98 (0.60)	0.94 (0.82)	0.96 (0.70)

<sup>a</sup>Quantified using COBAS TaqMan version 2.0 (LLOQ=20 IU/mL). <sup>b</sup>Quantified using quantitative RT-PCR method developed by Assembly Biosciences (LLOQ=135 U/mL). <sup>c</sup>Quantified using the Architect i2000SR assay (LLOQ=0.11 IU/mL). <sup>d</sup>Quantified using the Architect i2000SR assay (LLOQ=0.05 IU/mL). <sup>e</sup>Quantified using the Lumipulse G HBcAg assay (LLOQ=1 kU/mL).

ALT, alanine aminotransferase; AST, aspartate aminotransferase; BMI, body mass index; ETV, entecavir; FIB-4, Fibrosis-4; HBcAg, hepatitis B core antigen; HBeAg, hepatitis B e antigen; HBsAg, hepatitis B surface antigen; HBV, hepatitis B virus; INR, international normalized ratio; LLOQ, lower limit of quantification; PBO, placebo; pgRNA, pregenomic RNA; RT-PCR, reverse transcription polymerase chain reaction; SD, standard deviation; VBR, vebicorvir.

- Overall, HBV disease characteristics were similar between treatment groups at Baseline (Table 1)
- Most patients were Asian (24/25 [96%]) and female (17/25 [68%]), with ages ranging from 20–66 years and body mass index ranging from 18.4–32.3 kg/m<sup>2</sup>
- Of the 25 total patients, 11 were infected with HBV genotype B and 11 with genotype C
- In these untreated patients, the overall mean (standard deviation [SD]) HBV DNA and pgRNA at Baseline were 8.0 (0.93) log<sub>10</sub> IU/mL and 7.2 (1.03) log<sub>10</sub> U/mL, respectively
- Mean (SD) FIB-4 values at Baseline for VBR+ETV patients were 0.98 (0.60) and 0.94 (0.82) for PBO+ETV patients, corresponding to Ishak 0–1 fibrosis

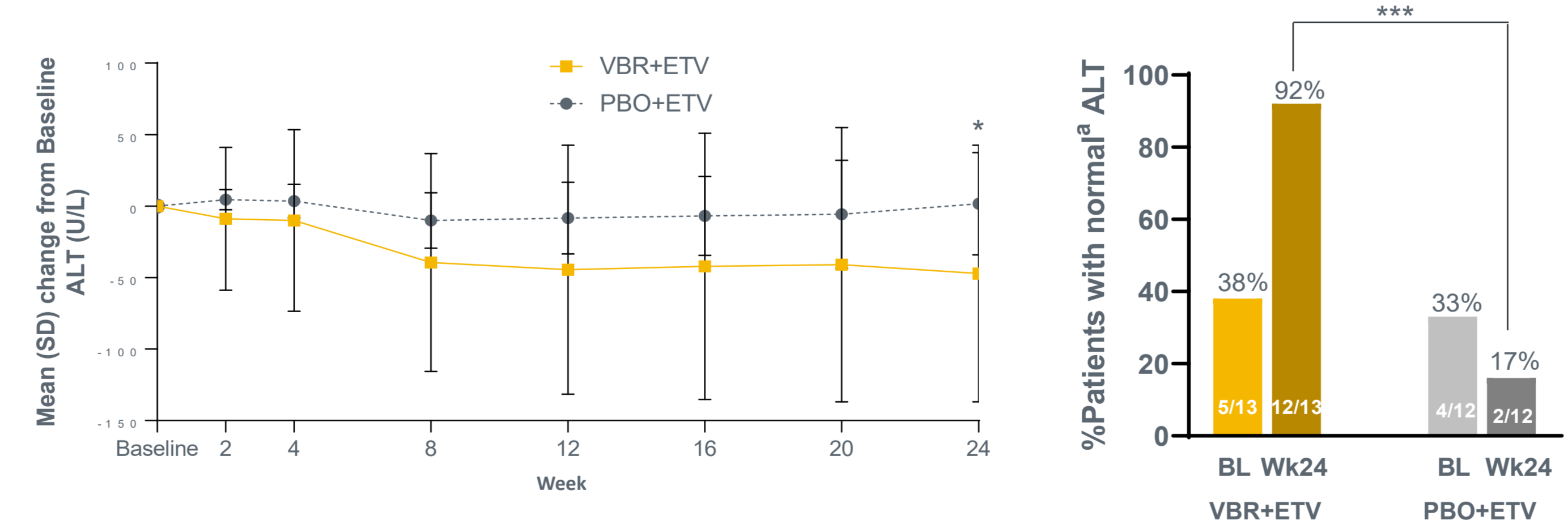
**Figure 3. Mean HBV DNA and pgRNA change from Baseline over 24 weeks**



