

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

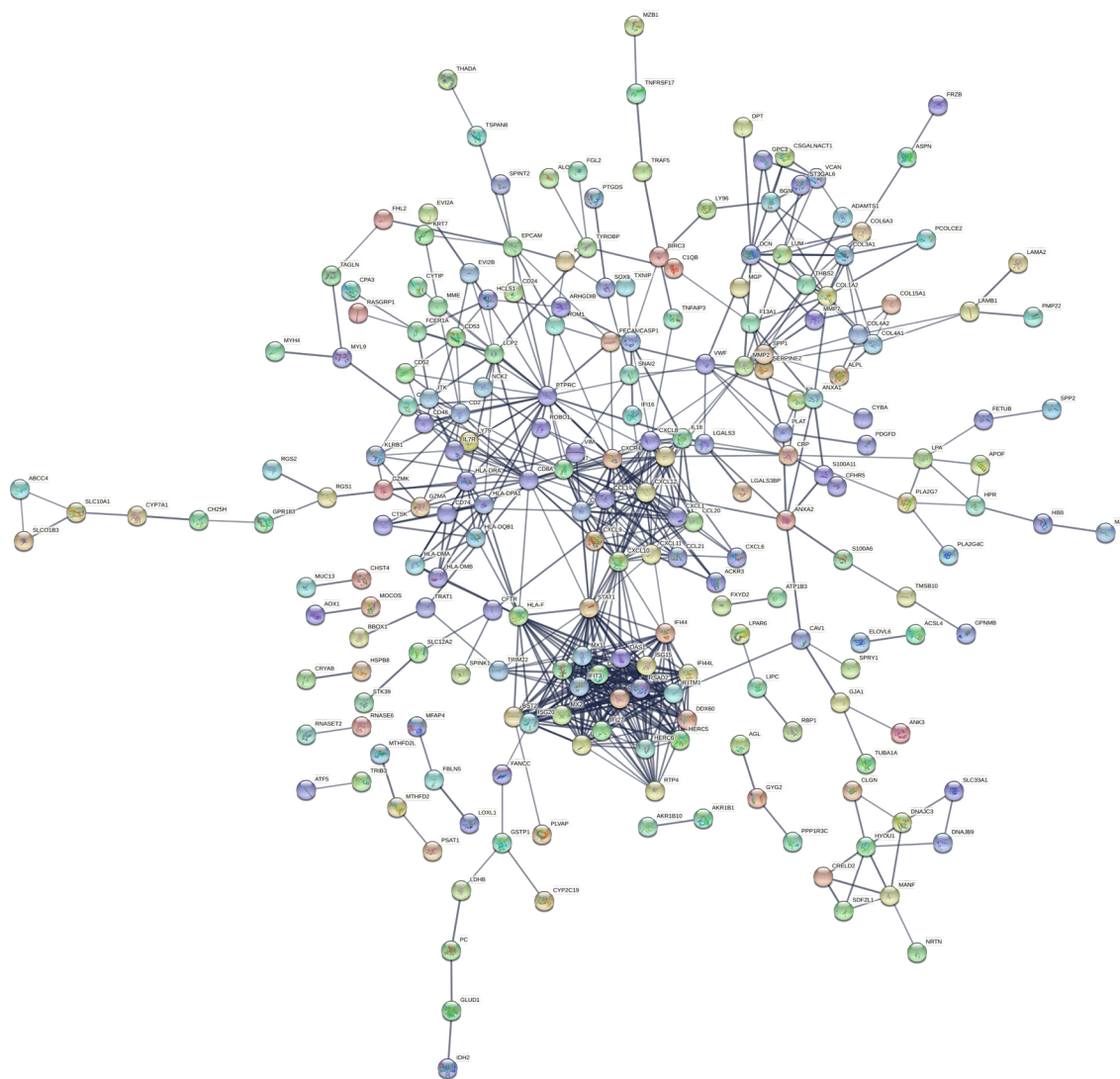


Figure S2 PPI network. PPI, protein-protein interaction.

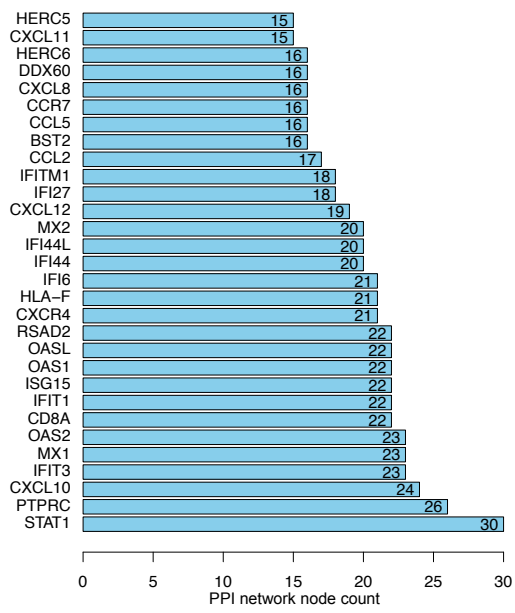


Figure S3 Top 30 genes in PPI network node count. PPI, protein-protein interaction.

