

## SUPPLEMENTARY TABLE

Supplementary Table 5. Patient characteristics in the TCGA RNA-seq and CGGA\_301 microarray data.

Characteristics	TCGA RNA-seq (n=697)	CGGA microarray (n=301)
<b>Gender</b>		
male	370	180
female	271	121
NA	56	0
<b>Age (year)</b>	47 ± 15	42 ± 12
<b>Tumor subtype</b>		
Classical	90	23
Mesenchymal	104	111
Proneural	248	86
Neural	115	81
NA	140	0
<b>WHO grade</b>		
Grade II	226	122
Grade III	249	51
Grade IV	167	128
NA	55	0
<b>Karnofsky Performance Score</b>	84 ± 14	NA
<b>IDH mutation status</b>		
Mut	442	134
WT	245	165
NA	10	2
<b>1p/19q Codeletion status</b>		
Codeletion	181	16
Non-codeletion	491	76
NA	25	209
<b>MGMT promoter status</b>		
Methylated	461	99
Unmethylated	162	187
NA	74	15

NA: Not Available; KPS: Karnofsky Performance Score; MGMT: O<sup>6</sup>-Methylguanine Methyltransferase.