

Supplemental Table 1 Patient characteristics in TCGA RNA-seq and CGGA_301 microarray data.

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

NA: Not Available; KPS: Karnofsky Performance Score; MGMT: O⁶-

Methylguanine Methyltransferase