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	500 510 520 530	CGGCAACCTCTCGGGCTTCTGGATTTCGGATCCCGGAATT
G0210463-1.seq(1>10379)	→	cggcaacctctcgggcttctggatttccgatccccggaatt
	540 550 560 570	AGATCTTGGCAGGATATATTGTGGTGTAACGTTCTCGCATG
G0210463-1.seq(1>10379)	→	agatcttggcaggatataattgtggtgtaacgttctcgcatg
	580 590 600 610	CCTGCAGGTCAACATGGTGGAGCACGACACGCTTGTCTACT
G0210463-1.seq(1>10379)	→	cctgcaggtcaacatggtggagcacgacacgcttgtctact
	620 630 640 650	CCAAAATATCAAAGATACAGTCTCAGAAGACCAAAGGGCA
G0210463-1.seq(1>10379)	→	ccaaaatatcaaagatacagtctcagaagaccaaggcca
	660 670 680 690	ATTGAGACTTTTCAACAAAGGGTAATATCCGGAAACCTCCT
G0210463-1.seq(1>10379)	→	attgagacttttcaacaaagggtaatatccggaaacctcct
	700 710 720 730	CGGATTCCATTGCCAGCTATCTGTCACTTTATTGTGAAGA
G0210463-1.seq(1>10379)	→	cggattccattgcccagctatctgtcactttattgtgaaga
	740 750 760 770	TAGTGGAAAAGGAAGGTGGCTCCTACAAATGCCATCATTGC
G0210463-1.seq(1>10379)	→	tagtggaaaaggaaggtggctcctacaaatgccatcattgc
	780 790 800 810 820	GATAAAGGAAAGGCCATCGTTGAAGATGCCTCTGCCGACAG
G0210463-1.seq(1>10379)	→	gataaaggaaaggccatcgttgaagatgcctctgccgacag
	830 840 850 860	TGGTCCCAAAGATGGACCCCCACCCACGAGGAGCATCGTGG
G0210463-1.seq(1>10379)	→	tggtcctcaaagatggacccccaccacgaggagcatcgtgg
	870 880 890 900	AAAAGAAGACGTTCCAACCACGTCTTCAAAGCAAGTGGAT
G0210463-1.seq(1>10379)	→	aaaagaagacgttccaaccacgtcttcaaagcaagtggat
	910 920 930 940	TGATGTGATATCTCCACTGACGTAAGGGATGACGCACAATC
G0210463-1.seq(1>10379)	→	tgatgtgatatctccactgacgtaagggatgacgcacaatc
	950 960 970 980	CCACTATCCTTCGCAAGACCCTTCCTCTATATAAGGAAGTT
G0210463-1.seq(1>10379)	→	ccactatccttcgcaagacccttcctctatataaggaagtt

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990 1000 1010 1020
CATTTCATTGGAGAGGAGAAAATAAACCATACACCACCA
G0210463-1.seq(1>10379) → catttcatttggagagggagaaaactaaaccatacaccacca

1030 1040 1050 1060
ACACAACCAAACCCACCACGCCCAATTGTTACACACCCGCT
G0210463-1.seq(1>10379) → acacaaccaaaccaccacgccaattgttacacacccgct

1070 1080 1090 1100
TGAAAAGAAAGTTTAAACAAATGGCCAAGGTGCGCGAGGTT
G0210463-1.seq(1>10379) → tgaaaagaaagtttaacaaatggccaaggtgcgcgaggtt

1110 1120 1130 1140
TACCAATCTTTTACAGACTCCACCACAAAACCTCTCATCCA
G0210463-1.seq(1>10379) → taccaatcttttacagactccaccacaaaaactctcatcca

1150 1160 1170 1180
AGATGAGGCTTATAGAAACATTCGCCCATCATGGAAAAC
G0210463-1.seq(1>10379) → agatgaggcttatagaaacattcgcccatcatggaaaaac

1190 1200 1210 1220 1230
ACAACTAGCTAACCCCTTACGCTCAAACGGTTGAAGCGGCT
G0210463-1.seq(1>10379) → acaactagctaaccccttacgctcaaacggttgaagcggct

1240 1250 1260 1270
AATGATCTAGAGGGGTTTCGGCATAGCCACCAATCCCTATAG
G0210463-1.seq(1>10379) → aatgatctagagggggttcggcatagccaccaatccctatag

1280 1290 1300 1310
CATTGAATTGCATACACATGCAGCCGCTAAGACCATAGAGA
G0210463-1.seq(1>10379) → cattgaattgcatacacatgcagccgctaagacccatagaga

1320 1330 1340 1350
ATAAACTTCTAGAGGTGCTTGGTTCCATCTACCACAAGAA
G0210463-1.seq(1>10379) → ataaacttctagaggtgcttggttccatctaccacaagaa

1360 1370 1380 1390
CCTGTTACATTTATGTTTCTTAAACCCAGAAAGCTAAACTA
G0210463-1.seq(1>10379) → cctgttacatTTATGTTTCTTAAACCCAGAAAGCTAAACTA

1400 1410 1420 1430
CATGAGAAGAAACCCGCGGATCAAGGACATTTTCCAAAATG
G0210463-1.seq(1>10379) → catgagaagaacccgcggatcaaggacatTTTCCAAAATG

1440 1450 1460 1470
TTGCCATTGAACCAAGAGACGTAGCCAGGTACCCCAAGGAA
G0210463-1.seq(1>10379) → ttgccattgaaccaagagacgtagccaggtaccccaaggaa

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	1480 1490 1500 1510	ACAATAATTGACAAACTCACAGAGATCACAACGGAAACAGC
G0210463-1.seq(1>10379)	→	acaataattgacaaactcacagagatcacaacggaaacagc
	1520 1530 1540 1550	ATACATTAGTGACACTCTGCACTTCTTGGATCCGAGCTACA
G0210463-1.seq(1>10379)	→	atacattagtgacactctgcacttcttggatccgagctaca
	1560 1570 1580 1590	TAGTGGAGACATTCCAAAAGTCCCAAATGCAAACATTG
G0210463-1.seq(1>10379)	→	tagtggagacattccaaaactgccccaaaattgcaaacattg
	1600 1610 1620 1630 1640	TATGCGACCTTAGTTCTCCCCGTTGAGGCAGCCTTTAAAAT
G0210463-1.seq(1>10379)	→	tatgcgaccttagttctccccgttgaggcagcctttaaaat
	1650 1660 1670 1680	GGAAAGCACTCACCCGAACATATACAGCCTCAAATACTTCG
G0210463-1.seq(1>10379)	→	ggaaagcactcaccggaacatatacagcctcaaatacttcg
	1690 1700 1710 1720	GAGATGGTTTCCAGTATATACCAGGCAACCATGGTGGCGGG
G0210463-1.seq(1>10379)	→	gagatggtttccagtatataccaggcaaccatggtggcggg
	1730 1740 1750 1760	GCATACCATCATGAATTCGCTCATCTACAATGGCTCAAAGT
G0210463-1.seq(1>10379)	→	gcataccatcatgaattcgctcatctacaatggctcaaagt
	1770 1780 1790 1800	GGGAAAGATCAAGTGGAGGGACCCCAAGGATAGCTTTCTCG
G0210463-1.seq(1>10379)	→	gggaaagatcaagtggagggaccccaaggatagctttctcg
	1810 1820 1830 1840	GACATCTCAATTACACGACTGAGCAGGTTGAGATGCACACA
G0210463-1.seq(1>10379)	→	gacatctcaattacacgactgagcaggttgagatgcacaca
	1850 1860 1870 1880	GTGACAGTACAGTTGCAGGAATCGTTTCGCGGCAAACCACTT
G0210463-1.seq(1>10379)	→	gtgacagtacagttgcaggaatcgtttcgcggcaaaccactt
	1890 1900 1910 1920	GTACTGCATCAGGAGAGGAGACTTGCTCACACCGGAGGTGC
G0210463-1.seq(1>10379)	→	gtactgcatcaggagaggagacttgctcacaccggaggtgc
	1930 1940 1950 1960	GCACTTTCGGCCAACCTGACAGGTACGTGATTCCACCACAG
G0210463-1.seq(1>10379)	→	gcactttcggccaacctgacaggtacgtgattccaccacag

