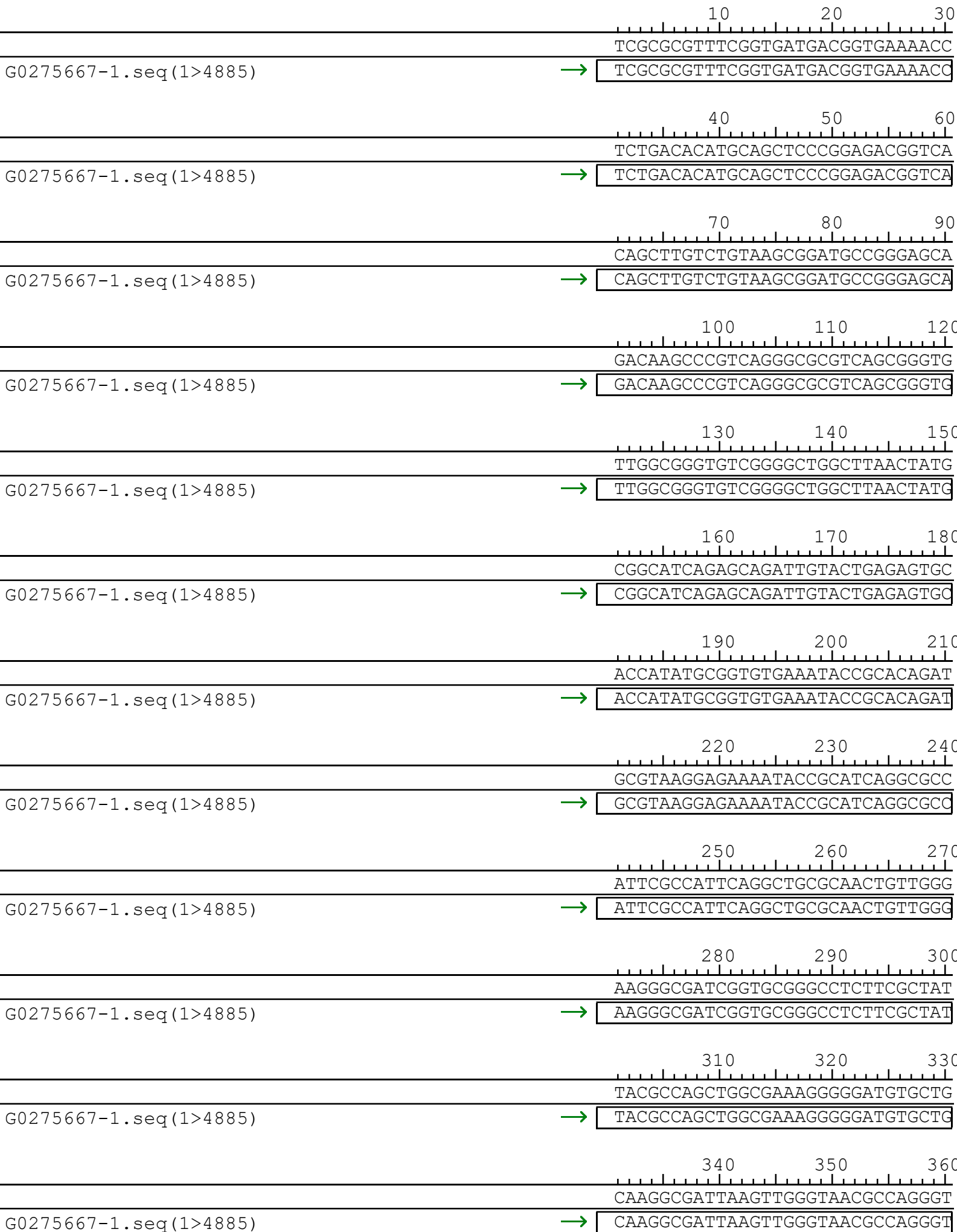
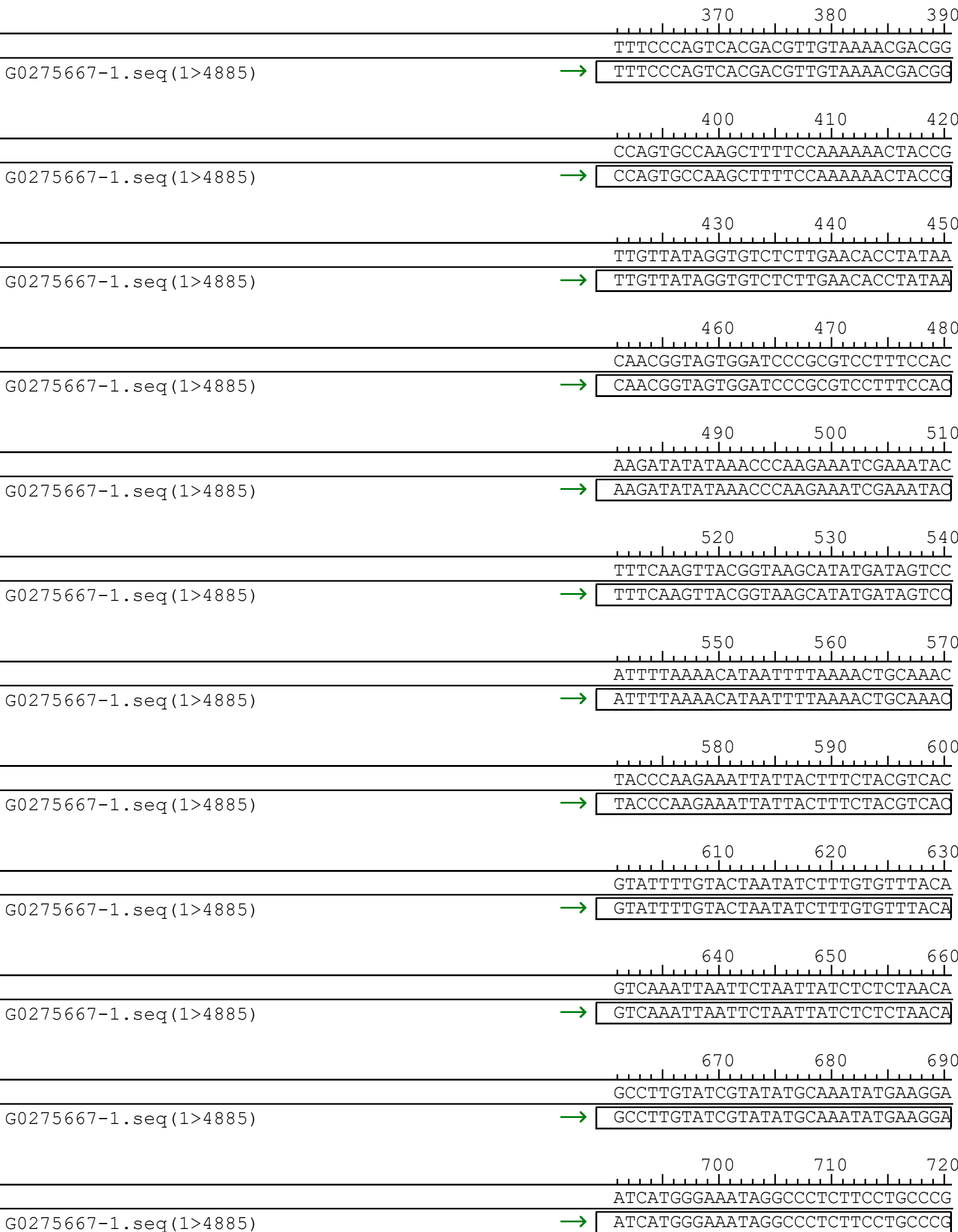
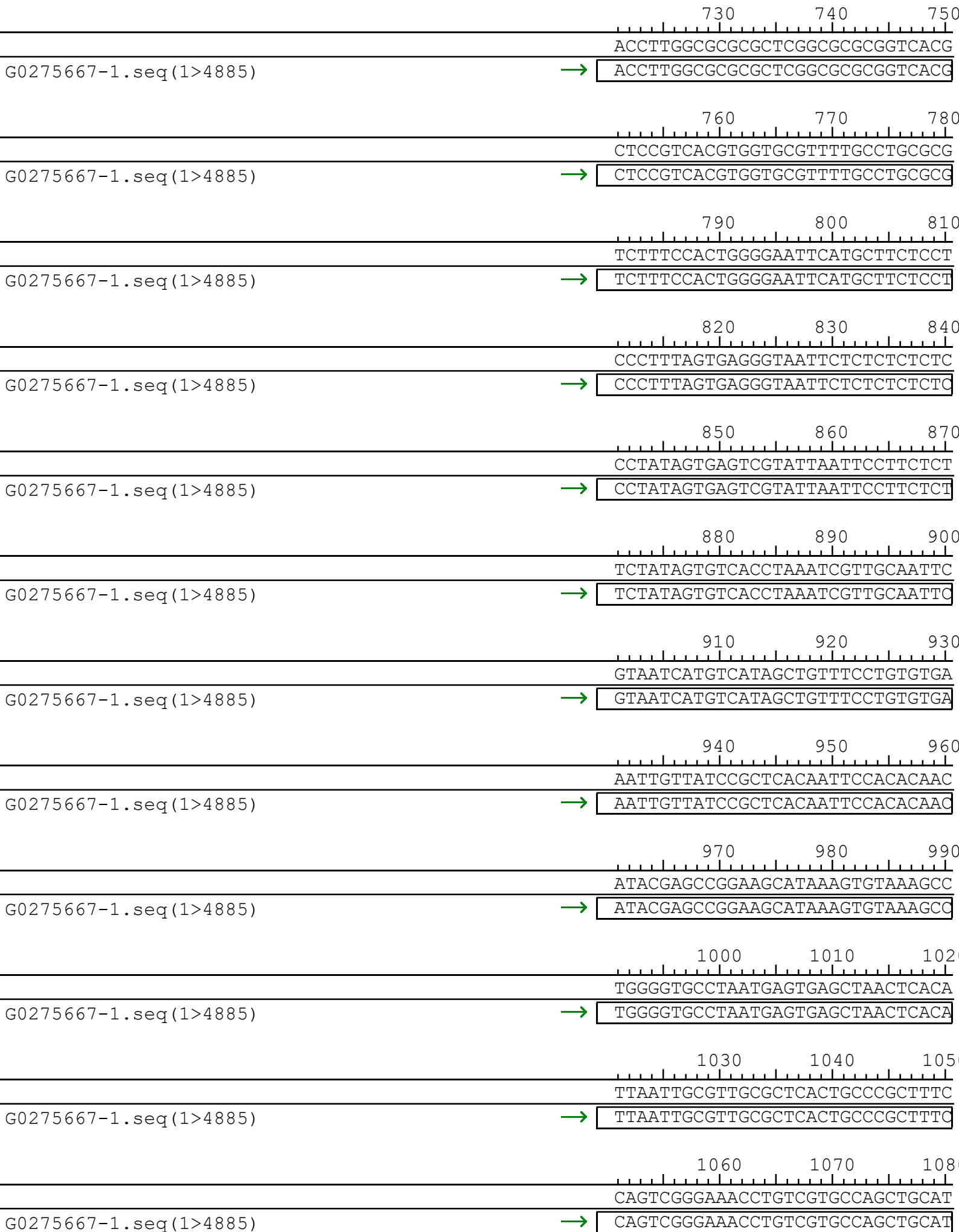


