Supplementary Fig. 4

**Fig. 4.** **RNA sequencing analysis of ATOX1-silencing and overexpressing Huh7 cells.** (A and B) The Volcano map of the RNA sequencing analysis of ATOX1-silencing and overexpressing Huh7 cells. Red dots show signiﬁcantly up-regulated genes and green dots show signiﬁcantly down-regulated genes. (C) The enriched pathways through KEGG analysis in the ATOX1 overexpression group. (D) GSEA analysis of differentially expressed genes derived from RNA sequencing data of ATOX1-silencing Huh7 cells. GSEA, Gene Set Enrichment Analysis; DEGs, differentially expressed genes; KEGG, Kyoto Encyclopedia of Genes and Genomes; OE-ATOX1, ATOX1 overexpression; si-NC, small interfering RNA targeting negative control; si-ATOX1, small interfering RNA targeting ATOX1.