

Figure S1. RNA was extracted from tumor samples and cDNA was prepared. The quantitative gene expression of SMAD2, OGT, RAB10 and CUL4b was performed using SYBR green approach. GAPDH was used as house keeping gene. Statistical significance was determined by Student's *t*-test, *** $p < 0.001$.

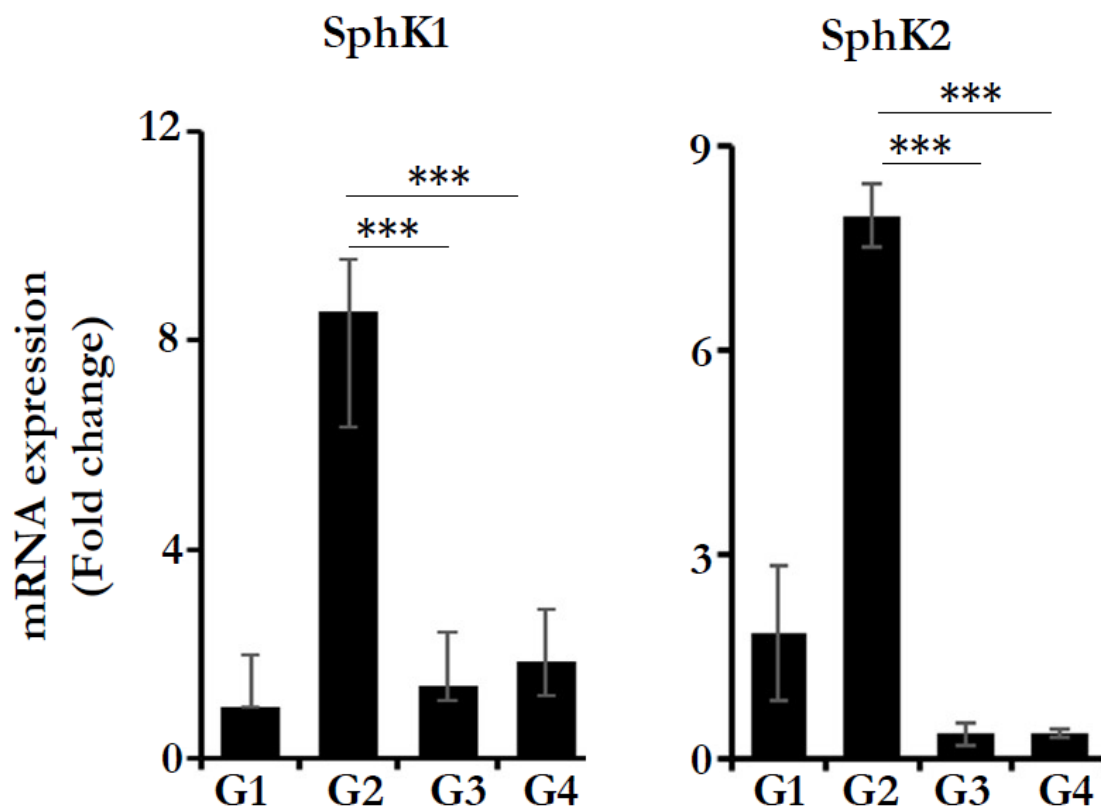


Figure S2. RNA was extracted from tumor samples and cDNA was prepared. The quantitative gene expression of SphK1 and SphK2 was performed using SYBR green approach. GAPDH was used as house keeping gene. Statistical significance was determined by Student's *t*-test, *** $p < 0.001$.