

Project: Untitled.SQD Contig 1

		10 20	GACGTTGTAAAACGACGGCCAGTGCCAAG
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→		GACGTTGTAAaAACGACGGCCAGTGCCAAG
G0257397-1.seq (1>2459)	→		GCCAAG
		30 40 50	CTTGCATGCCTGCAGGTCGACGCTCTCCC
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→		CTTGCATGCCTGCAGGTCGACGCTCTCCC
G0257397-1.seq (1>2459)	→		CTTGCATGCCTGCAGGTCGACGCTCTCCC
		60 70 80	TTATGCGACTCCTGCATTAGGAAGCAGCC
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→		TTATGCGACTCCTGCATTAGGAAGCAGCC
G0257397-1.seq (1>2459)	→		TTATGCGACTCCTGCATTAGGAAGCAGCC
		90 100 110	CAGTAGTAGGTTGAGGCCGTTGAGCACCG
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→		CAGTAGTAGGTTGAGGCCGTTGAGCACCG
G0257397-1.seq (1>2459)	→		CAGTAGTAGGTTGAGGCCGTTGAGCACCG
		120 130 140	CCGCCGCAAGGAATGGTGCATGCAAGGAG
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→		CCGCCGCAAGGAATGGTGCATGCAAGGAG
G0257397-1.seq (1>2459)	→		CCGCCGCAAGGAATGGTGCATGCAAGGAG
		150 160 170	ATGGCGCCCAACAGTCCCCCGGCCACGGG
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→		ATGGCGCCCAACAGTCCCCCGGCCACGGG
G0257397-1.seq (1>2459)	→		ATGGCGCCCAACAGTCCCCCGGCCACGGG
		180 190 200	GCCTGCCACCATACCCACGCCGAAACAAG
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→		GCCTGCCACCATACCCACGCCGAAACAAG
G0257397-1.seq (1>2459)	→		GCCTGCCACCATACCCACGCCGAAACAAG
		210 220 230	CGCTCATGAGCCCGAAGTGCGGAGCCCGA
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→		CGCTCATGAGCCCGAAGTGCGGAGCCCGA
G0257397-1.seq (1>2459)	→		CGCTCATGAGCCCGAAGTGCGGAGCCCGA
		240 250 260	TCTTCCCCATCGGTGATGTCGGCGATATA
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→		TCTTCCCCATCGGTGATGTCGGCGATATA
G0257397-1.seq (1>2459)	→		TCTTCCCCATCGGTGATGTCGGCGATATA
		270 280 290	GCGCCAGCAACCGCACCTGTGGCGCCGG
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→		GCGCCAGCAACCGCACCTGTGGCGCCGG
G0257397-1.seq (1>2459)	→		GCGCCAGCAACCGCACCTGTGGCGCCGG

Project: Untitled.SQD Contig 1

		300	310	
		TGATGCCGGCCACGATGCGTCCGGCGTAG		
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→	TGATGCCGGCCACGATGCGTCCGGCGTAG		
G0257397-1.seq (1>2459)	→	TGATGCCGGCCACGATGCGTCCGGCGTAG		
		320	330	340
		AGGATCCTCTAGAAAGAAGGATTACCTCT		
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→	AGGATCCTCTAGAAAGAAGGATTACCTCT		
G0257397-1.seq (1>2459)	→	AGGATCCTCTAGAAAGAAGGATTACCTCT		
		350	360	370
		AAACAAGTGTACCTGTGCATTCTGGGTAA		
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→	AAACAAGTGTACCTGTGCATTCTGGGTAA		
G0257397-1.seq (1>2459)	→	AAACAAGTGTACCTGTGCATTCTGGGTAA		
		380	390	400
		ACGACTCATAGGAGAGTTGTAAAAAAGTT		
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→	ACGACTCATAGGAGAGTTGTAAAAAAGTT		
G0257397-1.seq (1>2459)	→	ACGACTCATAGGAGAGTTGTAAAAAAGTT		
		410	420	430
		TCGGCCGGCGTATTGGGTGTTACGGAGCA		
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→	TCGGCCGGCGTATTGGGTGTTACGGAGCA		
G0257397-1.seq (1>2459)	→	TCGGCCGGCGTATTGGGTGTTACGGAGCA		
		440	450	460
		TTCACTAGGCAACCATGGTACTATTGTA		
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→	TTCACTAGGCAACCATGGTACTATTGTA		
G0257397-1.seq (1>2459)	→	TTCACTAGGCAACCATGGTACTATTGTA		
		470	480	490
		TACCATCTTAGTAGGAATGATTTGAGGT		
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→	TACCATCTTAGTAGGAATGATTTGAGGT		
G0257397-1.seq (1>2459)	→	TACCATCTTAGTAGGAATGATTTGAGGT		
		500	510	520
		TTATACCTACGATGAATGTGTGTCCTGTA		
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→	TTATACCTACGATGAATGTGTGTCCTGTA		
G0257397-1.seq (1>2459)	→	TTATACCTACGATGAATGTGTGTCCTGTA		
		530	540	550
		GGCTTGAGAGTTCAAGGAAGAAACATGCA		
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→	GGCTTGAGAGTTCAAGGAAGAAACATGCA		
G0257397-1.seq (1>2459)	→	GGCTTGAGAGTTCAAGGAAGAAACATGCA		
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1 (1>797)	→	GCA		
		560	570	580
		ATTATCTTTGCGAACCCAGGGCTGGTGAC		
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→	ATTATCTTTGCGAACCCAGGGCTGGTGAC		
G0257397-1.seq (1>2459)	→	ATTATCTTTGCGAACCCAGGGCTGGTGAC		
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1 (1>797)	→	ATTATCTTTGCGAACCCAGGGCTGGTGAC		

