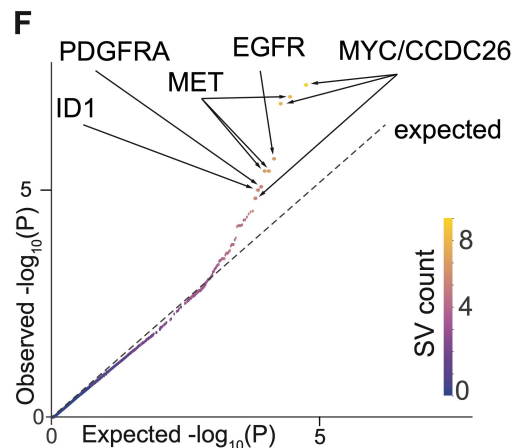
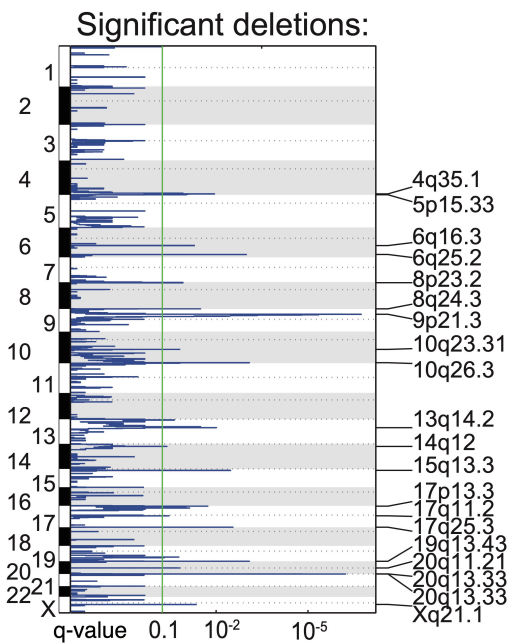
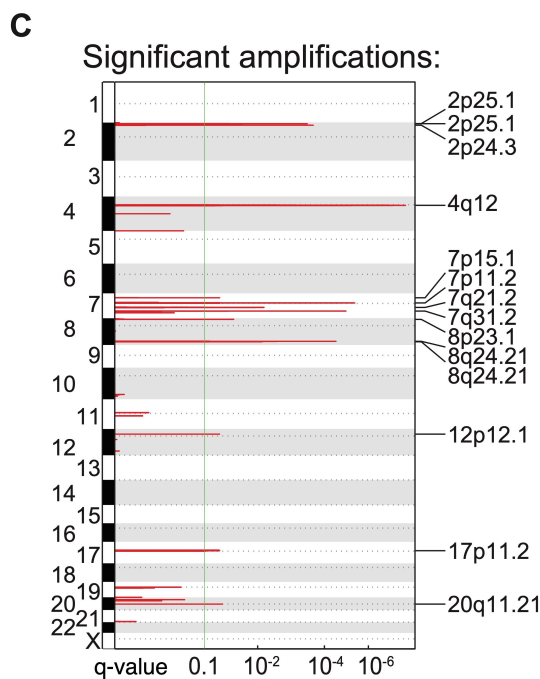
