

Supporting Information

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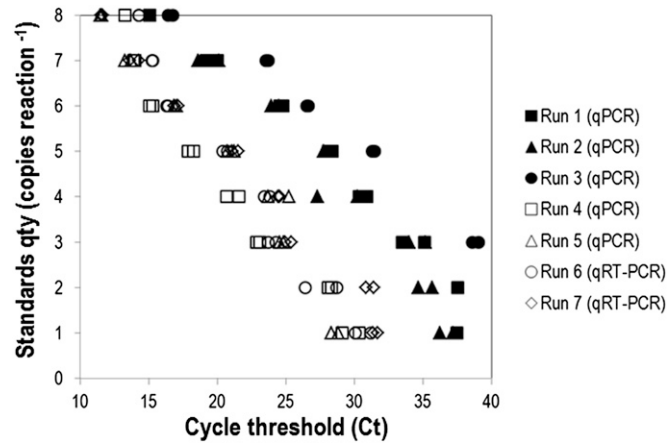


Fig. S1. Standard quantity compared with cycle threshold of quantitative PCR and quantitative RT-PCR analyses of LaCopCV prevalence, viral load, and transcription. Regression statistics are provided in Table S1.

Table S1. Linear regression statistics for quantitative PCR (qPCR) and quantitative RT-PCR (qRT-PCR) runs examining LaCopCV prevalence, viral load, and transcription

Run no.	Objective	R^2	m	b
1	qPCR	0.97	-0.28	12.6
2	qPCR	0.94	-0.32	13.5
3	qPCR	0.98	-0.23	12.1
4	qPCR	0.96	-0.38	12.2
5	qPCR	0.96	-0.37	12.3
6	qRT-PCR	0.97	-0.40	13.1
7	qRT-PCR	0.98	-0.32	11.6

Regression fit with the model $y = mx + b$, where m is the slope and b is the intercept.