

**Supplementary Table S2:** Overview of the coronavirus contigs obtained in positive samples

Location	Sample	Read count	Contigs									
Site 2 (Soysambu Conservancy)	SYRL007	145	Region	Contig 1								
			Range	(425bp)								
Site 3 (Mount Suswa)	SUOHC1001	1067	ID	ORF1ab								
				1-425								
				94.33%								
			Region	Contig 1								
			Range	(5329bp)								
			ID	ORF1ab								
				1-5329								
				98.45%								
	SUOHC1004	8673	Region	Contig 1	Contig 2	Contig 3			Contig 4	Contig 5		
			Range	(11893bp)	(809bp)	(1235bp)			(476bp)	(610bp)		
			ID	ORF1ab	S partial	E partial	M complete	N partial	N partial	N partial	ORFx complete	
				1-11893	1-809	1-39	49-729	740-1235	1-476	1-123	137-367	
	SUOHC1005	52196		93.06%	65.93%	75.00%	94.25%	83.83%	88.59%	84.85%	66.67%	
			Region	Contig 1	Contig 2	Contig 3			Contig 4		Contig 5	Contig 6
			Range	(14647bp)	(847bp)	(571bp)			(1235bp)		(476bp)	(612bp)
			ID	ORF1ab	S partial	NS3B partial	E partial	E partial	M complete	N partial	N partial	N partial
				1-14647	1-847	1-543	524-571	1-39	49-729	740-1235	1-476	1-123
				91.27%	64.89%	74.43%	75.00%	75.00%	94.25%	83.83%	88.59%	84.85%
	SUOHC1012	45	Region	Contig 1								
			Range	(476bp)								
			ID	N partial								
				1-476								
	SUOHC1013	34507		88.59%								
			Region	Contig 1	Contig 2	Contig 3			Contig 4	Contig 5		
			Range	(12065bp)	(525bp)	(1107bp)			(446bp)	(611bp)		
			ID	ORF1ab	NS3B partial	E partial	M complete	N partial	N partial	N partial	ORFx complete	
				1-12065	1-525	1-36	46-726	736-1107	1-446	1-123	137-367	
				93.09%	74.29%	72.73%	93.81%	83.74%	81.76%	81.82%	66.67%	
	SUOHC2001	1037	Region	Contig 1	Contig 2							
			Range	(8048bp)	(781bp)							
			ID	ORF1ab	S partial							
				1-8048	1-781							
	SUOHC2004	171		97.65%	65.42%							
			Region	Contig 1								
			Range	(865bp)								
			ID	ORF1ab								
				1-865								
				98.95%								
	SUOHC3001	22943	Region	Contig 1	Contig 2	Contig 3			Contig 4	Contig 5		
			Range	(14666bp)	(539bp)	(1235bp)			(476bp)	(608bp)		
			ID	ORF1ab	NS3B partial	E partial	M complete	N partial	N partial	N partial	ORFx complete	
				1-14666	1-539	1-39	49-729	740-1235	1-476	1-122	136-366	
	SUOHC3002	11893		91.41%	73.74%	75.00%	94.25%	83.83%	88.59%	87.10%	66.67%	
			Region	Contig 1	Contig 2	Contig 3			Contig 4		Contig 5	
			Range	(8048bp)	(809bp)	(557bp)			(1831bp)		(612bp)	
			ID	ORF1ab	S partial	NS3B partial	E partial	E partial	M complete	N partial	N partial	ORFx complete
				1-8048	1-809	1-548	549-557	1-39	49-729	740-1831	1-123	137-372
				97.76%	65.93%	74.30%	75.00%	75.00%	94.25%	79.40%	84.85%	66.67%
	SUOHC3004	12100	Region	Contig 1	Contig 2	Contig 3	Contig 4	Contig 5				Contig 6
			Range	(11908bp)	(970bp)	(485bp)	(1197bp)	(2274bp)				(443bp)
			ID	ORF1ab	S partial	S partial	S partial	S partial	NS3 complete	E complete	M complete	N partial
				1-11908	1-970	1-485	1-1197	1-353	352-1040	989-1216	1226-1096	1917-2274
				94.57%	62.97%	72.05%	79.35%	79.31%	70.37%	73.61%	94.25%	85.71%
			Region	Contig 1								
			Range	(413bp)								
			ID	ORF1ab								
				1-413								
				96.38%								
	SUOHC3007	226	Region	Contig 1								
			Range	(1133bp)								
			ID	ORF1ab								
				1-1133								
				98.67%								

Identities are provided according to the closely-related virus sequences and based on nucleotide BLAST results. (bp: base pairs).