

PRIMER NAME	SEQUENCE
P25_BAK1_F	TTGGTCTCAAATGGAACGAAGATTAATG
P26_BAK1_R	TTGGTCTCTAAGCTTATCTTGGACCCGAGGG
P30_Y304F_R	TTGGTCTCTTTAAAACTTTACCAAAACCAC
P32_Y363F_R	TTGGTCTCTGGAAAAACAAGCAATCTTTCCGG
P34_Y365F_R	TTGGTCTCTATGAAGGGATAAACAAGCAATC
P36_Y403F_R	TTGGTCTCTAAAAACGCAAGCCCTCTTGCAG
P38_Y443F_R	TTGGTCTCTTTGAAGTCCATGAGTTTTGCAA
P40_Y463F_R	TTGGTCTCTAGGAACTCAGGGGCTATATGAC
P42_Y530F_R	TTGGTCTCTAAGAAATTACCCTGAAGATCAA
P44_Y587F_R	TTGGTCTCTGGGAAGTTGAAATCTTGTCTGA
P29_Y304F_F	TTGGTCTCATTAAAGGACGGTTAGCTGATGG
P31_Y363F_F	TTGGTCTCATTCCCTACATGGCTAATGGAAG
P33_Y365F_F	TTGGTCTCATCATGGCTAATGGAAGTGTTC
P35_Y403F_F	TTGGTCTCATTTTACATGATCATTGCGACCC
P37_Y443F_F	TTGGTCTCATCAAAGACACACATGTGACAAC
P39_Y463F_F	TTGGTCTCATCCTTTCCACTGGAAAATCATC
P41_Y530F_F	TTGGTCTCATCTTAGACGAAGAAGTGGAGCA
P43_Y587F_F	TTGGTCTCATCCCAACCCACCATCCAGCCGT
P63_S612D_R	TTGGTCTCTAAGCTTATCTTGGACCATCGGG
P64_Y478F_F	TTGGTCTCATTGGAGTCATGCTTCTTGTAGCT
P65_Y478F_R	TTGGTCTCTCCAAACCCAAAGACATCGGTTT
P66_Y610F_R	TTGGTCTCTAAGCTTATCTTGGACCCGAGGGGAATTCGTT
P82_C408F_F	TTGGTCTCATCGACCCAAAGATTATTCATCG
P83_C408F_R	TTGGTCTCTTCGAAATGATCATGTAATAACG
P86_C408A_F	TTGGTCTCAGCCGACCCAAAGATTATTCATCG
P95_C408A_R	TTGGTCTCTCGGCATGATCATGTAATAACG
P90_C408S_F	TTGGTCTCACCGACCCAAAGATTATTCATCG
P91_C408S_R	TTGGTCTCTTCGGAATGATCATGTAATAACG
P58_S602A_R	TTATCTTGGACCCGAGGGGTATTCTGTTTTCGATCTGGGAAGTGGCATC
P60_T603A_R	TTATCTTGGACCCGAGGGGTATTCTGTTTTCGATCTGGGAAGCGGAATC
P61_S604A_R	TTATCTTGGACCCGAGGGGTATTCTGTTTTCGATCTGGGCAGTGGGAATC
P69_S602A/T603A_R	TTATCTTGGACCCGAGGGGTATTCTGTTTTCGATCTGGGAAGCGGCATCGCC
P70_S602A/S604A_R	TTATCTTGGACCCGAGGGGTATTCTGTTTTCGATCTGGGCAGTGGCATCGCC
P68_S602A/T602A/S603A	TTATCTTGGACCCGAGGGGTATTCTGTTTTCGATCTGGGCAGCGGCATCGCC
P111_S602D/T602D/S603D	TTGGTCTCTAAGCTTATCTTGGACCATCGGG
P22_Y403D_F	CTGCAAGAGGGCTTGCAGTTTACATGATCATTGCGAC
P22_Y403D_R	GTCGCAATGATCATGTAATCCGCAAGCCCTCTTGCAG
P74_GREEN_BAK1_F	GTTACTCCATGTACATGGTTTCATGTTACTTGCAATAGC
P75_GREEN_BAK1_R	TGACTCTAGAGGATCCTTATCTTGGACCCGAGGGGTATTC

P76_GREEN_R_S612A	TGACTCTAGAGGATCCTTATCTTGGACCCGCGGGGTATTC
P108_POPINM_WT_F	AAGTTCTGTTTCAGGGCCCGGACCACTTCTTTGATGTACCAGC
P109_POPINM_WT_R	ATGGTCTAGAAAAGCTTTATCTTGGACCCGAGGGGTA
P110_POPINM_S612A	ATGGTCTAGAAAAGCTTTATCTTGGACCCGCGGGGTA
P112_POPINM_S612D	ATGGTCTAGAAAAGCTTTATCTTGGACCATCGGGGTA
P128_D416N_F	CCCAAAGATTATTCATCGAAATGTGAAAGCTGCAAATATTTTGTG
P129_D416N_R	CAACAAAATATTTGCAGCTTTCACATTTTCGATGAATAATCTTTGGG
BS127_C408F-F	GGCTTGCGTATTTACATGATCATTTCGACCCAAAGATTATTCATCGAG
BS128_C408F-R	CTCGATGAATAATCTTTGGGTCGAAATGATCATGTAATACGCAAGCC
ERECTA_Y760F_F	GCACAAGGTTTAGCTTTTCTACACCATGACTG
ERECTA_Y760F_RC	CAGTCATGGTGTAGAAAAGCTAAACCTTGTGC
