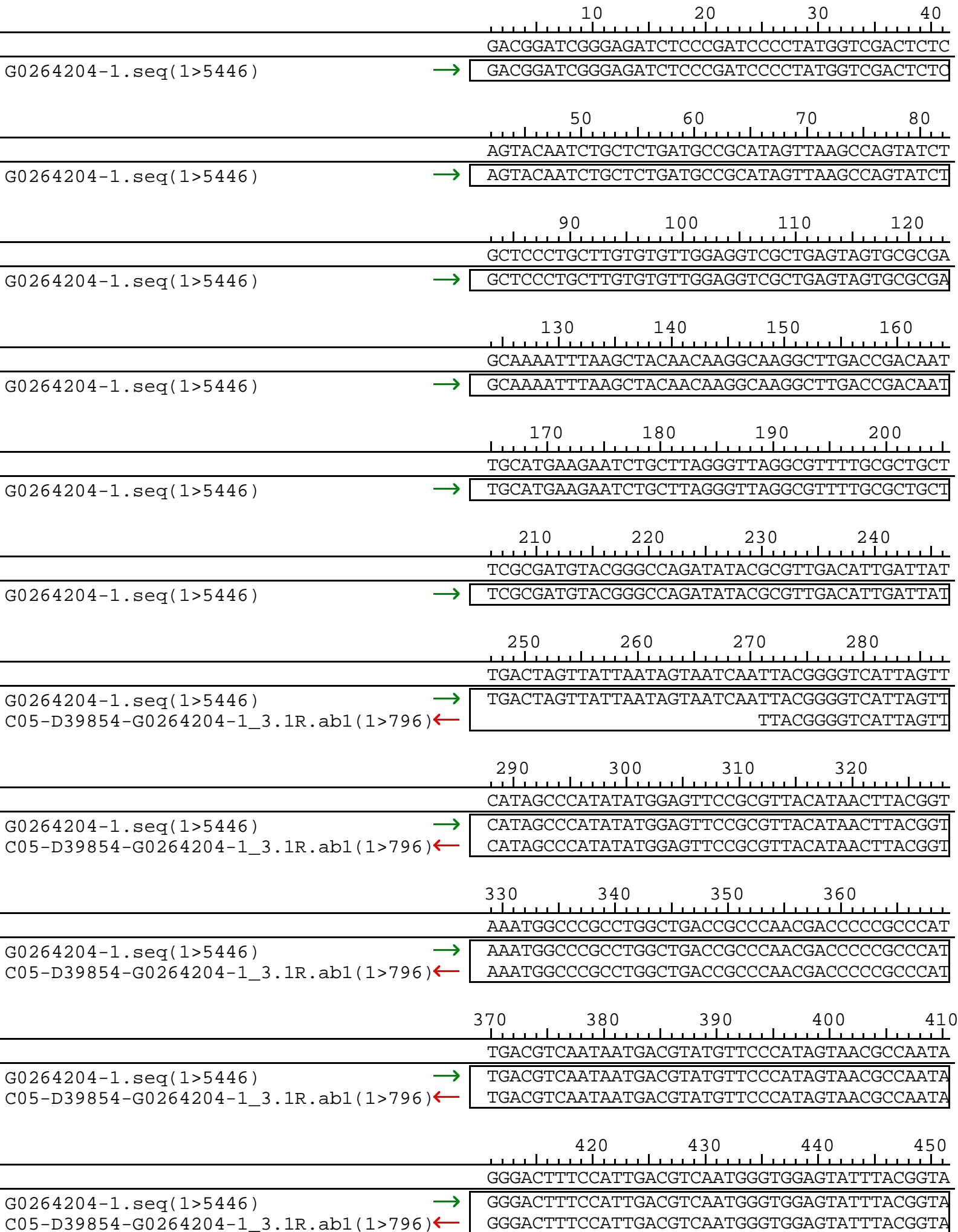


Project: G0264204-1.SQD Contig 1



Project: G0264204-1.SQD Contig 1

		460	470	480	490	
		AACTGCCCACTTGGCAGTACATCAAGTGTATCATATGCCAA				
G0264204-1.seq(1>5446)	→	AACTGCCCACTTGGCAGTACATCAAGTGTATCATATGCCAA				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	AACTGCCCACTTGGCAGTACATCAAGTGTATCATATGCCAA				
		500	510	520	530	
		GTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCGCC				
G0264204-1.seq(1>5446)	→	GTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCGCC				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	GTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCGCC				
		540	550	560	570	
		TGGCATTATGCCCAGTACATGACCTTATGGGACTTTCCTAC				
G0264204-1.seq(1>5446)	→	TGGCATTATGCCCAGTACATGACCTTATGGGACTTTCCTAC				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	TGGCATTATGCCCAGTACATGACCTTATGGGACTTTCCTAC				
		580	590	600	610	
		TTGGCAGTACATCTACGTATTAGTCATCGCTATTACCATGG				
G0264204-1.seq(1>5446)	→	TTGGCAGTACATCTACGTATTAGTCATCGCTATTACCATGG				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	TTGGCAGTACATCTACGTATTAGTCATCGCTATTACCATGG				
		620	630	640	650	
		TGATGCGGTTTTGGCAGTACATCAATGGGCGTGGATAGCGG				
G0264204-1.seq(1>5446)	→	TGATGCGGTTTTGGCAGTACATCAATGGGCGTGGATAGCGG				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	TGATGCGGTTTTGGCAGTACATCAATGGGCGTGGATAGCGG				
