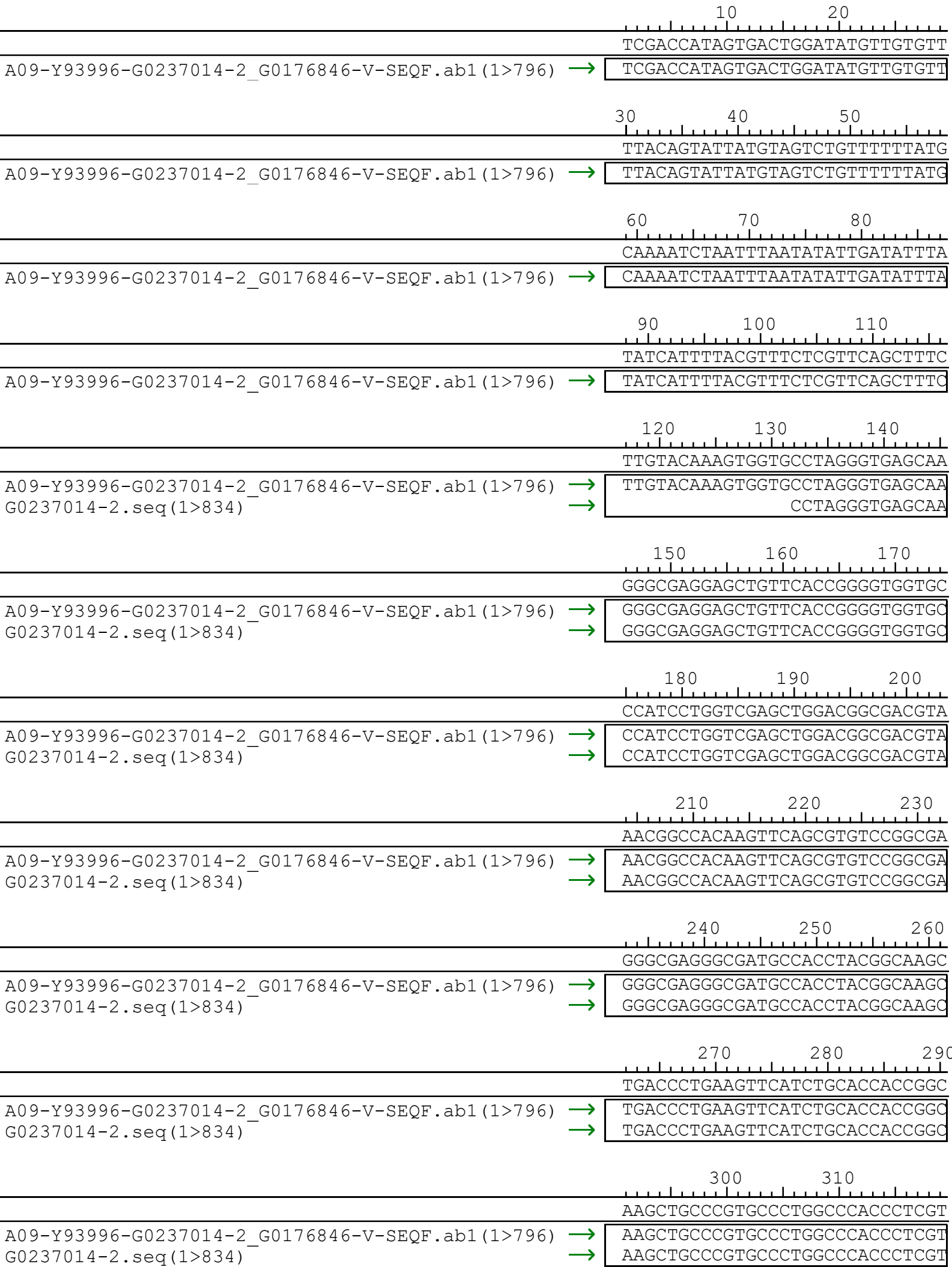


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		320	330	340	
		GACCACCCTGACCTGGGGCGTGCAGTGCT			
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	GACCACCCTGACCTGGGGCGTGCAGTGCT			
G0237014-2.seq (1>834)	→	GACCACCCTGACCTGGGGCGTGCAGTGCT			
		350	360	370	
		TCAGCCGCTACCCCGACCACATGAAGCAG			
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	TCAGCCGCTACCCCGACCACATGAAGCAG			
G0237014-2.seq (1>834)	→	TCAGCCGCTACCCCGACCACATGAAGCAG			
		380	390	400	
		CACGACTTCTTCAAGTCCGCCATGCCCGA			
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	CACGACTTCTTCAAGTCCGCCATGCCCGA			
G0237014-2.seq (1>834)	→	CACGACTTCTTCAAGTCCGCCATGCCCGA			
		410	420	430	
		AGGCTACGTCCAGGAGCGCACCATCTTCT			
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	AGGCTACGTCCAGGAGCGCACCATCTTCT			
G0237014-2.seq (1>834)	→	AGGCTACGTCCAGGAGCGCACCATCTTCT			
		440	450	460	
		TCAAGGACGACGGCAACTACAAGACCCGC			
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	TCAAGGACGACGGCAACTACAAGACCCGC			
G0237014-2.seq (1>834)	→	TCAAGGACGACGGCAACTACAAGACCCGC			
		470	480	490	
		GCCGAGGTGAAGTTCGAGGGCGACACCCT			
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	GCCGAGGTGAAGTTCGAGGGCGACACCCT			
G0237014-2.seq (1>834)	→	GCCGAGGTGAAGTTCGAGGGCGACACCCT			
		500	510	520	
		GGTGAACCGCATCGAGCTGAAGGGCATCG			
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	GGTGAACCGCATCGAGCTGAAGGGCATCG			
G0237014-2.seq (1>834)	→	GGTGAACCGCATCGAGCTGAAGGGCATCG			
		530	540	550	
		ACTTCAAGGAGGACGGCAACATCCTGGGG			
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	ACTTCAAGGAGGACGGCAACATCCTGGGG			
G0237014-2.seq (1>834)	→	ACTTCAAGGAGGACGGCAACATCCTGGGG			
E03-Y93996-G0237014-2_G0237014-2V-SEQR.ab1 (1>520)	←	TTCAAGGAGGACGGCAACATCCTGGGG			
		560	570	580	
		CACAAGCTGGAGTACAACATCAGCCA			
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	CACAAGCTGGAGTACAACATCAGCCA			
G0237014-2.seq (1>834)	→	CACAAGCTGGAGTACAACATCAGCCA			
E03-Y93996-G0237014-2_G0237014-2V-SEQR.ab1 (1>520)	←	CACAAGCTGGAGTACAACATCAGCCA			
		590	600		
		CAACGTCTATATCACCGCCGACAAGCAGA			
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	CAACGTCTATATCACCGCCGACAAGCAGA			
G0237014-2.seq (1>834)	→	CAACGTCTATATCACCGCCGACAAGCAGA			

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		590	600	CAACGTCTATATCACCGCCGACAAGCAGA	
