

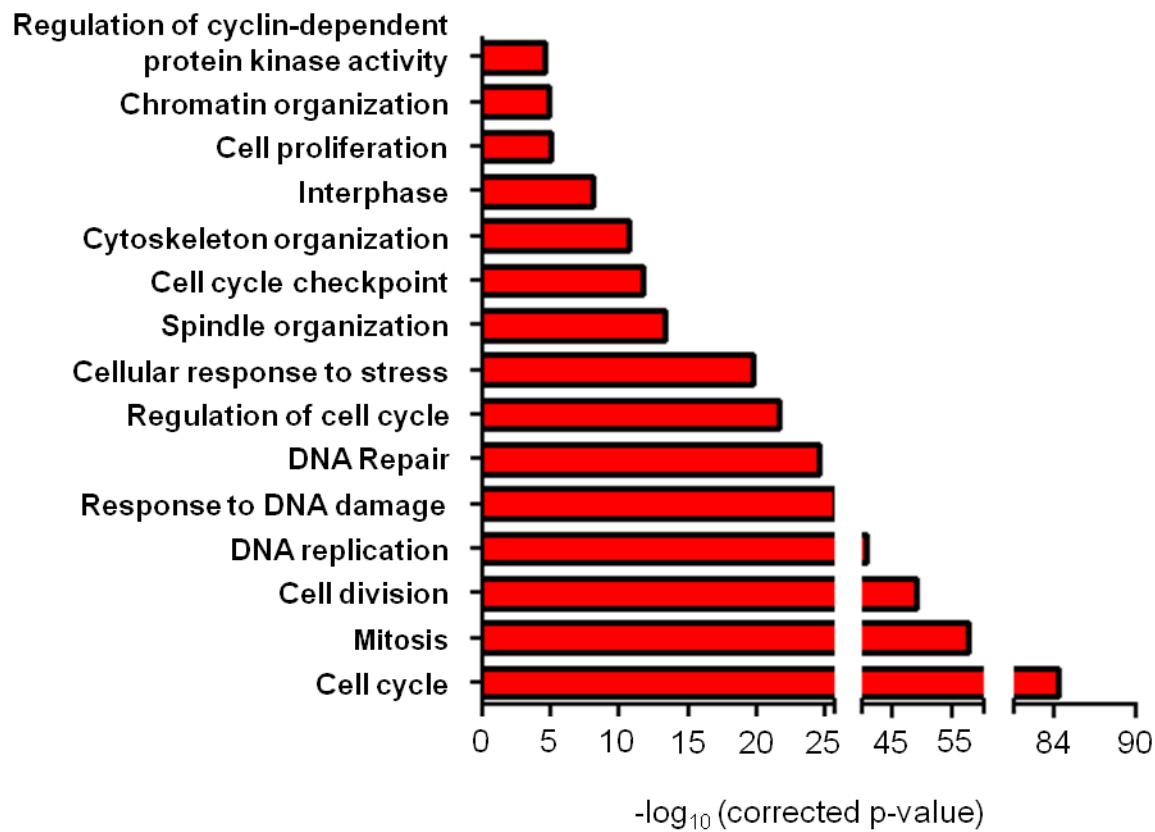


Figure S2. Genes whose expression was upregulated by ZNF750 silencing in HaCaT cells: Gene Ontology (GO) terms enrichment.

For the 381 genes that were upregulated in ZNF750 silenced cultures (shRNA-a) vs. controls ($p\text{-value} < 0.05$, $FC > 2$), a total of 59 GO terms were significantly enriched (Bonferroni-corrected $p\text{-value} < 0.001$). Selected GO terms are shown.