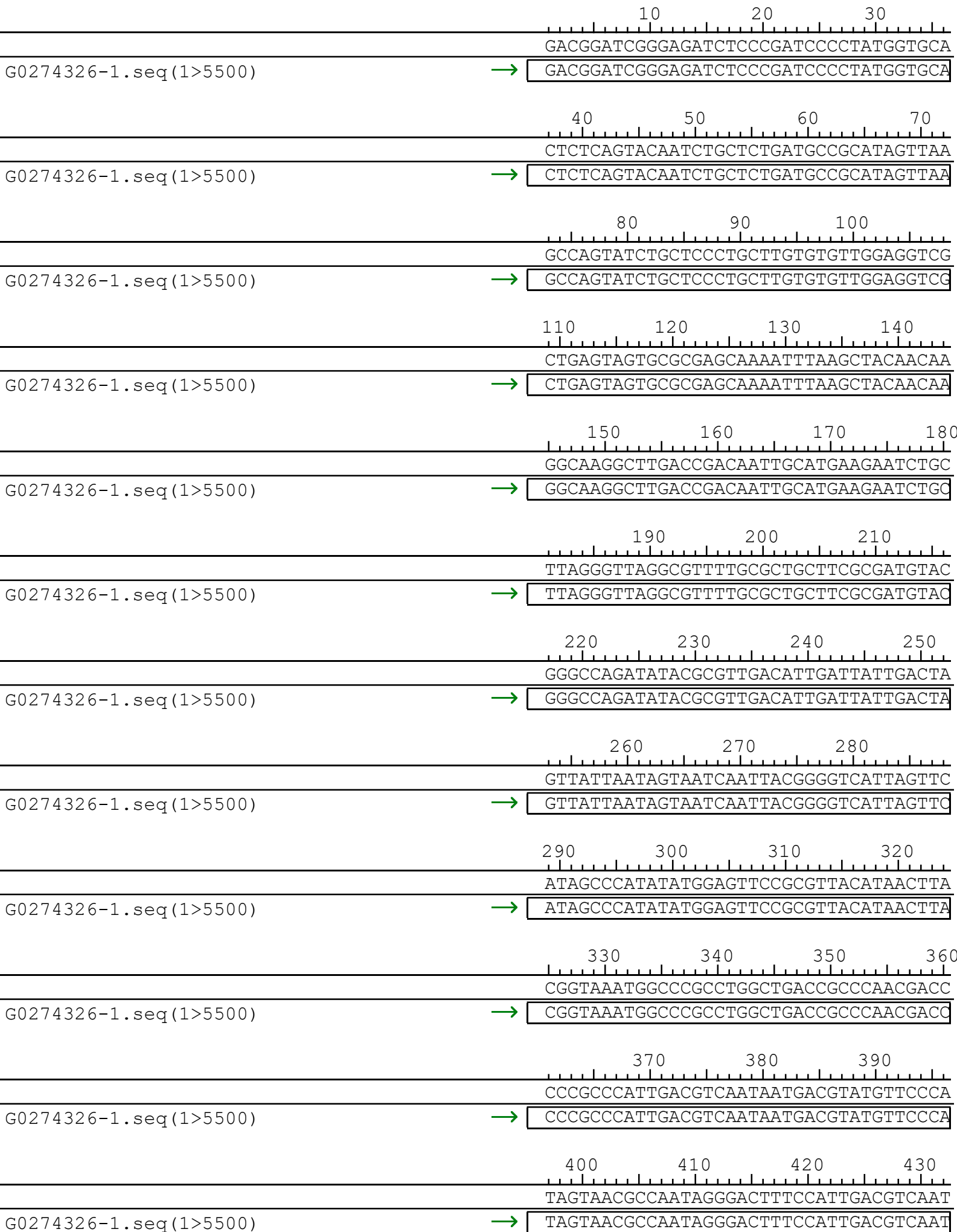


Project: Untitled.SQD Contig 1



Project: Untitled.SQD Contig 1

		440 450 460	GGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAG
G0274326-1.seq (1>5500)	→		GGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAG
		470 480 490 500	TACATCAAGTGTATCATATGCCAAGTACGCCCCCTA
G0274326-1.seq (1>5500)	→		TACATCAAGTGTATCATATGCCAAGTACGCCCCCTA
		510 520 530 540	TTGACGTCAATGACGGTAAATGGCCCGCCTGGCATT
G0274326-1.seq (1>5500)	→		TTGACGTCAATGACGGTAAATGGCCCGCCTGGCATT
		550 560 570	ATGCCCAGTACATGACCTTATGGGACTTTCCTACTT
G0274326-1.seq (1>5500)	→		ATGCCCAGTACATGACCTTATGGGACTTTCCTACTT
		580 590 600 610	GGCAGTACATCTACGTATTAGTCATCGCTATTACCA
G0274326-1.seq (1>5500)	→		GGCAGTACATCTACGTATTAGTCATCGCTATTACCA
		620 630 640	TGGTGATGCGGTTTTGGCAGTACATCAATGGGCGTG
G0274326-1.seq (1>5500)	→		TGGTGATGCGGTTTTGGCAGTACATCAATGGGCGTG
		650 660 670 680	GATAGCGGTTTGACTCACGGGGATTTCCAAGTCTCC
G0274326-1.seq (1>5500)	→		GATAGCGGTTTGACTCACGGGGATTTCCAAGTCTCC
		690 700 710 720	ACCCCATTTGACGTCAATGGGAGTTTGTTTTGGCACC
G0274326-1.seq (1>5500)	→		ACCCCATTTGACGTCAATGGGAGTTTGTTTTGGCACC
		730 740 750	AAAATCAACGGGACTTTCAAAATGTCGTAACAAC
G0274326-1.seq (1>5500)	→		AAAATCAACGGGACTTTCAAAATGTCGTAACAAC
		760 770 780 790	CCGCCCCATTGACGCAAATGGGCGGTAGGCGTGTAC
G0274326-1.seq (1>5500)	→		CCGCCCCATTGACGCAAATGGGCGGTAGGCGTGTAC
		800 810 820	GGTGGGAGGTCTATATAAGCAGAGCTCTCTGGCTAA
G0274326-1.seq (1>5500)	→		GGTGGGAGGTCTATATAAGCAGAGCTCTCTGGCTAA
		830 840 850 860	CTAGAGAACCCACTGCTTACTGGCTTATCGAAATTA
G0274326-1.seq (1>5500)	→		CTAGAGAACCCACTGCTTACTGGCTTATCGAAATTA
E06-E61664-G0274326-1_PEGFP-N-5.ab1 (1>792)	→		TGGCTTATCGAAATTA

Project: Untitled.SQD Contig 1

		870 880 890 900	ATACGACTCACTATAGGGAGACCCAAGCTGGCTAGT
G0274326-1.seq(1>5500)	→		ATACGACTCACTATAGGGAGACCCAAGCTGGCTAGT
E06-E61664-G0274326-1_PEGFP-N-5.ab1(1>792)	→		ATACGACTCACTATAGGGAGACCCAAGCTGGCTAGT