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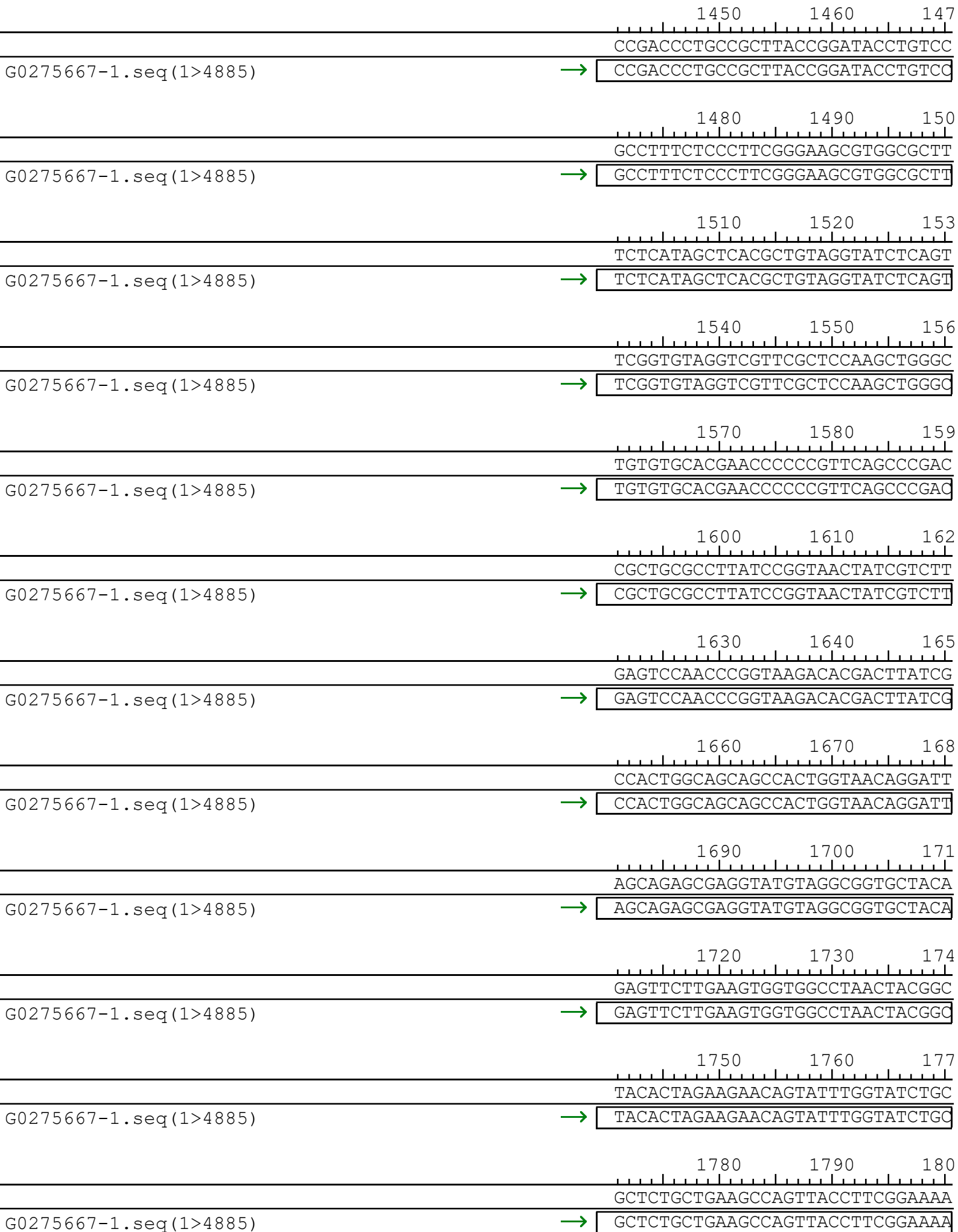
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		1090	1100	111
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G0275667-1.seq(1>4885)	→	TAATGAATCGGCCAACGCGCGGGGAGAGGC		
		1120	1130	114
		GGTTTGCCTATTGGGCGCTCTTCCGCTTCC		
G0275667-1.seq(1>4885)	→	GGTTTGCCTATTGGGCGCTCTTCCGCTTCC		
		1150	1160	117
		TCGCTCACTGACTCGCTGCGCTCGGTCGTT		
G0275667-1.seq(1>4885)	→	TCGCTCACTGACTCGCTGCGCTCGGTCGTT		
		1180	1190	120
		CGGCTGCGGCGAGCGGTATCAGCTCACTCA		
G0275667-1.seq(1>4885)	→	CGGCTGCGGCGAGCGGTATCAGCTCACTCA		
		1210	1220	123
		AAGGCGGTAATACGGTTATCCACAGAATCA		
G0275667-1.seq(1>4885)	→	AAGGCGGTAATACGGTTATCCACAGAATCA		
		1240	1250	126
		GGGGATAACGCAGGAAAGAACATGTGAGCA		
G0275667-1.seq(1>4885)	→	GGGGATAACGCAGGAAAGAACATGTGAGCA		
		1270	1280	129
		AAAGGCCAGCAAAAGGCCAGGAACCGTAAA		
G0275667-1.seq(1>4885)	→	AAAGGCCAGCAAAAGGCCAGGAACCGTAAA		
		1300	1310	132
		AAGGCCGCGTTGCTGGCGTTTTTCCATAGG		
G0275667-1.seq(1>4885)	→	AAGGCCGCGTTGCTGGCGTTTTTCCATAGG		
		1330	1340	135
		CTCCGCCCCCTGACGAGCATCACAAAAT		
G0275667-1.seq(1>4885)	→	CTCCGCCCCCTGACGAGCATCACAAAAT		
		1360	1370	138
		CGACGCTCAAGTCAGAGGTGGCGAAACCCG		
G0275667-1.seq(1>4885)	→	CGACGCTCAAGTCAGAGGTGGCGAAACCCG		
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G0275667-1.seq(1>4885)	→	ACAGGACTATAAAGATAACCAGGCGTTTCCC		
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		CCTGGAAGCTCCCTCGTGCGCTCTCCTGTT		
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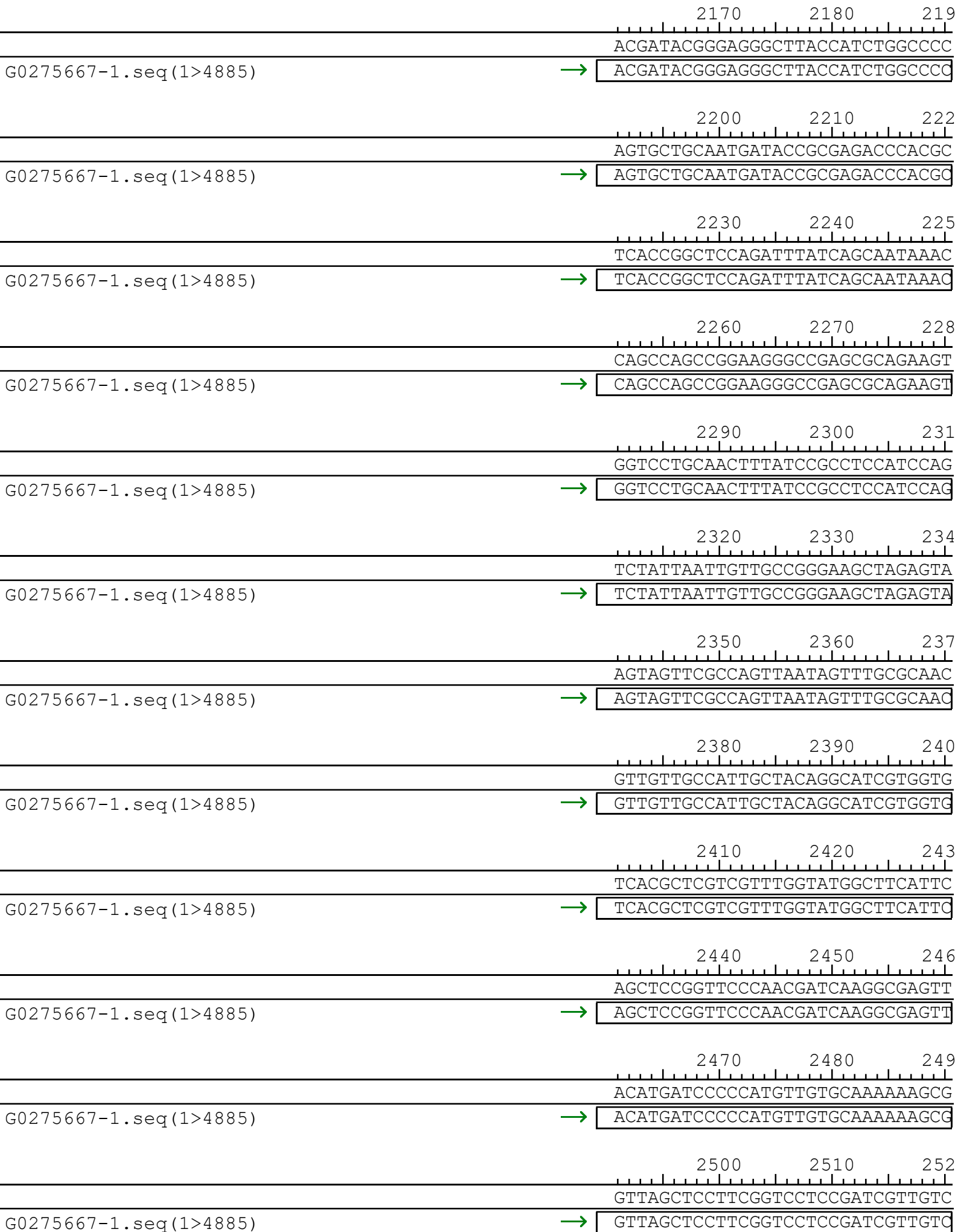
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		AGAGTTGGTAGCTCTTGATCCGGCAAAAAA		
G0275667-1.seq(1>4885)	→	AGAGTTGGTAGCTCTTGATCCGGCAAAAAA		
		1840	1850	186
		ACCACCGCTGGTAGCGGTGGTTTTTTTGT		
G0275667-1.seq(1>4885)	→	ACCACCGCTGGTAGCGGTGGTTTTTTTGT		
		1870	1880	189
		TGCAAGCAGCAGATTACGCGCAGAAAAAA		
G0275667-1.seq(1>4885)	→	TGCAAGCAGCAGATTACGCGCAGAAAAAA		
		1900	1910	192
		GGATCTCAAGAAGATCCTTTGATCTTTTCT		
G0275667-1.seq(1>4885)	→	GGATCTCAAGAAGATCCTTTGATCTTTTCT		
		1930	1940	195
		ACGGGGTCTGACGCTCAGTGGAAACGAAAAC		
G0275667-1.seq(1>4885)	→	ACGGGGTCTGACGCTCAGTGGAAACGAAAAC		
		1960	1970	198
		TCACGTTAAGGGATTTTGGTCATGAGATTA		
G0275667-1.seq(1>4885)	→	TCACGTTAAGGGATTTTGGTCATGAGATTA		
		1990	2000	201
		TCAAAAAGGATCTTCACCTAGATCCTTTTA		
G0275667-1.seq(1>4885)	→	TCAAAAAGGATCTTCACCTAGATCCTTTTA		
		2020	2030	204
		AATTAAAAATGAAGTTTTAAATCAATCTAA		
G0275667-1.seq(1>4885)	→	AATTAAAAATGAAGTTTTAAATCAATCTAA		
		2050	2060	207
		AGTATATATGAGTAACTTGGTCTGACAGT		
G0275667-1.seq(1>4885)	→	AGTATATATGAGTAACTTGGTCTGACAGT		
		2080	2090	210
		TACCAATGCTTAATCAGTGAGGCACCTATC		
G0275667-1.seq(1>4885)	→	TACCAATGCTTAATCAGTGAGGCACCTATC		
		2110	2120	213
		TCAGCGATCTGTCTATTTTCGTTTCATCCATA		
G0275667-1.seq(1>4885)	→	TCAGCGATCTGTCTATTTTCGTTTCATCCATA		
		2140	2150	216
		GTTGCCTGACTCCCCGTCGTGTAGATAACT		
G0275667-1.seq(1>4885)	→	GTTGCCTGACTCCCCGTCGTGTAGATAACT		