Project: Untitled.SQD Contig 1

		590	600	GGAATTTTCATAGTCAAGCTATCAGAGTA	
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→			GGAATTTTCATAGTCAAGCTATCAGAGTA	
G0257397-1.seq (1>2459)	→			GGAATTTTCATAGTCAAGCTATCAGAGTA	
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1 (1>797)	→			GGAATTTTCATAGTCAAGCtaTCAGAGTA	
		610	620	630	AAGAAGAGGAGCATGTCAAAGTACAATTA
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→				AAGAAGAGGAGCATGTCAAAGTACAATTA
G0257397-1.seq (1>2459)	→				AAGAAGAGGAGCATGTCAAAGTACAATTA
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1 (1>797)	→				AAGAAGAGGAGCATGTCAAAGTACAATTA
		640	650	660	GAGACAAATATATAGTCGCGTGGAGCCAA
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→				GAGACAAATATATAGTCGCGTGGAGCCAA
G0257397-1.seq (1>2459)	→				GAGACAAATATATAGTCGCGTGGAGCCAA
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1 (1>797)	→				GAGACAAATATATAGTCGCGTGGAGCCAA
		670	680	690	GAGCGGATTCCTCAGTCTCGTAGGTCTCT
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→				GAGCGGATTCCTCAGTCTCGTAGGTCTCT
G0257397-1.seq (1>2459)	→				GAGCGGATTCCTCAGTCTCGTAGGTCTCT
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1 (1>797)	→				GAGCGGATTCCTCAGTCTCGTAGGTCTCT
		700	710	720	TGACGACCGTTGATCTGCTTGATCTCGTC
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→				TGACGACCGTTGATCTGCTTGATCTCGTC
G0257397-1.seq (1>2459)	→				TGACGACCGTTGATCTGCTTGATCTCGTC
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1 (1>797)	→				TGACGACCGTTGATCTGCTTGATCTCGTC
		730	740	750	TCCCGAAAATGAAAATAGCTCTGCTAAGC
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→				TCCCGAAAATGAAAATAGCTCTGCTAAGC
G0257397-1.seq (1>2459)	→				TCCCGAAAATGAAAATAGCTCTGCTAAGC
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1 (1>797)	→				TCCCGAAAATGAAAATAGCTCTGCTAAGC
		760	770	780	TATTCTTCTCTTCGCCGAGCCTGAAGGC
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→				TATTCTTCTCTTCGCCGAGCCTGAAGGC
G0257397-1.seq (1>2459)	→				TATTCTTCTCTTCGCCGAGCCTGAAGGC
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1 (1>797)	→				TATTCTTCTCTTCGCCGAGCCTGAAGGC
		790	800	810	GTTACTAGGTTGCAGTCAATGCATTAATG
H04-F52109-G0257397-1_M13F-77.ab1 (1>796)	→				GTTACTAGGTTGC
G0257397-1.seq (1>2459)	→				GTTACTAGGTTGCAGTCAATGCATTAATG
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1 (1>797)	→				GTTACTAGGTTGCAGTCAATGCATTAATG
		820	830	840	CATTGCAGATGAGCTGTATCTGGAAGAGG
G0257397-1.seq (1>2459)	→				CATTGCAGATGAGCTGTATCTGGAAGAGG
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1 (1>797)	→				CATTGCAGATGAGCTGTATCTGGAAGAGG