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	1700	1800	1900	2000	ATCTTCCTCCCAAAGTTCACAACCTGCAAGAAGCCGATTCT	
G0210463-1.seq(1>10379)	→	atcttcctcccaaagttcacaactgcaagaagccgattct				
	2010	2020	2030	2040	2050	CAAGAAAACCTATGATGCAGCTCTTCTTGTATGTTAGGACAG
G0210463-1.seq(1>10379)	→	caagaaaactatgatgcagctcttcttgtatgtaggacag				
	2060	2070	2080	2090	TCAAGGTCGAAAAAATTGTGACATTTTGCCAAAGTCAGA	
G0210463-1.seq(1>10379)	→	tcaaggctcgaaaaaattgtgacatTTTTGCCAAAGTCAGA				
	2100	2110	2120	2130	CAATTAATTAATCATCTGACTTGGACAAATACTCTGCTGT	
G0210463-1.seq(1>10379)	→	caattaattaatcatctgacttggacaaataactctgctgt				
	2140	2150	2160	2170	GGAAGTGGTTTACTTAGTAAGCTACATGGAGTTCCTTGCCG	
G0210463-1.seq(1>10379)	→	ggaactggtttacttagtaagctacatggagttccttgccg				
	2180	2190	2200	2210	ATTTACAAGCTACCACCTGCTTCTCAGACACACTTTCTGGT	
G0210463-1.seq(1>10379)	→	atTTACAAGCTACCACCTGCTTCTCAGACACACTTTCTGGT				
	2220	2230	2240	2250	GGCTTGCTAACAAAGACCCTTGCACCGGTGAGGGCTTGGAT	
G0210463-1.seq(1>10379)	→	ggcttgctaacAAAGACCCTTGCACCGGTGAGGGCTTGGAT				
	2260	2270	2280	2290	ACAAGAGAAAAAGATGCAGCTGTTTGGTCTTGAGGACTACG	
G0210463-1.seq(1>10379)	→	acaagagaaaaagatgcagctgTTTGGTCTTGAGGACTACG				
	2300	2310	2320	2330	CGAAGTTAGTCAAAGCAGTTGATTTCCACCCGGTGGATTTT	
G0210463-1.seq(1>10379)	→	cgaagttagtCAAAGCAGTTGATTTCCACCCGGTGGATTTT				
	2340	2350	2360	2370	TCTTTCAAAGTGGAAACTTGGGACTTCAGATTCCACCCCTT	
G0210463-1.seq(1>10379)	→	tctttCAAAGTGGAAACTTGGGACTTCAGATTCCACCCCTT				
	2380	2390	2400	2410	GCAAGCGTGGAAAGCCTTCCGACCAAGGGAAGTGTCGGATG	
G0210463-1.seq(1>10379)	→	gcaagcgtGGAAAGCCTTCCGACCAAGGGAAGTGTCGGATG				
	2420	2430	2440	2450	2460	TAGAGGAAATGGAAAGTTTGTCTCAGATGGGGACCTGCTT
G0210463-1.seq(1>10379)	→	tagaggAAATGGAAAGTTTGTCTCAGATGGGGACCTGCTT				

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		2470	2480	2490	2500	
		GATTGCTTCACAAGAATGCCAGCTTATGCGGTAAACGCAGA				
G0210463-1.seq(1>10379)	→	gattgcttcacaagaatgccagcttatgcggtaaacgcaga				
		2510	2520	2530	2540	
		GGAAGATTTAGCTGCAATCAGGAAAACGCCCGAGATGGATG				
G0210463-1.seq(1>10379)	→	ggaagatttagctgcaatcaggaaaacgcccgagatggatg				
		2550	2560	2570	2580	
		TCGGTCAAGAAGTTAAAGAGCCTGCAGGAGACAGAAATCAA				
G0210463-1.seq(1>10379)	→	tcggtcaagaagttaagagcctgcaggagacagaaatcaa				
		2590	2600	2610	2620	
		TACTCAAACCCTGCAGAACTTTCCTCAACAAGCTCCACAG				
G0210463-1.seq(1>10379)	→	tactcaaaccctgcagaaactttcctcaacaagctccacag				
		2630	2640	2650	2660	
		GAAACACAGTAGGGAGGTGAAACACCAGGCCGCAAAGAAAG				
G0210463-1.seq(1>10379)	→	gaaacacagtagggaggtgaaacaccaggccgcaaagaaag				
		2670	2680	2690	2700	
		CTAAACGCCTAGCTGAAATCCAGGAGTCAATGAGAGCTGAA				
G0210463-1.seq(1>10379)	→	ctaaacgcctagctgaaatccaggagtcaatgagagctgaa				
		2710	2720	2730	2740	
		GGTGATGCCGAACCAAATGAAATAAGCGGGACGATGGGGGC				
G0210463-1.seq(1>10379)	→	ggtgatgccgaaccaaataagcgggacgatgggggc				
		2750	2760	2770	2780	
		AATACCAGCAACGCCGAACTTCCTGGCACGAATGATGCCA				
G0210463-1.seq(1>10379)	→	aataccagcaacgccgaacttctctggcacgaatgatgcc				
		2790	2800	2810	2820	
		GACAAGAACTCACACTCCAACCACTAAACCTGTCCCTGCA				
G0210463-1.seq(1>10379)	→	gacaagaactcacactccaaccactaaacctgtccctgca				
		2830	2840	2850	2860	2870
		AGGTGGGAAGATGCTTCATTCACAGATTCTAGTGTGGAAGA				
G0210463-1.seq(1>10379)	→	aggtgggaagatgcttcattcacagattctagtgtggaaga				
		2880	2890	2900	2910	
		GGAGCAGGTTAAACTCCTTGAAAAGAAACCGTTGAAACAG				
G0210463-1.seq(1>10379)	→	ggagcaggttaaactccttgaaaagaacccgttgaaacag				
		2920	2930	2940	2950	
		CGACGCAACAAGTCATCGAAGGACTTCCTTGAAACACTGG				
G0210463-1.seq(1>10379)	→	cgacgcaacaagtcacgaaggacttccttgaaacactgg				