<sup>a</sup>HBV DNA LLOQ=20 IU/mL; <sup>b</sup>HBV pgRNA LLOQ=135 U/mL. <sup>\*</sup>P<0.01; <sup>\*\*</sup>P<0.001. ETV, entecavir; HBV, hepatitis B virus; LLOQ, lower limit of quantification; PBO, placebo; pgRNA, pregenomic RNA; SD, standard deviation; VBR, vebicorvir.

- More profound HBV DNA and pgRNA reductions were observed with VBR+ETV compared to ETV monotherapy (Figure 3), as previously described<sup>6,7</sup>

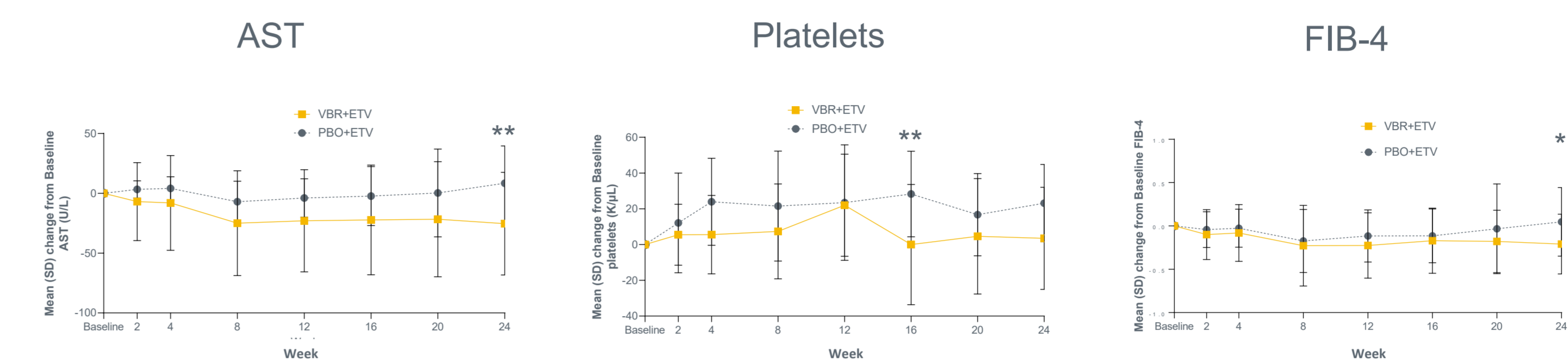
**Figure 4. ALT change from Baseline and normalization with VBR+ETV**



<sup>a</sup>Normal ALT consists of values less than ULN; ALT ULN is 25 U/L for females and 35 U/L for males per AASLD criteria. <sup>\*</sup>P<0.05; <sup>\*\*</sup>P<0.001. AASLD, American Association for the Study of Liver Diseases; ALT, alanine aminotransferase; BL, Baseline; ETV, entecavir; PBO, placebo; SD, standard deviation; ULN, upper limit of normal; VBR, vebicorvir; Wk, week.

- Most patients receiving VBR+ETV experienced a numerical decrease in ALT from Baseline at Week 24
- At Week 24, an overall mean reduction in ALT was observed with VBR+ETV but not PBO+ETV (Figure 4)
- At Baseline, 5/13 (38%) and 4/12 (33%) VBR+ETV and PBO+ETV patients had normal ALT; at Week 24, those proportions were 12/13 (92%) for VBR+ETV compared to 2/12 (17%) for PBO+ETV

