

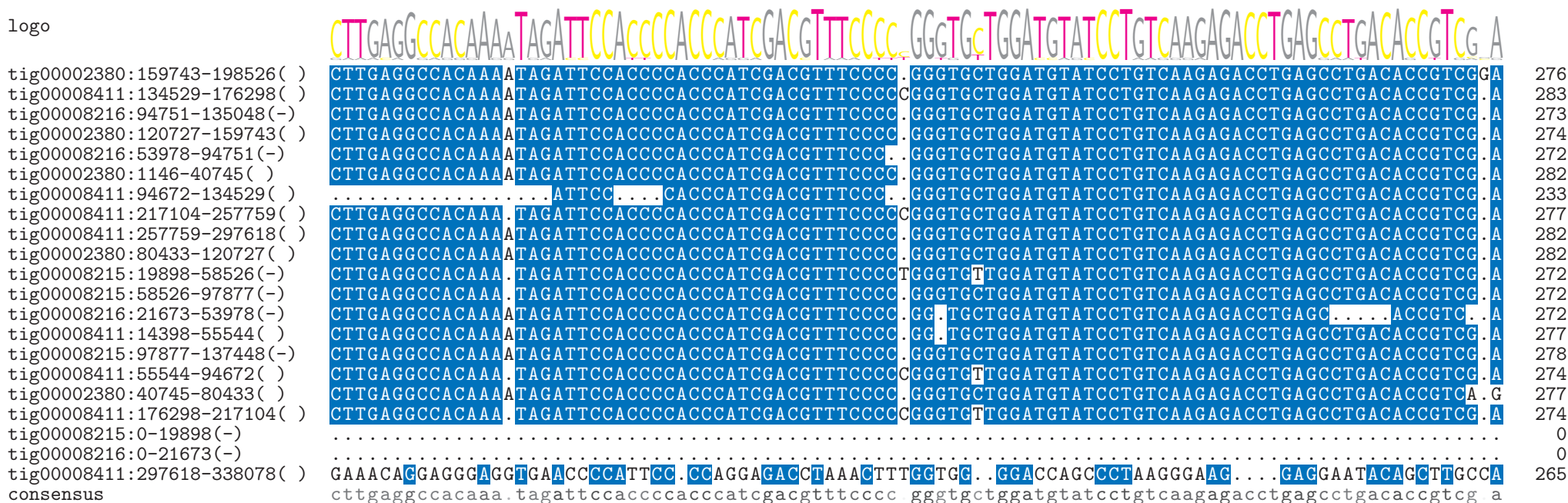
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tig00008411:297618-338078()
consensus

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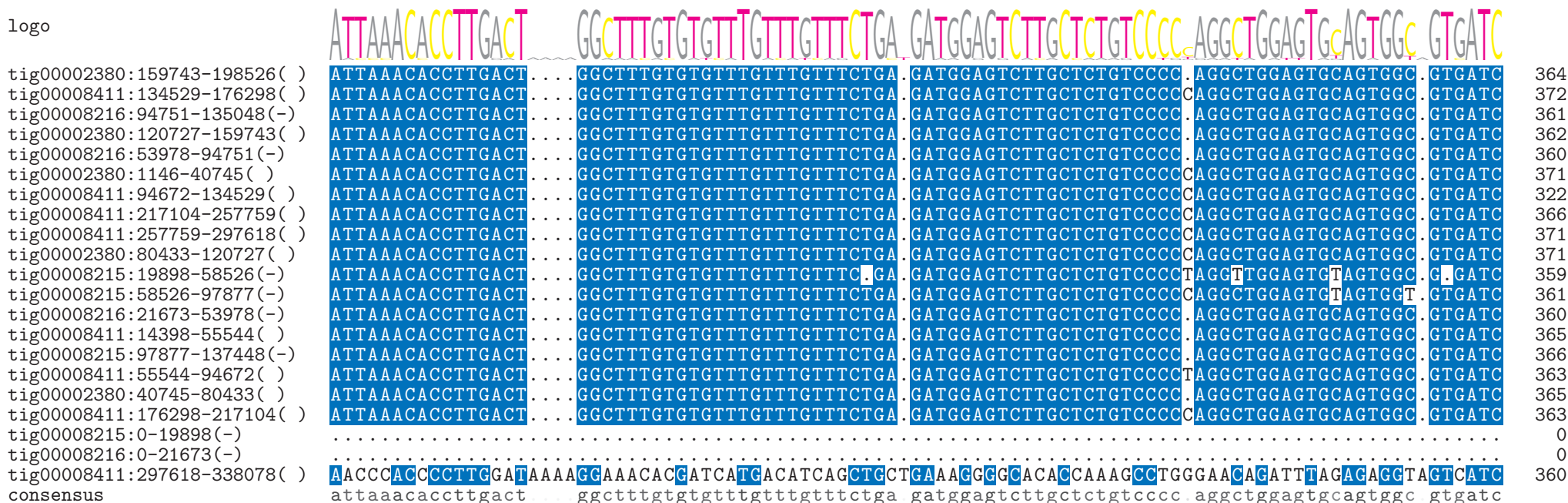
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consensus	aga	aac	ggt	att	aa	acac	tgt	gct	aat	gat	agt	gaa	agt	ga	aagacaaa	...gaaaggctatctat	ttt	gtggttagaataaagttgctcagt			

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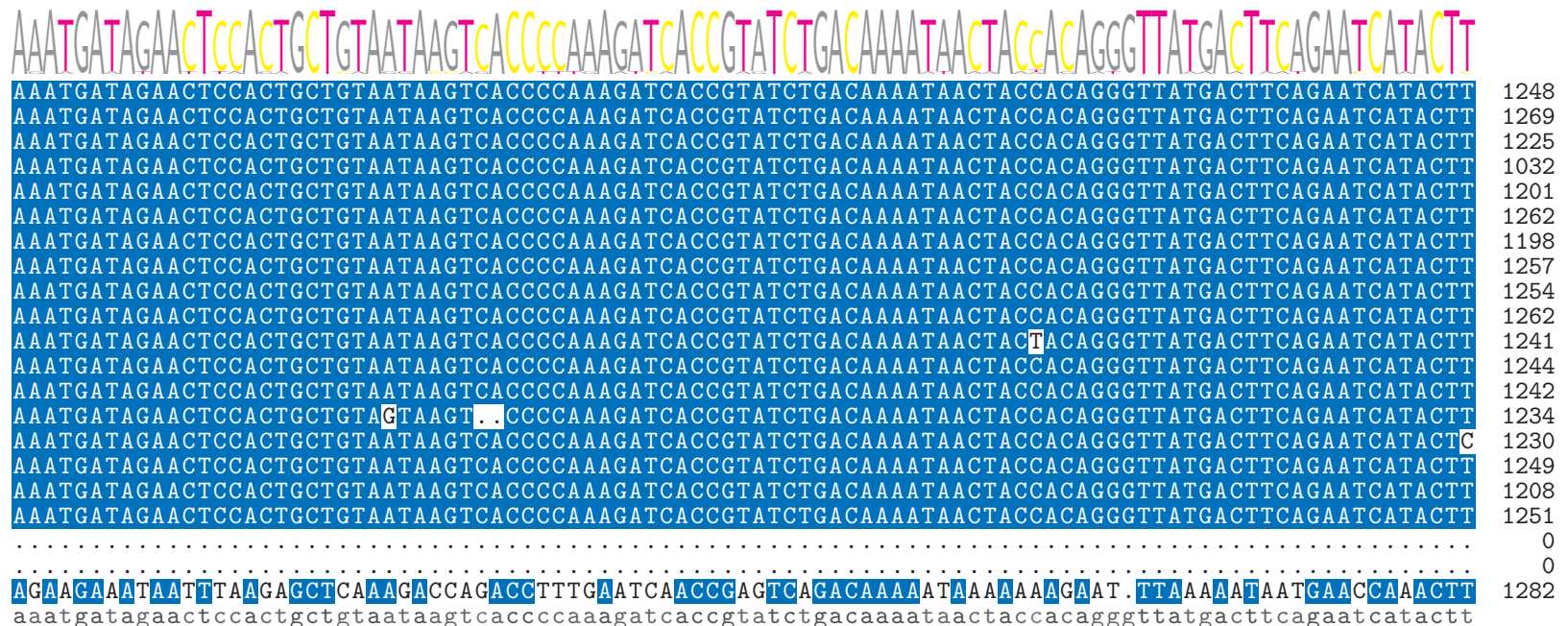
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tig00002380:80433-120727()	TCTCTTGATATTTACTTATGTAT	GTATTTATTTTTT	TAAATTT	ATTTCTCTTGAG	ACGGCG	TCTCGCTCTGTCGCCCAGGCTGGAGT	1348
tig00008215:19898-58526(-)	TCTCTTGATATTTACTTATGTAT	GTATTTATTTTTT	TAAATTT	ATTTCTCTTGAG	ACGGCG	TCTCGCTCTGTCGCCCAGGCTGGAGT	1325
tig00008215:58526-97877(-)	TCTCTTGATATTTACTTATGTAT	GTATTTATTTTTT	TAAATTT	ATTTCTCTTGAG	ACGGCG	TCTCGCTCTGTCGCCCAGGCTGGAGT	1329
tig00008216:21673-53978(-)	TCTCTTGATATTTACTTATGTAT	GTATTTATTTTTT	TAAATTT	ATTTCTCTTGAG	ACGGCG	TCTCGCTCTGTCGCCCAGGCTGGAGT	1328
tig00008411:14398-55544()	TCTCTTGATATTTACTTATGTAT	GTATTTATTTTTT	TAAATTT	ATTTCTCTTGAG	ACGGCG	TCTCGCTCTGTCGCCCAGGCTGGAGT	1320
