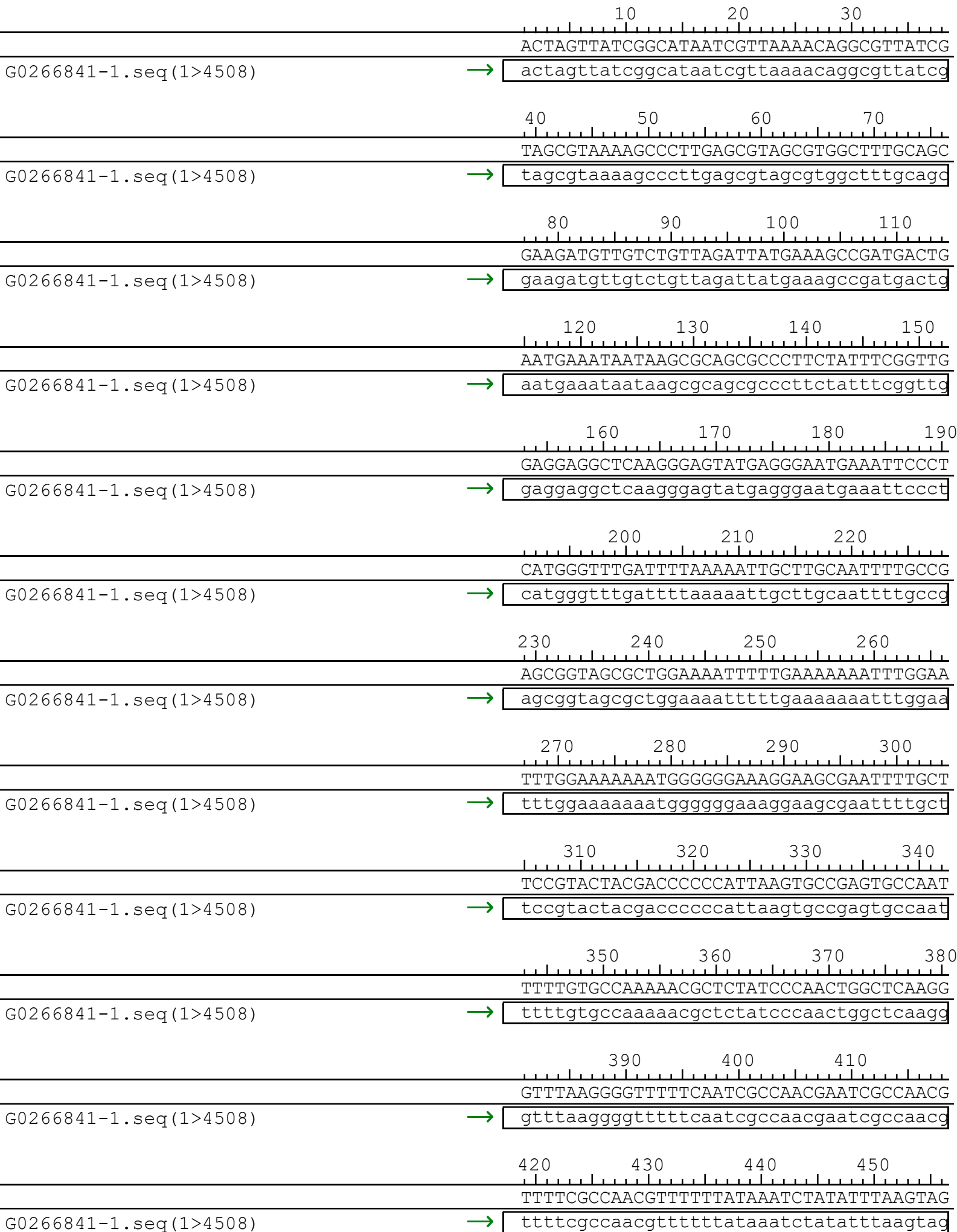
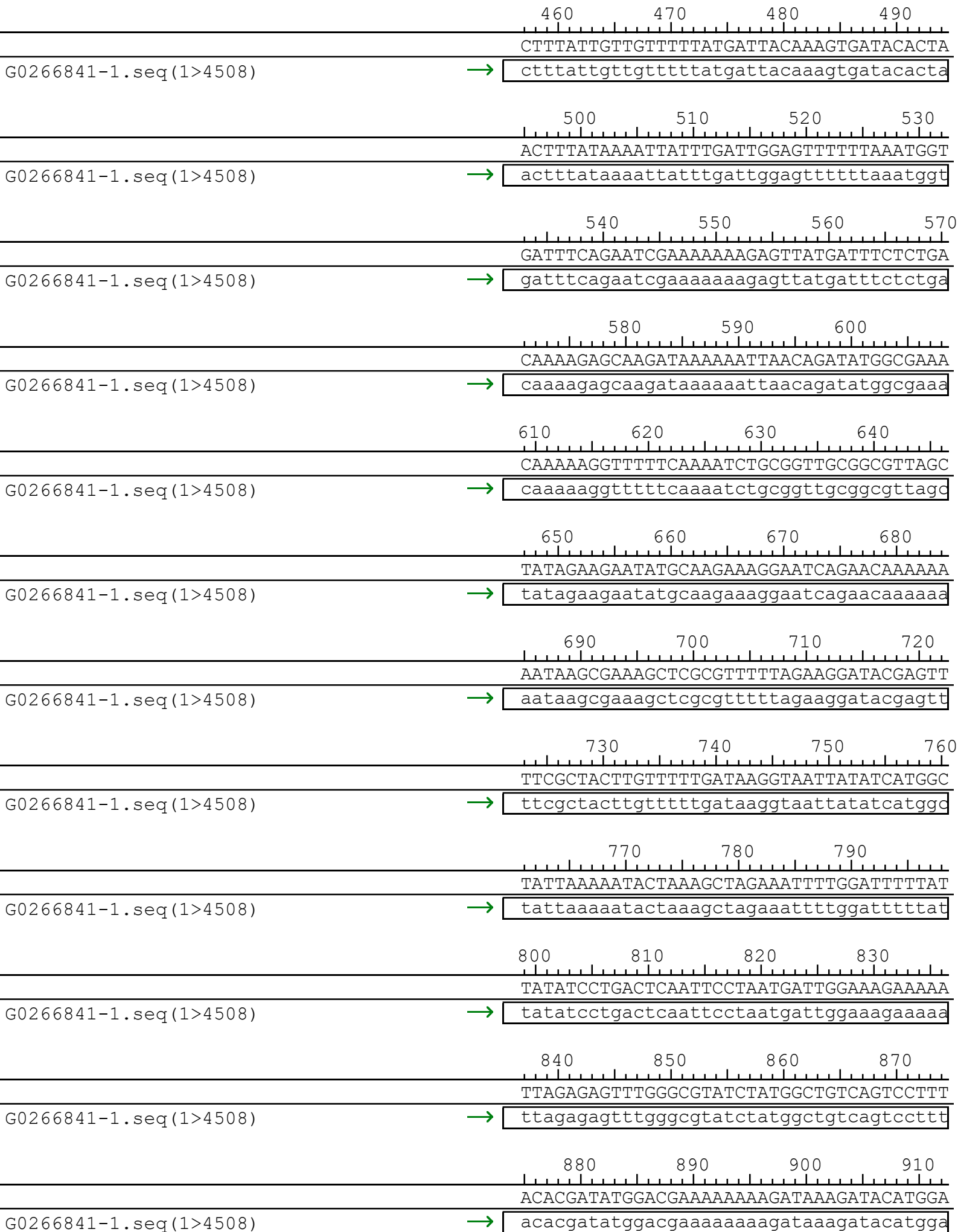


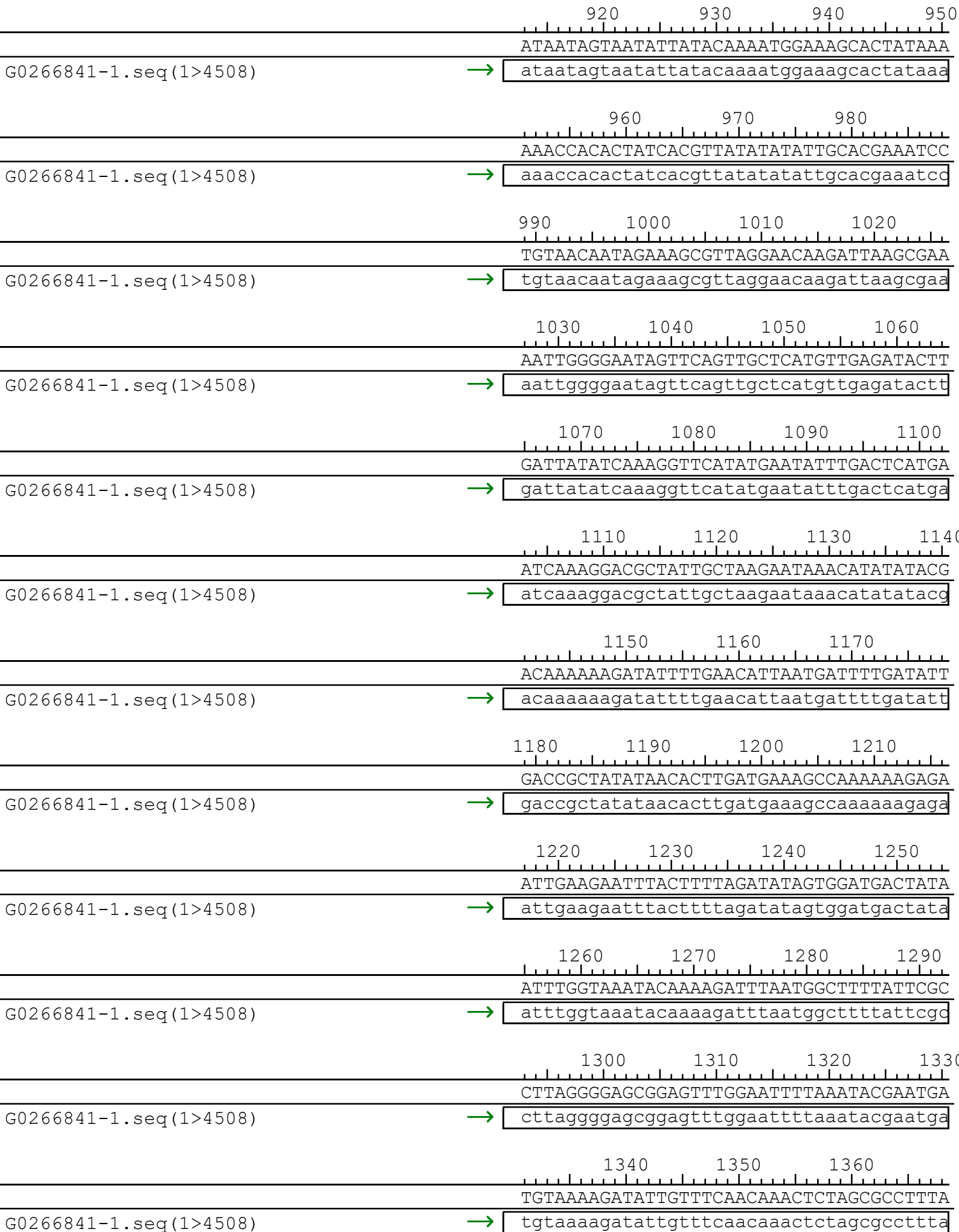
## Project: G0266841-1.SQD Contig 1



## Project: G0266841-1.SQD Contig 1



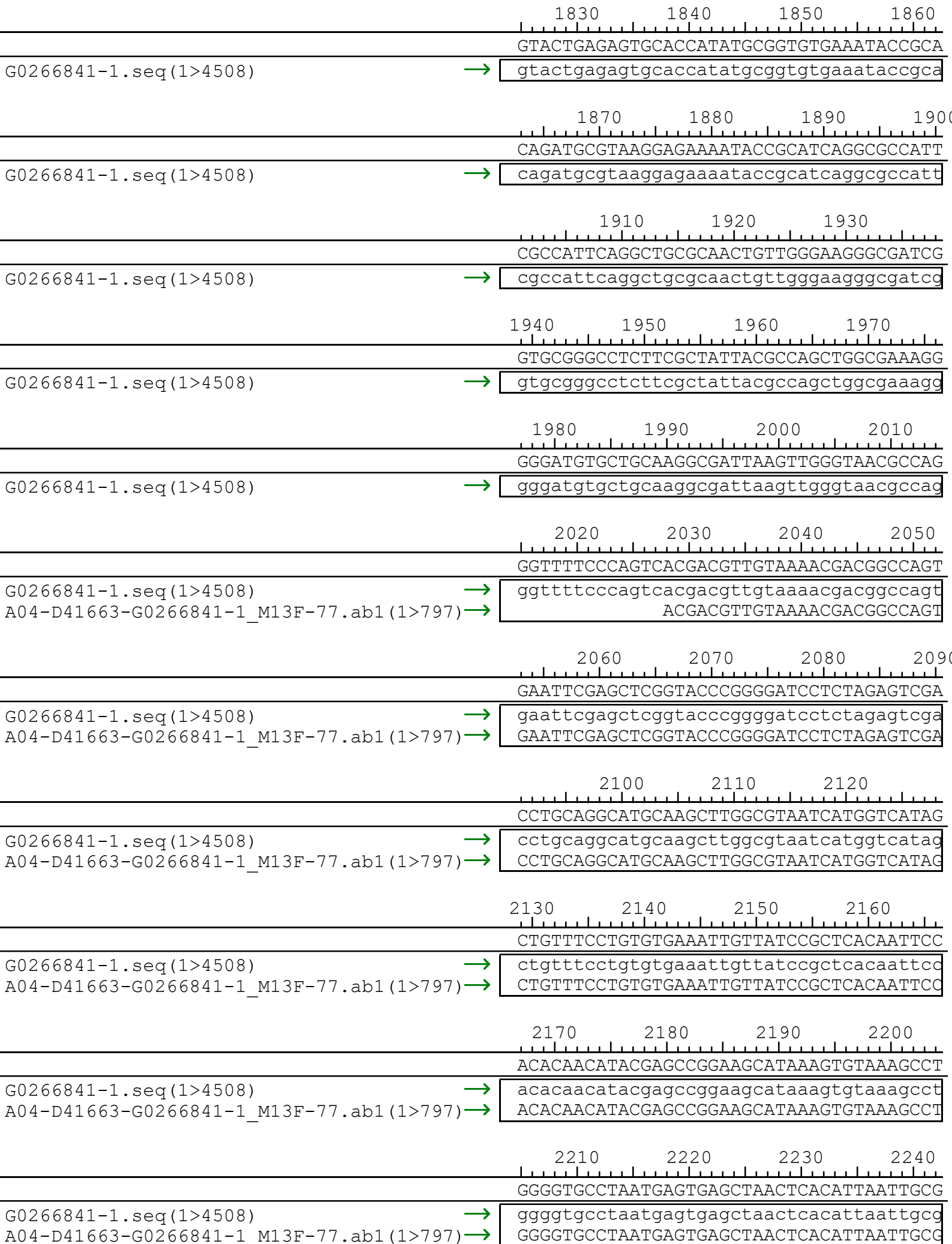
## Project: G0266841-1.SQD Contig 1



## Project: G0266841-1.SQD Contig 1

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G0266841-1.seq(1>4508)	→	1410 1420 1430 1440 GCAAGTTATGCAAAGTTCTTGATGCTGAAACGGGGGA gcaagttatgcaaagttcttgatgctgaaacggggga
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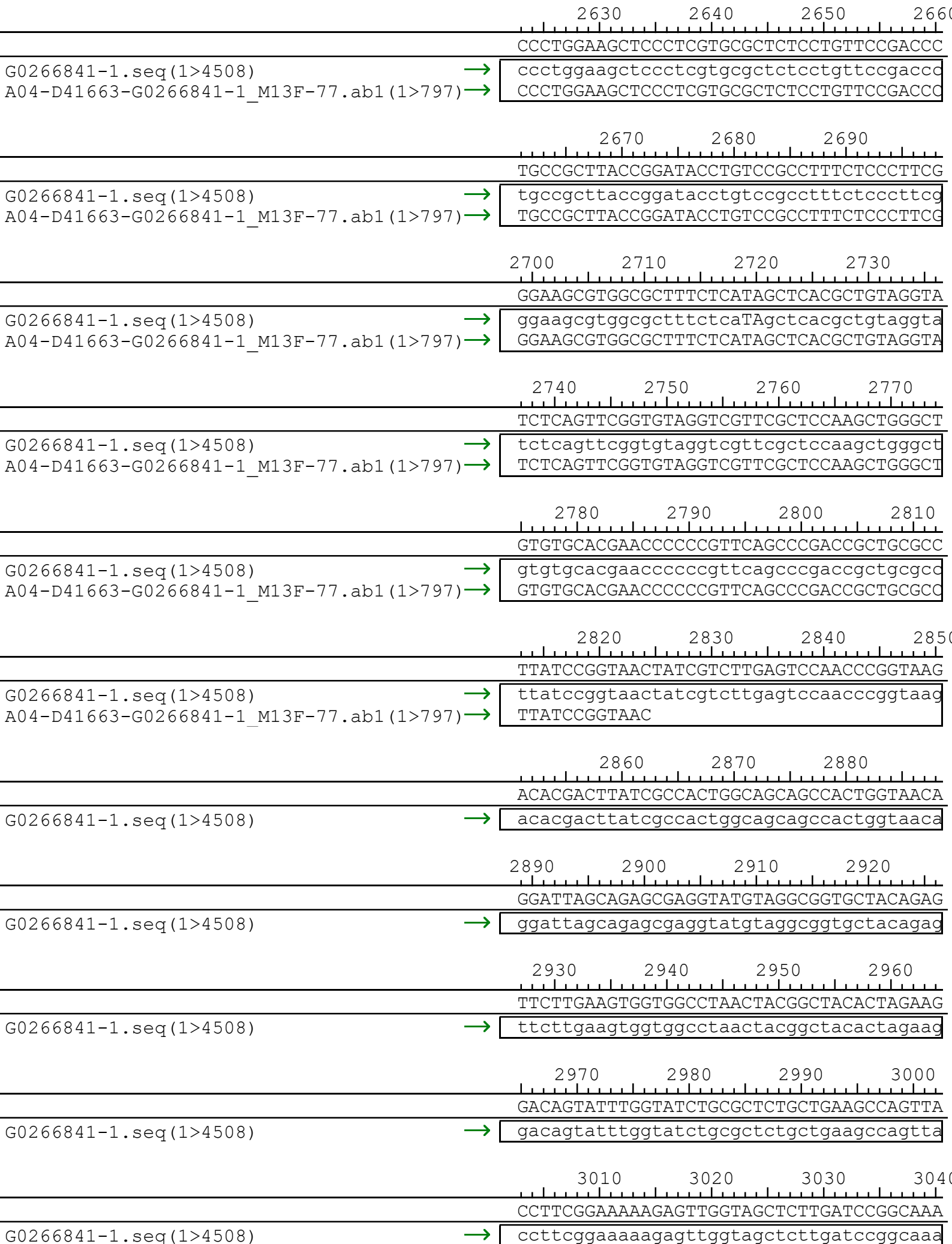
## Project: G0266841-1.SQD Contig 1



## Project: G0266841-1.SQD Contig 1

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		GTGCCAGCTGCATTAATGAATCGGCCAACGCGCGGGGA			
