

Supplemental Table Legend

Supplemental Table 1. Description and features of the four major human cancer sample types.

Supplemental Table 2. Copy number alterations present in a panel of 50 malignant glioma cell lines. These copy number data were generated on Affymetrix 6.0 SNP arrays by the Cancer Genome Project of the Wellcome Trust Sanger Institute.

Supplemental Table 3. Mean copy number values reported by the TCGA of the genomic deletions in primary tumor samples containing homozygous deletions of CDKN2A (a), PTEN (b), CDKN2C (c), and NF1 (d).