

Project: G0275879-2.SQD Contig 1

		10 20 30	CCATTCGCCATTCAGGCTGCGCAACTGTTGGGAAGG
G0275879-2.seq(1>4679)	→		CCATTCGCCATTCAGGCTGCGCAACTGTTGGGAAGG
		40 50 60 70	GCGATCGGTGCGGGCCTCTTCGCTATTACGCCAGCT
G0275879-2.seq(1>4679)	→		GCGATCGGTGCGGGCCTCTTCGCTATTACGCCAGCT
		80 90 100	GGCGAAAGGGGGATGTGCTGCAAGGCGATTAAGTTG
G0275879-2.seq(1>4679)	→		GGCGAAAGGGGGATGTGCTGCAAGGCGATTAAGTTG
		110 120 130 140	GGTAACGCCAGGGTTTCCCAGTCACGACGTTGTAA
G0275879-2.seq(1>4679)	→		GGTAACGCCAGGGTTTCCCAGTCACGACGTTGTAA
		150 160 170 180	AACGACGGCCAGTGCCAAGCTGATCTATACATTGAA
G0275879-2.seq(1>4679)	→		AACGACGGCCAGTGCCAAGCTGATCTATACATTGAA
		190 200 210	TCAATATTGGCAATTAGCCATATTAGTCATTGGTTA
G0275879-2.seq(1>4679)	→		TCAATATTGGCAATTAGCCATATTAGTCATTGGTTA
		220 230 240 250	TATAGCATAAATCAATATTGGCTATTGGCCATTGCA
G0275879-2.seq(1>4679)	→		TATAGCATAAATCAATATTGGCTATTGGCCATTGCA
		260 270 280	TACGTTGTATCTATATCATAATATGTACATTTATAT
G0275879-2.seq(1>4679)	→		TACGTTGTATCTATATCATAATATGTACATTTATAT
		290 300 310 320	TGGCTCATGTCCAATATGACCGCCATGTTGACATTG
G0275879-2.seq(1>4679)	→		TGGCTCATGTCCAATATGACCGCCATGTTGACATTG
		330 340 350 360	ATTATTGACTAGTTATTAATAGTAATCAATTACGGG
G0275879-2.seq(1>4679)	→		ATTATTGACTAGTTATTAATAGTAATCAATTACGGG
		370 380 390	GTCATTAGTTCATAGCCCATATATGGAGTTCGCGT
G0275879-2.seq(1>4679)	→		GTCATTAGTTCATAGCCCATATATGGAGTTCGCGT
		400 410 420 430	TACATAACTTACGGTAAATGGCCCGCTGGCTGACC
G0275879-2.seq(1>4679)	→		TACATAACTTACGGTAAATGGCCCGCTGGCTGACC

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		440 450 460	GCCCAACGACCCCCGCCATTGACGTCAATAATGAC
G0275879-2.seq (1>4679)	→		GCCCAACGACCCCCGCCATTGACGTCAATAATGAC
		470 480 490 500	GTATGTTCCCATAGTAACGCCAATAGGGACTTTCCA
G0275879-2.seq (1>4679)	→		GTATGTTCCCATAGTAACGCCAATAGGGACTTTCCA
		510 520 530 540	TTGACGTCAATGGGTGGAGTATTTACGGTAAACTGC
G0275879-2.seq (1>4679)	→		TTGACGTCAATGGGTGGAGTATTTACGGTAAACTGC
		550 560 570	CCACTTGGCAGTACATCAAGTGTATCATATGCCAAG
G0275879-2.seq (1>4679)	→		CCACTTGGCAGTACATCAAGTGTATCATATGCCAAG
		580 590 600 610	TCCGCCCCCTATTGACGTCAATGACGGTAAATGGCC
G0275879-2.seq (1>4679)	→		TCCGCCCCCTATTGACGTCAATGACGGTAAATGGCC
		620 630 640	CGCCTGGCATTATGCCCAGTACATGACCTTACGGGA
G0275879-2.seq (1>4679)	→		CGCCTGGCATTATGCCCAGTACATGACCTTACGGGA
		650 660 670 680	CTTTCCTACTTGGCAGTACATCTACGTATTAGTCAT
G0275879-2.seq (1>4679)	→		CTTTCCTACTTGGCAGTACATCTACGTATTAGTCAT
		690 700 710 720	CGCTATTACCATGGTGATGCGGTTTTGGCAGTACAC
G0275879-2.seq (1>4679)	→		CGCTATTACCATGGTGATGCGGTTTTGGCAGTACAC
		730 740 750	CAATGGGCGTGGATAGCGGTTTGACTCACGGGGATT
G0275879-2.seq (1>4679)	→		CAATGGGCGTGGATAGCGGTTTGACTCACGGGGATT
		760 770 780 790	TCCAAGTCTCACCACCCATTGACGTCAATGGGAGTTT
G0275879-2.seq (1>4679)	→		TCCAAGTCTCACCACCCATTGACGTCAATGGGAGTTT
		800 810 820	GTTTTGGCACCAAAATCAACGGGACTTTCCAAAATG
G0275879-2.seq (1>4679)	→		GTTTTGGCACCAAAATCAACGGGACTTTCCAAAATG
		830 840 850 860	TCGTAATAACCCCGCCCCGTTGACGCAAATGGGCGG
G0275879-2.seq (1>4679)	→		TCGTAATAACCCCGCCCCGTTGACGCAAATGGGCGG

