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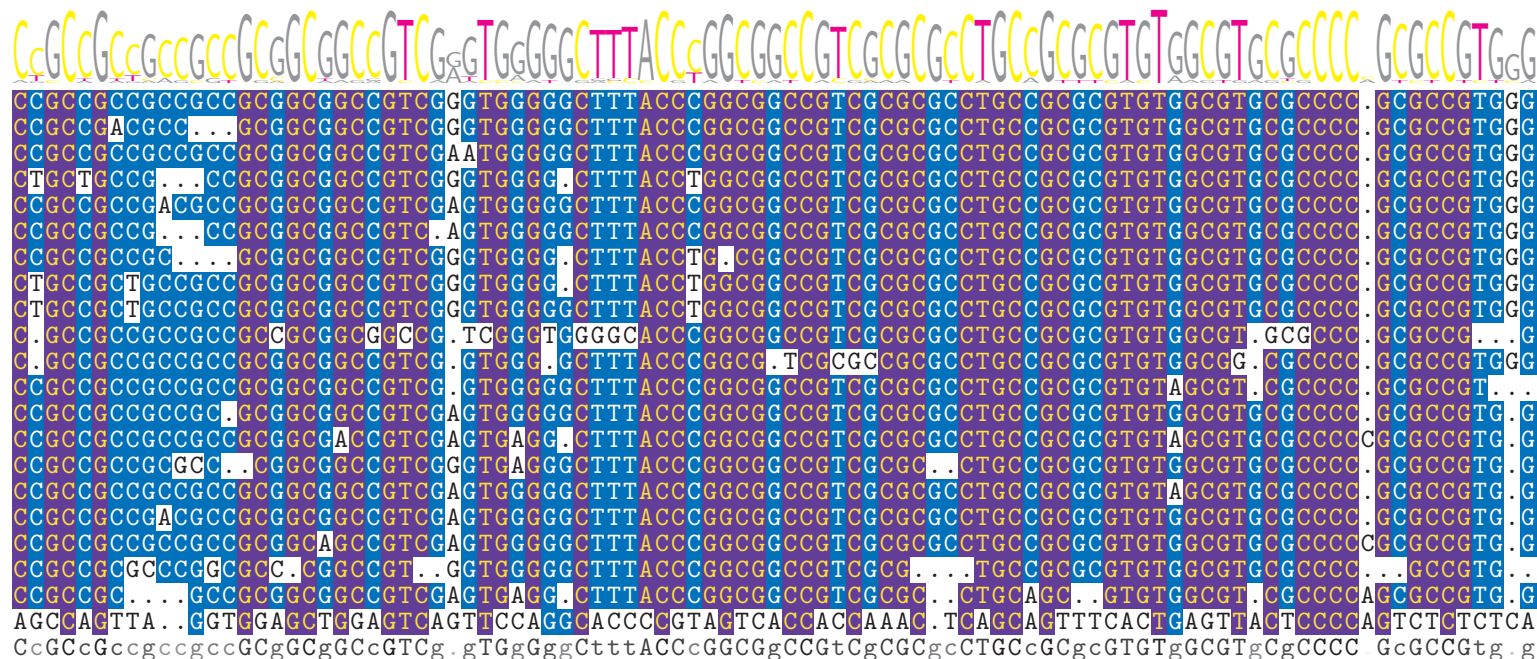
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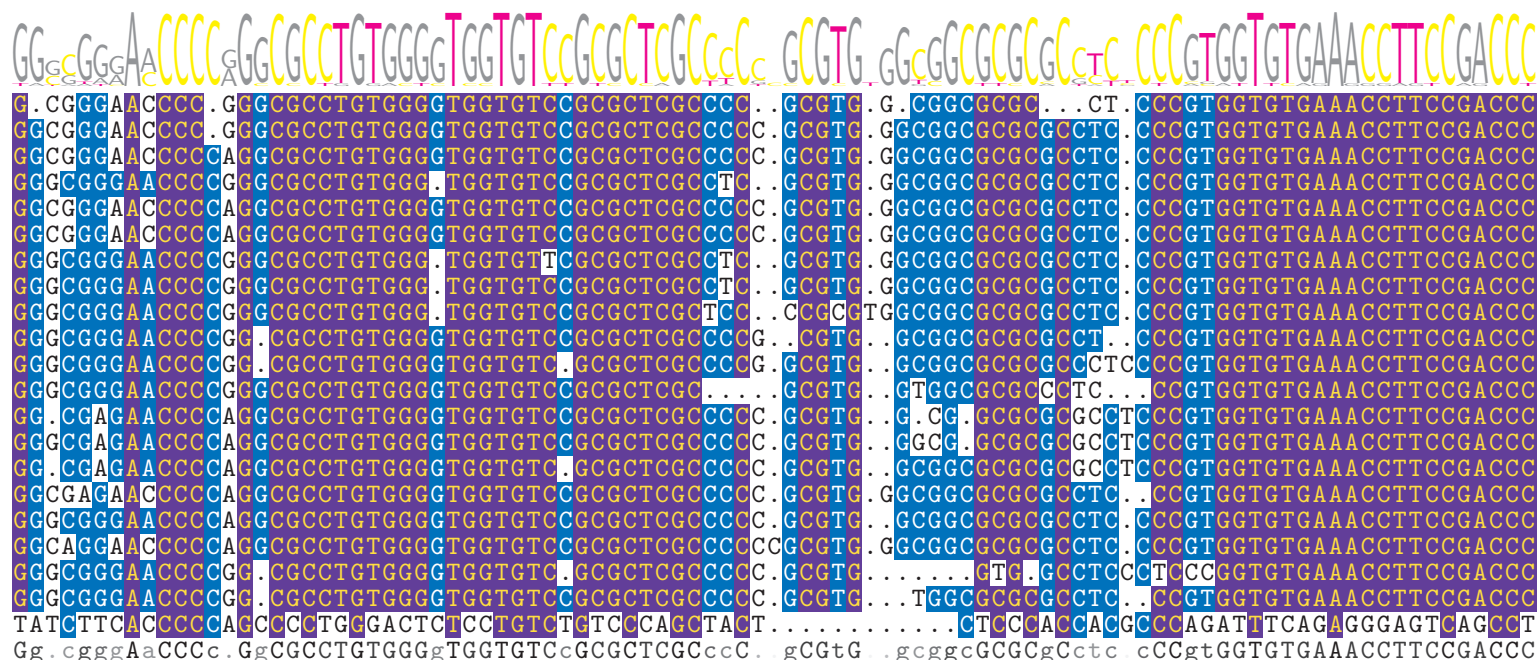
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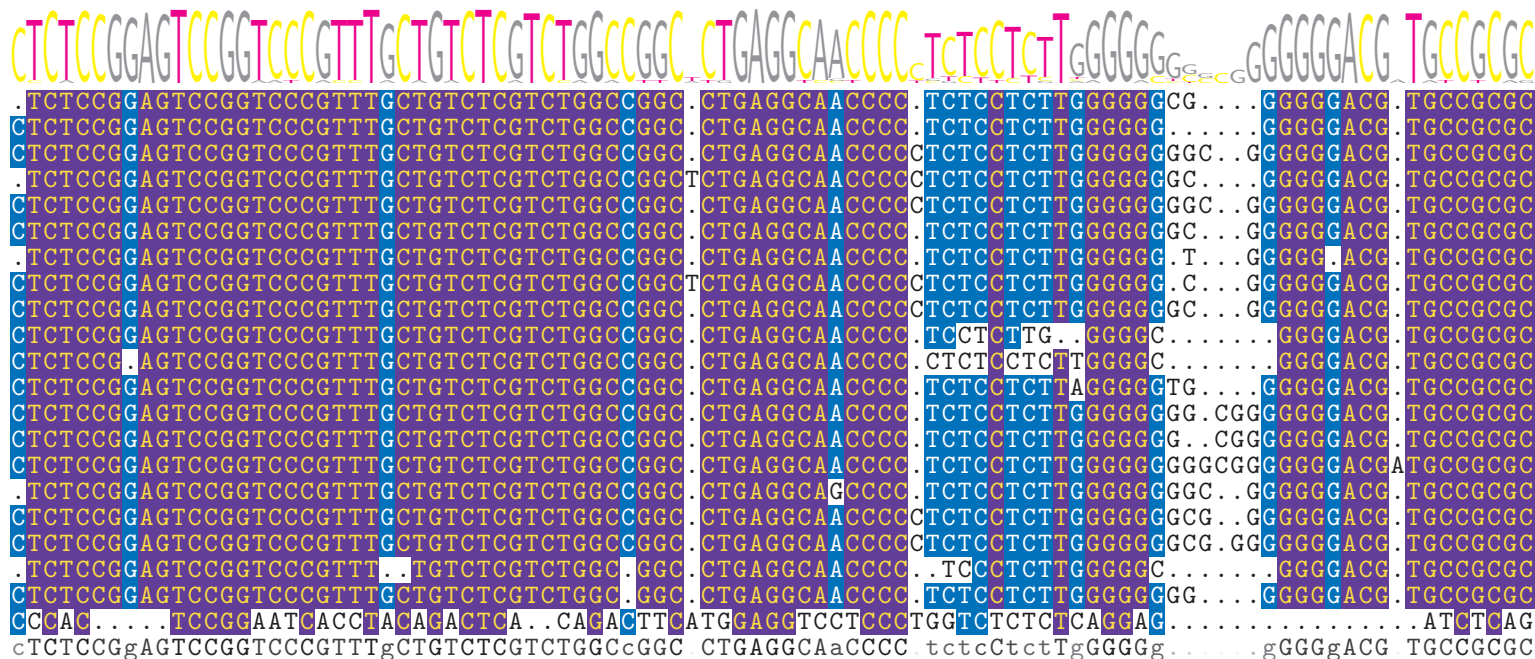
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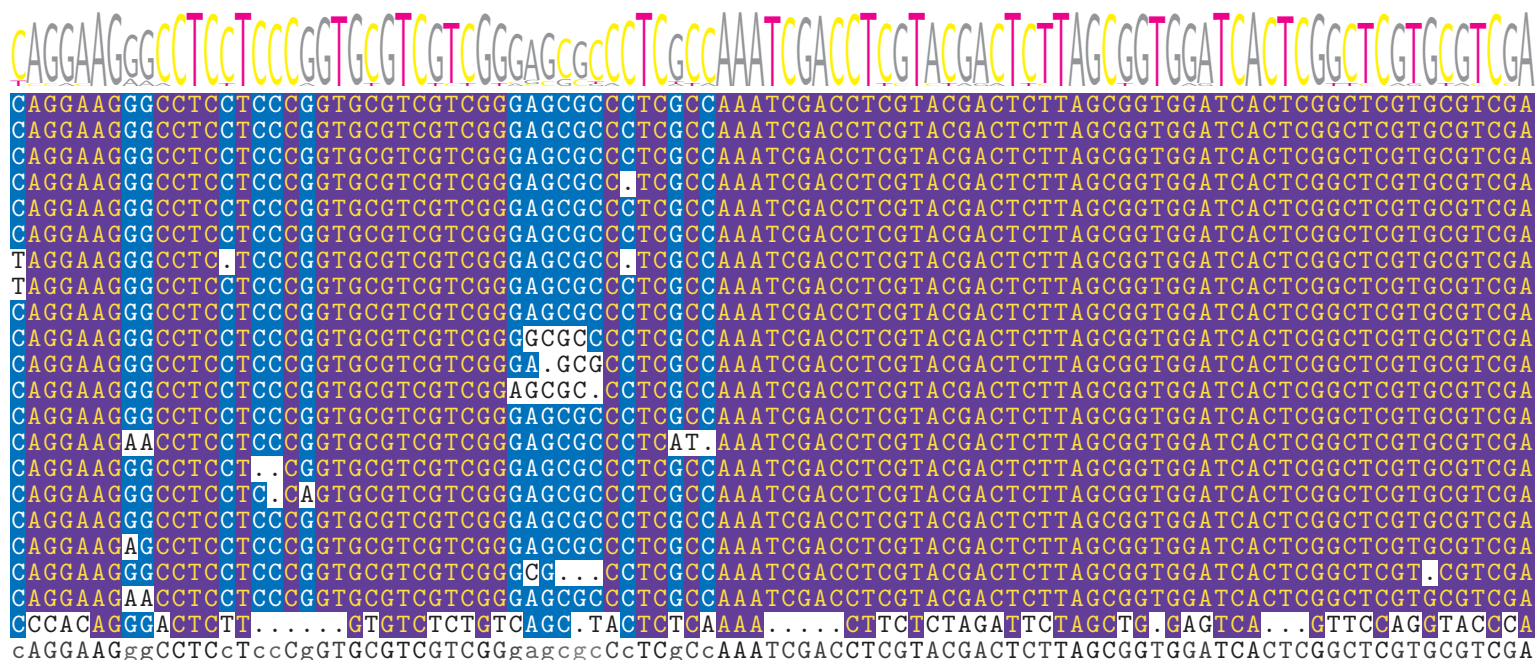
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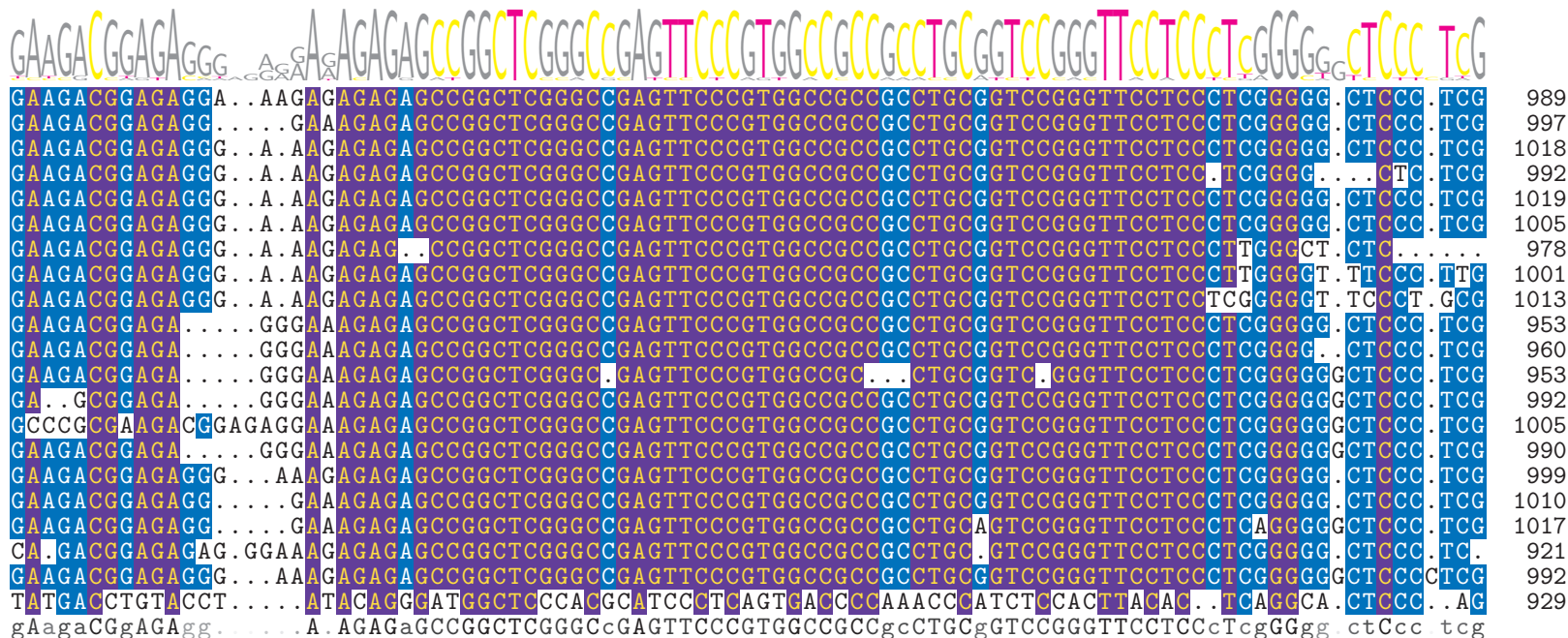
