Supplementary Table S4 (Ahn et al.)

Sequences of the primers used in this study.

Primer	Sequence[a]
JH1061	5'-ATGTCTATAAATATAAGAGACCCTCTTATAGTAA-3'
JH1062	5'-CAAAGTATAGAAGTTCCTGAGGTCTT-3'
JH1063	5'-CAATGGAGATATTCTCGGAGGT-3'
JH1064	5'-CAAAGGTTGTTCCAGTTGTAGCA-3'
JH1065	5'-CTAAAGTCTTCTTCCTCCGCA-3'
JH1066	5'-atggagaatatgggaactagag-3'
JH1067	5'-CCTCAGGAACTTCTATACTTTGgtgatgatagacccagatgttc-3'
JH1068	5'-TCACCTCCGAGAATATCTCCATTGgatcgttacaaacattcccgg-3'
JH1069	5'-TGCTACAACTGGAACAACCTTTGgcaaagaggtggtgagctatg-3'
JH1070	5'-ctagcgtttgcgtgcagc-3'
JH1071	5'-gatctcagatccttcttcactttgGTTATGGTGGATCCAGATGTTC-3'
JH1072	5'-ctttctaaaagaacacctgcactgGTTGGTGACTGATATCCCTGC-3'
JH1073	5'-gcacaacagatgctacgtttgGCAATGAGATTGTGTGTTACGAA-3'
JH1074	5'-caaagtgaagaaggatctgagatc-3'
JH1075	5'-cagtgcaggtgttcttttagaaag-3'
JH1076	5'-caaacgtagcatctgttgtgc-3'
JH1089[b]	5'-GAGGTACCATGTCTATAAATATAAGAGACCCTCTTATAGTAA-3'
JH1090[b]	5'-GATCTAGACTAAAGTCTTCTTCCTCCGCA-3'
JH1091[b]	5'-AAGGTACCatggagaatatgggaactagag-3'
JH1092[b]	5'-TCTCTAGActagcgtttgcgtgcagc-3'
JH1125	5'-CTCCCTCTGACAATTGTAGAAAACTG-3'
JH1126	5'-ttctctttgtgcgttaaagaagacg-3'
JH1127	5'-GTTCTGGCGCCACCCTGGT-3'
JH1128	5'-gtgatctctcgaagggatattaggaa-3'

JH1129	5’-CTGTCGAAACAATATAAACACGACAC-3’
JH1130	5’-ctgcctgaacagaacaaacacaaac-3’
JH1131	5’-GTGTCGTGTTTATATTGTTTCGACAGaagcaaagacgtgttatctttcct-3’
JH1132	5’-gtttgtgtttgttctgttcaggcagCTTGGCAGGCAAACAGTGTA-3’
JH1133	5’-ACCAGGGTGGCGCCAGAACttcaacactcgtaaatttgcggt-3’
JH1134	5’-ttcctaatatcccttcgagagatcacTTCAACACTCGCGAGTTTGCT-3’
JH1135[c]	5’-CTTGGCAGGCAAACAGTGTATGCACCAGGGTGGCGCCAGAAC-3’
JH1136[c]	5’-GTTCTGGCGCCACCCTGGTGCATACACTGTTTGCCTGCCAAG-3’
JH1137[c]	5’-aagcaaagacgtgttatctttcctaatatcccttcgagagatcac-3’
JH1138[c]	5’-gtgatctctcgaagggatattaggaaagataacacgtctttgctt-3’

[a] Oligonucleotide sequences designed from *FT* are denoted in uppercase, whereas those from *TFL1* are designated in lowercase. Synthetic oligonucleotides from JH1061 to JH1076 were for the generation of exon-swapping constructs, whereas those from JH1125 to JH1138 were to produce the segment-swapping constructs within the fourth exon.

[b] The nucleotides underlined denote the sequences for the two synthetic restriction sites, *Kpn* I and *Xba* I, used to facilitate subcloning of the final PCR products.

[c] Primers corresponding to both sense and antisense strand of segment B in the fourth exon.