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		2530	2540	255
		AGAAGTAAGTTGGCCGCAGTGTTATCACTC		
G0275667-1.seq (1>4885)	→	AGAAGTAAGTTGGCCGCAGTGTTATCACTC		
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		ATGGTTATGGCAGCACTGCATAATTCTCTT		
G0275667-1.seq (1>4885)	→	ATGGTTATGGCAGCACTGCATAATTCTCTT		
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		ACTGTCATGCCATCCGTAAGATGCTTTTCT		
G0275667-1.seq (1>4885)	→	ACTGTCATGCCATCCGTAAGATGCTTTTCT		
		2620	2630	264
		GTGACTGGTGAGTACTCAACCAAGTCATTC		
G0275667-1.seq (1>4885)	→	GTGACTGGTGAGTACTCAACCAAGTCATTC		
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G0275667-1.seq (1>4885)	→	TGAGAATAGTGTATGCGGCGACCGAGTTGC		
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		TCTTGCCCGGCGTCAATACGGGATAATACC		
G0275667-1.seq (1>4885)	→	TCTTGCCCGGCGTCAATACGGGATAATACC		
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		GCGCCACATAGCAGAACTTTAAAAGTGCTC		
G0275667-1.seq (1>4885)	→	GCGCCACATAGCAGAACTTTAAAAGTGCTC		
		2740	2750	276
		ATCATTGGAAAACGTTCTTCGGGGCGAAAA		
G0275667-1.seq (1>4885)	→	ATCATTGGAAAACGTTCTTCGGGGCGAAAA		
		2770	2780	279
		CTCTCAAGGATCTTACCGCTGTTGAGATCC		
G0275667-1.seq (1>4885)	→	CTCTCAAGGATCTTACCGCTGTTGAGATCC		
		2800	2810	282
		AGTTCGATGTAACCCACTCGTGCACCCAAC		
G0275667-1.seq (1>4885)	→	AGTTCGATGTAACCCACTCGTGCACCCAAC		
		2830	2840	285
		TGATCTTCAGCATCTTTTACTTTTACCAGC		
G0275667-1.seq (1>4885)	→	TGATCTTCAGCATCTTTTACTTTTACCAGC		
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		GTTTCTGGGTGAGCAAAAACAGGAAGGCAA		
G0275667-1.seq (1>4885)	→	GTTTCTGGGTGAGCAAAAACAGGAAGGCAA		

