Figure S1. Distribution and location of missense mutations in the top 10 mutated genes in gliomas. Schematic representation of the distribution and location of missense mutations in (A) MKI67, (B) CENPF, (C) BUB1, (D) KIF4A, (E) TOP2A, (F) KIF20A, (G) DLGAP5, (H) NUF2, (I) CEP55 and (J) BUB1B in GBM and LGG. MKI67, marker of proliferation Ki-67; CENPF, centromere protein F; BUB1, BUB1 mitotic checkpoint serine/threonine kinase; KIF4A, kinesin superfamily protein 4A; TOP2A, DNA topoisomerase IIα; KIF20A, kinesin superfamily protein 20A; DLGAP5, DLG associated protein 5; NUF2, NUF2 component of NDC80 kinetochore complex; CEP55, centrosomal protein 55; BUB1B, BUB1 mitotic checkpoint serine/threonine kinase B; GBM, glioblastoma multiforme; LGG, lower-grade gliomas.