

Figure S1. Reverse transcription-quantitative PCR analysis showing the expression levels of three differentially expressed genes in the RNA-seq analysis of FOXP3-overexpressing MDA-MB-231 cells vs. control cells. * $P < 0.05$, ** $P < 0.01$. PDCD4, programmed cell death 4; FOXP3, forkhead box P3; PGGHG, protein-glucosylgalactosylhydroxylysine glucosidase; SQOR, sulfide quinone oxidoreductase.

