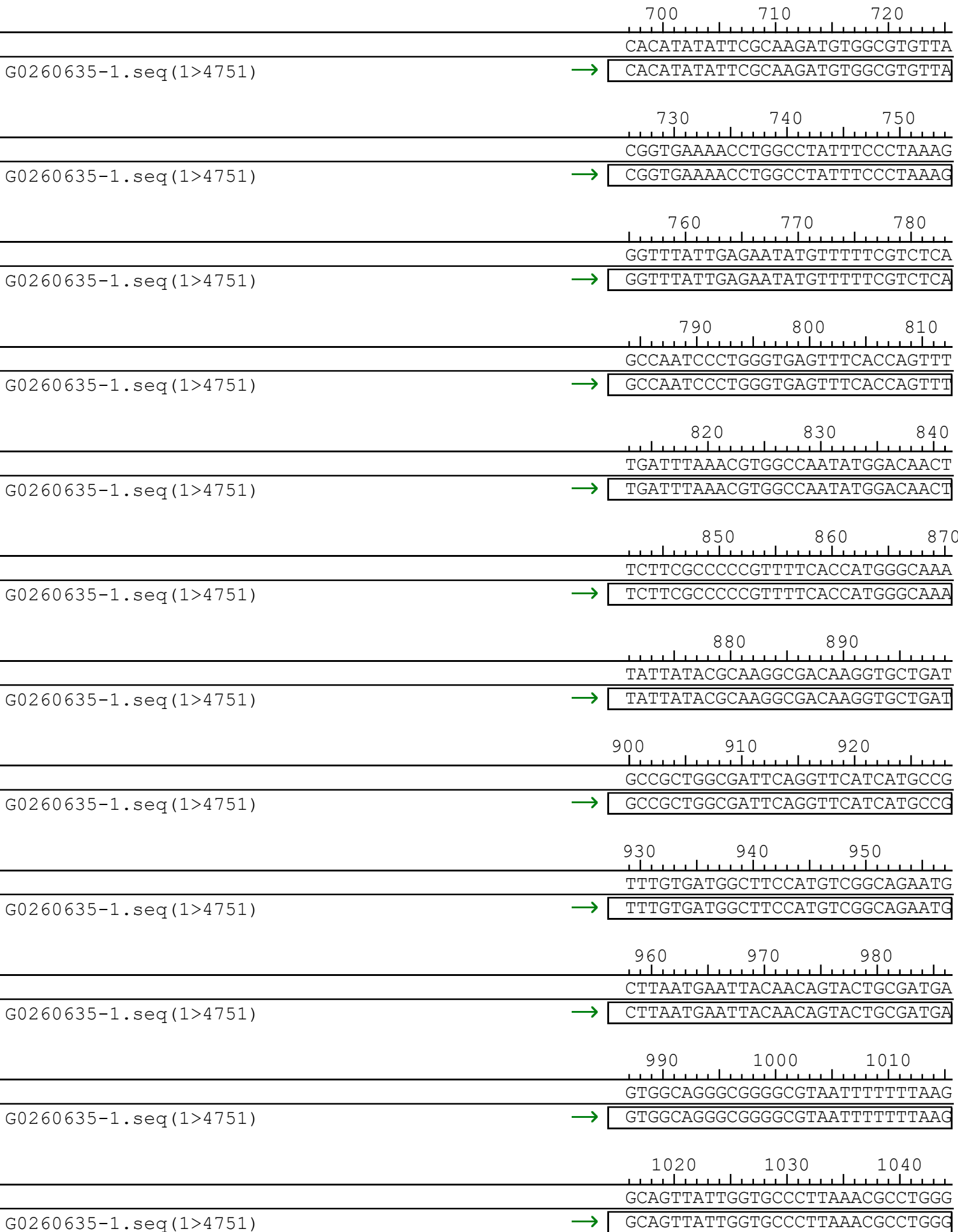
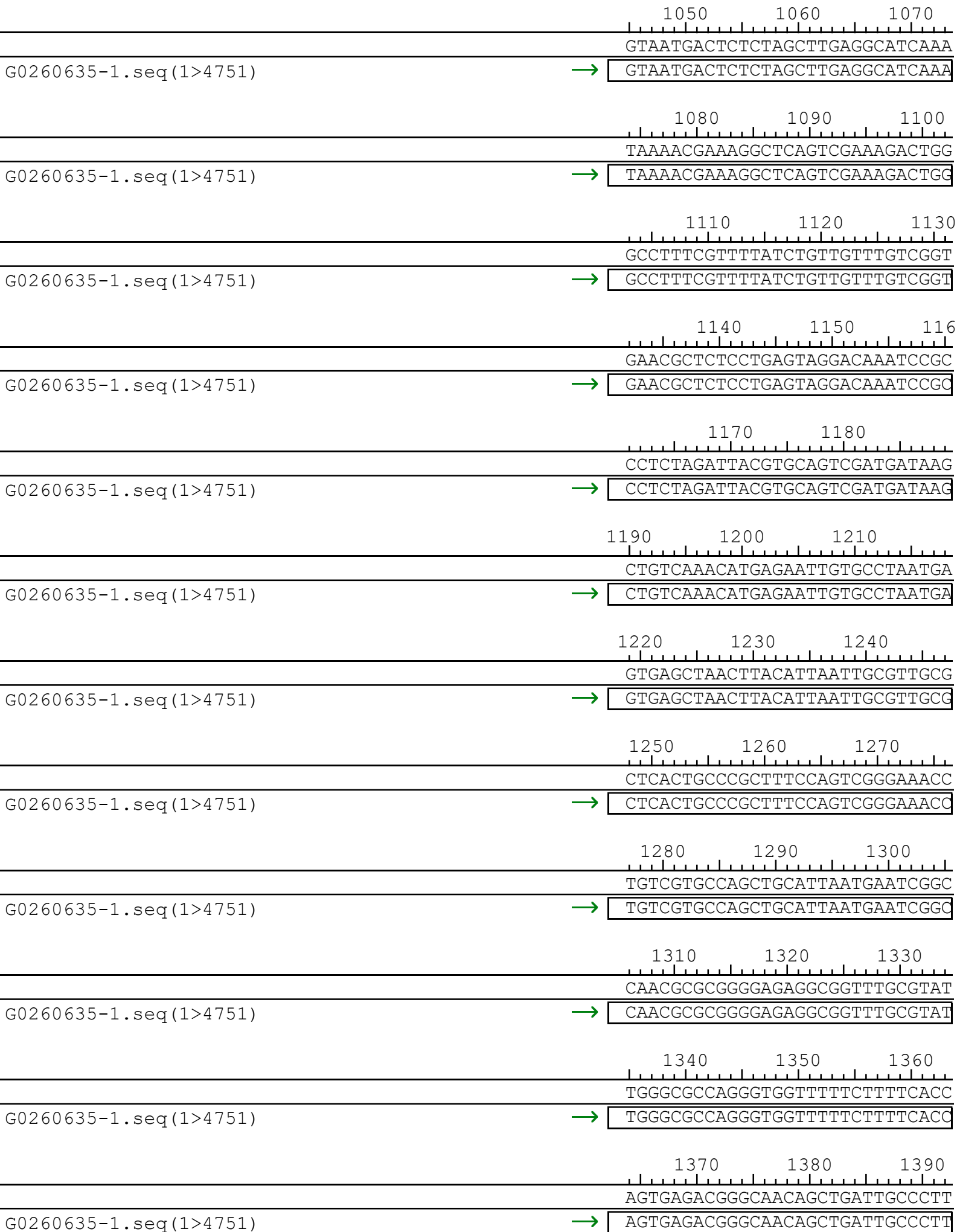


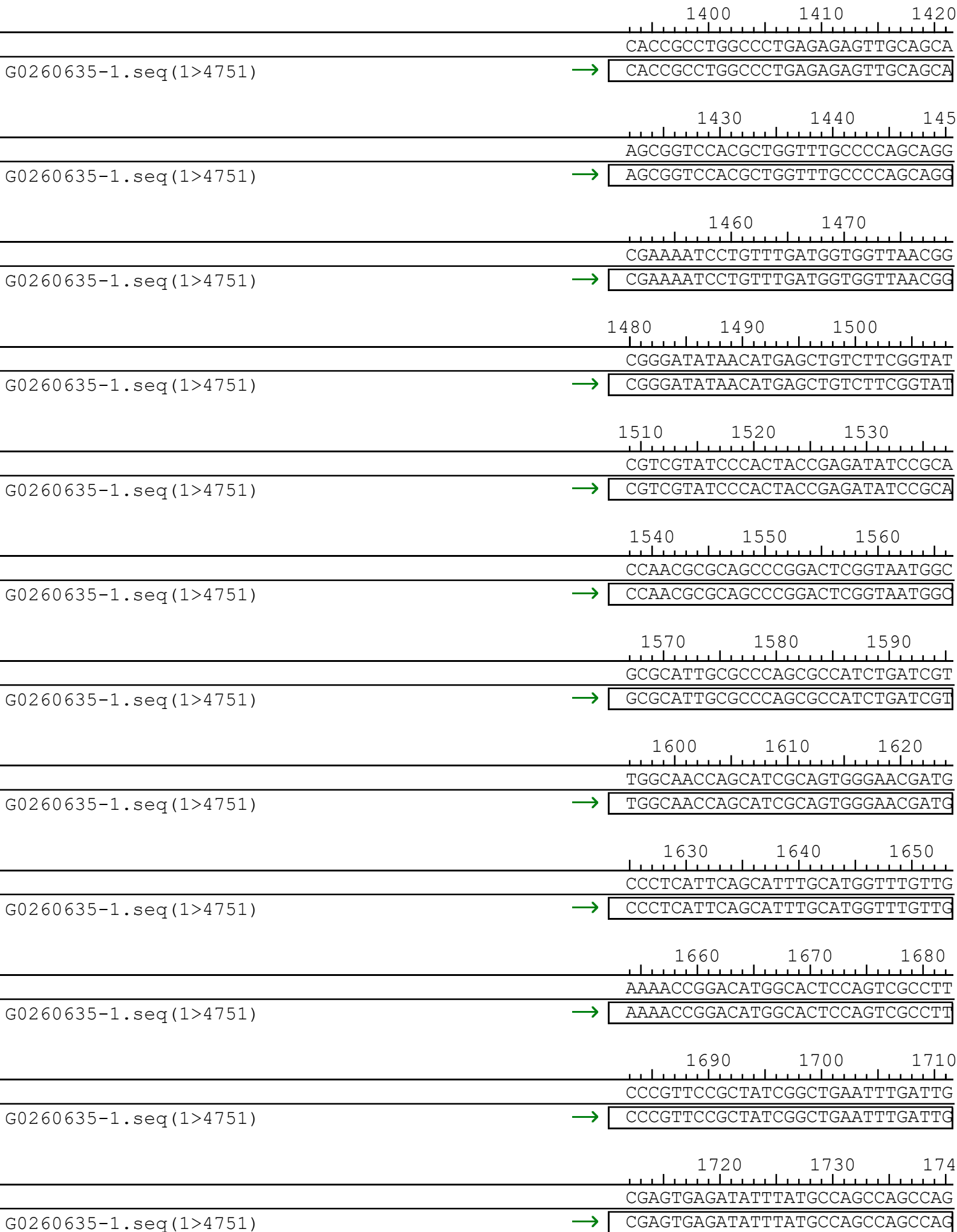
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	440	450	460	TGTACCTATAACCAGACCGTTCAGCTGGA
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	530	540	550	ATTCACATTCTTGCCCGCCTGATGAATGC
