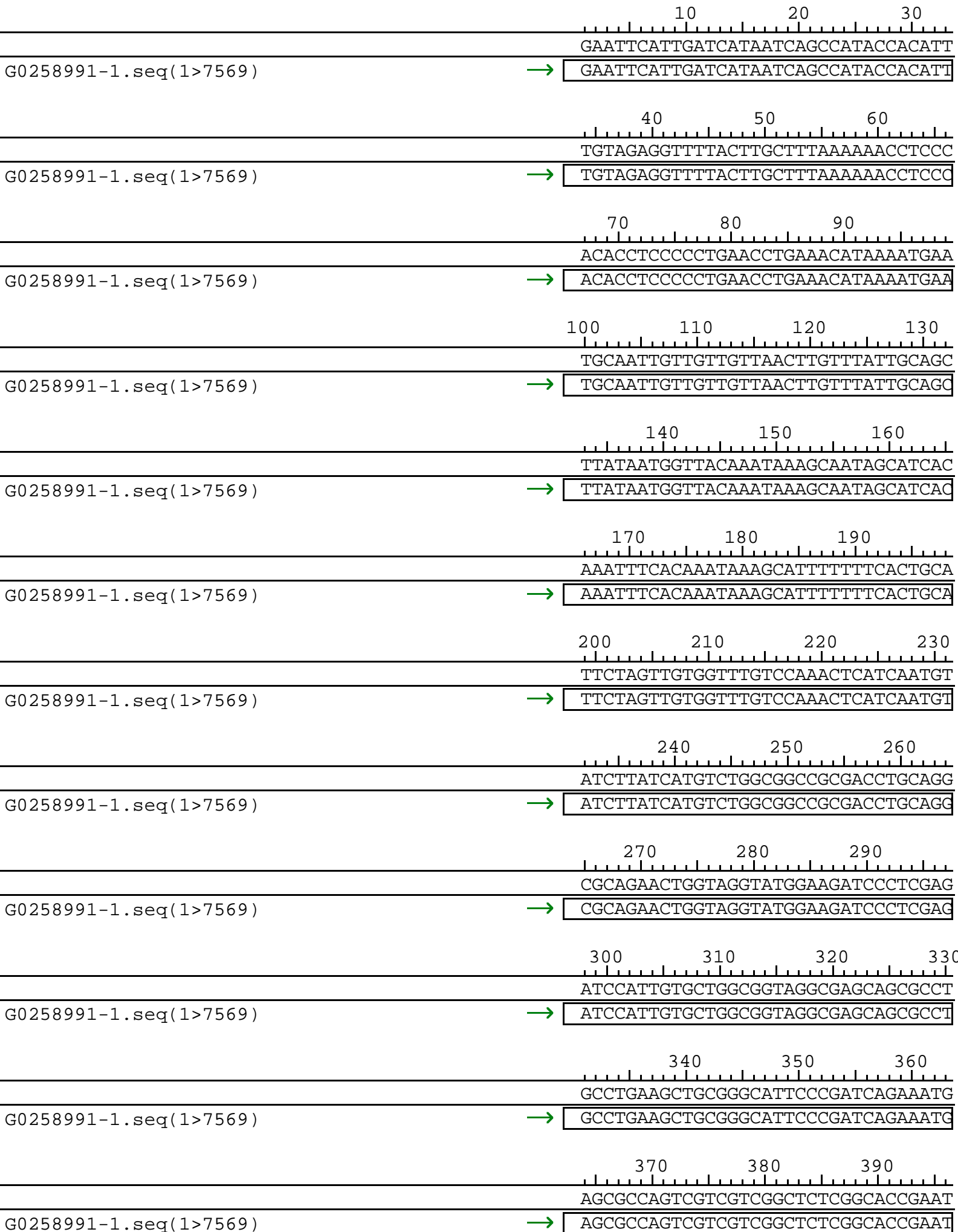


Project: Untitled.SQD Contig 1



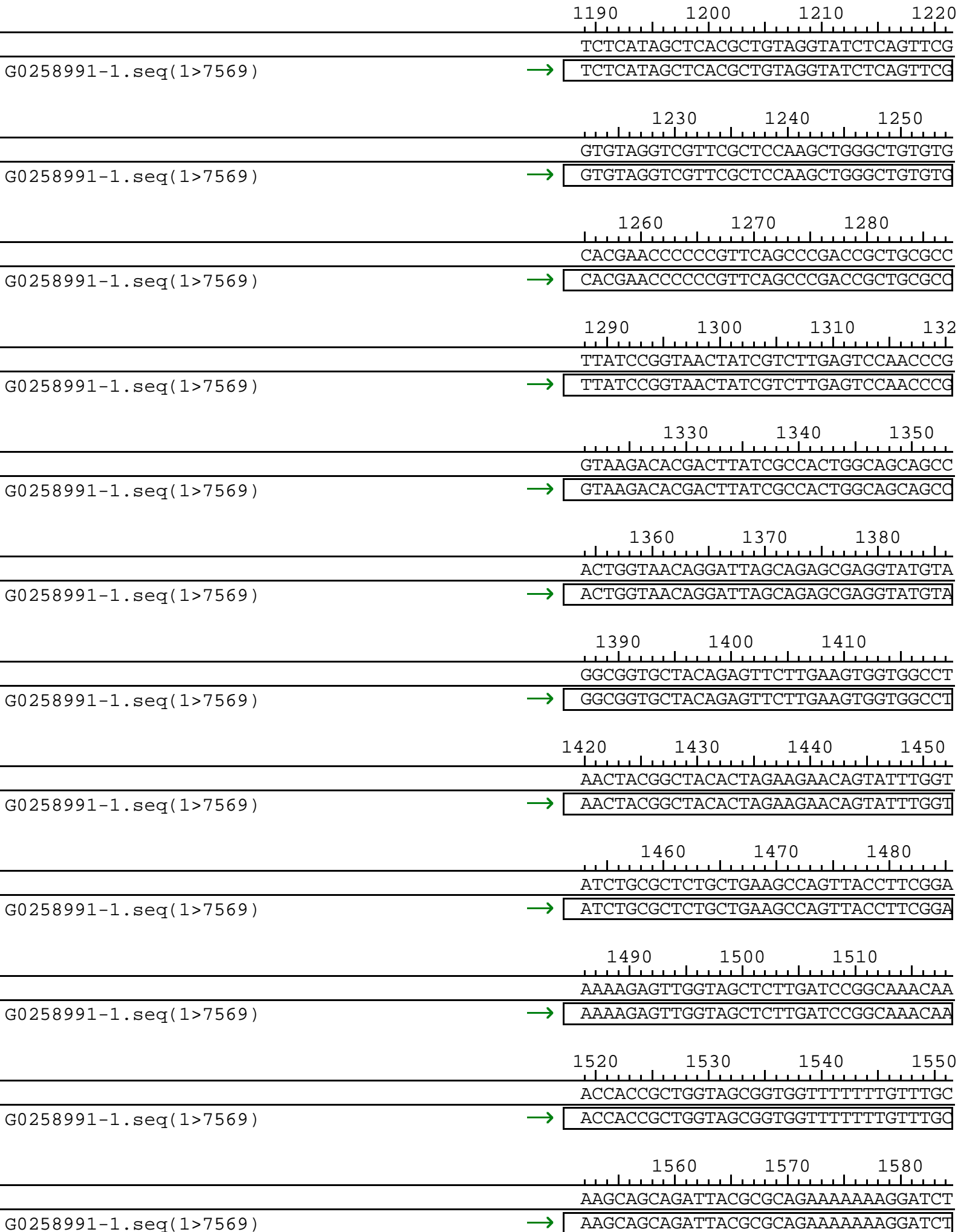
Project: Untitled.SQD Contig 1

		400	410	420	
		GCGTATGATTCTCCGCCAGCATGGCTTCGGCCA			
G0258991-1.seq(1>7569)	→	GCGTATGATTCTCCGCCAGCATGGCTTCGGCCA			
		430	440	450	460
		GTGCGTCGAGCAGCGCCCGCTTGTTCCTGAAGT			
G0258991-1.seq(1>7569)	→	GTGCGTCGAGCAGCGCCCGCTTGTTCCTGAAGT			
		470	480	490	
		GCCAGTAAAGCGCCGGCTGCTGAACCCCAACC			
G0258991-1.seq(1>7569)	→	GCCAGTAAAGCGCCGGCTGCTGAACCCCAACC			
		500	510	520	
		GTTCCGCCAGTTTGCGTGTTCGTCAGACCGTCTA			
G0258991-1.seq(1>7569)	→	GTTCCGCCAGTTTGCGTGTTCGTCAGACCGTCTA			
		530	540	550	560
		CGCCGACCTCGTTCAACAGGTCCAGGGCGGCAC			
G0258991-1.seq(1>7569)	→	CGCCGACCTCGTTCAACAGGTCCAGGGCGGCAC			
		570	580	590	
		GGATCACTGTATTCGGCTGCAACTTTGTCATGC			
G0258991-1.seq(1>7569)	→	GGATCACTGTATTCGGCTGCAACTTTGTCATGC			
		600	610	620	
		TTGACACTTTTATCACTGATAAACATAATATGTC			
G0258991-1.seq(1>7569)	→	TTGACACTTTTATCACTGATAAACATAATATGTC			
		630	640	650	660
		CACCAACTTATCAGTGATAAAGAATCCGCGCCA			
G0258991-1.seq(1>7569)	→	CACCAACTTATCAGTGATAAAGAATCCGCGCCA			
		670	680	690	
		GCACAATGGATCTCGAGGTCGAGGGATCTCTAG			
G0258991-1.seq(1>7569)	→	GCACAATGGATCTCGAGGTCGAGGGATCTCTAG			
		700	710	720	
		AGGATCCTCTACGCCGGACGCATCGTGGCCGGC			
G0258991-1.seq(1>7569)	→	AGGATCCTCTACGCCGGACGCATCGTGGCCGGC			
		730	740	750	
		ATCACCGGCGCCACAGGTGCGGTTGCTGGCGCC			
G0258991-1.seq(1>7569)	→	ATCACCGGCGCCACAGGTGCGGTTGCTGGCGCC			
		760	770	780	790
		TATATCGCCGACATCACCGATGGGGAAGATCGG			
G0258991-1.seq(1>7569)	→	TATATCGCCGACATCACCGATGGGGAAGATCGG			

Project: Untitled.SQD Contig 1

		800 810 820	GCTCGCCACTTCGGGCTCATGAGCGCTTGTTTC
G0258991-1.seq(1>7569)	→		GCTCGCCACTTCGGGCTCATGAGCGCTTGTTTC
		830 840 850	GGCGTGGGTATGGTGGCAGGCCCGTGGCCGGG
G0258991-1.seq(1>7569)	→		GGCGTGGGTATGGTGGCAGGCCCGTGGCCGGG
		860 870 880 890	GGACTGTTGGGCGCCATCTCCTTGCATGCACCA
G0258991-1.seq(1>7569)	→		GGACTGTTGGGCGCCATCTCCTTGCATGCACCA
		900 910 920	TTCCTTGCGGCGGCGGTGCTCAACGGCCTCAAC
G0258991-1.seq(1>7569)	→		TTCCTTGCGGCGGCGGTGCTCAACGGCCTCAAC
		930 940 950	CTACTACTGGGCTGCTTCCTAATGCAGGAGTCC
G0258991-1.seq(1>7569)	→		CTACTACTGGGCTGCTTCCTAATGCAGGAGTCC
		960 970 980 990	CATAAGGGAGAGCGTCGACCTCGGGCCGCGTTG