		910 920 930	TAAGCTTGGTACCGAGCTCGGATCCACTAGTCCAGT
G0274326-1.seq(1>5500)	→		TAAGCTTGGTACCGAGCTCGGATCCACTAGTCCAGT
E06-E61664-G0274326-1_PEGFP-N-5.ab1(1>792)	→		TAAGCTTGGTACCGAGCTCGGATCCACTAGTCCAGT
		940 950 960 970	GTGGTGGAATTCTGCAGATATCCAGCACAGTGGCGG
G0274326-1.seq(1>5500)	→		GTGGTGGAATTCTGCAGATATCCAGCACAGTGGCGG
E06-E61664-G0274326-1_PEGFP-N-5.ab1(1>792)	→		GTGGTGGAATTCTGCAGATATCCAGCACAGTGGCGG
		980 990 1000	CCGCTCGAGTCTAGAGGGCCCTTCGAAGGTAAGCCT
G0274326-1.seq(1>5500)	→		CCGCTCGAGTCTAGAGGGCCCTTCGAAGGTAAGCCT
E06-E61664-G0274326-1_PEGFP-N-5.ab1(1>792)	→		CCGCTCGAGTCTAGAGGGCCCTTCGAAGGTAAGCCT
		1010 1020 1030 1040	ATCCCTAACCCCTCTCCTCGGTCTCGATTCTACGCGT
G0274326-1.seq(1>5500)	→		ATCCCTAACCCCTCTCCTCGGTCTCGATTCTACGCGT
E06-E61664-G0274326-1_PEGFP-N-5.ab1(1>792)	→		ATCCCTAACCCCTCTCCTCGGTCTCGATTCTACGCGT
		1050 1060 1070 1080	ACCGGTCATCATCACCATCACCATTGAGTTTAAACC
G0274326-1.seq(1>5500)	→		ACCGGTCATCATCACCATCACCATTGAGTTTAAACC
E06-E61664-G0274326-1_PEGFP-N-5.ab1(1>792)	→		ACCGGTCATCATCACCATCACCATTGAGTTTAAACC
		1090 1100 1110	CGCTGATCAGCCTCGACTGTGCCTTCTAGTTGCCAG
G0274326-1.seq(1>5500)	→		CGCTGATCAGCCTCGACTGTGCCTTCTAGTTGCCAG
E06-E61664-G0274326-1_PEGFP-N-5.ab1(1>792)	→		CGCTGATCAGCCTCGACTGTGCCTTCTAGTTGCCAG
		1120 1130 1140 1150	CCATCTGTTGTTTGCCCCTCCCCCGTGCCTTCCTTG
G0274326-1.seq(1>5500)	→		CCATCTGTTGTTTGCCCCTCCCCCGTGCCTTCCTTG
E06-E61664-G0274326-1_PEGFP-N-5.ab1(1>792)	→		CCATCTGTTGTTTGCCCCTCCCCCGTGCCTTCCTTG
		1160 1170 1180	ACCCTGGAAGGTGCCACTCCCCTGTCCTTTCCTAA
G0274326-1.seq(1>5500)	→		ACCCTGGAAGGTGCCACTCCCCTGTCCTTTCCTAA
E06-E61664-G0274326-1_PEGFP-N-5.ab1(1>792)	→		ACCCTGGAAGGTGCCACTCCCCTGTCCTTTCCTAA
		1190 1200 1210 1220	TAAAATGAGGAAATTGCATCGCATTGTCTGAGTAGG
G0274326-1.seq(1>5500)	→		TAAAATGAGGAAATTGCATCGCATTGTCTGAGTAGG
E06-E61664-G0274326-1_PEGFP-N-5.ab1(1>792)	→		TAAAATGAGGAAATTGCATCGCATTGTCTGAGTAGG

