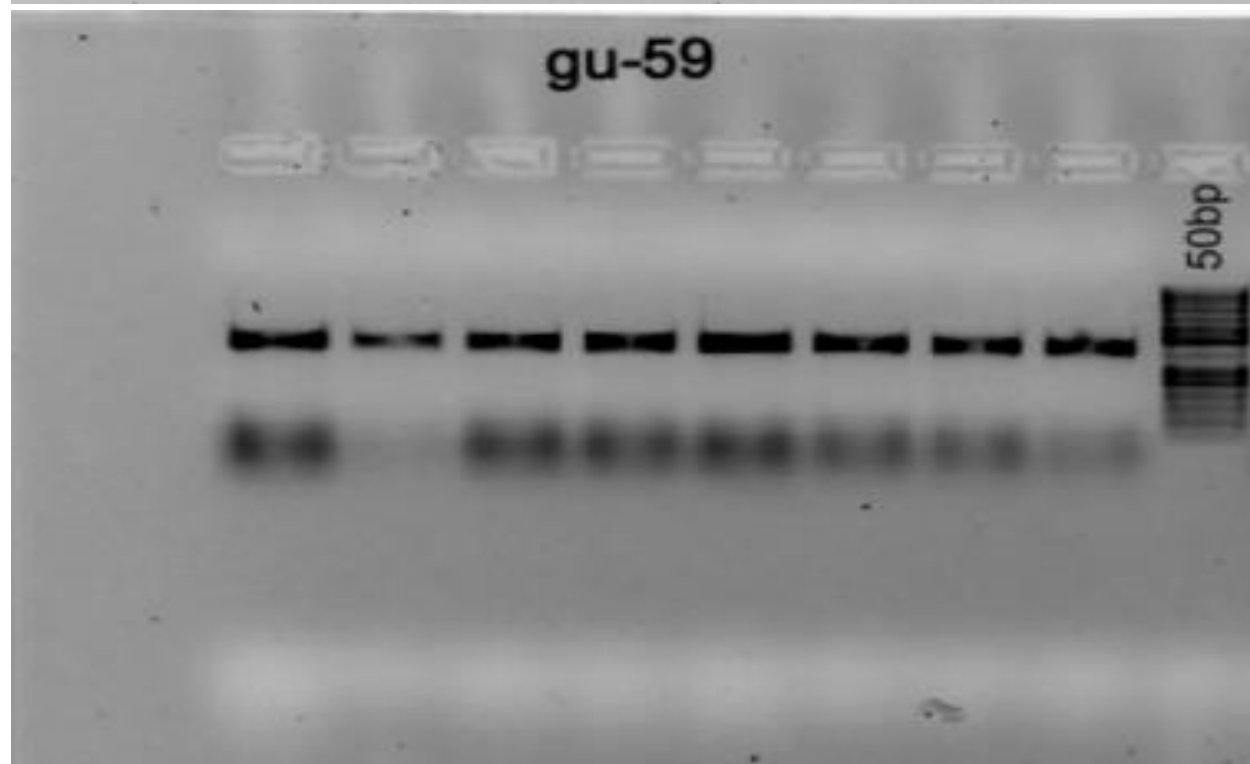
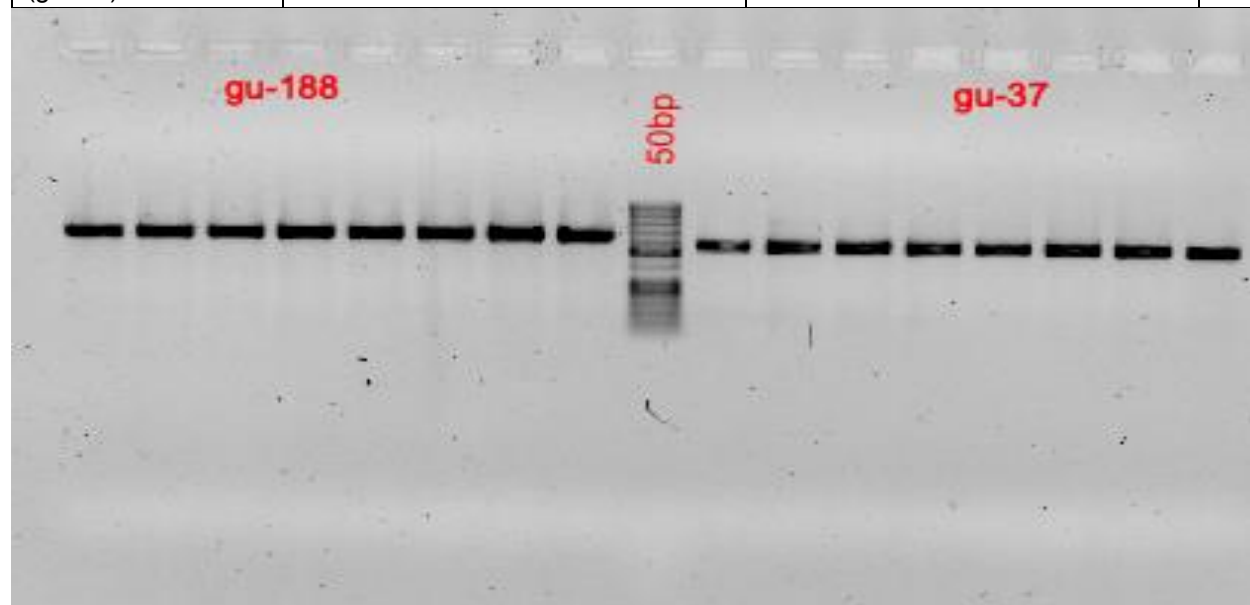