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tig00008215:19898-58526(-)	CGGGCCCTGCGTGGT	CACGCGTCGGGTGGCGGGGGG	...	AGGGG	GCGCGCCCGGCT	..	GAGAGAG	GGGAGGGCGCGCGCGCGCGCGCGCCCGC	...	GCCCGC	873
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tig00008216:21673-53978(-)	CGGGCCCTGCGTGGT	CACGCGTCGGGTGGCGGGGGG	...	GAGGGG	GCGCGCCCGGCT	..	GAGAGAGAT	GGGAGGGCGCGCGCGCGCGCGCGCCCGC	...	GCCCGC	905
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tig00008411:297618-338078()	GATTTCTGC	CACAGTCA	GCCCAGGCAC	CACAGACA	...	ACCTAGAC	ACTC	ACAGGCCTCATG	...	ACTATCTCC	844
consensus	cGGGCCcTGCGTGGT	CACGCGTCGGGTGGCGGGGGg	...	g.g	GCGGcCcGgCt	gagAGAGa	gGg	gggcgGCGccGCCGCC	...	cgC	

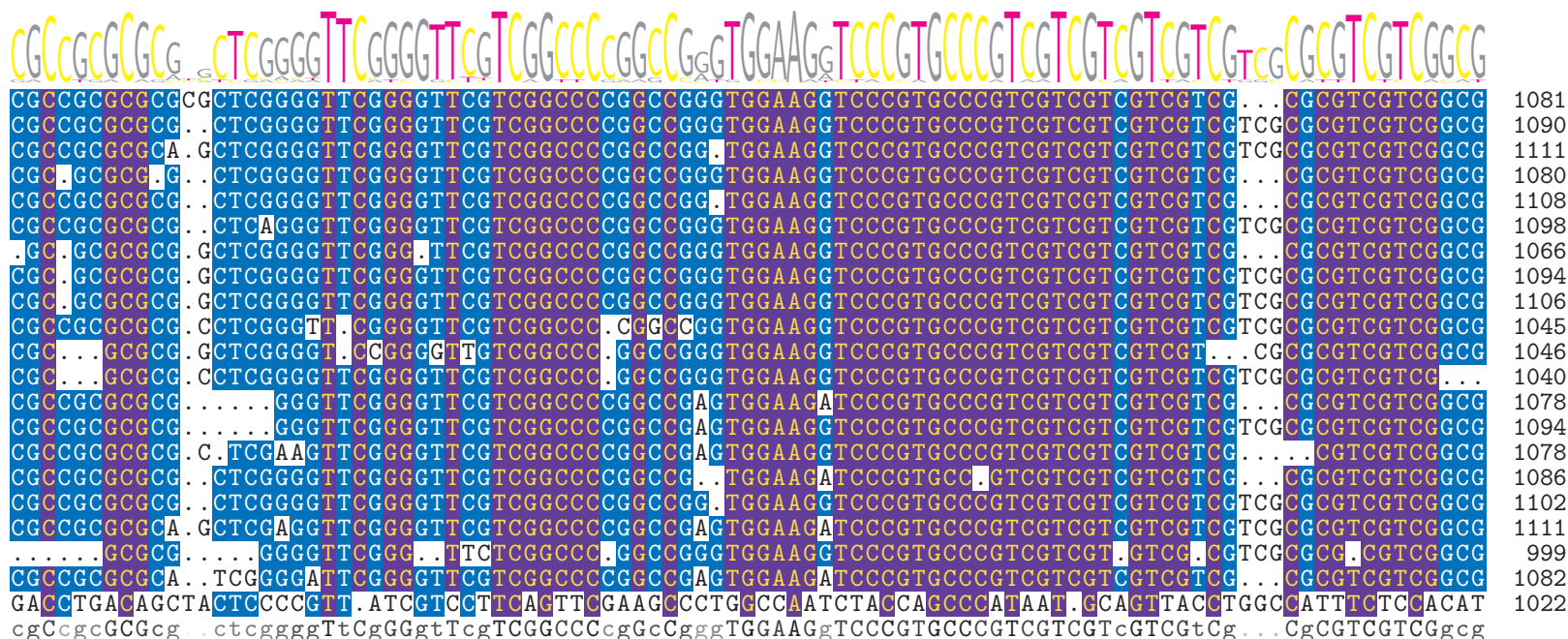
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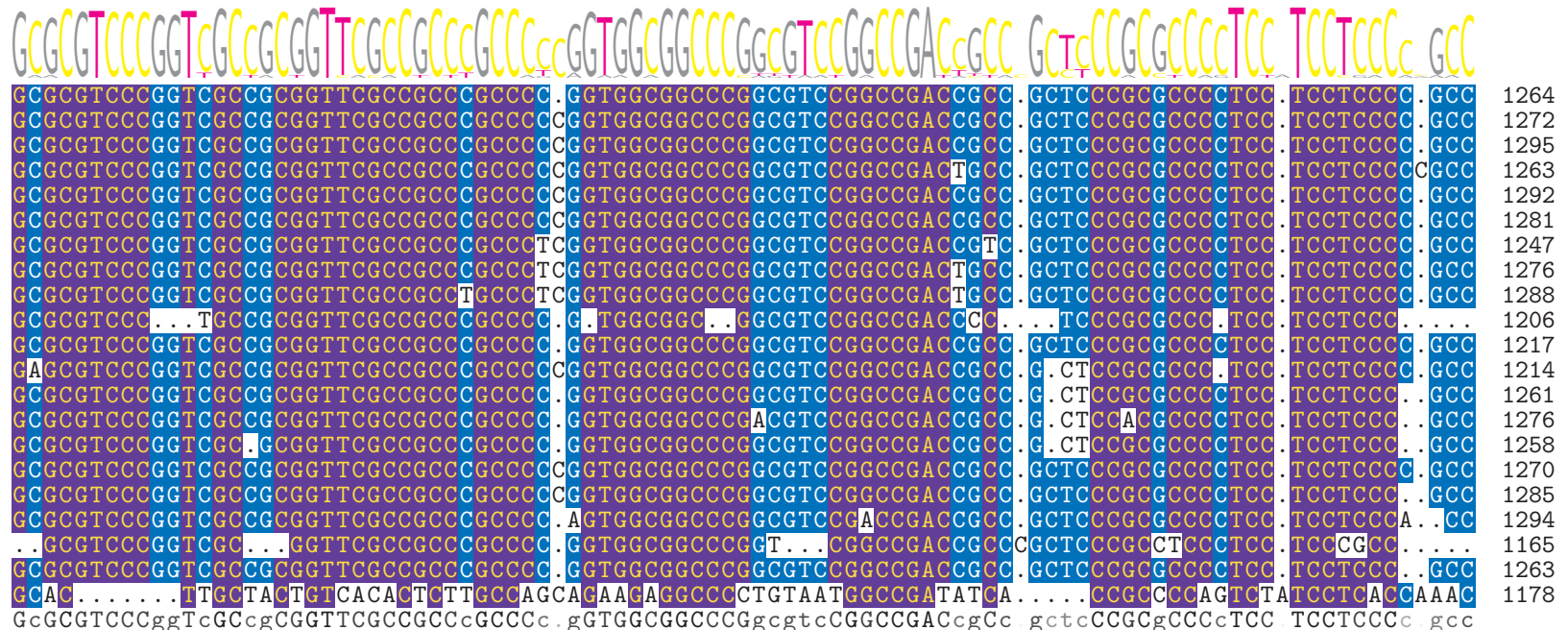


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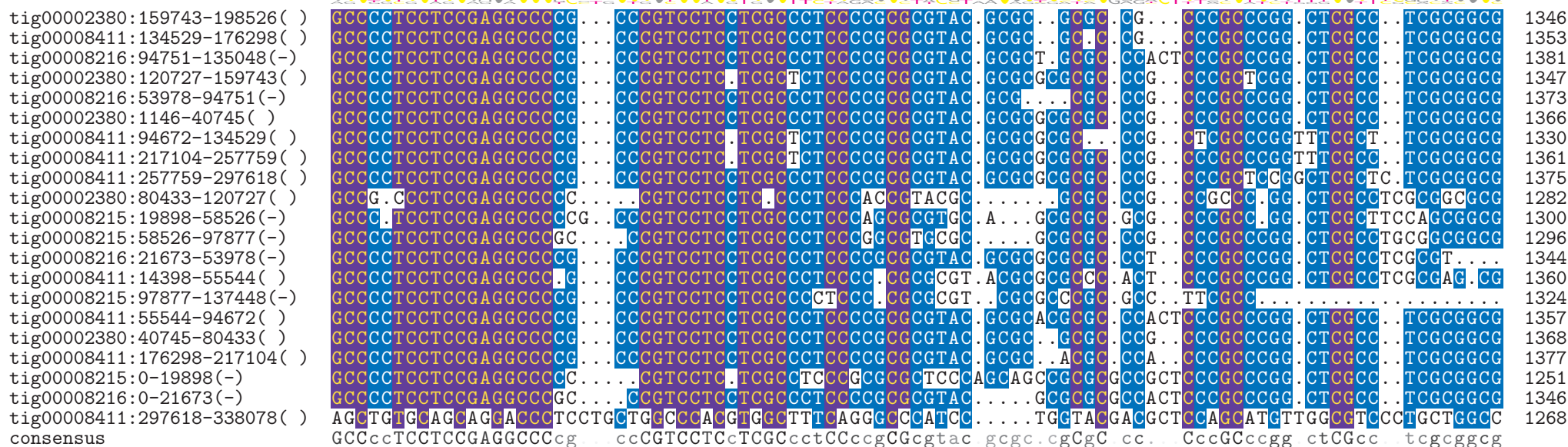
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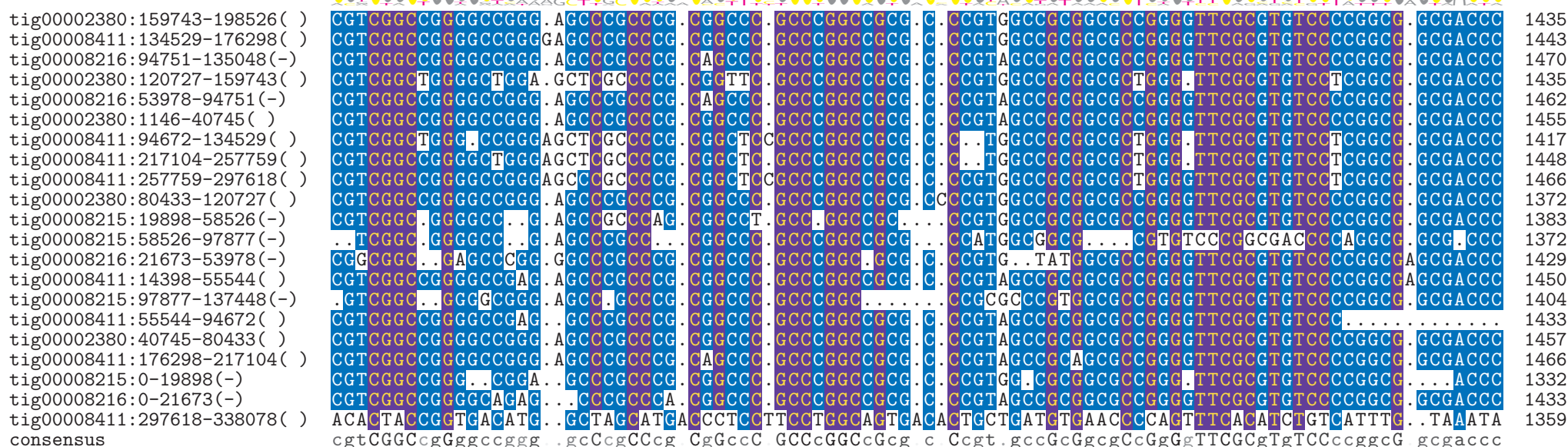