Project: Untitled.SQD Contig 1

		850	860	870
		TAAACCCGAAAACGCGTTTTATTCTTGTT		
G0257397-1.seq(1>2459)	→	TAAACCCGAAAACGCGTTTTATTCTTGTT		
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→	TAAACCCGAAAACGCGTTTTATTCTTGTT		
		880	890	
		GACATGGAGCTATTAATCACTAGAAGGC		
G0257397-1.seq(1>2459)	→	GACATGGAGCTATTAATCACTAGAAGGC		
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→	GACATGGAGCTATTAATCACTAGAAGGC		
		900	910	920
		ACTCTTTGCTGCTTGGACAAATGAACGTA		
G0257397-1.seq(1>2459)	→	ACTCTTTGCTGCTTGGACAAATGAACGTA		
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→	ACTCTTTGCTGCTTGGACAAATGAACGTA		
		930	940	950
		TCTTATCGAGATCCTGAACACCATTGTC		
G0257397-1.seq(1>2459)	→	TCTTATCGAGATCCTGAACACCATTGTC		
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→	TCTTATCGAGATCCTGAACACCATTGTC		
		960	970	980
		TCAACTCCGGAGCTGACATCGACACCAAC		
G0257397-1.seq(1>2459)	→	TCAACTCCGGAGCTGACATCGACACCAAC		
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→	TCAACTCCGGAGCTGACATCGACACCAAC		
		990	1000	1010
		GATCTTATATCCAGATTCGTCAGCTGTT		
G0257397-1.seq(1>2459)	→	GATCTTATATCCAGATTCGTCAGCTGTT		
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→	GATCTTATATCCAGATTCGTCAGCTGTT		
		1020	1030	1040
		TGATGATTCAGTAACGTTAAGTGGATCC		
G0257397-1.seq(1>2459)	→	TGATGATTCAGTAACGTTAAGTGGATCC		
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→	TGATGATTCAGTAACGTTAAGTGGATCC		
		1050	1060	1070
		GGTCGGCATCTACTCTATTCCTTTGCCCT		
G0257397-1.seq(1>2459)	→	GGTCGGCATCTACTCTATTCCTTTGCCCT		
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→	GGTCGGCATCTACTCTATTCCTTTGCCCT		
		1080	1090	1100
		CGGACGAGTGCTGGGGCGTCGGTTTCCAC		
G0257397-1.seq(1>2459)	→	CGGACGAGTGCTGGGGCGTCGGTTTCCAC		
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→	CGGACGAGTGCTGGGGCGTCGGTTTCCAC		
		1110	1120	1130
		TATCGGCGAGTACTTCTACACAGCCATCG		
G0257397-1.seq(1>2459)	→	TATCGGCGAGTACTTCTACACAGCCATCG		
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→	TATCGGCGAGTACTTCTACACAGCCATCG		

