

Primer		Sequence	T _M [°C]	Sub- fragment	Split marker fragment	Split marker fragment size [bp]
1	A_fw	TGCATAAGAACACATCAAACACACG	67.9	A	AB	2424
2	A_bw	GTTGTCGACCTGCAGCGTACGTGTCAAGCAATC GGGGTG	67.03			
3	B_fw	CACCCCGATTGCTTGACACGTACGCTGCAGGTC GACAAC	66.24	B		
4	B_bw	CGGTGAGAATGGCAAAGCTTATG	70			
5	C_fw	AAGCCCGATGCGCCAGAGTTG	68	C	CD	2360
6	C_bw	CCGTATCCTAAATGGGCAGTTGTAGTGGATCTG ATATCACCTA	54			
7	D_fw	TAGGTGATATCAGATCCACTACAACCTGCCATTT AGGATACGG	66.92	D		
8	D_bw	GGATGAGGATGACGACGATG	66.01			