Project: Untitled.SQD Contig 1

		1230 1240 1250 1260	TGTCATTCTATTCTGGGGGTGGGGTGGGGCAGGAC
G0274326-1.seq (1>5500)	→		TGTCATTCTATTCTGGGGGTGGGGTGGGGCAGGAC
E06-E61664-G0274326-1_PEGFP-N-5.ab1 (1>792)	→		TGTCATTCTATTCTGGGGGTGGGGTGGGGCAGGAC
		1270 1280 1290	AGCAAGGGGAGGATTGGGAAGACAATAGCAGGCAT
G0274326-1.seq (1>5500)	→		AGCAAGGGGAGGATTGGGAAGACAATAGCAGGCAT
E06-E61664-G0274326-1_PEGFP-N-5.ab1 (1>792)	→		AGCAAGGGGAGGATTGGGAAGACAATAGCAGGCAT
		1300 1310 1320 1330	GCTGGGGATGCGGTGGGCTCTATGGCTTCTGAGGCG
G0274326-1.seq (1>5500)	→		GCTGGGGATGCGGTGGGCTCTATGGCTTCTGAGGCG
E06-E61664-G0274326-1_PEGFP-N-5.ab1 (1>792)	→		GCTGGGGATGCGGTGGGCTCTATGGCTTCTGAGGCG
		1340 1350 1360	GAAAGAACCAGCTGGGGCTCTAGGGGTATCCCCAC
G0274326-1.seq (1>5500)	→		GAAAGAACCAGCTGGGGCTCTAGGGGTATCCCCAC
E06-E61664-G0274326-1_PEGFP-N-5.ab1 (1>792)	→		GAAAGAACCAGCTGGGGCTCTAGGGGTATCCCCAC
		1370 1380 1390 1400	GCGCCCTGTAGCGGCGCATTAAAGCGGCGGGTGTG
G0274326-1.seq (1>5500)	→		GCGCCCTGTAGCGGCGCATTAAAGCGGCGGGTGTG
E06-E61664-G0274326-1_PEGFP-N-5.ab1 (1>792)	→		GCGCCCTGTAGCGGCGCATTAAAGCGGCGGGTGTG
		1410 1420 1430 1440	GTGGTTACGCGCAGCGTGACCGCTACACTTGCCAGC
G0274326-1.seq (1>5500)	→		GTGGTTACGCGCAGCGTGACCGCTACACTTGCCAGC
E06-E61664-G0274326-1_PEGFP-N-5.ab1 (1>792)	→		GTGGTTACGCGCAGCGTGACCGCTACACTTGCCAGC
		1450 1460 1470	GCCCTAGCGCCCGCTCCTTTCGCTTTCCTCCCTTCC
G0274326-1.seq (1>5500)	→		GCCCTAGCGCCCGCTCCTTTCGCTTTCCTCCCTTCC
E06-E61664-G0274326-1_PEGFP-N-5.ab1 (1>792)	→		GCCCTAGCGCCCGCTCCTTTCGCTTTCCTCCCTTCC
		1480 1490 1500 1510	TTTCTCGCCACGTTTCGCCGGCTTTCCTCCCGTCAAGCT
G0274326-1.seq (1>5500)	→		TTTCTCGCCACGTTTCGCCGGCTTTCCTCCCGTCAAGCT
E06-E61664-G0274326-1_PEGFP-N-5.ab1 (1>792)	→		TTTCTCGCCACGTTTCGCCGGCTTTCCTCCCGTCAAGCT
		1520 1530 1540	CTAAATCGGGGCTCCCTTTAGGGTTCCGATTTAGT
G0274326-1.seq (1>5500)	→		CTAAATCGGGGCTCCCTTTAGGGTTCCGATTTAGT
E06-E61664-G0274326-1_PEGFP-N-5.ab1 (1>792)	→		CTAAATCGGGGCTCCCTTTAGGGTTCCGATTTAGT
		1550 1560 1570 1580	GCTTTACGGCACCTCGACCCCAAAAACTTGATTAG
G0274326-1.seq (1>5500)	→		GCTTTACGGCACCTCGACCCCAAAAACTTGATTAG
E06-E61664-G0274326-1_PEGFP-N-5.ab1 (1>792)	→		GCTTTACGGCACCTCGACCCCAAAAACTTGATTAG