Project: Untitled.SQD Contig 1

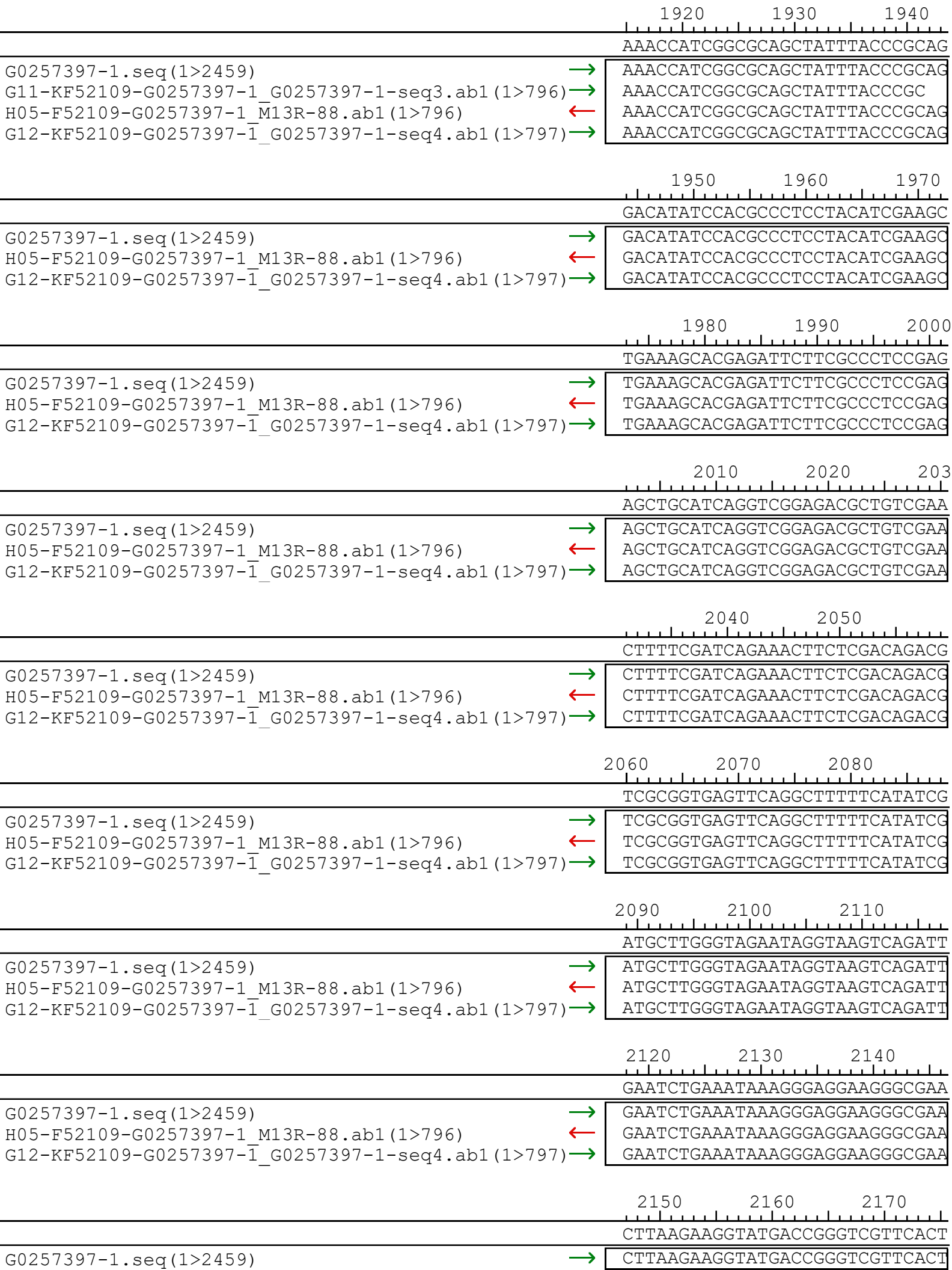
		1140 1150 1160	GTCCAGACGGCCGCGCTTCTGCGGGCGAT
G0257397-1.seq(1>2459)	→		GTCCAGACGGCCGCGCTTCTGCGGGCGAT
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→		GTCCAGACGGCCGCGCTTCTGCGGGCGAT
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		GCTTCTGCGGGCGAT
		1170 1180	TTGTGTACGCCCGACAGTCCCGGCTCCGG
G0257397-1.seq(1>2459)	→		TTGTGTACGCCCGACAGTCCCGGCTCCGG
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→		TTGTGTACGCCCGACAGTCCCGGCTCCGG
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		TTGTGTACGCCCGACAGTCCCGGCTCCGG
		1190 1200 1210	ATCGGACGATTGCGTTCGCATCGACCCTGC
G0257397-1.seq(1>2459)	→		ATCGGACGATTGCGTTCGCATCGACCCTGC
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→		ATCGGACGATTGCGTTCGCATCGACCCTGC
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		ATCGGACGATTGcgTcGcATCGACCCTGC
		1220 1230 1240	GCCCAAGCTGCATCATCGAAATTGCCGTC
G0257397-1.seq(1>2459)	→		GCCCAAGCTGCATCATCGAAATTGCCGTC
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→		GCCCAAGCTGCATCATCGAAATTGCCGTC
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		GCCCAAGCTGCATCATCGAAATTGCCGTC
		1250 1260 1270	AACCAAGCTCTGATAGAGTTGGTCAAGAC
G0257397-1.seq(1>2459)	→		AACCAAGCTCTGATAGAGTTGGTCAAGAC
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→		AACCAAGCTCTGATAGAGTTGGTCAAGAC
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		AACCAAGCTCTGATAGAGTTGGTCAAGAC
		1280 1290 1300	CAATGCGGAGCATATAACGCCCGGAGCCGC
G0257397-1.seq(1>2459)	→		CAATGCGGAGCATATAACGCCCGGAGCCGC
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→		CAATGCGGAGCATATAACGCCCGGAGCCGC
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		CAATGCGGAGCATATAACGCCCGGAGCCGC
		1310 1320 1330	GCGATCCTGCAAGCTCCGGATGCCTCCG
G0257397-1.seq(1>2459)	→		GCGATCCTGCAAGCTCCGGATGCCTCCG
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→		GCGATCCTGCAAGCTCCGGATGCCTCCG
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		GCGATCCTGCAAGCTCCGGATGCCTCCG
		1340 1350 1360	CTCGAAGTAGCGCGTCTGCTGCTCCATAC
G0257397-1.seq(1>2459)	→		CTCGAAGTAGCGCGTCTGCTGCTCCATAC
G10-KF52109-G0257397-1_G0257397-1-seq2.ab1(1>797)	→		CTCGAAGTAGC
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		CTCGAAGTAGCGCGTCTGCTGCTCCATAC
		1370 1380 1390	AAGCCAACCACGGCCTCCAGAAGAAGATG
G0257397-1.seq(1>2459)	→		AAGCCAACCACGGCCTCCAGAAGAAGATG
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		AAGCCAACCACGGCCTCCAGAAGAAGATG

