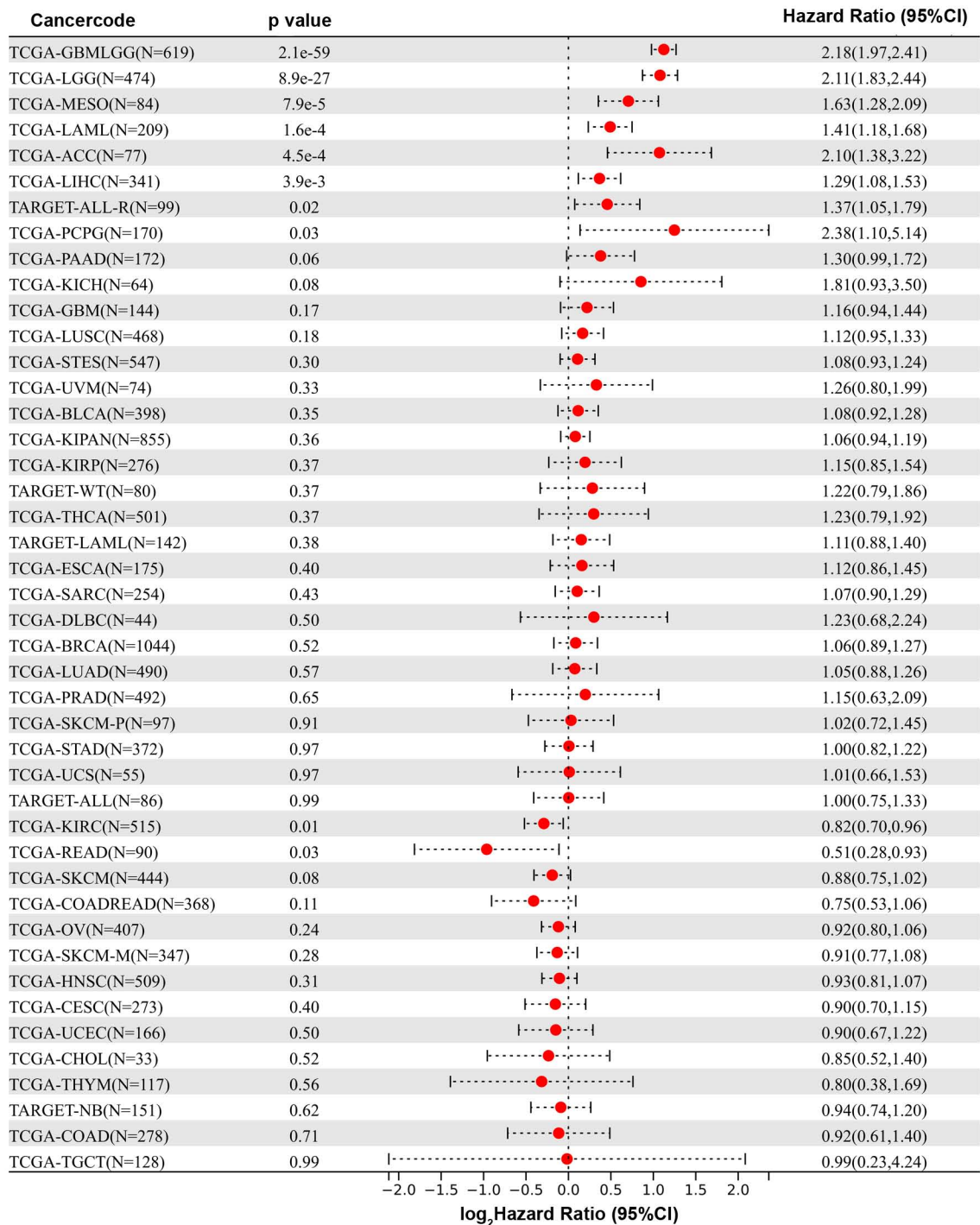
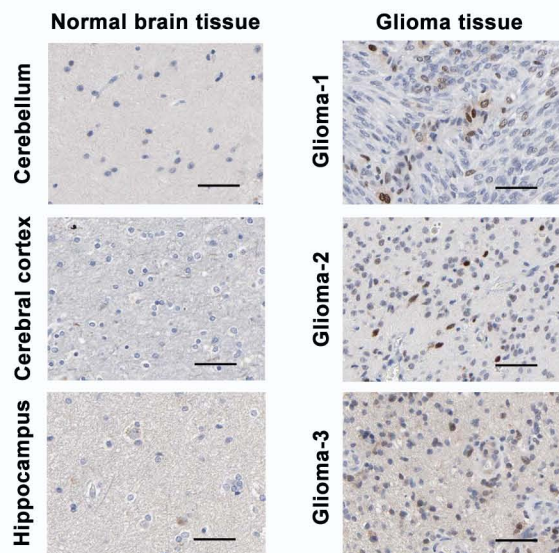


