

Supplemental Table 3 - NCBI accessions of sequences of representative nidoviruses used for phylogenetic analysis

Accession (protein)	Name	Abbreviation (used in Supp. Fig. 5)
ABQ39957.1	Asian leopard cat coronavirus Guangxi/F230/2006	
See KJ541759	Ball python nidovirus	
ABN10838	Bat coronavirus HKU4-1	Bat CoV HKU4
ABN10910	Bat coronavirus HKU9-1	Bat CoV HKU9
ABW87819	Beluga Whale coronavirus SW1	
P0C6V7	Berne virus	
YP_337905	Breda virus	
AAR14192	Equine arteritis virus	EAV
ADN95978.2	Fathead minnow nidovirus	FHMNV
YP_004070193	Feline infectious peritonitis virus	FIPV
AAF82690	Gill-associated virus	GAV
YP_007697629.1	Hana virus strain A4/CI/2004	
NP_073549	Human coronavirus 229E	Human CoV 229E
ABE97136	Human coronavirus NL63	Human CoV NL63
AAT84351	Human coronavirus OC43	Human CoV OC43
NP_066134	Infectious bronchitis virus	IBV
AAA85663 - 4	Lactate dehydrogenase-elevating virus	LDEV
AAA46273 - 4	Lelystad virus	
YP_007697636.1	Meno virus strain E9/CI/2004	
YP_001718610	Miniopterus bat coronavirus HKU8	Bat CoV HKU8
AAU06353	Murine hepatitis virus	
YP_004767305	Nam Dinh virus	
YP_005352862.1	Night-heron coronavirus HKU19	
NP_598309	Porcine epidemic diarrhea virus	PEDV
AAG13725 - 6	Porcine reproductive and respiratory syndrome virus	PRRS
AEU12347.2	Possam nidovirus isolate WPDV	Possun NV
YP_001552234	Rhinolophus bat coronavirus HKU2	Bat CoV HKU2
AAP94757	SARS coronavirus CUHK-AG03	SARS
ABG47077	Scotophilus bat coronavirus 512	Bat CoV 512
NP_742092	Simian hemorrhagic fever virus	SHFV
YP_005352845.1	Sparrow coronavirus HKU17	
YP_002308496.1	Thrush coronavirus HKU12-600	
YP_803213	White bream virus	