		660	670	680	690	
		TTTGACTCACGGGGATTTCCAAGTCTCCACCCCATTTGACGT				
G0264204-1.seq(1>5446)	→	TTTGACTCACGGGGATTTCCAAGTCTCCACCCCATTTGACGT				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	TTTGACTCACGGGGATTTCCAAGTCTCCACCCCATTTGACGT				
		700	710	720	730	
		CAATGGGAGTTTGTTTTGGCACCAAAATCAACGGGACTTTC				
G0264204-1.seq(1>5446)	→	CAATGGGAGTTTGTTTTGGCACCAAAATCAACGGGACTTTC				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	CAATGGGAGTTTGTTTTGGCACCAAAATCAACGGGACTTTC				
		740	750	760	770	
		CAAAATGTCGTAACAACCTCCGCCCCATTGACGCAAATGGGC				
G0264204-1.seq(1>5446)	→	CAAAATGTCGTAACAACCTCCGCCCCATTGACGCAAATGGGC				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	CAAAATGTCGTAACAACCTCCGCCCCATTGACGCAAATGGGC				
		780	790	800	810	820
		GGTAGGCGTGTACGGTGGGAGGTCTATATAAGCAGAGCTCT				
G0264204-1.seq(1>5446)	→	GGTAGGCGTGTACGGTGGGAGGTCTATATAAGCAGAGCTCT				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	GGTAGGCGTGTACGGTGGGAGGTCTATATAAGCAGAGCTCT				
		830	840	850	860	
		CTGGCTAACTAGAGAACCCACTGCTTACTGGCTTATCGAAA				
G0264204-1.seq(1>5446)	→	CTGGCTAACTAGAGAACCCACTGCTTACTGGCTTATCGAAA				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	CTGGCTAACTAGAGAACCCACTGCTTACTGGCTTATCGAAA				

