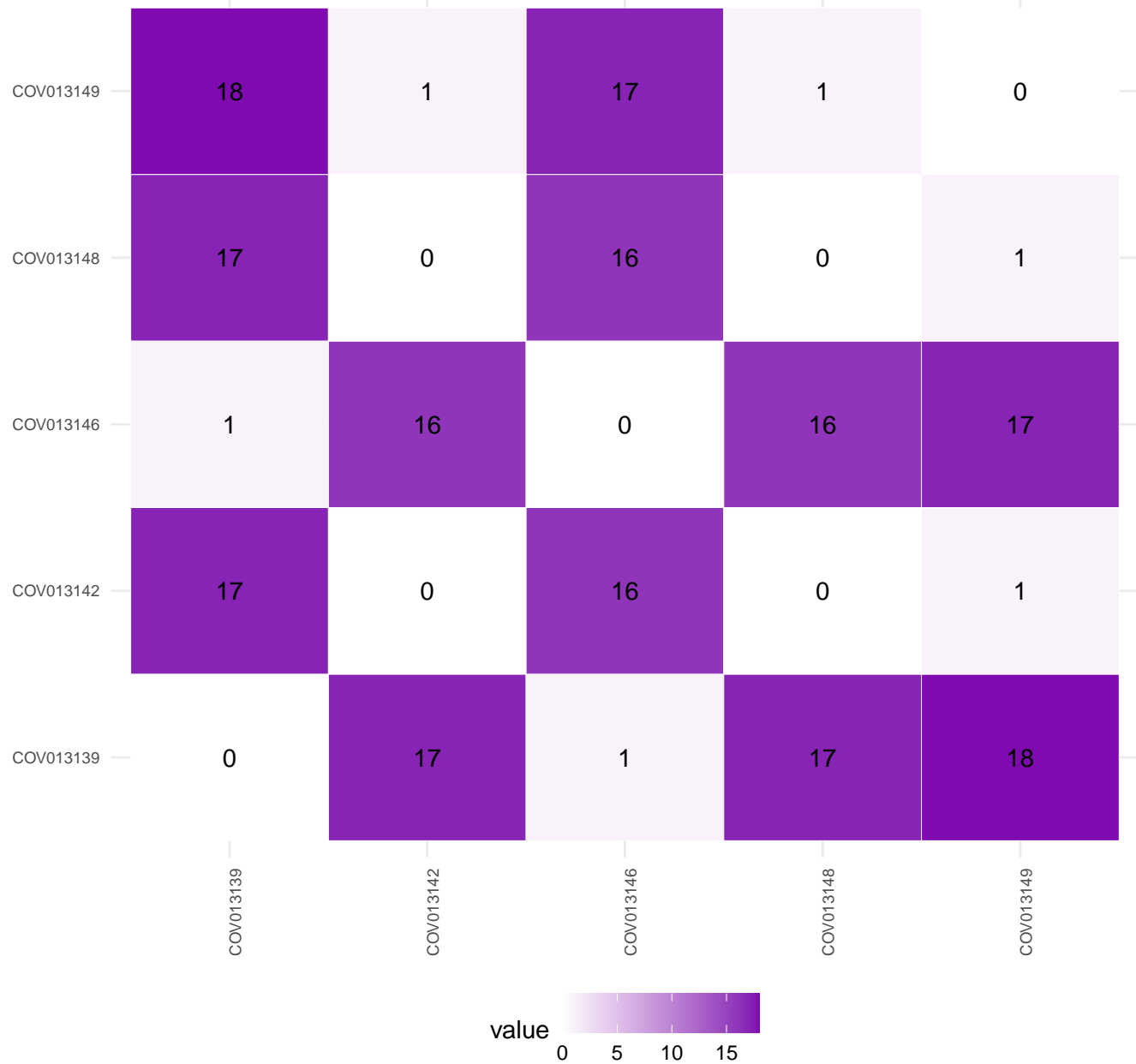
