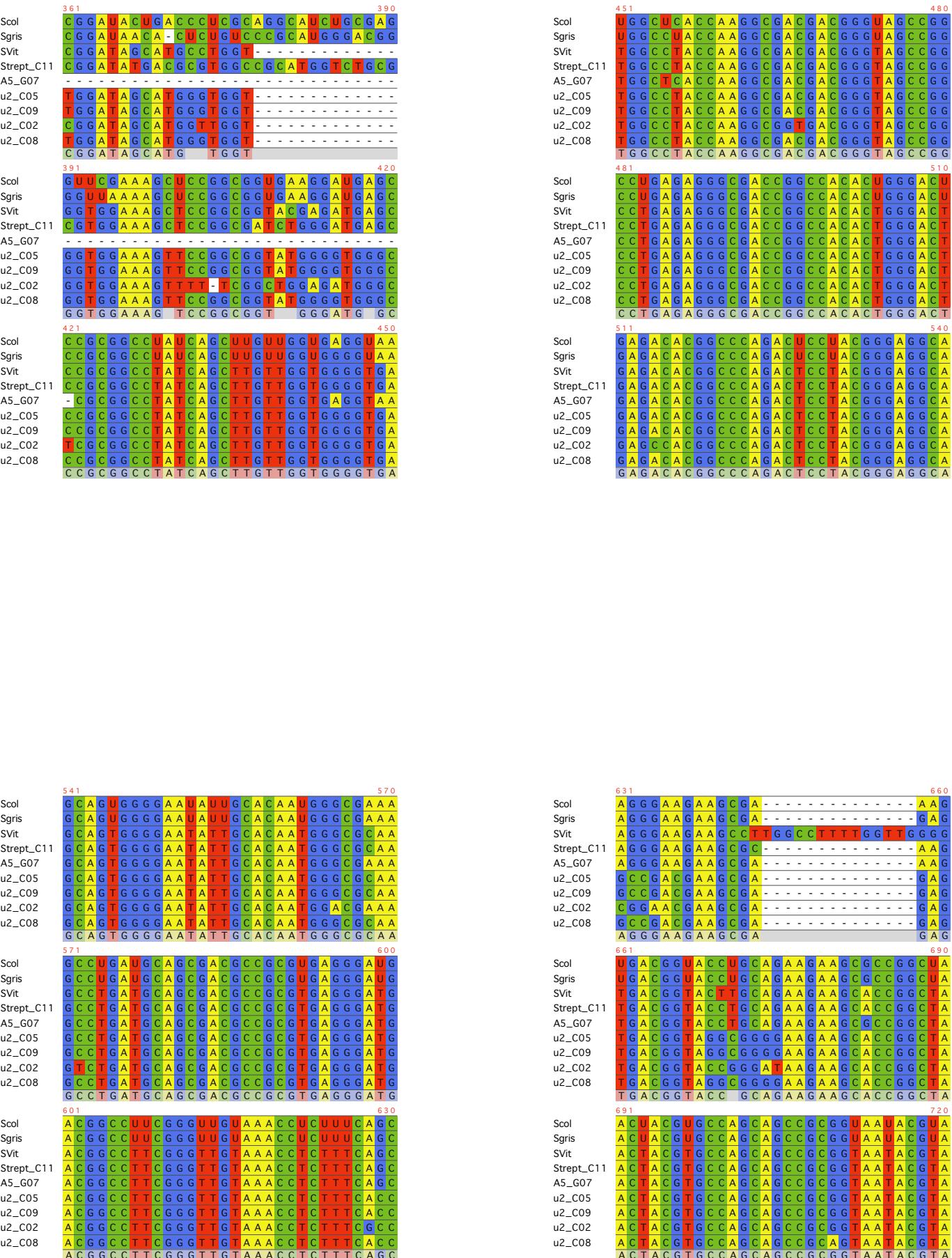


Scol	- - - - -	30
Sgris	- - - - -	
SVit	- - - - -	
Strept_C11	G A A C T C A A C A G C G T G C C A A A A T C A A C G C C	
A5_G07	- - - - -	
u2_C05	- - - - -	
u2_C09	- - - - -	
u2_C02	- - - - -	
u2_C08	- - - - -	
Scol	- - - - -	60
Sgris	- - - - -	
SVit	- - - - -	
Strept_C11	A G A T T A G T T G A T A C C C C C G G A C G G G G C C G C	
A5_G07	- - - - -	
u2_C05	- - - - -	
u2_C09	- - - - -	
u2_C02	- - - - -	
u2_C08	- - - - -	
Scol	- - - - -	90
Sgris	- - - - -	
SVit	- - - - -	
Strept_C11	G A A T T T G T T T G T G G T G T T G T G G T T C	
A5_G07	- - - - -	
u2_C05	- - - - -	
u2_C09	- - - - -	
u2_C02	- - - - -	
u2_C08	- - - - -	
Scol	- - - - -	120
Sgris	- - - - -	
SVit	- - - - -	
Strept_C11	C T T T G A A A C A C A T A T G C A A G G A C G C T G G T G	
A5_G07	- - - - -	
u2_C05	- - - - -	
u2_C09	- - - - -	
u2_C02	- - - - -	
u2_C08	- - - - -	
Scol	- - - - -	150
Sgris	- - - - -	
SVit	- - - - -	
Strept_C11	G G G G C C C C G A T T T T C C T C G G T G G C G G G A C C A	
A5_G07	- - - - -	
u2_C05	- - - - -	
u2_C09	- - - - -	
u2_C02	- - - - -	
u2_C08	- - - - -	
Scol	- - - - -	180
Sgris	- - - - -	
SVit	- - - - -	
Strept_C11	C C G C T C C T C C G C G T G T G T C A T C C C T G A T T A	
A5_G07	- - - - -	
u2_C05	- - - - -	
u2_C09	- - - - -	
u2_C02	- - - - -	
u2_C08	- - - - -	

Scol	- - - - -	210
Sgris	- - - - -	
SVit	- - - - -	
Strept_C11	C A G G G A A G C A T T C A C G G A G A G T T T G A T C C	
A5_G07	- - - - -	
u2_C05	- - - - -	
u2_C09	- - - - -	
u2_C02	- - - - -	
u2_C08	- - - - -	
Scol	- - - - -	240
Sgris	U G G C U C A G G A C G A A C G C U G G C G G C G U G C U U	
SVit	- - - - -	
Strept_C11	T G G C T C A G G A C G A A T G C T G G C G G C G T G C T T	
A5_G07	- - - - -	
u2_C05	- - - - -	
u2_C09	- - - - -	
u2_C02	- - - - -	
u2_C08	- - - - -	
Scol	- - - - -	270
Sgris	A A C A C A U G C A A G U C G A A C G A U G A A C C A C U U	
SVit	- - - - -	
Strept_C11	A A C A C A T G C A A G T C G A A C G G T G A A G C C T T T	
A5_G07	- - - - -	
u2_C05	- - - - -	
u2_C09	- - - - -	
u2_C02	- - - - -	
u2_C08	- - - - -	
Scol	- - - - -	300
Sgris	C G G U G G G G A U U A G U G G C G A A C G G G U G A U A	
SVit	- - - - -	
Strept_C11	C G G G T G G A T C A G T G G C G A A C G G G T G A G T A	
A5_G07	- - - - -	
u2_C05	- - - - -	
u2_C09	- - - - -	
u2_C02	- - - - -	
u2_C08	- - - - -	
Scol	- - - - -	330
Sgris	A C A C G U G G G C A A U C U G - C C C U U C A C U C U - G	
SVit	- - - - -	
Strept_C11	A C A C G T G G G C A A T C T G - C C T C G T A C T T T - G	
A5_G07	- - - - -	
u2_C05	- - - - -	
u2_C09	- - - - -	
u2_C02	- - - - -	
u2_C08	- - - - -	
Scol	- - - - -	360
Sgris	G G A C A A G C C C U G G A A A - C G G G G U C U A A U A C	
SVit	- - - - -	
Strept_C11	G G A T A A C C C G G G G A A A - C T C G G G C T A A T A C	
A5_G07	- - - - -	
u2_C05	- - - - -	
u2_C09	- - - - -	
u2_C02	- - - - -	
u2_C08	- - - - -	



Scol	GGGCGCAAGCGUUGUCGGAAUUAUUGGGC	721	Scol	GUCUGCAUCGAUACGGGCAAGGUAGGU	811	Scol	GUCUGCAUCGAUACGGGCAAGGUAGGU	840
Sgris	GGCGCGAACAGCGUUGUCGGAAUUAUUGGGC		Sgris	GUCUGCAUCGAUACGGGCAAGGUAGGU		Sgris	GUCUGCAUCGAUACGGGCAAGGUAGGU	
