Investigating heat shock protein dysregulation and immune cell dynamics in non-alcoholic steatohepatitis by integrated multi-RNA seq analysis

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Supporting Information

Cell type	Normal Diet Liver		HFD Diet Induced NASH Liver	
Monocyte	400	0.272	2135	0.419
Macrophage/Kupffer cell	843	0.574	1469	0.288
Mast cell	187	0.127	1087	0.213
Neutrophil	38	0.026	402	0.079

Table S1. Proportions of the CD16+ myeloid cells in normal and HFD diet mouse liver

Table S2. Proportions of monocyte-derived macrophages and Kupffer cell macrophages in normal and HFD mouse liver

Cell type	Normal Liver		HFD NASH Liver	
Monocyte-derived macrophage	85	0.101	1464	0.997
Kupffer cell	758	0.899	5	0.003



Figure S1. Gene expression patterns of the monocytes in normal and HFD-induced NASH liver. (A-D) The expression of monocyte/macrophage markers Itgam, Adgre1, CD68, and Csf1r in monocytes from normal (left) and HFD-induced NASH (right) liver. (E) The expression of lipid receptor Trem2 in the monocytes of normal (left) and HFD-induced NASH (right) liver. (F-G) The expression of monocyte/macrophage recruitment chemokine receptors Ccr2, Cx3cr1 in the monocytes of normal (left) and HFD-induced NASH (right) liver. (H) The expression of proinflammatory cytokine II1b in the monocytes of normal (left) and HFD- induced NASH (right) liver.



Figure S2. Gene expression patterns of Kupffer cells and monocyte-derived macrophages in normal and HFD-induced NASH livers. (A-C) The expression of Kupffer cell markers of Clec4f, Tim4, Vsig4 in Kupffer cells and MDMs of normal (left) and HFD-induced NASH (right) liver. (D-G) The expression of monocyte/macrophage markers Itgam, Adgre1, Cd68, and in Kupffer cells and MDMs of normal (left) and HFD-induced NASH (right) liver. (H-I) The expression of lipid receptors Trem2, Cd9 in Kupffer cells and MDMs of normal (left) and HFD-induced NASH (right) liver. (J-K) The expression of monocyte/macrophage recruitment chemokine receptors Ccr2, Cx3cr1 in Kupffer cells and MDMs of normal (left) and HFD-induced NASH (right) liver. (L) The expression of proinflammatory cytokine II1b in Kupffer cells and MDMs of normal (left) and HFD-induced NASH (right) liver.



Figure S3. Dysregulation of HSPs in the MDMs of HFD-induced NASH liver compared with normal livers. The red underlines mark the significantly upregulated HSPs in the monocyte of NASH liver. The remaining are significantly downregulated HSPs.