tig00008215:97877-137448(-)	TCT	TATTTACTTATGTAT	GTATTTATTTTTT	ATTTCTCTTGAG	1306
tig00008411:55544-94672()	TCTCTTGATATTTACTTATGTAT	GTATTTATTTTTT	TAAATTT	ATTTCTCTTGAG	ACGGCG	TCTCGCTCTGTCGCCCAGGCTGGAGT	1334
tig00002380:40745-80433()	TCTCTTGATATTTACTTATGTAT	GTATTTATTTTTT	TAAATTT	ATTTCTCTTGAG	ACGGCG	TCTCGCTCTGTCGCCCAGGCTGGAGT	1296
tig00008411:176298-217104()	TCTCTTGATATTTACTTATGTAT	GTATTTATTTTTT	TAAATTT	ATTTCTCTTGAG	ACGGCG	TCTCGCTCTGTCGCCCAGGCTGGAGT	1337
tig00008215:0-19898(-)	0
tig00008216:0-21673(-)	0
tig00008411:297618-338078()	TTGAGAAATGTAGGA	TTATGTA	AAAGGAACA	AAATCTATGAC	TTATTGGC	ATTTT	TTTTTGAGACAGAA	1376
consensus	tctcttgatatttactttatgtat	gtattttatTTTTT	aatttt	atTTctcttgag	acggcg	tctcgctctgtcgcccaggctggagt	

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tig00002380:159743-198526()	GCGATGGTGTGATCTCGGCTCACTGCAACCGCCACCTCCCTGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGG	.ACTACAGGTG	1428
tig00008411:134529-176298()	GCGATGGTGTGATCTCGGCTCACTGCAACCGCCACCTCCCTGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGG	.ACTACAGGTG	1449
tig00008216:94751-135048(-)	GCGATGGTGTGATCTCGGCTCACTGCAACCGCCACCTCCCTGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGG	.ACTACAGGTG	1339