G0258991-1.seq(1>7569)	→		CATAAGGGAGAGCGTCGACCTCGGGCCGCGTTG
		1000 1010 1020	CTGGCGTTTTTCCATAGGCTCCGCCCCCTGAC
G0258991-1.seq(1>7569)	→		CTGGCGTTTTTCCATAGGCTCCGCCCCCTGAC
		1030 1040 1050	GAGCATCACAAAATCGACGCTCAAGTCAGAGG
G0258991-1.seq(1>7569)	→		GAGCATCACAAAATCGACGCTCAAGTCAGAGG
		1060 1070 1080	TGGCGAAACCCGACAGGACTATAAAGATACCAG
G0258991-1.seq(1>7569)	→		TGGCGAAACCCGACAGGACTATAAAGATACCAG
		1090 1100 1110 1120	GCGTTTCCCCCTGGAAGCTCCCTCGTGCGCTCT
G0258991-1.seq(1>7569)	→		GCGTTTCCCCCTGGAAGCTCCCTCGTGCGCTCT
		1130 1140 1150	CCTGTTCCGACCCTGCCGCTTACCGGATACCTG
G0258991-1.seq(1>7569)	→		CCTGTTCCGACCCTGCCGCTTACCGGATACCTG
		1160 1170 1180	TCCGCCTTTCTCCCTTCGGGAAGCGTGGCGCTT
G0258991-1.seq(1>7569)	→		TCCGCCTTTCTCCCTTCGGGAAGCGTGGCGCTT

Project: Untitled.SQD Contig 1



Project: Untitled.SQD Contig 1

		1590 1600 1610	CAAGAAGATCCTTTGATCTTTTCTACGGGGTCT
G0258991-1.seq(1>7569)	→		CAAGAAGATCCTTTGATCTTTTCTACGGGGTCT
		1620 1630 1640 1650	GACGCTCAGTGAACGAAAACCTCACGTTAAGGG
G0258991-1.seq(1>7569)	→		GACGCTCAGTGAACGAAAACCTCACGTTAAGGG
		1660 1670 1680	ATTTTGGTCATGAGATTATCAAAAAGGATCTTC
G0258991-1.seq(1>7569)	→		ATTTTGGTCATGAGATTATCAAAAAGGATCTTC
		1690 1700 1710	ACCTAGATCCTTTTAAATTA AAAATGAAGTTTT
G0258991-1.seq(1>7569)	→		ACCTAGATCCTTTTAAATTA AAAATGAAGTTTT
		1720 1730 1740	AAATCAATCTAAAGTATATATGAGTAAACTTGG
G0258991-1.seq(1>7569)	→		AAATCAATCTAAAGTATATATGAGTAAACTTGG
		1750 1760 1770 1780	TCTGACAGTTACCAATGCTTAATCAGTGAGGCA
G0258991-1.seq(1>7569)	→		TCTGACAGTTACCAATGCTTAATCAGTGAGGCA
		1790 1800 1810	CCTATCTCAGCGATCTGTCTATTTTCGTTTCATCC
G0258991-1.seq(1>7569)	→		CCTATCTCAGCGATCTGTCTATTTTCGTTTCATCC
		1820 1830 1840	ATAGTTGCCTGACTCCCCGTCGTGTAGATAACT
G0258991-1.seq(1>7569)	→		ATAGTTGCCTGACTCCCCGTCGTGTAGATAACT
		1850 1860 1870 1880	ACGATACGGGAGGGCTTACCATCTGGCCCCAGT
G0258991-1.seq(1>7569)	→		ACGATACGGGAGGGCTTACCATCTGGCCCCAGT
		1890 1900 1910	GCTGCAATGATACCGCGAGACCCACGCTCACCG
G0258991-1.seq(1>7569)	→		GCTGCAATGATACCGCGAGACCCACGCTCACCG
		1920 1930 1940	GCTCCAGATTTATCAGCAATAAACCAGCCAGCC
G0258991-1.seq(1>7569)	→		GCTCCAGATTTATCAGCAATAAACCAGCCAGCC
		1950 1960 1970 1980	GGAAGGGCCGAGCGCAGAAGTGGTCCTGCAACT
G0258991-1.seq(1>7569)	→		GGAAGGGCCGAGCGCAGAAGTGGTCCTGCAACT