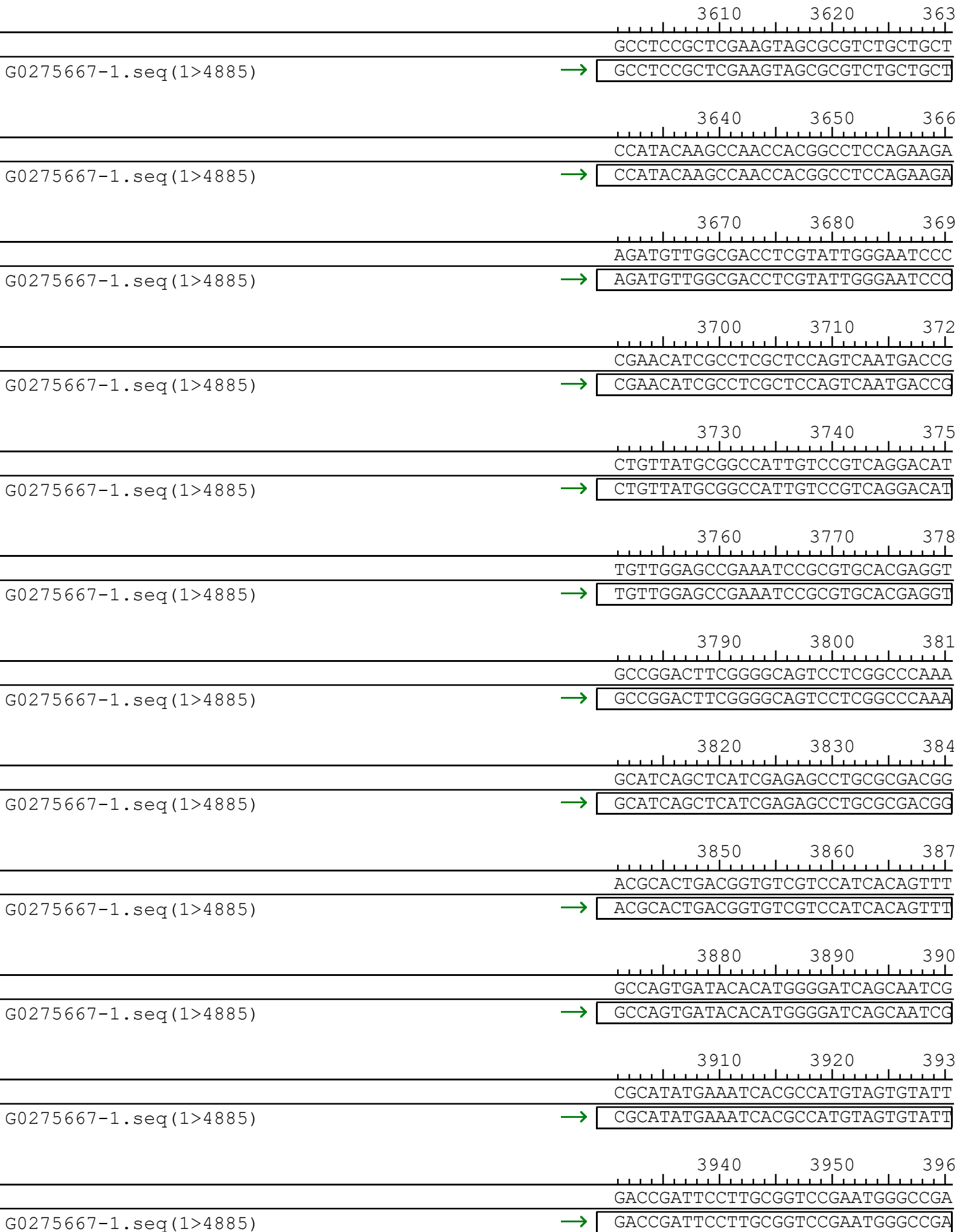
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		AATGCCGCAAAAAGGGAATAAGGGCGACA		
G0275667-1.seq (1>4885)	→	AATGCCGCAAAAAGGGAATAAGGGCGACA		
		2920	2930	294
		CGGAAATGTTGAATACTCATACTCTTCCTT		
G0275667-1.seq (1>4885)	→	CGGAAATGTTGAATACTCATACTCTTCCTT		
		2950	2960	297
		TTTCAATATTCAGGCTTACCTTTATGCTTC		
G0275667-1.seq (1>4885)	→	TTTCAATATTCAGGCTTACCTTTATGCTTC		
		2980	2990	300
		CGGCTCGTATGTTGTGTGGAATTGTGAGCG		
G0275667-1.seq (1>4885)	→	CGGCTCGTATGTTGTGTGGAATTGTGAGCG		
		3010	3020	303
		GATAACAATTTACACAGGAAACAGCTATG		
G0275667-1.seq (1>4885)	→	GATAACAATTTACACAGGAAACAGCTATG		
		3040	3050	306
		ACCATGATTACGCCAAGCTCTAGCTAGAGG		
G0275667-1.seq (1>4885)	→	ACCATGATTACGCCAAGCTCTAGCTAGAGG		
		3070	3080	309
		TCGACGGTATACAGACATGATAAGATACAT		
G0275667-1.seq (1>4885)	→	TCGACGGTATACAGACATGATAAGATACAT		
		3100	3110	312
		TGATGAGTTTGGACAAACCACAACCTAGAAT		
G0275667-1.seq (1>4885)	→	TGATGAGTTTGGACAAACCACAACCTAGAAT		
		3130	3140	315
		GCAGTGAAAAAATGCTTTATTTGTGAAAT		
G0275667-1.seq (1>4885)	→	GCAGTGAAAAAATGCTTTATTTGTGAAAT		
		3160	3170	318
		TTGTGATGCTATTGCTTTATTTGTAACCAT		
G0275667-1.seq (1>4885)	→	TTGTGATGCTATTGCTTTATTTGTAACCAT		
		3190	3200	321
		TATAAGCTGCAATAAACAAGTTGGGGTGGG		
G0275667-1.seq (1>4885)	→	TATAAGCTGCAATAAACAAGTTGGGGTGGG		
		3220	3230	324
		CGAAGAACTCCAGCATGAGATCCCCGCGCT		
G0275667-1.seq (1>4885)	→	CGAAGAACTCCAGCATGAGATCCCCGCGCT		