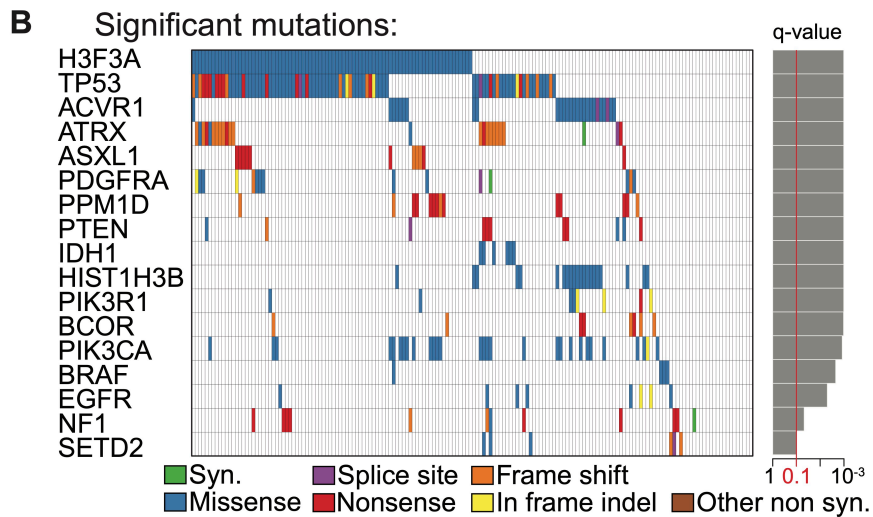
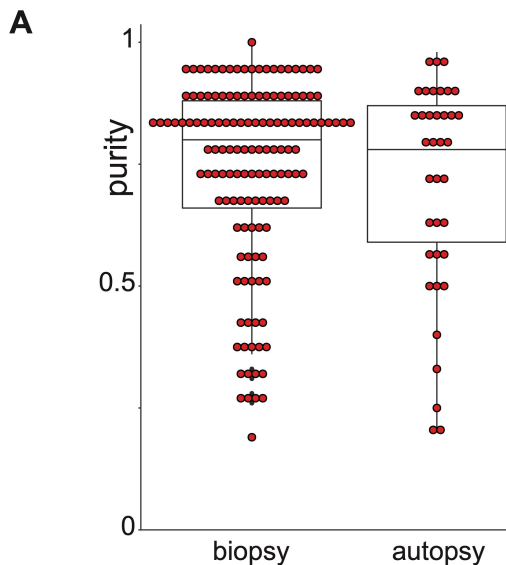
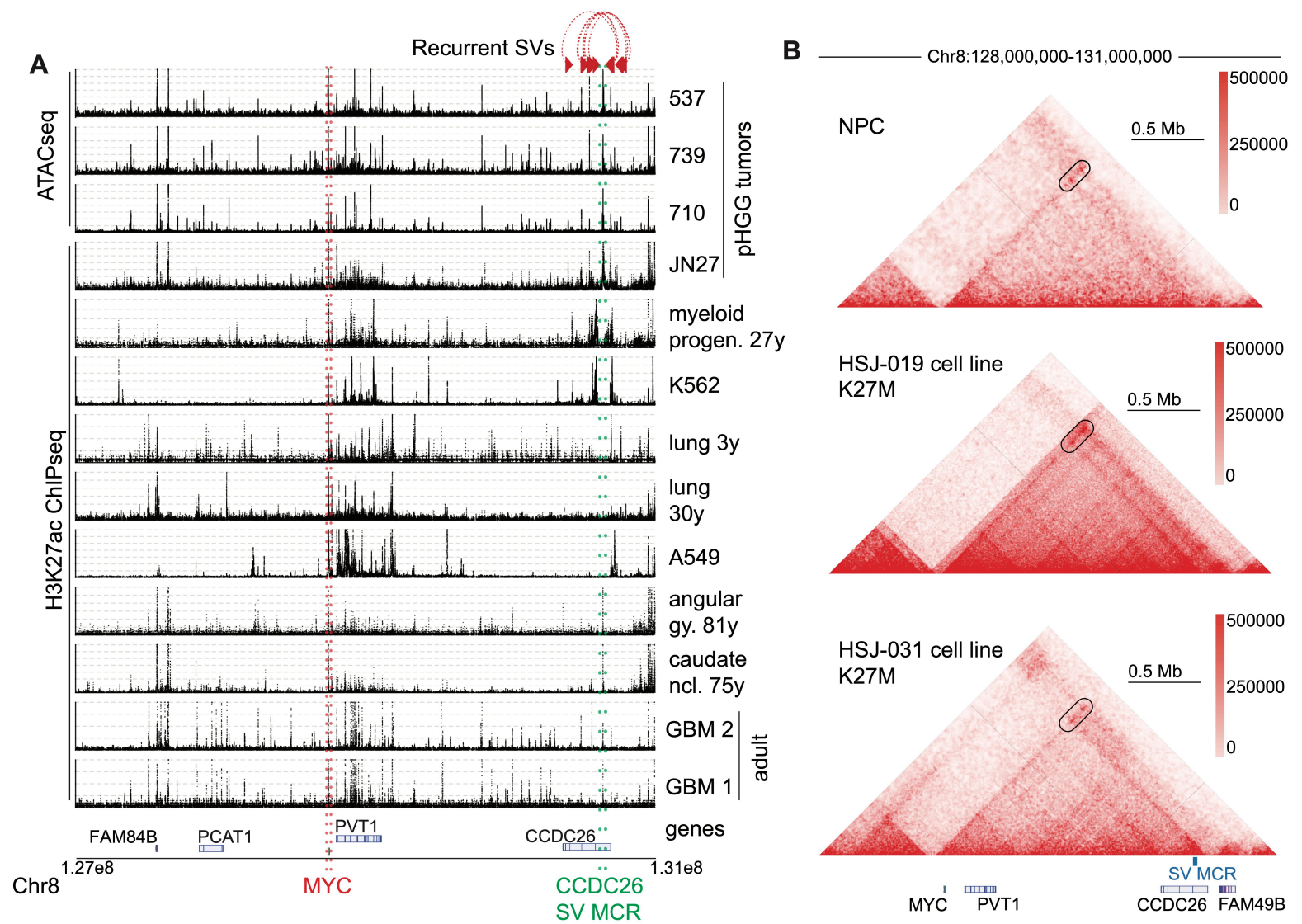