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G0260635-1.seq (1>4751)	→			TCATCCGGAATTTTCGTATGGCAATGAAAG
	590	600		ACGGTGAGCTGGTGATATGGGATAGTGTT
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	610	620	630	CACCCTTGTTACACCGTTTTCCATGAGCA
G0260635-1.seq (1>4751)	→			CACCCTTGTTACACCGTTTTCCATGAGCA
	640	650	660	AACTGAAACGTTTTTCATCGCTCTGGAGTG
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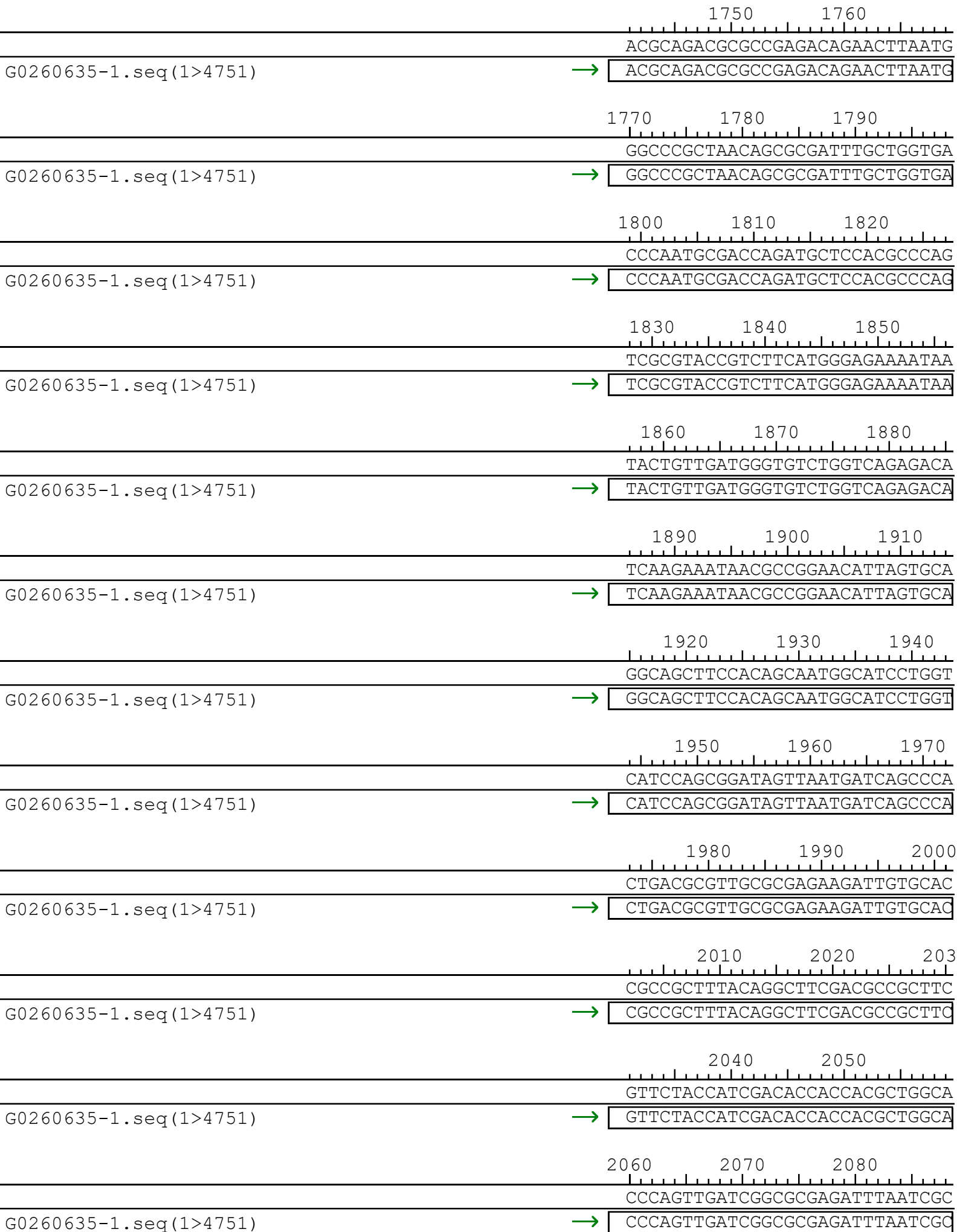
