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		370 380 390	CATGGTGATGCGGTTTTGGCAGTACATCAA
G0251720-1.seq(1>4716)	→		CATGGTGATGCGGTTTTGGCAGTACATCAA
		400 410 420	TGGGCGTGGATAGCGGTTTGGACTCACGGGG
G0251720-1.seq(1>4716)	→		TGGGCGTGGATAGCGGTTTGGACTCACGGGG
		430 440 450	ATTTCOAAGTCTCCACCCCATTTGACGTCAA
G0251720-1.seq(1>4716)	→		ATTTCOAAGTCTCCACCCCATTTGACGTCAA
		460 470 480	TGGGAGTTTGTTTTGGCACCAAATCAACG
G0251720-1.seq(1>4716)	→		TGGGAGTTTGTTTTGGCACCAAATCAACG
		490 500 510	GGACTTTCAAAATGTCGTAACAACCTCCGC
G0251720-1.seq(1>4716)	→		GGACTTTCAAAATGTCGTAACAACCTCCGC
		520 530 540	CCCATTGACGCAAATGGGCGGTAGGCGTGT
G0251720-1.seq(1>4716)	→		CCCATTGACGCAAATGGGCGGTAGGCGTGT
		550 560 570	ACGGTGGGAGGTCTATATAAGCAGAGCTGG
G0251720-1.seq(1>4716)	→		ACGGTGGGAGGTCTATATAAGCAGAGCTGG
		580 590 600	TTTAGTGAACCGTCAGATCCGCTAGCGCTA
G0251720-1.seq(1>4716)	→		TTTAGTGAACCGTCAGATCCGCTAGCGCTA
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		TAGCGCTA
		610 620 630	CCGTCGCCACCATGAACACCCCGGGAATT
G0251720-1.seq(1>4716)	→		CCGTCGCCACCATGAACACCCCGGGAATT
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		CCGTCGCCACCATGAACACCCCGGGAATT
		640 650 660	AACCTGATCAAGGAGGACATGCGCGTGAAG
G0251720-1.seq(1>4716)	→		AACCTGATCAAGGAGGACATGCGCGTGAAG
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		AACCTGATCAAGGAGGACATGCGCGTGAAG
		670 680 690	GTGCACATGGAGGGCAACGTGAACGGCCAC
G0251720-1.seq(1>4716)	→		GTGCACATGGAGGGCAACGTGAACGGCCAC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		GTGCACATGGAGGGCAACGTGAACGGCCAC

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		700 710 720	GCCTTCGTGATCGAGGGCGAGGGCAAGGGC
G0251720-1.seq(1>4716)	→		GCCTTCGTGATCGAGGGCGAGGGCAAGGGC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		GCCTTCGTGATCGAGGGCGAGGGCAAGGGC
		730 740 750	AAGCCCTACGAGGGCACCCAGACCGCCAAC
G0251720-1.seq(1>4716)	→		AAGCCCTACGAGGGCACCCAGACCGCCAAC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		AAGCCCTACGAGGGCACCCAGACCGCCAAC
		760 770 780	CTGACCGTGAAGGAGGGCGCCCCCTGCC
G0251720-1.seq(1>4716)	→		CTGACCGTGAAGGAGGGCGCCCCCTGCC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		CTGACCGTGAAGGAGGGCGCCCCCTGCC
		790 800 810	TTCAGCTACGACATCCTGACCACCGCCGTG
G0251720-1.seq(1>4716)	→		TTCAGCTACGACATCCTGACCACCGCCGTG
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		TTCAGCTACGACATCCTGACCACCGCCGTG
		820 830 840	CACTACGGCAACCGGGTGTTCACCAAGTAC
G0251720-1.seq(1>4716)	→		CACTACGGCAACCGGGTGTTCACCAAGTAC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		CACTACGGCAACCGGGTGTTCACCAAGTAC
		850 860 870	CCCGAGGACATCCCCGACTACTTCAAGCAG
G0251720-1.seq(1>4716)	→		CCCGAGGACATCCCCGACTACTTCAAGCAG
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		CCCGAGGACATCCCCGACTACTTCAAGCAG
		880 890 900	AGCTTCCCCGAGGGCTACAGCTGGGAGCGC
G0251720-1.seq(1>4716)	→		AGCTTCCCCGAGGGCTACAGCTGGGAGCGC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		AGCTTCCCCGAGGGCTACAGCTGGGAGCGC
		910 920 930	ACCATGACCTTCGAGGACAAGGGCATCTGC
G0251720-1.seq(1>4716)	→		ACCATGACCTTCGAGGACAAGGGCATCTGC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		ACCATGACCTTCGAGGACAAGGGCATCTGC
		940 950 960	ACCATCCGCAGCGACATCAGCCTGGAGGGC
G0251720-1.seq(1>4716)	→		ACCATCCGCAGCGACATCAGCCTGGAGGGC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		ACCATCCGCAGCGACATCAGCCTGGAGGGC
		970 980 990	GACTGCTTCTTCCAGAACGTGCGCTTCAAG
G0251720-1.seq(1>4716)	→		GACTGCTTCTTCCAGAACGTGCGCTTCAAG
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		GACTGCTTCTTCCAGAACGTGCGCTTCAAG