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		870	880	890	900
		TAGGCGTGTACGGTGGGAGGTCTATATAAGCAGAGC			
G0275879-2.seq(1>4679)	→	TAGGCGTGTACGGTGGGAGGTCTATATAAGCAGAGC			
		910	920	930	
		TCGTTTAGTGAACCGTCAGAATTGATCTACCATGGA			
G0275879-2.seq(1>4679)	→	TCGTTTAGTGAACCGTCAGAATTGATCTACCATGGA			
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→	CCATGGA			
		940	950	960	970
		CTACAAAGACGATGACGACAAGCTTGCGGCCGCGAA			
G0275879-2.seq(1>4679)	→	CTACAAAGACGATGACGACAAGCTTGCGGCCGCGAA			
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→	CTACAAAGACGATGACGACAAGCTTGCGGCCGCGAA			
		980	990	1000	
		TTCATCGATAGATCTGATATCGGTACCAGTCGACTC			
G0275879-2.seq(1>4679)	→	TTCATCGATAGATCTGATATCGGTACCAGTCGACTC			
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→	TTCATCGATAGATCTGATATCGGTACCAGTCGACTC			
		1010	1020	1030	1040
		TAGAGGATCCCGGGTGGCATCCCTGTGACCCCTCCC			
G0275879-2.seq(1>4679)	→	TAGAGGATCCCGGGTGGCATCCCTGTGACCCCTCCC			
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→	TAGAGGATCCCGGGTGGCATCCCTGTGACCCCTCCC			
		1050	1060	1070	1080
		CAGTGCCTCTCCTGGCCCTGGAAGTTGCCACTCCAG			
G0275879-2.seq(1>4679)	→	CAGTGCCTCTCCTGGCCCTGGAAGTTGCCACTCCAG			
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→	CAGTGCCTCTCCTGGCCCTGGAAGTTGCCACTCCAG			
		1090	1100	1110	
		TGCCCACCAGCCTTGTCTAATAAAAATTAAGTTGCA			
G0275879-2.seq(1>4679)	→	TGCCCACCAGCCTTGTCTAATAAAAATTAAGTTGCA			
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→	TGCCCACCAGCCTTGTCTAATAAAAATTAAGTTGCA			
		1120	1130	1140	1150
		TCATTTTGTCTGACTAGGTGTCCTTCTATAATATTA			
G0275879-2.seq(1>4679)	→	TCATTTTGTCTGACTAGGTGTCCTTCTATAATATTA			
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→	TCATTTTGTCTGACTAGGTGTCCTTCTATAATATTA			
		1160	1170	1180	
		TGGGGTGGAGGGGGTGGTATGGAGCAAGGGGCAAG			
G0275879-2.seq(1>4679)	→	TGGGGTGGAGGGGGTGGTATGGAGCAAGGGGCAAG			
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→	TGGGGTGGAGGGGGTGGTATGGAGCAAGGGGCAAG			
		1190	1200	1210	1220
		TTGGGAAGACAACCTGTAGGGCCTGCGGGGTCTATT			
G0275879-2.seq(1>4679)	→	TTGGGAAGACAACCTGTAGGGCCTGCGGGGTCTATT			
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→	TTGGGAAGACAACCTGTAGGGCCTGCGGGGTCTATT			