Project: Untitled.SQD Contig 1

		1990	2000	2010	
		TTATCCGCCTCCATCCAGTCTATTAATTGTTGC			
G0258991-1.seq(1>7569)	→	TTATCCGCCTCCATCCAGTCTATTAATTGTTGC			
		2020	2030	2040	
		CGGGAAGCTAGAGTAAGTAGTTCGCCAGTTAAT			
G0258991-1.seq(1>7569)	→	CGGGAAGCTAGAGTAAGTAGTTCGCCAGTTAAT			
		2050	2060	2070	
		AGTTTGCGCAACGTTGTTGCCATTGCTACAGGC			
G0258991-1.seq(1>7569)	→	AGTTTGCGCAACGTTGTTGCCATTGCTACAGGC			
		2080	2090	2100	2110
		ATCGTGGTGTACGCTCGTCGTTTGGTATGGCT			
G0258991-1.seq(1>7569)	→	ATCGTGGTGTACGCTCGTCGTTTGGTATGGCT			
		2120	2130	2140	
		TCATTCAGCTCCGGTTCCCAACGATCAAGGCCGA			
G0258991-1.seq(1>7569)	→	TCATTCAGCTCCGGTTCCCAACGATCAAGGCCGA			
		2150	2160	2170	
		GTTACATGATCCCCATGTTGTGCAAAAAAGCG			
G0258991-1.seq(1>7569)	→	GTTACATGATCCCCATGTTGTGCAAAAAAGCG			
		2180	2190	2200	2210
		GTTAGCTCCTTCGGTCCCTCCGATCGTTGTCAGA			
G0258991-1.seq(1>7569)	→	GTTAGCTCCTTCGGTCCCTCCGATCGTTGTCAGA			
		2220	2230	2240	
		AGTAAGTTGGCCGCAGTGTTATCACTCATGGTT			
G0258991-1.seq(1>7569)	→	AGTAAGTTGGCCGCAGTGTTATCACTCATGGTT			
		2250	2260	2270	
		ATGGCAGCACTGCATAATTCTCTTACTGTCATG			
G0258991-1.seq(1>7569)	→	ATGGCAGCACTGCATAATTCTCTTACTGTCATG			
		2280	2290	2300	2310
		CCATCCGTAAGATGCTTTTCTGTGACTGGTGAG			
G0258991-1.seq(1>7569)	→	CCATCCGTAAGATGCTTTTCTGTGACTGGTGAG			
		2320	2330	2340	
		TACTCAACCAAGTCATTCTGAGAATAGTGTATG			
G0258991-1.seq(1>7569)	→	TACTCAACCAAGTCATTCTGAGAATAGTGTATG			
		2350	2360	2370	
		CGGCGACCGAGTTGCTCTTGCCCGGCGTCAATA			
G0258991-1.seq(1>7569)	→	CGGCGACCGAGTTGCTCTTGCCCGGCGTCAATA			

Project: Untitled.SQD Contig 1

		2380	2390	2400	
		CGGGATAATACCGCGCCACATAGCAGAACTTTA			
G0258991-1.seq(1>7569)	→	CGGGATAATACCGCGCCACATAGCAGAACTTTA			
		2410	2420	2430	2440
		AAAGTGCTCATCATTGGAAAACGTTCTTCGGGG			
G0258991-1.seq(1>7569)	→	AAAGTGCTCATCATTGGAAAACGTTCTTCGGGG			
		2450	2460	2470	
		CGAAAAC TCTCAAGGATCTTACCGCTGTTGAGA			
G0258991-1.seq(1>7569)	→	CGAAAAC TCTCAAGGATCTTACCGCTGTTGAGA			
		2480	2490	2500	
		TCCAGTTCGATGTAACCCACTCGTGCACCCAAC			
G0258991-1.seq(1>7569)	→	TCCAGTTCGATGTAACCCACTCGTGCACCCAAC			
		2510	2520	2530	2540
		TGATCTTCAGCATCTTTTACTTTCACCAGCGTT			
G0258991-1.seq(1>7569)	→	TGATCTTCAGCATCTTTTACTTTCACCAGCGTT			
		2550	2560	2570	
		TCTGGGTGAGCAAAAACAGGAAGGCAAAATGCC			
G0258991-1.seq(1>7569)	→	TCTGGGTGAGCAAAAACAGGAAGGCAAAATGCC			
		2580	2590	2600	
		GCAAAAAGGGAATAAGGGCGACACGGAAATGT			
G0258991-1.seq(1>7569)	→	GCAAAAAGGGAATAAGGGCGACACGGAAATGT			
		2610	2620	2630	2640
		TGAATACTCATACTCTTCCTTTTCAATATTAT			
G0258991-1.seq(1>7569)	→	TGAATACTCATACTCTTCCTTTTCAATATTAT			
		2650	2660	2670	
		TGAAGCATT TATCAGGGTTATTGTCTCATGAGC			
G0258991-1.seq(1>7569)	→	TGAAGCATT TATCAGGGTTATTGTCTCATGAGC			
		2680	2690	2700	
		GGATACATATTTGAATGTATTTAGAAAAATAAA			
G0258991-1.seq(1>7569)	→	GGATACATATTTGAATGTATTTAGAAAAATAAA			
		2710	2720	2730	
		CAAATAGGGGTCCGCGCACATTTCCCCGAAAA			
G0258991-1.seq(1>7569)	→	CAAATAGGGGTCCGCGCACATTTCCCCGAAAA			
		2740	2750	2760	2770
		GTGCCACCTGACGTCTAAGAAACCATTATTATC			
G0258991-1.seq(1>7569)	→	GTGCCACCTGACGTCTAAGAAACCATTATTATC			