SVit	GGGTGCGAGCATTTGTCGGAAATTATTGGGC		SVit	GTCATCGATTCGATAACGGGCAAGGGCTAGAGTT		SVit	GTCATCGATTCGATAACGGGCAAGGGCTAGAGTT	
Strept_C11	GGGTGCGAGCATTTGTCGGAAATTATTGGGC		Strept_C11	GTCATCGATTCGATAACGGGCAAGGGCTAGAGTT		Strept_C11	GTCATCGATTCGATAACGGGCAAGGGCTAGAGTT	
A5_G07	GGCGCGAACAGCGTGTGGCGGAATTATTGGGC		A5_G07	TTCATCGATTCGATAACGGGCTAGCTAGAGTG		A5_G07	TTCATCGATTCGATAACGGGCTAGCTAGAGTG	
u2_C05	GGGTGCGAGCGTGTGGCGGAATTATTGGTC		u2_C05	GCGTGGTCGATACGGGCAAGACCTTGAGTT		u2_C05	GCGTGGTCGATACGGGCAAGACCTTGAGTT	
u2_C09	GGGTGCGAGCGTGTGGCGGAATTATTGGGC		u2_C09	GCGTGGTCGATACGGGCAAGACCTTGAGTT		u2_C09	GCGTGGTCGATACGGGCAAGACCTTGAGTT	
u2_C02	GGGTGCGAGCGTGTGGCGGAATTATTGGGC		u2_C02	GCGTGGTCGATACGGGCAAGACTTGAAGTT		u2_C02	GCGTGGTCGATACGGGCAAGACTTGAAGTT	
u2_C08	GGGTGCGAGCGTGTGGCGGAATTATTGGGC		u2_C08	GCGTGGTCGATACGGGCAAGGCTAGAGTT		u2_C08	GCGTGGTCGATACGGGCAAGGCTAGAGTT	
	GGGTGCGAGCGTGTGGCGGAATTATTGGGC	750		GGGTGCGGGTGGATGGATACGGGAGACTTGAAGTT			GGGTGCGGGTGGATGGATACGGGAGACTTGAAGTT	
Scol	GUAAAAGAGCUUCGUAGGGCGGUUUCUACGU	751	Scol	GGGUAGGGAGAUUCGUUUCGGGUAG	841	Scol	GGGUAGGGAGAUUCGUUUCGGGUAG	870
Sgris	GUAAAAGAGCUUCGUAGGGCGGUUUCUACGU		Sgris	GGGUAGGGAGAUUCGUUUCGGGUAG		Sgris	GGGUAGGGAGAUUCGUUUCGGGUAG	
SVit	GTAAGAGGOTCGTAGGGCGGGCTTGTGCGTC		SVit	GGGTAGGGAGATCGGAATTCCCTGGGTGTA		SVit	GGGTAGGGAGATCGGAATTCCCTGGGTGTA	
Strept_C11	GTAAGAGGOTCGTAGGGCGGGCTTGTGCGTC		Strept_C11	GGGTAGGGAGATCGGAATTCCCTGGGTGTA		Strept_C11	GGGTAGGGAGATCGGAATTCCCTGGGTGTA	
A5_G07	GTAAGAGGOTCGTAGGGCGGGCTTGTGCGTC		A5_G07	TGGTAGGGAGATTCGAAATTCCCTGGGTGTA		A5_G07	TGGTAGGGAGATTCGAAATTCCCTGGGTGTA	
u2_C05	GTAAGAGGOTCGTAGGGCGGGTTTGTGCGTC		u2_C05	GGGTAGGGAGTCTGGAATTCCCTGGGTGTA		u2_C05	GGGTAGGGAGTCTGGAATTCCCTGGGTGTA	
u2_C09	GTAAGAGGOTCGTAGGGCGGGTTTGTGCGTC		u2_C09	GGGTAGGGAGTCTGGAATTCCCTGGGTGTA		u2_C09	GGGTAGGGAGTCTGGAATTCCCTGGGTGTA	
u2_C02	GTAAGAGGOTCGTAGGGCGGGTTTGTGCGTC		u2_C02	GGGTAGGGAGTCTGGAATTCCCTGGGTGTA		u2_C02	GGGTAGGGAGTCTGGAATTCCCTGGGTGTA	
u2_C08	GTAAGAGGOTCGTAGGGCGGGTTTGTGCGTC		u2_C08	GGGTAGGGAGACTTGAATTCCCTGGGTGTA		u2_C08	GGGTAGGGAGACTTGAATTCCCTGGGTGTA	
	GTAAGAGGOTCGTAGGGCGGGTTTGTGCGTC	780		GCGTGGAGACTTGAATTCCCTGGGTGTA			GCGTGGAGACTTGAATTCCCTGGGTGTA	
Scol	GGUUGUGAAAGCCC-GGGGUUAAACCCCCGG	781	Scol	GGGUAAAUGCGCAGAUACAGGAGAACAA	871	Scol	GGGUAAAUGCGCAGAUACAGGAGAACAA	900
Sgris	GGAUUGUGAAAGCCC-GGGGUUAAACCCCCGG		Sgris	GGGUAAAUGCGCAGAUACAGGAGAACAA		Sgris	GGGUAAAUGCGCAGAUACAGGAGAACAA	
SVit	GGATGTGAAAGCCC-GGGGUUAAACCCCCGG		SVit	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		SVit	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
Strept_C11	GGATGTGAAAGCCC-GGGGUUAAACCCCCGG		Strept_C11	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		Strept_C11	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
A5_G07	GGGTGTGAAAGACC-GGGGUUAAACCCCCGG		A5_G07	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		A5_G07	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
u2_C05	GGCCGTGAAAAACCT-ACCGCTTAAACGGTGG		u2_C05	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		u2_C05	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
u2_C09	GGCCGTGAAAAACCT-ACCGCTTAAACGGTGG		u2_C09	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		u2_C09	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
