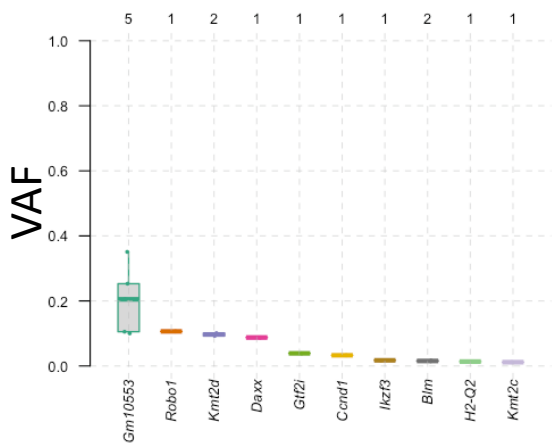
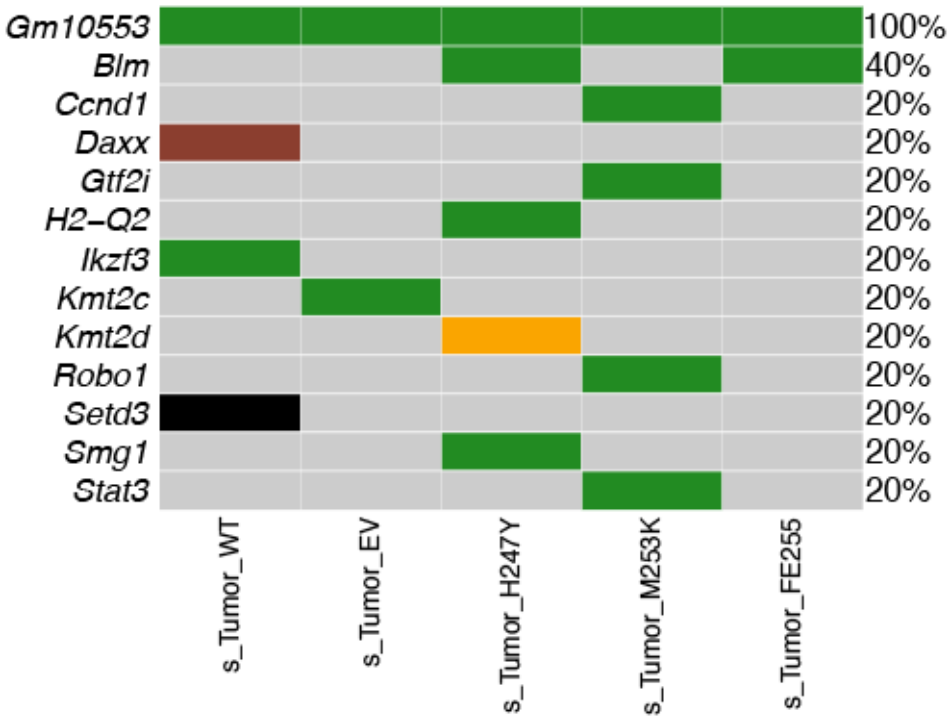


