**S1 Table. Identification of a diverse panel of GBM BTICs representative of the different mutational statuses of GBM patients.**

Common GBM molecular alterations including *MGMT* promoter methylation*, EGFR*, *PTEN*, *TP53*, *NF1*, *IDH1*, and *CDKN2A* mutation statuses for the eleven BTIC cultures used in this study. Mut indicates mutant and wt indicates wild-type. U indicates unmethylated and M indicated methylated. U/M indicates hemi-methylation. vIII indicates *EGFR* variant III, an activating deletion characteristic of GBM, het del indicates a heterozygous deletion, homo del indicates a homozygous deletion, N/A indicates not available.