Project: Untitled.SQD Contig 1

		1400 1410 1420	TTGGCGACCTCGTATTGGGAATCCCCGAA
G0257397-1.seq(1>2459)	→		TTGGCGACCTCGTATTGGGAATCCCCGAA
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		TTGGCGACCTCGTATTGGGAATCCCCGAA
		1430 1440 1450	CATCGCCTCGCTCCAGTCAATGACCGCTG
G0257397-1.seq(1>2459)	→		CATCGCCTCGCTCCAGTCAATGACCGCTG
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		CATCGCCTCGCTCCAGTCAATGACCGCTG
		1460 1470	TTATGCGGCCATTGTCCGTCAGGACATTG
G0257397-1.seq(1>2459)	→		TTATGCGGCCATTGTCCGTCAGGACATTG
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		TTATGCGGCCATTGTCCGTCAGGACATTG
		1480 1490 1500	TTGGAGCCGAAATCCGCGTGCACGAGGTG
G0257397-1.seq(1>2459)	→		TTGGAGCCGAAATCCGCGTGCACGAGGTG
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		TTGGAGCCGAAATCCGCGTGCACGAGGTG
		1510 1520 1530	CCGGA CTTCGGGGCAGTCCTCGGCCAAA
G0257397-1.seq(1>2459)	→		CCGGA CTTCGGGGCAGTCCTCGGCCAAA
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		CCGGA CTTCGGGGCAGTCCTCGGCCAAA
		1540 1550 1560	GCATCAGCTCATCGAGAGCCTGCGCGACG
G0257397-1.seq(1>2459)	→		GCATCAGCTCATCGAGAGCCTGCGCGACG
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		GCATCAGCTCATCGAGAGCCTGCGCGACG
		1570 1580 1590	GACGCACTGACGGTGTTCGTCCATCACAGT
G0257397-1.seq(1>2459)	→		GACGCACTGACGGTGTTCGTCCATCACAGT
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		GACGCACTGACGGTGTTCGTCCATCACAGT
		1600 1610 1620	TTGCCAGTGATACACATGGGGATCAGCAA
G0257397-1.seq(1>2459)	→		TTGCCAGTGATACACATGGGGATCAGCAA
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		TTGCCAGTGATACACATGGGGATCAGCAA
		1630 1640 1650	TCGCGCATATGAAATCACGCCATGTAGTG
G0257397-1.seq(1>2459)	→		TCGCGCATATGAAATCACGCCATGTAGTG
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		TCGCGCATATGAAATCACGCCATGTAGTG
		1660 1670 1680	TATTGACCGATTCCCTTGCGGTCCGAATGG
G0257397-1.seq(1>2459)	→		TATTGACCGATTCCCTTGCGGTCCGAATGG
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1(1>796)	→		TATTGACCGATTCCCTTGCGGTCCGAATGG

