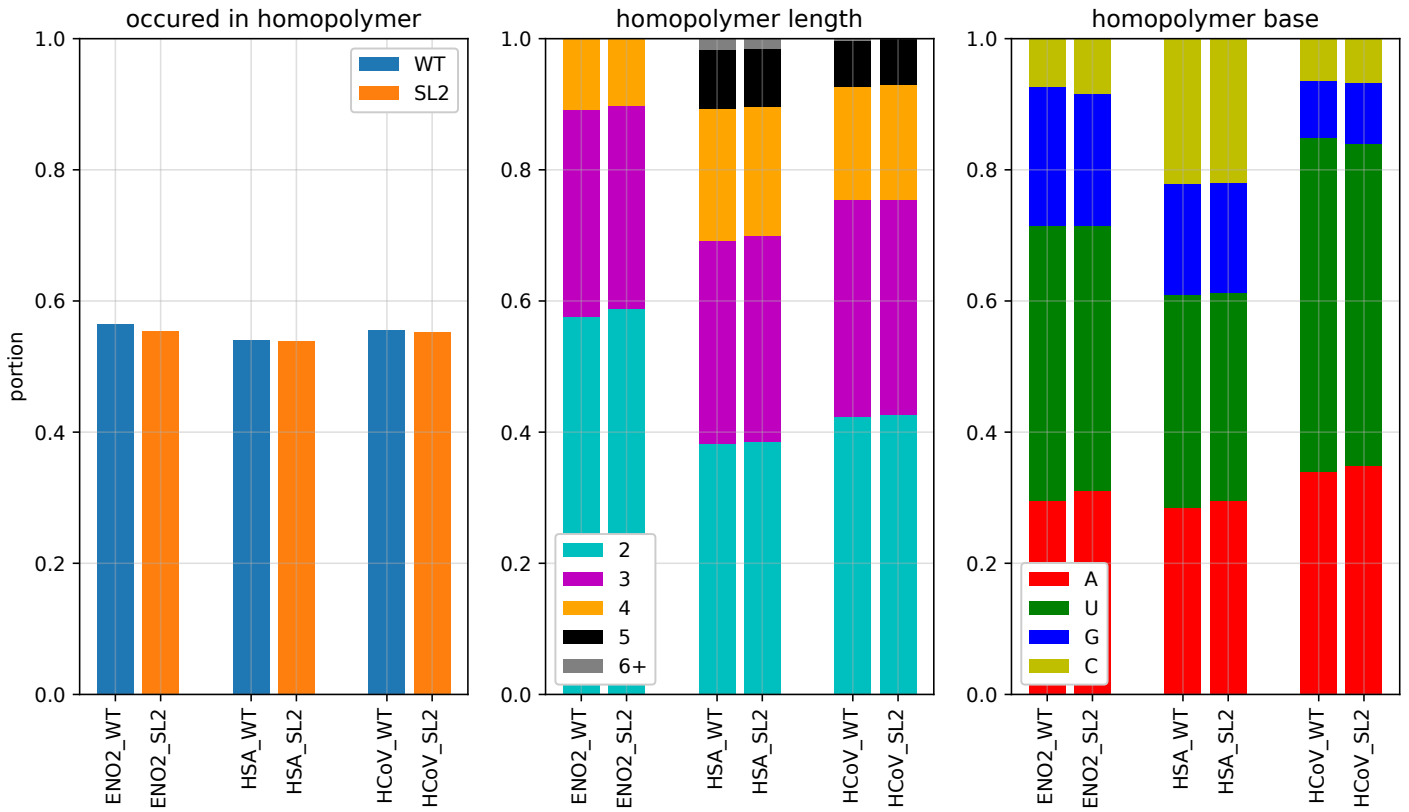


Statistics for deletions of size 1



Supplemental Figure 1 Statistics for all deletions of size 1 in all mappings against the yeast enolase 2, human and HCoV. **Left:** More than 53% of all size 1 deletions occurred in homopolymers (minimum stretch length of 2). **Middle:** A large proportion of the homopolymer stretches that coincide with a deletion are longer than 2 nucleotides. **Right:** A/U stretches appear more prone to causing deletion errors during basecalling, even with consideration of the G/C content of HCoV at 38.3%.