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		3250	3260	327
		GGAGGATCATCCAGCCGGCGTCCCGGAAAA		
G0275667-1.seq (1>4885)	→	GGAGGATCATCCAGCCGGCGTCCCGGAAAA		
		3280	3290	330
		CGATTCCGAAGCCCAACCTTTCATAGAAGG		
G0275667-1.seq (1>4885)	→	CGATTCCGAAGCCCAACCTTTCATAGAAGG		
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		CGGCGGTGGAATCGAAATCTCGTAGCACGT		
G0275667-1.seq (1>4885)	→	CGGCGGTGGAATCGAAATCTCGTAGCACGT		
		3340	3350	336
		GCTATTCCCTTTGCCCTCGGACGAGTGCTGG		
G0275667-1.seq (1>4885)	→	GCTATTCCCTTTGCCCTCGGACGAGTGCTGG		
		3370	3380	339
		GGCGTCGGTTTCCACTATCGGCGAGTACTT		
G0275667-1.seq (1>4885)	→	GGCGTCGGTTTCCACTATCGGCGAGTACTT		
		3400	3410	342
		CTACACAGCCATCGGTCCAGACGGCCGCGC		
G0275667-1.seq (1>4885)	→	CTACACAGCCATCGGTCCAGACGGCCGCGC		
		3430	3440	345
		TTCTGCGGGCGATTTGTGTACGCCGACAG		
G0275667-1.seq (1>4885)	→	TTCTGCGGGCGATTTGTGTACGCCGACAG		
		3460	3470	348
		TCCCGGCTCCGGATCGGACGATTGCGTCGC		
G0275667-1.seq (1>4885)	→	TCCCGGCTCCGGATCGGACGATTGCGTCGC		
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		ATCGACCCTGCGCCCAAGCTGCATCATCGA		
G0275667-1.seq (1>4885)	→	ATCGACCCTGCGCCCAAGCTGCATCATCGA		
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		AATTGCCGTCAACCAAGCTCTGATAGAGTT		
G0275667-1.seq (1>4885)	→	AATTGCCGTCAACCAAGCTCTGATAGAGTT		
		3550	3560	357
		GGTCAAGACCAATGCGGAGCATATACGCC		
G0275667-1.seq (1>4885)	→	GGTCAAGACCAATGCGGAGCATATACGCC		
		3580	3590	360
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G0275667-1.seq (1>4885)	→	GGAGCCGCGGCGATCCTGCAAGCTCCGGAT		