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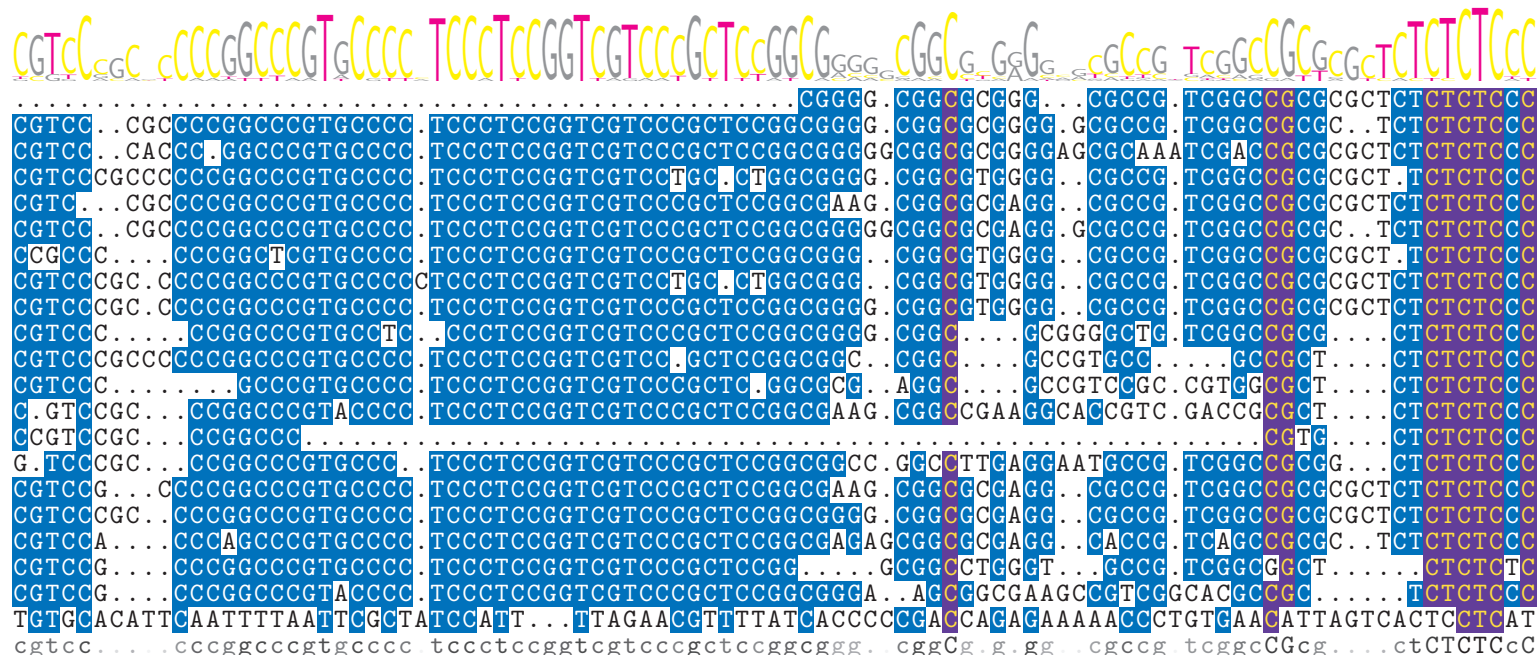
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tig00008215:19898-58526(-)	GCGGGAC	GCCGCGG	...	TGTGTC	GCCGTC	CGCG	GCC	GC	CTCG	GGGGCCCCGTCCCGAGCTTC	1423						
tig00008215:58526-97877(-)	GGGTCG	TCCGCG	TCGCG	CGCG	TCGCG	CGCG	CGCG	GC	CGCG	...	CGCG	CGCG	GGGGCCCCGTCCCGAGCTTC	1445						
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tig00008411:55544-94672()	GGC	1436						