Project: Untitled.SQD Contig 1



Project: Untitled.SQD Contig 1



Project: Untitled.SQD Contig 1

		2450	2460	2470	2480
		ACGTTGTCACTGAAGCGGGAAGGGACTGGCTGCTAT			
G0274326-1.seq (1>5500)	→	ACGTTGTCACTGAAGCGGGAAGGGACTGGCTGCTAT			
		2490	2500	2510	2520
		TGGGCGAAGTGCCGGGGCAGGATCTCCTGTCATCTC			
G0274326-1.seq (1>5500)	→	TGGGCGAAGTGCCGGGGCAGGATCTCCTGTCATCTC			
		2530	2540	2550	
		ACCTTGCTCCTGCCGAGAAAGTATCCATCATGGCTG			
G0274326-1.seq (1>5500)	→	ACCTTGCTCCTGCCGAGAAAGTATCCATCATGGCTG			
		2560	2570	2580	2590
		ATGCAATGCGGCGGCTGCATACGCTTGATCCGGCTA			
G0274326-1.seq (1>5500)	→	ATGCAATGCGGCGGCTGCATACGCTTGATCCGGCTA			
		2600	2610	2620	
		CCTGCCCATTCGACCACCAAGCGAAACATCGCATCG			
G0274326-1.seq (1>5500)	→	CCTGCCCATTCGACCACCAAGCGAAACATCGCATCG			
		2630	2640	2650	2660
		AGCGAGCACGTACTIONCGGATGGAAGCCGGTCTTGTCG			
G0274326-1.seq (1>5500)	→	AGCGAGCACGTACTIONCGGATGGAAGCCGGTCTTGTCG			
		2670	2680	2690	2700
		ATCAGGATGATCTGGACGAAGAGCATCAGGGGCTCG			
G0274326-1.seq (1>5500)	→	ATCAGGATGATCTGGACGAAGAGCATCAGGGGCTCG			
		2710	2720	2730	
		CGCCAGCCGAACTGTTCCGCCAGGCTCAAGGCGCGCA			
G0274326-1.seq (1>5500)	→	CGCCAGCCGAACTGTTCCGCCAGGCTCAAGGCGCGCA			
		2740	2750	2760	2770
		TGCCCGACGGCGAGGATCTCGTCGTGACCCATGGCG			
G0274326-1.seq (1>5500)	→	TGCCCGACGGCGAGGATCTCGTCGTGACCCATGGCG			
		2780	2790	2800	
		ATGCCTGCTTGCCGAATATCATGGTGGAAAATGGCC			
G0274326-1.seq (1>5500)	→	ATGCCTGCTTGCCGAATATCATGGTGGAAAATGGCC			
		2810	2820	2830	2840
		GCTTTTCTGGATTTCATCGACTGTGGCCGGCTGGGTG			
G0274326-1.seq (1>5500)	→	GCTTTTCTGGATTTCATCGACTGTGGCCGGCTGGGTG			
		2850	2860	2870	2880
		TGGCGGACCGCTATCAGGACATAGCGTTGGCTACCC			
G0274326-1.seq (1>5500)	→	TGGCGGACCGCTATCAGGACATAGCGTTGGCTACCC			