u2_C02	TGCCGTGAAAAACCT-GGGGUUAAACCTCCGG		u2_C02	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		u2_C02	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
u2_C08	GGCCGTGAAAAACCT-ACCGCTTAAACGGTGG		u2_C08	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		u2_C08	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
	GGGTGTGAAAAACCT-GGGGUUAAACCCCCGG	810		GGGTGAAATGCGCAGATATCAGGAGGAAACAA			GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
Scol	GGUUGUGAAAGCCC-GGGGUUAAACCCCCGG	871	Scol	GGUAAAUGCGCAGAUACAGGAGAACAA	900	Scol	GGUAAAUGCGCAGAUACAGGAGAACAA	930
Sgris	GGAUUGUGAAAGCCC-GGGGUUAAACCCCCGG		Sgris	GGGUAAAUGCGCAGAUACAGGAGAACAA		Sgris	GGGUAAAUGCGCAGAUACAGGAGAACAA	
SVit	GGATGTGAAAGCCC-GGGGUUAAACCCCCGG		SVit	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		SVit	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
Strept_C11	GGATGTGAAAGCCC-GGGGUUAAACCCCCGG		Strept_C11	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		Strept_C11	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
A5_G07	GGGTGTGAAAGACC-GGGGUUAAACCCCCGG		A5_G07	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		A5_G07	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
u2_C05	GGCCGTGAAAAACCT-ACCGCTTAAACGGTGG		u2_C05	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		u2_C05	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
u2_C09	GGCCGTGAAAAACCT-ACCGCTTAAACGGTGG		u2_C09	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		u2_C09	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
u2_C02	TGCCGTGAAAAACCT-GGGGUUAAACCTCCGG		u2_C02	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		u2_C02	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
u2_C08	GGCCGTGAAAAACCT-ACCGCTTAAACGGTGG		u2_C08	GGGTGAAATGCGCAGATATCAGGAGGAAACAA		u2_C08	GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
	GGGTGTGAAAAACCT-GGGGUUAAACCCCCGG	900		GGGTGAAATGCGCAGATATCAGGAGGAAACAA			GGGTGAAATGCGCAGATATCAGGAGGAAACAA	
Scol	ACUGACGCUAGGGAGCGAAGGGGUUAGGGAG	931	Scol	GCGUAAAACGGGUUAGGGCAACUAGGUAGGU	991	Scol	GCGUAAAACGGGUUAGGGCAACUAGGUAGGU	1020
Sgris	ACUGACGCUAGGGAGCGAAGGGGUUAGGGAG		Sgris	ACAUUCCACGGGUUAGGGCAACUAGGUAGGU		Sgris	ACAUUCCACGGGUUAGGGCAACUAGGUAGGU	
SVit	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		SVit	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		SVit	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
Strept_C11	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		Strept_C11	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		Strept_C11	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
A5_G07	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		A5_G07	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		A5_G07	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
u2_C05	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		u2_C05	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		u2_C05	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
u2_C09	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		u2_C09	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		u2_C09	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