Project: Untitled.SQD Contig 1

		2780	2790	2800	
		ATGACATTAACCTATAAAAATAGGCGTATCACG			
G0258991-1.seq(1>7569)	→	ATGACATTAACCTATAAAAATAGGCGTATCACG			
		2810	2820	2830	
		AGGCCCTGATGGCTCTTTGCGGCACCCATCGTT			
G0258991-1.seq(1>7569)	→	AGGCCCTGATGGCTCTTTGCGGCACCCATCGTT			
		2840	2850	2860	2870
		CGTAATGTTCCGTGGCACCGAGGACAACCCTCA			
G0258991-1.seq(1>7569)	→	CGTAATGTTCCGTGGCACCGAGGACAACCCTCA			
		2880	2890	2900	
		AGAGAAAATGTAATCACACTGGCTCACCTTCGG			
G0258991-1.seq(1>7569)	→	AGAGAAAATGTAATCACACTGGCTCACCTTCGG			
		2910	2920	2930	
		GTGGGCCTTTCTGCGTTTATAAGGAGACACTTT			
G0258991-1.seq(1>7569)	→	GTGGGCCTTTCTGCGTTTATAAGGAGACACTTT			
		2940	2950	2960	2970
		ATGTTTAAGAAGGTTGGTAAATTCCTTGCGGCT			
G0258991-1.seq(1>7569)	→	ATGTTTAAGAAGGTTGGTAAATTCCTTGCGGCT			
		2980	2990	3000	
		TTGGCAGCCAAGCTAGATCCGGCTGTGGAATGT			
G0258991-1.seq(1>7569)	→	TTGGCAGCCAAGCTAGATCCGGCTGTGGAATGT			
		3010	3020	3030	
		GTGTCAGTTAGGGTGTGGAAAGTCCCCAGGCTC			
G0258991-1.seq(1>7569)	→	GTGTCAGTTAGGGTGTGGAAAGTCCCCAGGCTC			
		3040	3050	3060	
		CCCAGCAGGCAGAAGTATGCAAAGCATGCATCT			
G0258991-1.seq(1>7569)	→	CCCAGCAGGCAGAAGTATGCAAAGCATGCATCT			
		3070	3080	3090	3100
		CAATTAGTCAGCAACCAGGTGTGGAAAGTCCCC			
G0258991-1.seq(1>7569)	→	CAATTAGTCAGCAACCAGGTGTGGAAAGTCCCC			
		3110	3120	3130	
		AGGCTCCCCAGCAGGCAGAAGTATGCAAAGCAT			
G0258991-1.seq(1>7569)	→	AGGCTCCCCAGCAGGCAGAAGTATGCAAAGCAT			
		3140	3150	3160	
		GCATCTCAATTAGTCAGCAACCATAGTCCCGCC			
G0258991-1.seq(1>7569)	→	GCATCTCAATTAGTCAGCAACCATAGTCCCGCC			

Project: Untitled.SQD Contig 1

		3170	3180	3190	3200
		CCTAACTCCGCCATCCCGCCCCTAACTCCGCC			
G0258991-1.seq(1>7569)	→	CCTAACTCCGCCATCCCGCCCCTAACTCCGCC			
		3210	3220	3230	
		CAGTTCCGCCATTCTCCGCCCATGGCTGACT			
G0258991-1.seq(1>7569)	→	CAGTTCCGCCATTCTCCGCCCATGGCTGACT			
		3240	3250	3260	
		AATTTTATTTATTTATGCAGAGGCCGAGGCCGC			
G0258991-1.seq(1>7569)	→	AATTTTATTTATTTATGCAGAGGCCGAGGCCGC			
		3270	3280	3290	3300
		CTCGGCCTCTGAGCTATTCCAGAAGTAGTGAGG			
G0258991-1.seq(1>7569)	→	CTCGGCCTCTGAGCTATTCCAGAAGTAGTGAGG			
		3310	3320	3330	
		AGGCTTTTTTGGAGGCCTAGGCTTTTTGCAAAA			
G0258991-1.seq(1>7569)	→	AGGCTTTTTTGGAGGCCTAGGCTTTTTGCAAAA			
		3340	3350	3360	
		GCTAGCTTGGGGCCACCGCTCAGAGCACCTTCC			
G0258991-1.seq(1>7569)	→	GCTAGCTTGGGGCCACCGCTCAGAGCACCTTCC			
		3370	3380	3390	
		ACCATGGCCACCTCAGCAAGTTCCCACTTGAAC			
G0258991-1.seq(1>7569)	→	ACCATGGCCACCTCAGCAAGTTCCCACTTGAAC			
		3400	3410	3420	3430
		AAAAACATCAAGCAAATGTACTTGTGCCTGCC			
G0258991-1.seq(1>7569)	→	AAAAACATCAAGCAAATGTACTTGTGCCTGCC			
		3440	3450	3460	
		CAGGGTGAGAAAGTCCAAGCCATGTATATCTGG			
G0258991-1.seq(1>7569)	→	CAGGGTGAGAAAGTCCAAGCCATGTATATCTGG			
		3470	3480	3490	
		GTTGATGGTACTGGAGAAGGACTGCGCTGCAA			
G0258991-1.seq(1>7569)	→	GTTGATGGTACTGGAGAAGGACTGCGCTGCAA			
		3500	3510	3520	3530
		ACCCGCACCCTGGACTGTGAGCCCAAGTGTGTA			
G0258991-1.seq(1>7569)	→	ACCCGCACCCTGGACTGTGAGCCCAAGTGTGTA			
		3540	3550	3560	
		GAAGAGTTACCTGAGTGAATTTTGATGGCTCT			
G0258991-1.seq(1>7569)	→	GAAGAGTTACCTGAGTGAATTTTGATGGCTCT			

Project: Untitled.SQD Contig 1

		3570 3580 3590	AGTACCTTTTCAGTCTGAGGGCTCCAACAGTGAC
G0258991-1.seq(1>7569)	→		AGTACCTTTTCAGTCTGAGGGCTCCAACAGTGAC
		3600 3610 3620 3630	ATGTATCTCAGCCCTGTTGCCATGTTTCGGGAC
G0258991-1.seq(1>7569)	→		ATGTATCTCAGCCCTGTTGCCATGTTTCGGGAC
		3640 3650 3660	CCCTTCCGCAGAGATCCAACAAGCTGGTGTTC
G0258991-1.seq(1>7569)	→		CCCTTCCGCAGAGATCCAACAAGCTGGTGTTC
		3670 3680 3690	TGTGAAGTTTTCAAGTACAACCGGAAGCCTGCA
G0258991-1.seq(1>7569)	→		TGTGAAGTTTTCAAGTACAACCGGAAGCCTGCA
		3700 3710 3720	GAGACCAATTTAAGGCACTCGTGTAACGGATA
G0258991-1.seq(1>7569)	→		GAGACCAATTTAAGGCACTCGTGTAACGGATA
		3730 3740 3750 3760	ATGGACATGGTGAGCAACCAGCACCCCTGGTTT
G0258991-1.seq(1>7569)	→		ATGGACATGGTGAGCAACCAGCACCCCTGGTTT
		3770 3780 3790	GGAATGGAACAGGAGTATACTCTGATGGGAACA
G0258991-1.seq(1>7569)	→		GGAATGGAACAGGAGTATACTCTGATGGGAACA
		3800 3810 3820	GATGGGCACCCTTTTGGTTGGCCTTCCAATGGC
G0258991-1.seq(1>7569)	→		GATGGGCACCCTTTTGGTTGGCCTTCCAATGGC
		3830 3840 3850 3860	TTTCTGGGCCCAAGGTCCGTATTACTGTGGT
G0258991-1.seq(1>7569)	→		TTTCTGGGCCCAAGGTCCGTATTACTGTGGT
		3870 3880 3890	GTGGGCGCAGACAAAGCCTATGGCAGGGATATC
G0258991-1.seq(1>7569)	→		GTGGGCGCAGACAAAGCCTATGGCAGGGATATC
		3900 3910 3920	GTGGAGGCTCACTACCGCGCCTGCTTGTATGCT
G0258991-1.seq(1>7569)	→		GTGGAGGCTCACTACCGCGCCTGCTTGTATGCT
		3930 3940 3950 3960	GGGGTCAAGATTACAGGAACAAATGCTGAGGTC
G0258991-1.seq(1>7569)	→		GGGGTCAAGATTACAGGAACAAATGCTGAGGTC

