

Figure S1. Phylogenetic tree generated using the partial sequences of TTCaV amplified by primer set 1 and the reference strains. The 13 TTCaV-positive samples identified in this study are marked with closed triangles

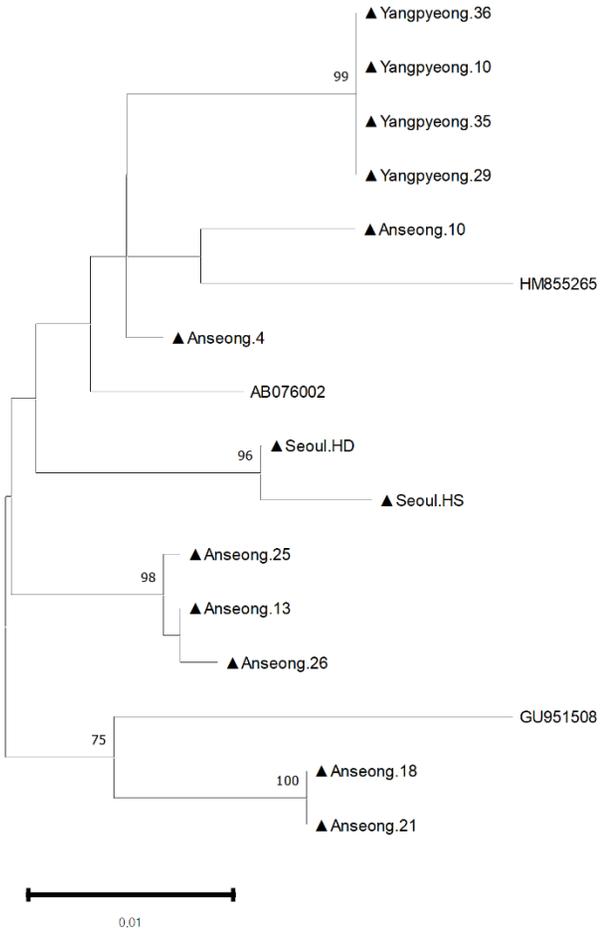


Table S1. Recombination region identified in MZ503912

Recombinant	Parental virus		Breakpoints ^a		Score for the seven detection methods in RDP4 ^b						
	Major	Minor	Begin	End	RDP	GENECONV	BootScan	MaxChi	Chimaera	SiScan	3Seq
MZ503912	KX827768	MZ503910	40	985	1.3×10^{-224}	1.1×10^{-226}	5.5×10^{-223}	1.2×10^{-60}	3.6×10^{-42}	4.8×10^{-83}	6.3×10^{-14}

^aThe breakpoints are based on the locations in the genome of MZ503912.

^bThe *p* value cut-off was 0.05. *p* < 0.05 indicated that the recombination events were significant.