

Additive hepatoprotective effects of semaglutide and resmetirom combination therapy in the GAN diet-induced obese and biopsy-confirmed mouse model of MASH

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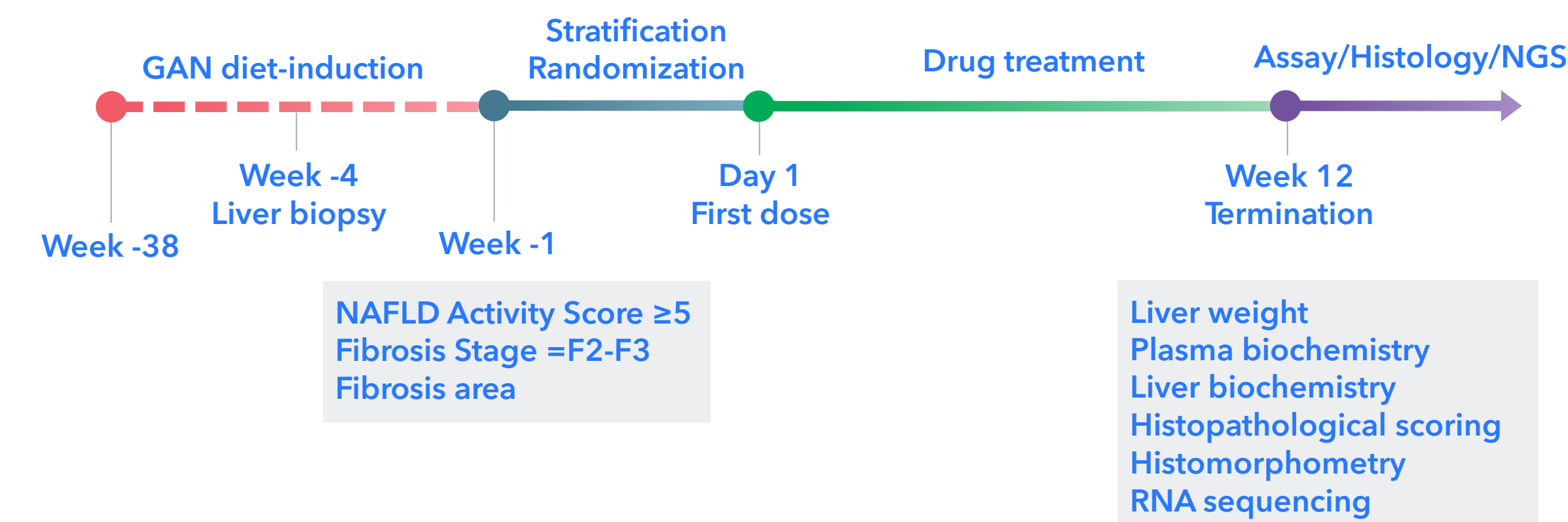
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Background & Aim

Resmetirom (THR- β receptor agonist) has recently been approved for the treatment of metabolic dysfunction-associated steatohepatitis (MASH) and semaglutide (GLP-1R agonist) is in late-stage clinical development for treatment of MASH. The present study aimed to characterize metabolic, biochemical, histological and transcriptomic outcomes of resmetirom and semaglutide combination treatment in the translational GAN diet-induced obese (DIO) and biopsy-confirmed mouse model of MASH with liver fibrosis.

1 Study outline



Group	Animal	Gender	Number of animals	Treatment	Administration route	Dosing frequency	Dosing concentration
1	LEAN-CHOW	Male	10	Vehicle	SC	QD	NA
2	DIO-MASH	Male	18	Vehicle	SC	QD	NA
3	DIO-MASH	Male	18	Semaglutide	SC	QD	30 nmol/kg
4	DIO-MASH	Male	18	Resmetirom	PO	QD	3 mg/kg
5	DIO-MASH	Male	17	Semaglutide + Resmetirom	SC + PO	QD + QD	30 nmol/kg + 3 mg/kg

Figure 1. Study outline. PO; per oral, SC; subcutaneous, QD; once daily, NA; not applicable, GAN; Gubra Amylin NASH, NGS; next generation sequencing.