Project: G0264204-1.SQD Contig 1

		870	880	890	900	
		TTAATACGACTCACTATAGGGAGACCCAAGCTTGGTACCGA				
G0264204-1.seq(1>5446)	→	TTAATACGACTCACTATAGGGAGACCCAAGCTTGGTACCGA				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	TTAATACGACTCACTATAGGGAGACCCAAGCTTGGTACCGA				
		910	920	930	940	
		GCTCGGATCCACTAGTAACGGCCGCCAGTGTGCTGGAATTC				
G0264204-1.seq(1>5446)	→	GCTCGGATCCACTAGTAACGGCCGCCAGTGTGCTGGAATTC				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	GCTCGGATCCACTAGTAACGGCCGCCAGTGTGCTGGAATTC				
		950	960	970	980	
		TGCAGATATCCATCACACTGGCGGCCGCTCGAGCATGCATC				
G0264204-1.seq(1>5446)	→	TGCAGATATCCATCACACTGGCGGCCGCTCGAGCATGCATC				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	TGCAGATATCCATCACACTGGCGGCCGCTCGAGCATGCATC				
		990	1000	1010	1020	
		TAGAGGGCCCTATTCTATAGTGTACCTAAATGCTAGAGCT				
G0264204-1.seq(1>5446)	→	TAGAGGGCCCTATTCTATAGTGTACCTAAATGCTAGAGCT				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	TAGAGGGCCCTATTCTATAGTGTACCTAAATGCTAGAGCT				
		1030	1040	1050	1060	
		CGCTGATCAGCCTCGACTGTGCCTTCTAGTTGCCAGCCATC				
G0264204-1.seq(1>5446)	→	CGCTGATCAGCCTCGACTGTGCCTTCTAGTTGCCAGCCATC				
C05-D39854-G0264204-1_3.1R.ab1(1>796)	←	CGCTGATCAGCCTCGACTGTGCCTTCTAGTTGCCAGCCATC				
		1070	1080	1090	1100	
		TGTTGTTTGCCCCTCCCCCGTGCCTTCCTTGACCCTGGAAG				
G0264204-1.seq(1>5446)	→	TGTTGTTTGCCCCTCCCCCGTGCCTTCCTTGACCCTGGAAG				
		1110	1120	1130	1140	
		GTGCCACTCCCACTGTCCTTTCCTAATAAAATGAGGAAATT				
G0264204-1.seq(1>5446)	→	GTGCCACTCCCACTGTCCTTTCCTAATAAAATGAGGAAATT				
		1150	1160	1170	1180	
		GCATCGCATTGTCTGAGTAGGTGTCATTCTATTCTGGGGGG				
G0264204-1.seq(1>5446)	→	GCATCGCATTGTCTGAGTAGGTGTCATTCTATTCTGGGGGG				
		1190	1200	1210	1220	1230
		TGGGGTGGGGCAGGACAGCAAGGGGGAGGATTGGGAAGACA				
G0264204-1.seq(1>5446)	→	TGGGGTGGGGCAGGACAGCAAGGGGGAGGATTGGGAAGACA				
		1240	1250	1260	1270	
		ATAGCAGGCATGCTGGGGATGCGGTGGGCTCTATGGCTTCT				
G0264204-1.seq(1>5446)	→	ATAGCAGGCATGCTGGGGATGCGGTGGGCTCTATGGCTTCT				
		1280	1290	1300	1310	
		GAGGCGGAAAGAACCAGCTGGGGCTCTAGGGGGTATCCCCA				
G0264204-1.seq(1>5446)	→	GAGGCGGAAAGAACCAGCTGGGGCTCTAGGGGGTATCCCCA				

Project: G0264204-1.SQD Contig 1

		1320 1330 1340 1350	CGCGCCCTGTAGCGGCGCATTAAAGCGCGGCGGGTGTGGTGG
G0264204-1.seq(1>5446)	→		CGCGCCCTGTAGCGGCGCATTAAAGCGCGGCGGGTGTGGTGG
		1360 1370 1380 1390	TTACGCGCAGCGTGACCGCTACACTTGCCAGCGCCCTAGCG
G0264204-1.seq(1>5446)	→		TTACGCGCAGCGTGACCGCTACACTTGCCAGCGCCCTAGCG
		1400 1410 1420 1430	CCCGCTCCTTTCGCTTTCCTTCCCTTCTTTCTCGCCACGTT
G0264204-1.seq(1>5446)	→		CCCGCTCCTTTCGCTTTCCTTCCCTTCTTTCTCGCCACGTT
		1440 1450 1460 1470	CGCCGGCTTTCCTCCGTCAAGCTCTAAATCGGGGGCTCCCTT
G0264204-1.seq(1>5446)	→		CGCCGGCTTTCCTCCGTCAAGCTCTAAATCGGGGGCTCCCTT
		1480 1490 1500 1510	TAGGGTTCGATTTAGTGCTTTACGGCACCTCGACCCCAA
G0264204-1.seq(1>5446)	→		TAGGGTTCGATTTAGTGCTTTACGGCACCTCGACCCCAA
		1520 1530 1540 1550	AAACTTGATTAGGGTGATGGTTCACGTAGTGGGCCATCGCC
G0264204-1.seq(1>5446)	→		AAACTTGATTAGGGTGATGGTTCACGTAGTGGGCCATCGCC
		1560 1570 1580 1590	CTGATAGACGGTTTTTCGCCCTTTGACGTTGGAGTCCACGT
G0264204-1.seq(1>5446)	→		CTGATAGACGGTTTTTCGCCCTTTGACGTTGGAGTCCACGT
		1600 1610 1620 1630 1640	TCTTTAATAGTGGACTCTTGTTCCAAACTGGAACAACACTC
G0264204-1.seq(1>5446)	→		TCTTTAATAGTGGACTCTTGTTCCAAACTGGAACAACACTC
		1650 1660 1670 1680	AACCCTATCTCGGTCTATTCTTTTGATTTATAAGGGATTTT
G0264204-1.seq(1>5446)	→		AACCCTATCTCGGTCTATTCTTTTGATTTATAAGGGATTTT
		1690 1700 1710 1720	GGGGATTTTCGGCCTATTGGTTAAAAAATGAGCTGATTTAAC
G0264204-1.seq(1>5446)	→		GGGGATTTTCGGCCTATTGGTTAAAAAATGAGCTGATTTAAC
		1730 1740 1750 1760	AAAAATTTAACGCGAATTAATTCTGTGGAATGTGTGTCAGT
G0264204-1.seq(1>5446)	→		AAAAATTTAACGCGAATTAATTCTGTGGAATGTGTGTCAGT
		1770 1780 1790 1800	TAGGGTGTGGAAAGTCCCCAGGCTCCCCAGGCAGGCAGAAG
G0264204-1.seq(1>5446)	→		TAGGGTGTGGAAAGTCCCCAGGCTCCCCAGGCAGGCAGAAG