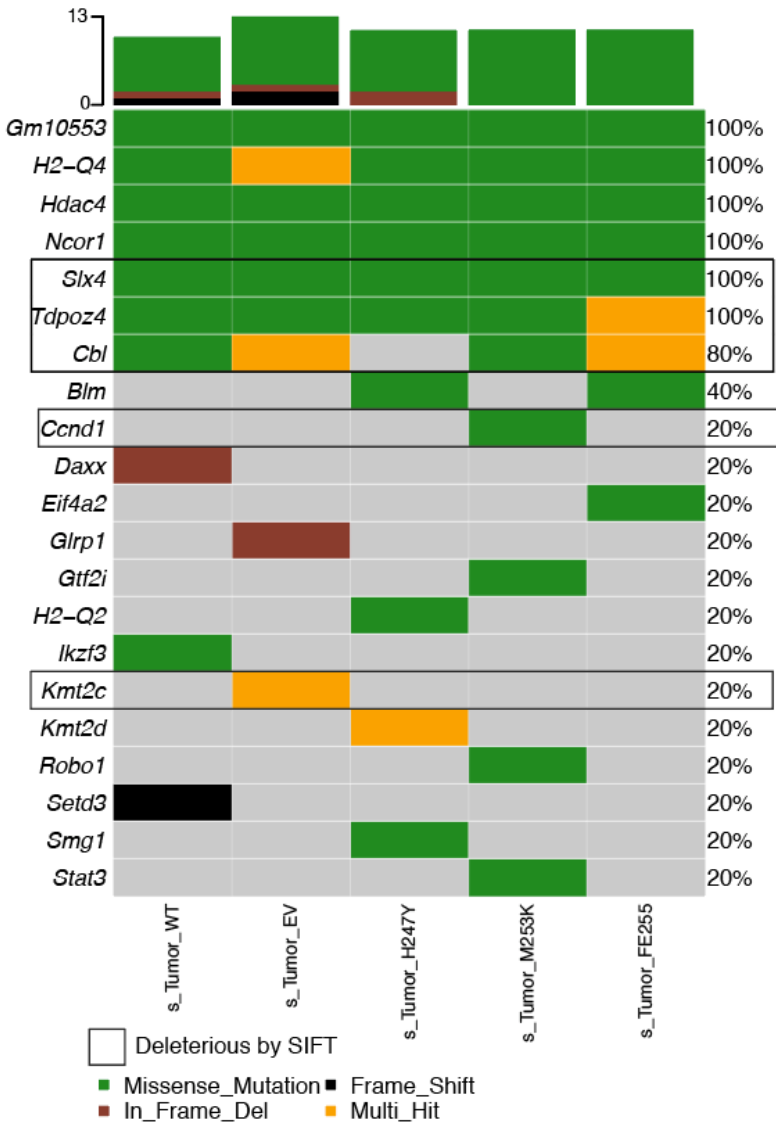
Mutations:

Tumor vs Pooled Normal (from core)



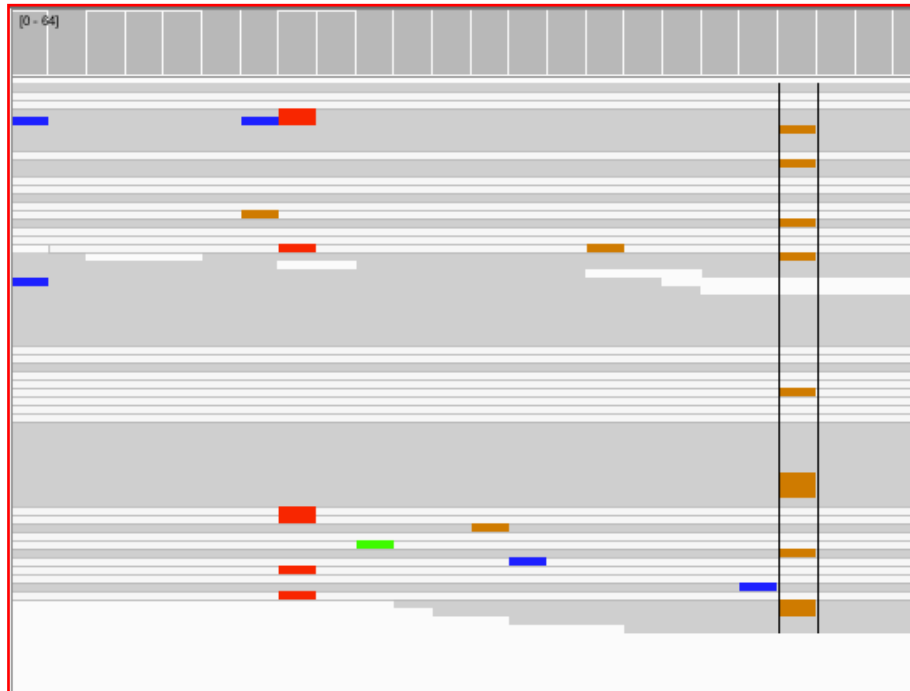
Mutations:

Tumor vs Matched Normal (organoids)

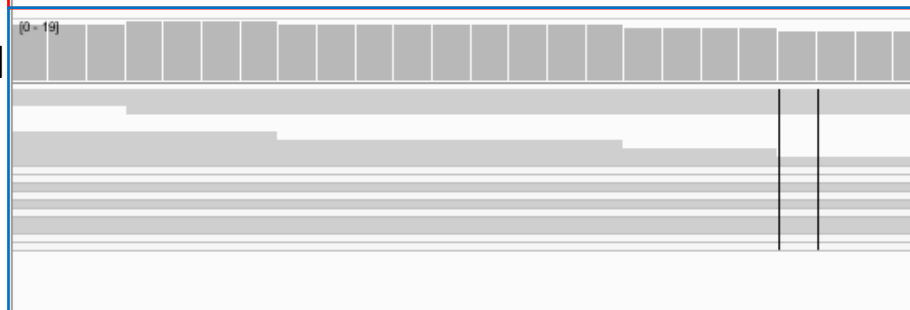


Filtered out mutations present in tumor samples (left) that existed in the pooled normal as well eg: SPOP (next slide)

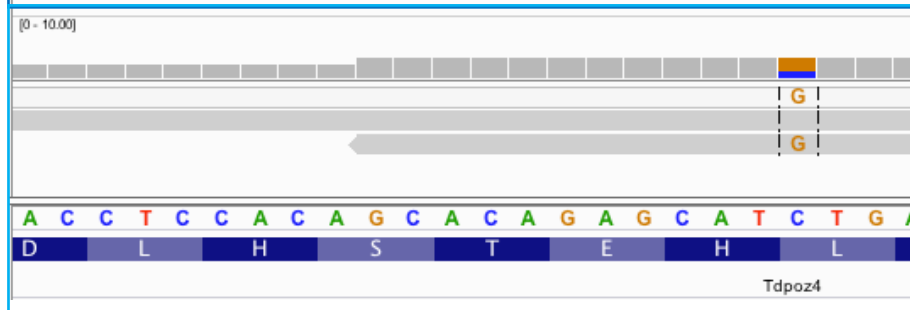
H247_Tumor



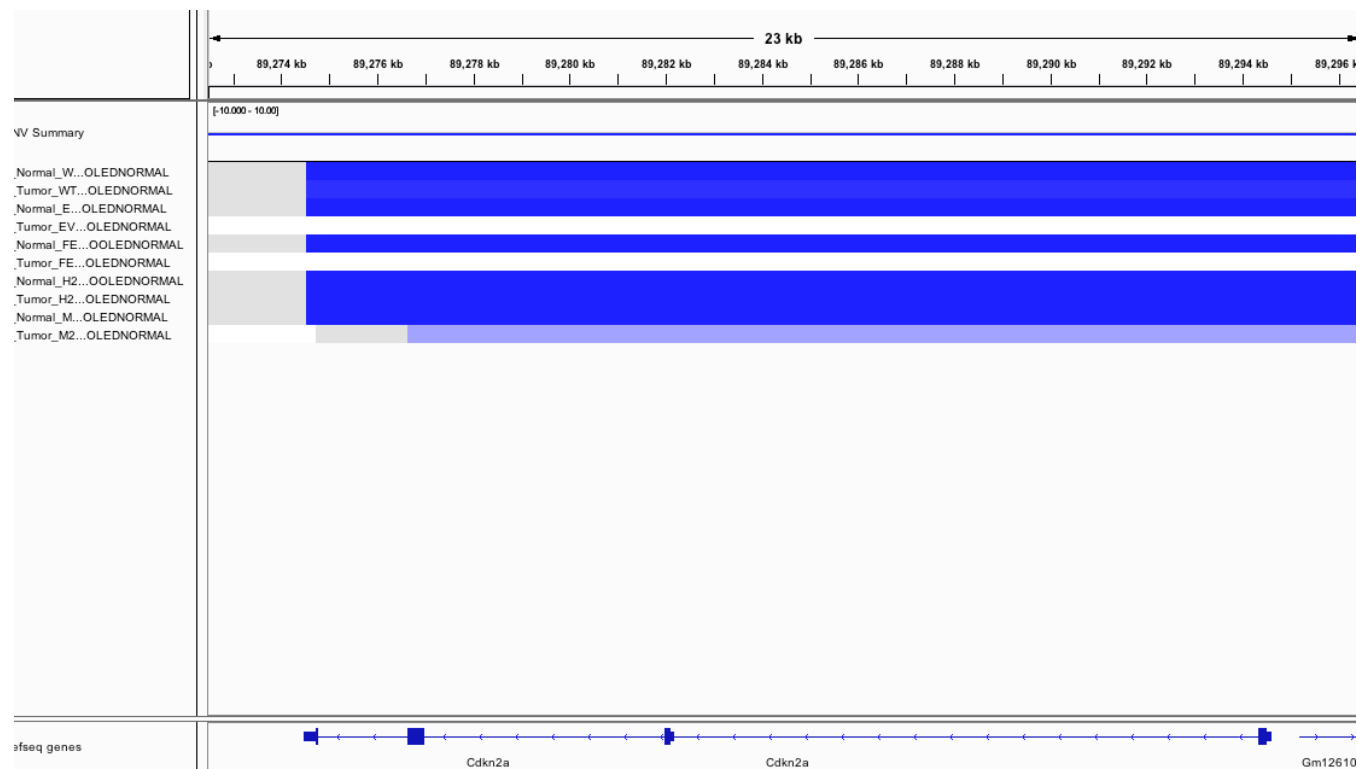