G0266841-1.seq (1>4508)	→	gtgccagctgcattaatgaatcggccaacgCGcgggga			
A04-D41663-G0266841-1_M13F-77.ab1 (1>797)	→	GTGCCAGCTGCATTAATGAATCGGCCAACGCGCGGGGA			
		2320	2330	2340	2350
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G0266841-1.seq (1>4508)	→	gaggCGgtttgcctattgggCGctcttccgcttcctcG			
A04-D41663-G0266841-1_M13F-77.ab1 (1>797)	→	GAGGCGGTTTGCCTATTGGGCGCTCTTCCGCTTCCTCG			
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A04-D41663-G0266841-1_M13F-77.ab1 (1>797)	→	CTCACTGACTCGCTGCGCTCGGTCGTTCCGGCTGCGGCG			
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G0266841-1.seq (1>4508)	→	agCGgtatcagctcactCAAaggCGgtaatacggttat			
A04-D41663-G0266841-1_M13F-77.ab1 (1>797)	→	AGCGGTATCAGCTCACTCAAAGGCGGTAATACGGTTAT			
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		CCACAGAATCAGGGGATAACGCAGGAAAGAACATGTGA			
G0266841-1.seq (1>4508)	→	ccacagaatcaggggataacgcaggaAagaacatgtga			
A04-D41663-G0266841-1_M13F-77.ab1 (1>797)	→	CCACAGAATCAGGGGATAACGCAGGAAAGAACATGTGA			
		2480	2490	2500	
		GCAAAGGCCAGCAAAGGCCAGGAACCGTAAAAAGGC			
G0266841-1.seq (1>4508)	→	gcaaagGCCagcaaagGCCaggaacCGtaaaaaggc			
A04-D41663-G0266841-1_M13F-77.ab1 (1>797)	→	GCAAAGGCCAGCAAAGGCCAGGAACCGTAAAAAGGC			
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G0266841-1.seq (1>4508)	→	cgCGttgctggCGTTTTTTCataggctccgCCCCctg			
A04-D41663-G0266841-1_M13F-77.ab1 (1>797)	→	CGCGTTGCTGGCGTTTTTTCATAGGCTCCGCCCCCTG			