Project: G0264204-1.SQD Contig 1

		1810 1820 1830 1840	TATGCAAAGCATGCATCTCAATTAGTCAGCAACCAGGTGTG
G0264204-1.seq(1>5446)	→		TATGCAAAGCATGCATCTCAATTAGTCAGCAACCAGGTGTG
		1850 1860 1870 1880	GAAAGTCCCAGGCTCCCAGCAGGCAGAAGTATGCAAAGC
G0264204-1.seq(1>5446)	→		GAAAGTCCCAGGCTCCCAGCAGGCAGAAGTATGCAAAGC
		1890 1900 1910 1920	ATGCATCTCAATTAGTCAGCAACCATAGTCCC GCCCTAAC
G0264204-1.seq(1>5446)	→		ATGCATCTCAATTAGTCAGCAACCATAGTCCC GCCCTAAC
		1930 1940 1950 1960	TCCGCCCATCCCGCCCCTAACTCCGCCAGTTCCGCCATT
G0264204-1.seq(1>5446)	→		TCCGCCCATCCCGCCCCTAACTCCGCCAGTTCCGCCATT
		1970 1980 1990 2000	CTCCGCCCCATGGCTGACTAATTTTTTTTATTTATGCAGAG
G0264204-1.seq(1>5446)	→		CTCCGCCCCATGGCTGACTAATTTTTTTTATTTATGCAGAG
		2010 2020 2030 2040 2050	GCCGAGGCCGCTCTGCCTCTGAGCTATTCCAGAAGTAGTG
G0264204-1.seq(1>5446)	→		GCCGAGGCCGCTCTGCCTCTGAGCTATTCCAGAAGTAGTG
		2060 2070 2080 2090	AGGAGGCTTTTTTGGAGGCCTAGGCTTTTGCAAAAAGCTCC
G0264204-1.seq(1>5446)	→		AGGAGGCTTTTTTGGAGGCCTAGGCTTTTGCAAAAAGCTCC
		2100 2110 2120 2130	CGGGAGCTTGTATATCCATTTTCGGATCTGATCAAGAGACA
G0264204-1.seq(1>5446)	→		CGGGAGCTTGTATATCCATTTTCGGATCTGATCAAGAGACA
		2140 2150 2160 2170	GGATGAGGATCGTTTCGCATGATTGAACAAGATGGATTGCA
G0264204-1.seq(1>5446)	→		GGATGAGGATCGTTTCGCATGATTGAACAAGATGGATTGCA
		2180 2190 2200 2210	CGCAGGTTCTCCGGCCGCTTGGGTGGAGAGGCTATTCCGGCT
G0264204-1.seq(1>5446)	→		CGCAGGTTCTCCGGCCGCTTGGGTGGAGAGGCTATTCCGGCT
		2220 2230 2240 2250	ATGACTGGGCACAACAGACAATCGGCTGCTCTGATGCCGCC
G0264204-1.seq(1>5446)	→		ATGACTGGGCACAACAGACAATCGGCTGCTCTGATGCCGCC
		2260 2270 2280 2290	GTGTTCCGGCTGTCAGCGCAGGGGCGCCCGGTTCTTTTTGT
G0264204-1.seq(1>5446)	→		GTGTTCCGGCTGTCAGCGCAGGGGCGCCCGGTTCTTTTTGT

