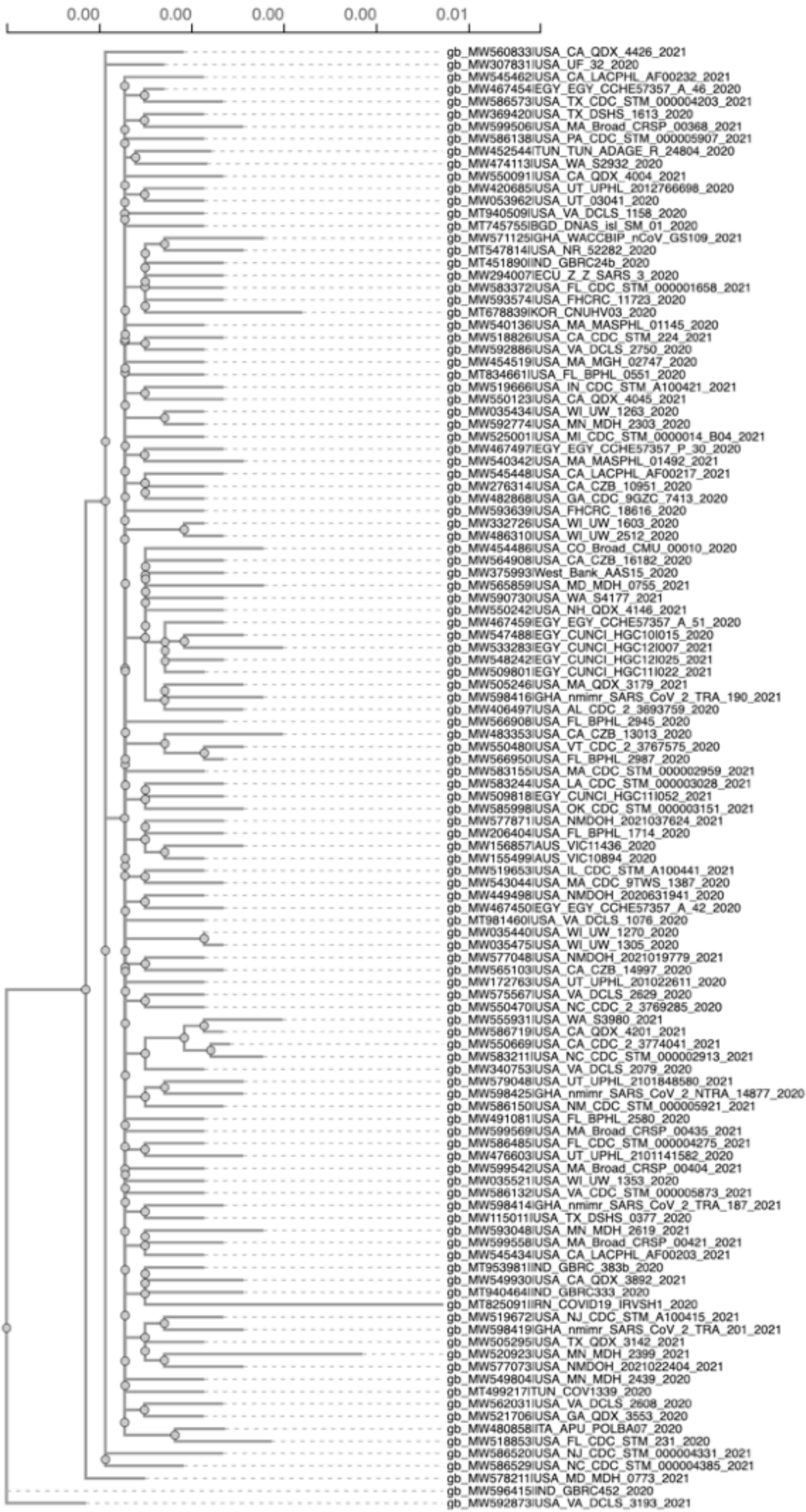


SARS-CoV-2 spike



Information content of the alignments

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