		2550	2560	2570	2580
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G0266841-1.seq (1>4508)	→	acgagcatcacAAAatcgacgctcaagtcagaggtgg			
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## Project: G0266841-1.SQD Contig 1



## Project: G0266841-1.SQD Contig 1

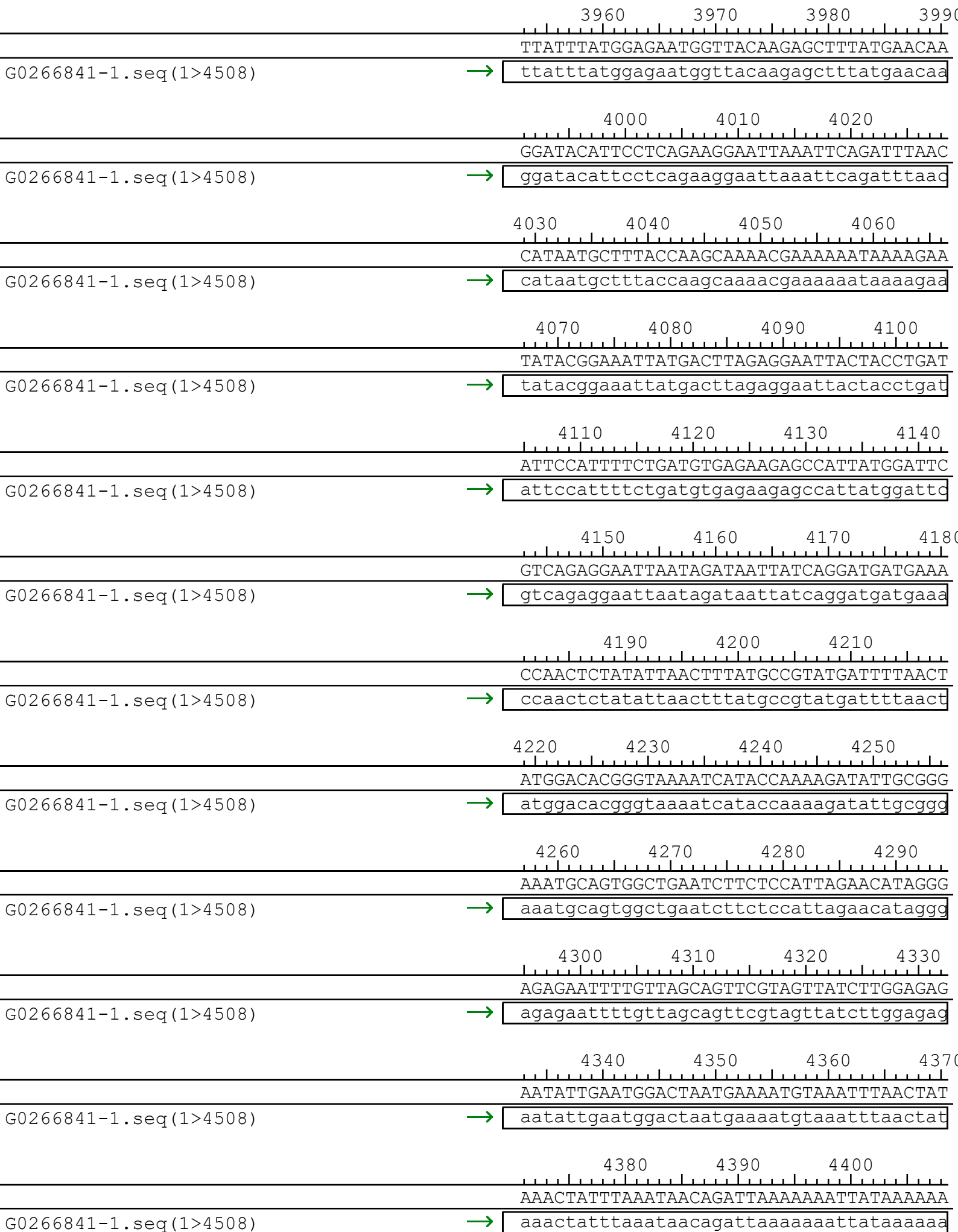
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		3080	3090	3100	3110	GCAGCAGATTACGCGCAGAAAAAAGGATCTCAAGAAG
G0266841-1.seq(1>4508)	→	gcagcagattacgcgcagaaaaaaggatctcaagaag				
		3120	3130	3140	3150	ATCCTTTGATCTTTTCTACGGGGTCTGACGCTCAGTGG
G0266841-1.seq(1>4508)	→	atcctttgatcttttctacggggctgacgctcagtgg				
		3160	3170	3180	3190	AACGAAAACCTCACGTTAAGGGATTTTGGTCATGAGATT