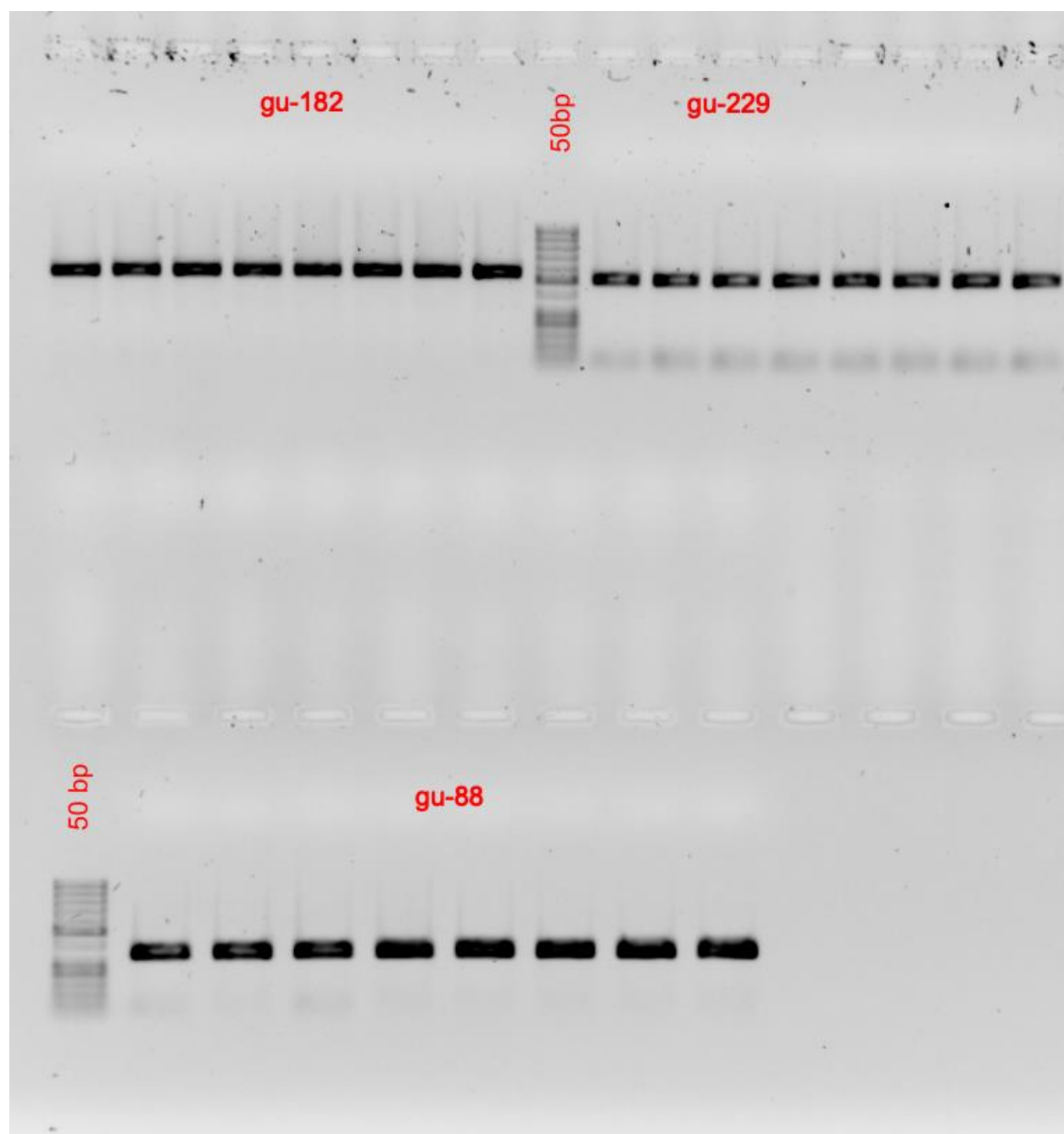


