Possible Occupational Infection of Healthcare Workers with Monkeypox Virus, Brazil

Appendix

Appendix Figure 1. Single nucleotide polymorphisms of the MPXV genomes from patient and HCW-1 with respect to the reference genome. Figure was rendered using the Snipit tool (https://github.com/aineniamh/snipit).

Appendix Figure 2. Maximum likelihood phylogenetic tree showing the 2 sequenced genomes in this study (red circles) along with the worldwide Nextclade subsampled dataset of 919 MPXV sequences.
Appendix Figure 3. Phylogenetic tree focusing on the MPXV Clade IIb.