E03-Y93996-G0237014-2_G0237014-2V-SEQR.ab1 (1>520)	←	CAACGTCTATATCACCGCCGACAAGCAGA			
		610	620	630	AGAACGGCATCAAGGCCAACTTCAAGATC
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	AGAACGGCATCAAGGCCAACTTCAAGATC			
G0237014-2.seq (1>834)	→	AGAACGGCATCAAGGCCAACTTCAAGATC			
E03-Y93996-G0237014-2_G0237014-2V-SEQR.ab1 (1>520)	←	AGAACGGCATCAAGGCCAACTTCAAGATC			
		640	650	660	CGCCACAACATCGAGGACGGCAGCGTGCA
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	CGCCACAACATCGAGGACGGCAGCGTGCA			
G0237014-2.seq (1>834)	→	CGCCACAACATCGAGGACGGCAGCGTGCA			
E03-Y93996-G0237014-2_G0237014-2V-SEQR.ab1 (1>520)	←	CGCCACAACATCGAGGACGGCAGCGTGCA			
		670	680	690	GCTCGCCGACCACTACCAGCAGAACACCC
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	GCTCGCCGACCACTACCAGCAGAACACCC			
G0237014-2.seq (1>834)	→	GCTCGCCGACCACTACCAGCAGAACACCC			
E03-Y93996-G0237014-2_G0237014-2V-SEQR.ab1 (1>520)	←	GCTCGCCGACCACTACCAGCAGAACACCC			
		700	710	720	CCATCGGCGACGGCCCCGTGCTGCTGCC
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	CCATCGGCGACGGCCCCGTGCTGCTGCC			
G0237014-2.seq (1>834)	→	CCATCGGCGACGGCCCCGTGCTGCTGCC			
E03-Y93996-G0237014-2_G0237014-2V-SEQR.ab1 (1>520)	←	CCATCGGCGACGGCCCCGTGCTGCTGCC			
		730	740	750	GACAACCACTACCTGAGCACCCAGTCCGC
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	GACAACCACTACCTGAGCACCCAGTCCGC			
G0237014-2.seq (1>834)	→	GACAACCACTACCTGAGCACCCAGTCCGC			
E03-Y93996-G0237014-2_G0237014-2V-SEQR.ab1 (1>520)	←	GACAACCACTACCTGAGCACCCAGTCCGC			
		760	770	780	CCTGAGCAAAGACCCCAACGAGAAGCGCG
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	CCTGAGCAAAGACCCCAACGAGAAGCGCG			
G0237014-2.seq (1>834)	→	CCTGAGCAAAGACCCCAACGAGAAGCGCG			
E03-Y93996-G0237014-2_G0237014-2V-SEQR.ab1 (1>520)	←	CCTGAGCAAAGACCCCAACGAGAAGCGCG			
		790	800	810	ATCACATGGTCCTGCTGGAGTTCGTGACC
A09-Y93996-G0237014-2_G0176846-V-SEQF.ab1 (1>796)	→	ATCACATGGTCCT			
G0237014-2.seq (1>834)	→	ATCACATGGTCCTGCTGGAGTTCGTGACC			
E03-Y93996-G0237014-2_G0237014-2V-SEQR.ab1 (1>520)	←	ATCACATGGTCCTGCTGGAGTTCGTGACC			
		820	830	840	GCCGCCGGGATCACTCTCGGCATGGACGA
G0237014-2.seq (1>834)	→	GCCGCCGGGATCACTCTCGGCATGGACGA			
E03-Y93996-G0237014-2_G0237014-2V-SEQR.ab1 (1>520)	←	GCCGCCGGGATCACTCTCGGCATGGACGA			

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