

Appendix 4. Genbank accession codes of viral sequences produced in this work

Virus	Sample ID	Tissue	Biosample	SRA	Type of file and sequencing	Lenght (nt)		
Lyssavirus rabies	321	CNS	SAMN53524318	SRR36385191	Fastq RNA-seq sequencing full genome	11922		
				SRR36253825	BAM RNA-seq sequencing	11922		
		GIT	SAMN53524319	SRR36253824	BAM RNA-seq sequencing	11922		
				SAMN53795949	SRR36365193	BAM Illumina Viral Surveillance Pannel v2	7863	
		Kidney	SAMN53524320	SRR36384303	Fastq Sanger sequencing Nucleoprotein gene	554		
				SRR36384314	Fastq Sanger sequencing Polymerase gene	218		
				SRR36253822	BAM RNA-seq sequencing	9914		
			SAMN53524321	SRR36384302	Fastq Sanger sequencing Nucleoprotein gene	568		
				SRR36384313	Fastq Sanger sequencing Polymerase gene	216		
	359	Fetus	SAMN53524321	SRR36253821	BAM RNA-seq sequencing	1189		
				SAMN53795950	SRR36365192	BAM Illumina Viral Surveillance Pannel v2	2948	
				SRR36384309	Fastq Sanger sequencing Nucleoprotein gene	576		
			SAMN53524322	SRR36384304	Fastq Sanger sequencing Polymerase gene	219		
				SRR36385190	Fastq RNA-seq sequencing full genome	11906		
	202	CNS	SAMN53524322	SRR36253820	BAM RNA-seq sequencing	11906		
				GIT	SAMN53524323	SRR36253819	BAM RNA-seq sequencing	611
				SRR36384310	Fastq Sanger sequencing Nucleoprotein gene	568		
				SRR36384305	Fastq Sanger sequencing Polymerase gene	219		
				SRR36385189	Fastq RNA-seq sequencing full genome	11927		
			SAMN53524325	SRR36253818	BAM RNA-seq sequencing	11927		
				SRR36253817	BAM RNA-seq sequencing	815		
				SRR36384301	Fastq Sanger sequencing Nucleoprotein gene	576		
				SRR36384308	Fastq Sanger sequencing Polymerase	216		
				SRR36385188	Fastq RNA-seq sequencing full genome	11905		
	27	CNS	SAMN53524328	SRR36384311	Fastq Sanger sequencing Nucleoprotein gene	574		
				SRR36384306	Fastq Sanger sequencing Polymerase gene	219		
				SRR36385187	Fastq RNA-seq sequencing full genome	11905		
	Pool11	CNS	SAMN53524328	SRR36253816	BAM RNA-seq sequencing	11905		
	Pool12	GIT	SAMN53524329	SRR36253815	BAM RNA-seq sequencing	2875		
				SRR36384312	Fastq Sanger sequencing Nucleoprotein gene	570		

	24	Anal Swab	SAMN53524330	SRR36384307 SRR36253823	Fastq Sanger sequencing BAM RNA-seq sequencing	Polymerase gene	218 8555	
Adeno-associated virus	321	CNS	SAMN53697126	SRR36330080	BAM RNA-seq sequencing		421	
		GIT	SAMN53697127	SRR36384661 SRR36330079	Fastq Sanger sequencing BAM RNA-seq sequencing		398 2731	
		Kidney	SAMN53697128	SRR36384660 SRR36330078	Fastq Sanger sequencing BAM RNA-seq sequencing		318 810	
		Fetus	SAMN53697129	SRR36330077	BAM RNA-seq sequencing		527	
		Hepatovirus	Pool12	GIT	SAMN53697130	SRR36330076	BAM RNA-seq sequencing	135
			Pool14	Fetus	SAMN53697131	SRR36330075	BAM RNA-seq sequencing	4204
Human papillomavirus	Pool6	GIT	SAMN53697132	SRR36330074	BAM RNA-seq sequencing		7761	
Mammalian orthoreovirus	321	GIT	SAMN53697133	SRR36332060	BAM RNA-seq sequencing L1 segment		2267	
				SRR36332059	BAM RNA-seq sequencing L2 segment		1673	
				SRR36332058	BAM RNA-seq sequencing L3 segment		2554	
				SRR36332057	BAM RNA-seq sequencing M1 segment		1594	
				SRR36332056	BAM RNA-seq sequencing M2 segment		1348	
				SRR36332055	BAM RNA-seq sequencing M3 segment		1634	
				SRR36332054	BAM RNA-seq sequencing S1 segment		1059	
				SRR36332053	BAM RNA-seq sequencing S2 segment		823	
				SRR36332052	BAM RNA-seq sequencing S3 segment		269	
				SRR36332051	BAM RNA-seq sequencing S4 segment		656	
Roseolovirus	321	CNS	SAMN53697134	SRR36330073	BAM RNA-seq sequencing		338	