u2_C02	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		u2_C02	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		u2_C02	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
u2_C08	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		u2_C08	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		u2_C08	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG	960		ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC			ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
Scol	ACUGACGCUAGGGAGCGAAGGGGUUAGGGAG	961	Scol	ACAUUACGGGUUAGGGCAACUAGGUAGGU	1021	Scol	ACAUUACGGGUUAGGGCAACUAGGUAGGU	1050
Sgris	ACUGACGCUAGGGAGCGAAGGGGUUAGGGAG		Sgris	ACAUUACGGGUUAGGGCAACUAGGUAGGU		Sgris	ACAUUACGGGUUAGGGCAACUAGGUAGGU	
SVit	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		SVit	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		SVit	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
Strept_C11	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		Strept_C11	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		Strept_C11	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
A5_G07	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		A5_G07	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		A5_G07	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
u2_C05	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		u2_C05	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		u2_C05	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
u2_C09	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		u2_C09	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		u2_C09	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
u2_C02	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		u2_C02	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		u2_C02	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
u2_C08	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG		u2_C08	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC		u2_C08	ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
	ACGTACGCTGAGGGAGCGAAGGGGUUAGGGAG	990		ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC			ACATTCACGGTGTGCGTGGCCGTAGGTGAGCTAAC	
Scol	CGAACAGGAUUAGAUACCGUAGGUAGGUAG	991	Scol	GCAUAAAGUG-CCCCGCCUUGGGAGGUAG	1051	Scol	GCAUAAAGUG-CCCCGCCUUGGGAGGUAG	1080
Sgris	CGAACAGGAUUAGAUACCGUAGGUAGGUAG		Sgris	GCAUAAAGGUUAGGUAGGUAGGUAGGUAG		Sgris	CGCAUAAAGGUUAGGUAGGUAGGUAGGUAG	
SVit	CGAACAGGAATTAGATACCCCTGGTAGTCCCAC		SVit	GCAATTAGGTT-CCCCGCCUUGGGAGGUAG		SVit	CGCAATTAGGTT-CCCCGCCUUGGGAGGUAG	
Strept_C11	CGAACAGGAATTAGATACCCCTGGTAGTCCCAC		Strept_C11	GCAATTAGGTTCCCCGCCUUGGGAGGUAG		Strept_C11	GCAATTAGGTTCCCCGCCUUGGGAGGUAG	
A5_G07	CGAACAGGAATTAGATACCCCTGGTAGTCCCAC		A5_G07	GCAATTAGGTTCCCCGCCUUGGGAGGUAG		A5_G07	GCAATTAGGTTCCCCGCCUUGGGAGGUAG	
u2_C05	CGAACAGGAATTAGATACCCCTGGTAGTCCCAC		u2_C05	GCAATTAGGTTCCCCGCCUUGGGAGGUAG		u2_C05	GCAATTAGGTTCCCCGCCUUGGGAGGUAG	
u2_C09	CGAACAGGAATTAGATACCCCTGGTAGTCCCAC		u2_C09	GCAATTAGGTTCCCCGCCUUGGGAGGUAG		u2_C09	GCAATTAGGTTCCCCGCCUUGGGAGGUAG	