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		1230 1240 1250 1260	GGGAACCAAGCTGGAGTGCAGTGGCACAATCTTGGC
G0275879-2.seq(1>4679)	→		GGGAACCAAGCTGGAGTGCAGTGGCACAATCTTGGC
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		GGGAACCAAGCTGGAGTGCAGTGGCACAATCTTGGC
		1270 1280 1290	TCACTGCAATCTCCGCCTCCTGGGTTCAAGCGATTC
G0275879-2.seq(1>4679)	→		TCACTGCAATCTCCGCCTCCTGGGTTCAAGCGATTC
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		TCACTGCAATCTCCGCCTCCTGGGTTCAAGCGATTC
		1300 1310 1320 1330	TCCTGCCTCAGCCTCCCGAGTTGTTGGGATTCCAGG
G0275879-2.seq(1>4679)	→		TCCTGCCTCAGCCTCCCGAGTTGTTGGGATTCCAGG
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		TCCTGCCTCAGCCTCCCGAGTTGTTGGGATTCCAGG
		1340 1350 1360	CATGCATGACCAGGCTCAGCTAATTTTTGTTTTTTTT
G0275879-2.seq(1>4679)	→		CATGCATGACCAGGCTCAGCTAATTTTTGTTTTTTTT
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		CATGCATGACCAGGCTCAGCTAATTTTTGTTTTTTTT
		1370 1380 1390 1400	GGTAGAGACGGGGTTTCACCATATTGGCCAGGCTGG
G0275879-2.seq(1>4679)	→		GGTAGAGACGGGGTTTCACCATATTGGCCAGGCTGG
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		GGTAGAGACGGGGTTTCACCATATTGGCCAGGCTGG
		1410 1420 1430 1440	TCTCCAACCTCCTAATCTCAGGTGATCTACCCACCTT
G0275879-2.seq(1>4679)	→		TCTCCAACCTCCTAATCTCAGGTGATCTACCCACCTT
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		TCTCCAACCTCCTAATCTCAGGTGATCTACCCACCTT
		1450 1460 1470	GGCCTCCCAAATTGCTGGGATTACAGGCGTGAACCA
G0275879-2.seq(1>4679)	→		GGCCTCCCAAATTGCTGGGATTACAGGCGTGAACCA
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		GGCCTCCCAAATTGCTGGGATTACAGGCGTGAACCA
		1480 1490 1500 1510	CTGCTCCCTTCCCTGTCCTTCTGATTTTAAAATAAC
G0275879-2.seq(1>4679)	→		CTGCTCCCTTCCCTGTCCTTCTGATTTTAAAATAAC
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		CTGCTCCCTTCCCTGTCCTTCTGATTTTAAAATAAC
		1520 1530 1540	TATACCAGCAGGAGGACGTCCAGACACAGCATAGGC
G0275879-2.seq(1>4679)	→		TATACCAGCAGGAGGACGTCCAGACACAGCATAGGC
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		TATACCAGCAGGAGGACGTCCAGACACAGCATAGGC
		1550 1560 1570 1580	TACCTGGCCATGCCCAACCGGTGGGACATTTGAGTT
G0275879-2.seq(1>4679)	→		TACCTGGCCATGCCCAACCGGTGGGACATTTGAGTT
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		TACCTGGCCATGCCCAACCGGTGGGACATTTGAGTT