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tig00002380:1146-40745()	GCGATGGTGTGATCTCGGCTCACTGCAACCGCCACCTCCCTGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGG	.ACTACAGGTG	1442
tig00008411:94672-134529()	GCGATGGTGTGATCTCGGCTCACTGCAACCGCCACCTCCCTGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGG	.ACTACAGGTG	1377
tig00008411:217104-257759()	GCGATGGTGTGATCTCGGCTCACTGCAACCGCCACCTCCCTGGGTTCAAGCGATTCTCCTGCCTCAGCTTCCCGAGTAGCTGGG	.ACTACAGGTG	1438
tig00008411:257759-297618()	GCGATGGTGTGATCTCGGCTCACTGCAACCGCCACCTCCCTGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGG	.ACTACAGGTG	1433
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tig00008216:21673-53978(-)	GCGATGGTGTGATCTCGGCTCACTGCAACCGCCACCTCCCTGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGG	.ACTACAGGTG	1422
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tig00008411:55544-94672()	GCGATGGTGTGATCTCGGCTCACTGCAACCGCCACCTCCCTGGGTTCAAGCGATTCTCCTGCCTCAGCTTCTCTGAGTAGCTGGG	.ACTACAGGTG	1428
tig00002380:40745-80433()	GCGATGGTGTGATCTCGGCTCACTGCAACCGCCACCTCCCTGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGG	.ACTACAGGTG	1391
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tig00008215:0-19898(-)	0
