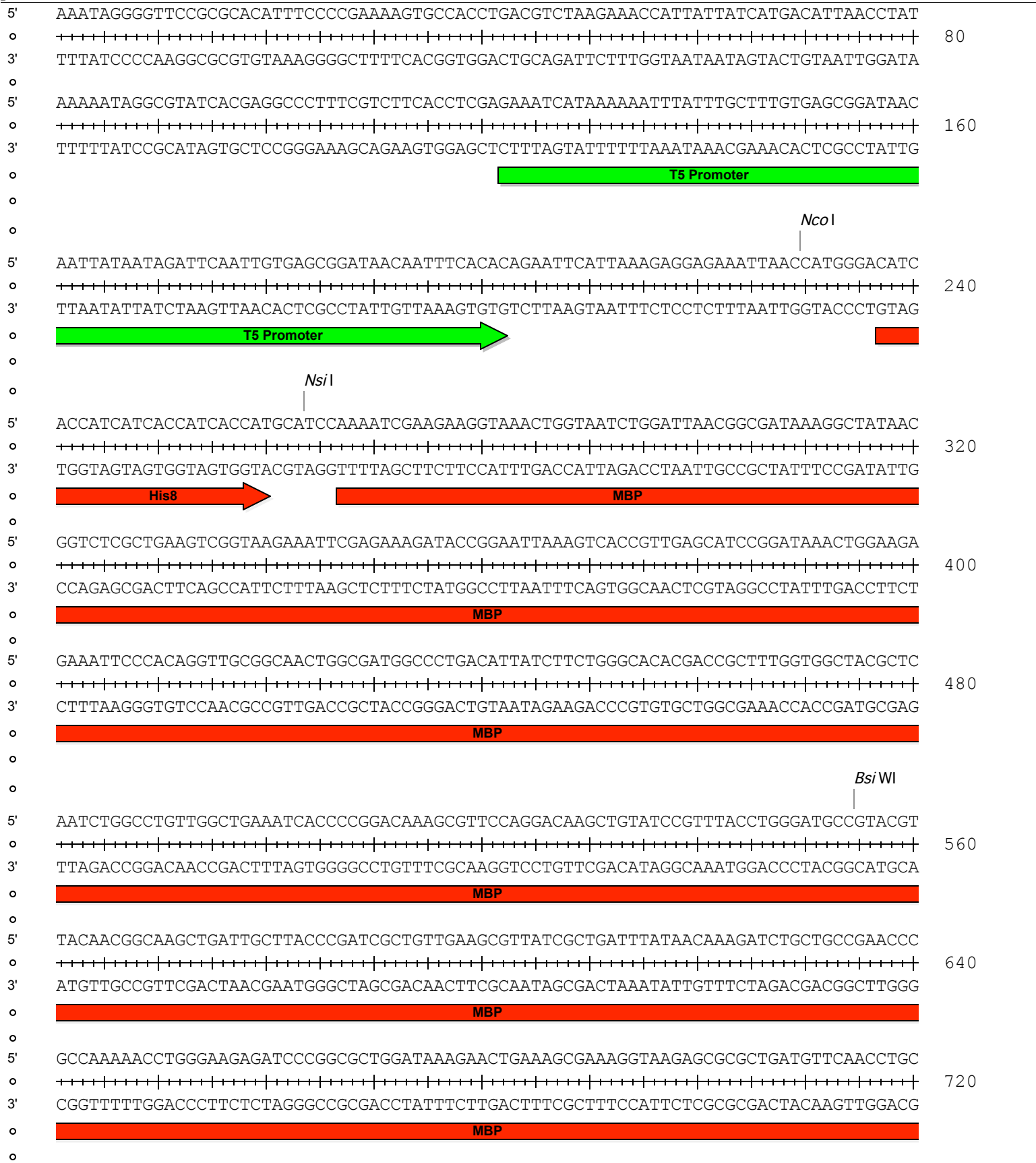


pVP33K



5' AAGAACCCTACTTCACCTGGCCGCTGATTGCTGCTGACGGGGTTATGCGTTCAAGTATGAAAACGGCAAGTACGACATG 800
 3' TTCTTGGCATGAAGTGGACCGGCGACTAACGACGACTGCCCCAATACGCAAGTTCATACTTTTGCCGTTTCATGCTGTAA
 MBP
 5' AAAGACGTGGGCGTGGATAACGCTGGCGCGAAAGCGGGTCTGACCTTCCTGGTTGACCTGATTAAAAACAAACACATGAA 880
 3' TTTCTGCACCCGCACCTATTGCGACCGCGCTTTTCGCCAGACTGGAAGGACCAACTGGACTAATTTTGTGTTGTGACTT
 MBP
 5' TGCAGACACCGATTACTCCATCGCAGAAGCTGCCTTTAATAAAGGCGAAACAGCGATGACCATCAACGGCCCGTGGGCAT 960
 3' ACGTCTGTGGCTAATGAGGTAGCGTCTTCGACGAAATTATTTCCGCTTTGTCGCTACTGGTAGTTGCCGGGCACCCGTA
 MBP
 5' GGTCCAACATCGACACCAGCAAAGTGAATTATGGTGTAACGGTACTGCCGACCTTCAAGGGTCAACCATCCAAACCGTTC 1040
 3' CCAGGTTGTAGCTGTGGTCGTTTCACTTAATACCACATTGCCATGACGGCTGGAAGTTCACAGTTGGTAGTTTGGCAAG
 MBP
 5' GTTGGCGTGCTGAGCGCAGGTATTAACGCCGCCAGTCCGAACAAAGAGCTGGCAAAAGAGTTCCTCGAAAACCTATCTGCT 1120
 3' CAACCGCACGACTCGCGTCCATAATTGCGGCGGTGAGGCTTGTTCGACCGTTTCTCAAGGAGCTTTTGATAGACGA
 MBP
 5' GACTGATGAAGGTCTGGAAGCGGTTAATAAAGACAAACCGCTGGGTGCCGTAGCGCTGAAGTCTTACGAGGAAGAGTTGG 1200
 3' CTGACTACTTCCAGACCTTCGCCAATTATTTCTGTTTGGCGACCCACGGCATCGCGACTTCAGAATGCTCCTTCTCAACC
 MBP
 5' CGAAAGATCCACGTATTGCCGCCACTATGGAAAACGCCAGAAAGGTGAAATCATGCCGAACATCCCGCAGATGTCCGCT 1280
 3' GCTTTCTAGGTGCATAACGGCGGTGATACCTTTTGGGGTCTTTCCACTTTAGTACGGCTTGTAGGGCGTCTACAGGCGA
