

# Hepatoprotective effects of the oral fatty acid synthase inhibitor TVB-3664 in the GAN diet-induced obese and biopsy-confirmed mouse model of MASH

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## Background & Aim

Fatty acid synthase (FASN) inhibitors is a promising drug class for the treatment of metabolic dysfunction-associated steatohepatitis (MASH). Accordingly, the oral FASN inhibitor denifanstat (TVB-2640) has recently been reported to improve NAFLD Activity Score (NAS) and fibrosis stage in a phase 2b clinical trial (FASCINATE-2) in MASH patients. The present study aimed to investigate the effect of TVB-3664, an oral FASN inhibitor, on metabolic, biochemical and histological endpoints in the translational Gubra Amylin NASH (GAN) diet-induced obese and biopsy-confirmed mouse model of MASH with liver fibrosis.

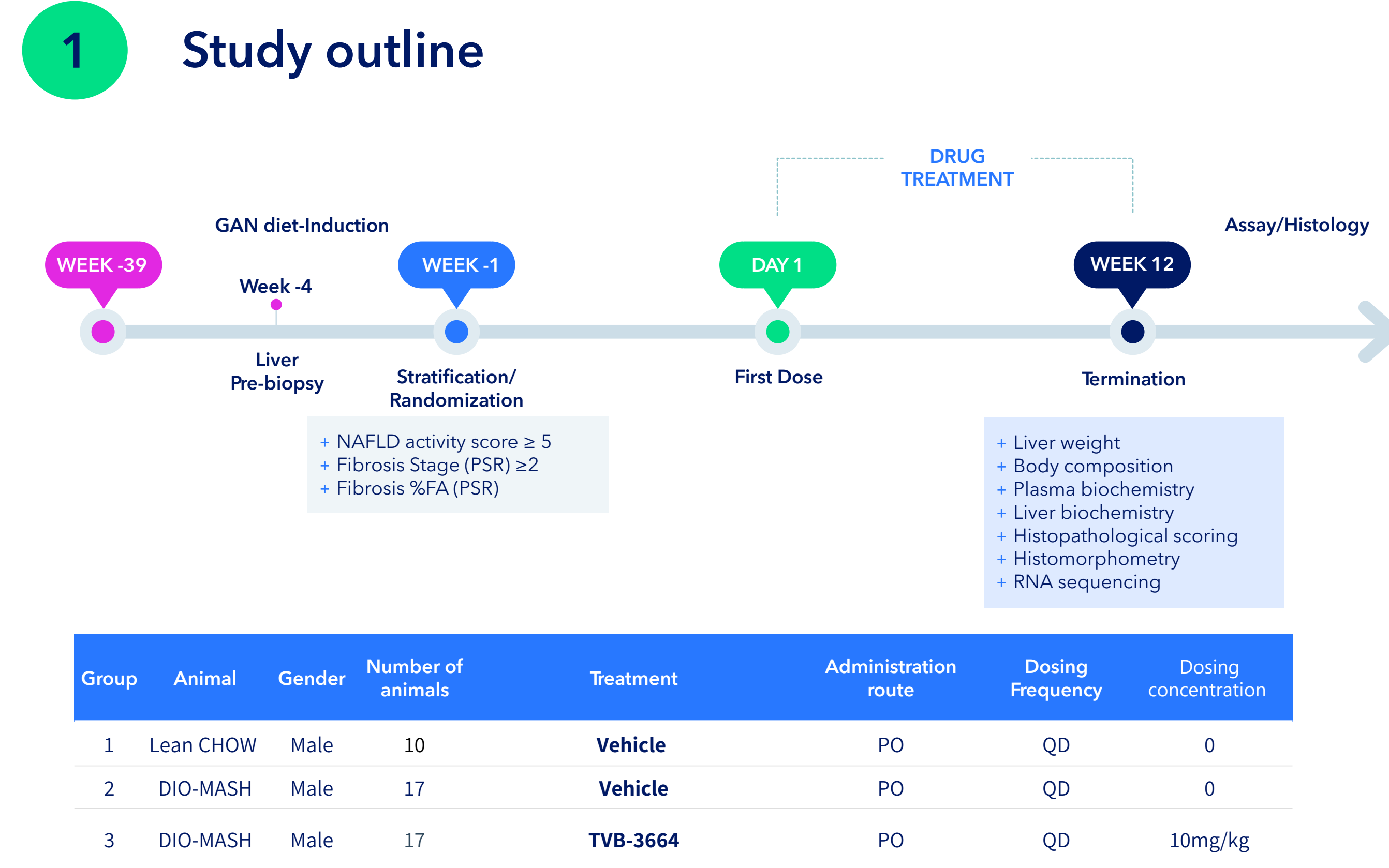


Figure 1. Study outline. PO; per oral, QD; Once daily; GAN; Gubra Amylin NASH.