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tig00008216:0-21673(-)	GCGGGG	GCCGCGG	...	TGTGTC	CGCG	GTCG	CGGCC	GC	CTCGGCT	CGCGGCCGCGC	...	CGCGCC	GGG	CCCCGTCCCGAGCTTC	1517					
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consensus	Gcgggac	gcCgCgg		tgtCgtCgcCgtcgCgCgccGC		ctCgggct	cgcggccgcgc	cgC	cgcgcc	gggccccgtcccgagcttc										

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tig00002380:1146-40745()	CGCGTCGGGG	GCGGGGCGG	CTCC	A...	CCGCCGCGT	CTCGGA	CCCCGTCCCC	CGAC	CTCC	CGGGGGAGAC	CGGGTTCGGG	CGT	GCGCGCC	1629									
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tig00008411:217104-257759()	CGCGT	TGGG	CGGG	TGG	CTCC	G...	CCGCCGCGT	CTCGGA	CCCCGTCCCC	CGAC	CTCC	CGGGGGAGAC	CGGGTTCGGG	CGT	GCGCGCC	1623							
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tig00008215:19898-58526(-)	...	GGCGG	CGCG	...	CCAT	...	CTCGGA	CCCCGTCCCC	CGAC	CTCC	CGGGGGAGAC	CGGGTTCGGG	CGT	GCGCGCC	1488								
tig00008215:58526-97877(-)	CGCGTCGGG	...	GCGGGCGG	CTC	...	CCGCCGCGT	CTCGGA	CCCCGTCCCC	CGAC	CTCC	CGGGG	GGAGG	CGGGT	...	GCGCGCC	1519							