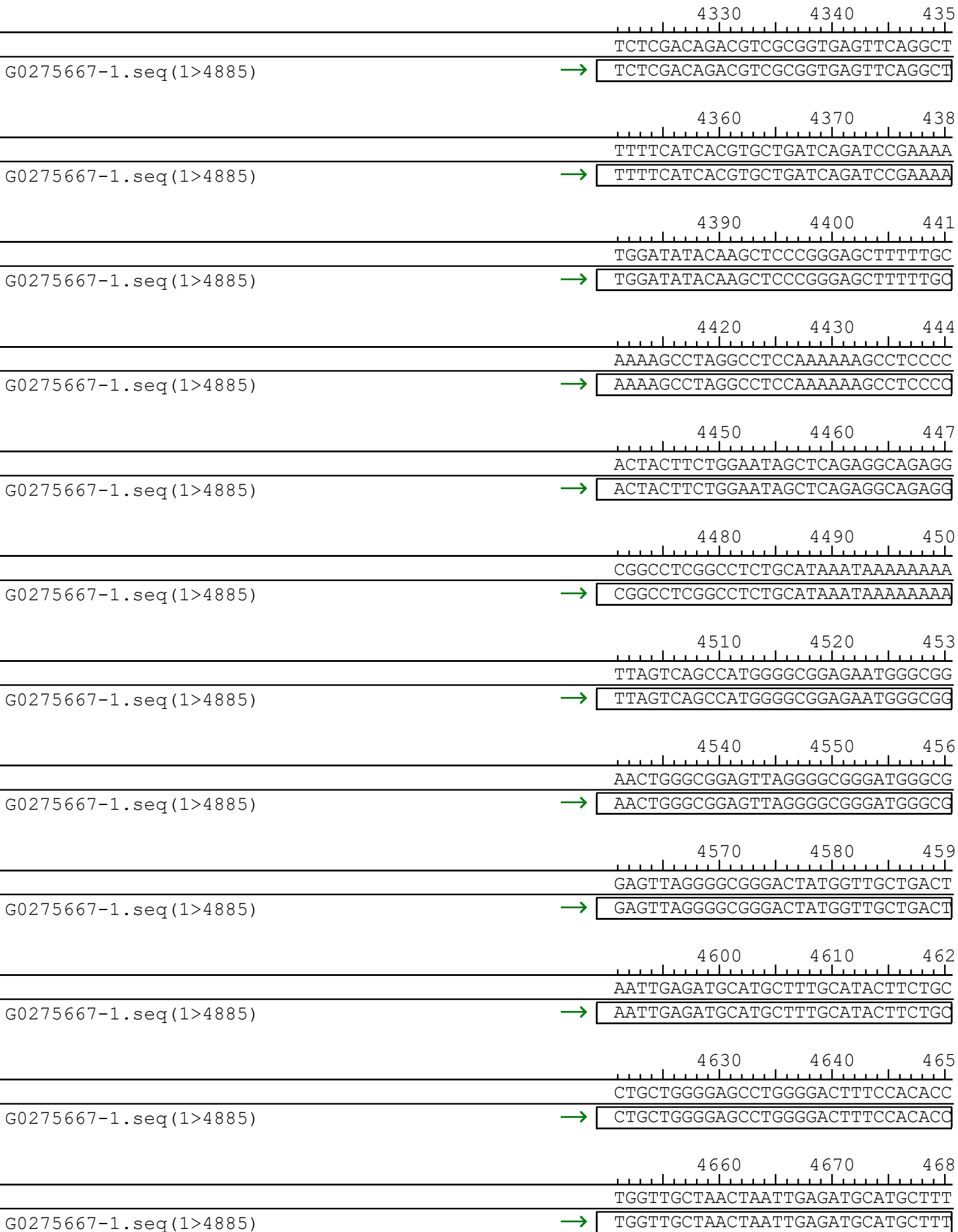
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G0275667-1.seq (1>4885)	→	ACCCGCTCGTCTGGCTAAGATCGGCCGCAG		
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G0275667-1.seq (1>4885)	→	CGATCGCATCCATGGCCTCCGCGACCGGCT		
		4030	4040	405
		GCAGAACAGCGGGCAGTTCGGTTTCAGGCA		
G0275667-1.seq (1>4885)	→	GCAGAACAGCGGGCAGTTCGGTTTCAGGCA		
		4060	4070	408
		GGTCTTGCAACGTGACACCCTGTGCACGGC		
G0275667-1.seq (1>4885)	→	GGTCTTGCAACGTGACACCCTGTGCACGGC		
		4090	4100	411
		GGGAGATGCAATAGGTCAGGCTCTCGCTGA		
G0275667-1.seq (1>4885)	→	GGGAGATGCAATAGGTCAGGCTCTCGCTGA		
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		ATTCCCCAATGTCAAGCACTTCCGGAATCG		
G0275667-1.seq (1>4885)	→	ATTCCCCAATGTCAAGCACTTCCGGAATCG		
		4150	4160	417
		GGAGCGCGGCCGATGCAAAGTGCCGATAAA		
G0275667-1.seq (1>4885)	→	GGAGCGCGGCCGATGCAAAGTGCCGATAAA		
		4180	4190	420
		CATAACGATCTTTGTAGAAACCATCGGCGC		
G0275667-1.seq (1>4885)	→	CATAACGATCTTTGTAGAAACCATCGGCGC		
		4210	4220	423
		AGCTATTTACCCGCAGGACATATCCACGCC		
G0275667-1.seq (1>4885)	→	AGCTATTTACCCGCAGGACATATCCACGCC		
		4240	4250	426
		CTCCTACATCGAAGCTGAAAGCACGAGATT		
G0275667-1.seq (1>4885)	→	CTCCTACATCGAAGCTGAAAGCACGAGATT		
		4270	4280	429
		CTTCGCCCTCCGAGAGCTGCATCAGGTCGG		
G0275667-1.seq (1>4885)	→	CTTCGCCCTCCGAGAGCTGCATCAGGTCGG		
		4300	4310	432
		AGACGCTGTCGAACTTTTCGATCAGAACT		
G0275667-1.seq (1>4885)	→	AGACGCTGTCGAACTTTTCGATCAGAACT		