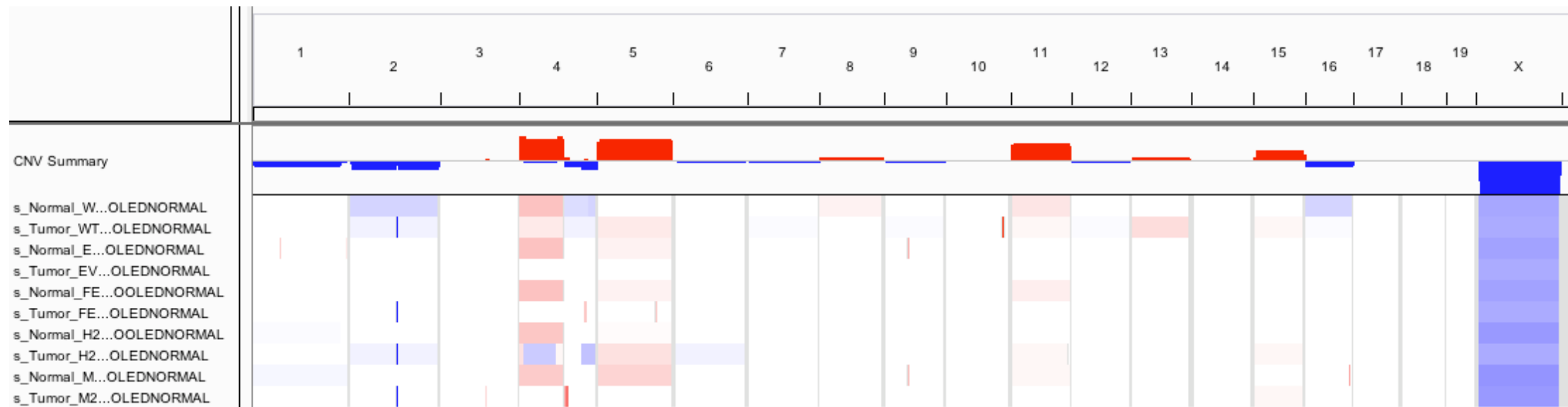
H247_Normal



Pooled Normal



Tdpoz4 (SPOP):
chr3:93797214; p.L307V



Almost all normal and tumor samples have this deletion in *cdkn2a* – potentially noise?



Super focal
amplification of Jun –
usually not the
pattern in other
cohorts?