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		1000 1010 1020 GGCACCAACTTCCCCCCAACGGCCCCGTG
G0251720-1.seq(1>4716)	→	GGCACCAACTTCCCCCCAACGGCCCCGTG
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→	GGCACCAACTTCCCCCCAACGGCCCCGTG
		1030 1040 1050 ATGCAGAAGAAGACCCTGAAGTGGGAGCCC
G0251720-1.seq(1>4716)	→	ATGCAGAAGAAGACCCTGAAGTGGGAGCCC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→	ATGCAGAAGAAGACCCTGAAGTGGGAGCCC
		1060 1070 1080 AGCACCGAGAAGCTGCACGTGCGCGACGGC
G0251720-1.seq(1>4716)	→	AGCACCGAGAAGCTGCACGTGCGCGACGGC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→	AGCACCGAGAAGCTGCACGTGCGCGACGGC
		1090 1100 1110 CTGCTGGTGGGCAACATCAACATGGCCCTG
G0251720-1.seq(1>4716)	→	CTGCTGGTGGGCAACATCAACATGGCCCTG
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→	CTGCTGGTGGGCAACATCAACATGGCCCTG
		1120 1130 1140 CTGCTGGAGGGCGGCGGCCACTACCTGTGC
G0251720-1.seq(1>4716)	→	CTGCTGGAGGGCGGCGGCCACTACCTGTGC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→	CTGCTGGAGGGCGGCGGCCACTACCTGTGC
		1150 1160 1170 GACTTCAAGACCACCTACAAGGCCAAGAAG
G0251720-1.seq(1>4716)	→	GACTTCAAGACCACCTACAAGGCCAAGAAG
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→	GACTTCAAGACCACCTACAAGGCCAAGAAG
		1180 1190 1200 GTGGTGCAGCTGCCCGACGCCCACTTCGTG
G0251720-1.seq(1>4716)	→	GTGGTGCAGCTGCCCGACGCCCACTTCGTG
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→	GTGGTGCAGCTGCCCGACGCCCACTTCGTG
		1210 1220 1230 GACCACCGCATCGAGATCCTGGGCAACGAC
G0251720-1.seq(1>4716)	→	GACCACCGCATCGAGATCCTGGGCAACGAC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→	GACCACCGCATCGAGATCCTGGGCAACGAC
		1240 1250 1260 AGCGACTACAACAAGGTGAAGCTGTACGAG
G0251720-1.seq(1>4716)	→	AGCGACTACAACAAGGTGAAGCTGTACGAG
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→	AGCGACTACAACAAGGTGAAGCTGTACGAG
		1270 1280 1290 CACGCCGTGGCCCGCTACAGCCCCCTGCC
G0251720-1.seq(1>4716)	→	CACGCCGTGGCCCGCTACAGCCCCCTGCC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→	CACGCCGTGGCCCGCTACAGCCCCCTGCC