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		4690	4700	471
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G0275667-1.seq (1>4885)	→	GCATACTTCTGCCTGCTGGAATATTATTGA		
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		AGCATTATCAGGGTTATTGTCTCATGAGC		
G0275667-1.seq (1>4885)	→	AGCATTATCAGGGTTATTGTCTCATGAGC		
		4750	4760	477
		GGATACATATTTGAATGTATTTAGAAAAAT		
G0275667-1.seq (1>4885)	→	GGATACATATTTGAATGTATTTAGAAAAAT		
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531)	→	AAAT		
		4780	4790	480
		AAACAAATAGGGGTTCGCGCACATTTCCC		
G0275667-1.seq (1>4885)	→	AAACAAATAGGGGTTCGCGCACATTTCCC		
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531)	→	AAACAAATAGGGGTTCGCGCACATTTCCC		
		4810	4820	483
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G0275667-1.seq (1>4885)	→	CGAAAAGTGCCACCTGACGTCTAAGAAACC		
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531)	→	CGAAAAGTGCCACCTGACGTCTAAGAAACC		
		4840	4850	486
		ATTATTATCATGACATTAACCTATAAAAAT		
G0275667-1.seq (1>4885)	→	ATTATTATCATGACATTAACCTATAAAAAT		
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531)	→	ATTATTATCATGACATTAACCTATAAAAAT		
		4870	4880	489
		AGGCGTATCACGAGGCCCTTTCGTCTCGCG		
G0275667-1.seq (1>4885)	→	AGGCGTATCACGAGGCCCTTTCGTCTCGCG		
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531)	→	AGGCGTATCACGAGGCCCTTTCGTCTCGCG		
		4900	4910	492
		CGTTTCGGTGATGACGGTGAAAACCTCTGA		
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531)	→	CGTTTCGGTGATGACGGTGAAAACCTCTGA		
		4930	4940	495
		CACATGCAGCTCCCGGAGACGGTCACAGCT		
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531)	→	CACATGCAGCTCCCGGAGACGGTCACAGCT		
		4960	4970	498
		TGTCTGTAAGCGGATGCCGGGAGCAGACAA		
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531)	→	TGTCTGTAAGCGGATGCCGGGAGCAGACAA		
		4990	5000	501
		GCCCGTCAGGGCGCGTCAGCGGGTGTGGC		
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531)	→	GCCCGTCAGGGCGCGTCAGCGGGTGTGGC		

