

Figure S1. (A) Somatic mutation landscapes of patients with advanced NSCLC with BM (n=26). (B) Venn diagram of genomic subtyping in the BM group and the non-BM group. (C) Somatic mutation landscapes of advanced NSCLC patients without BM (n=64). Mutated genes are ranked by mutant frequency. The bars on the right correspond to the mutation frequencies of each gene. TMB (mutations per Mb) is shown in the upper panel. Patient samples were arranged along the x-axis. NSCLC, non-small cell lung cancer; BM, brain metastases; TMB, tumor mutation burden.

