Fig. S3: Neighbor-joining trees based on Pairwise-Fst-values of the gtf gene variants defined as OTUs at 99% identity level, under omission of all OTUs that were unique to a single individual. This calculation was performed to test the impact of putative chimers on the tree topology.

AR = Argentina, BO = Bolivia, CA = California, CH = China, CO = Congo, DE = Germany, GE = Georgia, LO = Louisiana, PH = Philippines, PO = Poland, SO = South Africa, TU = Turkey