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		1590 1600 1610 1620	GCTTGCTTGGCACTGTCCTCTCATGCGTTGGGTCCA
G0275879-2.seq(1>4679)	→		GCTTGCTTGGCACTGTCCTCTCATGCGTTGGGTCCA
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		GCTTGCTTGGCACTGTCCTCTCATGCGTTGGGTCCA
		1630 1640 1650	CTCAGTAGATGCCTGTTGAATTGGGTACGCGGCCAG
G0275879-2.seq(1>4679)	→		CTCAGTAGATGCCTGTTGAATTGGGTACGCGGCCAG
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		CTCAGTAGATGCCTGTTGAATTGGGTACGCGGCCAG
		1660 1670 1680 1690	CTTGGCTGTGGAATGTGTGTCAGTTAGGGTGTGGAA
G0275879-2.seq(1>4679)	→		CTTGGCTGTGGAATGTGTGTCAGTTAGGGTGTGGAA
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		CTTGGCTGTGGAATGTGTGTCAGTTAGGGTGTGGAA
		1700 1710 1720	AGTCCCCAGGCTCCCCAGCAGGCAGAAGTATGCAA
G0275879-2.seq(1>4679)	→		AGTCCCCAGGCTCCCCAGCAGGCAGAAGTATGCAA
A06-D46343-G0275879-2_PEGFP-N-5.ab1(1>795)	→		AGTCCCCAGGCTCCCCAGCAGGCAGAAGTATG
		1730 1740 1750 1760	GCATGCATCTCAATTAGTCAGCAACCAGGTGTGGAA
G0275879-2.seq(1>4679)	→		GCATGCATCTCAATTAGTCAGCAACCAGGTGTGGAA
		1770 1780 1790 1800	AGTCCCCAGGCTCCCCAGCAGGCAGAAGTATGCAA
G0275879-2.seq(1>4679)	→		AGTCCCCAGGCTCCCCAGCAGGCAGAAGTATGCAA
		1810 1820 1830	GCATGCATCTCAATTAGTCAGCAACCATAGTCCCGC
G0275879-2.seq(1>4679)	→		GCATGCATCTCAATTAGTCAGCAACCATAGTCCCGC
		1840 1850 1860 1870	CCCTAACTCCGCCCATCCCGCCCCTAACTCCGCCCA
G0275879-2.seq(1>4679)	→		CCCTAACTCCGCCCATCCCGCCCCTAACTCCGCCCA
		1880 1890 1900	GTTCCGCCCATTTCTCCGCCCATGGCTGACTAATTT
G0275879-2.seq(1>4679)	→		GTTCCGCCCATTTCTCCGCCCATGGCTGACTAATTT
		1910 1920 1930 1940	TTTTTATTTATGCAGAGGCCGAGGCCGCTCGGCCT
G0275879-2.seq(1>4679)	→		TTTTTATTTATGCAGAGGCCGAGGCCGCTCGGCCT
		1950 1960 1970 1980	CTGAGCTATTCCAGAAGTAGTGAGGAGGCTTTTTTG
G0275879-2.seq(1>4679)	→		CTGAGCTATTCCAGAAGTAGTGAGGAGGCTTTTTTG

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		1990	2000	2010	
		GAGGCCTAGGCTTTTGCAAAAAGCTCCTCGAGGAAC			
G0275879-2.seq(1>4679)	→	GAGGCCTAGGCTTTTGCAAAAAGCTCCTCGAGGAAC			
		2020	2030	2040	2050
		TGAAAAACCAGAAAGTTAATTCCTATAGTGAGTCCG			
G0275879-2.seq(1>4679)	→	TGAAAAACCAGAAAGTTAATTCCTATAGTGAGTCCG			
		2060	2070	2080	
		TATTAAATTCGTAATCATGTCATAGCTGTTTCCTGT			
G0275879-2.seq(1>4679)	→	TATTAAATTCGTAATCATGTCATAGCTGTTTCCTGT			
		2090	2100	2110	2120
		GTGAAATTGTTATCCGCTCACAATTCCACACAACAT			
G0275879-2.seq(1>4679)	→	GTGAAATTGTTATCCGCTCACAATTCCACACAACAT			
		2130	2140	2150	2160
		ACGAGCCGGAAGCATAAAGTGTAAGCCTGGGGTGC			
G0275879-2.seq(1>4679)	→	ACGAGCCGGAAGCATAAAGTGTAAGCCTGGGGTGC			
		2170	2180	2190	
		CTAATGAGTGAGCTAACTCACATTAATTGCGTTGCG			
G0275879-2.seq(1>4679)	→	CTAATGAGTGAGCTAACTCACATTAATTGCGTTGCG			
		2200	2210	2220	2230
		CTCACTGCCCCTTTCCAGTCGGGAAACCTGTCTGTG			
G0275879-2.seq(1>4679)	→	CTCACTGCCCCTTTCCAGTCGGGAAACCTGTCTGTG			
		2240	2250	2260	
		CCAGCTGCATTAATGAATCGGCCAACGCGCGGGGAG			
G0275879-2.seq(1>4679)	→	CCAGCTGCATTAATGAATCGGCCAACGCGCGGGGAG			
		2270	2280	2290	2300
		AGGCGGTTTGCCTATTGGGCGCTCTTCCGCTTCCTC			
G0275879-2.seq(1>4679)	→	AGGCGGTTTGCCTATTGGGCGCTCTTCCGCTTCCTC			
		2310	2320	2330	2340
		GCTCACTGACTCGCTGCGCTCGGTTCGTTTCGGCTGCG			
G0275879-2.seq(1>4679)	→	GCTCACTGACTCGCTGCGCTCGGTTCGTTTCGGCTGCG			
		2350	2360	2370	
		GCGAGCGGTATCAGCTCACTCAAAGGCGGTAATACG			
G0275879-2.seq(1>4679)	→	GCGAGCGGTATCAGCTCACTCAAAGGCGGTAATACG			
		2380	2390	2400	2410
		GTTATCCACAGAATCAGGGGATAACGCAGGAAAGAA			
G0275879-2.seq(1>4679)	→	GTTATCCACAGAATCAGGGGATAACGCAGGAAAGAA			

