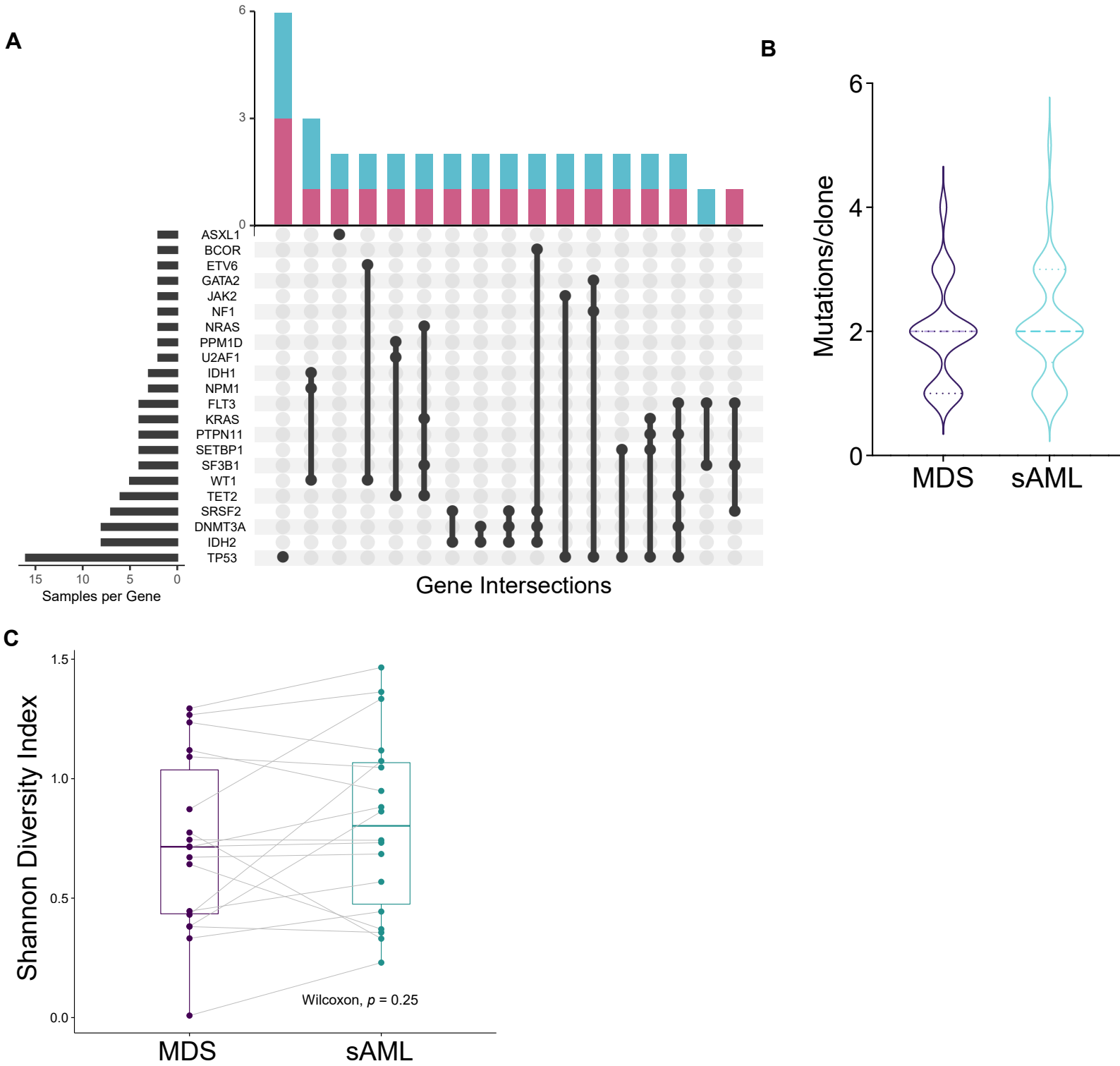## Supplementary Figure 2

Supplementary Figure 2
a, Upset plot depicting co-mutational occurrences in MDS and sAML samples. The number of times a gene was involved in a sample is listed on the left side of the plot. The number of times each gene combination was detected in a sample is listed at the top of the plot. **b**, Violin plot illustrating the number of mutations per clone in MDS and sAML samples. Center line represents median; distal lines represent interquartile range (IQR), 1.5 x IQR. **c**, Box plot portraying differences in clonal diversity for each patient as measured by the Shannon Diversity Index. (p = 0.25, Wilcoxon rank sum test, center line represents median, box represents interquartile range (IQR), whiskers, 1.5 x IQR).