## Project: Untitled.SQD Contig 1





## Project: Untitled.SQD Contig 1





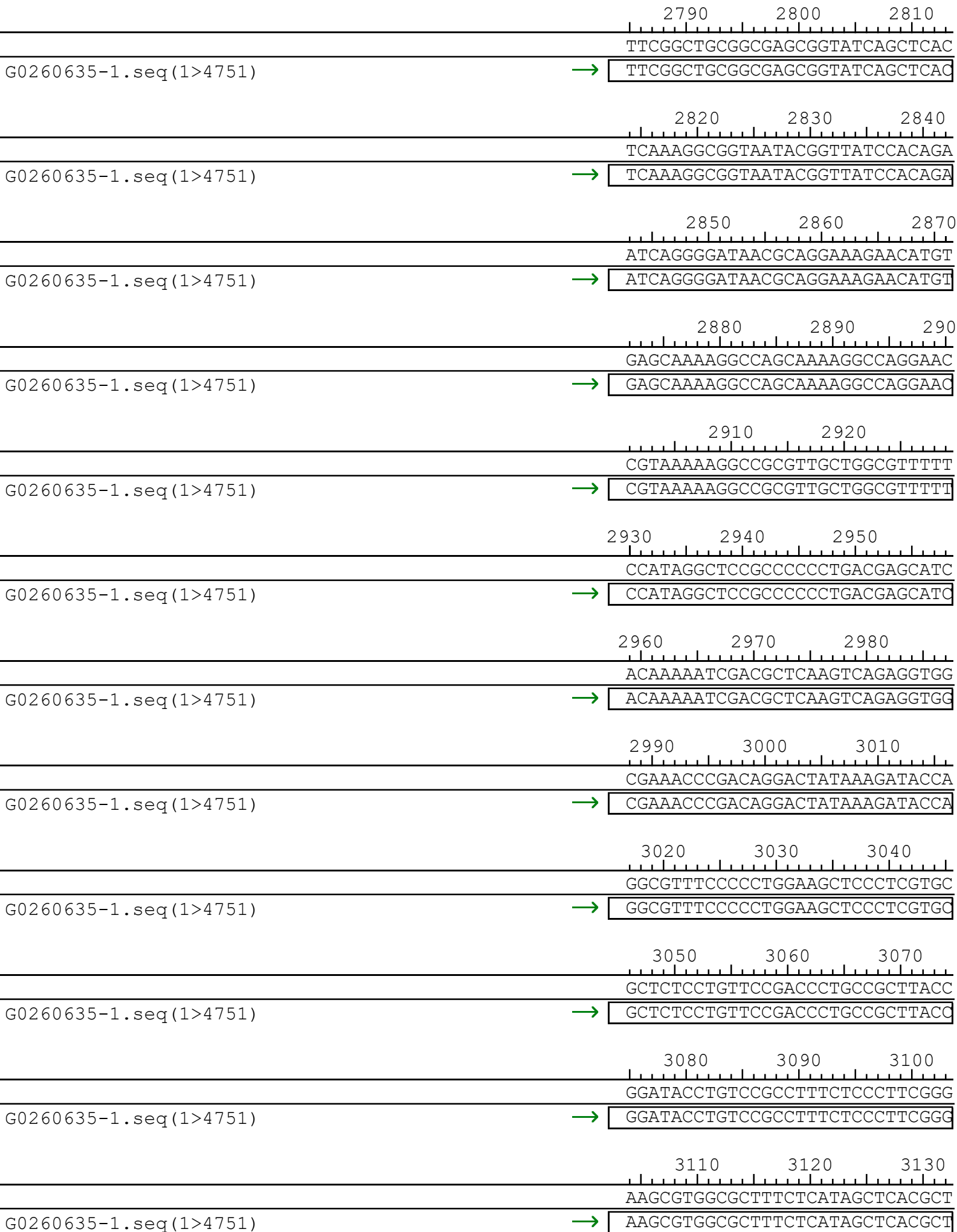
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		CACGCGGTTGGGAATGTAATTCAGCTCCG		
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		CCATCGCCGCTTCCACTTTTTCCCGCGTT		
G0260635-1.seq(1>4751)	→	CCATCGCCGCTTCCACTTTTTCCCGCGTT		