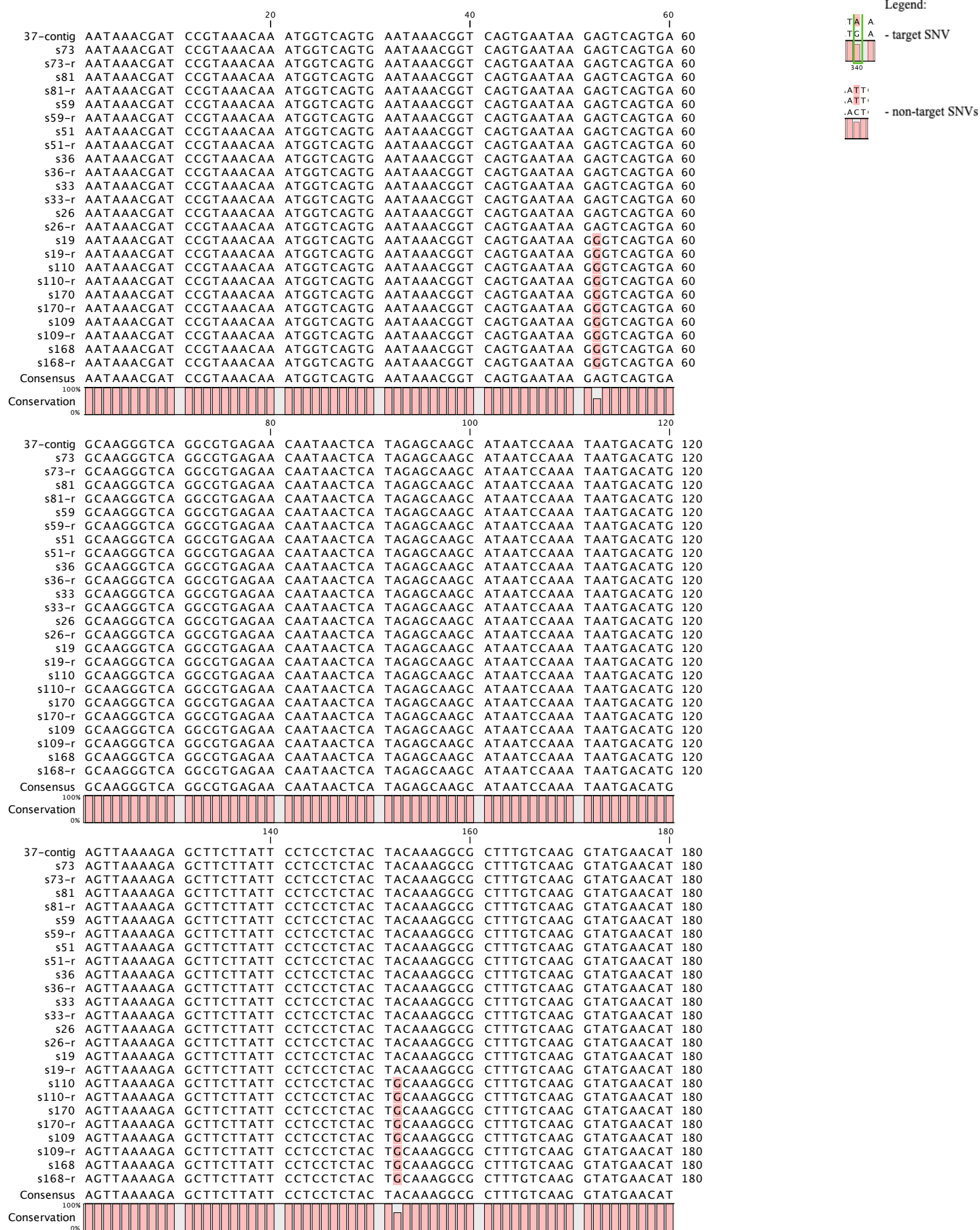
Supplementary Materials Figure S5. Validation of SNPs and deletions associated with early maturation by PCR and Sanger sequencing

