



Supplemental FIGURE 2. Expression levels of glioblastoma-associated lncRNAs correlate with the expressions of cell-cycle genes.

(A) Top 30 KEGG pathways significantly correlated with the expression of therapy-resistant lncRNAs. **(B)** Top 30 KEGG pathways significantly correlated with the expression of therapy reversed lncRNAs. Each column corresponds to a single lncRNA, and each row corresponds to a KEGG pathway with an overrepresentation of genes correlating with lncRNA ($p \leq 0.01$). The pathways were ranked by the sum of the negative log₁₀ p-value of each lncRNA for each pathway. Top 30 pathways are shown.