Supplementary Figure 1

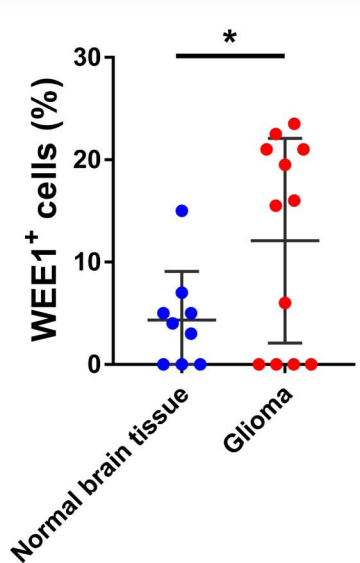


Supplementary Figure 2

A

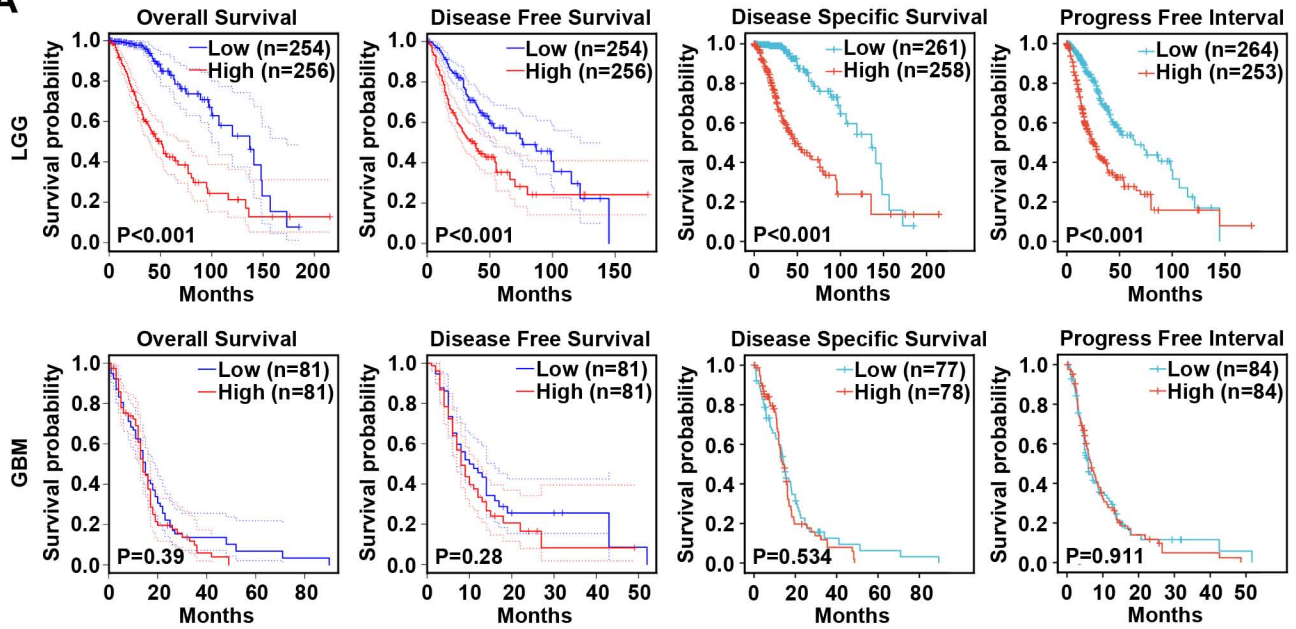


B

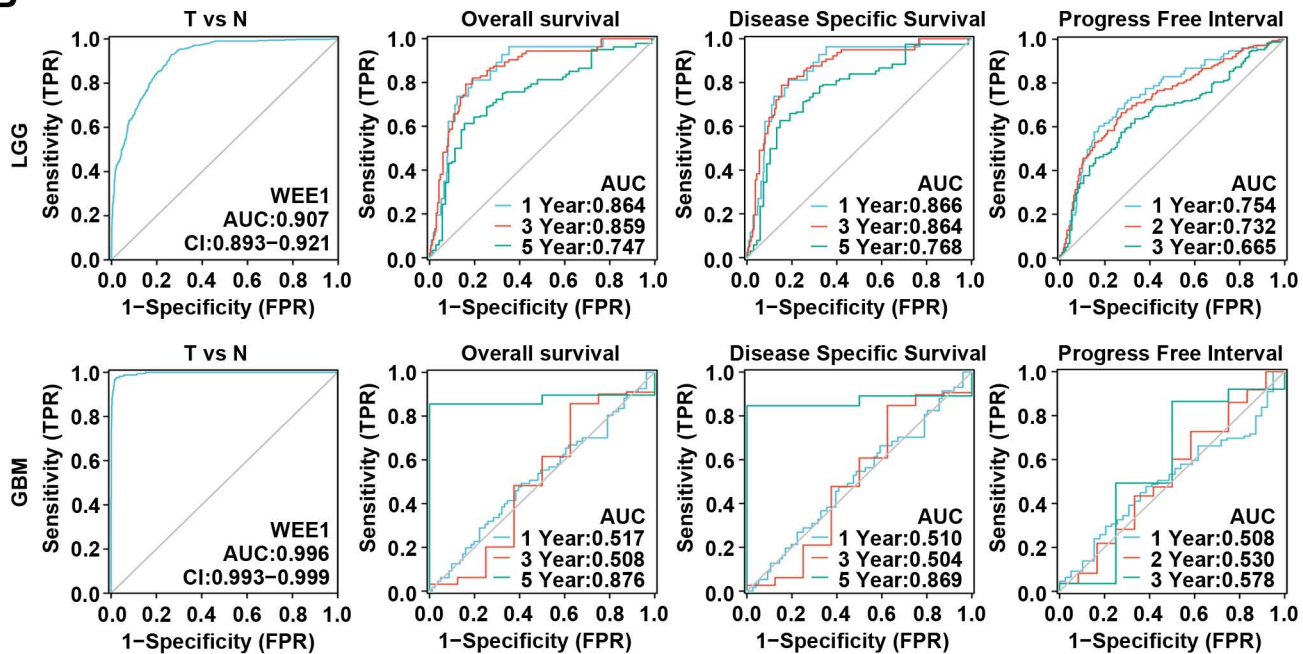


Supplementary Figure 3

A

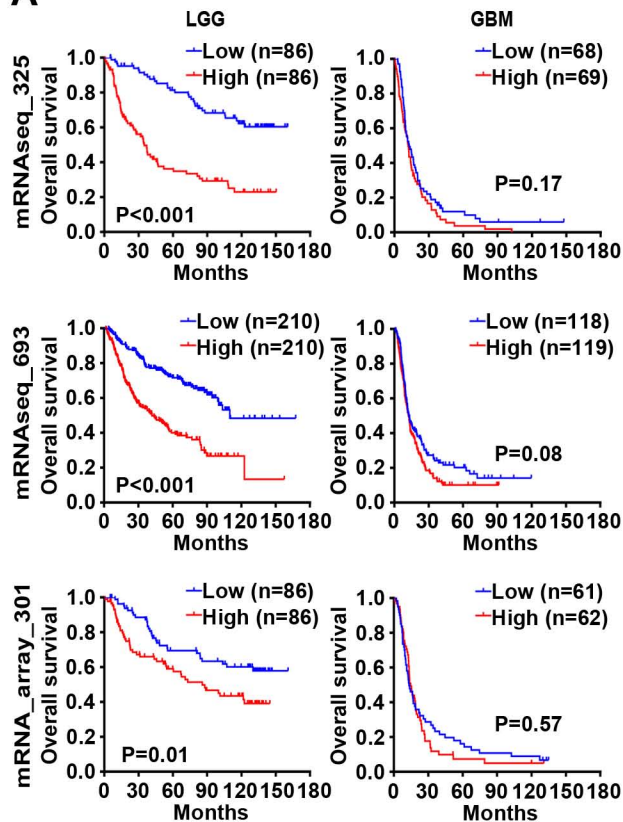


B

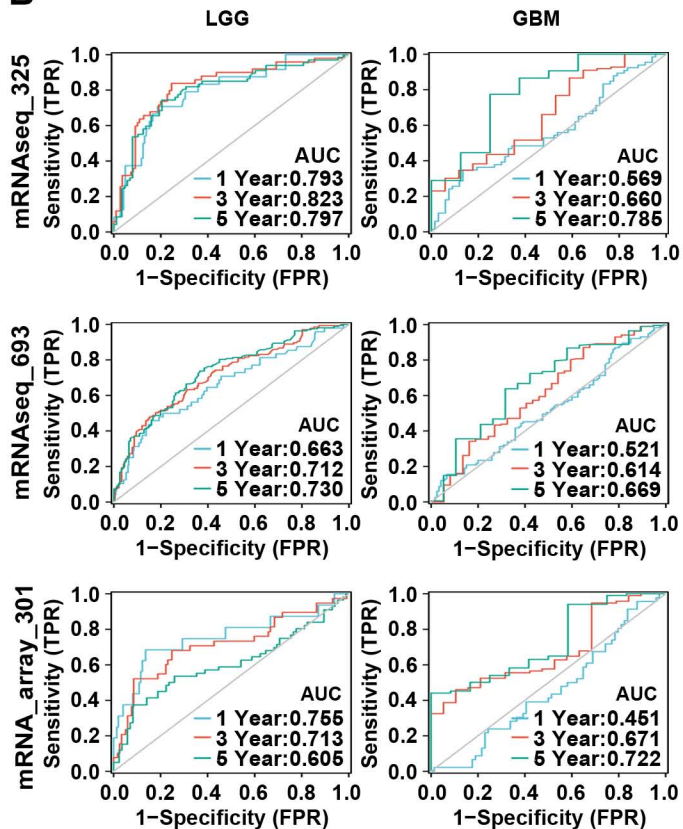