u2_C02	CGAACAGGAATTAGATACCCCTGGTAGTCCCAC		u2_C02	GCAATTAGGTTCCCCGCCUUGGGAGGUAG		u2_C02	GCAATTAGGTTCCCCGCCUUGGGAGGUAG	
u2_C08	CGAACAGGAATTAGATACCCCTGGTAGTCCCAC		u2_C08	GCAATTAGGTTCCCCGCCUUGGGAGGUAG		u2_C08	GCAATTAGGTTCCCCGCCUUGGGAGGUAG	
	CGAACAGGAATTAGATACCCCTGGTAGTCCCAC	990		CGAACAGGAATTAGGTTCCCCGCCUUGGGAGGUAG			CGAACAGGAATTAGGTTCCCCGCCUUGGGAGGUAG	

	1171	1200
Scol	A C C A A G G C U U G A C A U A C A C C G G - - A A A G G C A	
Sgris	A C C A A G G C U U G A C A U A C A C C G G - - A A A G G C A	
SVit	A C C A A G G C T T G A C A T G C G C G G G - - A C G G G T	
Strept_C11	A C C A A G G C T T G A C A T G C G C G G G - - A A A G C C	
A5_G07	- - - - -	
u2_C05	A C C T G G G T T T G A C A T G C A C A G G - - A T C C C G	
u2_C09	A C C T G G G T T T G A C A T G C A C A G G - - A T C C C G	
u2_C02	A C C T G G G C T T G A C A T G C G C C A G - - A C A T C C	
u2_C08	A C C T G G G T T T G A C A T G C A C A G G - - A T C C C G	
	A C C G G C T T G A C A T C A C G G A C	
	1201	1230
Scol	U C A G A G A G A U G G U G C C C C C C U U G U G G U C G G U G	
Sgris	U C A G A G A G A U G G U G C C C C C C U U G U G G U C G G U A	
SVit	C C A G A G A T G G G G T T T C C T T T T G G T C C G G T	
Strept_C11	C C A G A G A T G G G G - C C C C T T T T A G T C C G G T	
A5_G07	- - - - -	
u2_C05	G C A G A G A T G T C G G T T C C C T T G T G G C C T G T G	
u2_C09	G C A G A G A T G T C G G T T C C C T T G T G G C C T G T G	
u2_C02	C C A G A G A T G G G G C T T C C C T T G T G G G T T G G T	
u2_C08	G C A G A G A T G T C G G T T C C C T T G T G G C C T G T G	
	C A G A G A T G G G T T C C C C T T G G G C C G G T G	
	1231	1260
Scol	U A C A G G U G G U G C A U G G C U G U C G U C A G C U G	
Sgris	U A C A G G U G G U G C A U G G C U G U C G U C A G C U G	
SVit	- A C A G G T G G T G C A T G G C T G T C G T C A G C T C G	
Strept_C11	- A C A G G T G G T G C A T G G C T G T C G T C A G C T C G	
A5_G07	- - - - -	
u2_C05	T G C A G G T G G T G C A T G G C T G T C G T C A G C T C G	
u2_C09	T G C A G G T G G T G C A T G G C T G T C G T C A G C T C G	
u2_C02	T G C A G G T G G T G C A T G G C T G T C G T C A G C T C G	
u2_C08	T G C A G G T G G T G C A T G G C T G T C G T C A G C T C G	
	C A G G T G G T G C A T G G C T G T C G T C A G C T C G	

	1 2 6 1	1 2 9 0
Scol	U G U C G G U G A G A U G U U G G G U U A A G U C C C G C A A	
Sgris	U G U C G G U G A G A U G U U G G G U U A A G U C C C G C A A	
SVit	T G T C G T G A G A T G T T G G G T T A A G T C C C C G C A A	
Strept_C11	T G T C G T G A G A T G T T G G G T T A A G T C C C C G C A A	
A5_G07		
u2_C05	T G T C G T G A G A T G T T G G G T T A A G T C C C C G C A A	
u2_C09	T G T C G T G A G A T G T T G G G T T A A G T C C C C G C A A	
u2_C02	T G T C G T G A G A T G T T G G G T T A A G T C C C C G C A A	
u2_C08	T G T C G T G A G A T G T T G G G T T A A G T C C C C G C A A T G T C G T G A G A T G T T G G G T T A A G T C C C C G C A A	
1 2 9 1		1 3 2 0
Scol	C G A G C G C A A C C C U U G U C C C G U G U U R G C C A G C	
Sgris	C G A G C G C A A C C C U U G U U C U G U G U U R G C C A G C	
SVit	C G A G C G C A A C C C T T G T T C T G T G T T G C C A A G C	
