



Figure S1

Expression of *VEGFA*, *MYC* (A), and fucosylation-related genes (B) in HepG2 cells treated with mithramycin. *RPL4* was used as the reference gene, and data are shown as the average \pm S.D. (N = 3 for each condition). Student's t-test with Bonferroni correction, * $p < 0.05$, comparing with 0 μ M mithramycin in each gene.