2 Metabolic and biochemical parameters

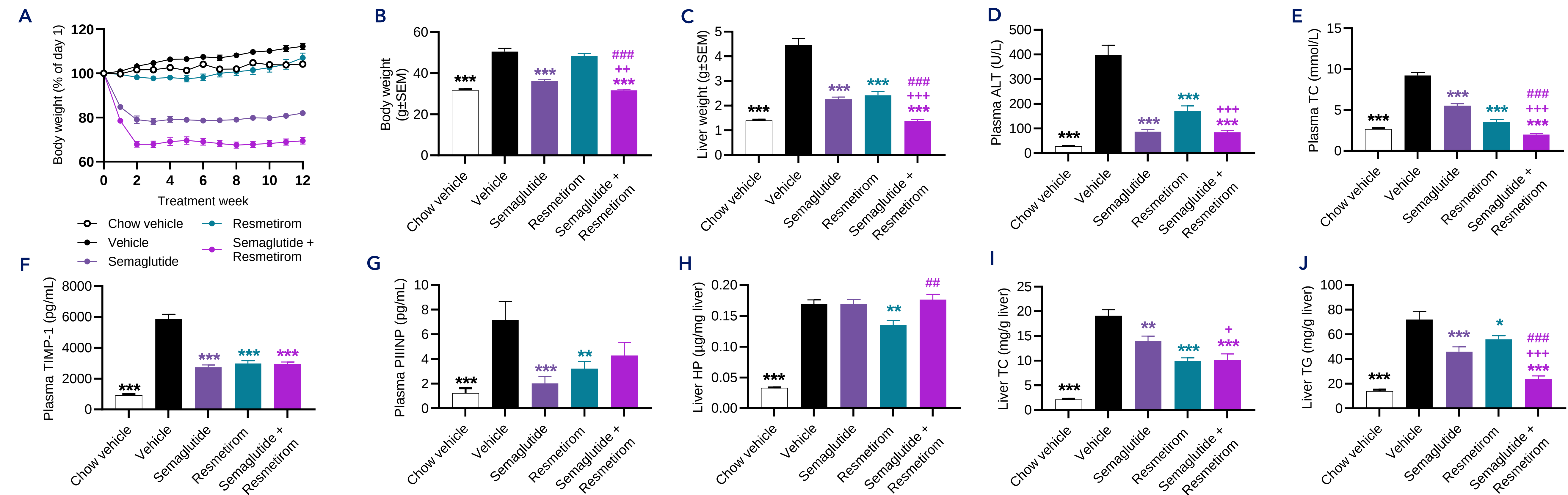


Figure 2. Semaglutide + Resmetirom combo treatment reduced weight loss, liver weight and biochemical parameters superior to mono treatments. (A) Relative body weight during study period. (B) Terminal body weight. (C) Terminal liver weight. (D) Terminal plasma alanine aminotransferase (ALT). (E) Terminal plasma total cholesterol. (F) Plasma tissue inhibitor of metalloproteinases (TIMP-1). (G) Plasma Type III Procollagen Peptide (PIINP). (H) Liver hydroxyproline (HP). (I) Liver triglycerides (TG). (J) Terminal liver total cholesterol (TC). * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ compared to Vehicle. #/ $p < 0.05$, ##/ $p < 0.01$, ###/ $p < 0.001$ compared to resmetirom (#) or semaglutide (+) (Dunnett's test one-factor linear model).