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		1300 1310 1320	AGCCAGGTGTGGTCCGGCGACAGCGGCGTG
G0251720-1.seq(1>4716)	→		AGCCAGGTGTGGTCCGGCGACAGCGGCGTG
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		AGCCAGGTGTGGTCCGGCGACAGCGGCGTG
		1330 1340 1350	TACAAGACTCGAGCTCAAGCTTCGAATTCT
G0251720-1.seq(1>4716)	→		TACAAGACTCGAGCTCAAGCTTCGAATTCT
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		TACAAGACTCGAGCTCAAGCTTCGAATTCT
		1360 1370 1380	GCAGTCGACGGTACCGCGGGCCCGGGATCC
G0251720-1.seq(1>4716)	→		GCAGTCGACGGTACCGCGGGCCCGGGATCC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		GCAGTCGACGGTACCGCGGGCCCGGGATCC
		1390 1400 1410	ACCGGATCTAGATAACTGATCATAATCAGC
G0251720-1.seq(1>4716)	→		ACCGGATCTAGATAACTGATCATAATCAGC
A03-E45006-G0251720-1_PEGFP-PEGFP-N-5.ab1(1>793)	→		ACCGG
		1420 1430 1440	CATACCACATTTGTAGAGGTTTTACTTGCT
G0251720-1.seq(1>4716)	→		CATACCACATTTGTAGAGGTTTTACTTGCT
		1450 1460 1470	TTAAAAAACCTCCCACACCTCCCCCTGAAC
G0251720-1.seq(1>4716)	→		TTAAAAAACCTCCCACACCTCCCCCTGAAC
		1480 1490 1500	CTGAAACATAAAATGAATGCAATTGTTGTT
G0251720-1.seq(1>4716)	→		CTGAAACATAAAATGAATGCAATTGTTGTT
		1510 1520 1530	GTTAACTTGTTTATTGCAGCTTATAATGGT
G0251720-1.seq(1>4716)	→		GTTAACTTGTTTATTGCAGCTTATAATGGT
		1540 1550 1560	TACAAATAAAGCAATAGCATCACAAATTTTC
G0251720-1.seq(1>4716)	→		TACAAATAAAGCAATAGCATCACAAATTTTC
		1570 1580 1590	ACAAATAAAGCATTTTTTTTCACTGCATTCT
G0251720-1.seq(1>4716)	→		ACAAATAAAGCATTTTTTTTCACTGCATTCT
		1600 1610 1620	AGTTGTGGTTTGTCCAACTCATCAATGTA
G0251720-1.seq(1>4716)	→		AGTTGTGGTTTGTCCAACTCATCAATGTA

		1630 1640 165	TCTTAACGCGTAAATTGTAAGCGTTAATAT
G0251720-1.seq(1>4716)	→		TCTTAACGCGTAAATTGTAAGCGTTAATAT
		1660 1670 168	TTTGTTAAAATTCGCGTTAAATTTTGTTA
G0251720-1.seq(1>4716)	→		TTTGTTAAAATTCGCGTTAAATTTTGTTA
		1690 1700 171	AATCAGCTCATTTTTTAACCAATAGGCCGA
G0251720-1.seq(1>4716)	→		AATCAGCTCATTTTTTAACCAATAGGCCGA
		1720 1730 174	AATCGGCAAATCCCTTATAAATCAAAGA
G0251720-1.seq(1>4716)	→		AATCGGCAAATCCCTTATAAATCAAAGA
		1750 1760 177	ATAGACCGAGATAGGGTTGAGTGTTGTTCC
G0251720-1.seq(1>4716)	→		ATAGACCGAGATAGGGTTGAGTGTTGTTCC
		1780 1790 180	AGTTTGGAACAAGAGTCCACTATTAAAGAA
G0251720-1.seq(1>4716)	→		AGTTTGGAACAAGAGTCCACTATTAAAGAA
		1810 1820 183	CGTGGACTCCAACGTCAAAGGGCGAAAAC
G0251720-1.seq(1>4716)	→		CGTGGACTCCAACGTCAAAGGGCGAAAAC
		1840 1850 186	CGTCTATCAGGGCGATGGCCCACTACGTGA
G0251720-1.seq(1>4716)	→		CGTCTATCAGGGCGATGGCCCACTACGTGA
		1870 1880 189	ACCATCACCCTAATCAAGTTTTTTGGGGTC
G0251720-1.seq(1>4716)	→		ACCATCACCCTAATCAAGTTTTTTGGGGTC
		1900 1910 192	GAGGTGCCGTAAAGCACTAAATCGGAACCC
G0251720-1.seq(1>4716)	→		GAGGTGCCGTAAAGCACTAAATCGGAACCC
		1930 1940 195	TAAAGGGAGCCCCGATTTAGAGCTTGACC
G0251720-1.seq(1>4716)	→		TAAAGGGAGCCCCGATTTAGAGCTTGACC
		1960 1970 198	GGGAAAGCCGGCGAACGTGGCGAGAAAGGA
G0251720-1.seq(1>4716)	→		GGGAAAGCCGGCGAACGTGGCGAGAAAGGA

