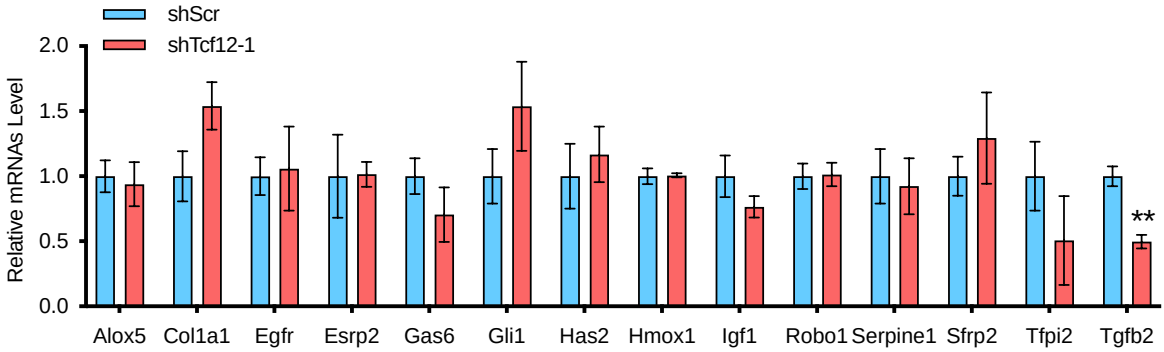


Figure S4. Analysis of potential melanoma-related target genes of TCF12: Expression analysis of a set of melanoma-related target genes with potential TCF12 binding sites in YUMM1.7 cells after TCF12 knockdown. Relative mRNA levels were quantified by RT-qPCR. Statistical significance is based on comparison with control shRNA (shScr) group. ** $p < 0.01$.