3 NAFLD Activity Score and Fibrosis Stage

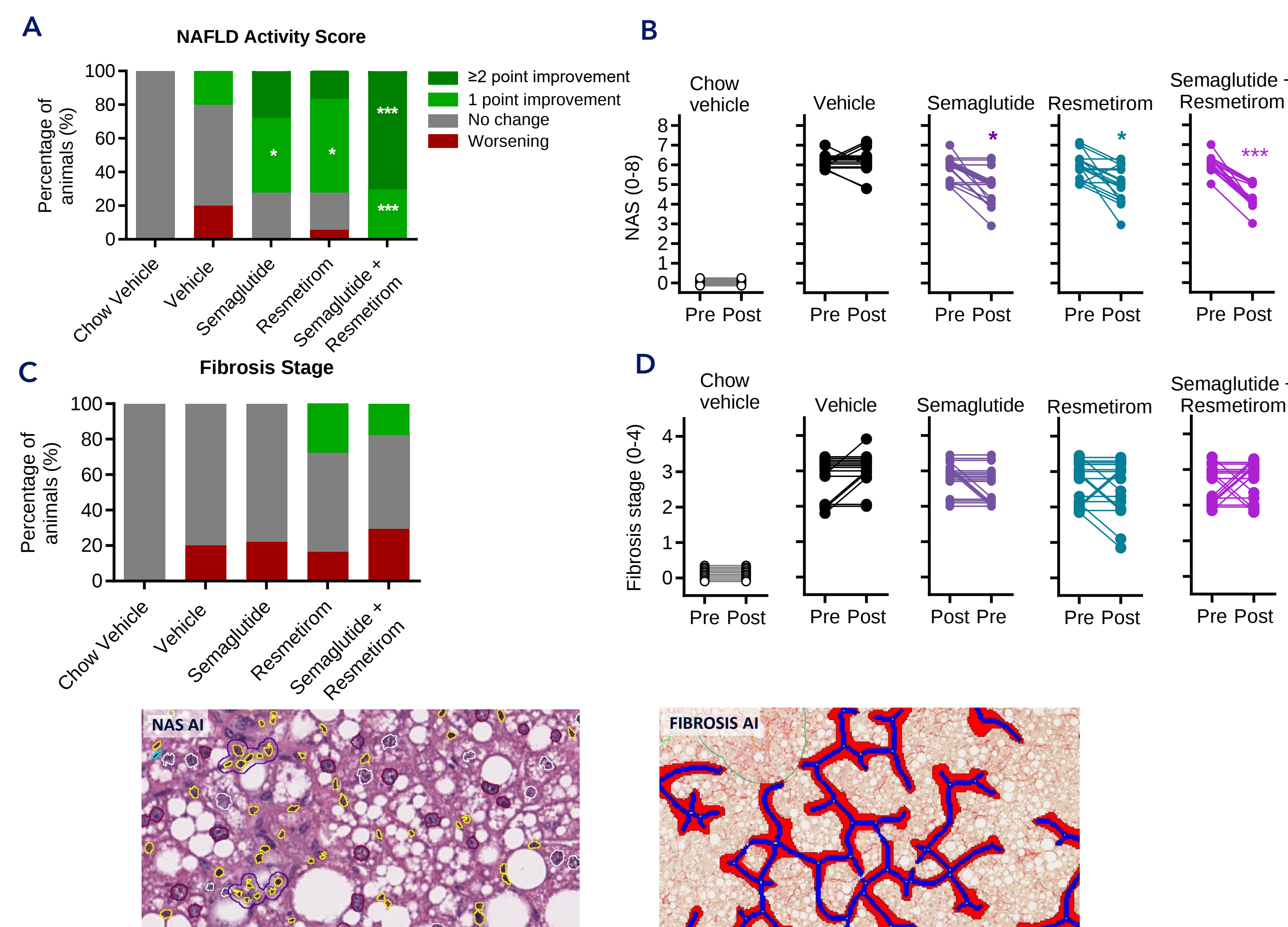


Figure 3. Semaglutide + Resmetirom combo treatment improved NAFLD Activity Score superior to mono treatments. Histopathological scores were determined by Gubra Histopathological Objective Scoring Technique (GHOST) deep learning-based image analysis. (A) NAFLD Activity Score (NAS). (B) Individual pre-post NAS. (C) Fibrosis stage. (D) Individual pre-post fibrosis stage. * $p < 0.05$, *** $p < 0.001$ compared to vehicle. (One-sided Fisher's exact test).

