

Additional file 12.

Table 2. Oligonucleotide compilation.

Plant species	Name	Type	Primer sequence (5'-3')	Motif
<i>Jatropha curcas</i>	Jc60D	Forward	GGAGGATCCYACGGACAAATTCGTRACCGTG	PPT and LTR 5' end
	Jc60FullR	Reverse	GTCTCTAGACATCCTCCCCACTCAAACCAG	
<i>Jatropha curcas</i>	Jc60AdjD	Forward	CCGTGACAGAGTGGTATCAGAGCAG	LTR 3' end and PBS
	Jc60R	Reverse	CGGTCTAGAGTCTCGCGCAGCACTGCGCCC	
<i>Jatropha curcas</i>	Jc77D	Forward	TGGTCTAGAAGCAATGGCGGAGTCTAGGCACC	Central region
	Jc77R	Reverse	CGAGGATCCAAGACACAGGGCGGTACTTTTGAC	
<i>Fragaria ananassa</i>	Fa87D	Forward	GAGTGTCTAGACCCGCTTTYGGGAGTGTATYATG	PPT and LTR 5' end
	Fa85R	Reverse	CTCTCCCCACCAAACAAGGTGTACG	
<i>Fragaria ananassa</i>	Fa92D	Forward	TTGAgGATCctTAAAAGTTGAAACTTAC	Full genomic retrozyme (BATT01039028.1)
	Fa92R	Reverse	CTTTTctaGAACAAAGATCATATTGTCA	
<i>Eucalyptus camaldulensis</i>	Ec72D	Forward	CACGGATCCGTGAATYCTCCCGTACYCGTGA	PPT and LTR 5' end
	Ec72R	Reverse	ACAWTCTAGACCACTCAATTCGCAACGCCCTCGTT	
<i>Citrus sinensis</i>	Cs70D	Forward	CTTAGGATCCTGTACGGACYGCAATCGTGGCACCC	PPT and LTR 5' end
	Cs70R	Reverse	GTCTTCTAGATTCTCCACCGCCATAACCATGC	