Project: G0264204-1.SQD Contig 1

	2300	2310	2320	2330	CAAGACCGACCTGTCCGGTGCCCTGAATGAACTGCAGGACG	
G0264204-1.seq(1>5446)	→	CAAGACCGACCTGTCCGGTGCCCTGAATGAACTGCAGGACG				
	2340	2350	2360	2370	AGGCAGCGCGGCTATCGTGGCTGGCCACGACGGGCGTTCCT	
G0264204-1.seq(1>5446)	→	AGGCAGCGCGGCTATCGTGGCTGGCCACGACGGGCGTTCCT				
	2380	2390	2400	2410	TGCGCAGCTGTGCTCGACGTTGTCACTGAAGCGGGAAGGGA	
G0264204-1.seq(1>5446)	→	TGCGCAGCTGTGCTCGACGTTGTCACTGAAGCGGGAAGGGA				
	2420	2430	2440	2450	2460	CTGGCTGCTATTGGGCGAAGTGCCGGGGCAGGATCTCCTGT
G0264204-1.seq(1>5446)	→	CTGGCTGCTATTGGGCGAAGTGCCGGGGCAGGATCTCCTGT				
	2470	2480	2490	2500	CATCTCACCTTGCTCCTGCCGAGAAAGTATCCATCATGGCT	
G0264204-1.seq(1>5446)	→	CATCTCACCTTGCTCCTGCCGAGAAAGTATCCATCATGGCT				
	2510	2520	2530	2540	GATGCAATGCGGCGGCTGCATACGCTTGATCCGCTACCTG	
G0264204-1.seq(1>5446)	→	GATGCAATGCGGCGGCTGCATACGCTTGATCCGCTACCTG				
	2550	2560	2570	2580	CCCATTGACCACCAAGCGAAACATCGCATCGAGCGAGCAC	
G0264204-1.seq(1>5446)	→	CCCATTGACCACCAAGCGAAACATCGCATCGAGCGAGCAC				
	2590	2600	2610	2620	GTACTCGGATGGAAGCCGGTCTTGTCGATCAGGATGATCTG	
G0264204-1.seq(1>5446)	→	GTACTCGGATGGAAGCCGGTCTTGTCGATCAGGATGATCTG				
	2630	2640	2650	2660	GACGAAGAGCATCAGGGGCTCGCGCCAGCCGAACTGTTTCGC	
G0264204-1.seq(1>5446)	→	GACGAAGAGCATCAGGGGCTCGCGCCAGCCGAACTGTTTCGC				
	2670	2680	2690	2700	CAGGCTCAAGGCGCGCATGCCCGACGGCGAGGATCTCGTCCG	
G0264204-1.seq(1>5446)	→	CAGGCTCAAGGCGCGCATGCCCGACGGCGAGGATCTCGTCCG				
	2710	2720	2730	2740	TGACCCATGGCGATGCCTGCTTGCCGAATATCATGGTGGAA	
G0264204-1.seq(1>5446)	→	TGACCCATGGCGATGCCTGCTTGCCGAATATCATGGTGGAA				
	2750	2760	2770	2780	AATGGCCGCTTTTCTGGATTCATCGACTGTGGCCGGCTGGG	
G0264204-1.seq(1>5446)	→	AATGGCCGCTTTTCTGGATTCATCGACTGTGGCCGGCTGGG				

