Hepatoprotective effects of a HSD17b13 inhibitor in the CDAA-HFD mouse model of advanced MASH with progressive fibrosis

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Background & Aim

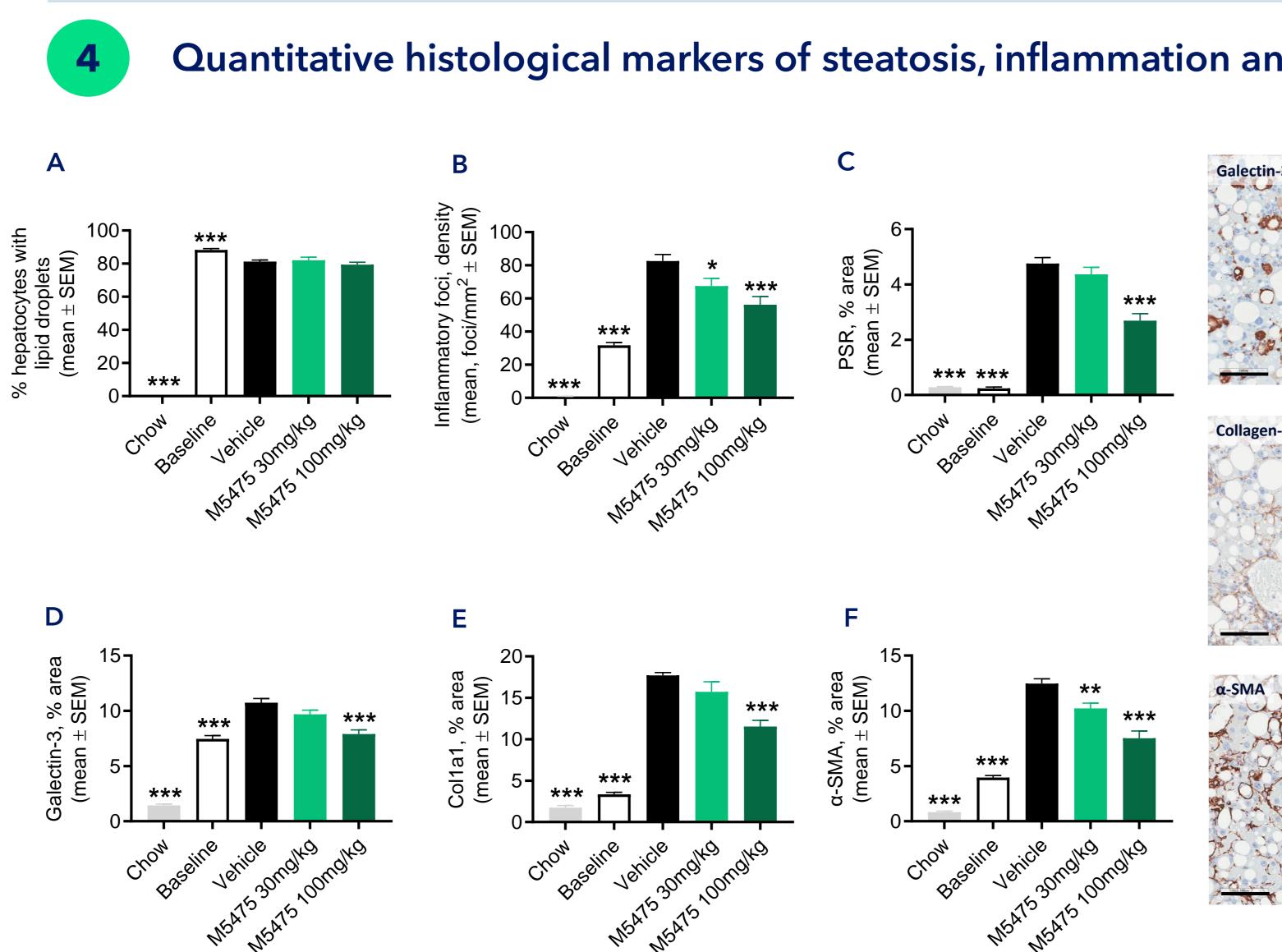
HSD17B13, a newly identified hepatocytespecific, lipid droplet-associated protein, has recently been reported to be closely associated with development and progression of MASLD/MASH. Furthermore, inhibition of HSD17B13 has shown to protect against liver fibrosis in preclinical models of MASH.