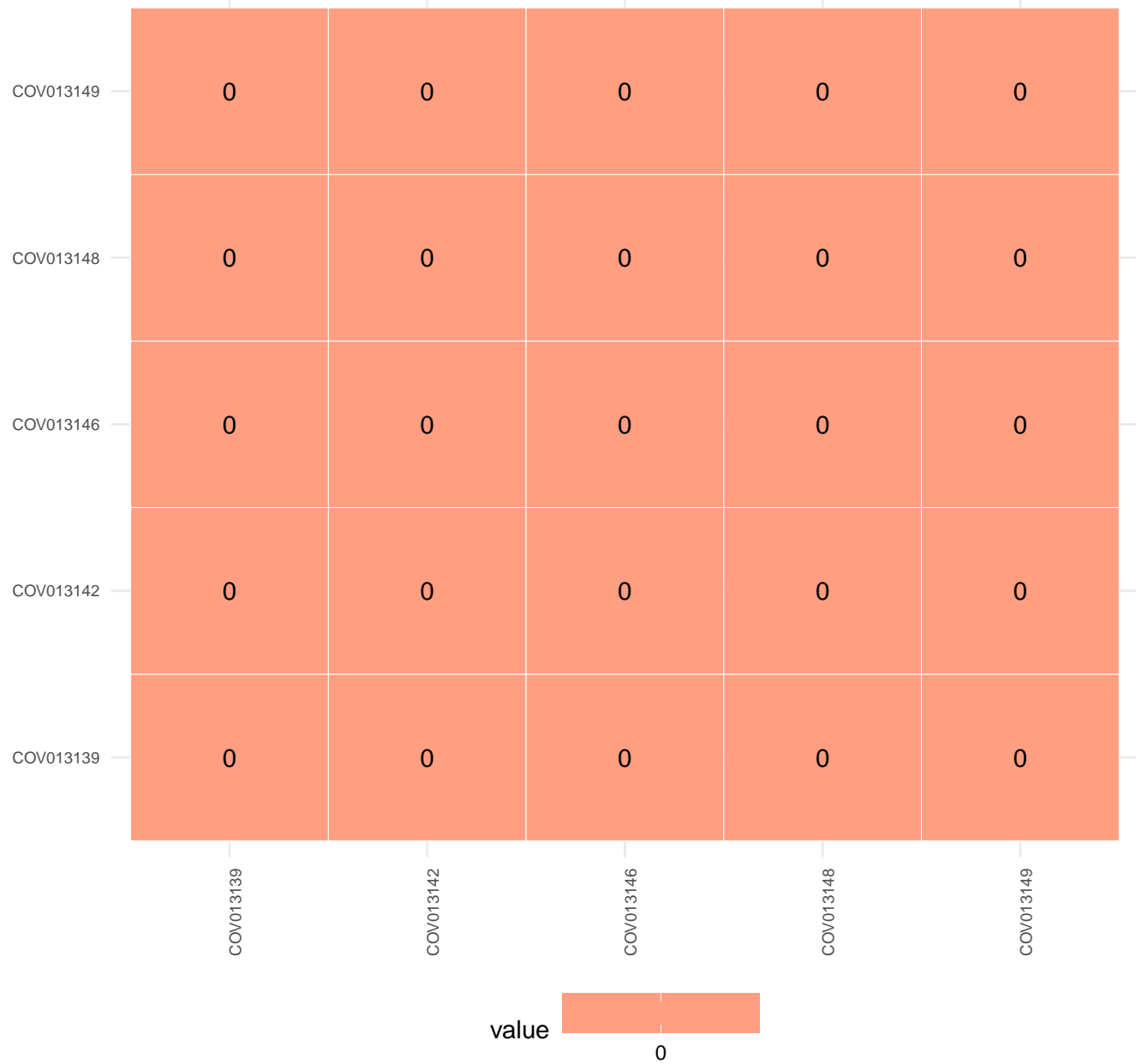