		2960	2970	2980	2990	
		ATTCTCAATTA	ATGCTGTTGGATT	CAAGGCGCTGGAAAT		
G0210463-1.seq(1>10379)	→	attcctcaattaaatgctgttggattcaaggcgctggaaat				
		3000	3010	3020	3030	
		TCAGAGGGATAGGAGT	GGAACAATGATCAT	GCCCATCACAG		
G0210463-1.seq(1>10379)	→	tcagagggataggagtggaacaatgatcatgccatcacag				
		3040	3050	3060	3070	
		AAATGGTCTCCGGGCT	GGAAAAGAGGACTT	CCCTGAAGGA		
G0210463-1.seq(1>10379)	→	aaatggtctccgggctggaaaagaggacttccctgaagga				
		3080	3090	3100	3110	
		ACTCCAAAAGAGTT	TGGCACGAGAATTG	TTCGCTATGAACAG		
G0210463-1.seq(1>10379)	→	actccaaaagagttggcacgagaattgttcgctatgaacag				
		3120	3130	3140	3150	
		AAGCCCTGCCACCAT	CCCTTTGGACCTG	CTTAGAGCCAGAG		
G0210463-1.seq(1>10379)	→	aagccctgccaccatccctttggacctgcttagagccagag				
		3160	3170	3180	3190	
		ACTACGGCAGTGAT	GTAAGAACAAGAGA	AATTGGTGCCATC		
G0210463-1.seq(1>10379)	→	actacggcagtgatgtaagaacaagagaattggtgccatc				
		3200	3210	3220	3230	
		ACAAAGACACAGG	CAACGAGTTGGGG	CGAATACTTGACAGG		
G0210463-1.seq(1>10379)	→	acaagacacaggcaacgagttggggcgaatacttgacagg				
		3240	3250	3260	3270	3280
		AAAGATAGAAAGCT	TAACTGAGAGGAA	AGTTGCGACTTGTG		
G0210463-1.seq(1>10379)	→	aaagatagaaagcttaactgagaggaaagttgcgacttgtg				
		3290	3300	3310	3320	
		TCATTCATGGAGCT	TGGAGTTCTGGAAA	AGTCATGCCATC		
G0210463-1.seq(1>10379)	→	tcattcatggagctggagttctggaaaaagtcatgccatc				
		3330	3340	3350	3360	
		CAGAAGGCATTGAG	AGAAATTGGCAAGG	GCTCGGACATCAC		
G0210463-1.seq(1>10379)	→	cagaaggcattgagagaaattggcaagggctcggacatcac				
		3370	3380	3390	3400	
		TGTAGTCCTGCCG	ACCAATGAACTGCG	GCTAGATTGGAGTA		
G0210463-1.seq(1>10379)	→	tgtagtcctgccgaccaatgaactgcggttagattggagta				
		3410	3420	3430	3440	
		AGAAAGTGCCTAAC	ACTGAGCCCTATAT	GTTCAAGACCTCT		
G0210463-1.seq(1>10379)	→	agaaagtgcctaactgagccctatatgttcaagacctct				

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		3450	3460	3470	3480	
		GAAAAGGCGTTAATTGGGGGAACAGGCAGCATAGTCATCTT				
G0210463-1.seq(1>10379)	→	gaaaaggcgttaattgggggaacaggcagcatagtcatctt				
		3490	3500	3510	3520	
		TGACGATTACTCAAAACTTCCTCCCGGTTACATAGAAGCCT				
G0210463-1.seq(1>10379)	→	tgacgattactcaaaacttcctcccggttacatagaagcct				
		3530	3540	3550	3560	
		TAGTCTGTTTCTACTCTAAAATCAAGCTAATCATTCTAACA				
G0210463-1.seq(1>10379)	→	tagtctgtttctactctaaaatcaagctaatacttctaaca				
		3570	3580	3590	3600	
		GGAGATAGCAGACAAAGCGTCTACCATGAAACTGCTGAGGA				
G0210463-1.seq(1>10379)	→	ggagatagcagacaaagcgtctaccatgaaactgctgagga				
		3610	3620	3630	3640	
		CGCCTCCATCAGGCATTTGGGACCAGCAACAGAGTACTTCT				
G0210463-1.seq(1>10379)	→	cgcctccatcaggcatttgggaccagcaacagagtacttct				
		3650	3660	3670	3680	3690
		CAAATACTGCCGATACTATCTCAATGCCACACACCCGCAAC				
G0210463-1.seq(1>10379)	→	caaaatactgccgatactatctcaatgccacacaccgcaac				
		3700	3710	3720	3730	
		AAGAAAGATCTTGCGAACATGCTTGGTGTCTACAGTGAGAG				
G0210463-1.seq(1>10379)	→	aagaaagatcttgcgaaacatgcttgggtgtctacagtgagag				
		3740	3750	3760	3770	
		AACGGGAGTCACCGAAATCAGCATGAGCGCCGAGTTCTTAG				
G0210463-1.seq(1>10379)	→	aacgggagtcaccgaaatcagcatgagcgccgagttcttag				
		3780	3790	3800	3810	
		AAGGAATCCCAACTTTGGTACCCTCGGATGAGAAGAGAAAG				
G0210463-1.seq(1>10379)	→	aaggaatcccaactttggtaccctcggatgagaagagaaaag				
		3820	3830	3840	3850	
		CTGTACATGGGCACCGGGAGGAATGACACGTTACATACGC				
G0210463-1.seq(1>10379)	→	ctgtacatgggcaccgggaggaatgacacgttcacatacgc				
		3860	3870	3880	3890	
		TGGATGCCAGGGGCTAACTAAGCCGAAGGTACAAATAGTGT				
G0210463-1.seq(1>10379)	→	tggatgccaggggctaactaagccgaaggtacaaatagtgt				
		3900	3910	3920	3930	
		TGGACCACAACACCCAAGTGTGTAGCGCGAATGTGATGTAC				
G0210463-1.seq(1>10379)	→	tggaccacaacacccaagtgtgtagcgcgaatgtgatgtac				

	3940 3950 3960 3970	ACGGCACTTTCTAGAGCCACCGATAGGATTCACTTCGTGAA
G0210463-1.seq(1>10379)	→	acggcacttttctagagccaccgataggattcacttcgtgaa
	3980 3990 4000 4010	CACAAGTGCAAATTCCTCTGCCTTCTGGGAAAAGTTGGACA
G0210463-1.seq(1>10379)	→	cacaagtgcaaattcctctgccttctgggaaaagttggaca
	4020 4030 4040 4050	GCACCCCTTACCTCAAGACTTTCCTATCAGTGGTGAGAGAA
G0210463-1.seq(1>10379)	→	gcacccttacctcaagactttcctatcagtggtgagagaa
	4060 4070 4080 4090 4100	CAAGCACTCAGGGAGTACGAGCCGGCAGAGGCAGAGCCAAT
G0210463-1.seq(1>10379)	→	caagcactcagggagtacgagccggcagagggcagagccaat
	4110 4120 4130 4140	TCAAGAGCCTGAGCCCCAGACACACATGTGTGTCGAGAATG
G0210463-1.seq(1>10379)	→	tcaagagcctgagccccagacacacatgtgtgtcgagaatg
	4150 4160 4170 4180	AGGAGTCCGTGCTAGAAGAGTACAAAGAGGAACTCTTGAA
G0210463-1.seq(1>10379)	→	aggagtccgtgctagaagagtacaagaggaactcttgaa
	4190 4200 4210 4220	AAGTTTGACAGAGAGATCCACTCTGAATCCCATGGTCATTC
G0210463-1.seq(1>10379)	→	aagtttgacagagagatccactctgaatcccatggtcattc
	4230 4240 4250 4260	AAACTGTGTCCAAACTGAAGACACAACCATTCAAGTTGTTTT
G0210463-1.seq(1>10379)	→	aaactgtgtcctaaactgaagacacaaccattcagttgTTTT
	4270 4280 4290 4300	CGCATCAACAAGCAAAAGATGAGACCCCTCCTCTGGGCGACT
G0210463-1.seq(1>10379)	→	cgcatacaagcaaaagatgagaccctcctctgggcgact
	4310 4320 4330 4340	ATAGATGCGCGGCTCAAGACCAGCAATCAAGAAACAAACTT
G0210463-1.seq(1>10379)	→	atagatgcgcggctcaagaccagcaatcaagaaacaaactt
	4350 4360 4370 4380	CCGAGAATTCTGAGCAAGAAGGACATTGGGGACGTTCTGT
G0210463-1.seq(1>10379)	→	ccgagaattcctgagcaagaaggacattggggacgttctgt
	4390 4400 4410 4420	TTTTAAACTACCAAAAAGCTATGGGTTTACCCAAAGAGCGT
G0210463-1.seq(1>10379)	→	ttttaaactaccaaaaagctatgggTTTACCCAAAGAGCGT