Supplementary Figure 4

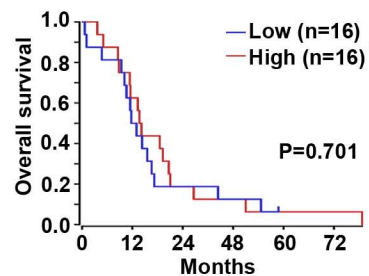
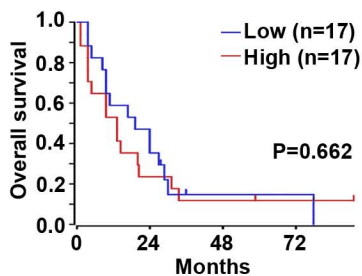
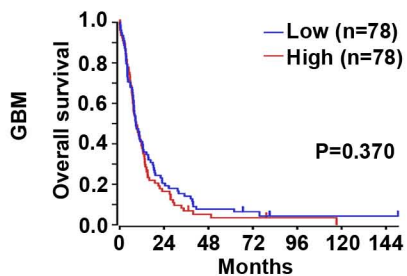
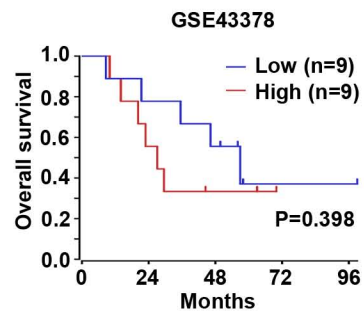
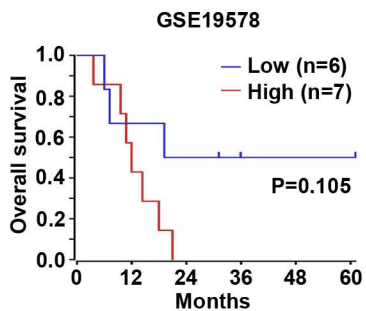
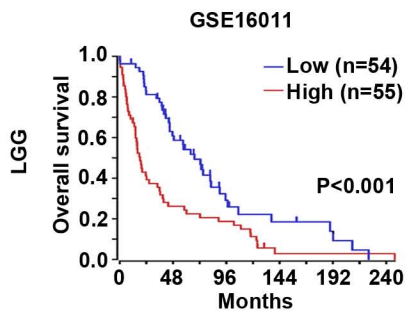
A