G0266841-1.seq(1>4508)	→	aacgaaaactcacgttaagggattttggatcatgagatt				
		3200	3210	3220	3230	ATCAAAAAGGATCTTCACCTAGATCCTTTTAAATTAAA
G0266841-1.seq(1>4508)	→	atcaaaaaggatcttcacctagatccttttaaatataa				
		3240	3250	3260	AATGAAGTTTTAAATCAATCTAAAGTATATATGAGTAA	
G0266841-1.seq(1>4508)	→	aatgaagttttaaatcaatctaaagtatatatgagtaa				
		3270	3280	3290	3300	ACTTGGTCTGACAGTTACCAATGCTTAATCAGTGAGGC
G0266841-1.seq(1>4508)	→	acttggctctgacagttaccaatgcttaatcagtgaggc				
		3310	3320	3330	3340	ACCTATCTCAGCGATCTGTCTATTTCGTTTCATCCATAG
G0266841-1.seq(1>4508)	→	acctatctcagcgatctgtctatttcgttcatccatag				
		3350	3360	3370	3380	TGCTGACTCCCGTTCGTGTAGATAACTAGTGTTTCGT
G0266841-1.seq(1>4508)	→	ttgctgactcccgctcgtgtagataactagtgttcgt				
		3390	3400	3410	3420	GAATACATGTTATAATAACTATAACTAATAACGTAACG
G0266841-1.seq(1>4508)	→	gaatacatgttataataactataactaataacgtaacg				
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G0266841-1.seq(1>4508)	→	tgactggcaagagatatttttaaaacaatgaataggtt				
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G0266841-1.seq(1>4508)	→	tacacttacttttagttttatggaaatgaaagatcatat				



## Project: G0266841-1.SQD Contig 1

	3500	3510	3520	3530	CATATATAATCTAGAATAAAATTAACTAAAATAATTAT
G0266841-1.seq(1>4508)	→	catatataatctagaataaaattaactaaaataattat			
	3540	3550	3560	3570	TATCTAGATAAAAAATTTAGAAGCCAATGAAATCTATA
G0266841-1.seq(1>4508)	→	tatctagataaaaaatTTAGAAGCCAATGAAATCTATA			
	3580	3590	3600	3610	AATAAACTAAATTAAGTTTATTTAATTAACAACCTATGG