Project: Untitled.SQD Contig 1

		2890	2900	2910	
		GTGATATTGCTGAAGAGCTTGGCGGCGAATGGGCTG			
G0274326-1.seq (1>5500)	→	GTGATATTGCTGAAGAGCTTGGCGGCGAATGGGCTG			
		2920	2930	2940	2950
		ACCGCTTCCTCGTGCTTTACGGTATCGCCGCTCCCG			
G0274326-1.seq (1>5500)	→	ACCGCTTCCTCGTGCTTTACGGTATCGCCGCTCCCG			
		2960	2970	2980	
		ATTTCGAGCGCATCGCCTTCTATCGCCTTCTTGACG			
G0274326-1.seq (1>5500)	→	ATTTCGAGCGCATCGCCTTCTATCGCCTTCTTGACG			
		2990	3000	3010	3020
		AGTTCTTCTGAGCGGGACTCTGGGGTTCGAAATGAC			
G0274326-1.seq (1>5500)	→	AGTTCTTCTGAGCGGGACTCTGGGGTTCGAAATGAC			
		3030	3040	3050	3060
		CGACCAAGCGACGCCAACCTGCCATCACGAGATTT			
G0274326-1.seq (1>5500)	→	CGACCAAGCGACGCCAACCTGCCATCACGAGATTT			
		3070	3080	3090	
		CGATTCCACCGCCGCCTTCTATGAAAGGTTGGGCTT			
G0274326-1.seq (1>5500)	→	CGATTCCACCGCCGCCTTCTATGAAAGGTTGGGCTT			
		3100	3110	3120	3130
		CGGAATCGTTTTCCGGGACGCCGGCTGGATGATCCT			
G0274326-1.seq (1>5500)	→	CGGAATCGTTTTCCGGGACGCCGGCTGGATGATCCT			
		3140	3150	3160	
		CCAGCGCGGGATCTCATGCTGGAGTTCTTCGCCCA			
G0274326-1.seq (1>5500)	→	CCAGCGCGGGATCTCATGCTGGAGTTCTTCGCCCA			
		3170	3180	3190	3200
		CCCCAACTTGTTTATTGCAGCTTATAATGGTTACAA			
G0274326-1.seq (1>5500)	→	CCCCAACTTGTTTATTGCAGCTTATAATGGTTACAA			
		3210	3220	3230	3240
		ATAAAGCAATAGCATCACAAATTTACAAATAAAGC			
G0274326-1.seq (1>5500)	→	ATAAAGCAATAGCATCACAAATTTACAAATAAAGC			
		3250	3260	3270	
		ATTTTTTTCACCTGCATTCTAGTTGTGGTTTGTCCAA			
G0274326-1.seq (1>5500)	→	ATTTTTTTCACCTGCATTCTAGTTGTGGTTTGTCCAA			
		3280	3290	3300	3310
		ACTCATCAATGTATCTTATCATGTCTGTATACCGTC			
G0274326-1.seq (1>5500)	→	ACTCATCAATGTATCTTATCATGTCTGTATACCGTC			

