

Additional file 2

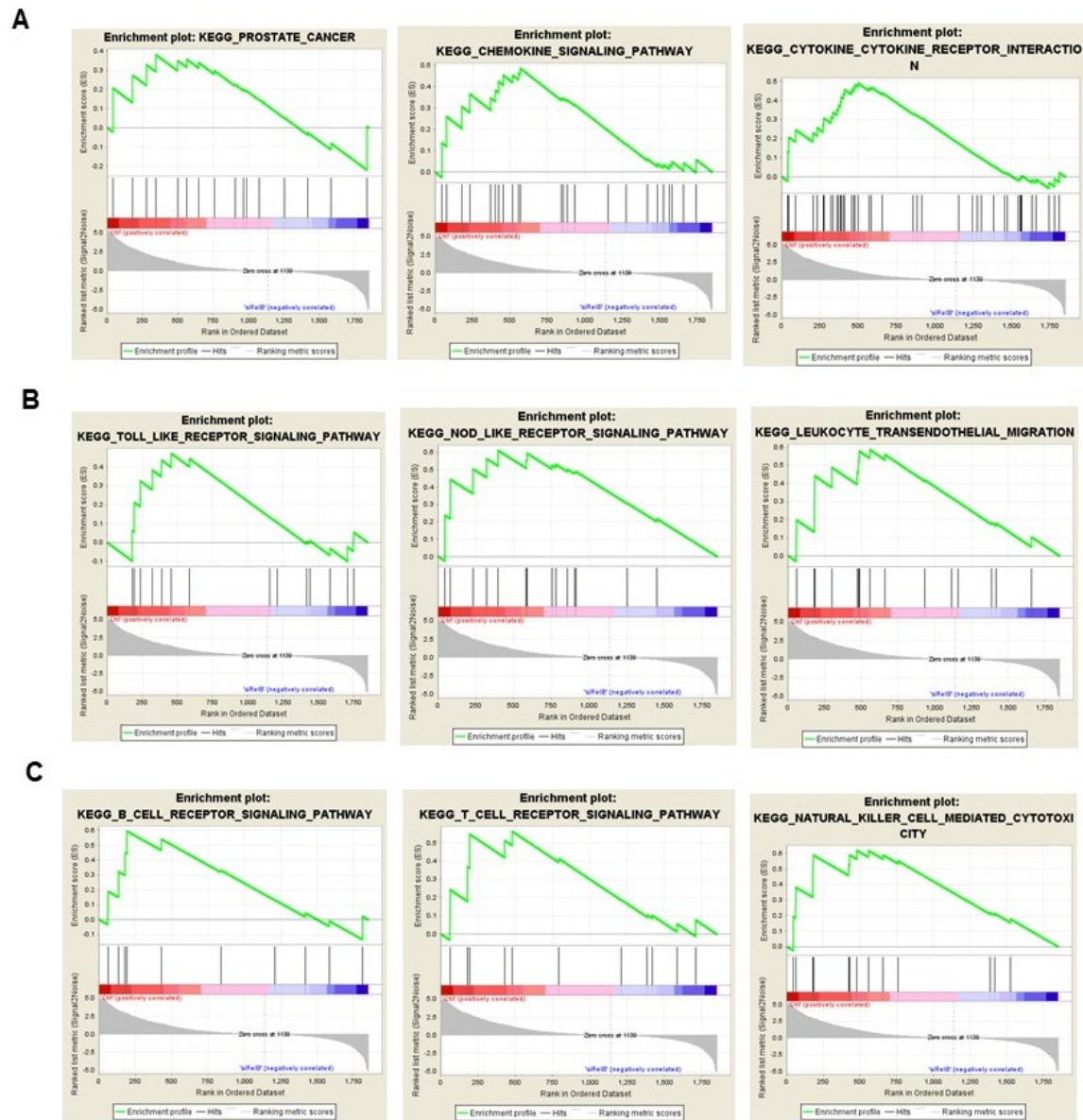


Fig. S2. Comparison of mRNA expression profiles in RelB-silenced PC-3 cells vs. control cells. Total RNA was extracted from shRelB and shCtrl PC-3 cells. The mRNA expression profiles were examined by RNA-Seq. **a** KEGG pathway enrichment analysis was performed to assess the association of altered mRNA profiles with cytokine/chemokine-activated signalling pathways. **b-c** The altered mRNA profiles are also related to oncogenic receptor-mediated signalling pathways and the immune response.