Project: Untitled.SQD Contig 1

		1690	1700	1710
		GCCGAACCCGCTCGTCTGGCTAAGATCGG		
G0257397-1.seq (1>2459)	→	GCCGAACCCGCTCGTCTGGCTAAGATCGG		
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1 (1>796)	→	GCCGAACCCGCTCGTCTGGCTAAGATCGG		
		1720	1730	1740
		CCGCAGCGATCGCATCCATGGCCTCCGCG		
G0257397-1.seq (1>2459)	→	CCGCAGCGATCGCATCCATGGCCTCCGCG		
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1 (1>796)	→	CCGCAGCGATCGCATCCATGGCCTCCGCG		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	AGCGATCGCATCCATGGCCTCCGCG		
		1750	1760	
		ACCGGCTGCAGAACAGCGGGCAGTTCGGT		
G0257397-1.seq (1>2459)	→	ACCGGCTGCAGAACAGCGGGCAGTTCGGT		
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1 (1>796)	→	ACCGGCTGCAGAACAGCGGGCAGTTCGGT		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	ACCGGCTGCAGAACAGCGGGCAGTTCGGT		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	GCTGCAGAACAGCGGGCAGTTCGGT		
		1770	1780	1790
		TTCAGGCAGGTCTTGCAACGTGACACCCT		
G0257397-1.seq (1>2459)	→	TTCAGGCAGGTCTTGCAACGTGACACCCT		
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1 (1>796)	→	TTCAGGCAGGTCTTGCAACGTGACACCCT		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	TTCAGGCAGGTCTTGCAACGTGACACCCT		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	TTCAGGCAGGTCTTGCAACGTGACACCCT		
		1800	1810	1820
		GTGCACGGCGGGAGATGCAATAGGTCAGG		
G0257397-1.seq (1>2459)	→	GTGCACGGCGGGAGATGCAATAGGTCAGG		
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1 (1>796)	→	GTGCACGGCGGGAGATGCAATAGGTCAGG		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	GTGCACGGCGGGAGATGCAATAGGTCAGG		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	GTGCACGGCGGGAGATGCAATAGGTCAGG		
		1830	1840	1850
		CTCTCGCTGAATTCCCCAATGTCAAGCAC		
G0257397-1.seq (1>2459)	→	CTCTCGCTGAATTCCCCAATGTCAAGCAC		
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1 (1>796)	→	CTCTCGCTGAATTCCCCAATGTCAAGCAC		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	CTCTCGCTGAATTCCCCAATGTCAAGCAC		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	CTCTCGCTGAATTCCCCAATGTCAAGCAC		
		1860	1870	1880
		TTCCGGAATCGGGAGCGCGGCCGATGCAA		
G0257397-1.seq (1>2459)	→	TTCCGGAATCGGGAGCGCGGCCGATGCAA		
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1 (1>796)	→	TTCCGGAATCGGGAGCGCGGCCGATGCAA		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	TTCCGGAATCGGGAGCGCGGCCGATGCAA		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	TTCCGGAATCGGGAGCGCGGCCGATGCAA		
		1890	1900	1910
		AGTGCCGATAAACATAACGATCTTTGTAG		
G0257397-1.seq (1>2459)	→	AGTGCCGATAAACATAACGATCTTTGTAG		
G11-KF52109-G0257397-1_G0257397-1-seq3.ab1 (1>796)	→	AGTGCCGATAAACATAACGATCTTTGTAG		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	AGTGCCGATAAACATAACGATCTTTGTAG		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	AGTGCCGATAAACATAACGATCTTTGTAG		