 MBP
 MBP Forward
 5' TTCTGGTATGCCGTGCGTACTGCGGTGATCAACGCCGCCAGCGGTCGTCAGACTGTGCGATGAAGCCCTGAAAGACGCGCA 1360
 3' AAGACCATACGGCACGCATGACGCCACTAGTTGCGGCGGTGCGCCAGCAGTCTGACAGCTACTTCGGGACTTCTGCGCGT
 MBP
 MBP Forward
 5' GACTTTAATTAACTGGAAGTTCTGTTCCAGGGGCCCTGTTGCCAGGTTGCTGTGAAAACCTGTACTTccaggcgatcg 1440
 3' CTGAAATTAATTGGACCTTCAAGACAAGTCCCCGGGACAACGGGTCCAACGACACTTTTGGACATGAAGgtccgctagc
 3CP Cleavage Tetra-Cysteine Motif TEV Cleavage

Genomic map of the Barnase gene region on chromosome 1. The map shows the Barnase gene structure with exons and introns, and the BarCat cassette. The Barnase gene is transcribed from the left, with the 5' end at the top. The BarCat cassette is located downstream of the Barnase gene. The map includes sequence coordinates from 1520 to 2080. The Barnase gene is shown as a red bar with a black arrow indicating the direction of transcription. The BarCat cassette is shown as a blue bar. The sequence is shown in 5' to 3' orientation.

Sequence coordinates: 1520, 1600, 1680, 1760, 1840, 1920, 2000, 2080.

Gene structure: Barnase (red bar), BarCat Cassette (blue bar), CAT (red bar).











Transcription direction: 5' to 3' (indicated by the black arrow).

[illegible]

pVP33K

Hin dIII

[illegible]

