



S5 Fig. Oncoplot of genomic alterations in colorectal cancer patients for all 34 mutated genes. The vertically stacked bar plot on the top shows tumor mutational burden (TMB) for each patient with different colors representing different DNA sample types. The horizontal clustered bar plot on the right denotes mutation prevalence for each sample type. ACMP, Axen Cancer Master Panel; ACPI, Axen Cancer Panel 1; AIM, additionally identified mutation; MSI, microsatellite instability; MSI-L, microsatellite instability-low; MSS, microsatellite stable; PD, progressive disease; SNUBH_V1, SNUBH Pan-Cancer Panel version 1; SNUBH_V2, SNUBH Pan-Cancer Panel version 2; TS500, TruSight Oncology 500; VAF, variant allele frequency.