



**Supplement Figure S1. Comparative quantification of viral genome by qRT-PCR in TOCs (A,C) and OOCs (B,D) after inoculation with AMPV- (A,B) or NDV- (C,D) viral strains** Per group, three to five rings were collected at 1, 24, 48, 72, and 96hpi and processed for viral quantification. Normalised data are presented as mean 40 - Ct. Error bars represent standard deviations (SD). Small letters indicate significant differences between compared viral strains at the same time point for each virus. Two-sample t-test. *P*-value < 0.05. Graphs represent data of one representative experiment.