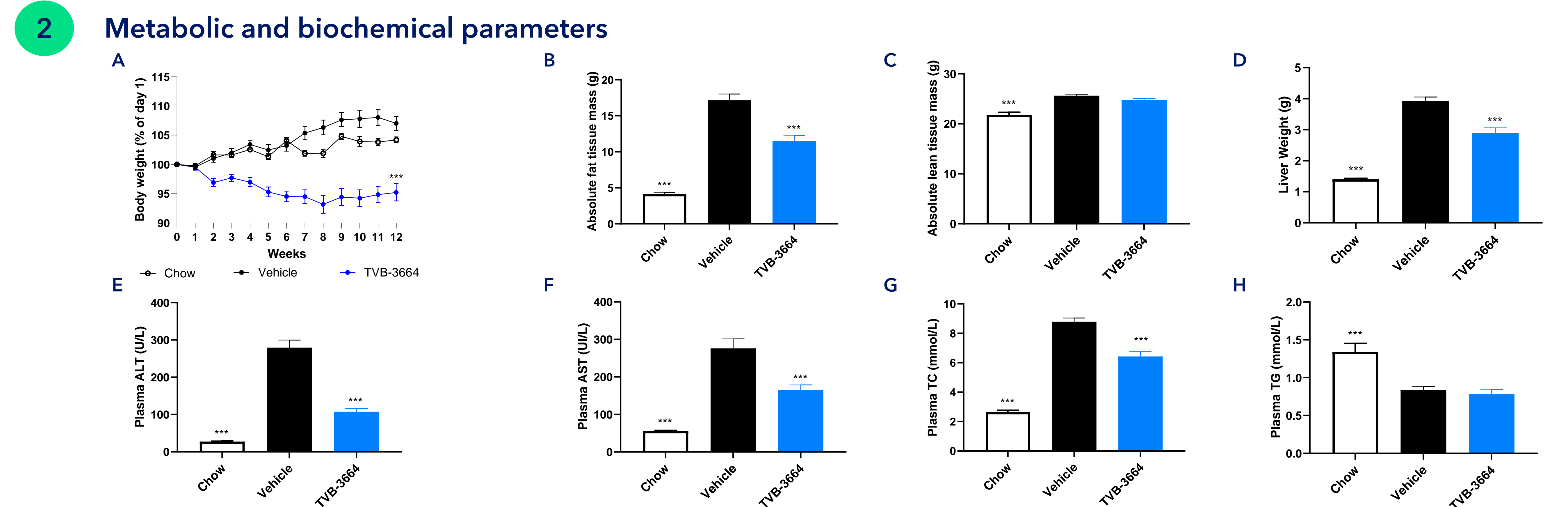


Figure 2. Metabolic and biochemical effects of TVB-3664. (A) Relative body weight during study period. (B) Terminal absolute fat tissue mass. (C) Terminal absolute lean tissue mass. (D) Terminal liver weight. (E) plasma alanine aminotransferase (ALT). (F) Terminal plasma aspartate aminotransferase (AST). (G) Terminal plasma total cholesterol (TC). (H) Terminal plasma triglycerides (TG). \*\*\*p<0.001 compared to vehicle control (Dunnett's test one-factor linear model).

