

SUPPLEMENTARY FIG. S7. Expression levels of ROR α , SOD2, and GPx1 in human NASH patients: Database-based gene expression analysis. The public datasets were obtained from GEO site at the NCBI gene expression (www.ncbi.nlm.nih.gov/geo). The data processed as median-normalized signal intensity value. Significances were analyzed by Mann–Whitney U test, and the positive correlation coefficients (r) were calculated by Pearson correlation test.