Primer list

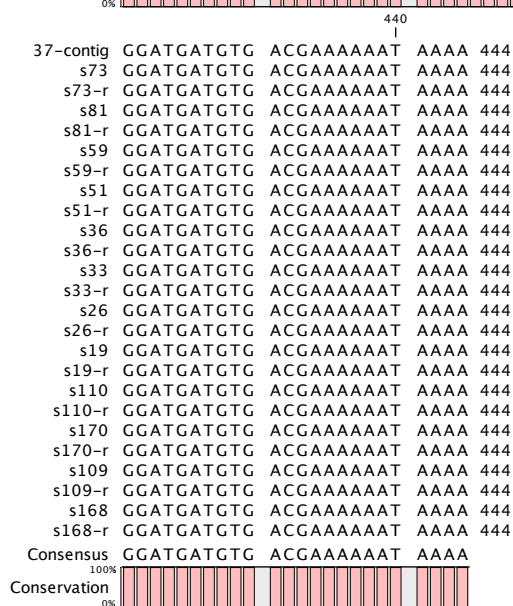
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contig_229:330388 (gu229)	CCAAGTGTCAATTCTCCGTTAGT	GGTCTGATCTGTCGTGCATATAG	500
contig_88:4931265 (gu88)	GCGTGGAGTTTGTAGTGGACTATG	GTGCTCTCTCTAAGGCTGAAAC	347
contig_182:841660 (gu182)	GACTCTCCATCATAGGGTTTACTT	AGTCCTGTCTCTCCTGGTAAT	550
contig_37:351180 (gu37)	CGAACAAACGGTCAGCAAATAA	AAATGAAGCACGTGAAGCTAATC	553
contig_188:3026232 (gu188)	GCCAGTCCAATAGGCTAACA	CCTGCCTGACTCAGATAAGAAC	638