### 3 NAFLD Activity Score and Fibrosis Stage

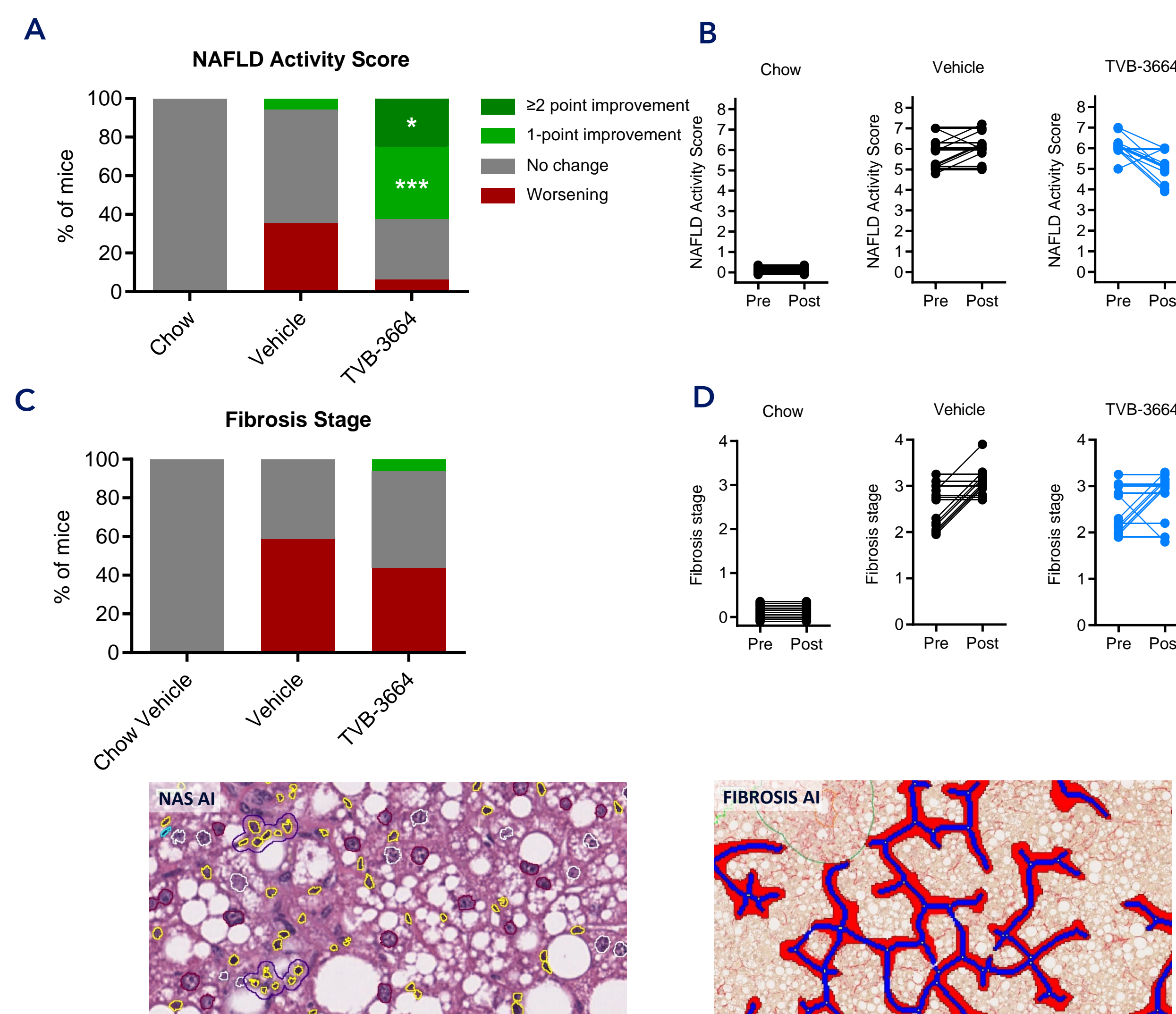


Figure 3. TVB-3664 improves NAFLD Activity Score in GAN DIO-MASH mice. Histopathological scores were determined by Gubra Histopathological Objective Scoring Technique (GHOST) deep learning-based image analysis. (A) NAFLD Activity Score (NAS). (B) Individual pre-post NAS. (C) Fibrosis stage. (D) Individual pre-post fibrosis stage. \*p<0.05, \*\*\*p<0.001 compared to vehicle control (one-sided Fisher's exact test).