Project: Untitled.SQD Contig 1

		3970	3980	3990	
		ATGCCTGCCCAGTGGGAGTTCCAAATAGGACCC			
G0258991-1.seq(1>7569)	→	ATGCCTGCCCAGTGGGAGTTCCAAATAGGACCC			
		4000	4010	4020	
		TGTGAAGGAATCCGCATGGGAGATCATCTCTGG			
G0258991-1.seq(1>7569)	→	TGTGAAGGAATCCGCATGGGAGATCATCTCTGG			
		4030	4040	4050	
		GTGGCCCGTTTCATCTTGCATCGAGTATGTGAA			
G0258991-1.seq(1>7569)	→	GTGGCCCGTTTCATCTTGCATCGAGTATGTGAA			
		4060	4070	4080	4090
		GACTTTGGGGTAATAGCAACCTTTGACCCCAAG			
G0258991-1.seq(1>7569)	→	GACTTTGGGGTAATAGCAACCTTTGACCCCAAG			
		4100	4110	4120	
		CCCATTCTGGGAACTGGAATGGTGCAGGCTGC			
G0258991-1.seq(1>7569)	→	CCCATTCTGGGAACTGGAATGGTGCAGGCTGC			
		4130	4140	4150	
		CATACCAACTTTAGCACCAAGGCCATGCGGGAG			
G0258991-1.seq(1>7569)	→	CATACCAACTTTAGCACCAAGGCCATGCGGGAG			
		4160	4170	4180	4190
		GAGAATGGTCTGAAGCACATCGAGGAGGCCATC			
G0258991-1.seq(1>7569)	→	GAGAATGGTCTGAAGCACATCGAGGAGGCCATC			
		4200	4210	4220	
		GAGAACTAAGCAAGCGGCACCGGTACCACATT			
G0258991-1.seq(1>7569)	→	GAGAACTAAGCAAGCGGCACCGGTACCACATT			
		4230	4240	4250	
		CGAGCCTACGATCCCAAGGGGGCCTGGACAAT			
G0258991-1.seq(1>7569)	→	CGAGCCTACGATCCCAAGGGGGCCTGGACAAT			
		4260	4270	4280	4290
		GCCCGTCGTCTGACTGGGTTCCACGAAACGTCC			
G0258991-1.seq(1>7569)	→	GCCCGTCGTCTGACTGGGTTCCACGAAACGTCC			
		4300	4310	4320	
		AACATCAACGACTTTTCTGCTGGTGTGCGCAAT			
G0258991-1.seq(1>7569)	→	AACATCAACGACTTTTCTGCTGGTGTGCGCAAT			
		4330	4340	4350	
		CGCAGTGCCAGCATCCGCATTCCCCGGACTGTC			
G0258991-1.seq(1>7569)	→	CGCAGTGCCAGCATCCGCATTCCCCGGACTGTC			

Project: Untitled.SQD Contig 1

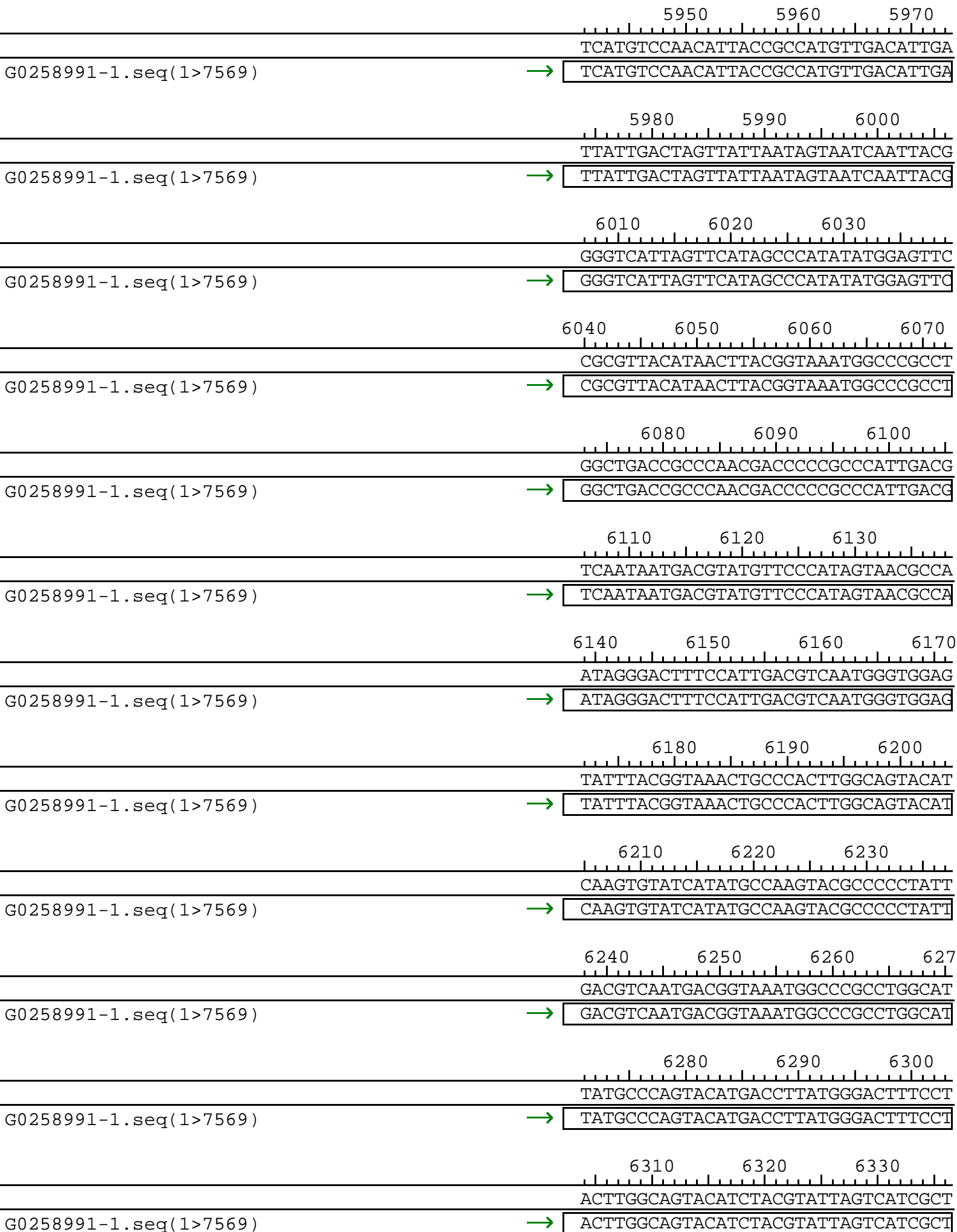
		4760	4770	4780	
		TTTGCTCAGAAGAAATGCCATCTAGTGATGATG			
G0258991-1.seq(1>7569)	→	TTTGCTCAGAAGAAATGCCATCTAGTGATGATG			
		4790	4800	4810	
		AGGCTACTGCTGACTCTCAACATTCTACTCCTC			
G0258991-1.seq(1>7569)	→	AGGCTACTGCTGACTCTCAACATTCTACTCCTC			
		4820	4830	4840	4850
		CAAAAAGAAGAGAAAGGTAGAAGACCCCAAGG			
G0258991-1.seq(1>7569)	→	CAAAAAGAAGAGAAAGGTAGAAGACCCCAAGG			
		4860	4870	4880	
		ACTTTCCTTCAGAATTGCTAAGTTTTTTGAGTC			
G0258991-1.seq(1>7569)	→	ACTTTCCTTCAGAATTGCTAAGTTTTTTGAGTC			
		4890	4900	4910	
		ATGCTGTGTTTAGTAATAGAACTCTTGCTTGCT			
G0258991-1.seq(1>7569)	→	ATGCTGTGTTTAGTAATAGAACTCTTGCTTGCT			
		4920	4930	4940	4950
		TTGCTATTTACACCACAAAGGAAAAAGCTGCAC			
G0258991-1.seq(1>7569)	→	TTGCTATTTACACCACAAAGGAAAAAGCTGCAC			
		4960	4970	4980	
		TGCTATACAAGAAAATTATGGAAAAATATTCTG			
G0258991-1.seq(1>7569)	→	TGCTATACAAGAAAATTATGGAAAAATATTCTG			
		4990	5000	5010	
		TAACCTTTATAAGTAGGCATAACAGTTATAATC			
G0258991-1.seq(1>7569)	→	TAACCTTTATAAGTAGGCATAACAGTTATAATC			
		5020	5030	5040	
		ATAACATACTGTTTTTTCTTACTCCACACAGGC			
G0258991-1.seq(1>7569)	→	ATAACATACTGTTTTTTCTTACTCCACACAGGC			
		5050	5060	5070	5080
		ATAGAGTGTCTGCTATTAATAACTATGCTCAA			
G0258991-1.seq(1>7569)	→	ATAGAGTGTCTGCTATTAATAACTATGCTCAA			
		5090	5100	5110	
		AATTGTGTACCTTTAGCTTTTTTAATTTGTAAAG			
G0258991-1.seq(1>7569)	→	AATTGTGTACCTTTAGCTTTTTTAATTTGTAAAG			
		5120	5130	5140	
		GGGTTAATAAGGAATATTTGATGTATAGTGCCT			
G0258991-1.seq(1>7569)	→	GGGTTAATAAGGAATATTTGATGTATAGTGCCT			