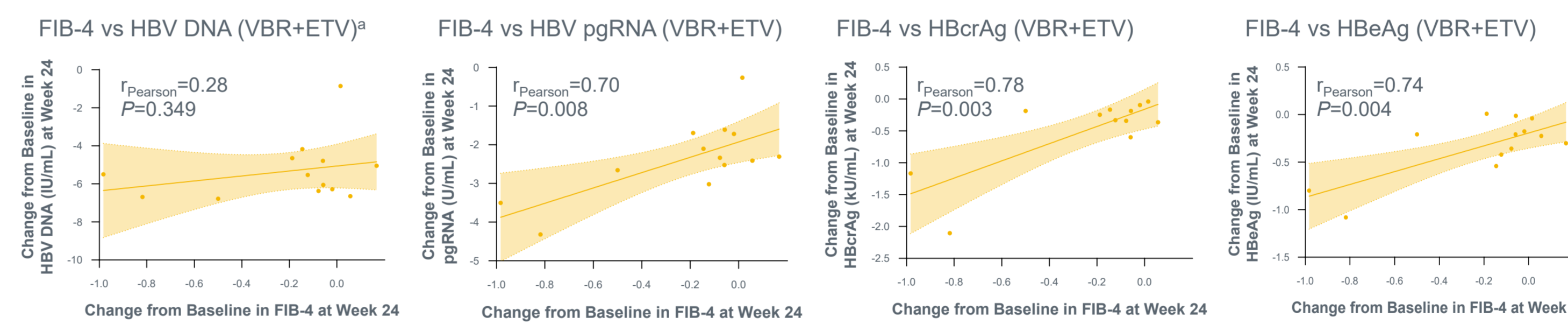
**Figure 5. AST, platelets, and FIB-4 mean changes from Baseline**



<sup>\*</sup>P<0.05; <sup>\*\*</sup>P<0.01. AST, aspartate aminotransferase; ETV, entecavir; FIB-4, fibrosis-4; PBO, placebo; SD, standard deviation; VBR, vebicorvir.

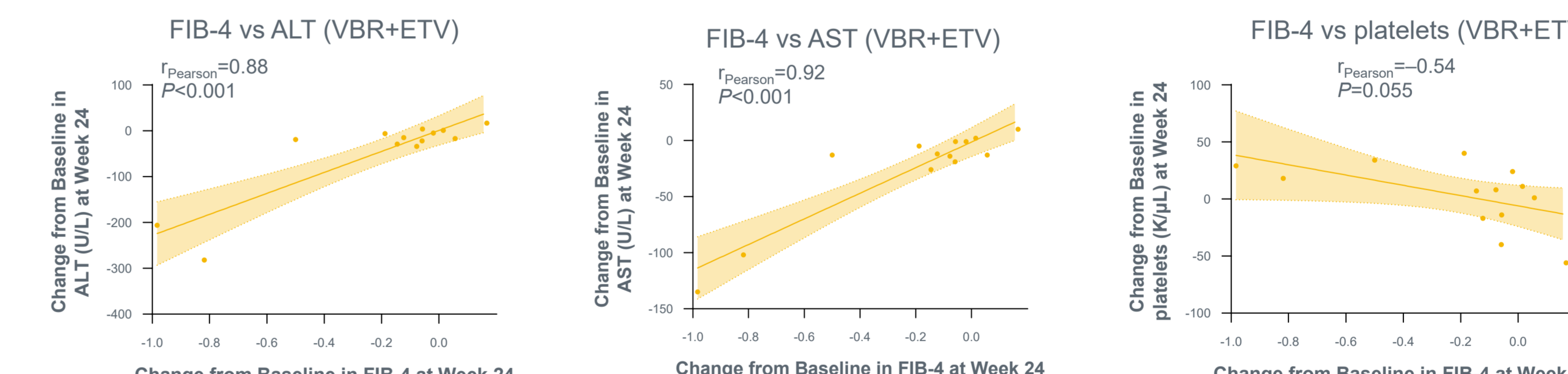
- At Week 24, an overall mean reduction in AST was observed with VBR+ETV but not with PBO+ETV (Figure 5)
- PBO+ETV recipients experienced a mean increase in platelet count of 23.2 K/μL over the treatment interval, which was numerically greater than the 3.5 K/μL mean increase observed in patients receiving VBR+ETV for 24 weeks
- While patients entering the study had a similar mean FIB-4 index at Baseline, the mean change from Baseline was significantly greater for VBR+ETV (-0.21) vs PBO+ETV (0.05; P=0.034) at Week 24
- Overall, treatment with VBR+ETV resulted in statistically significant improvements in HBV DNA, pgRNA, ALT, AST, and FIB-4 compared to PBO+ETV
- To further explore these findings, additional correlation analyses were performed to assess relationships between FIB-4 and ALT, AST, platelets, and viral parameters