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tig00008411:297618-338078()	GCAGCGGCATGATCTTGGCTCACTGAACCTCCACCTCCTTGA.TTCAAGTAATTCTCATGCCTCAGCCTCCCAAGTATCTGAG.ATTACAGGCG		1469
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tig00002380:1146-40745()	CCCGCCACCACGCCAGCTAATCTTT	.ATACTTTTAATA	GAGACGGGG	..TTTCACCGTGT	CGGCCCGGATGGTCTCGATCTCTTGACCTCG	..T	1532
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tig00008215:0-19898(-)							0
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tig00008216:94751-135048(-)	GACCCGCCCCGCTCGGCCTCCCAAAGTGCTGGGATGACAGGCGTGAGCCACTGAGCCCGG	.CCTTCTCTTGACGTTTAAACTATGAAGTCAGTCC	1523
tig00002380:120727-159743()	GACCCGCCCCGCTCGGCCTCCCAAAGTGCTGGGATGACAGGCGTGAGCCACTGAGCCCGG	.CCTTCTCTTGACGTTTAAACTATGAAGTCAGTCC	1250
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tig00008215:0-19898(-)	0
tig00008216:0-21673(-)	0
tig00008411:297618-338078()	GATCCTCCCACCTTGGCCTCCCAAAGTGCTGGGATTACTGGCATGAGCCACGGTGGTGGCTGACTTATTGGCATTCTAACAC	1645
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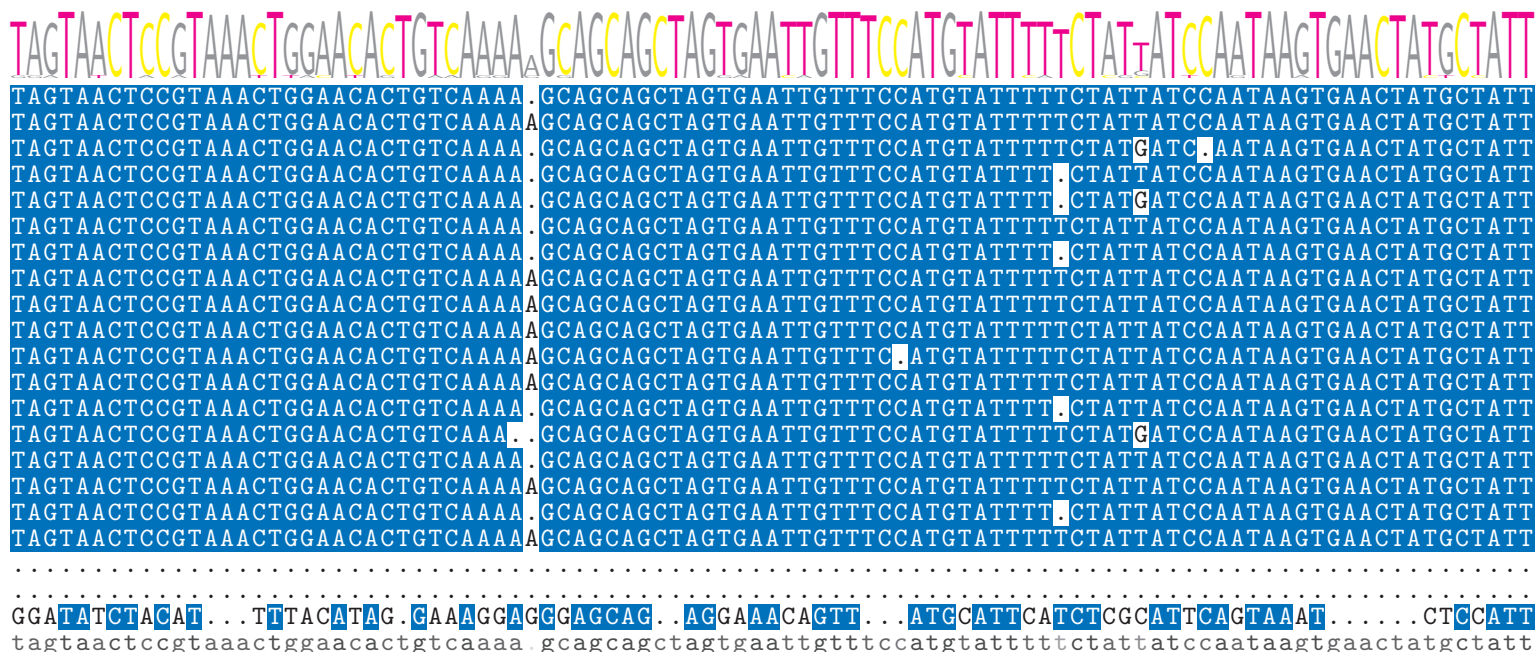
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tig00008411:217104-257759()	AGAGAAACGCAATAAATGTCAACGGTGAGG.	ATGGTGTTGAGG.	CAGAAGTAGGACCACACTTTTCTATCTTATTCAGTTGATAACAATATGA	1714
