

Supplemental Results

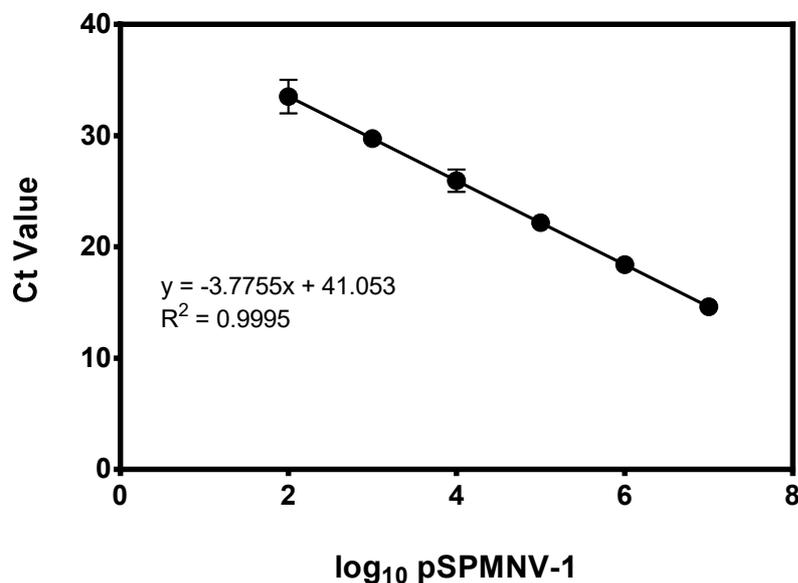


Figure S1. Representative standard curve for determining murine norovirus concentration in samples quantified using RT-qPCR. The plasmid pSPMNV-1 was used in qPCR analysis of MNV-1 containing samples to determine virus concentration. Serial dilutions starting from 10^7 and ending at 10^2 were used to generate the standard curve. Linear regression analysis of this curve resulted in an $R^2 = 0.9995$, which was typical for all RT-qPCR runs.

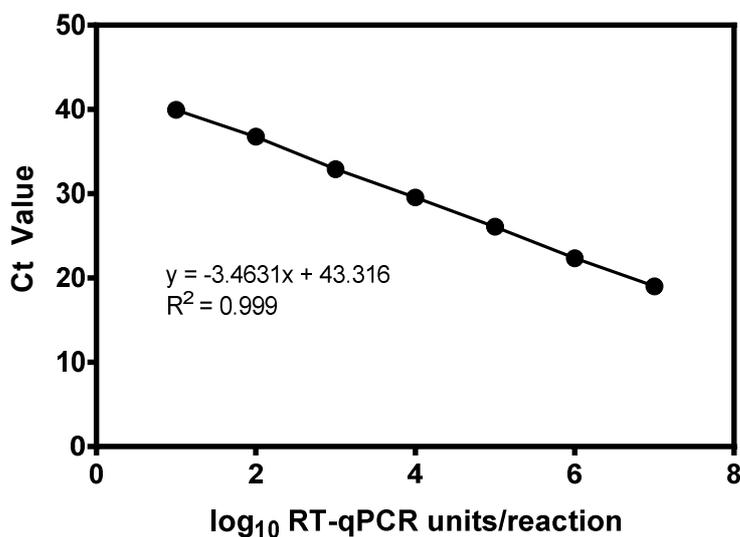


Figure S2. Representative standard curve for RT-qPCR quantification of MS2 viral genome in samples. The viral RNA was used for the determination of MS2 in the samples. The amplification signals showed a strong linearity ($R^2 = 0.999$) from 10^7 to 10^1 RT-qPCR Units.