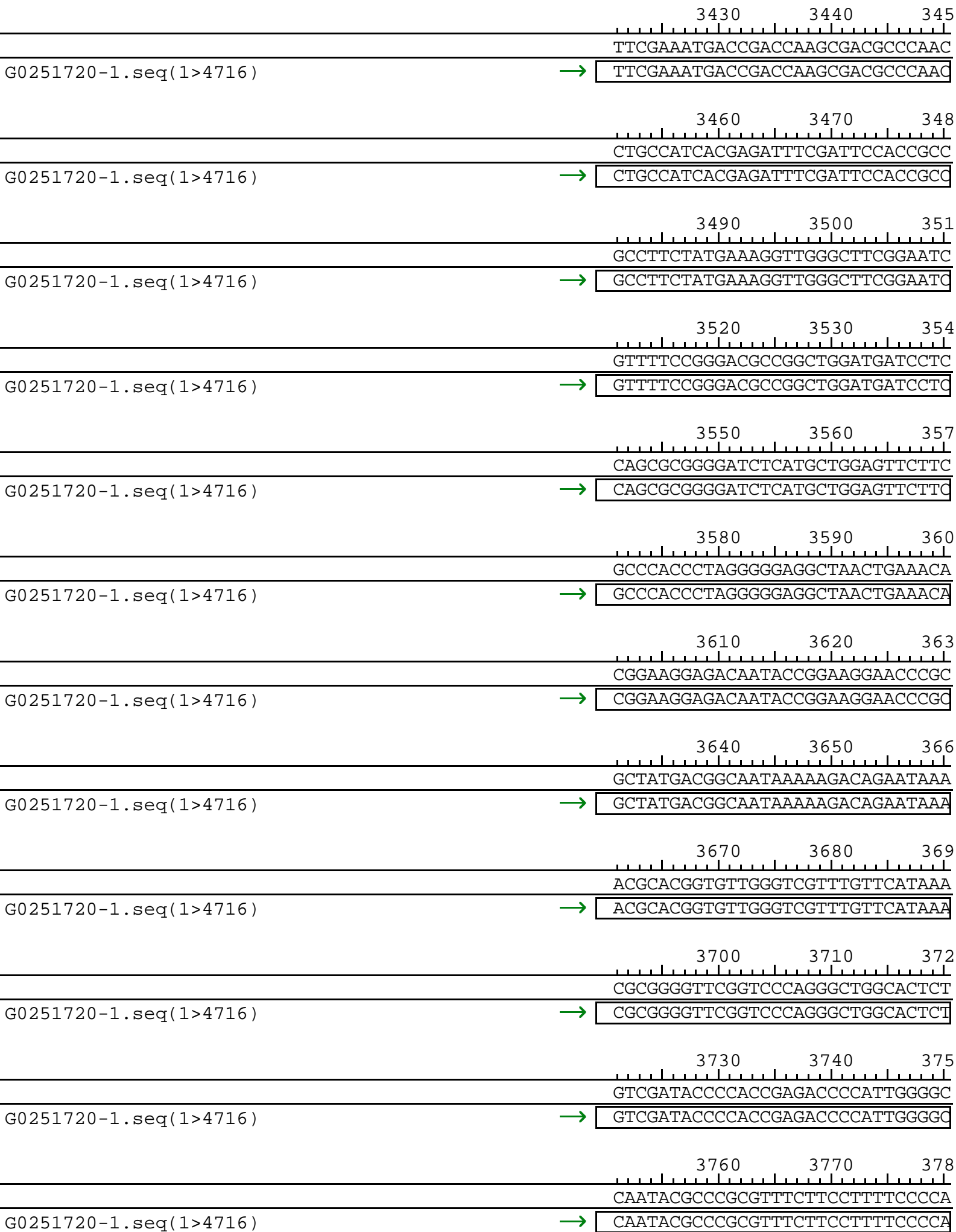
		1990 2000 201
		AGGGAAGAAAGCGAAAGGAGCGGGCGCTAG
G0251720-1.seq(1>4716)	→	AGGGAAGAAAGCGAAAGGAGCGGGCGCTAG
		2020 2030 204
		GGCGCTGGCAAGTGTAGCGGTCACGCTGCC
G0251720-1.seq(1>4716)	→	GGCGCTGGCAAGTGTAGCGGTCACGCTGCC
		2050 2060 207
		CGTAACCACCACACCCGCCGCGCTTAATGC
G0251720-1.seq(1>4716)	→	CGTAACCACCACACCCGCCGCGCTTAATGC
		2080 2090 210
		GCCGCTACAGGGCGCGTCAGGTGGCACTTT
G0251720-1.seq(1>4716)	→	GCCGCTACAGGGCGCGTCAGGTGGCACTTT
		2110 2120 213
		TCGGGGAAATGTGCGCGGAACCCCTATTTG
G0251720-1.seq(1>4716)	→	TCGGGGAAATGTGCGCGGAACCCCTATTTG
		2140 2150 216
		TTTATTTTTCTAAATACATTCAAATATGTA
G0251720-1.seq(1>4716)	→	TTTATTTTTCTAAATACATTCAAATATGTA
		2170 2180 219
		TCCGCTCATGAGACAATAACCCTGATAAAT
G0251720-1.seq(1>4716)	→	TCCGCTCATGAGACAATAACCCTGATAAAT
		2200 2210 222
		GCTTCAATAATATTGAAAAGGAAGAGTCC
G0251720-1.seq(1>4716)	→	GCTTCAATAATATTGAAAAGGAAGAGTCC
		2230 2240 225
		TGAGGCGGAAAGAACCAGCTGTGGAATGTG
G0251720-1.seq(1>4716)	→	TGAGGCGGAAAGAACCAGCTGTGGAATGTG
		2260 2270 228
		TGTCAGTTAGGGTGTGGAAAGTCCCCAGGC
G0251720-1.seq(1>4716)	→	TGTCAGTTAGGGTGTGGAAAGTCCCCAGGC
		2290 2300 231
		TCCCAGCAGGCAGAAGTATGCAAAGCATG
G0251720-1.seq(1>4716)	→	TCCCAGCAGGCAGAAGTATGCAAAGCATG
		2320 2330 234
		CATCTCAATTAGTCAGCAACCAGGTGTGGA
G0251720-1.seq(1>4716)	→	CATCTCAATTAGTCAGCAACCAGGTGTGGA

