

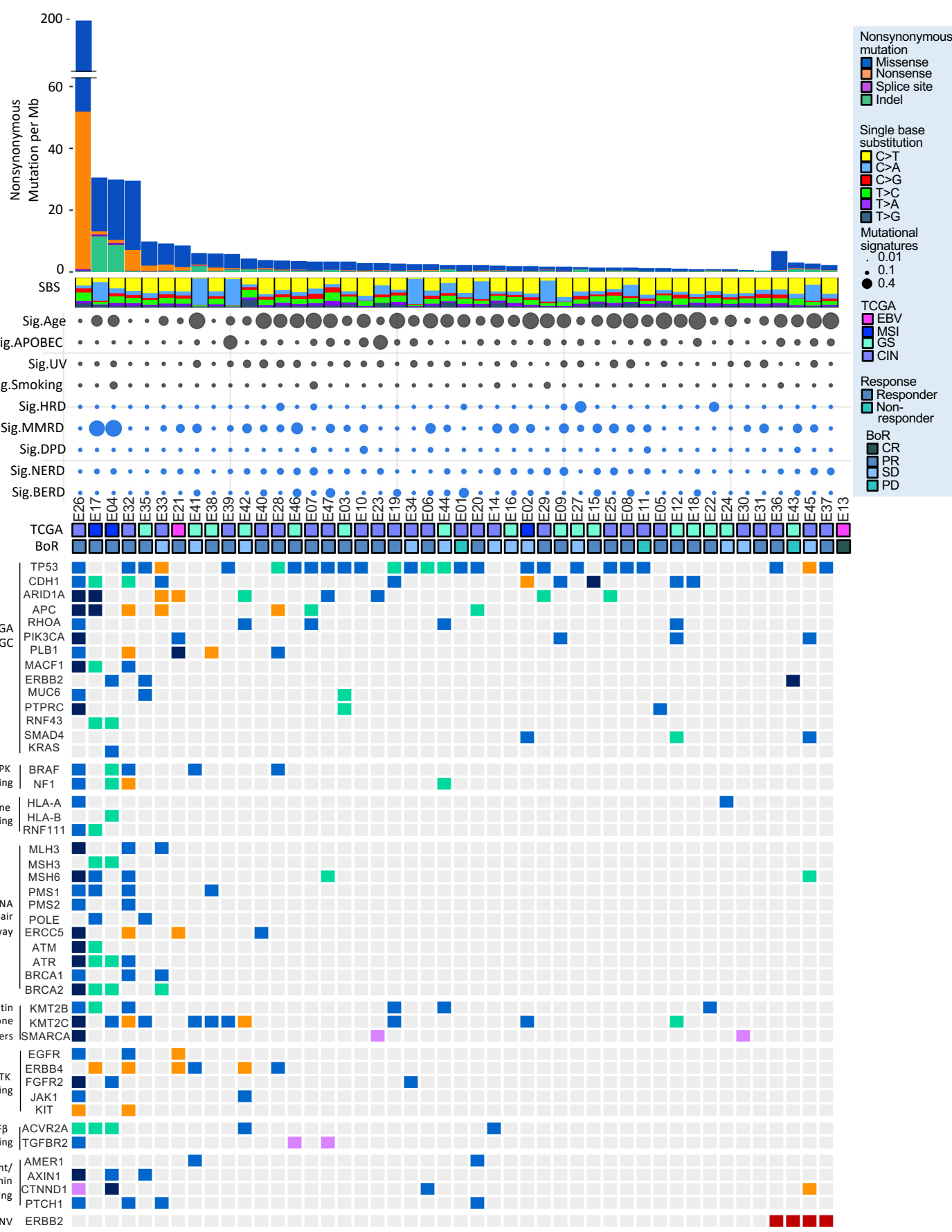
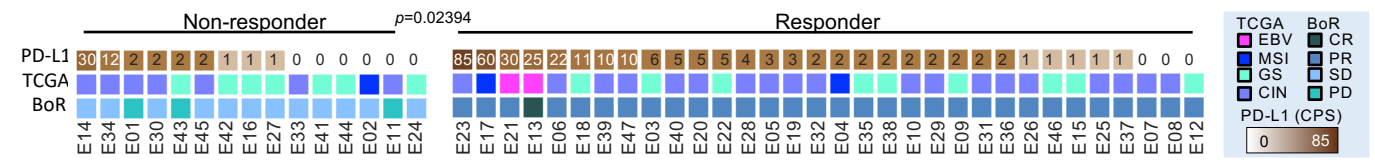
A**B**

Figure S2. (A) CoMut plot demonstrating TCGA subtype, tumor mutational burden, mutational signature enrichment and mutational profiles of key driver genes for all patient samples at baseline in the reported clinical trial. (B) PD-L1 expression for each patient sample at baseline in the reported clinical trial.