5'	ATTACGTGCAGTCGATGATAAGCTGTCAAACATGAGAATTGTGCCTAATGAGTGAGCTAACTTACATTAATTGCGTTGCG	
o	+++++	
3'	TAATGCACGTCAGCTACTATTTCGACAGTTTGTACTCTTAACACGGATTACTCACTCGATTGAATGTAATTAACGCAACGC	3840
o		
5'	CTCACTGCCCCGCTTTCCAGTCGGGAAACCTGTCGTGCCAGCTGCATTAATGAATCGGCCAACGCGGGGAGAGGCGGTT	
o	+++++	
3'	GAGTGACGGGCGAAAGGTCAGCCCTTTGGACAGCACGGTCGACGTAATTACTTAGCCGGTTGCGCGCCCCCTCTCCGCCAA	3920
o		
o		
5'	TGCGTATTGGGCGCCAGGGTGGTTTTTCTTTTCACCACTGAGACGGGCAACAGCTGATTGCCCTTCACCGCCTGGCCCTG	
o	+++++	
3'	ACGCATAACCCGCGGTCCCAACAAAAGAAAAGTGGTCACTCTGCCCGTTGTCGACTAACGGGAAGTGGCGGACCGGGAC	4000
o		
o		
5'	AGAGAGTTGCAGCAAGCGGTCCACGCTGGTTTGCCCCAGCAGGCGAAAATCCTGTTTGATGGTGGTTAACGGCGGGATAT	
o	+++++	
3'	TCTCTCAACGTCGTTTCGCCAGGTGCGACCAAACGGGGTCGTCCGCTTTTAGGACAACTACCACCAATTGCCGCCCTATA	4080
o		
o		
5'	AACATGAGCTGTCTTCGGTATCGTCGTATCCCACTACCGAGATATCCGCACCAACGCGCAGCCCGGACTCGGTAATGGCG	
o	+++++	
3'	TTGTACTCGACAGAAGCCATAGCAGCATAGGGTGATGGCTCTATAGGCGTGGTTGCGCGTCGGGCCTGAGCCATTACCGC	4160
o		
o		
5'	CGCATTGCGCCCAGCGCCATCTGATCGTTGGCAACCAGCATCGCAGTGGGAACGATGCCCTCATTACAGCATTTCATGGT	
o	+++++	
3'	GCGTAACGCGGGTCGCGGTAGACTAGCAACCGTTGGTCGTAGCGTCACCCTTGCTACGGGAGTAAGTCGTAAACGTACCA	4240
o		
o		
5'	TTGTTGAAAACCGGACATGGCACTCCAGTCGCCTTCCCGTTCCGCTATCGGCTGAATTTGATTGCGAGTGAGATATTTAT	
o	+++++	
3'	AACAACTTTTGGCCTGTACCGTGAGGTGAGCGGAAGGGCAAGGCGATAGCCGACTTAACTAACGCTCACTCTATAAATA	4320
o		
o		
5'	GCCAGCCAGCCAGACGCAGACGCGCCGAGACAGAACTTAATGGGCCCCGCTAACAGCGCGATTGCTGGTGACCCAATGCG	
o	+++++	
3'	CGGTCGGTCGGTCTGCGTCTGCGCGGCTCTGTCTTGAATTACCCGGGCGATTGTCGCGCTAAACGACCACTGGGTACGC	4400
o		
o		
5'	ACCAGATGCTCCACGCCCAGTCGCGTACCGTCTTCATGGGAGAAAATAATACTGTTGATGGGTGTCTGGTCAGAGACATC	
o	+++++	
3'	TGGTCTACGAGGTGCGGGTCAGCGCATGGCAGAAGTACCCTCTTTTATTATGACAACTACCCACAGACCAGTCTCTGTAG	4480
o		
o		
5'	AAGAAATAACGCCGGAACATTAGTGCAGGCAGCTTCCACAGCAATGGCATCCTGGTCATCCAGCGGATAGTTAATGATCA	
o	+++++	
3'	TTCTTTATTGCGGCCTTGTAATCACGTCCGTGCAAGGTGTCGTTACCGTAGGACCAGTAGGTGCGCTATCAATTACTAGT	4560
o		
o		
5'	GCCCCACTGACGCGTTGCGCGAGAAGATTGTGCACCGCCGCTTTACAGGCTTCGACGCGGCTTCGTTCTACCATCGACACC	
o	+++++	
3'	CGGGTGACTGCGCAACGCGCTCTTCTAACACGTGGCGGCGAAATGTCCGAAGCTGCGGCGAAGCAAGATGGTAGCTGTGG	4640
o		
o		

The diagram illustrates the genomic organization of the *LacIq* gene and its regulatory elements. The *LacIq* gene is shown as a series of exons (represented by black boxes) and introns (represented by lines). The *LacIq* Promoter & UTR region is highlighted in green, and the pBR322 Origin is shown in brown. The map includes coordinates for various features and a scale bar.