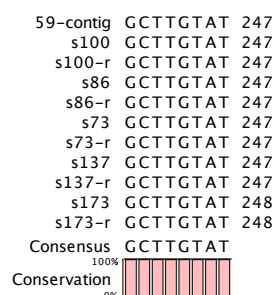
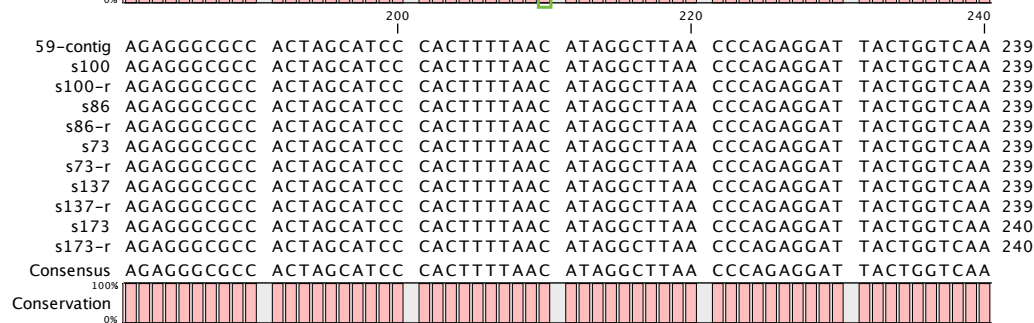
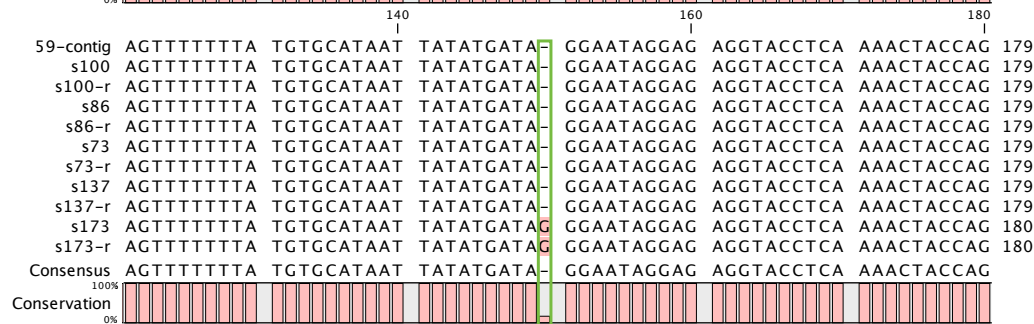
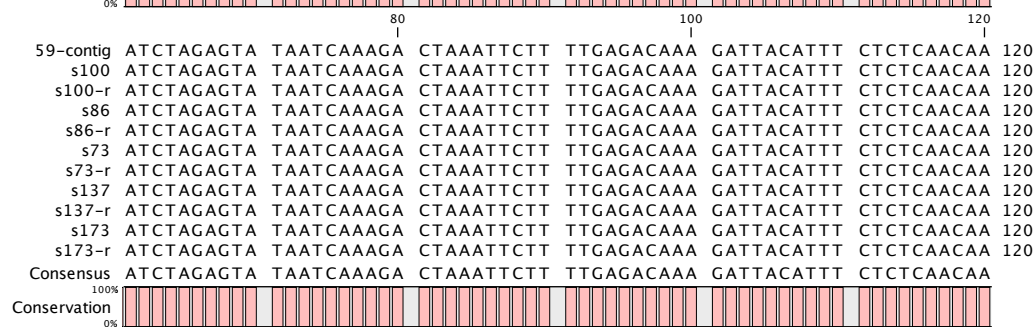
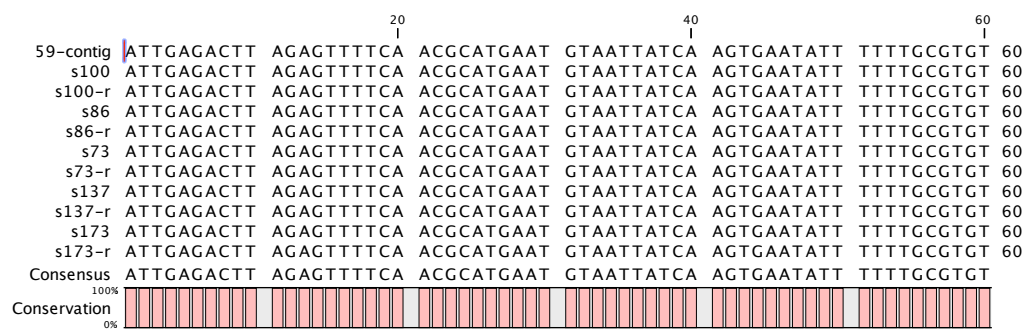


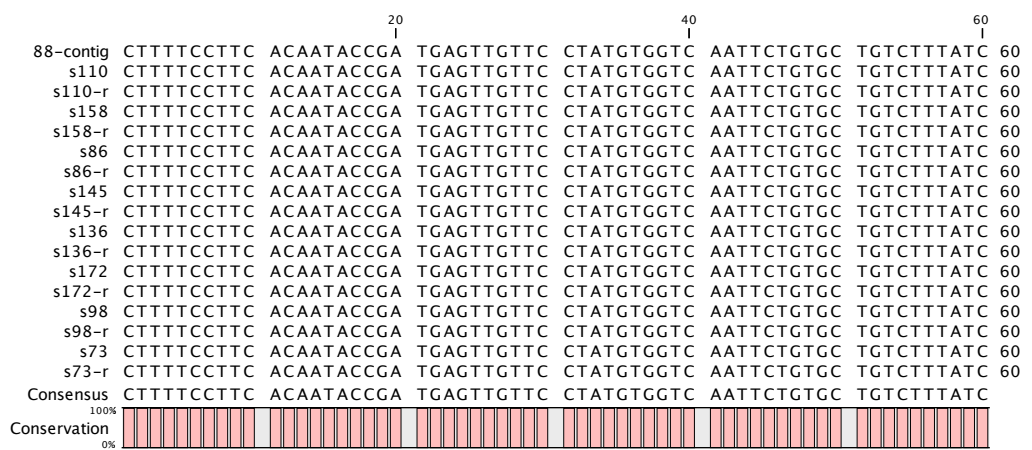












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s86-r AGAGG 245  
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s145-r AGAGG 245  
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s136-r AGAGG 245  
s172 AGAGG 245  
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s73-r AGAGG 245

