

EBmac0679	4H	94.50	Forward Sequence (5' -> 3')	ATTGGAGCGGATTAGGAT	(AC)22	0,8	148
			Reverse Sequence (5' -> 3')	CCCTATGTCATGTAGGAGATG			
Bmag0223	5H	86.88	Forward Sequence (5' -> 3')	TTAGTCACCCTCAACGGT	(AG)16	0.82	127
			Reverse Sequence (5' -> 3')	CCCCTAACTGCTGTGATG			
scssr02306	5H	6.13	Forward Sequence (5' -> 3')	TGCCTTGTTTATGTAATATCTTG	(AT)6 (CA)7	0.67	
			Reverse Sequence (5' -> 3')	GGCGTAAATAAGAGTGTCTTCAG			
Bmac0303	5H	53.10	Forward Sequence (5' -> 3')	CCTCCAAGATTAGATCTCTCTC	(AG)13(AC)21	0.68	138
			Reverse Sequence (5' -> 3')	CCGTATATTTAAGAAATGGTGA			
Bmac0040	6H	113.19	Forward Sequence (5' -> 3')	AGCCCGATCAGATTTACG	(AC)20	0.89	236
			Reverse Sequence (5' -> 3')	TTCTCCCTTTGGTCCTTG			
Bmac0316	6H	7.16	Forward Sequence (5' -> 3')	ATGGTAGAGGTCCCAACTG	(AC)19	0.69	135
			Reverse Sequence (5' -> 3')	ATCACTGCTGTGCCTAGC			
Bmac0127	6H	62.27	Forward Sequence (5' -> 3')	AACTATGTCCAGTCGTTCC	(AC)26	0,83	118
			Reverse Sequence (5' -> 3')	CTTGTCGTATCATCTTATTCAGA			
Bmag0007	7H	20.62	Forward Sequence (5' -> 3')	TGAAGGAAGAATAAACAACCAACA	(AG)16(AC)16	0.76	185
			Reverse Sequence (5' -> 3')	TCCCCTATTAAGTGACGGTGTG			
Bmag0217	7H	79.06	Forward Sequence (5' -> 3')	AATGCTCAAATATCTATCATGAA	(AG)19	0,69	196
			Reverse Sequence (5' -> 3')	GGGGCTGTCACAAGTATATAG			
Bmag0135	7H	147.51	Forward Sequence (5' -> 3')	ACGAAAGAGTTACAACGGATA	(AG)10GG(AG)12	0,87	161
			Reverse Sequence (5' -> 3')	GTTTACCACAGATCTACAGGTG			