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tig00008411:14398-55544()	CGCGTCGGGG	GCGGGGCGG	CTC	A...	CCGCCGCGT	CTCGGA	CCCCGTCCCC	CGAC	CTCC	CGCGGGGGAGAC	CGGGTTCGA	GGCGT	CA	GCGCC	1626								
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tig00008411:55544-94672()	...	GG	CTCC	A...	CCGCCGCGT	CTCGGA	CCCCGTCCCC	CGAC	CTCC	CGGGGGAGAC	CGGGTTCGGG	CGT	GCGCGCC	1506									
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tig00008411:297618-338078()	CTCACTAT	GCTCTTCAT	GCTCTG	GATAA	AAGT	GTC	CA	TATAA	AATT	GCCCC	ATTA	AAAGT	C	ATGAAT	GGAGT	GATT	TTTA	G	TATAT	TTT	G	TGT	1547
consensus	cgcgtcgggg	cggggcgG	CTCc	...	cgcgcgCgtC	CTCGGA	accGtCCC	cc	gacctcc	...	gggggagacgggttcggG	cgt	cgGcGcc										

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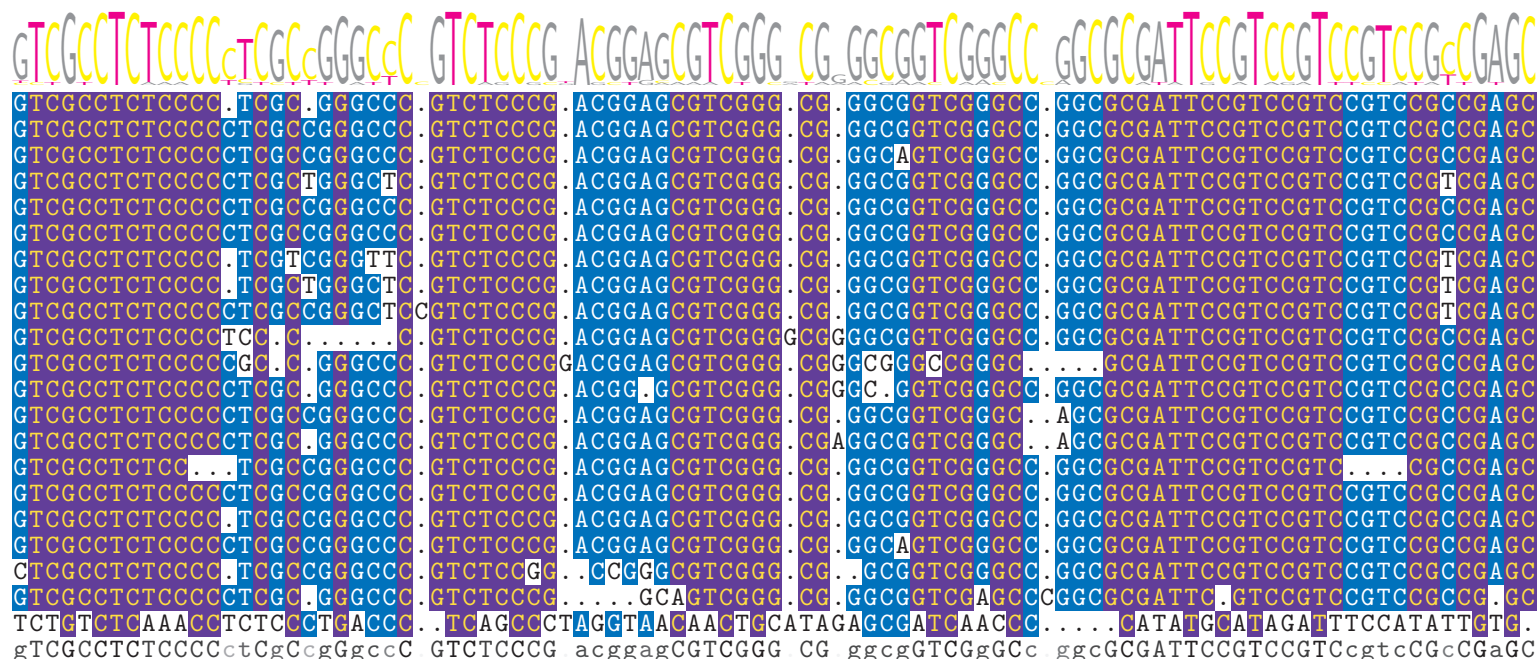
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consensus

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