B

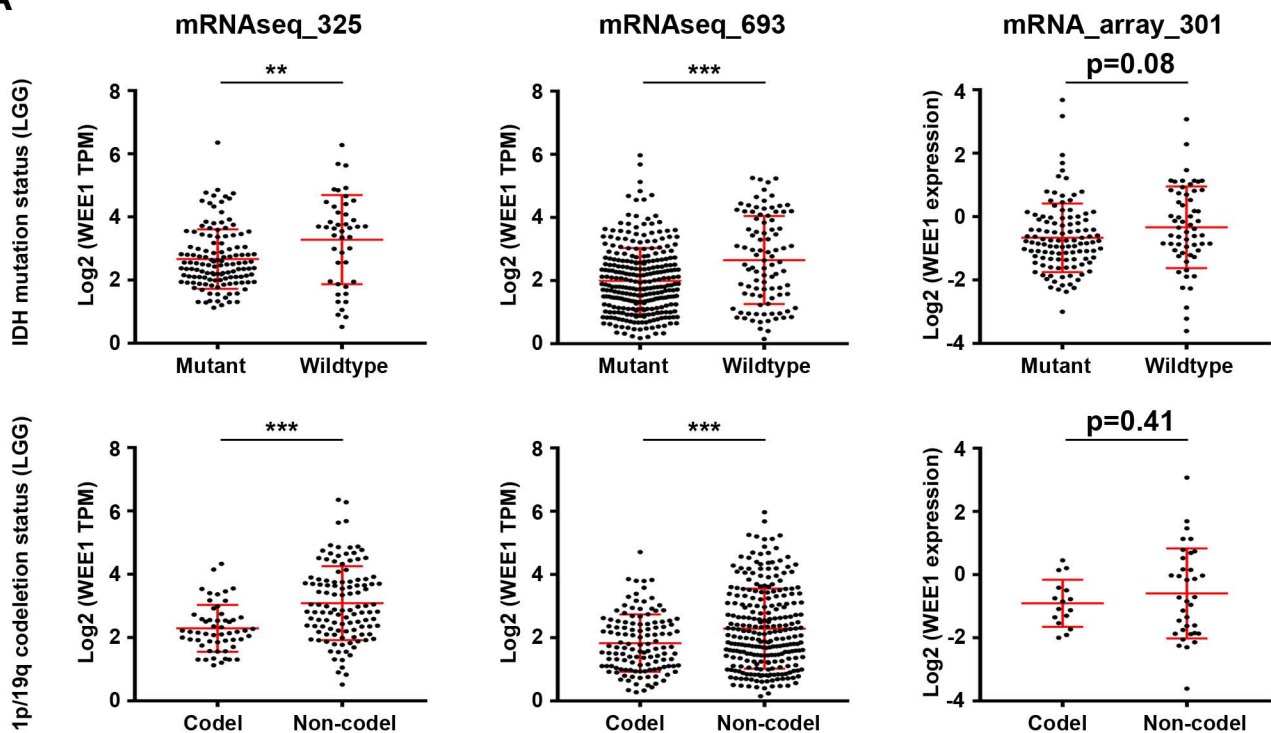


Supplementary Figure 5

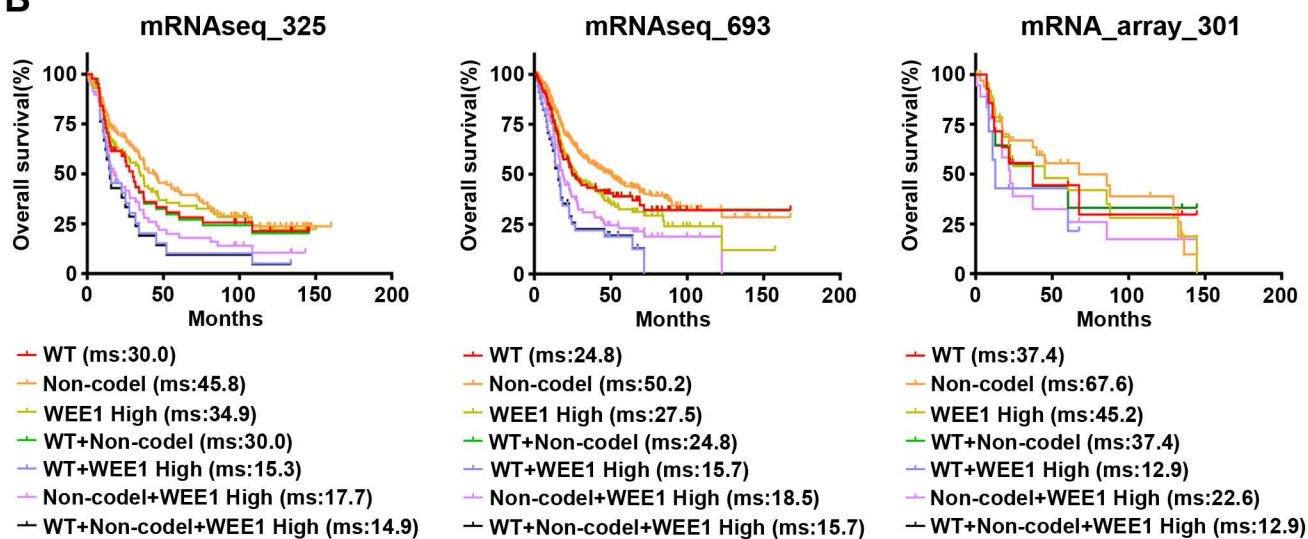


Supplementary Figure 6

A

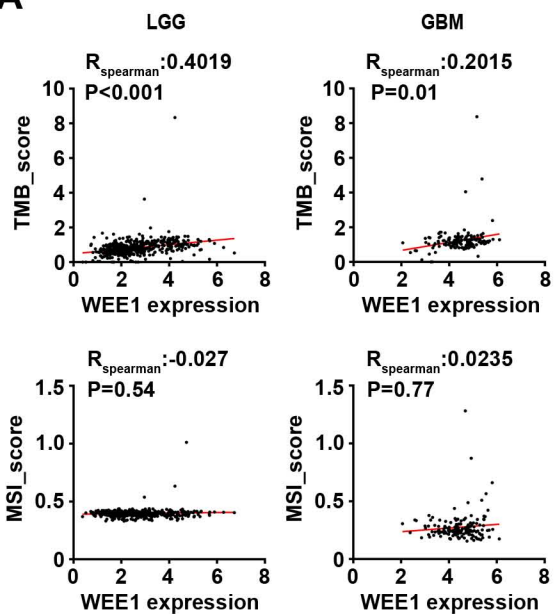


B

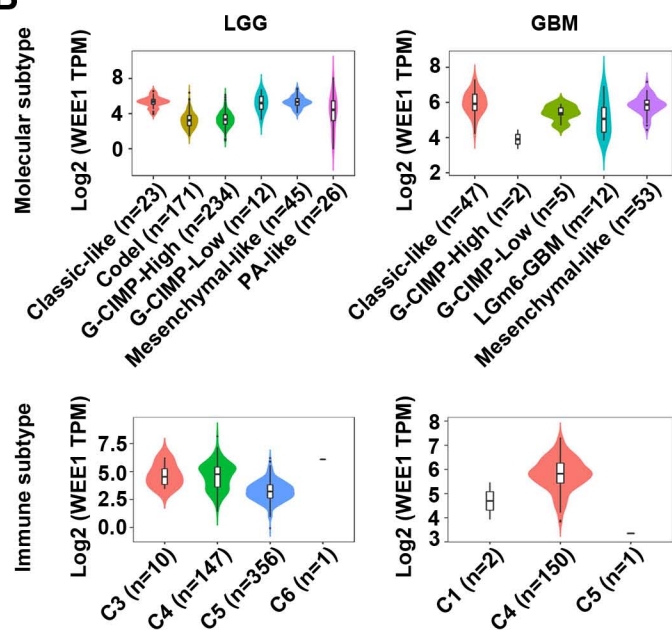


Supplementary Figure 7

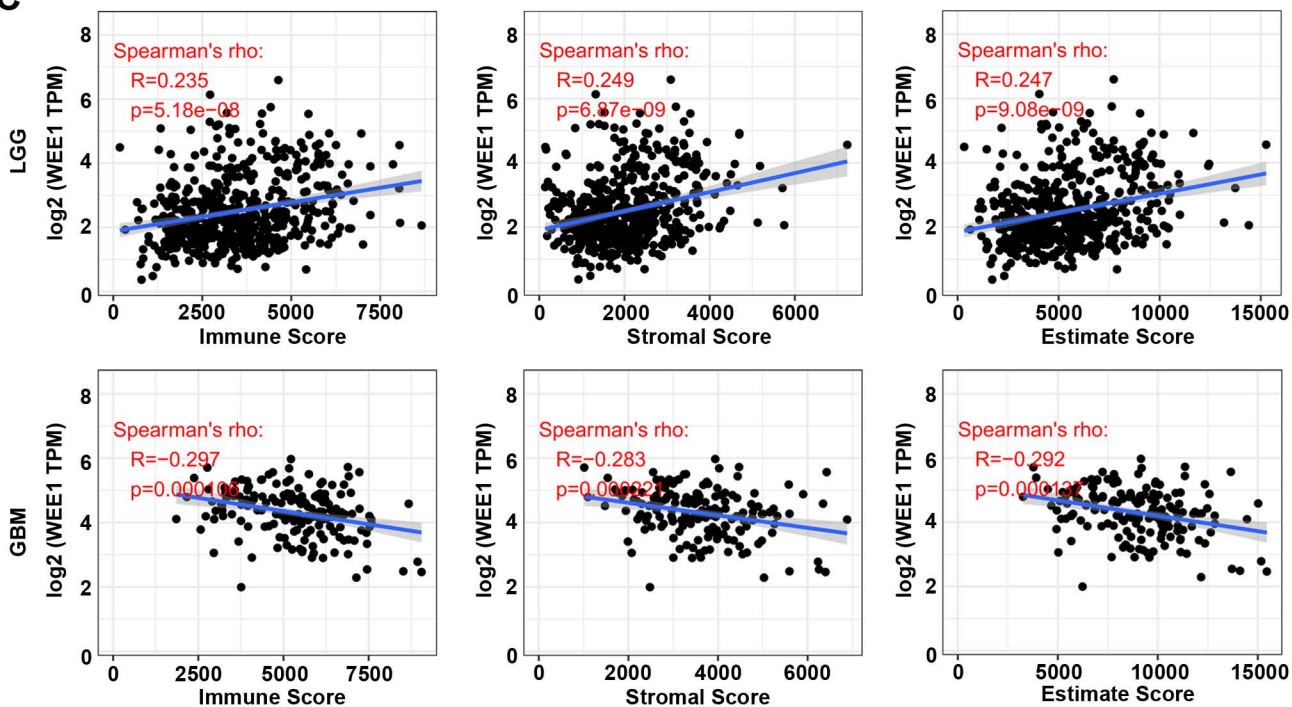
A



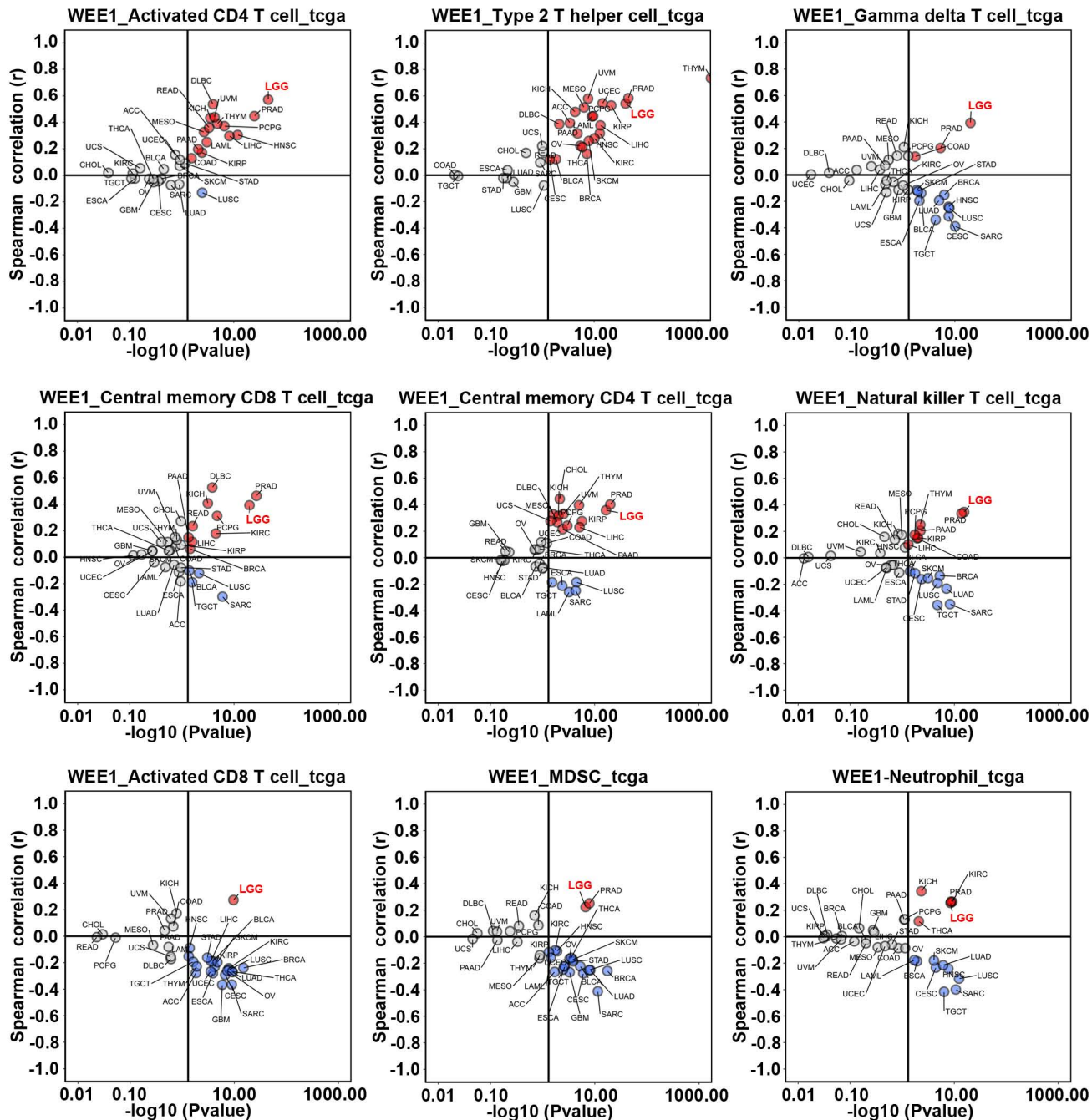
B



C

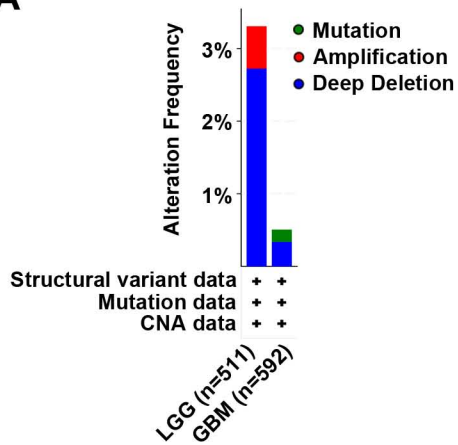