		2350	2360	237
		AAGTCCCCAGGCTCCCCAGCAGGCAGAAGT		
G0251720-1.seq(1>4716)	→	AAGTCCCCAGGCTCCCCAGCAGGCAGAAGT		
		2380	2390	240
		ATGCAAAGCATGCATCTCAATTAGTCAGCA		
G0251720-1.seq(1>4716)	→	ATGCAAAGCATGCATCTCAATTAGTCAGCA		
		2410	2420	243
		ACCATAGTCCCGCCCCTAACTCCGCCATC		
G0251720-1.seq(1>4716)	→	ACCATAGTCCCGCCCCTAACTCCGCCATC		
		2440	2450	246
		CCGCCCTAACTCCGCCAGTTCCGCCAT		
G0251720-1.seq(1>4716)	→	CCGCCCTAACTCCGCCAGTTCCGCCAT		
		2470	2480	249
		TCTCCGCCCATGGCTGACTAATTTTTTTT		
G0251720-1.seq(1>4716)	→	TCTCCGCCCATGGCTGACTAATTTTTTTT		
		2500	2510	252
		ATTTATGCAGAGGCCGAGGCCGCTCGGCC		
G0251720-1.seq(1>4716)	→	ATTTATGCAGAGGCCGAGGCCGCTCGGCC		
		2530	2540	255
		TCTGAGCTATTCCAGAAGTAGTGAGGAGGC		
G0251720-1.seq(1>4716)	→	TCTGAGCTATTCCAGAAGTAGTGAGGAGGC		
		2560	2570	258
		TTTTTTGGAGGCCTAGGCTTTTGCAAAGAT		
G0251720-1.seq(1>4716)	→	TTTTTTGGAGGCCTAGGCTTTTGCAAAGAT		
		2590	2600	261
		CGATCAAGAGACAGGATGAGGATCGTTTCG		
G0251720-1.seq(1>4716)	→	CGATCAAGAGACAGGATGAGGATCGTTTCG		
		2620	2630	264
		CATGATTGAACAAGATGGATTGCACGCAGG		
G0251720-1.seq(1>4716)	→	CATGATTGAACAAGATGGATTGCACGCAGG		
		2650	2660	267
		TTCTCCGGCCGCTTGGGTGGAGAGGCTATT		
G0251720-1.seq(1>4716)	→	TTCTCCGGCCGCTTGGGTGGAGAGGCTATT		
		2680	2690	270
		CGGCTATGACTGGGCACAACAGACAATCGG		
G0251720-1.seq(1>4716)	→	CGGCTATGACTGGGCACAACAGACAATCGG		