### 4 Histological markers of steatosis, inflammation and fibrosis

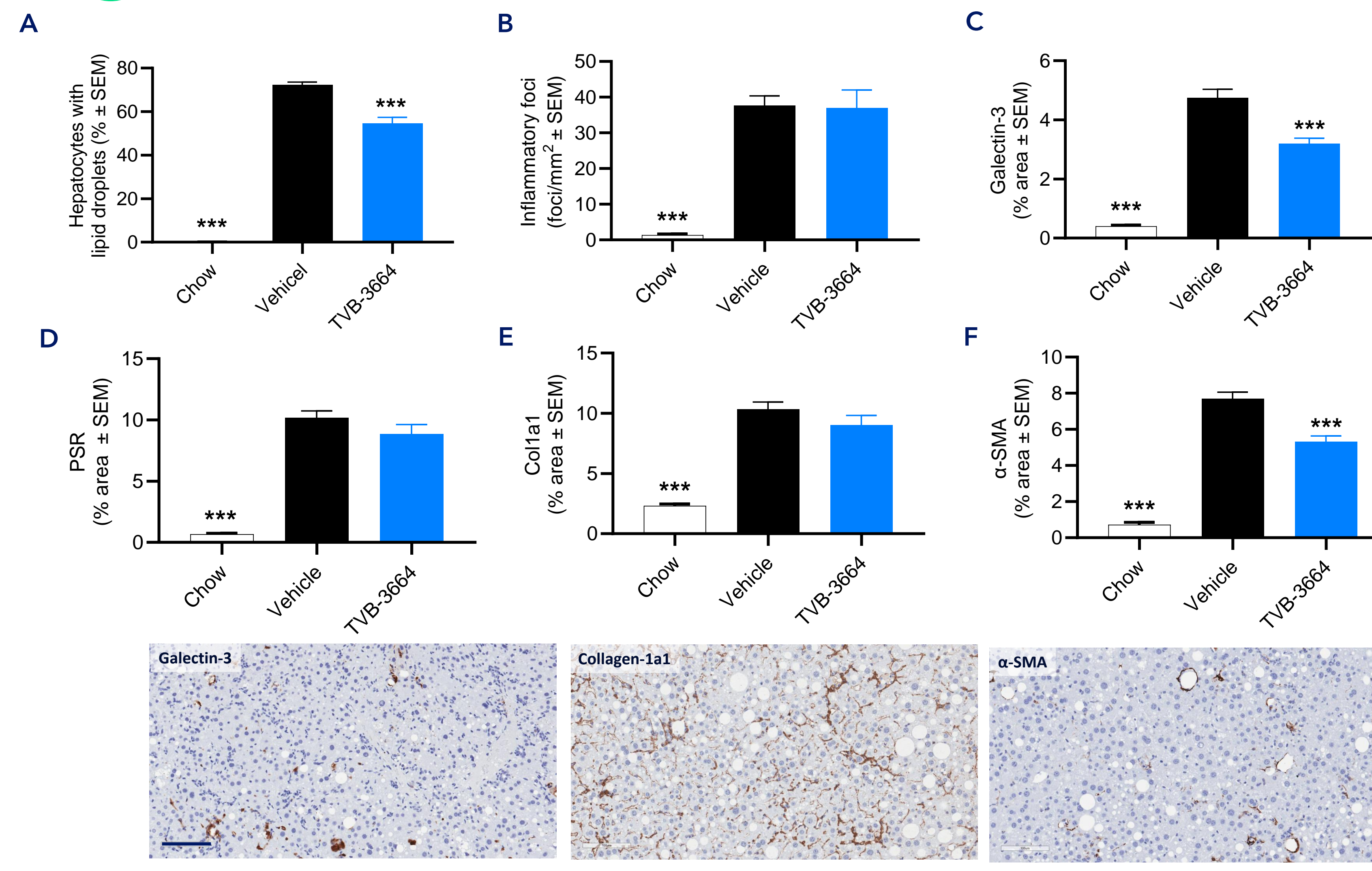


Figure 4. TVB-3664 improves quantitative histological endpoints. Histomorphometric assessments were performed by GHOST deep learning-based image analysis on scoring-associated variables (panels A-B) and conventional IHC image analysis (panels C-F). (A) % hepatocytes with lipid droplets. (B) Number of inflammatory foci. (C) % area of galectin-3 (steatosis-adjusted). (D) % area of PSR (steatosis-adjusted). (E) % area of collagen-1a1 (Col1a1, steatosis-adjusted) (F) % area of alpha-smooth muscle actin (alpha-SMA, marker of stellate cell activation; steatosis-adjusted). \*\*\*p<0.001 compared to vehicle control (Dunnett's test one-factor linear model). Bottom panels: Representative photomicrographs of galectin-3, Col1a1 and alpha-SMA (scale bar, 100 µm).