**Figure 6. Associations between FIB-4 and HBV parameters in VBR+ETV patients at Week 24**



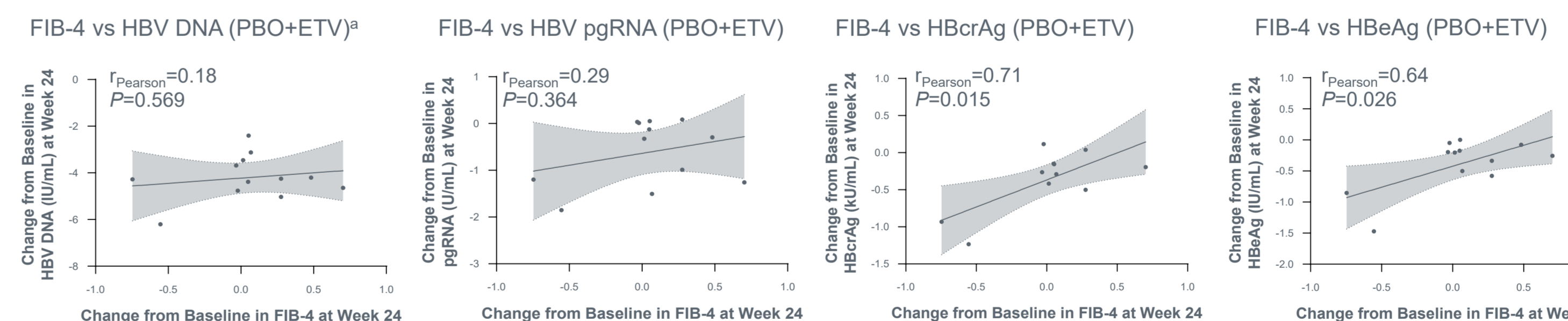
<sup>r</sup><sub>Pearson</sub>=0.18, P=0.566. ETV, entecavir; FIB-4, fibrosis-4; HBcAg, hepatitis B core antigen; HBeAg, hepatitis B e antigen; HBV, hepatitis B virus; pgRNA, pregenomic RNA; <sup>r</sup><sub>Pearson</sub>, Pearson's correlation coefficient; <sup>r</sup><sub>Spearman</sub>, Spearman's correlation coefficient; VBR, vebicorvir.

**Figure 7. Associations between FIB-4 and ALT, AST, and platelets in VBR+ETV patients at Week 24**



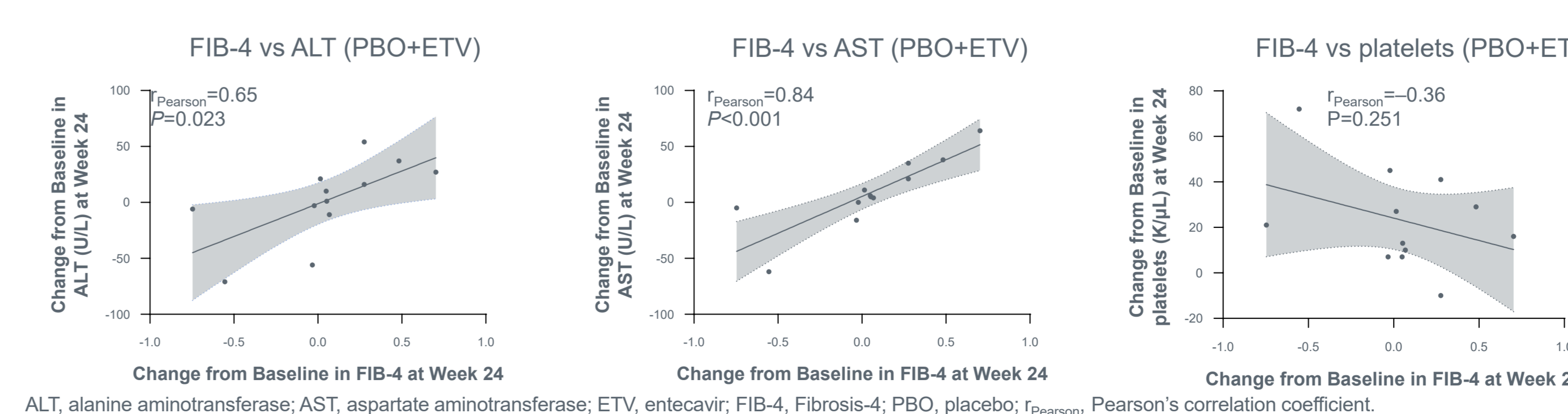
ALT, alanine aminotransferase; AST, aspartate aminotransferase; ETV, entecavir; FIB-4, Fibrosis-4; <sup>r</sup><sub>Pearson</sub>, Pearson's correlation coefficient; VBR, vebicorvir.