		2710 2720 273	CTGCTCTGATGCCGCCGTGTTCCGGCTGTC
G0251720-1.seq(1>4716)	→		CTGCTCTGATGCCGCCGTGTTCCGGCTGTC
		2740 2750 276	AGCGCAGGGGCGCCCGGTTCTTTTGTCAA
G0251720-1.seq(1>4716)	→		AGCGCAGGGGCGCCCGGTTCTTTTGTCAA
		2770 2780 279	GACCGACCTGTCCGGTGCCCTGAATGAACT
G0251720-1.seq(1>4716)	→		GACCGACCTGTCCGGTGCCCTGAATGAACT
		2800 2810 282	GCAAGACGAGGCAGCGCGGCTATCGTGGCT
G0251720-1.seq(1>4716)	→		GCAAGACGAGGCAGCGCGGCTATCGTGGCT
		2830 2840 285	GGCCACGACGGGCGTTCCTTGCGCAGCTGT
G0251720-1.seq(1>4716)	→		GGCCACGACGGGCGTTCCTTGCGCAGCTGT
		2860 2870 288	GCTCGACGTTGTCACTGAAGCGGGAAGGGA
G0251720-1.seq(1>4716)	→		GCTCGACGTTGTCACTGAAGCGGGAAGGGA
		2890 2900 291	CTGGCTGCTATTGGGCGAAGTGCCGGGGCA
G0251720-1.seq(1>4716)	→		CTGGCTGCTATTGGGCGAAGTGCCGGGGCA
		2920 2930 294	GGATCTCCTGTCATCTCACCTTGCTCCTGC
G0251720-1.seq(1>4716)	→		GGATCTCCTGTCATCTCACCTTGCTCCTGC
		2950 2960 297	CGAGAAAGTATCCATCATGGCTGATGCAAT
G0251720-1.seq(1>4716)	→		CGAGAAAGTATCCATCATGGCTGATGCAAT
		2980 2990 300	GCGGCGGCTGCATACGCTTGATCCGGCTAC
G0251720-1.seq(1>4716)	→		GCGGCGGCTGCATACGCTTGATCCGGCTAC
		3010 3020 303	CTGCCCATTCGACCACCAAGCGAAACATCG
G0251720-1.seq(1>4716)	→		CTGCCCATTCGACCACCAAGCGAAACATCG
		3040 3050 306	CATCGAGCGAGCACGTA CTGGATGGAAGC
G0251720-1.seq(1>4716)	→		CATCGAGCGAGCACGTA CTGGATGGAAGC