### 5 Liver transcriptome analysis

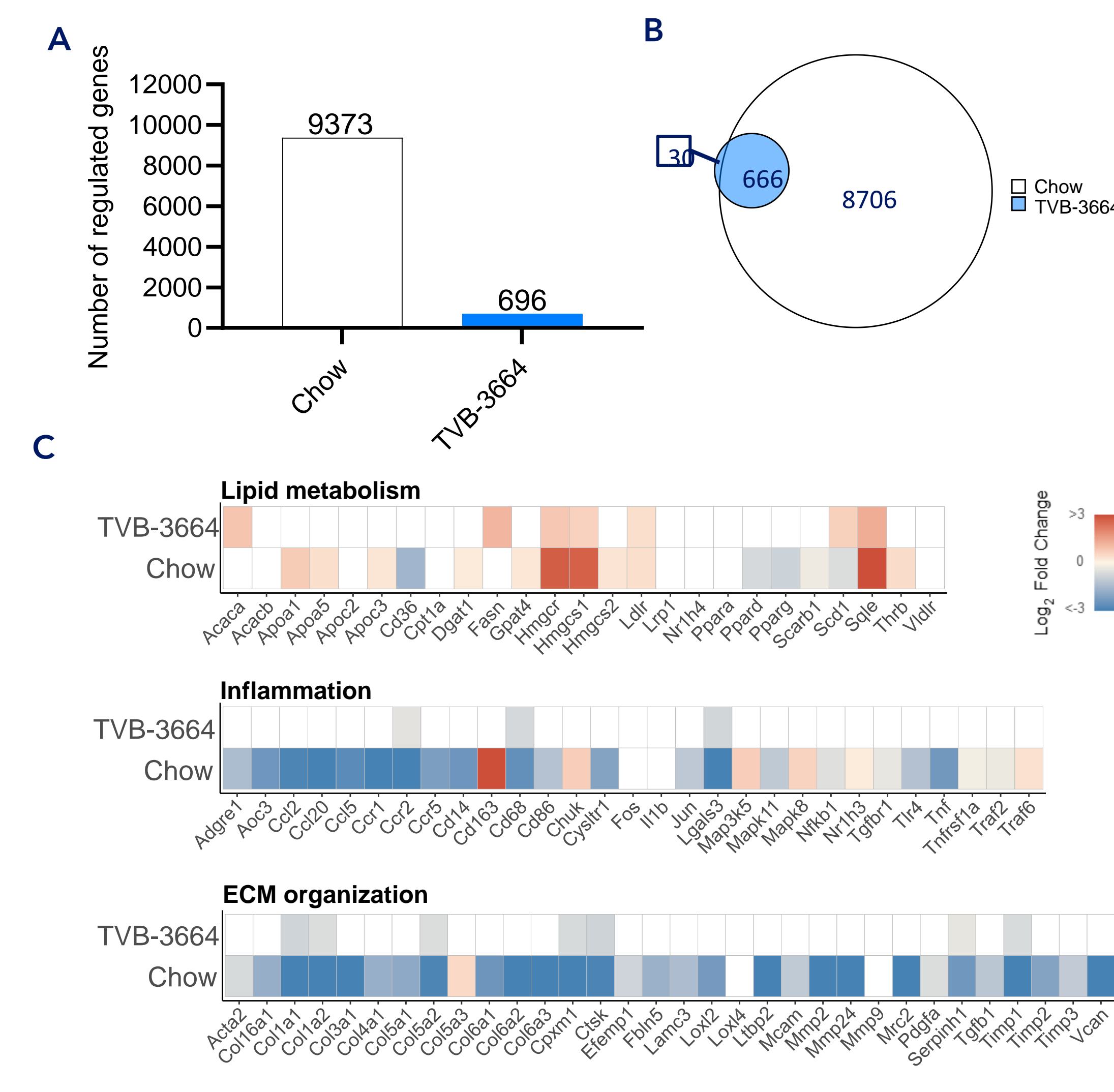


Figure 5. Hepatic transcriptome signatures following TVB-3664 therapy. (A) Total number of differentially expressed genes. (B) Venn diagram on overlapping gene expression signatures. (C) Regulation of gene expression markers of hepatic lipid metabolism, inflammation and extracellular matrix (ECM). Log<sub>2</sub> fold change is indicated for significantly regulated genes (p<sub>adj</sub> < 0.05 after correcting for multiple testing). Red and blue colours indicate up- and down-regulation, respectively, compared to Vehicle.

## Conclusion

- + TVB-3664 improves body weight and adiposity, while conserving lean tissue mass.
- + TVB-3664 improves hepatomegaly, plasma liver enzymes and cholesterol levels.
- + TVB-3664 improves NAFLD Activity Score, but not Fibrosis Stage.
- + Benefits on NAS are supported by quantitative histological markers for steatosis and inflammation.
- + TVB-3664 reduces quantitative histological marker for fibrogenesis.
- + TVB-3664 minimally regulates hepatic transcriptomic profile, predominantly improving gene expression markers of hepatic lipid metabolism and fibrogenesis.
- + Longer treatment intervention might be needed for demonstrating anti-fibrotic action for TVB-3664 in GAN DIO-MASH mouse model.