Project: G0264204-1.SQD Contig 1

	2790	2800	2810	2820	TGTGGCGGACCGCTATCAGGACATAGCGTTGGCTACCCGTG	
G0264204-1.seq(1>5446)	→	TGTGGCGGACCGCTATCAGGACATAGCGTTGGCTACCCGTG				
	2830	2840	2850	2860	2870	ATATTGCTGAAGAGCTTGGCGGCGAATGGGCTGACCGCTTC
G0264204-1.seq(1>5446)	→	ATATTGCTGAAGAGCTTGGCGGCGAATGGGCTGACCGCTTC				
	2880	2890	2900	2910	CTCGTGCTTTACGGTATCGCCGCTCCCGATTTCGAGCGCAT	
G0264204-1.seq(1>5446)	→	CTCGTGCTTTACGGTATCGCCGCTCCCGATTTCGAGCGCAT				
	2920	2930	2940	2950	CGCCTTCTATCGCCTTCTTGACGAGTTCTTCTGAGCGGGAC	
G0264204-1.seq(1>5446)	→	CGCCTTCTATCGCCTTCTTGACGAGTTCTTCTGAGCGGGAC				
	2960	2970	2980	2990	TCTGGGGTTCGAAATGACCGACCAAGCGACGCCCAACCTGC	
G0264204-1.seq(1>5446)	→	TCTGGGGTTCGAAATGACCGACCAAGCGACGCCCAACCTGC				
	3000	3010	3020	3030	CATCACGAGATTTTCGATTCCACCGCCGCCTTCTATGAAAGG	
G0264204-1.seq(1>5446)	→	CATCACGAGATTTTCGATTCCACCGCCGCCTTCTATGAAAGG				
	3040	3050	3060	3070	TTGGGCTTCGGAATCGTTTTCCGGGACGCCGGCTGGATGAT	
G0264204-1.seq(1>5446)	→	TTGGGCTTCGGAATCGTTTTCCGGGACGCCGGCTGGATGAT				
	3080	3090	3100	3110	CCTCCAGCGCGGGGATCTCATGCTGGAGTTCTTCGCCACC	
G0264204-1.seq(1>5446)	→	CCTCCAGCGCGGGGATCTCATGCTGGAGTTCTTCGCCACC				
	3120	3130	3140	3150	CCAACTTGTTTATTGCAGCTTATAATGGTTACAAATAAAGC	
G0264204-1.seq(1>5446)	→	CCAACTTGTTTATTGCAGCTTATAATGGTTACAAATAAAGC				
	3160	3170	3180	3190	AATAGCATCACAATTTACAAATAAAGCATTFTTTTCACT	
G0264204-1.seq(1>5446)	→	AATAGCATCACAATTTACAAATAAAGCATTFTTTTCACT				
	3200	3210	3220	3230	GCATTCTAGTTGTGGTTTGTCCAAACTCATCAATGTATCTT	
G0264204-1.seq(1>5446)	→	GCATTCTAGTTGTGGTTTGTCCAAACTCATCAATGTATCTT				
	3240	3250	3260	3270	3280	ATCATGTCTGTATACCGTCGACCTCTAGCTAGAGCTTGGCG
G0264204-1.seq(1>5446)	→	ATCATGTCTGTATACCGTCGACCTCTAGCTAGAGCTTGGCG				

Project: G0264204-1.SQD Contig 1

		3290	3300	3310	3320	
		TAATCATGGTCATAGCTGTTTCCTGTGTGAAATTGTTATCC				
G0264204-1.seq(1>5446)	→	TAATCATGGTCATAGCTGTTTCCTGTGTGAAATTGTTATCC				
		3330	3340	3350	3360	
		GCTCACAATTCCACACAACATACGAGCCGGAAGCATAAAGT				
G0264204-1.seq(1>5446)	→	GCTCACAATTCCACACAACATACGAGCCGGAAGCATAAAGT				
		3370	3380	3390	3400	
		GTAAAGCCTGGGGTGCCTAATGAGTGAGCTAACTCACATTA				
G0264204-1.seq(1>5446)	→	GTAAAGCCTGGGGTGCCTAATGAGTGAGCTAACTCACATTA				
		3410	3420	3430	3440	
		ATTGCGTTGCGCTCACTGCCCGCTTTCAGTCGGGAAACCT				
G0264204-1.seq(1>5446)	→	ATTGCGTTGCGCTCACTGCCCGCTTTCAGTCGGGAAACCT				
		3450	3460	3470	3480	
		GTCGTGCCAGCTGCATTAATGAATCGGCCAACGCGCGGGGA				
G0264204-1.seq(1>5446)	→	GTCGTGCCAGCTGCATTAATGAATCGGCCAACGCGCGGGGA				
		3490	3500	3510	3520	
		GAGGCGTTTGCATATTGGGCGCTCTTCCGCTTCCTCGCTC				
G0264204-1.seq(1>5446)	→	GAGGCGTTTGCATATTGGGCGCTCTTCCGCTTCCTCGCTC				
		3530	3540	3550	3560	
		ACTGACTCGCTGCGCTCGGTCGTTCCGGCTGCGGCGAGCGGT				
G0264204-1.seq(1>5446)	→	ACTGACTCGCTGCGCTCGGTCGTTCCGGCTGCGGCGAGCGGT				
		3570	3580	3590	3600	
		ATCAGCTCACTCAAAGGCGGTAATACGGTTATCCACAGAAT				
G0264204-1.seq(1>5446)	→	ATCAGCTCACTCAAAGGCGGTAATACGGTTATCCACAGAAT				
		3610	3620	3630	3640	
		CAGGGGATAACGCAGGAAAGAACATGTGAGCAAAAGGCCAG				
G0264204-1.seq(1>5446)	→	CAGGGGATAACGCAGGAAAGAACATGTGAGCAAAAGGCCAG				
		3650	3660	3670	3680	3690
		CAAAAGGCCAGGAACCGTAAAAAGGCCGCGTTGCTGGCGTT				
G0264204-1.seq(1>5446)	→	CAAAAGGCCAGGAACCGTAAAAAGGCCGCGTTGCTGGCGTT				
		3700	3710	3720	3730	
		TTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAATC				
G0264204-1.seq(1>5446)	→	TTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAATC				
		3740	3750	3760	3770	
		GACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAA				
G0264204-1.seq(1>5446)	→	GACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAA				

