



Figure S1: Bean plot of Cq values of 1×10^1 copies/reaction of RNA standards measured in porcine sapelovirus (PSV), porcine teschovirus (PTV) and porcine astrovirus type 3 (PoAstV-3) RT-qPCR assays. Grey lines represent individual data points; polygons represent the estimated density of the data. Horizontal black line indicates the cut-off Cq value which was arbitrarily set to 36.5. The cut-off Cq threshold was set to be located between the highest and second highest Cq values in case of all three viruses.