	4430	4440	4450	4460	
	ATTCTTTTTTCCCAAGAGGTCTGGGAAGCTTGTGCCACGA				
G0210463-1.seq(1>10379)	→	attcctttttcccaagaggtctgggaagcttgtgcccacga			
	4470	4480	4490	4500	4510
	AGTACAAAGCAAGTACCTCAGCAAGTCAAAGTGCAACTTGA				
G0210463-1.seq(1>10379)	→	agtacaaagcaagtacctcagcaagtcaaagtgcaacttga			
	4520	4530	4540	4550	
	TCAATGGGACTGTGAGACAGAGCCCAGACTTCGATGAAAAT				
G0210463-1.seq(1>10379)	→	tcaatgggactgtgagacagagcccagacttcgatgaaaat			
	4560	4570	4580	4590	
	AAGATTATGGTATTCTCAAGTCGCAGTGGGTCACAAAGGT				
G0210463-1.seq(1>10379)	→	aagattatggtattcctcaagtcgcagtgggtcaciaaaggt			
	4600	4610	4620	4630	
	GGAAAACTAGGTCTACCCAAGATTAAGCCAGGTCAAACCA				
G0210463-1.seq(1>10379)	→	ggaaaaactaggtctacccaagattaagccaggtcaaacca			
	4640	4650	4660	4670	
	TAGCAGCCTTTTACCAGCAGACTGTGATGCTTTTGGAACT				
G0210463-1.seq(1>10379)	→	tagcagccttttaccagcagactgtgatgctttttggaact			
	4680	4690	4700	4710	
	ATGGCTAGGTACATGCGATGGTTCAGACAGGCTTTCAGCC				
G0210463-1.seq(1>10379)	→	atggctaggtacatgcgatggttcagacaggctttccagcc			
	4720	4730	4740	4750	
	AAAAGAAGTCTTCATAAACTGTGAGACCACGCCAGATGACA				
G0210463-1.seq(1>10379)	→	aaaagaagtcttcataaactgtgagaccacgccagatgaca			
	4760	4770	4780	4790	
	TGTCTGCATGGCCTTGAACAACCTGGAATTTTCAGCAGACCT				
G0210463-1.seq(1>10379)	→	tgtctgcatggccttgaacaactggaatttcagcagacct			
	4800	4810	4820	4830	
	AGCTTGGCTAATGACTACACAGCTTTCGACCAGTCTCAGGA				
G0210463-1.seq(1>10379)	→	agcttggctaataactgactacacagctttcgaccagtctcagga			
	4840	4850	4860	4870	
	TGGAGCCATGTTGCAATTTGAGGTGCTCAAAGCCAAACACC				
G0210463-1.seq(1>10379)	→	tggagccatgttgcaatTTGAGGTGCTCAAAGCCAAACACC			
	4880	4890	4900	4910	4920
	ACTGCATACCAGAGGAAATCATTCAGGCATACATAGATATT				
G0210463-1.seq(1>10379)	→	actgcataaccagaggaaatcattcaggcatacatagatatt			

4930 4940 4950 4960
AAGACTAATGCACAGATTTTCCTAGGCACGTTATCAATTAT
G0210463-1.seq(1>10379) → aagactaatgcacagatTTTCCTAGGCACGTTATCAATTAT

4970 4980 4990 5000
GCGCCTGACTGGTGAAGGTCCCACCTTTTGATGCAAACACTG
G0210463-1.seq(1>10379) → gcgctgactggtgaaggtcccactTTTGATGCAAACACTG

5010 5020 5030 5040
AGTGCAACATAGCTTACACCCATACAAAGTTTGACATCCCA
G0210463-1.seq(1>10379) → agtgcaacatagctTACACCCATACAAAGTTTGACATCCCA

5050 5060 5070 5080
GCCGGAAGTCTCAAGTTTATGCAGGAGACGACTCCGCACT
G0210463-1.seq(1>10379) → gccggaactgctcaagTTTATGCAGGAGACGACTCCGCACT

5090 5100 5110 5120
GGACTGTGTTCCAGAAGTGAAGCATAGTTTCCACAGGCTTG
G0210463-1.seq(1>10379) → ggactgtgTTCCAGAAGTGAAGCATAGTTTCCACAGGCTTG

5130 5140 5150 5160
AGGACAAATTACTCCTAAAGTCAAAGCCTGTAATCACGCAG
G0210463-1.seq(1>10379) → aggacaaATTACTCCTAAAGTCAAAGCCTGTAATCACGCAG

5170 5180 5190 5200
CAAAGAAGGGCAGTTGGCCTGAGTTTGTGGTTGGCTGAT
G0210463-1.seq(1>10379) → caaagaAGGGCAGTTGGCCTGAGTTTGTGGTTGGCTGAT

5210 5220 5230 5240
CACACCAAAGGGGTGATGAAAGACCCAATTAAGCTCCATG
G0210463-1.seq(1>10379) → cacacCAAAGGGGTGATGAAAGACCCAATTAAGCTCCATG

5250 5260 5270 5280
TTAGCTTAAAATTGGCTGAAGCTAAGGGTGAAGTCAAGAAA
G0210463-1.seq(1>10379) → ttagctTAAAATTGGCTGAAGCTAAGGGTGAAGTCAAGAAA

5290 5300 5310 5320 5330
TGTCAAGATTCCTATGAAATTGATCTGAGTTATGCCTATGA
G0210463-1.seq(1>10379) → tgtcaagATTCCTATGAAATTGATCTGAGTTATGCCTATGA

5340 5350 5360 5370
CCACAAGGACTCTCTGCATGACTTGTTCGATGAGAAACAGT
G0210463-1.seq(1>10379) → ccacaAGGACTCTCTGCATGACTTGTTCGATGAGAAACAGT

5380 5390 5400 5410
GTCAGGCACACACTCACTTGCAGAACACTAATCAAGTCA
G0210463-1.seq(1>10379) → gtcaggCACACACTCACTTGCAGAACACTAATCAAGTCA

		5420 5430 5440 5450	GGGAGAGGCACTGTCTCACTTTCCCGCCTCAGAACTTTCT
G0210463-1.seq(1>10379)	→		gggagaggcactgtctcactttcccgctcagaaactttct
		5460 5470 5480 5490	TTAACCGTTAAGTTACCTTAGAGATTGAATAAGATGGATA
G0210463-1.seq(1>10379)	→		ttaaccgttaagttaccttagagatttgaataagatggata
		5500 5510 5520 5530	TTCTCATCAGTAGTTGAAAAGTTTAGGTTATTCTAGGACT
G0210463-1.seq(1>10379)	→		ttctcatcagtagtttgaaaagtttaggttattctaggact
		5540 5550 5560 5570	TCCAAATCTTTAGATTCAGGACCTTTGGTAGTACATGCAGT
G0210463-1.seq(1>10379)	→		tccaaatctttagattcaggacctttggtagtacatgcagt
		5580 5590 5600 5610	AGCCGGAGCCGTAAGTCCACAGCCCTAAGGAAGTTGATCC
G0210463-1.seq(1>10379)	→		agccggagccgtaagtccacagccctaaggaagttgatcc
		5620 5630 5640 5650	TCAGACACCCAACATTCACCGTGCATACACTCGGTGTCCCT
G0210463-1.seq(1>10379)	→		tcagacacccaacattcaccgtgcatacactcgggtgtccct
		5660 5670 5680 5690	GACAAGGTGAGTATCAGAAGTAGAGGCATACAGAAGCCAGG
G0210463-1.seq(1>10379)	→		gacaaggtgagtatcagaactagaggcatacagaagccagg
		5700 5710 5720 5730 5740	ACCTATTCCTGAGGGCAACTTCGCAATCCTCGATGAGTATA
G0210463-1.seq(1>10379)	→		acctattcctgagggcaacttcgcaatcctcgatgagtata
		5750 5760 5770 5780	CTTTGGACAACACCACAAGGAAGTCTAACCAGGCCTTTTT
G0210463-1.seq(1>10379)	→		ctttggacaacaccacaaggaactctaaccaggcacttttt
		5790 5800 5810 5820	GCTGACCCTTATCAGGCACCGGAGTTTAGCCTAGAGCCCCA
G0210463-1.seq(1>10379)	→		gctgacccttatcaggcaccggagtttagcctagagcccca
		5830 5840 5850 5860	CTTCTACTTGAAACATCATTTTCGAGTCCGAGGAAAGTGG
G0210463-1.seq(1>10379)	→		cttctacttgaaacatcatttfcgagttccgaggaaagtgg
		5870 5880 5890 5900	CAGATTTGATAGCTGGCTGTGGCTTCGATTTTCGAGACCAAC
G0210463-1.seq(1>10379)	→		cagatttgatagctggctgtggcttcgatttfcgagaccaac