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		2420 2430 2440	CATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCG
G0275879-2.seq(1>4679)	→		CATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCG
		2450 2460 2470 2480	TAAAAAGGCCGCGTTGCTGGCGTTTTTCCATAGGCT
G0275879-2.seq(1>4679)	→		TAAAAAGGCCGCGTTGCTGGCGTTTTTCCATAGGCT
		2490 2500 2510 2520	CCGCCCCCTGACGAGCATCACAAAATCGACGCTC
G0275879-2.seq(1>4679)	→		CCGCCCCCTGACGAGCATCACAAAATCGACGCTC
		2530 2540 2550	AAGTCAGAGGTGGCGAAACCCGACAGGACTATAAAG
G0275879-2.seq(1>4679)	→		AAGTCAGAGGTGGCGAAACCCGACAGGACTATAAAG
		2560 2570 2580 2590	ATACCAGGCGTTTCCCCCTGGAAGCTCCCTCGTGCG
G0275879-2.seq(1>4679)	→		ATACCAGGCGTTTCCCCCTGGAAGCTCCCTCGTGCG
		2600 2610 2620	CTCTCCTGTTCCGACCCTGCCGCTTACCGGATACCT
G0275879-2.seq(1>4679)	→		CTCTCCTGTTCCGACCCTGCCGCTTACCGGATACCT
		2630 2640 2650 2660	GTCCGCCTTTCTCCCTTCGGGAAGCGTGGCGCTTTC
G0275879-2.seq(1>4679)	→		GTCCGCCTTTCTCCCTTCGGGAAGCGTGGCGCTTTC
		2670 2680 2690 2700	TCATAGCTCACGCTGTAGGTATCTCAGTTCGGTGTAA
G0275879-2.seq(1>4679)	→		TCATAGCTCACGCTGTAGGTATCTCAGTTCGGTGTAA
		2710 2720 2730	GGTCGTTTCGCTCCAAGCTGGGCTGTGTGCACGAACC
G0275879-2.seq(1>4679)	→		GGTCGTTTCGCTCCAAGCTGGGCTGTGTGCACGAACC
		2740 2750 2760 2770	CCCCGTTTCAGCCCGACCGCTGCGCCTTATCCGGTAA
G0275879-2.seq(1>4679)	→		CCCCGTTTCAGCCCGACCGCTGCGCCTTATCCGGTAA
		2780 2790 2800	CTATCGTCTTGAGTCCAACCCGGTAAGACACGACTT
G0275879-2.seq(1>4679)	→		CTATCGTCTTGAGTCCAACCCGGTAAGACACGACTT
		2810 2820 2830 2840	ATCGCCACTGGCAGCAGCCACTGGTAACAGGATTAG
G0275879-2.seq(1>4679)	→		ATCGCCACTGGCAGCAGCCACTGGTAACAGGATTAG

