

*Additional file 4:* Phylogenetic trees depicting bootstrap values

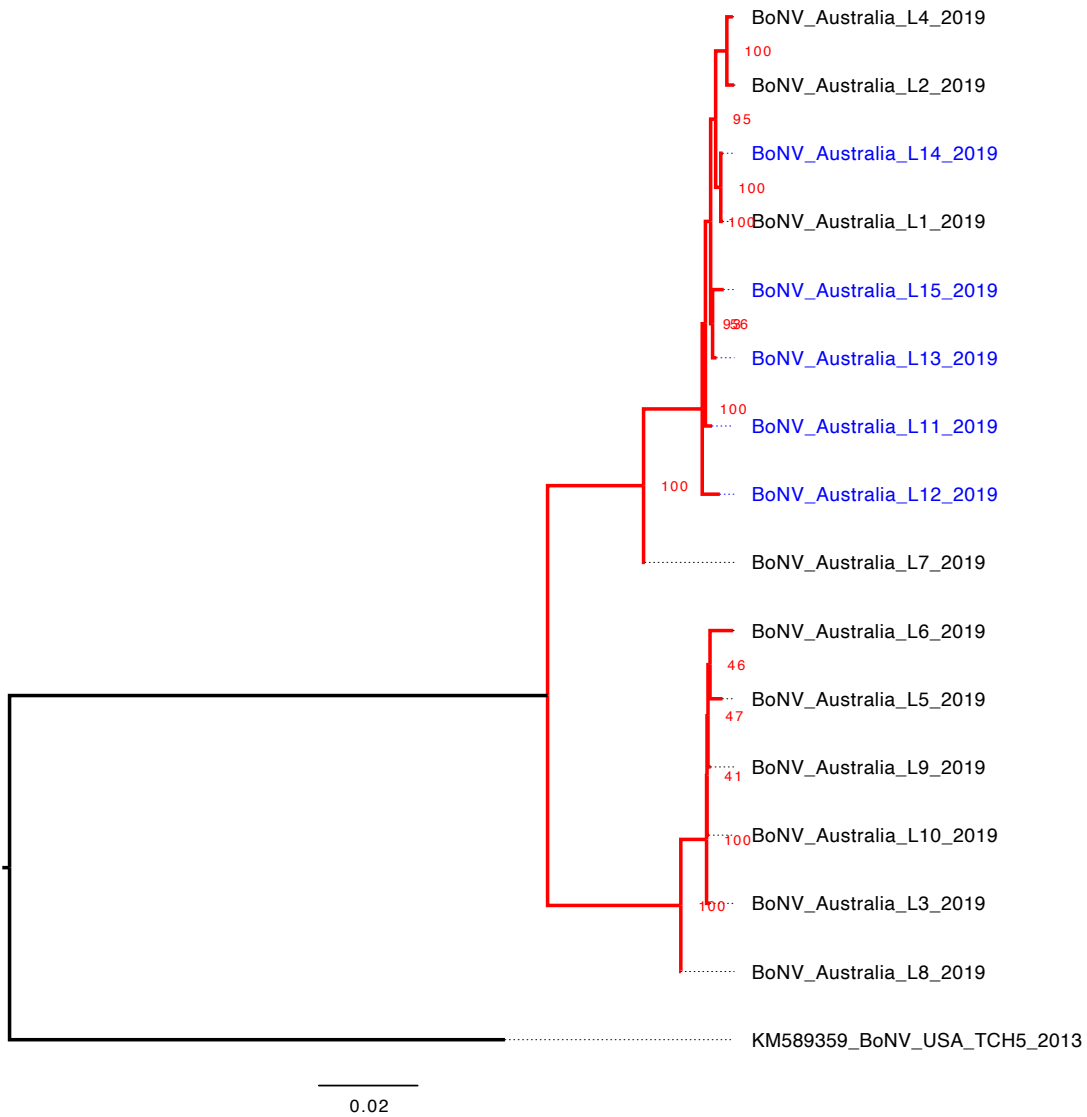


Fig. Maximum likelihood phylogenetic tree of bovine nidovirus (using whole genome).