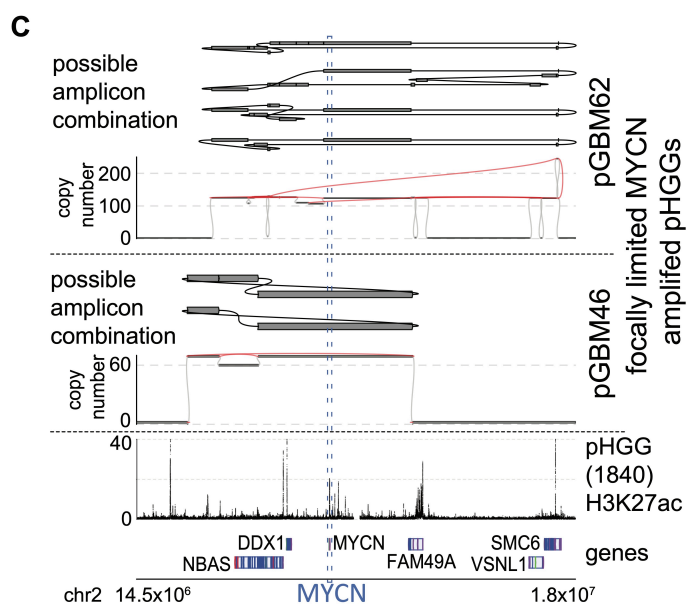
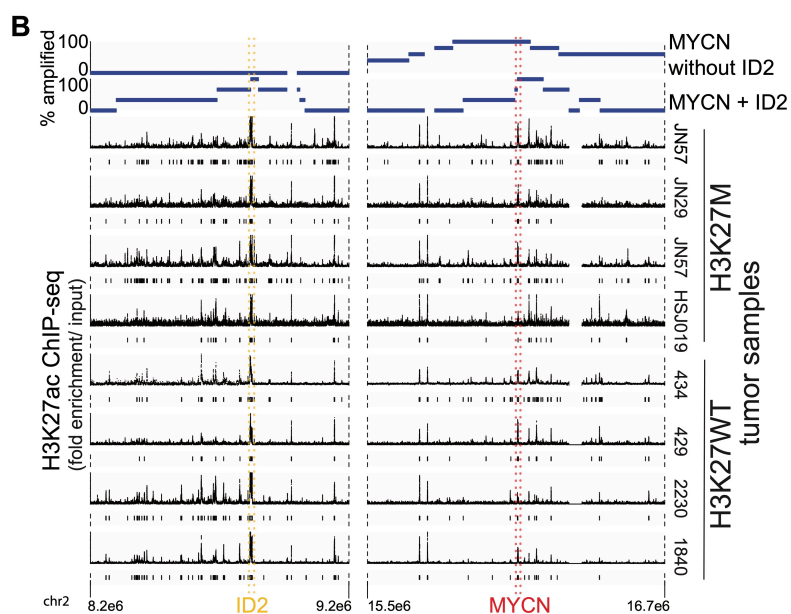
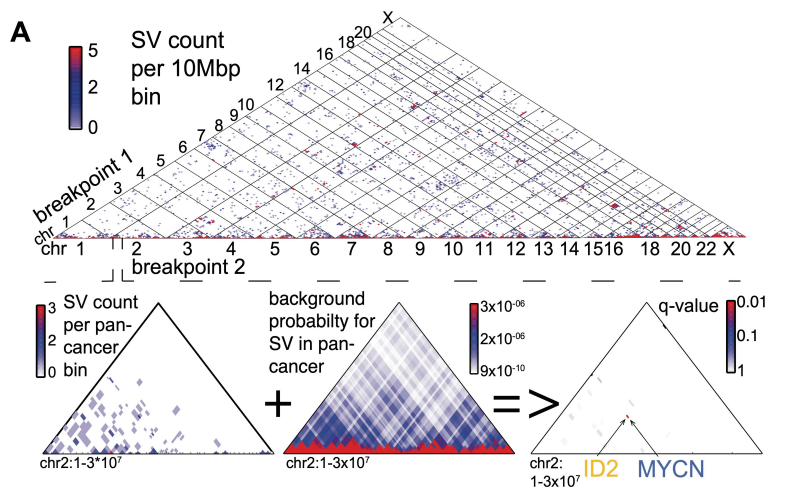