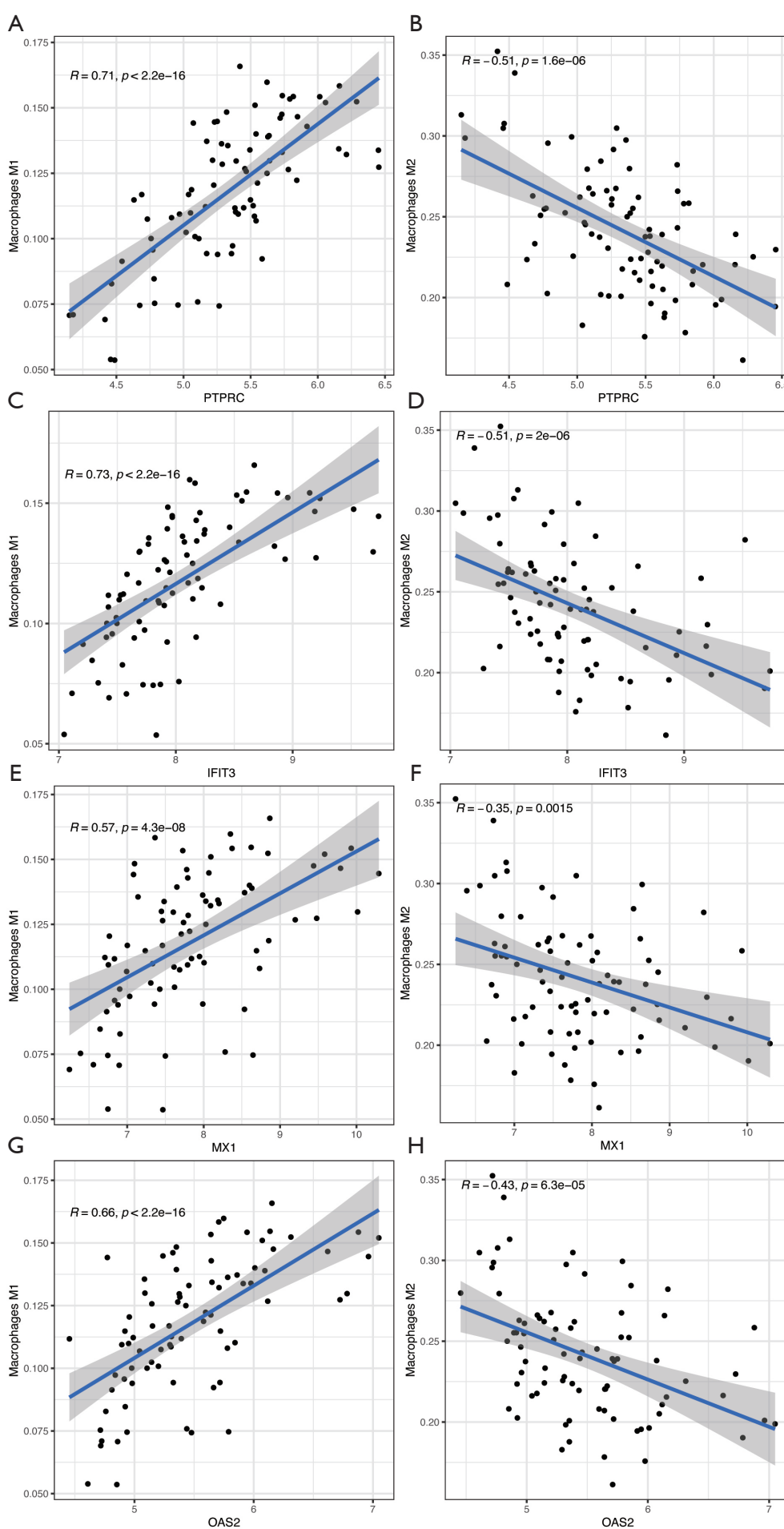


Figure S4 Correlation of M1 and M2 macrophages with PTPRC (A,B), IFIT3 (C,D), MX1 (E,F), and OAS2 (G,H), respectively.