The present study aimed to evaluate effects of a HSD17b13 inhibitor following interventional therapy in the non-obese choline-deficient Lamino-acid defined high-fat diet (CDAA-HFD) mouse model of advanced MASH with progressive fibrosis.

Methods

C57BL/6JRj mice were fed chow or CDAA-HFD (45 kcal% fat, 0.1% methionine, 1% cholesterol, 28 kcal% fructose) for 3 weeks prior to treatment start (*i.e.* before onset of fibrosis). Animals were randomized into treatment groups based on body weight. A baseline group (n=12) was terminated at study start. CDAA-HFD fed mice (n=9-12 per group) were administered (PO) HSD17b13 inhibitor (M5475, 30 mg/kg or 100 mg/kg) or vehicle for 9 weeks. Chow-fed mice (n=8) served as normal controls. Terminal endpoints included plasma biomarkers, liver biochemistry, NAFLD Activity Score (NAS), fibrosis stage, quantitative liver histology.

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M5475	5	



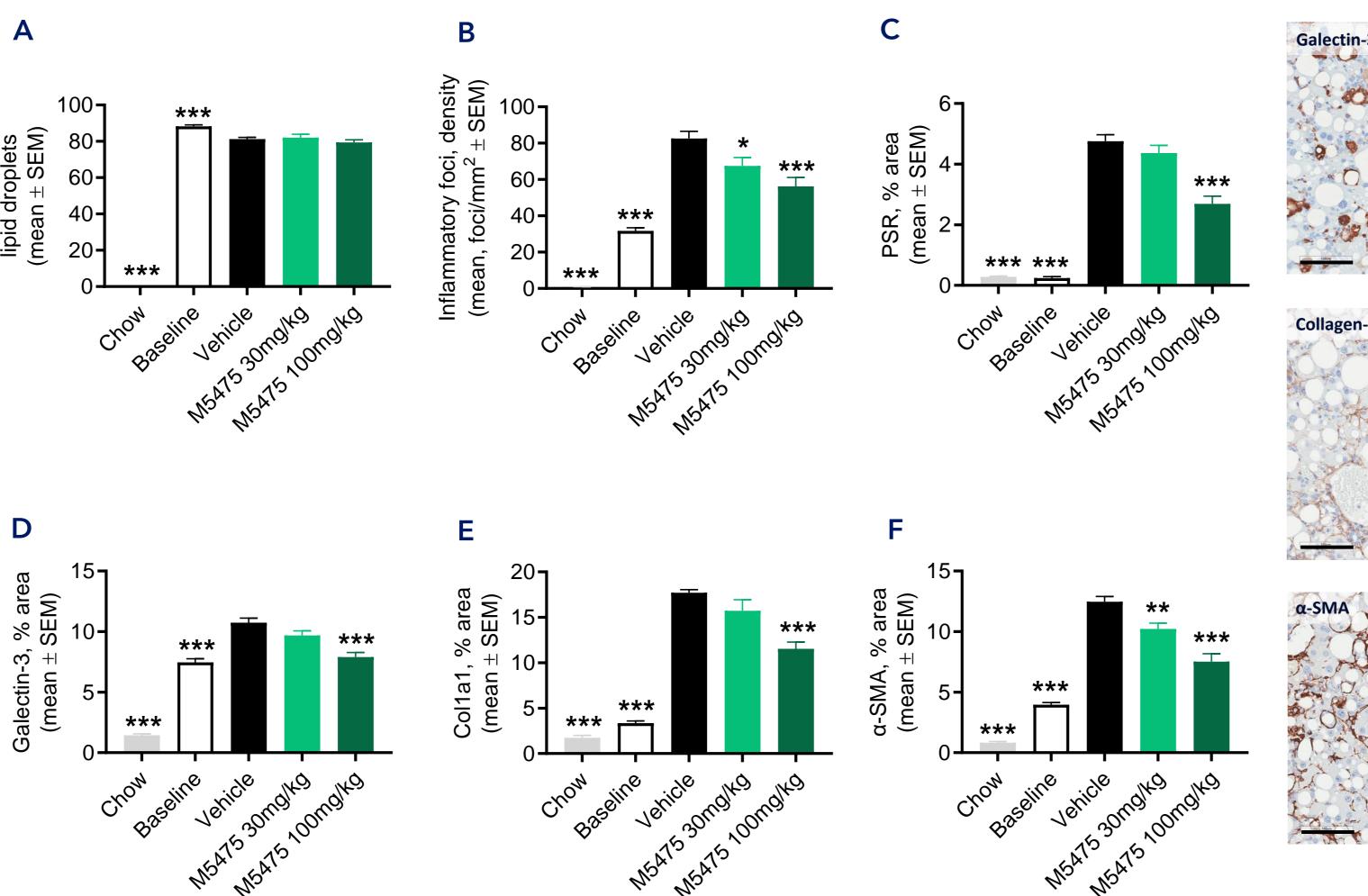
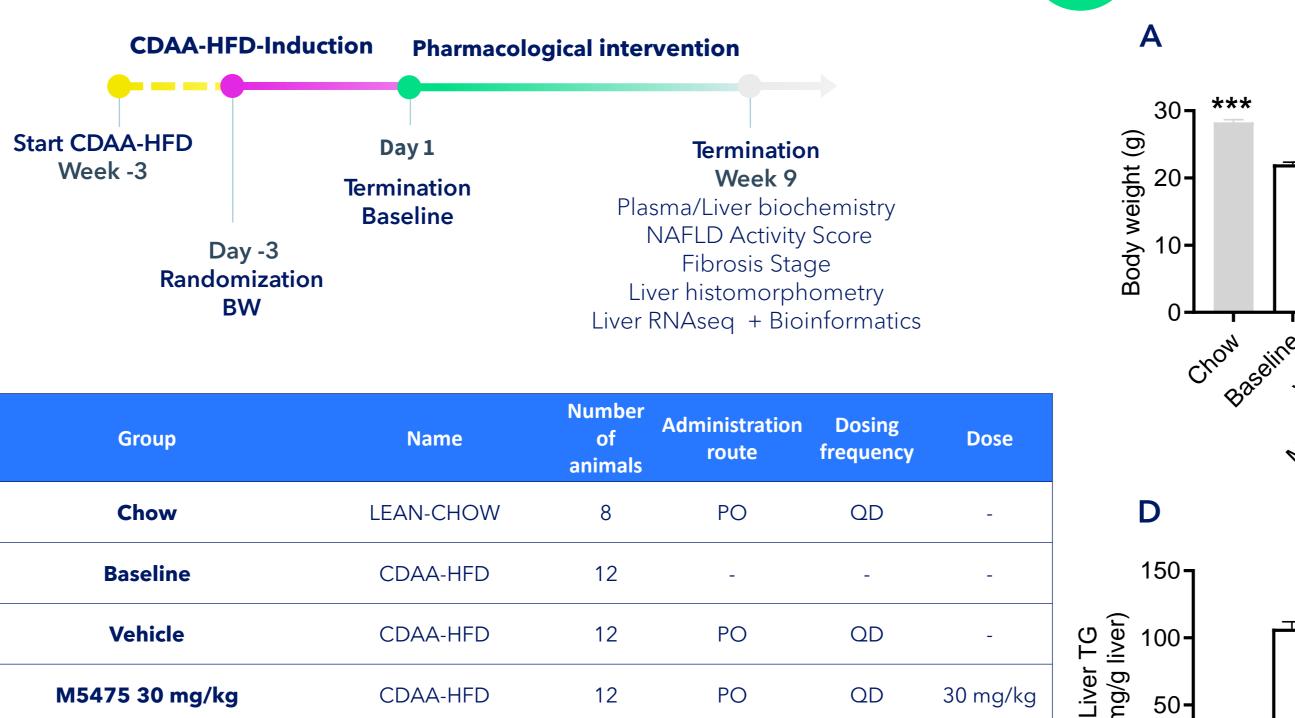
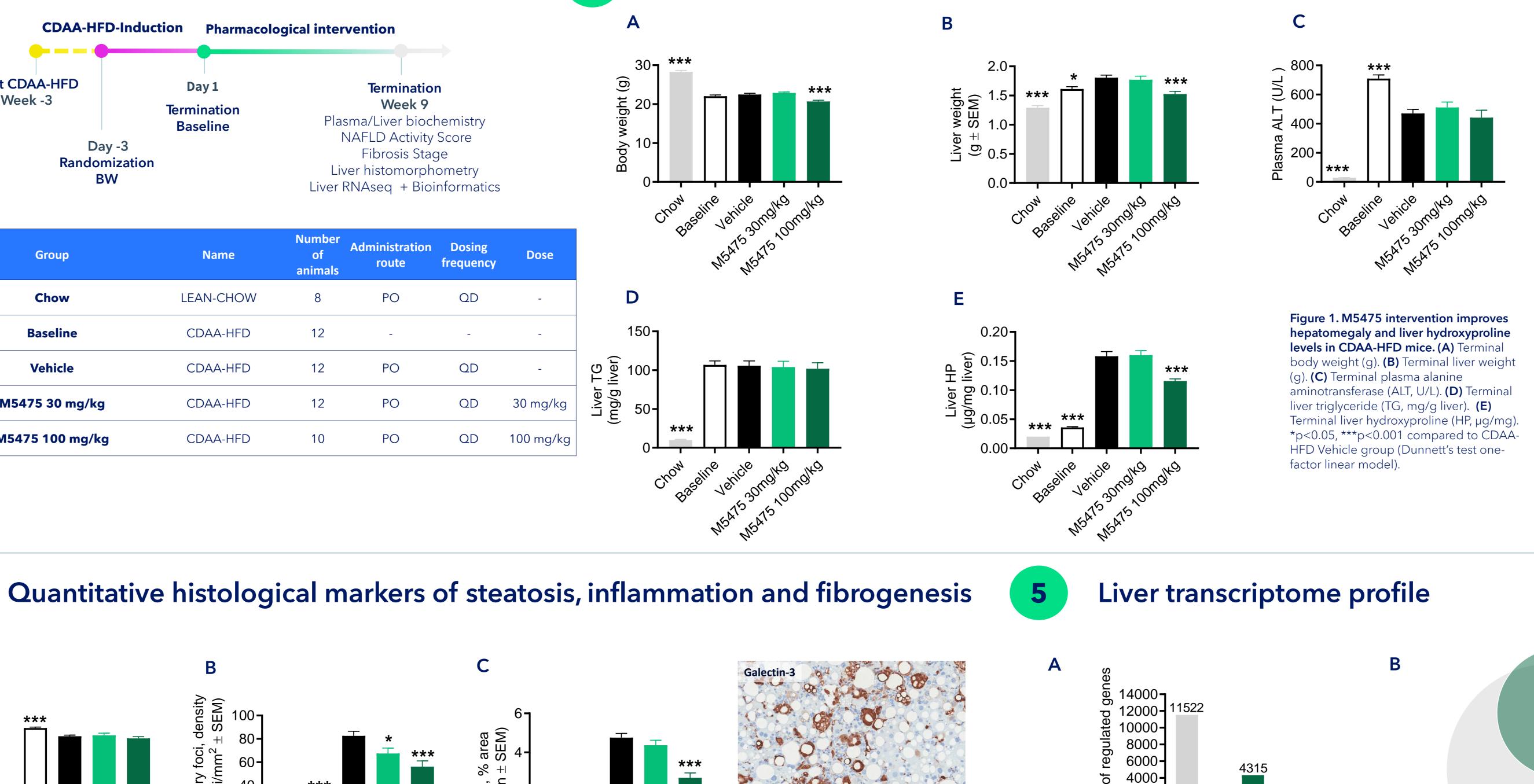