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		2850	2860	2870	2880
		CAGAGCGAGGTATGTAGGCGGTGCTACAGAGTTCTT			
G0275879-2.seq(1>4679)	→	CAGAGCGAGGTATGTAGGCGGTGCTACAGAGTTCTT			
		2890	2900	2910	
		GAAGTGGTGGCCTAACTACGGCTACACTAGAAGAAC			
G0275879-2.seq(1>4679)	→	GAAGTGGTGGCCTAACTACGGCTACACTAGAAGAAC			
		2920	2930	2940	2950
		AGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTAC			
G0275879-2.seq(1>4679)	→	AGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTAC			
		2960	2970	2980	
		CTTCGGAAAAAGAGTTGGTAGCTCTTGATCCGGCAA			
G0275879-2.seq(1>4679)	→	CTTCGGAAAAAGAGTTGGTAGCTCTTGATCCGGCAA			
		2990	3000	3010	3020
		ACAAACCACCGCTGGTAGCGGTGGTTTTTTTTGTTTG			
G0275879-2.seq(1>4679)	→	ACAAACCACCGCTGGTAGCGGTGGTTTTTTTTGTTTG			
		3030	3040	3050	3060
		CAAGCAGCAGATTACGCGCAGAAAAAAGGATCTCA			
G0275879-2.seq(1>4679)	→	CAAGCAGCAGATTACGCGCAGAAAAAAGGATCTCA			
		3070	3080	3090	
		AGAAGATCCTTTGATCTTTTCTACGGGGTCTGACGC			
G0275879-2.seq(1>4679)	→	AGAAGATCCTTTGATCTTTTCTACGGGGTCTGACGC			
		3100	3110	3120	3130
		TCAGTGGAACGAAACTCACGTTAAGGGATTTTGGT			
G0275879-2.seq(1>4679)	→	TCAGTGGAACGAAACTCACGTTAAGGGATTTTGGT			
		3140	3150	3160	
		CATGAGATTATCAAAAAGGATCTTCACCTAGATCCT			
G0275879-2.seq(1>4679)	→	CATGAGATTATCAAAAAGGATCTTCACCTAGATCCT			
		3170	3180	3190	3200
		TTTAAATTAAAAATGAAGTTTTAAATCAATCTAAAG			
G0275879-2.seq(1>4679)	→	TTTAAATTAAAAATGAAGTTTTAAATCAATCTAAAG			
		3210	3220	3230	3240
		TATATATGAGTAACTTGGTCTGACAGTTACCAATG			
G0275879-2.seq(1>4679)	→	TATATATGAGTAACTTGGTCTGACAGTTACCAATG			
		3250	3260	3270	
		CTTAATCAGTGAGGCACCTATCTCAGCGATCTGTCT			
G0275879-2.seq(1>4679)	→	CTTAATCAGTGAGGCACCTATCTCAGCGATCTGTCT			

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		3710	3720	3730	3740
		ACTCATGGTTATGGCAGCACTGCATAATTCTCTTAC			
G0275879-2.seq(1>4679)	→	ACTCATGGTTATGGCAGCACTGCATAATTCTCTTAC			
		3750	3760	3770	3780
		TGTCATGCCATCCGTAAGATGCTTTTCTGTGACTGG			
G0275879-2.seq(1>4679)	→	TGTCATGCCATCCGTAAGATGCTTTTCTGTGACTGG			
		3790	3800	3810	
		TGAGTACTCAACCAAGTCATTCTGAGAATAGTGTAT			
G0275879-2.seq(1>4679)	→	TGAGTACTCAACCAAGTCATTCTGAGAATAGTGTAT			
		3820	3830	3840	3850
		GCGGCGACCGAGTTGCTCTTGCCCGGCGTCAATACG			
G0275879-2.seq(1>4679)	→	GCGGCGACCGAGTTGCTCTTGCCCGGCGTCAATACG			
		3860	3870	3880	
		GGATAATACCGCGCCACATAGCAGAACTTTAAAAGT			
G0275879-2.seq(1>4679)	→	GGATAATACCGCGCCACATAGCAGAACTTTAAAAGT			
		3890	3900	3910	3920
		GCTCATCATTGGAAAACGTTCTTCGGGGCGAAAAC			
G0275879-2.seq(1>4679)	→	GCTCATCATTGGAAAACGTTCTTCGGGGCGAAAAC			
		3930	3940	3950	3960
		CTCAAGGATCTTACCGCTGTTGAGATCCAGTTCGAT			
G0275879-2.seq(1>4679)	→	CTCAAGGATCTTACCGCTGTTGAGATCCAGTTCGAT			
		3970	3980	3990	
		GTAACCCACTCGTGCACCCAACCTGATCTTCAGCATC			
G0275879-2.seq(1>4679)	→	GTAACCCACTCGTGCACCCAACCTGATCTTCAGCATC			
		4000	4010	4020	4030
		TTTTACTTTTACCAGCGTTTCTGGGTGAGCAAAAAC			
G0275879-2.seq(1>4679)	→	TTTTACTTTTACCAGCGTTTCTGGGTGAGCAAAAAC			
		4040	4050	4060	
		AGGAAGGCAAAAATGCCGCAAAAAGGGAATAAGGGC			
G0275879-2.seq(1>4679)	→	AGGAAGGCAAAAATGCCGCAAAAAGGGAATAAGGGC			
		4070	4080	4090	4100
		GACACGGAATGTTGAATACTCATACTCTTCCTTTT			
G0275879-2.seq(1>4679)	→	GACACGGAATGTTGAATACTCATACTCTTCCTTTT			
		4110	4120	4130	4140
		TCAATATTATTGAAGCATTATCAGGGTTATTGTCT			
G0275879-2.seq(1>4679)	→	TCAATATTATTGAAGCATTATCAGGGTTATTGTCT			

