

Figure S1 Classification of infiltrating immune cells between fibrosis and non-fibrosis tissues by PCA analysis.



Figure S2 PPI network. PPI, protein-protein interaction.



Figure S3 Top 30 genes in PPI network node count. PPI, proteinprotein interaction.



Figure S4 Correlation of M1 and M2 macrophages with PTPRC (A,B), IFIT3 (C,D), MX1 (E,F), and OAS2 (G,H), respectively.

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