		5150	5160	5170	5180
		TGACTAGAGATCATAATCAGCCATAACCACATTT			
G0258991-1.seq(1>7569)	→	TGACTAGAGATCATAATCAGCCATAACCACATTT			
		5190	5200	5210	
		GTAGAGGTTTTACTTGCTTTAAAAAACCTCCCA			
G0258991-1.seq(1>7569)	→	GTAGAGGTTTTACTTGCTTTAAAAAACCTCCCA			
		5220	5230	5240	
		CACCTCCCCCTGAACCTGAAACATAAAATGAAT			
G0258991-1.seq(1>7569)	→	CACCTCCCCCTGAACCTGAAACATAAAATGAAT			
		5250	5260	5270	528
		GCAATTGTTGTTGTTAACTTGTTTATTGCAGCT			
G0258991-1.seq(1>7569)	→	GCAATTGTTGTTGTTAACTTGTTTATTGCAGCT			
		5290	5300	5310	
		TATAATGGTTACAAATAAAGCAATAGCATCACA			
G0258991-1.seq(1>7569)	→	TATAATGGTTACAAATAAAGCAATAGCATCACA			
		5320	5330	5340	
		AATTTACAAATAAAGCATTTTTTTTCCTACTGCAT			
G0258991-1.seq(1>7569)	→	AATTTACAAATAAAGCATTTTTTTTCCTACTGCAT			
		5350	5360	5370	
		TCTAGTTGTGGTTTGTCCAACTCATCAATGTA			
G0258991-1.seq(1>7569)	→	TCTAGTTGTGGTTTGTCCAACTCATCAATGTA			
		5380	5390	5400	5410
		TCTTATCATGTCTGGATCTAGCTTCGTGTCAAG			
G0258991-1.seq(1>7569)	→	TCTTATCATGTCTGGATCTAGCTTCGTGTCAAG			
		5420	5430	5440	
		GACGGTGAAGTGCAGTGAATAATAAAATGTGTGT			
G0258991-1.seq(1>7569)	→	GACGGTGAAGTGCAGTGAATAATAAAATGTGTGT			
		5450	5460	5470	
		TTGTCCGAAATACGCGTTTTGAGATTTCTGTCC			
G0258991-1.seq(1>7569)	→	TTGTCCGAAATACGCGTTTTGAGATTTCTGTCC			
		5480	5490	5500	5510
		CCGACTAAATTCATGTGCGCGGATAGTGGTGT			
G0258991-1.seq(1>7569)	→	CCGACTAAATTCATGTGCGCGGATAGTGGTGT			
		5520	5530	5540	
		TATCGCCGATAGAGATGGCGATATTGGAAAAAT			
G0258991-1.seq(1>7569)	→	TATCGCCGATAGAGATGGCGATATTGGAAAAAT			

Project: Untitled.SQD Contig 1

		5550 5560 5570	CGATATTTGAAAATATGGCATATTGAAAATGTC
G0258991-1.seq(1>7569)	→		CGATATTTGAAAATATGGCATATTGAAAATGTC
		5580 5590 5600 5610	GCCGATGTGAGTTTCTGTGTAACTGATATCGCC
G0258991-1.seq(1>7569)	→		GCCGATGTGAGTTTCTGTGTAACTGATATCGCC
		5620 5630 5640	ATTTTTCCAAAGTGATTTTTGGGCATACGCGA
G0258991-1.seq(1>7569)	→		ATTTTTCCAAAGTGATTTTTGGGCATACGCGA
		5650 5660 5670	TATCTGGCGATAGCGCTTATATCGTTTACGGGG
G0258991-1.seq(1>7569)	→		TATCTGGCGATAGCGCTTATATCGTTTACGGGG
		5680 5690 5700	GATGGCGATAGACGACTTTGGTGACTTGGGCGA
G0258991-1.seq(1>7569)	→		GATGGCGATAGACGACTTTGGTGACTTGGGCGA
		5710 5720 5730 5740	TTCTGTGTGTCGCAAATATCGCAGTTTCGATAT
G0258991-1.seq(1>7569)	→		TTCTGTGTGTCGCAAATATCGCAGTTTCGATAT
		5750 5760 5770	AGGTGACAGACGATATGAGGCTATATCGCCGAT
G0258991-1.seq(1>7569)	→		AGGTGACAGACGATATGAGGCTATATCGCCGAT
		5780 5790 5800	AGAGGCGACATCAAGCTGGCACATGGCCAATGC
G0258991-1.seq(1>7569)	→		AGAGGCGACATCAAGCTGGCACATGGCCAATGC
		5810 5820 5830 5840	ATATCGATCTATACATTGAATCAATATTGGCCA
G0258991-1.seq(1>7569)	→		ATATCGATCTATACATTGAATCAATATTGGCCA
		5850 5860 5870	TTAGCCATATTATTCATTGGTTATATAGCATAA
G0258991-1.seq(1>7569)	→		TTAGCCATATTATTCATTGGTTATATAGCATAA
		5880 5890 5900	ATCAATATTGGCTATTGGCCATTGCATACGTTG
G0258991-1.seq(1>7569)	→		ATCAATATTGGCTATTGGCCATTGCATACGTTG
		5910 5920 5930 5940	TATCCATATCATAAATATGTACATTTATATTGGC
G0258991-1.seq(1>7569)	→		TATCCATATCATAAATATGTACATTTATATTGGC

