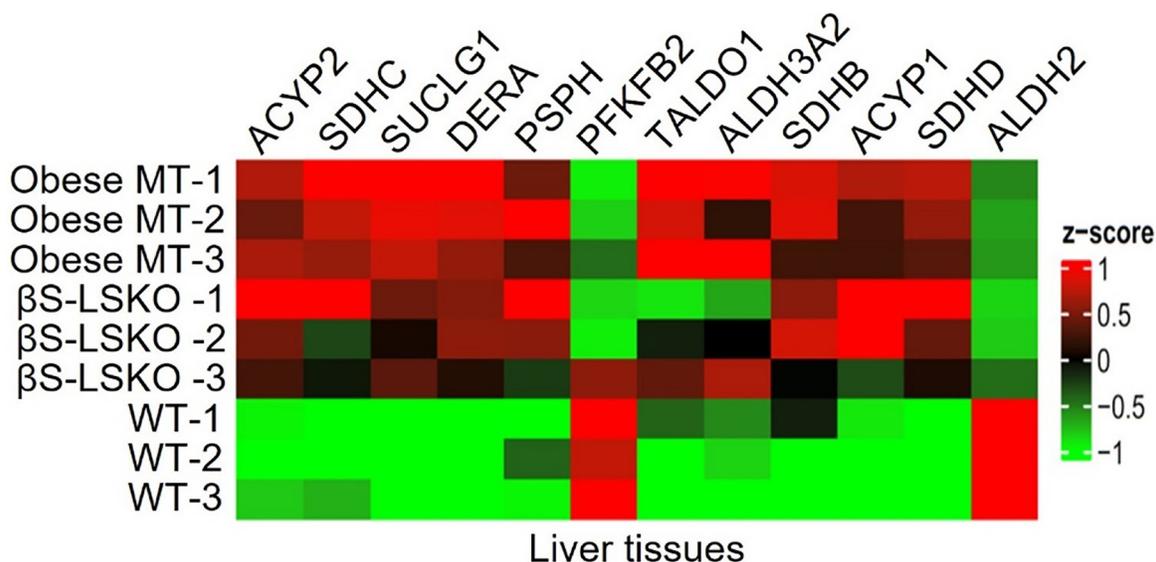


Mechanistically based blood proteomic markers in the TGF- β pathway stratify risk of hepatocellular cancer in patients with cirrhosis

SUPPLEMENTARY MATERIALS



Supplementary Figure 1: Altered glucose metabolic processes in the liver tissues from obese mice. Heatmap indicate relative expression level of glucose metabolic genes in liver tissue from mouse via RNA-seq analysis in the mentioned groups. Abbreviations: WT: Wild type; Obese MT: Obese mouse tissues; β S-LSKO: liver-specific knockout of *Sptbn1*. ($p < 0.05$).