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		2270	2280	2290
		CACGCGGGAAACGGTCTGATAAGAGACAC		
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		2300	2310	232
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## Project: Untitled.SQD Contig 1

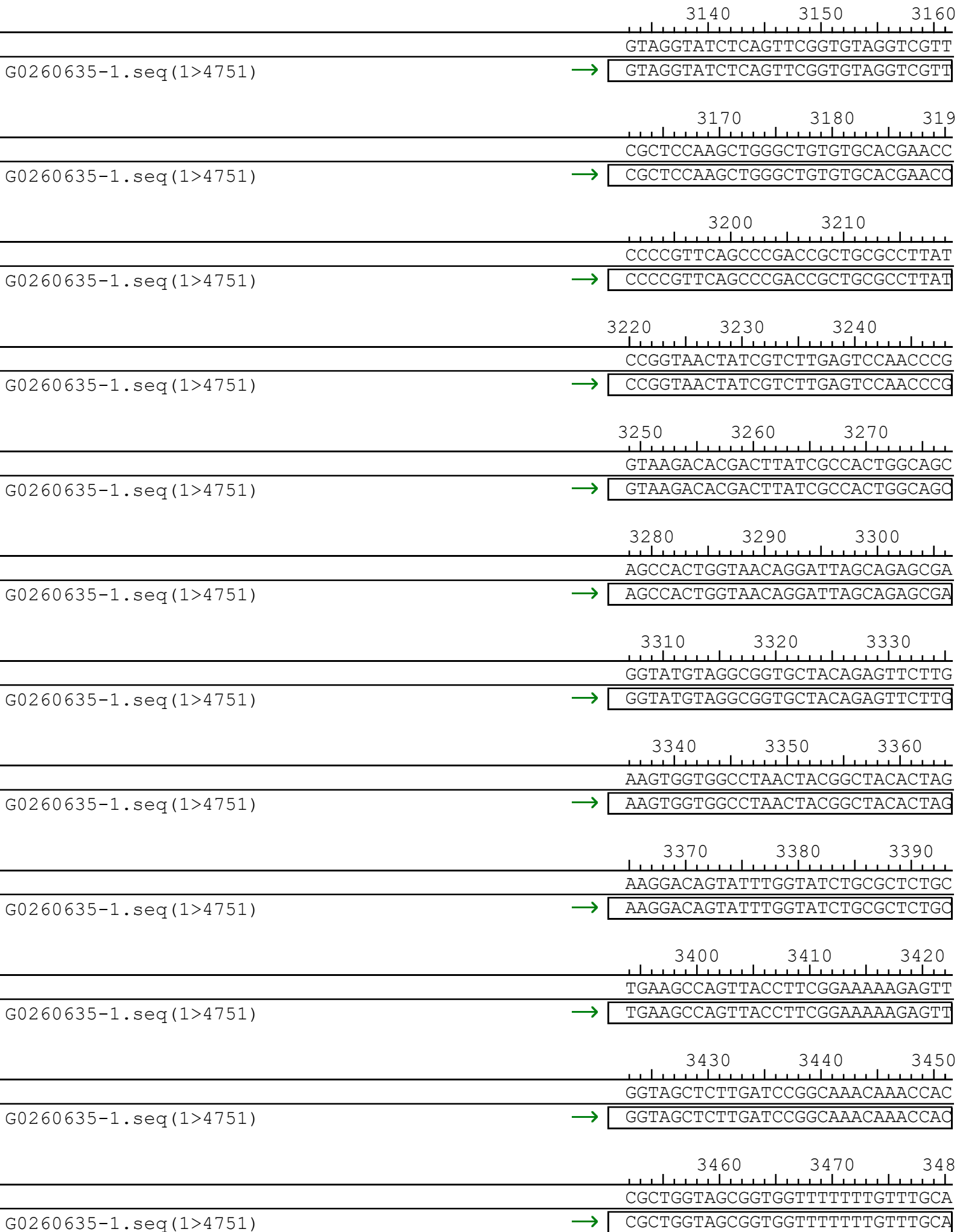
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G0260635-1.seq(1>4751)	→	TCTGACACATGCAGCTCCCGGAGACGGTC		