Project: Untitled.SQD Contig 1

		3320 3330 3340	GACCTCTAGCTAGAGCTTGGCGTAATCATGGTCATA
G0274326-1.seq (1>5500)	→		GACCTCTAGCTAGAGCTTGGCGTAATCATGGTCATA
		3350 3360 3370 3380	GCTGTTTCCTGTGTGAAATTGTTATCCGCTCACAAT
G0274326-1.seq (1>5500)	→		GCTGTTTCCTGTGTGAAATTGTTATCCGCTCACAAT
		3390 3400 3410 3420	TCCACACAACATACGAGCCGGAAGCATAAAGTGTAA
G0274326-1.seq (1>5500)	→		TCCACACAACATACGAGCCGGAAGCATAAAGTGTAA
		3430 3440 3450	AGCCTGGGGTGCCTAATGAGTGAGCTAACTCACATT
G0274326-1.seq (1>5500)	→		AGCCTGGGGTGCCTAATGAGTGAGCTAACTCACATT
		3460 3470 3480 3490	AATTGCGTTGCGCTCACTGCCCGCTTTCAGTCGGG
G0274326-1.seq (1>5500)	→		AATTGCGTTGCGCTCACTGCCCGCTTTCAGTCGGG
		3500 3510 3520	AAACCTGTCGTGCCAGCTGCATTAATGAATCGGCCA
G0274326-1.seq (1>5500)	→		AAACCTGTCGTGCCAGCTGCATTAATGAATCGGCCA
		3530 3540 3550 3560	ACGCGCGGGGAGAGGCGGTTTGCATTTGGGCGCTC
G0274326-1.seq (1>5500)	→		ACGCGCGGGGAGAGGCGGTTTGCATTTGGGCGCTC
		3570 3580 3590 3600	TTCCGCTTCCTCGCTCACTGACTCGCTGCGCTCGGT
G0274326-1.seq (1>5500)	→		TTCCGCTTCCTCGCTCACTGACTCGCTGCGCTCGGT
		3610 3620 3630	CGTTCGGCTGCGGCGAGCGGTATCAGCTCACTCAA
G0274326-1.seq (1>5500)	→		CGTTCGGCTGCGGCGAGCGGTATCAGCTCACTCAA
		3640 3650 3660 3670	GGCGGTAATACGGTTATCCACAGAATCAGGGGATAA
G0274326-1.seq (1>5500)	→		GGCGGTAATACGGTTATCCACAGAATCAGGGGATAA
		3680 3690 3700	CGCAGGAAAGAACATGTGAGCAAAGGCCAGCAAAA
G0274326-1.seq (1>5500)	→		CGCAGGAAAGAACATGTGAGCAAAGGCCAGCAAAA
		3710 3720 3730 3740	GGCCAGGAACCGTAAAAAGGCCGCGTTGCTGGCGTT
G0274326-1.seq (1>5500)	→		GGCCAGGAACCGTAAAAAGGCCGCGTTGCTGGCGTT

Project: Untitled.SQD Contig 1

		3750 3760 3770 3780	TTTCCATAGGCTCCGCCCCCTGACGAGCATCACAA
G0274326-1.seq (1>5500)	→		TTTCCATAGGCTCCGCCCCCTGACGAGCATCACAA
		3790 3800 3810	AAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGAC
G0274326-1.seq (1>5500)	→		AAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGAC
		3820 3830 3840 3850	AGGACTATAAAGATAACCAGGCGTTTCCCCCTGGAAG
G0274326-1.seq (1>5500)	→		AGGACTATAAAGATAACCAGGCGTTTCCCCCTGGAAG
		3860 3870 3880	CTCCCTCGTGCGCTCTCCTGTTCCGACCCTGCCGCT
G0274326-1.seq (1>5500)	→		CTCCCTCGTGCGCTCTCCTGTTCCGACCCTGCCGCT
		3890 3900 3910 3920	TACCGGATACCTGTCCGCCTTCTCCCTTCGGGAAG
G0274326-1.seq (1>5500)	→		TACCGGATACCTGTCCGCCTTCTCCCTTCGGGAAG
		3930 3940 3950 3960	CGTGCGCTTCTCATAGCTCACGCTGTAGGTATCT
G0274326-1.seq (1>5500)	→		CGTGCGCTTCTCATAGCTCACGCTGTAGGTATCT
		3970 3980 3990	CAGTTCGGTGTAGGTCGTTTCGCTCCAAGCTGGGCTG
G0274326-1.seq (1>5500)	→		CAGTTCGGTGTAGGTCGTTTCGCTCCAAGCTGGGCTG
		4000 4010 4020 4030	TGTGCACGAACCCCCGTTTCAGCCCAGCCGCTGCGC
G0274326-1.seq (1>5500)	→		TGTGCACGAACCCCCGTTTCAGCCCAGCCGCTGCGC
		4040 4050 4060	CTTATCCGGTAACTATCGTCTTGAGTCCAACCCGGT
G0274326-1.seq (1>5500)	→		CTTATCCGGTAACTATCGTCTTGAGTCCAACCCGGT
		4070 4080 4090 4100	AAGACACGACTTATCGCCACTGGCAGCAGCCACTGG
G0274326-1.seq (1>5500)	→		AAGACACGACTTATCGCCACTGGCAGCAGCCACTGG
		4110 4120 4130 4140	TAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGC
G0274326-1.seq (1>5500)	→		TAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGC
		4150 4160 4170	TACAGAGTTCTTGAAGTGGTGGCCTAACTACGGCTA
G0274326-1.seq (1>5500)	→		TACAGAGTTCTTGAAGTGGTGGCCTAACTACGGCTA