Project: G0264204-1.SQD Contig 1

		3780	3790	3800	3810	
		AGATACCAGGCGTTTCCCCCTGGAAGCTCCCTCGTGCGCTC				
G0264204-1.seq(1>5446)	→	AGATACCAGGCGTTTCCCCCTGGAAGCTCCCTCGTGCGCTC				
		3820	3830	3840	3850	
		TCCTGTTCCGACCCTGCCGCTTACCGGATACCTGTCCGCCT				
G0264204-1.seq(1>5446)	→	TCCTGTTCCGACCCTGCCGCTTACCGGATACCTGTCCGCCT				
		3860	3870	3880	3890	
		TTCTCCCTTCGGAAGCGTGGCGCTTCTCAATGCTCACGC				
G0264204-1.seq(1>5446)	→	TTCTCCCTTCGGAAGCGTGGCGCTTCTCAATGCTCACGC				
		3900	3910	3920	3930	
		TGTAGGTATCTCAGTTCGGTGTAGGTCGTTTCGCTCCAAGCT				
G0264204-1.seq(1>5446)	→	TGTAGGTATCTCAGTTCGGTGTAGGTCGTTTCGCTCCAAGCT				
		3940	3950	3960	3970	
		GGGCTGTGTGCACGAACCCCCGTTTCAGCCCGACCGCTGCC				
G0264204-1.seq(1>5446)	→	GGGCTGTGTGCACGAACCCCCGTTTCAGCCCGACCGCTGCC				
		3980	3990	4000	4010	
		CCTTATCCGGTAACTATCGTCTTGAGTCCAACCCGGTAAGA				
G0264204-1.seq(1>5446)	→	CCTTATCCGGTAACTATCGTCTTGAGTCCAACCCGGTAAGA				
		4020	4030	4040	4050	
		CACGACTTATCGCCACTGGCAGCAGCCACTGGTAACAGGAT				
G0264204-1.seq(1>5446)	→	CACGACTTATCGCCACTGGCAGCAGCCACTGGTAACAGGAT				
		4060	4070	4080	4090	4100
		TAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTTCTTGA				
G0264204-1.seq(1>5446)	→	TAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTTCTTGA				
		4110	4120	4130	4140	
		AGTGGTGGCCTAACTACGGCTACACTAGAAGGACAGTATTT				
G0264204-1.seq(1>5446)	→	AGTGGTGGCCTAACTACGGCTACACTAGAAGGACAGTATTT				
		4150	4160	4170	4180	
		GGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGGAAAAAG				
G0264204-1.seq(1>5446)	→	GGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGGAAAAAG				
		4190	4200	4210	4220	
		AGTTGGTAGCTCTTGATCCGGCAAACAACACCACCGCTGGTA				
G0264204-1.seq(1>5446)	→	AGTTGGTAGCTCTTGATCCGGCAAACAACACCACCGCTGGTA				
		4230	4240	4250	4260	
		GCGGTGGTTTTTTTTGTTTGAAGCAGCAGATTACGCGCAGA				
G0264204-1.seq(1>5446)	→	GCGGTGGTTTTTTTTGTTTGAAGCAGCAGATTACGCGCAGA				

