

* – diagnostic positions

	1
AF059893_ITS1_A1	GGTTGCCAGCCGCCGGCATGGGGCTGTGCTCCGAAAACCAAAGTCTGCTGTGGGGTGGGTAGGGTATGGGGGCTCAGCCCCC
AF059899_ITS1_A2
AF059897_ITS1_A3
AF059900_ITS1
	84
AF059893_ITS1_A1	CGCCTCGCCCATCTCTCGGCGCGGGTGTCTCGGTCTTAGCCCGGTTCCCCGCTATTCTTTTGCTAAGGGTTGCACCCGAC
AF059899_ITS1_A2
AF059897_ITS1_A3
AF059900_ITS1T.....
	*
	167
AF059893_ITS1_A1	CGGCTCCATCCCCTTTCCCCGTTAGCCACGGCCACATGGCGCACCTATGGGCAGGTGAGTCGGCCGCTACCGAAGGGGACTGG
AF059899_ITS1_A2
AF059897_ITS1_A3
AF059900_ITS1
	250
AF059893_ITS1_A1	GGGTGTCCGGTGAACCGGGACTTCCCGAAA TGTAA TCCCATTTATAAGCGGC TTGAGTATCGCCCAGTATCCTCGCTCGGCAC
AF059899_ITS1_A2C.....
AF059897_ITS1_A3C.....T.....
AF059900_ITS1A.....
	*
	333
AF059893_ITS1_A1	CGGGAACCCAGTCAACCGCTCTGCGCCCCGGCGCAGGCGGGGGTTTAA TGTCTCCCCCAGCCCTCACGGCGCTTCGRCGACGG
AF059899_ITS1_A2G.....
AF059897_ITS1_A3G.....
AF059900_ITS1G.....
	416
AF059893_ITS1_A1	CGGCGGC GGGTGAGCACCCGGAGGCCCCCCAACCTGAAACAACACTTGTCTTGAACTATGGCCTCTCACTCTGGCGAAGTGCGG
AF059899_ITS1_A2
AF059897_ITS1_A3
AF059900_ITS1AAC.C.....
	*** *
	499
AF059893_ITS1_A1	CGGGGGGAAAGGAGGGCAACCTCCCCAAC TCCGTCTAGCAACTAGCCTCTGTGTACAAAAAAAAAGAGTA
AF059899_ITS1_A2
AF059897_ITS1_A3C.....
AF059900_ITS1-