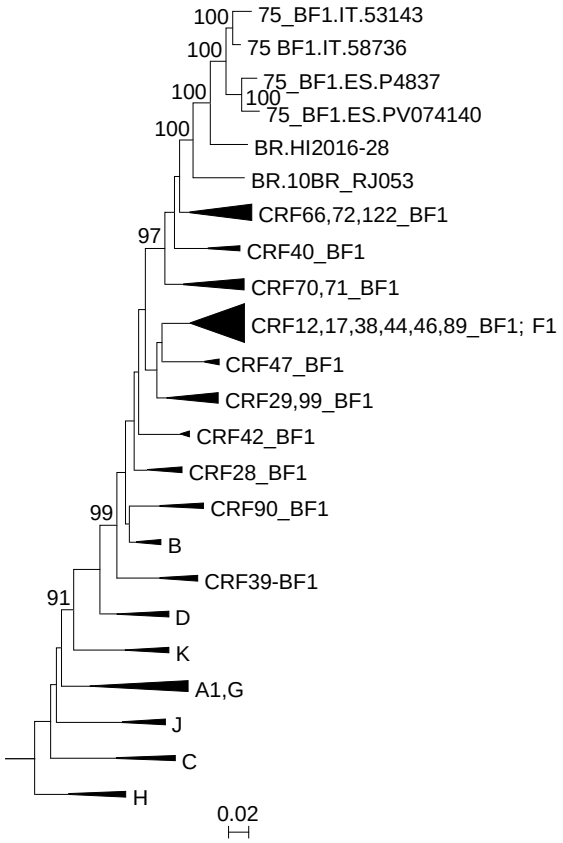
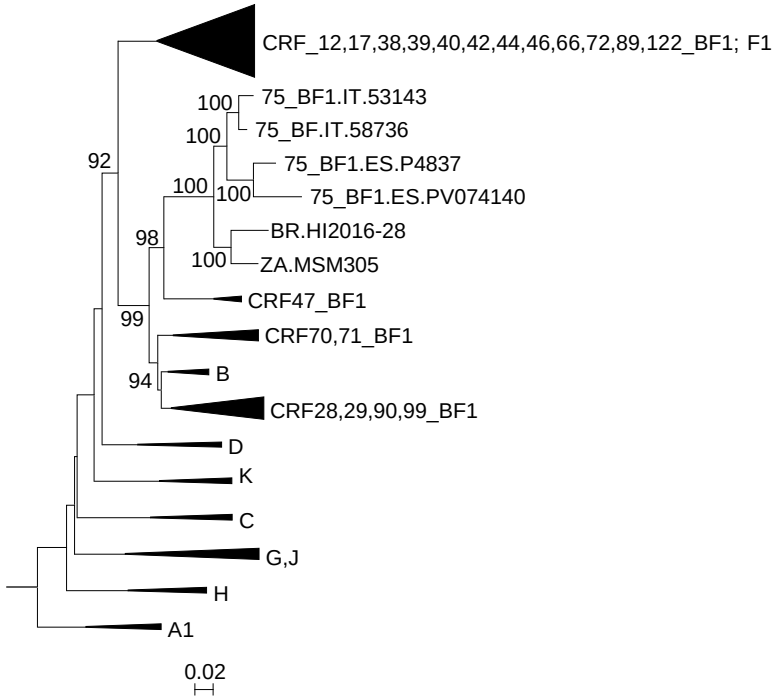
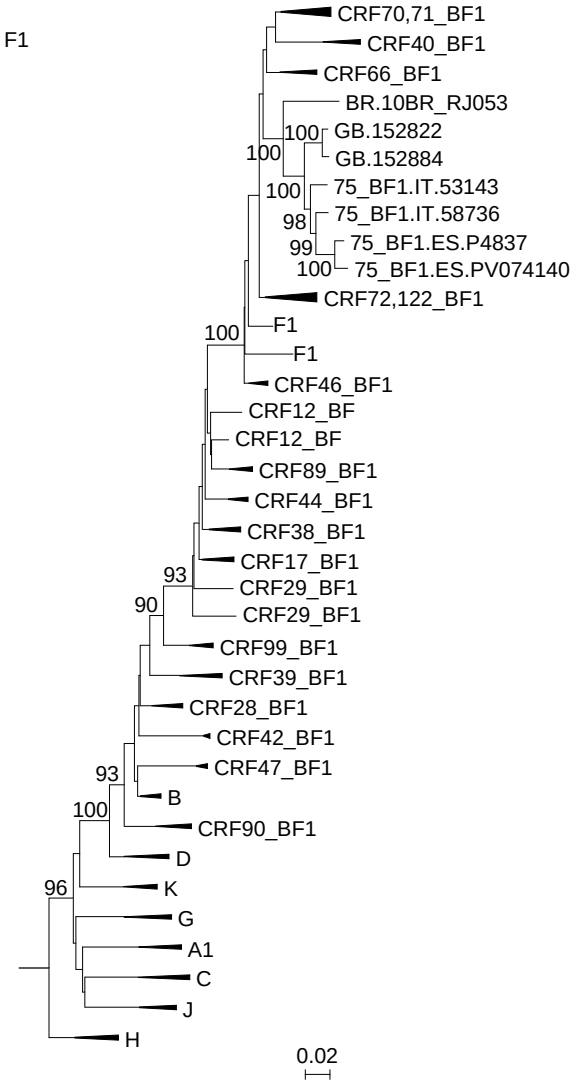
(a) HXB2 pos. 551-1660,2200-2398, 2505-3784,4901-7801



(b) HXB2 pos. 6225-8795



(c) HXB2 pos. 2253-5093



Supplementary Figure 5. Phylogenetic trees showing the relationship of BF1 viruses from (a) Brazil (10BR_RJ053 and HI2016-28), (b) South Africa (MSM305), and (c) United Kingdom (15822 and 15884) with the Spanish-Italian CRF75_BF1 cluster; in (b), the relationship between MSM305 and HI2016-28 is also seen. HXB2 positions of the analyzed segments are shown above each tree. In (a) fragments differing in subtype between 10BR_RJ053 or HI2016-28 and CRF75_BF1 were removed. For viewing purposes, clades of CRF_BF and subtype references are compressed. Trees were constructed with IQ-Tree and only node UFB values ≥90% are shown, except for the nodes of compressed clades, whose UFB values (in all cases >95%) are omitted. Trees were rooted with SIVcpzMB66.