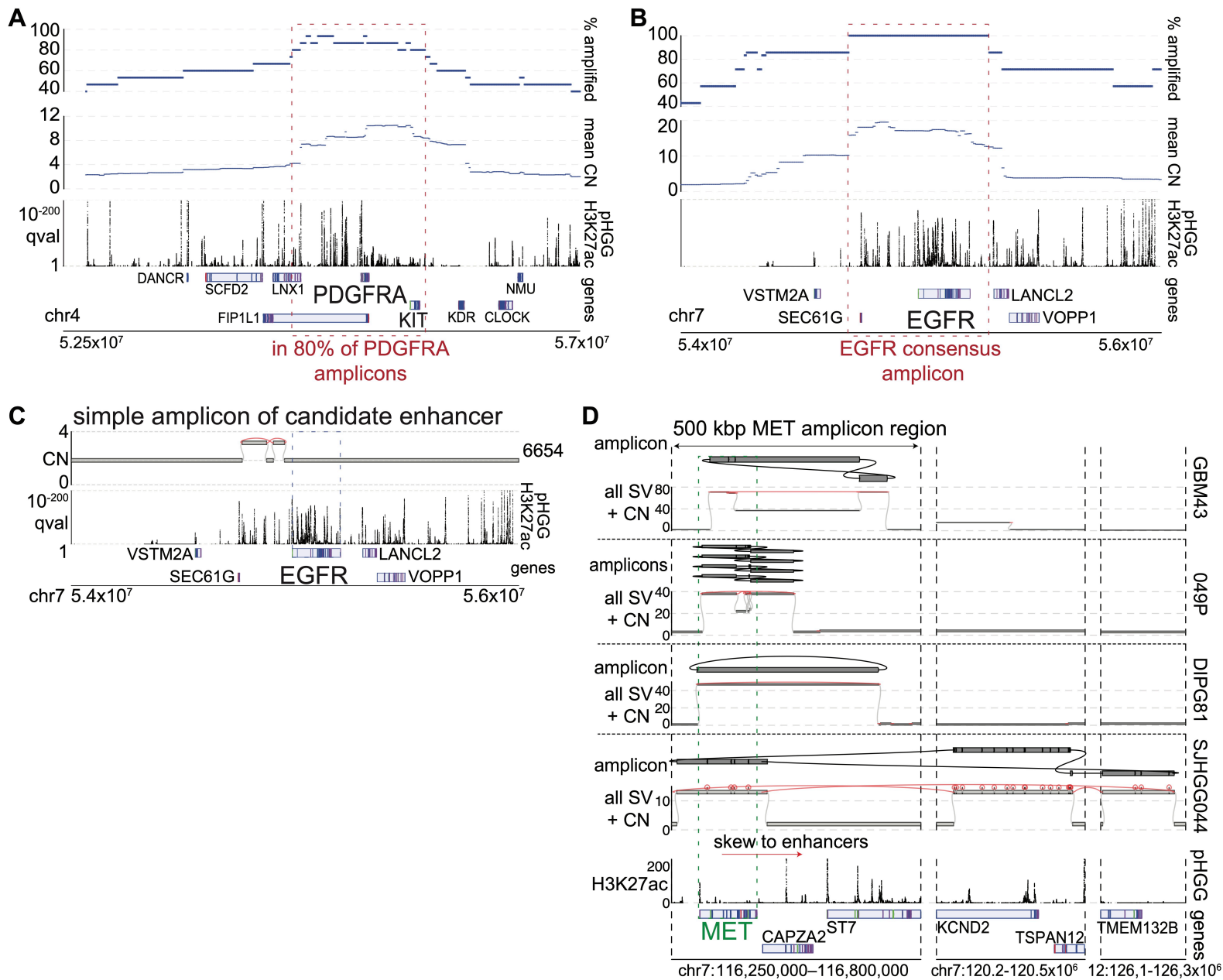
Extended Data Figure 1. Sample characteristics and significantly recurrent variants.