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tig00002380:80433-120727()	AGAGAAACGCAATAAATGTCAACGGTGAGG.	ATGGTGTTGAGG.	CAGAAGTAGGACCACACTTTT.CCTATCTTATTCAGTTGATAACAATATGA	1718
tig00008215:19898-58526(-)	AGAGAAACGCAATAAATGTCAACGG G GAGG.	ATGGTGTTGAGG.	CAGAAGTAGGACT T ACACTTTTCTATCTTATTCAGTTGATAACAATATGA	1690
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tig00002380:40745-80433()	AGAGAAACGCAATAAATGTCAACGGTGAGG G	ATGGTGTTG G GG.	CAGA .. GTAGGACCACACTTTT.CCTATCTTATTCAGTTGATAACAATATGA	1577
tig00008411:176298-217104()	AGAGAAACGCAATAAATGTCAACGGTGAGG.	ATGGTGTTGAGG.	CAGAAGTAGGACCACACTTTTCTATCTTATTCAGTTGATAACAATATGA	1708
tig00008215:0-19898(-)	0
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tig00008411:297618-338078()	AGA AG AA .. GAG AA AG TAA TCAAC TT GGA AA.. ACG T TTT GAGAA CAT AAT TCAGG .. AAACTG T CCTCA AT CTTG C TAA AGAG GGT .. CGA C ATG ..			1733
consensus	agagaaacgcaataaatgtcaacggtgagg	atggtgttgagg	cagaagtaggaccacactttttcctatcttattcagttgataacaatatga	

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tig00008411:94672-134529()	CCTAGGTAGTAATTTCTATGTGCCTACTTATACACG	AGTACAAA	GAGTAAAACAGAGA	CTGCTAAATTAAGGGTACGTGAAGTTCTTCA	1641	
tig00008411:217104-257759()	CCTAGGTAGTAATTTCTATGTGCCTACTTATACACG	AGTACAAA	GAGTAAAACAGAGAGA	CTGCTAAATTAAGGGTACGTGAAGTTCTTCA	1809	
