



Fig S1. Nested PCR for gG-qPCR_{positive} samples

Nineteen samples (No.8-1, through 8-19) have been initially detected to be positive by gG-qPCR and further confirmed by nested PCR. The amplicon is 359 bp long. No. 8-1 through 8-19 have been loaded. M; 100 bp DNA ladder. Target bands are identified in No.8-14 and 8-17.

Table S1. Sequences of the primers for conventional PCR and sequencing

Primer set	Primer	Sequence (5'-3')
UL27-LAMP _{1st}	UL27-LAMP_Seq_F1	ACGTCATCATGCAGAACTCG
	UL27-LAMP_Seq_R1	GGTTGGAGAGGAAGGAGGAC
UL27-LAMP _{nested}	UL27-LAMP_Seq_F1	ACGTCATCATGCAGAACTCG
	UL27-LAMP_Seq_R2	GGCCCTCGAAGAAGGAGTAG
gG-qPCR _{1st}	gG-qPCR_Seq_F1	GACCCCGCTACTGCTAC
	gG-qPCR_Seq_R1	CCCACCAGGATCTCGTAGTC
gG-qPCR _{nested}	gG-qPCR_Seq_F2	GCCGAYGTGCACAGACAT
	gG-qPCR_Seq_R1	CCCACCAGGATCTCGTAGTC
UL29-qPCR _{1st}	UL29-qPCR_Seq_F1	CAGTAGCGCAGGATCTGGTTG
	UL29-qPCR_Seq_R1	GCGTGATGGACCTCTTYAACA
UL29-qPCR _{nested}	UL29-qPCR_Seq_F2	CAGGATCTGGTTGGCCATGTAG
	UL29-qPCR_Seq_R1	GCGTGATGGACCTCTTYAACA
US5-6 _{1st}	BV_US5-6_PCR_F1	CGTTTCCTCCCGTGGACTTC
	BV_US5-6_PCR_R1	GTCTGGAACGGGTTCTCCAC
US5-6 _{nested}	BV_US5-6_PCR_F2	CTGACCCTGGCCGCCATG
	BV_US5-6_PCR_R2	GGTCCGCTCTCTGCTCCAGCG
US5-6 _{Seq}	BV_US5-6_Seq_addF	TAAGTGGGAGGGGATAAGG
	BV_US5-6_Seq_addR	CCGGCTTTTAAAGGACAAA