4 Histological markers of steatosis, inflammation and fibrosis

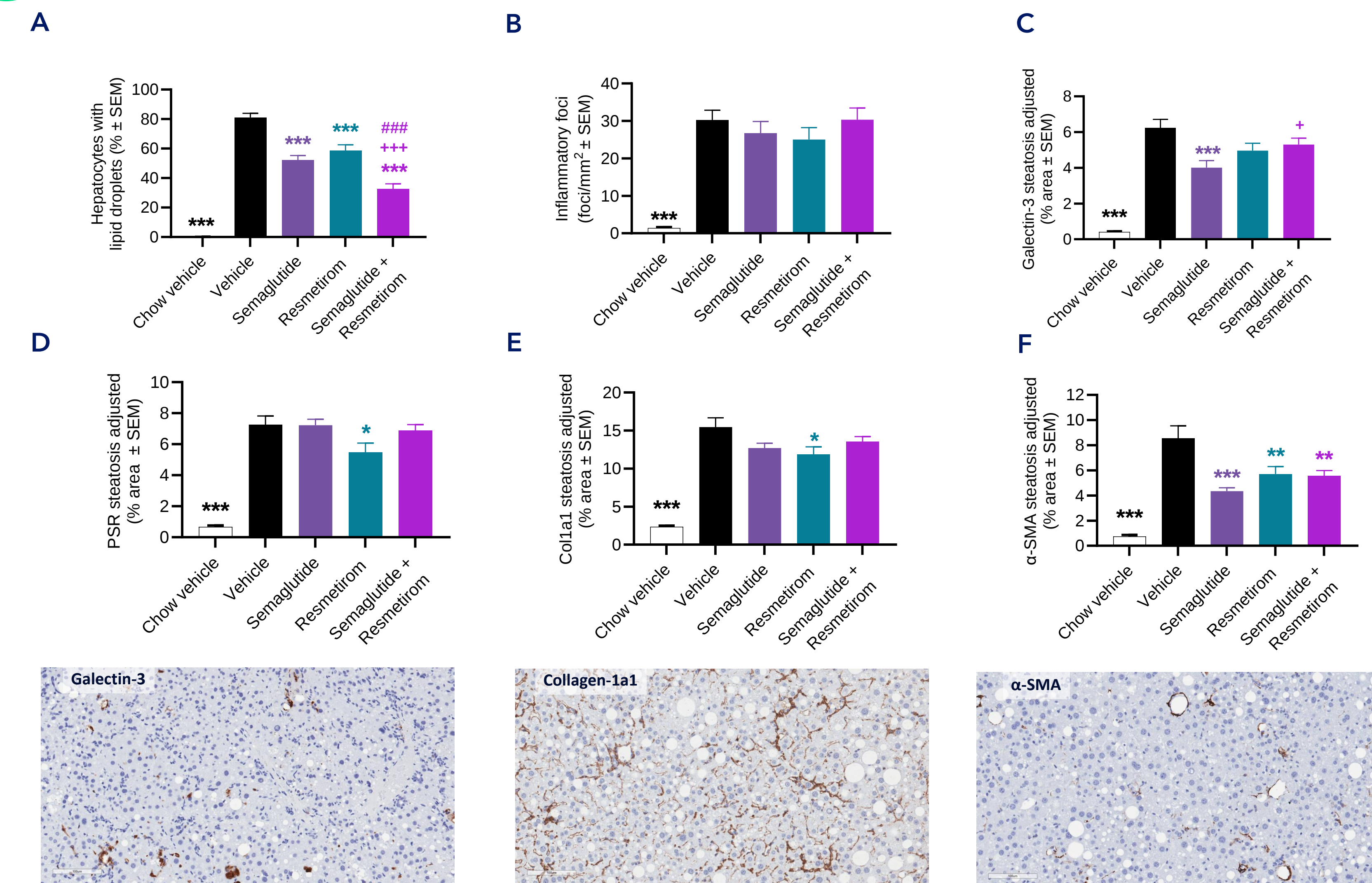


Figure 4. Semaglutide + Resmetirom combo treatment improved quantitative histological marker of steatosis superior to mono treatments. Histomorphometric assessments were performed by GHOST deep learning-based image analysis on scoring-associated variables (panels A-F) and conventional IHC image analysis (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 α -smooth muscle actin (α -SMA, marker of stellate cell activation, steatosis adjusted). Mean $\pm$ SEM. * $p < 0.05$, *** $p < 0.001$ compared to vehicle. #/ $p < 0.05$, ##/ $p < 0.01$, ###/ $p < 0.001$ compared to resmetirom (#) or semaglutide (+) (Dunnett's test one-factor linear model). Bottom panels: Representative photomicrographs of galectin-3, α -SMA and Col1a1 (scale bar, 100 μ m).