Consensus AGAGG

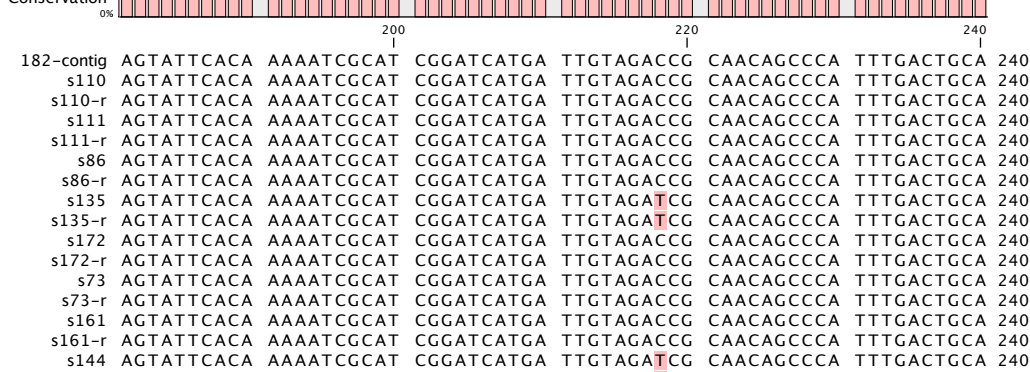
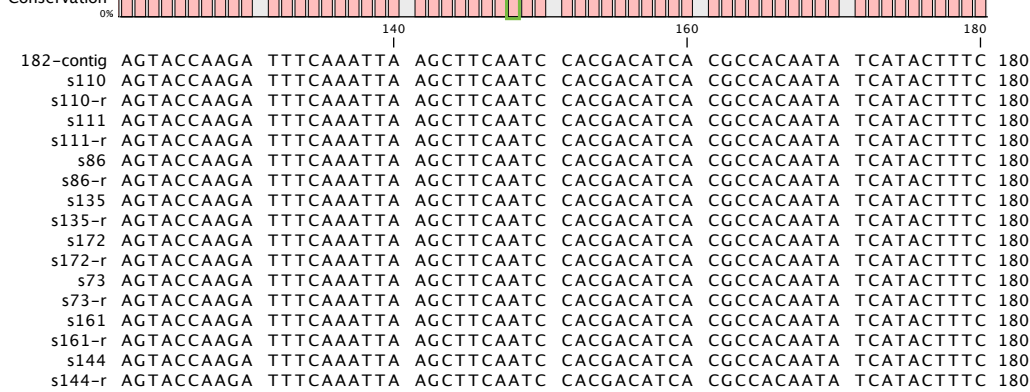
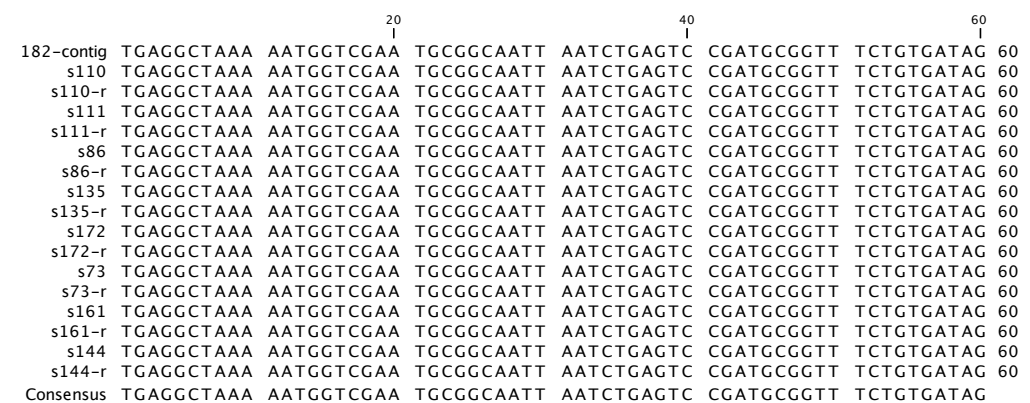
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Conservation

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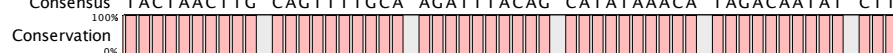




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Consensus	TTTAACAAAA	GGGACTATAA	CAGCAGTATG	ATCACAACTG	CATTTTGCAA	CCTAGATAAG	



		320		340		
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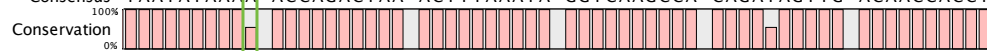
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