

TR19I

GATTGACGGCACCCACACAG CGTGCCACCTTT CGATTCC CGATTCC CGATTCC CGATTCC
CCAATCC TCAATCC TCAATCC TCAATCC TCAATCC TCAATCC TCAATCC
CCACCACGCATGC ACCACGGCACAACGTCCTGG

TR54I

CTGCATCGGCCTGATCGTGG CCGGGGTGATCGGGCTGAAGCTGGTGGGGTAGT
CGTGGTCAGCCCTG CGTGATCAGCCCTG CGTGGTCAGCCC
CACTGCGTGGGCCTGCCTCCCAAGATTGCTATGCAAACCTTGGGCCCCACTTCGTAGCAGC
CTATCCCCCGCCTTCGGCGCGCCCCCTTAACAGAAGGGGGCTCTGCTCCAG
TGGGGTGTGCGAGGTGTGGGGCATAGTTTGCTAGGC GCCGCCA GCCGCCA GCCGCCA
GCCGCCAATCC CCAATCC CCAATCC CCAATCC CCAATCC CCAATCC CCA CCGGCT
ACAGCCGGTACGCCACCGCTTT

TR05II/TR06II

GTGCAGCACCAGCCAAAGGCAGTTGCATCAGACCCAGCGCCGCAGATGCTGTCCCGAT
TCCCGAT TCCCGAT TCCCGAT TCCCGAT TCCCGAT TCCCGAT TCCCGAT TCCCGAT
TCCCAAT CCCCGAT TCCCAAT CCCCAAT CCCCAAT CCCCAAT CCCCAAT CCCCAAT
CCCCAAT CCCCA GCCTATGACCTTCCGCGAACCCGTCGAC

TR67II

AGCTCGCAACTGCTT TTCCCGA TTCCCGA TTCCCGA TTCCCGA TTCCCGA TTCCCGA
TTCCCGA TTCCCGA TTCCCGA TTCCCGA ATCCCGA TTCCCGA TTCC
AGCCCCAATATGCGAAACTTCCACGATGACAATCGTTCTCATCGCGTTCGCCTTGTATC

TR18II

CCGGCATT TTTT GGAATCG GGAATCG GGAATCG GGAATCG GGAATCG GGAATCG
GGAATC GGAATCG GGAATCG GAGAGGCG GGAATCG GAGAGGCG
GGAATCGCAGAAGCGGGGTGTCGCTCTTGTCTACCTATCCCCATTCCAATTCCCGATTCC
CAGCCCTCAAAGCATCCGGCTGATCGTGAA

FIG S1 Nucleotide sequence in *Xanthomonas arboricola* pv. pruni strain CFBP 3894 of TR loci developed by Cesbron et al. [33] that were excluded from the analysis (TR19I and TR54I) or for which new primer combinations were designed (other loci).

Color blocks indicate TRs differing in nucleotide sequence. Underlined regions correspond to primers designed by Cesbron et al. [33]. For TR05II and TR06II, the bold and italicized regions correspond to TR05II and TR06II, respectively. Plain and dashed underlined regions correspond to primers designed by Cesbron et al. [33] for TR05II and TR06II, respectively.