Supplementary Figure 8

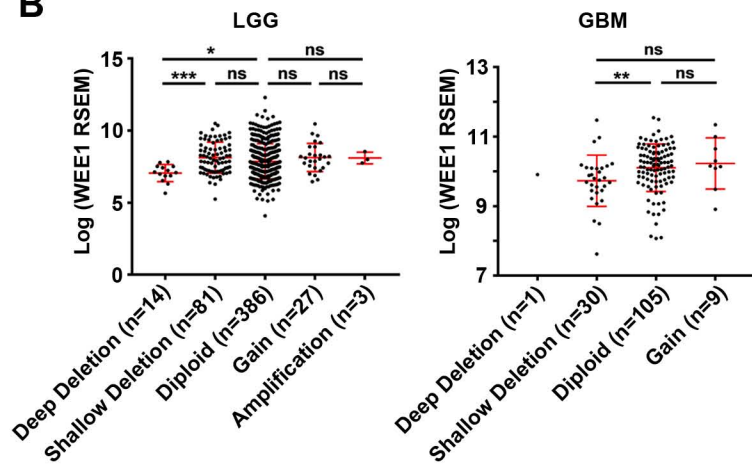


Supplementary Figure 9

A

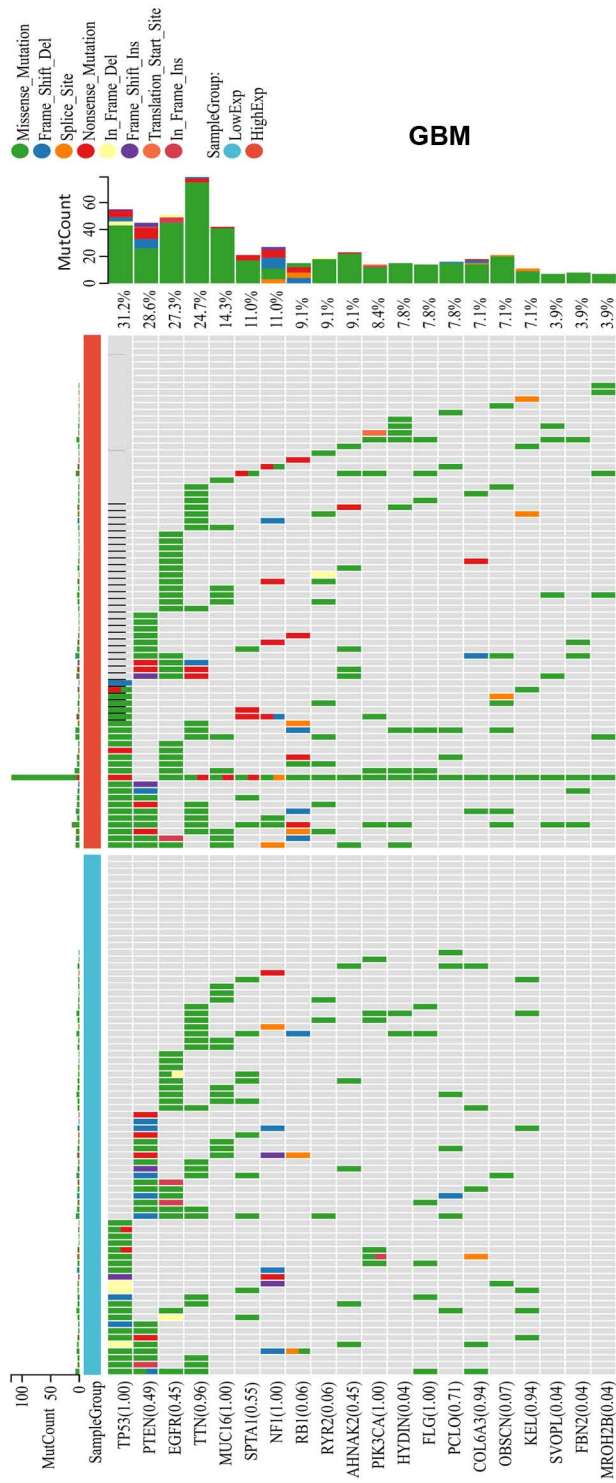


B



Supplementary Figure 10

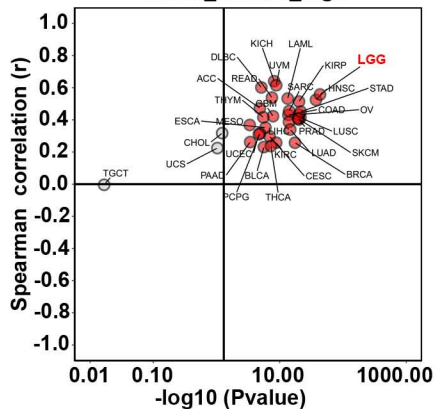
A



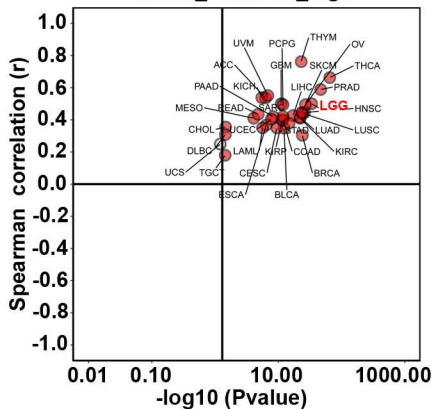
Supplementary Figure 11

A

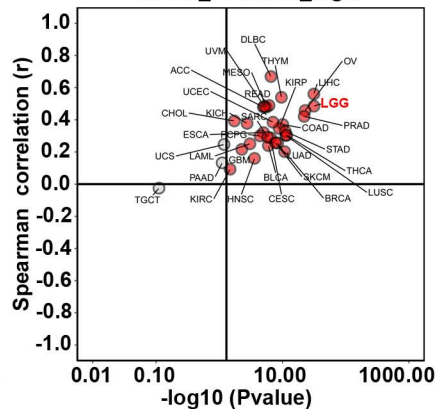
WEE1_DNMT1_tcga



WEE1_DNMT3A_tcga

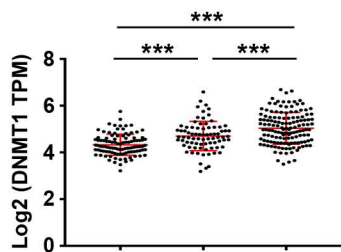