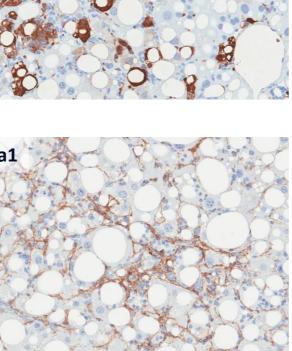
Figure 3. M5475 intervention improves quantitative histological markers of inflammation and fibrosis in CDAA-HFD mice. Histomorphometric assessments were performed by GHOST deep learning-based image analysis on scoring-associated variables and conventional IHC image analysis. (A) % hepatocytes with lipid droplets. (B) Number of inflammatory foci. (C) % area of PSR. (D) % area of galectin-3. (E) % area of collagen-1a1. (F) % area of alpha-smooth muscle actin (α-SMA). Mean ± SEM. *p<0.05, **p<0.001, ***p<0.001 compared to CDAA-HFD Vehicle group (Dunnett's test one-factor linear model). Right panels: Representative galectin-3, collagen 1a1 and α -SMA photomicrographs (scale bar, 100 μ m).

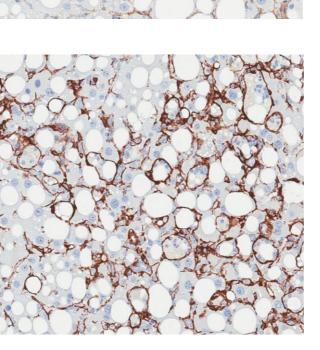
Study Outline





Metabolic and biochemical parameters





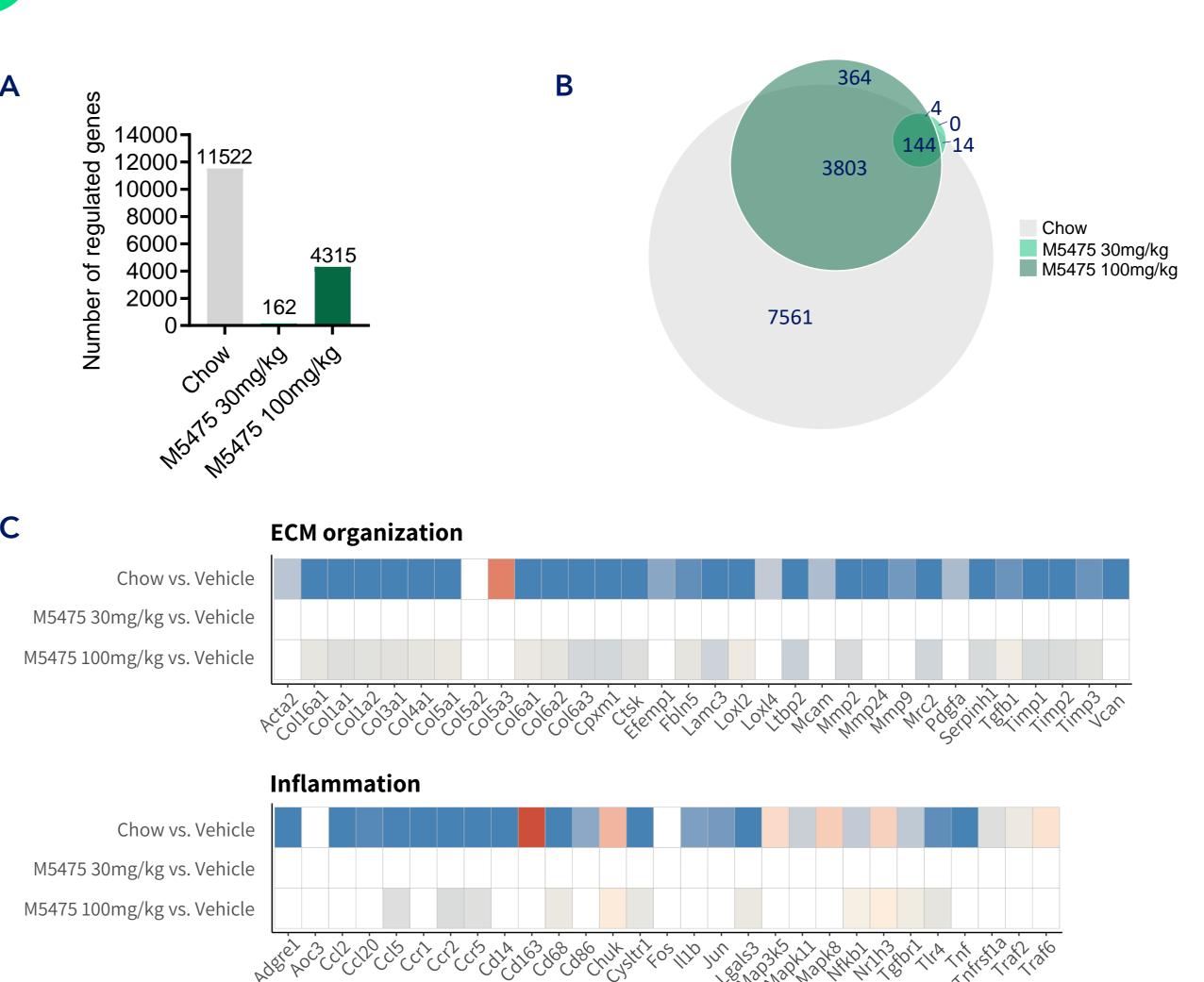


Figure 4. M5475 intervention marginally influences candidate gene expression markers of extracellular matrix (ECM) and inflammation in CDAA-HFD mice. (A) Total number of differentially expressed genes (DEGs) compared to CDAA-HFD Vehicle controls. (B) Total number of DEGs compared to CDAA-HFD Vehicle controls. (C) Regulation of candidate genes associated with ECM organization and inflammation (log2-fold change compared to CDAA-HFD vehicle group). Color gradients indicate significantly upregulated (red) and downregulated (blue) genes, respectively (p<0.05). White boxes indicate genes not significantly regulated (p>0.05).