Genomic Map Details:

- LacIq Gene Structure:** The gene is composed of several exons and introns. The exons are represented by black boxes, and the introns are represented by lines. The gene is located on the positive strand.
- LacIq Promoter & UTR:** This region is highlighted in green and includes the promoter and untranslated regions (UTR) of the *LacIq* gene. It is located upstream of the first exon.
- pBR322 Origin:** The pBR322 Origin is shown in brown and is located downstream of the *LacIq* gene. It is a common feature of plasmid vectors used in molecular biology.
- Coordinates:** The map includes coordinates for various features, such as the start and end of exons, introns, and the promoter region. These coordinates are provided in base pairs (bp).
- Scale Bar:** A scale bar is provided at the bottom of the map, indicating the distance in base pairs (bp) between various features.

pVP33K

5'	GGGATAACGCAGGAAAGAACATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAAGGCCGCGTTGCTGGCGTTT	
o	+++++	5520
3'	CCCTATTGCGTCCTTTCTTGTACACTCGTTTTCCGGTCGTTTTCCGGTCCTTGGCATTTTTCCGGCGCAACGACCGCAAA	
o	pBR322 Origin	
o		
5'	TTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATA	
o	+++++	5600
3'	AAGGTATCCGAGGCGGGGGGACTGCTCGTAGTGTTTTTAGCTGCGAGTTTCAGTCTCCACCGCTTTGGGCTGTCTTGATAT	
o	pBR322 Origin	
o		
5'	AAGATACCAGGCGTTTCCCCCTGGAAGCTCCCTCGTGCGCTCTCCTGTTCCGACCTGCCGCTTACCGGATACCTGTCCG	
o	+++++	5680
3'	TTCTATGGTCCGCAAAGGGGGACCTTCGAGGGAGCACGCGAGAGGACAAGGCTGGGACGGCGAATGGCCTATGGACAGGC	
o	pBR322 Origin	
o		
5'	CCTTTCTCCCTTCGGGAAGCGTGGCGCTTTTCTCATAGCTCACGCTGTAGGTATCTCAGTTCGGTGTAGGTCGTTTCGCTCC	
o	+++++	5760
3'	GGAAAGAGGGAAGCCCTTCGCACCGCGAAAGAGTATCGAGTGCGACATCCATAGAGTCAAGCCACATCCAGCAAGCGAGG	
o	pBR322 Origin	
o		
5'	AAGCTGGGCTGTGTGCACGAACCCCCCGTTCAGCCCGACCGCTGCGCCTTATCCGGTAACTATCGTCTTGAGTCCAACCC	
o	+++++	5840
3'	TTTCGACCCGACACACGTGCTTGGGGGGCAAGTCGGGCTGGCGACGCGGAATAGGCCATTGATAGCAGAACTCAGGTTGGG	
o	pBR322 Origin	
o		
5'	GGTAAGACACGACTTATCGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAG	
o	+++++	5920
3'	CCATTCTGTGCTGAATAGCGGTGACCGTCGTCGGTGACCATTGTCTTAATCGTCTCGCTCCATACATCCGCCACGATGTC	
o	pBR322 Origin	
o		
5'	AGTTCTTTGAAGTGGTGGCCTAACTACGGCTACACTAGAAGGACAGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTACC	
o	+++++	6000
3'	TCAAGAACTTCACCACCGGATTGATGCCGATGTGATCTTCCTGTCTATAAACCATAGACGCGAGACGACTTCGGTCAATGG	
o	pBR322 Origin	
o		
5'	TTTCGAAAAAGAGTTGGTAGCTCTTGATCCGGCAAACAAACCACCGCTGGTAGCGGTGGTTTTTTTGTGTTGCAAGCAGCA	
o	+++++	6080
3'	AAGCCTTTTTCTCAACCATCGAGAACTAGGCCGTTTGTGTTGGTGGCGACCATCGCCACCAAAAAAACAACGTTTCGTCTGT	
o	pBR322 Origin	
o		
5'	GATTACGCGCAGAAAAAAAGGATCTCAAGAAGATCCTTTGATCTTTTCTACGGGGTCTGACGCTCAGTGGAACGAAAACT	
o	+++++	6160
3'	CTAATGCGCGTCTTTTTTTCCTAGAGTTCTTCTAGGAACTAGAAAAGATGCCCCAGACTGCGAGTCACCTTGCTTTTGA	
o	pBR322 Origin	
o		
5'	CACGTTAAGGGATTTTGGTCATGAGATTATCAAAAAGGATCTTCACCTAGATCCTTTTAAATTAAAAATGAAGTTTTAAA	
o	+++++	6240
3'	GTGCAATTCCTTAAACCAGTACTCTAATAGTTTTTCTAGAAAGTGATCTAGGAAAATTTAATTTTTACTTCAAAATTT	
o	pBR322 Origin	
o		