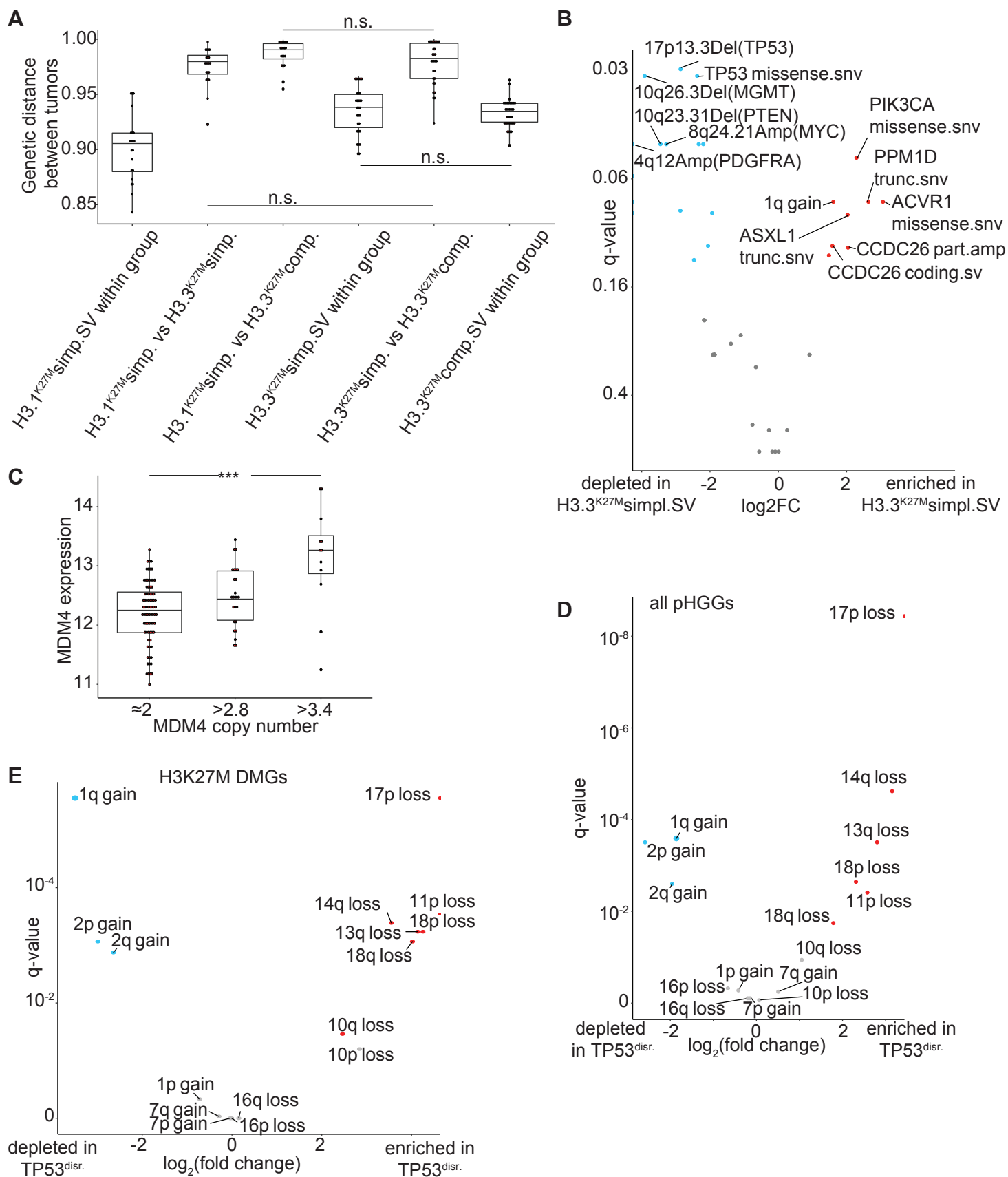
Extended Data Figure 2. Lineage specificity of the enhancer peak in CCDC26.



