

Template Alignment: ConstructC_primer47

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 template sequence ConstructC_primer47

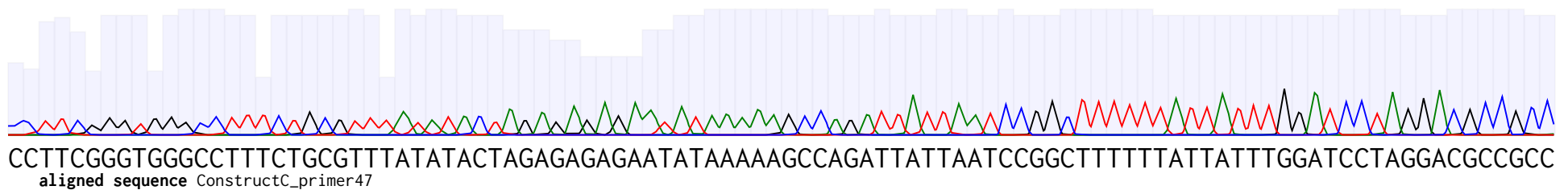
CCTTCGGGTGGGCCTTTCTGCGTTTATATACTAGAGAGAGAATATAAAAAGCCAGATTATTAATCCGGCTTTTTTATTATTTGGATCctaggacgccgcc

aligned sequence ch168 + C Revers complement

Bidirectional terminator

deriv...tion

pDG3661



AAGCCAGCTTAAACCCAGCTCAATGAGCTGGGTTTTTTGTTTGTTAAAAATGAAGAAGAACTGTGAAGCGTATTTAACTAGTAAATCTAAGACATATC
 template sequence ConstructC_primer47

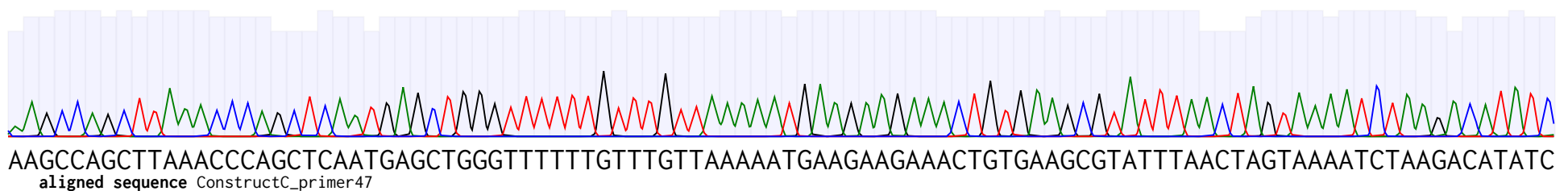
aagccagcttaaaccagctcaatgagctgggTTTTTTGTTTGTTAAAAATGAAGAAGAACTGTGAAGCGTATTTAACTAGTAAATCTAAGACATATC

aligned sequence ch168 + C Revers complement

derived from Bacillus subtilis rrn0; terminates lacZtranscription

derived f...yE front

pDG3661



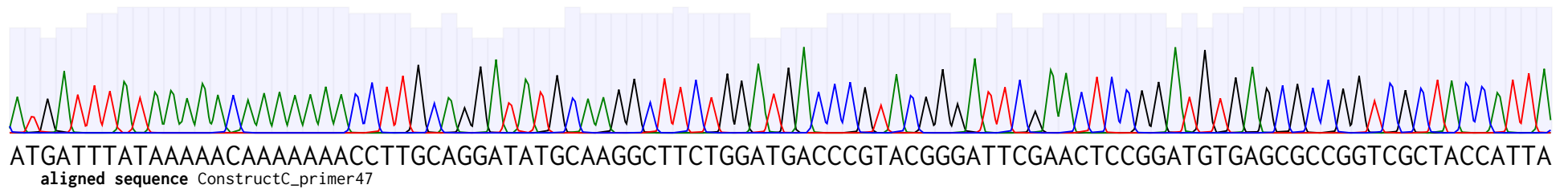
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template sequence ConstructC_primer47

atgattttataaaaaacaaaaaaccttgcaggatatgcaaggcttctggatgaccggtacgggattctgaactccggatgtgagcgccggtcgctaccatta

