

Supplemental figure: S1. Logarithm of copy number of HearSNPV and HearMNPV genomes determined by qPCR of progeny occlusion bodies.

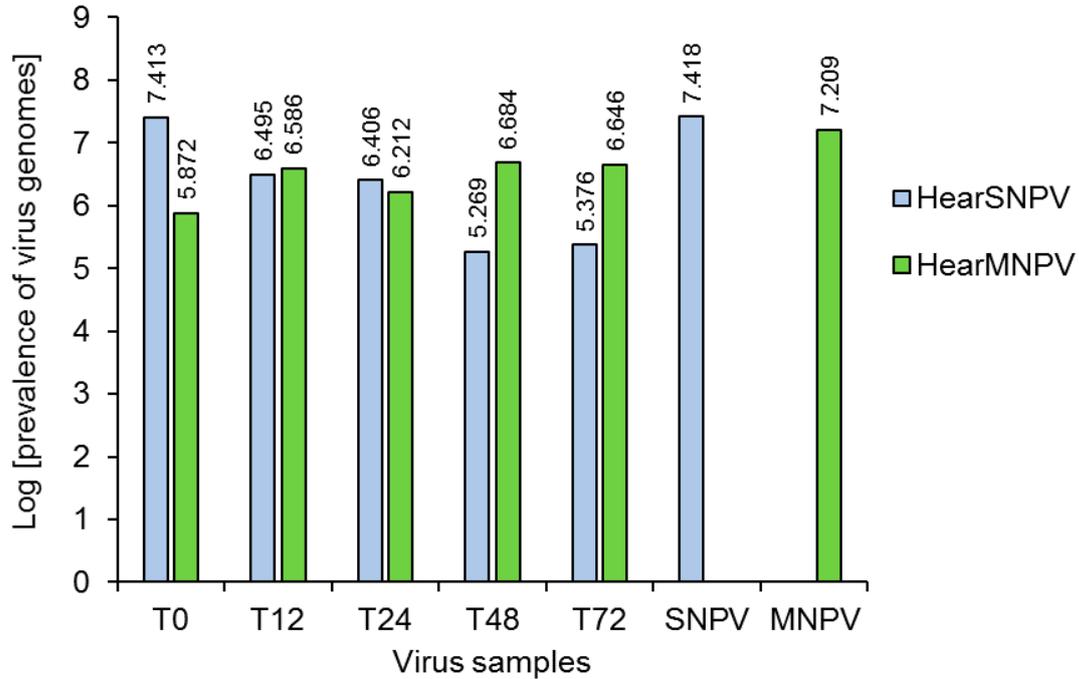


Figure S1. Logarithm of copy number of HearSNPV (blue columns) and HearMNPV (green columns) genomes determined by qPCR of progeny occlusion bodies obtained after initial infection of *Helicoverpa armigera* larvae with an LC₉₀ concentration of HearMNPV followed at intervals of 0 h (T0), 12 h (T12), 24 h (T24), 48 h (T48) and 72 h (T72) by superinfection with an LC₉₀ concentration of HearSNPV occlusion bodies. Values above columns indicate logarithms of the prevalence of virus genomes (copy number).