5 Hepatic transcriptomic profile

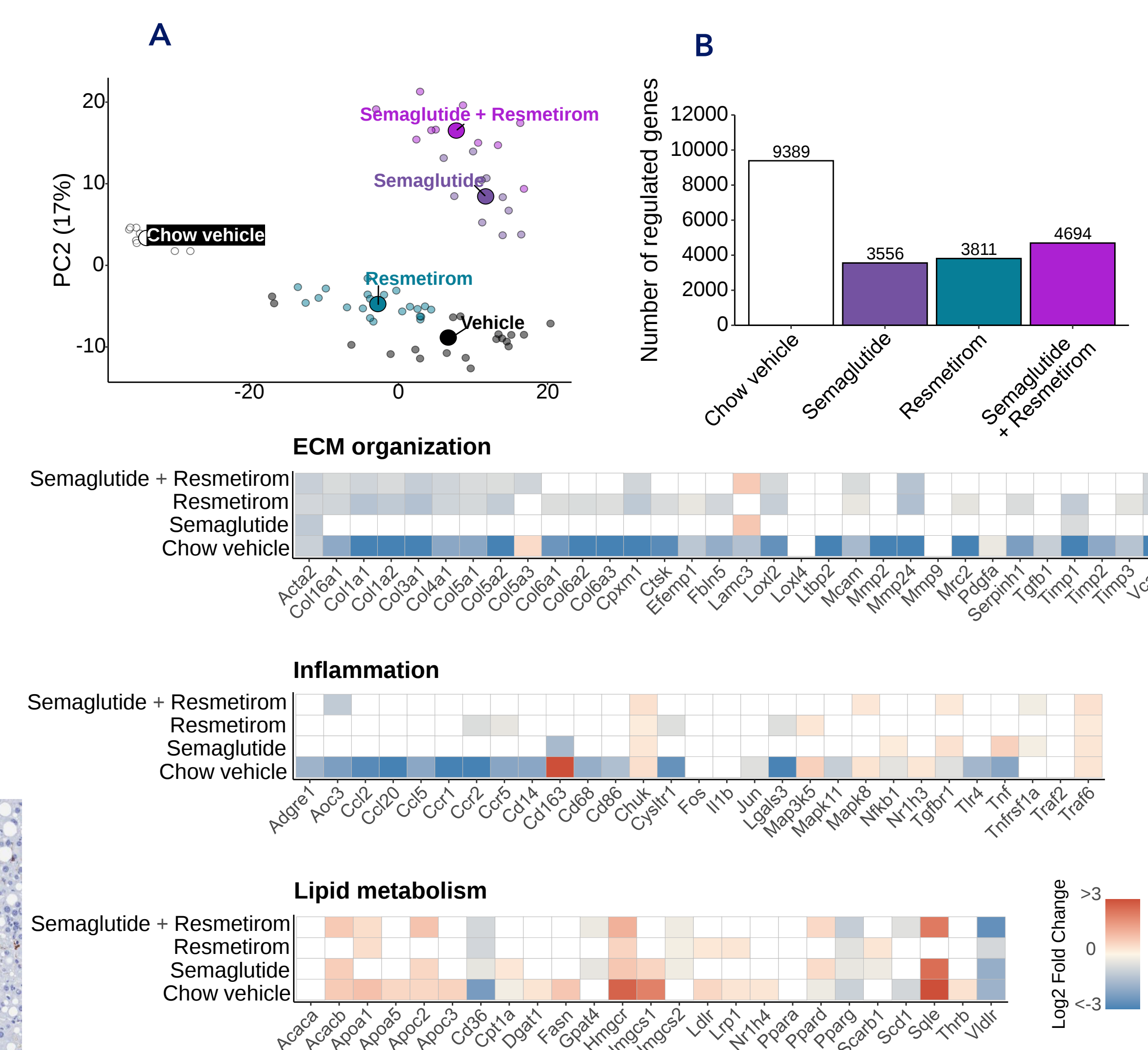


Figure 5. Semaglutide + Resmetirom combo treatment increased number of differentially expressed genes. (A) Principal component analysis (PCA) of samples based on top 500 most variable gene expression levels. (B) Differentially expressed genes. (C) Regulation of hepatic extracellular matrix (ECM), inflammation and lipid metabolism candidate genes (log2-fold change compared to corresponding DIO-NASH vehicle control mice). Blue and red colour gradients indicate significantly ($p < 0.05$) down-regulated and up-regulated gene expression, respectively. White boxes indicate genes not significantly regulated ($p > 0.05$).

Conclusion

- + Semaglutide + Resmetirom combo treatment promote superior weight loss to mono treatments.
- + Semaglutide + Resmetirom combo treatment improved hepatomegaly and plasma/liver lipids superior to mono treatments.
- + Semaglutide + Resmetirom combo treatment improved NAS (2-point) superior to mono treatments.
- + Semaglutide + Resmetirom combo treatment reduces steatosis superior to mono treatments.
- + Only resmetirom mono treatment reduces markers of fibrosis.
- + Semaglutide + Resmetirom combo treatment amplified hepatic transcriptomic profile.

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