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G0260635-1.seq(1>4751)	→	ACAGCTTGTCTGTAAGCGGATGCCGGGAG		
		2560	2570	2580
		CAGACAAGCCCGTCAGGGCGCGTCAGCGG		
G0260635-1.seq(1>4751)	→	CAGACAAGCCCGTCAGGGCGCGTCAGCGG		
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G0260635-1.seq(1>4751)	→	ACCCAGTCACGTAGCGATAGCGGAGTGTA		
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		AAATACCGCATCAGGCGCTCTTCCGCTTC		
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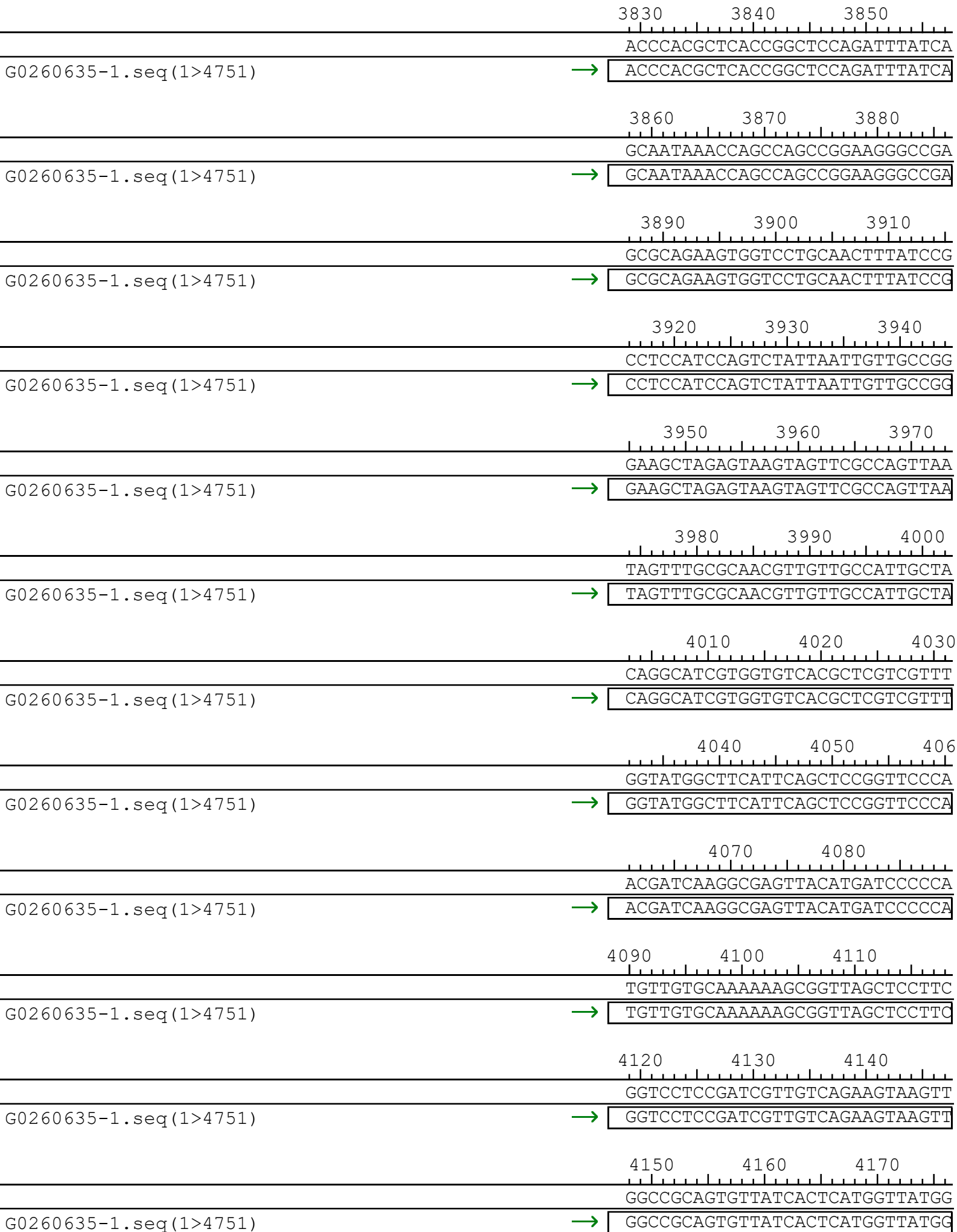
## Project: Untitled.SQD Contig 1



## Project: Untitled.SQD Contig 1



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G0260635-1.seq (1>4751)	→	GGGGTCTGACGCTCAGTGGAACGAAACT			
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G0260635-1.seq (1>4751)	→	AGTTACCAATGCTTAATCAGTGAGGCACC			
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G0260635-1.seq (1>4751)	→	CCATAGTTGCCTGACTCCCCGTCGTGTAG			
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		4410	4420	4430	
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## Project: Untitled.SQD Contig 1

