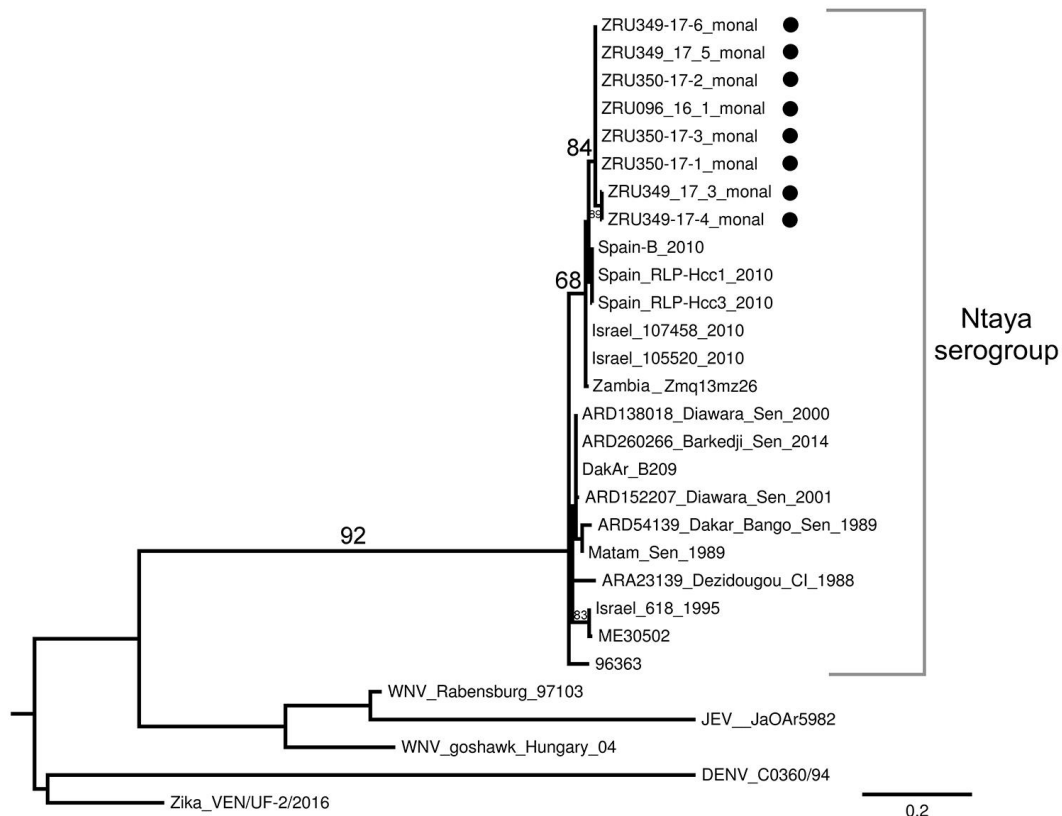


# Bagaza Virus in Himalayan Monal Pheasants, South Africa, 2016–2017

## Appendix

**Appendix Table.** Summary of Bagaza virus cultures used for electron microscopy, using cell line BHK (BSR), in investigation of Bagaza virus in Himalayan monal pheasants (*Lophophorus impejanus*), Pretoria, South Africa, 2017

Sample no.	Passage no.	BHK (BSR) passage no.
ZRU350_17_1	3	47
ZRU350_17_2	3	73
ZRU349_17_6	4	35



**Appendix Figure.** Maximum likelihood phylogram of Bagaza virus isolated in samples from Himalayan monal pheasants, South Africa, 2016–2017. Phylogram represents partial (166 nt) nonstructural coding gene 5 (NS5; taxa = 29). Bootstrap support with values of >60 indicated on branches. Black circles indicate strains sequenced in this study. Scale bar indicates nucleotide substitutions per site.