

Figure S5

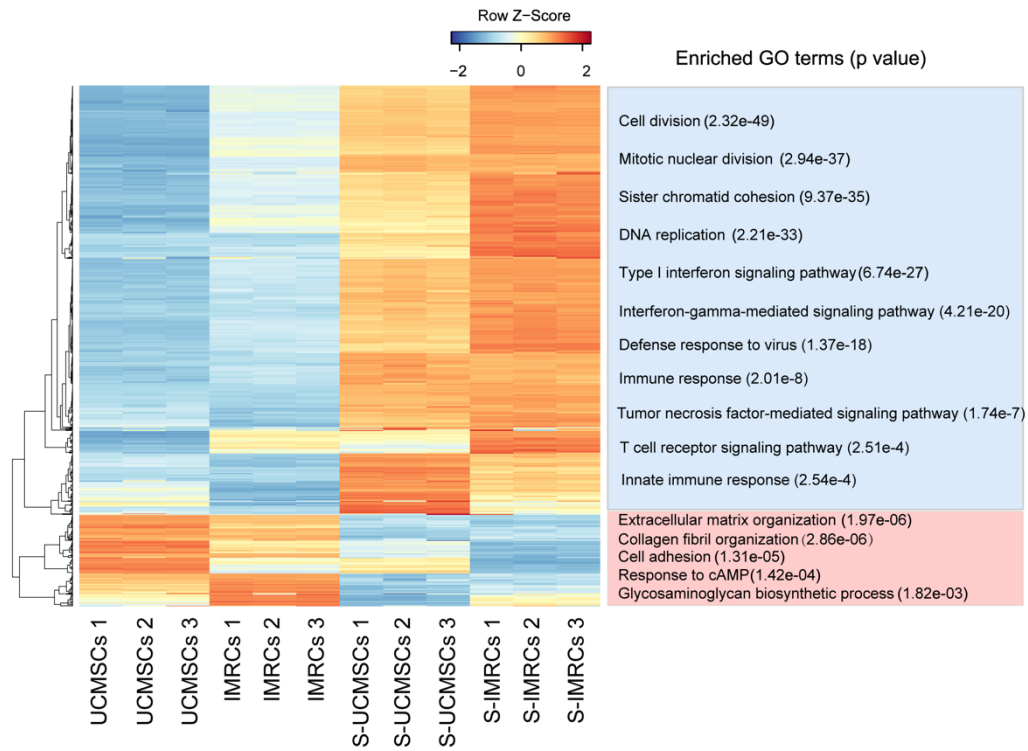


Fig. S5 The differentially expressed genes in UCMSCs and IMRCs, before and after IFN- γ stimulation. In total, 763 differentially expressed genes were found in UCMSCs and IMRCs before and after IFN- γ stimulation. The enriched Gene Ontology (GO) terms and corresponding p values are shown.