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	5910	5920	5930	5940	TCACCGGAAGAAGGGCACTTAGAGATCACTGGCATATTCAA
G0210463-1.seq(1>10379)	→	tcaccggaagaagggcacttagagatcactggcatattcaa			
	5950	5960	5970	5980	AGGGCCCCTACTCGGAAAGGTGATAGCCATTGATGAGGAGT
G0210463-1.seq(1>10379)	→	agggcccctactcggaaaggtgatagccattgatgaggagt			
	5990	6000	6010	6020	CTGAGACAACACTGTCCAGGCATGGTGTGAGTTTGTTAAG
G0210463-1.seq(1>10379)	→	ctgagacaacactgtccaggcatgggtgtgagtttgtttaag			
	6030	6040	6050	6060	CCCTGCCAAGTGACGGGACTTGAGTTCAAAGTAGTCACTAT
G0210463-1.seq(1>10379)	→	ccctgccaagtgacgggacttgagttcaaagtagtcactat			
	6070	6080	6090	6100	TGTGTCTGCCGCACCAATAGAGGAAATTGGCCAGTCCACAG
G0210463-1.seq(1>10379)	→	tgtgtctgccgcaccaatagaggaaattggccagtccacag			
	6110	6120	6130	6140	CTTTCTACAACGCTATCACCAGGTCAAAGGGATTGACATAT
G0210463-1.seq(1>10379)	→	ctttctacaacgctatcaccaggtcaaagggattgacatat			
	6160	6170	6180	6190	GTCCGCGCAGGGCCATAGGCTGACCGCTCCGGTCAATTCTG
G0210463-1.seq(1>10379)	→	gtccgcgcagggccataggctgaccgctccggccaattctg			
	6200	6210	6220	6230	AAAAAGTGTACATAGTATTAGGTCTATCATTGCTTTAGTT
G0210463-1.seq(1>10379)	→	aaaaagtgtacatagtattaggtctatcatttgctttagtt			
	6240	6250	6260	6270	TCAATTACCTTTCTGCTTTCTAGAAATAGCTTACCCACGT
G0210463-1.seq(1>10379)	→	tcaattacctttctgctttctagaaatagcttaccacgt			
	6280	6290	6300	6310	CGGTGACAACATTCACAGCTTGCCACACGGAGGAGCTTACA
G0210463-1.seq(1>10379)	→	cggtgacaacattcacagcttgccacacggaggagcttaca			
	6320	6330	6340	6350	GAGACGGCACCAAGCAATCTTGTACAACCTCCCAAATCTA
G0210463-1.seq(1>10379)	→	gagacggcaccaagcaatcttgtacaactcccaaactta			
	6360	6370	6380	6390	GGGTCACGAGTGAGTCTACACAACGGAAAGAACGCAGCATT
G0210463-1.seq(1>10379)	→	gggtcacgagtgagtctacacaacggaaagaacgcagcatt			

	6400 6410 6420 6430	TGCTGCCGTTTTGCTACTGACTTTGCTGATCTATGGAAGTA
G0210463-1.seq(1>10379)	→	tgctgccgttttgctactgactttgctgatctatggaagta
	6440 6450 6460 6470	AATACATATCTCAACGCAATCATACTTGTGCTTGTGGTAAC
G0210463-1.seq(1>10379)	→	aatacatatctcaacgcaatcatacttgtgcttgtggtaac
	6480 6490 6500 6510	AATCATAGCAGTCATTAGCACTTCCTTAGTGAGGACTGAAC
G0210463-1.seq(1>10379)	→	aatcatagcagtcattagcacttccttagtgaggactgaac
	6520 6530 6540 6550 6560	CTTGTGTCATCAAGATTACTGGGGAATCAATCACAGTGTTG
G0210463-1.seq(1>10379)	→	cttgtgtcatcaagattactggggaatcaatcacagtgttg
	6570 6580 6590 6600	GCTTGCAAAC TAGATGCAGAAACCATAAGGGCCATTGCCGA
G0210463-1.seq(1>10379)	→	gcttgcaaactagatgcagaaaccataagggccattgccga
	6610 6620 6630 6640	TCTCAAGCCACTCTCCGTTGAACGGTTAAGTTTCCATTGAT
G0210463-1.seq(1>10379)	→	tctcaagccactctccgttgaacggttaagtttccattgat
	6650 6660 6670 6680	ACTCGAAAGAGGTCAGCACCAGCTAGCATCGATTGGCGCGC
G0210463-1.seq(1>10379)	→	actcgaagaggtcagcaccagctagcatcgattggcgcgc
	6690 6700 6710 6720	CTGGAGCGGCCGCGTCGACCCGCGATAAGCTTGATAGGGCC
G0210463-1.seq(1>10379)	→	ctggagcggccgcgctcgaccgcgataagcttgatagggcc
	6730 6740 6750 6760	ATTGCCGATCTCAAGCCACTCTCCGTTGAACGGTTAAGTTT
G0210463-1.seq(1>10379)	→	attgccgatctcaagccactctccgttgaacggttaagttt
	6770 6780 6790 6800	CCATTGATACTCGAAAGATGTCAGCACCAGCTAGCACAAACA
G0210463-1.seq(1>10379)	→	ccattgatactcgaagatgtcagcaccagctagcacaaaca
	6810 6820 6830 6840	CAGCCCATAGGGTCAACTACCTCAACTACCACAAAACTGC
G0210463-1.seq(1>10379)	→	cagcccataggggtcaactacctcaactaccacaaaaactgc
	6850 6860 6870 6880	AGGCGCAACTCCTGCCACAGCTTCAGGCCTGTTACCATCC
G0210463-1.seq(1>10379)	→	aggcgcaactcctgccacagcttcaggcctgttacccatcc

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6890 6900 6910 6920
 CGGATGGGGATTTCTTTAGTACAGCCCCTGCCATAGTAGCC
 G0210463-1.seq(1>10379) → cggatggggatttcttttagtacagcccctgccatagtagcc

6930 6940 6950 6960 6970
 AGCAATGCTGTCGCAACAAATGAGGACCTCAGCAAGATTGA
 G0210463-1.seq(1>10379) → agcaatgctgctcgcaacaaatgaggacctcagcaagattga

6980 6990 7000 7010
 GGCTATTTGGAAGGACATGAAGGTGCCACAGACACTATGG
 G0210463-1.seq(1>10379) → ggctatTTGGAAGGACATGAAGGTGCCACAGACACTATGG

7020 7030 7040 7050
 CACAGGCTGCTTGGGACTTAGTCAGACACTGTGCTGATGTA
 G0210463-1.seq(1>10379) → cacaggctgcttgggacttagtcagacactgtgctgatgta
 F09-E17216-G0210463-1_M13F.ab1(1>794) ← CACTGTGCTGATGTA

7060 7070 7080 7090
 GGATCATCCGCTCAAACAGAAATGATAGATACAGGTCCCTA
 G0210463-1.seq(1>10379) → ggatcatccgctcaaacagaaatgatagatacagggtcccta
 F09-E17216-G0210463-1_M13F.ab1(1>794) ← GGATCATCCGCTCAAACAGAAATGATAGATACAGGTCCCTA