derived from Bacillus subtilis rrnB; terminates transcription from amyE front

pDG3661

aligned sequence ch168 + C Revers complement

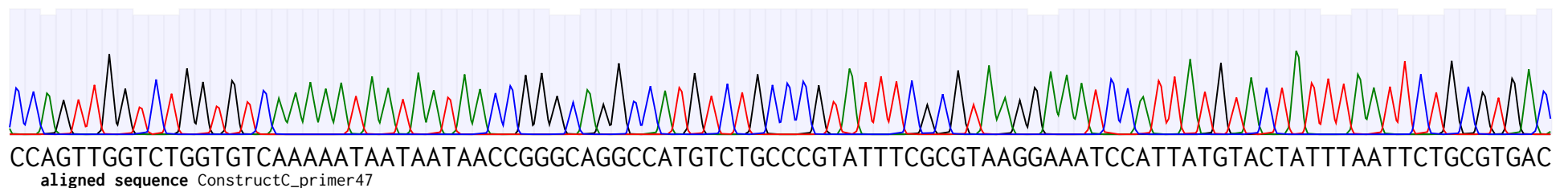


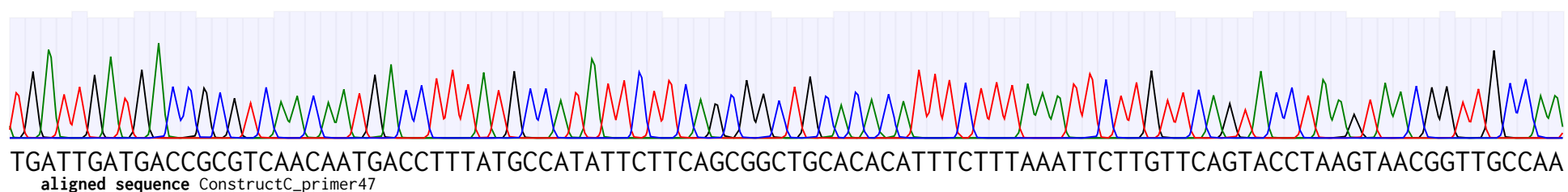
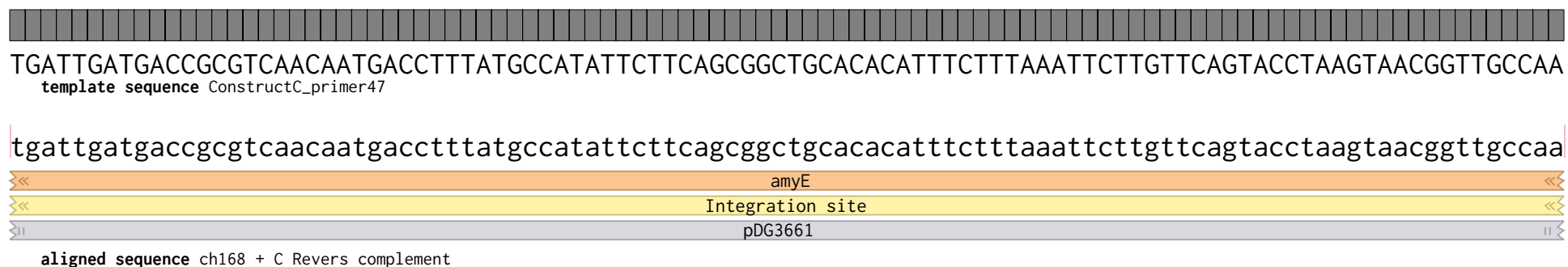
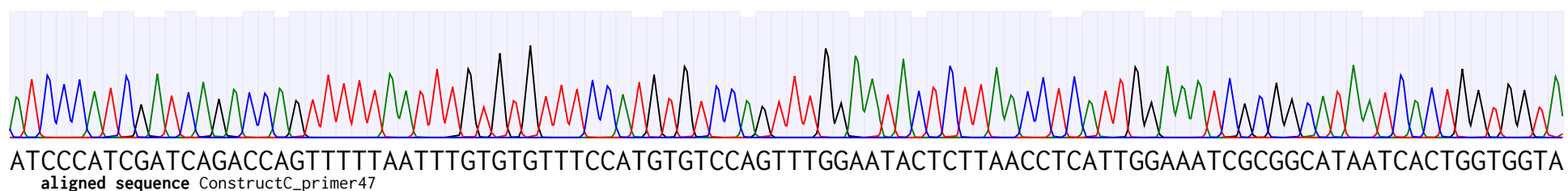
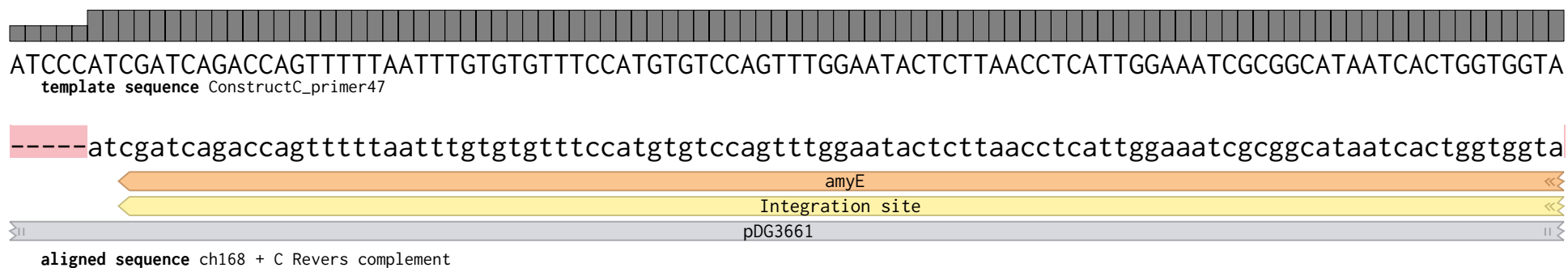
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template sequence ConstructC_primer47

