

siGABPA_VS_siNC_kegg_up_down

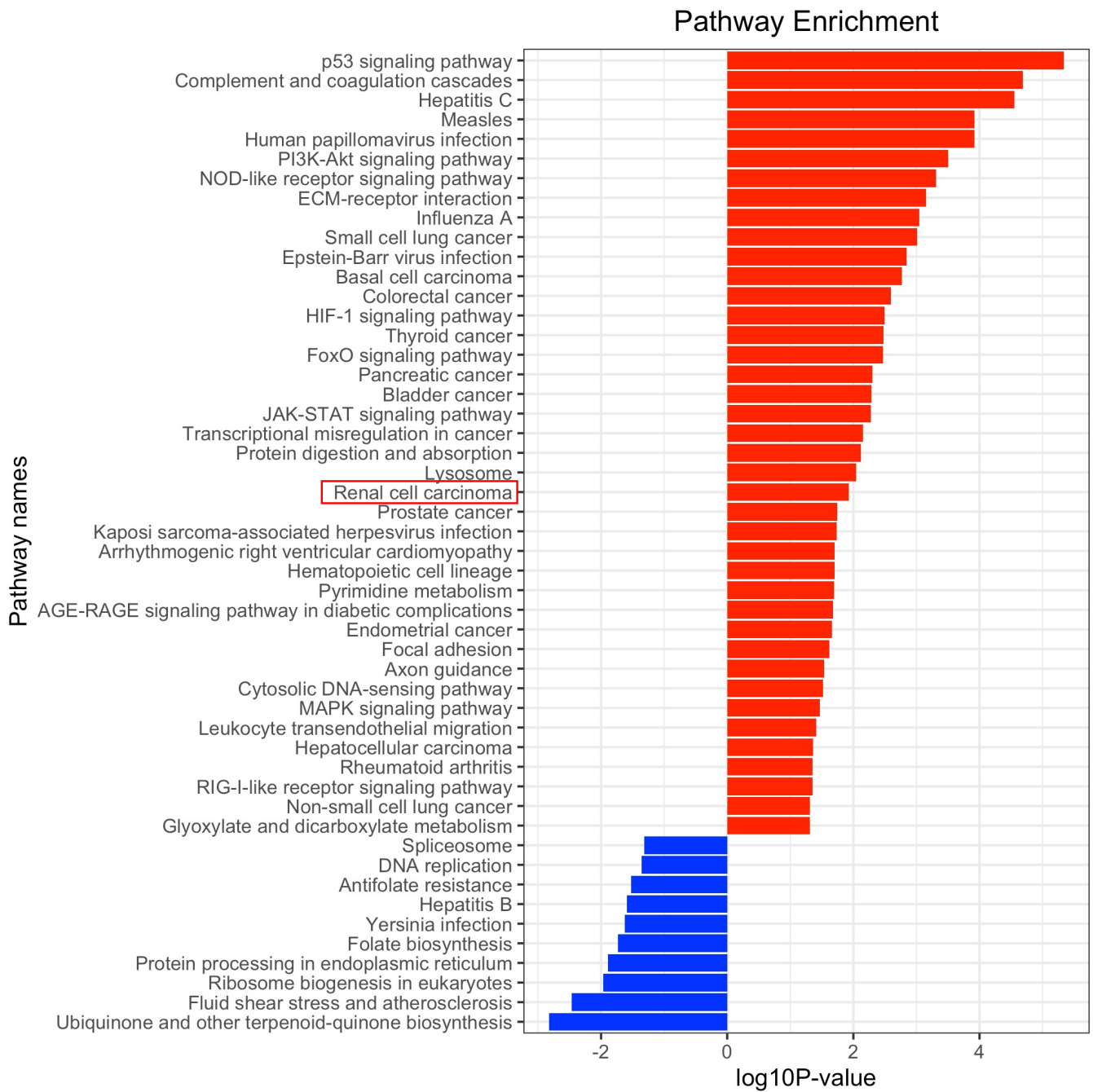


Figure S6. KEGG analyses of RNA seq data from GABPA-depleted 786-O cells show the enrichments of RCC pathways (Marked in RED)