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		3070 3080 309	CGGTCTTGTCGATCAGGATGATCTGGACGA
G0251720-1.seq(1>4716)	→		CGGTCTTGTCGATCAGGATGATCTGGACGA
		3100 3110 312	AGAGCATCAGGGGCTCGCGCCAGCCGA
G0251720-1.seq(1>4716)	→		AGAGCATCAGGGGCTCGCGCCAGCCGA
		3130 3140 315	GTTCGCCAGGCTCAAGGCGAGCATGCCCGA
G0251720-1.seq(1>4716)	→		GTTCGCCAGGCTCAAGGCGAGCATGCCCGA
		3160 3170 318	CGGCGAGGATCTCGTCGTGACCCATGGCGA
G0251720-1.seq(1>4716)	→		CGGCGAGGATCTCGTCGTGACCCATGGCGA
		3190 3200 321	TGCCTGCTTGCCGAATATCATGGTGGAAAA
G0251720-1.seq(1>4716)	→		TGCCTGCTTGCCGAATATCATGGTGGAAAA
		3220 3230 324	TGGCCGCTTTTCTGGATTCATCGACTGTGG
G0251720-1.seq(1>4716)	→		TGGCCGCTTTTCTGGATTCATCGACTGTGG
		3250 3260 327	CCGGCTGGGTGTGGCGGACCGCTATCAGGA
G0251720-1.seq(1>4716)	→		CCGGCTGGGTGTGGCGGACCGCTATCAGGA
		3280 3290 330	CATAGCGTTGGCTACCCGTGATATTGCTGA
G0251720-1.seq(1>4716)	→		CATAGCGTTGGCTACCCGTGATATTGCTGA
		3310 3320 333	AGAGCTTGCGGGCGAATGGGCTGACCGCTT
G0251720-1.seq(1>4716)	→		AGAGCTTGCGGGCGAATGGGCTGACCGCTT
		3340 3350 336	CCTCGTGCTTTACGGTATCGCCGCTCCCGA
G0251720-1.seq(1>4716)	→		CCTCGTGCTTTACGGTATCGCCGCTCCCGA
		3370 3380 339	TTCGCAGCGCATCGCCTTCTATCGCCTTCT
G0251720-1.seq(1>4716)	→		TTCGCAGCGCATCGCCTTCTATCGCCTTCT
		3400 3410 342	TGACGAGTTCTTCTGAGCGGGACTCTGGGG
G0251720-1.seq(1>4716)	→		TGACGAGTTCTTCTGAGCGGGACTCTGGGG



		3790	3800	381
		CCCCACCCCCCAAGTTCGGGTGAAGGCCCA		
G0251720-1.seq(1>4716)	→	CCCCACCCCCCAAGTTCGGGTGAAGGCCCA		
		3820	3830	384
		GGGCTCGCAGCCAACGTCGGGGCGGCAGGC		
G0251720-1.seq(1>4716)	→	GGGCTCGCAGCCAACGTCGGGGCGGCAGGC		
		3850	3860	387
		CCTGCCATAGCCTCAGGTTACTCATATATA		
G0251720-1.seq(1>4716)	→	CCTGCCATAGCCTCAGGTTACTCATATATA		
		3880	3890	390
		CTTTAGATTGATTTAAACTTCATTTTTAA		
G0251720-1.seq(1>4716)	→	CTTTAGATTGATTTAAACTTCATTTTTAA		
		3910	3920	393
		TTTAAAAGGATCTAGGTGAAGATCCTTTTT		
G0251720-1.seq(1>4716)	→	TTTAAAAGGATCTAGGTGAAGATCCTTTTT		
		3940	3950	396
		GATAATCTCATGACCAAATCCCTTAACGT		
G0251720-1.seq(1>4716)	→	GATAATCTCATGACCAAATCCCTTAACGT		
		3970	3980	399
		GAGTTTTTCGTTCCACTGAGCGTCAGACCCC		
G0251720-1.seq(1>4716)	→	GAGTTTTTCGTTCCACTGAGCGTCAGACCCC		
		4000	4010	402
		GTAGAAAAGATCAAAGGATCTTCTTGAGAT		
G0251720-1.seq(1>4716)	→	GTAGAAAAGATCAAAGGATCTTCTTGAGAT		
		4030	4040	405
		CCTTTTTTTCTGCGCGTAATCTGCTGCTTG		
G0251720-1.seq(1>4716)	→	CCTTTTTTTCTGCGCGTAATCTGCTGCTTG		
		4060	4070	408
		CAAACAAAAAACCACCGCTACCAGCGGTG		
G0251720-1.seq(1>4716)	→	CAAACAAAAAACCACCGCTACCAGCGGTG		
		4090	4100	411
		GTTTGTTTGCCGGATCAAGAGCTACCAACT		
G0251720-1.seq(1>4716)	→	GTTTGTTTGCCGGATCAAGAGCTACCAACT		
		4120	4130	414
		CTTTTTCCGAAGGTAAGTGGCTTCAGCAGA		
G0251720-1.seq(1>4716)	→	CTTTTTCCGAAGGTAAGTGGCTTCAGCAGA		