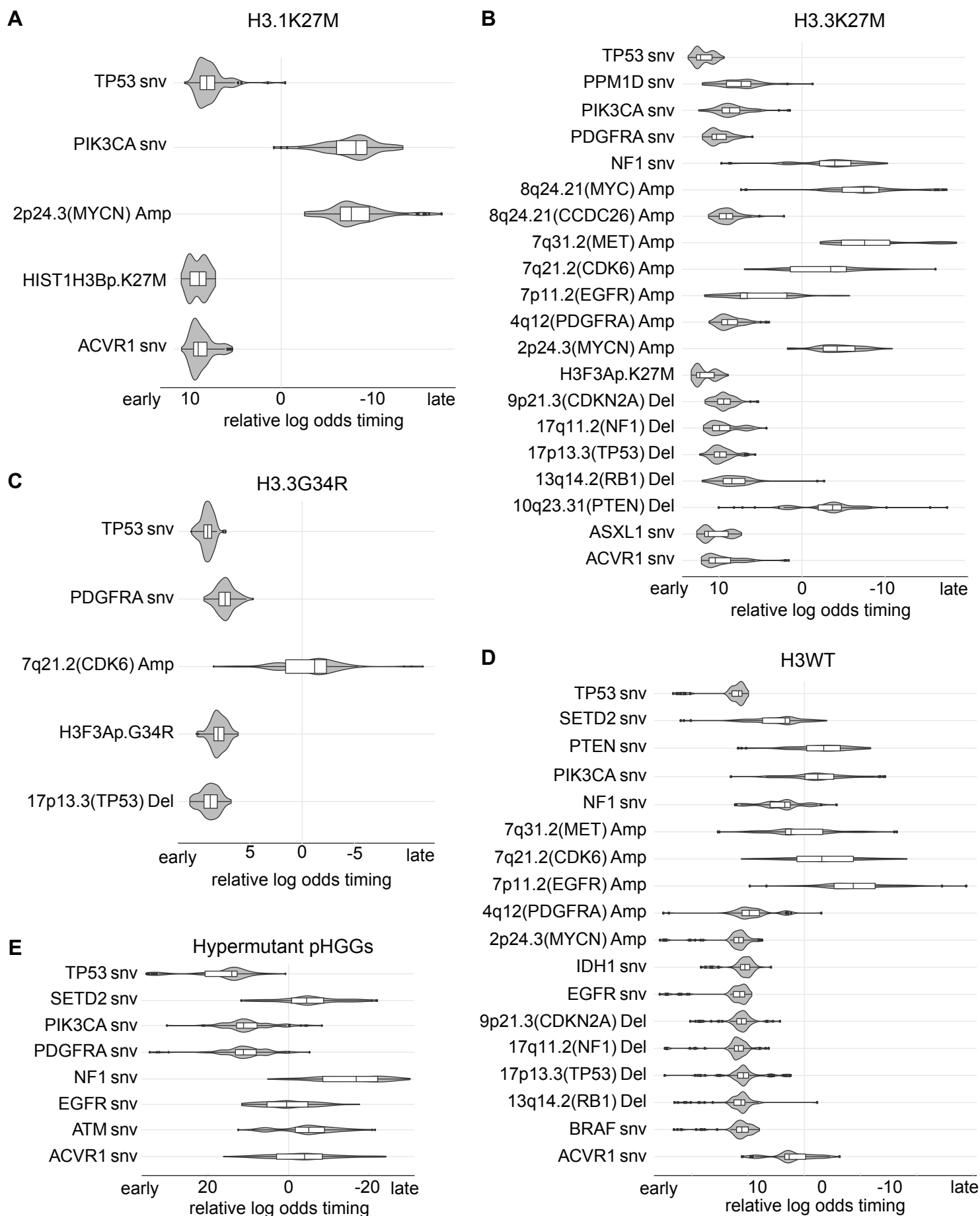
Extended Data Figure 3. Significantly recurrent juxtaposition between MYCN and ID2.



Extended Data Figure 4. Structures of recurrent RTK amplicons.



Extended Data Figure 5. Associations between SV-defined groups.



Extended Data Figure 6. Timing analysis of somatic variant acquisition in histone mutation-defined pHGG subgroups.