WEE1_DNMT3B_tcga

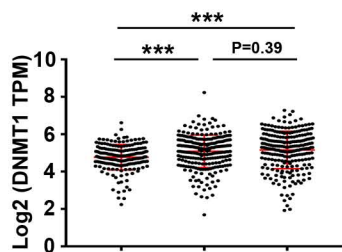


B

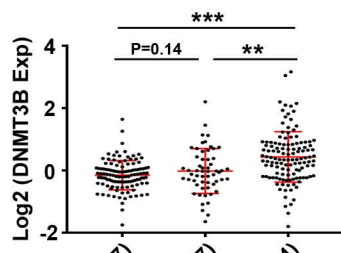
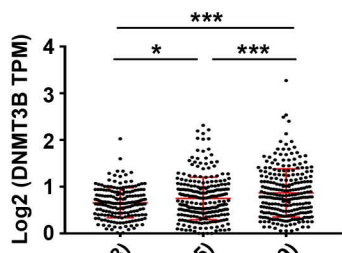
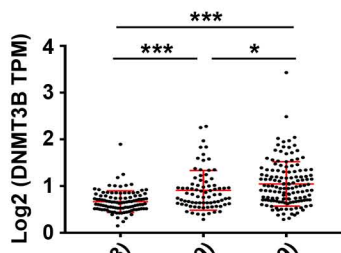
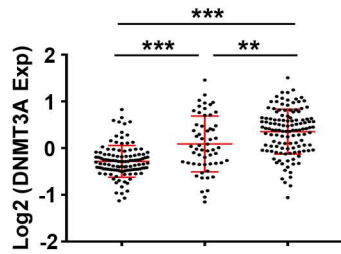
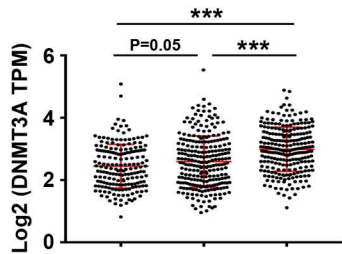
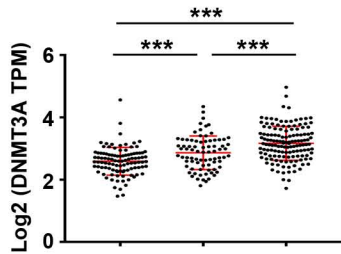
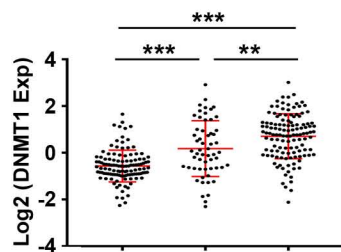
CGGA (mRNAseq_325)