**Figure 8. Associations between FIB-4 and HBV parameters in PBO+ETV patients at Week 24**



<sup>r</sup><sub>Spearman</sub>=0.11; P=0.729. ETV, entecavir; FIB-4, fibrosis-4; HBcAg, hepatitis B core antigen; HBeAg, hepatitis B e antigen; HBV, hepatitis B virus; PBO, placebo; pgRNA, pregenomic RNA; <sup>r</sup><sub>Pearson</sub>, Pearson's correlation coefficient; <sup>r</sup><sub>Spearman</sub>, Spearman's correlation coefficient.

**Figure 9. Associations between FIB-4 and ALT, AST, and platelets in PBO+ETV patients at Week 24**



ALT, alanine aminotransferase; AST, aspartate aminotransferase; ETV, entecavir; FIB-4, Fibrosis-4; PBO, placebo; <sup>r</sup><sub>Pearson</sub>, Pearson's correlation coefficient.

- At Week 24, the deeper reductions in HBV DNA, pgRNA, ALT, and AST that were observed with VBR+ETV compared to PBO+ETV were reflected in greater reductions in the composite FIB-4 Index
- For VBR+ETV, significant, strong correlations were observed between FIB-4 and pgRNA (<sup>r</sup><sub>Pearson</sub>=0.70; P=0.008), HBcAg (<sup>r</sup><sub>Pearson</sub>=0.78; P=0.003), HBeAg (<sup>r</sup><sub>Pearson</sub>=0.74; P=0.004), ALT (<sup>r</sup><sub>Pearson</sub>=0.88; P<0.001), and AST (<sup>r</sup><sub>Pearson</sub>=0.92; P<0.001), (Figures 6 and 7)
- For PBO+ETV, significant, strong correlations were observed between FIB-4 and HBcAg (<sup>r</sup><sub>Pearson</sub>=0.71; P=0.015) and AST (<sup>r</sup><sub>Pearson</sub>=0.84; P<0.001), while there were moderate correlations between FIB-4 and HBeAg (<sup>r</sup><sub>Pearson</sub>=0.64; P=0.026) and ALT (<sup>r</sup><sub>Pearson</sub>=0.65; P=0.023) (Figures 8 and 9)
- FIB-4 was not significantly correlated with platelets in either treatment arm at Week 24
- When examining both Pearson's and Spearman's correlation coefficients, there were no significant correlations between FIB-4 and HBV DNA for either treatment group at Week 24

## Limitations of this analysis

- The small sample size and 24-week treatment duration limit interpretation and generalizability. Studies of longer treatment duration and larger sample size will be required to confirm the observations

## Conclusions

- VBR+ETV demonstrated potent antiviral activity compared to PBO+ETV in treatment-naïve, HBeAg positive patients as demonstrated by:
  - More rapid and deeper declines in HBV DNA and pgRNA
  - Greater decrease and more rapid normalization of ALT and AST
- These changes were reflected in a reduction in FIB-4 for VBR+ETV recipients and an increase for patients receiving PBO+ETV. The difference in FIB-4 between the treatment arms at Week 24 was statistically significant
- When assessing associations between viral and host factors with FIB-4 at the end of the 24-week treatment period:
  - For VBR+ETV, strong positive correlations were observed between FIB-4 and pgRNA, HBcAg, HBeAg, ALT, and AST,
  - For PBO+ETV, strong positive correlations were observed between FIB-4 and HBcAg and AST and moderate correlations between FIB-4 and HBeAg and ALT
- The increased antiviral potency of VBR+ETV compared to PBO+ETV resulted in greater reductions in viral replication and hepatic inflammation and significantly greater improvement in FIB-4
- Regimens providing deeper levels of viral suppression and greater reductions in hepatic inflammation may further improve long-term clinical outcomes

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