5020 5030 504
GGGTGTCGGGGCTGGCTTAACTATGCGGCA
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531) → GGGTGTCGGGGCTGGCTTAACTATGCGGCA

5050 5060 507
TCAGAGCAGATTGTACTGAGAGTGCACCAT
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531) → TCAGAGCAGATTGTACTGAGAGTGCACCAT

5080 5090 510
ATGCGGTGTGAAATACCGCACAGATGCGTA
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531) → ATGCGGTGTGAAATACCGCACAGATGCGTA

5110 5120 513
AGGAGAAAATACCGCATCAGGCGCCATTCC
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531) → AGGAGAAAATACCGCATCAGGCGCCATTCC

5140 5150 516
CCATTCAGGCTGCGCAACTGTTGGGAAGGG
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531) → CCATTCAGGCTGCGCAACTGTTGGGAAGGG

5170 5180 519
CGATCGGTGCGGGCCTCTTCGCTATTACGC
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531) → CGATCGGTGCGGGCCTCTTCGCTATTACGC

5200 5210 522
CAGCTGGCGAAAGGGGGATGTGCTGCAAGG
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531) → CAGCTGGCGAAAGGGGGATGTGCTGCAAGG

5230 5240 525
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A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531) → CGATTAAGTTGGGTAACGCCAGGGTTTTCC

5260 5270 528
CAGTCACGACGTTGTAAAACGACGGCCAGT
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531) → CAGTCACGACGTTGTAAAACGACGGCCAGT

5290
GCCAAGCTTTTCCAAAA
A08-D46341-G0275667-1_PUC57-1.8K-SEQF.ab1 (1>531) → GCCAAGCTTTTCCAAAA