CGGA (mRNAseq_693)



CGGA (mRNA_array_301)

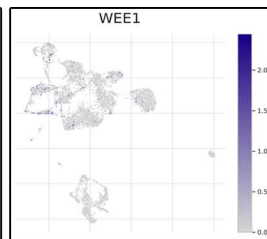
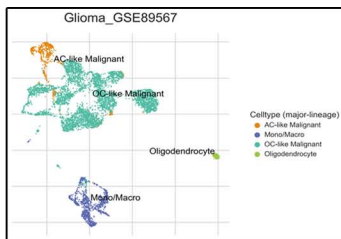
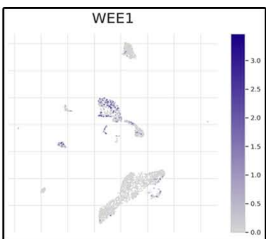
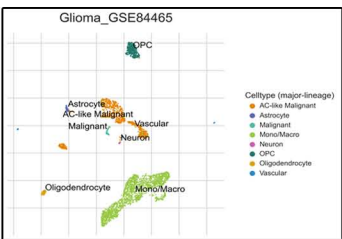
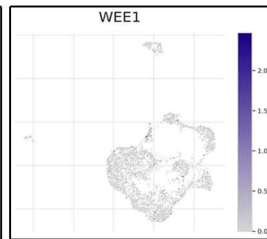
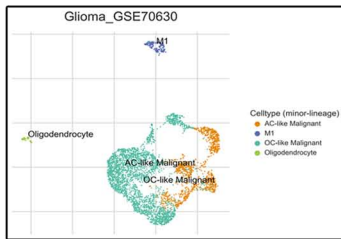
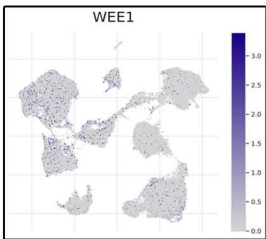
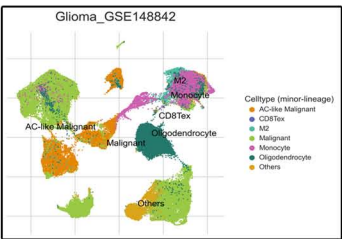
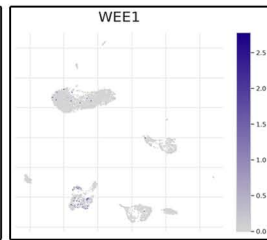
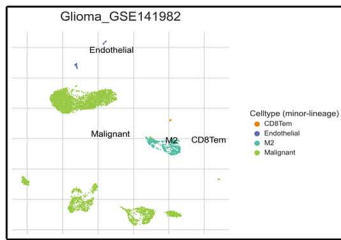
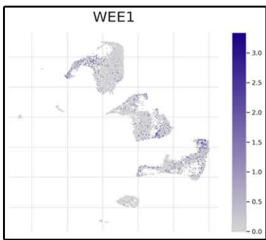
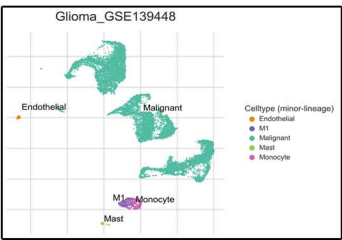
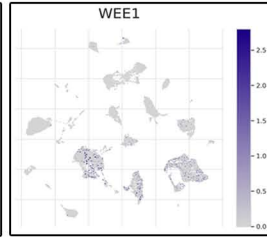
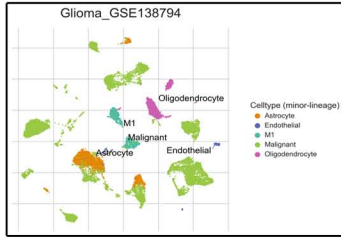
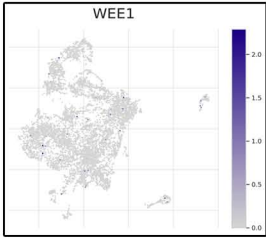
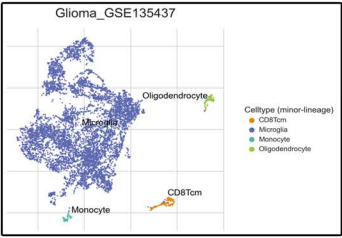
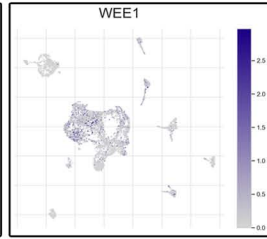
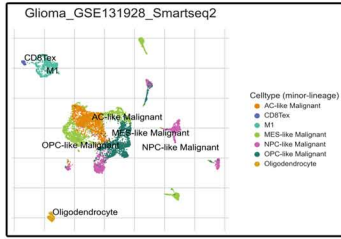
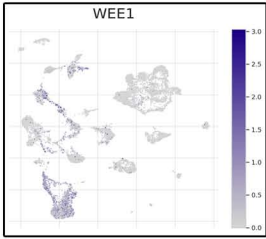
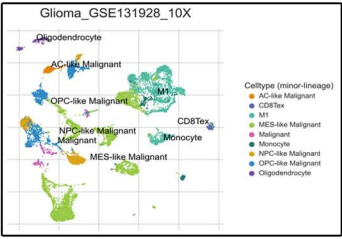
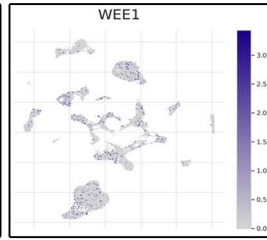
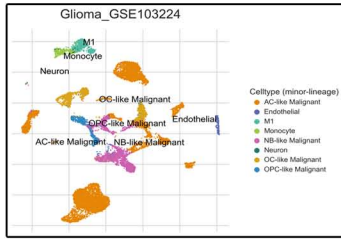
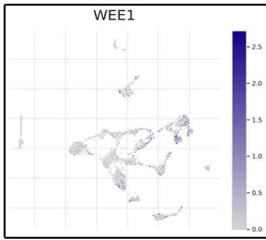
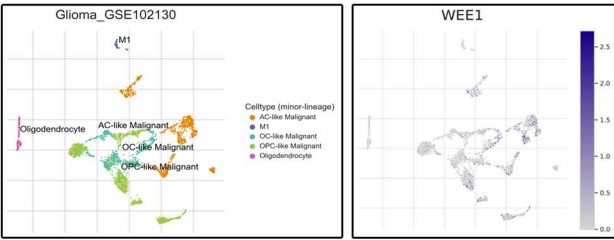