Project: Untitled.SQD Contig 1

		4180	4190	4200	4210
		CACTAGAAGAACAGTATTTGGTATCTGCGCTCTGCT			
G0274326-1.seq (1>5500)	→	CACTAGAAGAACAGTATTTGGTATCTGCGCTCTGCT			
		4220	4230	4240	
		GAAGCCAGTTACCTTCGGAAAAAGAGTTGGTAGCTC			
G0274326-1.seq (1>5500)	→	GAAGCCAGTTACCTTCGGAAAAAGAGTTGGTAGCTC			
		4250	4260	4270	4280
		TTGATCCGGCAAACAACCACCGCTGGTAGCGGTGG			
G0274326-1.seq (1>5500)	→	TTGATCCGGCAAACAACCACCGCTGGTAGCGGTGG			
		4290	4300	4310	4320
		TTTTTTTGT TTGCAAGCAGCAGATTACGCGCAGAAA			
G0274326-1.seq (1>5500)	→	TTTTTTTGT TTGCAAGCAGCAGATTACGCGCAGAAA			
		4330	4340	4350	
		AAAAGGATCTCAAGAAGATCCTTTGATCTTTTCTAC			
G0274326-1.seq (1>5500)	→	AAAAGGATCTCAAGAAGATCCTTTGATCTTTTCTAC			
		4360	4370	4380	4390
		GGGTCTGACGCTCAGTGGAACGAAAAC TACGTTA			
G0274326-1.seq (1>5500)	→	GGGTCTGACGCTCAGTGGAACGAAAAC TACGTTA			
		4400	4410	4420	
		AGGGATTTTGGTCATGAGATTATCAAAAAGGATCTT			
G0274326-1.seq (1>5500)	→	AGGGATTTTGGTCATGAGATTATCAAAAAGGATCTT			
		4430	4440	4450	4460
		CACCTAGATCCTTTTAAATTAAAAATGAAGTTTAA			
G0274326-1.seq (1>5500)	→	CACCTAGATCCTTTTAAATTAAAAATGAAGTTTAA			
		4470	4480	4490	4500
		ATCAATCTAAAGTATATATGAGTAAACTTGGTCTGA			
G0274326-1.seq (1>5500)	→	ATCAATCTAAAGTATATATGAGTAAACTTGGTCTGA			
		4510	4520	4530	
		CAGTTACCAATGCTTAATCAGTGAGGCACCTATCTC			
G0274326-1.seq (1>5500)	→	CAGTTACCAATGCTTAATCAGTGAGGCACCTATCTC			
		4540	4550	4560	4570
		AGCGATCTGTCTATTTTCGTTTCATCCATAGTTGCCTG			
G0274326-1.seq (1>5500)	→	AGCGATCTGTCTATTTTCGTTTCATCCATAGTTGCCTG			
		4580	4590	4600	
		ACTCCCCGTCGTGTAGATAACTACGATACGGGAGGG			
G0274326-1.seq (1>5500)	→	ACTCCCCGTCGTGTAGATAACTACGATACGGGAGGG			

Project: Untitled.SQD Contig 1

	4610	4620	4630	4640	CTTACCATCTGGCCCCAGTGCTGCAATGATACCGCG
G0274326-1.seq (1>5500)	→	CTTACCATCTGGCCCCAGTGCTGCAATGATACCGCG			
	4650	4660	4670	4680	AGACCCACGCTCACCGGCTCCAGATTTATCAGCAAT
G0274326-1.seq (1>5500)	→	AGACCCACGCTCACCGGCTCCAGATTTATCAGCAAT			
	4690	4700	4710		AAACCAGCCAGCCGGAAGGGCCGAGCGCAGAAGTGG
G0274326-1.seq (1>5500)	→	AAACCAGCCAGCCGGAAGGGCCGAGCGCAGAAGTGG			
	4720	4730	4740	4750	TCCTGCAACTTTATCCGCCTCCATCCAGTCTATTAA
G0274326-1.seq (1>5500)	→	TCCTGCAACTTTATCCGCCTCCATCCAGTCTATTAA			
	4760	4770	4780		TTGTTGCCGGAAGCTAGAGTAAGTAGTTCGCCAGT
G0274326-1.seq (1>5500)	→	TTGTTGCCGGAAGCTAGAGTAAGTAGTTCGCCAGT			
	4790	4800	4810	4820	TAATAGTTTGCGCAACGTTGTTGCCATTGCTACAGG
G0274326-1.seq (1>5500)	→	TAATAGTTTGCGCAACGTTGTTGCCATTGCTACAGG			
	4830	4840	4850	4860	CATCGTGGTGTCACGCTCGTCGTTTGGTATGGCTTC
G0274326-1.seq (1>5500)	→	CATCGTGGTGTCACGCTCGTCGTTTGGTATGGCTTC			
	4870	4880	4890		ATTCAGCTCCGGTTCCCAACGATCAAGGCGAGTTAC
G0274326-1.seq (1>5500)	→	ATTCAGCTCCGGTTCCCAACGATCAAGGCGAGTTAC			
	4900	4910	4920	4930	ATGATCCCCATGTTGTGCAAAAAGCGGTTAGCTC
G0274326-1.seq (1>5500)	→	ATGATCCCCATGTTGTGCAAAAAGCGGTTAGCTC			
	4940	4950	4960		CTTCGGTCCCTCCGATCGTTGTCAGAAGTAAGTTGGC
G0274326-1.seq (1>5500)	→	CTTCGGTCCCTCCGATCGTTGTCAGAAGTAAGTTGGC			
	4970	4980	4990	5000	CGCAGTGTTATCACTCATGGTTATGGCAGCACTGCA
G0274326-1.seq (1>5500)	→	CGCAGTGTTATCACTCATGGTTATGGCAGCACTGCA			
	5010	5020	5030	5040	TAATTCTCTTACTGTCATGCCATCCGTAAGATGCTT
G0274326-1.seq (1>5500)	→	TAATTCTCTTACTGTCATGCCATCCGTAAGATGCTT			