Strept_C11	C G A G C G C A A C C C T T G T T C T G T G T T G C C A A G C	
A5_G07	- - - - -	
u2_C05	C G A G C G C A A C C C T T A T C C T A T G T T G C C A A G C	
u2_C09	C G A G C G C A A C C C T T A T C C T A T G T T G C C A A G C	
u2_C02	C G A G C G C A A C C C T T A T C C T A T G T T G C C A A G C	
u2_C08	C G A G C G C A A C C C T T A T C C T A T G T T G C C A A G C C G A G C G C A A C C C T T C C T G T T G C C A A G C	
1 3 2 1		1 3 5 0
Scol	A A G C C C U U C - - G G G G U G U U U G G G G A C U C A C G	
Sgris	A U G C C C U U C - - G G G G U G A U G G G G A C U C A C A	
SVit	A - - - - - - - - G G T G G G G A C T C A C A	
Strept_C11	A C G C T C C T T C G G G G G T G G T G G G G A C T C A C A	
A5_G07	- - - - -	
u2_C05	G - - - - - - - - G G C G G G G G A C T C G T T G	
u2_C09	G - - - - - - - - G G C G G G G G A C T C G T T G	
u2_C02	G - - - - - - - - G C C G G G G G A C T C G T G	
u2_C08	G - - - - - - - - G G C G G G G G A C T C G T G G G G G G G A C T C	

	1 3 5 1	1 3 8 0
Scol	G G A G A C C - G C C G G G G U C A A C U C G G A G G A A G	
Sgris	G G A G A C U - G C C G G G G U C A A C U C G G A G G A A G	
SVit	G G A G A G C T - G C C G G G G T C A A C T C G G A G G A A G	
Strept_C11	G G A G A C T T G G C C G G G G T C A A C T C G G A G G A A G	
A5_G07	- - - - -	
u2_C05	G G A G A C T T - G C C G G G G T C A A C T C G G A G G A A G	
u2_C09	G G A G A C T - G C C G G G G T C A A C T C G G A G G A A G	
u2_C02	G G A G A C C - G C C G G G G T C A A C T C G G A G G A A G	
u2_C08	G G A G A C T - G C C G G G G T C A A C T C G G A G G A A G G G A G A C T G C C G G G G T C A A C T C G G A G G A A G	
	1 3 8 1	1 4 1 0
Scol	G U G G G G A C G A C G U C A A G U C A U C A U G C C C C U	
Sgris	G U G G G G A C G A C G U C A A G U C A U C A U G C C C C U	
SVit	G T G G G G A C G A C G T C A A G T C A T C A T G C C C C T	
Strept_C11	T G G G G G A C G A C G T C A A G T C A T C A T G C C C C T	
A5_G07	- - - - -	
u2_C05	G T G G G G A T G A C G T C A A G T C A T C A T G C C C C T	
u2_C09	G T G G G G A T G A C G T C A A G T C A T C A T G C C C C T	
u2_C02	G T G G G G A T G A C G T C A A G T C A T C A T G C C C C T	
u2_C08	G T G G G G A T G A C G T C A A G T C A T C A T G C C C C T G T G G G G A G A C G T C A A G T C A T C A T G C C C C T	
	1 4 1 1	1 4 4 0
Scol	U A U G U C U U U G G G C U G C A C A C G U G C U A C A A U G	
Sgris	U A U G U C U U U G G G C U G G C A C A C G U G C U A C A A U G	
SVit	T A T G T C T T G G G C T G C A C A C G T G C T A C A A T G	
Strept_C11	T A T G T C T T G G G C T G C A C A C G T G C T A C A A T G	
A5_G07	- - - - -	
u2_C05	T A T G T C C A G G G C T T C A C A C A T G C T A C A A T G	
u2_C09	T A T G T C C A G G G C T T C A C A C A T G C T A C A A T G	
u2_C02	T A T G T C C A G G G C T T C A C A C A T G C T A C A A T G	
u2_C08	T A T G T C C A G G G C T T C A C A C A T G C T A C A A T G T A T G T C G G G C T C A C A C G T G C T A C A A T G	

Scol	1441	G C G G U A C A A U G A G C U G C G A U A C C G C A A G G	1470	A A G U C G G A G U C G C U A G U A A U C G C A G A U C A G	1560
Sgris		G C G G U A C A A U G A G C U G C G A U U G C C G C A A G G		A A G U C G G A G U U G C U A G U A A U C G C A G A U C A G	