LGG_Grade II (n=103)
LGG_Grade III (n=79)
GBM_Grade IV (n=139)

LGG_Grade II (n=188)
LGG_Grade III (n=235)
GBM_Grade IV (n=249)

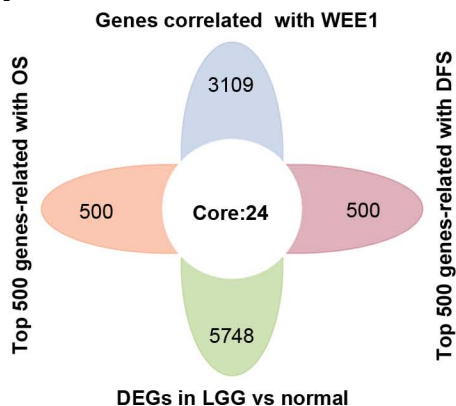
LGG_Grade II (n=117)
LGG_Grade III (n=57)
GBM_Grade IV (n=124)

Supplementary Figure 12

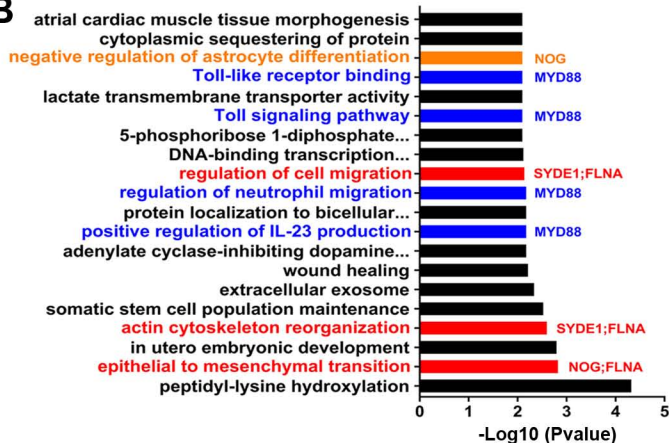


Supplementary Figure 13

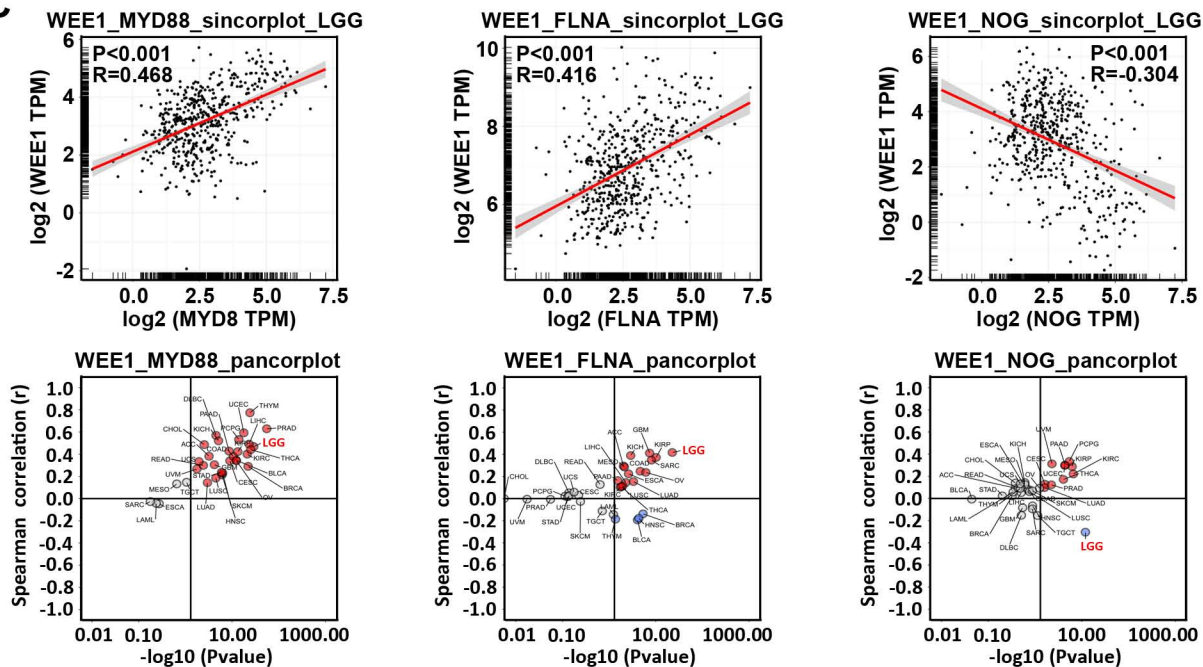
A



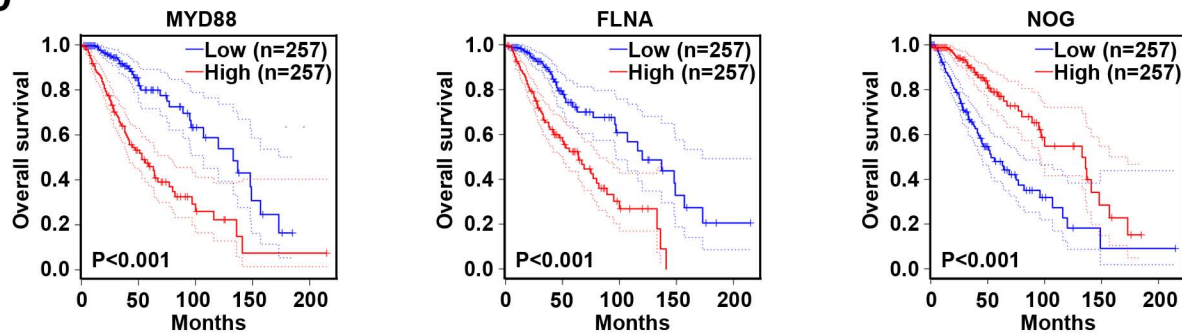
B



C



D



Supplementary Figure 14