Project: Untitled.SQD Contig 1

		2150	2160	2170
		CTTAAGAAGGTATGACCGGGTCGTTCACT		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	CTTAAGAAGGTATGACCGGGTCGTTCACT		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	CTTAAGAAGGTATGACCGGGTCGTTCACT		
		2180	2190	2200
		TACCTTGCTTGACAAACGCACAAGTTATC		
G0257397-1.seq (1>2459)	→	TACCTTGCTTGACAAACGCACAAGTTATC		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	TACCTTGCTTGACAAACGCACAAGTTATC		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	TACCTTGCTTGACAAACGCACAAGTTATC		
		2210	2220	2230
		GTGCACCAAGCAGCAGATGATAATAATGT		
G0257397-1.seq (1>2459)	→	GTGCACCAAGCAGCAGATGATAATAATGT		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	GTGCACCAAGCAGCAGATGATAATAATGT		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	GTGCACCAAGCAGCAGATGATAATAATGT		
		2240	2250	2260
		CCTCGTTCCTGTCTGCTAATAAGAGTCAC		
G0257397-1.seq (1>2459)	→	CCTCGTTCCTGTCTGCTAATAAGAGTCAC		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	CCTCGTTCCTGTCTGCTAATAAGAGTCAC		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	CCTCGTTCCTGTCTGCTAATAAGAGTCAC		
		2270	2280	2290
		ACTTCGAGCGCCGCCGCTACTGCTTACAA		
G0257397-1.seq (1>2459)	→	ACTTCGAGCGCCGCCGCTACTGCTTACAA		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	ACTTCGAGCGCCGCCGCTACTGCTTACAA		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	ACTTCGAGCGCCGCCGCTACTGCTTACAA		
		2300	2310	232
		GTGGGCTGATCTGACCAGTTGCCTAAATG		
G0257397-1.seq (1>2459)	→	GTGGGCTGATCTGACCAGTTGCCTAAATG		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	GTGGGCTGATCTGACCAGTTGCCTAAATG		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	GTGGGCTGATCTGACCAGTTGCCTAAATG		
		2330	2340	
		AACCATCTTGTCAAACGACACAAATTTTG		
G0257397-1.seq (1>2459)	→	AACCATCTTGTCAAACGACACAAATTTTG		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	AACCATCTTGTCAAACGACACAAATTTTG		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	AACCATCTTGTCAAACGACACAAATTTTG		
		2350	2360	2370
		TGATCCGCCTGGACGACTAAACCAAATA		
G0257397-1.seq (1>2459)	→	TGATCCGCCTGGACGACTAAACCAAATA		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	TGATCCGCCTGGACGACTAAACCAAATA		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	TGATCCGCCTGGACGACTAAACCAAATA		
		2380	2390	2400
		GCATTGATGTGTTGACCTCCACTAGCTCC		
G0257397-1.seq (1>2459)	→	GCATTGATGTGTTGACCTCCACTAGCTCC		
H05-F52109-G0257397-1_M13R-88.ab1 (1>796)	←	GCATTGATGTGTTGACCTCCACTAGCTCC		
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797)	→	GCATTGATGTGTTGACCTCCACTAGCTCC		

2410 2420 2430
AGCCAAGCCCAAAAATGCTCCTTCAATAT
G0257397-1.seq (1>2459) → AGCCAAGCCCAAAAATGCTCCTTCAATAT
H05-F52109-G0257397-1_M13R-88.ab1 (1>796) ← AGCCAAGCCCAAAAATGCTCCTTCAATAT
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797) → AGCCAAGCCCAAAAATGCTCCTTCAATAT

2440 2450 2460
CATCTTCTGTCGACTCTAGAGGATCCCCG
G0257397-1.seq (1>2459) → CATCTTCTGTCGACTCTAGAGGATCCCCG
H05-F52109-G0257397-1_M13R-88.ab1 (1>796) ← CATCTTCTGTCGACTCTAGAGGATCCCCG
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797) → CATCTTCTGTCGACTCTAGAGGATCCCCG

2470 2480 2490
GGTACCGAGCTCGAATTCGTAATCATGGT
G0257397-1.seq (1>2459) → GGTACCGAGCTCGAATT
H05-F52109-G0257397-1_M13R-88.ab1 (1>796) ← GGTACCGAGCTCGAATTCGTAATCATGGT
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797) → GGTACCGAGCTCGAATTCGTAATCATGGT

2500 2510 2520
CATAGCTGTTTCCTGTGTGAAATTGTTAT
H05-F52109-G0257397-1_M13R-88.ab1 (1>796) ← CATAGCTGTTTCCTGTG
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797) → CATAGCTGTTTCCTGTGTGAAATTGTTAT

2530 2540
CCGCTCACAATTCCACAC
G12-KF52109-G0257397-1_G0257397-1-seq4.ab1 (1>797) → CCGCTCACAATTCCACAC