Project: Untitled.SQD Contig 1



Project: Untitled.SQD Contig 1

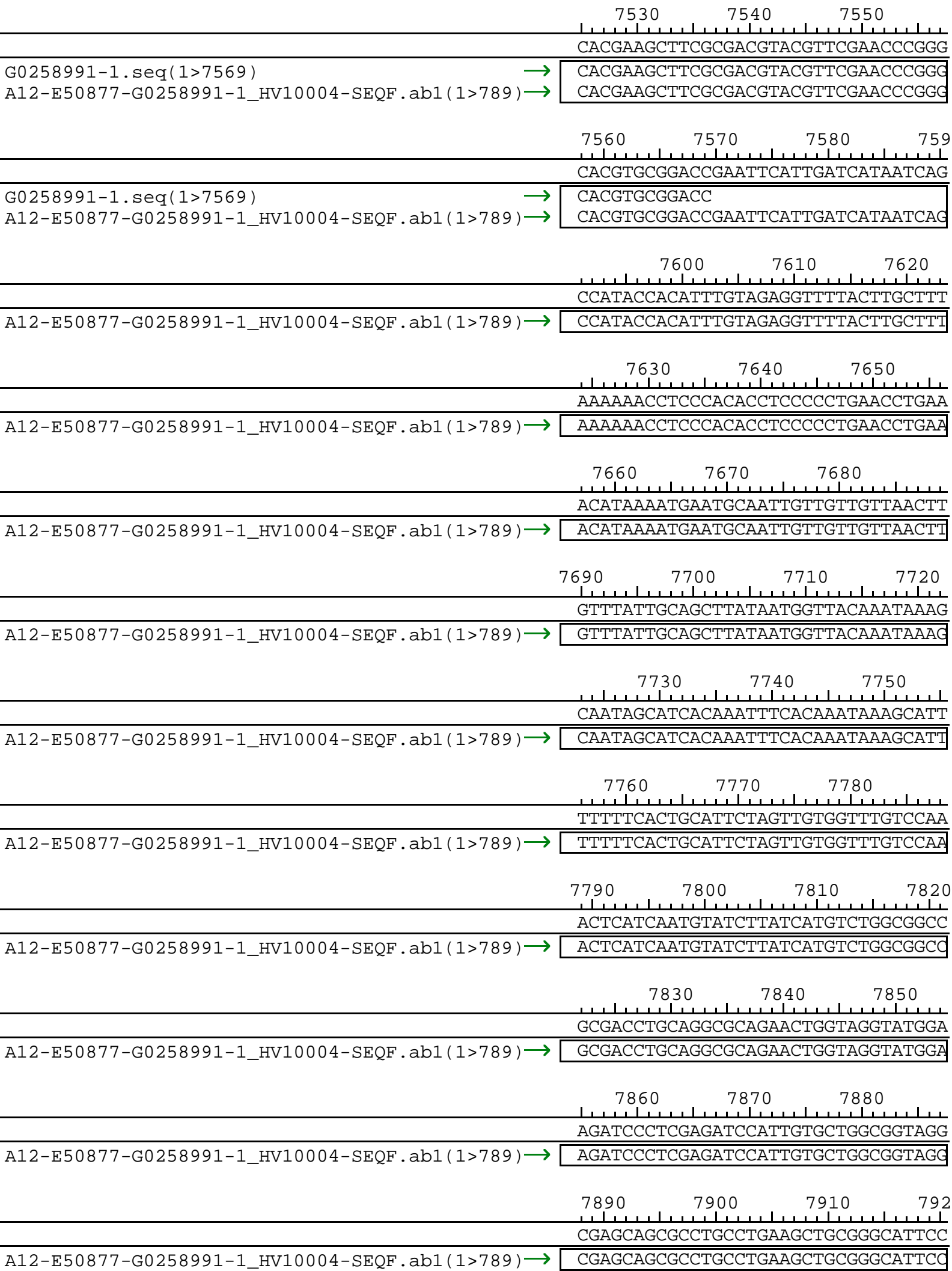
		6340	6350	6360	ATTACCATGGTGATGCGGTTTTGGCAGTACATC	
G0258991-1.seq(1>7569)	→	ATTACCATGGTGATGCGGTTTTGGCAGTACATC				
		6370	6380	6390	6400	AATGGGCGTGGATAGCGGTTTGACTCACGGGGA
G0258991-1.seq(1>7569)	→	AATGGGCGTGGATAGCGGTTTGACTCACGGGGA				
		6410	6420	6430	TTTCCAAGTCTCACCCATTGACGTCAATGGG	
G0258991-1.seq(1>7569)	→	TTTCCAAGTCTCACCCATTGACGTCAATGGG				
		6440	6450	6460	AGTTTGTTTTGGCACCAAATCAACGGGACTTT	
G0258991-1.seq(1>7569)	→	AGTTTGTTTTGGCACCAAATCAACGGGACTTT				
		6470	6480	6490	6500	CAAATGGTTCGTAACAACCTCCGCCCCATTGACG
G0258991-1.seq(1>7569)	→	CAAATGGTTCGTAACAACCTCCGCCCCATTGACG				
		6510	6520	6530	CAAATGGGCGGTAGGCGTGTACGGTGGGAGGTC	
G0258991-1.seq(1>7569)	→	CAAATGGGCGGTAGGCGTGTACGGTGGGAGGTC				
		6540	6550	6560	TATATAAGCAGAGCTCGTTTAGTGAACCGTCAG	
G0258991-1.seq(1>7569)	→	TATATAAGCAGAGCTCGTTTAGTGAACCGTCAG				
		6570	6580	6590	6600	ATCGCCTGGAGACGCCATCCACGCTGTTTGGAC
G0258991-1.seq(1>7569)	→	ATCGCCTGGAGACGCCATCCACGCTGTTTGGAC				
		6610	6620	6630	CTCCATAGAAGACACCGGACCGATCCAGCCTC	
G0258991-1.seq(1>7569)	→	CTCCATAGAAGACACCGGACCGATCCAGCCTC				
		6640	6650	6660	CGCGGCCGGGAACGGTGCATTGGAACGCGGATT	
G0258991-1.seq(1>7569)	→	CGCGGCCGGGAACGGTGCATTGGAACGCGGATT				
		6670	6680	6690	CCCCGTGCCAAGAGTGACGTAAGTACCGCCTAT	
G0258991-1.seq(1>7569)	→	CCCCGTGCCAAGAGTGACGTAAGTACCGCCTAT				
		6700	6710	6720	6730	AGAGTCTATAGGCCACCCCCTTGGCTTCTTAT
G0258991-1.seq(1>7569)	→	AGAGTCTATAGGCCACCCCCTTGGCTTCTTAT				

