

Primer positions in the attached sequence, analyzed in pDRWN32 :

positive intron 2 fp (CAG TCC TTC GCT CTT GAA TCG TTG C)50966  
positive intron 2 rp (GAG GAA GGT GAA ACG AGG CGG AGA)51076  
positive intron 4 fp (CTC TGT TCT CAG ACA GTA ATA TCT)81034  
positive intron 4 rp (GAG AAC AAT CAC ACT GGT AGA G) 81156  
positive intron 15-1 fp (CAA GAC TTC TGT GTC CAC CTT GGC) 257181  
positive intron 15-1 rp (AAT GCC AGG AGT TGA GCT GA) 257312  
positive intron 15-2 fp (GAC CAG CCT ATC CAA CAT AGT GA) 258106  
positive intron 15-2 rp (GTG CAC CTC CAC CTC AAC TCA CTG) 258247  
intron 4 forward 1 (CCT TCG TGT GAA TGA CCA TCT ATT CC) 90965  
intron-4 reverse 1 (GTT TTC CTG AGA TGT TAT ATT GAA ACG T) 91062  
intron 4 forward 2 (GAC CCT GTT CAT AGT TGC AGA TAC A) 93592  
intron 4 reverse 2 (TCA TAA TGA ACA ACT TAG ACA AAC GGA G) 93759  
intron-4 forward 3 (CAA CAG CAG TAA AGT CAG AAG CCA G) 94899  
intron-4 reverse 3 (GGG TAT TGC ACT GTG TGG GAA AGA) 95053  
intron-4 forward 4 (GCA CAT TCG AAG TCT CAC TGC TCA CA)96572  
intron-4 reverse 4 (CTG AAC TCC TTT GTA ATT CAG GCA AC) 96647  
intron-4 forward 6 (CAG CGG GTG GTT CAA ATG CAG AAA) 104335  
intron-4 reverse 6 (ACA GGG TCT CAA CTA GGA GGG AAA) 104490  
intron 18 forward (CAT GTG TGG GCA CCT CAA CCA AAT) 276059  
intron 18 reverse (TTA CAG GGC CAT ACA AAC GAG CTG) 276173