

Table 1: Primer sequences and description of the genomic region targeted for qPCR.

Primer	Sequence	Description
SeIV-1q	F: 5'- TGTGAAGTTAGACACGCATGGAA-3' R: 5'-CGACTTGTGCTACTCTCTTCATCAA-3'	Amplifies a 97-bp fragment in the RNA-dependent RNA polymerase (RdRp) region from SeIV-1.
SeIV-2q	F :5'-CCGCTCGCTTATTGAAACGT-3' R: 5'-CATGAGACAGCTGGAATTGGAA-3'	Amplifies a 78-bp fragment in the RNA-dependent RNA polymerase (RdRp) region from SeIV-2.
qATPsynthase	F: 5'-GTTGCTGGTCTGGTGGGATT-3' R: 5'-AGGCCTCAGACACCATTGAAA-3'	Amplifies a 72-bp fragment in ATP-synthase subunit C gene from <i>S. exigua</i> .

Carballo, A.; Murillo, R.; Jakubowska, A.; Herrero, S.; Williams, T.; Caballero, P. Co-infection with iflaviruses influences the insecticidal properties of *Spodoptera exigua* multiple nucleopolyhedrovirus occlusion bodies: Implications for the production and biosecurity of baculovirus insecticides. *PLoS One* 2017, 12, e0177301:1-14. <https://doi.org/10.1371/journal.pone.0177301.t001>