		6740	6750	6760	
		GCATGCTATACTGTTTTGGCTTGGGGTCTATA			
G0258991-1.seq(1>7569)	→	GCATGCTATACTGTTTTGGCTTGGGGTCTATA			
		6770	6780	6790	
		CACCCCCGCTTCCTCATGTTATAGGTGATGGTA			
G0258991-1.seq(1>7569)	→	CACCCCCGCTTCCTCATGTTATAGGTGATGGTA			
		6800	6810	6820	6830
		TAGCTTAGCCTATAGGTGTGGGTATTGACCAT			
G0258991-1.seq(1>7569)	→	TAGCTTAGCCTATAGGTGTGGGTATTGACCAT			
		6840	6850	6860	
		TATTGACCACTCCCCTATTGGTGACGATACTTT			
G0258991-1.seq(1>7569)	→	TATTGACCACTCCCCTATTGGTGACGATACTTT			
		6870	6880	6890	
		CCATTACTAATCCATAACATGGCTCTTTGCCAC			
G0258991-1.seq(1>7569)	→	CCATTACTAATCCATAACATGGCTCTTTGCCAC			
		6900	6910	6920	6930
		AACTCTCTTTATTGGCTATATGCCAATACTG			
G0258991-1.seq(1>7569)	→	AACTCTCTTTATTGGCTATATGCCAATACTG			
		6940	6950	6960	
		TCCTTCAGAGACTGACACGGACTCTGTATTTTT			
G0258991-1.seq(1>7569)	→	TCCTTCAGAGACTGACACGGACTCTGTATTTTT			
		6970	6980	6990	
		ACAGGATGGGGTCTCATTTATTATTTACAAATT			
G0258991-1.seq(1>7569)	→	ACAGGATGGGGTCTCATTTATTATTTACAAATT			
		7000	7010	7020	
		CACATATAACAACACCACCGTCCCCAGTGCCCGC			
G0258991-1.seq(1>7569)	→	CACATATAACAACACCACCGTCCCCAGTGCCCGC			
		7030	7040	7050	7060
		AGTTTTTATTAAACATAACGTGGGATCTCCACG			
G0258991-1.seq(1>7569)	→	AGTTTTTATTAAACATAACGTGGGATCTCCACG			
		7070	7080	7090	
		CGAATCTCGGGTACGTGTTCCGGACATGGGCTC			
G0258991-1.seq(1>7569)	→	CGAATCTCGGGTACGTGTTCCGGACATGGGCTC			
		7100	7110	7120	
		TTCTCCGGTAGCGGCGGAGCTTCTACATCCGAG			
G0258991-1.seq(1>7569)	→	TTCTCCGGTAGCGGCGGAGCTTCTACATCCGAG			

Project: Untitled.SQD Contig 1

		7130	7140	7150	7160
		CCCTGCTCCCATGCCTCCAGCGACTCATGGTCC			
G0258991-1.seq(1>7569)	→	CCCTGCTCCCATGCCTCCAGCGACTCATGGTCC			
		7170	7180	7190	
		CTCGGCAGCTCCTTGCTCCTAACAGTGGAGGCC			
G0258991-1.seq(1>7569)	→	CTCGGCAGCTCCTTGCTCCTAACAGTGGAGGCC			
		7200	7210	7220	
		AGACTTAGGCACAGCACGATGCCACCACCACC			
G0258991-1.seq(1>7569)	→	AGACTTAGGCACAGCACGATGCCACCACCACC			
		7230	7240	7250	7260
		AGTGTGCCGCACAAGGCCGTGGCGGTAGGGTAT			
G0258991-1.seq(1>7569)	→	AGTGTGCCGCACAAGGCCGTGGCGGTAGGGTAT			
		7270	7280	7290	
		GTGTCTGAAAATGAGCTCGGGGAGCGGGCTTGC			
G0258991-1.seq(1>7569)	→	GTGTCTGAAAATGAGCTCGGGGAGCGGGCTTGC			
		7300	7310	7320	
		ACCGCTGACGCATTTGGAAGACTTAAGGCAGCG			
G0258991-1.seq(1>7569)	→	ACCGCTGACGCATTTGGAAGACTTAAGGCAGCG			
		7330	7340	7350	
		GCAGAAGAAGATGCAGGCAGCTGAGTTGTTGTG			
G0258991-1.seq(1>7569)	→	GCAGAAGAAGATGCAGGCAGCTGAGTTGTTGTG			
		7360	7370	7380	7390
		TTCTGATAAGAGTCAGAGGTAACTCCCGTTGCC			
G0258991-1.seq(1>7569)	→	TTCTGATAAGAGTCAGAGGTAACTCCCGTTGCC			
		7400	7410	7420	
		GTGCTGTTAACGGTGGAGGGCAGTGTAGTCTGA			
G0258991-1.seq(1>7569)	→	GTGCTGTTAACGGTGGAGGGCAGTGTAGTCTGA			
		7430	7440	7450	
		GCAGTACTCGTTGCTGCCGCGCGGCCACCAGA			
G0258991-1.seq(1>7569)	→	GCAGTACTCGTTGCTGCCGCGCGGCCACCAGA			
		7460	7470	7480	7490
		CATAATAGCTGACAGACTAACAGACTGTTTCCTT			
G0258991-1.seq(1>7569)	→	CATAATAGCTGACAGACTAACAGACTGTTTCCTT			
A12-E50877-G0258991-1_HV10004-SEQF.ab1(1>789)	→	GCTGACAGACTAACAGACTGTTTCCTT			
		7500	7510	7520	
		TCCATGGGTCTTTTCTGCAGTCACCGTCCTTGA			
G0258991-1.seq(1>7569)	→	TCCATGGGTCTTTTCTGCAGTCACCGTCCTTGA			
A12-E50877-G0258991-1_HV10004-SEQF.ab1(1>789)	→	TCCATGGGTCTTTTCTGCAGTCACCGTCCTTGA			

Project: Untitled.SQD Contig 1



	7930 7940 7950 CGATCAGAAATGAGCGCCAGTCGTCGTCGGCTC
A12-E50877-G0258991-1_HV10004-SEQF.ab1 (1>789) →	CGATCAGAAATGAGCGCCAGTCGTCGTCGGCTC
	7960 7970 7980 TCGGCACCGAATGCGTATGATTCTCCGCCAGCA
A12-E50877-G0258991-1_HV10004-SEQF.ab1 (1>789) →	TCGGCACCGAATGCGTATGATTCTCCGCCAGCA
	7990 8000 8010 TGGCTTCGGCCAGTGCCTCGAGCAGCGCCCGCT
A12-E50877-G0258991-1_HV10004-SEQF.ab1 (1>789) →	TGGCTTCGGCCAGTGCCTCGAGCAGCGCCCGCT
	8020 8030 8040 8050 TGTTCTGAAGTGCCAGTAAAGCGCCGGCTGCT
A12-E50877-G0258991-1_HV10004-SEQF.ab1 (1>789) →	TGTTCTGAAGTGCCAGTAAAGCGCCGGCTGCT
	8060 8070 8080 GAACCCCAACCGTTCCGCCAGTTTGCGTGTCC
A12-E50877-G0258991-1_HV10004-SEQF.ab1 (1>789) →	GAACCCCAACCGTTCCGCCAGTTTGCGTGTCC
	8090 8100 8110 TCAGACCGTCTACGCCGACCTCGTTCAACAGGT
A12-E50877-G0258991-1_HV10004-SEQF.ab1 (1>789) →	TCAGACCGTCTACGCCGACCTCGTTCAACAGGT
	8120 8130 8140 8150 CCAGGGCGGCACGGATCACTGTATTCGGCTGCA
A12-E50877-G0258991-1_HV10004-SEQF.ab1 (1>789) →	CCAGGGCGGCACGGATCACTGTATTCGGCTGCA
	8160 8170 8180 ACTTTGTCATGCTTGACACTTTTATCACTGATAA
A12-E50877-G0258991-1_HV10004-SEQF.ab1 (1>789) →	ACTTTGTCATGCTTGACACTTTTATCACTGATAA
	8190 8200 8210 ACATAATATGTCCACCAACTTATCAGTGATAAA
A12-E50877-G0258991-1_HV10004-SEQF.ab1 (1>789) →	ACATAATATGTCCACCAACTTATCAGTGATAAA
	8220 8230 8240 825 GAATCCGCGCCAGCACAATGGATCTCGAGGTCC
A12-E50877-G0258991-1_HV10004-SEQF.ab1 (1>789) →	GAATCCGCGCCAGCACAATGGATCTCGAGGTCC
	AGGG
A12-E50877-G0258991-1_HV10004-SEQF.ab1 (1>789) →	AGGG