7100 7110 7120 7130
 TTCCAACGGCATCAGCAGAGCTAGACTGGCAGCAGCAATTA
 G0210463-1.seq(1>10379) → ttccaacggcatcagcagagctagactggcagcagcaatta
 F09-E17216-G0210463-1_M13F.ab1(1>794) ← TTCCAACGGCATCAGCAGAGCTAGACTGGCAGCAGCAATTA

7140 7150 7160 7170
 AAGAGGTGTGCACACTTAGGCAATTTTGCATGAAGTATGCT
 G0210463-1.seq(1>10379) → aagaggtgtgcacacttaggcaatTTTGCATGAAGTATGCT
 F09-E17216-G0210463-1_M13F.ab1(1>794) ← AAGAGGTGTGCACACTTAGGCAATTTTGCATGAAGTATGCT

7180 7190 7200 7210
 CCAGTGGTATGGAAGTGGATGTTAACTAACAACAGTCCACC
 G0210463-1.seq(1>10379) → ccagtggatggaactggatgTTAACTAACAACAGTCCACC
 F09-E17216-G0210463-1_M13F.ab1(1>794) ← CCAGTGGTATGGAAGTGGATGTTAACTAACAACAGTCCACC

7220 7230 7240 7250
 TGCTAACTGGCAAGCACAAGGTTTCAAGCCTGAGCACAAAT
 G0210463-1.seq(1>10379) → tgctaaactggcaagcacaaggTTTCAAGCCTGAGCACAAAT
 F09-E17216-G0210463-1_M13F.ab1(1>794) ← TGCTAACTGGCAAGCACAAGGTTTCAAGCCTGAGCACAAAT

7260 7270 7280 7290
 TCGCTGCATTTCGACTTCTTCAATGGAGTCACCAACCCAGCT
 G0210463-1.seq(1>10379) → tcgctgcattTCGACTTCTTCAATGGAGTCACCAACCCAGCT
 F09-E17216-G0210463-1_M13F.ab1(1>794) ← TCGCTGCATTTCGACTTCTTCAATGGAGTCACCAACCCAGCT

7300 7310 7320 7330
 GCCATCATGCCCAAAGAGGGGCTCATCCGGCCACCGTCTGA
 G0210463-1.seq(1>10379) → gccatcatgcccaaagaggggctcatccggccaccgtctga

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		7300	7310	7320	7330	
		----- ----- ----- -----				
		GCCATCATGCCCAAAGAGGGGCTCATCCGGCCACCGTCTGA				
F09-E17216-G0210463-1_M13F.ab1 (1>794)	←	GCCATCATGCCCAAAGAGGGGCTCATCCGGCCACCGTCTGA				
		7340	7350	7360	7370	7380
		----- ----- ----- ----- -----				
		AGCTGAAATGAATGCTGCCCAAAGTCTGCCTTTGTGAAGA				
G0210463-1.seq (1>10379)	→	agctgaaatgaatgctgcccAAAGTCTGCCTTTGTGAAGA				
F09-E17216-G0210463-1_M13F.ab1 (1>794)	←	AGCTGAAATGAATGCTGCCCAAAGTCTGCCTTTGTGAAGA				
		7390	7400	7410	7420	
		----- ----- ----- -----				
		TTACAAAGGCCAGGGCACAATCCAACGACTTTGCCAGCCTA				
G0210463-1.seq (1>10379)	→	ttacaaaggccagggcacaatccaacgactttgccagccta				
F09-E17216-G0210463-1_M13F.ab1 (1>794)	←	TTACAAAGGCCAGGGCACAATCCAACGACTTTGCCAGCCTA				
		7430	7440	7450	7460	
		----- ----- ----- -----				
		GATGCAGCTGTCACTCGAGGTCGTATCACTGGAACAACAAC				
G0210463-1.seq (1>10379)	→	gatgcagctgtcactcgaggtcgtatcactggaacaacaac				
F09-E17216-G0210463-1_M13F.ab1 (1>794)	←	GATGCAGCTGTCACTCGAGGTCGTATCACTGGAACAACAAC				
		7470	7480	7490	7500	
		----- ----- ----- -----				
		CGCTGAGGCTGTTGTCACTCTACCACCACCATAACTACGTC				
G0210463-1.seq (1>10379)	→	cgctgaggctgttgtcactctaccaccaccataactacgtd				
F09-E17216-G0210463-1_M13F.ab1 (1>794)	←	CGCTGAGGCTGTTGTCACTCTACCACCACCATAACTACGTC				
		7510	7520	7530	7540	
		----- ----- ----- -----				
		TACATAACCGACGCCTACCCAGTTTCATAGTATTTTCTGG				
G0210463-1.seq (1>10379)	→	tacataaccgacgcctacccagtttcatagtatTTTCTGG				
F09-E17216-G0210463-1_M13F.ab1 (1>794)	←	TACATAACCGACGCCTACCCAGTTTCATAGTATTTTCTGG				
		7550	7560	7570	7580	
		----- ----- ----- -----				
		TTTGATTGTATGAATAATATAAATAAAAAAAAAAAAACTAGT				
G0210463-1.seq (1>10379)	→	tttgattgtatgaataatataaataaaaaaaaaaaaaactagt				
F09-E17216-G0210463-1_M13F.ab1 (1>794)	←	TTTGATTGTATGAATAATATAAATAAAAAAAAAAAAACTAGT				
		7590	7600	7610	7620	
		----- ----- ----- -----				
		GGGTACCGCGAATTTCCCCGATCGTTCAAACATTTGGCAAT				
G0210463-1.seq (1>10379)	→	gggtaccgCGAATTTCCCCGATCGTTCAAACATTTGGCAAT				
F09-E17216-G0210463-1_M13F.ab1 (1>794)	←	GGGTACCGCGAATTTCCCCGATCGTTCAAACATTTGGCAAT				
		7630	7640	7650	7660	
		----- ----- ----- -----				
		AAAGTTTCTTAAGATTGAATCCTGTTGCCGGTCTTGCGATG				
G0210463-1.seq (1>10379)	→	aaagtttcttaagattgaatcctgTTGCCGGTCTTGCGATG				
F09-E17216-G0210463-1_M13F.ab1 (1>794)	←	AAAGTTTCTTAAGATTGAATCCTGTTGCCGGTCTTGCGATG				
		7670	7680	7690	7700	
		----- ----- ----- -----				
		ATTATCATATAATTTCTGTTGAATTACGTTAAGCATGTAAT				
G0210463-1.seq (1>10379)	→	attatcatataatTTTCTGTTGAATTACGTTAAGCATGTAAT				
F09-E17216-G0210463-1_M13F.ab1 (1>794)	←	ATTATCATATAATTTCTGTTGAATTACGTTAAGCATGTAAT				

