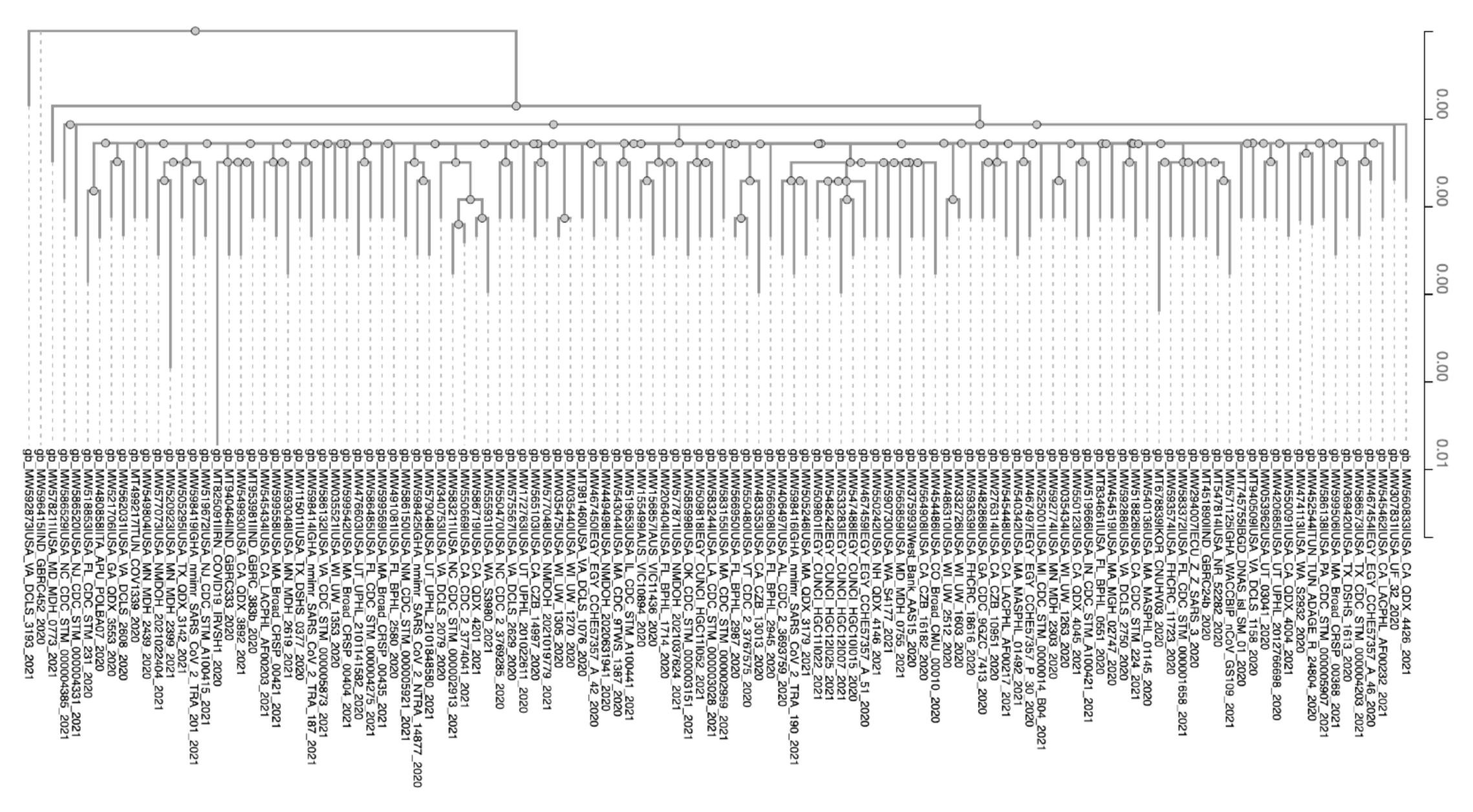
SARS-CoV-2 spike



Information content of the alignments

	WNV NS3	HIV-1 env	SARS-CoV-2 spike
<u>Sequences</u>	19	16	118
Codons	619	288	1273
Tree Length MG94 model, subs/site	0.67	0.20	0.134