Project: Untitled.SQD Contig 1

		5050	5060	5070	TTCTGTGACTGGTGAGTACTCAACCAAGTCATTCTG	
G0274326-1.seq (1>5500)	→	TTCTGTGACTGGTGAGTACTCAACCAAGTCATTCTG				
		5080	5090	5100	5110	AGAATAGTGTATGCGGCGACCGAGTTGCTCTTGCCC
G0274326-1.seq (1>5500)	→	AGAATAGTGTATGCGGCGACCGAGTTGCTCTTGCCC				
		5120	5130	5140	GGCGTCAATACGGGATAATACCGCGCCACATAGCAG	
G0274326-1.seq (1>5500)	→	GGCGTCAATACGGGATAATACCGCGCCACATAGCAG				
		5150	5160	5170	5180	AACTTTAAAAGTGCTCATCATTGGAAAACGTTCTTC
G0274326-1.seq (1>5500)	→	AACTTTAAAAGTGCTCATCATTGGAAAACGTTCTTC				
		5190	5200	5210	5220	GGGGCGAAAAC TCTCAAGGATCTTACCGCTGTTGAG
G0274326-1.seq (1>5500)	→	GGGGCGAAAAC TCTCAAGGATCTTACCGCTGTTGAG				
		5230	5240	5250	ATCCAGTTCGATGTAACCCACTCGTGCACCCAAC TG	
G0274326-1.seq (1>5500)	→	ATCCAGTTCGATGTAACCCACTCGTGCACCCAAC TG				
		5260	5270	5280	5290	ATCTTCAGCATCTTTTACTTTTACCAGCGTTTCTGG
G0274326-1.seq (1>5500)	→	ATCTTCAGCATCTTTTACTTTTACCAGCGTTTCTGG				
		5300	5310	5320	GTGAGCAAAAACAGGAAGGCAAAATGCCGCAAAAAA	
G0274326-1.seq (1>5500)	→	GTGAGCAAAAACAGGAAGGCAAAATGCCGCAAAAAA				
		5330	5340	5350	5360	GGGAATAAGGGCGACACGGAAATGTTGAATACTCAT
G0274326-1.seq (1>5500)	→	GGGAATAAGGGCGACACGGAAATGTTGAATACTCAT				
		5370	5380	5390	5400	ACTCTTCCTTTTTCAATATTATTGAAGCATT TATCA
G0274326-1.seq (1>5500)	→	ACTCTTCCTTTTTCAATATTATTGAAGCATT TATCA				
		5410	5420	5430	GGGTTATTGTCTCATGAGCGGATACATATTTGAATG	
G0274326-1.seq (1>5500)	→	GGGTTATTGTCTCATGAGCGGATACATATTTGAATG				
		5440	5450	5460	5470	TATTTAGAAAAATAAACAAATAGGGGTTCCGCGCAC
G0274326-1.seq (1>5500)	→	TATTTAGAAAAATAAACAAATAGGGGTTCCGCGCAC				

5480 5490 5500

ATTTC~~CCCG~~AAAAGTGCCACCTGACGTC

G0274326-1.seq (1>5500)



ATTTC~~CCCG~~AAAAGTGCCACCTGACGTC