		7710	7720	7730	7740	
		AATTAACATGTAATGCATGACGTTATTTATGAGATGGGTTT				
G0210463-1.seq(1>10379)	→	aattaacatgtaatgcatgacgttatTTATGAGATGGGTTT				
F09-E17216-G0210463-1_M13F.ab1(1>794)	←	AATTAACATGTAATGCATGACGTTATTTATGAGATGGGTTT				
		7750	7760	7770	7780	7790
		TTATGATTAGAGTCCCGCAATTATACATTTAATACGCGATA				
G0210463-1.seq(1>10379)	→	ttatgattagagtcccgcaattataacatttaatacgcgata				
F09-E17216-G0210463-1_M13F.ab1(1>794)	←	TTATGATTAGAGTCCCGCAATTATACATTTAATACGCGATA				
		7800	7810	7820	7830	
		GAAAACAAAATATAGCGCGCAAACCTAGGATAAATTATCGCG				
G0210463-1.seq(1>10379)	→	gaaaacaaaatatagcgcgcaaactaggataaattatcgcg				
F09-E17216-G0210463-1_M13F.ab1(1>794)	←	GAAAACAAAATATAGCGCGCAAACCTAGGATAAATTATCGCG				
		7840	7850	7860	7870	
		CGCGGTGTCATCTATGTTACTAGATCGAATTCCTGGCCGT				
G0210463-1.seq(1>10379)	→	cgcggtgtcatctatgTTACTAGATCGAATTCCTGGCCGT				
		7880	7890	7900	7910	
		CGTTTTACAACGTCGTGACTGGGAAAACCTGGCGTTACCC				
G0210463-1.seq(1>10379)	→	cgTTTTACAACGTCGTGACTGGGAAAACCTGGCGTTACCC				
		7920	7930	7940	7950	
		AACTTAATCGCCTTGCAGCACATCCCCCTTTCGCCAGCTGG				
G0210463-1.seq(1>10379)	→	aacttaatcgCCTTGCAGCACATCCCCCTTTCGCCAGCTGG				
		7960	7970	7980	7990	
		CGTAATAGCGAAGAGGCCCGCACCGATCGCCCTTCCCAACA				
G0210463-1.seq(1>10379)	→	cgtaatagcgaagaggccCGCACCGATCGCCCTTCCCAACA				
		8000	8010	8020	8030	
		GTTGCGCAGCCTGAATGGCGAATGGCGCCTGATGAGCTCCA				
G0210463-1.seq(1>10379)	→	gTTGCGCAGCCTGAATGGCGAATGGCGCCTGATGAGCTCCA				
		8040	8050	8060	8070	
		GCTTTTGTTCCTTTAGTGAGGGTTAATTCCGAGCTTGGCG				
G0210463-1.seq(1>10379)	→	gctTTTGTTCCTTTAGTGAGGGTTAATTCCGAGCTTGGCG				
		8080	8090	8100	8110	
		TAATCATGGTCATAGCTGTTTCTGTGTGAAATTGTTATCC				
G0210463-1.seq(1>10379)	→	taatcatggTCATAGCTGTTTCTGTGTGAAATTGTTATCC				
		8120	8130	8140	8150	
		GCTCACAATTCCACACAACATACGAGCCGGAAGCATAAAGT				
G0210463-1.seq(1>10379)	→	gctcacaATTCCACACAACATACGAGCCGGAAGCATAAAGT				

	8160	8170	8180	8190	8200
	GTAAGCCTGGGGTGCCTAATGAGTGAGCTAACTCACATTA				
G0210463-1.seq(1>10379)	→	gtaaagcctgggggtgcctaataatgagtgagctaactcacatta			
	8210	8220	8230	8240	
	ATTGCGTTGCGCTCACTGCCCGCTTTCAGTCGGGAAACCT				
G0210463-1.seq(1>10379)	→	attgcgttgcgctcactgcccgctttccagtcgggaaacct			
	8250	8260	8270	8280	
	GTCGTGCCAGCTGCATTAATGAATCGGCCAACGCGCGGGGA				
G0210463-1.seq(1>10379)	→	gtcgtgccagctgcattaatgaatcggccaacgcgcgggga			
	8290	8300	8310	8320	
	GAGGCGTTTGCATATTGGGCGCTCTTCCGCTTCTCGCTC				
G0210463-1.seq(1>10379)	→	gaggcggtttgcataattgggcgctcttccgcttctcgctc			
	8330	8340	8350	8360	
	ACTGACTCGCTGCGCTCGGTCTGTTCCGGCTGCGGCGAGCGGT				
G0210463-1.seq(1>10379)	→	actgactcgctgcgctcggctcggtcggctgcggcgagcgggt			
	8370	8380	8390	8400	
	ATCAGCTCACTCAAAGGCGGTAATACGTTTATCCACAGAAT				
G0210463-1.seq(1>10379)	→	atcagctcactcaaaggcggtaatacggttatccacagaat			
	8410	8420	8430	8440	
	CAGGGGATAACGCAGGAAAGAACATGAAGGCCTTGACAGGA				
G0210463-1.seq(1>10379)	→	caggggataacgcaggaagaacatgaaggccttgacagga			
	8450	8460	8470	8480	
	TATATTGGCGGGTAAACTAAGTCGCTGTATGTGTTTGTGTTG				
G0210463-1.seq(1>10379)	→	tatattggcgggtaactaagtcgctgtatgtgtttgtgttg			
	8490	8500	8510	8520	
	AGATCTCATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACC				
G0210463-1.seq(1>10379)	→	agatctcatgtgagcaaaaggccagcaaaaggccaggaacc			
	8530	8540	8550	8560	
	GTAAAAGGCCGCGTTGCTGGCGTTTTTCCATAGGCTCCGC				
G0210463-1.seq(1>10379)	→	gtaaaaaggccgcttgctggcgtttttccataggctccgc			
	8570	8580	8590	8600	8610
	CCCCCTGACGAGCATCACAAAATCGACGCTCAAGTCAGAG				
G0210463-1.seq(1>10379)	→	ccccctgacgagcatcacaaaatcgacgctcaagtcagag			
	8620	8630	8640	8650	
	GTGGCGAAACCCGACAGGACTATAAAGATAACCAGGCGTTTC				
G0210463-1.seq(1>10379)	→	gtggcgaaacccgacaggactataaagataaccaggcgttttc			

		8660	8670	8680	8690	
		CCCCTGGAAGCTCCCTCGTGCCTCTCCTGTTCCGACCTG				
G0210463-1.seq(1>10379)	→	cccctggaagctccctcgtgcgctctcctgttccgaccctg				
		8700	8710	8720	8730	
		CCGCTTACCGGATACCTGTCCGCCTTCTCCCTTCGGGAAG				
G0210463-1.seq(1>10379)	→	ccgcttaccggatacctgtccgcctttctcccttcgggaag				
		8740	8750	8760	8770	
		CGTGCGCTTCTCATAGCTCACGCTGTAGGTATCTCAGTT				
G0210463-1.seq(1>10379)	→	cgtggcgctttctcatagctcacgctgtaggtatctcagtt				
		8780	8790	8800	8810	
		CGGTGTAGGTCGTTTCGCTCCAAGCTGGGCTGTGTGCACGAA				
G0210463-1.seq(1>10379)	→	cggtgtaggtcgttcgctccaagctgggctgtgtgcacgaa				
		8820	8830	8840	8850	
		CCCCCGTTTCAGCCCGACCGCTGCGCCTTATCCGGTAACTA				
G0210463-1.seq(1>10379)	→	cccccgttcagcccgaccgctgcgcttatccggtaacta				
		8860	8870	8880	8890	
		TCGTCTTGAGTCCAACCCGGTAAGACACGACTTATCGCCAC				
G0210463-1.seq(1>10379)	→	tcgtcttgagtccaacccggtaagacacgacttatcgccac				
		8900	8910	8920	8930	
		TGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTAT				
G0210463-1.seq(1>10379)	→	tggcagcagccactggtaacaggattagcagagcgaggtat				
		8940	8950	8960	8970	
		GTAGGCGGTGCTACAGAGTTCTTGAAGTGGTGGCCTAACTA				
G0210463-1.seq(1>10379)	→	gtaggcggtgctacagagttcttgaagtggccttaacta				
		8980	8990	9000	9010	9020
		CGGCTACACTAGAAGGACAGTATTTGGTATCTGCGCTCTGC				
G0210463-1.seq(1>10379)	→	cggctacactagaaggacagtatttggatatctgcgctctgc				
		9030	9040	9050	9060	
		TGAAGCCAGTTACCTTCGGAAAAAGAGTTGGTAGCTCTTGA				
G0210463-1.seq(1>10379)	→	tgaagccagttaccttcggaaaaagagttggtagctcttga				
		9070	9080	9090	9100	
		TCCGGCAAACAAACCACCGCTGGTAGCGGTGGTTTTTTTGT				
G0210463-1.seq(1>10379)	→	tccggcaaacaaaccaccgctggtagcggcggtttttttgt				
		9110	9120	9130	9140	
		TTGCAAGCAGCAGATTACGCGCAGAAAAAAGGATCTCAAG				
G0210463-1.seq(1>10379)	→	ttgcaagcagcagattacgcgcagaaaaaaggatctcaag				

