

Figure S2

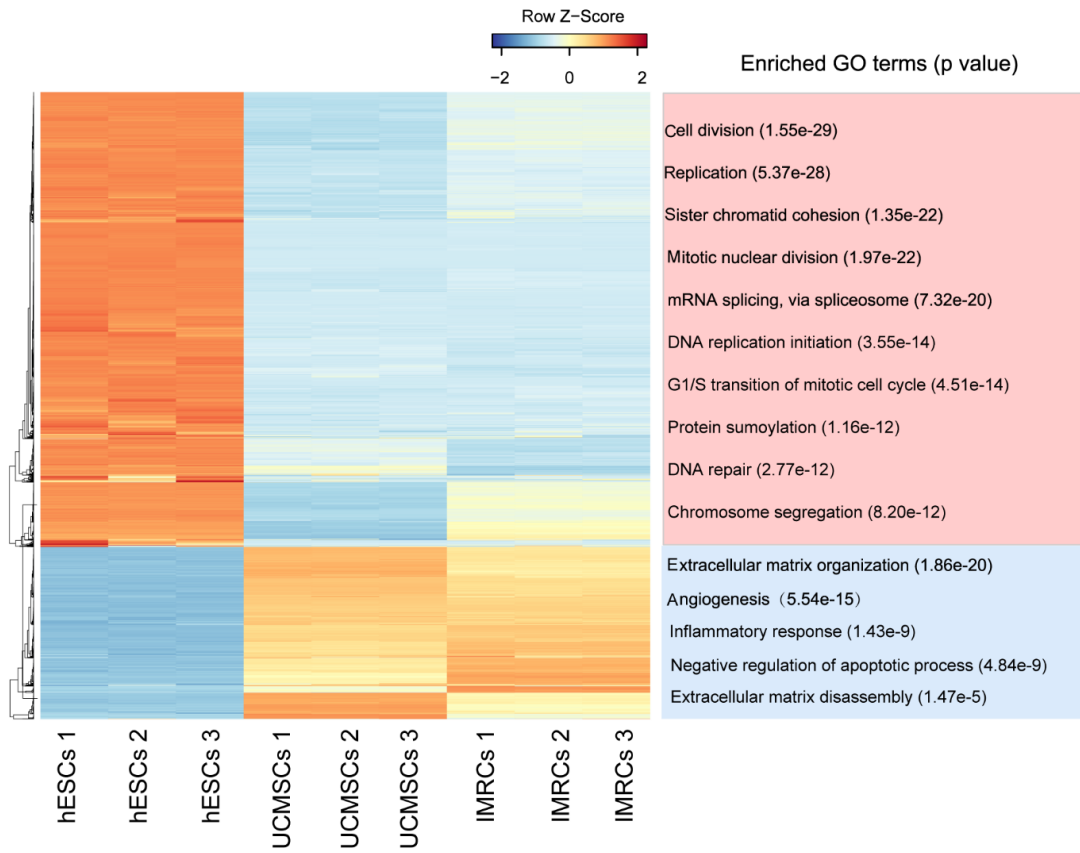


Fig. S2 The differentially expressed genes among hESCs and UCMSCs, IMRCs. In total, 4,730 differentially expressed genes were found in hESCs compared to UCMSCs and IMRCs. The enriched Gene Ontology (GO) terms and corresponding p values are shown.