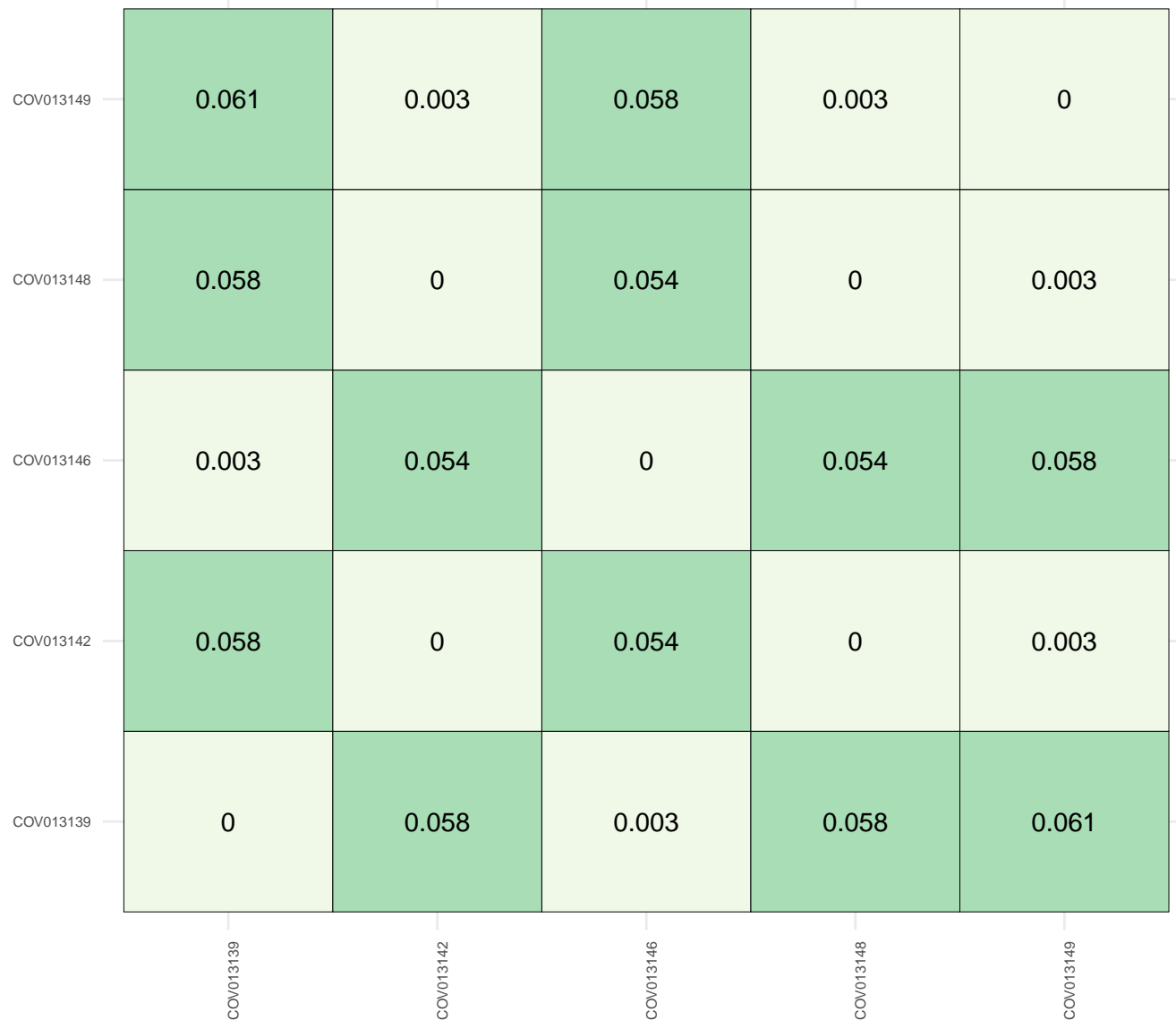
Number of Nucleotide differences



Number of indel differences



Percentage of differences



Sequence
dissimilarity, %



[0,0.05)



[0.05,0.1)