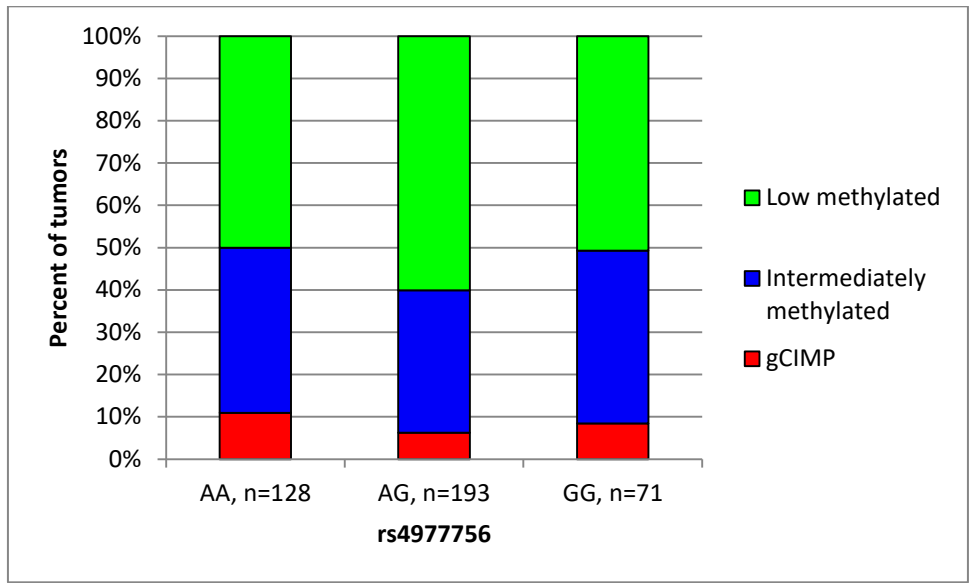


A



B

S2 Figure. Association between rs4977756 and DNA methylation pattern of the tumor. Distribution of DNA methylation patterns between genotypes in (A) 77 glioma tumors, and (B) 392 TCGA glioblastoma tumors. $P_{AA \text{ vs. } AG/GG} = 4.41 \times 10^{-4}$ and 0.23 in (A) and (B), respectively.