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		4150	4160	4170	CATGAGCGGATACATATTTGAATGTATTTAGAAAAA	
G0275879-2.seq (1>4679)	→	CATGAGCGGATACATATTTGAATGTATTTAGAAAAA				
		4180	4190	4200	4210	TAAACAAATAGGGGTTCGCGCACATTTCCCCGAAA
G0275879-2.seq (1>4679)	→	TAAACAAATAGGGGTTCGCGCACATTTCCCCGAAA				
		4220	4230	4240	AGTGCCACCTGACGCGCCCTGTAGCGGCGCATTAAG	
G0275879-2.seq (1>4679)	→	AGTGCCACCTGACGCGCCCTGTAGCGGCGCATTAAG				
		4250	4260	4270	4280	CGCGGCGGGTGTGGTGGTTACGCGCAGCGTGACCGC
G0275879-2.seq (1>4679)	→	CGCGGCGGGTGTGGTGGTTACGCGCAGCGTGACCGC				
		4290	4300	4310	4320	TACACTTGCCAGCGCCCTAGCGCCCGCTCCTTTCGC
G0275879-2.seq (1>4679)	→	TACACTTGCCAGCGCCCTAGCGCCCGCTCCTTTCGC				
		4330	4340	4350	TTTCTTCCCTTCCTTTCTCGCCACGTTCGCCGGCTT	
G0275879-2.seq (1>4679)	→	TTTCTTCCCTTCCTTTCTCGCCACGTTCGCCGGCTT				
		4360	4370	4380	4390	TCCCCGTCAAGCTCTAAATCGGGGGCTCCCTTTAGG
G0275879-2.seq (1>4679)	→	TCCCCGTCAAGCTCTAAATCGGGGGCTCCCTTTAGG				
		4400	4410	4420	GTTCCGATTTAGTGCTTTACGGCACCTCGACCCCAA	
G0275879-2.seq (1>4679)	→	GTTCCGATTTAGTGCTTTACGGCACCTCGACCCCAA				
		4430	4440	4450	4460	AAAAC TTGATTAGGGTGATGGTTCACGTAGTGGGCC
G0275879-2.seq (1>4679)	→	AAAAC TTGATTAGGGTGATGGTTCACGTAGTGGGCC				
		4470	4480	4490	4500	ATCGCCCTGATAGACGGTTTTTCGCCCTTTGACGTT
G0275879-2.seq (1>4679)	→	ATCGCCCTGATAGACGGTTTTTCGCCCTTTGACGTT				
		4510	4520	4530	GGAGTCCACGTTCTTTAATAGTGGACTCTTGTTCCA	
G0275879-2.seq (1>4679)	→	GGAGTCCACGTTCTTTAATAGTGGACTCTTGTTCCA				
		4540	4550	4560	4570	AACTGGAACAACACTCAACCCTATCTCGGTCTATTC
G0275879-2.seq (1>4679)	→	AACTGGAACAACACTCAACCCTATCTCGGTCTATTC				

4580 4590 4600
TTTGTATTATAAGGGATTTGCCGATTTTCGGCCTA
G0275879-2.seq (1>4679) → TTTGTATTATAAGGGATTTGCCGATTTTCGGCCTA

4610 4620 4630 4640
TTGGTTAAAAAATGAGCTGATTTAACAAAAATTTAA
G0275879-2.seq (1>4679) → TTGGTTAAAAAATGAGCTGATTTAACAAAAATTTAA

4650 4660 4670
CGCGAATTTTAACAAAATATTAACGCTTACAATTT
G0275879-2.seq (1>4679) → CGCGAATTTTAACAAAATATTAACGCTTACAATTT