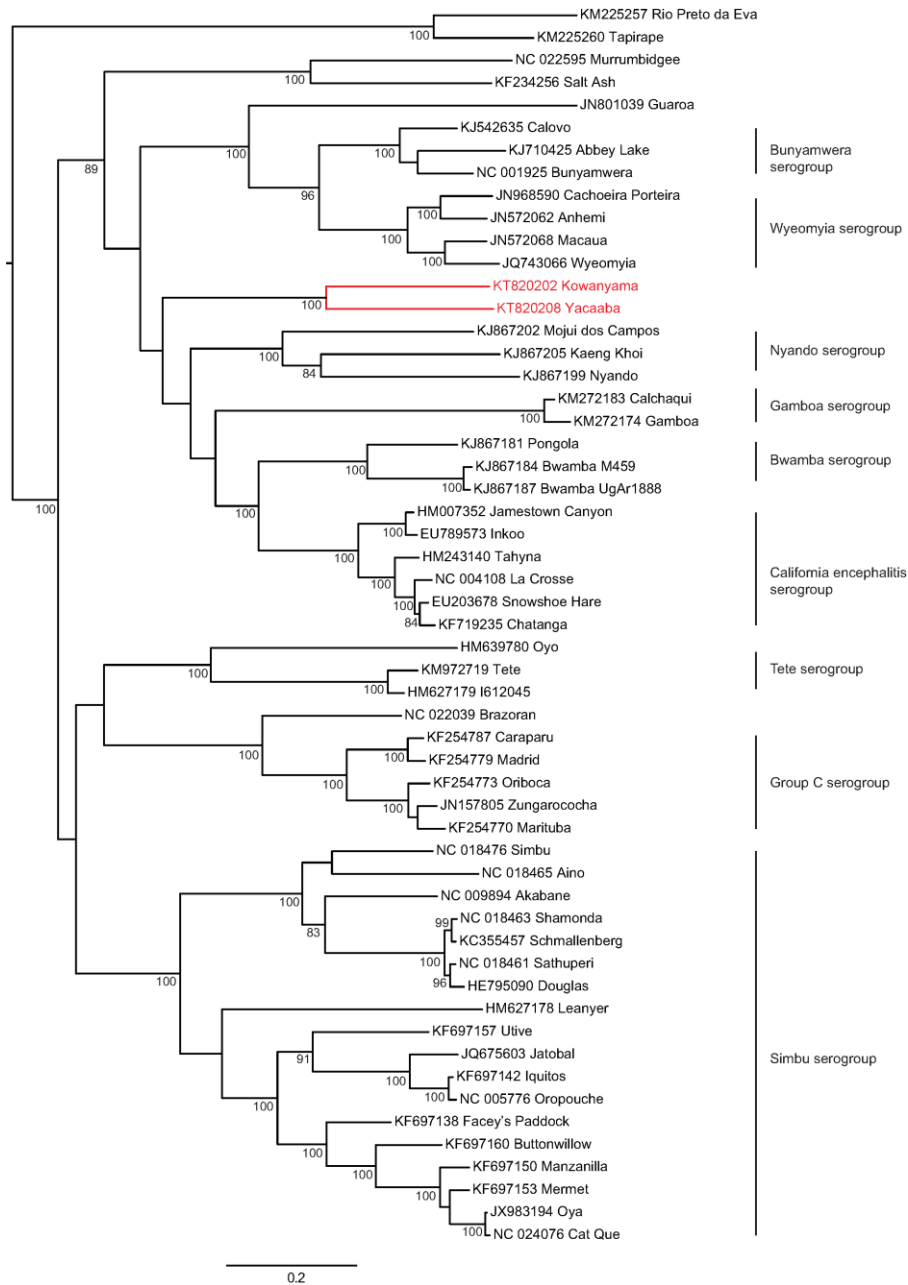


# Genetic Characterization of Archived Bunyaviruses and their Potential for Emergence in Australia

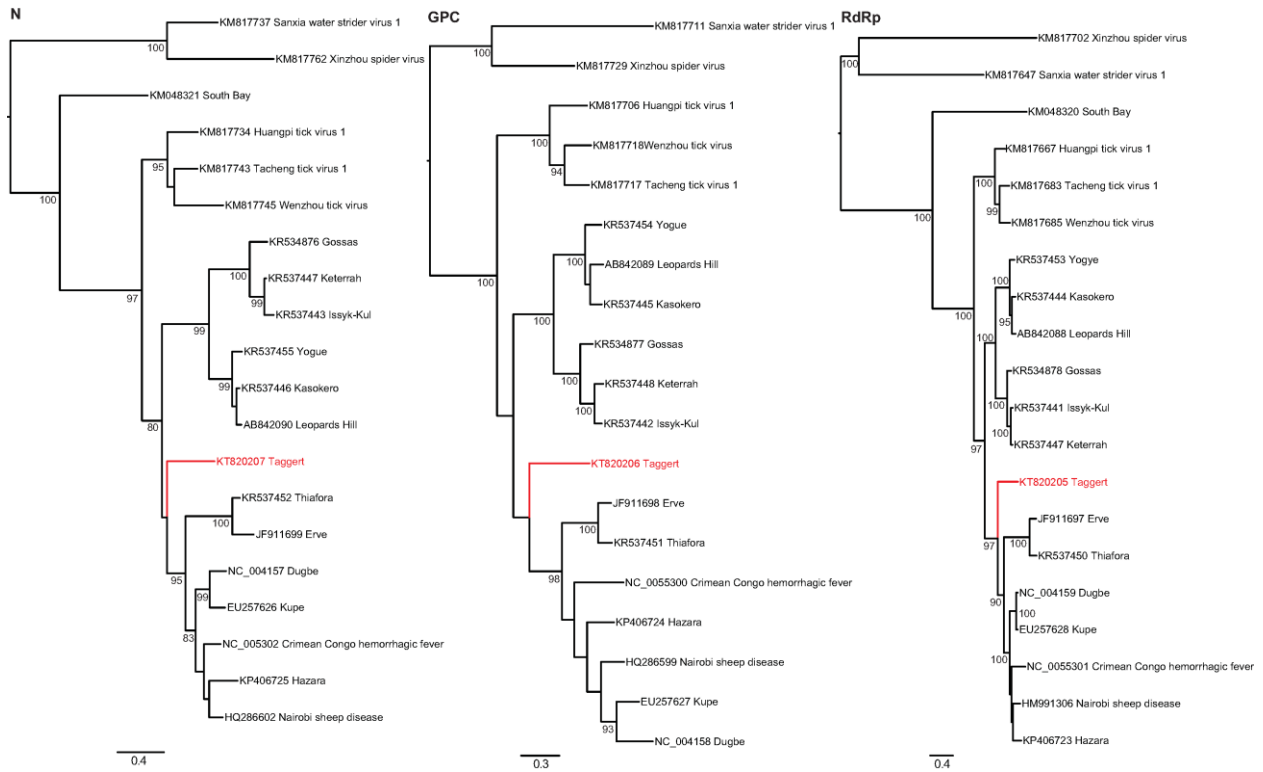
## Technical Appendix



**Technical Appendix Figure 1.** Phylogenetic tree of the orthobunyaviruses based on the S segment ORF. The relationship of KOWV and YACV to other orthobunyaviruses using the predicted ORF sequence with a maximum likelihood model. GenBank accession numbers are shown.



**Technical Appendix Figure 2.** Phylogenetic tree of the orthobunyaviruses based on the L segment ORF. The relationship of KOWV and YACV to other orthobunyaviruses using the predicted ORF sequence with a maximum likelihood model. GenBank accession numbers are shown.



**Technical Appendix Figure 3.** Phylogenetic trees of the nairoviruses based on the complete S, M and L ORFs. The relationship of TAGV to other orthobunyaviruses using the predicted ORF sequence with a maximum likelihood model. GenBank accession numbers are shown.