TCAAGCAAAAGCAAAAGGCT	CCATTGGCGGGTGGACGTTGA	TCCATCGCGAGCACGGA
	.GC	GG
	.GC	GA
ATCATTC	T	GACGA.T
ATCA.GCTTC	CTCTC.C.	GGCC
ATCA.GCTTC	CACC.C.	GGCC
ATCA.GCTTC	cccc.c.	AGCC
AATGA.GTT.	.GCC	GG.T.TCC
AATGA.GGTCCA.	.GCTCC.C.	GG.T.TAGCC
ATATGAA.GTAC	.TCTC.C.	GGGT.TT.T
AATGAA.G.C.CAC	.GCCC.C.	GG.T.TGCGC
A.CTC.TCAGGCGCAAC	.AG.CAAC.AC.	G.TCCTCCC
	GCGC.C GCGC.C A.TC.A.TTC A.TC.A.GC.TTC A.TC.A.GC.TTC A.TC.A.GC.TTC A.TC.A.GC.TTC A.TC.A.GC.TTC A.TC.A.GC.TTC A.TC.A.GC.TTC A.TC.A.GC.TTC A.ATG.A.G.TT	

## Additional file 7.

## TaqMan qPCR assay design for sensitive detection and quantification of A. astaci.

Target sites of primers and probe are highlighted by arrows and line, respectively. Parentheses contain GenBank accession numbers. Primers of the TaqMan qPCR and qPCR/MCA assays are identical except the deleted nucleotide at the very 5' end of the reverse primer used in qPCR/MCA (Figure 5a).