		4150	4160	417	GCGCAGATACCAAATACTGTTCTTCTAGTG
G0251720-1.seq(1>4716)	→				GCGCAGATACCAAATACTGTTCTTCTAGTG
		4180	4190	420	TAGCCGTAGTTAGGCCACCACCTTCAAGAAC
G0251720-1.seq(1>4716)	→				TAGCCGTAGTTAGGCCACCACCTTCAAGAAC
		4210	4220	423	TCTGTAGCACCGCCTACATACCTCGCTCTG
G0251720-1.seq(1>4716)	→				TCTGTAGCACCGCCTACATACCTCGCTCTG
		4240	4250	426	CTAATCCTGTTACCAGTGGCTGCTGCCAGT
G0251720-1.seq(1>4716)	→				CTAATCCTGTTACCAGTGGCTGCTGCCAGT
		4270	4280	429	GGCGATAAGTCGTGTCTTACCGGGTTGGAC
G0251720-1.seq(1>4716)	→				GGCGATAAGTCGTGTCTTACCGGGTTGGAC
		4300	4310	432	TCAAGACGATAGTTACCGGATAAGGCGCAG
G0251720-1.seq(1>4716)	→				TCAAGACGATAGTTACCGGATAAGGCGCAG
		4330	4340	435	CGGTCGGGCTGAACGGGGGGTTTCGTGCACA
G0251720-1.seq(1>4716)	→				CGGTCGGGCTGAACGGGGGGTTTCGTGCACA
		4360	4370	438	CAGCCCAGCTTGGAGCGAACGACCTACACC
G0251720-1.seq(1>4716)	→				CAGCCCAGCTTGGAGCGAACGACCTACACC
		4390	4400	441	GAACTGAGATACCTACAGCGTGAGCTATGA
G0251720-1.seq(1>4716)	→				GAACTGAGATACCTACAGCGTGAGCTATGA
		4420	4430	444	GAAAGCGCCACGCTTCCCGAAGGGAGAAAG
G0251720-1.seq(1>4716)	→				GAAAGCGCCACGCTTCCCGAAGGGAGAAAG
		4450	4460	447	GCGGACAGGTATCCGGTAAGCGGCAGGGTC
G0251720-1.seq(1>4716)	→				GCGGACAGGTATCCGGTAAGCGGCAGGGTC
		4480	4490	450	GGAACAGGAGAGCGCACGAGGGAGCTTCCA
G0251720-1.seq(1>4716)	→				GGAACAGGAGAGCGCACGAGGGAGCTTCCA

4510 4520 453
GGGGGAAACGCCTGGTATCTTTATAGTCCT
G0251720-1.seq(1>4716) → GGGGGAAACGCCTGGTATCTTTATAGTCCT

4540 4550 456
GTCGGGTTTCGCCACCTCTGACTTGAGCGT
G0251720-1.seq(1>4716) → GTCGGGTTTCGCCACCTCTGACTTGAGCGT

4570 4580 459
CGATTTTTGTGATGCTCGTCAGGGGGCGG
G0251720-1.seq(1>4716) → CGATTTTTGTGATGCTCGTCAGGGGGCGG

4600 4610 462
AGCCTATGGAAAAACGCCAGCAACGCGGCC
G0251720-1.seq(1>4716) → AGCCTATGGAAAAACGCCAGCAACGCGGCC

4630 4640 465
TTTTTACGGTTCCCTGGCCTTTTGCTGGCCT
G0251720-1.seq(1>4716) → TTTTTACGGTTCCCTGGCCTTTTGCTGGCCT

4660 4670 468
TTTGCTCACATGTTCTTTCCCTGCGTTATCC
G0251720-1.seq(1>4716) → TTTGCTCACATGTTCTTTCCCTGCGTTATCC

4690 4700 471
CCTGATTCTGTGGATAACCGTATTACCGCC
G0251720-1.seq(1>4716) → CCTGATTCTGTGGATAACCGTATTACCGCC

ATGCAT
G0251720-1.seq(1>4716) → ATGCAT