		9150	9160	9170	9180	
		AAGATCCTTTGATCTTTTCTACGGGGTCTGACGCTCAGTGG				
G0210463-1.seq(1>10379)	→	aagatcctttgatcttttctacggggctgacgctcagtgg				
		9190	9200	9210	9220	
		AACGAAAACCTCACGTTAAGGGATTTTGGTCATGGTTACAAC				
G0210463-1.seq(1>10379)	→	aacgaaaactcacgttaagggatTTTGGTCATGGTTACAAC				
		9230	9240	9250	9260	
		CAATTAACCAATTCTGATTAGAAAACTCATCGAGCATCAA				
G0210463-1.seq(1>10379)	→	caattaaccaattctgattagaaaaactcatcgagcatcaa				
		9270	9280	9290	9300	
		ATGAAACTGCAATTTATTCATATCAGGATTATCAATACCAT				
G0210463-1.seq(1>10379)	→	atgaaactgcaatTTATTCATATCAGGATTATCAATACCAT				
		9310	9320	9330	9340	
		ATTTTGGAAAAAGCCGTTTCTGTAATGAAGGAGAAAACTCA				
G0210463-1.seq(1>10379)	→	atTTTGGAAAAAGCCGTTTCTGTAATGAAGGAGAAAACTCA				
		9350	9360	9370	9380	
		CCGAGGCAGTTCATAGGATGGCAAGATCCTGGTATCGGTC				
G0210463-1.seq(1>10379)	→	ccgaggcagTTCATAGGATGGCAAGATCCTGGTATCGGTC				
		9390	9400	9410	9420	9430
		TGCGATTCCGACTCGTCCAACATCAATACAACCTATTAATT				
G0210463-1.seq(1>10379)	→	tgcgattccgactCGTCCAACATCAATACAACCTATTAATT				
		9440	9450	9460	9470	
		TCCCCTCGTCAAAAATAAGGTTATCAAGTGAGAAATCACCA				
G0210463-1.seq(1>10379)	→	tcccctcgtcaaaaataaggttatcaagtgagaaatcacca				
		9480	9490	9500	9510	
		TGAGTGACGACTGAATCCGGTGAGAATGGCAAAGTTTATG				
G0210463-1.seq(1>10379)	→	tgagtgacgactgaatccggtgagaatggcaaaagTTTATG				
		9520	9530	9540	9550	
		CATTTCTTTCCAGACTTGTTCAACAGGCCAGCCATTACGCT				
G0210463-1.seq(1>10379)	→	catttctTTTCCAGACTTGTTCAACAGGCCAGCCATTACGCT				
		9560	9570	9580	9590	
		CGTCATCAAAATCACTCGCATCAACCAACCGTTATTCATT				
G0210463-1.seq(1>10379)	→	cgtcatcaaaatcactcgcATCAACCAACCGTTATTCATT				
		9600	9610	9620	9630	
		CGTGATTGCGCCTGAGCGAGACGAAATACGCGATCGCTGTT				
G0210463-1.seq(1>10379)	→	cgtgattgCGCCTGAGCGAGACGAAATACGCGATCGCTGTT				

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	9640 9650 9660 9670	AAAAGGACAATTACAAACAGGAATCGAATGCAACCGGCGCA
G0210463-1.seq(1>10379)	→	aaaaggacaattacaaacaggaatcgaatgcaaccggcgca
	9680 9690 9700 9710	GGAACACTGCCAGCGCATCAACAATATTTTCACCTGAATCA
G0210463-1.seq(1>10379)	→	ggaacactgccagcgcatcaacaatattttcacctgaatca
	9720 9730 9740 9750	GGATATTCTTCTAATACCTGGAATGCTGTTTTCCCTGGGAT
G0210463-1.seq(1>10379)	→	ggatattcttctaataacctggaatgctgttttccctgggat
	9760 9770 9780 9790	CGCAGTGGTGAGTAACCATGCATCATCAGGAGTACGGATAA
G0210463-1.seq(1>10379)	→	cgcagtggtgagtaaccatgcatcatcaggagtacggataa
	9800 9810 9820 9830 9840	AATGCTTGATGGTCGGAAGAGGCATAAATTCCGTCAGCCAG
G0210463-1.seq(1>10379)	→	aatgcttgatggtcggaagaggcataaattccgtcagccag
	9850 9860 9870 9880	TTTAGTCTGACCATCTCATCTGTAACATCATTGGCAACGCT
G0210463-1.seq(1>10379)	→	tttagtctgaccatctcatctgtaacatcattggcaacgct
	9890 9900 9910 9920	ACCTTTGCCATGTTTCAGAAACAACCTCTGGCGCATCGGGCT
G0210463-1.seq(1>10379)	→	acctttgccatgtttcagaaacaactctggcgcatcgggct
	9930 9940 9950 9960	TCCCATACAATCCATAGATTGTGCGACCTGATTGCCCGACA
G0210463-1.seq(1>10379)	→	tcccatacaatccatagattgtgacacctgattgcccgaca
	9970 9980 9990 10000	TTATCGCGAGCCATTTATACCCATATAAATCAGCATCCAT
G0210463-1.seq(1>10379)	→	ttatcgcgagcccatttatacccatataaatacagcatccat
	10010 10020 10030 10040	GTTGGAATTTAATCGCGGCTGGAGCAAGACGTTTCCCGTT
G0210463-1.seq(1>10379)	→	gttggaatttaatcgcggctggagcaagacgtttcccgtt
	10050 10060 10070 10080	GAATATGGCTCATAACACCCCTTGTATTACTGTTTATGTAA
G0210463-1.seq(1>10379)	→	gaatatggctcataaacacccttgtattactgtttatgtaa
	10090 10100 10110 10120	GCAGACAGTTTTATTGTTTCATGATGATATATTTTTATCTTG
G0210463-1.seq(1>10379)	→	gcagacagttttattgtttcatgatgatataatTTTTATCTTG

10130 10140 10150 10160
TGCAATGTAACATCAGAGATTTTGAGACACAACGTGGCTTT
G0210463-1.seq(1>10379) → tgcaatgtaacatcagagatTTTgagacacaacgtggcttt

10170 10180 10190 10200
GTTGAATAAAATCGAACTTTTGCTGAGTTGAAGGATCAGATC
G0210463-1.seq(1>10379) → gttgaataaaatcgaactTTTgctgagttgaaggatcagatc

10210 10220 10230 10240 10250
ACGCATCTTCCCGACAACGCAGACCGTTCCGTGGCAAAGCA
G0210463-1.seq(1>10379) → acgcatacttcccgacaacgcagaccgTTCCGTGGCAAAGCA

10260 10270 10280 10290
AAAGTTCAAAATCACCAACTGGTCCACCTACAACAAAGCTC
G0210463-1.seq(1>10379) → aaagttcaaaatcaccaactGGTCCACCTACAACAAAGCTC

10300 10310 10320 10330
TCATCAACCGTGGCTCCCTCACTTTCTGGCTGGATGATGGG
G0210463-1.seq(1>10379) → tcatcaaccgTGGCTCCCTCACTTTCTGGCTGGATGATGGG

10340 10350 10360 10370
GCGATTCAGGCGATCCCCATCCAACAGCCC GCCGTGAGCG
G0210463-1.seq(1>10379) → gcgattcaggcgatCCCCATCCAACAGCCC GCCGTGAGCG

GGCT
G0210463-1.seq(1>10379) → ggct