Project: G0264204-1.SQD Contig 1

	4760	4770	4780	4790	GCCATTGCTACAGGCATCGTGGTGTACAGCTCGTCGTTTGG	
G0264204-1.seq(1>5446)	→	GCCATTGCTACAGGCATCGTGGTGTACAGCTCGTCGTTTGG				
	4800	4810	4820	4830	TATGGCTTCATTCAGCTCCGGTTCCCAACGATCAAGGCGAG	
G0264204-1.seq(1>5446)	→	TATGGCTTCATTCAGCTCCGGTTCCCAACGATCAAGGCGAG				
	4840	4850	4860	4870	TTACATGATCCCCCATGTTGTGCAAAAAAGCGTTAGCTCC	
G0264204-1.seq(1>5446)	→	TTACATGATCCCCCATGTTGTGCAAAAAAGCGTTAGCTCC				
	4880	4890	4900	4910	4920	TTCGGTCTCCGATCGTTGTGAGAAGTAAGTTGGCCGCAGT
G0264204-1.seq(1>5446)	→	TTCGGTCTCCGATCGTTGTGAGAAGTAAGTTGGCCGCAGT				
	4930	4940	4950	4960	GTTATCACTCATGGTTATGGCAGCACTGCATAATTCTCTTA	
G0264204-1.seq(1>5446)	→	GTTATCACTCATGGTTATGGCAGCACTGCATAATTCTCTTA				
	4970	4980	4990	5000	CTGTCATGCCATCCGTAAGATGCTTTTCTGTGACTGGTGAG	
G0264204-1.seq(1>5446)	→	CTGTCATGCCATCCGTAAGATGCTTTTCTGTGACTGGTGAG				
	5010	5020	5030	5040	TACTCAACCAAGTCATTCTGAGAATAGTGTATGCGGCGACC	
G0264204-1.seq(1>5446)	→	TACTCAACCAAGTCATTCTGAGAATAGTGTATGCGGCGACC				
	5050	5060	5070	5080	GAGTTGCTCTTGCCCGGCGTCAATACGGGATAATACCGCGC	
G0264204-1.seq(1>5446)	→	GAGTTGCTCTTGCCCGGCGTCAATACGGGATAATACCGCGC				
	5090	5100	5110	5120	CACATAGCAGAACTTTAAAAGTGCTCATCATTGAAAACGT	
G0264204-1.seq(1>5446)	→	CACATAGCAGAACTTTAAAAGTGCTCATCATTGAAAACGT				
	5130	5140	5150	5160	TCTTCGGGGCGAAAACCTCTCAAGGATCTTACCGCTGTTGAG	
G0264204-1.seq(1>5446)	→	TCTTCGGGGCGAAAACCTCTCAAGGATCTTACCGCTGTTGAG				
	5170	5180	5190	5200	ATCCAGTTCGATGTAACCCACTCGTGCACCCAACCTGATCTT	
G0264204-1.seq(1>5446)	→	ATCCAGTTCGATGTAACCCACTCGTGCACCCAACCTGATCTT				
	5210	5220	5230	5240	CAGCATCTTTTACTTTCACCAGCGTTTCTGGGTGAGCAAAA	
G0264204-1.seq(1>5446)	→	CAGCATCTTTTACTTTCACCAGCGTTTCTGGGTGAGCAAAA				

	5250	5260	5270	5280	ACAGGAAGGCAAAATGCCGCAAAAAGGGAATAAGGGCGAC	
G0264204-1.seq(1>5446)	→	ACAGGAAGGCAAAATGCCGCAAAAAGGGAATAAGGGCGAC				
	5290	5300	5310	5320	5330	ACGGAAATGTTGAATACTCATACTCTTCCTTTTCAATATT
G0264204-1.seq(1>5446)	→	ACGGAAATGTTGAATACTCATACTCTTCCTTTTCAATATT				
		5340	5350	5360	5370	ATTGAAGCATTTATCAGGGTTATTGTCTCATGAGCGGATAC
G0264204-1.seq(1>5446)	→	ATTGAAGCATTTATCAGGGTTATTGTCTCATGAGCGGATAC				
		5380	5390	5400	5410	ATATTTGAATGTATTTAGAAAAATAAACAAATAGGGGTTC
G0264204-1.seq(1>5446)	→	ATATTTGAATGTATTTAGAAAAATAAACAAATAGGGGTTC				
		5420	5430	5440	GCGCACATTTCCCCGAAAAGTGCCACCTGACGTC	
G0264204-1.seq(1>5446)	→	GCGCACATTTCCCCGAAAAGTGCCACCTGACGTC				