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tig00008216:21673-53978(-)	CCTAGGTAGTAATTTCTATGCTACTTATACACG	AGTACAAA	GAGTAAAACAGAGA	CTGCTAAATTAAGGGTACGTGAAGTTCTTCA	1790	
tig00008411:14398-55544()	CCTAGGTAGTAATTTCTATGTGCCTACTTATACACG	AGTACAAA	GAGTAAAACAGAGAGA	CTGCTAAATTAAGGGTACGTGAAGTTCTTCA	1781	
tig00008215:97877-137448(-)	CCTAGGTAGTAATTTCTATGTGCCTACTTATACACG	AGTACAAA	GAGTAAAACAGAGA	CTGCTAAATTAAGGGTACGTGAAGTTCTTCA	1762	
tig00008411:55544-94672()	CCTAGGTAGTAATTTCTATGTGCCTACTTATACACG	AGTACAAA	GAGTAAAACAGAGAGA	CTGCTAAATTAAGGGTACGTGAAGTTCTTCA	1799	
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tig00008411:176298-217104()	CCTAGGTAGTAATTTCTATGTGCCTACTTATACACG	AGTACAAA	GAGTAAAACAGAGAGA	CTGCTAAATTAAGGGTACGTGAAGTTCTTCA	1802	
tig00008215:0-19898(-)	0	
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tig00008411:297618-338078()	CTGATACAGAAATCTCAGAGAACTCCGTGAGACTGAAGGTAGACA	GTGTAAGCAAAATGC	TTGCTTTC	TTGTGTGGCTC	TGATTAGCATCA	1823
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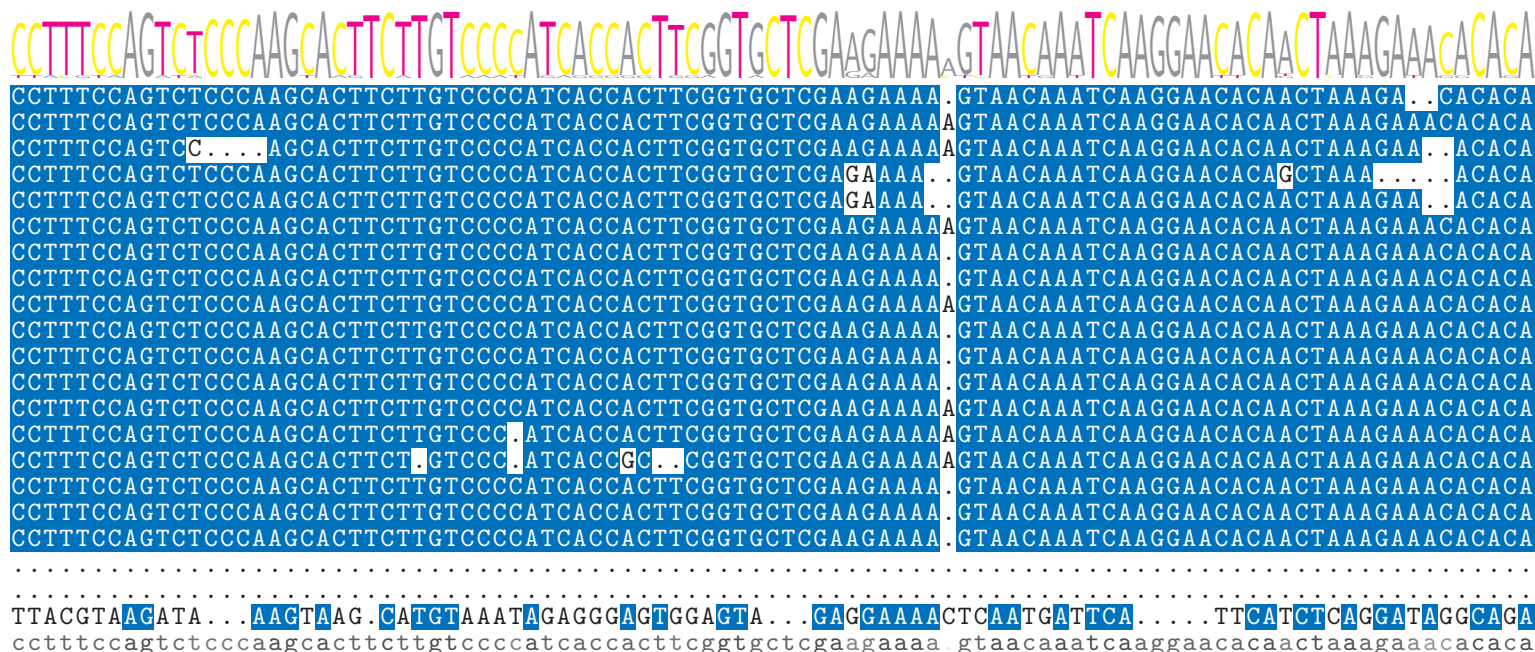
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consensus



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