

Supplementary data table 1: New *F8* amplimers and primers for DNA mutation screening by CSGE.

Primer Number	Primer sequence (5' - 3')	GenBank AY769950 Nucleotide range	Amplimer Size (bp)	Annealing T. (°C)
14A 14A ₂ *	GACCTGTGATATAATGATAC GAAAAAGTCTCATAT TTGGC	94035-94054 94553-94534	518	55
14B ₁ * 14B	CCTTGGTTTGCAGACAGAAC TGTATTATCAGTACCTGCTG	94414-944433 94806-94787	392	58
14C 14C ₂ *	AGCAACAGAGTTGAAGAAAC CTAATATATTTTGCAGACT	94704-94723 95203-95184	499	55
14D ₁ * 14D	AACAAAACCTCCAATAATTC AGAGTTCCTTCCATGAGTCC	95104-95123 95454-95435	350	55
14E 14E ₂ *	CCCCATTCCACCAGATGCAC ATCTTGAAGTACTGGAGCAT	95352-95371 95853-95834	501	58
14F ₁ * 14F	TACATACAGTGAAGTGGCACT GACCACTGGGTTGAGGTGTC	95744-95763 96125-96106	381	58
14G 14G ₂ *	CACGCAACGTTAGTAAGAGAG GCCAACCTCTCTTTGATCAC	96024-96043 96483-96464	460	58
14H ₁ * 14H	TATAGAAAGAAAATTCTGG CAGGTCTGTTTGCTTCATTC	96364-96383 96727-96708	364	50
14J 14J ₂ *	CCCTACGGAACTAGCAATG TCTTCATTTCAACTGATATG	96621-96640 97123-97104	503	55
14K ₁ * 14K	AGGACTGAAAGGCTGTGCTC AAGAGTTTCAAGACACCTTG	96994-97013 97379-97360	386	58
17A 17B*	TGTCATTCTGGAATCTACTGA CAC TCC CAC AGA TAT ACT CT	121165-121185 121656-121637	491	55
18A* 18B	AGAGTATATCTGTGGGAGTG CTTAAGAGCATGGAGCTTGT	121637-121656 122049-122030	412	55-58
26A 26A ₂ *	GGTTTAATCCTGGACTACTG GCACAAAGGTAGAAGGCAAG	187995-188014 188351-188332	356	55-58

New amplimers were performed using primers from Williams et al [7], and this study*.