off-target cross-reactivity: DNA all spacers (n=29880) .00 0.75 target_type dsDNA 0.50 human CoVs ssRNA (-) ssRNA (+) 0.25 0.00 |_pneumophila s_pneumoniae pneumoniae parechovirus enterovirus oarainfluenza resp_syncytia parainfluenza parainfluenza oarainfluenza influenza

% spacers aligned with ...4 mismatches