Hazard ratio for Low-grade glioma astrocytoma 9.4e + 1390.011 * (N=151)(1.1e+32 - 8,1e+247) Macrophage_M1 (N=151)< 0.001 (2.3e+47 4.4e+93) 0.037 * Macrophage_M2 (N=151)(N=151)< 0.001 * 9.9e-29) (N=151)reference 0.003 ** T_cell_CD4_non_regulateNy=151) 0.451 -248 - 1.2e+1<u>1</u>0 (N=151)reference T cell regulatory Tregs (N=151) reference 4.0e-07 Myeloid dendritic cell (N=151) LGG, BRAF V600E 1.4 0.161 molecular subtype < 0.001 ' GG, BRAF V600E, BUHN22/E < 0.001 ' (N=4) GG:FGFR (3.9e+02 - 4.9e+05 0.004 ** (N=2) KIAA1549-BRAFe-01 N=70 GG, KIAA1549-BRATE 11.6 0.227 (N=1) MYB/MYBL1 reference (N=2) NF1-germline reference (N=1) NF1-somatic_{reference} (KGG, other MAPK reference (N=10) RTK reference (N=5) RTK, CDKN2A/Brence GG. To be classified.7e+00 0.102 (N=4) (GG, wildtype 7.7e-01 - 1.8e+0 (N=29) (3.3e-01 - 7.2e+0 0.574 3.3e-01-72e+0 0.925 (6.4e-01-1.5e+0)# Events: 15; Global p-value (Log-Rank): 0.018329

1e+43

1e-39

1e+125

1e+207

B cell

Monocyte

Neutrophil

T cell CD8

NK_cell

CD274

AIC: 110.45; Concordance Index; 0.92