G0266841-1.seq(1>4508)	→	aataaaactaaattaagtttattttaattaacaactatgg			
	3620	3630	3640		ATATAAAATAGGTACTAATCAAAATAGTGAGGAGGATA
G0266841-1.seq(1>4508)	→	atataaaataggtactaatcaaaatagtgaggaggata			
	3650	3660	3670	3680	TATTTGAATACATACGAACAAATTAATAAAGTGAAAAA
G0266841-1.seq(1>4508)	→	tatttgaatacacatACGAACAAATTAATAAAGTGAAAAA			
	3690	3700	3710	3720	AATACTTCGGAAACATTTAAAAAATAACCTTATTGGTA
G0266841-1.seq(1>4508)	→	aatacttcggaaacattTTAAAAAATAACCTTATTGGTA			
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G0266841-1.seq(1>4508)	→	cttacatgtttggatcaggagttgagagtggactaaaa			
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G0266841-1.seq(1>4508)	→	ccaatagtgatcttgactTTTTAGTCGTCGTATCTGA			
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G0266841-1.seq(1>4508)	→	accattgacagatcaaagTAAAGAAATACTTATACAAA			
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G0266841-1.seq(1>4508)	→	aaattagacctatTTTCAAAAAAATAGGAGATAAAAGC			
	3880	3890	3900	3910	AACTTACGATATATTGAATTAACAATTATTATTTCAGCA
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## Project: G0266841-1.SQD Contig 1



4410 4420 4430 4440  
ATTGAAAAAATGGTGGAAACACTTTTTTCAATTTTTTT  
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4450 4460 4470 4480  
GTTTTATTATTTAATATTTGGGAAATATTCATTCTAAT  
G0266841-1.seq(1>4508) → gttttattatttaaatatttgggaaatattcattctaatt

4490 4500  
TGGTAATCAGATTTTAGAAAAC  
G0266841-1.seq(1>4508) → tggtaatcagatTTTAGAAAAC