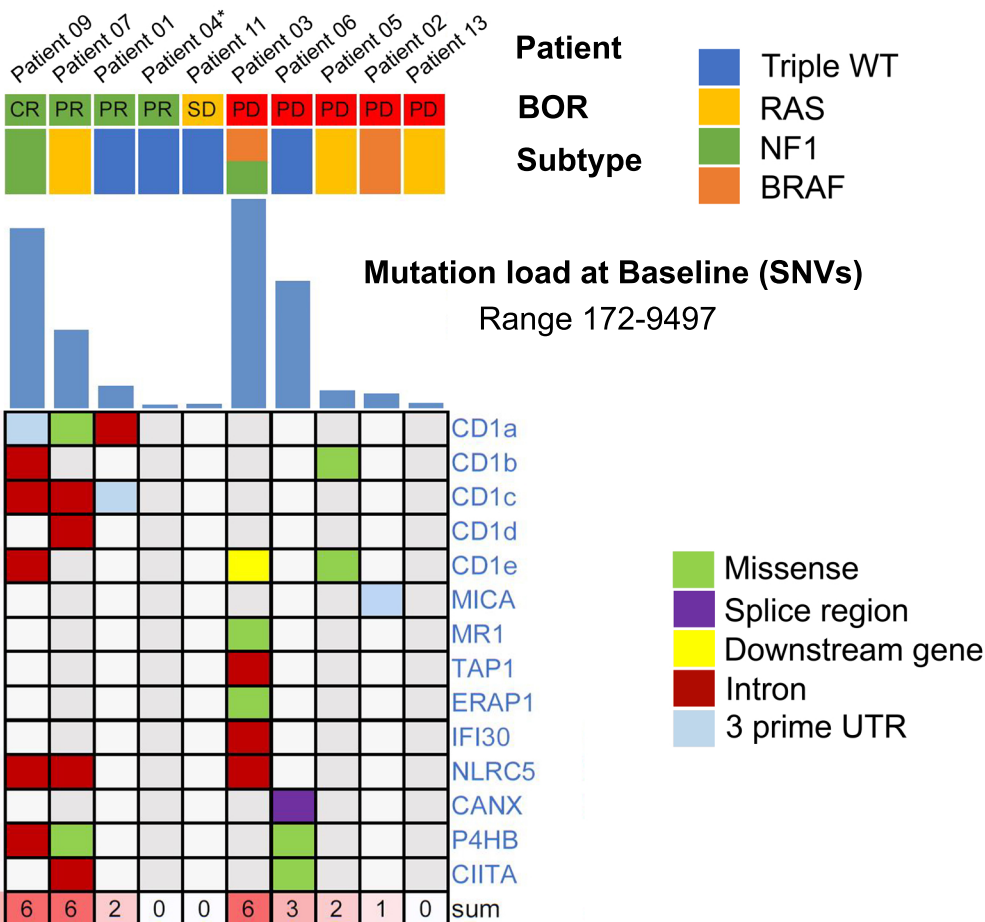# Supplementary figure 5

Genomic features of evaluable patients at baseline

Baseline genomic characteristics, sorted by best overall response (BOR) and mutation load (number of single nucleotide variants). The figure depicts variants detected in 29 genes known to be involved in antigen processing and presentation (not including HLA alleles) (non-mutated genes are not shown) (Kelly and Trowsdale, Immunogenetics, 2019).

*Patient 4 only had post-treatment biopsy available.