Figure S4

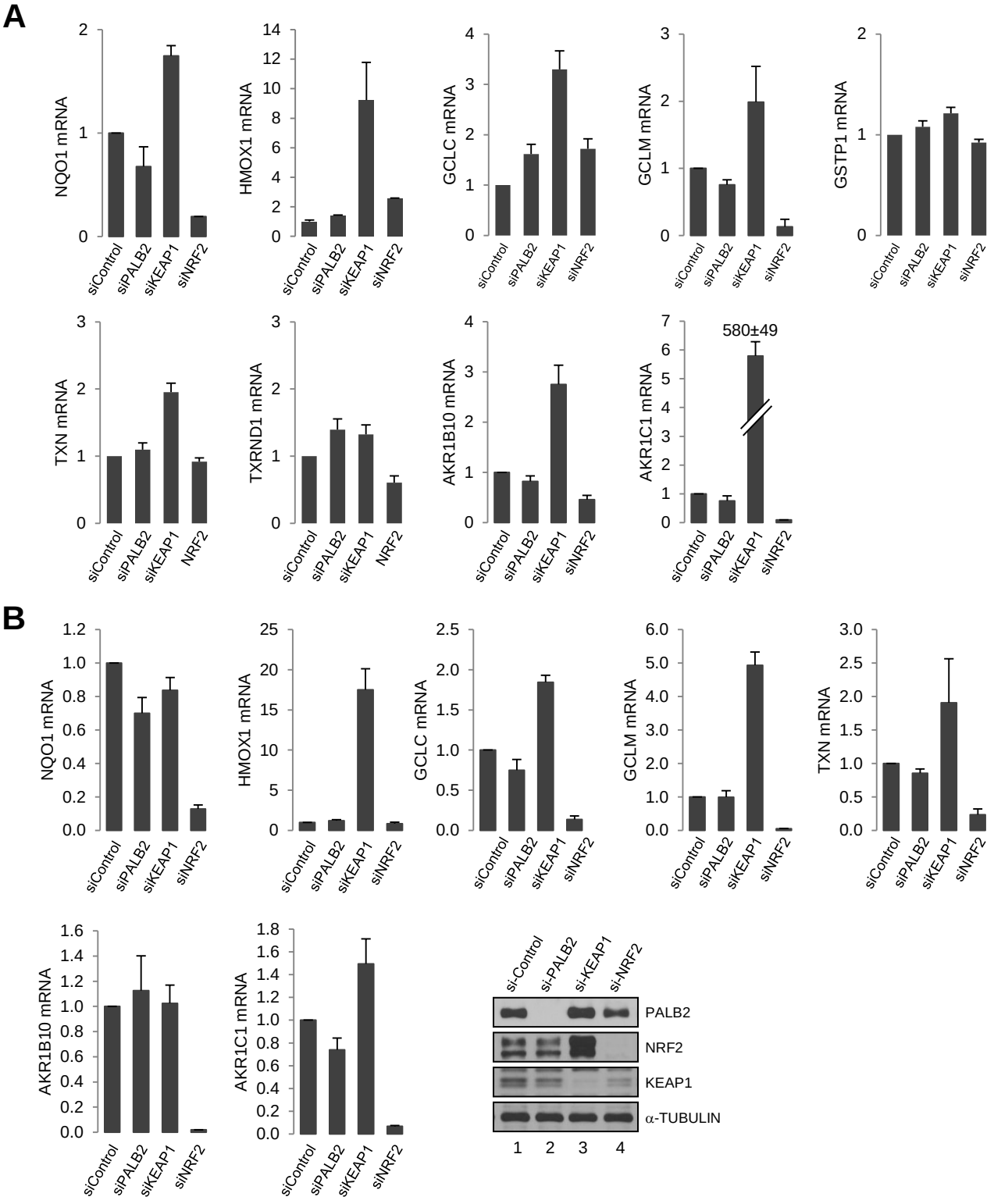

Figure S4. PALB2 depletion reduces expression of a subset of NRF2 downstream genes. (**A**) U2OS cells were treated with control, PALB2, KEAP1 and NRF2 siRNAs for 48 hour. Total RNAs were extracted and mRNA levels were determined by qRT-PCR. Knockdown of protein levels is as shown Fig. 6A. (**B**) T98G cells were treated with the siRNAs for 48hr, and expression levels of NRF2 downstream genes were analyzed by qRT-PCR. PALB2, KEAP1 and NRF2 protein levels were analyzed by Western blotting.