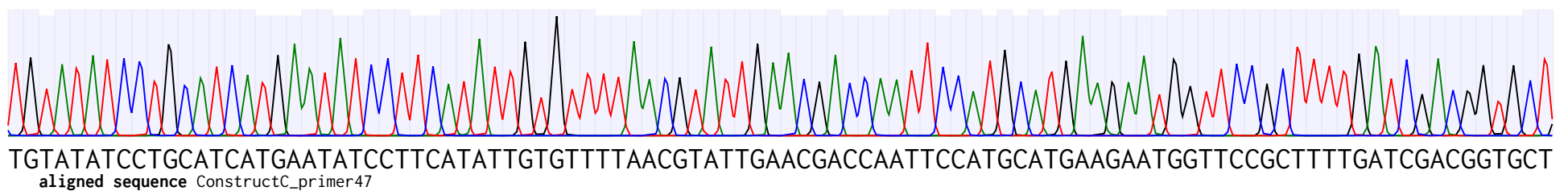
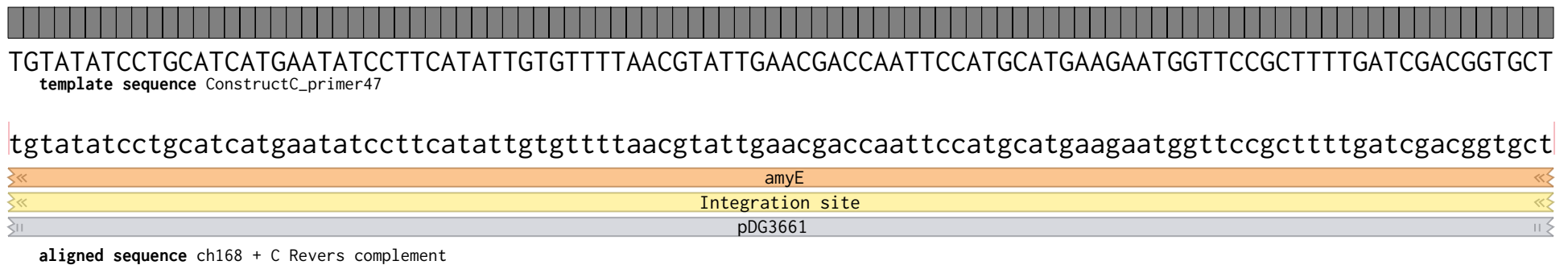
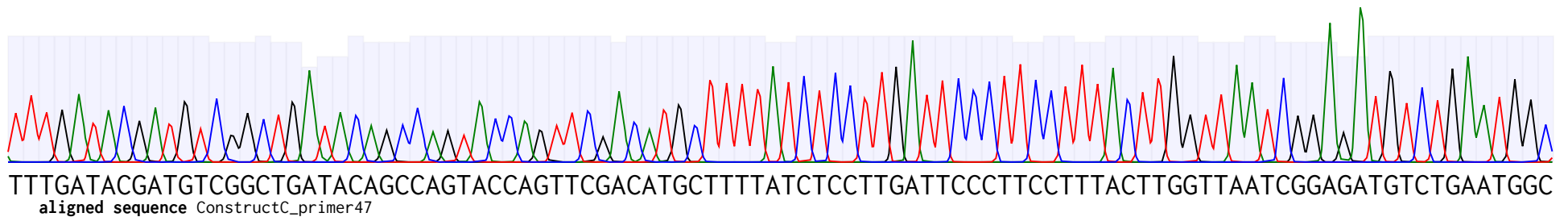
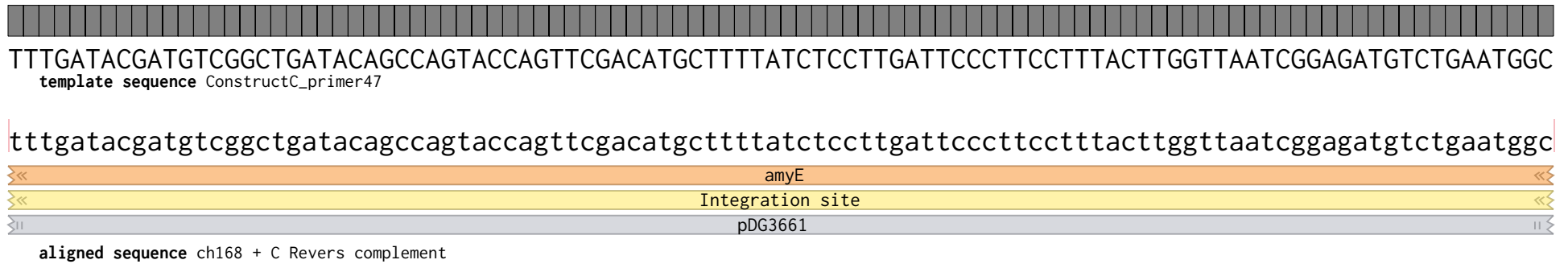
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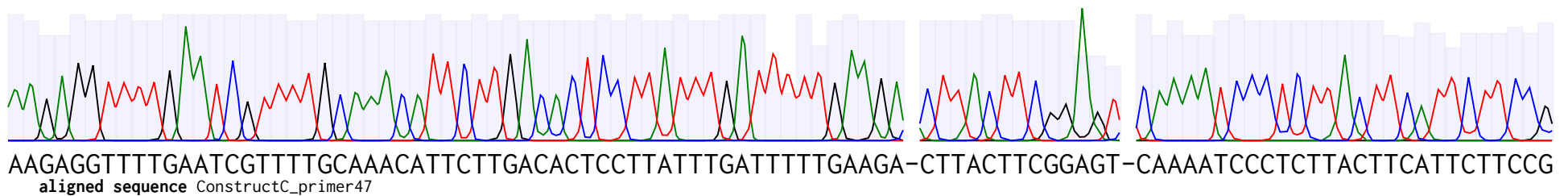
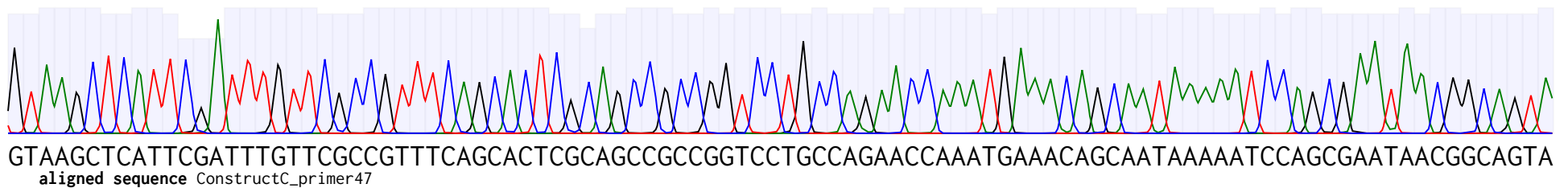
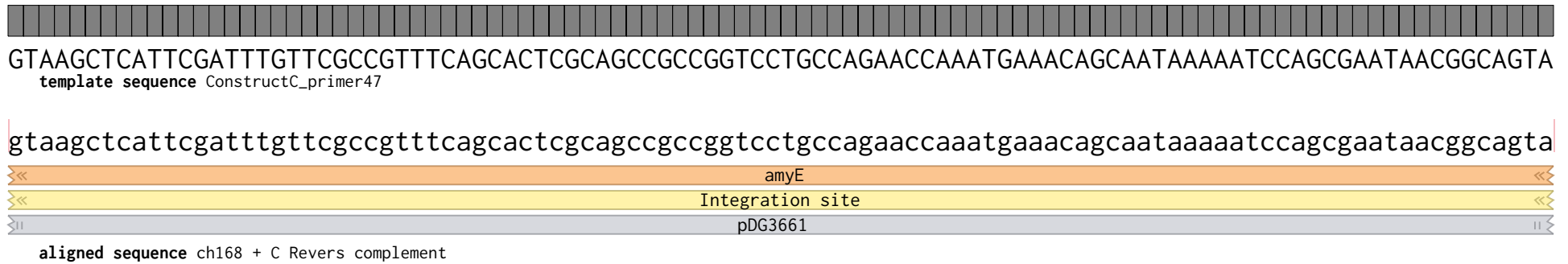
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aligned sequence ch168 + C Revers complement











CTTCCTCCTTTCAAACGATGTG-AAAACCTGGAGAATTTGTAASTMTA
template sequence ConstructC_primer47

cttcctcctttcaaaccgatgtgaagactggagaatTTTgtttaacgc
Chromosome 2
aligned sequence ch168 + C Revers complement