SVit		G C G G T A C A G T G A G C T G C G A T A C C G T G A G G		A A G T C G G A G T C T C T A G T A A T C G C A G A T C A G	
Strept_C11		G C G G T A C A A T G A G C T G C G A T A C C G T G A G G		A A G T C G G A G T C T C T A G T A A T C G C A G A T C A G	
A5_G07	- - - - -	- - - - -	- - - - -	- - - - -	- - - - -
u2_C05	1471	G C T G G T A C A G T G G G T T G C G A T A C C G T G A G G	1500	A A G T C G G A G T C G C T A G T A A T C G C A G A T C A G	1561
u2_C09		G C T G G T A C A G T G G G T T G C G A T A C C G T G A G G		A A G T C G G A G T C G C T A G T A A T C G C A G A T C A G	
u2_C02		G C T G G T A C A G A G G G T T G C G A T A T C G C G A G G		A A G T C G G A G T C G C T A G T A A T C G C A G A T C A G	
u2_C08		G C T G G T A C A G T G G G T T G C G A T A C C G T G A G G		A A G T C G G A G T C G C T A G T A A T C G C A G A T C A G	
		G C G G T A C A G T G T G C G A T A C C G T G A G G		A A G T C G G A G T C G C T A G T A A T C G C A G A T C A G	
Scol	1501	U G G A G C G A A U C U C A A A A G C C G G U C U C A G U	1530	C A U U G C U C G C G G U G A A U A C G U U C C C G G G C C U	1590
Sgris		C G G A G C G A A U C U C A A A A G C C G G U C U C A G U		C A U U G C U C G G G U G A A U A C G U U C C C G G G C C U	
SVit		T G G A G C G A A T C T C T A A A A G C C G G T C T C A G T		C A T T G C T G C G G G T G A A T A C G T T C C C G G G C C T	
Strept_C11		T G G A G C G A A T C T C T A A A A G C C G G T C T C A G T		C A T T G C T G C G G G T G A A T A C G T T C C C G G G C C T	
A5_G07	- - - - -	- - - - -	- - - - -	- - - - -	- - - - -
u2_C05	1531	T G G A G C G A A T C C C T G A A A G C T G G T C T C A G T	1561	C A A C G C T G C G G T G A A T A C G T T C C C G G G C C T	1591
u2_C09		T G G A G C G A A T C C C T G A A A G C T G G T C T C A G T		C A A C G C T G C G G T G A A T A T G T T C C C G G G C C T	
u2_C02		T G G A G C G A A T C C C T T A A A G C C G G T C T C A G T		C A A C G C T G C G G T G A A T A C G T T C C C G G G C C T	
u2_C08		T G G A G C G A A T C C C T G A A A G C T G G T C T C A G T		C A A C G C T G C G G T G A A T A C G T T C C C G G G C C T	
		T G G A G C G A A T C T C G A A C T C G A C T T G C G T G		C A G C T G C G G T G A A T A C G T T C C C G G G C C T	
Scol	1591	U G U A C A C A C C G C C C G U C A C G U C A C G A A A G U	1620	T G T A C A C A C C G C C C G T C A C G T C A T C G A A A G T	1620
Sgris		U G U A C A C A C C G C C C G U C A C G U C A C G A A A G U		T G T A C A C A C C G C C C G T C A C G T C A C G A A A G T	
SVit		T G G A T T G G G G T C T G C G A A C T C G A C T T G C G T G		T G T A C A C A C C G C C C G T C A C G T C A C G A A A G T	
Strept_C11		T G G A T T G G G G T C T G C G A A C T C G A C T T G C G T G		T G T A C A C A C C G C C C G T C A C G T C A C G A A A G T	
A5_G07	- - - - -	- - - - -	- - - - -	- - - - -	- - - - -
u2_C05	1621	T C G G A T C G G C A G T C T G C G A A C T C G A C T T G C G T G	1650	Scol	1711
u2_C09		T C G G A T C G G C A G T C T G C G A A C T C G A C T T G C G T G		Sgris	1740
u2_C02		T C G G A T C G G C A G T C T G C G A A C T C G A C T T G C G T G		SVit	
