Supplemental Figure 3. *DGKα modulates several oncogene-related pathways.* A, In both U87 and U251 cells, mRNA levels of FDPS1, FDPS2, HMGCR, and SCD were quantified by qRT-PCR in response to DGKα knockdown via siRNA and log-scale fold expression changes in comparison to control siRNA are shown. B, c-Myc was over-expressed through plasmid transfection after DGKα knockdown via siRNA and cell proliferation was assessed in U87 cells. (*, P<0.05 and **, P<0.01 Student t test).