Figure S1 Weighted co-expression network construction and identification
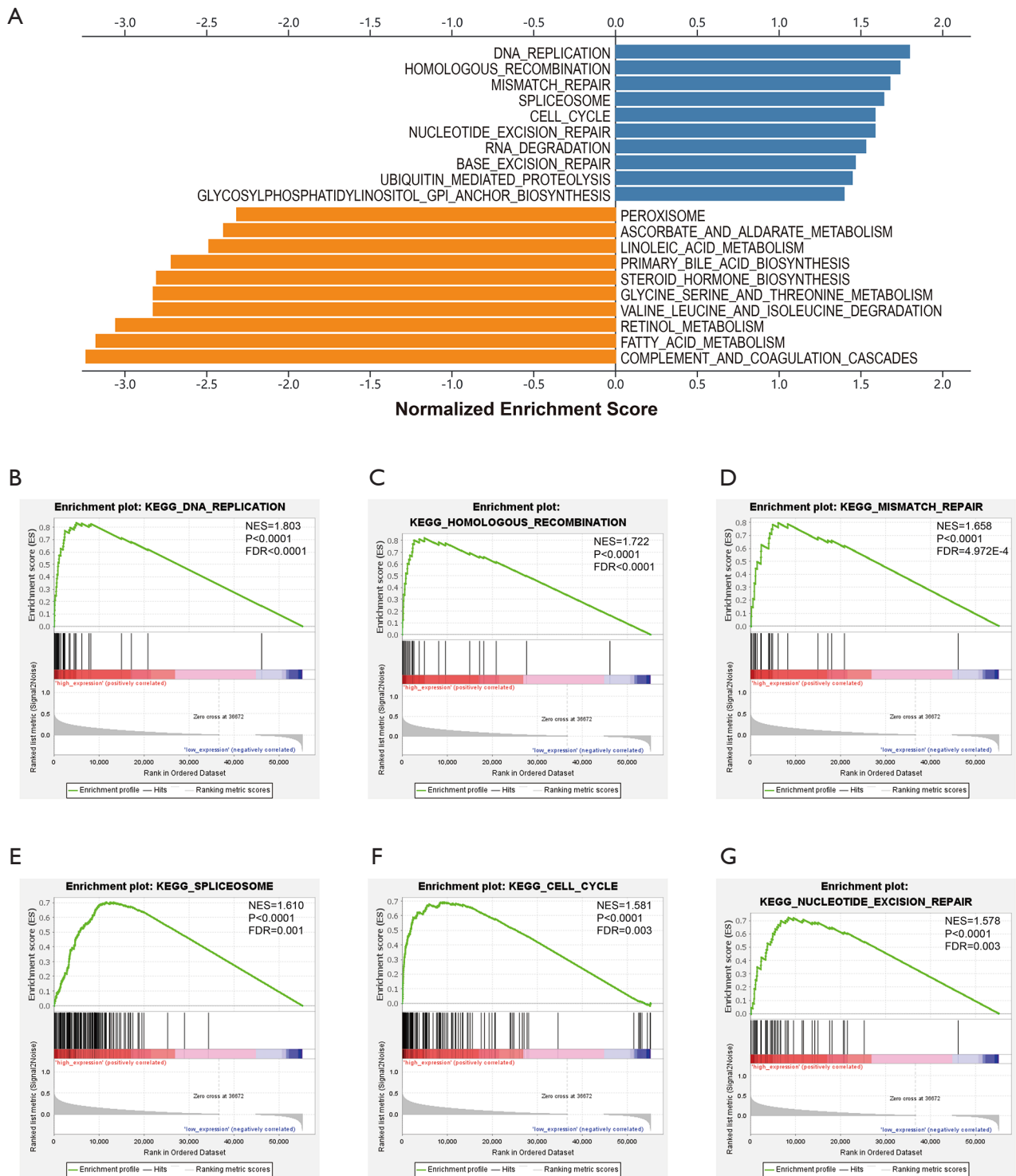
Figure S2 Gene set enrichment analysis results enriched in phenotype of high NIFK-AS1 expression group.