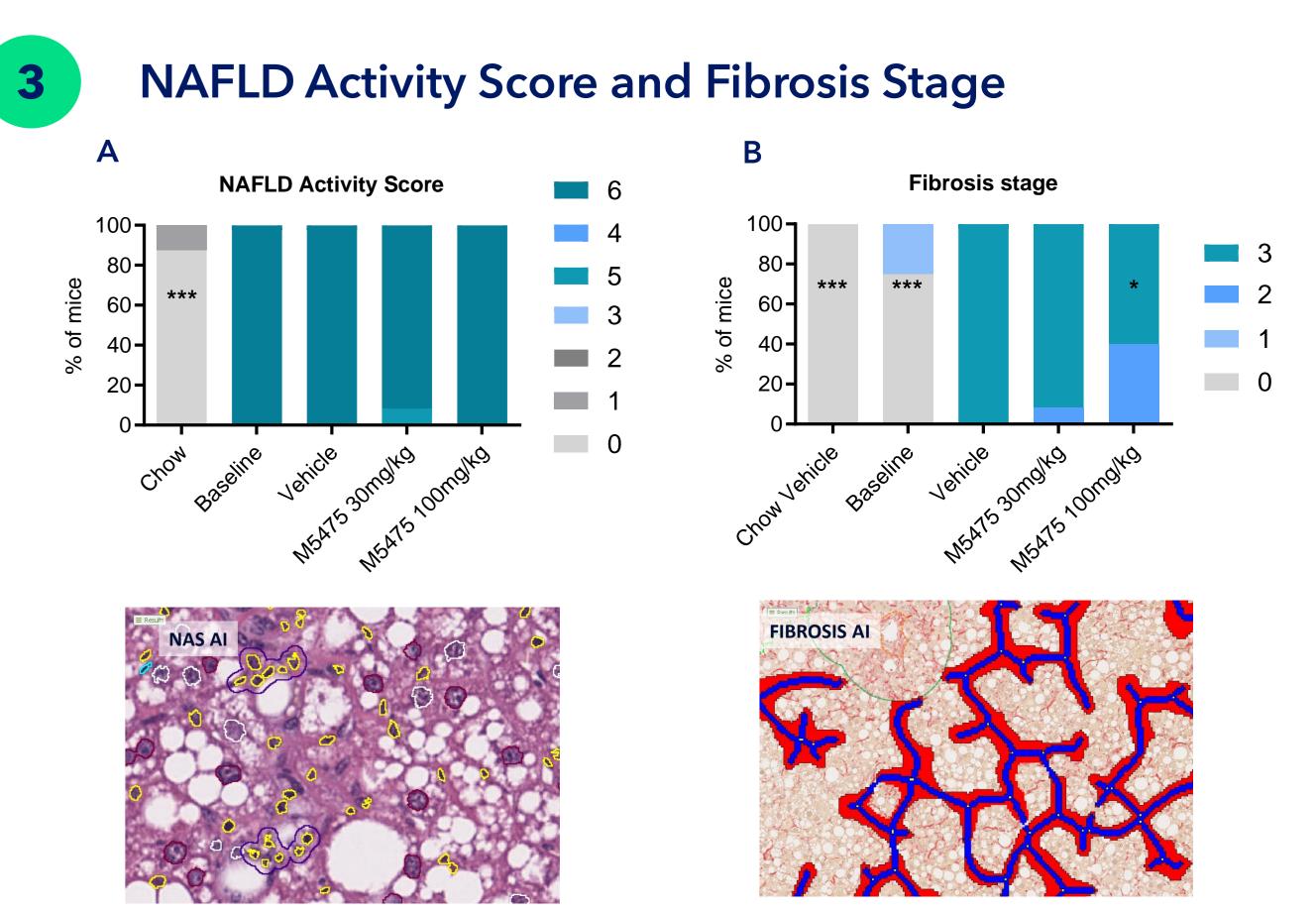


Figure 2. M5475 improves fibrosis scores in CDAA-HFD mice. Histopathological scores were determined by Gubra Histopathological Objective Scoring Technique (GHOST) deep learning-based image analysis. **(A)** NAFLD Activity Score (NAS). **(B)** Fibrosis Stage.). *p<0.05, ***p<0.001 compared to CDAA-HFD Vehicle group (one-sided Fisher's exact test with Bonferroni correction). Bottom panels: Representative HE and PSR photomicrographs used for GHOST evaluation.

Conclusion

Outcomes of HSD17b13 inhibitor (M5475) therapy in CDAA-HFD mice:

- Improved hepatomegaly and liver hydroxyproline levels with marginal body weight loss
- No effect on NAFLD Activity Score
- Reduced fibrosis stage
- Reduced quantitative histological markers of inflammation and fibrosis
- Marginal effects on candidate genes involved in ECM remodelling and immune system

In conclusion, M5475 elicits anti-inflammatory action and prevents progression of liver fibrosis in the non-obese CDAA-HFD mouse model of MASH.



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