u2_C08		T C G G A T C G G C A G T C T G C G A A C T C G A C T T G C G T G		Strept_C11	
		T C G G A T C A C A C C C G A A G G C C C A		A5_G07	- - - - -
Scol	1651	A A C C C C U U G G U G G G A G G G A G C U G U C G A A G G U	1680	u2_C05	- - - - -
Sgris		A A C C C C U U G G U G G G A G G G A G C U G U C G A A G G U		u2_C09	- - - - -
SVit		- - - - -		u2_C02	- - - - -
Strept_C11		A C A C T C G T G G T G G G G G A G C C G T C G A A G G T		u2_C08	- - - - -
A5_G07	- - - - -	- - - - -	- - - - -		
u2_C05	1681	G G G A C U G G G G G A U U G G G A C G A A G G U C G U A A C A	1710	Scol	1741
u2_C09		G G G A C U G G G G G A U U G G G A C G A A G G U C G U A A C A		Sgris	1770
u2_C02		G G G A C C A G G G A T T G G G A C G A A G T C G T A A C A		SVit	
u2_C08		G G G A C C A G G G A T T G G G A C G A A G T C G T A A C A		Strept_C11	
		G G G A C C A G G G A T T G G G A C G A A G T C G T A A C A		A5_G07	- - - - -
Scol	1711	A G G U A G C C G U A C C G G A A G G U G G G C U G G A U	1740	u2_C05	- - - - -
Sgris		A G G U A G C C G U A C C G G A A G G U G G G C U G G A U		u2_C09	- - - - -
SVit		- - - - -		u2_C02	- - - - -
Strept_C11		A G G T A G C C G T A C C G G A A G G T G C G G C T G G A T		u2_C08	- - - - -
A5_G07	- - - - -	- - - - -	- - - - -		
u2_C05	1741	- - - - -		Scol	1771
u2_C09		- - - - -		Sgris	1800
u2_C02		- - - - -		SVit	
u2_C08		- - - - -		Strept_C11	
		- - - - -		A5_G07	- - - - -
Scol	1771	- - - - -		u2_C05	- - - - -
Sgris		- - - - -		u2_C09	- - - - -
SVit		- - - - -		u2_C02	- - - - -
Strept_C11		C C G G A G C G A A C G T C T C C G G G T G G T G C T C A		u2_C08	- - - - -
A5_G07	- - - - -	- - - - -	- - - - -		
u2_C05	1771	- - - - -			
u2_C09		- - - - -			
u2_C02		- - - - -			
u2_C08		- - - - -			

	1801	1830		1891	1920
Scol	- - -	- - -	Scol	- - -	- - -
Sgris	- - -	- - -	Sgris	- - -	- - -
SVit	- - -	- - -	SVit	- - -	- - -
Strept_C11	T G G G T G G A A C G T T G A T T A T T C G G C A C C T G G		Strept_C11	T C G C T C T C G G G T C C G G G T G C C G G G C G C G C T	
A5_G07	- - -	- - -	A5_G07	- - -	- - -
u2_C05	- - -	- - -	u2_C05	- - -	- - -
u2_C09	- - -	- - -	u2_C09	- - -	- - -
u2_C02	- - -	- - -	u2_C02	- - -	- - -
u2_C08	- - -	- - -	u2_C08	- - -	- - -
	1831	1860		1921	1950
Scol	- - -	- - -	Scol	- - -	- - -
Sgris	- - -	- - -	Sgris	- - -	- - -
SVit	- - -	- - -	SVit	- - -	- - -
Strept_C11	T C A C T G A C G G T G A C T G C T A G T A C T T G C A C C C		Strept_C11	G T T G G G T C C T G A G G G C A C G G C C G T G A G T G T	
A5_G07	- - -	- - -	A5_G07	- - -	- - -
u2_C05	- - -	- - -	u2_C05	- - -	- - -
u2_C09	- - -	- - -	u2_C09	- - -	- - -
u2_C02	- - -	- - -	u2_C02	- - -	- - -
u2_C08	- - -	- - -	u2_C08	- - -	- - -
	1861	1890		1951	1962
Scol	- - -	- - -	Scol	- - -	- - -
Sgris	- - -	- - -	Sgris	- - -	- - -
SVit	- - -	- - -	SVit	- - -	- - -
Strept_C11	C C C T T C T T C G G G G G G T G C G T G G A A C G C G G T		Strept_C11	C G C G G T T G G C C T	
A5_G07	- - -	- - -	A5_G07	- - -	- - -
u2_C05	- - -	- - -	u2_C05	- - -	- - -
u2_C09	- - -	- - -	u2_C09	- - -	- - -
u2_C02	- - -	- - -	u2_C02	- - -	- - -
u2_C08	- - -	- - -	u2_C08	- - -	- - -