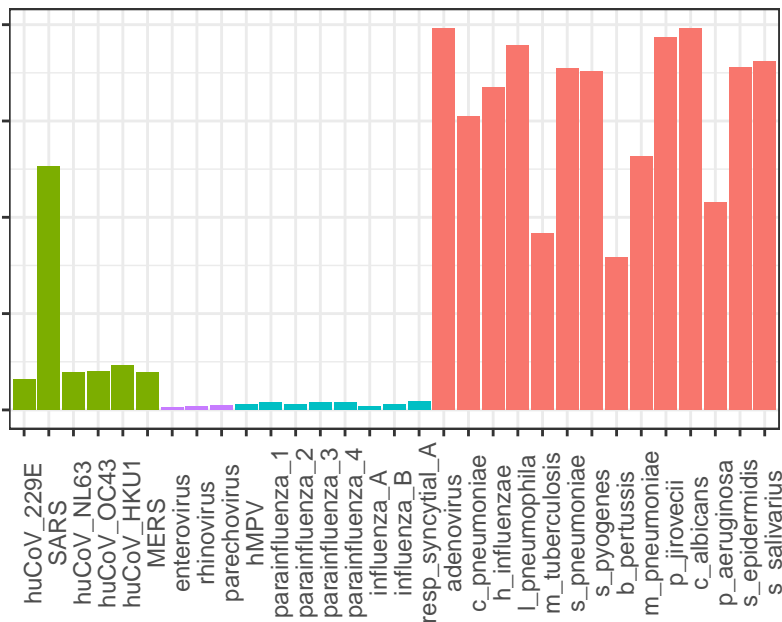


off-target cross-reactivity: DNA

all spacers (n=29880)

% spacers aligned with ...4 mismatches



target_type

